

* = intron location

A = editing site

adar (CG12598):

CAGCACCTTGTGGGGATGCACGGATATTT^A¹GTCCTCACGAAAACGACACTGG
TGTTGATAAAC

caps (CG33653):

GACACAAGGTCTAAGCGGTAAACTT^A¹TGTCTGTGCTAGAGTCGACTTTGTC

caID (CG4894):

CAGCGGAATGTGTTATGAAAATTTT^A¹GCATATGGTTTTGTGTTACAT^A²ATGG
TGCATATCT^A³^A⁴GAAATGGATGGAATTTATTAGATTTTATTAGATTTTACAAT
TGTAGTTAT^A⁵GG*GGCGATAAGTACTGCACTCTCCC

cpx (CG32490):

AACTGAAAAATCAAAT^A¹GAAACGCAAGTA^A²^A³TGAGCTAAAAACTCAAATA
GAGGGAAAAATGTGTCATGCAGTGA

da5 (CG32975):

CTACTTCCGATGCGGTGCCATTGCTGG*GT^A¹CATATTTCAATTGCATAATGTT
TATGGTAGCTTCATCCGTTGTGTCAACGATTTT^A²^A³TATTAAATTATCATCATC
GAAATGC

TGACAGACTGTGCCTTATCATATTCACAATGTTT^A⁴CA^A⁵TATT^A⁶GCCACAAT
^A⁷GCTGTACTACT^A⁸TC^A⁹GCACCACATATTATTGTCTCGT

dop (CG18314):

AACTGCTCTGGCTGAAA^A¹TTT^A²TCGAACCCT^A³GTCAT^A⁴TGATGTCATTTCGC
ATT^A⁵GTCTTTGCATTCTGGGTGTCCTGGCTGCCATGGATTCTG

TGCTTAACCATTTGTTGTAAGACT^A⁶AGGGCCGTTTGCAAGCAGAGCT^A⁸^A⁹TC
GGGTTGGACCCAGATGACT^A¹⁰GAGTTAGATTTAGATTTTCATTCTCC

eag (CG10952):

CTGGATACCGAGAAG*GTACTAAACT^A²TTGTCCGAAAGATATGAAGGCTGAC
TATGTGTTTCATCTAAATCGCAAAGT^A⁵TTT^A⁴ACGAGCATCCGGC^A⁵TTTCGTCT
GGCTCGGATGGTTGTC

CTCAGACATCGACTGATTTTTTCGCA^A⁶GGTGGCCGATGTGAAGCGCGAAAAA

para (CG9907):

CAATTTACGACTGAAAACCTTTTC^A¹ATTAATTGA^A²^A³ATAAATATTTTGAAAC
AGCTGTTATCACTATG

rdl (CG10537):

GTCAATTTTGACCGATCCTCGTTT^A¹GCGTAT^A²GAAAACGACCTGGTGTAGAA
AACT

CTACATACCCTCTGGACTGATCGTT^A³TTATATCATGGGTATCATTTTGGCTAT
GGGTATCATTTTGGCTCAATCGC^A⁴ATGCAACGCCGGCGCGTGTGGCGT

GTCGGCTACATGGCAAACGAATTCA^A⁵^A⁶TGCGAAAACAAAGATTTATGGCG
AT

sbd (CG6798):

GCCCAGCGATTCGGGCGAAAAG*GTT^A¹CGTT^A²TGTATTTGATACTCGTGTC

shab (CG1066):

TCGTTTCAATACCGGAA^A³CATTTTGGTGGGCGGGTATTACAATGACAACCTGT
TGGCTACGGGGACATCT^A⁴TCCCACAACCTGCACTGGGAAAGGTTATTGGT^A⁵C

CTGTGTGTTGCATATGCGGTGTTCTGGT^A⁶^A⁷TCGCTTTGCCTATTCCCATCATC
GTTAACAATTTTGCTGAATTTTATAAGAATCAG

stn (CG40306):

GTTTACTATGCCCGCCACGCAGGTCTCCCAT^A¹CCACCGTGCCTCGGTGAGT
GTCCAGGATTCGGATGGCGATGAACCGC

syt (CG3139):

TTGAAGAAGATGGACGTGGGCGGATTGTCTG*ATCCATATGTGAAAATTGCA
ATCATGCAAAAATGGCAAACGTTTGAAAAAGA^A²GAAGACAAGT^A³TCAAAAA
ATGCACCCTCAACCCTTACTATAATGAGTCGTTCTCATTGAAAGTACCATTG
AACAAAT^A⁴CAA*AAAATCTGTCTCGTTGTGACCGTCGTGGACTACGATCGTA
TTGGCACCTCCGAACCCATCGGCCGCTG

unc (CG2999):

ACGCAATTGACGCAGTGTTTCGATGTTGTT^A¹GCAAACCTCGAGTGTCCCTGATC
CAGAAATTTGGA