

Table S2. KEGG categories by GeneSetAnalysis in *Drosophila* Schneider S2 cells exposed to C16-Yammamarin. Fold change was defined as the ratio of the normalized value of T2 to that of control (the levels of fold change are coded with shades of red, green and the lighter colors corresponding to more than 1.5, less than 1/1.5 and within 1.5-fold, respectively).

Map Category	Map Name	KEGG map	Number of Included Genes	Description	Symbol	KEGG entry	Fold Change (T2/Ctr)	Gene Name
Nucleotide Metabolism	Purine metabolism	dme00230	119	Adenylyl cyclase 76E	<i>Ac76E</i>	dme:Dmel_CG7978	1.256	1628231_at
				recombination-defective	<i>rec</i>	dme:Dmel_CG31293	1.251	1625964_at
				Adenylyl cyclase 35C	<i>Ac13E</i>	dme:Dmel_CG9210	1.181	1636219_at
				PAPS synthetase	<i>Paps</i>	dme:Dmel_CG8363	1.18	1634522_s_at
				discs large 1	<i>dlg1</i>	dme:Dmel_CG1725	1.104	1627882_at
				abnormal wing discs	<i>awd</i>	dme:Dmel_CG2210	1.088	1637980_at
				domino	<i>dom</i>	dme:Dmel_CG9696	1.061	1636034_at
				DNA polymerase alpha 50kD	<i>DNApol-alpha50</i>	dme:Dmel_CG7108	1.061	1633854_at
				Helicase at 25E	<i>Hel25E</i>	dme:Dmel_CG7269	1.037	1629620_s_at
				spaghetti squash	<i>sqh</i>	dme:Dmel_CG3595	1.035	1624065_at
				RNA polymerase III 128kD subunit	<i>Rpl1128</i>	dme:Dmel_CG8344	1.035	1628131_at
				Myosin light chain cytoplasmic	<i>Mlc-c</i>	dme:Dmel_CG3201	1.034	1631832_at
				rutabaga	<i>rut</i>	dme:Dmel_CG9533	1.032	1628594_at
				crinkled	<i>ck</i>	dme:Dmel_CG7595	1.031	1626841_s_at
				Dead box protein 73D	<i>Dbp73D</i>	dme:Dmel_CG9680	1.03	1639810_at
				DNA polymerase epsilon	<i>DNApol-epsilon</i>	dme:Dmel_CG6768	1.03	1626355_at
				Helicase 89B	<i>Hel89B</i>	dme:Dmel_CG4261	1.029	1627970_at
				DEAD box protein 45A	<i>Dbp45A</i>	dme:Dmel_CG12759	1.028	1625634_at
				okra	<i>okr</i>	dme:Dmel_CG3736	1.016	1628198_at
				discs large 1	<i>dlg1</i>	dme:Dmel_CG1725	1.014	1635382_at
				polychaetoid	<i>pvl</i>	dme:Dmel_CG31349	1.008	1637322_at
				RNA polymerase II 215kD subunit	<i>Rpl215</i>	dme:Dmel_CG1554	1.006	1640764_at
				cut up	<i>cutp</i>	dme:Dmel_CG6998	1.005	1634599_s_at
				TER94	<i>TER94</i>	dme:Dmel_CG2331	1.002	1628008_a_at
				maternal expression at 31B	<i>me31B</i>	dme:Dmel_CG4916	1.002	1628017_a_at
				Phosphodiesterase 6	<i>Pde6</i>	dme:Dmel_CG8279	1	1629322_at
				polychaetoid	<i>pvl</i>	dme:Dmel_CG31349	0.996	1637428_a_at
				Nucleoplasmmin	<i>Nlp</i>	dme:Dmel_CG7917	0.98	1635608_at
				cut up	<i>cutp</i>	dme:Dmel_CG6998	0.979	1623073_s_at
				Rm62	<i>Rm62</i>	dme:Dmel_CG10279	0.978	1623475_s_at
				DNA polymerase alpha 180kD	<i>DNApol-alpha180</i>	dme:Dmel_CG6349	0.975	1633249_at
				polychaetoid	<i>pvl</i>	dme:Dmel_CG31349	0.973	1630791_a_at
				Helicase	<i>Hlc</i>	dme:Dmel_CG1666	0.972	1638310_at
				lethal (1) G0007	<i>l(1)G0007</i>	dme:Dmel_CG32604	0.97	1623333_a_at
				smallminded	<i>smid</i>	dme:Dmel_CG8571	0.97	1624482_at
				adenosine 2	<i>ade2</i>	dme:Dmel_CG9127	0.969	1629194_s_at
				DEAD/DEAH RNA helicase 1	<i>Dhh1</i>	dme:Dmel_CG6539	0.968	1629281_at
				zipper	<i>zip</i>	dme:Dmel_CG15792	0.964	1627840_a_at
				cutlet	<i>cutlet</i>	dme:Dmel_CG33122	0.961	1627253_at
				RecQ4	<i>RecQ4</i>	dme:Dmel_CG7487	0.956	1623031_a_at
				RNA polymerase II 140kD subunit	<i>Rpl140</i>	dme:Dmel_CG3180	0.953	1625962_at
				Phosphoribosylamidotransferase	<i>Prat</i>	dme:Dmel_CG2867	0.948	1627868_at
				haywire	<i>hay</i>	dme:Dmel_CG8019	0.947	1641306_at
				Ribonucleoside diphosphate reductase large subunit	<i>RnrL</i>	dme:Dmel_CG5371	0.947	1631007_at
				Ribonucleoside diphosphate reductase small subunit	<i>RnrS</i>	dme:Dmel_CG8975	0.946	1635409_at
				Mi-2	<i>Mi-2</i>	dme:Dmel_CG8103	0.943	1640441_at
				RIC3	<i>RIC3</i>	dme:Dmel_CG5313	0.941	1637166_at
				brahma	<i>brm</i>	dme:Dmel_CG5942	0.94	1636831_s_at
				kismet	<i>kis</i>	dme:Dmel_CG3696	0.939	1636513_a_at
				Origin recognition complex subunit 5	<i>Orc5</i>	dme:Dmel_CG7833	0.938	1633660_at
				Chromodomain-helicase-DNA-binding protein	<i>Chd1</i>	dme:Dmel_CG3733	0.938	1639816_at
				no mitochondrial derivative	<i>nmd</i>	dme:Dmel_CG5395	0.932	1635448_at
				Dead-box-1	<i>Dxb1</i>	dme:Dmel_CG9054	0.932	1639264_at
				Adenylate kinase-2	<i>Ak2</i>	dme:Dmel_CG3140	0.93	1637064_at
				diadenosine tetraphosphate hydrolase	<i>Apf</i>	dme:Dmel_CG31713	0.929	1635473_at
				spasmin	<i>spas</i>	dme:Dmel_CG5977	0.927	1628019_a_at
				DNA polymerase alpha 73kD	<i>DNApol-alpha73</i>	dme:Dmel_CG5923	0.924	1628372_a_at
				Pyruvate kinase	<i>Pyk</i>	dme:Dmel_CG7070	0.921	1628027_a_at
				Imitation SWI	<i>Iswi</i>	dme:Dmel_CG8625	0.92	1641309_s_at
				twister	<i>twi</i>	dme:Dmel_CG10210	0.92	1624956_at
				Proteasome 26S subunit subunit 4 ATPase	<i>Pros26.4</i>	dme:Dmel_CG5289	0.918	1630788_at
				Myo28B1	<i>Myo28B1</i>	dme:Dmel_CG6976	0.915	1629784_a_at
				burgandy	<i>bur</i>	dme:Dmel_CG9242	0.911	1635718_at
				Adenosine deaminase-related growth factor A	<i>AdgfA</i>	dme:Dmel_CG5992	0.911	1624054_at
				Adenylate kinase-3	<i>Ak3</i>	dme:Dmel_CG6612	0.91	1630986_s_at
				XNP	<i>XNP</i>	dme:Dmel_CG4548	0.909	1630882_s_at
				Magi	<i>Magi</i>	dme:Dmel_CG30388	0.906	1638176_at
				Adenine phosphoribosyltransferase	<i>Aprt</i>	dme:Dmel_CG18315	0.906	1628492_at
				dunce	<i>dnc</i>	dme:Dmel_CG32498	0.905	1624686_a_at
				mutagen-sensitive 301	<i>msu301</i>	dme:Dmel_CG7972	0.894	1638294_at
				RNA polymerase II 18kD subunit	<i>Rpl18</i>	dme:Dmel_CG1163	0.891	1636644_at
				RNA polymerase I 135kD subunit	<i>Rpl135</i>	dme:Dmel_CG4033	0.891	1627924_at
				domino	<i>dom</i>	dme:Dmel_CG9696	0.889	1633331_at
				stardust	<i>std</i>	dme:Dmel_CG32717	0.889	1631568_a_at
				homolog of RecQ	<i>RecQ5</i>	dme:Dmel_CG4879	0.888	1628264_a_at
				RNA polymerase I subunit	<i>Rpl1</i>	dme:Dmel_CG10122	0.887	1629356_at
				Germ line transcription factor 1	<i>Gnf1</i>	dme:Dmel_CG1119	0.886	1623086_at
				Marcoal1	<i>Marcoal1</i>	dme:Dmel_CG3753	0.88	1626531_at
				spindle B	<i>spn-B</i>	dme:Dmel_CG3325	0.879	1630018_at
				belie	<i>bel</i>	dme:Dmel_CG9748	0.878	1640256_at
				Myosin heavy chain-like	<i>Mhcl</i>	dme:Dmel_CG31045	0.878	1632231_a_at
				c12.2	<i>c12.2</i>	dme:Dmel_CG12149	0.876	1639233_at
				lethal (2) 35Df	<i>l(2)35Df</i>	dme:Dmel_CG4152	0.875	1636986_at
				lodestar	<i>lds</i>	dme:Dmel_CG2684	0.873	1634695_at
				DNA-polymerase-delta	<i>DNApol-delta</i>	dme:Dmel_CG5949	0.872	1641438_at
				maleless	<i>mle</i>	dme:Dmel_CG11680	0.872	1629507_a_at
				discs large 1	<i>dlg1</i>	dme:Dmel_CG1725	0.869	1624021_a_at

				domino	<i>dom</i>	dme:Dmel CG9696	0.869	1628160 a at
				Xeroderma pigmentosum D	<i>Xpd</i>	dme:Dmel CG9433	0.869	1629170 s at
				dilute class unconventional myosin	<i>dicdm</i>	dme:Dmel CG2146	0.865	1635040 a at
				spindle A	<i>spn-A</i>	dme:Dmel CG7948	0.859	1640738 a at
				Lap1	<i>Lap1</i>	dme:Dmel CG10255	0.859	1639189 at
				Myosin 95E	<i>Myo95E</i>	dme:Dmel CG5501	0.856	1631844 at
				Inverted repeat-binding protein	<i>Irbp</i>	dme:Dmel CG5247	0.854	1640431 at
				rasberry	<i>ras</i>	dme:Dmel CG1799	0.848	1627916 s at
				Ac3	<i>Ac3</i>	dme:Dmel CG1506	0.837	1625072 s at
				belphegor	<i>bor</i>	dme:Dmel CG6815	0.837	1628099 at
				Dier-2	<i>Dcr-2</i>	dme:Dmel CG6493	0.836	1641449 at
				roadblock	<i>robl</i>	dme:Dmel CG10751	0.833	1625632 at
				Dlic2	<i>Dlic2</i>	dme:Dmel CG1938	0.829	1638127 s at
				lethal (3) 70Da	<i>l(3)70Da</i>	dme:Dmel CG6760	0.817	1632434 at
				adenosine 3	<i>ade3</i>	dme:Dmel CG31628	0.816	1637037 at
				centrosomin	<i>cm</i>	dme:Dmel CG4832	0.806	1635619 a at
				NEM-sensitive fusion protein 2	<i>Nsf2</i>	dme:Dmel CG33101	0.8	1639023 at
				spastin	<i>spas</i>	dme:Dmel CG5977	0.8	1629123 at
				NTase	<i>NTase</i>	dme:Dmel CG3059	0.798	1630857 s at
				Ku80	<i>Ku80</i>	dme:Dmel CG18801	0.797	1625759 at
				Rsl	<i>Rsl</i>	dme:Dmel CG2173	0.796	1631198 at
				jaguar	<i>jar</i>	dme:Dmel CG5695	0.794	1623185 s at
				klarsicht	<i>klar</i>	dme:Dmel CG17046	0.79	1628155 at
				Adenosine deaminase	<i>Ada</i>	dme:Dmel CG11994	0.749	1637806 at
				ABC transporter expressed in trachea	<i>Atet</i>	dme:Dmel CG2969	0.722	1629559 s at
				Sulfonyleurea receptor	<i>Sur</i>	dme:Dmel CG5772	0.719	1632337 at
				mutagen-sensitive 304	<i>ms304</i>	dme:Dmel CG7347	0.705	1638933 at
				Rad51C	<i>Rad51C</i>	dme:Dmel CG2412	0.697	1629547 at
				spindle E	<i>spn-E</i>	dme:Dmel CG3158	0.69	1636886 at
				Adenyl cyclase 78C	<i>Ac78C</i>	dme:Dmel CG10564	0.689	1628412 at
				kurz	<i>kz</i>	dme:Dmel CG3228	0.662	1623496 at
				Adenylate kinase-1	<i>Akl1</i>	dme:Dmel CG17146	0.627	1631377 a at
				Ribosomal protein L27	<i>Rpl27</i>	dme:Dmel CG4759	1.19	1632335 at
				Ribosomal protein L37A	<i>Rpl37A</i>	dme:Dmel CG5827	1.171	1639957 s at
				Ribosomal protein S20	<i>Rps20</i>	dme:Dmel CG15693	1.164	1641614 at
				Ribosomal protein S19a	<i>Rps19a</i>	dme:Dmel CG4464	1.162	1637448 s at
				Ribosomal protein L17	<i>Rpl17</i>	dme:Dmel CG3203	1.16	1628028 s at
				Ribosomal protein L35A	<i>Rpl35A</i>	dme:Dmel CG2099	1.153	1623995 at
				Ribosomal protein S27A	<i>Rps27A</i>	dme:Dmel CG5271	1.152	1624906 at
				Ribosomal protein S3	<i>Rps3</i>	dme:Dmel CG6779	1.152	1632269 at
				Ribosomal protein S12	<i>Rps12</i>	dme:Dmel CG11271	1.141	1627364 s at
				Ribosomal protein S18	<i>Rps18</i>	dme:Dmel CG8900	1.14	1624983 s at
				Ribosomal protein L14	<i>Rpl14</i>	dme:Dmel CG6253	1.139	1623914 at
				Ribosomal protein L29	<i>Rpl29</i>	dme:Dmel CG10071	1.139	1631819 at
				Ribosomal protein L21	<i>Rpl21</i>	dme:Dmel CG12775	1.137	1636715 at
				Ribosomal protein S25	<i>Rps25</i>	dme:Dmel CG6684	1.137	1627774 a at
				Ribosomal protein S28b	<i>Rps28b</i>	dme:Dmel CG2998	1.136	1624792 at
				Ribosomal protein S13	<i>Rps13</i>	dme:Dmel CG13389	1.134	1635303 at
				Ribosomal protein L35	<i>Rpl35</i>	dme:Dmel CG4111	1.134	1626209 a at
				Ribosomal protein S4	<i>Rps4</i>	dme:Dmel CG11276	1.132	1639949 s at
				stubarita	<i>sta</i>	dme:Dmel CG14792	1.13	1638847 s at
				Ribosomal protein S3A	<i>Rps3A</i>	dme:Dmel CG2168	1.129	1638439 a at
				Ribosomal protein L.P2	<i>RplP2</i>	dme:Dmel CG4918	1.129	1623150 at
				Ribosomal protein L.P0	<i>RplP0</i>	dme:Dmel CG7490	1.128	1637546 at
				Ribosomal protein L4	<i>Rpl4</i>	dme:Dmel CG5502	1.128	1627988 at
				Ribosomal protein L37a	<i>Rpl37a</i>	dme:Dmel CG9091	1.126	1628061 at
				Ribosomal protein S17	<i>Rps17</i>	dme:Dmel CG3922	1.121	1624529 at
				Ribosomal protein L7	<i>Rpl7</i>	dme:Dmel CG4897	1.121	1629107 at
				Ribosomal protein L13	<i>Rpl13</i>	dme:Dmel CG4651	1.119	1623836 at
				Ribosomal protein L27A	<i>Rpl27A</i>	dme:Dmel CG15442	1.118	1639377 at
				Ribosomal protein S27	<i>Rps27</i>	dme:Dmel CG10423	1.117	1631296 at
				Ribosomal protein L22	<i>Rpl22</i>	dme:Dmel CG7434	1.115	1639312 at
				Ribosomal protein L.P1	<i>RplP1</i>	dme:Dmel CG4087	1.115	1623606 at
				Ribosomal protein L6	<i>Rpl6</i>	dme:Dmel CG11522	1.114	1641355 a at
				Ribosomal protein L18A	<i>Rpl18A</i>	dme:Dmel CG6510	1.112	1636056 at
				Ribosomal protein S8	<i>Rps8</i>	dme:Dmel CG7808	1.112	1626931 a at
				Ribosomal protein L36A	<i>Rpl36A</i>	dme:Dmel CG7424	1.11	1623262 at
				string of pearls	<i>sop</i>	dme:Dmel CG5920	1.108	1632483 at
				Ribosomal protein L9	<i>Rpl9</i>	dme:Dmel CG6141	1.107	1630634 s at
				Ribosomal protein L31	<i>Rpl31</i>	dme:Dmel CG1821	1.107	1628928 s at
				Ribosomal protein S16	<i>Rps16</i>	dme:Dmel CG4046	1.106	1639270 at
				Ribosomal protein L19	<i>Rpl19</i>	dme:Dmel CG2746	1.106	1630455 at
				Ribosomal protein L32	<i>Rpl32</i>	dme:Dmel CG7939	1.105	1625337 s at
				Ribosomal protein L8	<i>Rpl8</i>	dme:Dmel CG1263	1.105	1628775 s at
				Ribosomal protein S30	<i>Rps30</i>	dme:Dmel CG15697	1.104	1623997 s at
				Ribosomal protein S6	<i>Rps6</i>	dme:Dmel CG10944	1.103	1626924 a at
				Ribosomal protein L40	<i>Rpl40</i>	dme:Dmel CG2960	1.102	1633277 at
				Ribosomal protein S26	<i>Rps26</i>	dme:Dmel CG10305	1.102	1632220 s at
				overgrown hematopoietic organs at 23B	<i>oho23B</i>	dme:Dmel CG2986	1.101	1623644 s at
				Ribosomal protein L13A	<i>Rpl13A</i>	dme:Dmel CG1475	1.101	1631759 at
				Ribosomal protein S23	<i>Rps23</i>	dme:Dmel CG8415	1.099	1638832 at
				Ribosomal protein L35	<i>Rpl35</i>	dme:Dmel CG4111	1.097	1627206 at
				Ribosomal protein L18	<i>Rpl18</i>	dme:Dmel CG8615	1.096	1629400 at
				Ribosomal protein L36	<i>Rpl36</i>	dme:Dmel CG7622	1.095	1624019 s at
				Ribosomal protein S5a	<i>Rps5a</i>	dme:Dmel CG8922	1.095	1630031 at
				Ribosomal protein L23	<i>Rpl23</i>	dme:Dmel CG3661	1.093	1634764 at
				Ribosomal protein S10b	<i>Rps10b</i>	dme:Dmel CG14206	1.088	1635528 s at
				Ribosomal protein S14b	<i>Rps14b</i>	dme:Dmel CG1527	1.086	1624337 at
				Ribosomal protein L11	<i>Rpl11</i>	dme:Dmel CG7726	1.086	1630813 at
				Ribosomal protein L26	<i>Rpl26</i>	dme:Dmel CG6846	1.086	1628603 at
				Ribosomal protein L34a	<i>Rpl34a</i>	dme:Dmel CG6090	1.084	1641721 a at
				Ribosomal protein L12	<i>Rpl12</i>	dme:Dmel CG3195	1.082	1640446 s at
				Ribosomal protein L24	<i>Rpl24</i>	dme:Dmel CG9282	1.073	1641600 at
				Ribosomal protein S24	<i>Rps24</i>	dme:Dmel CG3751	1.073	1630093 at
				Ribosomal protein L7A	<i>Rpl7A</i>	dme:Dmel CG3314	1.072	1639893 s at
				Ribosomal protein L30	<i>Rpl30</i>	dme:Dmel CG10652	1.068	1634313 s at
				Ribosomal protein L39	<i>Rpl39</i>	dme:Dmel CG3997	1.065	1626964 at
				Ribosomal protein L23A	<i>Rpl23A</i>	dme:Dmel CG7977	1.054	1634917 at
				Ribosomal protein L12	<i>Rpl12</i>	dme:Dmel CG3195	0.962	1627312 at
				Ribosomal protein L35	<i>Rpl35</i>	dme:Dmel CG4111	0.948	1635917 at
				Ribosomal protein S30	<i>Rps30</i>	dme:Dmel CG15697	0.864	1625943 at
				Glycerol 3 phosphate dehydrogenase	<i>Gpdh</i>	dme:Dmel CG9042	1.148	1634893 at
				Glycerol 3 phosphate dehydrogenase	<i>Gpdh</i>	dme:Dmel CG9042	1.098	1636311 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	1.097	1631833 at
				retinal degeneration A	<i>rdgA</i>	dme:Dmel CG10966	1.06	1641349 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	1.059	1638829 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	1.032	1636471 at
Lipid Metabolism	Glycerophospholipid metabolism	dme00564	46					

				CTP:phosphocholine cytidylyltransferase 1	<i>Cct1</i>	dme:Dmel_CG1049	1.026	1629930_s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	1.006	1628932_at
				Glycerol 3 phosphate dehydrogenase	<i>Gpdh</i>	dme:Dmel_CG9042	0.982	1616608_a_at
				Phosphoethanolamine cytidylyltransferase	<i>Pect</i>	dme:Dmel_CG5547	0.979	1641183_a_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.965	1634258_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.96	1624020_at
				CTP:phosphocholine cytidylyltransferase 2	<i>Cct2</i>	dme:Dmel_CG18330	0.958	1628589_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.952	1632973_at
				Glycerol 3 phosphate dehydrogenase	<i>Gpdh</i>	dme:Dmel_CG9042	0.933	1625949_at
				Spingosine kinase 2	<i>Sk2</i>	dme:Dmel_CG32484	0.926	1624594_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.92	1628921_s_at
				GXIVsPLA2	<i>GXIVsPLA2</i>	dme:Dmel_CG17035	0.917	1625027_a_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.904	1637723_at
				Diacyl glycerol kinase epsilon	<i>Dgkepsilon</i>	dme:Dmel_CG8657	0.902	1626289_at
				GDP dissociation inhibitor	<i>Gdi</i>	dme:Dmel_CG4422	0.899	1641129_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.888	1626650_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.888	1628562_s_at
				wunen-2	<i>wun2</i>	dme:Dmel_CG8805	0.872	1641598_at
				wunen	<i>wun</i>	dme:Dmel_CG8804	0.857	1631573_a_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.854	1631742_s_at
				Rab escort protein	<i>Rep</i>	dme:Dmel_CG8432	0.847	1631388_at
				CDP diglyceride synthetase	<i>Cds1</i>	dme:Dmel_CG7962	0.845	1627711_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.843	1640268_at
				Dihydroxyacetone phosphate acyltransferase	<i>Dhap-at</i>	dme:Dmel_CG4625	0.825	1632807_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.822	1639308_at
				tafazzin	<i>tafazzin</i>	dme:Dmel_CG8766	0.812	1626481_a_at
				rasp	<i>rasp</i>	dme:Dmel_CG11495	0.81	1638008_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.81	1630536_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.809	1640554_at
				Phospholipase D	<i>Pld</i>	dme:Dmel_CG12110	0.806	1627315_s_at
				easily shocked	<i>ees</i>	dme:Dmel_CG3525	0.792	1640655_s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.772	1631500_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.761	1636765_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.745	1627953_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.74	1622976_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.731	1625283_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.715	1634052_s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.703	1633152_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel_CG32491	0.695	1638041_at
				lethal (2) k05713	<i>l(2)k05713</i>	dme:Dmel_CG8256	0.556	1631474_s_at
				Sin3A	<i>Sin3A</i>	dme:Dmel_CG8815	1.062	1630165_s_at
				ATP synthase-gamma chain	<i>ATPsyn-gamma</i>	dme:Dmel_CG7610	1.049	1627018_s_at
				ATP synthase-beta	<i>ATPsyn-beta</i>	dme:Dmel_CG11154	1.044	1633673_a_at
				Vha100-2	<i>Vha100-2</i>	dme:Dmel_CG18617	1.035	1631359_s_at
				RhoGAP71E	<i>RhoGAP71E</i>	dme:Dmel_CG32149	1.029	1637051_at
				bellwether	<i>bw</i>	dme:Dmel_CG3612	1.025	1636548_at
				NADH:ubiquinone reductase 42kD subunit precursor	<i>ND42</i>	dme:Dmel_CG6343	-1.025	1639585_a_at
				ATP synthase-gamma chain	<i>ATPsyn-gamma</i>	dme:Dmel_CG7610	0.999	1632838_at
				lethal (2) 06225	<i>l(2)06225</i>	dme:Dmel_CG6105	0.972	1629647_at
				Vacuolar H[+] ATPase G-subunit	<i>Vha13</i>	dme:Dmel_CG6213	0.968	1627006_at
				Vacuolar H[+] ATPase 26kD E subunit	<i>Vha26</i>	dme:Dmel_CG1088	0.967	1629065_s_at
				Vacuolar H[+] ATPase 16kD subunit	<i>Vha16</i>	dme:Dmel_CG3161	0.957	1632117_s_at
				Cytochrome c oxidase subunit Va	<i>Co1a</i>	dme:Dmel_CG14724	0.945	1639962_a_at
				toutais	<i>tau</i>	dme:Dmel_CG10897	0.944	1640433_a_at
				lethal (3) neo18	<i>l(3)neo18</i>	dme:Dmel_CG9762	0.943	1633698_at
				lethal (1) G0230	<i>l(1)G0230</i>	dme:Dmel_CG2968	0.937	1637737_at
				Nucleosome remodeling factor - 38kD	<i>Nurf-38</i>	dme:Dmel_CG4634	0.932	1637982_at
				Oligomycin sensitivity-conferring protein	<i>Oscp</i>	dme:Dmel_CG4307	0.932	1631750_a_at
				ATP synthase, subunit b	<i>ATPsyn-b</i>	dme:Dmel_CG8189	0.931	1628573_a_at
				ATP synthase-beta	<i>ATPsyn-beta</i>	dme:Dmel_CG11154	0.927	1630984_at
				RhoGAP71E	<i>RhoGAP71E</i>	dme:Dmel_CG32149	0.924	1640295_a_at
				ATPase coupling factor 6	<i>ATPsyn-Cf6</i>	dme:Dmel_CG4412	0.92	1641328_at
				Sin3A	<i>Sin3A</i>	dme:Dmel_CG8815	0.919	1628903_s_at
				mitochondrial acyl carrier protein 1	<i>mtacp1</i>	dme:Dmel_CG9160	0.916	1634955_at
				oxen	<i>ox</i>	dme:Dmel_CG8764	0.913	1633973_at
				mitochondrial acyl carrier protein 1	<i>mtacp1</i>	dme:Dmel_CG9160	0.906	1624776_a_at
				Vacuolar H[+] ATPase 55kD B subunit	<i>Vha55</i>	dme:Dmel_CG17369	0.903	1625131_s_at
				bloated tubules	<i>blot</i>	dme:Dmel_CG3897	0.901	16266839_s_at
				lethal (2) 351d	<i>l(2)351d</i>	dme:Dmel_CG13240	0.899	1631318_at
				mitochondrial acyl carrier protein 1	<i>mtacp1</i>	dme:Dmel_CG9160	0.897	1624715_at
				lalafel	<i>llf</i>	dme:Dmel_CG9351	0.887	1637302_s_at
				Vha100-1	<i>Vha100-1</i>	dme:Dmel_CG1709	0.887	1632395_s_at
				lethal (2) 351d	<i>l(2)351d</i>	dme:Dmel_CG13240	0.886	1630062_a_at
				Vacuolar H[+] ATPase 44kD C subunit	<i>Vha44</i>	dme:Dmel_CG8048	0.885	1635352_s_at
				Vacuolar H[+] ATPase SFD subunit	<i>VhaSFD</i>	dme:Dmel_CG17332	0.882	1633931_a_at
				Rieske iron-sulfur protein	<i>RFeSP</i>	dme:Dmel_CG7361	0.875	1631856_a_at
				VhaM9.7-2	<i>VhaM9.7-2</i>	dme:Dmel_CG7625	0.872	1623653_at
				stunted	<i>sun</i>	dme:Dmel_CG9032	0.861	1624845_a_at
				Succinate dehydrogenase B	<i>SdhB</i>	dme:Dmel_CG3283	0.86	1640632_at
				Vacuolar H[+] ATPase 14kD subunit	<i>Vha14</i>	dme:Dmel_CG8210	0.856	1639788_at
				cyclope	<i>cype</i>	dme:Dmel_CG14028	0.855	1638416_at
				NADH:ubiquinone reductase 75kD subunit precursor	<i>ND75</i>	dme:Dmel_CG2286	0.822	1638593_a_at
				NADH:ubiquinone reductase 23kD subunit precursor	<i>ND23</i>	dme:Dmel_CG3944	0.813	1627220_at
				Succinyl coenzyme A synthetase flavoprotein subunit	<i>Scs-fp</i>	dme:Dmel_CG17246	0.791	1629086_s_at
				VhaM9.7-1	<i>VhaM9.7-1</i>	dme:Dmel_CG11589	0.76	1639067_at
				jumeau	<i>jumu</i>	dme:Dmel_CG4029	0.757	1631242_at
				nejire	<i>nej</i>	dme:Dmel_CG15319	1.03	1622925_at
				porcupine	<i>por</i>	dme:Dmel_CG6205	0.976	1636527_at
				Phospholipase C at 21C	<i>Plc21C</i>	dme:Dmel_CG4574	0.976	1630007_s_at
				basket	<i>bsk</i>	dme:Dmel_CG5680	0.97	1626404_a_at
				basket	<i>bsk</i>	dme:Dmel_CG5680	0.969	1634580_at
				Smad on X	<i>Smox</i>	dme:Dmel_CG2262	0.963	1629290_at
				armadillo	<i>arm</i>	dme:Dmel_CG11579	0.955	1629020_s_at
				frizzled 2	<i>fr2</i>	dme:Dmel_CG9739	0.95	1624219_s_at
				nemo	<i>nmo</i>	dme:Dmel_CG7892	0.945	1624300_s_at
				cAMP-dependent protein kinase I	<i>Pka-C1</i>	dme:Dmel_CG4379	0.943	1631055_at
				Protein phosphatase 2A at 29B	<i>Pp2A-29B</i>	dme:Dmel_CG17291	0.942	1625851_s_at
				shaggy	<i>sgg</i>	dme:Dmel_CG2621	0.94	1630774_s_at
				lin-19-like	<i>lin19</i>	dme:Dmel_CG1877	0.935	1624407_s_at
				no receptor potential A	<i>norpA</i>	dme:Dmel_CG3620	0.929	1636576_s_at

				Casein kinase II alpha subunit	<i>Ckl1alpha</i>	dme:Dmel	CG17520	0.929	1640051 s at
				arrow	<i>arr</i>	dme:Dmel	CG5912	0.917	1634211 at
				Calcium/calmodulin-dependent protein kinase II	<i>CaMKII</i>	dme:Dmel	CG18069	0.915	1640534 s at
				microtubule star	<i>mts</i>	dme:Dmel	CG7109	0.909	1640841 at
				APC-like	<i>Apc</i>	dme:Dmel	CG1451	0.902	1628548 at
				groucho	<i>gro</i>	dme:Dmel	CG8384	0.897	1636141 s at
				frizzled 2	<i>frz</i>	dme:Dmel	CG9739	0.892	1635583 s at
				basket	<i>bsk</i>	dme:Dmel	CG5680	0.892	1625533 at
				Rho-kinase	<i>rok</i>	dme:Dmel	CG9774	0.891	1626306 at
				Van Gogh	<i>Vang</i>	dme:Dmel	CG8075	0.887	1635743 at
				Presenilin	<i>Psn</i>	dme:Dmel	CG18803	0.886	1624165 a at
				point	<i>point</i>	dme:Dmel	CG4003	0.861	1635279 at
				cAMP-dependent protein kinase I	<i>Pka-C1</i>	dme:Dmel	CG4379	0.857	1632409 a at
				Calcineurin B2	<i>CanB2</i>	dme:Dmel	CG11217	0.851	1627067 at
				naked cuticle	<i>nkd</i>	dme:Dmel	CG11614	0.851	1630361 at
				Protein phosphatase 2B at 14D	<i>Pp2B-14D</i>	dme:Dmel	CG9842	0.844	1626781 at
				Cyclin D	<i>CycD</i>	dme:Dmel	CG9096	0.807	1627295 s at
				TGF-beta activated kinase 1	<i>Tak1</i>	dme:Dmel	CG18492	0.802	1636514 at
				supernumerary limbs	<i>slmb</i>	dme:Dmel	CG3412	0.774	1635960 at
				division abnormally delayed	<i>dally</i>	dme:Dmel	CG4974	0.771	1628489 at
				Calcineurin A at 14F	<i>CanA-14F</i>	dme:Dmel	CG9819	0.706	1623931 s at
				no receptor potential A	<i>norpA</i>	dme:Dmel	CG3620	0.697	1630119 s at
				dishvelled	<i>dsh</i>	dme:Dmel	CG18361	0.661	1628788 a at
				cAMP-dependent protein kinase	<i>Pka-C3</i>	dme:Dmel	CG6117	0.587	1635928 a at
				Type III alcohol dehydrogenase	<i>T3dh</i>	dme:Dmel	CG3425	1.19	1628907 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.097	1631833 at
				retinal degeneration A	<i>rdgA</i>	dme:Dmel	CG10966	1.06	1641349 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.059	1638829 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.032	1636471 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.006	1628932 at
				Formaldehyde dehydrogenase	<i>Fdh</i>	dme:Dmel	CG6598	0.996	1636740 at
				Aldehyde dehydrogenase	<i>Aldh</i>	dme:Dmel	CG3752	0.995	1634096 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.965	1634258 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.96	1624020 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.952	1632973 at
				Sphingosine kinase 2	<i>Sk2</i>	dme:Dmel	CG32484	0.926	1624594 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.92	1628921 s at
				Ect3	<i>Ect3</i>	dme:Dmel	CG1132	0.909	1624517 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.904	1637723 at
				Diacyl glycerol kinase epsilon	<i>Dgkepsilon</i>	dme:Dmel	CG8657	0.902	1626289 at
				nriyu	<i>nriyu</i>	dme:Dmel	CG1216	0.896	1628487 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1626650 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1628562 s at
				UDP-glucose-glycoprotein glycosyltransferase	<i>Ugt</i>	dme:Dmel	CG6850	0.887	1624465 at
				wunen-2	<i>wun2</i>	dme:Dmel	CG8805	0.872	1641598 at
				Glycerol kinase	<i>Gyk</i>	dme:Dmel	CG18374	0.862	1638285 a at
				wunen	<i>wun</i>	dme:Dmel	CG8804	0.857	1631573 a at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.854	1631742 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.843	1640268 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.822	1639308 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.81	1630536 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.809	1640554 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.772	1631500 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.761	1636765 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.745	1627953 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.74	1622976 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.731	1625283 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.715	1634052 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.703	1633152 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.695	1638041 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.097	1631833 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.059	1638829 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.032	1636471 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.006	1628932 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.965	1634258 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.96	1624020 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.952	1632973 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.92	1628921 s at
				Ceramidase	<i>C'Dase</i>	dme:Dmel	CG1471	0.91	1639396 s at
				Ect3	<i>Ect3</i>	dme:Dmel	CG1132	0.909	1624517 at
				Sphingosine-1-phosphate lyase	<i>Sply</i>	dme:Dmel	CG8946	0.908	1624818 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.904	1637723 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1626650 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1628562 s at
				wunen-2	<i>wun2</i>	dme:Dmel	CG8805	0.872	1641598 at
				wunen	<i>wun</i>	dme:Dmel	CG8804	0.857	1631573 a at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.854	1631742 s at
				Serine palmitoyltransferase subunit I	<i>Spt-1</i>	dme:Dmel	CG4016	0.853	1625352 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.843	1640268 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.822	1639308 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.81	1630536 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.809	1640554 at
				infertile crescent	<i>ifc</i>	dme:Dmel	CG9078	0.809	1625414 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.772	1631500 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.761	1636765 at
				brain washing	<i>bwa</i>	dme:Dmel	CG13969	0.761	1625491 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.745	1627953 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.74	1622976 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.731	1625283 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.715	1634052 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.703	1633152 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.695	1638041 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.097	1631833 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.059	1638829 s at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.032	1636471 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	1.006	1628932 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.965	1634258 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.96	1624020 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.952	1632973 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.92	1628921 s at
				GXIVsPLA2	<i>GXIVsPLA2</i>	dme:Dmel	CG17035	0.917	1625027 a at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.904	1637723 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1626650 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel	CG32491	0.888	1628562 s at
				wunen-2	<i>wun2</i>	dme:Dmel	CG8805	0.872	1641598 at
				Platelet-activating factor acetylhydrolase alpha	<i>Paf-AHalpha</i>	dme:Dmel	CG8962	0.871	1627403 s at
				wunen	<i>wun</i>	dme:Dmel	CG8804	0.857	1631573 a at

				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.854	1631742 s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.843	1640268 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.822	1639308 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.81	1630536 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.809	1640554 at
				Phospholipase D	<i>Pld</i>	dme:Dmel CG12110	0.806	1627315 s_at
				Lissencephaly-1	<i>Lis-1</i>	dme:Dmel CG8440	0.78	1624854 s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.772	1631500 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.761	1636765 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.745	1627953 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.74	1622976 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.731	1625283 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.715	1634052 s_at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.703	163152 at
				modifier of mdg4	<i>mod(mdg4)</i>	dme:Dmel CG32491	0.695	1638041 at
Xenobiotics Biodegradation and Metabolism	Metabolism of xenobiotics by cytochrome P450	dme00980	27	Glutathione S transferase S1	<i>GstS1</i>	dme:Dmel CG8938	1.727	1623957 s_at
				Type III alcohol dehydrogenase	<i>T3dh</i>	dme:Dmel CG3425	1.19	1628907 at
				Glutathione S transferase D1	<i>GstD1</i>	dme:Dmel CG10045	1.151	1628353 at
				GST-containing FLYWCH zinc-finger protein	<i>gdf</i>	dme:Dmel CG33546	1.037	1640913 at
				UDP-glycosyltransferase 35a	<i>Ugt35a</i>	dme:Dmel CG6644	1.025	1627662 at
				Glutathione S transferase D3	<i>GstD3</i>	dme:Dmel CG4381	1.02	1635701 at
				Juvenile hormone epoxide hydrolase 2	<i>Jheh2</i>	dme:Dmel CG15102	1.007	1627773 a_at
				Formaldehyde dehydrogenase	<i>Fdh</i>	dme:Dmel CG6598	0.996	1636740 at
				Glutathione S transferase D7	<i>GstD7</i>	dme:Dmel CG4371	0.974	1624793 at
				GST-containing FLYWCH zinc-finger protein	<i>gdf</i>	dme:Dmel CG33546	0.933	1625757 at
				Microsomal glutathione S-transferase-like	<i>Mgstl</i>	dme:Dmel CG1742	0.928	1631058 s_at
				Glutathione S transferase E6	<i>GstE6</i>	dme:Dmel CG17530	0.856	1625744 at
				GleAT-P	<i>GleAT-P</i>	dme:Dmel CG6207	0.834	1624164 s_at
				Ugt58Fa	<i>Ugt58Fa</i>	dme:Dmel CG4414	0.824	1632070 at
				Glutathione S transferase E2	<i>GstE2</i>	dme:Dmel CG17523	0.816	1641429 at
				Glutathione S transferase E3	<i>GstE3</i>	dme:Dmel CG17524	0.808	1637129 at
				GleAT-S	<i>GleAT-S</i>	dme:Dmel CG3881	0.797	1628402 a_at
				Glutathione S transferase D9	<i>GstD9</i>	dme:Dmel CG10091	0.785	1636174 at
				Aldehyde dehydrogenase type III	<i>Aldh-III</i>	dme:Dmel CG11140	0.756	1627861 s_at
				Glutathione S transferase D2	<i>GstD2</i>	dme:Dmel CG4181	0.75	1630258 at
				Glutathione S transferase D10	<i>GstD10</i>	dme:Dmel CG18548	0.741	1627890 at
				Glutathione S transferase D5	<i>GstD5</i>	dme:Dmel CG12242	0.736	1634152 at
				Glutathione S transferase E9	<i>GstE9</i>	dme:Dmel CG17534	0.704	1628657 at
				Glutathione S transferase E7	<i>GstE7</i>	dme:Dmel CG17531	0.673	1640065 at
				Ugt86D	<i>Ugt86D</i>	dme:Dmel CG6658	0.642	1641242 at
				Glutathione S transferase D4	<i>GstD4</i>	dme:Dmel CG11512	0.609	1626253 at
				Glutathione S transferase D6	<i>GstD6</i>	dme:Dmel CG4423	0.412	1626136 at
				Glutathione S transferase S1	<i>GstS1</i>	dme:Dmel CG8938	1.727	1623957 s_at
				Glutathione S transferase D1	<i>GstD1</i>	dme:Dmel CG10045	1.151	1628353 at
				SP1029	<i>SP1029</i>	dme:Dmel CG11956	1.148	1639232 s_at
				GST-containing FLYWCH zinc-finger protein	<i>gdf</i>	dme:Dmel CG33546	1.037	1640913 at
				Glutathione S transferase D3	<i>GstD3</i>	dme:Dmel CG4381	1.02	1635701 at
				slamdance	<i>sda</i>	dme:Dmel CG3518	0.996	1623503 at
				Glutathione S transferase D7	<i>GstD7</i>	dme:Dmel CG4371	0.974	1624793 at
				Isocitrate dehydrogenase	<i>Idh</i>	dme:Dmel CG7176	0.934	1634314 s_at
				GST-containing FLYWCH zinc-finger protein	<i>gdf</i>	dme:Dmel CG33546	0.933	1625757 at
				Microsomal glutathione S-transferase-like	<i>Mgstl</i>	dme:Dmel CG1742	0.928	1631058 s_at
				Glutathione Synthetase	<i>GS</i>	dme:Dmel CG6835	0.905	1627321 s_at
				Glutamate-cysteine ligase catalytic subunit	<i>Gclc</i>	dme:Dmel CG2259	0.895	1625362 at
				Glutathione S transferase E6	<i>GstE6</i>	dme:Dmel CG17530	0.856	1625744 at
wolfam syndrome 1	<i>wfs1</i>	dme:Dmel CG4917	0.841	1626657 a_at				
Glutathione S transferase E2	<i>GstE2</i>	dme:Dmel CG17523	0.816	1641429 at				
Glutathione S transferase E3	<i>GstE3</i>	dme:Dmel CG17524	0.808	1637129 at				
Glutathione S transferase D9	<i>GstD9</i>	dme:Dmel CG10091	0.785	1636174 at				
Zwischenferment	<i>Zw</i>	dme:Dmel CG12529	0.785	1625995 a_at				
Glutamate-cysteine ligase modifier subunit	<i>Gclm</i>	dme:Dmel CG4919	0.771	1640552 at				
Glutathione S transferase D2	<i>GstD2</i>	dme:Dmel CG4181	0.75	1630258 at				
Glutathione S transferase D10	<i>GstD10</i>	dme:Dmel CG18548	0.741	1627890 at				
Glutathione S transferase D5	<i>GstD5</i>	dme:Dmel CG12242	0.736	1634152 at				
Glutathione S transferase E9	<i>GstE9</i>	dme:Dmel CG17534	0.704	1628657 at				
Glutathione S transferase E7	<i>GstE7</i>	dme:Dmel CG17531	0.673	1640065 at				
Glutathione S transferase D4	<i>GstD4</i>	dme:Dmel CG11512	0.609	1626253 at				
Glutathione S transferase D6	<i>GstD6</i>	dme:Dmel CG4423	0.412	1626136 at				
abnormal wing discs	<i>awd</i>	dme:Dmel CG2210	1.088	1637980 at				
Nucleotide Metabolism	Pyrimidine metabolism	dme00240	26	DNA polymerase alpha 50kD	<i>DNApol-alpha50</i>	dme:Dmel CG7108	1.061	1633854 at
				RNA polymerase III 128kD subunit	<i>RplIII28</i>	dme:Dmel CG8344	1.035	1628131 at
				DNA polymerase epsilon	<i>DNApol-epsilon</i>	dme:Dmel CG6768	1.03	1626355 at
				RNA polymerase II 215kD subunit	<i>RplII215</i>	dme:Dmel CG1554	1.006	1640764 at
				DNA polymerase alpha 180kD	<i>DNApol-alpha180</i>	dme:Dmel CG6349	0.975	1633249 at
				Deoxyuridine triphosphatase	<i>dUTPase</i>	dme:Dmel CG4584	0.97	1634637 a_at
				RNA polymerase II 140kD subunit	<i>RplII140</i>	dme:Dmel CG3180	0.953	1625962 at
				Ribonucleoside diphosphate reductase large subunit	<i>RnrL</i>	dme:Dmel CG5371	0.947	1631007 at
				Ribonucleoside diphosphate reductase small subunit	<i>RnrS</i>	dme:Dmel CG8975	0.946	1635409 at
				pyd3	<i>pyd3</i>	dme:Dmel CG3027	0.945	1630038 at
				Dihydroorotate dehydrogenase	<i>Dhod</i>	dme:Dmel CG9741	0.943	1623792 a_at
				Rhythmically expressed gene 3	<i>Reg-3</i>	dme:Dmel CG2194	0.937	1625686 s_at
				diadenosine tetraphosphate hydrolase	<i>Apf</i>	dme:Dmel CG31713	0.929	1635473 at
				DNA polymerase alpha 73kD	<i>DNApol-alpha73</i>	dme:Dmel CG5923	0.924	1628372 a_at
				Thioredoxin reductase-1	<i>Trxr-1</i>	dme:Dmel CG2151	0.912	1641487 a_at
				rudimentary-like	<i>r-l</i>	dme:Dmel CG3593	0.895	1637799 at
				RNA polymerase II 18kD subunit	<i>Rpl18</i>	dme:Dmel CG1163	0.891	1636644 at
				RNA polymerase I 135kD subunit	<i>Rpl135</i>	dme:Dmel CG4033	0.891	1627924 at
				RNA polymerase I subunit	<i>Rpl1</i>	dme:Dmel CG10122	0.887	1629356 at
				DNA-polymerase-delta	<i>DNApol-delta</i>	dme:Dmel CG5949	0.872	1641438 at
				RluA-2	<i>RluA-2</i>	dme:Dmel CG6187	0.802	1632649 at
				lethal (2) k01209	<i>l(2)k01209</i>	dme:Dmel CG4798	0.776	1636958 s_at
				rudimentary	<i>r</i>	dme:Dmel CG18572	0.749	1641267 at
				Collapsin Response Mediator Protein	<i>CRMP</i>	dme:Dmel CG1411	0.737	1639109 a_at
				Collapsin Response Mediator Protein	<i>CRMP</i>	dme:Dmel CG1411	0.67	1635381 a_at
				Type III alcohol dehydrogenase	<i>T3dh</i>	dme:Dmel CG3425	1.19	1628907 at
				fructose-1,6-bisphosphatase	<i>fbp</i>	dme:Dmel CG31692	1.039	1641066 s_at
				Formaldehyde dehydrogenase	<i>Fdh</i>	dme:Dmel CG6598	0.996	1636740 at
				Aldehyde dehydrogenase	<i>Aldh</i>	dme:Dmel CG3752	0.995	1634096 at
				Aldolase	<i>Ald</i>	dme:Dmel CG6058	0.994	1633473 s_at
Acetyl Coenzyme A synthase	<i>AcCoAS</i>	dme:Dmel CG9390	0.981	1641148 a_at				
Phosphoglyceromutase	<i>Pgfy78</i>	dme:Dmel CG1721	0.975	1627227 at				

				Glyceraldehyde 3 phosphate dehydrogenase 2	<i>Gapdh2</i>	dme:Dmel CG8893	0.951	1627915 a at
				Enolase	<i>Eno</i>	dme:Dmel CG17654	0.941	1634065 s at
				Hexokinase A	<i>Hex-A</i>	dme:Dmel CG3001	0.935	162638 a at
				Phosphoglycerate kinase	<i>Pgk</i>	dme:Dmel CG3127	0.926	1639766 at
				lethal (2) 44DEa	<i>l(2)44DEa</i>	dme:Dmel CG8732	0.925	1629961 s at
				DDB1	<i>DDB1</i>	dme:Dmel CG7769	0.922	1633951 at
				Pyruvate kinase	<i>Pyk</i>	dme:Dmel CG7070	0.921	1628027 a at
				Glyceraldehyde 3 phosphate dehydrogenase 1	<i>Gapdh1</i>	dme:Dmel CG12055	0.894	1640907 at
				Triose phosphate isomerase	<i>Tpi</i>	dme:Dmel CG2171	0.863	1637816 s at
				Glyceraldehyde 3 phosphate dehydrogenase 2	<i>Gapdh2</i>	dme:Dmel CG8893	0.854	1632667 s at
				Aldolase	<i>Ald</i>	dme:Dmel CG6058	0.851	1631993 s at
				Acylphosphatase 2	<i>Acyp2</i>	dme:Dmel CG18505	0.85	1629363 at
				lethal (1) G0334	<i>l(1)G0334</i>	dme:Dmel CG7010	0.829	1635253 a at
				Phosphoglucose isomerase	<i>Pgi</i>	dme:Dmel CG8251	0.823	1638550 s at
				Phosphofructokinase	<i>Pfk</i>	dme:Dmel CG4001	0.797	1634739 a at
				Phosphogluconate mutase	<i>Pgm</i>	dme:Dmel CG5165	0.769	1633803 at
				Aldehyde dehydrogenase type III	<i>Aldh-III</i>	dme:Dmel CG11140	0.756	1627861 s at
				Ecdysone-inducible gene L3	<i>ImpL3</i>	dme:Dmel CG10160	0.735	1635227 at
				Inositol 1,4,5-triphosphate kinase 1	<i>IP3K1</i>	dme:Dmel CG4026	1.443	1634619 at
				Phosphatidylinositol 3 kinase 59F	<i>Pi3K59F</i>	dme:Dmel CG5373	1.166	1625678 at
				Phosphatidylinositol 3 kinase 68D	<i>Pi3K68D</i>	dme:Dmel CG11621	1.095	1641006 s at
				retinal degeneration A	<i>rdgA</i>	dme:Dmel CG10966	1.06	1641349 at
				Ipk2	<i>Ipk2</i>	dme:Dmel CG13688	1.008	1628514 at
				Calmodulin	<i>Cam</i>	dme:Dmel CG8472	1.005	1623682 a at
				Phospholipase C at 21C	<i>Ptc21C</i>	dme:Dmel CG4574	0.976	1630007 s at
				immune response deficient 5	<i>ird5</i>	dme:Dmel CG4201	0.957	1625974 at
				loki	<i>lok</i>	dme:Dmel CG10895	0.946	1625038 s at
				no receptor potential A	<i>norpA</i>	dme:Dmel CG3620	0.929	1636576 s at
				Sphingosine kinase 2	<i>Sk2</i>	dme:Dmel CG32484	0.926	1624594 at
				Cyclin-dependent kinase 4	<i>Cdk4</i>	dme:Dmel CG5072	0.925	1639347 s at
				Diacyl glycerol kinase epsilon	<i>Dgkepsilon</i>	dme:Dmel CG8657	0.902	1626289 at
				synaptotagmin	<i>synj</i>	dme:Dmel CG6562	0.887	1631427 s at
				Inositol 1,4,5-triphosphate kinase 2	<i>IP3K2</i>	dme:Dmel CG1630	0.882	1636354 s at
				skittles	<i>skit</i>	dme:Dmel CG9985	0.882	1624275 at
				Inositol polyphosphate 1-phosphatase	<i>lpp</i>	dme:Dmel CG3028	0.874	1626217 at
				small wing	<i>sl</i>	dme:Dmel CG4200	0.867	1628815 at
				Inositol 1,4,5-tris-phosphate receptor	<i>Itpr-r83A</i>	dme:Dmel CG1063	0.86	1639461 a at
				Pt4K1alpha	<i>Pt4K1alpha</i>	dme:Dmel CG2929	0.86	1627996 a at
				Phosphatidylinositol 3 kinase 68D	<i>Pi3K68D</i>	dme:Dmel CG11621	0.855	1625235 at
				CDP diglyceride synthetase	<i>CdsA</i>	dme:Dmel CG7962	0.845	1627711 at
				four wheel drive	<i>fwd</i>	dme:Dmel CG7004	0.839	1634509 s at
				Neurexin IV	<i>Nrx-IV</i>	dme:Dmel CG6827	0.771	1639459 a at
				no receptor potential A	<i>norpA</i>	dme:Dmel CG3620	0.697	1630119 s at
				Leucyl-tRNA synthetase	<i>Aats-leu</i>	dme:Dmel CG7479	1.01	1623460 at
				mitochondrial alanine-tRNA synthetase	<i>Aats-ala-m</i>	dme:Dmel CG4633	1	1636795 at
				mitochondrial ribosomal protein L39	<i>mRpl39</i>	dme:Dmel CG17166	0.976	1625665 at
				Asparaginyl-tRNA synthetase	<i>Aats-asn</i>	dme:Dmel CG10687	0.956	1624475 at
				Tyrosyl-tRNA synthetase	<i>Aats-tyr</i>	dme:Dmel CG4561	0.956	1632334 at
				Prolyl-tRNA synthetase	<i>Aats-pro</i>	dme:Dmel CG12186	0.943	1629758 at
				Tryptophanyl-tRNA synthetase	<i>Aats-try</i>	dme:Dmel CG9735	0.948	1629619 at
				Isoleucyl-tRNA synthetase	<i>Aats-ile</i>	dme:Dmel CG11471	0.942	1631106 s at
				Aspartyl-tRNA synthetase	<i>Aats-asp</i>	dme:Dmel CG3821	0.94	1629026 at
				Cysteinyl-tRNA synthetase	<i>Aats-cys</i>	dme:Dmel CG8431	0.939	1625878 at
				Seryl-tRNA synthetase	<i>Aats-ser</i>	dme:Dmel CG4938	0.934	1636797 at
				Glutamyl-prolyl-tRNA synthetase	<i>Aats-glp</i>	dme:Dmel CG5394	0.928	1626523 a at
				Lysoyl-tRNA synthetase	<i>Aats-lys</i>	dme:Dmel CG12141	0.921	1622945 at
				Threonyl-tRNA synthetase	<i>Aats-thr</i>	dme:Dmel CG5353	0.889	1640984 s at
				Tryptophanyl-tRNA synthetase	<i>Aats-try</i>	dme:Dmel CG9735	0.88	1638511 at
				Arginyl-tRNA synthetase	<i>Aats-arg</i>	dme:Dmel CG9020	0.875	1627148 at
				Alanyl-tRNA synthetase	<i>Aats-ala</i>	dme:Dmel CG13391	0.874	1639644 at
				Glycyl-tRNA synthetase	<i>Aats-gly</i>	dme:Dmel CG6778	0.857	1636948 a at
				Glutamyl-tRNA synthetase	<i>Aats-gln</i>	dme:Dmel CG10506	0.845	1640088 at
				Phenylalanyl-tRNA synthetase	<i>Aats-phe</i>	dme:Dmel CG13348	0.843	1636872 at
				Histidyl-tRNA synthetase	<i>Aats-his</i>	dme:Dmel CG6335	0.815	1631877 a at
				Valyl-tRNA synthetase	<i>Aats-val</i>	dme:Dmel CG4062	0.798	1638259 s at
				Methionyl-tRNA synthetase	<i>Aats-met</i>	dme:Dmel CG31322	0.795	1632235 at
				Tryptophanyl-tRNA synthetase	<i>Aats-try</i>	dme:Dmel CG9735	0.772	1635547 a at
				pole hole	<i>phl</i>	dme:Dmel CG2845	1.01	1627470 at
				downstream of receptor kinase	<i>drk</i>	dme:Dmel CG6033	0.989	1625475 s at
				broad	<i>br</i>	dme:Dmel CG11491	0.967	1636931 at
				capuccino	<i>capu</i>	dme:Dmel CG3399	0.964	1637398 a at
				rhubomb	<i>rho</i>	dme:Dmel CG1004	0.957	1635462 at
				Ras oncogene at 85D	<i>Ras85D</i>	dme:Dmel CG9375	0.955	1624899 at
				pointed	<i>pnt</i>	dme:Dmel CG17077	0.927	1630010 a at
				Son of sevenless	<i>Sos</i>	dme:Dmel CG7793	0.912	1633335 at
				egghead	<i>egh</i>	dme:Dmel CG9659	0.892	1634558 s at
				squid	<i>sqd</i>	dme:Dmel CG16901	0.887	1637716 a at
				female sterile (1) K10	<i>fs(1)K10</i>	dme:Dmel CG3218	0.87	1623086 at
				brainiac	<i>brn</i>	dme:Dmel CG4934	0.864	1625107 at
				squid	<i>sqd</i>	dme:Dmel CG16901	0.847	1628435 at
				encore	<i>enc</i>	dme:Dmel CG10847	0.844	1632178 at
				anterior open	<i>aop</i>	dme:Dmel CG3166	0.835	1627394 s at
				Downstream of raf1	<i>Dsorf1</i>	dme:Dmel CG15793	0.816	1626006 at
				egghead	<i>egh</i>	dme:Dmel CG9659	0.812	1631621 s at
				spire	<i>spir</i>	dme:Dmel CG10076	0.798	1631571 a at
				gurken	<i>grk</i>	dme:Dmel CG17610	0.761	1627656 at
				spire	<i>spir</i>	dme:Dmel CG10076	0.667	1635630 a at
				spire	<i>spir</i>	dme:Dmel CG10076	0.666	1637962 at
				broad	<i>br</i>	dme:Dmel CG11491	0.645	1635691 at
				TBP-associated factor 5	<i>Taf5</i>	dme:Dmel CG7704	1.048	1627331 at
				TBP-associated factor 6	<i>Taf6</i>	dme:Dmel CG32211	0.985	1626563 at
				TBP-associated factor 1	<i>Taf1</i>	dme:Dmel CG17603	0.983	1632256 at
				Transcription factor IIIFalpha	<i>TfIIIFalpha</i>	dme:Dmel CG10281	0.93	1640495 at
				Transcription-factor-IIA-S	<i>TfIIA-S</i>	dme:Dmel CG5163	0.917	1623370 at
				TBP-associated factor 13	<i>Taf13</i>	dme:Dmel CG10756	0.917	1628630 at
				TBP-associated factor 2	<i>Taf2</i>	dme:Dmel CG6711	0.91	1629974 at
				TATA binding protein	<i>Tbp</i>	dme:Dmel CG9874	0.906	1637663 at
				Transcription factor TFIIIFbeta	<i>TfIIIFbeta</i>	dme:Dmel CG6538	0.902	1633349 at
				TBP-associated factor 4	<i>Taf4</i>	dme:Dmel CG5444	0.896	1624037 s at
				Transcription factor IIIb	<i>TfIIIB</i>	dme:Dmel CG5193	0.895	1634172 at
				TBP-associated factor 12	<i>Taf12</i>	dme:Dmel CG17358	0.895	1625702 s at
				TBP-related factor	<i>Trf</i>	dme:Dmel CG7562	0.881	1624550 at
				TBP-associated factor 4	<i>Taf4</i>	dme:Dmel CG5444	0.858	1640309 s at
				enhancer of yellow 1	<i>e(y)1</i>	dme:Dmel CG6474	0.856	1633139 at
				TBP-associated factor 10b	<i>Taf10b</i>	dme:Dmel CG3069	0.852	1630803 at
				Transcription factor IIIBeta	<i>TfIIIBeta</i>	dme:Dmel CG1276	0.851	1634616 at
				TBP-associated factor 10	<i>Taf10</i>	dme:Dmel CG2859	0.82	1635508 at

				TBP-associated factor 7	<i>Taf7</i>	dme:Dmel CG2670	0.814	1640551_at
				Transcription factor IIIEalpha	<i>TllEalpha</i>	dme:Dmel CG10415	0.759	1639648_at
				Mgat2	<i>Mgat2</i>	dme:Dmel CG7921	1.028	1633061_at
				Alpha 3 glucosyltransferase	<i>Alg10</i>	dme:Dmel CG32076	0.996	1636806_at
				peptide O-xylosyltransferase	<i>oxt</i>	dme:Dmel CG32300	0.986	1637482_at
				Polypeptide N-acetylgalactosaminyltransferase 35A	<i>Pgant35A</i>	dme:Dmel CG7480	0.966	1632680_at
				UDP-GlcNAc:alpha-3-D-mannoside-beta-1,2-N-acetylglucosaminyltransferase 1	<i>Mgat1</i>	dme:Dmel CG13431	0.924	1623607_at
				O-glycosyltransferase	<i>Ogt</i>	dme:Dmel CG10392	0.911	1626219_s_at
				UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylgalactosaminyltransferase 2	<i>GalNAc-T2</i>	dme:Dmel CG6394	0.906	1629117_at
				Oligosaccharyltransferase 48kD subunit	<i>Ost48</i>	dme:Dmel CG9022	0.902	1625055_at
				Cog7	<i>Cog7</i>	dme:Dmel CG31040	0.879	1631831_at
				polypeptide GalNAc transferase 3	<i>pgant3</i>	dme:Dmel CG4445	0.866	1637584_at
				alpha Mannosidase II	<i>alpha-Man-II</i>	dme:Dmel CG18802	0.864	1626112_at
				brother of tout-velu	<i>btv</i>	dme:Dmel CG15110	0.863	1636990_at
				UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylgalactosaminyltransferase 2	<i>GalNAc-T2</i>	dme:Dmel CG6394	0.857	1637736_a_at
				alpha Mannosidase I	<i>alpha-Man-I</i>	dme:Dmel CG32684	0.854	1629744_a_at
				tout-velu	<i>tv</i>	dme:Dmel CG10117	0.819	1637442_at
				lethal (2) neighbor of tid	<i>l2not</i>	dme:Dmel CG4084	0.816	1641244_at
				polypeptide GalNAc transferase 6	<i>pgant6</i>	dme:Dmel CG2103	0.801	1636337_s_at
				beta-4-galactosyltransferase 7	<i>beta4GalT7</i>	dme:Dmel CG11780	0.799	1632917_at
				polypeptide GalNAc transferase 5	<i>pgant5</i>	dme:Dmel CG31651	0.783	1635173_at
				Glutamate oxaloacetate transaminase 2	<i>Got2</i>	dme:Dmel CG4233	0.957	1629055_a_at
				Phosphoribosylamidotransferase	<i>Prat</i>	dme:Dmel CG2867	0.948	1627868_at
				Glutamine:fructose-6-phosphate aminotransferase 2	<i>Gfat2</i>	dme:Dmel CG1345	0.916	1635153_at
				burgundy	<i>bur</i>	dme:Dmel CG9242	0.911	1635718_at
				Glutamine synthetase 1	<i>Gs1</i>	dme:Dmel CG2718	0.908	1632852_s_at
				Glutathione Synthetase	<i>GS</i>	dme:Dmel CG6835	0.905	1627321_x_at
				GS1-like	<i>Gs1l</i>	dme:Dmel CG15441	0.895	1633149_at
				Glutamate-cysteine ligase catalytic subunit	<i>Gclt</i>	dme:Dmel CG2259	0.895	1625362_at
				Glutamate dehydrogenase	<i>Gdh</i>	dme:Dmel CG5320	0.883	1640835_a_at
				Glutamate oxaloacetate transaminase 1	<i>Got1</i>	dme:Dmel CG8430	0.861	1631817_a_at
				Glutamyl-tRNA synthetase	<i>Aats-ghn</i>	dme:Dmel CG10506	0.845	1640088_at
				wolfram syndrome 1	<i>wfs1</i>	dme:Dmel CG4917	0.841	1626657_a_at
				Glutamate-cysteine ligase modifier subunit	<i>Gclm</i>	dme:Dmel CG4919	0.771	1640552_at
				GS1-like	<i>Gs1l</i>	dme:Dmel CG15441	0.757	1632028_a_at
				rudimentary	<i>r</i>	dme:Dmel CG18572	0.749	1641267_at
				GS1-like	<i>Gs1l</i>	dme:Dmel CG15441	0.732	1623430_at
				no extended memory	<i>nemy</i>	dme:Dmel CG8772	0.705	1634658_a_at
				Glutamine synthetase 2	<i>Gs2</i>	dme:Dmel CG1743	0.662	1630653_a_at
				Inositol 1,4,5-triphosphate kinase 1	<i>IP3K1</i>	dme:Dmel CG4026	1.443	1634619_at
				Phosphatidylinositol 3 kinase 59F	<i>Pi3K59F</i>	dme:Dmel CG5373	1.166	1625678_at
				ljk2	<i>ljk2</i>	dme:Dmel CG13688	1.008	1628514_at
				Phospholipase C at 21C	<i>Ple21C</i>	dme:Dmel CG4574	0.976	1630007_s_at
				immune response deficient 5	<i>ird5</i>	dme:Dmel CG4201	0.957	1625974_at
				loki	<i>lok</i>	dme:Dmel CG10895	0.946	1625038_s_at
				no receptor potential A	<i>norpA</i>	dme:Dmel CG3620	0.929	1636576_s_at
				Multiple inositol polyphosphate phosphatase 2	<i>Mipp2</i>	dme:Dmel CG4317	0.927	1630756_at
				Cyclin-dependent kinase 4	<i>Cdk4</i>	dme:Dmel CG5072	0.925	1639347_s_at
				synaptotagmin	<i>synj</i>	dme:Dmel CG6562	0.887	1631427_s_at
				Inositol 1,4,5-triphosphate kinase 2	<i>IP3K2</i>	dme:Dmel CG1630	0.882	1634634_s_at
				skittles	<i>sktl</i>	dme:Dmel CG9985	0.882	1624275_at
				Inositol polyphosphate 1-phosphatase	<i>lpp</i>	dme:Dmel CG3028	0.874	1626217_at
				small wing	<i>sl</i>	dme:Dmel CG4200	0.867	1628815_at
				Pi4KIIalpha	<i>Pi4KIIalpha</i>	dme:Dmel CG2929	0.86	1627996_a_at
				four wheel drive	<i>fwd</i>	dme:Dmel CG7004	0.839	1634509_s_at
				Neurexin IV	<i>Nrx-IV</i>	dme:Dmel CG6827	0.771	1639459_a_at
				no receptor potential A	<i>norpA</i>	dme:Dmel CG3620	0.697	1630119_s_at
				Suppressor of cytokine signaling at 36E	<i>SoCS36E</i>	dme:Dmel CG15154	1.669	1637703_a_at
				nejire	<i>nej</i>	dme:Dmel CG15319	1.03	1622925_at
				Signal-transducer and activator of transcription protein at 92E	<i>Stat92E</i>	dme:Dmel CG4257	0.995	1638827_s_at
				downstream of receptor kinase	<i>drk</i>	dme:Dmel CG6033	0.989	1625475_s_at
				Suppressor of variegation 2-10	<i>Suvvar2-10</i>	dme:Dmel CG8068	0.989	1627362_a_at
				Suppressor of variegation 2-10	<i>Suvvar2-10</i>	dme:Dmel CG8068	0.96	1632783_a_at
				hopscotch	<i>hop</i>	dme:Dmel CG1594	0.924	1639072_at
				Son of sevenless	<i>Sos</i>	dme:Dmel CG7793	0.912	1633355_at
				sprouty	<i>sty</i>	dme:Dmel CG1921	0.893	1635819_at
				Suppressor of Cytokine Signaling at 44A	<i>SoCS4A</i>	dme:Dmel CG2160	0.89	1633403_at
				Signal transducing adaptor molecule	<i>Stam</i>	dme:Dmel CG6521	0.876	1635260_at
				sprouty	<i>sty</i>	dme:Dmel CG1921	0.859	1635677_a_at
				corkscrew	<i>csw</i>	dme:Dmel CG3954	0.836	1625775_a_at
				Suppressor of Cytokine Signaling at 16D	<i>SoCS16D</i>	dme:Dmel CG8146	0.825	1636122_at
				Cyclin D	<i>CycD</i>	dme:Dmel CG9096	0.807	1627295_s_at
				Sprouty-related protein with EVH-1 domain	<i>Sprel</i>	dme:Dmel CG10155	0.752	1639243_at
				Synaptobrevin	<i>Syb</i>	dme:Dmel CG12210	1.199	1626835_at
				Syntaxin 8	<i>Syx8</i>	dme:Dmel CG4109	0.964	1623564_at
				Synaptobrevin	<i>Syb</i>	dme:Dmel CG12210	0.919	1637089_at
				ubsnap	<i>usnp</i>	dme:Dmel CG11173	0.917	1625742_at
				Syntaxin protein 24	<i>Snap24</i>	dme:Dmel CG9474	0.914	1635597_at
				Synaptobrevin	<i>Syb</i>	dme:Dmel CG12210	0.905	1635944_a_at
				Syntaxin 17	<i>Syx17</i>	dme:Dmel CG7452	0.9	1628202_at
				Syntaxin 1A	<i>Syx1A</i>	dme:Dmel CG31136	0.896	1624043_at
				Syntaxin 7	<i>Syx7</i>	dme:Dmel CG5081	0.895	1636575_s_at
				Syntaxin 5	<i>Syx5</i>	dme:Dmel CG4214	0.882	1625607_s_at
				Syntaxin 18	<i>Syx18</i>	dme:Dmel CG13626	0.876	1626506_at
				Syntaxin 16	<i>Syx16</i>	dme:Dmel CG1467	0.853	1635421_at
				Syntaxin 6	<i>Syx6</i>	dme:Dmel CG7736	0.804	1639075_a_at
				Syntaxin 4	<i>Syx4</i>	dme:Dmel CG2715	0.77	1638808_at
				lethal (1) G0155	<i>l1G0155</i>	dme:Dmel CG1515	0.763	1640976_at
				Syntaxin 13	<i>Syx13</i>	dme:Dmel CG11278	0.691	1635186_at
				UDP-glycosyltransferase 35a	<i>Ugt35a</i>	dme:Dmel CG6644	1.025	1627662_at
				sugarless	<i>sgl</i>	dme:Dmel CG10072	0.989	1629352_at
				abstrakt	<i>abs</i>	dme:Dmel CG14637	0.982	1623353_at
				Hexokinase A	<i>Hex-A</i>	dme:Dmel CG3001	0.935	1625638_a_at
				reptin	<i>rept</i>	dme:Dmel CG9750	0.858	1630792_at
				pitchoone	<i>pit</i>	dme:Dmel CG6375	0.843	1625758_s_at
				GleAT-P	<i>GleAT-P</i>	dme:Dmel CG6207	0.834	1624164_s_at
				Ugt58Fa	<i>Ugt58Fa</i>	dme:Dmel CG4414	0.824	1632070_at
				Phosphoglucose isomerase	<i>Pgi</i>	dme:Dmel CG8251	0.823	1638550_s_at
				GleAT-S	<i>GleAT-S</i>	dme:Dmel CG3881	0.797	1628402_a_at
				prune	<i>pu</i>	dme:Dmel CG3461	0.796	1637479_at