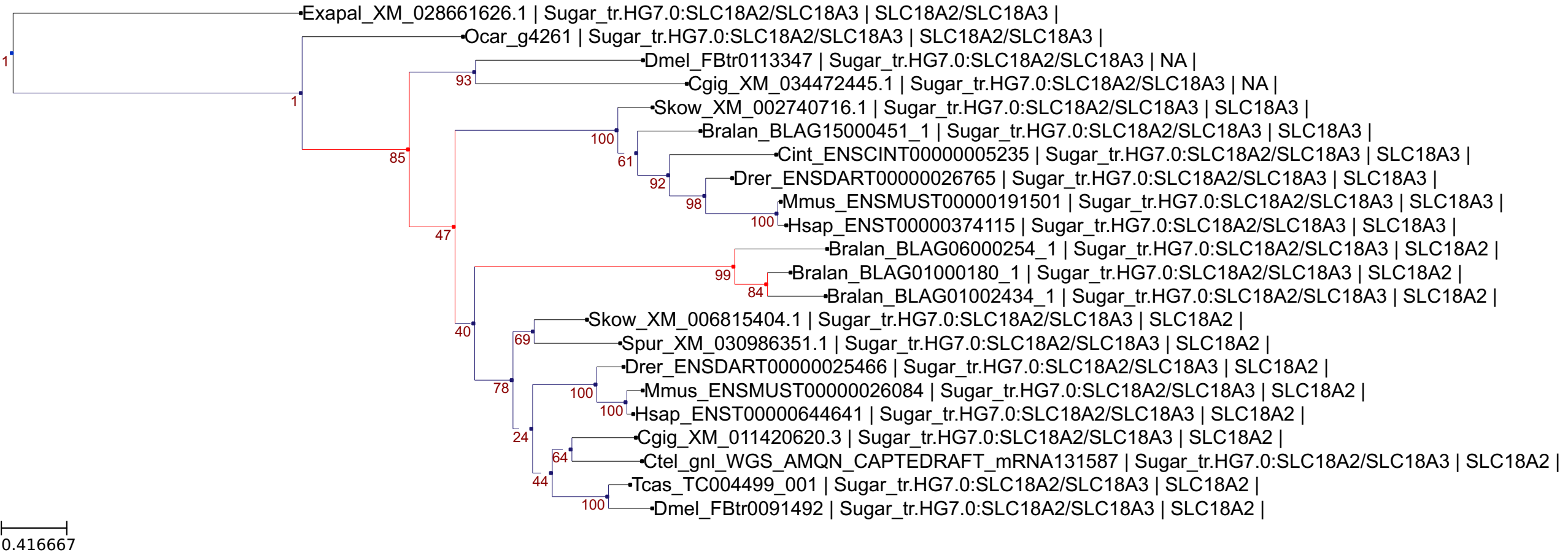
Exapal_XM_028661626.1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2/SLC18A3 |
Ocar_g4261 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2/SLC18A3 |
Dmel_FBtr0113347 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | NA |
Cgig_XM_034472445.1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | NA |
Skow_XM_002740716.1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Bralan_BLAG15000451_1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Cint_ENSCINT00000005235 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Drer_ENSDART00000026765 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Mmus_ENSMUST00000191501 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Hsap_ENST00000374115 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A3 |
Bralan_BLAG06000254_1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Bralan_BLAG01000180_1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Bralan_BLAG01002434_1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Skow_XM_006815404.1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Spur_XM_030986351.1 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Drer_ENSDART00000025466 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Mmus_ENSMUST00000026084 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Hsap_ENST00000644641 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Cgig_XM_011420620.3 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA131587 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Tcas_TC004499_001 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
Dmel_FBtr0091492 | Sugar_tr.HG7.0:SLC18A2/SLC18A3 | SLC18A2 |
1
1
93
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