Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA226021 | SSF.HG3.4:like:SLC5A7 | NA |
Cgig_XM_034444952.1 | SSF.HG3.0:like:SLC5A7 | NA |
Cgig_XM_034445498.1 | SSF.HG3.0:like:SLC5A7 | NA |
Cgig_XM_020074163.2 | SSF.HG3.0:like:SLC5A7 | NA |
Cgig_XM_020074166.2 | SSF.HG3.0:like:SLC5A7 | NA |
Bralan_BLAG06000305_1 | SSF.HG3.1:like:SLC5A7 | NA |
Bralan_BLAG19000287_1 | SSF.HG3.1:like:SLC5A7 | NA |
Spur_XM_030973201.1 | SSF.HG3.1:like:SLC5A7 | NA |
Cgig_XM_011442834.3 | SSF.HG3.1:like:SLC5A7 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA212834 | SSF.HG3.1:like:SLC5A7 | NA |
Skow_XM_006823263.1 | SSF.HG3.1:like:SLC5A7 | NA |
Skow_XM_006823264.1 | SSF.HG3.1:like:SLC5A7 | NA |
Cgig_XM_011443978.3 | SSF.HG3.5:like:SLC5A7 | NA |
Bralan_BLAG15000595_1 | SSF.HG3.2:like:SLC5A7 | NA |
Cgig_XM_020063410.2 | SSF.HG3.2:like:SLC5A7 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA157516 | SSF.HG3.2:like:SLC5A7 | NA |
Lampan_Lpan_C416583_01 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Vmul_sbMO426248 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Coemet_Cmet_C605004_01 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Horcal_Hcv1.av93.c9.g773.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1234570 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1234569 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Mlei_ML0438.g13.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238970 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Mlei_ML0412.g3.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Mlei_ML0438.g12.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb349115 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb349116 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb343989 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb343990 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Vmul_Simscaffold7356_01 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Coemet_Cmet_scaffold12643_01 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Lampan_Lpan_10749_01 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238461 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238458 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238460 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238457 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1238459 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Horcal_Hcv1.av93.c4.g250.i3 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Horcal_Hcv1.av93.c4.g250.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Horcal_Hcv1.av93.c4.g250.i2 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Horcal_Hcv1.av93.c4.g250.i4 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Mlei_ML0897.g14.i1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bolinf_Binf_sb1236402 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb347297 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb347299 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb347296 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Beraby_Baby_sb347298 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Skow_XM_002735497.2 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000236_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000239_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000232_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000226_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000231_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Bralan_BLAG14000227_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cint_ENSCINT00000008788 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cint_ENSCINT00000008776 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Drer_ENSDART00000147201 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Drer_ENSDART00000165960 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Hsap_ENST00000264047 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Mmus_ENSMUST00000095712 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Dmel_FBtr0083668 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Tcas_TC000935_001 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cgig_XM_034455816.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cgig_XM_011415184.3 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cgig_XM_011456113.3 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Cgig_XM_034457634.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
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Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA183218 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA131610 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA214938 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA120487 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA113629 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA217363 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_790185.5 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_011675698.2 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_030995007.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_030999145.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_789477.4 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_030982576.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spur_XM_030982597.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Dgig_XM_028551979.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Dgig_XM_028551982.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Exapal_XM_021053799.2 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Exapal_XM_028656748.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nvec_v1g95565 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nvec_NVE22942 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nvec_v1g89183 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spis_XP_022780800_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spis_XP_022780084_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spis_XP_022780106_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Spis_XP_022780993_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Adig_XM_015925466.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Adig_XM_015925458.1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g6283.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Morvir_scaffold64.g44.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Hvul_g32592_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Morvir_scaffold35.g28.t2 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Morvir_scaffold44.g60.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Morvir_scaffold552.g1.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Hvul_g22297_1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g8966.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g13544.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g19547.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g13543.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g19546.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
Nemnom_g13542.t1 | SSF.HG3.3:SLC5A7 | SLC5A7 |
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