Spur_XM_030975468.1 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Bralan_BLAG05000481_1 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Tcas_TC014251_001 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Dmel_FBtr0071675 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Cgig_XM_011424007.3 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA99982 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Cint_ENSCINT00000032280 | Ion_trans.HG13.0:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Scil_scict015220.1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Plajan_Pjan_scaffold2798_01 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Plajan_Pjan_scaffold2799_01 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Spur_XM_030998944.1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Skow_XM_006824545.1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Dgig_XM_028553262.1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Spis_XP_022794596_1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Nvec_v1g81021 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Exapal_XM_021061007.2 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Coemet_Cmet_C605664_01 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Vmul_SimC390722 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Mlei_ML0544.g19.i1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Horcal_Hcv1.av93.c6.g785.i1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Beraby_Baby_sb339525 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Mlei_ML3061.g7.i1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Vmul_Simscaffold16384_01 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Lampan_Lpan_C412377_01 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Horcal_Hcv1.av93.c6.g471.i1 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Horcal_Hcv1.av93.c6.g471.i2 | Ion_trans.HG13.1:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Plajan_Pjan_C539342_01 | Ion_trans.HG13.6:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Plajan_Pjan_scaffold22462_01 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Bralan_BLAG90000227_3 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Cint_ENSCINT00000024878 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Morvir_scaffold41.g81.t1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Hvul_g8592_1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Dgig_XM_028544658.1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Dgig_XM_028544660.1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Nvec_v1g20862 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Hhon_g06663.t1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
HoiH23_PIH23_007180-RA | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
TrH2_TrispH2_002895-RA | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Tadh_TriadT13081 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Skow_XM_002734089.1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Cgig_XM_034481366.1 | Ion_trans.HG13.2:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Emue_Em0023g424a.t1 | Ion_trans.HG13.4:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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Nemnom_g5791.t1 | Ion_trans.HG13.4:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
Morvir_scaffold37.g50.t1 | Ion_trans.HG13.4:like:CNGA1/CNGA2/CNGA3/CNGA4 | NA |
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