Hvul_g29345_1 | Ras.HG9.10:like:RAB29/RAB32/RAB38 | NA |
Cint_ENSCINT00000027450 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Tetwil_g12088.t2 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Lampan_Lpan_C415081_01 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Horcal_Hcv1.av93.c8.g296.i1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Horcal_Hcv1.av93.c8.g296.i2 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Bralan_BLAG01000082_1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Spur_XM_786252.5 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA208667 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Hvul_g17787_1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Nemnom_g3289.t1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Morvir_scaffold21.g140.t2 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Dgig_XM_028554142.1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Adig_XM_015903635.1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Spis_XP_022781357_1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Nvec_v1g202545 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Exapal_XM_021041012.2 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Plajan_Pjan_C560037_01 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Ocar_g4624 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Cgig_XM_034475056.1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Cgig_XM_034477295.1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Tadh_TriadT56305 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
TrH2_TrispH2_001078-RA | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Hhon_g05105.t1 | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
HoiH23_PIH23_003149-RA | Ras.HG9.0:like:RAB29/RAB32/RAB38 | NA |
Dgig_XM_028547321.1 | Ras.HG9.1:like:RAB29/RAB32/RAB38 | NA |
Adig_XM_015918071.1 | Ras.HG9.1:like:RAB29/RAB32/RAB38 | NA |
Spis_XP_022801655_1 | Ras.HG9.1:like:RAB29/RAB32/RAB38 | NA |
Nvec_v1g64028 | Ras.HG9.1:like:RAB29/RAB32/RAB38 | NA |
Exapal_XM_021037365.2 | Ras.HG9.1:like:RAB29/RAB32/RAB38 | NA |
Spis_XP_022801649_1 | Ras.HG9.11:like:RAB29/RAB32/RAB38 | NA |
Exapal_XM_021057674.2 | Ras.HG9.2:like:RAB29/RAB32/RAB38 | NA |
Adig_XM_015918075.1 | Ras.HG9.2:like:RAB29/RAB32/RAB38 | NA |
Spis_XP_022801644_1 | Ras.HG9.2:like:RAB29/RAB32/RAB38 | NA |
Spis_XP_022801651_1 | Ras.HG9.2:like:RAB29/RAB32/RAB38 | NA |
Emue_Em0005g1604a.t1 | Ras.HG9.12:like:RAB29/RAB32/RAB38 | NA |
Gracom_Gcom_C548110_01 | Ras.HG9.13:like:RAB29 | NA |
Cgig_XM_034471109.1 | Ras.HG9.3:like:RAB29 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA110893 | Ras.HG9.3:like:RAB29 | NA |
Tetwil_g9141.t2 | Ras.HG9.3:like:RAB29 | NA |
Aque_Aqu2.1.37182_001 | Ras.HG9.3:like:RAB29 | NA |
Tcas_TC031107_001 | Ras.HG9.3:like:RAB29 | NA |
Spur_XM_011679160.2 | Ras.HG9.3:like:RAB29 | NA |
Bralan_BLAG04000272_1 | Ras.HG9.3:like:RAB29 | NA |
Gracom_Gcom_scaffold17373_01 | Ras.HG9.3:like:RAB29 | NA |
Ocar_g7191 | Ras.HG9.3:like:RAB29 | NA |
Plajan_Pjan_scaffold22226_01 | Ras.HG9.3:like:RAB29 | NA |
Plajan_Pjan_scaffold22227_01 | Ras.HG9.3:like:RAB29 | NA |
Nvec_v1g245354 | Ras.HG9.3:like:RAB29 | NA |
Exapal_XM_021036222.2 | Ras.HG9.3:like:RAB29 | NA |
Exapal_XM_021057124.1 | Ras.HG9.3:like:RAB29 | NA |
Dgig_XM_028542754.1 | Ras.HG9.3:like:RAB29 | NA |
Spis_XP_022791009_1 | Ras.HG9.3:like:RAB29 | NA |
Adig_XM_015902886.1 | Ras.HG9.3:like:RAB29 | NA |
Coemet_Cmet_C592315_01 | Ras.HG9.4:like:RAB29 | NA |
Lampan_Lpan_6207_01 | Ras.HG9.4:like:RAB29 | NA |
Mlei_ML0191.g16.i1 | Ras.HG9.4:like:RAB29 | NA |
Bolinf_Binf_sb1252303 | Ras.HG9.4:like:RAB29 | NA |
Bolinf_Binf_sb1252304 | Ras.HG9.4:like:RAB29 | NA |
Plajan_Pjan_C537534_01 | Ras.HG9.4:like:RAB29 | NA |
Ocar_g9839 | Ras.HG9.4:like:RAB29 | NA |
Morvir_scaffold410.g16.t1 | Ras.HG9.4:like:RAB29 | NA |
Aque_Aqu2.1.42743_001 | Ras.HG9.4:like:RAB29 | NA |
Emue_Em0001g237a.t1 | Ras.HG9.4:like:RAB29 | NA |
Tetwil_g24168.t2 | Ras.HG9.4:like:RAB29 | NA |
Mmus_ENSMUST00000027693 | Ras.HG9.5:RAB29 | RAB29 |
Hsap_ENST00000235932 | Ras.HG9.5:RAB29 | RAB29 |
Plajan_Pjan_C545245_01 | Ras.HG9.5:RAB29 | RAB29 |
Plajan_Pjan_C543934_01 | Ras.HG9.5:RAB29 | RAB29 |
Aque_Aqu2.1.42742_001 | Ras.HG9.5:RAB29 | RAB29 |
Tetwil_g24167.t1 | Ras.HG9.5:RAB29 | RAB29 |
Spur_XM_011679006.2 | Ras.HG9.5:RAB29 | RAB29 |
Spur_XM_030986395.1 | Ras.HG9.5:RAB29 | RAB29 |
Bralan_BLAG05000531_1 | Ras.HG9.5:RAB29 | RAB29 |
Skow_XM_006825056.1 | Ras.HG9.5:RAB29 | RAB29 |
Scil_scict021640.4 | Ras.HG9.5:RAB29 | RAB29 |
Gracom_Gcom_scaffold14818_01 | Ras.HG9.5:RAB29 | RAB29 |
Cgig_XM_011437731.3 | Ras.HG9.5:RAB29 | RAB29 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA101663 | Ras.HG9.5:RAB29 | RAB29 |
Dgig_XM_028542239.1 | Ras.HG9.5:RAB29 | RAB29 |
Nvec_v1g200891 | Ras.HG9.5:RAB29 | RAB29 |
Spis_XP_022784498_1 | Ras.HG9.5:RAB29 | RAB29 |
Adig_XM_015917531.1 | Ras.HG9.5:RAB29 | RAB29 |
Adig_XM_015898282.1 | Ras.HG9.5:RAB29 | RAB29 |
Exapal_XM_021049409.2 | Ras.HG9.5:RAB29 | RAB29 |
Exapal_XM_021053777.2 | Ras.HG9.5:RAB29 | RAB29 |
Nvec_v1g84449 | Ras.HG9.5:RAB29 | RAB29 |
Nvec_v1g84100 | Ras.HG9.5:RAB29 | RAB29 |
Nvec_v1g238852 | Ras.HG9.5:RAB29 | RAB29 |
HoiH23_PIH23_004704-RA | Ras.HG9.6:like:RAB32/RAB38 | NA |
Hhon_g06701.t1 | Ras.HG9.6:like:RAB32/RAB38 | NA |
Tadh_wf_g7292.t1 | Ras.HG9.6:like:RAB32/RAB38 | NA |
TrH2_TrispH2_009168-RA | Ras.HG9.6:like:RAB32/RAB38 | NA |
TrH2_TrispH2_009166-RA | Ras.HG9.14:like:RAB32/RAB38 | NA |
Tadh_TriadT58409 | Ras.HG9.15:like:RAB32/RAB38 | NA |
Tadh_TriadT58410 | Ras.HG9.8:like:RAB32/RAB38 | NA |
TrH2_TrispH2_009167-RA | Ras.HG9.8:like:RAB32/RAB38 | NA |
Dgig_XM_028550205.1 | Ras.HG9.16:like:RAB32/RAB38 | NA |
Lampan_Lpan_C402363_01 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Horcal_Hcv1.av93.c6.g486.i1 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Horcal_Hcv1.av93.c6.g486.i2 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Coemet_Cmet_scaffold3167_01 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Vmul_Simscaffold14481_01 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Mlei_ML3061.g6.i1 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Beraby_Baby_sb322520 | Ras.HG9.7:like:RAB32/RAB38 | NA |
Drer_ENSDART00000060336 | Ras.HG9.17:like:RAB32/RAB38 | NA |
Hsap_ENST00000367495 | Ras.HG9.9:RAB32/RAB38 | RAB32 |
Mmus_ENSMUST00000019974 | Ras.HG9.9:RAB32/RAB38 | RAB32 |
Drer_ENSDART00000152970 | Ras.HG9.9:RAB32/RAB38 | NA |
Drer_ENSDART00000132831 | Ras.HG9.9:RAB32/RAB38 | NA |
Drer_ENSDART00000101408 | Ras.HG9.9:RAB32/RAB38 | NA |
Drer_ENSDART00000140186 | Ras.HG9.9:RAB32/RAB38 | RAB38 |
Mmus_ENSMUST00000107256 | Ras.HG9.9:RAB32/RAB38 | RAB38 |
Hsap_ENST00000243662 | Ras.HG9.9:RAB32/RAB38 | RAB38 |
Cint_ENSCINT00000013519 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Adig_XM_015908829.1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Spis_XP_022788100_1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Exapal_XM_021045065.2 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Nvec_NVE4434 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Bralan_BLAG16000533_1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Skow_XM_002733749.2 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Spur_XM_030973980.1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Aque_Aqu2.1.24987_001 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Tetwil_g3723.t1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Tadh_TriadT27795 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
TrH2_TrispH2_009169-RA | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
HoiH23_PIH23_004706-RA | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Hhon_g05679.t1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Cgig_XM_034459276.1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA64412 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Dmel_FBtr0089931 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Tcas_TC014165_001 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Gracom_Gcom_scaffold18210_01 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Dgig_XM_028550201.1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Ocar_g1837 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Plajan_Pjan_scaffold6372_01 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Lampan_Lpan_12689_01 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Horcal_Hcv1.av93.c11.g656.i1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Vmul_Simscaffold8396_01 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Mlei_ML0332.g43.i1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Beraby_Baby_sb334127 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Nemnom_g11503.t1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Morvir_scaffold36.g96.t1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Nemnom_g11501.t1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Hvul_g22057_1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
Hvul_g32603_1 | Ras.HG9.9:RAB32/RAB38 | RAB32/RAB38 |
0.416667