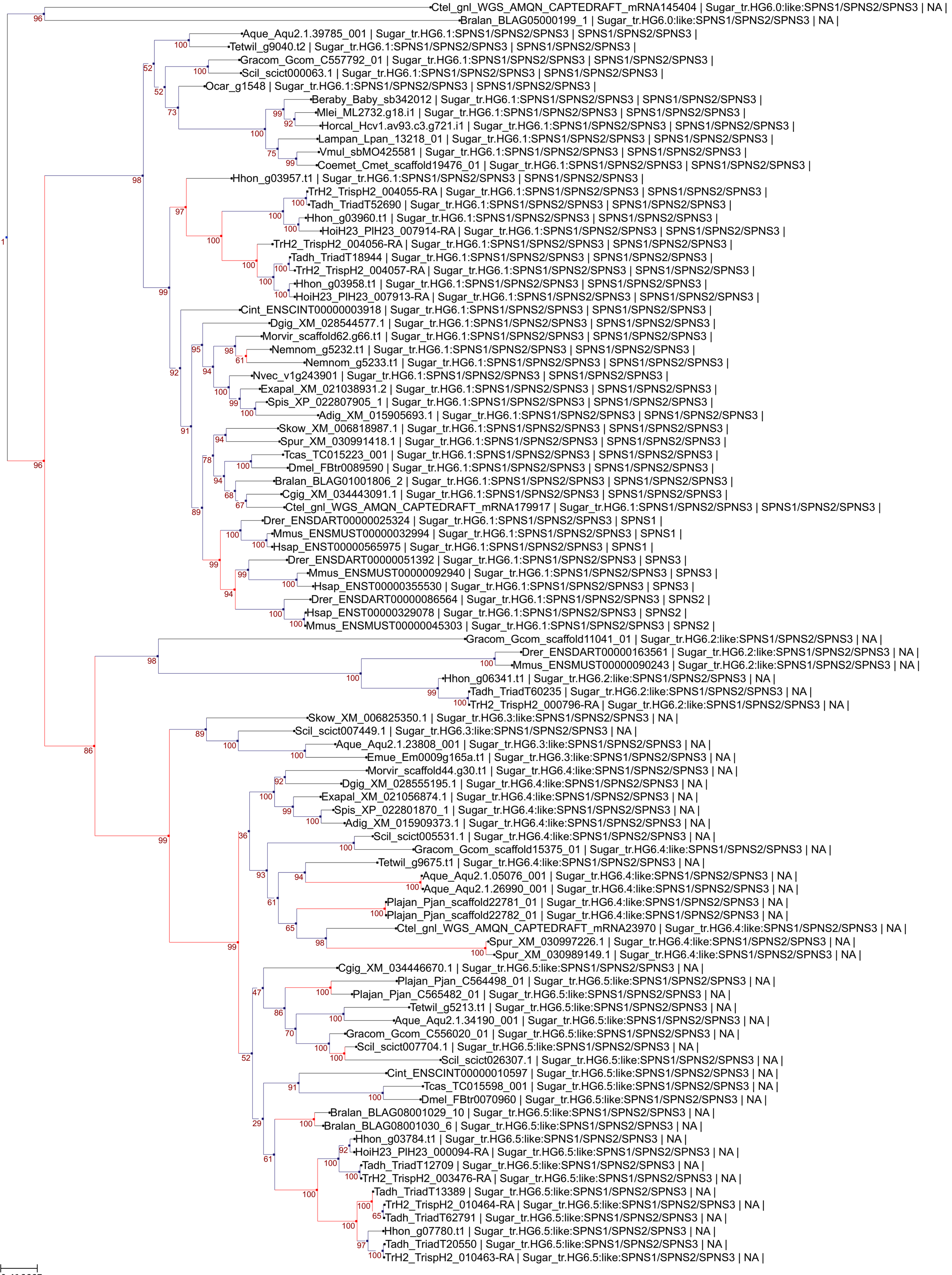
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA145404 | Sugar_tr.HG6.0:like:SPNS1/SPNS2/SPNS3 | NA |
Bralan_BLAG05000199_1 | Sugar_tr.HG6.0:like:SPNS1/SPNS2/SPNS3 | NA |
Aque_Aqu2.1.39785_001 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Tetwil_g9040.t2 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Gracom_Gcom_C557792_01 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Scil_scict000063.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Ocar_g1548 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Beraby_Baby_sb342012 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Mlei_ML2732.g18.i1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Horcal_Hcv1.av93.c3.g721.i1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Lampan_Lpan_13218_01 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Vmul_sbMO425581 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Coemet_Cmet_scaffold19476_01 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Hhon_g03957.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
TrH2_TrispH2_004055-RA | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Tadh_TriadT52690 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Hhon_g03960.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
HoiH23_PlH23_007914-RA | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
TrH2_TrispH2_004056-RA | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Tadh_TriadT18944 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
TrH2_TrispH2_004057-RA | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Hhon_g03958.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
HoiH23_PlH23_007913-RA | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Cint_ENSCINT00000003918 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Dgig_XM_028544577.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Morvir_scaffold62.g66.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Nemnom_g5232.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Nemnom_g5233.t1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Nvec_v1g243901 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Exapal_XM_021038931.2 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Spis_XP_022807905_1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Adig_XM_015905693.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Skow_XM_006818987.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Spur_XM_030991418.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Tcas_TC015223_001 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Dmel_FBtr0089590 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Bralan_BLAG01001806_2 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Cgig_XM_034443091.1 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA179917 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1/SPNS2/SPNS3 |
Drer_ENSDART00000025324 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1 |
Mmus_ENSMUST00000032994 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1 |
Hsap_ENST00000565975 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS1 |
Drer_ENSDART00000051392 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS3 |
Mmus_ENSMUST00000092940 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS3 |
Hsap_ENST00000355530 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS3 |
Drer_ENSDART00000086564 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS2 |
Hsap_ENST00000329078 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS2 |
Mmus_ENSMUST00000045303 | Sugar_tr.HG6.1:SPNS1/SPNS2/SPNS3 | SPNS2 |
Gracom_Gcom_scaffold11041_01 | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
Drer_ENSDART00000163561 | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
Mmus_ENSMUST00000090243 | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
Hhon_g06341.t1 | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
Tadh_TriadT60235 | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
TrH2_TrispH2_000796-RA | Sugar_tr.HG6.2:like:SPNS1/SPNS2/SPNS3 | NA |
Skow_XM_006825350.1 | Sugar_tr.HG6.3:like:SPNS1/SPNS2/SPNS3 | NA |
Scil_scict007449.1 | Sugar_tr.HG6.3:like:SPNS1/SPNS2/SPNS3 | NA |
Aque_Aqu2.1.23808_001 | Sugar_tr.HG6.3:like:SPNS1/SPNS2/SPNS3 | NA |
Emue_Em0009g165a.t1 | Sugar_tr.HG6.3:like:SPNS1/SPNS2/SPNS3 | NA |
Morvir_scaffold44.g30.t1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Dgig_XM_028555195.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Exapal_XM_021056874.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Spis_XP_022801870_1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Adig_XM_015909373.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Scil_scict005531.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Gracom_Gcom_scaffold15375_01 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Tetwil_g9675.t1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Aque_Aqu2.1.05076_001 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Aque_Aqu2.1.26990_001 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Plajan_Pjan_scaffold22781_01 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Plajan_Pjan_scaffold22782_01 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA23970 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Spur_XM_030997226.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Spur_XM_030989149.1 | Sugar_tr.HG6.4:like:SPNS1/SPNS2/SPNS3 | NA |
Cgig_XM_034446670.1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Plajan_Pjan_C564498_01 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Plajan_Pjan_C565482_01 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tetwil_g5213.t1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Aque_Aqu2.1.34190_001 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Gracom_Gcom_C556020_01 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Scil_scict007704.1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Scil_scict026307.1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Cint_ENSCINT00000010597 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tcas_TC015598_001 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Dmel_FBtr0070960 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Bralan_BLAG08001029_10 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Bralan_BLAG08001030_6 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Hhon_g03784.t1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
HoiH23_PlH23_000094-RA | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tadh_TriadT12709 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
TrH2_TrispH2_003476-RA | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tadh_TriadT13389 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
TrH2_TrispH2_010464-RA | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tadh_TriadT62791 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Hhon_g07780.t1 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
Tadh_TriadT20550 | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
TrH2_TrispH2_010463-RA | Sugar_tr.HG6.5:like:SPNS1/SPNS2/SPNS3 | NA |
0.416667