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Suberin and Cutin as biomarkers for shifts in plant community composition following land use changes on the Tibetan Plateau

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Background

There is still a debate if the Kobresia biome on the Tibetan Plateau is human induced or climate driven. Grazing exclosure experiments show a fast replacement of Cyperaceae by Poaceae. This points to a zoo-anthropogenic influenced plant composition. We aimed to find biomarker to distinguish between different grasses and herbs dominating under different conditions.

Why Suberin and Cutin?

-Material

- Plants and soils from Kobresia dominated grasslands in a montane (near Xinghai) and an alpine (north of Lhasa) area
- 20 x 20 m fences inside Yak pastures, fenced 1997 (Reting) and 2002 (Xinghai).
- Monitoring of plant community composition inside exclosures and adjacent grazed area to indicate changes in vegetation structure after fencing



Suberin and cutin signatures are used as biomarkers for determining soil organic matter (SOM) sources.

Previous studies succeeded in differentiating between closely related tree species, e.g. fir (*Abies alba*), spruce (*Picea abies*) and douglas fir (*Pseudotsuga menziesii*) (Spielvogel 2010). However, signatures of different grasses and herbs have not been analyzed in detail, and it is unknown if they are identifiable in soil.

Suberin and cutin...

- ... are ubiquitous in plants.
- ... are important components of hydrophobic layers in plant cell walls.
- ... play in an important role as barriers controlling gas-, water-, and nutrient transport in plants.
- ...are characteristic for roots (suberin) and leaves (cutin), respectively.





- Identification of characteristic plants for both treatments
- Analysis of root and shoot samples from indicator plants: among others Kobresia pygmaea (Cyperaceae, grazed), Leymus spec. (Poaceae, ungrazed)
- Analysis of soil samples taken inside and outside the grazing excluding fences

Fig 5: Distribution area of *Kobresia pygmaea* and study sides ——



Fig 6: Leymus and Kobresia pygmaea





Fig 3: Example for suberin and cutin composition, respectively

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Polyhydroxy-fatty acids

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Fig 4: Examples for common suberin and cutin monomers

Method

Samples (5-20 g, depending on carbon content)

- 1. Extraction (double deionized water) ------ water soluble polar compounds
- 2. Extraction CH_2CI_2 , MeOH \longrightarrow "free lipids"

Alkaline hydrolysis:

Teflon-lined bombs 100 C/3h

(1 M methanolic KOH)

Filtration, Residue:

Addition of internal recovery standard

Fig 7: Chromatograms of Kobresia (Cyperaceae) and Leymus (Poaceae) – shoots and roots

- Chromatograms of plant materials show clear differences between grass species
 Signatures of roots and shoots are also different
- The suberin signature of Kobresia roots is characterized by several long-chain fatty acids > 26 C-atoms that are missing in Leymus roots

Fig 8: Chromatograms of soils sampled inside and outside the grazing excluding fence (montane area).

• Depth increment 0-5 cm (= dense root layer) mirrors root pattern

 Fatty acid signature of soils from areas of different altitude but similar vegetation did not differ

- The long-chain fatty acids signature typical for Kobresia roots decreased distinctly seven years after fencing
- Di-carboxylic acids seemed to have longer turnover times than ω -Hydroxy fatty acids

Conclusion

Hydrolysable aliphatic lipids derived from suberin and cutin are well suitable to distinguish between different grasses and herbs indicating diverse grazing pressure. Cutin and suberin are compounds with a high diagnostic value for vegetation history of grasslands due to the preservation of species specific long-chain aliphatic lipids.

Reference

Pollard M., Beisson F, Li Y., Ohlrogge J.B. (2008). Building lipid barriers: biosynthesis of cutin and suberin. Trends in Plant Science Vol 13 No. 5. Spielvogel S., Prietzel J., Kögel-Knabner I. (2010). Lignin phenols and cutin- and suberin-derived aliphatic monomers as biomarkers for stand history, SOM source, and turn over. Geochimica et Cosmochimica Acta 74, Suppl. 1, A983-A983.

