

St. Louis River estuary as possible Dreissena veliger source to western Lake Superior

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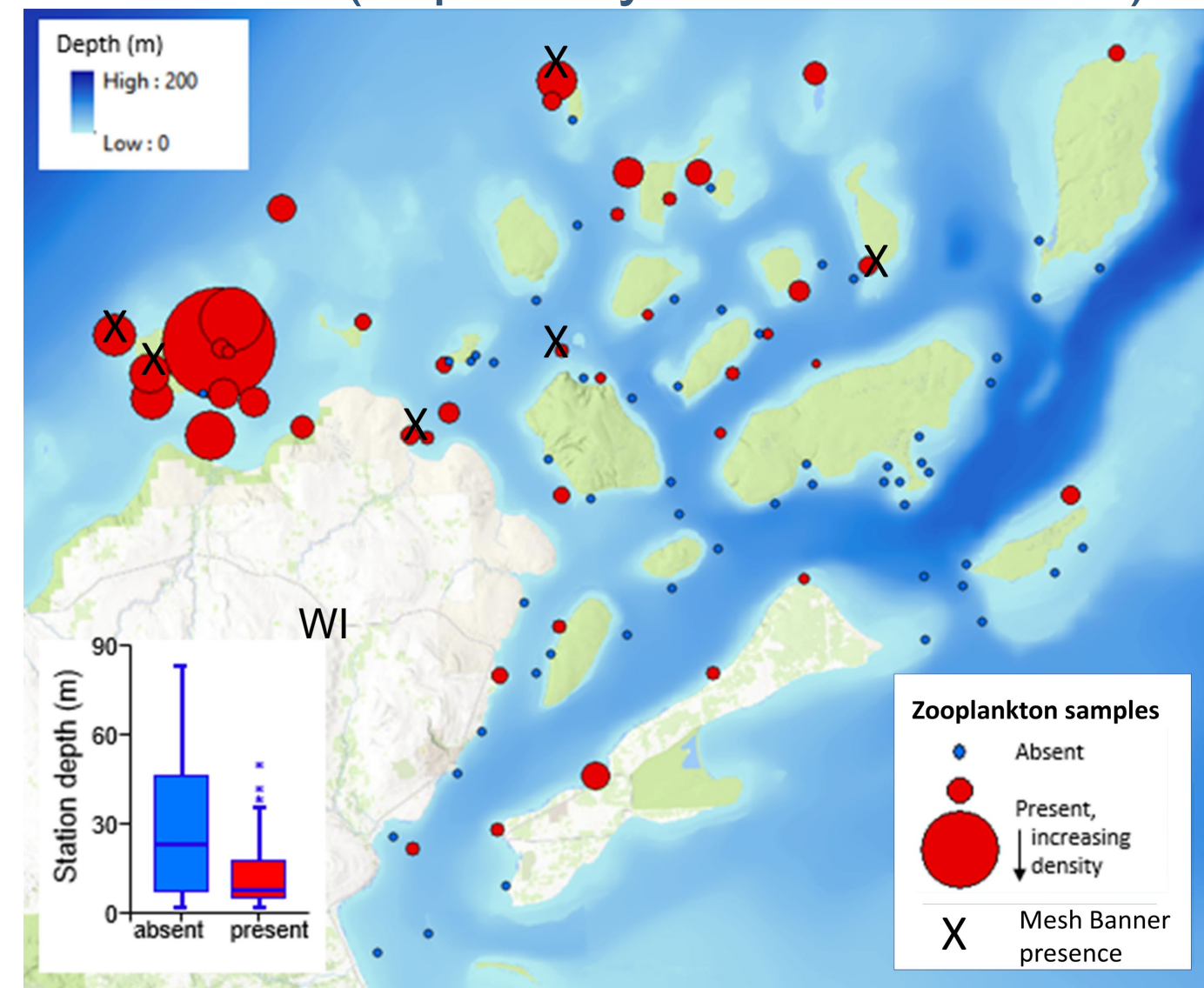
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Background

In 2017 EPA conducted early detection case study in western Lake Superior around the Apostle Islands (APIS) addressing concerns over adult Dreissena finds on shipwrecks & native mussels by Nat'l Park Service

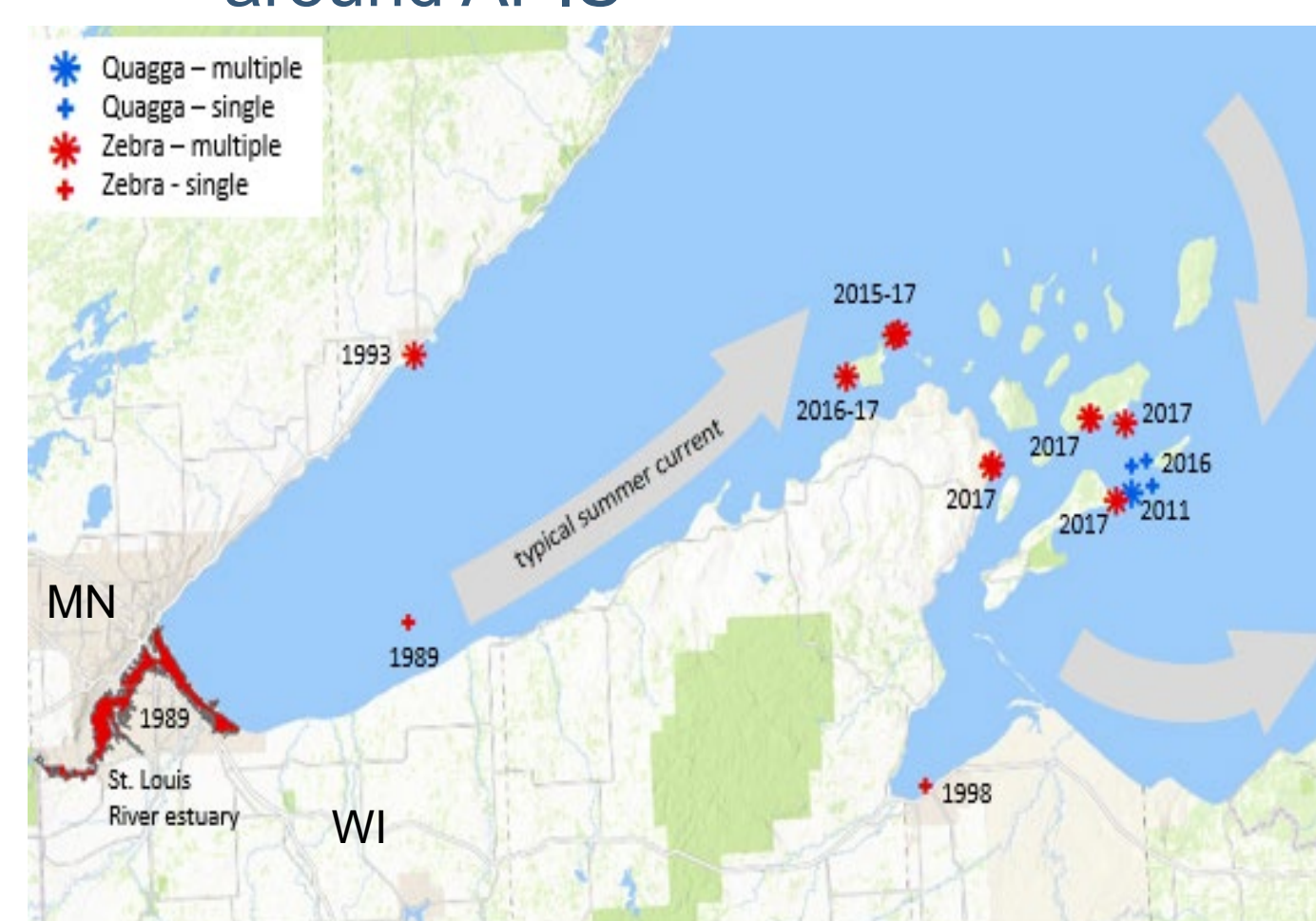
Results:

- No settled juvenile or adult Dreissena on passive gears
- Dreissena veligers present in 44% of zooplankton samples albeit in low densities
- Finds primarily along NW side of islands (especially around Sand Is.)



Conclusions:

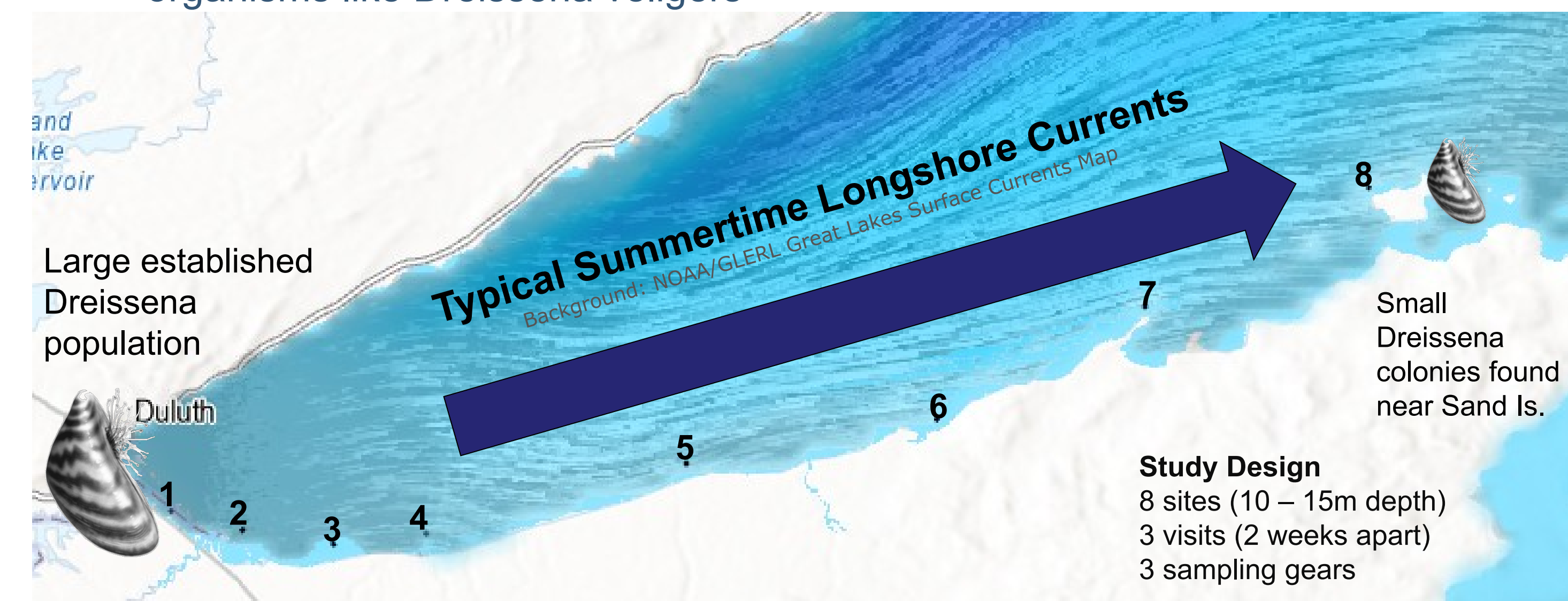
- Low veliger densities suggest adult Dreissenid populations are low and detected veligers may not have originated in APIS
- APIS detections point to possible transportation from longshore surface currents
 - St. Louis River Estuary (SLRE) has largest and most established Dreissena population in Lake Superior
 - Typical summer surface currents transport water from SLRE around APIS



Survey Overview, Design, & Methods (2019 W. Lake Superior south shore survey)

Objectives: Follow up 2017 survey by determining concentration gradient and detection of Dreissena veligers along south shore (SLRE to APIS)

- Evaluate if SLRE is a potential veliger source of APIS Dreissena and determine if a gradient of decreasing detection exists
- Evaluate modified methods for increased probability in detecting low abundance organisms like Dreissena veligers



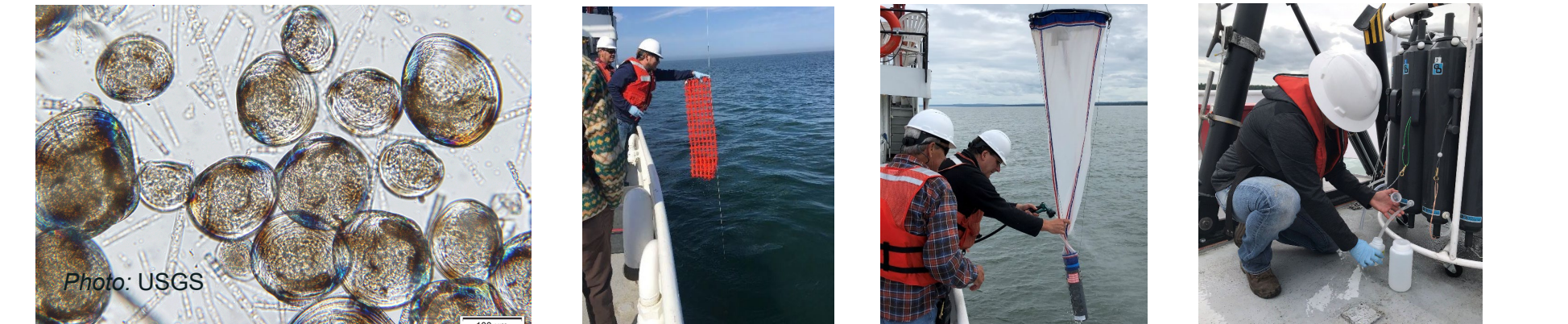
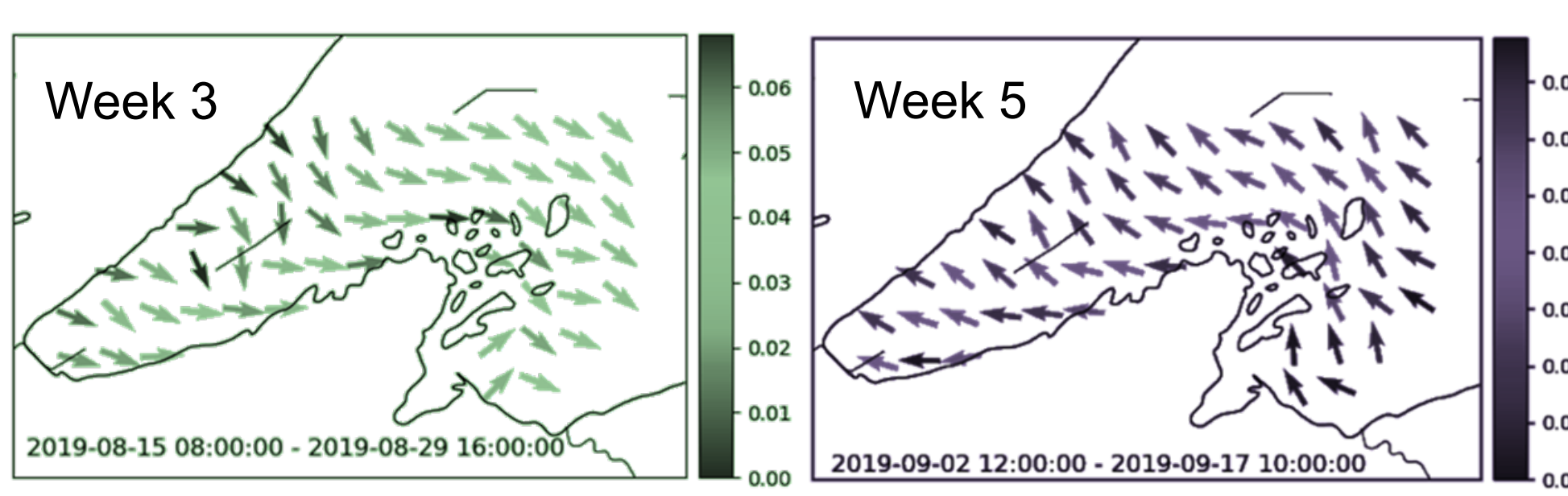
Study Design
8 sites (10 – 15m depth)
3 visits (2 weeks apart)
3 sampling gears

Trip			
	Week 1 (Aug. 12, 14)	Week 3 (Aug 29)	Week 5 (Sept 16, 17)
	deploy banners, collect zooplankton tows & eDNA	collect zooplankton tows & eDNA	retrieve banners, collect zooplankton tows & eDNA
Gear	2(2'x4') mesh banners (N = 16)	Zooplankton tow 64µm mesh (N = 24)	eDNA (N = 48)
Sampling modifications	2, larger, more rigid banners. Sampled 1m below surface & 2m above bottom	Composited 4 tows per sample. Saved decanted EtOH for DNA analysis	Increased water volume sampled to 1L. Sampled 3m below surface & 2m above benthos
Processing & analysis	qPCR targeting Dreissena w/ Genus and species level markers	Zoops: full enumeration Zoop EtOH: qPCR targeting Dreissena	Filtered samples & qPCR targeting Dreissena using general and species-specific markers

Environmental Conditions

Surface current patterns leading up to each sampling week (time averaged velocity; 2wks prior to sampling)

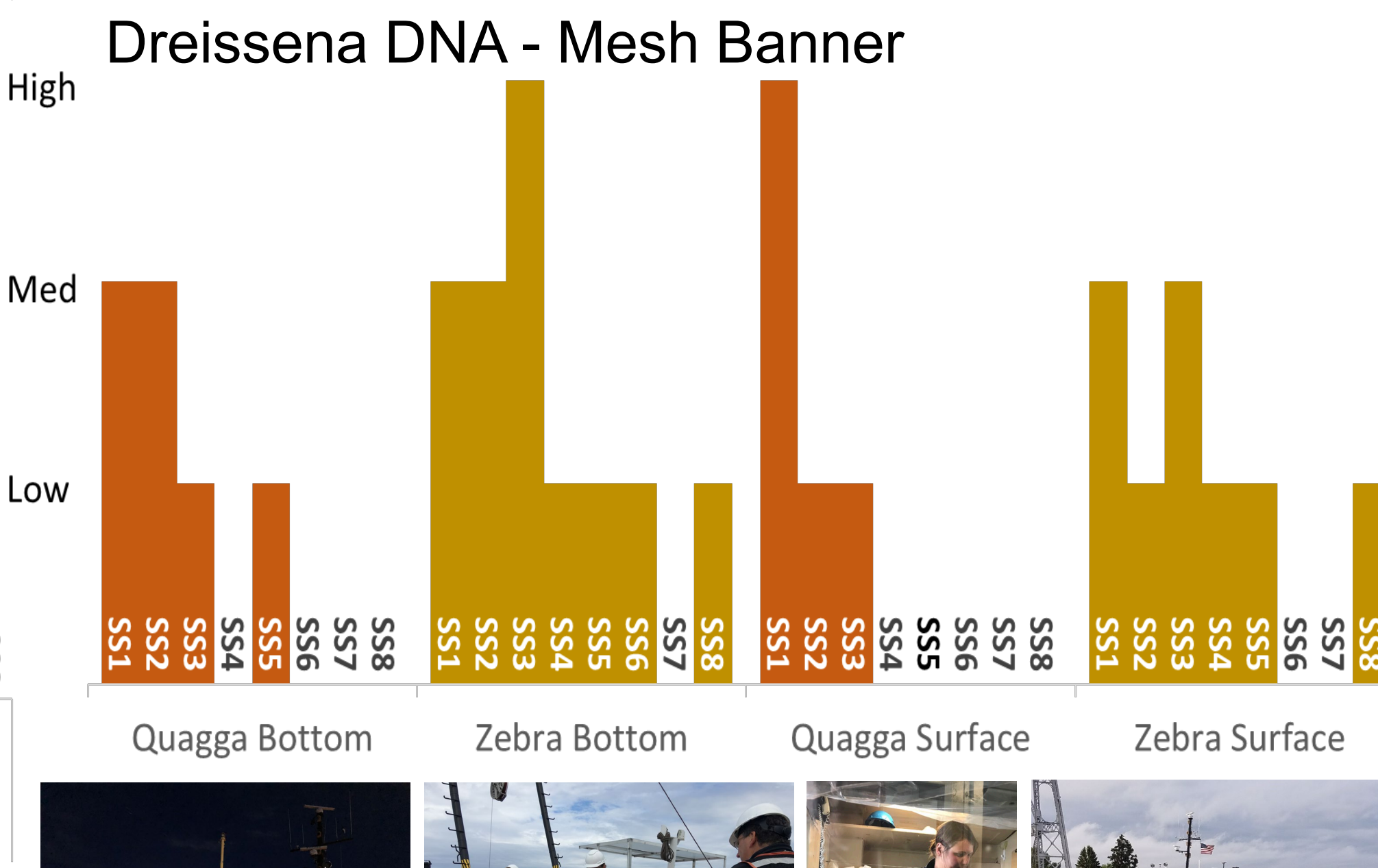
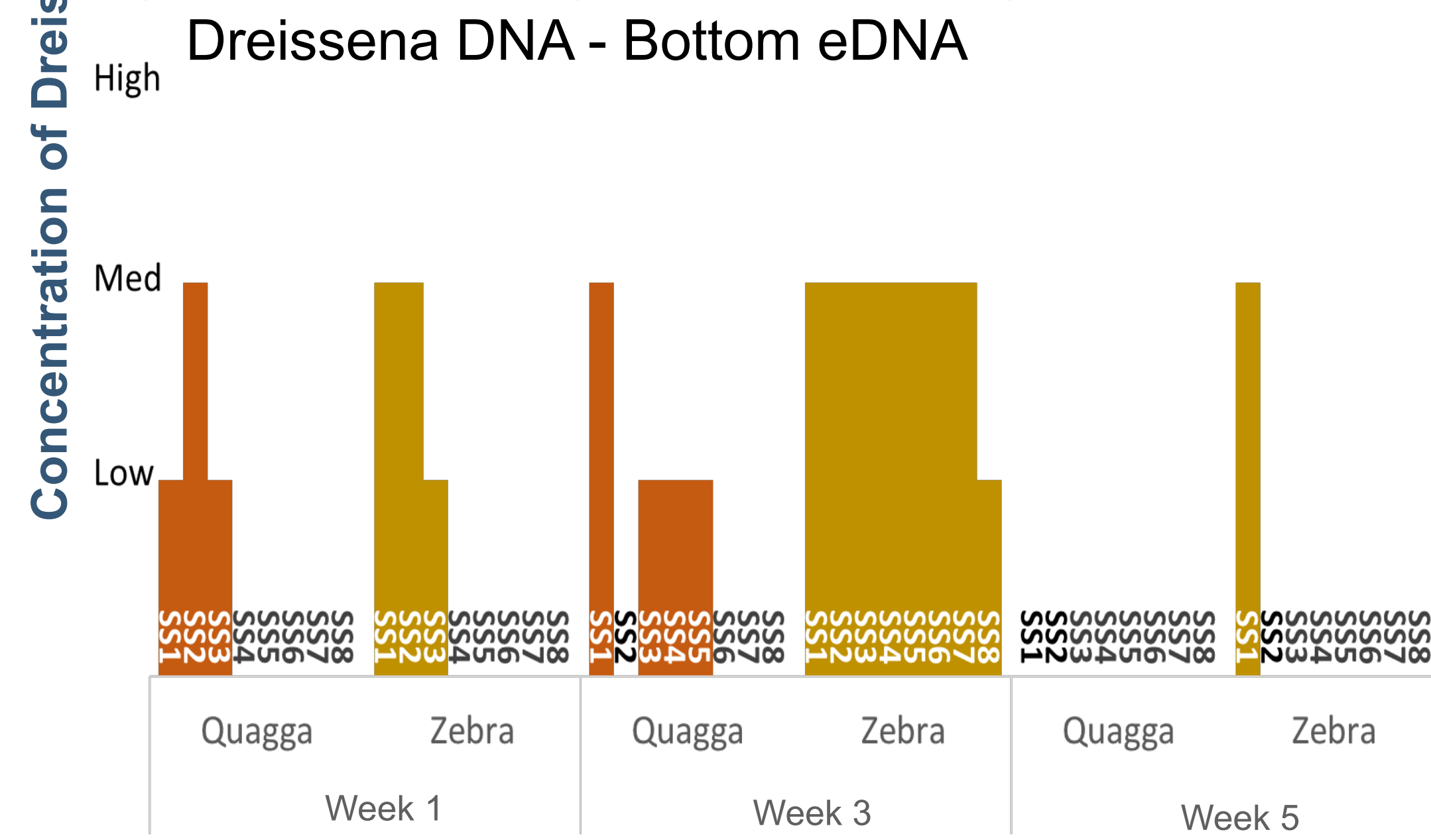
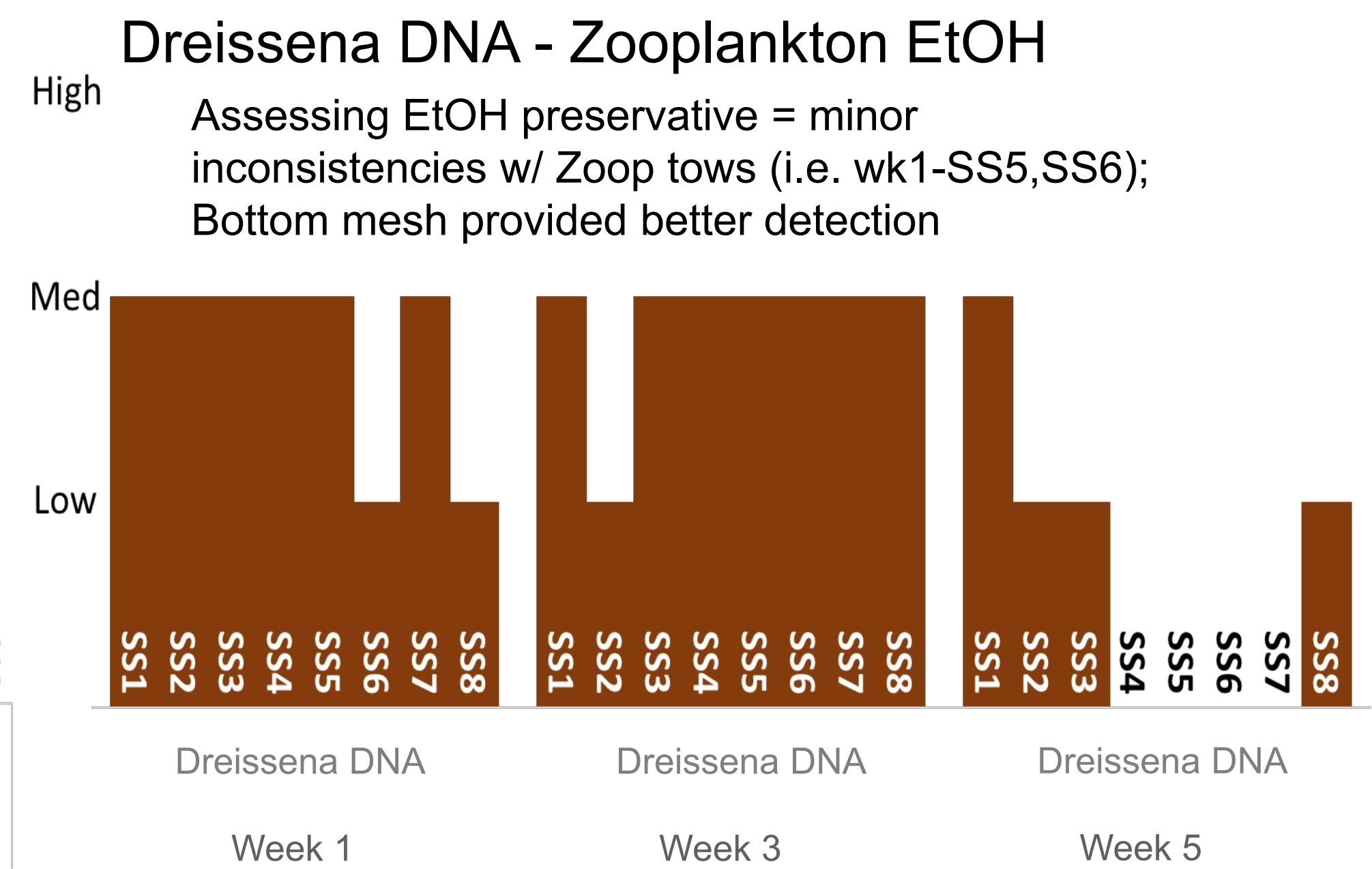
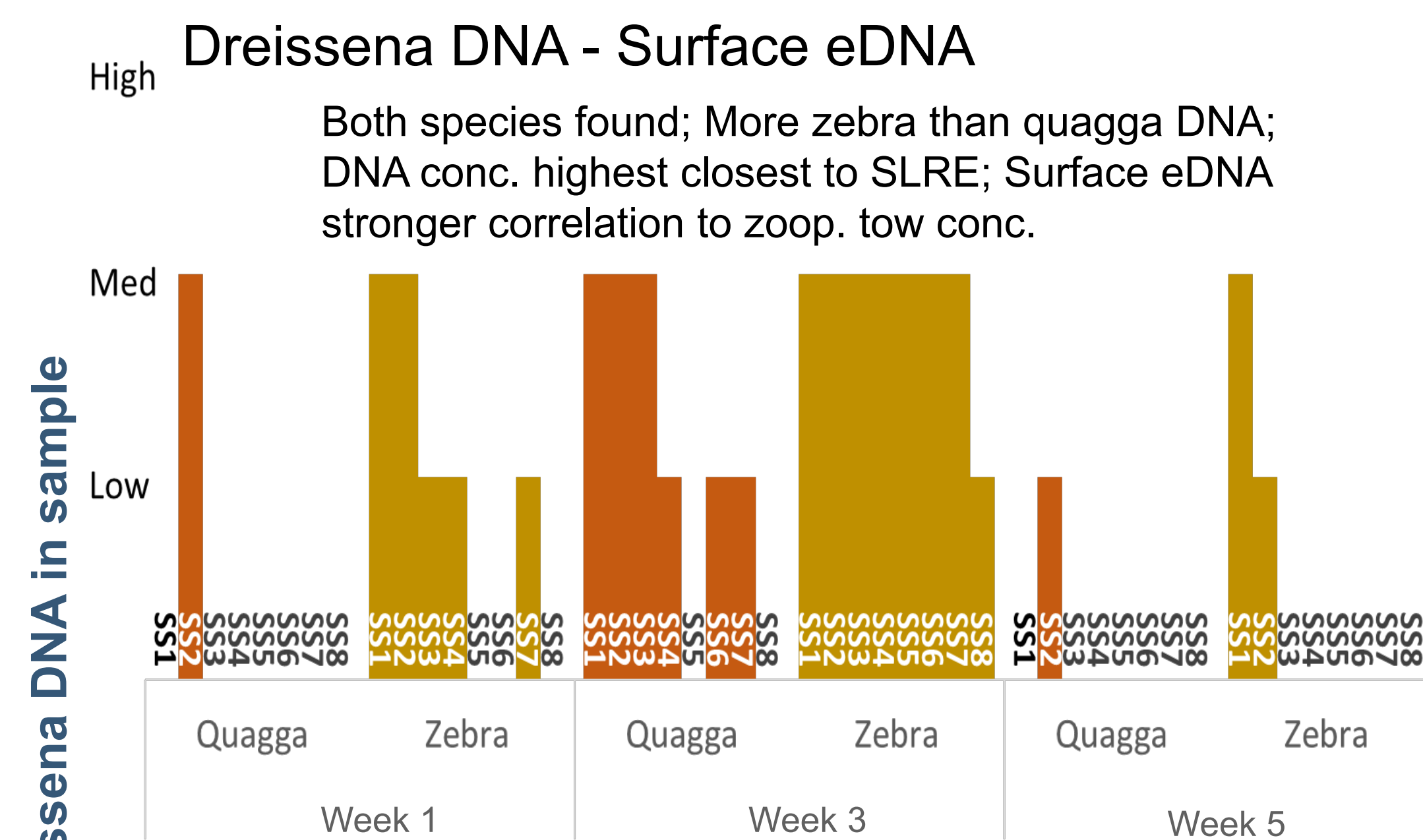
- Week 1 – SLRE to APIS
- Week 3 – SLRE to APIS
- Week 5 – APIS to SLRE



Findings (qPCR results)

PCR based DNA Concentration:

- Detection of target DNA with qPCR (45 total cycles)
- Fewer PCR cycles to detect target = higher DNA concentration in sample

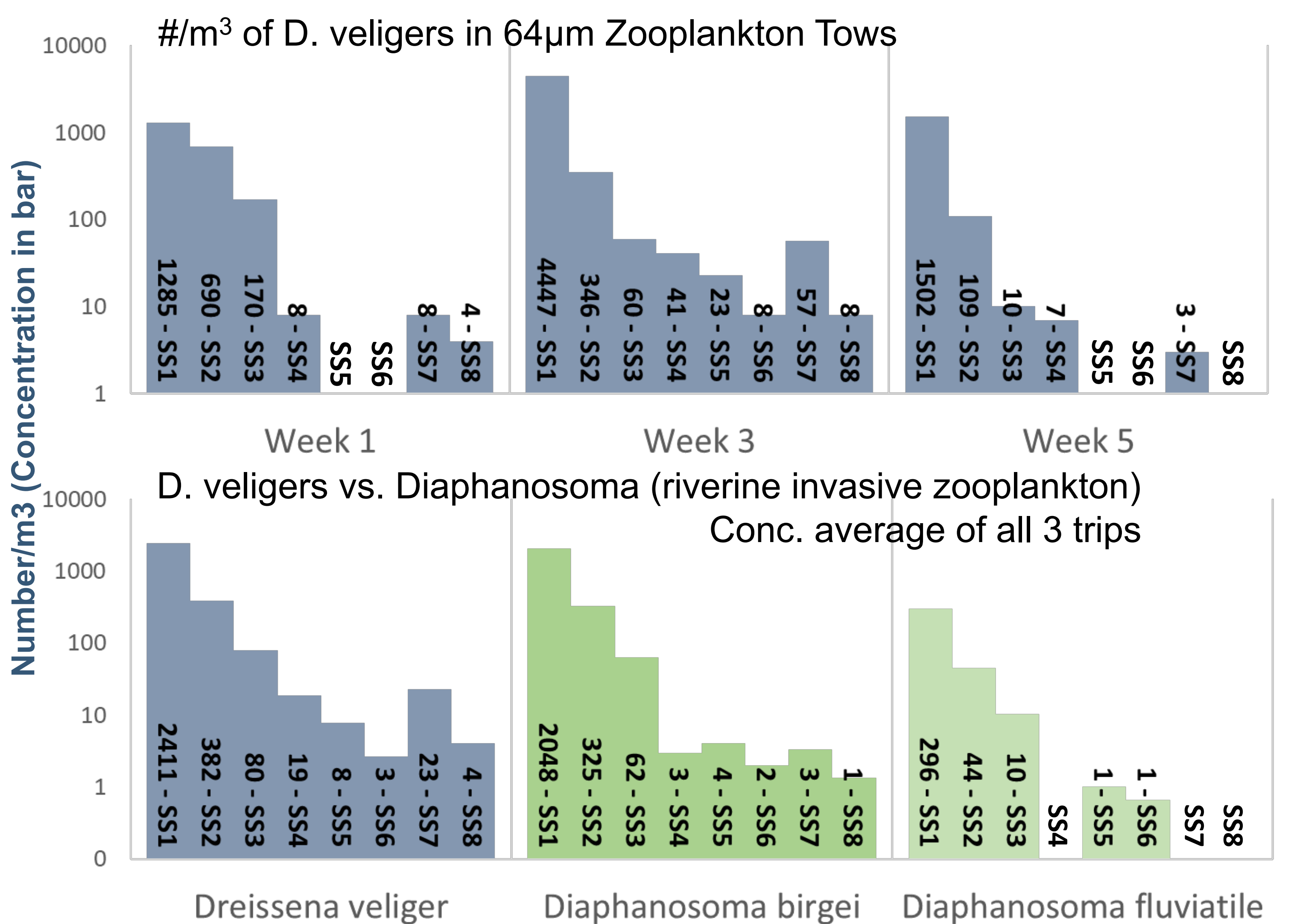


DNA concentration: High=DNA detected < 29 PCR cycles; Med=DNA detected 29-37 cycles; Low=DNA detected 38-40 cycles



Findings (Zooplankton Tows)

Veliger concentration: decreasing concentration of veligers from SLRE to APIS; Pattern similar to non-open-lake SLRE species like D. birgei



Conclusions

- Consistent decrease in D. veliger concentration along south shore in zooplankton enumeration
- Diaphanosoma species (invasive to SLRE – riverine) follow similar trends as D. veligers
- Zebra mussel eDNA found more consistently and in greater concentration w/ eDNA
- Surface eDNA more consistent with zoop. tows than bottom eDNA
- Environmental factors play large role in detection (wk5 currents & eDNA)

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