

<b>Name</b>	<b>Symbol</b>	<b>Primer AS T3</b>	<b>Primer S T7</b>
CG10041	CG10041	TCAAGCGTTTTGAGGAAAAC	CAACTGTGTGCGACCACGAAG
CG10225	RanBP3	GTTGTCGATGGCCGTTAGTC	CAAGTGCAGGTGAAGACAGC
CG10576	CG10576	TGCTTTGCTTTTTCGTTGC	TTAAGCACACGGTTCTGCTC
CG1058	rpk	GGCGCTTCAGATTGGTAAAG	TAGGCCCCGAAAACGTTAGTG
CG11155	CG11155	CTTCGTTTTCGCGTCTATG	TCGACAATGCTGGACTATGC
CG11212	Ptr	CGTCGTCTAAATCGCTCCTC	TCACCGCTCACATCTGCTAC
CG11263	CG11263	TAAGAGTTGGAACCCGAAGC	GATGGAAACGAGAGCTTTGC
CG11486	CG11486	TGCGTCCATGATCAGGTTAG	GACGCCCACTTCTATGCAC
CG11584	CG11584	TGTAGGTCACCTGCTGTTGC	AATTGCCATTCCAGTTCCTG
CG11897	CG11897	AACATAATCCTGGCGCAGAC	GCTCAGCTTTATCCCCTGC
CG12052	lola	TGTAGGGGCGAGTAGGGACAC	TAGATCCCTCGACCATTTTCG
CG12224	CG12224	GTTAATCCGCAGCAGCTTTC	GCCAGCTACTTTTCGTCAGG
CG12367	Hen1	CAGCGTTCCTCCTCTTGTTT	AAATTCTGCAGGGGAATGTG
CG12952	sage	GGAGCGCAGTAAGTCAAAGG	AGTTTTGGTGCCCGTATCAG
CG1303	agt	ACTTTTCATCGGCCAGAAGC	AAACACCCAGATTCGTTTGG
CG13083	CG13083	TGGTGTGACGGATTTTTGTG	TGAGGATCTGGACTCCATCC
CG13848	pinta	CATAGAAATCGGCGTTCTCC	CTGCTGCTAAAGCTCGATCC
CG13890	CG13890	CAGCTGTGCGATTCATTCAC	ATATGCTGCCCCTCATCTTG
CG13937	CG13937	CGTAGAGCTCCAGCATTTC	TGCGTGAGGCACAGAACTAC
CG14394	NijC	TACAATGTGGCAAGCAAGTG	CATTCTGTTCGTGGTCATCG
CG14485	swi2	CACAATCCAAGGAACTCGAC	CGGGACAATCCCCACTAC
CG14590	CG14590	CTCCAAGCCAGCTTCGTTAC	CACGAGTCAATGGAATGCAG
CG14869	CG6107	ACTGGCCAATGCGATTGTAG	GTGAGGGTGTGGAGAAGAGG
CG15400	CG15400	ATATAGTGGCGAGGCCAAAG	TTCTCAGCGTTCATTTGGAG
CG15892	CG15892	AGTCATAGCGGGTGCCTTC	ATGTCTCGGATTGGGATGAG
CG17077	pnt	CGGTCCACGAAATGAACTC	ACTGTCAATGGCAGCGGTAG
CG17104	CG17104	GGCTGCCTCTTCCTTTACTG	TGCCATTCTGCTACTGATCG
CG17134	CG17134	GCCAGGGTGAAGACAGTACC	GCAGTGTTAAGGCCGAGAAG
CG17533	GstE8	TTGAGGAGGGTACCAAATC	GAGGCTAGTCCCTCCAGTTCG
CG1780	ldgf4	GAACCTGTCGTGGCCAGTG	TCGATGTGCCCGCTATTATC

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CG17803	CG17803	TGTTGCGCGTATAGAGATCG	CTACCGCATCACATTTGCTC
CG18188	Damm	CGTTCGACCTCAGCCTTAAC	GCAGTTCCTCAGGATTCTC
CG18255	Strn-Mlck	GCGCTGATAGCGATAGTTCC	TTATCGGGGTGAAAACGAAG
CG18278	CG18278	GTGTTGTTCCATGCATCCTG	TGGCGGGAAGTACTTGAATC
CG18410	Ude	GTCTTCTTGCCAGCTTTTC	CGAGATCCGGACTACCAGAG
CG1851	Ady43A	TTGTTGTGCGCTATCTGAGG	CTCTCGGTGCCTTACAAACC
CG2056	spirit	GAATTCGGGACTGAACTGG	GAGCCACCCAGTCAGGTG
CG2184	Mlc2	TTCATCTTGCCGTGTGTAGC	TGATCGACGACAAGAACCAG
CG2488	phr6-4	TCTCCTCGAAAGACGACTCC	GCATGGAGGAATCCCTAAG
CG2999	unc-13	ACCCATAAAGCGCTTGTGAC	GTCAAGGCCTCCAGAGACAG
CG30488	CG30488	TGACTGTTTCATCATGCATCC	TGCGAGCTTGATAACCAGAAG
CG3074	Swim	CAACTTCACCGAGTGGAACC	GTGGACCCAATCATCTCCTG
CG3081	CG3081	CACCACCAGAACCCTACCAC	CATCCAATCATGTGCGAAAC
CG31098	CG10634	TGCTCTCAAACATGGTCAGC	GAACCGGCAGCAAGAACTAC
CG31262	CG31262	ACCACAACCTGCTGTGACGAC	ATCGGTGGCTTTTCATTCTG
CG31436	CG31436	CTTCCAAAGACCCGATTGAC	AGGCAGAGGGACACAAGAAG
CG31839	nimB2	CTGCAGATGCAGAAGTTTGG	CAATTCGCGGGATCAGTTC
CG31875	CG31875	TTTTTGTTGGCATCAACAG	CACGCAAGGAGAAAACGAGAC
CG3200	Reg-2	TCCTCGCCGTATTTCTTGAC	TGCACAACCTTCTCCAACCAC
CG32055	CG32055	CTTTCGACGATTCCTTGAGC	GGGAATATTTTACGCGTTGC
CG3240	Rad1	AATCAGTGTTGAGCAAAGGAA	GCAAGTCTCCGAAGATGGTC
CG32444	CG32444	TTGGGGGTTGGACTTTATTG	GGGAGGTGGTACAGAAGTCG
CG32491	mod(mdg4)	AGGACTTGGTCGCATTGTTT	GTCTTTCCGTCGAGTTGTGC
CG32625	CG32625	TCCATTAATCGCTTGCCTTC	TGAAGTTGACGAAGGGGAAG
CG32788	Crg-1	AGTTGTTGCCGCTGCTTTT	TACTTCCTCCCGCTCCTACC
CG33048	Mocs1	CTTTGTTGCCAAACAAGCAC	AGCCGAAAAACAACCTCCTG
CG33140	CG33140	ATTGGCTTTGGTTGATCTGG	AACGGGTCCGATACCTTTTC
CG33302	Cpr31A	AGGCCGTCTTCTTGTTCTCC	CACCTATGCCGGCTATCATC
CG33397	CG33397	GTTAATCCGCAGCAGCTTTC	GCCAGCTACTTTCGTC AAGG
CG34002	CG34002	TCACCTTCACCACCATCATC	GTCACCAAAGTGCGGAAAAG

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CG34215	CG34215	GGTAATCTGCGTTTGGGTTT	ATTTGAAGGCTCCAAGATGC
CG3649	CG3649	TTGGTGATGCTTCTGTTTCG	GCATCGTGTGGTGCCTTAC
CG40178	CG40178	AAAACCAGTTGTGGCACTTC	ATGATTACGGTGCCGAAAAG
CG40300	<i>ago-03</i>	GTGCTTGAATGTAGGCATCG	TTTTCTCCATTGACTCTG
CG40452	<i>Snap25</i>	AAACCATGAGGACCGTTTTG	TCTAATGAAGCGCGGATAGC
CG41320	CG41320	CCAGTTGTGGCACTTCTCTG	GACATTGCATTTTATCAGAAGTTATCA
CG4174	CG4174	AGTTTGGCAGGATTCGATTG	GTTGTAAGTGCCTGCCTTGG
CG42516	CG42516	TCGGTCTGATCGTCTCTTC	TGTGCACGCACATTTCTAAC
CG43079	<i>nrm</i>	TTTAGGGTTGCCATTTTTGG	TGCTCAAAGACCCAAGGAAC
CG43103	CG43103	AAACCGAATGTGTCGTATTTTT	CCAAGCCGGAGGCTAGTAGT
CG43114	CG43114	TTCGCCTTTATTCCGATTTG	CTGACAACCTCATCAGCATGGA
CG43117	CG43117	TGCCATTATTAGAAATGTGTTTT	CGCACAATGAAGCTAAGCAC
CG43144	CG43144	TTGTCTGTTGGTTGTCAGTCG	GATGGCTCAGGCTTTCCTC
CG43646	CG43646	CAGGGTTCGGGTATAGCTTG	TTGGCACAGATTTCACTTCG
CG43676	CG43676	TTTATTGTCCTGTAATTTCTTTCTGA	ATGCACTCCGCATCACTGTA
CG4379	<i>Pka-C1</i>	ACTCCTTGGCACACTTCTCG	GTGTCATGATCGTCCAGCAC
CG45050	CG43674	CGTTCCATCTTTGCTCCTTC	TGGACAAGAGCATTGAGCAC
CG4570	CG4570	AGCTTGGCTAGCAGAACTCG	CAGCTTGCTCGCCATAAAAC
CG4594	CG4594	GCTCCCATAACCTTCTGCAC	TGGATTGCCCTCTATGGAAC
CG5229	<i>chm</i>	GCTGGAATCACTTCCCTCAG	GGATCCACCAGTTCGGAAAC
CG5671	<i>Pten</i>	ATCTTCCTCGCCATCTTCG	CTGAAGCGTCCGAAAAAGAG
CG6134	<i>spz</i>	GCAGCAGGAAGGGATCTTG	CTTTGTCTTCCCCGATTCC
CG6579	<i>atilla</i>	TTGCAGGTGTAGCAGCTCTC	CGGCCATCAAGTGTTATCAG
CG6658	<i>Ugt86Di</i>	TGGTACTGCCACCAGTTCAG	GTGCCCCACATAATGCTACC
CG6713	<i>Nos</i>	GATTACGGCAGCCAAAGAAG	TTCGCTGACTGTGAATACGG
CG7178	<i>wupA</i>	TTCATCAAGGCGTTTCAGATG	AGAAACCCGACTGGTCCAAG
CG7420	CG7420	CATTTTCGGTGCTGTTGTTG	AGCCATCACGTTGACCAAAC
CG7590	<i>scyl</i>	CGTATACTCCGACTGATGG	GGACCTGATGTCCAAGAAGG
CG8023	<i>eIF4E-3</i>	CCTCAACGGTGTTGAAGCTG	AAGGTTTCGTTTATCGCATTG
CG8573	<i>su(Hw)</i>	ATTTGTCAATGCGCAAGATG	ATTATGGAACGCACGGATTG

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CG8589	<i>tej</i>	CACAGCATAAGCTGGAATGG	CAAATGGCATGGATAACTGC
CG8768	CG8768	TTTCGGAGGGCGTGATTC	CGCTGTTGTGGGTCATGG
CG8780	<i>tey</i>	CAAGGCGTACTTCCTCTTGC	CTGCATAATGCGGGGAAG
CG8800	CG8800	GAAAGTTCGCATGGAAAACC	TTGAGAAGATATTCGGCTTATCG
CG9214	<i>Tob</i>	TGGCACTTATTGGCCCTATC	CAATGTGGACTGCGGTAGTG

Name	Symbol	Primer AS T7	Primer S T3
CG10102	CR10102/ <i>Arc3</i>	TACTCCTCGTGCTGCTCCTC	CTCATCCGCGAACACTTCTC
CG10382	<i>wrapper</i>	GCTCAACACTTCCGCTTTTC	GATGTGGCTCTGGTGGACTC
CG11357	CG11357	TCGACAGAGCATCAGATTGG	AAATCATCGCCGATTTTAGC
CG11797	<i>Obp56a</i>	ACTTGGAGGCGGTATCACAC	CCCTTAATCTGAGCGACGAG
CG11883	CG11883	AAGGGCAATGATGAGGTCAC	CAGGGAACCTCTGACGAGAC
CG12256	CG12256	CAAACCTTTCCAAGGCACAG	CATGAACGGAGTAACTTATTTTGTG
CG12287	<i>pdm2</i>	GACCAGAGAGTTTGGCTTGG	TTGAGGACATCGGGAAAGTC
CG14121	<i>l(3)S147910/ver</i>	TGCATTGTTTTATGCCTTCG	TGGATTTTAATCAGAGTTTCGAG
CG15739	CG15739	TCCGCATAGTAGTCCGGTATG	TCGACGTGGACTTCAATCTG
CG15784	CG15784	AGCGCTGCTGTCTTTTCTTC	CACCAAAAGGACCAAGAAGC
CG16756	CG16756	CGTTTCGGCAATATCGTTG	TGCTCTCGAATTGTGAGTGG
CG16928	<i>mre11</i>	TCGCATTAACATGCAGGAC	TCCTGTTTCCCTGGAGATTC
CG16947	CG16947	GCTTGTAGTGAGCGCATCC	CGCCGAATATCGCTAAATG
CG1725	<i>dlg1</i>	GCTCCTCCTCATCCTGACTG	GGATTCGGATACGGACTCG
CG1763	<i>nod</i>	ACCTAGCAGCTCCAGCAGAG	CAGCAGCAGCTCAAAGTACG
CG18024	<i>SoxN</i>	CAGAGCTGTATTCCCGCTTC	TGCATTGAGCAGCCTAACTG
CG18657	<i>NetA</i>	GATCGAAGTGGAATGGCTTG	GCATTCCGGACTTTGTGAAC
CG1925	<i>mus205</i>	CACGATCATTGCCAGTATG	ATTCGGCAAATCCAGATGAG
CG2706	<i>fs(1)Yb</i>	CCGAATCAATCCATTTTTGG	TGTTGCTCCAAGTCTGAAC
CG2914	<i>Ets21C</i>	GTCGTAGTTTCCCGCAGAAG	AATCCAGCTGTGGCAGTTTC
CG30196	CG30196	ATTCTCATTGGAACGCTTCG	CGAGAGTCTGGATGGCAGTG
CG3100	<i>b6</i>	AAATTCATTGCCACAAGC	TACAGACGCGTGGAAGATTG

<b>Name</b>	<b>Symbol</b>	<b>Primer AS T7</b>	<b>Primer S T3</b>
CG32261	<i>Gr64a</i>	GTGCCACTGTCTGTGTGGTC	CGTGCACTTTTGTGTGGAAC
CG32364	CG32364	GAAACAGCCCTTTTGTCTGG	CGAGGAATCACAGTTGTTCTCC
CG3289	<i>Ptpa</i>	CACTGCCCCAGATGAAGG	GGCAATTGCGAAGAAGGTG
CG33960	<i>Sema-2b</i>	ACACCGGCGACTGTACTACC	AGGACGTTTCGTGATTTTTTCG
CG34406	CG44002	GATATAGCCGCGTTCTCGAC	TATGCCACCATGTGTGAAG
CG42330	<i>Dscam4</i>	GCATTATTGTGTGCCGTGAC	TAGGTCCCTTCGTGGAAATG
CG42614	<i>scrib</i>	ACTCGGTTTCGTCTTGGATG	CGATGAGGATGATGATGACG
CG4398	CG4398	GATTTGCGCTTGAACCTCTGC	TGAGACTGGTGTCTGCAATTC
CG4432	<i>PGRP-LC</i>	AATGCGGGGTTATCCTGTC	TGTTCCAATGCGAAACGTAG
CG4565	CG4565	CGCAGCTGTGGTTCAGGTAG	ATGGACGAATCTGAAACAGC
CG5249	<i>Blimp-1</i>	ACACATAGGGCCTGTCTGTTT	TCCAATCTGGGACAGAATCC
CG7080	CG7080	CTTGTAGCAGCCCTCCACTC	CTTTGGTTAGCTGCCAGGAC
CG7080	CG7080	CTTGTAGCAGCCCTCCACTC	CTTTGGTTAGCTGCCAGGAC
CG7201	CG7201	CCTCCACTACGCCATAGAGC	AAGGTGTTAAGGCGATGCAG
CG7577	<i>ppk20</i>	GGCCTGGCAGATACTTACG	ATTCGGTCAAACCGACAGAG
CG7734	<i>shn</i>	TGTTTGAGGAGAGCGATGTG	ACGGCCAAGCATTACAAAAC
CG7897	<i>gp210</i>	ATCAACAATGGCGAAAAAGG	TTCGCCGAGGAAGTACAAAC
CG7906	CG7906	GGCACAAATGGACAGTGATG	ACTACGGAAGCCGCAACTAC
CG8773	CG8773	TTCGTCTGTACAAAAGCGATG	ACAAGTGAACCCGGCACTAC
CG8799	<i>l(2)03659</i>	AGCCAAACTAATGCCACCTG	CAGTCCGAAGAGCTGGGTAG
CG8945	CG8945	ATTTGGCCTTGAAGTTGTGC	TTGTGCTCTTGATCGTGGTC
CG9333	<i>Oseg5</i>	AGTGGCTTGATGGATATGTGC	CGCGAATCTGCAAGAAAAAC
CG9460	<i>Spn42De</i>	GCTGCCTGTAGAGGGACATC	CGTGTTCACTTCTGCTGCTC