

Technical Support for Fish Tissue Monitoring for Implementation of EPA's 2016 Selenium Criterion

Draft

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List of Acronyms

CW	Cold water
CWA	Clean Water Act
EC₁₀	10% effect concentration
μM	Micron
MDL	Method detection limit
MPCA	Minnesota Pollution Control Agency
NAMC-SWG	North American Metals Council-Selenium Work Group
NCCA	National Coastal Condition Assessment
NOAA	National Oceanic and Atmospheric Administration
NPDES	National Pollutant Discharge Elimination System
QA/QC	Quality assurance/quality control
QL	Quantitation limit
SETAC	Society of Environmental Toxicology and Chemistry
SSD	Species sensitivity distribution
TMDL	Total maximum daily load
TSM	Technical support materials
TTF	Trophic transfer factor
USEPA	United States Environmental Protection Agency
USFWS	United States Fish and Wildlife Service
USGS	United States Geological Survey
WQC	Water quality criterion
WQS	Water quality standards
WW	Warm water
ww	Wet weight

Definitions

Anadromous fish

Fish with a life cycle that is divided between fresh and saltwater, including fish migrating to spawn in freshwater. Migrations should be cyclical, predictable, and cover more than 100 km (FishBase 2016).

Asynchronous spawners

Eggs are released in batches over a period of time that can last days or even months (Murua and Saborido-Rey 2003).

Fecundity

The physiological maximum potential reproductive output of an individual (usually female) over its lifetime (Bradshaw and McMahon 2008).

Gravid

Having the body distended with ripe eggs (FishBase 2016).

Indeterminate fecundity

Potential annual fecundity is not fixed before the onset of spawning and eggs can develop at any time during the spawning season (FishBase 2016).

Iteroparous

Producing offspring in successive batches, for example annual or seasonal batches, as is the case in most fishes (FishBase 2016).

Oocyte

Female sex cell which develops into an ovum. Oogonia become oocytes when meiosis begins, and specialized cells surround each oocyte to form a follicle. The oocyte undergoes maturation in preparation for spawning as an egg (modified from FishBase 2016).

Potamodromous

Fish species that spend their whole life in freshwater, but generally migrate for spawning purposes, typically back to a natal upstream tributary from a mainstream river or between connected lake and river systems. Migrations should be cyclical and predictable and cover more than 100 km (FishBase 2016).

Semelparous

Producing all offspring at one time, such as in most salmon. Usually these fish die after reproduction (FishBase 2016).

Site

In the context of site-specific criteria, a “site” may be a state, region, watershed, water body, or segment of a water body. A “site” for fish sampling is a specific water body segment.

Synchronous spawners

Eggs are released in a single episode during each breeding season (Murua and Saborido-Rey 2003).

Vitellogenesis

The process by which the yolk is formed and accumulated in the ovum. This is also the period when nutrients stored in the liver are transferred to the developing oocytes in the ovary or ovaries (FishBase 2016)

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1.0 Introduction

1.1 EPA's National CWA section 304(a) Recommended Chronic Aquatic Life Selenium Criterion in Freshwater

In 2016, the United States Environmental Protection Agency (EPA) updated its national Clean Water Act (CWA) section 304(a) recommended chronic aquatic life criterion for selenium in freshwater systems to reflect the latest scientific information. This information indicates that toxicity to aquatic life is driven by dietary exposures and that the reproductive life-stages of egg-laying vertebrates are the most sensitive to the toxic effects of selenium. The criterion has four criterion elements: (1) a fish egg-ovary criterion element; (2) a fish whole-body and/or muscle criterion element; (3) a water column criterion element (one value for lentic and one value for lotic aquatic systems); and (4) a water column intermittent criterion element (to account for potential chronic effects from short-term exposures to high concentrations in lentic and lotic aquatic systems) (see Table 1). Under EPA's 2016 CWA 304(a) recommended selenium criterion, the fish tissue criterion elements have primacy over water column elements, except where there are no fish, where fish tissue data do not meet state or tribal quality assurance procedures, or for water bodies with new discharges where selenium concentrations in fish tissue might not have stabilized (USEPA 2021a). EPA also recommends that the egg-ovary tissue criterion element has primacy over whole-body and muscle tissue criterion elements.

Toxicity data indicate that the selenium concentration in fish eggs and ovaries is the most robust and consistent measurement endpoint directly tied to adverse reproductive effects in aquatic organisms. Toxicity to developing embryos and larvae is directly linked to egg selenium concentration (USEPA 2021a). EPA derived the whole-body, muscle tissue, and water column elements from the egg-ovary element so that states and authorized tribes could more readily implement water quality criteria (WQC) based on EPA's national CWA section 304(a) recommended selenium criterion. The assessment of the available data on chronic selenium exposure for fish, invertebrates, and amphibians indicated that a criterion element derived from fish is expected to be protective of the aquatic community, since other taxa appear to be less sensitive to selenium than fish. EPA did not develop an acute criterion for selenium when it updated the chronic criterion because, although selenium may cause acute toxicity at high concentrations, the most deleterious effects on aquatic organisms are due to selenium's bioaccumulative properties. The chronic effects of bioaccumulated selenium occur at lower concentrations than acute effects.

In the case of bioaccumulative compounds like selenium, acute toxicity studies do not address risks that result from chronic exposure to chemicals via the diet (i.e., through the food web pathway). Such studies also do not account for the accumulation kinetics of many bioaccumulative compounds, such as selenium, and may underestimate effects from long-term accumulation in some types of aquatic systems. As described in EPA's *2021 Revision to: Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016* (hereafter referred to as *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016*), EPA also

included an intermittent exposure criterion element to provide protection from the most significant effects of selenium toxicity, reproductive toxicity, by protecting against selenium bioaccumulation in the aquatic ecosystem resulting from short-term, high concentration exposure events (USEPA 2021a). EPA recommends, as stated in the *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016*, that states and authorized tribes¹ adopt into their water quality standards (WQS) a selenium criterion that includes all four criterion elements. For more information see EPA’s *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016*, which can be found at <https://www.epa.gov/system/files/documents/2021-08/selenium-freshwater2016-2021-revision.pdf>.

Table 1: Summary of the Recommended Freshwater Selenium Ambient Chronic Water Quality Criterion for Protection of Aquatic Life.

Media Type	Fish Tissue ¹		Water Column ⁴	
	Egg-ovary ²	Fish Whole-body or Muscle ³	Monthly Average Exposure	Intermittent Exposure ⁵
Magnitude	15.1 mg/kg dry weight	8.5 mg/kg dry weight whole-body or 11.3 mg/kg dry weight muscle (skinless, boneless fillet)	1.5 µg/L in lentic aquatic systems 3.1 µg/L in lotic aquatic systems	$WQC_{int} = \frac{WQC_{30-day} - C_{bkgrnd}(1 - f_{int})}{f_{int}}$
Duration	Instantaneous measurement ⁶	Instantaneous measurement ⁶	30 days	Number of days/month with an elevated concentration
Frequency	Not to be exceeded	Not to be exceeded	Not more than once in three years on average	Not more than once in three years on average

1. Fish tissue elements are expressed as steady-state.
2. Egg-ovary supersedes any whole-body, muscle, or water column element when fish egg-ovary concentrations are measured, except as noted in footnote 4 below.
3. Fish whole-body or muscle tissue supersedes water column element when both fish tissue and water concentrations are measured, except as noted in footnote 4 below.
4. Water column values are based on dissolved total selenium in water and are derived from fish tissue values via bioaccumulation modeling. When selenium inputs are increasing, water column values are the applicable criterion element in the absence of steady-state condition fish tissue data.
5. Where WQC_{30-day} is the water column monthly element for either lentic or lotic waters; C_{bkgrnd} is the average background selenium concentration; and f_{int} is the fraction of any 30-day period during which elevated selenium concentrations occur, with f_{int} assigned a value ≥ 0.033 (corresponding to 1 day).
6. Fish tissue data provide instantaneous point measurements that reflect integrative accumulation of selenium over time and space in fish population(s) at a given site.

1 Throughout this document and in the [CWA](#), the term “states” means the fifty states, the District of Columbia, the Commonwealth of Puerto Rico, the United States Virgin Islands, Guam, American Samoa, and the Commonwealth of the Northern Mariana Islands. The term “authorized tribe” means those federally recognized Indian tribes with authority to administer a CWA WQS program.

1.2 Selenium Technical Support Materials

EPA has prepared a four-volume set of documents to provide recommendations to states, authorized tribes, and other agencies for implementing WQC based on the national CWA section 304(a) recommended selenium criterion (USEPA 2021a). These four documents constitute the Technical Support Materials for EPA's *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016* (USEPA 2021a). Each document of the set focuses on a specific aspect of implementation of the national CWA section 304(a) recommended selenium criterion. Together, these four EPA documents provide information that will assist states and authorized tribes with adopting WQC based on EPA CWA section 304(a) recommended selenium criterion and implementing it in various CWA programs.

- 1) *Technical Support for Adopting and Implementing EPA's Selenium 2016 Criterion in Water Quality Standards, Draft*: provides recommendations for the adoption and implementation of the national CWA section 304(a) recommended selenium criterion, including the various flexibilities available to states and tribes using WQS tools.
- 2) *Technical Support for Fish Tissue Monitoring for Implementation of EPA's 2016 Selenium Criterion, Draft*: provides an overview on how to establish or enhance existing fish tissue monitoring programs to facilitate implementation of the fish tissue-based criterion elements in the national CWA section 304(a) recommended selenium criterion.
- 3) *Frequently Asked Questions: Implementing Water Quality Standards Based on EPA's 2016 Recommended Selenium Criterion in Clean Water Act Section 402 NPDES Permits, Draft*: is intended to help NPDES permit writers understand what permitting guidance (i.e., state or tribal implementation procedures) may be appropriate to implement state and authorized tribal WQS based on EPA's CWA section 304(a) recommended selenium criterion. This set of FAQs also provides recommendations on how to establish water quality-based effluent limits (WQBELs) in NPDES permits.
- 4) *Frequently Asked Questions (FAQs): Implementing the 2016 Selenium Criterion in Clean Water Act Sections 303(d) and 305(b) Assessment, Listing, and Total Maximum Daily Load (TMDL) Programs, Draft*: provides information on how to complete assessments, list impaired waters, and develop TMDLs based on EPA approved WQS that adhere to EPA's national CWA section 304(a) recommended selenium criterion, including all four elements.

1.3 Document Overview

This document, *Technical Support for Fish Tissue Monitoring for Implementation of EPA's 2016 Selenium Criterion, Draft*, examines technical considerations for developing a robust sampling program to characterize selenium concentrations in fish tissue for a variety of CWA implementation programs (e.g. CWA section 303(d) listing and TMDLs). Some aspects of this document can also be used to support the development of site-specific water column criterion elements. Site-specific water column criterion elements can be developed by performing a site-specific water column translation from the fish tissue elements of the criterion. See Appendix K

of *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016* (USEPA 2021a) for site-specific approaches for translating between fish tissue and water column selenium concentrations.

This document is intended to assist states and authorized tribes in planning enhancements to their monitoring programs, not to limit the fish tissue data that can be used for waterbody assessments with the recommended criterion. The recommended criterion applies to all species present in a waterbody. EPA’s regulations require that states and authorized tribes evaluate all existing and readily available water quality-related data for assessment decisions (40 C.F.R. 130.7(b)(5)), which for this criterion would include selenium data and information for any fish species. For additional information about assessment decisions for this criterion, please refer to the *Frequently Asked Questions (FAQs): Implementing the 2016 Selenium Criterion in Clean Water Act Sections 303(d) and 305(b) Assessment, Listing, and Total Maximum Daily Load (TMDL) Programs, Draft* (USEPA 2021b).

This document does not specifically address the site-specific modification of the fish tissue criterion elements, which should be developed using fish assemblage data and the recalculation procedure (USEPA 2013). States and authorized tribes that wish to develop site-specific fish tissue criterion elements should engage their EPA regional office early in the process to ensure the development and use of sound scientific analyses.

2.0 Monitoring Strategy

When considering enhancements to monitoring strategies for CWA implementation, EPA recommends that agencies first review their existing fish tissue monitoring programs and determine how best to incorporate collection of fish tissue samples for the implementation of the aquatic life selenium criterion into the monitoring program. For example, existing fish tissue monitoring programs will likely be collecting data to assess the risk of fish consumption to human health. These monitoring programs may be designed with different objectives than monitoring activities designed to assess attainment of an aquatic life criterion. These differences need to be considered and reconciled when using an existing fish tissue monitoring program for assessing the fish tissue criterion elements of the selenium aquatic life criterion.

The following sections review study design and sampling considerations regarding fish tissue types, sample types, target species, and spatial and temporal variability. These topics should be considered when collecting data to assess fish tissue concentrations against the fish tissue criterion elements of the state’s or authorized tribe’s selenium aquatic life criterion or collecting data to translate a fish tissue criterion element into a site-specific water column criterion element.

The relationship between fish sampling locations and timing, species’ habits and natural history, and the selenium source(s) should be understood and accounted for during the design of sampling plans for criterion implementation. The fish tissue sampling methods should be designed to characterize the variability of selenium in the target population, including areas of high selenium bioaccumulation. The criterion establishes population level protection that is measured by the population mean concentration. Therefore, sampling should be designed to determine the mean selenium concentration in the fish tissue of the population. EPA encourages

early discussions with EPA regions to assure that studies capture the appropriate data. Detailed field collection procedures and sampling design considerations can be found in *Guidance for Assessing Chemical Contaminant Data for Use in Fish Advisories, Vol 1: Fish Sampling and Analysis* (EPA's Fish Advisory Guidance Volume 1) (USEPA 2000), the *Field Sampling Plan for the National Study of Chemical Residues in Lake Fish Tissue* (USEPA 2002a), and EPA's National Aquatic Resource Survey Field Operations Manuals for rivers and streams and Great Lakes and coastal waters (USEPA 2019a, 2019b, 2015). Appendix A of this document presents egg and ovary collection and sample preparation methods.

2.1 Tissue Type

EPA recommends sampling egg-ovary tissue for both assessment and site-specific water column criterion element development purposes, when possible. From the toxicological standpoint, the most representative measure of exposure to a toxic substance is its concentration at the site of toxic action. In fish (the aquatic species most sensitive to selenium), the most ecologically relevant sites of toxic action are the mature reproductive tissues of adults (ovaries) or early life stages (vitellogenic eggs/larvae). This was a major point of consensus of the 2009 Society of Environmental Toxicology and Chemistry (SETAC) Pellston workshop on selenium risk assessment (Chapman et al. 2009). Therefore, the national CWA section 304(a) recommended selenium criterion is based on reproductive effects in fish, as represented by selenium concentrations in egg-ovary tissue.

While egg-ovary tissue of adult female fish is the most direct reflection of reproductive toxicity in fish, samples of muscle or whole-body tissue from adult fish serve as robust alternatives and may be collected for implementation purposes.

Collection of fish samples for egg-ovary analysis poses special challenges as only gravid female fish can be sampled. Due to this constraint, the decision of what tissue type to collect should be made based on the following considerations:

- *Temporal*: Most fish species that are synchronous spawners, spawn in the spring; whereas fish tissue collection for fish consumption advisories typically occur in the late summer or early fall, when contaminant loads in the edible portion of the fish are highest. Spring sampling may also be challenging in states or tribal lands where rivers and streams have high flows due to storm run-off and spring snow melt. Timing the collection of mature eggs from asynchronous spawners can also be challenging, as these species can have eggs in various stages of development at once.
- *Spatial*: Some fish species migrate to upstream areas to spawn and these areas may be harder to access than higher order downstream segments that are inhabited during non-spawning seasons.
- *Size*: It is difficult to collect and analyze egg-ovary tissue samples from small fish species (e.g., certain species in the family Cyprinidae or Cyprinodontidae) due to the logistics of the collection and the small amount of tissue available for analysis (number of eggs or biomass).

Due to these concerns, states and authorized tribes have considerable discretion when selecting the fish tissue type to collect in their sampling protocols. The flexibility provided by being able to collect data from multiple fish tissue types allows for leveraging existing monitoring capacity. A number of the species that are good target species for selenium sampling could also be commonly collected as muscle (fillet) samples in state and tribal fish tissue monitoring programs for other contaminants (e.g., trout, salmon, bass, sunfish) (see section 2.3 for more information on target species). The whole-body tissue criterion element also simplifies the collection and processing of small fish species that may be the dominant trophic level in lower order stream networks.

When developing a new or modifying an existing fish tissue monitoring strategy, states or authorized tribes may want to consider funding and staff resources, opportunities to work with other fish tissue monitoring programs, existing information on the spawning habits and size of target species, and potential population level effects associated with using lethal sampling techniques when deciding what fish tissue type to sample. To keep the public, stakeholders, and EPA well informed, it is good practice for monitoring programs to describe in their sampling protocols why they are sampling a particular tissue type. Similar considerations might also be evaluated when selecting a tissue to sample for the development of a site-specific water column criterion element. If possible, EPA recommends that egg-ovary tissue be sampled to conduct a site-specific water column translation. Sampling considerations associated with different types of fish tissue are presented in Table 2.

Table 2. Sampling Considerations Associated with Different Types of Fish Tissue

Issue	Egg-ovary	Whole-body	Muscle/Fillet	Comments
Criterion Hierarchy Considerations	Has primacy in the criterion; supersedes other fish tissue criterion elements and the water column criterion elements	Supersedes water column criterion elements; equal consideration to muscle tissue	Supersedes water column criterion elements; equal consideration to whole-body tissue	While selenium concentrations in all three tissue types are significantly correlated to reproductive toxicity effects seen in fish, egg-ovary concentrations have the strongest relationship.
Ease of collection	Difficult	Easy	Easy – except on small fish	Egg-ovary samples are only collected from gravid females; there are seasonal and logistical considerations, and species-specific sampling windows. See Appendices A and B.
Consistency with existing state & tribal methods	Not typically collected	Sometimes collected	Primary tissue collected	Whole-body samples might be collected in special cases, such as for certain human populations that consume whole fish or for ecological risk assessments.
Sample availability	Limited – only from gravid females	Always	Always	For water bodies with small sized species at top trophic levels, whole-body may be the only option due to issues collecting a sufficient mass of muscle tissue.
Ability to make composite sample	Yes	Yes	Yes	Compositing can be used to reduce the overall cost of an analytical program, primarily by reducing the number of samples that must be analyzed to represent an average concentration. Compositing can also ensure that enough mass is available for chemical analyses. However, by compositing samples, information on the range of selenium concentrations in individual organisms is lost.
Ability to test individual sample	Yes, on larger species; may be difficult on small species	Yes, on larger species; may be difficult on small species	Yes, on larger species; may be difficult on small species	Individual samples are valuable when sampling from waters known or suspected to be impacted by selenium discharges (see section 3.2.2), however the need to analyze multiple individual samples versus a few composite samples can make them more resource intensive to prepare and expensive to analyze.

2.1.1 Egg-ovary Tissue Sample

Selenium concentrations in all three tissue types are significantly correlated to reproductive toxicity effects seen in fish; however, egg-ovary concentrations have the strongest relationship. EPA recommends sampling fish egg-ovary tissue for assessment of the selenium aquatic life criterion or for development of a site-specific water column criterion element, when possible. Egg-ovary tissue refers to mature eggs, pre-spawn ovary tissue that contains mature eggs, or both. As an oocyte grows into a mature egg, it passes through several stages of development (i.e., oogenesis, primary oocyte growth, cortical alveolus stage, vitellogenesis, maturation, and ovulation) (Tyler and Sumpter 1996). During this egg development process, the oocyte size increases dramatically as the yolk is developed. For example, the diameter of an undeveloped oocyte of the rainbow trout is around 20 μm and the fully developed egg is about 4 mm (Nagahama 1983). Selenium is transferred from an adult female fish to her eggs during vitellogenesis. Eggs should not be collected until after this transfer has occurred. Appendix A of this document presents egg and ovary collection and sample preparation methods.

Adult female fish must be collected during the late vitellogenic or pre-ovulatory periods of oogenesis for the tissue concentrations to be scientifically and toxicologically meaningful. The egg-ovary sample should represent the potential selenium load available to eggs and larvae through maternal transfer. Egg-ovary tissue from pre-spawn, reproductively mature (also called “gravid” or “vitellogenic”) females is the preferable tissue to collect because it will give the most accurate representation of potential selenium hazard to reproduction. Egg-ovary tissue data provide point measurements that reflect integrative dietary accumulation, transfer, and deposition of selenium over time and space in female fish at a given site. Research has shown that selenium concentrations in egg-ovary tissue is strongly correlated with selenium in the maternal diet; the selenium is transferred from the adult female to her eggs during vitellogenesis (Janz et al. 2010).

When using egg-ovary tissue for the implementation of the selenium criterion, states and authorized tribes should carefully consider the difficulty in timing egg-ovary sampling with spawning periods. Timing errors related to fish reproduction may result in data that falsely indicate the selenium criterion is being met. Ovary tissue sampled from a female that is not gravid will not be representative of the selenium concentrations of this tissue for a gravid individual and is not appropriate for comparison to the egg-ovary criterion. A female is typically gravid for a very small window of time for most synchronous species, and may occur in the spring or early summer, or in the fall to early winter (see Appendix B). The timing of the spawning season will depend on the species, geography, and a number of environmental cues (e.g., temperature, flow, photoperiod). In northern latitudes or higher elevations, spawning may occur slightly later than in southern latitudes or lower elevations. EPA recommends that states and authorized tribes consult with local fish biologists, who may be in other state or tribal agencies (e.g., Departments of Natural Resources, Departments of Fish and Game), when designing sampling plans.

Reproductively mature females of most fish species, except indeterminate spawning species and viviparous species (i.e., live bearing), will produce eggs that can be sampled for selenium. Ovary tissue of synchronous spawners (e.g., species in the genus *Oncorhynchus*) typically contain

oocytes that are all in the same stage of development. Fish species that spawn multiple times per season (asynchronous, e.g., some species in the family Cyprinidae) have variable cycles of oogenesis and commonly have ovary tissue that contains oocytes and eggs at all stages of development (Nagahama 1983). In these asynchronous spawners, egg maturation may occur well before, immediately prior to, or during the spawning season. For example, *Lepomis cyanellus* (green sunfish) can spawn multiple times per season (Osmundson and Skorupa 2011, Chapman et al. 2010). Thus, special care should be taken when sampling asynchronous species for egg-ovary tissue, as the pre-spawn window can be hard to predict.

Given these concerns, EPA has the following recommendations when sampling female asynchronous spawners: 1) if the fish is too small to easily sample the egg-ovary tissue, the whole body should be sampled (including the eggs) and the selenium concentration should be compared to the whole-body criterion element; 2) if fish are sampled during the reproductive season and they are large enough to easily sample the egg-ovary tissue, this tissue should be sampled (the 75% rule does not apply to egg-ovary composite samples, see section 2.2.1); and 3) muscle tissue should not be sampled during the reproductive season as selenium may be depleted from this tissue during this time.

The egg-ovary tissue criterion element has primacy over all other criterion elements of the selenium water quality criterion. Most states and authorized tribes do not currently collect egg-ovary tissue as part of their regular monitoring programs. EPA recognizes that many states and authorized tribes may not have the resources to augment their existing monitoring programs to include egg-ovary tissue collection. While egg-ovary remains the preferable tissue type, whole-body or muscle tissue samples can be used as an alternative.

2.1.2 Whole-body and Muscle Tissue Samples

Whole-body and muscle tissue samples may be collected as an alternative to egg-ovary tissue. Selenium concentrations in these tissues will provide representative information on selenium bioaccumulation and ecological exposure at almost any time of the year (except pre- and post-spawn windows for females). However, there will likely be some variation across seasons due to dietary composition, temperature, depuration of selenium from tissue during vitellogenesis prior to spawning, and other factors. If the sex of the fish can be determined, it is preferable to use male fish for muscle or whole-body samples since the selenium levels in their tissues are stable, regardless of reproductive state. The only time of year that should be avoided for collecting whole-body or muscle tissue samples from female fish is directly pre- and post-spawn because they could have reduced selenium concentrations in their tissues due to the recent transfer of selenium to eggs (USEPA 2021a). The exception is small-bodied fish, particularly asynchronous spawners, where the collection of egg-ovary tissue is impractical. In this instance, the whole body should be collected (with the eggs if the fish is female) and compared to the whole-body criterion element.

Summer and fall may be prime periods for whole-body and muscle tissue collection due to the engorgement of populations to replenish fat and energy reserves post-spawn and for over-wintering. Winter tissue collection is discouraged, except in subtropical regions where metabolic

changes due to lower water temperatures do not occur. Whole-body and muscle tissue data provide point measurements that reflect integrative dietary accumulation and deposition of selenium in fish tissues over time and space in fish population(s) at a given site.

EPA is aware that some states and authorized tribes use muscle plugs in their monitoring programs as an alternative to collecting muscle fillets. States or authorized tribes that utilize plugs should be aware of certain considerations. Contaminant concentrations can vary considerably depending on where the plug is taken from a fish. Plugs should be collected from a descaled meaty portion of the dorsal muscle tissue, between the dorsal fin and lateral line (USEPA 2019a). Waddell and May (1995) found that selenium concentrations in plugs from this location were significantly correlated to adjacent muscle tissue. Studies with mercury have also shown that this location results in a sample that has homogeneous concentrations and concentrations that were similar to mean concentrations for a muscle fillet (Cizdziel et al. 2002). Plugs provide very small tissue masses (about a gram of tissue per fish) and may not provide enough biomass for reanalysis or quality assurance/quality control (QA/QC) analysis. This may lead to difficulty confirming the quality of the sample analysis. In addition, relatively small individuals may not recover from a muscle plug biopsy punch. Care should be taken to ensure that there is enough tissue for the analytical method. In addition, states and authorized tribes may want to establish species-specific conversion factors or regressions at the start of their sampling program that quantify the relationship between the muscle plug concentration and the muscle fillet concentration so that selenium concentrations from plugs can be appropriately compared to the muscle tissue criterion element. Currently, EPA is conducting a study to better define the relationship between selenium concentrations in muscle plugs and muscle fillets. Soon EPA will have more information about the representativeness of muscle plugs and the analytical techniques for processing muscle plugs.

States or authorized tribes might choose to use whole-body or muscle tissue samples because egg-ovary tissue is not available year-round, or because existing monitoring programs can incorporate such selenium analysis into their existing fish tissue monitoring strategies. It is reasonable to collect whole-body or muscle tissue samples to gather data when adding selenium fish tissue sampling into an existing monitoring program or sampling for selenium in areas without known sources. If data indicates the need for a site-specific criterion or criterion element, agencies may want to expand their sampling to include egg-ovary tissue. States or authorized tribes might also choose to use whole-body samples because small-bodied species are the most appropriate to sample in a particular situation (Beatty and Russo 2014). In some low order streams only small-bodied species may be available for sampling. Collecting whole-body samples in these situations will allow for enough tissue mass for selenium analyses.

A specific case where sampling whole-body or muscle tissue is recommended over sampling egg-ovary is for sampling juvenile Pacific (smolt) salmonids. Anadromous fish species like salmon start their lives in freshwater, then as juveniles (e.g., smolts) migrate to the ocean, where they stay until adulthood before migrating back into freshwater to spawn. Notable among these species are the Coho, Chum, and Chinook salmon, and marine adapted Rainbow Trout (steelhead). Adult anadromous females (in the genus *Oncorhynchus* – except steelhead and Brown Trout) stop eating prior to re-entering freshwater environments as part of the

physiological modifications required for the migratory spawning process, and thus, lack exposure to freshwater selenium sources. They are also semelparous (except steelhead), meaning they die after spawning so there is no post-spawn residual exposure. Since adults of these species are not residents of the water body, the selenium concentrations will not be representative of localized freshwater selenium sources (see section 6.4.1 of EPA's *Aquatic Life Ambient Water Quality Criterion for Selenium–Freshwater 2016*) (USEPA 2021a). Given this lifecycle, it is not appropriate to sample adult semelparous anadromous fish for assessment of freshwater water bodies, as their selenium concentrations will not be representative of selenium exposure from where they are sampled. Instead, the juvenile fish from these species should be sampled, so they will reflect local selenium concentrations. An exception are landlocked variants of Striped Bass that cannot migrate out to sea, or hybrids (e.g., “wipers” which are striped bass-white bass crosses) in the Midwest. Adult fish in these landlocked populations may be representative of localized freshwater selenium concentrations, and thus appropriate for sampling. Although more uncertain, some studies indicate that selenium might affect endpoints such as juvenile growth and survival (Hamilton et al. 1990, DeForest and Adams 2011), therefore the most appropriate tissue to sample for Pacific anadromous salmon smolt is the whole body.

2.2 Sample Type

States and authorized tribes have flexibility in the type of sample that is collected to represent an instantaneous measurement of selenium in a fish population at a given site (see criterion duration footnote 6). Samples can include composites of multiple fish or the collection of individual fish in the population. The field sampling and analytical considerations for both sample types are described below.

2.2.1 Composite Samples

Composite samples are homogeneous mixtures of one type of tissue (e.g., egg-ovary, whole-body, or muscle tissue) from two or more individual organisms of the same species collected at a particular place and time and analyzed as a single sample. Composite samples can be useful for collecting enough tissue from small fish species to perform the appropriate analyses. Composite samples also allow for the analysis of additional target analytes if fish tissue samples are being collected as part of a broader fish tissue monitoring effort. Because chemical analytical costs are usually higher than field costs, using composite samples may be a cost-effective way to represent average selenium tissue concentrations in target species sample populations, by reducing the number of individual chemical analytical samples that are needed to characterize concentrations in the sample population (Patil et al. 2011). Composite sampling may also help with the issue of determining how to incorporate a sample with a concentration below the method detection limit (MDL) into an average, as the composite represents a physical averaging of the samples (USEPA 2011). Composite samples have also been shown to provide equivalent estimates of the mean compared to individual samples (Zhou et al. 2018, USEPA 1995). However, with composite samples, extreme contaminant concentration values for individual organisms are attenuated (Patil et al. 2011). Information from each individual sample is lost, which may mean losing information about spatial or temporal trends. Also, if a set number of fish are being analyzed,

compositing those fish rather than analyzing them individually will result in fewer data points, which can potentially lead to having less power in statistical analyses. However, this will be dependent on the data set and knowledge about the underlying sampling distribution. Hitt and Smith 2015 found that composites of two to four fish did not decrease power relative to individual samples when the sampling distribution was known (but power did decrease when an empirical sampling distribution was constructed instead of being known), because the composites had greater precision for estimating the mean.

Current EPA guidance on fish tissue monitoring recommends collecting three to ten individuals for a composite sample for each target species, as availability allows (USEPA 2000). EPA's Fish Advisory Guidance Volume 1 (USEPA 2000) also recommends collecting at least two composite samples at each site, and encourages a third, in order to properly estimate the site variance. An alternative approach may be to collect a greater number (five or greater) of smaller composites (two to three fish), which would increase sample size and statistical power, but still minimize resource expense compared to individual sampling (USEPA 2018, Hitt and Smith 2015). Section 6.1.2.7.1 of EPA's Fish Advisory Guidance Volume 1 ("Guidelines for Determining Sample Sizes") maintains that it is not possible to recommend a single set of sample size requirements for all fish contaminant monitoring studies (USEPA 2000). Rather, EPA presents a more general approach to sample size determination that is both scientifically defensible and cost-effective. Table 6-1 in that guidance shows the varying precision achieved by using additional numbers of individuals per composite and additional replicate composite samples. The data suggest that greater precision in the estimated standard error is gained by increasing the number of replicate composite samples than by increasing the number of fish per composite (USEPA 2000).

In EPA's National Lake Fish Tissue Study, composites were generally required to include five fish (USEPA 2002a). This composite size represented a reasonable number of fish that also satisfied statistical requirements for this study. If a state does not have previous data available for its sampling location, EPA recommends, in most waters, composites of five fish be used for fish tissue monitoring for CWA implementation of the selenium criterion. EPA recognizes that sometimes it might not be possible to collect a five-fish composite. In these cases, EPA encourages the state or authorized tribe to get as close to five fish as possible in the composite.

When the monitoring objectives include a desired level of statistical power, states and authorized tribes should consider additional information to determine the appropriate number of individuals per composite sample and number of replicate composite samples. Site-specific estimations of the population variance of selenium concentrations, fisheries management considerations, and statistical power considerations could be used to inform the sample size number to reach the desired outcome for the monitoring objectives. For example, fewer replicate composite samples and/or fewer individuals per composite sample may be required if the variance of the selenium concentration in the fish population at a site is small. In this case, it would not be cost-effective to use sample sizes that are larger than required to achieve the desired statistical power (Patil et al. 2011). Alternatively, in some instances a state or tribe may want to collect composites containing more than five fish or collect more replicates to have a more precise estimate of the mean. This may be desired at sites that are expected to have high variability.

The spatial variability of a site should also be considered when collecting composite samples. If a site is particularly large with high variability in selenium throughout, the site may need to be divided into subsites and composites collected from within each subsite to appropriately quantify the selenium impacts at that site.

EPA recommends that fish used in a composite sample meet the following specifications (USEPA 2000):

- All the same species.
- Of similar size so that the smallest individual in a composite is no less than 75% of the total length (size) of the largest individual (the “75% rule”; this “75% rule” does not apply to fish from which egg-ovary samples are collected).
- Collected at the same time (i.e., collected as close to the same time as possible but all samples should be collected within a week of each other).
- Collected in sufficient numbers to provide a composite homogenate tissue sample of at least 20 grams wet weight for selenium analysis.

EPA’s Fish Advisory Guidance Volume 1 (USEPA 2000) recommends including an equal number of fish in each composite sample and collecting two to three composite samples. However, when sampling fish in waters potentially impacted by selenium, the number of composite replicates may be determined on a case-by-case basis. This decision would primarily be based on the amount of variation in selenium concentration expected at the site and the number of individuals of the target species present at the site. States and authorized tribes who want to ensure the highest level of statistical power may want to collect multiple smaller composites to increase their sample size, which will increase their ability to detect differences from the criterion (Hitt and Smith 2015, USEPA 2018).

As species have different selenium bioaccumulation potentials and different sensitivities to selenium (USEPA 2021a), it is not scientifically defensible to create a composite sample that consists of more than one species. Compositing individuals that are the same genus, but not the same species is not appropriate. Accurate taxonomic identification is essential to prevent the mixing of species in a sample.

EPA recognizes that, in contrast to other bioaccumulative contaminants in fish, selenium concentrations are generally conserved across fish size (May et al. 2008). However, data on this topic is limited and variation in selenium concentrations might still be introduced by sampling fish of varying sizes. In particular, variation may be introduced if the species changes trophic levels or feeding ranges over the course of its development. Therefore, EPA recommends following the “75% rule” for the sizes of individual specimens included within a composite when sampling whole-body or muscle tissue (this 75% rule recommendation does not apply to fish collected for egg-ovary samples).

Individual organisms used in a composite sample ideally should be collected at the same time (if possible) so that temporal changes in contaminant concentrations are minimized. A best practice is to collect all organisms included in the composite sample within a week of each other so that

the composite sample accurately reflects the selenium concentration of fish in that water body at that time.

EPA recommends collecting a tissue sample of at least 20 grams wet weight (ww) for analysis, for both composite samples and individual samples. When creating composite samples from muscle tissue or egg-ovary tissue, an equal mass of homogenized tissue from each fish should be combined to create the composite, and then 20 grams ww should be sampled from the composite for analysis. When creating composite samples from whole bodies, all fish included in a composite should meet the 75% rule. While the specific amount of tissue needed for analysis will be dependent on the laboratory and analytical method used for analysis, 20 grams ww is a reasonable estimate of the amount of tissue needed for typical selenium analyses. This mass allows for 5 grams of tissue for the selenium analysis, 5 grams of tissue for a matrix spike sample, 5 grams of tissue for dry weight analysis, and then a final 5 grams of tissue available in case there is a problem with one of the other analyses and a procedure needs to be repeated. In addition, a sample of this size allows for a quality control sample to be processed, which can assure homogeneity of the tissue sample. If after these analyses enough remaining tissue mass is available, agencies may want to retain tissue from the individual fish, in case future analyses are wanted on the individuals. Most fish tissue samples for selenium analysis are processed as wet tissue, resulting in a selenium concentration based on wet weight. As the national CWA section 304(a) recommended selenium criterion is based on a dry weight selenium concentration, an analysis of the moisture content of the tissue needs to be performed and the wet weight concentration needs to be converted to a dry weight concentration. See Appendix C for more information on how to perform a wet weight to dry weight conversion. Monitoring agencies typically collect composite samples for other analytes when sampling for fish consumption advisories. If agencies currently discard or archive the composite homogenates that are in excess of their current analytical needs, the excess tissue could be used, if adequate in mass, for selenium analysis. Agencies could also collect additional tissue mass and add selenium as an analyte to their sampling protocol.

EPA recognizes that if a state or authorized tribe collects muscle plugs they will likely be collecting sample masses of less than 20g ww per fish. States and authorized tribes should assure that the mass of tissue they are collecting and the analytical methods that they are using will allow for accurate quantification of selenium. Compositing muscle plugs (one per fish from multiple fish) may be one way to achieve sufficient mass for analysis.

EPA also recommends collecting an equal number of fish in each composite, as this simplifies the statistical methods needed to analyze the results from this analysis. With equal numbers of fish, the arithmetic average of the replicate composite measurements is an unbiased estimator of the population mean. When unequal numbers are used, the arithmetic average is no longer unbiased. Instead, a weighted average of the composite measurements is calculated, where the weight for each composite reflects the number of fish in each composite sample. Oftentimes fish are lost or damaged prior to compositing. When several fish are damaged or lost, the allocation of the remaining fish to composites may be reconfigured to allow equal numbers of fish in composites. During this reconfiguration process, a sampler may be faced with the choice of either making composites composed of an equal number of fish or to follow the 75% rule. EPA

recommends adhering with the 75% rule over having an equal number of fish in each composite if both parameters cannot be met. If an equal number of fish cannot be included in each composite, care should be taken to adjust the statistical procedures to account for the unequal allocations (USEPA 2000).

2.2.2 Individual Samples

An individual sample is a discrete sample from a single fish, and can be an egg-ovary, whole-body, or muscle (fillet) tissue sample. Use of composite samples for selenium fish tissue monitoring is acceptable, but there are some instances where collecting individual fish may be desirable.

Analysis of individual fish samples may be of interest when collecting data for a site-specific study or for statistically evaluating patterns in selenium concentrations over time or space. Analysis of individual fish may also be important when evaluating Se concentrations in critical species, where understanding potential toxicity at the individual level is important. Analysis of individual samples also allows for the evaluation of spatial and temporal differences among individuals of a species of similar size or across the population of a species residing in a specific water body.

For water bodies or segments that are known to be impacted by selenium, individual samples describe the range of variability within a population including characterizing extreme values. Individual samples may also provide better information about selenium source-exposure relationships in large water bodies. Individual samples may also allow for the identification of fish that are migrant or transient in a population, since that fish may have a higher or lower concentration of selenium than other fish in the area. If studies are being conducted to monitor trends, EPA recommends sampling fish of the same species at each location and at each time interval so that data are spatially and temporally comparable.

If using individual samples to calculate mean selenium fish tissue concentrations, all fish should be the same species and collected from the same water body (or site for large water bodies) within the same sampling period (ideally within the same week). The fish should be of similar size (within the 75% rule) and the samples should be of the same tissue type. EPA recommends targeting at least five fish (per site or per location for larger sites) for individual analysis to achieve measurements of a reasonable statistical power (see discussion of statistical power in the “Composite Sample” discussion above). Greater or fewer fish samples may be needed based on the variation of selenium at a particular site. Those entities desiring greater statistical power for their analyses should collect additional fish samples. In the event that collecting at least five individuals of one species is not possible, fewer specimens may be collected, but the statistical power of the analysis will be affected (Hitt and Smith 2015). A power analysis should be conducted in these situations to assure that enough data is being collected to detect a difference. As with composite samples, EPA recommends collecting 20 grams ww as a minimum tissue mass per individual fish for analysis and QA/QC.

With individual fish samples, as well as with fish samples collected to be a part of a composite sample, EPA recommends documenting and reporting additional information about the fish

samples to assist with interpretation of the data. Useful information to document includes the species, length, age (if can be estimated), weight, and sex of the fish samples. Documenting information about the sampling site and day can also be useful for data interpretation. Samplers should note the location of collection by GPS if possible, sampling date and weather. Samplers may want to also note flow rate and other characteristics about the site that may affect the concentration of selenium at the sampling sites.

2.3 Target Species

The two main factors to consider when selecting a target fish species to sample for the implementation of the national CWA section 304(a) recommended selenium criterion are (1) a species' toxicological sensitivity to selenium, and (2) a species' bioaccumulation potential for selenium. In addition, it is important to consider a species' potential for exposure to selenium. EPA recommends that states and authorized tribes create a priority list of target species for sampling teams. This list should identify the primary target species and alternatives species if the primary species is not present or not present in sufficient numbers for sampling. If data cannot be acquired for a desired target species, data from any fish species can be used to make assessment decisions, as the criterion was developed for use with any fish species.

A species' toxicological sensitivity to selenium, for the purposes of implementing the national CWA section 304(a) recommended selenium criterion, is defined as a species' or a surrogate species' EC₁₀ (10% effect concentration). An EC₁₀ is the concentration of a chemical that is estimated to result in a 10 percent effect in a measured chronic endpoint (e.g., growth, reproduction, or survival). For the national CWA section 304(a) recommended selenium criterion, a species' EC₁₀ is the concentration of selenium within egg-ovary tissue that results in a 10% effect on a reproductive endpoint for that species. Based on the best available and acceptable reproductive-effect studies, as well as extensive analyses, EPA developed a species sensitivity distribution (SSD) to support the derivation of the national CWA section 304(a) recommended selenium criterion based on species' EC₁₀s (See Table 3.2 in USEPA 2021a). This SSD for the national CWA section 304(a) selenium criterion indicates that the four most sensitive genera for fish reproductive effects (in decreasing order) are *Acipenser*, *Lepomis*, *Salmo*, and *Oncorhynchus*.

The bioaccumulation potential of a species is largely determined by its dietary composition and the exposure of its prey items to selenium. Consumption of benthic invertebrates tends to drive greater selenium bioaccumulation than consumption of plankton (Schneider et al. 2015, Simmons and Wallschläger 2005). Amongst benthic organisms, the consumption of mollusks tends to drive greater selenium bioaccumulation than consumption of other benthic invertebrates (Luoma and Presser 2009, Stewart et al. 2004). Mollusks, such as mussels and clams, accumulate selenium to a much greater extent than planktonic crustaceans and insects due to higher ingestion rates of both suspended particulate-bound selenium (algae) and dissolved selenium from the water column through filter feeding. Mollusks also have a lower selenium elimination rate (Johns et al. 1988, Reinfelder et al. 1997). The greater bioaccumulation of selenium in benthic organisms suggests that bottom feeding fish may have higher selenium levels, at least for the lifecycle that ties their energy needs to food webs with benthic organisms. Other studies (Saiki et al. 1993, Saiki and Lowe 1987) have shown that detritivores may also be exposed to high levels of dietary selenium, as high concentrations of selenium were measured in detritus. The

bioaccumulation potentials of organisms at higher trophic levels (such as piscivores) are dependent on its food chain's cumulative exposure to and bioaccumulation of selenium. Trophic transfer factors (TTFs) provide a numeric representation of bioaccumulation between a consumer and its diet. A composite TTF ($TTF^{\text{composite}}$), which is the product of TTFs at each trophic level of a consumer's food chain, represents the overall TTF for a higher trophic level organism. In addition, composite TTFs account for the proportion of different food sources in a consumer's diet. Evaluations of TTFs may be helpful in determining which species to target for sampling. More information about TTFs can be found in Appendix B, Section 3.0 of the *Aquatic Life Ambient Water Quality Criterion for Selenium – Freshwater 2016* (USEPA 2021a). In addition, an explanation of how composite TTFs are calculated is included in Appendix E.

As both a species' sensitivity to selenium and bioaccumulation of selenium influences whether it will be impacted by selenium, states and authorized tribes should consider both when selecting a target species and designing fish tissue sampling plans. EPA recommends the following prioritization scheme for selecting a target species.

1. Sample the species from within the four most sensitive genera that has the greatest bioaccumulation potential.
2. If no species are present from the four most sensitive genera, sample the species with the greatest bioaccumulation potential.
3. If no species are present from the four most sensitive genera and if all species have similar bioaccumulation potential, sample the species from within the most sensitive genera present at the site (sensitive according to the SSD from USEPA 2021a).

States and authorized tribes should begin with targeting the fish species from within the four most sensitive genera that have the greatest bioaccumulation potential. As described above, the SSD for the national CWA section 304(a) recommended selenium criterion determined that the four most sensitive genera for fish reproductive effects (in decreasing order) are *Acipenser*, *Lepomis*, *Salmo*, and *Oncorhynchus*. When selecting a fish species to sample from within these genera, states and authorized tribes should consider the diet and exposure of all the species at the site that are within those four most sensitive genera and select the species that has the greatest potential to bioaccumulate selenium. For example, if a site has multiple species of *Lepomis* present (e.g., Bluegill and Redear Sunfish), the state or authorized tribe should sample the species that has the greatest bioaccumulation potential (Redear Sunfish). In the San Francisco estuary, sturgeon are monitored not only because they are sensitive to the toxic effects of selenium (low EC_{10}), but also because their primary prey at that site (clams) bioaccumulate selenium very efficiently. As a result, sturgeon receive large doses of selenium and may be more likely to bioaccumulate selenium to levels of concern than another species. Table 3 provides a summary of these genera (highlighting information for the specific species tested), their relative sensitivity, their general habitat type (warm water [WW] or cold water [CW]), and their estimated relative bioaccumulation potential based on consideration of typical diet and trophic level. Fish that consume primarily benthic organisms will tend to exhibit greater selenium bioaccumulation than fish that feed higher in the water column at the same trophic level (Schneider et al. 2015, Simmons and Wallschläger 2005). Table 3 also provides a representative list of species, that are within the same genus as the tested sensitive species, and that could be considered as surrogates for tissue collection. A more comprehensive list of these surrogate species, along with details about relevant characteristics and occurrence is presented in Appendix D.

If no species from the four most sensitive genera are present at the site, then the state or tribe should target the fish species with the greatest bioaccumulation potential. As stated above, bioaccumulation potential will be predominantly determined by the diet of the species.

Composite trophic transfer factors can provide a numeric representation of bioaccumulation between a consumer and its diet and help inform the decision about which species to sample.

If all species that are present at a site have similar diets and bioaccumulation potential, then the state or tribe should target the most sensitive species. This will be the species with the lowest EC₁₀. If the EC₁₀ of a particular species is unavailable, sensitivity can be estimated from the EC₁₀ of a closely related taxon.

In addition to sensitivity and bioaccumulation potential, there are a number of factors that should be considered when identifying appropriate fish species for collection. The following summarizes some key points for consideration:

1. White sturgeon (and available surrogates) are distributed in large river systems in the US and among the most sensitive genera (based on available data). These species are typically sampled by specialized monitoring programs (e.g., USFWS and NOAA-NMFS). Coordination with these existing programs may provide for expanded sampling opportunities or the use of existing selenium fish tissue data. Several species and specific populations of species within the *Acipenser* genus are federally listed species under the Endangered Species Act and may not be appropriate to sample. USFWS, NOAA-NMFS, and appropriate state agencies should be consulted before sampling any federal or state listed threatened or endangered species.
2. Bluegill, and the related sunfish species in the genus *Lepomis* are widely distributed in WW habitats, while trout species (particularly Rainbow and Brown trout) are widely distributed in CW habitats. These species are frequently targeted by monitoring programs in states and tribal lands with WW and CW habitats, respectively, offering a possible opportunity for leveraging these existing sampling programs for the collection of tissue for selenium analysis.
3. Smaller WW and CW systems (e.g., wadeable streams), which are not typically targeted by state fish tissue contaminant monitoring programs, are often dominated by cyprinid (minnow) species and may represent a source of fish tissue for selenium sampling in water bodies where other species may not be present. Some of these species are shown in Table 3 and a broader range of these species are shown in Appendix D. Given the large number of minnow genera and species and the diversity of their trophic strategies and habitats, the sensitivity and bioaccumulation potential for individual members of this diverse group must be considered when evaluating a candidate species for consideration in tissue sampling. However, state biomonitoring programs typically sample these waters for IBI metrics, and their expertise and sampling program could be leveraged to target species and obtain representative samples.
4. Although generalizations can be made about the potential for bioaccumulation within fish species, when developing a sampling plan, the potential for bioaccumulation should be considered for the specific area being evaluated. Potential for bioaccumulation within any given species can vary significantly with location-specific factors, including prey type and availability and the nature of selenium distribution in the environment.

5. There are a number of species for which toxicity data are not available, but for which dietary information allows us to characterize the potential for selenium bioaccumulation. Some of these species are collected as target species in state monitoring programs and could be considered to characterize selenium tissue concentrations in the absence of sensitive species, if available data indicates these species are bioaccumulative.

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Table 3. Recommended Fish Target Species for Collection Based on Selenium Sensitivity and Bioaccumulation Potential

Sensitivity Ranking (Based on Egg-Ovary EC ₁₀)	Genus, Common Name	Habitat (WW/CW)	Bioaccumulation			Additional Representative Surrogate Species**
			Expected Relative Bioaccumulation Potential	Adult Diet	Trophic Level	
1	<i>Acipenser</i> , White Sturgeon*	WW	Medium-High	Invertivore Piscivore Molluscivore	TL3/TL4	Shortnose Sturgeon*, Lake Sturgeon*
2	<i>Lepomis</i> , Bluegill	WW	Medium	Invertivore Piscivore	TL3	Pumpkinseed*, Redear Sunfish*, Green Sunfish*, Redbreast Sunfish*, Longear Sunfish, Warmouth, Orangespotted Sunfish, Redspotted Sunfish, Bantam Sunfish, Northern Longear Sunfish, Spotted Sunfish
3	<i>Salmo</i> , Brown Trout*	CW	Medium-High	Invertivore Piscivore Molluscivore	TL3/TL4	None
4	<i>Oncorhynchus</i> , Rainbow Trout	CW	Medium-High	Invertivore Piscivore	TL3/TL4	None
	<i>Oncorhynchus</i> , Cutthroat trout	CW	Medium	Invertivore	TL3	None
5	<i>Micropterus</i> Largemouth Bass	WW	Medium - High	Invertivore Piscivore	TL4	Smallmouth Bass, Spotted Bass, Redeye Bass
Not Ranked	<i>Pimephales</i> Fathead Minnow***	WW	Low-Medium	Herbivore Invertivore	TL2/TL3	WW: Satinfish Shiner, Red Shiner, Golden Shiner*, Bull Chub*, Creek Chub, Roundtail Chub, Thicklip Chub*, Striped Shiner, Central Stoneroller, Blacknose Dace, Speckled Dace, Cutlips Minnow*, River Darter*, Arkansas Darter*
						CW: Redside Shiner, Peamouth Chub, Mottled Sculpin

WW = warm water, CW = cold water, TL = trophic level

* Fish species that consume mollusks (clams, mussels, snails) as part of their diet and are anticipated to have relatively high bioaccumulation potential. For brown trout, molluscivory is incidental and will likely only be significant on a site-specific basis where mollusks are abundant.

** Species are surrogates for sensitivity based on taxonomic relationships.

*** Fathead minnow is a surrogate for small bodied fish species inhabiting Wadeable streams.

Along with sensitivity and bioaccumulation, it is also important to consider how a species' habitat preferences, feeding regimes, and/or home ranges will affect their selenium exposure. Species with smaller home ranges may be at risk of greater selenium exposure if the elevated selenium is localized to their home range or their prey's home range. Species with larger home ranges may not represent local selenium exposure. States and authorized tribes should target species with home ranges that closely match the site being evaluated, so that the fish reflect exposure to selenium at that particular site. Local fish biologists may be able to help states and authorized tribes identify species' habitat preferences, feeding regimes, and home ranges. If possible, migratory species and highly mobile species should be avoided. Highly mobile fish species such as potamodromous and anadromous species could travel back and forth between areas with low and elevated selenium concentrations, resulting in variable selenium tissue concentrations (Beatty and Russo 2014). It is possible that typical selenium exposure concentrations of migratory adults would be lower than concentrations at rearing grounds, therefore these fish may not reflect selenium concentrations that other fish species would experience at their time of spawning. Given this issue, resident species should be the first choice when selecting a target species. Recently stocked fish should also be avoided, regardless of species, since their residence time before sampling may be too short to provide a representative sample. EPA recommends speaking with local fish and game authorities to understand stocking frequency in waters where fish are being sampled for selenium analysis.

If migratory or highly mobile species must be sampled, then EPA recommends that sampling plans account for the life history of these species so that the correct locations for sampling within a watershed are selected. Before sampling these species, knowledge about the migratory extent of these species and the time spent in each location should be known so that the influence of selenium at each habitat location can be accounted for. EPA recommends consulting local fish biologists (state or academic) for information about the migratory patterns of local fish populations. If Pacific anadromous species are selected as target species to be used for sampling, EPA recommends that states and authorized tribes sample the whole-body of juveniles (smolts) and compare the concentration to the whole-body criterion element. This recommendation is due to the lack of selenium exposure to adult salmonids from freshwater prior to reproduction (see section 6.4.1.1 in USEPA 2021a).

One or more species that are sensitive to selenium (e.g., Bluegill, Rainbow Trout, Brown Trout) are commonly present in water bodies. However, if they or species that potentially bioaccumulate high levels of selenium are not available in sufficient numbers to sample, then other species that are available in sufficient numbers may be used for fish tissue monitoring, including species known to be tolerant to selenium. These selenium tissue samples can still be compared to the appropriate tissue criterion elements (see Table 1), which are designed to be protective of the entire aquatic community and can still be used to make an assessment decision. In addition, if fish tissue data were collected for an alternate purpose from other fish species than the ones recommended in this section, that data can also be used for an assessment decision.

As there are a large number of considerations involved in the selection of a target species, EPA recommends that sampling teams develop a sampling priority plan before going into the field. This plan should start by identifying the species and factors that will increase selenium exposure at a site. From there, an initial target species should be selected and any decisions about sampling locations can be made. The plan can then identify a prioritized list of species that the state or authorized tribe will sample if the initial target species is not present in sufficient numbers. EPA

recommends that the plan also include what type of sampling gear is required for the collection of each species. In addition, the plan should specify if collection permits are required for fish tissue sampling, and if so, include instructions for acquiring them before the sampling event. Having a clear plan will ensure that the fish collected will provide the most representative data of selenium conditions at the site.

States and authorized tribes may want to limit the number of target species that are sampled within their state or tribal waters. The use of a limited number of target species allows for the comparison of fish contaminant data among sites over a broad geographic area. It is difficult to compare contaminant monitoring results within a state or tribe or among states or tribes unless the data are from the same species because of differences in habitat, food preferences, and rate of contaminant uptake among various fish species. However, it is impracticable to sample the same species at every site. Limiting the number of species collected across a state allows for the comparison of contaminant data from across a state or region. Given this, EPA recommends that states and authorized tribes evaluate the range of sensitive species with high bioaccumulation potential across their state or tribal waters and determine which ones may be able to be sampled at multiple locations within the state or tribal lands.

If a state or authorized tribe is sampling fish tissue for the development of a site-specific water column criterion element, they may want to expand their sampling to include multiple species to better understand the system at their site. Sampling species that have high bioaccumulation potential, as well as species with high sensitivity to selenium, could provide a more complete picture of selenium dynamics at the site.

Lastly, care should be taken to avoid sampling threatened or endangered species when selecting a target species. For example, although *Acipenser*, *Salmo*, and *Oncorhynchus* are three of the four most sensitive genera, many species within these genera are threatened or endangered and thus are not suitable for sampling. Before sampling from these genera and other genera with federally listed species, EPA recommends speaking to local fish biologists to ensure that the target species is not threatened or endangered and to confirm that the population is healthy enough to withstand the sampling pressure. When conducting monitoring to ensure the protection of a federally listed species, states and authorized tribes should target surrogate species that have similar taxonomy (preferably at the genus level), diets, and trophic levels. Species with similar taxonomy, diets, and trophic levels should have similar selenium sensitivity and bioaccumulation as the threatened or endangered species. If a taxonomically similar surrogate is not present, then states and authorized tribes should target a species with a similar diet and trophic level.

2.4 Sampling Locations

Several factors should be considered when selecting where fish should be sampled (to be analyzed either individually or as a composite) to accurately characterize the concentration of selenium at a site of interest. The spatial extent of the site needs to be defined and the factors that may affect selenium variability throughout the site need to be identified so that they can be considered in the design of the sampling plan. The selection of a site and how its boundaries are defined will be influenced by the objective of the monitoring and by past monitoring activities (see section 3.0 *Leveraging Existing Fish Tissue Monitoring Programs and Sample Design*). The

factors informing the site definition and the subsequent selection of sampling locations will vary, and may include:

- monitoring objectives (e.g., assessment, site-specific selenium studies);
- water body type and site hydrology (lotic vs. lentic);
- water body size;
- selenium sources and location of sources;
- aquatic habitat variability; and
- physical barriers to fish movement.

These factors (and potentially others) will influence the definition of the site boundaries, decisions about where fish are collected from within the site, and decisions about how many fish need to be collected at the site. Discussed below are some of the factors that EPA recommends considering with site and sample location selection, but particular situations may warrant the consideration of other important factors.

2.4.1 Water Body Type

Selenium concentrations and bioaccumulation patterns are different in lotic (flowing waters such as rivers and streams) versus lentic (very slow moving or still waters such as lakes and reservoirs) environments. Water residence time is typically shorter in lotic systems than in lentic systems, and subsequently, aquatic organisms living in lentic systems tend to bioaccumulate proportionately more selenium than organisms living in lotic systems (ATSDR 2003; EPRI 2006; Luoma and Rainbow 2005; Orr et al. 2006; Simmons and Wallschlägel 2005). In addition, lentic water bodies tend to have greater reducing conditions (conditions that lead to reduction reactions and reduced ionic species of selenium such as selenite) which create an environment where selenium accumulates in sediment more readily and may also lead to higher bioavailability in the water column (Luoma and Rainbow 2008, Simmons and Wallschlägel 2005). Benthic organisms in lentic systems can then be exposed to higher concentrations of selenium in the sediment, leading to increased bioaccumulation potential in other organisms feeding on the benthic organisms (Simmons and Wallschlägel 2005, Orr et al. 2006). Hillwalker et al. (2006), for example, found that the body burden concentrations of selenium in insects within similar taxa were up to seven times greater in lentic systems than lotic systems within the same watershed.

When sampling fish, consideration should be given to the different flow characteristics of the site that is being sampled along with the locations where fish are feeding and obtaining their selenium body burdens. Some areas of a lotic site may have lentic characteristics and vice versa. For example, some rivers may have slow moving pools or backwaters that have characteristics similar to lentic environments. Human-made lakes and reservoirs may have some features that are intermediate between typical lotic and lentic systems. For example, reservoirs tend to be longer and narrower than natural lakes, and generally have a shorter water retention time than a natural lake of comparable volume (Thornton et al. 1990). When sampling sites, attempts should

be made to sample all habitat types to appropriately characterize the range and distribution of selenium concentrations at a site.

2.4.2 Water Body Size

Generally speaking, the variability of selenium within resident fish populations would be expected to be low when the spatial and temporal variability of selenium concentrations across all compartments of the ecosystem are low (e.g., water column, sediment, etc.). As the area of a site increases, the spatial and temporal variability is expected to also increase, thus increasing the number of samples needed to characterize the selenium concentrations in the resident fish populations. If the water body is sufficiently large that sub-populations are expected to be present (potentially applicable to large reservoirs), it could be advantageous for the sub-populations to be represented separately in the dataset.

2.4.3 Site-specific Studies for Water Column Translations

If a site-specific water column criterion element is being developed, a study should be designed that captures data which appropriately reflects the site (e.g., captures spatial, temporal, and habitat variability). To support a site-specific water column translation, data beyond what is necessary for other CWA implementation purposes is typically required (e.g., additional sampling locations, sampling times, species of fish, and/or selenium measured in multiple matrices). The extent of the sampling and type of data collected will depend on the size and complexity of the site. It will also depend on whether there are any discharges of selenium into the site. A “site” may be a state, region, watershed, water body, segment of water body, category of water (e.g., ephemeral stream), etc. Regardless of how the site is defined, the site-specific water column translation should be derived to provide adequate protection of aquatic life for the entire site, including both areas upstream and downstream of a discharge if one is present at the site. To assure protection for aquatic life throughout the entire site, fish should be sampled from locations where selenium is expected to bioaccumulate the most (areas of the site with more lentic properties and areas where selenium may be elevated due to source contributions). In addition to sampling from the area of greatest exposure, agencies may want to sample fish from various locations in the site to understand the dynamics of selenium within that site. With that knowledge, the site-specific water column translation can be designed to be protective of the most vulnerable fish community. When the area of interest is a segment of a water body, it is important to understand how the segment is characterized in the state or tribal WQS, the representativeness of the partial segment to the regulatory segment as a whole, as well as its relation to downstream segments that may support more sensitive fish species (e.g., lower EC₁₀ or threatened and endangered species) than the immediate area of interest. In these situations, the sampling may include fish populations in the immediate area of interest and the downstream water body.

Additional information related to sampling fish tissue to support a site-specific water column translation can be found in Appendix K of *Aquatic Life Ambient Water Quality Criterion for Selenium – Freshwater 2016* (USEPA 2021a) and in *Technical Support for Adopting and Implementing EPA’s 2016 Selenium Criterion in Water Quality Standards, Draft* (USEPA 2021c). One example of a method for conducting a site-specific water column translation can be

found in the *Draft Translation of Selenium Tissue Criterion Elements to Site-Specific Water Column Criterion Elements for California Version 1, August 8, 2018* (USEPA 2018). EPA recommends engaging with EPA regional staff early in the development of a site-specific water column translation to discuss study design and data needs.

2.4.4 Point Sources

When selecting sampling locations, samplers should consider where and how selenium is entering a water body and determine whether exposure is relatively equal throughout the water body or if some sections of the water body have greater exposure. The sampling objectives will provide direction on how known point sources (e.g., discharge, tributary with elevated selenium, irrigation return canal, groundwater discharge) and any associated mixing zones should be taken into consideration when collecting fish tissue samples. When the objective is to collect data to support assessment decisions, the goal is to measure the mean selenium concentration in the target population throughout the sampling reach. Therefore, when a point source is located within the defined sampling reach, there is no reason to avoid sampling fish from areas of incomplete mixing resulting from a discharge or tributary. Given the mobility of many fish taxa, it is reasonable to expect that fish freely move in and out of areas of incomplete mixing when the conditions do not elicit an avoidance response (e.g., due to chemical or temperature gradients). In some discharge situations, fish can be attracted to the effluent and spend a significant portion of their time in the area of incomplete mixing. Also, depending on their life history, some fish species have a limited mobility range and may spend more time in the area of incomplete mixing if it overlaps with their territory, breeding grounds, or feeding grounds. Ultimately, jurisdictions should consider and document if there are any reasons to avoid tissue collection from a location adjacent to a known point source prior to sampling (e.g., conditions are not representative of the rest of the segment).

When the sampling objective is to characterize the contribution of selenium that a known point source is making to the water body, samples collected upstream and downstream of the point source should be assessed independently (i.e., not composited or averaged). The downstream sampling reach should be large enough to include samples collected within and downstream of areas of incomplete mixing to characterize the range of bioaccumulation potential in the tissue samples as the water column concentrations decrease. One way to do this would be to collect fish and water samples at regular intervals from the discharge to observe how both decrease downstream of the discharge. It is important to understand the hydrology in the system as this will influence the range and direction of transport of selenium from the discharge source to other portions of the water body/site. For more information on considerations related to selenium sources and the locations of those sources, see section 2.1 in *Aquatic Life Ambient Water Quality Criterion for Selenium—Freshwater 2016* (USEPA 2021a).

3.0 Leveraging Existing Fish Tissue Monitoring Programs and Sample Designs

3.1 Augmenting Existing Fish Tissue Monitoring Programs

Many states and authorized tribes have existing fish tissue monitoring programs that can be leveraged to collect fish tissue data to assess against the fish tissue criterion elements of the national CWA section 304(a) 2016 recommended selenium criterion. In 2010, forty-five states monitored chemical contaminants in fish tissue to assess risks to human health. Twenty-eight states identified selenium as a contaminant in their human health monitoring programs (USEPA 2011). These states can potentially modify their current programs to not only assess human health risks, but also assess attainment of the aquatic life selenium criterion. The design of an agency's existing fish tissue monitoring program will likely drive its approach to selenium monitoring. Agencies should evaluate how current sampling and analytical protocols can be modified to meet both the objectives of monitoring for risk to human health and aquatic life protection. Case studies are provided in the following sections as examples of programs that might have the capacity and framework to augment their existing monitoring strategies to include fish tissue monitoring for the selenium aquatic life criteria.

3.1.1 Consistency with Existing Programs

EPA recommends evaluating current fish tissue monitoring programs to determine how they can be augmented to implement the national CWA section 304(a) recommended selenium criterion. To the extent possible within a state or tribal program, EPA recommends that fish tissue monitoring for the selenium aquatic life criterion should be consistent with the state's current fish tissue monitoring practices regarding spatial and temporal considerations, species collected, and sample type collected. In this way, logistical modifications to a state's fish tissue monitoring program can be minimized. Muscle tissue is the most common type of sample collected and analyzed by monitoring programs. Less frequently states and authorized tribes collect and analyze whole-body samples. To maximize efficiency, a portion of these samples can be submitted for selenium analysis. States or authorized tribes will realize cost efficiencies by choosing to use whole-bodies or fillets that are already being collected for an existing monitoring program.

When utilizing existing sampling programs that are designed for human health protection to assess selenium levels for protection of aquatic life, states and authorized tribes should recognize the potential limitations of these data. That data may not represent areas most likely to be contaminated by selenium, most relevant time periods for sampling, or most appropriate species. Where deviations from existing state or tribal programs are necessary due to spatial or temporal considerations or species/sample type, these can potentially be accommodated by leveraging expertise and logistical assistance from other agencies with existing fish tissue monitoring programs. Since the selenium criterion applies to ecological risk and not human health, monitoring agencies could evaluate their target species list and determine if they include appropriate species for assessing selenium risk to aquatic life. See the discussion in section 2.3

about selecting target species and see *Aquatic Life Ambient Water Quality Criterion for Selenium – Freshwater* (USEPA 2021a) for more information about species' sensitivity and bioaccumulation potential.

A survey conducted in 2010 reported that 40 state agencies conduct fish sampling at regular intervals, and several conduct statewide, rotating basin sampling programs over a multi-year period (USEPA 2011). Many states and authorized tribes may be able to utilize their current tissue sampling programs to monitor for the selenium criterion as well. Agencies could monitor state- or basin-wide, and track progress in individual basins relative to other areas. Regular yearly sampling could be conducted, with intensified sampling in targeted basins as needed (see Table 4 for several documents that provide guidance for sampling and survey designs). Several states use a probabilistic survey design to select sampling sites. This type of sampling design can produce estimates that represent the condition of the whole watershed, and an estimate of random spatial variability (USEPA 2000). Probability-based sampling provides the basis for estimating the resource (i.e., fish population(s)) extent and condition, for characterizing trends in resource extent or condition, and for representing spatial patterns, all with known certainty (USEPA 2009). Additional or more targeted sampling approaches may be needed in areas where elevated selenium is associated with a known point of discharge. The case study below presents the Kansas Department of Health and the Environment's (KDHE) fish tissue monitoring program, which uses several designs for selecting sites. Based on the information available, it is likely that a state or authorized tribe with a similar program could take advantage of their current sampling strategy to perform screening level selenium analysis throughout their state or tribal area. Where selenium is already a primary parameter of interest, the state or authorized tribe may have the data to support more intensive studies in certain water bodies.

CASE STUDY: The Kansas Department of Health and the Environment

The Kansas Department of Health and the Environment (KDHE) currently collects fish samples annually from 30-50 fixed and rotating stations. The KDHE selects sites based on targeted, census, and probability-based sampling designs. Specific sub-program objectives determine the numbers, species, and sizes of fish collected from a particular water body, and the tissues and parameters of interest (http://www.kdheks.gov/befs/fish_tissue_monitoring.htm).

Highlights (KDHE 2013):

- Whole fish, muscle, muscle plugs, or other specific tissues were collected for different programs.
- Selenium was a primary parameter of interest.
- Specific tissues (such as egg-ovary) were analyzed for specific chemicals of concern known to accumulate in certain organs.

http://www.kdheks.gov/environment/qmp/download/Fish_Tissue_Part_III.pdf

This program is an example of an existing fish sampling program that could be enhanced to collect data for the implementation of the selenium criterion.

3.1.2 Temporal Considerations

Various temporal considerations will influence fish tissue monitoring strategies for selenium. These can include considerations related to the ecology of the fish (e.g. species' spawning season) or to abiotic environmental factors (e.g. weather conditions and river flows). Temporal considerations will influence decisions regarding which tissue type is sampled: egg-ovary, whole-body, or muscle tissue. For example, most fish species that are synchronous spawners spawn in the spring, making spring the prime season to sample egg-ovary tissues, yet sampling for health advisories is typically done in the fall when concentrations of contaminants are highest in muscle tissue (fillets). Agencies will need to consider their resources and determine which tissue type they would like to sample and at what time of year. If agencies plan to sample egg-ovary tissue, they should plan to sample right before spawning. If an agency plans to conserve resources and sample for both fish consumption advisories and the selenium criterion at the same time, whole-body or muscle tissue should be sampled outside of species-specific pre- and post-spawning windows. In this case, muscle (or whole-body) tissue can be composited and evaluated for the selenium aquatic life criterion in addition to contaminants of interest for fish consumption advisories. If the agency has information indicating that there may be seasonal differences in whole-body or muscle tissue concentrations, then agencies should plan to sample during the

season when the highest selenium concentrations are expected. Agencies may want to sample spring spawners in late summer or fall to avoid the potential for underestimating selenium body burdens. Selenium body burdens can be decreased directly post-spawn due to the selenium depuration from whole-body or muscle tissue via the maternal transfer of selenium to eggs and the subsequent release of eggs to the environment. If sex can be determined in the field, agencies may want to target male fish to avoid this possibility.

For egg-ovary tissue sampling, EPA recommends that agencies with fish tissue monitoring responsibilities consult with local fisheries biologists to determine the appropriate time for sampling specific species in their region in order to capture the specimens in their pre-spawning phase. These regional experts will be familiar with the local species and are able to use their best professional judgment to determine which species are appropriate for egg-ovary sampling and the appropriate sampling period based on spawning season.

3.1.3 Spatial Considerations

Monitoring agencies generally target high-use fishing areas, areas of special concern, and areas of suspected contamination (such as water bodies where fish advisories have been issued), when selecting sites for sampling fish tissue (USEPA 2011). As current fish tissue monitoring programs are typically designed to specifically address the risk to human health from fish consumption, these programs predominantly sample locations where fishing is common. This may lead to mostly sampling lakes and higher-order streams. States and authorized tribes using this sampling design should consider if these existing programs will adequately capture water bodies impacted by point and non-point sources of selenium and potential areas of selenium contamination. If not, agencies may want to modify sampling designs to target such areas for sampling.

Some states and authorized tribes may incorporate fish tissue sampling for selenium into a statistical survey designed to understand the distribution of tissue concentrations across the state or tribal lands. The underlying geology of a region may produce elevated selenium concentrations in certain areas and make nearby waterbodies prone to selenium bioaccumulation, particularly if anthropogenic activities increase the release of selenium into the system. This should be kept in mind when selecting sites, and when analyzing data from these areas (Beatty and Russo 2014) (See USGS map of selenium concentrations in soils and stream sediments: <https://mrdata.usgs.gov/geochem/doc/averages/se/usa.html>)).

Additional sampling locations may need to be added to a current monitoring program that are outside of areas that are typically targeted due to fishing use when sampling for the assessment of the selenium aquatic life criterion. For example, mine runoff may elevate selenium concentrations in headwater streams, which may not be normally targeted for fish tissue monitoring. Agencies should also consider a species' home range in relation to the location of a known selenium source (e.g., the migratory patterns of a certain species versus the location of a power plant on a reservoir). It is also important to consider the relationship of an upstream source to downstream habitats, particularly when downstream habitats have characteristics that will lead to greater selenium bioaccumulation (e.g., lentic systems). Monitoring plans may need

to be adjusted to reflect the species of fish available in a water body (e.g., small streams), temporal issues (e.g., spring flood/safety, low flow), and the types of appropriate sampling gear.

The monitoring strategy in EPA's Fish Advisory Guidance Volume 1 (USEPA 2000) discusses two tiers of studies used to identify locations where fish consumption advisories may be needed. Information from these studies may be utilized to develop selenium specific monitoring programs for the assessment of the aquatic life criterion. Tier 1 studies are screening studies that evaluate a large number of sites for chemical contamination with few samples per site. These would be most useful for water bodies, regions, or states where there are no known or expected selenium problems. Screening studies can help states and authorized tribes identify those sites where selenium concentrations are elevated relative to other water bodies. Information from screening studies can be used to prioritize water bodies for future monitoring, thus enabling resources to be used more efficiently. For example, water bodies with fish having low selenium concentrations may be monitored less frequently in the future, while water bodies with fish having selenium concentrations at or near the tissue criterion elements may be prioritized for more frequent or more intensive monitoring. In addition, data collected during these screening studies can be used to inform assessment determinations for the waters where the samples were collected.

Tier 2 studies are intensive studies of areas identified as potential problems in screening studies. The purpose of a Tier 2 study is to determine the magnitude of chemical contamination in sensitive fish species, and to assess the geographic extent of the contamination. If a Tier 2 study is being conducted for selenium, fish species from a sensitive genus with high bioaccumulation potential should be sampled either in addition to or in place of sensitive species. Agencies will typically use Tier 2 studies to determine the overall magnitude and variability of a specific contaminant that was found at elevated levels during a Tier 1 study. In many areas, selenium sources have been well characterized; in these areas a screening study may not provide any additional information that would change the course of the investigation. At these sites, it may be most logical to move directly to an intensive study designed to capture the magnitude and geographical extent of the selenium contamination in fish tissue. These studies may be helpful as a basis for developing a site-specific water column criterion element, if necessary.

3.2 Existing Resources and Information

3.2.1 Available Expertise

Within each state or authorized tribe, the agency that develops the WQS and the agency that typically conducts fish sampling may not be the same. When designing sampling plans to assess the selenium aquatic life criterion, agencies with experience in the development and execution of fish sampling programs can be consulted to aid in designing an effective fish tissue monitoring plan. State agencies should also determine whether there is any overlap in current sampling efforts. Various state (e.g., Department of Natural Resources) and federal agencies (e.g., National Oceanic and Atmospheric Administration - National Marine Fisheries Service, United States Fish and Wildlife Service [USFWS], United States Geological Survey [USGS], EPA)

have expertise in fish sampling, biology, and ecology, and may be able to provide assistance with designing a sampling plan.

All states and most authorized tribes and interstate commissions have established biological assessment programs, and most have fisheries biologists and managers. This should provide the capacity to establish or modify existing fish tissue monitoring programs to facilitate implementation of the fish tissue-based criterion elements in the national CWA section 304(a) recommended selenium criterion. In addition to individual state and tribal agencies and local expertise, federal (e.g., USFWS) and state resource agency collaborations could be used as necessary to fill in data gaps and provide supporting data. By using all available resources for information and expertise, monitoring agencies should be able to:

- Identify potential sites/locations, water bodies, and watersheds for selenium sampling beyond the coverage of current monitoring programs
- Design an appropriate monitoring strategy (including selection of tissue type and sample type (i.e., individual or composite samples))
- Select target species
- Identify pre-spawning periods
- Procure analytical support

The case study below presents Minnesota's Fish Contaminant Monitoring Program, which is implemented through a collaborative partnership of four state agencies to maximize available expertise. Based on the available information, a state or authorized tribe with a similar collaborative program could take advantage of their joint resources to devise the most efficient approach for adding selenium to their current monitoring strategy. They could also use their extensive database to determine where to conduct more intensive studies.

CASE STUDY: Minnesota's Fish Contaminant Monitoring Program

Minnesota's Fish Contaminant Monitoring Program is implemented through a partnership of Minnesota Departments of Natural Resources, Health, and Agriculture and the Minnesota Pollution Control Agency (MPCA). The data are used to issue fish consumption advisories, identify impaired waters, research mercury cycling, and document long term trends for PCBs and mercury.

Highlights (MPCA 2008, P. McCann, personal communication, May 7, 2018):

- Approximately 130 lakes and river sites are sampled annually.
- The Fish Contaminant Monitoring Program database contains over 52,000 data records.
- As of 2016, the program has sampled 1,410 lakes of the estimated 5,500 fishing lakes in the state.

This program is a robust example of how interagency cooperation can maximize available expertise, resources, and cost effectiveness.

<https://www.pca.state.mn.us/sites/default/files/p-p2s4-05.pdf>

3.2.2 Existing Guidance

In 2000, EPA published guidance related to monitoring of contaminants in fish called *Guidance for Assessing Chemical Contaminant Data for Use in Fish Advisories, Vol 1: Fish Sampling and Analysis* (USEPA 2000). EPA's Fish Advisory Guidance Volume 1 discusses study design considerations and the major protocols that must be specified for fish collection, such as site selection, analyte screening values, sampling times, sampling type, and QA/QC.

EPA's Fish Advisory Guidance Volume 1 also contains information on monitoring strategies, field procedures, sample number, sample collection, and sample handling which can be helpful to state and tribal programs monitoring for implementation of the fish tissue components of EPA's aquatic life selenium criterion recommendations. EPA's Fish Advisory Guidance Volume 1 provides useful information on the collection of whole-body and muscle tissue samples. Specifically, section 7.2.2 of EPA's Fish Advisory Guidance Volume 1 (USEPA 2000) includes detailed directions for preparing muscle and whole-body samples. The limitation to this guidance is that it was developed specifically for assessing human health risks associated with consumption of fish and shellfish. As a result, there are aspects of implementing the aquatic life selenium fish tissue-based criterion that are not specifically addressed by EPA's Fish Advisory Guidance Volume 1 (e.g., fish egg-ovary sampling).

Data collected through monitoring for criterion assessment will be used differently than data collected for fish advisories. In the waterbody criterion assessment context, once a criterion

element is exceeded, the water body is considered impaired (and placed on the state's or authorized tribe's CWA section 303(d) list), and a likely next step would be additional monitoring for a TMDL (to identify sources) or site-specific criterion. Data from intensive fish advisory studies, like Tier 2 studies described in EPA's Fish Advisory Guidance Volume 1, might help to support TMDL development for those waters where one or more of the fish tissue criterion elements are exceeded.

In addition to EPA's Fish Advisory Guidance Volume 1, EPA and other stakeholders have produced numerous documents on bioassessment techniques. Specific sections of these documents contain information that may be helpful for developing guidelines for sampling fish for selenium fish tissue analysis, particularly for species like cyprinids which are not typically targeted by state monitoring programs. For example, *Rapid Bioassessment Protocols for Use in Streams and Wadeable Rivers: Periphyton, Benthic Macroinvertebrates, and Fish - Second Edition Chapter 3* (Barbour et al. 1999) provides guidance and information on the elements of biomonitoring, including seasonality and methods for fish collections. A selection of recommended documents for additional guidance is presented in Table 4.

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Table 4: Recommended Documents for Additional Guidance

Title	Author	Link
<i>Guidance for Assessing Chemical Contaminant Data for Use in Fish Advisories, Vol 1: Fish Sampling and Analysis</i> ¹	USEPA 2000	https://www.epa.gov/sites/production/files/2015-06/documents/volume1.pdf
<i>Rapid Bioassessment Protocols for Use in Streams and Wadeable Rivers: Periphyton, Benthic Macroinvertebrates, and Fish - Second Edition</i>	Barbour et al. 1999	https://www.epa.gov/sites/production/files/2019-02/documents/rapid-bioassessment-streams-rivers-1999.pdf
<i>Field Sampling Plan for the National Study of Chemical Residues in Lake Fish Tissue</i> ¹	USEPA 2002a	http://www.epa.gov/sites/production/files/2015-07/documents/fish-study-fieldplan.pdf
<i>The National Study of Chemical Residues in Lake Fish Tissue (Final Report)</i> ¹	USEPA 2009	https://nepis.epa.gov/Exe/ZyPDF.cgi/P1005P2Z.PDF?Dockey=P1005P2Z.PDF
<i>Concepts and Approaches for the Bioassessment of Non-Wadeable Streams and Rivers</i>	Flotemersch et al. 2006	https://nepis.epa.gov/Exe/ZyPDF.cgi/600006KV.PDF?Dockey=600006KV.PDF
<i>Guidance on Choosing a Sampling Design for Environmental Data Collection</i>	USEPA 2002b	http://www.epa.gov/sites/production/files/2015-06/documents/g5s-final.pdf
<i>Spatially Balanced Survey Designs for Natural Resources. Design and Analysis of Long-Term Ecological Monitoring Studies</i>	Olsen et al. 2012	https://www.cambridge.org/core/books/design-and-analysis-of-longterm-ecological-monitoring-studies/spatially-balanced-survey-designs-for-natural-resources/F06C6F53022E46D694D1233782D5F274
<i>Spatially Balanced Sampling of Natural Resources</i>	Stevens and Olsen 2004	https://cfpub.epa.gov/ncer_abstracts/index.cfm/fuseaction/display.files/fileID/13339
<i>Application of Global Grids in Environmental Sampling</i>	Olsen et al. 1998	https://archive.epa.gov/nheerl/arm/web/html/abolse_n98.html
<i>National Rivers and Streams Assessment 2018/19: Field Operations Manual – Non-Wadeable</i> ¹	USEPA 2019a	https://www.epa.gov/sites/production/files/2019-05/documents/nrsa_1819_fom_nonwadeable_version_1.2.pdf
<i>National Rivers and Streams Assessment 2018/19: Field Operations Manual – Wadeable</i> ¹	USEPA 2019b	https://www.epa.gov/sites/production/files/2019-05/documents/nrsa_1819_fom_wadeable_version_1.2_0.pdf
<i>National Coastal Condition Assessment 2015 Field Operations Manual</i> ¹	USEPA 2015	https://www.epa.gov/sites/production/files/2016-03/documents/national_coastal_condition_assessment_2015_field_operation_manual_version_1.0_1.pdf
<i>Biomonitoring of Environmental Status and Trends (BEST) Program: Field Procedures for Assessing the Exposure of Fish to Environmental Contaminants</i>	Schmitt et al. 1999	https://www.cerc.usgs.gov/pubs/center/pdfDocs/91116.pdf

¹Fishing sampling in these references are designed specifically for assessing risk to human health through fish consumption.

3.2.3 Using Existing Data to Enhance Selenium Monitoring

EPA recommends considering and utilizing all available data, as appropriate, to inform and enhance selenium monitoring. According to EPA's 2010 Fish Advisory Survey Report, 28 states identify selenium as a contaminant in their fish monitoring program (USEPA 2011). Several states have conducted extensive statewide assessments and could have existing selenium data. Other organizations may also have selenium data available. For example, the Ohio River Valley Water Sanitation Commission collects fish tissue samples for selenium analysis as part of their Fish Consumption Advisory Program, and has data available online (<http://www.orsanco.org/fish-tissue>). National scale data sources for selenium in fish tissue samples include EPA's 2008-2009 National Rivers and Streams Assessment (available at <https://www.epa.gov/national-aquatic-resource-surveys/national-rivers-and-streams-assessment-2008-2009-results>) and the National Listing of Fish Advisories Fish Tissue Search database (available at <https://fishadvisoryonline.epa.gov/FishTissue.aspx>). EPA also has concentration data available from one hundred paired mercury and selenium fish fillet samples collected in 2007 (available at <http://www.epa.gov/sites/production/files/2015-07/mercury-finaldata2012.xlsx>). Sample sites for this 2007 study were randomly selected from U.S. locations where mercury advisories for fish consumption were in place at the time of sampling. Available data can be used to conserve limited resources by providing baseline information which can inform future collections by indicating which areas may and may not need additional monitoring.

4.0 Sample Analysis

4.1 Analytical Chemistry

Fish tissue sampling to support implementation of the national CWA section 304(a) recommended selenium criterion will need to address many of the same analytical concerns as those associated with other tissue monitoring programs. Various researchers have shown that analytical results on the same population of fish can differ between studies and even within studies. These uncertainties inherent in any sampling program can be minimized through a rigorous study design, clear data quality objectives, meticulous QA/QC protocols, and careful execution of the monitoring program in the field. Standardized methods should be followed in the field to ensure the appropriate samples (have been handled, preserved, and shipped according to protocol) are analyzed in the laboratory (Beatty and Russo 2014). Consistent analytical methods and procedures should be used across implementation programs that are utilizing fish tissue data. Analytical methods should be selected that are sufficiently sensitive to address study objectives (e.g., methods with detection limits below the selenium fish tissue criterion elements after allowing for conversion to dry weight concentrations) and minimize the number of values that are below the MDL. Results should be reported to the appropriate significant figures for the precision of the analytical method.

Laboratories should be selected based on relevant laboratory accreditations, strong QA/QC protocols, and experience with using analytical methods for selenium and the fish tissue matrix. Samples should be prepared in accordance with the tissue type. Section 7.2.2 of EPA's Fish

Advisory Guidance Volume 1 (USEPA 2000) includes detailed directions for preparing muscle and whole-body samples. Appendix A of this document includes directions for preparing egg and ovary samples. EPA does not have approved 40 CFR Part 136 methods for measuring selenium in fish tissue at this time. However, states and authorized tribes are not required to use EPA-approved methods for monitoring and assessment of criteria attainment or criteria development. Additionally, in the case of pollutants or pollutant parameters for which there are not approved methods under 40 CFR Part 136 or methods are not otherwise required under 40 CFR Chapter I, subchapter N or O, monitoring for activities related to permit applications, permit limits, or permit compliance reports shall be conducted according to a test procedure specified in the permit for such pollutants or pollutant parameters.² In the assessment of criteria attainment and establishment of lists of waters not attaining criteria, however, states are required to assemble and evaluate all existing and readily-available water quality-related data and information (40 CFR 130.7(b)(5)). If a state or authorized tribe has additional statutes concerning data acceptability or laboratory accreditation programs, then the fish tissue analytical methods implemented by the state or authorized tribe should be in compliance with these statutes.

Before selecting a method for analysis and a laboratory to conduct those analyses, states and authorized tribes should discuss with laboratories their MDLs for detecting selenium in fish tissue using a particular analytical method. States and authorized tribes should confirm whether those MDLs are for wet weight or dry weight and assure that they are sensitive enough for the assessment of the selenium criterion or for site-specific study purposes. Table 5 presents several analytical procedures for measuring selenium in solids and biota with MDLs that are sufficiently sensitive for comparison to the tissue criterion elements. Exact MDLs and quantitation limits (QL) for these methods are not provided, as those values are laboratory and project specific, however, all the methods listed below should be sensitive down to a selenium concentration of at least 1.5 mg/kg dw. States and authorized tribes should decide which value they want the laboratory to use for reporting, whether they would like it to be equal to the MDL, QL, or some alternative value that they have confidence in using for regulatory decisions. See section 4.2 of this document for discussion about evaluating data that is below the MDL or falls in between the MDL and the QL. Furthermore, some of the analytical methods and procedures identified in Table 5 do not include specific QC requirements and acceptance limits. Therefore, jurisdictions will need to work closely with the laboratory to establish appropriate requirements so that the data meet the monitoring objectives.

² The standard conditions of an NPDES permit (40 CFR 122.41 and 122.4(i)) require, when available, permittees use test procedures specified in 40 CFR Part 136.

Table 5: List of Test Procedures for Total Selenium in Solids and Biota

Method	Digestion / Preparation in reference method?	Links to Methods
EPA Method 6020B ¹ – Inductively Coupled Plasma - Mass Spectrometry (ICP - MS)	No – Recommended: 3052 (total), or 3050B (total recoverable)	https://www.epa.gov/esam/epa-method-6020b-sw-846-inductively-coupled-plasma-mass-spectrometry https://www.epa.gov/sites/production/files/2015-12/documents/3052.pdf https://www.epa.gov/sites/production/files/2015-06/documents/epa-3050b.pdf
EPA Method 7742 ¹ – Atomic Absorption, Borohydride Reduction	No- Recommended: 3052 (total), or 3050B (total recoverable)	https://www.epa.gov/sites/production/files/2015-12/documents/7742.pdf (See links for digestion methods above)
USGS I-9020-05 – Determination of Elements in Natural-water, Biota, Sediment, and Soil Samples using Collision /Reaction Cell ICP - MS	No – References 3052 (total) Recommended: 3052 (total), or 3050B (total recoverable)	https://pubs.usgs.gov/tm/2006/tm5b1/PDF/TM5-B1.pdf (See links for digestion methods above)
NOAA 140.1 - Graphite Furnace-Atomic Absorption for the Analysis of Trace Metals in Marine Animal Tissues	Yes – Teflon Bomb	https://www.nemi.gov/methods/method_summary/7185/
EPA Method 200.8, Rev 5.4 ^{1,2} – Determinations of Trace Elements in Waters by ICP- MS (1994a)	Yes - Section 11.3 May also use: 3052 (total), or 3050B (total recoverable)	https://www.epa.gov/sites/production/files/2015-08/documents/method_200-8_rev_5-4_1994.pdf

¹ These EPA methods are not included in 40 CFR Part 136 for fish tissue analysis. EPA does not currently have any 40 CFR Part 136 methods for analyzing parameters in fish tissue.

² Tissue samples must be digested before using this method.

Tissue samples should be homogenized and digested prior to analysis using strong acid and either a closed-vessel microwave digestion or an open-vessel heated digestion procedure. If samples are to be dried before homogenization and digestion, freeze drying is a good drying technique to use to minimize selenium losses from the sample. However, undried tissues may be homogenized and digested, and a dry weight conversion can be determined using a separate aliquot of the homogenized tissue. The suitability of a given technique should be discussed with the individual laboratory given its capabilities and preference. The laboratory and the agency submitting the samples should mutually decide on a technique that meets the purposes of the monitoring. Care should be taken to use a process that will minimize the loss of volatile

selenium. Reference materials, analytical duplicates, and matrix spike samples are recommended to determine the applicability of the selected digestion and analysis procedures.

The North American Metals Council-Selenium Work Group (NAMC-SWG) has published comprehensive discussions of analytical concerns relevant to selenium; Ohlendorf et al. (2008) and Ohlendorf et al. (2011). An additional NAMC-SWG document, Ralston et al. (2008), presents guidance on analytical methods for selenium. Inductively coupled plasma mass spectrometry is the typical method used for analyzing selenium in tissue and other matrices; however, this method is sensitive to interferences. When using this method, these potential interferences should be addressed. Alternative methods for analyzing selenium are discussed in D'Ulivo (1997), Ohlendorf et al. (2008), and Ralston et al. (2008). States and authorized tribes should choose an analytical method that is sufficiently sensitive to implement its water quality standard for selenium or meet the site-specific study objectives.

States and authorized tribes can also adapt methods for analyzing selenium in water to measure selenium in fish tissue, as long as the fish tissue samples are appropriately digested. In particular, EPA Method 200.8, Rev 5.4, *Determination of Trace Elements in Waters and Wastes by Inductively Coupled Plasma-Mass Spectrometry* (1994a) can easily be adapted to tissue analyses by the addition of an appropriate digestion procedure. Additional information regarding analytical methods for water samples can be found in Appendix L of the *Aquatic Life Ambient Water Quality Criterion for Selenium – Freshwater 2016* (USEPA 2021a). Complete descriptions of analytical methods appropriate for analyzing selenium in different media can be found in the National Environmental Methods Index at <http://www.nemi.gov>.

4.2 Data Analysis

EPA's regulations require that states assemble and evaluate all existing and readily available water quality-related data and information for use in assessing water quality and developing their CWA section 303(d) lists. How a state or authorized tribe plans to evaluate data should be reflected in their assessment methodology. An assessment methodology constitutes the decision process that a state or authorized tribe employs to determine the use attainment status of waters in their jurisdiction. States and authorized tribes should evaluate if there is a need to update their existing assessment methodology to account for how they plan to analyze their selenium tissue data. The methodology should describe how selenium data and information are evaluated and used to make water quality attainment determinations, including data quality, quantity, and representativeness considerations. The methodology should also include any statistically based procedures used during the assessment.

When sufficient data are available, jurisdictions may consider use of the recommended statistical approaches in EPA's Fish Advisory Guidance Volume 1 (USEPA 2000). When comparing contaminants to the criterion, the guidance recommends use of a one-sample t-test to statistically compare the mean of all fish tissue data for a single species and single tissue type to the applicable criterion. If there are concerns with meeting critical assumptions underlying the t-test or another parametric test (e.g., normality), then a nonparametric test could be used. A

nonparametric t-test alternative, such as a one-sample Wilcoxon Signed Rank test, can be used to evaluate if the median selenium tissue concentration is greater than the applicable criterion.

Intensive studies may include the collection of fish tissue data from several locations within a region of interest or for multiple time periods (e.g., seasons or years) from a single location, or a combination of both. Data from intensive studies such as these may be used to perform spatial or temporal analyses to provide information on selenium variability in a target species population. EPA's Fish Advisory Guidance Volume 1 provides recommended statistical approaches for comparing contaminants measured at different locations or over time (See Appendix N of USEPA 2000). EPA recommends that states and authorized tribes consult a statistician to determine the specific statistical tests needed for a particular dataset and choose a method best suited to how they express their WQS. Consulting a statistician at the time of the study design may also be useful for assuring that the appropriate data are collected to answer the desired question.

When making assessment decisions, states and authorized tribes should consider how they will address potential data quality concerns, such as the use of analytical results that are below the method detection limit (MDL) and/or analytical results that are in between the MDL and the quantitation limit (QL). These results can be largely avoided with proper quality assurance project planning. The collection of sufficient tissue mass and use of a sufficiently sensitive analytical method will provide results with a minimal number of values below the MDL and between the MDL and QL. However, if a state or authorized tribe is using a dataset that includes values below the MDL or in between the MDL and the QL, it should decide how it will evaluate these values.

There are various conventions to deal with these measurements and the state or authorized tribe has the flexibility to determine which is appropriate for their given situation. EPA notes that identifying and developing approaches to statistically analyze datasets containing non-quantified chemical concentration values (i.e., "censored data") is an active area of research and no one method can be universally recommended (for more information see: Helsel 2005, Pleil 2016, and, Singh and Nocerino 2002). EPA's Fish Advisory Guidance Volume 1 (USEPA 2000) recommends using one-half of the MDL for values below the MDL in calculating mean values (section 9.1.2). The guidance also recommends that measurements that fall between the MDL and the QL be assigned a value of the MDL plus one-half the difference between the MDL and the QL. EPA notes, however, that these conventions provide a biased estimate of the average concentration (Gilbert 1987) and, where the computed average is close to the criterion, might suggest an impairment when one does not exist or, conversely, suggest no impairment when one does exist. As an alternative to this convention, some states, authorized tribes, and laboratories may choose to apply what is called a "J" flag to any results reported at or above the MDL, but below the QL. The "J" flag would indicate that the chemical is present, but the reported value is an estimate of the true concentration since it was detected below the QL. Some states and authorized tribes may choose to use these "J" flagged values for data analysis. EPA used this convention for the National Study of Chemical Residues in Lake Fish Tissue, including all the "J" flagged data in analyses of the fish tissue data (USEPA 2009).

States or authorized tribes can also calculate the average of a dataset that includes values below the MDL using other statistical methods (e.g., sample median and trimmed means) (Gilbert 1987). Singh and Nocerino (2002) have published a review of several methods for data reporting and analyzed the potential bias each can introduce into the calculation of the mean.

One approach that a state or authorized tribe could take to ascertain the consequence of what value is used to quantify samples below the MDL is to conduct a sensitivity analysis. In a sensitivity analysis, the state or authorized tribe would compute the mean concentration by first using the value of the MDL to quantify samples below the MDL and then using a zero value for samples below the MDL. For example, if the MDL is 1.5 mg/kg dw, first the mean would be calculated with all values below the MDL being assigned the value of 1.5 mg/kg dw. Then the mean would be recalculated with the value of 0.0 mg/kg dw being assigned to all values below the MDL. If both calculated means are above or below the criterion, it is clear that the choice of how to quantify samples below the MDL does not affect the decision. However, if one calculated mean is below the criterion and the other is above, it is clear that the choice of how to quantify samples below the MDL does affect the decision, and a state or authorized tribe may want to use a more sophisticated approach such as the ones presented in *Robust Estimation of Mean and Variance Using Environmental Datasets with Below Detection Limit Observations* (Singh and Nocerino 2002).

All data handling conventions have advantages and disadvantages. A state or authorized tribe should understand the consequences of which convention it uses, especially if the choice makes a difference as to whether a water body is considered impaired or not. Furthermore, a state or authorized tribe should be clear about which approach it used. The selected approach must be consistent with the state's EPA-approved WQS and should generally adhere to any published assessment method associated with them. For further discussion on handling values below the MDL, see USEPA 2000 (section 9.1) and USEPA 2010 (section 4.3.1). In general, states or authorized tribes should not have issues with measurements of selenium in fish tissue being below the MDL when a sufficient mass of tissue is collected as the method sensitivities are low enough and all fish should have selenium concentrations higher than those MDLs. Similarly, with sufficient sample mass and appropriate analytical methods, it is unlikely that many states and authorized tribes will have to deal with selenium measurements between the MDL and QL for fish tissue.

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Appendix A: Egg and Ovary Sample Preparation

Scope

This guidance is for egg and ovary collection from freshwater fish. The egg extraction method is excerpted and adapted from a more comprehensive guidance, *Standard operating procedure for evaluating selenium-induced deformities in early life stages of freshwater fish* (Janz and Muscatello 2008), that includes gamete collection, embryo incubations and evaluation of selenium-induced deformities in freshwater fish. The ovary dissection method was compiled from peer-reviewed literature.

1. Field collection and handling of adult fish

“Spawning adults can be collected in the field using a wide variety of techniques, including fish traps (e.g., hoop or trap nets), electrofishing or angling in areas close to spawning areas. Gillnets are also effective in capturing fish during spawning migrations, but it is essential to monitor these nets constantly to remove fish immediately after capture. If possible, the use of passive capture methods (e.g., hoop or trap nets) is recommended since this is the least stressful capture technique of those listed above. Trap nets are usually set up in creeks, streams or narrows in lakes, although successful fish capture can also occur when these nets are set perpendicular to shore in lentic habitats. Trap or hoop nets can be purchased from fisheries suppliers, or even constructed in creeks and streams using chicken wire, baling wire and reinforcing bar.” (Janz and Muscatello 2008)

Fish should be held in livewells until adult female fish are selected for egg collection.

2. Egg collection procedures

Fish should be carefully observed for signs of physical damage, mortality or other sources of stress. Since any handling of the fish will remove the protective body layer of slime, fish should be handled as little as possible using dip nets and soft material gloves. Adult fish for egg collection should be randomly selected from livewells.

“Eggs should not be in contact with water; thus, it is imperative to dry the area surrounding the urogenital opening with paper towels. All the material used for egg collection should be carefully cleaned and dried. Precautions to avoid fecal, blood or urine contamination should be taken. [Eggs] must be kept covered to avoid direct sun exposure . [Egg collection] should proceed after recording weight and length [of the gravid female]. Gentle pressure from behind the pectoral fins towards the anus is applied to express the eggs. This process needs to be repeated several times. Check that eggs are released ‘clean’ (e.g., without feces) before starting collection to avoid contamination of the entire egg batch. Eggs are

individually collected into pre-cleaned stainless steel bowls and kept covered in a cool place until use. Collected eggs should be closely inspected and eggs with adhered feces, urine or blood discarded by using a clean plastic pipette.” (Janz and Muscatello 2008)

Eggs are then weighed to the nearest gram using a top-loading digital scale, frozen for storage, and shipped for laboratory analysis when appropriate. An individual or composite homogenate tissue sample of 20 grams ww should be collected for analysis of selenium.

3. Ovary dissection procedures

Fish designated for ovary collection should be humanely euthanized, and necropsy procedures should commence immediately following euthanasia (Wolf et al. 2004). The fish should be placed in right lateral recumbency on a piece of aluminum foil. The left body wall should be removed by using fine dissecting instruments (Wolf et al. 2004). To identify female specimens for ovary collection, sex is determined by macroscopic inspection when the body cavity is opened. The ovaries are paired organs suspended from the dorsal wall, with color ranging from clear to white to yellow-orange. A yellow-orange color is indicative of a ripening or ripe adult specimen. Further, increased blood flow during the reproductive season causes the ovaries to become highly vascularized and appear reddish. In cross-section, the ovaries are round to elliptical and contain a central cavity (lumen). In young fish, the texture of the ovaries varies from smooth to slightly granular. The ovarian texture in a ripe fish will be highly granular (Fisheries Information Network 2006). If inspection of the ovaries reveals that the specimen is immature or developing, it is not recommended that the eggs/ovarian tissue be used for tissue monitoring for selenium.

After confirmation that the specimen is a ripe female, the ovaries should be excised by severing the oviducts and mesenteric attachments. All gonads are dissected in a caudal to cranial direction (Wolf et al. 2004). Ovaries are then weighed to the nearest gram using a top-loading digital scale, frozen for storage, and shipped for laboratory analysis when appropriate (Orr et al. 2012). An individual or composite homogenate sample of 20 grams ww of tissue should be collected for analysis of selenium.

4. Storing fish eggs and ovaries

“Eggs and ovaries should be kept frozen until analysis. After collection, samples should be kept in a container with ice or freezer packs until transfer to a freezer (-20°C) for storage” (Janz and Muscatello 2008). It is recommended to transfer the samples collected from each individual female into sealed resealable plastic storage bags to “prevent water (from ice melting) entering the sample” (Janz and Muscatello 2008). Recommendations for the storage, preservation and holding time for egg and ovary samples are equivalent to other tissue samples. Samples should be frozen at -20°C in plastic, borosilicate glass, quartz or PTFE bottles. The recommended maximum holding time is six months but can be up to two years for most trace metals, including selenium (USEPA 2000).

5. Laboratory preparation of egg and tissue samples for metal analysis

“Egg and tissue samples should be thawed, and wet weight recorded for each individual sample. To prevent cross contamination between samples, a plastic foil (e.g., parafilm®) should be placed on the scale and replaced after each weighing. Samples are oven dried at 60°C until constant weight is recorded. It is required to record the moisture content for each individual sample in order to express analytical data on a dry weight basis. Trace element (e.g., selenium) analysis is routinely performed using hydride generation atomic absorption spectrophotometry (HG-AAS) or inductively coupled plasma-mass spectrometry (ICP-MS) and reported on a dry-weight basis.” (Janz and Muscatello 2008)

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Appendix B: Spawning Seasons for Example Fish Assemblages from Select U.S. Watersheds

This appendix contains spawning season calendars for fish assemblages from selected watersheds in six different areas of the United States. The calendars are intended to provide examples of spawning periods for fish species commonly collected in those areas. EPA recommends that monitoring agencies use all available locally relevant resources to determine the appropriate time to collect fish for the purpose of implementing the selenium criterion, including contacting their local natural resources or fish and game agency.

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Spawning Seasons for Example Fish Assemblages in the Merrimack River, MA and NH Watershed

Family	Scientific Name	Common Name	Spawning Season
Atherinopsidae	<i>Menidia menidia</i>	Atlantic Silverside	April through August
Catostomidae	<i>Catostomus commersonii</i>	White Sucker	March through July
Centrarchidae	<i>Ambloplites rupestris</i>	Rock Bass	April through July
Centrarchidae	<i>Enneacanthus obesus</i>	Banded Sunfish	April through July
Centrarchidae	<i>Lepomis auritus</i>	Redbreast Sunfish	April through July
Centrarchidae	<i>Lepomis gibbosus</i>	Pumpkinseed	June through August
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	May through August
Centrarchidae	<i>Micropterus dolomieu</i>	Smallmouth Bass	April through June
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	April through June
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	April through July
Clupeidae	<i>Dorosoma cepedianum</i>	Gizzard Shad	March through August
Cyprinidae	<i>Carassius auratus</i>	Goldfish	March through August
Cyprinidae	<i>Cyprinus carpio</i>	Common Carp	April through August
Cyprinidae	<i>Luxilus cornutus</i>	Common Shiner	May through July
Cyprinidae	<i>Notemigonus crysoleucas</i>	Golden Shiner	May through July
Cyprinidae	<i>Notropis atherinoides</i>	Emerald Shiner	May through June
Cyprinidae	<i>Notropis bifrenatus</i>	Bridle Shiner	May through August
Cyprinidae	<i>Notropis hudsonius</i>	Spottail Shiner	May through September
Cyprinidae	<i>Rhinichthys atratulus</i>	Blacknose Dace	April through July
Cyprinidae	<i>Rhinichthys cataractae</i>	Longnose Dace	April through June
Cyprinidae	<i>Semotilus atromaculatus</i>	Creek Chub	March through June
Cyprinidae	<i>Semotilus corporalis</i>	Fallfish	April through May
Esocidae	<i>Esox lucius</i>	Northern Pike	March through May
Esocidae	<i>Esox niger</i>	Chain Pickerel	March through May
Fundulidae	<i>Fundulus diaphanus</i>	Banded Killifish	April through August
Fundulidae	<i>Fundulus heteroclitus</i>	Mummichog	June through July
Gadidae	<i>Lota lota</i>	Burbot	January through April
Gasterosteidae	<i>Apeltes quadracus</i>	Fourspine Stickleback	April through May
Gasterosteidae	<i>Gasterosteus aculeatus</i>	Threespine Stickleback	March through June
Gasterosteidae	<i>Pungitius pungitius</i>	Ninespine Stickleback	April through August
Ictaluridae	<i>Ameiurus catus</i>	White Catfish	May through July
Ictaluridae	<i>Ameiurus natalis</i>	Yellow Bullhead	May through June
Ictaluridae	<i>Ameiurus nebulosus</i>	Brown Bullhead	April through June
Ictaluridae	<i>Ictalurus punctatus</i>	Channel Catfish	April through September
Ictaluridae	<i>Noturus gyrinus</i>	Tadpole Madtom	May through July
Ictaluridae	<i>Noturus insignis</i>	Margined Madtom	June through July
Moronidae	<i>Morone americana</i>	White Perch	May through June
Percidae	<i>Etheostoma fusiforme</i>	Swamp Darter	April through May
Percidae	<i>Etheostoma olmstedii</i>	Tessellated Darter	March through May
Percidae	<i>Perca flavescens</i>	Yellow Perch	May through July
Percidae	<i>Sander vitreus</i>	Walleye	April through May
Salmonidae	<i>Oncorhynchus mykiss</i>	Rainbow Trout	April through June
Salmonidae	<i>Salmo trutta</i>	Brown Trout	October through February
Salmonidae	<i>Salvelinus fontinalis</i>	Brook Trout	September through November

(Scarola 1973, Page and Burr 1991)

Spawning Seasons for Example Fish Assemblages in the Delaware River, DE Watershed

Family	Scientific Name	Common Name	Spawning Season
Aphredoderidae	<i>Aphredoderus sayanus</i>	Pirate Perch	April through May
Atherinopsidae	<i>Membras martinica</i>	Rough Silverside	May through August
Atherinopsidae	<i>Menidia peninsulae</i>	Tidewater Silverside	May through August
Atherinopsidae	<i>Menidia menidia</i>	Atlantic Silverside	April through August
Catostomidae	<i>Catostomus commersonii</i>	White Sucker	March through May
Catostomidae	<i>Erimyzon oblongus</i>	Creek Chubsucker	March through May
Centrarchidae	<i>Acantharchus pomotis</i>	Mud Sunfish	May through June
Centrarchidae	<i>Enneacanthus chaetodon</i>	Blackbanded Sunfish	May through July
Centrarchidae	<i>Enneacanthus gloriosus</i>	Bluespotted Sunfish	May through September
Centrarchidae	<i>Enneacanthus obesus</i>	Banded Sunfish	June through September
Centrarchidae	<i>Lepomis auritus</i>	Redbreast Sunfish	May through June
Centrarchidae	<i>Lepomis gibbosus</i>	Pumpkinseed	May through August
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	May through August
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	April through June
Centrarchidae	<i>Pomoxis annularis</i>	White Crappie	April through June
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	May through June
Clupeidae	<i>Dorosoma cepedianum</i>	Gizzard Shad	April through June
Cyprinidae	<i>Carassius auratus</i>	Goldfish	June through July
Cyprinidae	<i>Cyprinus carpio</i>	Common Carp	May through July
Cyprinidae	<i>Hybognathus nuchalis</i>	Mississippi Silvery Minnow	April through May
Cyprinidae	<i>Notemigonus crysoleucas</i>	Golden Shiner	April through July
Cyprinidae	<i>Cyprinella analostana</i>	Satinfin Shiner	March through July
Cyprinidae	<i>Notropis bifrenatus</i>	Bridle Shiner	March through August
Cyprinidae	<i>Notropis chalybaeus</i>	Ironcolor Shiner	April through May
Cyprinidae	<i>Notropis hudsonius</i>	Spottail Shiner	April through July
Cyprinidae	<i>Rhinichthys atratulus</i>	Blacknose Dace	May through June
Esocidae	<i>Esox americanus americanus</i>	Redfin Pickerel	February through March
Fundulidae	<i>Fundulus diaphanus</i>	Banded Killifish	April through August
Fundulidae	<i>Fundulus heteroclitus</i>	Mummichog	April through September
Fundulidae	<i>Fundulus majalis</i>	Striped Killifish	April through September
Fundulidae	<i>Lucania parva</i>	Rainwater Killifish	May through July
Ictaluridae	<i>Ameiurus catus</i>	White Catfish	April through July
Ictaluridae	<i>Ameiurus nebulosus</i>	Brown Bullhead	May through July
Ictaluridae	<i>Ictalurus punctatus</i>	Channel Catfish	May through July
Ictaluridae	<i>Noturus gyrinus</i>	Tadpole Madtom	May through July
Moronidae	<i>Morone americana</i>	White Perch	April through June
Percidae	<i>Etheostoma fusiforme</i>	Swamp Darter	April through May
Percidae	<i>Etheostoma olmstedii</i>	Tessellated Darter	March through May
Percidae	<i>Perca flavescens</i>	Yellow Perch	March through April
Poeciliidae	<i>Gambusia affinis</i>	Mosquitofish	May through August
Umbridae	<i>Umbra pygmaea</i>	Eastern Mudminnow	April through June

(Wang and Kernehan 1979, Page and Burr 1991)

Spawning Seasons for Example Fish Assemblages in the Cahaba River, AL Watershed

Family	Scientific Name	Common Name	Spawning Season
Amiidae	<i>Amia calva</i>	Bowfin	March through June
Atherinopsidae	<i>Labidesthes sicculus</i>	Brook Silverside	June through August
Catostomidae	<i>Carpionodes cyprinus</i>	Quillback	March through September
Catostomidae	<i>Carpionodes velifer</i>	Highfin Carpsucker	May through July
Catostomidae	<i>Erimyzon oblongus</i>	Creek Chubsucker	March through May
Catostomidae	<i>Erimyzon sucetta</i>	Lake Chubsucker	March through April
Catostomidae	<i>Erimyzon tenuis</i>	Sharpfin Chubsucker	March through April
Catostomidae	<i>Hypentelium etowanum</i>	Alabama Hog Sucker	April through June
Catostomidae	<i>Ictiobus bubalus</i>	Smallmouth Buffalo	March through April
Catostomidae	<i>Minytrema melanops</i>	Spotted Sucker	April through May
Catostomidae	<i>Moxostoma carinatum</i>	River Redhorse	April
Catostomidae	<i>Moxostoma duquesnii</i>	Black Redhorse	April through May
Catostomidae	<i>Moxostoma erythrurum</i>	Golden Redhorse	April through June
Catostomidae	<i>Moxostoma poecilurum</i>	Blacktail Redhorse	April
Centrarchidae	<i>Ambloplites ariommus</i>	Shadow Bass	May through October
Centrarchidae	<i>Centrarchus macropterus</i>	Flier	February through May
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	March through May
Centrarchidae	<i>Lepomis marginatus</i>	Dollar Sunfish	May through August
Centrarchidae	<i>Lepomis megalotis</i>	Longear Sunfish	May through August
Centrarchidae	<i>Lepomis microlophus</i>	Redear Sunfish	March through May; September through November
Centrarchidae	<i>Lepomis miniatus</i>	Redspotted Sunfish	March through September
Centrarchidae	<i>Micropterus coosae</i>	Redeye Bass	May through July
Centrarchidae	<i>Micropterus dolomieu</i>	Smallmouth Bass	March through May
Centrarchidae	<i>Micropterus punctulatus</i>	Spotted Bass	April through May
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	April through June
Centrarchidae	<i>Pomoxis annularis</i>	White Crappie	April through June
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	February through May
Clupeidae	<i>Dorosoma cepedianum</i>	Gizzard Shad	April through May
Clupeidae	<i>Dorosoma petenense</i>	Threadfin Shad	April through August
Cottidae	<i>Cottus caroliniae</i>	Banded Sculpin	January through March
Cyprinidae	<i>Campostoma oligolepis</i>	Largescale Stoneroller	April through May
Cyprinidae	<i>Cyprinella callistia</i>	Alabama Shiner	March through May
Cyprinidae	<i>Cyprinella trichroistia</i>	Tricolor Shiner	June through July
Cyprinidae	<i>Cyprinella venusta</i>	Blacktail Shiner	March through October
Cyprinidae	<i>Hybognathus nuchalis</i>	Mississippi Silvery Minnow	March through April
Cyprinidae	<i>Hybopsis winchelli</i>	Clear Chub	February through April
Cyprinidae	<i>Luxilus chrysocephalus</i>	Striped Shiner	April through August
Cyprinidae	<i>Lythrurus bellus</i>	Pretty Shiner	April through June
Cyprinidae	<i>Macrhybopsis storeriana</i>	Silver Chub	May through August
Cyprinidae	<i>Notemigonus crysoleucas</i>	Golden Shiner	April through July
Cyprinidae	<i>Notropis ammophilus</i>	Orangefin Shiner	April through October
Cyprinidae	<i>Notropis asperifrons</i>	Burrhead Shiner	April through June
Cyprinidae	<i>Notropis atherinoides</i>	Emerald Shiner	May through July
Cyprinidae	<i>Notropis baileyi</i>	Rough Shiner	May through October
Cyprinidae	<i>Notropis buccatus</i>	Silverjaw Minnow	March through June
Cyprinidae	<i>Notropis candidus</i>	Silverside Shiner	June through September
Cyprinidae	<i>Notropis chrosomus</i>	Rainbow Shiner	May through June

Family	Scientific Name	Common Name	Spawning Season
Cyprinidae	<i>Notropis edwardraneyi</i>	Fluvial Shiner	May through June
Cyprinidae	<i>Notropis stilbius</i>	Silverstripe Shiner	March through August
Cyprinidae	<i>Notropis texanus</i>	Weed Shiner	February through October
Cyprinidae	<i>Notropis uranoscopus</i>	Skygazer Shiner	May through July
Cyprinidae	<i>Notropis volucellus</i>	Mimic Shiner	April through August
Cyprinidae	<i>Opsopoeodus emiliae</i>	Pugnose Minnow	April through September
Cyprinidae	<i>Phenacobius catostomus</i>	Riffle Minnow	April through May
Cyprinidae	<i>Pimephales notatus</i>	Bluntnose Minnow	April through August
Cyprinidae	<i>Pimephales vigilax</i>	Bullhead Minnow	May through August
Cyprinidae	<i>Semotilus atromaculatus</i>	Creek Chub	April through May
Cyprinidae	<i>Semotilus thoreauianus</i>	Dixie Chub	April through May
Elassomatidae	<i>Elassoma zonatum</i>	Banded Pygmy Sunfish	March through April
Esocidae	<i>Esox americanus</i>	Redfin Pickerel	April through May
Esocidae	<i>Esox niger</i>	Chain Pickerel	April through October
Fundulidae	<i>Fundulus olivaceus</i>	Blackspotted Topminnow	March through September
Hiodontidae	<i>Hiodon tergisus</i>	Mooneye	April through May
Ictaluridae	<i>Ameiurus melas</i>	Black Bullhead	May through August
Ictaluridae	<i>Ameiurus natalis</i>	Yellow Bullhead	April through June
Ictaluridae	<i>Ameiurus nebulosus</i>	Brown Bullhead	April through August
Ictaluridae	<i>Ictalurus furcatus</i>	Blue Catfish	April through June
Ictaluridae	<i>Ictalurus punctatus</i>	Channel Catfish	April through July
Ictaluridae	<i>Noturus funebris</i>	Black Madtom	May through June
Ictaluridae	<i>Noturus gyrinus</i>	Tadpole Madtom	May through September
Ictaluridae	<i>Pylodictis olivaris</i>	Flathead Catfish	June through July
Lepisosteidae	<i>Lepisosteus oculatus</i>	Spotted Gar	May through July
Lepisosteidae	<i>Lepisosteus osseus</i>	Longnose Gar	April through August
Moronidae	<i>Morone chrysops</i>	White Bass	February through March
Percidae	<i>Ammocrypta beanii</i>	Naked Sand Darter	March through October
Percidae	<i>Etheostoma meridianum</i>	Southern Sand Darter	April through June
Percidae	<i>Etheostoma chlorosomum</i>	Bluntnose Darter	April
Percidae	<i>Etheostoma jordani</i>	Greenbreast Darter	April through May
Percidae	<i>Etheostoma nigrum</i>	Johnny Darter	March through May
Percidae	<i>Etheostoma parvipinne</i>	Goldstripe Darter	March through April
Percidae	<i>Etheostoma ramseyi</i>	Alabama Darter	March through May
Percidae	<i>Etheostoma rupestre</i>	Rock Darter	March through April
Percidae	<i>Etheostoma stigmaeum</i>	Speckled Darter	March through May
Percidae	<i>Etheostoma swaini</i>	Gulf Darter	March through April
Percidae	<i>Percina kathae</i>	Mobile Logperch	April through June
Percidae	<i>Percina maculata</i>	Blackside Darter	March through June
Percidae	<i>Percina nigrofasciata</i>	Blackbanded Darter	May through June
Percidae	<i>Percina vigil</i>	Saddleback Darter	February through April
Percidae	<i>Sander vitreus</i>	Walleye	March through April
Sciaenidae	<i>Aplodinotus grunniens</i>	Freshwater Drum	May through June

(Boschung and Mayden 2004)

Spawning Seasons for Example Fish Assemblages in the Chicago River, IL Watershed

Family	Scientific Name	Common Name	Spawning Season
Amiidae	<i>Amia calva</i>	Bowfin	March through June
Catostomidae	<i>Catostomus commersonii</i>	White Sucker	April through May
Centrarchidae	<i>Ambloplites rupestris</i>	Rock Bass	May through July
Centrarchidae	<i>Lepomis cyanellus</i>	Green Sunfish	June through August
Centrarchidae	<i>Lepomis humilis</i>	Orangespotted Sunfish	May through July
Centrarchidae	<i>Lepomis gibbosus</i>	Pumpkinseed	May through July
Centrarchidae	<i>Lepomis gulosus</i>	Warmouth	May through August
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	May through August
Centrarchidae	<i>Micropterus dolomieu</i>	Smallmouth Bass	April through June
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	April through June
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	May through July
Clupeidae	<i>Dorosoma cepedianum</i>	Gizzard Shad	May through July
Cyprinidae	<i>Campostoma anomalum</i>	Central Stoneroller	April through July
Cyprinidae	<i>Carassius auratus</i>	Goldfish	May through June
Cyprinidae	<i>Cyprinella spiloptera</i>	Spotfin Shiner	May through August
Cyprinidae	<i>Cyprinus carpio</i>	Common Carp	May through August
Cyprinidae	<i>Hybopsis dorsalis</i>	Bigmouth Shiner	May through June
Cyprinidae	<i>Notemigonus crysoleucas</i>	Golden Shiner	May through August
Cyprinidae	<i>Notropis atherinoides</i>	Emerald Shiner	April through August
Cyprinidae	<i>Notropis hudsonius</i>	Spottail Shiner	June through July
Cyprinidae	<i>Notropis stramineus</i>	Sand Shiner	May through July
Cyprinidae	<i>Pimephales notatus</i>	Bluntnose Minnow	May through August
Cyprinidae	<i>Pimephales promelas</i>	Fathead Minnow	May through August
Cyprinidae	<i>Semotilus atromaculatus</i>	Creek Chub	April through June
Cyprinodontidae	<i>Fundulus notatus</i>	Blackstripe Topminnow	May through August
Esocidae	<i>Esox americanus</i>	Grass Pickerel	May through June; November
Esocidae	<i>Esox lucius</i>	Northern Pike	March through May
Gobiidae	<i>Neogobius melanostomus</i>	Round Goby	April through May
Ictaluridae	<i>Ameiurus melas</i>	Black Bullhead	May through June
Ictaluridae	<i>Ameiurus natalis</i>	Yellow Bullhead	May through June
Ictaluridae	<i>Ictalurus punctatus</i>	Channel Catfish	April through August
Moronidae	<i>Morone americana</i>	White Perch	May through June
Moronidae	<i>Morone chrysops</i>	White Bass	April through June
Moronidae	<i>Morone mississippiensis</i>	Yellow Bass	April through May
Percidae	<i>Etheostoma nigrum</i>	Johnny Darter	April through June
Percidae	<i>Sander vitreus</i>	Walleye	April through May
Percidae	<i>Perca flavescens</i>	Yellow Perch	May through July
Umbridae	<i>Umbra limi</i>	Central Mudminnow	April through May

(Auer 1982, Page and Burr 1991)

Spawning Seasons for Example Fish Assemblages in the Truckee and Carson River, NV Watersheds

Family	Scientific Name	Common Name	Spawning Season
Centrarchidae	<i>Micropterus dolomieu</i>	Smallmouth Bass	April through July
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	April through July
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	May through August
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	May through July
Ictaluridae	<i>Ictaluridae</i>	Catfish species	June through July
Moronidae	<i>Morone saxatilis</i>	Striped Bass*	April through June
Moronidae	<i>Morone chrysops</i>	White Bass	April through June
Percidae	<i>Sander vitreus</i>	Walleye	January through April
Salmonidae	<i>Oncorhynchus mykiss</i>	Rainbow Trout	March through May
Salmonidae	<i>Salmo trutta</i>	Brown Trout	January through March
Salmonidae	<i>Prosopium williamsoni</i>	Mountain Whitefish	October through December

* This population of striped bass is landlocked and cannot migrate out to sea.

(Nevada Division of Environmental Protection 2006)

Spawning Seasons for Example Fish Assemblages in the Rio Grande and Colorado River, TX Watersheds

Family	Scientific Name	Common Name	Spawning Season
Amiidae	<i>Amia calva</i>	Bowfin	March through June
Anguillidae	<i>Anguilla rostrata</i>	American Eel	February through June
Catostomidae	<i>Ictiobus bubalus</i>	Smallmouth Buffalo	March through September
Catostomidae	<i>Ictiobus cyprinellus</i>	Bigmouth Buffalo	April through May
Catostomidae	<i>Ictiobus niger</i>	Black Buffalo	April through May
Centrarchidae	<i>Lepomis macrochirus</i>	Bluegill	April through September
Centrarchidae	<i>Lepomis cyanellus</i>	Green Sunfish	April through August
Centrarchidae	<i>Lepomis megalotis</i>	Longear Sunfish	May through June
Centrarchidae	<i>Lepomis auritus</i>	Redbreast Sunfish	April through October
Centrarchidae	<i>Lepomis microlophus</i>	Redear Sunfish	May through July
Centrarchidae	<i>Lepomis gulosus</i>	Warmouth	March through October
Centrarchidae	<i>Micropterus salmoides</i>	Largemouth Bass	February through May
Centrarchidae	<i>Micropterus dolomieu</i>	Smallmouth Bass	April through May
Centrarchidae	<i>Micropterus punctulatus</i>	Spotted Bass	April through June
Centrarchidae	<i>Micropterus treculii</i>	Guadalupe Bass	March through June
Centrarchidae	<i>Pomoxis nigromaculatus</i>	Black Crappie	March through May
Centrarchidae	<i>Pomoxis annularis</i>	White Crappie	March through May
Cichlidae	<i>Herichthys cyanoguttatus</i>	Rio Grande Cichlid	March through August
Clupeidae	<i>Dorosoma cepedianum</i>	Gizzard Shad	April through June
Clupeidae	<i>Dorosoma petenense</i>	Threadfin Shad	April through September
Cyprinidae	<i>Ctenopharyngodon idella</i>	Grass Carp	April through July
Cyprinidae	<i>Cyprinus carpio</i>	Common Carp	March through June
Cyprinidae	<i>Cyprinella lutrensis</i>	Red Shiner	April through September
Cyprinidae	<i>Cyprinella venusta</i>	Blacktail Shiner	April through September
Cyprinidae	<i>Notropis amabilis</i>	Texas Shiner	February through September
Cyprinidae	<i>Notemigonus crysoleucas</i>	Golden Shiner	April through July
Cyprinidae	<i>Pimephales promelas</i>	Fathead Minnow	May through September
Esocidae	<i>Esox niger</i>	Chain Pickerel	December through February
Ictaluridae	<i>Ictalurus furcatus</i>	Blue Catfish	April through May
Ictaluridae	<i>Ictalurus punctatus</i>	Channel Catfish	April through June
Ictaluridae	<i>Pylodictis olivaris</i>	Flathead Catfish	June through July
Ictaluridae	<i>Ameiurus melas</i>	Black Bullhead	April through June
Ictaluridae	<i>Ameiurus natalis</i>	Yellow Bullhead	May through July
Lepisosteidae	<i>Atractosteus spatula</i>	Alligator Gar	April through May
Lepisosteidae	<i>Lepisosteus oculatus</i>	Spotted Gar	April through June
Lepisosteidae	<i>Lepisosteus osseus</i>	Longnose Gar	April through July
Lepisosteidae	<i>Lepisosteus platostomus</i>	Shortnose Gar	May through July
Moronidae	<i>Morone chrysops</i>	White Bass	March through May
Moronidae	<i>Morone mississippiensis</i>	Yellow Bass	April through June
Moronidae	<i>Morone saxatilis</i>	Striped Bass*	February through April
Percidae	<i>Sander vitreus</i>	Walleye	February through April
Polyodontidae	<i>Polyodon spathula</i>	Paddlefish	February through June

Family	Scientific Name	Common Name	Spawning Season
Salmonidae	<i>Oncorhynchus mykiss</i>	Rainbow Trout	November through February
Sciaenidae	<i>Aplodinotus grunniens</i>	Freshwater Drum	April through June
Sciaenidae	<i>Sciaenops ocellatus</i>	Red Drum	August through October

* This population of striped bass is landlocked and cannot migrate out to sea.
(Hendrickson and Cohen 2015, Texas Parks and Wildlife Department 2016)

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Appendix C: Conversion of Wet to Dry Tissue Weight

Conversion of Wet to Dry Tissue Weight

Selenium data in fish tissues can be reported in either dry weight or wet weight concentrations. It is essential that exposure assessors be aware of this difference so that they may ensure consistency between units when comparing data. If the contaminant concentration is measured in wet weight of fish, then the concentration must be converted to dry weight units in order to be compared to the selenium criterion, which is expressed in dry weight (USEPA 2021). Wet weight may be converted to dry weight, and vice versa, using the following equations:

$$WW = DW \times [1 - (\text{percent moisture}/100)] \text{ (USEPA 2011)}$$

$$DW = WW / [1 - (\text{percent moisture}/100)] \text{ (USEPA 2011)}$$

Measurements reported as wet weight can be converted to equivalent dry weights using available percent moisture data for the relevant species and tissue type. If percent moisture data is unavailable for a fish species, percent moisture data for a similar species (i.e., same genus or, if unavailable, same family) may be used. Table C-1 lists percent moisture of some species by tissue type (USEPA 2021). Percent moisture can vary within species; therefore, the data in Table C-1 should generally be used when dealing with historical data. When using field collected data, measuring % moisture within the field collected sample will provide the most accurate measurement of % moisture, thus giving more accurate conversions between dry weight and wet weight data.

Table C-1. Percent moisture by species and tissue type

Scientific Name	Common Name	Average % Moisture	% Moisture by Tissue			Reference
			Whole -body	Muscle	Egg-ovary	
<i>Cyprinus carpio</i>	Common Carp	75.64 ^a		75.81 ^b		^a USEPA 2014; ^b Chatakondi et al. 1995
<i>Rhinichthys cataractae</i>	Longnose Dace	73.25				USEPA 2014
<i>Rhinichthys atratulus</i>	Blacknose Dace	73.75				USEPA 2014
<i>Semotilus atromaculatus</i>	Creek Chub	76.71				USEPA 2014
<i>Pimephales promelas</i>	Fathead Minnow	76.64 ^a			75.3 ^b	USEPA 2014; ^b USEPA 2015
<i>Pimephales notatus</i>	Bluntnose Minnow	74.8				USEPA 2014
<i>Nocomis micropogon</i>	River Chub	75.2				USEPA 2014
<i>Ictalurus punctatus</i>	Channel Catfish			81.22 ^a 78.43 ^b		^a Pinkney 2003; ^b May et al. 2009

Scientific Name	Common Name	Average % Moisture	% Moisture by Tissue			Reference
			Whole -body	Muscle	Egg-ovary	
<i>Ictalurus melas</i>	Black Bullhead	76.82				USEPA 2014
<i>Pylodictis olivaris</i>	Flathead Catfish			75.97		May et al. 2009
<i>Catostomus commersonii</i>	White Sucker	77.37				USEPA 2014
<i>Coregonus clupeaformis</i>	Lake Whitefish			80		Rieberger 1992
<i>Oncorhynchus kisutch</i>	Coho Salmon			80		Rieberger 1992
<i>Oncorhynchus mykiss</i>	Rainbow Trout			77.54	61.2	USEPA 2021
<i>Sander canadensis</i>	Sauger	77				USEPA 2014
<i>Perca flavescens</i>	Yellow Perch	73.98				USEPA 2014
<i>Micropterus salmoides</i>	Largemouth Bass	75.74 ^a		79.06 ^b 78.53 ^c		^a USEPA 2014; ^b Pinkney 2003, ^c May et al. 2009
<i>Micropterus dolomieu</i>	Smallmouth Bass	74.22				USEPA 2014
<i>Pomoxis annularis</i>	White Crappie			80.57		May et al. 2009
<i>Pomoxis nigromaculatus</i>	Black Crappie			79.75		May et al. 2009
<i>Lepomis macrochirus</i>	Bluegill		74.8	80.09	76	USEPA 2021
<i>Ambloplites rupestris</i>	Rock Bass	74.95				USEPA 2014
<i>Esox lucius</i>	Northern Pike			78		Rieberger 1992
<i>Pylodictis olivaris</i>	Flathead Catfish				58.97	May et al. 2009
<i>Scaphirhynchus platyrhynchus</i>	Shovelnose Sturgeon			77.13	47.18	May et al. 2009

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Appendix D: Extended List of Potential Target Species for Monitoring of Selenium in Fish Tissue

This appendix is intended for use with the recommendations in section 2.3 and Table 3. This appendix provides detailed information for the target species identified in Table 3, as well as additional species that may be considered appropriate for monitoring in certain situations (e.g., when species in Table 3 are unavailable for collection).

The tables in this appendix are generally organized by taxon (with the exception of a group of molluscivores) and includes nomenclature, distribution within US states, basic habitat information (warmwater [WW] or cool or coldwater [CW]), presence in waterbodies (e.g., lotic, lentic), adult diet, and adult trophic level. Information is presented by the following groupings:

1. Sturgeon in the family Acipenseridae
2. Sunfish and other genera in the family Centrarchidae
3. Trout and other genera in the family Salmonidae
4. Freshwater molluscivores & related genera (Catostomidae)
5. Minnows in the family Cyprinidae
6. Darters in the family Percidae
7. Sculpin in the family Scorpiionidae

Information sources for these species include NatureServe (<https://explorer.natureserve.org/Search>), and the USGS NAS - Nonindigenous Aquatic Species Database (<https://nas.er.usgs.gov/>). Users of this Appendix should consult with both as they examine available information to make decisions about target species in state and tribal waters as they develop sampling plans.

1. Sturgeon

The family Acipenseridae is comprised of twenty-seven species in four genera, with two genera (*Acipenser* and *Scaphyrhynchus*) occurring in the US. Sturgeon in the genus *Acipenser* (e.g., white sturgeon) include three freshwater species. Sturgeons are long-lived, late maturing bottom feeding fishes inhabiting large river systems and estuaries. Independent monitoring for these species is generally discouraged since most populations are under pressure from habitat loss and other stressors and coordination with federal agencies (US Fish and Wildlife Service or NOAA's National Marine Fishery Service) is therefore recommended prior to developing sampling plans that may include these species.

Table D-1. Species in the family Acipenseridae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Genus <i>Acipenser</i>			
White Sturgeon <i>(Acipenser transmontanus)</i>	US: AK, AZ, CA, ID, MT, OR, WA	WW Lotic Estuarine	Invertivore Piscivore Molluscivore TL3/TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=300 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100679/Acipenser_transmontanus			
Shortnose Sturgeon* <i>(Acipenser brevirostrum)</i>	US: CT, DC, DE, FL, GA, MA, MD, ME, NC, NH, NJ, NY, PA, RI, SC, VA	WW Lotic Estuarine	Invertivore Molluscivore TL3
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105033/Acipenser_brevirostrum			
Lake Sturgeon* <i>(Acipenser fluvescens)</i>	US: AL, AR, GA, IA, IL, IN, KS, KY, MI, MN, MO, NC, ND, NE, NY, OH, PA, SD, TN, VT, WI, WV	WW Lentic Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=299 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104232/Acipenser_fluvescens			
Other Sturgeon:			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Shovelnose Sturgeon <i>(Scaphirhynchus platorynchus)</i>	US: AL, AR, IA, IL, IN, KS, KY, LA, MN, MO, MS, MT, ND, NE, NM, OH, OK, PA, SD, TN, TX, WI, WV, WY	WW Lotic Estuarine	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103361/Scaphirhynchus_platorynchus			

* Species documented to consume invasive zebra mussels and quagga mussels in the genus *Dreissena* by the Army Corps of Engineers (Kirk, 2001).

2. Sunfish (genus *Lepomis*), and other genera in the family Centrarchidae

The bluegill is a species of freshwater fish and a member of the sunfish family Centrarchidae of the order Perciformes. It is native to North America and lives in streams, rivers, lakes, and ponds. The centrarchid family comprises 38 species of fish and includes many recreational and sportfish familiar to North Americans, including the rock bass, largemouth bass, pumpkinseed, and crappies. This family typically inhabits medium to large warmwater river systems and all sizes of lentic waterbodies. All species in the family are native to only North America.

Table D-2. Species in the family Centrarchidae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Genus <i>Lepomis</i>			
Bluegill (<i>Lepomis macrochirus</i>)	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=385 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101764/Lepomis_macrochirus			
Pumpkinseed* (<i>Lepomis gibbosus</i>)	US: AL, AZ, CA, CO, CT, DC, DE, GA, IA, ID, IL, IN, KY, MA, MD, ME, MI, MN, MT, NC, ND, NE, NH, NJ, NV, NY, OH, OR, PA, RI, SC, TN, TX, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=382 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105048/Lepomis_gibbosus			
Redear Sunfish* (<i>Lepomis microlophus</i>)	US: AL, AR, AZ, DC, DE, FL, GA, IA, IL, IN, KS, KY, LA, MI, MO, MS, NC, NE, NM, NV, OH, OK, OR, PA, SC, TN, TX, VA, VT, WV	WW Lentic Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=390 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100707/Lepomis_microlophus			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Green sunfish <i>(Lepomis cyanellus)</i>	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NJ, NM, NN, NV, NY, OH, OK, OR, PA, SC, SD, TN, TX, UT, VA, WA, WI, WV, WY	WW Lentic Lotic	Piscivore Invertivore TL3/4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=380 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103917/Lepomis_cyanellus https://pubs.usgs.gov/pp/1803/pdf/pp1803.pdf			
Redbreast Sunfish* <i>(Lepomis auritus)</i>	US: AL, AR, CT, DC, DE, FL, GA, KY, LA, MA, MD, ME, MS, NC, NH, NJ, NY, OK, PA, RI, SC, TN, TX, VA, VT, WV	WW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=379 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101339/Lepomis_auritus			
Longear Sunfish <i>(Lepomis megalotis)</i>	US: AL, AR, FL, GA, IA, IL, IN, KS, KY, LA, MD, MN, MO, MS, NC, NJ, NM, OH, OK, PA, TN, TX, VA, WI, WV	WW Lentic Lotic	Piscivore Invertivore TL3/4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=388 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.885331/Lepomis_megalotis			
Warmouth <i>(Lepomis gulosus)</i>	US: AL, AR, AZ, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MD, MI, MN, MO, MS, NC, NJ, NM, NV, NY, OH, OK, OR, PA, SC, TN, TX, VA, WA, WI, WV	WW Lentic Lotic	Piscivore Invertivore TL3/4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=376 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102803/Lepomis_gulosus			
Orangespotted Sunfish <i>(Lepomis humilis)</i>	US: AL, AR, CO, FL, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, ND, NE, OH, OK, PA, SD, TN, TX, WI, WV	WW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=383			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103679/Lepomis_humilis			
Redspotted Sunfish (<i>Lepomis miniatus</i>)	US: AL, AR, IL, IN, KY, LA, MO, MS, OK, TN, TX	WW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=391			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105029/Lepomis_miniatus			
Bantam Sunfish (<i>Lepomis symmetricus</i>)	US: AR, IL, IN, KY, LA, MO, MS, OK, TN, TX	WW Lentic Lotic	Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103718/Lepomis_symmetricus			
Northern Longear Sunfish (<i>Lepomis peltastes</i>)	US: IL, IN, MI, MN, NY, OH, PA	WW Lentic Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.883965/Lepomis_peltastes			
Spotted Sunfish (<i>Lepomis punctatus</i>)	US: FL, GA, NC, SC, TN	WW Lentic Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100708/Lepomis_punctatus			
Other Centrarchids			
Largemouth Bass (<i>Micropterus salmoides</i>)	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Piscivore TL4

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=401 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101622/Micropterus_salmoides			
Smallmouth Bass <i>(Micropterus dolomieu)</i>	US: AL, AR, AZ, CA, CO, CT, DC, DE, GA, IA, ID, IL, IN, KS, KY, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Piscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=396 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104786/Micropterus_dolomieu			
Spotted Bass <i>(Micropterus punctulatus)</i>	US: AL, AR, AZ, CA, FL, GA, IA, IL, IN, KS, KY, LA, MO, MS, NC, NE, NM, NV, OH, OK, PA, TN, TX, VA, WV	WW Lentic Lotic	Piscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=397 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.872252/Micropterus_punctulatus			
White Crappie <i>(Pomoxis annularis)</i>	US: AL, AR, AZ, CA, CO, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=408 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106200/Pomoxis_annularis			
Black Crappie <i>(Pomoxis nigromaculatus)</i>	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Piscivore TL3

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=409 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103134/Pomoxis_nigromaculatus			
Rock Bass <i>(Ambloplites rupestris)</i>	US: AL, AR, AZ, CT, DC, DE, GA, IA, IL, IN, KS, KY, MA, MD, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NY, OH, OK, PA, RI, SC, SD, TN, TX, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Molluscivore Piscivore TL3/TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=373 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105635/Ambloplites_rupestris			

* Species documented to consume invasive zebra mussels and quagga mussels in the genus *Dreissena* by the Army Corps of Engineers (Kirk, 2001).

3. Salmonids, including brown, rainbow, cutthroat trout and whitefish

Trout is the common name for species of freshwater fish belonging to the genera *Oncorhynchus*, *Salmo* and *Salvelinus* in the family Salmonidae. Trout are considered cold water fish and are usually found in clear streams, rivers and lakes with temperatures not exceeding 60°F (16°C).

Table D-3. Species in the family Salmonidae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Family Salmonidae			
Brown Trout (<i>Salmo trutta</i>)	US: AL, AR, AZ, CA, CO, CT, DC, DE, GA, IA, ID, IL, IN, KY, MA, MD, ME, MI, MN, MO, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SD, TN, UT, VA, VT, WA, WI, WV, WY	CW Lentic Lotic	Invertivore Molluscivore Piscivore TL3/TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=931			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103603/Salmo_trutta			
Rainbow Trout (<i>Oncorhynchus mykiss</i>)	US: AK, AL, AR, AZ, CA, CO, CT, DE, GA, HI, IA, ID, IL, IN, KS, KY, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	CW Lentic Lotic	Invertivore Piscivore TL3/TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=910			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105164/Oncorhynchus_mykiss			
Cutthroat Trout (<i>Oncorhynchus clarkii</i>)	US: AK, AR, AZ, CA, CO, ID, MD, MT, ND, NM, NN, NV, OR, UT, WA, WY	CW Lentic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=890			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103888/Oncorhynchus_clarkii			
Dolly Varden (<i>Salvelinus malma</i>)	US: AK, NV, NM, WA, WY	CW Lentic Lotic	Invertivore Molluscivore Piscivore

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
			TL3/TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=941			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104555/Salvelinus_malma			
Brook Trout (<i>Salvelinus fontinalis</i>)	US: AK, AR, AZ, CA, CO, CT, DE, GA, IA, ID, IL, IN, KY, MA, MD, ME, MI, MN, MT, NC, ND, NE, NH, NJ, NM, NV, NY, OH, OR, PA, RI, SC, SD, TN, UT, VA, VT, WA, WI, WV, WY	CW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=939			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103972/Salvelinus_fontinalis			
Mountain Whitefish (<i>Prosopium williamsoni</i>)	US: CA, CO, ID, MT, NV, OR, UT, WA, WY	CW Lentic Lotic	Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=924			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104696/Prosopium_williamsoni			
Round Whitefish (<i>Prosopium cylindraceum</i>)	US: AK, CT, IL, ME, MI, MN, NH, NY, VT, WI	CW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=921			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102380/Prosopium_cylindraceum			
Lake Whitefish* (<i>Coregonus clupeiformes</i>)	US: AK, ID, IL, IN, ME, MI, MN, MT, ND, NH, NV, NY, OH, PA, SD, VT, WA, WI	CW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=887			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105498/Coregonus_clupeaformis			

* Species documented to consume invasive zebra mussels and quagga mussels in the genus *Dreissena* by the Army Corps of Engineers (Kirk, 2001).

4. Freshwater Molluscivores and Related Genera

Molluscivorous fish feed either preferentially or opportunistically on a variety of mollusks (e.g., clams, mussels and snails) in freshwater systems. Although taxonomically diverse, physiologically these fish are adapted to feed on mollusks due to the presence of teeth or plates on the lower (and in some species upper) pharyngeal jaws, as well as mouth gape and jaw muscle structure that accommodates feeding on mollusks (Eastman 1977). A study by the Army Corps of Engineers (Kirk, 2001) documents at least 17 species* of North American fish that consume invasive zebra mussels and quagga mussels in the genus *Dreissena*. Several of these species are sunfish in the genus *Lepomis*; they are presented in the table addressing sunfish. Molluscivores may have elevated exposure to selenium, as mollusks bioaccumulate more selenium than other classes of aquatic invertebrates. These taxa typically inhabit larger warmwater lentic and lotic systems.

Table D-4. Molluscivores and related genera that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Freshwater Molluscivores			
Freshwater Drum* (<i>Aplodinotus grunniens</i>)	US: AL, AR, CO, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, MT, NC, ND, NE, NM, NY, OH, OK, PA, SD, TN, TX, VA, VT, WI, WV, WY	WW Lentic Lotic	Invertivore Piscivore Molluscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=946 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100338/Aplodinotus_grunniens			
White Perch* (<i>Morone americana</i>)	US: CO, CT, DC, DE, GA, IN, MA, MD, ME, MI, NC, NE, NH, NJ, NY, PA, RI, VA, VT, WI	WW Lentic Lotic Estuarine	Invertivore Molluscivore Piscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=777 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100436/Morone_americanana			
White bass* (<i>Morone chrysops</i>)	US: AL, AR, AZ, CA, CO, DC, FL, GA, IA, IL, IN, KS, KY, LA, MD, MI, MN, MO, MS, MT, NC, ND, NE, NM, NV, NY, OH, OK, PA, SC, SD, TN, TX, UT, VA, WI, WV	WW Lentic Lotic	Piscivore Invertivore Molluscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=779			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100951/Morone_chrysops			
Round Goby* <i>(Neogobius melanostomus)</i>	US: IL, IN, MI, MN, NY, OH, PA, WI	WW Lentic Lotic	Piscivore Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=713			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100501/Neogobius_melanostomus			
Brown Bullhead* <i>(Ameiurus nebulosus)</i>	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, NC, ND, NE, NH, NJ, NM, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, VA, VT, WA, WI, WV	WW Lentic Lotic	Omnivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=734			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103081/Ameiurus_nebulosus			
Yellow Perch* <i>(Perca flavescens)</i>	US: AL, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SC, SD, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Invertivore Piscivore Molluscivore TL4
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=820			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102985/Perca_flavescens			
Common carp* <i>(Cyprinus carpio)</i>	US: AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NN, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Omnivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=4			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105636/Cyprinus_carpio			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Catostomidae			
Smallmouth Buffalo (<i>Ictiobus bubalus</i>)	US: AL, AR, AZ, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, MT, NC, ND, NE, NM, OH, OK, PA, SD, TN, TX, WI, WV	WW Lentic Lotic	Herbivore Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/factsheet.aspx?SpeciesID=361 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105191/Ictiobus_bubalus			
Black Buffalo (<i>Ictiobus niger</i>)	US: AL, AR, AZ, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, NC, ND, NE, OH, OK, PA, SD, TN, TX, WI, WV	WW Lentic Lotic	Herbivore Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=363 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101227/Ictiobus_niger			
White sucker* (<i>Catostomus commersoni</i>)	US: AL, AR, CO, CT, DC, DE, GA, IA, IL, IN, KS, KY, MA, MD, ME, MI, MN, MO, MT, NC, ND, NE, NH, NJ, NM, NN, NY, OH, OK, PA, RI, SC, SD, TN, UT, VA, VT, WI, WV, WY	WW Lentic Lotic	Herbivore Molluscivore Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=346 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.833297/Catostomus_commersonii			
Largescale Sucker (<i>Catostomus macrocheilus</i>)	US: ID, MT, NV, OR, WA	CW Lentic Lotic	Herbivore Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.1098871/Catostomus_macrocheilus			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Greater Redhorse* <i>(Moxostoma valencienni)</i>	US: IL, IN, KY, MI, MN, ND, NY, OH, VT, WI	WW Lentic Lotic	Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101488/Moxostoma_valenciennesi			
Shorthead Redhorse <i>(Moxostoma macrolepidotum)</i>	US: DC, DE, IA, IL, IN, KS, MD, MI, MN, MO, MS, MT, NC, ND, NE, NY, OH, OK, PA, SC, SD, TX, VA, VT, WI, WV, WY	WW Lentic Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=366 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.791411/Moxostoma_macrolepidotum			
River Redhorse <i>(Moxostoma carinatum)</i>	US: AL, AR, FL, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, NC, NY, OH, OK, PA, TN, VA, WI, WV	WW Lentic	Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106031/Moxostoma_carinatum			
Golden redhorse <i>(Moxostoma erythrurum)</i>	US: AL, AR, DC, GA, IA, IL, IN, KS, KY, MD, MI, MN, MO, MS, NC, ND, NY, OH, OK, PA, SD, TN, TX, VA, WI, WV	WW Lentic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=365 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100778/Moxostoma_erythrurum			
Silver Redhorse <i>(Moxostoma anisurum)</i>	US: AL, AR, GA, IA, IL, IN, KY, MI, MN, MO, MS, ND, NY, OH, PA, TN, VA, VT, WI, WV	WW Lentic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=2912 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100712/Moxostoma_anisurum			

* Species documented to consume invasive zebra mussels and quagga mussels in the genus *Dreissena* by the Army Corps of Engineers (Kirk, 2001).

5. Minnows (*Cyprinidae*)

The family Cyprinidae (carps and minnows) is naturally distributed throughout most of the world and is the largest family of freshwater fishes with about 2,010 species in 210 genera. About 300 species in 50 genera are native to North America (Canada, Mexico, United States; Nelson, 2006). Cyprinids exhibit considerable variation in morphology, diet, and habitat use, and are often the only fish taxa (along with darters and sculpins) occurring in small order streams. Although cyprinids are not typically considered monitoring targets for contaminant analysis in their tissues, they are routinely collected as part of state biomonitoring programs that use the fish index of biotic integrity to assess stream health in wadeable streams.

EPA recommends that fish tissue monitoring programs collaborate with state or tribal biomonitoring programs to leverage expertise, experience and resources to collect cyprinids and related species in watersheds located in geographic areas of elevated selenium where anthropogenic activities may introduce selenium to surface waters if other more sensitive species are not present.

Table D-5. Species in the family Cyprinidae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Fathead Minnow (<i>Pimephales promelas</i>)	US: AL, AR, AZ, CA, CO, CT, DE, GA, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NM, NN, NV, NY, OK, OR, PA, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	WW Lentic Lotic	Herbivore Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=621			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102599/Pimephales_promelas			
Bluntnose Minnow (<i>Pimephales notatus</i>)	US: AL, AR, CT, DC, GA, IA, IL, IN, KS, KY, LA, MA, MD, MI, MN, MO, MS, NC, ND, NE, NJ, NY, OH, OK, PA, SD, TN, VA, VT, WI, WV	WW Lentic Lentic	Herbivore Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=620			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103436/Pimephales_notatus			
Bullhead Minnow (<i>Pimephales vigilax</i>)	US: AL, AR, CO, GA, IA, IL, IN, KS, KY, LA, MN, MO, MS, NE, NM, OH, OK, PA, SD, TN, TX, VA, WI,	WW Lentic Lentic	Herbivore Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=623			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106123/Pimephales_vigilax			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Cutlip Minnow (<i>Exoglossum maxilllingua</i>)	US: CT, DC, DE, MD, NC, NJ, NY, PA, VA, VT, WV	WW Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=530 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102719/Exoglossum_maxilllingua			
Suckermouth Minnow (<i>Phenacobius mirabilis</i>)	US: AL, AR, CO, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, NE, NM, OH, OK, SD, TN, TX, VA, WI, WV, WY	WW Lotic	Herbivore Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=617 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104716/Phenacobius_mirabilis			
Blackstripe Topminnow (<i>Fundulus notatus</i>)	US: AL, AR, IA, IL, IN, KS, KY, LA, MI, MO, MS, OH, OK, TN, TX, WI	WW Lentic Lotic	Herbivore Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=690 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100269/Fundulus_notatus			
Starhead Topminnow (<i>Fundulus dispar</i>)	US: AL, AR, FL, IA, IL, IN, KY, LA, MI, MO, MS, OK, TN, WI	WW Lentic Lotic	Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105342/Fundulus_dispar			
Western Blacknose Dace (<i>Rhinichthys obtusus</i>)	US: AL, GA, IA, IL, IN, KS, KY, MI, MN, MO, MS, NC, ND, NE, NY, OH, PA, SC, SD, TN, VA, WI, WV	CW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.790464/Rhinichthys_obtusum			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Eastern Blacknose Dace (<i>Rhinichthys atratulus</i>)	US: CT, DC, DE, GA, MA, MD, ME, NC, NH, NJ, NY, PA, RI, VA, VT, WV	CW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=637			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.828296/Rhinichthys_atratulus			
Longnose Dace (<i>Rhinichthys cataractae</i>)	US: CO, CT, DC, DE, GA, IA, ID, IL, IN, MA, MD, ME, MI, MN, MT, NC, ND, NE, NH, NJ, NM, NV, NY, OH, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY	CW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=638			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101847/Rhinichthys_cataractae			
Finescale Dace (<i>Chrosomus neogaeus</i>)	US: ME, MI, MN, ND, NE, NH, NY, SD, VT, WI, WY	WW Lentic Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=2556			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102927/Chrosomus_neogaeus			
Speckled Dace (<i>Rhinichthys osculus</i>)	US: AZ, CA, CO, ID, NM, NN, NV, OR, UT, WA, WY	CW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=640			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100335/Rhinichthys_osculus			
Satinfin Shiner (<i>Cyprinella analostana</i>)	US: DC, DE, MD, NC, NJ, NY, PA,	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=516			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106108/Cyprinella_analostana			
Red Shiner (<i>Cyprinella lutrensis</i>)	US: AL, AR, AZ, CO, GA, IA, IL, IN, KS, KY, LA, MN, MO, MS, NC, ND, NE, NM, NN, NV, OK, SD, TN, TX,	WW Lentic Lotic	Invertivore TL3

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=518			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105504/Cyprinella_lutrensis			
Bigmouth Shiner (<i>Notropis dorsalis</i>)	US: CO, IA, IL, IN, KS, MI, MN, MO, ND, NE, NY, OH, PA, SD, TN, WI, WV, WY	WW Lentic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=593			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104308/Notropis_dorsalis			
Chub Shiner (<i>Notropis potteri</i>)	US: AR, LA, OK, TX	WW Lentic Lotic	Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=606			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105179/Notropis_potteri			
Sand Shiner (<i>Notropis stramineus</i>)	US: AR, AZ, CO, IA, IL, IN, KS, KY, MI, MN, MO, MT, ND, NE, NM, NN, NY, OH, OK, PA, SD, TN, TX, UT, VA, VT, WI, WV, WY	WW Lentic Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=600			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104717/Notropis_stramineus			
Redside Shiner (<i>Richardsonius balteatus</i>)	US: AZ, CO, ID, MT, NV, OR, UT, WA, WY	CW Lentic Lotic	Herbivore Invertivore Molluscivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=644			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100279/Richardsonius_balteatus			
Thicklip Chub (<i>Cyprinella labrosa</i>)	US: NC, SC, VA	WW Lentic	Invertivore Molluscivore TL3

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101215/Cyprinella_labrosa			
Streamline Chub (<i>Erimystax dissimilis</i>)	US: AL, IN, KY, NY, OH, PA, TN, VA, WV	WW Lotic	Invertivore Molluscivore TL3
Map: & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106034/Erimystax_dissimilis			
Shoal Chub (<i>Macrhybopsis hyostoma</i>)	US: AL, AR, IA, IL, IN, KS, KY, LA, MN, MO, MS, NE, OH, OK, TN, TX, WI, WV	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106278/Macrhybopsis_hyostoma			
Silver Chub (<i>Macrhybopsis storeriana</i>)	US: AL, AR, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, ND, NE, NY, OH, OK, PA, SD, TN, TX, WI, WV	WW Lotic	Invertivore Molluscivore TL3
Map: Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101653/Macrhybopsis_storeriana			
River Chub (<i>Nocomis micropogon</i>)	US: AL, DC, GA, IL, IN, KY, MD, MI, NC, NY, OH, PA, SC, TN, VA, WV	WW Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=577 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101786/Nocomis_micropogon			
Bull Chub* (<i>Nocomis raneyi</i>)	US: NC, VA	WW Lotic	Herbivore Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101374/Nocomis_raneyi			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Peamouth <i>(Mylocheilus caurinus)</i>	US: ID, MT, OR, WA	CW Lentic Lotic	Invertivore Molluscivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=2349 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100544/Mylocheilus_caurinus			
Creek Chub <i>(Semotilus atromaculatus)</i>	US: AL, AR, CO, CT, DC, DE, FL, GA, IA, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NY, OH, OK, PA, SC, SD, TN, TX, UT, VA, VT, WI, WV, WY	WW Lotic	Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=649 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104867/Semotilus_atromaculatus			
Central Stoneroller <i>(Campostoma anomalum)</i>	US: AR, CO, CT, GA, IA, IL, IN, KS, KY, LA, MD, MI, MN, MO, MS, NC, ND, NE, NM, NY, OH, OK, PA, SC, SD, TN, TX, VA, WI, WV, WY	WW Lotic	Herbivore
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=506 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.844144/Campostoma_anomalum			
Largescale Stoneroller <i>(Campostoma oligolepis)</i>	US: AL, AR, GA, IA, IL, IN, KY, MN, MO, MS, ND, OK, VA, WI	WW Lotic	Herbivore
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=507 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102552/Campostoma_oligolepis			
Sacramento Splittail <i>(Pogonichthys macrolepidotus)</i>	US: CA	WW Lotic Estuarine	Herbivore Invertivore

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105438/Pogonichthys_macrolepidotus			

* Species documented to consume invasive zebra mussels and quagga mussels in the genus *Dreissena* by the Army Corps of Engineers (Kirk, 2001).

6. Darters (*Percidae*)

Darters are small, perch-like fish in the family Percidae and are found in freshwater streams in North America. Darters typically occur in riverine systems, inhabiting cold to cool streams and small river systems in North America. Species distributions range from single watersheds in one state to multiple watersheds in several states. Darters are typically benthic omnivores, practicing herbivory as well as preying on invertebrates and for some species, small fish and fish eggs as well.

Table D-6. Species in the family Percidae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Greenside Darter (<i>Etheostoma blennioides</i>)	US: AL, AR, DC, GA, IL, IN, KS, KY, MD, MI, MO, MS, NC, NY, OH, OK, PA, TN, VA, WV	WW Lotic Lentic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=808			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.790349/Etheostoma_blennioides			
Arkansas Darter (<i>Etheostoma cragini</i>)	US: AR, CO, KS, MO, OK	WW Lotic	Invertivore Molluscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=810			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103800/Etheostoma_cragini			
Iowa Darter (<i>Etheostoma exile</i>)	US: CO, IA, IL, IN, MI, MN, MT, ND, NE, NM, NY, OH, PA, SD, UT, WI, WY	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=812			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100441/Etheostoma_exile			
Fantail Darter (<i>Etheostoma flabellare</i>)	US: AL, AR, DC, IA, IL, IN, KS, KY, MD, MI, MN, MO, MS, NC, NY, OH, OK, PA, SC, TN, VA, VT, WI, WV	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.832912/Etheostoma_flabellare			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Least Darter (<i>Etheostoma microperca</i>)	US: AR, IA, IL, IN, KS, KY, MI, MN, MO, OH, OK, WI	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103616/Etheostoma_microperca			
Johnny Darter (<i>Etheostoma nigrum</i>)	US: AL, AR, CO, IA, IL, IN, KS, KY, MD, MI, MN, MO, MS, NC, ND, NE, NY, OH, OK, PA, SD, TN, UT, VA, WI, WV, WY	WW Lotic Lentic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=814 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.100152/Etheostoma_nigrum			
Tessellated Darter (<i>Etheostoma olmstedii</i>)	US: CT, DC, DE, FL, GA, MA, MD, NC, NH, NJ, NY, PA, RI, SC, VA, VT, WV	WW Lotic Lentic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=816 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106063/Etheostoma_olmstedii			
Cypress Darter (<i>Etheostoma proeliare</i>)	US: AL, AR, FL, IL, KY, LA, MO, MS, OK, TN, TX	WW Lotic Lentic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101593/Etheostoma_proeliare			
Redline Darter (<i>Etheostoma rufilineatum</i>)	US: AL, GA, KY, MS, NC, TN, VA	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=2886 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103471/Etheostoma_rufilineatum			
Orangebelly Darter (<i>Etheostoma radiosum</i>)	US: AR, OK, TX	WW Lotic	Invertivore TL3

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.1156842/Etheostoma_radiosum			
Orangethroat Darter (<i>Etheostoma spectabile</i>)	US: AR, CO, IA, IL, IN, KS, KY, MI, MO, NE, OH, OK, TN, TX, WY	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102592/Etheostoma_spectabile			
Speckled Darter (<i>Etheostoma stigmaeum</i>)	US: AL, AR, FL, GA, KY, LA, MO, MS,	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.1006953/Etheostoma_stigmaeum			
Gulf Darter (<i>Etheostoma swaini</i>)	US: AL, FL, GA, KY, LA, MS, TN	WW Lotic	Invertivore TL3
Map & Info: Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102176/Etheostoma_swaini			
Variegate Darter (<i>Etheostoma variatum</i>)	US: IN, KY, NY, OH, PA, VA, WV	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=3333 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102018/Etheostoma_variatum			
Banded Darter (<i>Etheostoma zonale</i>)	US: AL, AR, GA, IA, IL, IN, KS, KY, MD, MI, MN, MO, NC, NY, OH,	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=818 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106576/Etheostoma_zonale			
River Darter (<i>Percina shumardi</i>)	US: AL, AR, GA, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, ND, OH, OK, PA, TN, TX, WI, WV	WW Lotic	Invertivore TL3

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=826 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101870/Percina_shumardi			
Slenderhead Darter (<i>Percina phoxocephala</i>)	US: AL, AR, IA, IL, IN, KS, KY, MN, MO, MS, OH, OK, SD, TN, WI, WV	WW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.104090/Percina_phoxocephala			
Shield Darter (<i>Percina peltata</i>)	US: DC, DE, MD, NJ, NY, PA, VA, WV	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=2775 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.105028/Percina_peltata			
Blackbanded Darter (<i>Percina nigrofasciata</i>)	US: AL, FL, GA, LA, MS, NC, SC, TN	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=824 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.102848/Percina_nigrofasciata			
Blackside Darter (<i>Percina maculata</i>)	US: AL, AR, IA, IL, IN, KS, KY, LA, MI, MN, MO, MS, ND, NE, NY, OH, OK, PA, SD, TN, TX, VA, WI, WV	WW Lotic	Invertivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=823 Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.106566/Percina_maculata			

7. Sculpins (*Scorpionidae*)

Sculpins are members of the family Scorpionidae (Scorpionfish), and most species in the Northern Hemisphere are saltwater fishes. Most freshwater sculpins in the US are in the genus *Cottus*, and are small benthic predators consuming mainly invertebrates. Sculpins typically prefer cooler, headwater streams, but can be found in larger warmer streams in some states.

Table D-7. Species in the family Scorpionidae that may be sampled for implementation of the selenium criterion.

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Mottled Sculpin (<i>Cottus bairdii</i>)	US: AL, AZ, CO, DE, GA, IA, ID, IL, IN, KY, MD, MI, MN, MO, MS, MT, NC, NM, NV, NY, OH, OR, PA, SC, TN, UT, VA, VT, WA, WI, WV, WY	CW/WW Lotic Lentic	Herbivore Invertivore Piscivore TL3
Map: https://nas.er.usgs.gov/queries/FactSheet.aspx?SpeciesID=502			
Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.819868/Cottus_bairdii			
Paiute Sculpin (<i>Cottus beldingii</i>)	US: CA, CO, ID, NV, OR, UT, WA, WY	CW/WW Lotic	Herbivore Molluscivore Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101884/Cottus_beldingii			
Banded Sculpin (<i>Cottus carolinae</i>)	US: AL, AR, GA, IL, IN, KS, KY, MO, MS, NC, OK, TN, VA	CW/WW Lotic	Herbivore Invertivore Piscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.819914/Cottus_carolinae			
Slimy Sculpin (<i>Cottus cognatus</i>)	US: AK, CT, IA, ID, IL, IN, MA, ME, MI, MN, MT, NH, NJ, NY, PA, VA, VT, WA, WI, WV	CW Lotic Lentic	Herbivore Invertivore Piscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.101449/Cottus_cognatus			

Name (Common, Scientific)	Distribution	Habitat (WW/CW) Lentic/Lotic	Adult Diet/ Trophic Level
Shorthead Sculpin (<i>Cottus confusus</i>)	US: ID, NV, OR, WA	CW Lotic	Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.905574/Cottus_confusus			
Riffle Sculpin (<i>Cottus gulosus</i>)	US: CA, OR, WA	CW Lotic	Invertivore Molluscivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103055/Cottus_gulosus			
Spoonhead Sculpin (<i>Cottus ricei</i>)	US: IL, MI, MN, MT, NY, OH, PA, WI	CW Lotic Lentic	Omnivore Invertivore TL3
Map & Info: https://explorer.natureserve.org/Taxon/ELEMENT_GLOBAL.2.103689/Cottus_ricei			

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Appendix E: Calculation of Composite Trophic Transfer Factors

Derivation of Trophic Transfer Factor (TTF) Values

The parameter $TTF^{composite}$ (composite trophic transfer factor) in Equation 1 quantitatively represents all dietary pathways of selenium exposure for a particular fish species within an aquatic system. The parameter is derived from species-specific TTF values representing the food web characteristics of the aquatic system and the proportion of species consumed. It is possible to differentiate bioaccumulative potential for different predator species and food webs by modeling different exposure scenarios. For example, where a fish species of interest is a trophic level 4 predator that primarily consumes trophic level 3 fish, the term $TTF^{composite}$ can be represented as the product of all TTF parameters that includes the additional trophic level given as:

$$TTF^{composite} = TTF^{TL4} \times TTF^{TL3} \times TTF^{TL2}$$

(Equation 1)

where:

TTF^{TL2}	=	the trophic transfer factor of the trophic level 2 species
TTF^{TL3}	=	the trophic transfer factor of the trophic level 3 species
TTF^{TL4}	=	the trophic transfer factor of the trophic level 4 species
$TTF^{composite}$	=	the product of all the trophic transfer factors

The consumption of more than one species of organism at the same trophic level can also be modeled by expressing the TTF at a particular trophic level as the weighted average of the TTF s of all species consumed given as:

$$\overline{TTF}^{TLx} = \sum_i (TTF_i^{TLx} \times w_i)$$

(Equation 2)

where:

TTF_i^{TLx}	=	the trophic transfer factor of the i^{th} species at a particular trophic level
w_i	=	the proportion of the i^{th} species consumed

Figure 1 below describes five example food web scenarios and the formulation of $TTF^{composite}$ to model selenium bioaccumulation in each of them.

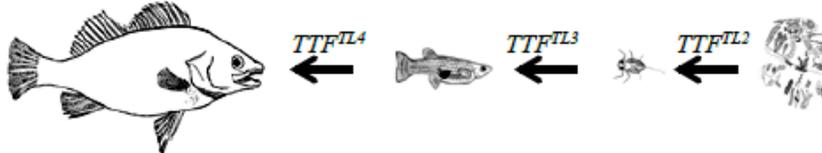
A) Three trophic levels (simple):

$$TTF^{composite} = TTF^{TL3} \times TTF^{TL2}$$



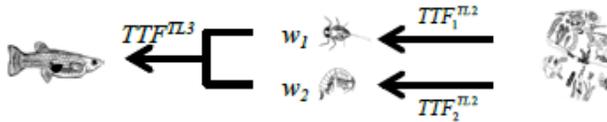
B) Four trophic levels (simple):

$$TTF^{composite} = TTF^{TL4} \times TTF^{TL3} \times TTF^{TL2}$$



C) Three trophic levels (mix within trophic levels):

$$TTF^{composite} = TTF^{TL3} \times \left[(TTF_1^{TL2} \times w_1) + (TTF_2^{TL2} \times w_2) \right]$$



D) Three trophic levels (mix across trophic levels):

$$TTF^{composite} = (TTF^{TL3} \times w_1) + (TTF^{TL3} \times TTF^{TL2} \times w_2)$$



E) Four trophic levels (mix across trophic levels):

$$TTF^{composite} = \left[(TTF^{TL4} \times TTF^{TL3} \times w_1) + (TTF^{TL4} \times w_2) \right] \times TTF^{TL2}$$

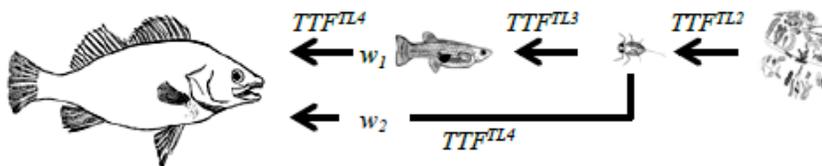


Figure 1. Example aquatic system scenarios and the derivation of the equation parameter $TTF^{composite}$.