

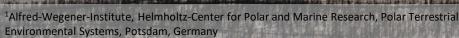
Identification and characterization of vegetation loss during the last 50,000 years in Beringia

Does the loss of the Pleistocene steppe tundra induce plant taxa loss?



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CONTEXT



- With ongoing climate change: extinction events
 - Impact all Eukaryota groups worldwide¹
 - Plants are also impacted (even in the arctic) 2 + similar proportion than mammals 3
- Pleistocene / Holocene: major climatic change
 - Loss of Pleistocene steppe-tundra⁴ + extinction event (megafauna)⁵
 - Extinction is not supposed to be group specific² + megafauna: keystone taxa⁶
 - No extinction of plant reported so far⁷ (due to methodological biases)

AIMS:

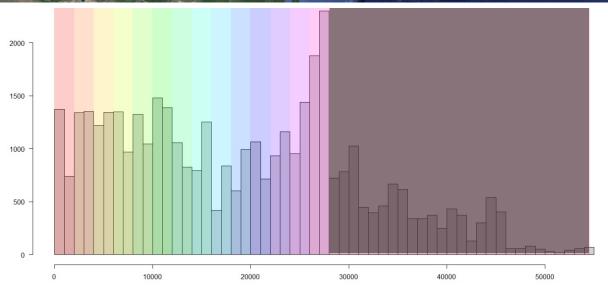
- Improve detection of rare taxa with sedaDNA proxies from lake sediments
- Identify potential extirpation and extinction events
- Characterise the potential loss

work in progress

MATERIAL





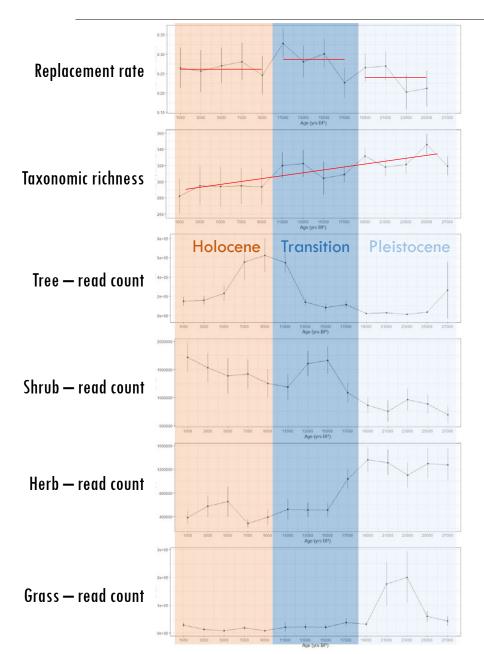


Cores	Age
Bolshoe Toko	35kyrs BP
Levinson Lessing	55kyrs BP
Ilerney 16KP	54kyrs BP
Ilerney EN18208	28kyrs BP
Bilyakh	50kyrs BP
Lama	50kyrs BP
Kyutyunda	50kyrs BP
E5	30kyrs BP
Emanda	50kyrs BP
Rauchuagytgyn	30kyrs BP

9 lake sediment cores Previous Pleistocene steppe-tundra Covering last \sim 50,000 years Investigate the last \sim 28,000 years

Work in progress MAIN RESULTS





- ~60% of plant taxa are present in every time slice -> core community
- Composition shift: Pleistocene steppe-tundra to Holocene taiga
- Steady decrease in plant richness
- Stable Pleistocene / unstable transition / less stable Holocene

121 taxa absent from modern time slice -> extirpated Average extinction rate per time slice: $0.95 \text{ E/MSY} > \text{background extinction rate } (0.05 \text{ to } 0.35 \text{ E/MSY})^{9}$



Reappearance rate

3 extinction events Last Glacial maximum Pleistocene / Holocene Mid Holocene

Match megafauna extinction events⁸



Does the loss of the Pleistocene steppe tundra induce plant taxa loss?

- Subcontinental plant taxa loss identified and quantified
- Happen at the transition to the Pleistocene to Holocene
- In parallel to steppe tundra disappearance & megafauna extinction



THANK YOU







- After DNA extraction + metabarcoding (trnL g/h) + OBITools 3 pipeline: work with all ASVs >90%
- DNA databases: modern taxa. To investigate extinction, we look at taxa not present in databases:
- 100% ASVs: assigned at 100% to modern taxon / Candidate ASVs: assigned 90%>ASV>100%: to modern taxon
- Work with ASVs with sufficient reads (>100 reads)
- From ASVs co-occurrence patterns: build communities. Assume that taxa are part of communities
- Each community has only unique taxa merge unique assignments per community to identify 474 potential taxa
- We kept most of 100% signal and reduce the candidate ASVs one with a stringent method
- Potential extinct taxa are in the candidate portion absent from the modern time slice

	Starting (min 10 samples)		>100 reads		Community > 5 ASVs			Different assignments				
Туре	Total	100%	candidates	Total	100%	candidates	Total	100%	candidates	Total	100%	candidates
Total	21977	556	21424	5302	475	4827	4957	409	4548	474	340	134
Percent		2.5	97.5		8.9	91		8.3	91.7		72	28

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