Dgig_XM_028548112.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Gracom_Gcom_C559810_01 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Scil_scict024262.3 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Hhon_g00852.t1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
HoiH23_PlH23_007623-RA | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
TrH2_TrispH2_006597-RA | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Tadh_TriadT28406 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Plajan_Pjan_scaffold18838_01 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Plajan_Pjan_scaffold18839_01 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Mlei_ML0447.g3.i1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Vmul_SimC380560 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Coemet_Cmet_scaffold9606_01 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Aque_Aqu2.1.41948_001 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Tetwil_g7119.t2 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Emue_Em0013g124a.t1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Emue_Em0735g4a.t1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Nvec_v1g164975 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Exapal_XM_021057010.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Spis_XP_022790526_1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Adig_XM_015914434.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Nemnom_g4878.t1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Hvul_g15090_1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Spur_XM_030986265.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | NA |
Spur_XM_030986269.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Skow_NM_001164974.1 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Bralan_BLAG03002176_5 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Cint_ENSCINT00000006561 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Tcas_TC001270_003 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Dmel_FBtr0112810 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Cgig_XM_011442715.3 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA93575 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1/EBF2/EBF3/EBF4 |
Mmus_ENSMUST00000110288 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF4 |
Hsap_ENST00000609451 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF4 |
Drer_ENSDART00000193255 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF2 |
Hsap_ENST00000520164 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF2 |
Mmus_ENSMUST00000022637 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF2 |
Drer_ENSDART00000177303 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF3 |
Drer_ENSDART00000155989 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF3 |
Hsap_ENST00000440978 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF3 |
Mmus_ENSMUST00000210774 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF3 |
Hsap_ENST00000313708 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1 |
Mmus_ENSMUST00000101326 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1 |
Drer_ENSDART00000158634 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1 |
Drer_ENSDART00000179622 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1 |
Drer_ENSDART00000140184 | HLH.HG3.0:EBF1/EBF2/EBF3/EBF4 | EBF1 |
0.416667