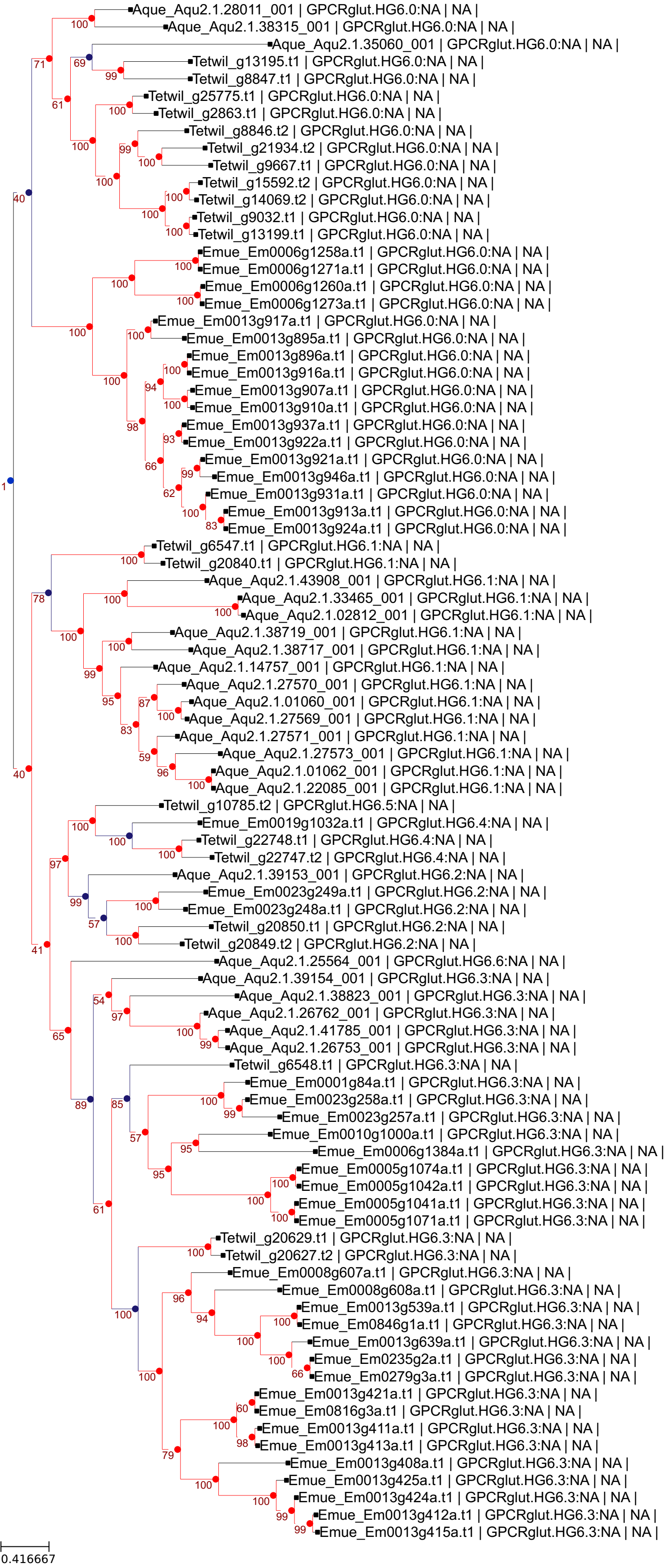
Aque_Aqu2.1.28011_001 | GPCRglut.HG6.0:NA | NA |
Aque_Aqu2.1.38315_001 | GPCRglut.HG6.0:NA | NA |
Aque_Aqu2.1.35060_001 | GPCRglut.HG6.0:NA | NA |
Tetwil_g13195.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g8847.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g25775.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g2863.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g8846.t2 | GPCRglut.HG6.0:NA | NA |
Tetwil_g21934.t2 | GPCRglut.HG6.0:NA | NA |
Tetwil_g9667.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g15592.t2 | GPCRglut.HG6.0:NA | NA |
Tetwil_g14069.t2 | GPCRglut.HG6.0:NA | NA |
Tetwil_g9032.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g13199.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0006g1258a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0006g1271a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0006g1260a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0006g1273a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g917a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g895a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g896a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g916a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g907a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g910a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g937a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g922a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g921a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g946a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g931a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g913a.t1 | GPCRglut.HG6.0:NA | NA |
Emue_Em0013g924a.t1 | GPCRglut.HG6.0:NA | NA |
Tetwil_g6547.t1 | GPCRglut.HG6.1:NA | NA |
Tetwil_g20840.t1 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.43908_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.33465_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.02812_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.38719_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.38717_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.14757_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.27570_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.01060_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.27569_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.27571_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.27573_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.01062_001 | GPCRglut.HG6.1:NA | NA |
Aque_Aqu2.1.22085_001 | GPCRglut.HG6.1:NA | NA |
Tetwil_g10785.t2 | GPCRglut.HG6.5:NA | NA |
Emue_Em0019g1032a.t1 | GPCRglut.HG6.4:NA | NA |
Tetwil_g22748.t1 | GPCRglut.HG6.4:NA | NA |
Tetwil_g22747.t2 | GPCRglut.HG6.4:NA | NA |
Aque_Aqu2.1.39153_001 | GPCRglut.HG6.2:NA | NA |
Emue_Em0023g249a.t1 | GPCRglut.HG6.2:NA | NA |
Emue_Em0023g248a.t1 | GPCRglut.HG6.2:NA | NA |
Tetwil_g20850.t1 | GPCRglut.HG6.2:NA | NA |
Tetwil_g20849.t2 | GPCRglut.HG6.2:NA | NA |
Aque_Aqu2.1.25564_001 | GPCRglut.HG6.6:NA | NA |
Aque_Aqu2.1.39154_001 | GPCRglut.HG6.3:NA | NA |
Aque_Aqu2.1.38823_001 | GPCRglut.HG6.3:NA | NA |
Aque_Aqu2.1.26762_001 | GPCRglut.HG6.3:NA | NA |
Aque_Aqu2.1.41785_001 | GPCRglut.HG6.3:NA | NA |
Aque_Aqu2.1.26753_001 | GPCRglut.HG6.3:NA | NA |
Tetwil_g6548.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0001g84a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0023g258a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0023g257a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0010g1000a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0006g1384a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0005g1074a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0005g1042a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0005g1041a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0005g1071a.t1 | GPCRglut.HG6.3:NA | NA |
Tetwil_g20629.t1 | GPCRglut.HG6.3:NA | NA |
Tetwil_g20627.t2 | GPCRglut.HG6.3:NA | NA |
Emue_Em0008g607a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0008g608a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g539a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0846g1a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g639a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0235g2a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0279g3a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g421a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0816g3a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g411a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g413a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g408a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g425a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g424a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g412a.t1 | GPCRglut.HG6.3:NA | NA |
Emue_Em0013g415a.t1 | GPCRglut.HG6.3:NA | NA |
0.416667