



I would like to acknowledge and thank the Yukon First Nations and Alaska ancestral lands I traveled through and collected samples from. I look forward to continue focusing on research reciprocity with the people living in these communities and sharing of

COLLEGE OF FISHERIES AND OCEAN SCIENCES

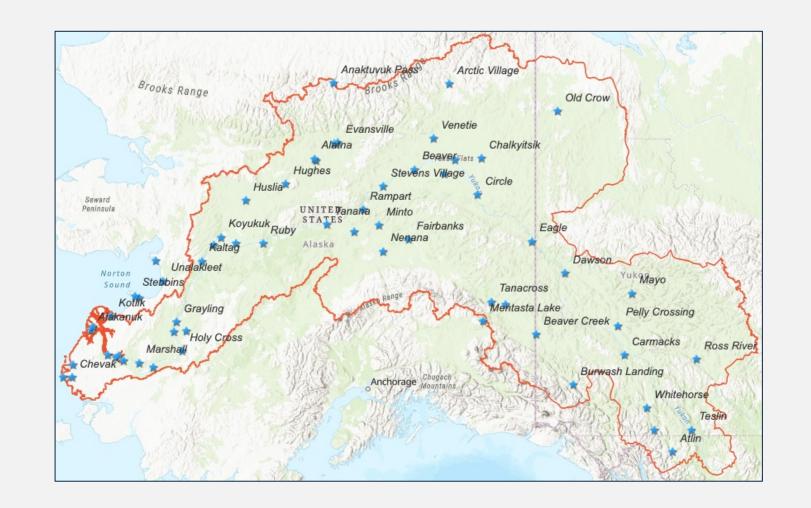
University of Alaska Fairbanks

Significance

Aquatic diversity of the Yukon River is an important indicator of watershed health in interior Alaska, USA and Yukon, CAN. Presence absence data of anadromous species may also be used to describe how Subarctic and Arctic marine environments are changing. communities in this watershed are remote and rely on anadromous and resident fish populations to support traditional food and cultural practices.

Question 1

Are eDNA field protocols effective in the interior subarctic given higher sediment loads due to glacially influenced rivers?



watershed boundary. Blue stars represent member communities of the Yukon River Inter-Tribal Watershed Council.

Below. Right, Dams in the watershed in orange. Left, Known Chinook salmon spawning sites marked by fish. Figures above and below provided by the YRITWC.



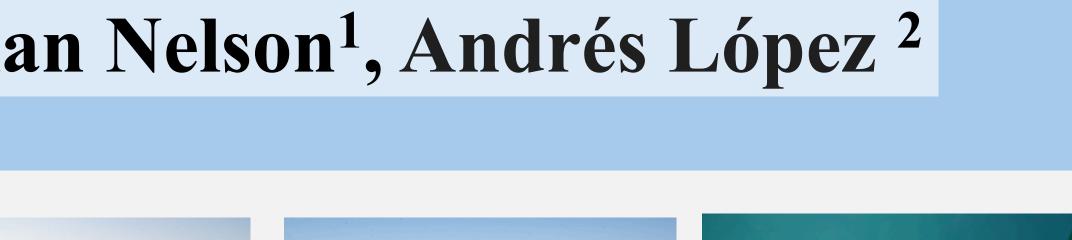


What happens on the Yukon River leaves genetic traces;

analysis of eDNA samples from a thousand-mile canoe expedition



Lillian Nelson¹, Andrés López²

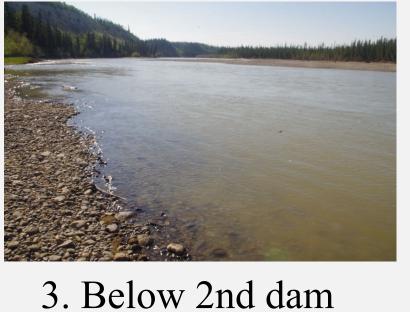




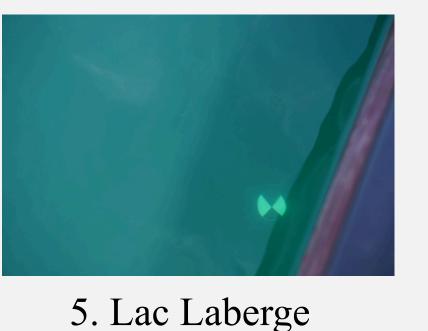
1. Marsh Lake

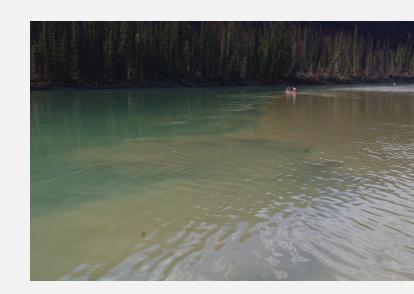


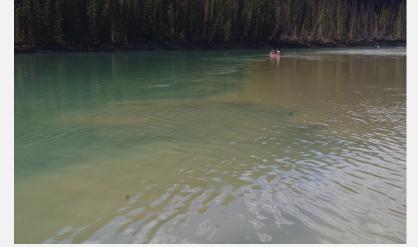
2. Below 1st dam





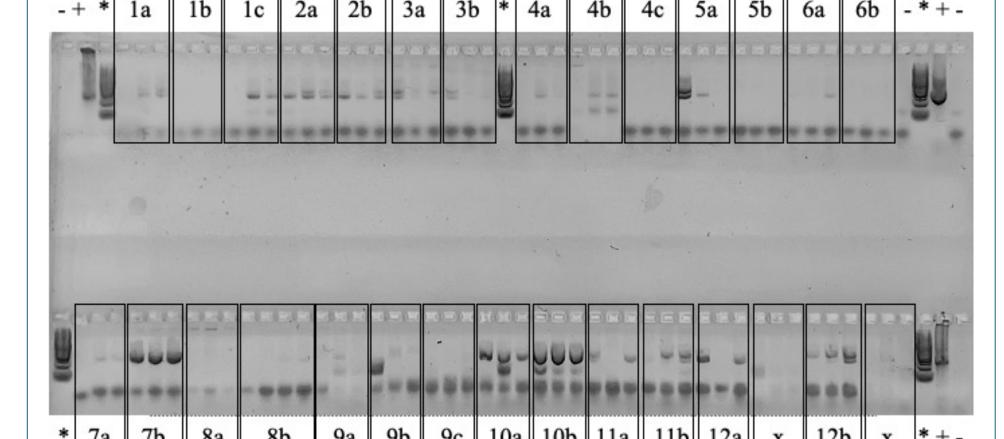


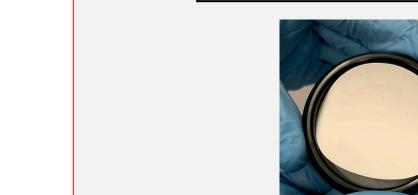




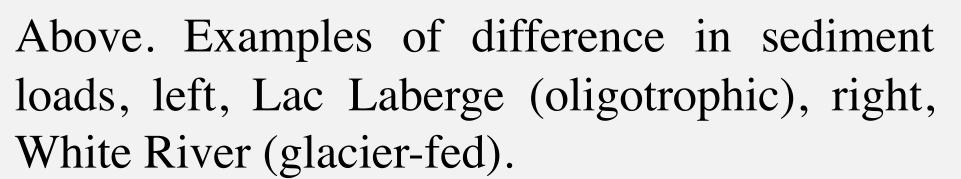


6. Teslin River









Question 2

Results

Sample Site

2. Below 1st dam, main

Below 2nd dam,

. Marsh Lake

flow of Yukon

Whitehorse

4. Takhini River

5. Lac Lagerge

6. Teslin River

8. Pelly River

9. White River

flow of Yukon

10. Charley River

11. Porcupine River

12. Mile 950, main

7. Big Salmon River

Mean | DNA | per

Sample (ng/μL)

1.75

1.16

2.80

1.09

4.53

2.62

3.05

0.59

1.08

1.34

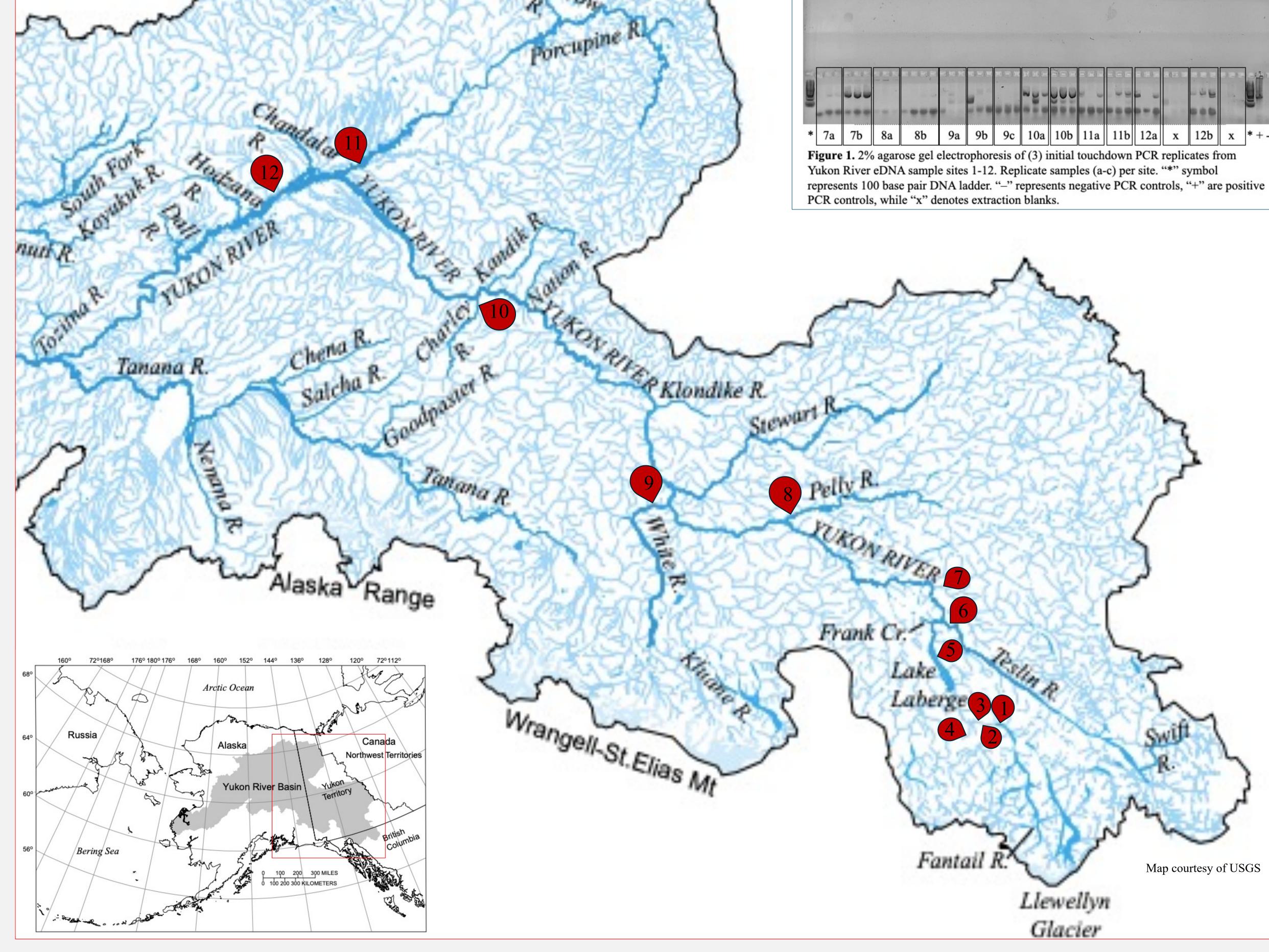
0.88

What fish species are present in the mainstream and tributaries of the Yukon River?

A continuation of this experiment is processing of extracted samples for preparation of DNA sequencing libraries targeting a short segment of mitochondrial genomes. Mitochondrial DNA sequences obtained from river samples will be compared against reference databases to generate a fish assemblage of sample sites.

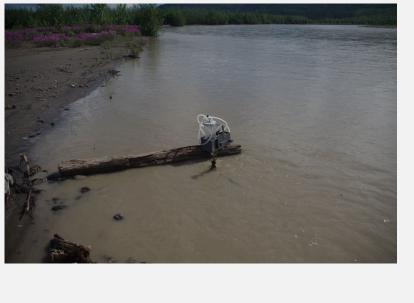
Acknowledgements

Thank you to Maggie Harings and Maris Goodwin for sharing their methods. Special thank you to Goodr and Blender for sponsoring all our sunglasses needs. Huge shout out to Upper Yukon River Expedition '22 members for a magical summer floating the river!











10. Charley River





7. Big Salmon River 8. Pelly River

9. White River

11. Porcupine River

12. Mile 950