Dmel_FBtr0077381 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Nemnom_g17822.t1 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA161525 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Cgig_XM_034472122.1 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Cint_ENSCINT00000031952 | RRM.HG22.0:ZRSR2 | ZRSR2 |
HoiH23_PIH23_006403-RA | RRM.HG22.0:ZRSR2 | ZRSR2 |
Hsap_ENST00000307771 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Mmus_ENSMUST00000049506 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Spur_XM_011669566.2 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Nvec_v1g209221 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Aque_Aqu2.1.35041_001 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Lampan_Lpan_C415919_01 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Horcal_Hcv1.av93.c9.g303.i1 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Horcal_Hcv1.av93.c9.g303.i2 | RRM.HG22.0:ZRSR2 | ZRSR2 |
Lampan_Lpan_3283_01 | RRM.HG22.1:like:U2AF1/U2AF1L4/U2AF1L5 | NA |
Gracom_Gcom_scaffold17890_01 | RRM.HG22.1:like:U2AF1/U2AF1L4/U2AF1L5 | NA |
Scil_scict006247.1 | RRM.HG22.1:like:U2AF1/U2AF1L4/U2AF1L5 | NA |
Plajan_Pjan_C548739_01 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Lampan_Lpan_15439_01 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Vmul_Simscaffold6205_01 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Mlei_ML1054.g26.i1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Horcal_Hcv1.av93.c10.g629.i1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Coemet_Cmet_scaffold18377_01 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Lampan_Lpan_C405361_01 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Horcal_Hcv1.av93.c2.g1015.i1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Beraby_Baby_sb345329 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Beraby_Baby_sb345330 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Horcal_Hcv1.av93.c3.g100.i1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Mlei_ML0549.g3.i1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Vmul_sbMO387626 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Hhon_g04150.t1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
HoiH23_PIH23_008429-RA | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Tadh_TriadT33482 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
TrH2_TrispH2_007582-RA | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Tcas_TC031227_001 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Dmel_FBtr0078133 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Aque_Aqu2.1.29920_001 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Aque_Aqu2.1.39941_001 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Emue_Em0006g1245a.t1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Tetwil_g21394.t2 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Hvul_g27802_1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Nemnom_g17474.t1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Morvir_scaffold354.g16.t1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Dgig_XM_028546124.1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Spis_XP_022790376_1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Adig_XM_015925000.1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Nvec_v1g113937 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Exapal_XM_021057870.1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Cint_ENSCINT00000008425 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Cgig_XM_011419100.3 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA164217 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Skow_XM_006813064.1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Spur_XM_030978336.1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Spur_XM_784098.5 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Bralan_BLAG04000869_1 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L4/U2AF1L5 |
Hsap_ENST00000412391 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1L4 |
Mmus_ENSMUST00000043273 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1L4 |
Mmus_ENSMUST00000167042 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1L4 |
Drer_ENSDART00000021191 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1/U2AF1L5 |
Hsap_ENST00000610664 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1L5 |
Hsap_ENST00000291552 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1 |
Mmus_ENSMUST00000014684 | RRM.HG22.2:U2AF1/U2AF1L4/U2AF1L5 | U2AF1 |
0.416667