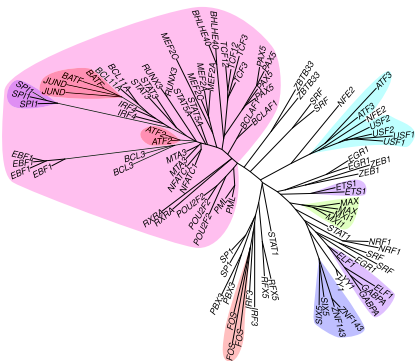
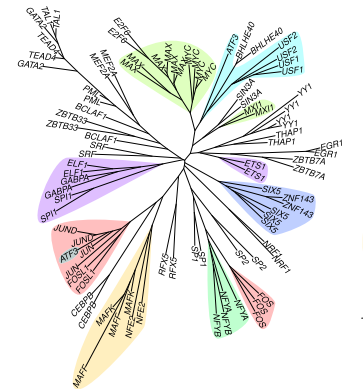


GM12878



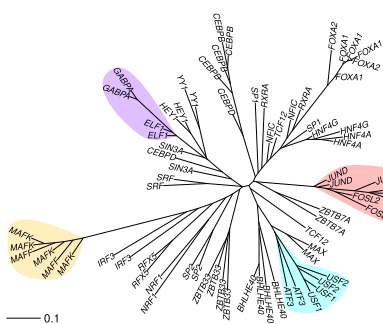
0.1

K562



0.1

HepG2



0.1

### Known cofactors

- AP-1
- MAF/NFE2
- MAX/MYC/MXI
- NFY

### Putative cofactors

- ATF3/USF
- SIX5/ZNF143

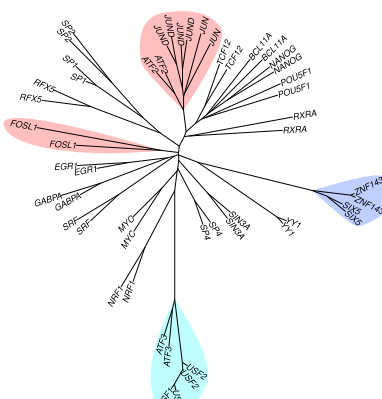
### Conserved binding domain

- ETS family

### Functional pathway

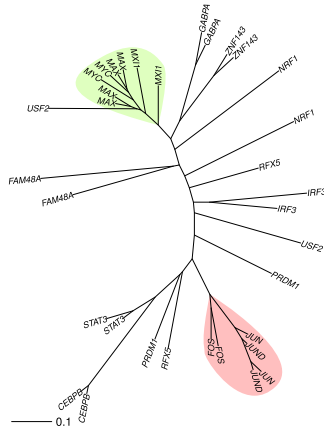
- Lymphocyte differentiation

H1-hESC



0.1

HeLa-S3



0.1