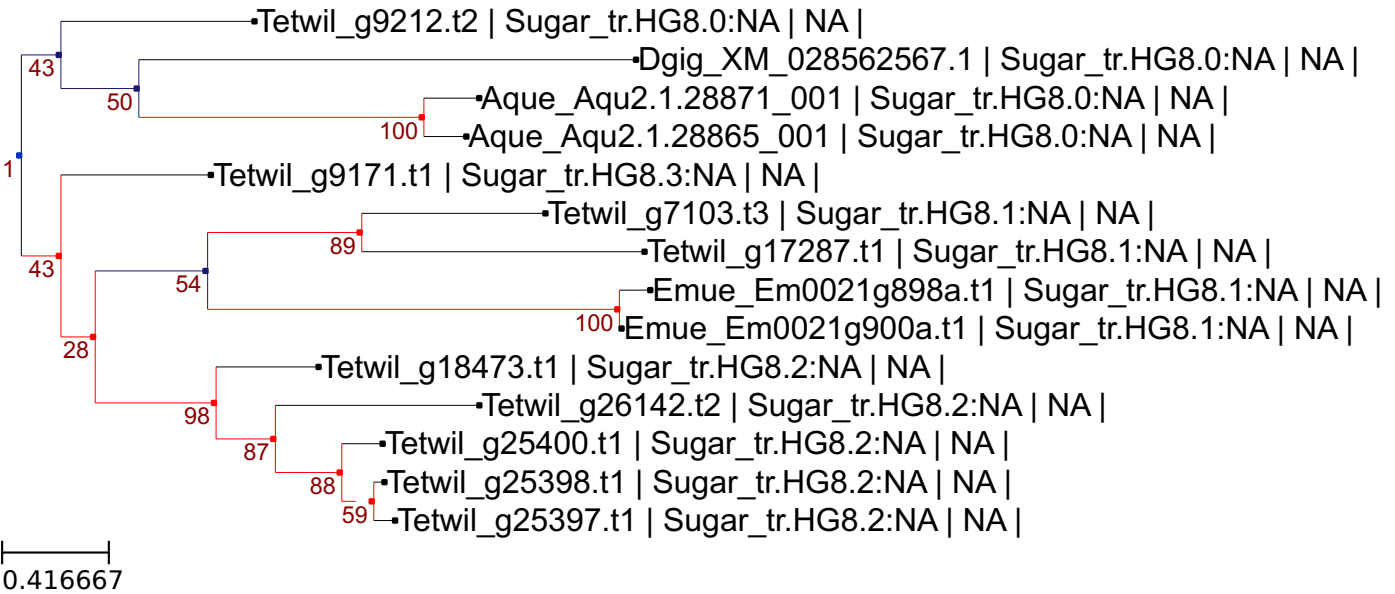
Tetwil_g9212.t2 | Sugar_tr.HG8.0:NA | NA |
Dgig_XM_028562567.1 | Sugar_tr.HG8.0:NA | NA |
Aque_Aqu2.1.28871_001 | Sugar_tr.HG8.0:NA | NA |
Aque_Aqu2.1.28865_001 | Sugar_tr.HG8.0:NA | NA |
Tetwil_g9171.t1 | Sugar_tr.HG8.3:NA | NA |
Tetwil_g7103.t3 | Sugar_tr.HG8.1:NA | NA |
Tetwil_g17287.t1 | Sugar_tr.HG8.1:NA | NA |
Emue_Em0021g898a.t1 | Sugar_tr.HG8.1:NA | NA |
Emue_Em0021g900a.t1 | Sugar_tr.HG8.1:NA | NA |
Tetwil_g18473.t1 | Sugar_tr.HG8.2:NA | NA |
Tetwil_g26142.t2 | Sugar_tr.HG8.2:NA | NA |
Tetwil_g25400.t1 | Sugar_tr.HG8.2:NA | NA |
Tetwil_g25398.t1 | Sugar_tr.HG8.2:NA | NA |
Tetwil_g25397.t1 | Sugar_tr.HG8.2:NA | NA |
43
50
100
1
43
89
54
100
28
98
87
88
59
0.416667