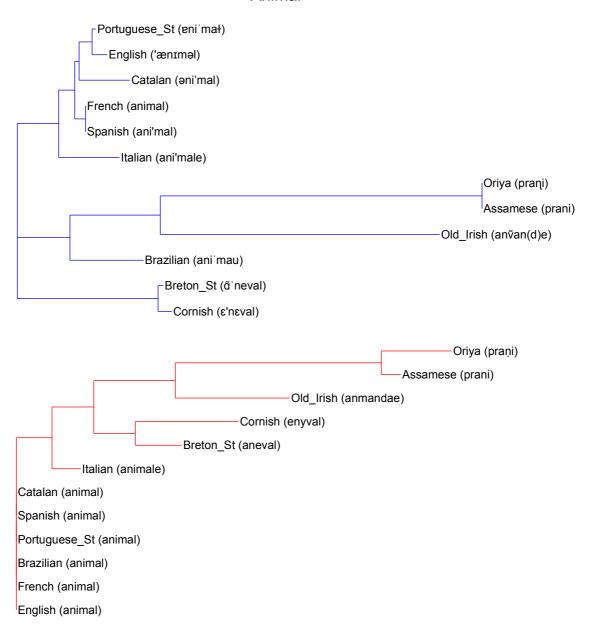
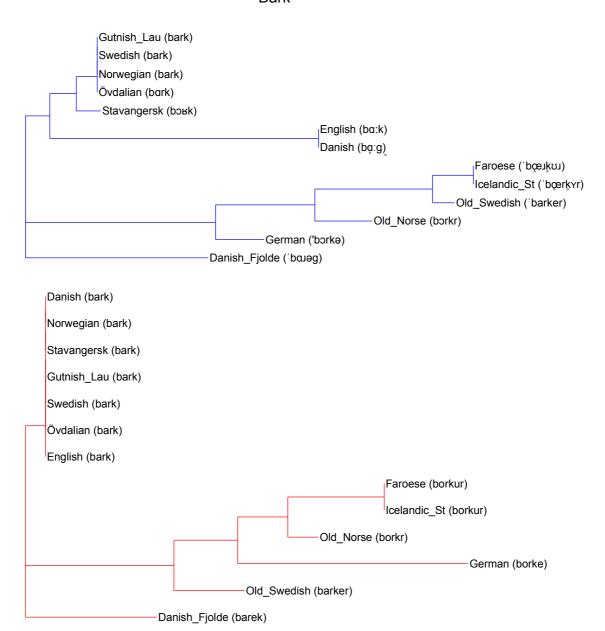
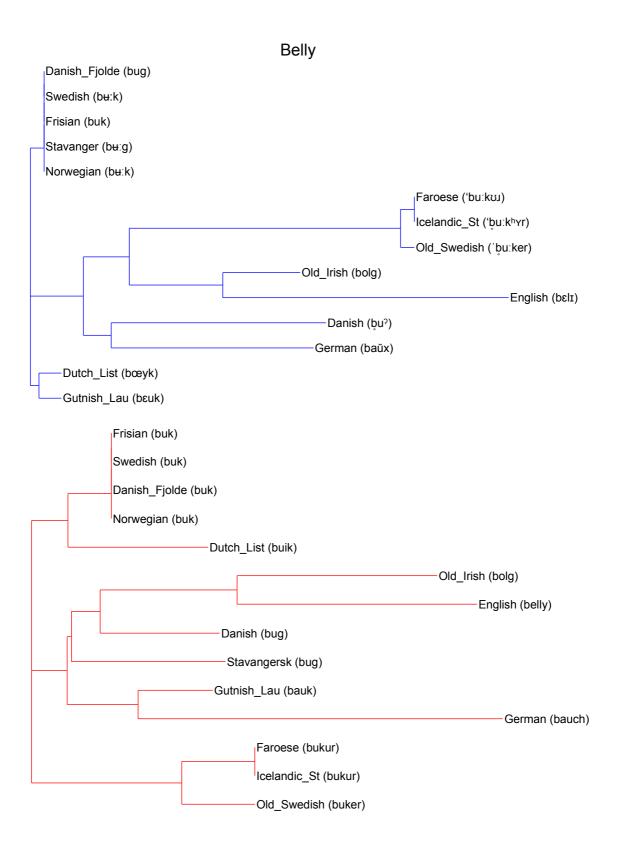
Phylogenetic trees obtained using the SCA phonetic distance (List, 2012) are in blue and using the normalized Levenshtein distance are in red. The Neighbor-Joining (Saitou et Nei, 1987) algorithm was applied to infer the trees from the distance matrices. All the data have been taken from the IELex database (http://ielex.mpi.nl/) (Dunn *et al.* 2011).



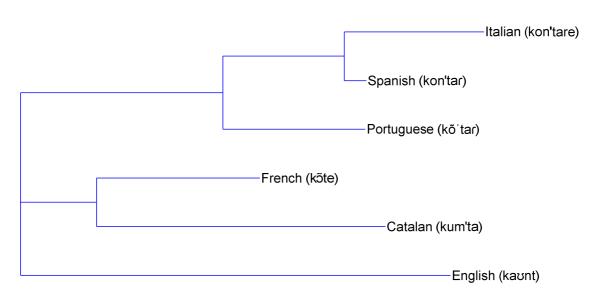


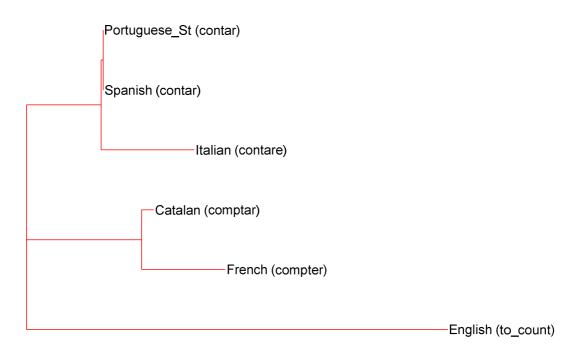
#### Bark

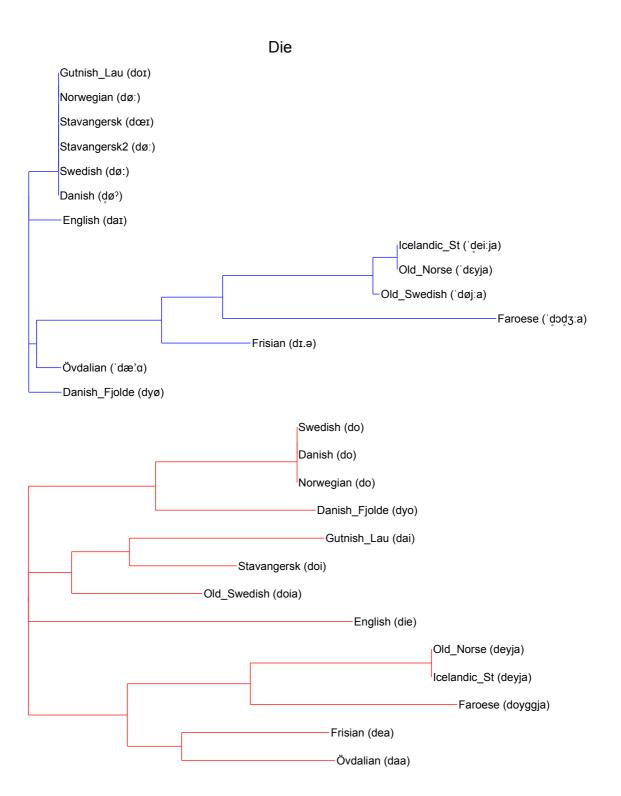




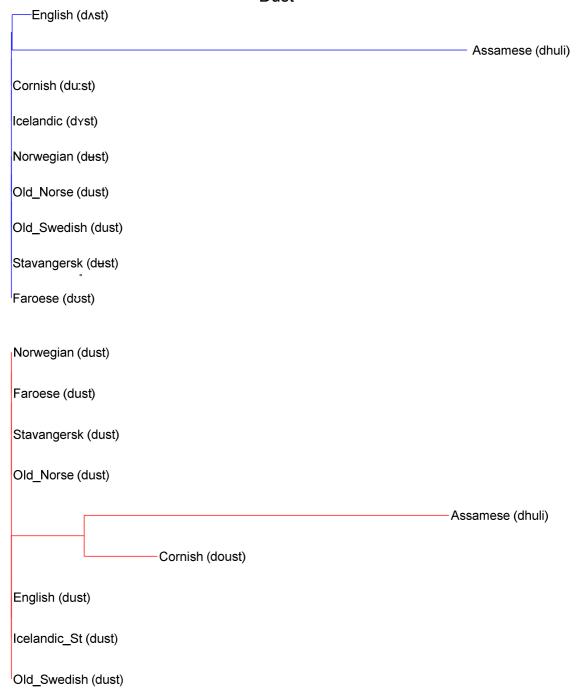


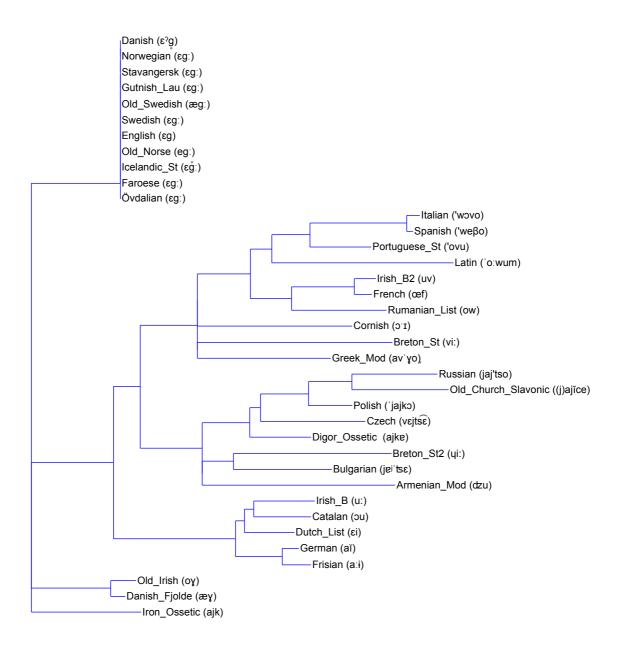


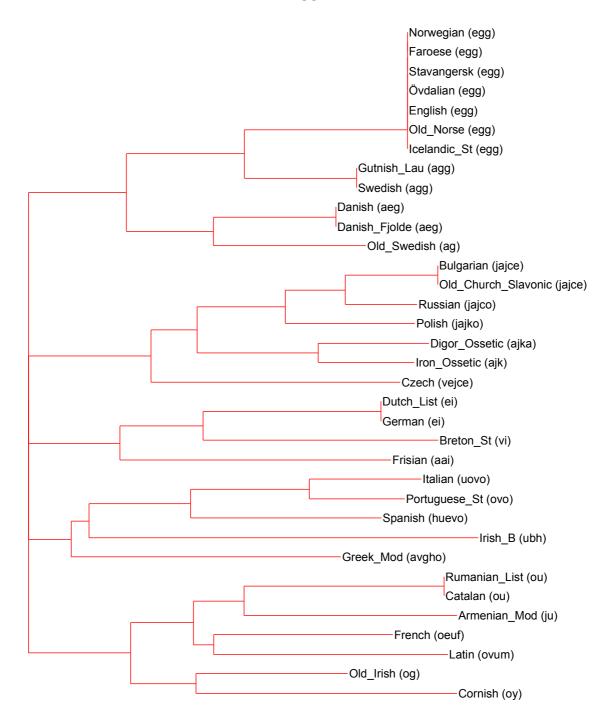


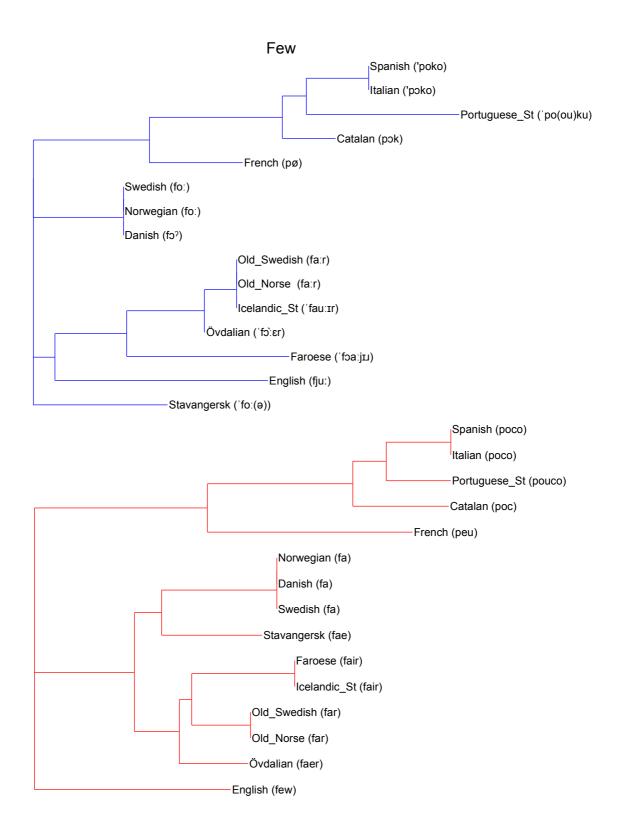


#### Dust

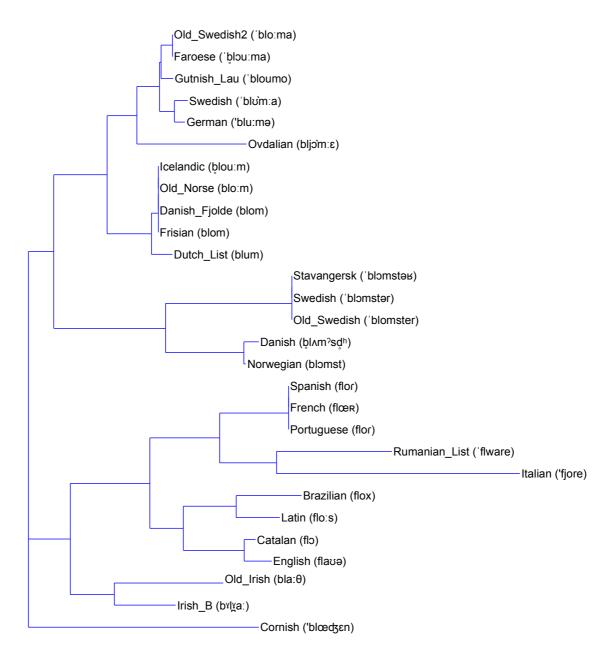


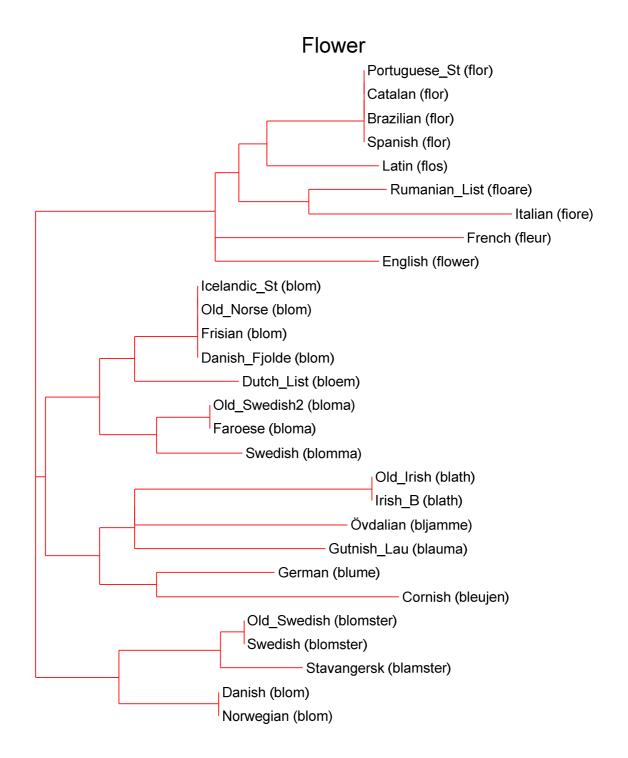




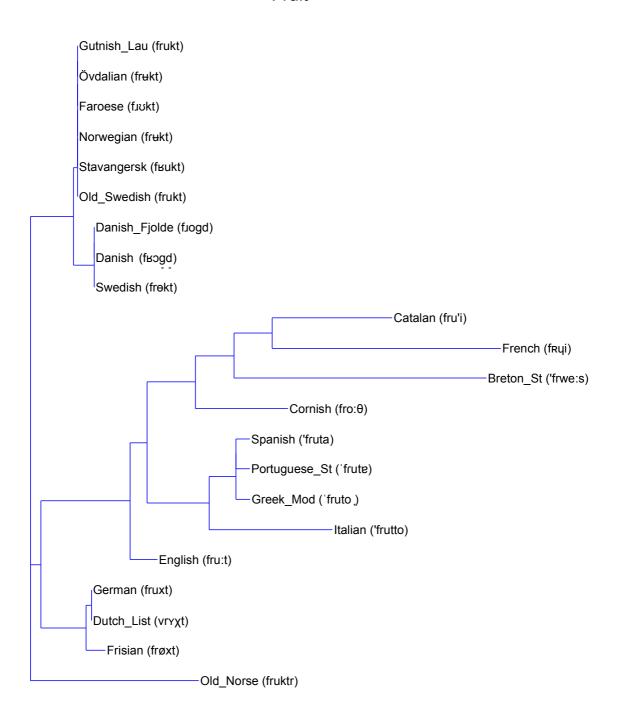


#### Flower

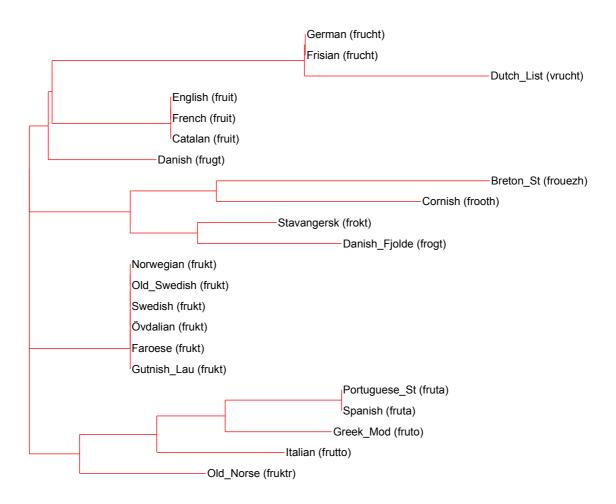




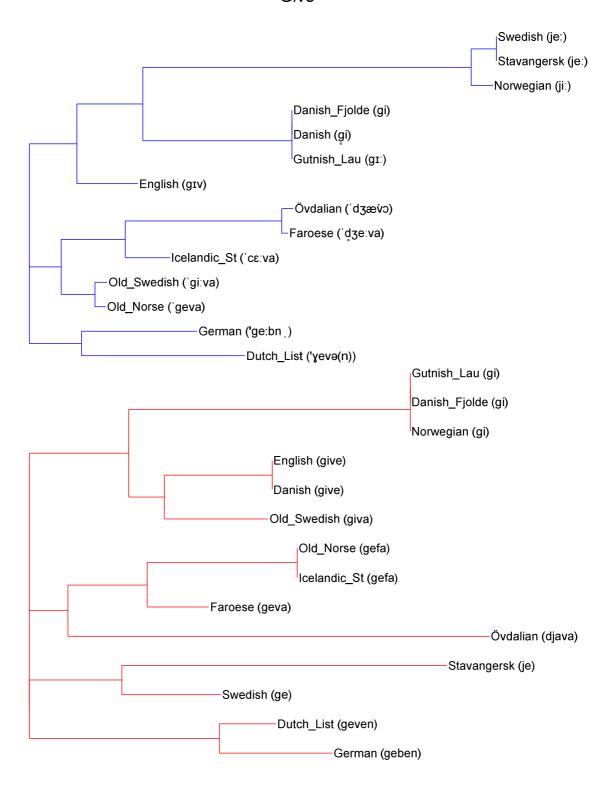
### Fruit



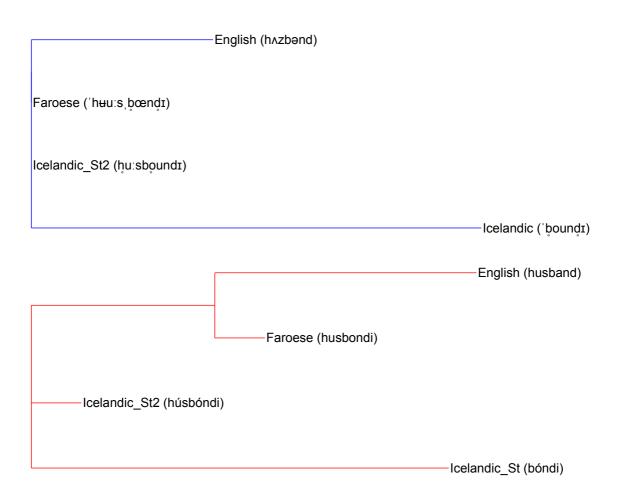
Fruit



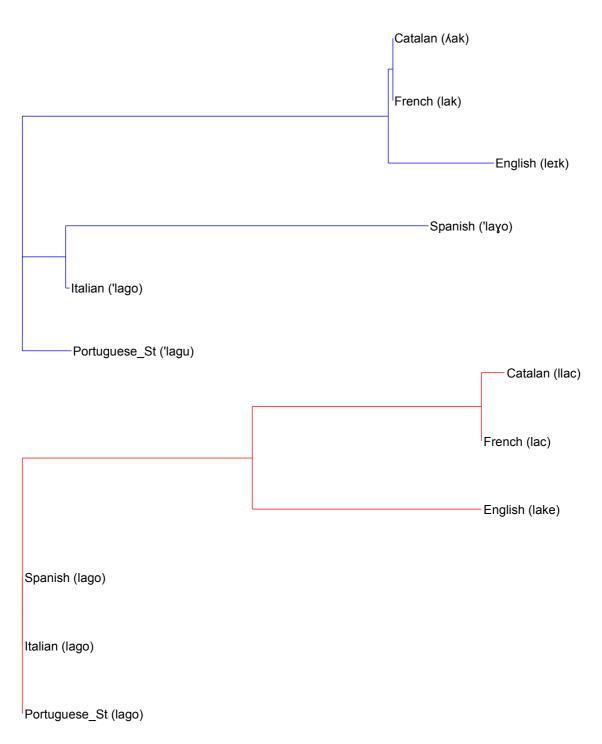
#### Give



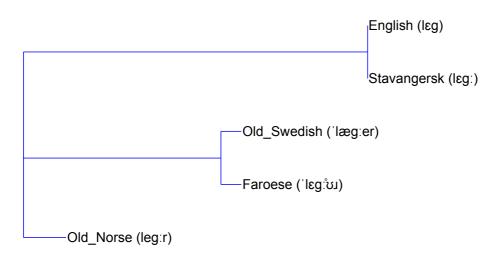
### Husband

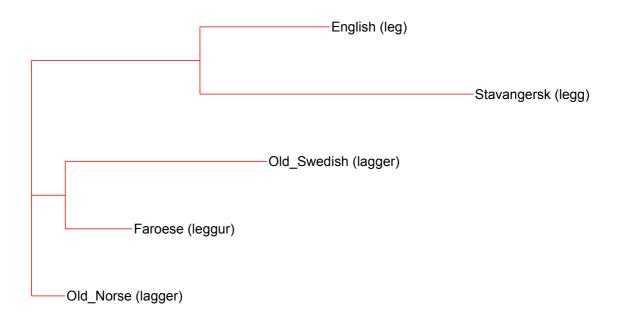


### Lake

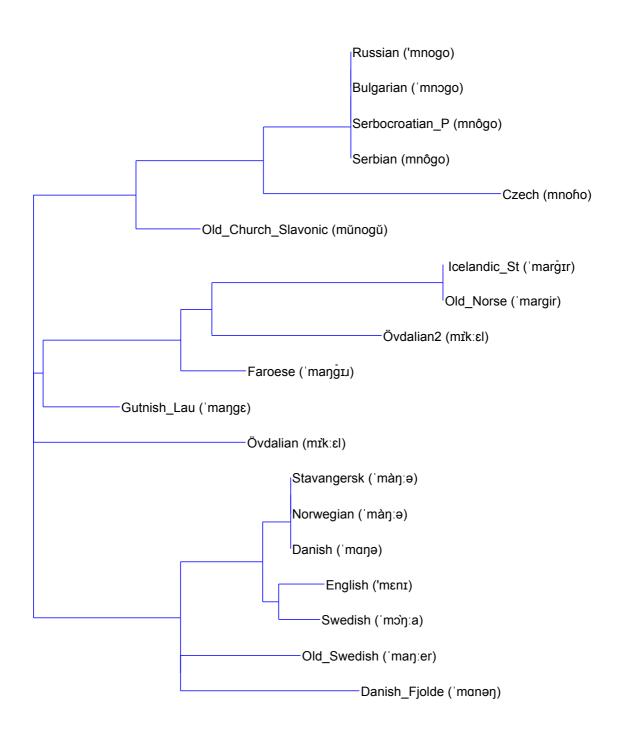




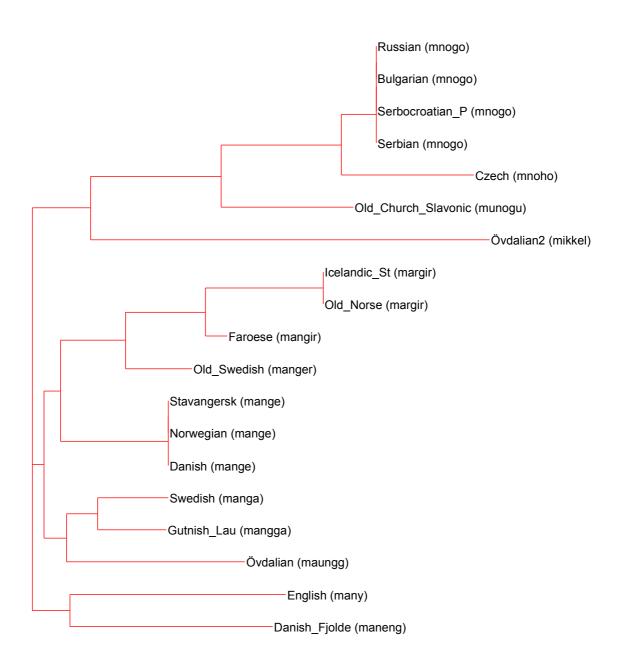




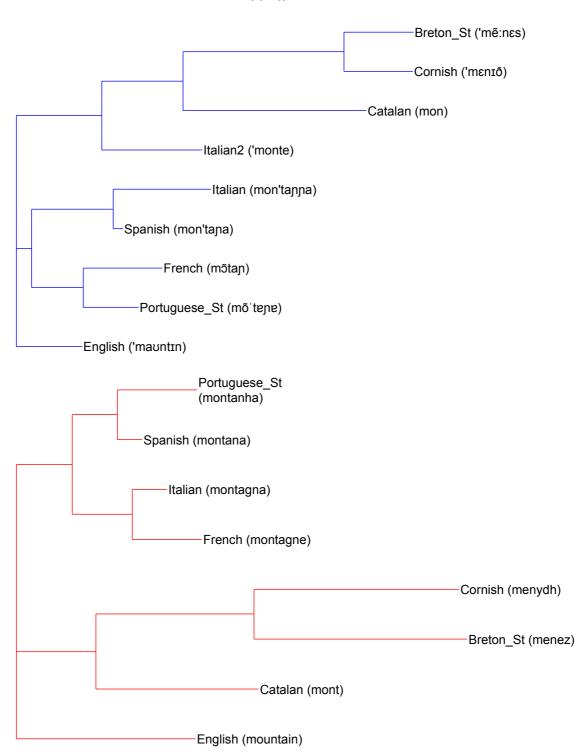
# Many

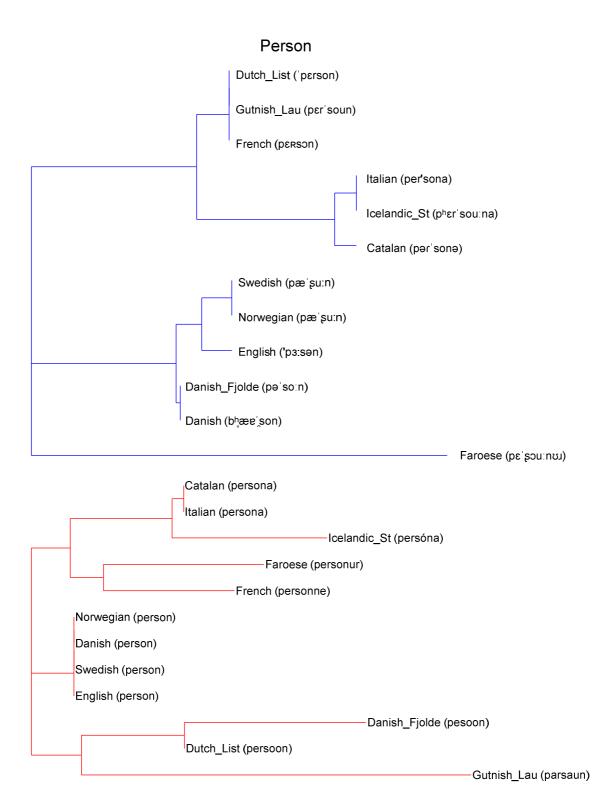


## Many

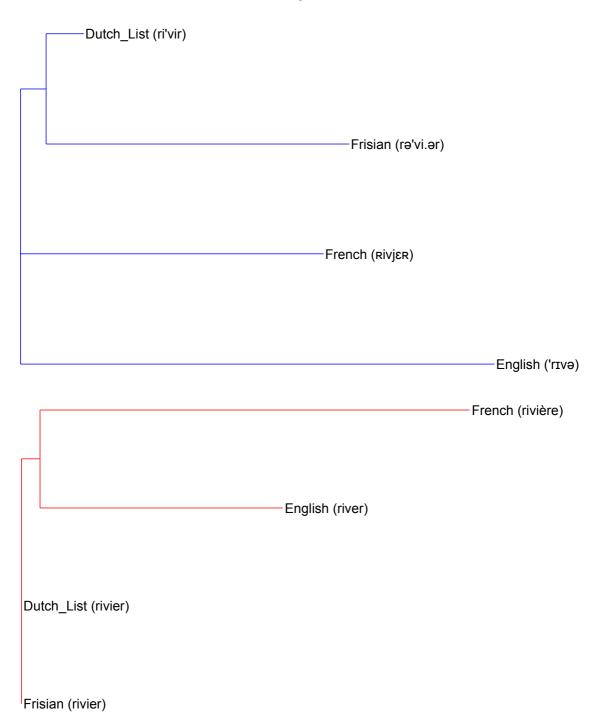


### Mountain

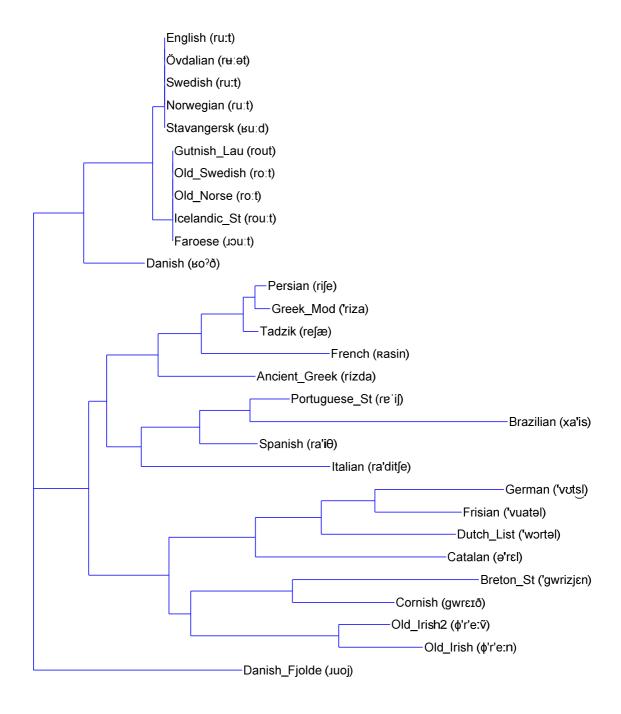




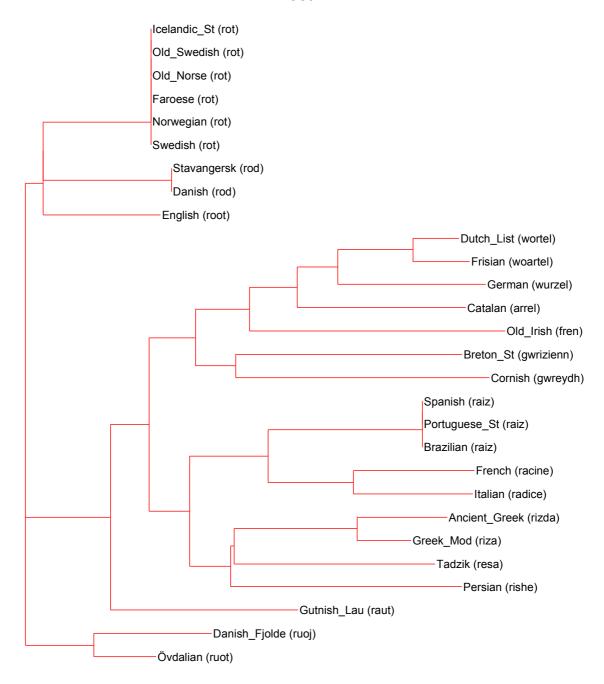
## River

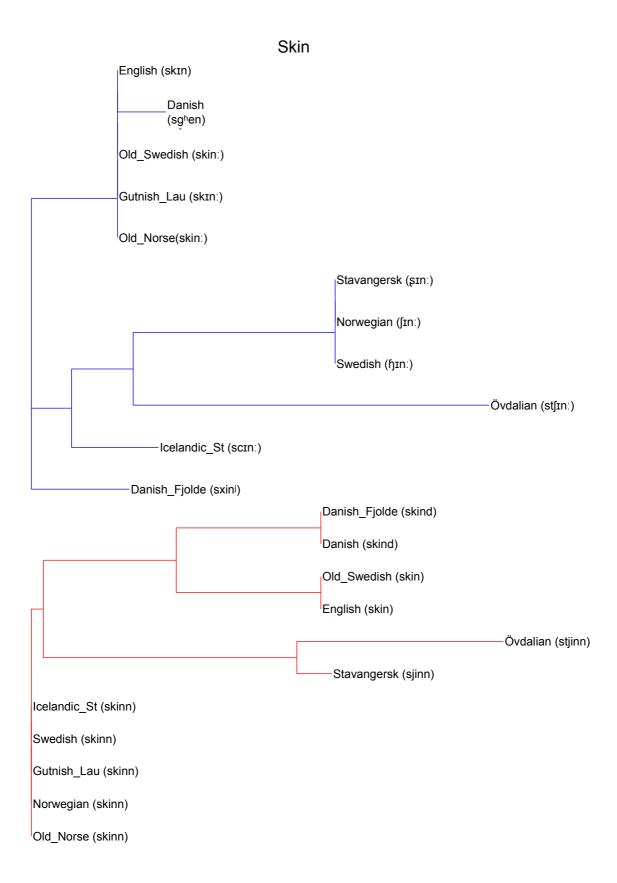


## Root

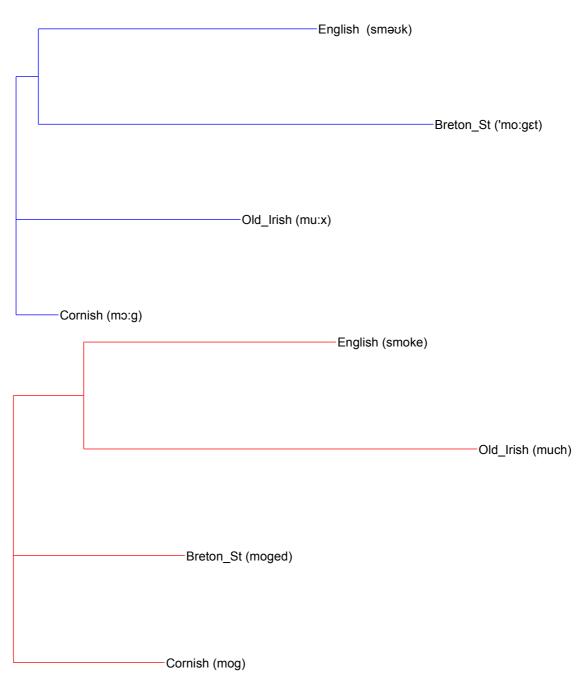


#### Root

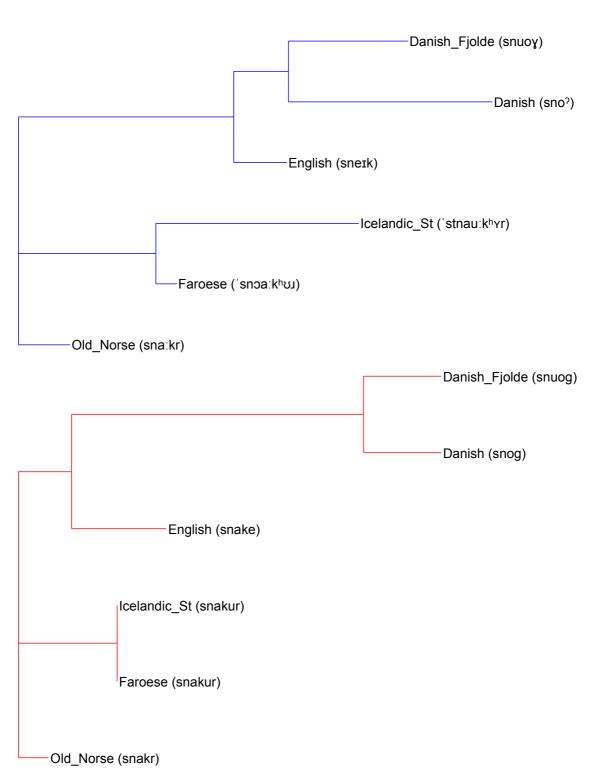




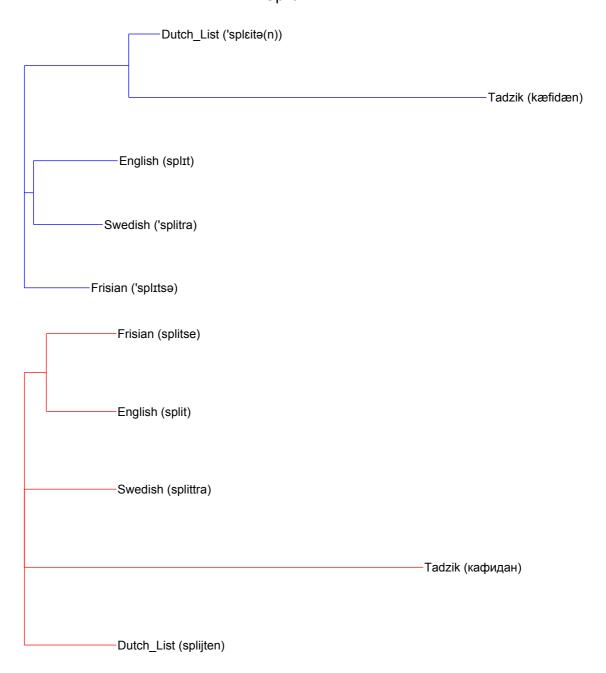
### Smoke

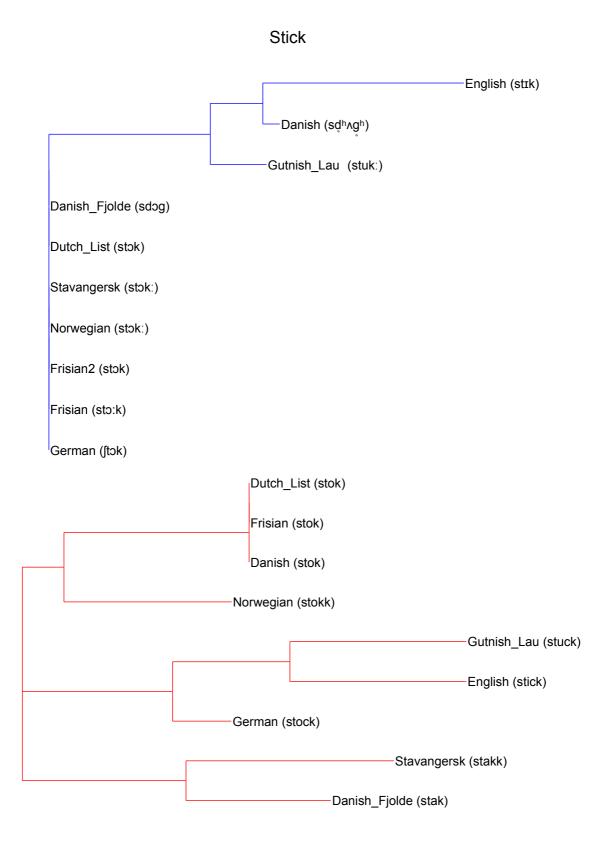






## Split





## They

