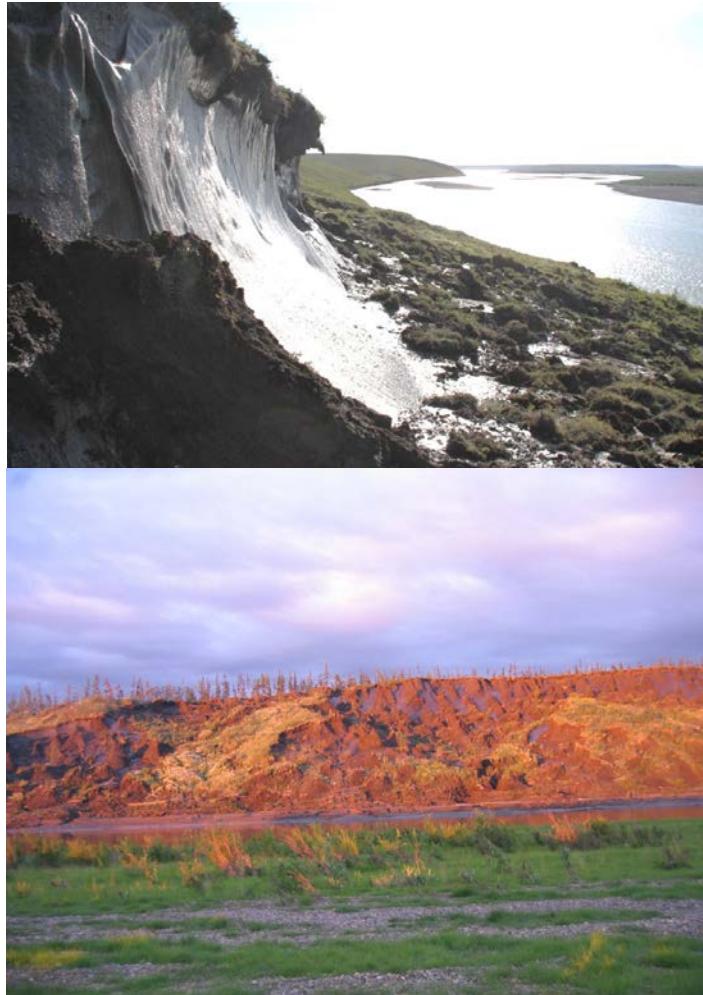


The polygenetic hypothesis of Yedoma origin – comparing grain-size distributions of Alaskan and Siberian Yedoma

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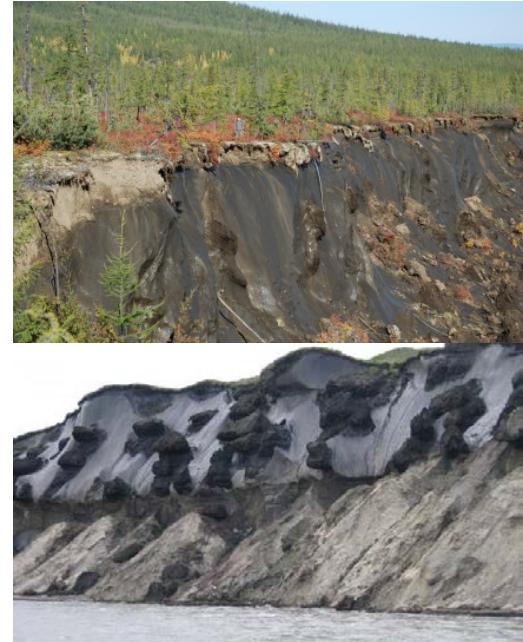
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Objectives and backgrounds

Hypothesis of Yedoma genesis

- 1) alluvial genesis,
- 2) ice-sheet-dammed basin deposits
- 3) deltaic formation
- 4) proluvial and slope deposits
- 5) marine-estuarine-lagoon formation
- 6) cryogenic-aeolian deposits
- 7) nival deposits
- 8) polygenetic origins.



Main goals of this study

- Identification and interpretation of characteristic patterns in grain-sizes distribution data sets
- find out common features and differences
- Developing a site-specific interpretation of past depositional processes to understand Yedoma formation



Locations of study sites in Siberia and Alaska



Introduction

Study sites

Methods

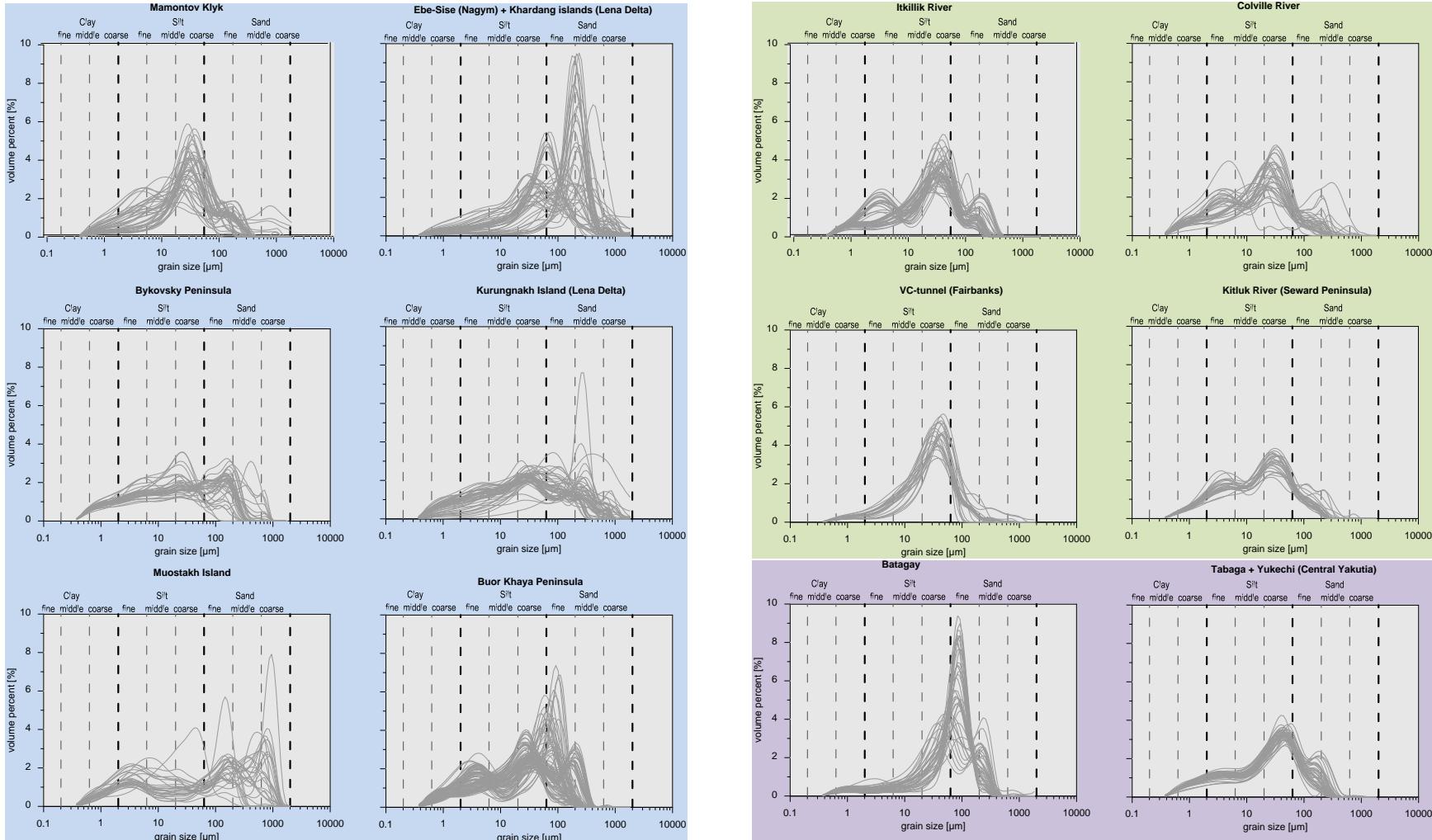
Results

Conclusions



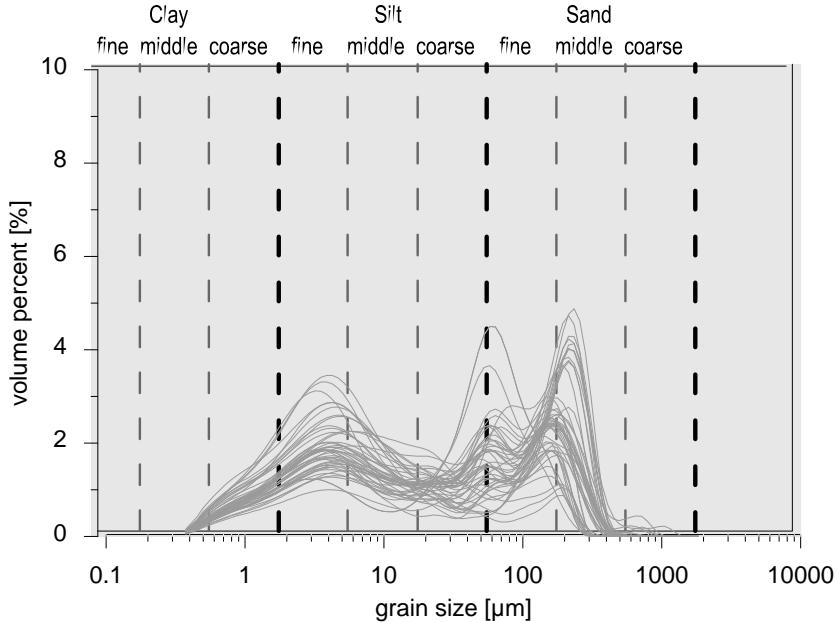
Grain size analysis and End-member modelling analysis

- Laser diffraction particle analyzer (Coulter LS 200) - 92 size channels between 0.375 and 2000 μm
- **780 samples from 17 Yedoma sites** in Alaska and Yakutia studied between 1998 and 2014
- **180 reference samples** from low centered polygons, flood plain deposits and loess deposits
- **End-member modelling analysis (EMMA)** to unmix poly-modal grain-size distributions of each site
- Into characteristic grain-size subpopulations after Dietze et al. (2012)

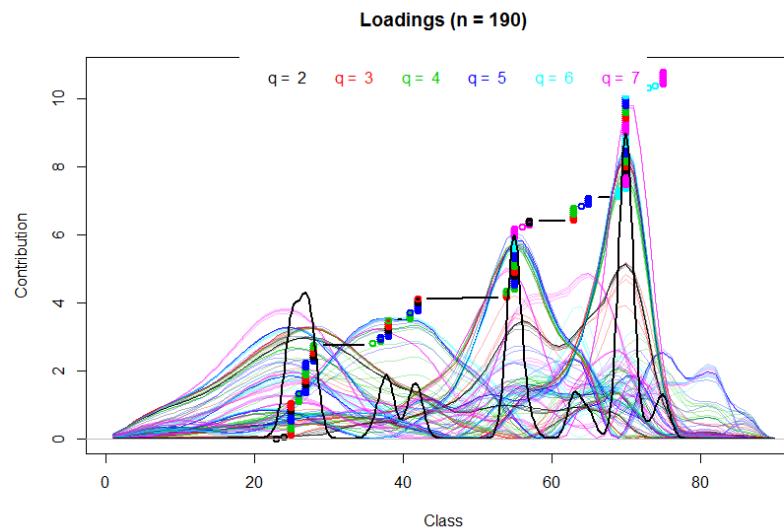


Grain size analysis and End-member modelling analysis (software package EMMAgeo in open source R)

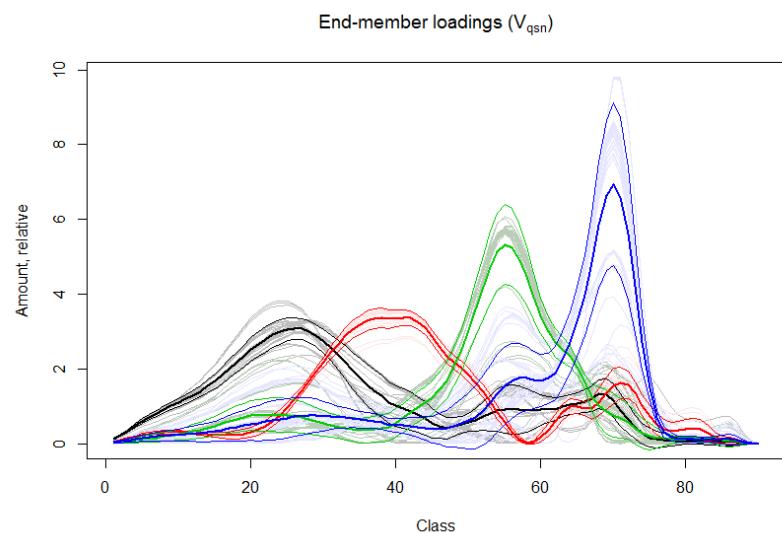
1. Measured grain size distribution



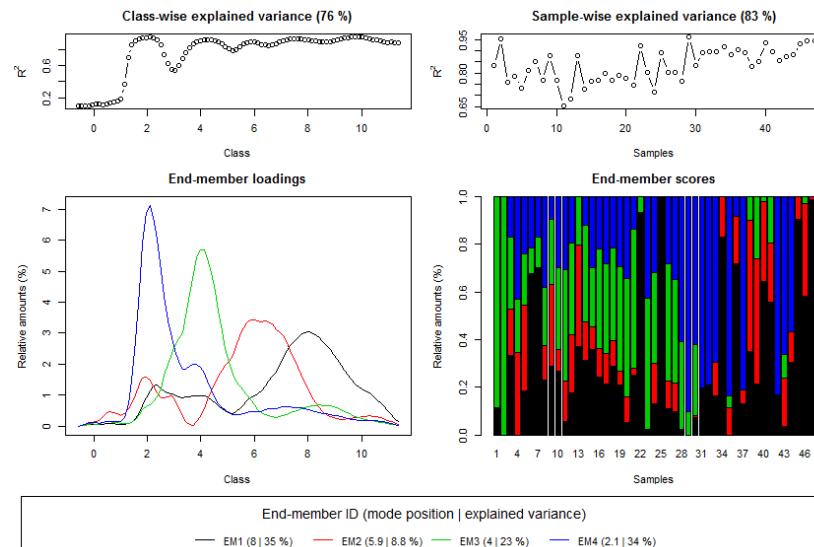
2. Identification of robust endmembers from all similarly likely endmembers (explained variance >50%)



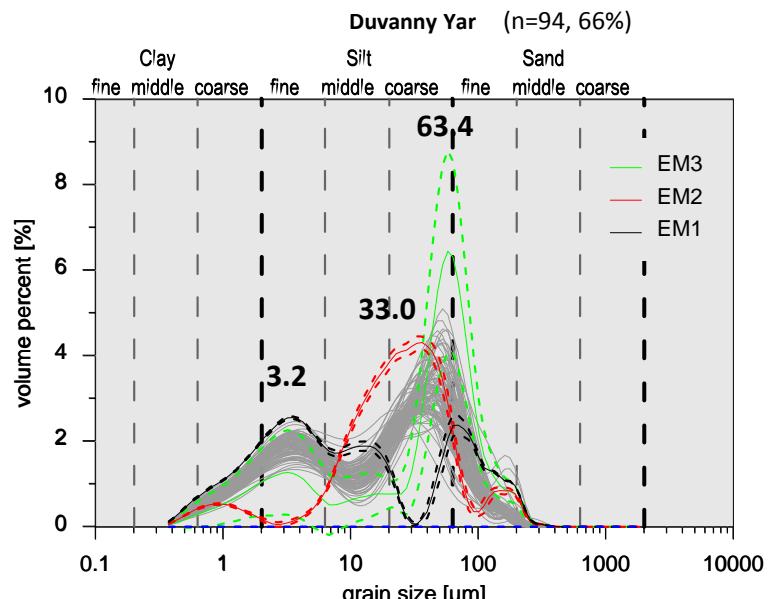
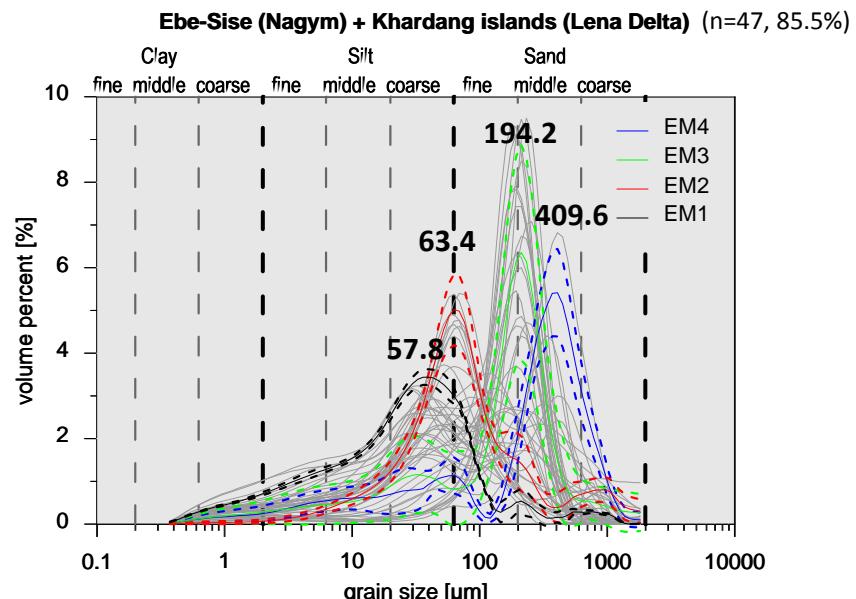
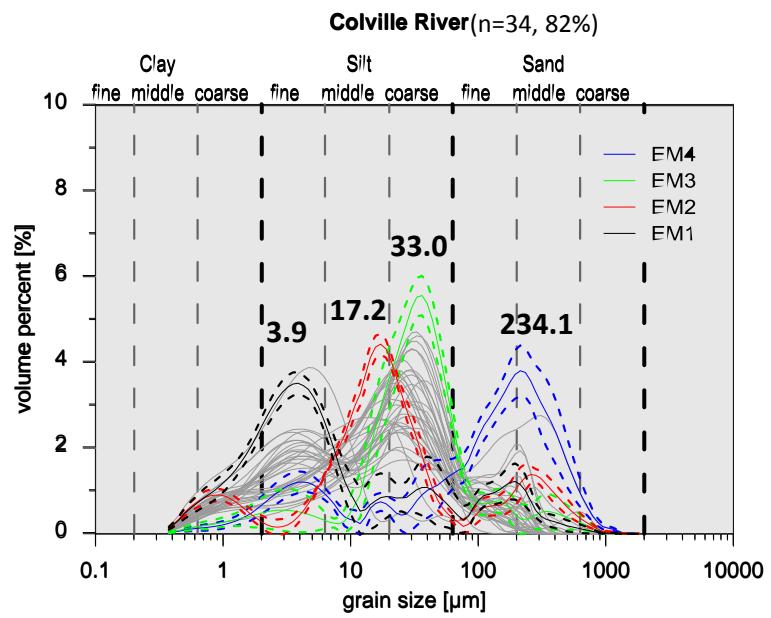
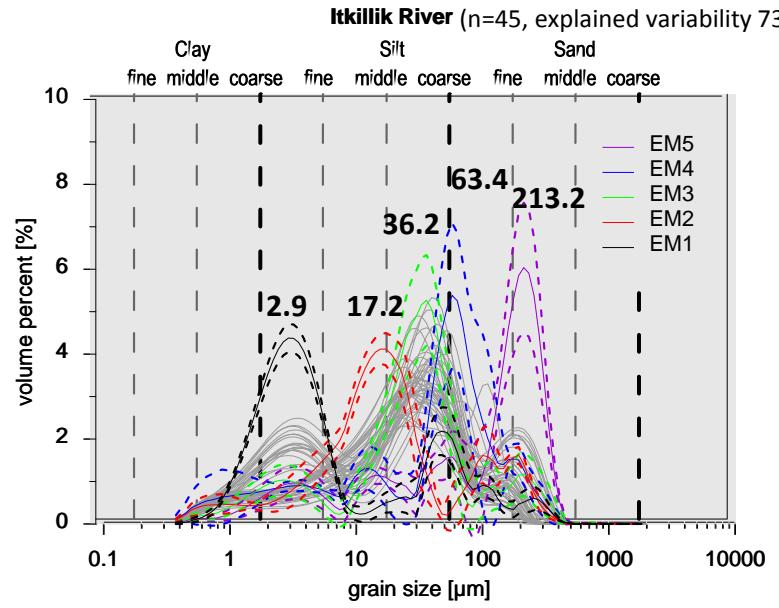
3. Mean loadings and spread of robust endmembers



4. Scores of mean robust endmembers and explained variances

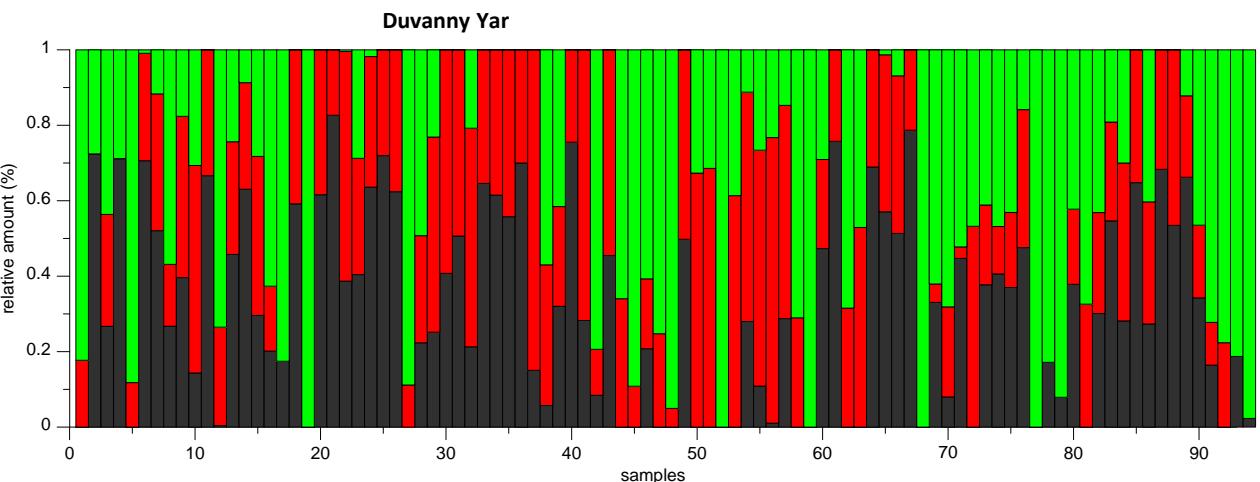
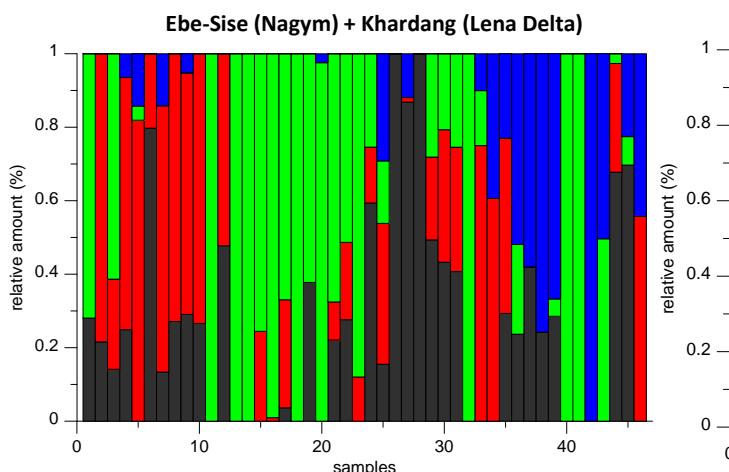
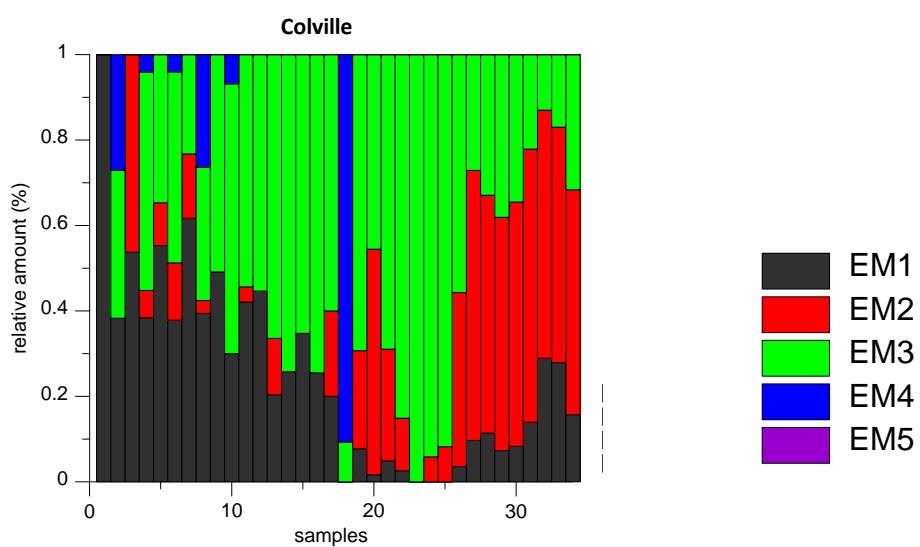
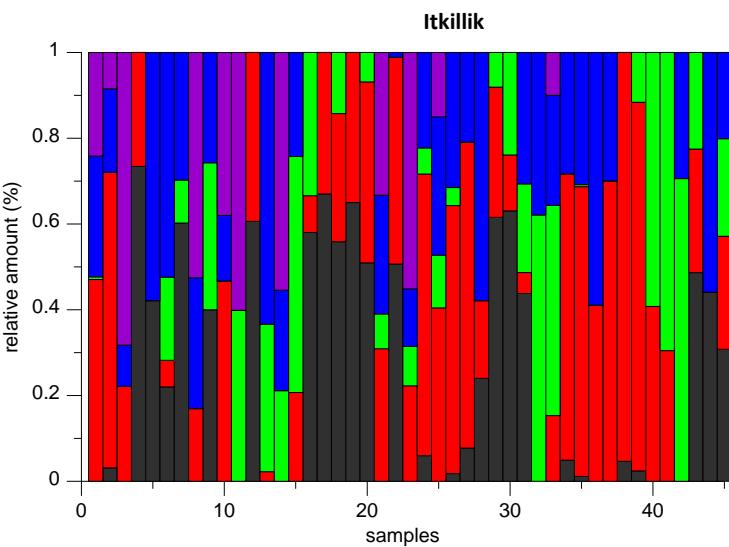


Results of robust end-member modeling (gray curves - original grain-size distributions), end-member loadings (contribution of grain-size classes to each end-member) dashed lines -standard deviation for each end-member)

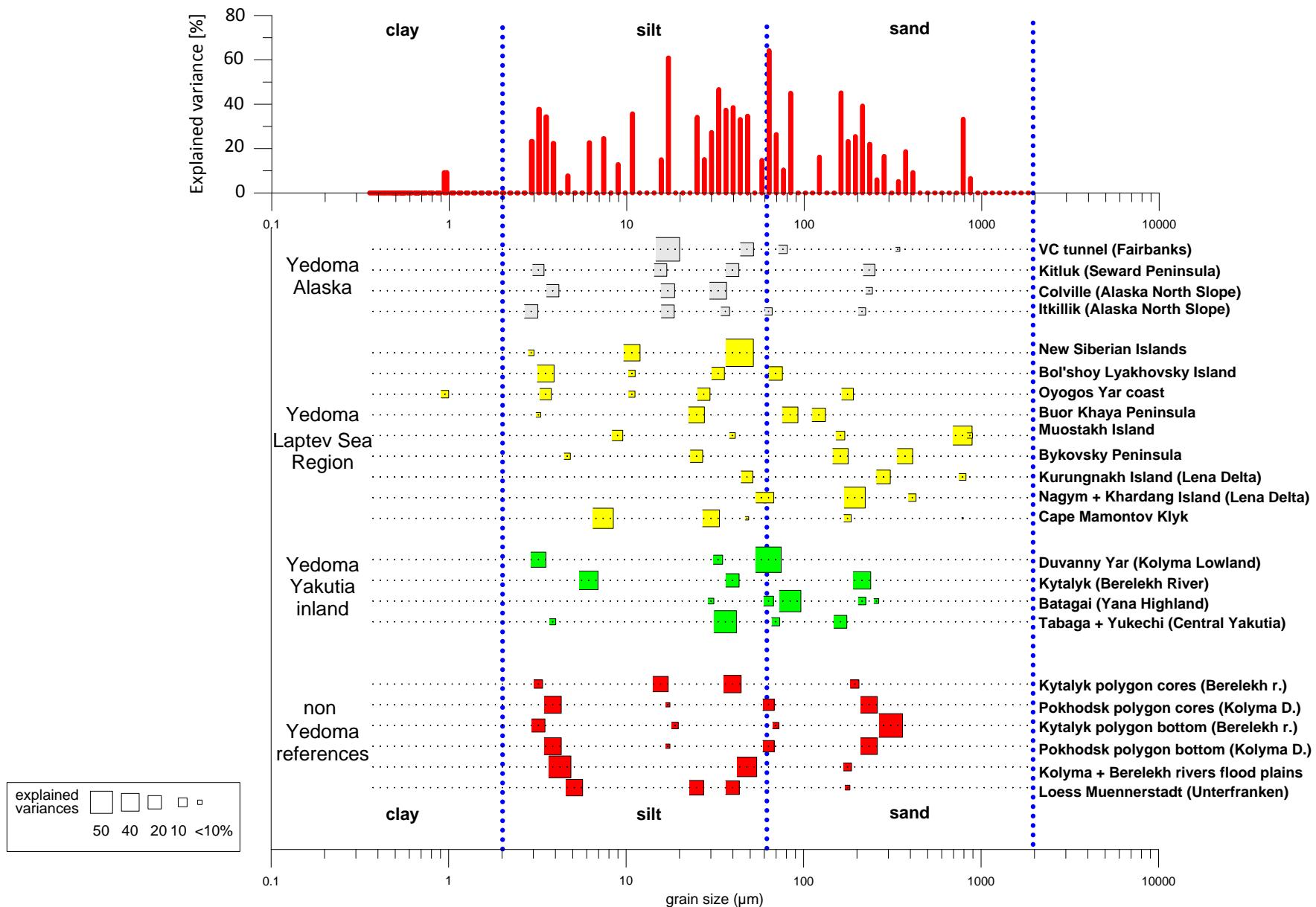


Distribution of robust end-member loadings for each sample

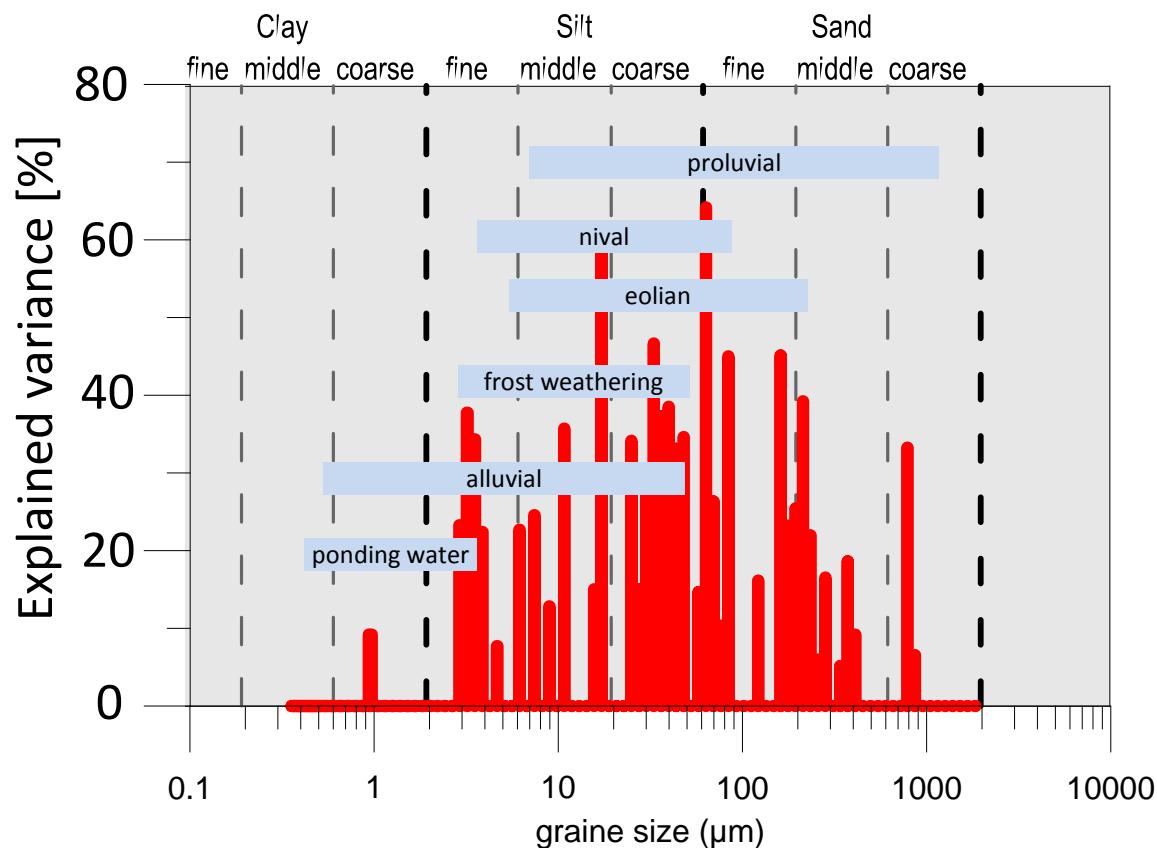
mean scores (i.e. relative contribution of an end-member to each sample)



Position of the robust endmembers and its explained variances



Expected processes contributing to Yedoma formation

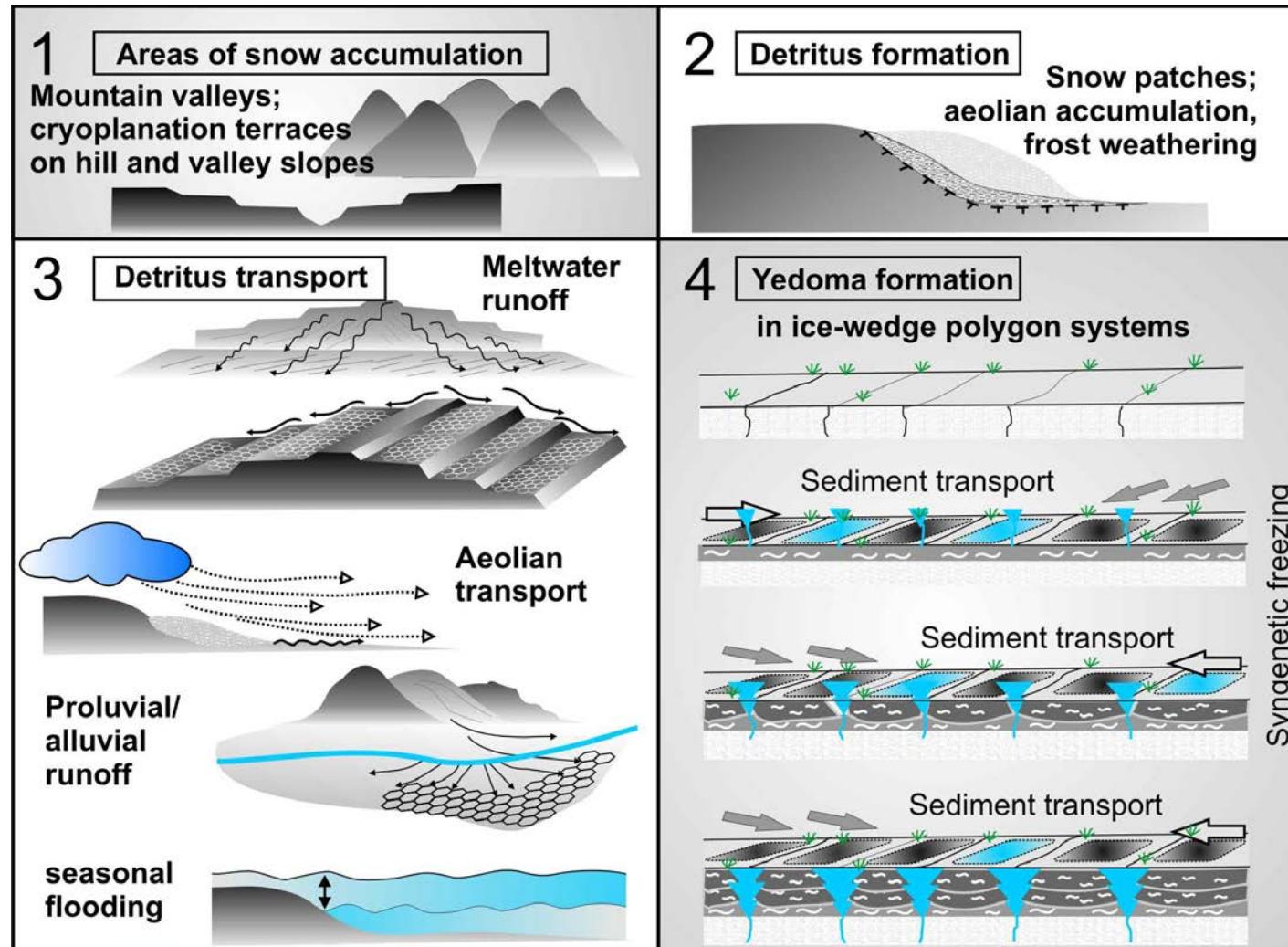


Stacked overall explained variance of robust end-members in Yedoma deposits (n=780, 17 sites)



- **Clay:** Ponding water
- **Silt:** Eolian, alluvial (main mode at 10 μm , range 0.3-30 μm , Frost weathering: Schwamborn et al. (2012) Formation of up to 10 % of <63 μm fraction after 100-150 freeze-thaw cycles of fine-sand samples (63-125 μm)
- **Sand:** Running water (i.e. melt water runoff, braided streams), Mass movements (i.e. cryoturbation, solifluction)

The cryolithogenic concept of polygenetic Yedoma Formation



Stages of Yedoma genesis based on a cryolithogenic concept. Note: If the re-transportation of loess (also called secondary loess) is included in the loess concept, the loess and the polygenetic concepts are very similar.