Ocar_g8825 | GPCRsecr.HG7.0:like:GPR143 | NA |
Gracom_Gcom_C556098_01 | GPCRsecr.HG7.0:like:GPR143 | NA |
Gracom_Gcom_scaffold18474_01 | GPCRsecr.HG7.0:like:GPR143 | NA |
Scil_scict008876.2 | GPCRsecr.HG7.0:like:GPR143 | NA |
Bralan_BLAG04000307_1 | GPCRsecr.HG7.0:like:GPR143 | NA |
Dgig_XM_028548069.1 | GPCRsecr.HG7.0:like:GPR143 | NA |
Dgig_XM_028548067.1 | GPCRsecr.HG7.0:like:GPR143 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA206680 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Tcas_TC012209_001 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Bralan_BLAG01001311_4 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Hsap_ENST00000467482 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Mmus_ENSMUST00000026383 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Cgig_XM_011425403.3 | GPCRsecr.HG7.1:GPR143 | GPR143 |
Hhon_g06973.t1 | GPCRsecr.HG7.1:GPR143 | GPR143 |
HoiH23_PlH23_000385-RA | GPCRsecr.HG7.1:GPR143 | GPR143 |
Tadh_TriadT59750 | GPCRsecr.HG7.1:GPR143 | GPR143 |
TrH2_TrispH2_000330-RA | GPCRsecr.HG7.1:GPR143 | GPR143 |
95
40
100
28
75
100
1
39
36
69
100
95
48
100
77
98
0.416667