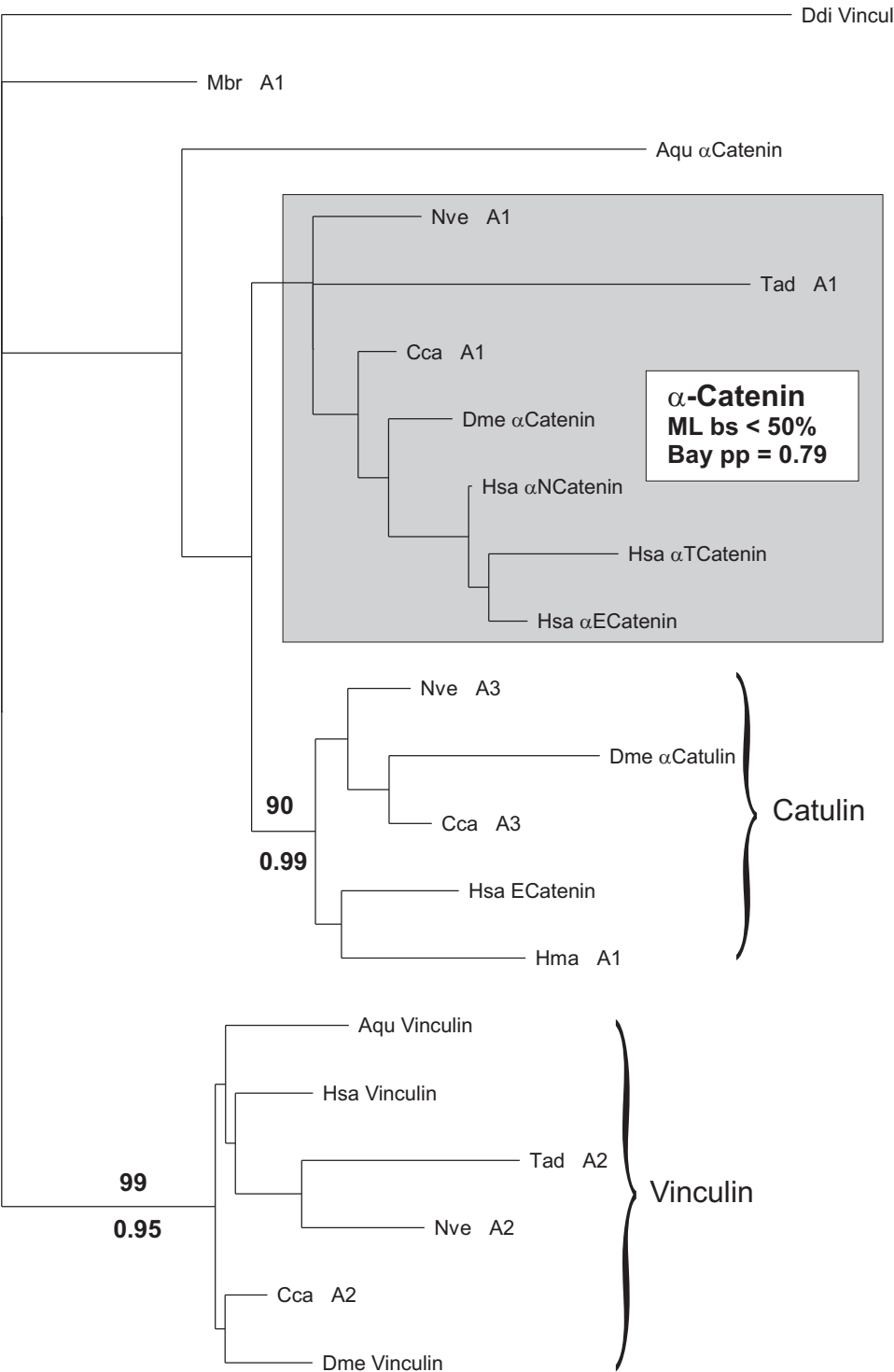
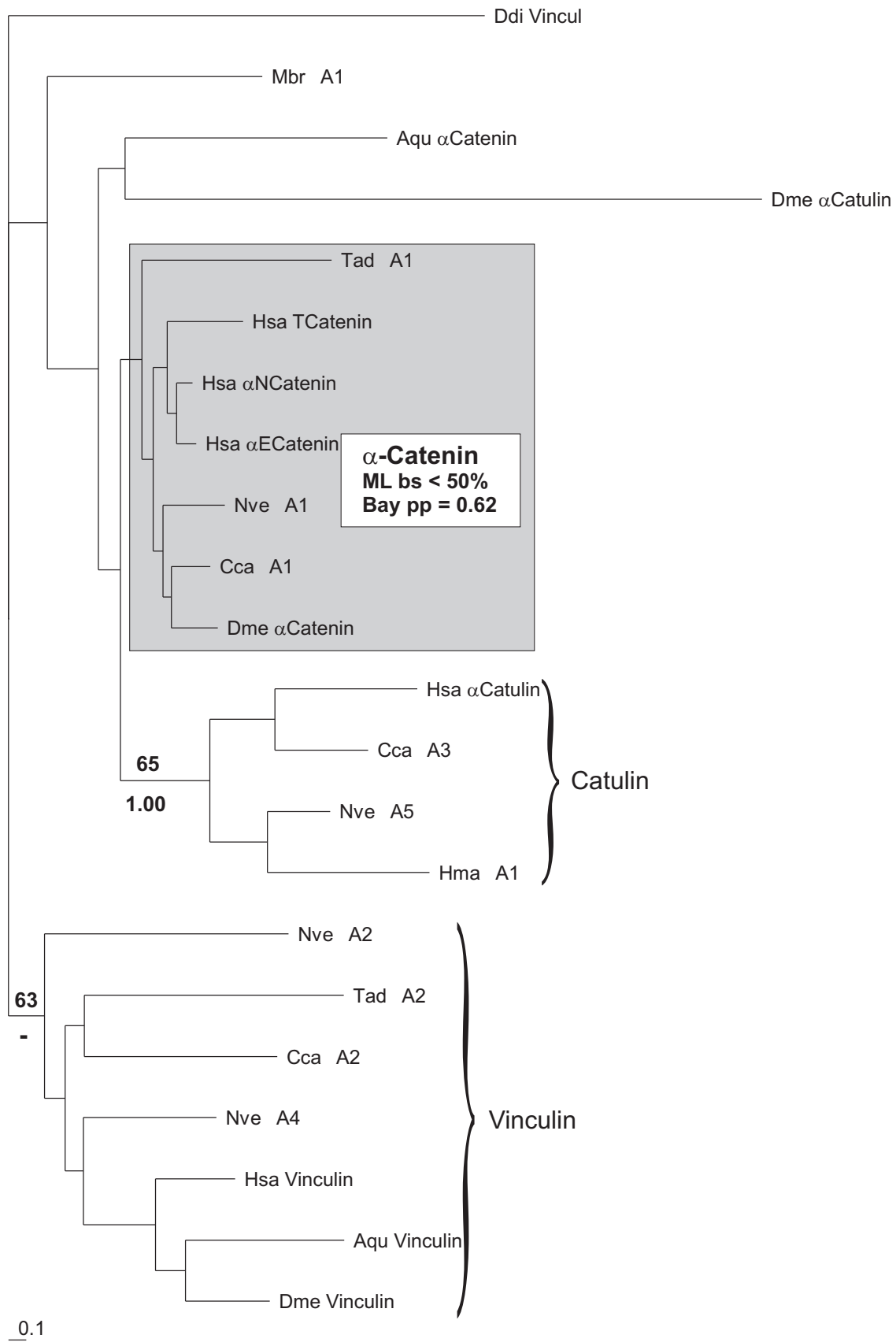


α -Catenin partition A

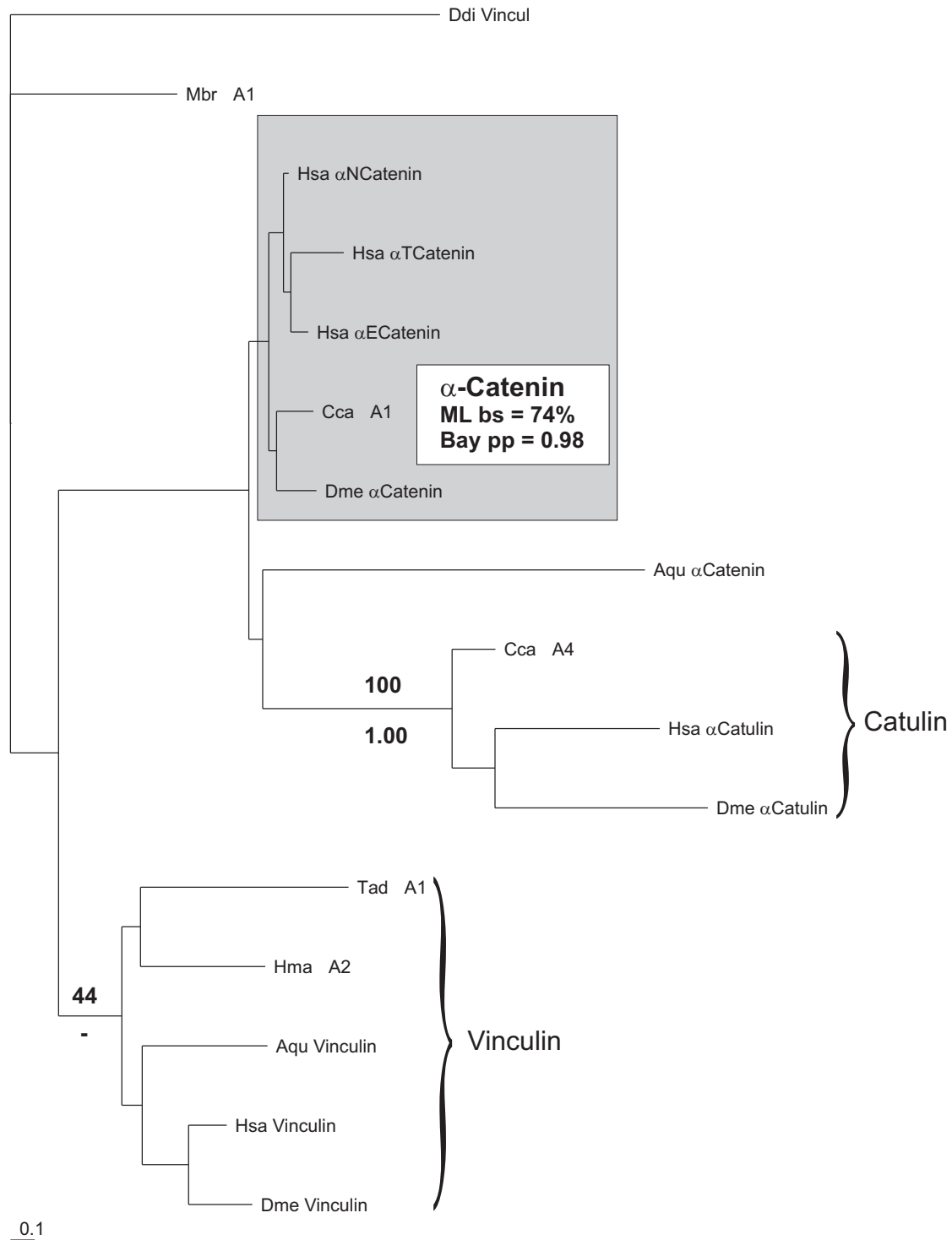


0.1

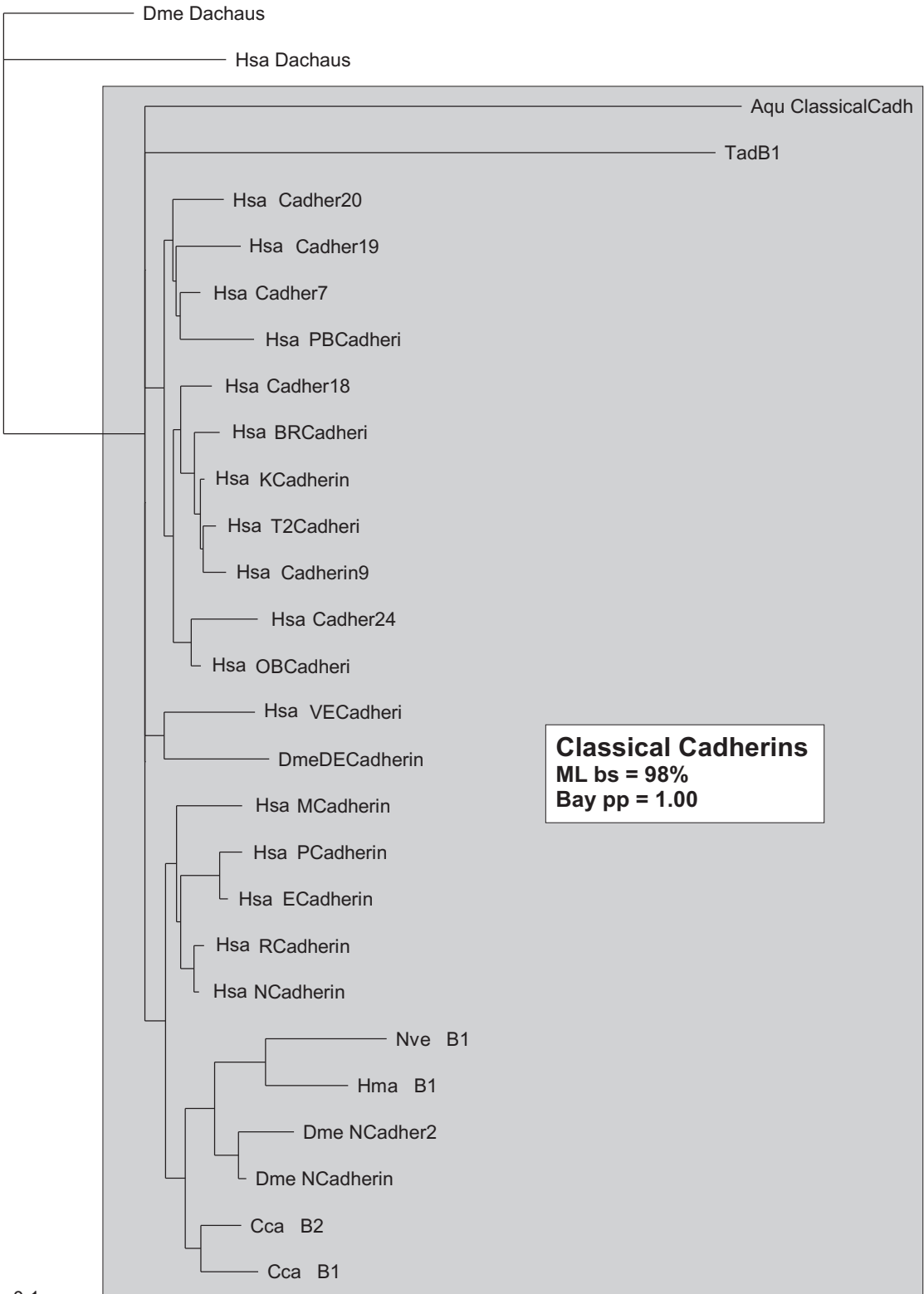
α -Catenin partition B



α -Catenin partition C

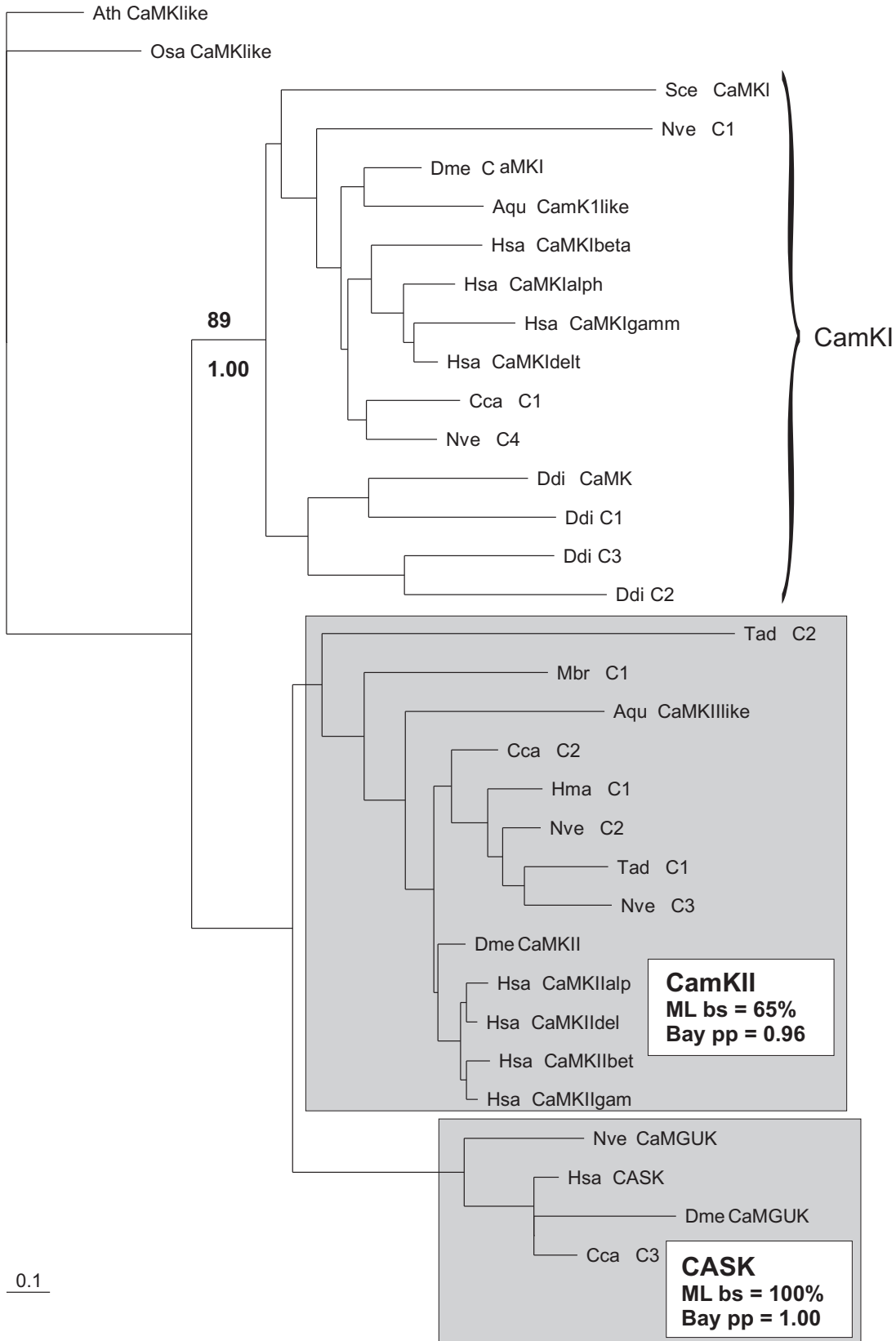


Classical Cadherins

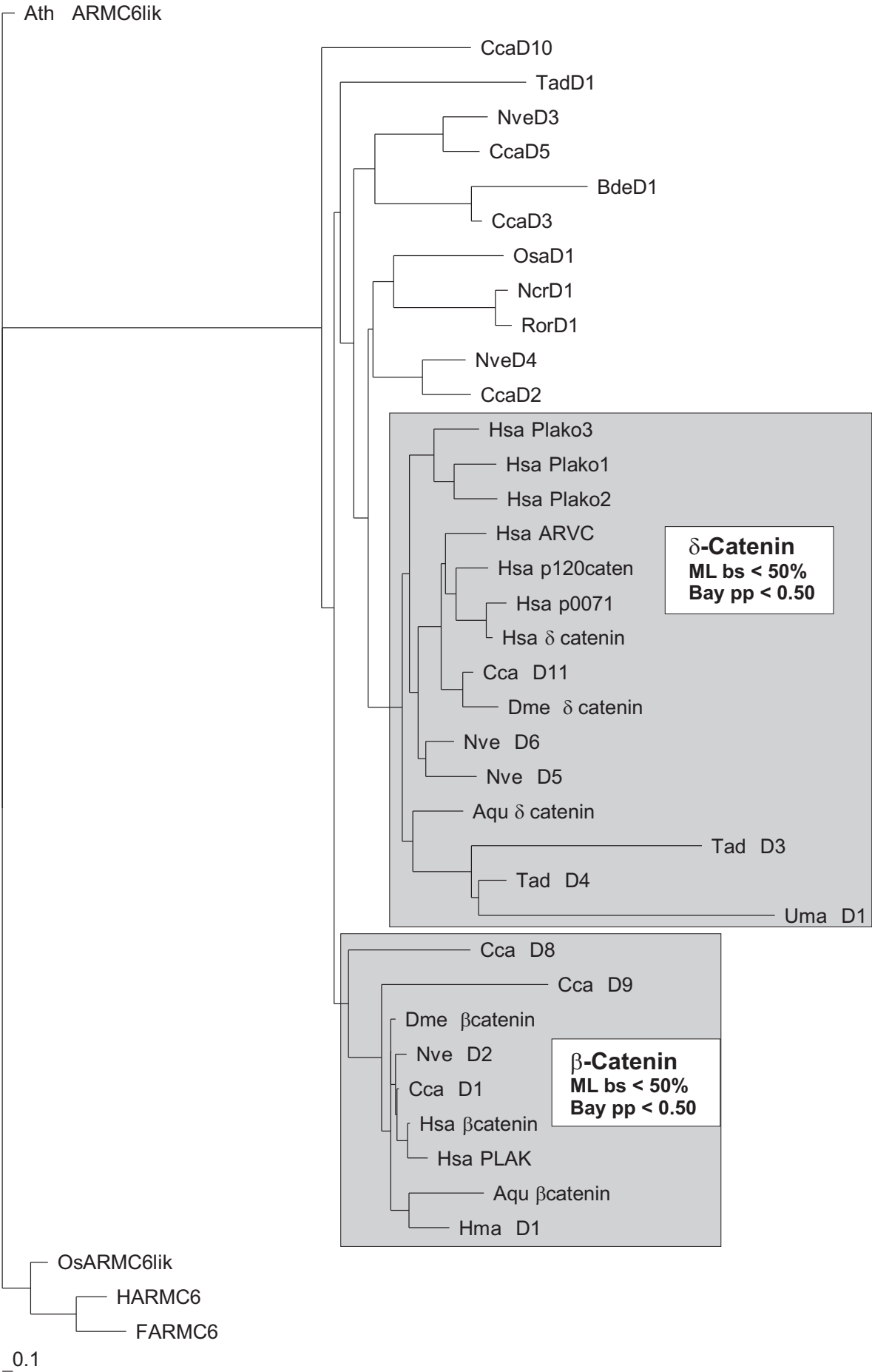


_0.1

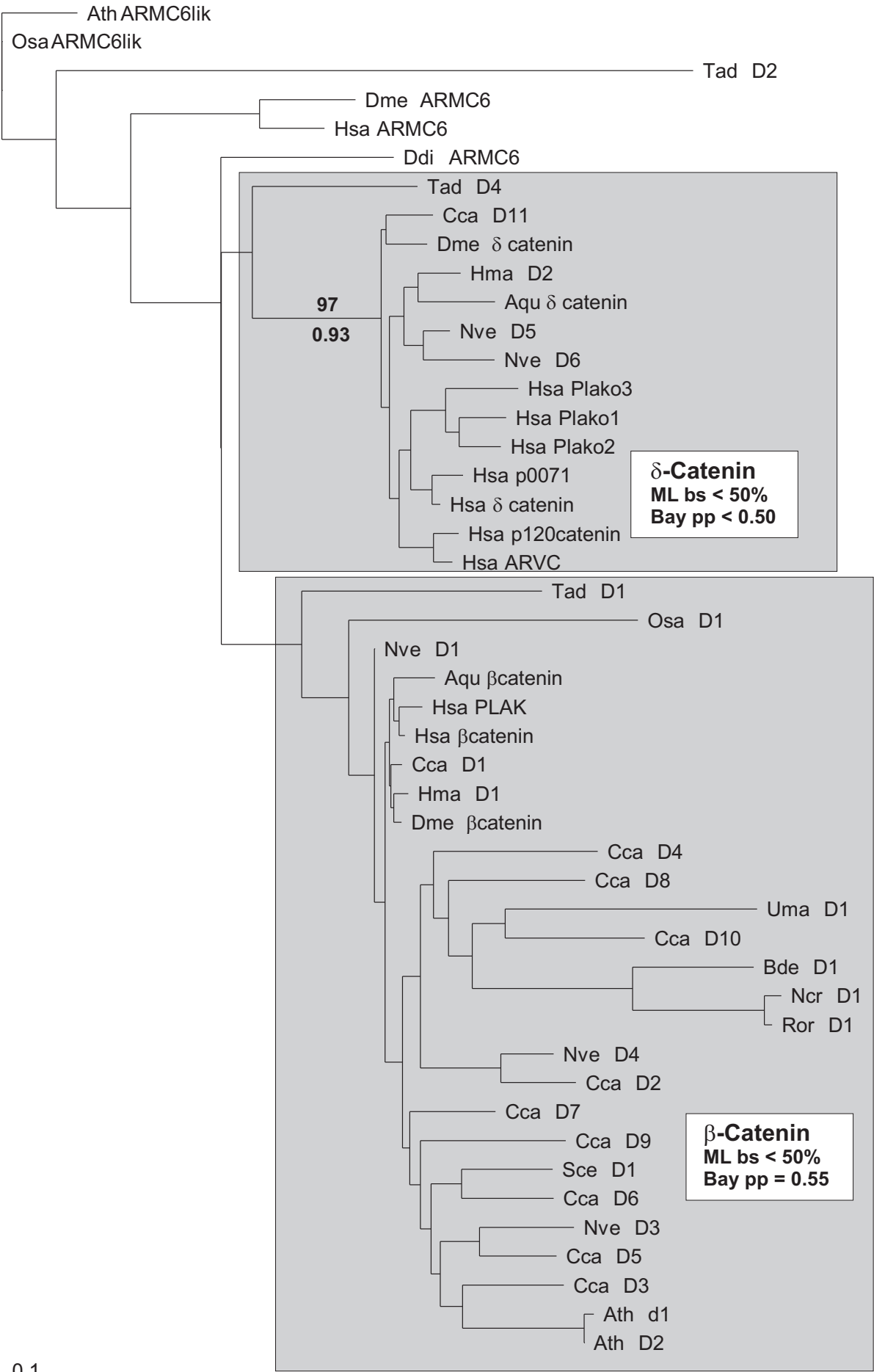
CamKII



Catenins partition A

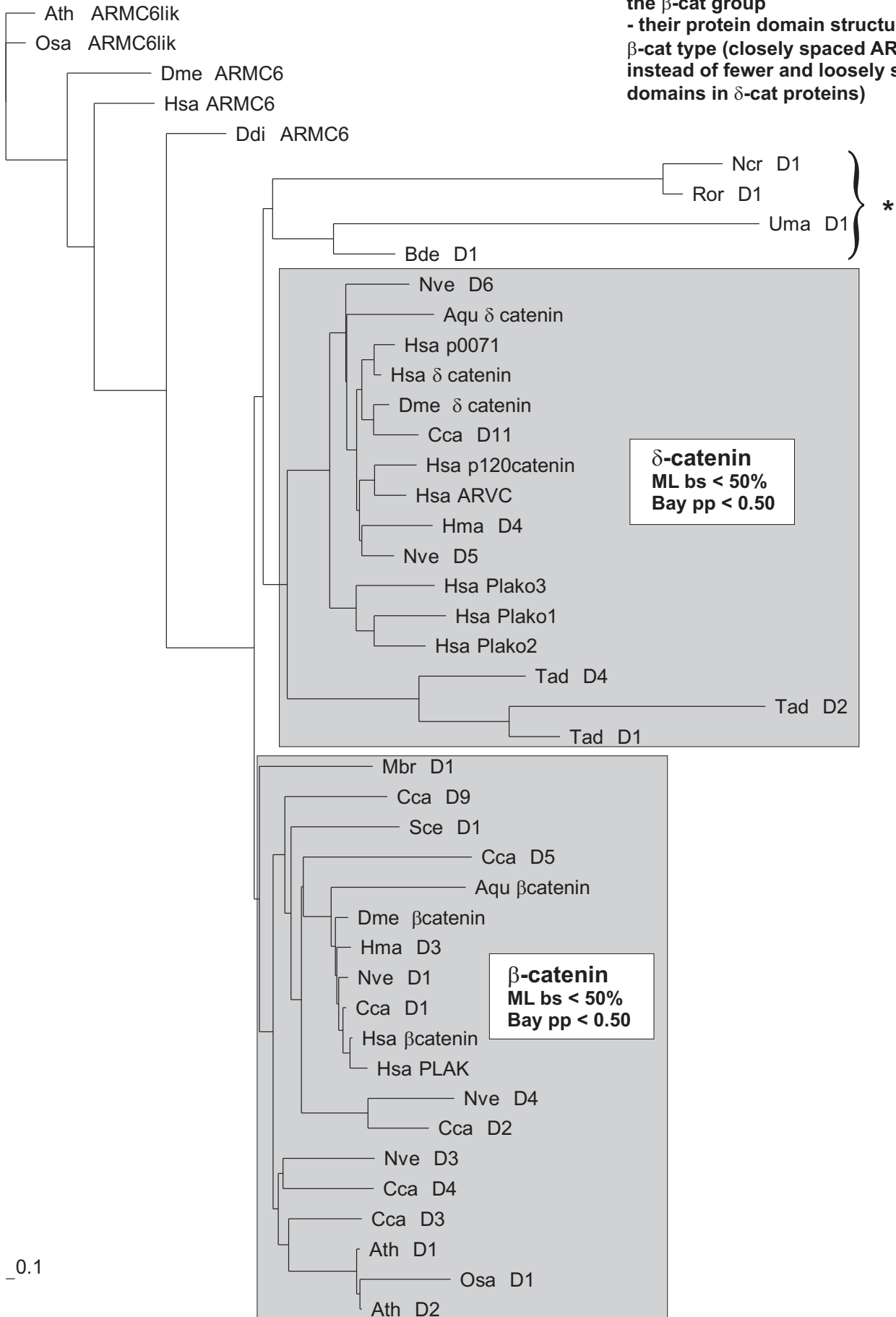


Catenins partition B

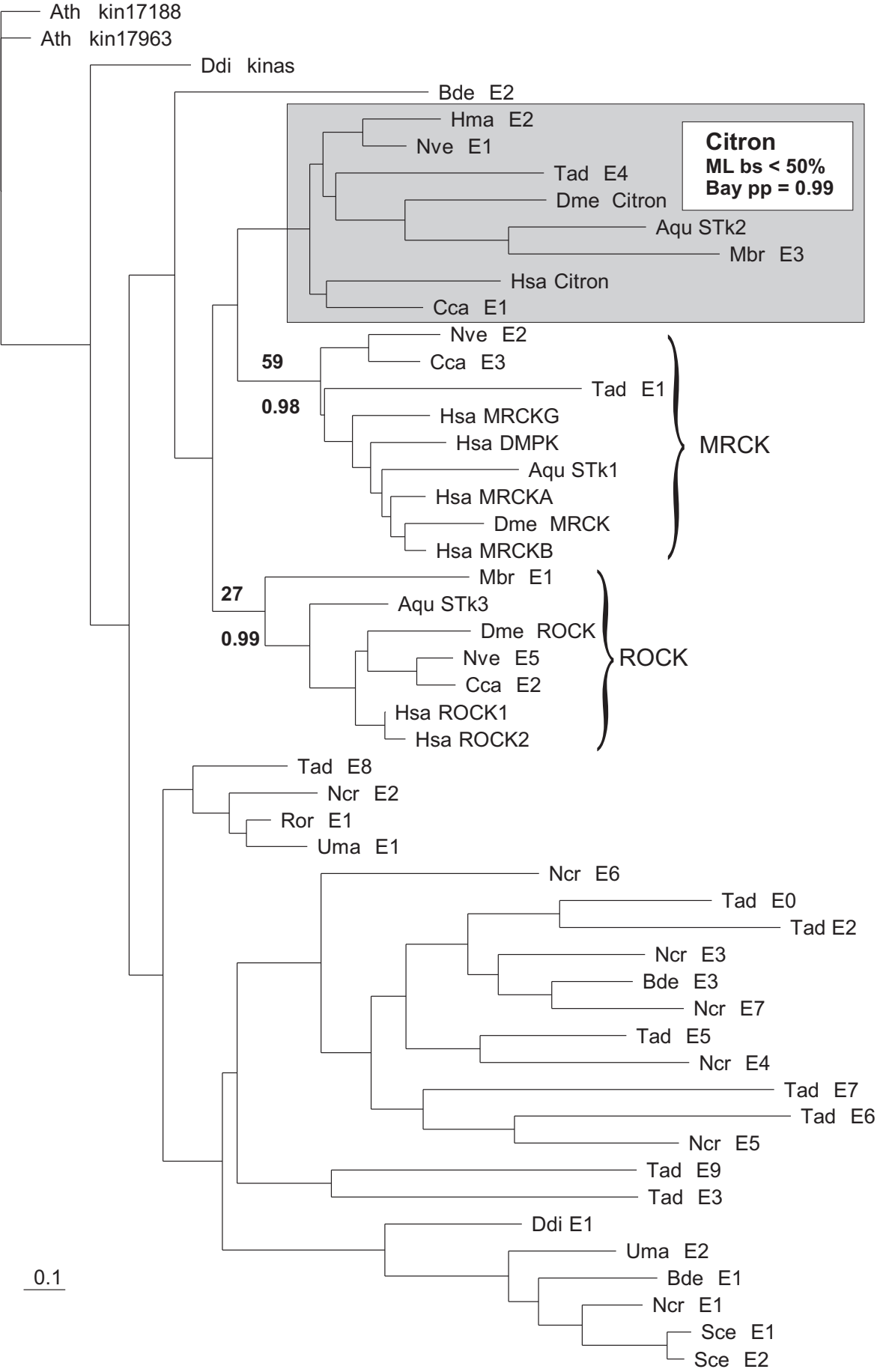


Catenins partition C

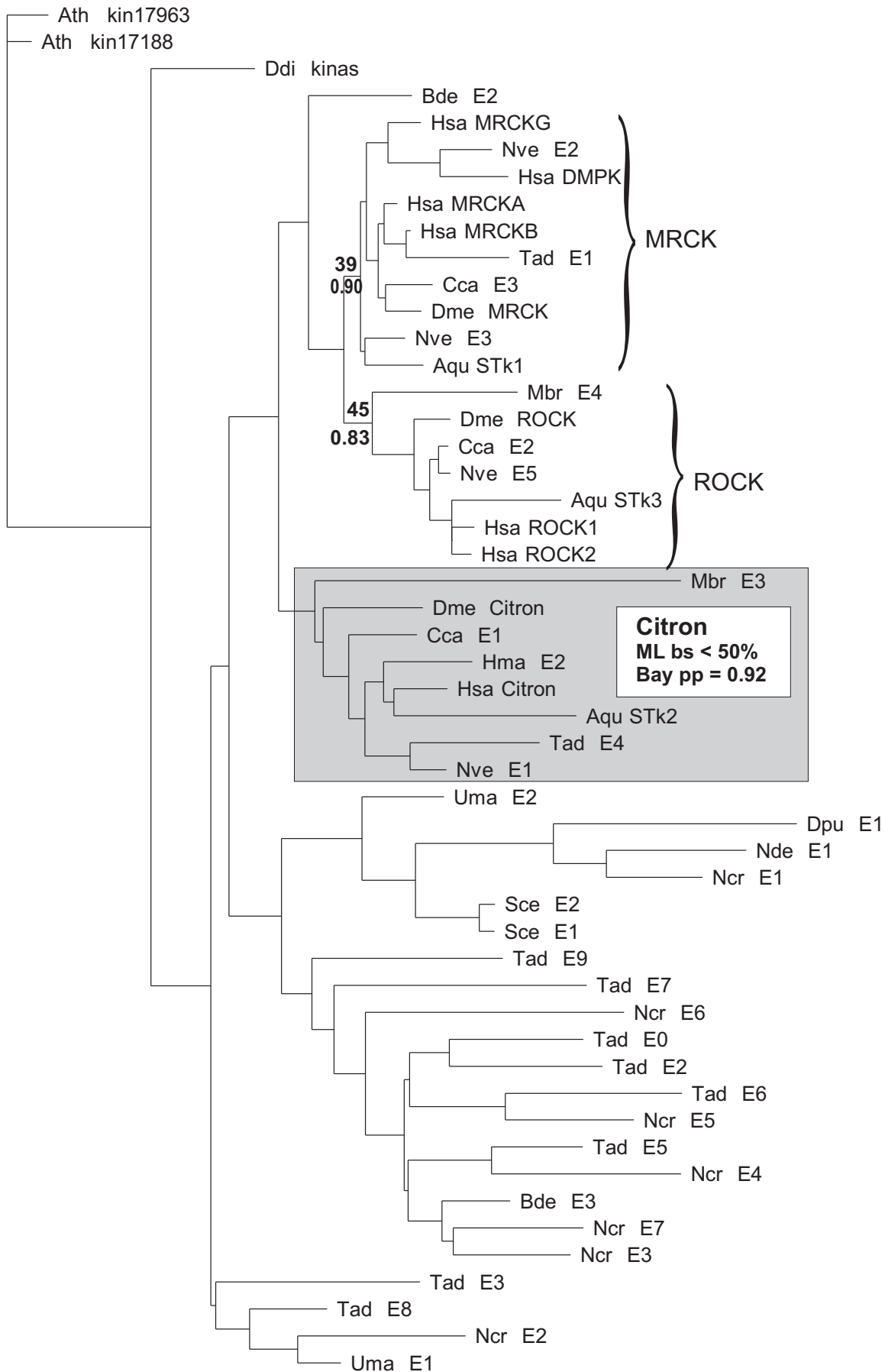
* These genes are not considered part of the δ -cat group because :
 - in partition B they are placed within the β -cat group
 - their protein domain structure is of the β -cat type (closely spaced ARM domains, instead of fewer and loosely spaced ARM domains in δ -cat proteins)



Citron partition A

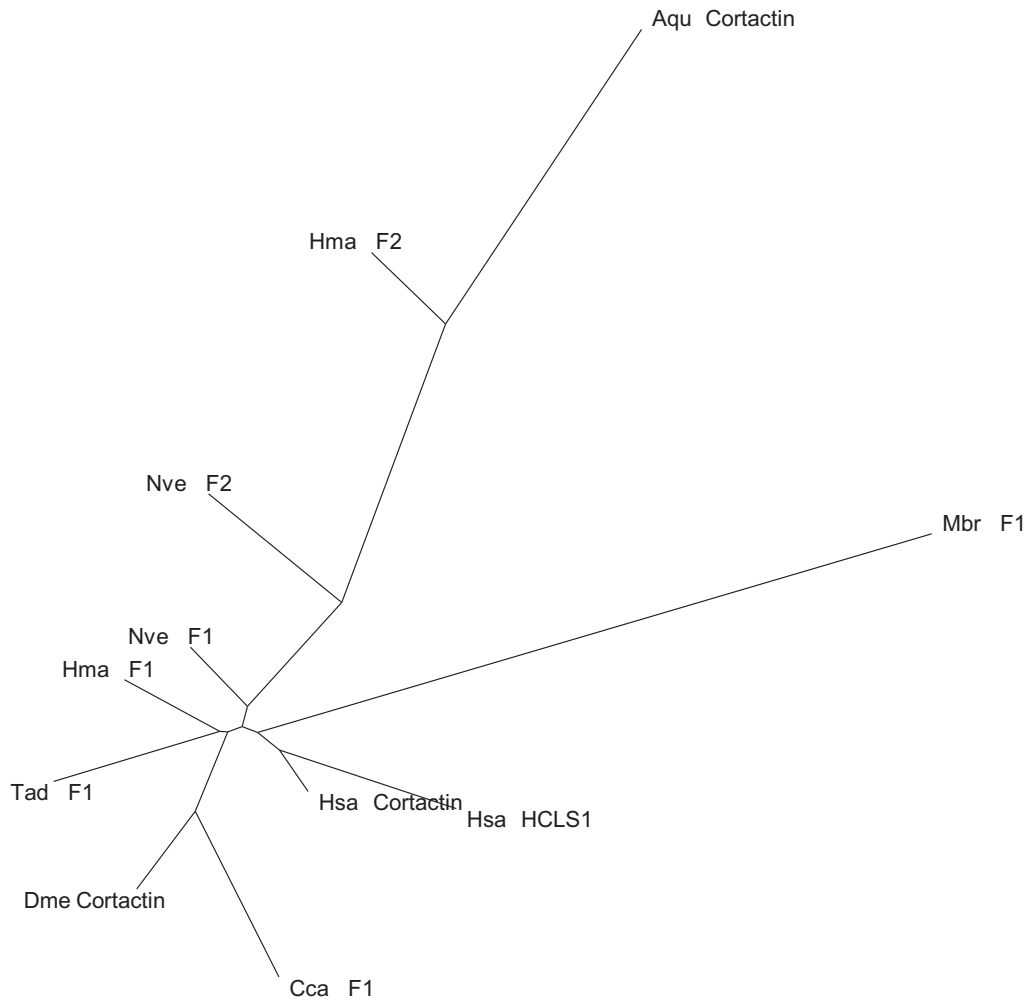


Citron partition B



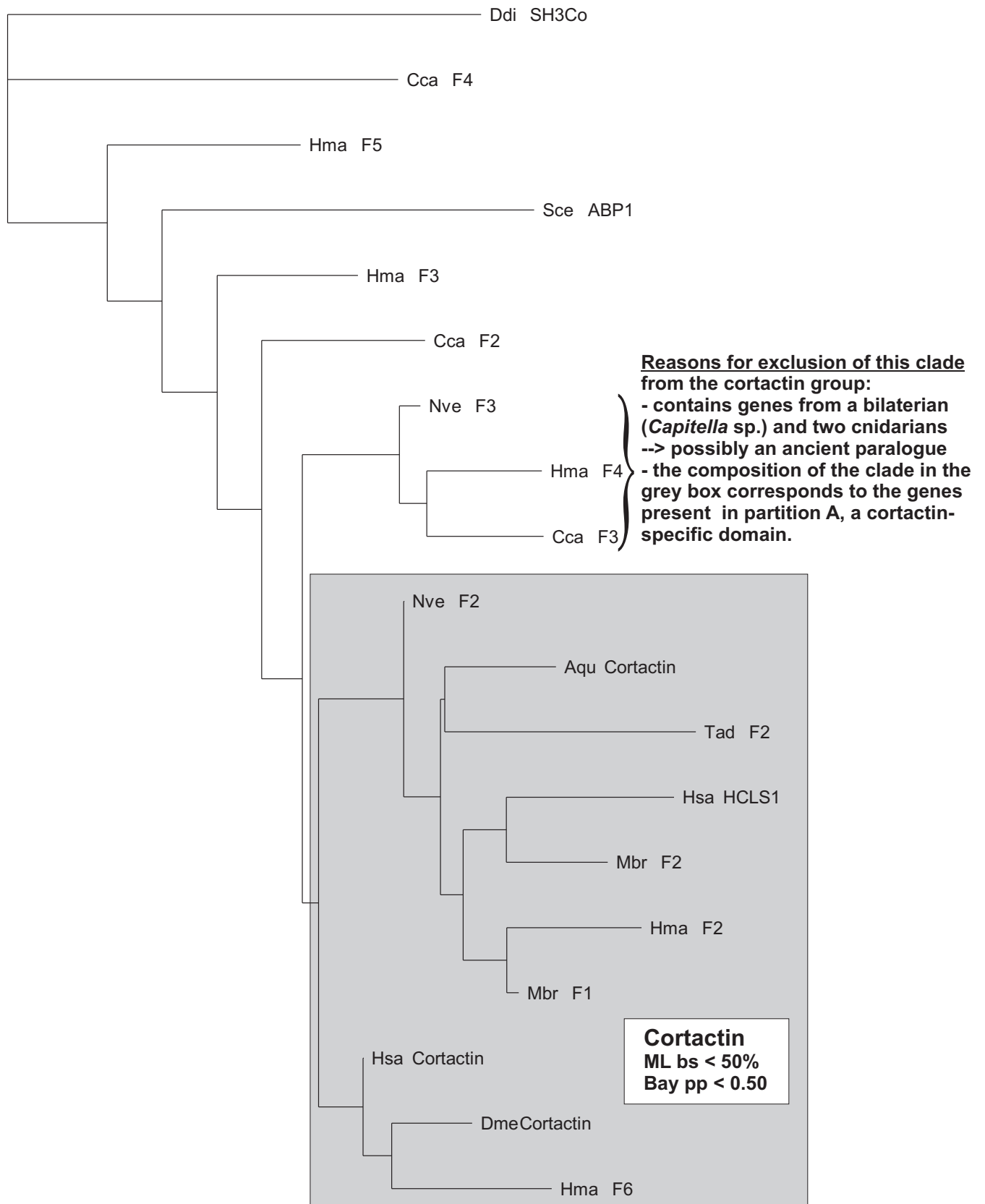
Cortactin partition A

This part of cortactin has no similarity to other proteins



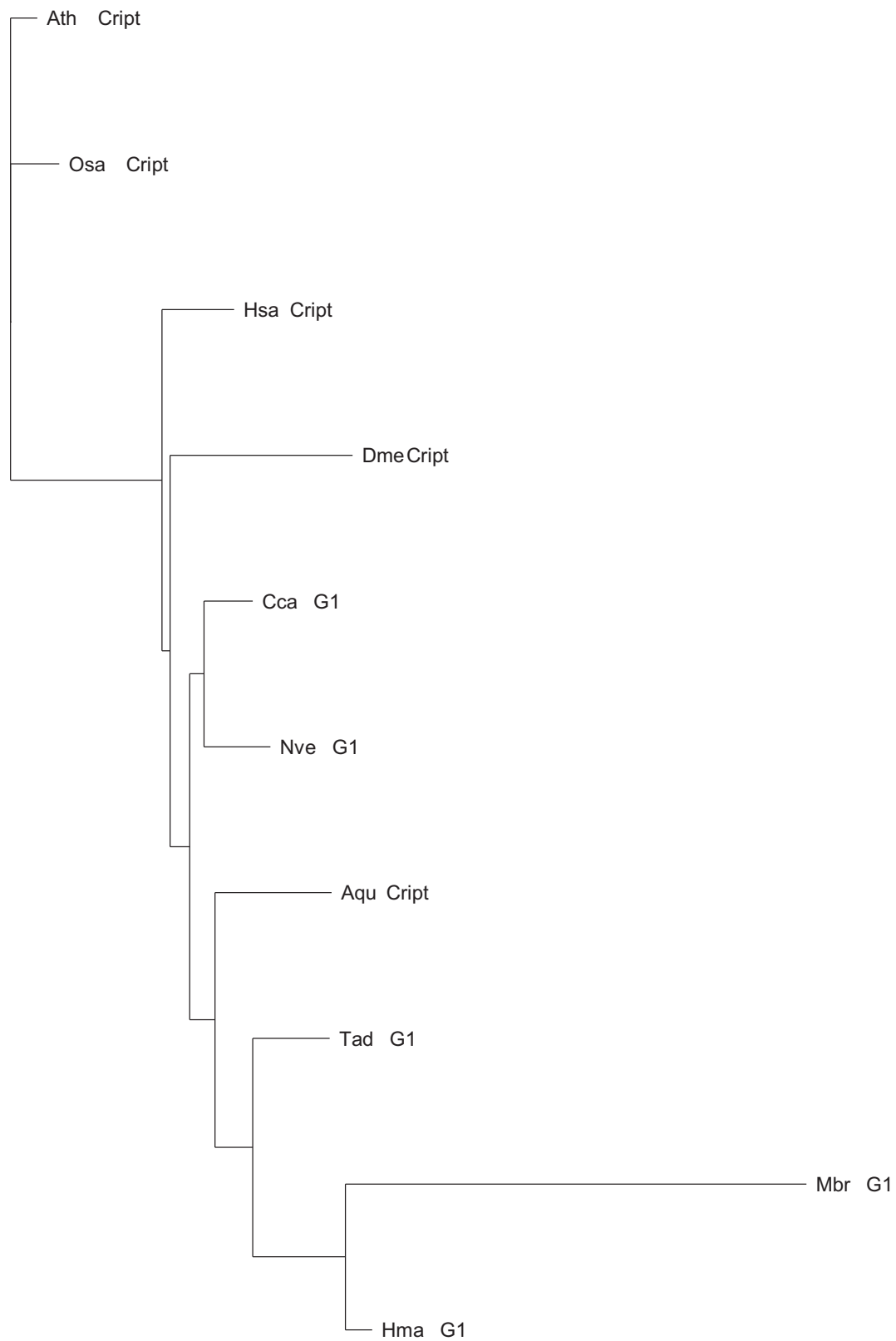
0.1

Cortactin partition B



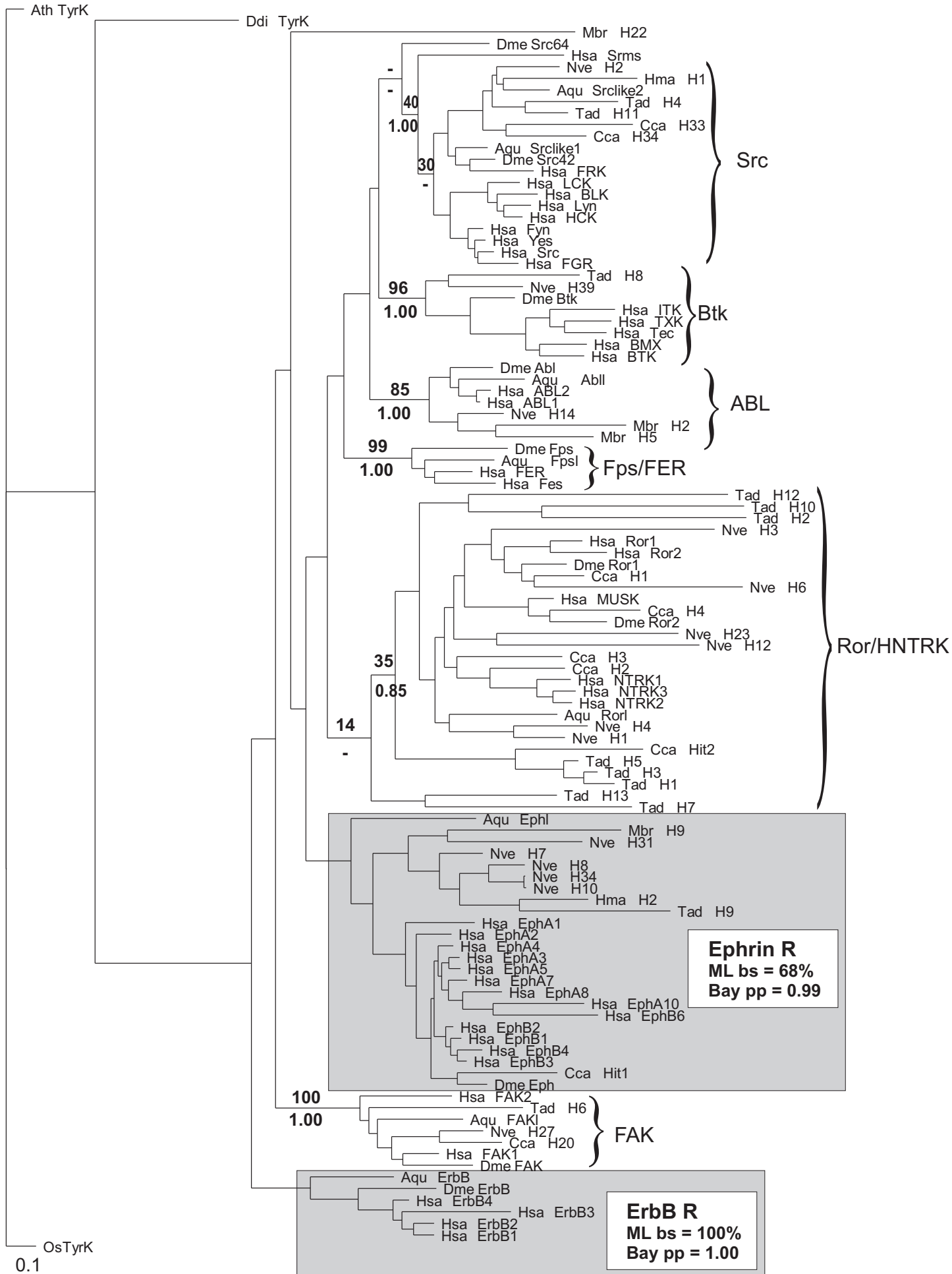
CRIPT

CRIPT has no similarity to other proteins

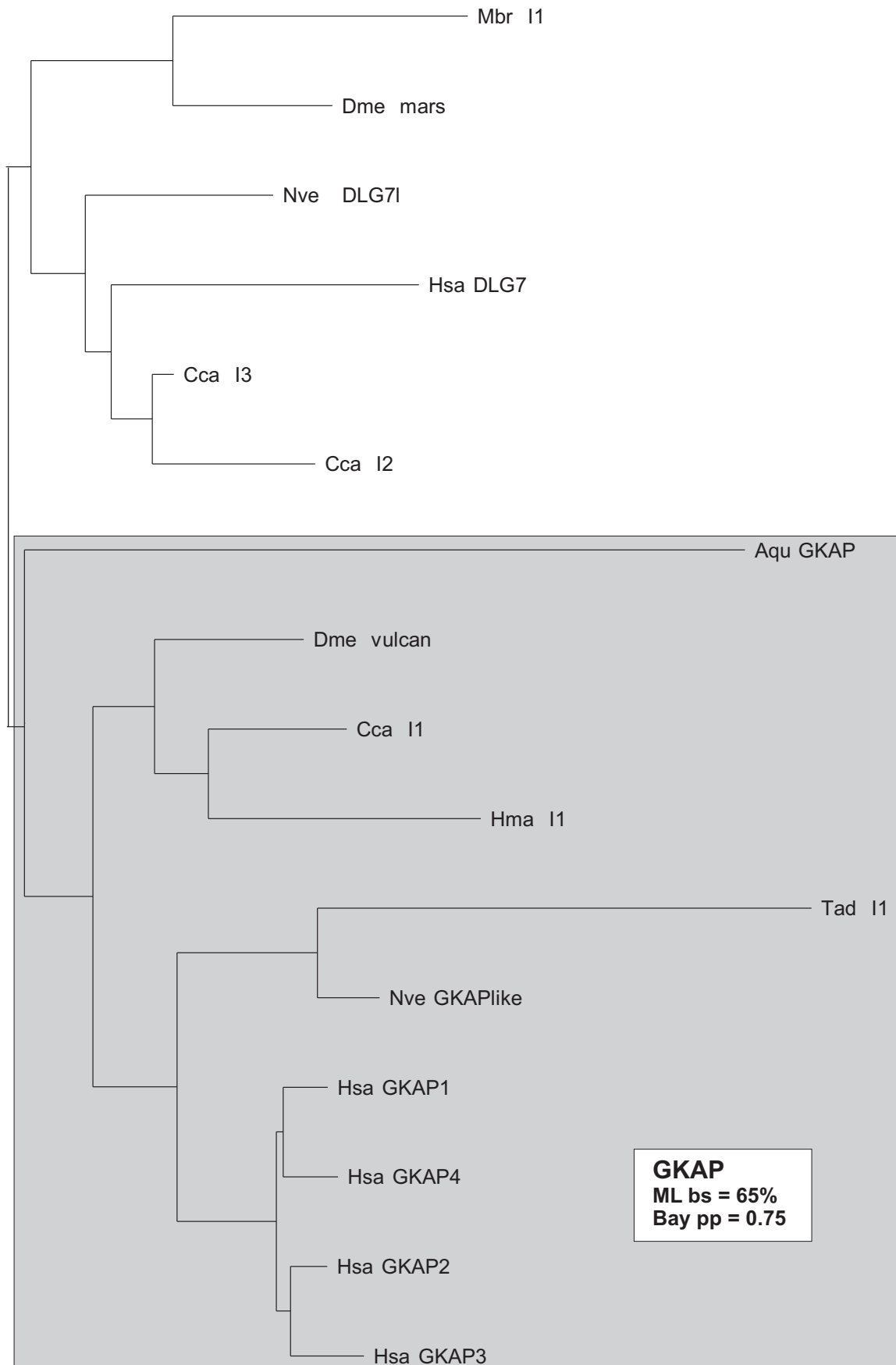


0.1

Ephrin R / ErbB R

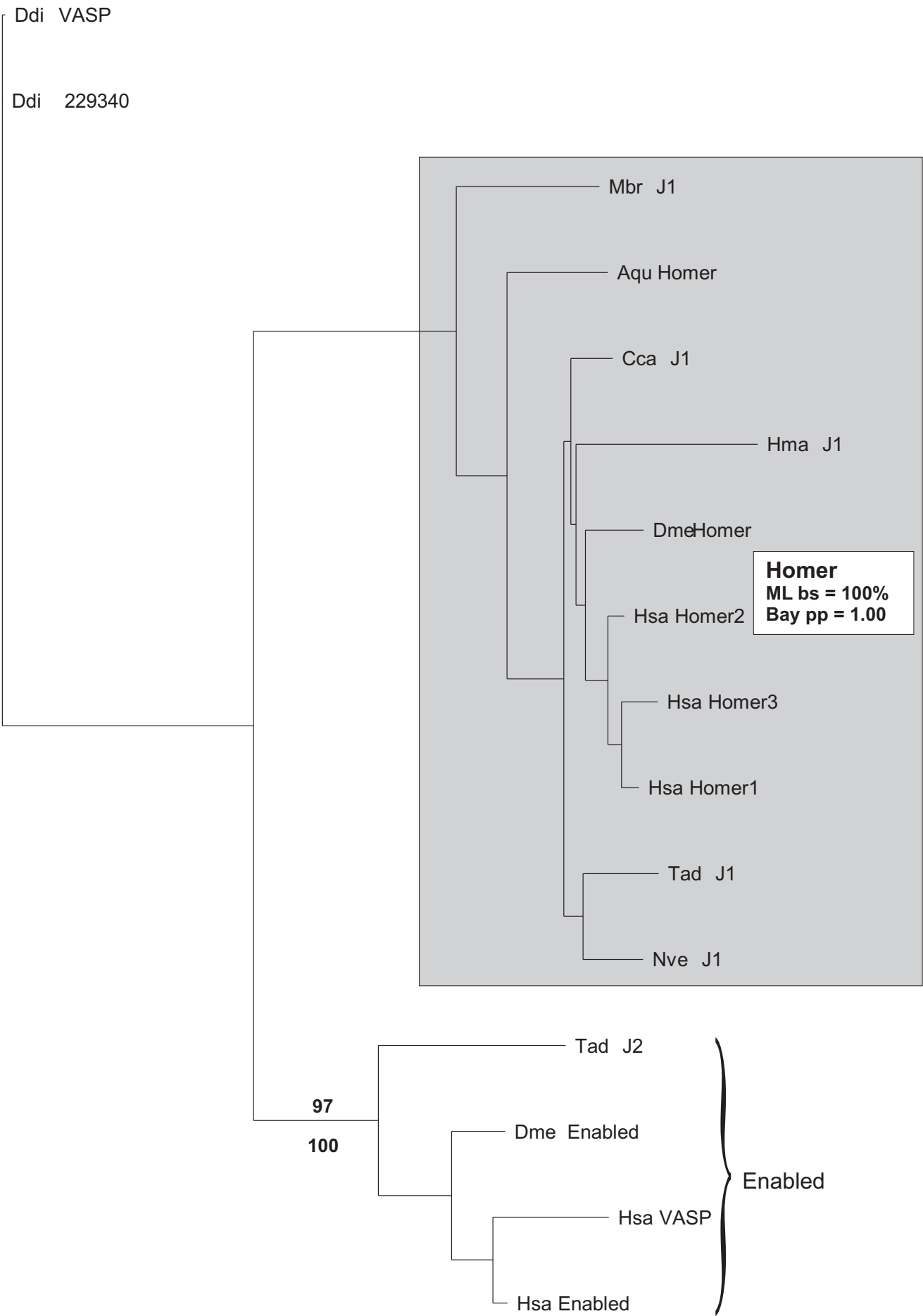


GKAP

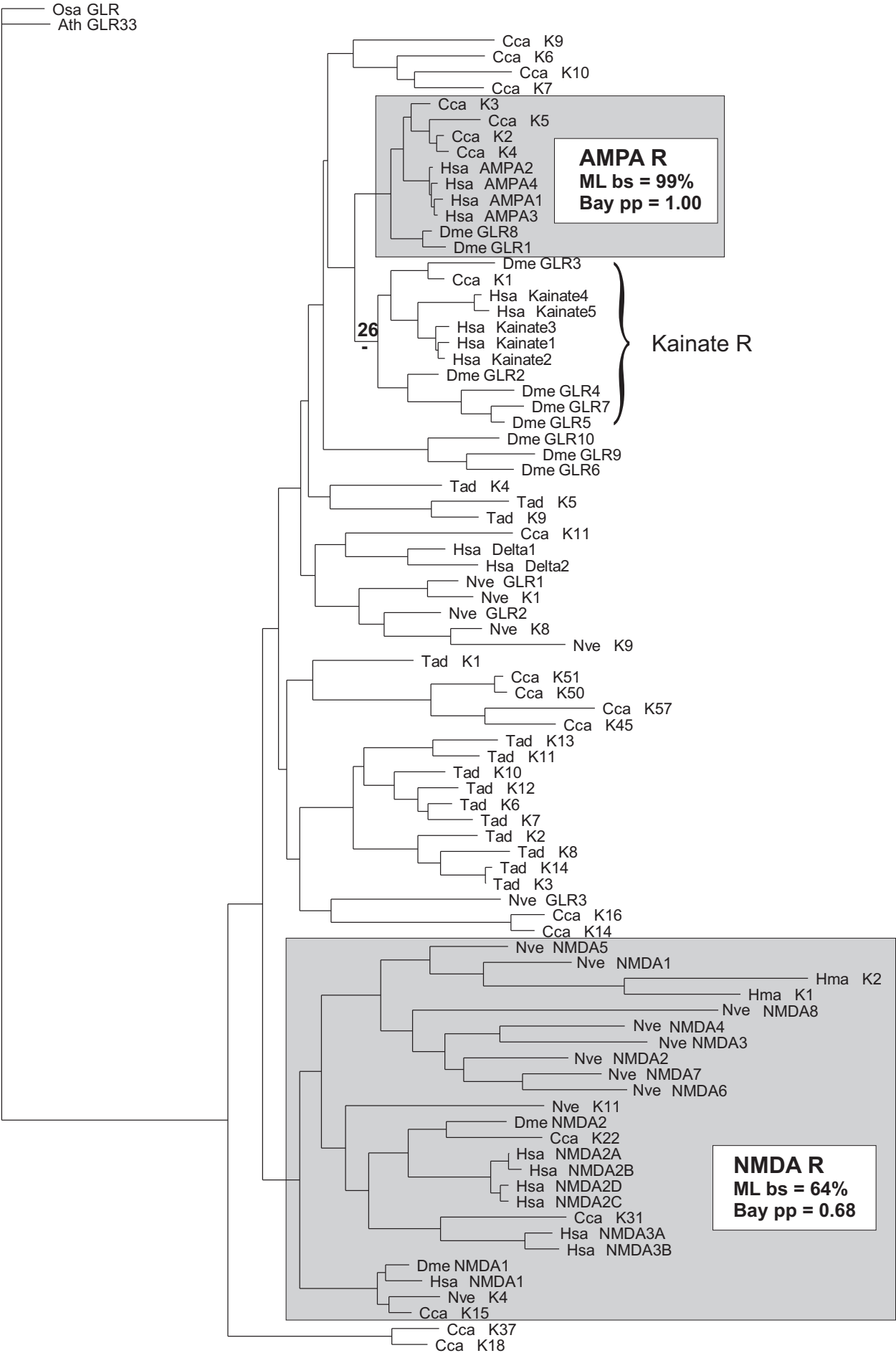


0.1

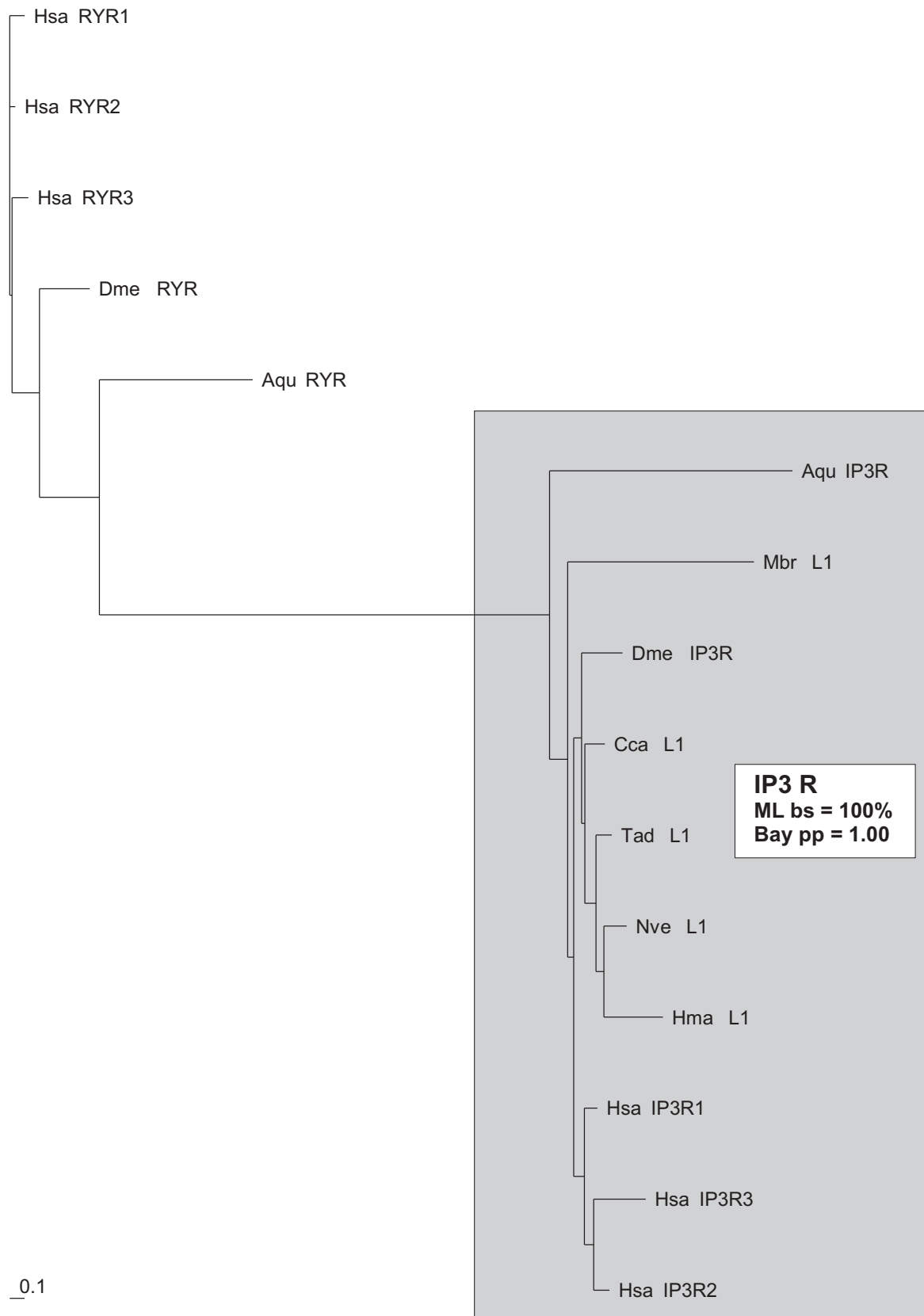
Homer



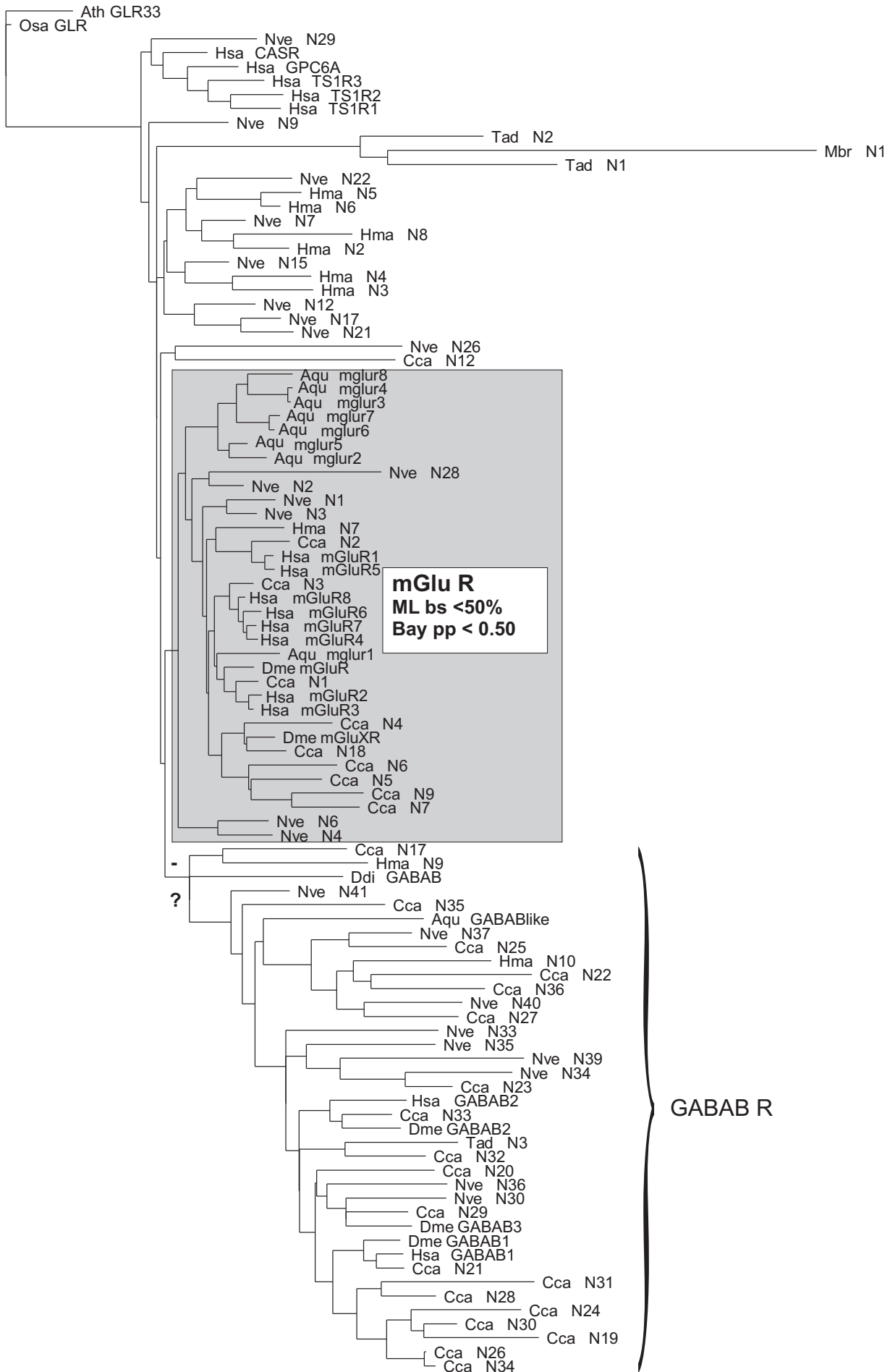
Ionotropic Glutamate Receptors



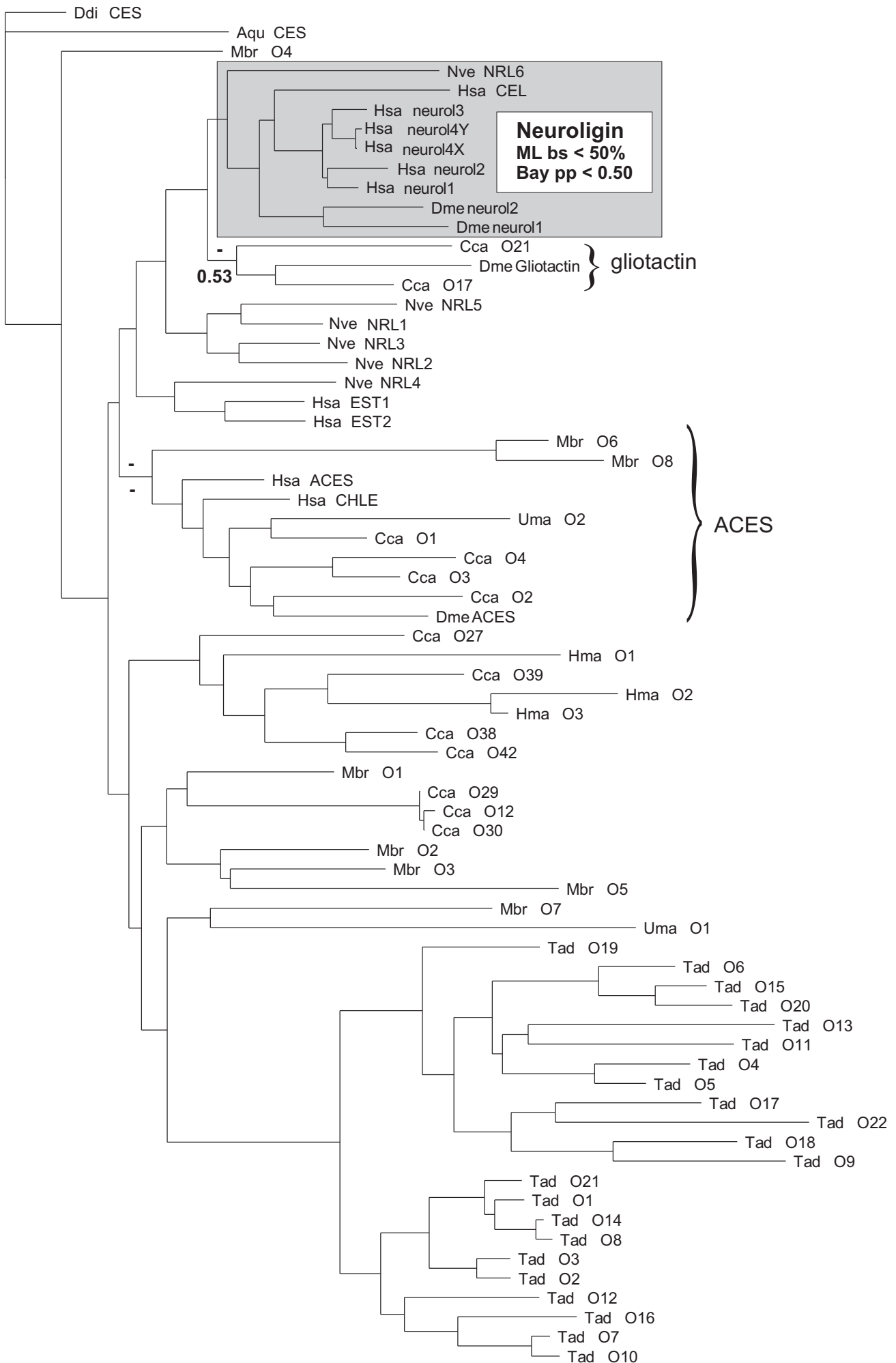
IP3 Receptor



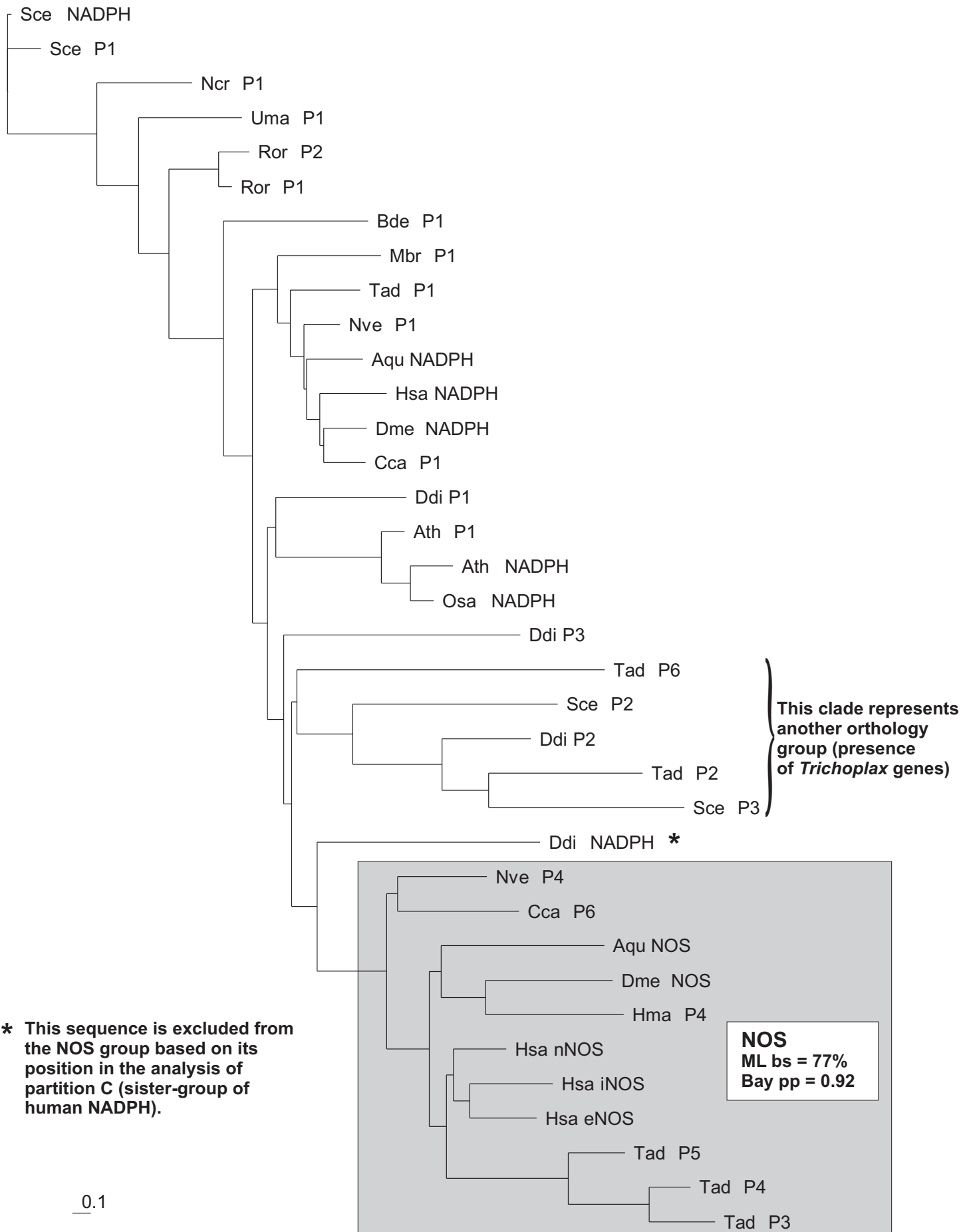
Metabotropic Glutamate Receptors



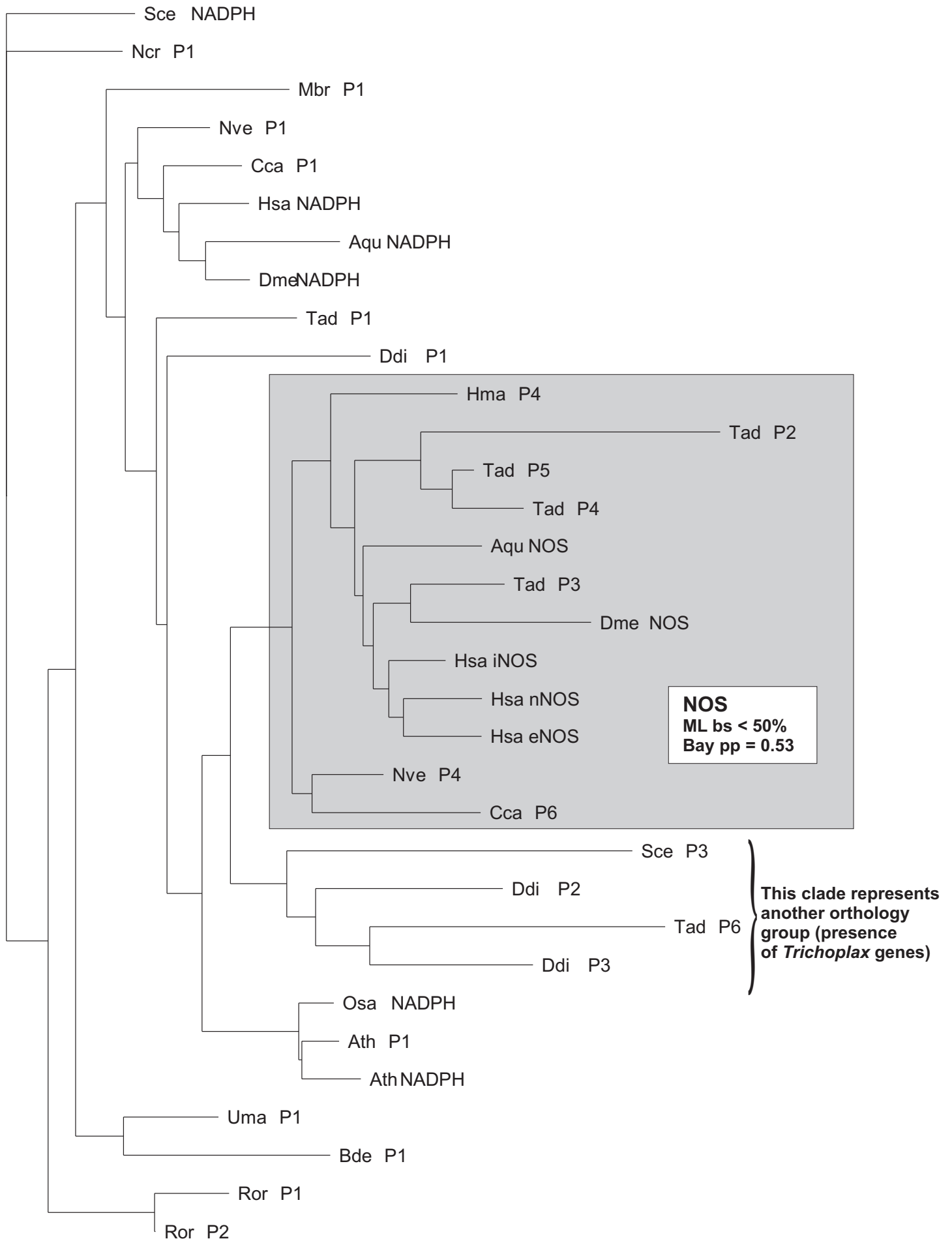
Neuroligin



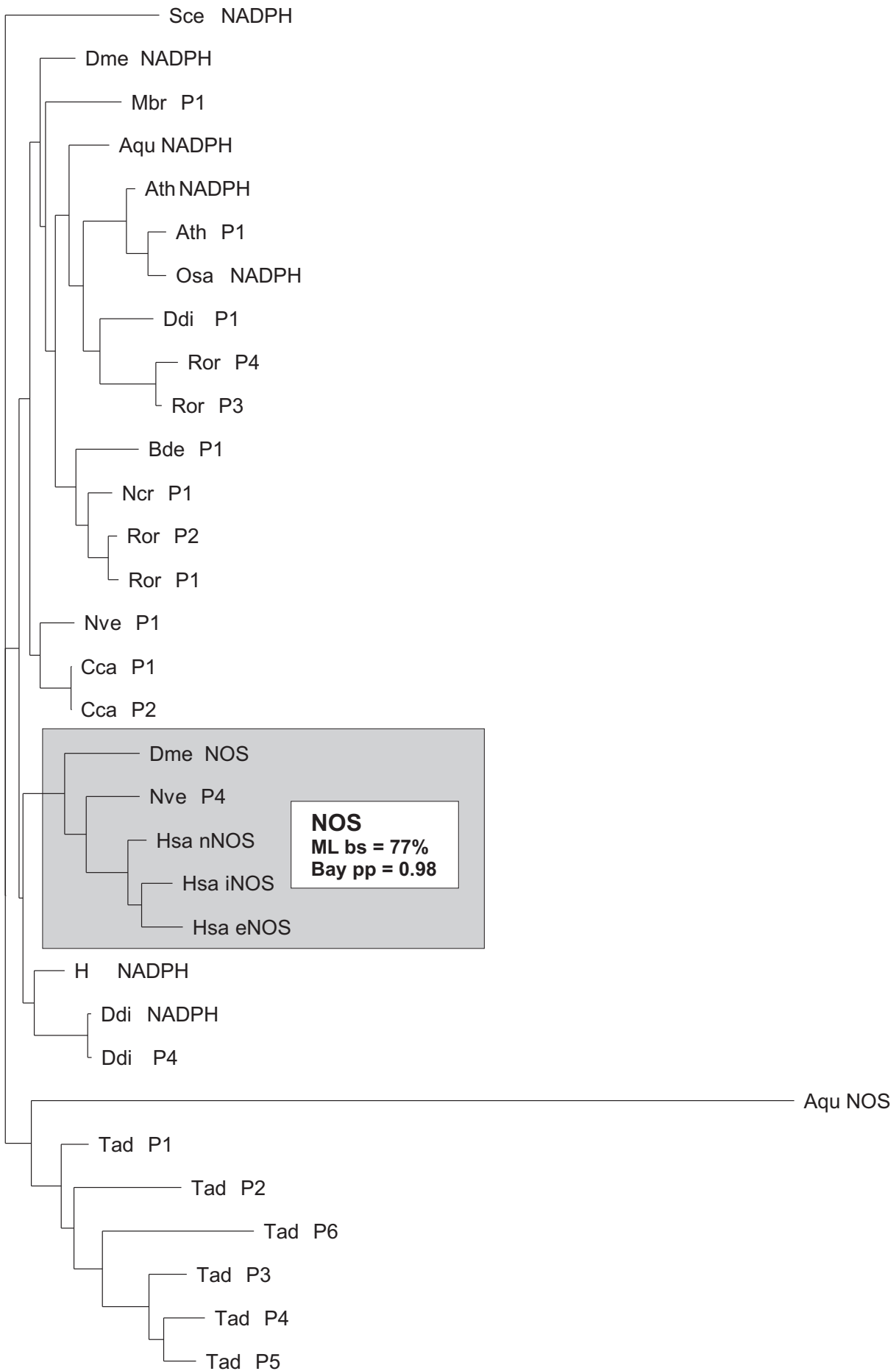
NOS partition A



NOS partition B



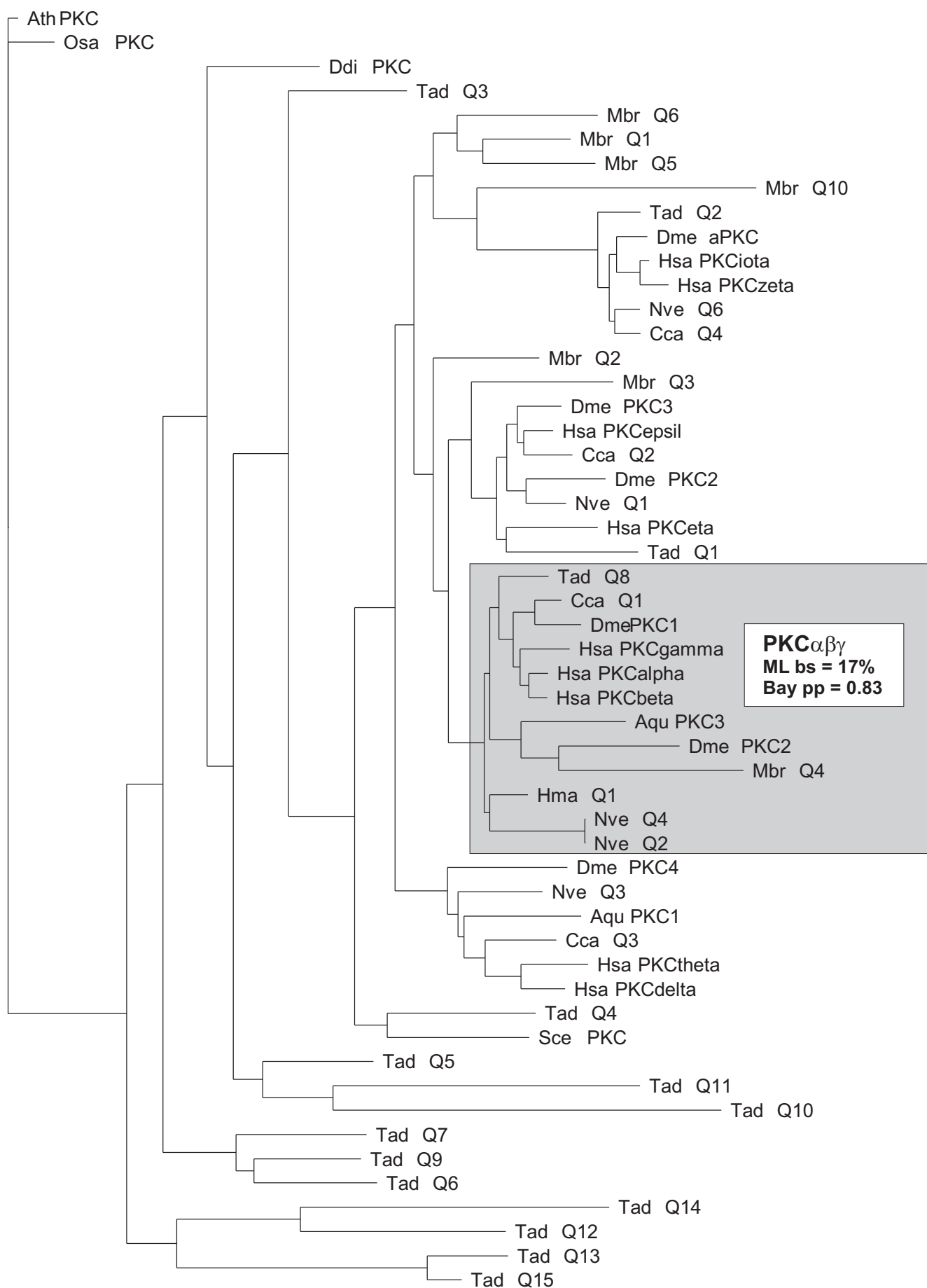
NOS partition C



PDZ domain

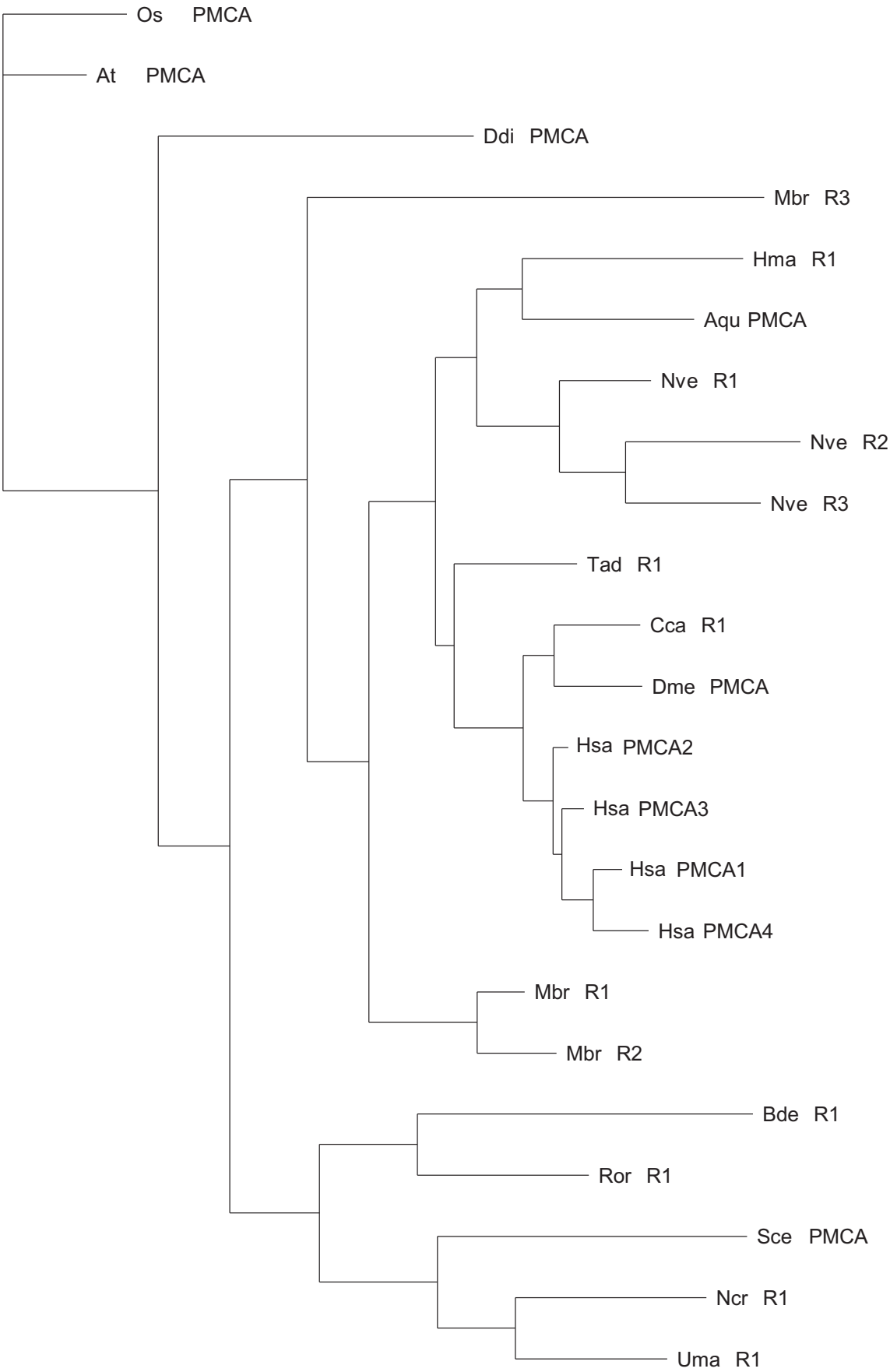


Protein Kinase C

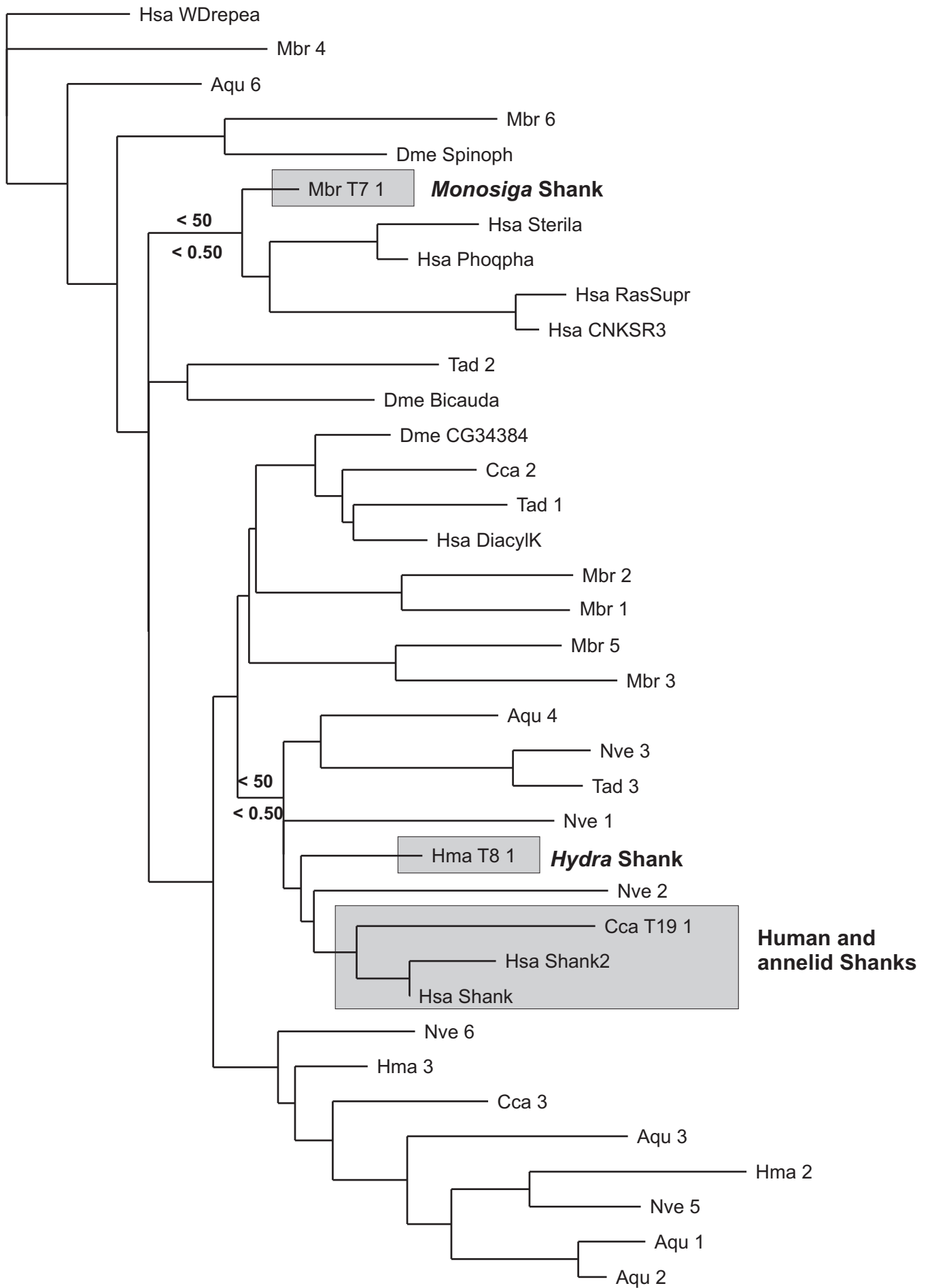


PMCA

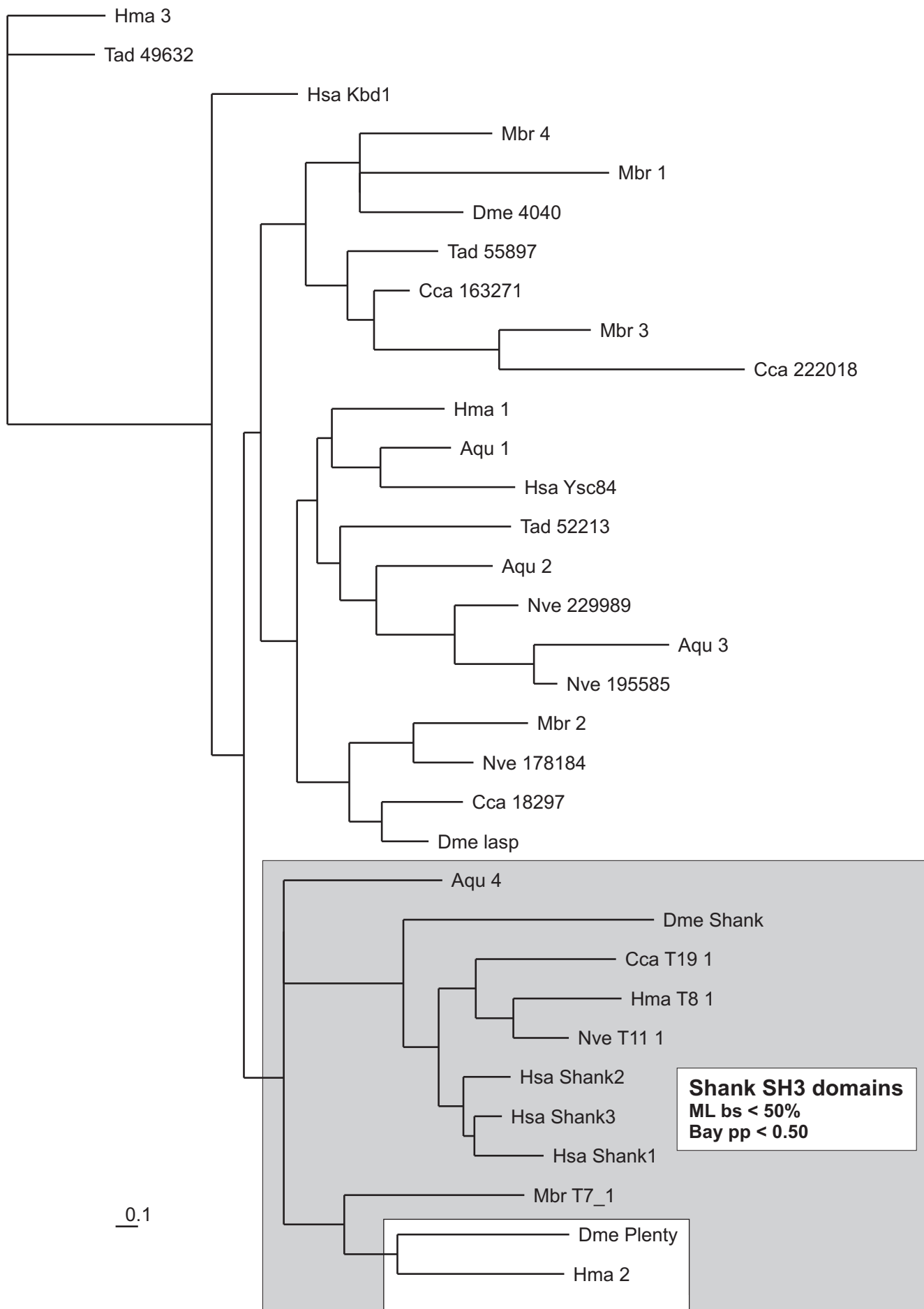
PMCA has no similarity to any other protein.



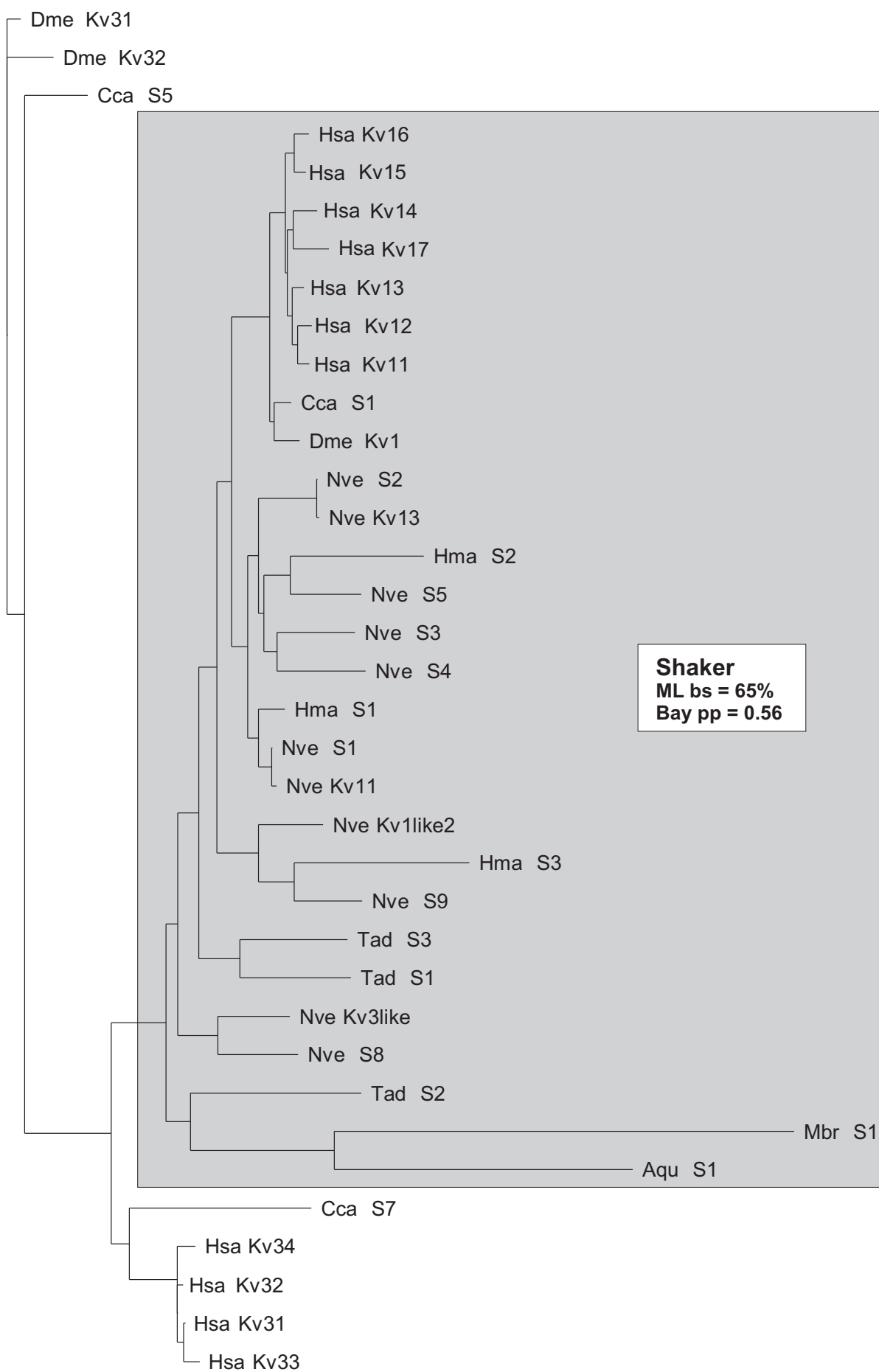
SAM domain



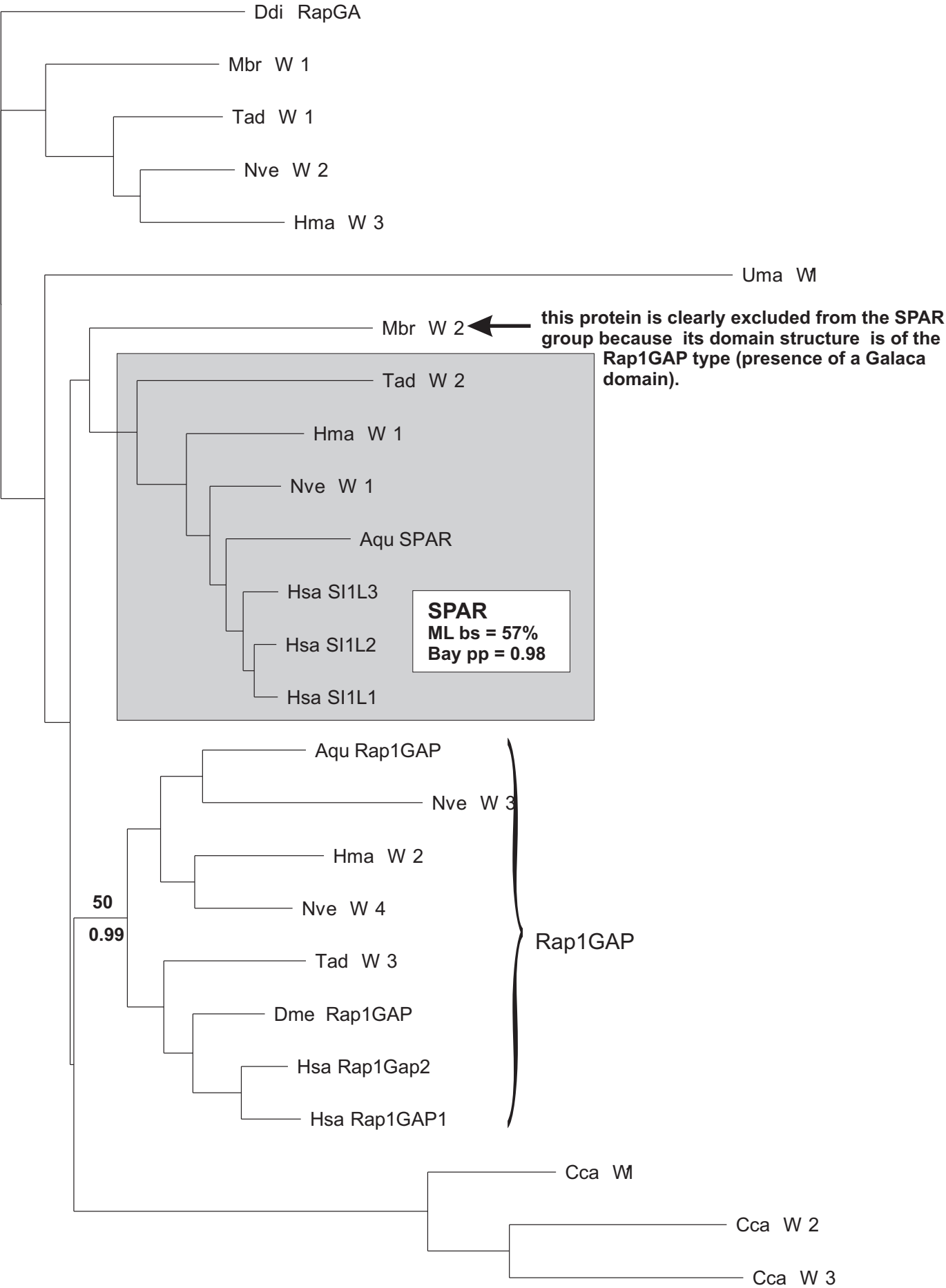
SH3 domain



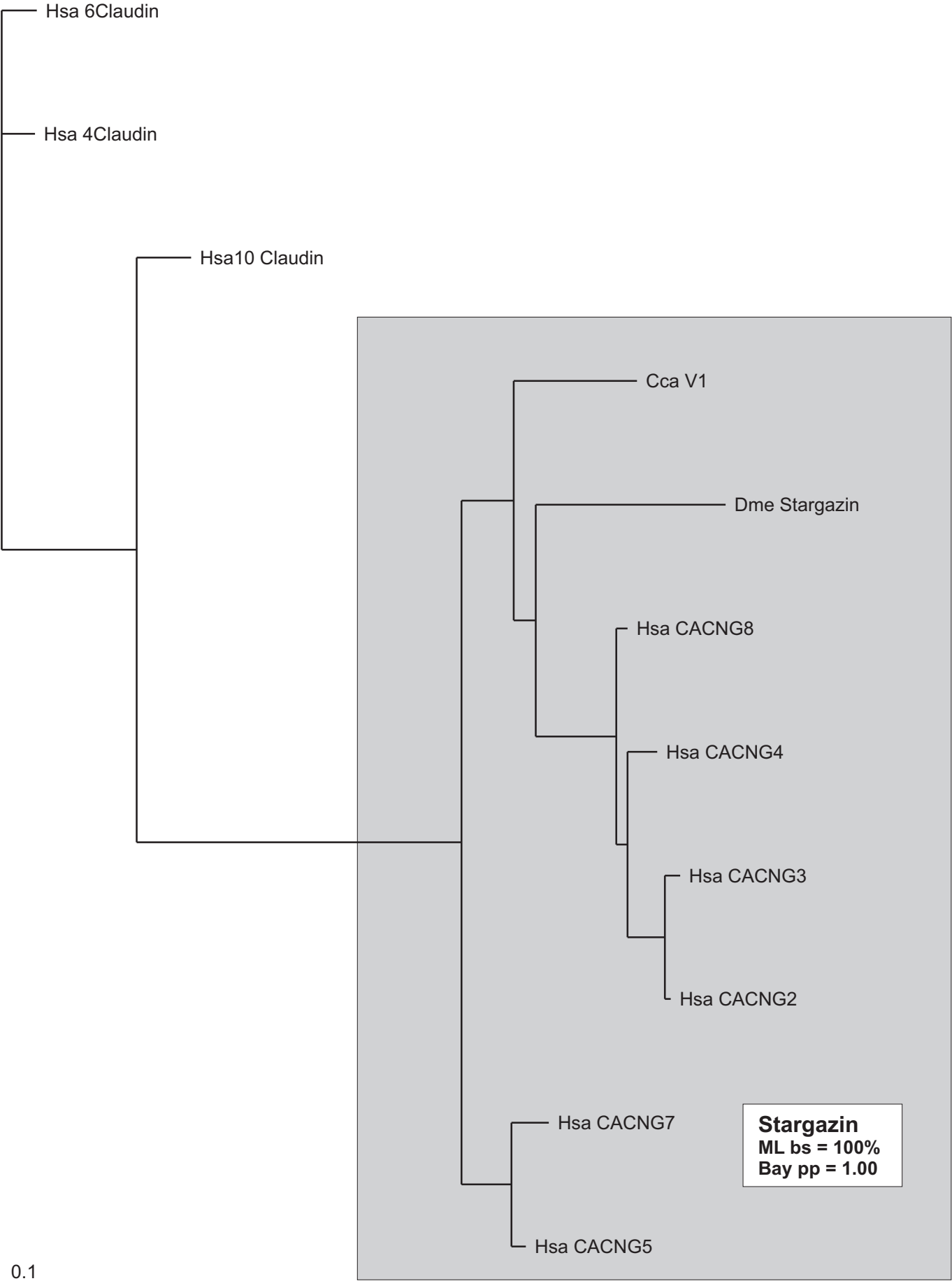
K⁺ channel Shaker



SPAR



Stargazin



SynGAP

