-Horcal_Hcv1.av93.c4.g863.i1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A14/SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1/SLC2A1/SLC2A11/SLC2A14/SLC2A1/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A1 -MIei_ML1322.g3.i1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A14/SLC2A1/SLC2A14/SLC2A14/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A14/SLC2A11/SLC2A14/SLC2A14/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A9/SLC2A11/SLC2A14/SLC2A11/SLC2A14/SLC2A14/SLC2A1/SLC2 Beraby_Baby_sb351465 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1/SLC2A1/SLC2A11/SLC2A14/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A1 1001Beraby_Baby_sb351466 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A1/SLC2A14/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A1/SLC2A3/SLC2A4/SLC2A7/SLC2A9/SLC2A9/SLC2A9/SLC2A1/SL -Vmul_sbMO441809 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1 -Coemet_Cmet_scaffold19043_01 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 |
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AP000350.4/SLC2A1/SLC2A1/SLC2A14/SLC2A14/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9 | Beraby_Baby_sb348744 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A3/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A11/SLC2A14/SLC2A14/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A14/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A11/SLC2A1/S 56 Beraby_Baby_sb348745 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A4/SLC2A5/SLC2A5/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A9/SLC2A1 -MIei ML3593.g13.i2 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1/SLC Bolinf_Binf_sb1266520 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A1/SLC2A11/SLC2A14/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A11/SLC2A1/SLC2A1/SLC2A11/SLC2A14/SLC2A1/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A9/SLC2A1/SLC2A ⁵¹Bolinf Binf sb1266522 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 |
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1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Morvir_scaffold59.g27.t1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | 100 -Dgig_XM_028538010.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Adig_XM_015923942.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | Spis XP 022780136 1 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Nvec v1g191325 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | •Nvec_v1g100165 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Exapal_XM_028663020.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | →Exapal_XM_021058016.2 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000003472 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | --Cgig_XM_011437962.3 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | -Cgig_XM_020068273.2 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 •Tcas_TC033126_001 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | •Tcas_TC013486_001 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | -Tcas TC014313 001 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | -Skow_XM_006819844.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | Spur_XM_030987096.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | 1001 Spur_XM_030987133.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | _•Dmel_FBtr0333501 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | 100 -Tcas TC032400 001 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | -Cgig_XM_034464756.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 | -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA161693 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1/SLC2A14/SLC2A2/SLC2A3/SLC2A4 -Mmus_ENSMUST00000018710 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A4 | 100¹-Hsap_ENST00000317370 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A4 | -Drer ENSDART00000078639 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A2 | -Mmus_ENSMUST00000029240 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A2 | Hsap_ENST00000314251 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A2 | -Drer_ENSDART00000019804 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Drer_ENSDART00000025414 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1 | Tool-Drer_ENSDART00000137458 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1 | Hsap_ENST00000426263 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1 | 100 Mmus_ENSMUST00000030398 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A1 | -Drer_ENSDART00000175075 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A14/SLC2A3 | -Drer_ENSDART00000174467 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A14/SLC2A3 | -Mmus_ENSMUST00000032476 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A14/SLC2A3 | Hsap ENST00000539924 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A14/SLC2A3 | 1001 Hsap_ENST00000075120 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A14/SLC2A3 | -Cgig_XM_011418018.3 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA170630 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA219782 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A5/SLC2A7/SLC2A9 | -Dgig_XM_028540655.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA129046 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A5/SLC2A7/SLC2A9 -Tcas_TC012857_001 |
MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Dmel_FBtr0088777 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Dmel_FBtr0085966 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | 100 -Dmel_FBtr0088778 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Dmel_FBtr0088779 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Spur_XM_790455.5 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 -Skow_XM_006819843.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 -Skow_XM_002736074.2 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | 100 -Skow_XM_006818168.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Spur_XM_011662934.2 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Spur_XM_030981616.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | Spur_XM_030987134.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | 1001-Spur_XM_030987098.1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 Bralan_BLAG15000493_1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 | -Bralan_BLAG11000682_1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A7/SLC2A9 100 -Bralan_BLAG04001664_1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A5/SLC2A7/SLC2A9 | 100¹—•Bralan_BLAG09000823_1 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A5/SLC2A5/SLC2A7/SLC2A9 | --Hsap_ENST00000400906 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A7 | 100 → Mmus_ENSMUST00000059893 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A7 | -Drer_ENSDART00000147087 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A5 | ___Mmus_ENSMUST00000030826 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A5 | ¹⁰⁰Ū-Hsap_ENST00000377424 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A5 | -Cint_ENSCINT00000014757 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11/SLC2A9 --Hsap_ENST00000264784 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A9 | 100 Mmus_ENSMUST00000067886 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A9 | Drer_ENSDART00000075286 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A9 | Too Drer ENSDART00000043658 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | SLC2A9 | -Drer_ENSDART00000144847 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | Drer_ENSDART00000103983 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | Drer_ENSDART00000092333 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Drer ENSDART00000190465 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11 | -Hsap ENST00000611880 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11 | -Hsap ENST00000433835 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | AP000350.4/SLC2A11 | -Cint ENSCINT00000008151 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000024732 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint ENSCINT00000029029 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | •Cint_ENSCINT00000006802 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000029030 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000015967 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000036736 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint ENSCINT00000032435 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint ENSCINT00000036341 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint_ENSCINT00000028297 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint ENSCINT00000027509 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | T001 Cint ENSCINT00000006481 | MFS_1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Cint ENSCINT00000028908 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | ¹⁰⁰1—Cint ENSCINT00000028907 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | Cint ENSCINT00000019981 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | 1001—•Cint ENSCINT00000028909 | MFS 1.HG4.0:AP000350.4/SLC2A1/SLC2A2/SLC2A3/SLC2A4/SLC2A5/SLC2A7/SLC2A9/SLC2A11/SLC2A14 | NA | -Plajan_Pjan_scaffold21737_01 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | Ocar g407 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | HoiH23 PIH23 010441-RA | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | ¹⁰⁰ Hhon g08440.t1 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | *Tadh TriadT64243 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | TrH2 TrispH2 008388-RA | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | •Scil_scict022989.4 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | 100 Gracom Gcom scaffold15187 01 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA ¹⁰⁰lGracom Gcom scaffold15188 01 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | -Tetwil g24923.t2 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | -Tetwil_g25435.t2 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | Tetwil_g6688.t2 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | 100 - Tetwil g11851.t2 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | Tetwil g11852.t1 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | Ague Agu2.1.01742 001 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | ¹⁰⁰¹Aque Aqu2.1.19218_001 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | Aque_Aqu2.1.07069_001 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | _-Aque_Aqu2.1.18435_001 | MFS_1.HG4.3:like:SLC2A10/SLC2A12 | NA | ⁷⁹¹Aque Aqu2.1.19215 001 | MFS 1.HG4.3:like:SLC2A10/SLC2A12 | NA | -Hhon g00151.t1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | *Tadh TriadT58068 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | ¹⁰⁰¹•TrH2 TrispH2 000540-RA | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | •Drer ENSDART00000053531 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A12 | -Mmus ENSMUST00000042261 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A12 | 100 - Hsap ENST00000275230 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A12 | 100 99 -Drer ENSDART00000167938 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10 | --Mmus_ENSMUST00000029196 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10 | Hsap ENST00000359271 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10 | •Ctel gnl WGS AMQN CAPTEDRAFT mRNA228172 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | Gracom_Gcom_C549996_01 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Morvir scaffold206.g20.t1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Nemnom g18935.t1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Hvul g13867 1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Bralan_BLAG02000333_1 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | •Cgig_XM_034457498.1 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Spur XM 030983482.1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Skow XM 006812512.1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | Plajan_Pjan_scaffold11208_01 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | Plajan_Pjan_scaffold11210_01 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | 531—•Plajan_Pjan_scaffold11211_01 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | •Dgig XM 028550578.1 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Nvec NVE3004 | MFS 1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | -Exapal_XM_028658294.1 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 →Spis_XP_022796206_1 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | --Adig_XM_015919022.1 | MFS_1.HG4.4:SLC2A10/SLC2A12 | SLC2A10/SLC2A12 | Lampan Lpan C416373 01 | MFS 1.HG4.5:like:SLC2A13 | NA | Coemet Cmet scaffold19402 01 | MFS 1.HG4.5:like:SLC2A13 | NA | -Mlei ML1022.g60.i1 | MFS 1.HG4.5:like:SLC2A13 | NA | -Beraby_Baby_sb324500 | MFS_1.HG4.5:like:SLC2A13 | NA | Horcal_Hcv1.av93.c1.g360.i2 |
MFS_1.HG4.5:like:SLC2A13 | NA | ¹⁰⁰ Horcal Hcv1.av93.c1.g360.i1 | MFS_1.HG4.5:like:SLC2A13 | NA | -Vmul SimC377798 | MFS 1.HG4.10:like:SLC2A13 | NA | •Coemet Cmet scaffold5805 01 | MFS 1.HG4.9:like:SLC2A13 | NA | →Vmul_SimC401016 | MFS_1.HG4.9:like:SLC2A13 | NA | -Horcal Hcv1.av93.c8.g441.i1 | MFS 1.HG4.8:like:SLC2A13 | NA | -Lampan Lpan C384495 01 | MFS 1.HG4.8:like:SLC2A13 | NA | 100 -Coemet Cmet C603766 01 | MFS 1.HG4.7:like:SLC2A13 | NA | _fVmul_SimC403394 | MFS_1.HG4.7:like:SLC2A13 | NA | 100¹Vmul sbMO456567 | MFS_1.HG4.7:like:SLC2A13 | NA | -Lampan_Lpan_C416607_01 | MFS_1.HG4.7:like:SLC2A13 | NA | -Bolinf Binf sb1169233 | MFS 1.HG4.7:like:SLC2A13 | NA | -Mlei_ML0830.g25.i1 | MFS_1.HG4.7:like:SLC2A13 | NA | -Horcal_Hcv1.av93.c5.g790.i1 | MFS_1.HG4.7:like:SLC2A13 | NA | •Coemet_Cmet_scaffold7061_01 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Horcal_Hcv1.av93.c3.g545.i1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Bolinf Binf sb1299895 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | -Beraby_Baby_sb350948 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | Aque_Aqu2.1.41105_001 | MFS_1.HG4.6:SLC2A13 | NA | 100 - Aque Aqu2.1.41106_001 | MFS_1.HG4.6:SLC2A13 | NA | -Gracom_Gcom_C555504_01 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | →Scil_scict014873.3 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | •Aque_Aqu2.1.32928_001 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | Emue_Em0006g1034a.t1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Emue Em0006g1040a.t1 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | -Hvul_g17124_1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | •Nemnom_g18719.t1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | 100 -Morvir_scaffold137.g7.t1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | •Morvir scaffold322.g27.t1 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | •Dgig XM 028542580.1 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | •Dgig_XM_028542582.1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | •Nvec_v1g228429 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | 100 -Exapal_XM_021053766.2 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Spis XP 022805589 1 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | -Adig XM 015912455.1 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | --Spis_XP_022805578_1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | Spis_XP_022805579_1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | ¹⁰⁰ → Spis_XP_022805078_1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | Cgig_XM_034449836.1 | MFS_1.HG4.6:SLC2A13 | NA | -Cgig XM 034449700.1 | MFS 1.HG4.6:SLC2A13 | NA | -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA175747 | MFS_1.HG4.6:SLC2A13 | NA | -Skow_XM_006820920.1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Cgig_XM_011440134.3 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | -Ctel gnl WGS AMQN CAPTEDRAFT mRNA168828 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | -Cint ENSCINT00000002435 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | •Bralan_BLAG05101597_1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 Spur_XM_030985399.1 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | ¹⁰⁰¹-Spur_XM_793355.4 | MFS_1.HG4.6:SLC2A13 | SLC2A13 | Drer ENSDART00000154979 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | -Hsap ENST00000280871 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | ¹⁰⁰→Mmus ENSMUST00000109283 | MFS 1.HG4.6:SLC2A13 | SLC2A13 | •Ctel gnl WGS AMQN CAPTEDRAFT mRNA63779 | MFS 1.HG4.1:like:SLC2A6/SLC2A8 | NA | -Beraby_Baby_sb335635 | MFS_1.HG4.1:like:SLC2A6/SLC2A8 | NA | -Bolinf Binf sb1213686 | MFS 1.HG4.1:like:SLC2A6/SLC2A8 | NA | •Horcal Hcv1.av93.c5.g58.i1 | MFS 1.HG4.1:like:SLC2A6/SLC2A8 | NA | 100 Lampan_Lpan_4654_01 | MFS_1.HG4.1:like:SLC2A6/SLC2A8 | NA | -Vmul_Simscaffold11049_01 | MFS_1.HG4.1:like:SLC2A6/SLC2A8 | NA | -Coemet_Cmet_scaffold8678_01 | MFS_1.HG4.1:like:SLC2A6/SLC2A8 | NA | •Tcas TC033857 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC013484 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel FBtr0082011 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas TC015433 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC015095_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC033468 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 -Tcas TC015809 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC014877_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel FBtr0071013 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Dmel FBtr0087310 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas TC030899 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC012760_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel FBtr0072239 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel FBtr0088073 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 1001-Dmel FBtr0088075 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC000121_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC030889_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC013653_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 •Tcas_TC012566_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 -Tcas TC012830 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Dmel FBtr0113197 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas TC003228 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 34 -Tcas_TC034087_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | Dmel FBtr0075942 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Dmel FBtr0086903 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Dmel FBtr0089674 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel_FBtr0083340 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Dmel FBtr0076315 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 62 •Tcas TC034243 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas TC012291 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC012289_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC012290_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC000733_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 -Tcas TC031339 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC008119_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC033626 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tcas_TC007235_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC003287 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC003288 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC003289 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 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MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | --Tcas_TC012232_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 1001-Tcas_TC012233_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC012234 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC012235_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 1001-Tcas TC012230 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | rTcas_TC001413_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | Tcas_TC034222_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 911-Tcas TC001414 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC012228 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas TC012229 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tcas_TC012231_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tetwil g23102.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tetwil g23103.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tetwil g23106.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tetwil_g23104.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰¹_-Tetwil_g23105.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | Emue Em0003g1642a.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰ Emue Em0003g1673a.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Emue Em0005g1623a.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | Emue_Em0005g1617a.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Emue_Em0005g1624a.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Emue Em0018g415a.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Aque Aqu2.1.36131 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Tetwil g25976.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 1001-Tetwil g25979.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Aque_Aqu2.1.36154_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Aque Aqu2.1.37781 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Aque Aqu2.1.20811 001 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | --Aque_Aqu2.1.20812_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ₋-Aque_Aqu2.1.20808_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰L•Aque_Aqu2.1.39896_001 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | •Tetwil g3878.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | retwil g13157.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | --Tetwil_g13158.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | _*Tetwil_g18468.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 991-Tetwil_g15755.t2 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 -Tetwil g18470.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 Tetwil g18467.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | TrH2 TrispH2 007925-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰Tadh TriadT31909 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -HoiH23 PIH23 005098-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 - Hhon g01114.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | *Tadh TriadT32030 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰TrH2 TrispH2 007926-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 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SLC2A6/SLC2A8 | 100 + Hhon g01115.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | Tadh TriadT58298 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 ¹⁰⁰TrH2 TrispH2 002820-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -HoiH23 PIH23 003802-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 99¹--Hhon_g07604.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | *Tadh TriadT38377 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰ TrH2 TrispH2 007932-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -Hhon g01122.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | ¹⁰⁰1—HoiH23 PIH23 008932-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | TrH2_TrispH2_007931-RA | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | -HoiH23 PIH23 008933-RA | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 + Hhon g01121.t1 | MFS 1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | rhoiH23_PIH23_008936-RA | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | 100 Hhon g01119.t1 | MFS_1.HG4.2:SLC2A6/SLC2A8 | SLC2A6/SLC2A8 | TrH2_TrispH2_007928-RA | 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