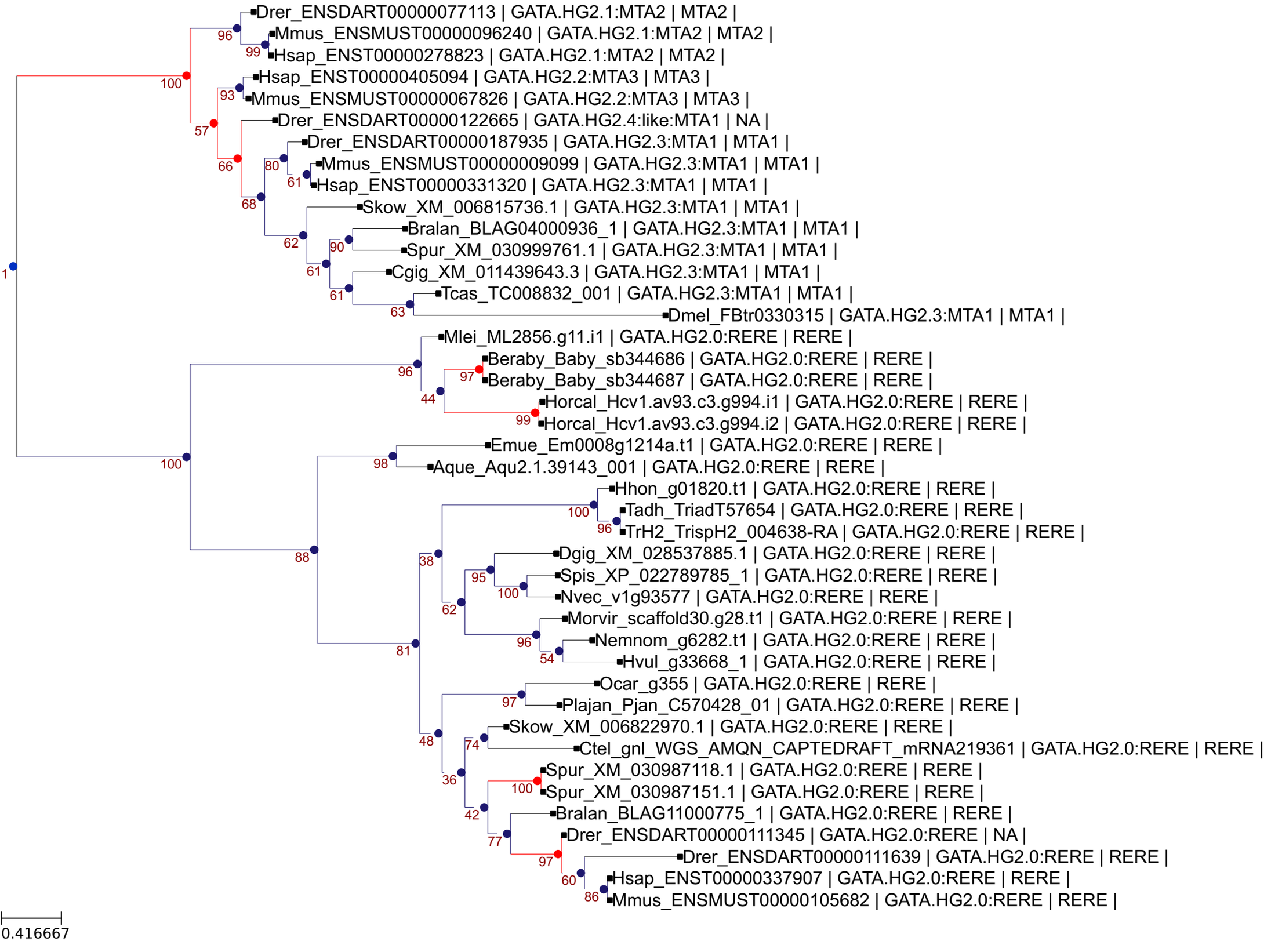
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Mmus_ENSMUST00000096240 | GATA.HG2.1:MTA2 | MTA2 |
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Hsap_ENST00000405094 | GATA.HG2.2:MTA3 | MTA3 |
Mmus_ENSMUST00000067826 | GATA.HG2.2:MTA3 | MTA3 |
Drer_ENSDART00000122665 | GATA.HG2.4:like:MTA1 | NA |
Drer_ENSDART00000187935 | GATA.HG2.3:MTA1 | MTA1 |
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Hsap_ENST00000331320 | GATA.HG2.3:MTA1 | MTA1 |
Skow_XM_006815736.1 | GATA.HG2.3:MTA1 | MTA1 |
Bralan_BLAG04000936_1 | GATA.HG2.3:MTA1 | MTA1 |
Spur_XM_030999761.1 | GATA.HG2.3:MTA1 | MTA1 |
Cgig_XM_011439643.3 | GATA.HG2.3:MTA1 | MTA1 |
Tcas_TC008832_001 | GATA.HG2.3:MTA1 | MTA1 |
Dmel_FBtr0330315 | GATA.HG2.3:MTA1 | MTA1 |
Mlei_ML2856.g11.i1 | GATA.HG2.0:RERE | RERE |
Beraby_Baby_sb344686 | GATA.HG2.0:RERE | RERE |
Beraby_Baby_sb344687 | GATA.HG2.0:RERE | RERE |
Horcal_Hcv1.av93.c3.g994.i1 | GATA.HG2.0:RERE | RERE |
Horcal_Hcv1.av93.c3.g994.i2 | GATA.HG2.0:RERE | RERE |
Emue_Em0008g1214a.t1 | GATA.HG2.0:RERE | RERE |
Aque_Aqu2.1.39143_001 | GATA.HG2.0:RERE | RERE |
Hhon_g01820.t1 | GATA.HG2.0:RERE | RERE |
Tadh_TriadT57654 | GATA.HG2.0:RERE | RERE |
TrH2_TrispH2_004638-RA | GATA.HG2.0:RERE | RERE |
Dgig_XM_028537885.1 | GATA.HG2.0:RERE | RERE |
Spis_XP_022789785_1 | GATA.HG2.0:RERE | RERE |
Nvec_v1g93577 | GATA.HG2.0:RERE | RERE |
Morvir_scaffold30.g28.t1 | GATA.HG2.0:RERE | RERE |
Nemnom_g6282.t1 | GATA.HG2.0:RERE | RERE |
Hvul_g33668_1 | GATA.HG2.0:RERE | RERE |
Ocar_g355 | GATA.HG2.0:RERE | RERE |
Plajan_Pjan_C570428_01 | GATA.HG2.0:RERE | RERE |
Skow_XM_006822970.1 | GATA.HG2.0:RERE | RERE |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA219361 | GATA.HG2.0:RERE | RERE |
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Spur_XM_030987151.1 | GATA.HG2.0:RERE | RERE |
Bralan_BLAG11000775_1 | GATA.HG2.0:RERE | RERE |
Drer_ENSDART00000111345 | GATA.HG2.0:RERE | NA |
Drer_ENSDART00000111639 | GATA.HG2.0:RERE | RERE |
Hsap_ENST00000337907 | GATA.HG2.0:RERE | RERE |
Mmus_ENSMUST00000105682 | GATA.HG2.0:RERE | RERE |
0.416667