WQX 3.0 Benthic Macroinvertebrate Best Practices

WQX User Call August 27, 2020



Background

- Convened Biological Workgroup to determine requirements
- Goals:
 - Make datasets more user-friendly
 - Promote consistency when submitting data
 - Maximize the extent data can be reused
- Created a best practices document addressing new and important data elements





Assemblage Sampled Name & Biological Intent

- Assemblage is the group of organisms that cooccur in the same trophic or spatial relationships, such as benthic macroinvertebrates
 - Used to derive metadata requirements
 - Required
- Biological intent is the goal of the assessment such as species density or population census
 - Helps users understand the data to expect
 - Required

Habitat Selection Method

- Targeted vs. Non-Targeted
- Most productive habitat is sampled with a targeted method
 - More likely to capture all taxa
- Non-targeted uses a systematic sampling technique
 - Often more quantitatively representative
- Conditionally required
- New field in WQX 3.0





Collection Effort

- Time-Constrained
 - Collection Duration Measure and Units
- Area-Constrained
 - Collection Area Measure and Units
 - Needed along with Proportion Sample Processed to calculate total abundance
 - New fields in WQX 3.0
- Constrained by staff or equipment
 - Gear Procedure Measure and Units
 - New fields in WQX 3.0
- Optional but recommended

Sample Collection Equipment Name & Net Mesh Size

- Equipment such as D-frame net can impact what taxa or life stages are captured
 - Optional but recommended
- Net mesh size and net surface area used in sampling can also impact what taxa or life stages are captured
 - Used with net type such as non-tow
 - Optional but recommended





Characteristic Name

- Characteristic name is the parameter being monitored
- Usually a count with the value of the count being the result
- Required

Target Count

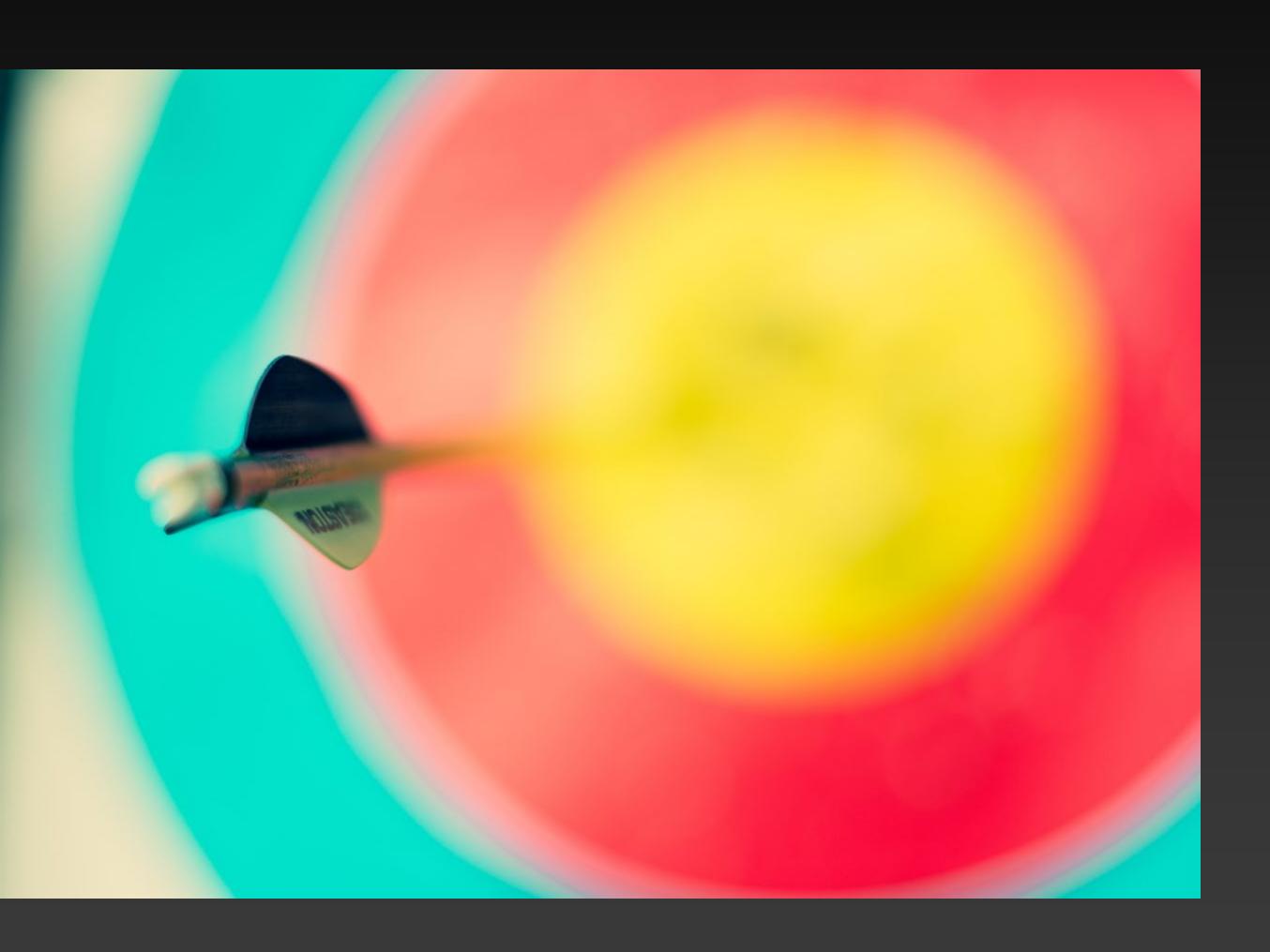
- Sorter will count a certain number of individual specimens and stop
- Useful in determining data comparability or if the data user needs to randomly subset the data before use
- Conditionally required
- New field in WQX 3.0



Proportion Sample Processed

- Prior to sorting the sample is placed in a sampling grid where a sorter chooses a subset of the sample for identification
- This information is important to know when aggregating data
- Conditionally required
- New field in WQX 3.0

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Target Taxonomic Levels

- Organizations identify specimens to a certain level of taxonomic resolution and life stage
- Data users need to know this information to determine if a life stage or taxon was not encountered or was deliberately excluded
- Also important to know whether identification is done to the genus level or the family level
- Should be specified in Sample Collection
 Method

Subject Taxonomic Name & Subject Taxonomic Name User Supplied

- Subject taxonomic name is the name of the specimen identified
 - Required
- Subject taxonomic name user supplied is for organizations who use a preferred name that may be different than WQX domain values
 - Used with subject taxonomic name user supplied reference text
 - Optional
 - New field in WQX 3.0





Uploading QA/QC Documents

- QAPP outlines data and measurement quality objectives of a monitoring program and contains metrics to quantify data quality
- Data users use the QAPP to screen for comparable protocols and comparable data quality
- Optional but recommended to upload at the Project or Activity level

Status

- New data elements included with WQX 3.0
- EN users can still submit to WQX 2.2, but must move to WQX 3.0 to use new data elements
- WQX Web users have already been migrated to WQX 3.0
- Best practices document is available at: https://www.epa.gov/sites/production/files/2020-08/documents/wqx-benthics-best-practices-guide.pdf

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