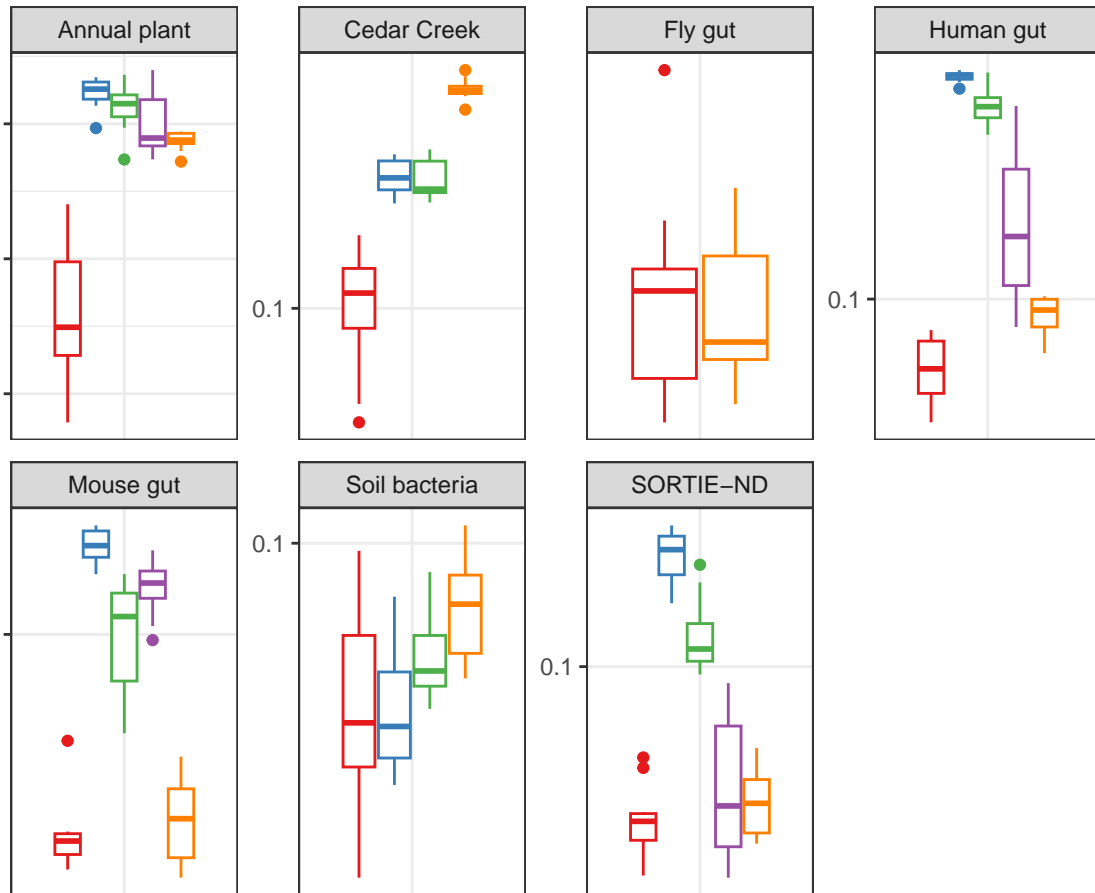


Mean absolute abundance error
scaled by 95% quantile abundance in dataset



Experimental design

- Mixed
- Doublets only
- Doublets + triplets only
- 1-dropouts + 2-dropouts only
- Singlets + 1-dropouts, then mixed

Dataset