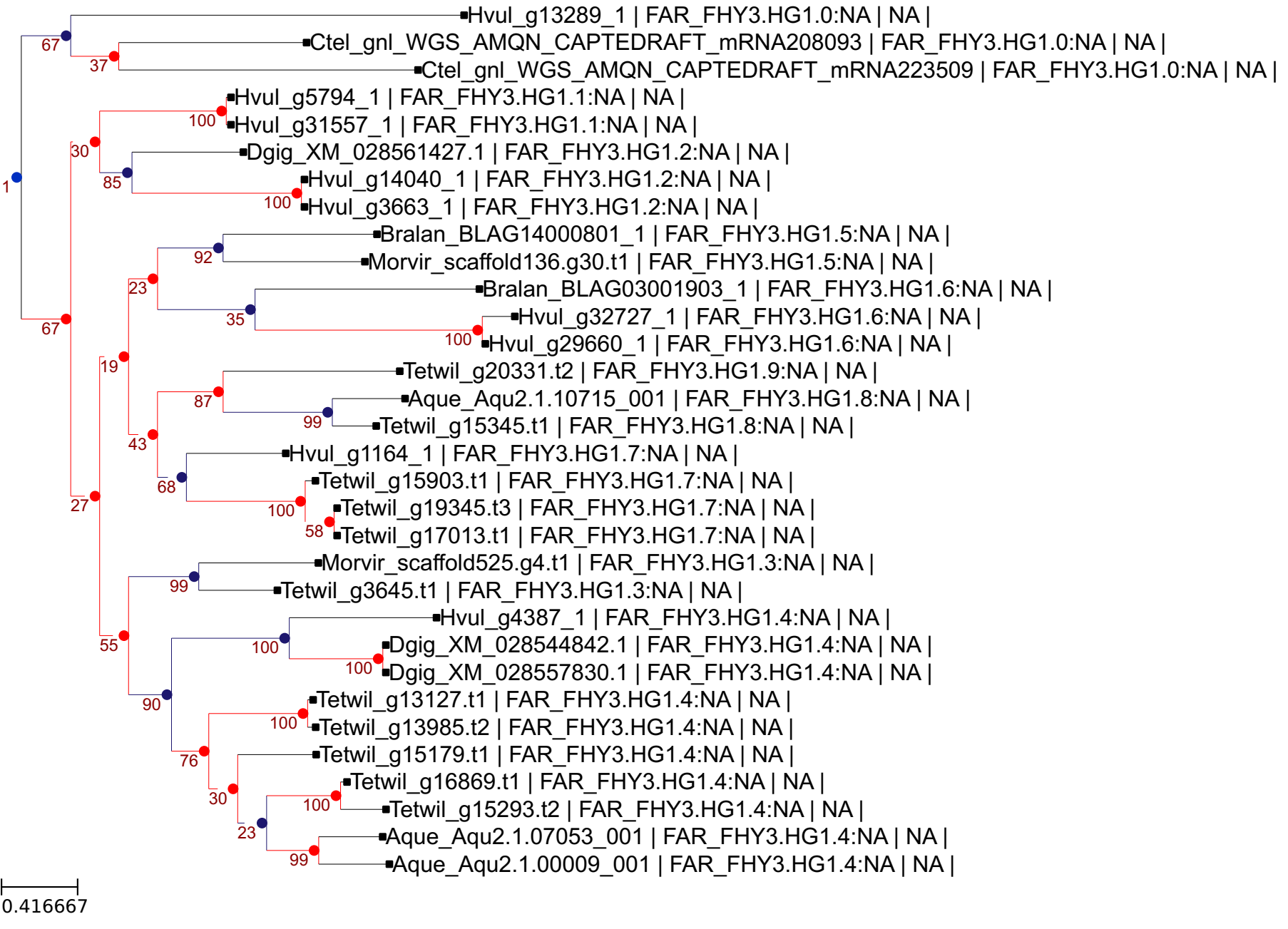
Hvul_g13289_1 | FAR_FHY3.HG1.0:NA | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA208093 | FAR_FHY3.HG1.0:NA | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA223509 | FAR_FHY3.HG1.0:NA | NA |
Hvul_g5794_1 | FAR_FHY3.HG1.1:NA | NA |
Hvul_g31557_1 | FAR_FHY3.HG1.1:NA | NA |
Dgig_XM_028561427.1 | FAR_FHY3.HG1.2:NA | NA |
Hvul_g14040_1 | FAR_FHY3.HG1.2:NA | NA |
Hvul_g3663_1 | FAR_FHY3.HG1.2:NA | NA |
Bralan_BLAG14000801_1 | FAR_FHY3.HG1.5:NA | NA |
Morvir_scaffold136.g30.t1 | FAR_FHY3.HG1.5:NA | NA |
Bralan_BLAG03001903_1 | FAR_FHY3.HG1.6:NA | NA |
Hvul_g32727_1 | FAR_FHY3.HG1.6:NA | NA |
Hvul_g29660_1 | FAR_FHY3.HG1.6:NA | NA |
Tetwil_g20331.t2 | FAR_FHY3.HG1.9:NA | NA |
Aque_Aqu2.1.10715_001 | FAR_FHY3.HG1.8:NA | NA |
Tetwil_g15345.t1 | FAR_FHY3.HG1.8:NA | NA |
Hvul_g1164_1 | FAR_FHY3.HG1.7:NA | NA |
Tetwil_g15903.t1 | FAR_FHY3.HG1.7:NA | NA |
Tetwil_g19345.t3 | FAR_FHY3.HG1.7:NA | NA |
Tetwil_g17013.t1 | FAR_FHY3.HG1.7:NA | NA |
Morvir_scaffold525.g4.t1 | FAR_FHY3.HG1.3:NA | NA |
Tetwil_g3645.t1 | FAR_FHY3.HG1.3:NA | NA |
Hvul_g4387_1 | FAR_FHY3.HG1.4:NA | NA |
Dgig_XM_028544842.1 | FAR_FHY3.HG1.4:NA | NA |
Dgig_XM_028557830.1 | FAR_FHY3.HG1.4:NA | NA |
Tetwil_g13127.t1 | FAR_FHY3.HG1.4:NA | NA |
Tetwil_g13985.t2 | FAR_FHY3.HG1.4:NA | NA |
Tetwil_g15179.t1 | FAR_FHY3.HG1.4:NA | NA |
Tetwil_g16869.t1 | FAR_FHY3.HG1.4:NA | NA |
Tetwil_g15293.t2 | FAR_FHY3.HG1.4:NA | NA |
Aque_Aqu2.1.07053_001 | FAR_FHY3.HG1.4:NA | NA |
Aque_Aqu2.1.00009_001 | FAR_FHY3.HG1.4:NA | NA |
0.416667