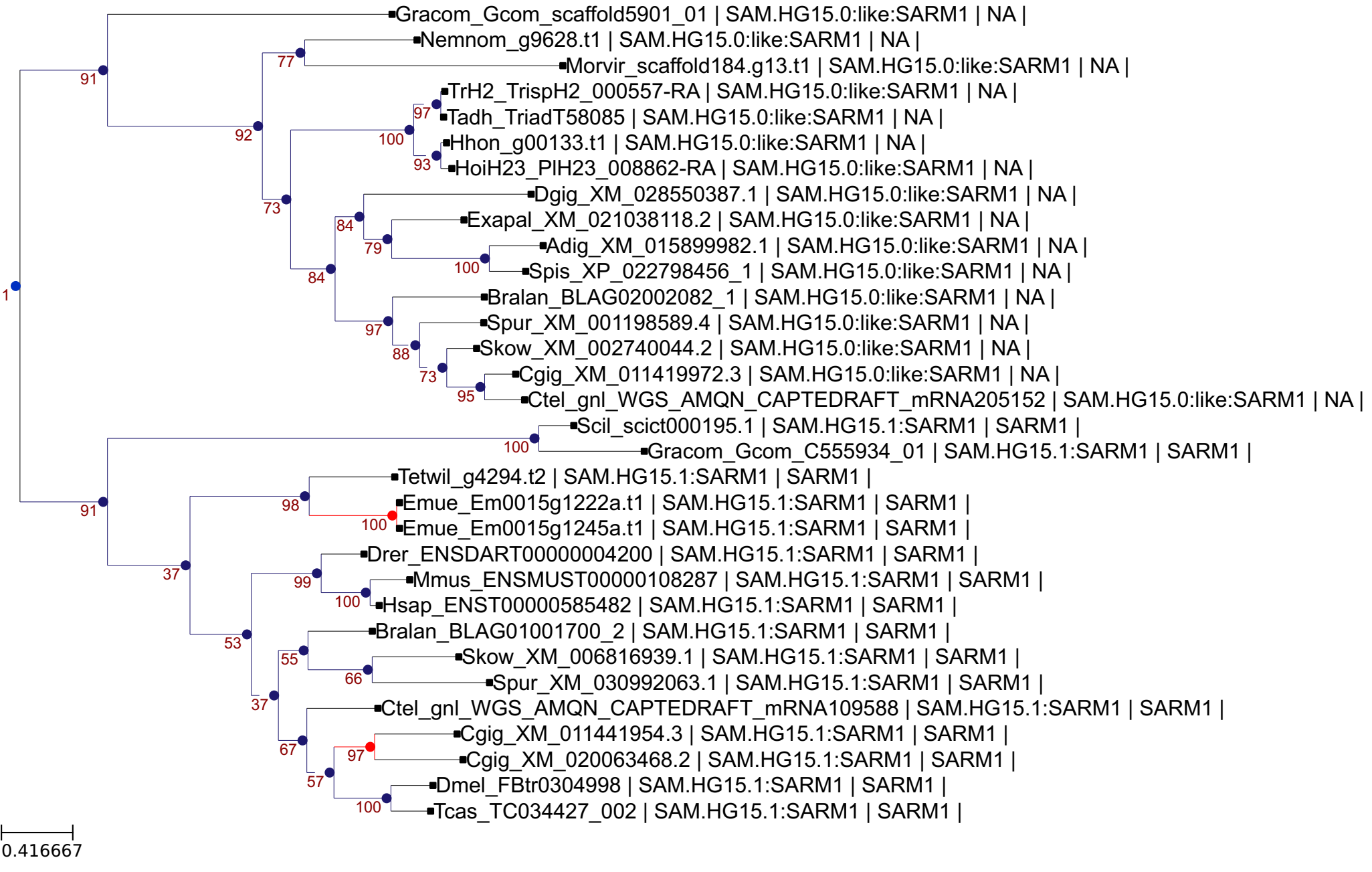
Gracom_Gcom_scaffold5901_01 | SAM.HG15.0:like:SARM1 | NA |
Nemnom_g9628.t1 | SAM.HG15.0:like:SARM1 | NA |
Morvir_scaffold184.g13.t1 | SAM.HG15.0:like:SARM1 | NA |
TrH2_TrispH2_000557-RA | SAM.HG15.0:like:SARM1 | NA |
Tadh_TriadT58085 | SAM.HG15.0:like:SARM1 | NA |
Hhon_g00133.t1 | SAM.HG15.0:like:SARM1 | NA |
HoiH23_PlH23_008862-RA | SAM.HG15.0:like:SARM1 | NA |
Dgig_XM_028550387.1 | SAM.HG15.0:like:SARM1 | NA |
Exapal_XM_021038118.2 | SAM.HG15.0:like:SARM1 | NA |
Adig_XM_015899982.1 | SAM.HG15.0:like:SARM1 | NA |
Spis_XP_022798456_1 | SAM.HG15.0:like:SARM1 | NA |
Bralan_BLAG02002082_1 | SAM.HG15.0:like:SARM1 | NA |
Spur_XM_001198589.4 | SAM.HG15.0:like:SARM1 | NA |
Skow_XM_002740044.2 | SAM.HG15.0:like:SARM1 | NA |
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Scil_scict000195.1 | SAM.HG15.1:SARM1 | SARM1 |
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Tetwil_g4294.t2 | SAM.HG15.1:SARM1 | SARM1 |
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