Chrtob_KOO53356.1 | Acetyltransf_1.HG21.0:NA | NA |
Amobb2_AMBB2_12850.mRNA.1 | Acetyltransf_1.HG21.0:NA | NA |
Symmic_Smic32268 | Acetyltransf_1.HG21.0:NA | NA |
Bremin_symbB.v1.2.035503.t1 | Acetyltransf_1.HG21.0:NA | NA |
Auraur_scaffold1747.g7.t1 | Acetyltransf_1.HG21.0:NA | NA |
Cvel_2874.t1-p1 | Acetyltransf_1.HG21.0:NA | NA |
Cmer_CML264CT | Acetyltransf_1.HG21.0:NA | NA |
Malcal_MALCA_01344.mRNA.1 | Acetyltransf_1.HG21.0:NA | NA |
Nutlon_comp27604_c0_seq1.p1 | Acetyltransf_1.HG21.0:NA | NA |
Manpla_Gene.3169 | Acetyltransf_1.HG21.0:NA | NA |
Spur_XP_030838534.1 | Acetyltransf_1.HG21.0:NA | NA |
Amobb2_AMBB2_01859.mRNA.2 | Acetyltransf_1.HG21.0:NA | NA |
Jakbah_JAKBA_06360.mRNA.1 | Acetyltransf_1.HG21.0:NA | NA |
Spimul_7639_TR4022_c0_g1 | Acetyltransf_1.HG21.0:NA | NA |
Spimul_7641_TR4022_c0_g2 | Acetyltransf_1.HG21.0:NA | NA |
Gonavo_comp74879_c0_seq1_6_ORF5_432 | Acetyltransf_1.HG21.0:NA | NA |
Gthe_EKX47137 | Acetyltransf_1.HG21.0:NA | NA |
Ttra_KNC52965 | Acetyltransf_1.HG21.0:NA | NA |
Symmic_Smic7772 | Acetyltransf_1.HG21.0:NA | NA |
Bremin_symbB.v1.2.026895.t1 | Acetyltransf_1.HG21.0:NA | NA |
Bremin_symbB.v1.2.026895.t2 | Acetyltransf_1.HG21.0:NA | NA |
Azfi_s0010.g012296 | Acetyltransf_1.HG21.0:NA | NA |
Sacu_v1.1_s0019.g007840 | Acetyltransf_1.HG21.0:NA | NA |
Sacu_v1.1_s0093.g019046 | Acetyltransf_1.HG21.0:NA | NA |
Azfi_s0038.g026357 | Acetyltransf_1.HG21.0:NA | NA |
Azfi_s0002.g001623 | Acetyltransf_1.HG21.0:NA | NA |
Sacu_v1.1_s0014.g006281 | Acetyltransf_1.HG21.0:NA | NA |
Sacu_v1.1_s0267.g026963 | Acetyltransf_1.HG21.0:NA | NA |
Sacu_v1.1_s0118.g021205 | Acetyltransf_1.HG21.0:NA | NA |
Azfi_s0147.g053069 | Acetyltransf_1.HG21.0:NA | NA |
Smoe_EFJ22800 | Acetyltransf_1.HG21.0:NA | NA |
Smoe_EFJ30840 | Acetyltransf_1.HG21.0:NA | NA |
Marpol_PTQ33977 | Acetyltransf_1.HG21.0:NA | NA |
Ppat_Pp3c22_13550V3.1 | Acetyltransf_1.HG21.0:NA | NA |
Ppat_Pp3c19_14710V3.1 | Acetyltransf_1.HG21.0:NA | NA |
Sphfal_Sphfalx02G123700.1.p | Acetyltransf_1.HG21.0:NA | NA |
Sphfal_Sphfalx01G061300.1.p | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA61054.1 | Acetyltransf_1.HG21.0:NA | NA |
Atri_ERN05326 | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA59504.1 | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA32396.1 | Acetyltransf_1.HG21.0:NA | NA |
Atha_AT2G30090.1 | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA45856.1 | Acetyltransf_1.HG21.0:NA | NA |
Mgut_XP_012856193.1 | Acetyltransf_1.HG21.0:NA | NA |
Sbic_KXG29625 | Acetyltransf_1.HG21.0:NA | NA |
Bdis_KQJ93656 | Acetyltransf_1.HG21.0:NA | NA |
Sbic_EER90127 | Acetyltransf_1.HG21.0:NA | NA |
Bdis_KQK16841 | Acetyltransf_1.HG21.0:NA | NA |
Sbic_EER97861 | Acetyltransf_1.HG21.0:NA | NA |
Sbic_EER93401 | Acetyltransf_1.HG21.0:NA | NA |
Bdis_PNT74088 | Acetyltransf_1.HG21.0:NA | NA |
Bdis_KQK13054 | Acetyltransf_1.HG21.0:NA | NA |
Sbic_EER92674 | Acetyltransf_1.HG21.0:NA | NA |
Bdis_PNT77946 | Acetyltransf_1.HG21.0:NA | NA |
Atri_ERM93938 | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA44135.1 | Acetyltransf_1.HG21.0:NA | NA |
Acoe_PIA33185.1 | Acetyltransf_1.HG21.0:NA | NA |
Mgut_XP_012846660.1 | Acetyltransf_1.HG21.0:NA | NA |
Mgut_XP_012856464.1 | Acetyltransf_1.HG21.0:NA | NA |
Atha_AT5G67430.1 | Acetyltransf_1.HG21.0:NA | NA |
Atha_AT2G23060.1 | Acetyltransf_1.HG21.0:NA | NA |
Atha_AT4G37580.1 | Acetyltransf_1.HG21.0:NA | NA |
0.416667