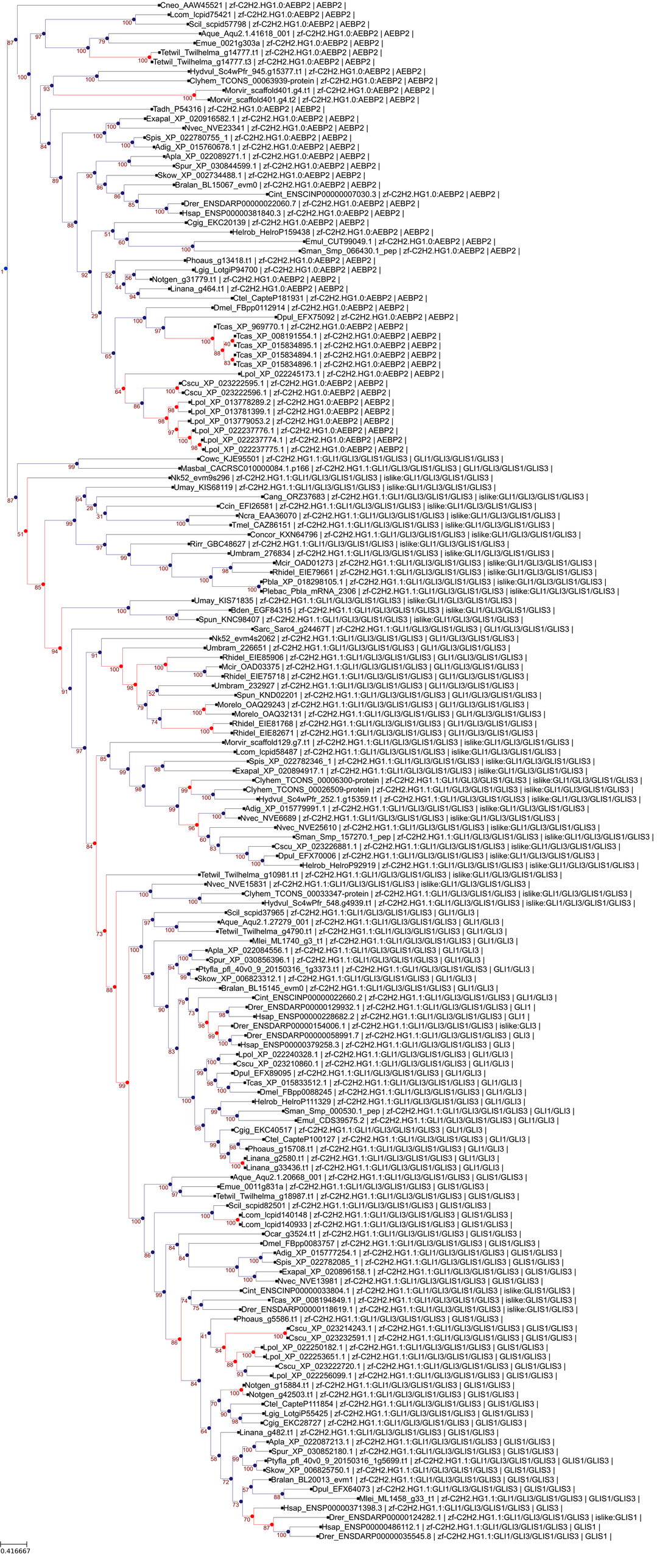
Cneo_AAW45521 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
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Scil_scpid57798 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Aque_Aqu2.1.41618_001 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Emue_0021g303a | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tetwil_Twilhelma_g14777.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tetwil_Twilhelma_g14777.t3 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Hydvul_Sc4wPfr_945.g15377.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Clyhem_TCONS_00063939-protein | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Morvir_scaffold401.g4.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Morvir_scaffold401.g4.t2 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tadh_P54316 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Exapal_XP_020916582.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Nvec_NVE23341 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Spis_XP_022780755_1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Adig_XP_015760678.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Apla_XP_022089271.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Spur_XP_030844599.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Skow_XP_002734488.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Bralan_BL15067_evm0 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Cint_ENSCINP00000007030.3 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Drer_ENSDARP00000022060.7 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Hsap_ENSP00000381840.3 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Cgig_EKC20139 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Helrob_HelroP159438 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Emul_CUT99049.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Sman_Smp_066430.1_pep | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Phoaus_g13418.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Lgig_LotgiP94700 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Notgen_g31779.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Linana_g464.t1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Ctel_CapteP181931 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Dmel_FBpp0112914 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Dpul_EFX75092 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tcas_XP_969770.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tcas_XP_008191554.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tcas_XP_015834895.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tcas_XP_015834894.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Tcas_XP_015834896.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Lpol_XP_022245173.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Cscu_XP_023222595.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Cscu_XP_023222596.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Lpol_XP_013778289.2 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
Lpol_XP_013781399.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
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Lpol_XP_022237776.1 | zf-C2H2.HG1.0:AEBP2 | AEBP2 |
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Masbal_CACRSC010000084.1.p166 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3/GLIS1/GLIS3 |
Nk52_evm9s296 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Umay_KIS68119 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Cang_ORZ37683 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Ccin_EFI26581 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Ncra_EAA36070 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Tmel_CAZ86151 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Concor_KXN64796 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Rirr_GBC48627 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Umbram_276834 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Mcir_OAD01273 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Rhidel_EIE79661 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Pbla_XP_018298105.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Plebac_Pbla_mRNA_2306 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Umay_KIS71835 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Bden_EGF84315 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Spun_KNC98407 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Sarc_Sarc4_g24467T | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3/GLIS1/GLIS3 |
Nk52_evm4s2062 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3/GLIS1/GLIS3 |
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Spun_KND02201 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3/GLIS1/GLIS3 |
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Rhidel_EIE81768 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3/GLIS1/GLIS3 |
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Morvir_scaffold129.g7.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Lcom_lcpid58487 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Spis_XP_022782346_1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Exapal_XP_020894917.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Clyhem_TCONS_00006300-protein | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Clyhem_TCONS_00026509-protein | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Hydvul_Sc4wPfr_252.1.g15359.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Adig_XP_015779991.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Nvec_NVE6689 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
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Dpul_EFX70006 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
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Tetwil_Twilhelma_g10981.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Nvec_NVE15831 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Clyhem_TCONS_00033347-protein | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Hydvul_Sc4wPfr_548.g4939.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLI1/GLI3/GLIS1/GLIS3 |
Scil_scpid37965 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Aque_Aqu2.1.27279_001 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
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Spur_XP_030856396.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Ptyfla_pfl_40v0_9_20150316_1g3373.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
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Bralan_BL15145_evm0 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
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Hsap_ENSP00000228682.2 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1 |
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Hsap_ENSP00000379258.3 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI3 |
Lpol_XP_022240328.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Cscu_XP_023210860.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Dpul_EFX89095 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
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Sman_Smp_000530.1_pep | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Emul_CDS39575.2 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
Cgig_EKC40517 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLI3 |
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Tetwil_Twilhelma_g18987.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLIS3 |
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Spis_XP_022782085_1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLIS3 |
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Nvec_NVE13981 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLI1/GLIS3 |
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Tcas_XP_008194849.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLIS1/GLIS3 |
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Phoaus_g5586.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
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Lpol_XP_022250182.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Lpol_XP_022253651.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Cscu_XP_023222720.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Lpol_XP_022256099.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Notgen_g15884.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Notgen_g42503.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Ctel_CapteP111854 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Lgig_LotgiP55425 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Cgig_EKC28727 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Linana_g482.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Apla_XP_022087213.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Spur_XP_030852180.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Ptyfla_pfl_40v0_9_20150316_1g5699.t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Skow_XP_006825750.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Bralan_BL20013_evm1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Dpul_EFX64073 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
Mlei_ML1458_g33_t1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1/GLIS3 |
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Drer_ENSDARP00000124282.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | islike:GLIS1 |
Hsap_ENSP00000486112.1 | zf-C2H2.HG1.1:GLI1/GLI3/GLIS1/GLIS3 | GLIS1 |
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