Exapal XM 021042392.2 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Nvec v1g198777 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA Nvec v1g200012 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | 47 Nvec v1g225210 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | TrH2_TrispH2_004645-RA | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Tadh TriadT57663 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA HoiH23 PIH23 009268-RA | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | ■Hhon g01812.t1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA 100 Hhon g02170.t1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA -■HoiH23_PIH23_005798-RA | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | -■Hhon g09503.t1 | 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TriadT57425 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | -HoiH23 PIH23 007897-RA | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Hhon g01955.t1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | ¹⁰⁰ Hhon g10338.t1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Gracom_Gcom_scaffold17564_01 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Scil scict003827.1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Gracom Gcom C537522 01 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Gracom Gcom C560256 01 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Gracom Gcom scaffold16083 01 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Scil_scict020810.1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | Gracom 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GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | -Aque Aqu2.1.06178 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | -Aque_Aqu2.1.25207_001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA 98 100 Aque Aqu2.1.25208 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA -Aque Aqu2.1.11053 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA 400 Aque Aqu2.1.13316 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA Aque Aqu2.1.25214 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | 400 Aque Aqu2.1.44682 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA -Aque Aqu2.1.14054 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | ¹⁰⁰ Aque Aqu2.1.14052 001 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA 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GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | NA | -■Hsap_ENST00000333493 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG3 | -•Mmus ENSMUST00000051259 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG3 | •Drer ENSDART00000188170 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | NA | •Drer ENSDART00000114592 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG1/ADGRG5 | -Hsap_ENST00000565976 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG1 | 100 Mmus ENSMUST00000093271 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG1 | -- Hsap ENST00000340339 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG5 | *Mmus_ENSMUST00000166802 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG5 | --Hsap ENST00000370652 | GPCRsecr.HG2.2:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6 | ADGRG4 | -•Mmus_ENSMUST00000153784 | 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GPCRsecr.HG2.3:like:ADGRG7 | NA | Spur XM 030990112.1 | GPCRsecr.HG2.3:like:ADGRG7 | NA 100 Spur XM 030990793.1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 Ctel gnl WGS AMQN CAPTEDRAFT mRNA212970 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA227623 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Ctel gnl WGS AMQN CAPTEDRAFT mRNA70548 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ┌─■Ctel gnl WGS AMQN CAPTEDRAFT mRNA137798 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Ctel gnl WGS AMQN CAPTEDRAFT mRNA95355 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Plajan Pjan scaffold19632 01 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Plajan Pjan scaffold5466_01 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Ocar g5868 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Ocar g4585 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Ocar g5546 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.22167 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 -Aque_Aqu2.1.22166_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil_g11433.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g6918.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ■Tetwil g19081.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 Aque Aqu2.1.28798_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.22161 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.37163 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.37185 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Aque Aqu2.1.36489 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Aque_Aqu2.1.40350_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.11575 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.27151 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA -Aque Aqu2.1.28045 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Aque Aqu2.1.36490 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque_Aqu2.1.43876_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Aque_Aqu2.1.28038_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 Aque Aqu2.1.28039 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0021g592a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g21944.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Aque Aqu2.1.26221 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 - Aque_Aqu2.1.26223_001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g19153.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Aque Aqu2.1.42300 001 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g23149.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil_g27520.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g25302.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g9930.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g19770.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g10250.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 100 ■Tetwil g22041.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil_g18728.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | *Tetwil g18729.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g18730.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | r-Tetwil g7524.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -Tetwil g7527.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g7525.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil g6587.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 99 Tetwil g19891.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | rulu = Tetwil g16941.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Tetwil_g7526.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | -■Tetwil_g16940.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ■Tetwil g16939.t2 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0006g43a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA Emue Em0006g44a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue_Em0006g45a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA Emue Em0006g42a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA ■Emue Em0016g893a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0016g906a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0557g2a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ■Emue Em0001g688a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0018g260a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 Emue Em0018g259a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0009g911a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ■Emue Em0003g1493a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0009g870a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0009g931a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Emue Em0024g169a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | ■Emue Em0009g903a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | 100 Emue Em0009g893a.t1 | GPCRsecr.HG2.3:like:ADGRG7 | NA | Dgig_XM_028539298.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | •Dgig XM 028537690.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028556151.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | 100 100 Dgig XM 028543541.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028563059.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig_XM_028551470.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028551484.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028560906.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | 100 Dgig XM 028551516.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028543693.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig_XM_028556156.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | _ Dgig XM 028543564.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | 100 - Dgig XM 028556137.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028562295.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | 100 - Dgig XM 028556131.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | •Dgig_XM_028563398.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig_XM_028562293.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | 95 Dgig XM 028562762.1 | GPCRsecr.HG2.4:like:ADGRG7 | NA | Dgig XM 028552237.1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Nemnom g5067.t1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Morvir_scaffold267.g15.t2 | GPCRsecr.HG2.6:like:ADGRG7 | NA | •Adig XM 015906351.1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | 100 Spis XP_022801811_1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Nvec v1g238736 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Exapal XM 021048071.2 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Exapal_XM_028661585.1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Exapal_XM_021053364.2 | GPCRsecr.HG2.6:like:ADGRG7 | NA | Exapal XM 028661600.1 | GPCRsecr.HG2.6:like:ADGRG7 | NA | 72 Morvir scaffold311.g2.t1 | GPCRsecr.HG2.10:like:ADGRG7 | NA | Nemnom g15058.t1 | GPCRsecr.HG2.9:like:ADGRG7 | NA | Morvir_scaffold21.g77.t2 | GPCRsecr.HG2.9:like:ADGRG7 | NA | •Morvir_scaffold22.g24.t1 | GPCRsecr.HG2.9:like:ADGRG7 | NA | Morvir scaffold1040.g1.t1 | GPCRsecr.HG2.9:like:ADGRG7 | NA | Morvir scaffold353.g17.t3 | GPCRsecr.HG2.9:like:ADGRG7 | NA | Nemnom g5106.t1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Nemnom_g15841.t1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | 100 Nemnom g5202.t1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Nemnom g11786.t1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | 100 Nemnom g8779.t1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g1761 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g33666_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA -■Hvul g4685 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g17569 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g4758 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g7572 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g7582_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g7591 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g7583 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g447 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g33161 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g33148_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g33159 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g3327 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g15777 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g17053 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -•Hvul_g15557_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -Hvul g17048 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | 100 - Hvul g17042 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g21975 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | ─■Hvul g2597 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -Hvul_g26506_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g12669_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | _-Hvul_g26505_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g2598_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g11612 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul_g30234_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | ---Hvul_g3365_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Hvul g812 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -Hvul g11611 1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -•Hvul_g27125_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | -Hvul_g27127_1 | GPCRsecr.HG2.8:like:ADGRG7 | NA | Dgig XM 028555939.1 | GPCRsecr.HG2.11:like:ADGRG7 | NA | •Dgig XM 028547803.1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Exapal XM 021040556.2 | GPCRsecr.HG2.7:like:ADGRG7 | NA | ■Nvec v1g199785 | GPCRsecr.HG2.7:like:ADGRG7 | NA | 100 Spis_XP_022796933_1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | -Adig XM 015920411.1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Morvir scaffold33.g72.t2 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Hvul g19529 1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Nemnom g12653.t1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Morvir_scaffold476.g13.t1 | GPCRsecr.HG2.7:like:ADGRG7 | NA | Bralan BLAG07001106 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Skow XM 006823449.1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 Cgig XM 011425807.3 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cgig XM 011440923.3 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA122466 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Bralan BLAG19000174 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | 95 Bralan BLAG13000080 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Bralan BLAG17000514 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | 100 ■Bralan BLAG17000088 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Bralan_BLAG02002080_1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Bralan_BLAG04000221_1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 ■Bralan BLAG17000572 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 Bralan BLAG16000207 2 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Bralan BLAG90000455 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | _■Bralan_BLAG90000456_1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 ■Bralan BLAG90000454 1 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | •Drer ENSDART00000188437 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | -Hsap ENST00000273352 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | 100 -Mmus_ENSMUST00000023437 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint_ENSCINT00000017701 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000003422 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 Cint ENSCINT00000025358 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000037295 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000014593 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | -Cint_ENSCINT00000025518 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000006340 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | 75 Cint ENSCINT00000020231 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000023760 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000032398 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint_ENSCINT00000009663 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Cint ENSCINT00000009670 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 Cint ENSCINT00000012957 | GPCRsecr.HG2.5:ADGRG7 | ADGRG7 | Spur XM 030984230.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Hvul g9482 1 | GPCRsecr.HG2.5:ADGRG7 | NA | -Cint_ENSCINT00000036783 | GPCRsecr.HG2.5:ADGRG7 | NA | Cint ENSCINT00000001727 | GPCRsecr.HG2.5:ADGRG7 | NA | ■Mlei ML0350.g2.i1 | GPCRsecr.HG2.5:ADGRG7 | NA | Horcal Hcv1.av93.c3.g807.i1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Horcal_Hcv1.av93.c3.g807.i2 | GPCRsecr.HG2.5:ADGRG7 | NA | Tcas_TC000764_001 | GPCRsecr.HG2.5:ADGRG7 | NA | Tcas TC033064 001 | GPCRsecr.HG2.5:ADGRG7 | NA | Tcas TC001376 001 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 •Dmel FBtr0085737 | GPCRsecr.HG2.5:ADGRG7 | NA | •Dmel FBtr0085739 | GPCRsecr.HG2.5:ADGRG7 | NA | Tcas TC016199 001 | GPCRsecr.HG2.5:ADGRG7 | NA | Tcas TC034763 001 | GPCRsecr.HG2.5:ADGRG7 | NA | ■Tcas TC009282 001 | GPCRsecr.HG2.5:ADGRG7 | NA | ■Tcas TC009526 001 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030972703.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030978818.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 031000100.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ⁶⁴ Spur XM 791682.5 | GPCRsecr.HG2.5:ADGRG7 | NA | -Spur XM 030988911.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988629.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _ Spur_XM_030988152.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 84 Spur_XM_030996579.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011664823.2 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030990758.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030990453.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030973713.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030994279.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030987829.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988269.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ¹⁰⁰ Spur XM 030994300.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030973701.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030989817.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011666426.2 | GPCRsecr.HG2.5:ADGRG7 | NA | ¹⁰⁰ **■**Spur XM 030990454.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030974267.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030974407.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Cint ENSCINT00000018311 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030986587.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030974607.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_031000105.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011683783.2 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030975569.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 782944.5 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030992369.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030991278.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 -Spur XM 030996477.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030996977.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030996569.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030987485.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030987086.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ■Spur XM 030983803.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011680752.2 | GPCRsecr.HG2.5:ADGRG7 | NA | ¹⁰⁰ Spur XM 030990497.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030990460.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ¹⁰⁰ Spur_XM_030988103.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030990200.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988101.1 | GPCRsecr.HG2.5:ADGRG7 | NA | -Spur XM 030988455.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988100.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur_XM_030988102.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030994385.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ■Spur XM 030994727.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 - Spur_XM_030994384.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 001199941.2 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030994726.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030994378.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030995526.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _=Spur XM 030994382.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 99 - Spur XM 030995448.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_003726486.3 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030995769.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _ Spur XM 030996764.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 99 -Spur XM 011662845.2 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011674797.2 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030985019.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030975118.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM_030976170.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030999927.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _Spur_XM_030999911.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur_XM_030985243.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030999880.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030985240.1 | GPCRsecr.HG2.5:ADGRG7 | NA | ¹⁰⁰ Spur XM 030999886.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030985248.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030984798.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030985246.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030994347.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030999937.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _ Spur XM 030984784.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur_XM_030984785.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030995931.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030996434.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030995901.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030996367.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030984255.1 | GPCRsecr.HG2.5:ADGRG7 | NA _ Spur XM 030979144.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030979143.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030981221.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030982487.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030981505.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030982486.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030995884.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030989737.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030989740.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030988961.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 77 Spur XM 030988962.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030996041.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030987529.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030987531.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 Spur XM 030987532.1 | GPCRsecr.HG2.5:ADGRG7 | NA Spur XM 030978191.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030977553.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030978242.1 | GPCRsecr.HG2.5:ADGRG7 | NA 100 100 100 Spur XM 030978227.1 | GPCRsecr.HG2.5:ADGRG7 | NA | -Spur XM 030986358.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 - Spur XM 030978949.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030985199.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 775291.2 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030989078.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 99 **-**Spur_XM_030989399.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030989432.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988838.1 | GPCRsecr.HG2.5:ADGRG7 | NA | -Spur XM 030973648.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030973608.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 93 Spur_XM_030973676.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030973652.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988840.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 99 Spur XM 030973125.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988836.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030989003.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988833.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 LSpur XM 030973587.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030996558.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030997151.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _=Spur_XM_030997161.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030997451.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030983779.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030983778.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 011663400.2 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur_XM_030983619.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur_XM_030983617.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030983628.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030987908.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 100 Spur XM 030987909.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _=Spur_XM_030987527.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 85 Spur XM 030994248.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030988194.1 | GPCRsecr.HG2.5:ADGRG7 | NA | _=Spur XM 030996945.1 | GPCRsecr.HG2.5:ADGRG7 | NA 96 Spur XM 030995663.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030993108.1 | GPCRsecr.HG2.5:ADGRG7 | NA | Spur XM 030993803.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 90 Spur XM 030993808.1 | GPCRsecr.HG2.5:ADGRG7 | NA | 0.416667

─■Dgig_XM_028537776.1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | ─■Adig_XM_015922279.1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA | ■Spis XP 022805477 1 | GPCRsecr.HG2.0:like:ADGRG1/ADGRG2/ADGRG3/ADGRG4/ADGRG5/ADGRG6/ADGRG7 | NA |