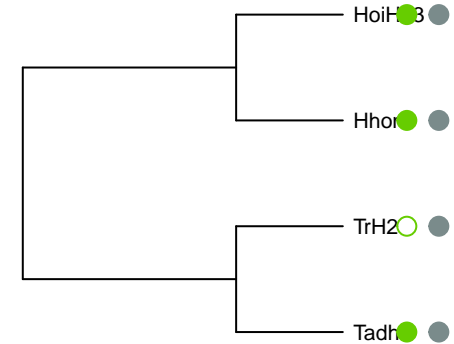
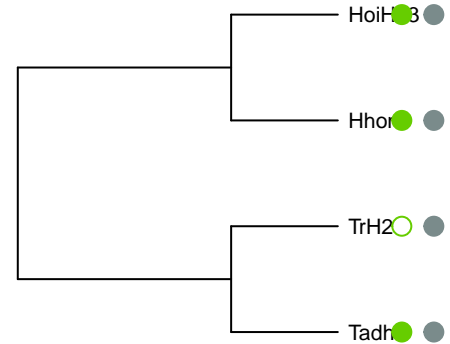


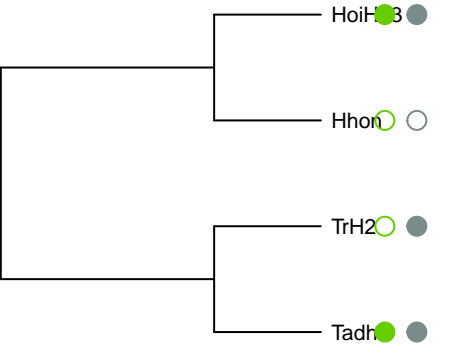
Homeodomains.HG1.5:HHEX



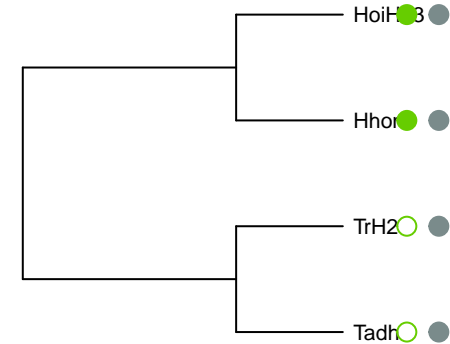
Homeodomains.HG6.11:POU3F1/POU3F2/POU3F3/P



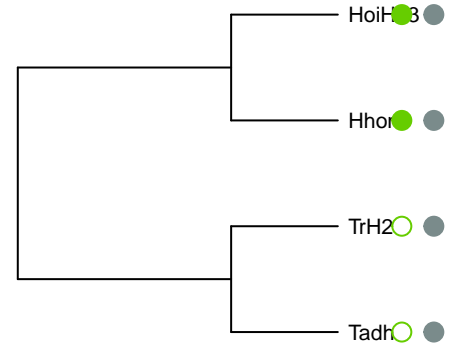
zf-C2H2.HG3.3:like:MTF1/ZNF76/ZNF143



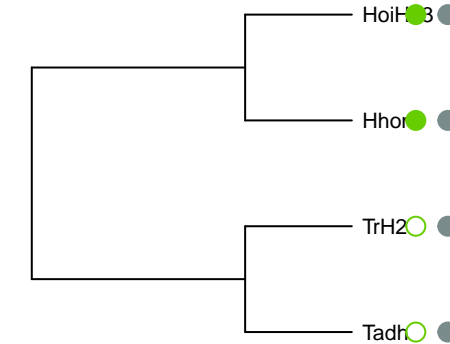
SAND.HG1.10:DEAF1



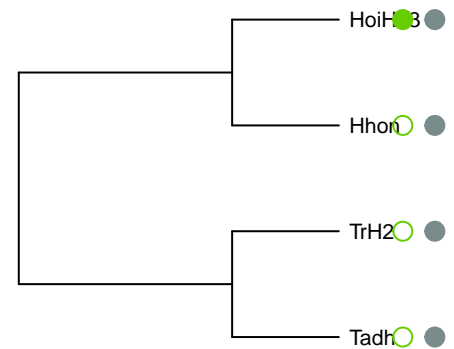
HLH.HG1.39:MYC/MYCL/MYCN



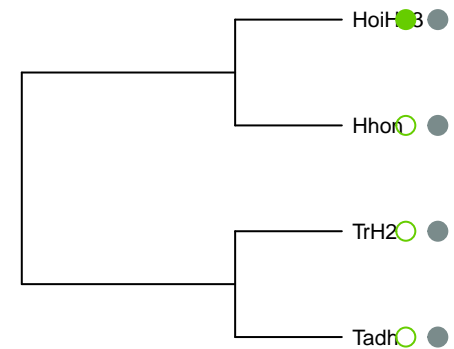
RFX.HG1.7:RFX4/RFX6



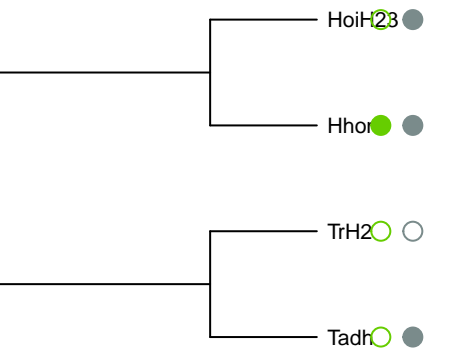
ACH2/MAF/MAFA/MAFB/MAFF/MAFG/MAFK/NFE2/



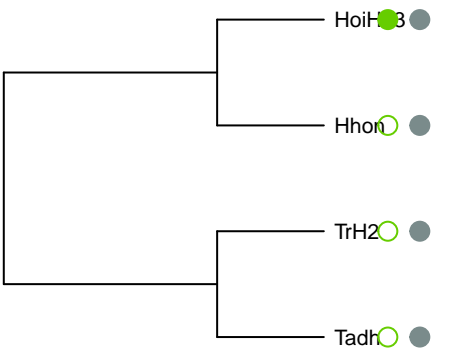
Myb.HG1.1:DNAJC1/DNAJC2



DBD_Tnp_MULE.Unclassified



F_related_matrix_associated_actin_dependent_reg



HMGbox_Sox.HG1.3:LEF1/TCF7/TCF7L1/TCF7L2/426/ZNF524/ZNF559-ZNF177/ZNF628/ZNF883,zf-C

