Spis10051_1 | Pkinase.HG44.0:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Cgig_XM_011455711.3 | Pkinase.HG44.0:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Scil_scict005854.1 | Pkinase.HG44.0:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Hvul_g19975_1 | Pkinase.HG44.1:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Hvul_g11450_1 | Pkinase.HG44.1:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Hvul_g31388_1 | Pkinase.HG44.2:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Hvul_g17937_1 | Pkinase.HG44.2:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Scil_scict015250.1 | Pkinase.HG44.2:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Cint_ENSCINT00000024269 | Pkinase.HG44.2:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Cint_ENSCINT00000033046 | Pkinase.HG44.2:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Cint_ENSCINT00000011085 | Pkinase.HG44.3:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Plajan_Pjan_scaffold22549_01 | Pkinase.HG44.3:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Ocar_g3278 | Pkinase.HG44.3:like:AL451136.1/LIMK1/LIMK2/TESK1/TESK2 | NA |
Aque_Aqu2.1.20503_001 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Emue_Em0015g935a.t1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Horcal_Hcv1.av93.c5.g559.i1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Beraby_Baby_sb336763 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Beraby_Baby_sb336764 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Bolinf_Binf_sb1240458 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Vmul_Simscaffold11383_01 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Coemet_Cmet_C603744_01 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Plajan_Pjan_C562264_01 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Ocar_g2918 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Gracom_Gcom_scaffold18860_01 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Scil_scict005417.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Gracom_Gcom_scaffold18670_01 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Scil_scict003773.2 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Nemnom_g5070.t1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Morvir_scaffold300.g11.t1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Dgig_XM_028559405.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Dgig_XM_028562411.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Adig_XM_015921951.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Spis_XP_022792074_1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Exapal_XM_028661939.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Nvec_v1g95939 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Tcas_TC015763_001 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Dmel_FBtr0073660 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Cgig_XM_011450584.3 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA165760 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Cint_ENSCINT00000005959 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Spur_XM_030992386.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Skow_XM_006822297.1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Bralan_BLAG01001300_1 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1/LIMK2 |
Drer_ENSDART00000093185 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK2 |
Hsap_ENST00000340552 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK2 |
Mmus_ENSMUST00000101638 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK2 |
Drer_ENSDART00000004686 | Pkinase.HG44.4:LIMK1/LIMK2 | NA |
Drer_ENSDART00000163441 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1 |
Mmus_ENSMUST00000015137 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1 |
Hsap_ENST00000418310 | Pkinase.HG44.4:LIMK1/LIMK2 | LIMK1 |
Morvir_scaffold51.g21.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Nemnom_g12583.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Horcal_Hcv1.av93.c9.g446.i1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Horcal_Hcv1.av93.c9.g446.i2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Mlei_ML0482.g4.i1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Lampan_Lpan_C419817_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Coemet_Cmet_scaffold8597_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Vmul_sbMO457685 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Vmul_Simscaffold8493_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244706 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244708 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244704 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244705 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244703 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bolinf_Binf_sb1244707 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Ocar_g2964 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Adig_XM_015912691.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Spis_XP_022797046_1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Exapal_XM_021038038.2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Nvec_v1g240002 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Dgig_XM_028548737.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
HoiH23_PlH23_005552-RA | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Hhon_g02301.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
TrH2_TrispH2_009585-RA | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Tadh_TriadT28807 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Plajan_Pjan_scaffold2648_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Gracom_Gcom_scaffold18627_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Scil_scict003325.3 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Gracom_Gcom_scaffold4_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Scil_scict020939.2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Plajan_Pjan_scaffold11945_01 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Ocar_g1762 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Tetwil_g25054.t2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Tetwil_g9258.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Aque_Aqu2.1.17966_001 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Aque_Aqu2.1.17965_001 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Aque_Aqu2.1.01105_001 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Spur_XM_011673120.2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Spur_XM_030996082.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Drer_ENSDART00000134347 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Spur_XM_030997409.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Skow_XM_002739466.2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | NA |
Tetwil_g9006.t2 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Emue_Em0011g231a.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Aque_Aqu2.1.26628_001 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Tetwil_g14750.t1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Skow_XM_006826020.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Cgig_XM_034458497.1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA133079 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Dmel_FBtr0083716 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Tcas_TC012191_001 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Bralan_BLAG13101162_1 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK1/TESK2 |
Drer_ENSDART00000133022 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | TESK1 |
Drer_ENSDART00000146680 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | TESK1 |
Hsap_ENST00000336395 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | TESK1 |
Mmus_ENSMUST00000060864 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | TESK1 |
Drer_ENSDART00000087643 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK2 |
Mmus_ENSMUST00000045542 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK2 |
Hsap_ENST00000372086 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK2 |
Hsap_ENST00000671898 | Pkinase.HG44.5:AL451136.1/TESK1/TESK2 | AL451136.1/TESK2 |
0.416667