>DroMel seq1357_2R_18937351_18937843 2R 18937351 18937843

CAAAAGGACAC GGACACGCAT CCTGGCATTG TTTTCAATTC ATTTACGCGT CGACTCAAGT

CGCTCGCCCA CACATATGTT GGGAAAAATAA TCGAAAAACG GTTTGTCCCT TGGCATGAGG

TCCTTTTTGC TGCGCAGCTA GATTTGCAGT CCCTTGGCGC ATAGCACATG CCATTGATTA
9.2

CCATGTGTGG GAATTAGCCT TGCGAAAAAAA GAAGTTAGTT AAAATCACAC ATAAAGAGGC

AATGCCAATG GCTGTAGGTC CCTGATCCTC CGTCCTGGCA GAGAGCTCAC ACGTCCTGCA

GCGATCCTTC CACCCTCTGC ACGTTCTAAA TTTAGCCGCT GGATTTATGA CCCCTGTCAT
7.3

TTGAGCCGCT CTTCCAGTAC TTCTTAAAAAA ACCCCTGGAA TTACATGCAG TCGAAGAAGG

AAGAGAGTGG AGGAGTCCAA TGAGTCGAAG GAGCCCAATG AGTCGAATGA GTCGGAGAGG

GGGCGTTGAC AGT

>DroMel seq1396_2R_19526211_19526451 2R 19526211 19526451

CAGAGGCCAT CGTGTCATCC AAGATTATGA TTTTGACAGG CATCACGACG ATATGACATA

TGCACGAATG TCTCTTCGAT CTATTTTAT AGGAGAGATA TCTGTCAGCA CATGTTTGTA

ACCCTACACT CGTGCCCTGC CTATAAATAA AAAAGTTAAG ATCACAATTT TAAGGAGATC

TTTGTTATAA TATATTGATT AATTTATCTC GAAATACCTA CAATTCTTAT CCCCATTAGC

C

>DroMel seq1397_2R_19526657_19527033 2R 19526657 19527033

>DroMel seq2300_3R_10415381_10415780 3R 10415381 10415780

TTGTTAATAT CTTAACTTT CAGGATATAA CATGCTAAGT GTTATATTAT ATAACATTAA

AGTTATTTCT GCCAGTGTAA CGTTGCAGAG TCACTTGCTC CGCAGAGAGC CGCCACTCAA

7.3

AAGATCAACC CTCGATGGCA GACTCTCGCC TTTATGGACT ACGAAATGTG CAATTTCTGC

TCGGGTACCG AGTTTTACGA TTTTTCCATA GGTCCCAATA TGCCGGGCAT CTGTGTGGTA

10.6

GGGGTTATTT GGTTGTTGGC CCAGTATCGA TGGCAGGCCC GATTTTAGAC CCAGTCCACT

TAGGAATTCC ACGGACAGAC CACATGGACT CTCTAGCATT TTGCCACTTC AGTTGTTGCT

11.0

CAGTTTCAGT TTTTTTTTTC CTAGTTGCAC TGGGGAAAAA

>DroMel seq257_2R_5025930_5026695 2R 5025930 5026695

AAGTCCACGA CAATCAACAT GCATCGGAGC ACAATAAAAA GGCAAAAGTG GAAAGGAAGA
CTAGGCAAGA CCGAGAAGAG TGACAAAGAG CGAGGGCGGA AGTGCGGCGA GGACAAGGCT
GATCTGCGAT GAACACCAGG TAGAAGGTAA GTGTATCCAC CGCACAGATT TAGCCAAAAC
AGTCCAAAAC ATATGTATGA AAGCAACTAA AAGAAACGGC GATCAAAACA AACAAATGTT
TGGTACCAGT GTACCACAGA TGTAGAAACC AGACGCTTTC CAGACTATGG CAGCACCCCA
AGAAACTCTT GGAAATACTA CTCATTGCTA CACCGCACAT ATGACGTACA CTTGGAGAAA
CAAGCACATT CCAAACAATG CGTATGTACC TAGTTATTTC ATTACAAGGA ATTGAATTTA
ATGTACCCAA AGTTAAGCAG TAGCATTTAG TATTTATGTT TTCTGGATAT ATTAACTGAA

TGAATAGGAA ACTTGTACCT AAATCCATTT CAGATCCGTG CCATTTCTTC CAGTGCACGA
GCGGCTTTCC CATGTGTGT TGCTGCGTGG TAGCCACTCA CACAGCGGCA AAACACGACG
CCTTCGCATT TAAAATTGAT TAAAACTAAA AACACTCGAT GCCGACGCCG TGCCTTGTTT
CGTTTTCAGT GCGATCGCAG ACATTGTGCA AAACGGTTCA CAGTTCTCTC CTCGCGACTG
CAGGTGATCG AGTTTTTTC GTTTGAGTGC ACTAAAAAAA AACAGA

>DroMel seq2820_3R_17227574_17227958 3R 17227574 17227958

>DroMel seq2821_3R_17228198_17228402 3R 17228198 17228402

TGTGGCATAT GTGTATATTT TCCATATGTT AAAGTACATT ATTTTACGCA TTTACTGAAG 8.0

AAACTTCGTC TAAGTTAATA TATTTTCAGA CAGTTCTTAT GGAAATTATA AATATCATTA

ATGCCTCGAT CTCTATTTGT TGTTTACAGT ACAGAAACGA TTTTTGGCCG TGTATCCCAC 7.7

CACTCGACTA GTGCATTTAG TGCTG

>DroMel seq2837_3R_17392100_17392639 3R 17392100 17392639

TGCAGTGAAT GCAACTTGTA AATCTGTCAC GCGTCCCAGT AAATCATCCT GCATAAGTAA

TGTGCAGTTT AGCCCGGAGA ACACACCCAG AATTCGAAAA CGATACTCCC CTCGAACCAC
9.2

TTTGTCACTC GAGATACTCG AAATACCCGA GGAGACACTG CGGGATACTC CGGAATACTC
7.8

AAAGATACTC CTCTGACTGC AACAGATGCC GGAATATAAT AAAGTCGGGG GAGTGTGAAA
8.6

TTGTGTGCTC TGCGGCGACT GAAGGCGGTC CGCGGCTAAT CTATGACAGA AAACATGTAA
10.7

TCGGAGCGAA ATGCCCAAA AGGCCTGGCC CCTGGCACTA AAGTTGCTCT GGGTGGCGAT

TCGGCGGCCA ATTAAAATCC CAATCAGGGC TCTACTGTAA TACATTGTGC GTCGAATTTA

GTTGTTCAAA TTAATTATGG TACACGGGAA ATAAAAATGA GTTCGTAGAG CGTAGAAAAA

>DroMel seq6531_2L_21853080_21853669 2L 21853080 21853669

GAACGAACCG TCGAAAAAGA AAGAGCCTCA GAGAAAATCT CATGGACTGA AAACCATAAA
TTTGATAATT GACTTTCCAC CAAAGGAAAA GAAGCCAAAG GAACCGAAAA ATTTCTATTG
GCTTACACAA TCGATGAAAC ACATAAATCT TAAGTGAGGG AATTTAGAAA ATCCAGCCAC
CTTAAAATAGA CAACTGCAAA ACGGAAATGT ACATCATAAA AATATTAAAA ATTCTCTGTC
ATTCTTCTAA TATGGTCTTT TTGGACATTC TTGGAAAAGAT AAAAGTCGAT GCCGGTTGGC
CGCCGTTCCA GTGCATTAGA GATCGCGTTT AAAAATGGTAC CCCGATGGGC TGTCAAGGCC
GTAAAATCTAG AATAGGGGCA CTCTTTGCTG CCCTCTCGTG CAGCTCATGT ATTTTCATAA
TTTTCATATT TCTCCGTGGC TCAGCAGCTG CGCGCCTGCG CTAAAGTGTC TGACGGATGC
GGCGCATTCG ATTTCGTTCT GCCAAAATTAG CCGGGCCCCG TGATTTATGT CGTCCTTCTG
8.4
GCTCTTTGGC CGTTTGTCC TTTTAGCCAG TCCGCGTGAC TCGAGCGTTG