

## **NOTES** on vegetation - **TrophCost**



## Removed observations:

- Plants from the BEs matrix that were not identified up to genus level were removed (i.e. those only until family level):
  - Brassicaceae\_sp, Asteraceae\_sp, Caryophyllaceae\_sp,
     Orchidaceae\_sp
- "Baumkeimling sp" observation
- "Unknown" observation
- Tree species:
  - Acer sp
  - Betula pendula (Hänge-Birke)
  - Carpinus betulus (Heinbuche)
  - Fraxinus excelsior (Gewöhnliche Esche)
  - Pinus sylvestris (Wald-Kiefer, Föhre, Forle)
  - Populus\_tremula (Zitter-Pappel, Espe)
  - Prunus\_avium (Süßkirsche)
  - Prunus sp
  - Prunus\_spinosa (Schlehdorn)
  - Quercus robur (Stiel-Eiche)
  - Tilia sp (Linden)
- Shrubs species:
  - Juniperus communis (Gewöhnlicher Wacholder)
  - Crataegus\_sp (Weißdorne)
- Fern species:
  - Ophioglossum vulgatum (Gewöhnliche Natternzunge)



Medicago × varia (Bastard-Luzerne) is a hybrid between the two sister species Medicago sativa (DE=Luzerne, EN=Alfalfa) and Medicago falcata (DE=Sichelklee, EN=sickle clover). Additionally, M. falcata is very rare in BB, especially in the Oder region (Odergebiet); mostly M. falcata x M. sativa (Richert & Brauner, 2018). Moreover, Medicago x varia is synonym of:

- Medicago sativa subsp. hemicycla (Grossh.) C.R.Gunn
- Medicago sativa var. pauciflora (Ledeb.) Urb.
- Medicago sativa subsp. varia (Martyn) Arcang.
- Medicago sativa var. varia (Martyn) Urb.)

Therefore, for practical purpose, Medicago x varia was consider synonym of Medicago sativa aggr.

Only in BW (Ebert et al. 2005) both plants were listed.

Whenever an interaction was repeated between Medicago x varia and Medicago sativa aggr., the highest interaction strength was consider. In BB (Richert & Brauner 2018) only Medicago x varia was listed.



if Potentilla tabernaemontani = P. verna (http://powo.science.kew.org) and P. neumanniana = P. tabernaemontani = Frühlings Fingerkraut (Ebert 2005, BW)

and P. neumanniana = Frühlings Fingerkraut (Richert & Brauner 2018)

then P. neumanniana = P. verna = Frühlings Fingerkraut

(https://www.floraweb.de/xsgl/artenhome.xsgl?suchnr=20053&)



Valeriana officinalis includes all following subspecies and aggregations that appear in both books. \*\*\*Whenever an interaction was repeated between any of the plants, the highest interaction strength was consider.

- (Ebert 2005, BW):
  - Valeriana officinalis ssp. officinalis (Echter Arznei-Baldrian)
  - Valeriana officinalis ssp. excelsa (Syn. = Valeriana procurrens;
     Kriechender Arznei-Baldrian)
  - Valeriana officinalis ssp. tenuifolia (Syn. = Valeriana wallrothi,
     Valeriana pratensis; Schmalblättriger Arznei-Baldrian)
  - Valeriana officinalis agg. (Arznei-Baldrian)
- And (Richert & Brauner 2018, BB):
  - Valeriana officinalis s.l. (Arznei-Baldrian)



Vicia sativa aggr. included all subspecies of V. sativa i.e:

- Vicia sativa (Futter-Wicke, Saat-Wicke)
- Vicia sativa ssp. nigra = Vicia angustifolia (Schmalblättrige Wicke; Richert & Brauner 2018, BB)
- Whenever a species interaction was repeated between any of the plants subspecies, the highest interaction strength was consider.
- For red list data, whenever two or more subsp. of a plant where present, the worst i.e. most endangered score was considered and a note was added to comment column.
- Taraxacum officinale auct. = Taraxacum sect. Ruderalia in Richert & Brauner ⇒
  should we consider it as Taraxacum sp. ????????? ⇒ ask Klaus! ♥ TODO



Conflicting cases in Richert und Brauner: does not have \* to \*\*\*, they are missing, e.g:

- Echinops banaticus in Maniola jurtina
- Vicia spec. in Polyommatus coridon
  - >>> Consider them as 0.2 (\*)



To check manually the species which are only identified up to family level do the following:

- 1) in the file interactions unmatched.csv
- add filters to every column
- highlight in red all genes in the plant\_genus column that are identified until family levels in the BE's list
- highlight in red all species in the plant\_species column which are spec. or sp.
- 2) in the file trophic-link-matrix
- for each butterfly species, search and check each red column.
- fill in with 0.1s

