Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA191774 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Bralan_BLAG12000697_1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Spur_XM_030972106.1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Skow_XM_006825598.1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Hsap_ENST00000308436 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Hsap_ENST00000496634 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Dgig_XM_028549603.1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Morvir_scaffold20.g79.t1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Spis_XP_022810505_1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Spis_XP_022801046_1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Exapal_XM_021057353.2 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Nvec_v1g238029 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Gracom_Gcom_scaffold18805_01 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Emue_Em0015g1151a.t1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Emue_Em0015g1135a.t1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Plajan_Pjan_scaffold23395_01 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Ocar_g1413 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Lampan_Lpan_9847_01 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
Mlei_ML0888.g9.i1 | RRM.HG40.0:AP002990.1/TUT1 | AP002990.1/TUT1 |
65
51
48
37
100
1
85
40
100
61
89
65
68
100
44
88
58
100
0.416667