

<i>CG5755</i>	DMv3 (-6)	2L CG5755 protein_coding 1 simulans:chr2L:1 sechellia:contig_1645:+1 yakuba:chr2R:-1 erecta:contig_7039:-1	GAAGTCTACCACATCTGTCATTACAGACTCGATTTTAAACTATAA GAAGTCTACCACATCTGTCATTACAGACTCGAT----- GCAGTCTACCACATCTGTCATTACAGACTCGAT----- GAAGTCTACCACATCTGTCATTACAGACTCGAT----- GAAGTCTACCACATCTGTCATTACAGACTCGAT----- GAAGTCTACCACATCTGTCATTACAGACTCGAT----- ***** **
<i>CG8186</i>	DRE (-29), INR (+12)	willistoni:contig_9942:-1 ananassae:contig_18824:-1 pseudoscura:Ch3:+1 persimilis:contig_9194:+1 2R CG8186 protein_coding 1 simulans:chr2R:+1 sechellia:contig_653:+1 erecta:contig_7056:-1 yakuba:chr2R:-1 grimshawi:contig_21591:-1 mojavensis:contig_10114:+1 virilis:contig_14923:-1	TAT-TGATTGTC--GGTGACCAATCGATTTCCAGATTTG--CAACCTGGTGGTTC TAT-CGATTTAT--GGAATGGGCTTA-----TCAACACTGGTGCCAGTCA-----TTCC TAT-CGATTTCT--TGCAGTTGCTAG-----ACAGCACTG-----CAAATTC TAT-CGATTTCT--TGCAGTTGCTAG-----ACAGCACTG-----CAAATTC TAT-CGATTTGC--TACGGGCGAGTAG-----CCAACACTG-----TCATCGTCAATTTCC TAT-CGATTTGC--TACGGGCGAGTAG-----CCAACACTG-----TCATCGTCAATTTCC TAT-CGATTTGC--TACGGGCGAGTAG-----CCAACACTG-----TCATCGTCAATTTCC TAT-CGATTTGC--TACGGGCGAGTAG-----CCAACACTG-----TCATCGTCAATTTCC TAT-CGATTTGC--TACTGGCAGTAG-----CCAACACTG-----CCATCGTCAATTTCC TAT-CGATTTGC--TACGGGCGAGTAG-----CCAACACTG-----CCATCGTCAATTTCC TGCATGATTCGTGTCAAACTTTTA-----CCAACACTGAAATTTATCA-----GTCC -----TCACTTTC--TG--AGGCTTAG-----TCAACACTGTGTTTATCA-----GTCC TATCTAAGTTAC--TG--TGGCTTAA-----CCAACACTGTTATTTATCA-----GTCC * * * * *
<i>CG7975</i>	INR (-19)	mojavensis:contig_10184:-1 pseudoscura:Ch3:+1 persimilis:contig_2161:-1 willistoni:contig_6314:-1 ananassae:contig_18793:+1 yakuba:chr2R:-1 erecta:contig_7049:+1 2R CG7975 protein_coding 1 simulans:chr2R:+1 sechellia:contig_2003:+1	TCATAGC-AATCGGTGCTGTATGGCATTITACT---ATTAITTAGCATGTCTGAAAT GAATCCC-AGTAACCCCTGC---AAACTTCAAT---CAAATTCAG-----TCACAAATC GAATCCC-AGTAACCCCTGC---AAACTTCAAT---CAAATTCAG-----TCACAAATC GAAACCCAAACACCTTACATCCAAATTAACA---ATCTTCTGTC-----TTTAAATA CTGGACG-GACTACTGACATTT---CCTTGCCACAAAAACACTCTG-----CTTAAAT GAAACTC-AGCCTGATGTTCTACCTTCGAA---AATGTTCAG-----TCGAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG GAAACTC-AGCAACTGATGCTCTACCTTCGAA---AAGCTTCAG-----TTAAAGCG * * * * *
<i>CG5150</i>	DMv5 (-36)	grimshawi:contig_20566:-1 mojavensis:contig_11553:-1 virilis:contig_17426:+1 willistoni:contig_6203:+1 pseudoscura:ChXR_group6:-1 ananassae:contig_19344:-1 yakuba:chr3L:+1 erecta:contig_6553:+1 3L CG5150 protein_coding 1 sechellia:contig_868:+1	-----GABCTATCATCA-CTTGAATTAATTGCAGCGCTAAA-----AGCGCT -----TCGTATCATCA-CTTCAAAATAGTAGTCTCGGTAA-----CGTGGT TCAGCGGTCTCAATCATCA-CTTCAAAATAGTAGTCTCGGTAA-----CGTGGT -----GTCTG-CGG-----AAT-----GGGT CC--ACTGCCCTATCGACT-CCTCACACGAGCGG-CAC--CTGATCTGGATGGGGG-- TCGA-----CTG-CACCAGT-----GGAT-- CCAACTGCCCTATCGACT-CCTCACACGAGCGG-CAC--CTGAT-----GGTA-- TC-----CC-GATCACACGAGCTG-CAC--CTGAT-----GGTA-- CCAACTGCCCTATCGACT-CCTCACACGAGCTG-CAC--CTGAT-----GGTA-- CCAACTGCCCTATCGACT-CCTCACACGAGCTG-CAC--CTGAT-----GGTA-- CCAACTGCCCTATCGACT-CCTCACACGAGCTG-CAC--CTGAT-----GGTA-- * * * * *
<i>CG12362</i>	INR (+4)	grimshawi:contig_20602:-1 pseudoscura:ChXR_group6:+1 mojavensis:contig_11641:-1 3L CG12362 protein_coding 1 simulans:chr3L:+1 sechellia:contig_48:+1 yakuba:chr3L:-1 erecta:contig_6564:+1 virilis:contig_17463:+1 ananassae:contig_19353:+1 willistoni:contig_8074:-1	TTTTGAGATGTTCTTTCATCTGTCTGTCTGTGGAACATAAAG----- -TACAGTATCGTTTCGCTACTAAA---AAGTGTCAA----- -ACTCACAACATTTAATTACATCGA---AAGTGTCAA---TATCAAGCAACACAGAATAA -CTTGACAATTAACAAAGTAGTTTGTCTTCAGTA---GAAAAA----- -CTTGACAAGCACTAACAAGTAGTTTGTCTTCAGTA---GAAAAA----- -CTTGACAAGCACTAACAAGTAGTTTGTCTTCAGTA---GAAAAA----- -CTTGACAAGTAACTAAGTAGTTTGTCTTCAGTA---GAAAAA----- -ACTCACAACATTTAATTACATCA---AAGTGTCAA---TATCAAGCAACACAGAATAA -ACACACAACATTTACATCACTCGA---AAGTGTCAA---TATCAAGCAACACAGAATAA -ACACACAACATTTACATCACTCGA---AAGTGTCAA---TATCAAGCAACACAGAATAA * * * * *
<i>CG7815</i>	TATA (-64)	erecta:contig_6579:-1 yakuba:chr3L:-1 3L CG7815 protein_coding 1 simulans:chr3L:-1 sechellia:contig_149:-1	AAGCGGCTACCAGAAAGTTTCCCTCT-----TTCCAAAGTAGTTGGATAT ---ACCGATCTACCACACTTATTCCTTAAATTTTGTGTGCACACAGTAGTTGGGATAT -----TTTTCTTGACATTTTTCTGCACAAAAGTAGTTGGGATAT -----ACTTTTTCTTACATTTTTCTGCACAAAAGTAGTTGGGATAT -----ACTTTTTCTTACATTTTTCTGCACAAAAGTAGTTGGGATAT * * * * *
		erecta:contig_6579:-1 yakuba:chr3L:-1 3L CG7815 protein_coding 1 simulans:chr3L:-1 sechellia:contig_149:-1	ATAAGTAGATTTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT ATGAGTAACGTGTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT ATAAGTAGATTTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT ATAAGTAGATTTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT ATAAGTAGATTTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT ATAAGTAGATTTTTCTGTCGCAAAAACCTAAAATAAGTCAGAAAACCGATCC-----GT * * * * *

<p><i>CG6050</i></p>	<p>DMv4 (-16, DMv3 (-14), DMv2 (-15)</p>	<p>simulans:chr2R:+1 willistoni:contig_6315:+1 mojavensis:contig_18277:+1 virilis:contig_18897:-1 pseudoobscura:Ch3:+1 persimilis:contig_2861:+1 ananassae:contig_18823:+1 2R CG6050 protein_coding 1 sechellia:contig_608:+1 yakuba:chr2R:-1 erecta:contig_7868:-1</p>	<pre> -----TGGC-----GGG---G ATTACTATCGATACCTAG-----AGCA-----GCAGCACATTTTGTGAG -----AATGCTAATATTACAGTC---TGGCAACTCTCSATAGCAGCA----- -----AACACTTAAATTATCCGATC---TGGCAGCACCGCTGAGCAGCACATTTTAA---G -----CTATCGATAAAATTTGATCAGGTTGGCAGTTC--ASTGGGCAGCACATTTGTAG---A -----CTATCGATAAAATTTGATCAGGTTGGCAGTTC--ASTGGGCAGCACATTTGTAG---G -----CTATCGATAAAATTTAAGCTAGA---TGGCAACTCTGTC--CGCAGCACATTTTCAG---G -----CTATCGGACAACCTCCACCATG---TGGCAACTCTGTC--CGCAGCACACATTTTCAG---G -----CTATCGGACAACCTCCACGAGG---TGGCAACTCTGTC--CGCAGCACACATTTTCAG---G -----CTATCGAACAACCTACACGAGG---TGGCAGTCTGTC--AGCACAGCACATTTTCAG---G -----CTATCGAACAACCTCCACGAGG---TGGCAGTCTGTC--CGCAGCACACATTTTCAG---G *** </pre>
<p><i>CG9920</i></p>	<p>INR (-13)</p>	<p>virilis:contig_17264:-1 grimshawi:contig_18124:-1 pseudoobscura:Ch2:+1 persimilis:contig_348:-1 erecta:contig_6454:-1 yakuba:chr3R:+1 3R CG9920 protein_coding -1 sechellia:contig_232:+1 simulans:chr3R:+1 mojavensis:contig_11042:+1 ananassae:contig_19618:+1 willistoni:contig_8648:-1</p>	<pre> CTTAACCACAATCCAGCATACCTCAGTCTGTTTCTGGCATTGCAACT-----G CCAGTTTTTCA-----TATTGCTCAGTCTGTTTCCGGCAATTGAACT----- TACATTTGTAGGACTAGGAAAAATCAGTCTCCAGTCGAGAGTGTACC-----A TACATTTGTAGGACTAGGAAAAATCAGTCTCCAGTCGAGAGTGTACC-----A -----TTTCAGTCAGTCAACAGCTGGAAAGTATAGG-----A -----TTTCCCTCAGTCTACAGCTGGAAAGTATAGG-----A -----TATTCCCTCAGTCTACAGCTAGAAAGTATAGG-----A -----TATTCCCTCAGTCTACAGCTAGAAAGTATAGG-----A -----TATTCCCTCAGTCTACAGCTAGAAAGTATAGG-----A -----TATTCCCTCAGTCTACAGCTAGAAAGTATAGG-----A -----TTATAGCACTCCAC-GTAAACGCTGAAAAAATTAATAAAAAATAA -----AAACATCGGTGCA---ATAAAAAAAGT-----A -----GTAACAAACATTTATAAGT-----A ** </pre>