

GENE	NCBI ACCESSION #	FORWARD PRIMER	REVERSE PRIMER
Ablim3	NM_001191698.1	GCATGACCATCTCAGAGTTCGA	ACAGCCTCCGCCTAGAACAG
Adora2a	NM_053294	GTCTGGTCCCTCACGCAGA	CACCTGTCCACCAAGCCATTG
Akap8	NM_053855.1	ATTGGCCAGGAGCATTTCTTC	TGGTGCTGTGCAGGAATCAG
Aqp4	NM_001142366.1	CTCGTGGCTGTCTGGGAACT	GGCAGCAAGATAATGGACCACTA
AUF1	NM024404	GGCAGAGGGAGCGAAGATC	CCCAGCTAAGGCCTCCTATAAAC
Arpp19	NM_031660	TGCAGCCCCGGATAAGAC	CTAGCAACAAGGGATGGTTTCC
b-glucuronidase	Y00717	AGAGGTTACGGTTCAGTGC	TCCCCCTTGTGTTGGTTCA
Brn-4	NM_017252	GGCTGATTCATCCACAGGAAG	TTCAGTACGCCCTTGACACT
C-myb	XR_086102.2	TGTGGGAAGACAGAGGACCAG	ACATGACCAGAGTCCGAGCTG
CaMKIIa	NM_012920	AGGATGAAGACACCAAAGTGC	GGAAATGACAGCCTTTGAACC
CaMKIIb	NM_001042354	ACCTCATTTGAGCCTGAAGC	CGATCCACACCACTATCCTG
CART	NM_017110	GCCAAGTCCCATGTGTGAC	CACCCCTTACAAGCACTTCA
Ccrn4l	NM_138526.1	CGCAAAGATCCCCTGTATAG	GAGGCTGGAGGAAGCAAAGTG
Cdh4	NM_009867.2 (mouse)	GCAATCAGCACTCTTCAACCC	CAATGAAGCATCCTGTGCTGAG
Clk4	NM_001013041	TTTGGAGCATAGGCTGCATTTC	GAGTATCCGCTCCATCATTGC
Ctip2	NM_001108057	AAATCCCAGGTGTCCATTGC	CTAAGTGCCTATCAGCGTGAGATC
D1R	NM012546	CCAGAAAGGAAGGGTAGGAGTTG	GCATGCACCTACCTTTCCAATC
D2R	NM012547	ACCCGGACCTCCCTTAAGAC	TGAACACACCCGAGAACAATGG
D3R	NM017140	TCACGTTCCTAATACCCACTGTCAAG	TTCAGGGCACTGTTACATAGC
DARPP32	NM_138521	ATTCGGATGTCTGCCCTTATTG	TGCATTCTGGCACCAAAG
Dpysl3	NM_012934.1	CGCCACAATGCGTTTGC	CCAGTGAGGAACCTCCATCAG
Dream	NM032462	AGGGCTTGACCAGCTACAG	CCGTGGGACATTCGTTCTTG
DRRF	XM001075745	AATGCGTCTCAGACAATCTTCTGT	CCCAAGAGTGAAGAGCACACAA
Dynorphin	NM_019374.3	CACCTGATCAGCCAGAAGC	GATCACAAGGAGGCAAGCTG
Ebf1	NM_053820	CAGTACTCCCTGTATCAAAGCC	CCCATCAAAGAAATTGTCCCC
Enkephalin	NM_017139	GGACTCGCTAAATGCAGCTA	GTGTGCATGCCAGGAAGTTG
Epha4	NM_001162411.1	CATGCTTAGAATGCCTGTGAATG	TGGTCTCCGGCTCTCGTATATC
Fbxl16	NM_001009504.1	CTGACGCTGGTCTAGAGGTGATG	GCCTCGGTGAAATCGTTACAG
FoxP1	NM_001034131	CAGCCACGTGGAAGAATGC	ACGTTCTCTACTCGCACAAAACAC
Gnb4	NM_001013910	CCCTTGCTGCGGTTAAGC	TCCGGATAGGATAAAGTCAACAC
Gpr158	NM_001170326.1	ACAAAGTAGTACTCCTCCGGTTATCTC	GCAGGAGGCAGAGGAGTAAAGAG
Grina	NM_153308	TGAGCCCAGAAGAATATGTGTTTG	GGCCAATGATGGTGAGAATGTATAG
Ikaros	XM223600	GAGGTATTAGAATGGTGTGAGGTTTG	GCTGAGAACCTCTAGCTAATTGACTTG
Mag	NM_017190.4	TCAAGTCCCGCACACAAGTG	AGAAGCAGGGTGCAGTTTCG
Map1b	NM_019217.1	ACAGAATCCGATGGGAAAAC	GGTGTGCCAGATTGTGTAGG
MBP	NM_017026.2	CTCTGGCCACTTCTCACTTTAGG	TTCTAAGAGGCAGGGAATGGTTAC
Mobp	NM_012720.1	AACAGCCAGCACGAAGCAG	GTGTTACCAGAACCTAGGAGCTCTG
Mog	NM_022668.2	TCCAAGAACAGGGACAATAACAAAC	CCGTGCAGAAGTCGAGAATCTC
NKB	NM_019162	TGATCTCTCTCTGCTACCTCCAC	CCCTGTCTTTATGATGCAGTCC
NK3R / NKB-R	NM_017053	CCAACTACTGCCGCTTCCA	GAAATGTTGCTTGGACCTTCT
NM23B	NM_031833.2	CCGATCTCTTTCTCGGCACTC	TCGTGGGGATTCTGCATTC
Plp1	NM_030990	TGCTGTGCCTGTGTACATTTACTTC	ATCAGCGCAGAGACTGCCTATAC
Rapgef5	NM_001047915.1	GGAGCTAGCGGCTGTTATTGC	TCTGTACCGGTTACGTGTTTGTGTC
RARbV1	XM001059459	CGTCTGCCAGGACAAGTCATC	TCGGAAAAAGCCCTTGCA
RXRg	NM031765	AAGCAGCTCTTACCCTTGTTG	TGCCCCGAGCAGAATGAC
SLC1A2h, rat Est	FQ082741.1 - Est	AAGTCCGAGCTGGACACCAT	ACTGCGTCTTGGTCATTTCGA
Tf	NM_001013110.1	CAGCAGCAATGGGCAGATAG	GGCATCTGCTTCTCCATTAC
Trim28	NM_053916.1	TCTGCCGTGTATGCCAGAAG	GGGAAGGGAGGTGACAATCC
Tubb2b	NM_001013886.2	CCTCTGTGTAGTGGCCCTTTG	GTCAGATCCGGACCATTGGG
Tubb3	NM_139254.2	TCCGCTCATGGTAGCAGACA	CGACAATGAAGCCCTCTACGA
Ube2v1	NM_001110345.2	TAAGATAGAGTGTGGCCTAAGTATCC	ACTGCTCACGCCACTCATATTG
Znf521/Evi3	NM_001107403	TCCTGGCTACATTTGAAGCTTTG	TCAACCGATTCAAGGATTTGG
S01		available upon request	
S02	CK598023.1	TGAACAAGGCTAATGTCCGAAAG	CCAGGGCATAGAATAGGAAAACAC
S03	NW_047512.2	CAGTGGGCAGTGATGTTAGCA	CACGATTGGTTTTGGAAATGG
S04	CK477547.1 cDNA	TAATAGCCACGTGAGAGGCAAAC	GCAGCATGTCAGCCTTCATTC
S05	NM_001109161.2	TCCAAGATTGCAAGGATTCC	CCCGTCACTTCTGAGGTTGTC
S06	FM098268.1	GCCGTTGGTAAATGAATAGATTCTG	GGTGGTGTAGGCACACATGTAGAC
S07	NM_001108951.1	TGTGCCTTGCACTCAATCACTAC	CTGGAACGTATCAGCATTGTC
S08	CB744760.1 cDNA	AATTTGAGAACAAGCCGAGCTT	AAAGGCGTTC AAGGTGAGA
S09	NW_047696.2	TCTCACAGCTTGCATGTCCTAAC	TCGTGCTGTGGA AAAACC
S10 Map1b	NM_019217.1	CCAGTGGATCAGTGACAGTTTTG	AAGAACCATAGCCAGCCTTCAG
S11	XR_086213.1	TCTTCTACATGTTGACCTCCAGTTG	CACAGTAGCCAAAACCATCAAATG