

Spalt +

Name	Symbol	RNAi strain	Phenotype	Molecular	Molecular
Age 24h/48h					
CG1058	<i>rpk</i>	1058R-1	+	Ligand-gated sodium channel	Neu
CG10965	<i>Corp</i>	102751-KK	wS	Apoptotic signaling pathway in response to DNA damage	CD
CG11086	<i>Gadd45</i>	BL-35023	+	JNK cascade	CS
CG11155	CG11155	BL-31991	+	Glutamate receptor	Neu
CG11584	CG11584	-----	-----	-----	CG
CG11897	CG11897	28260-GD	+	ABC transporter	Tra
CG12177	CG12177	17018-GD	+	Inosine/uridine-preferring nucleoside hydrolase	M
CG1221	<i>miple</i>	1221R-1	+	Pleiotrophin/Midkine heparin-binding growth factor	CS
CG12224	CG12224	26497-GD	+	Oxidoreductase	RedOx
CG12367	<i>Hen1</i>	31920-GD	+	RNA methylation	R
CG13041	CG13041	109222-KK	S-P//S-P	Retinin-like	CGh
CG13679	CG13679	49149-GD	V+/S	-----	CG
CG13937	CG13937	3373-GD	+	Carbohydrate sulfotransferase	M
CG14907	CG14907	39851-GD	+	-----	CG
CG1512	<i>Cul-2</i>	100540-KK	sS	Ubiquitin protein ligase	P
CG15400	CG15400	7261-GD	+	Glucose-6-phosphatase	M
CG15784	CG15784	15784R-3	nW/sN//S-P	-----	CG
CG15892	CG15892	BL-42945	+	-----	CG
CG16857	CG16857	BL-31974	+	Fibronectin, type III; Immunoglobulin I-set	CA
CG17104	CG17104	17104R-3	+	-----	CG
CG17530	<i>GstE6</i>	17530R-1	+	Glutathione S-transferase	RedOx
CG17533	<i>GstE8</i>	26497-GD	PL-nW//S	Glutathione transferase	RedOx
CG17574	CG17574	102094-KK	+	-----	CG
CG17803	CG17803	BL-29367	+	C2H2-type/integrase DNA-binding domain	D
CG18188	<i>Damm</i>	109520-KK	sF//+	Cysteine-type endopeptidase	P
CG18278	CG18278	BL-28520	+	N-acetylglucosamine-6-sulfatase	M

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<i>CG18455</i>	<i>Optix</i>	18455R-2	S-P	Sequence-specific DNA binding	D
<i>CG1851</i>	<i>Ady43A</i>	33133-GD	+	Carbohydrate kinase PfkB	M
<i>CG18522</i>	<i>CG18522</i>	100330-KK	+	Oxidoreductase	RedOx
<i>CG18543</i>	<i>mtrm</i>	BL-35430	+	Meiotic cell cycle	CDiv
<i>CG2184</i>	<i>Mlc2</i>	BL-36694	EPL//S-P	Microfilament motor	Cyt
<i>CG2560</i>	<i>Cpr11A</i>	103321-KK	V+	Cuticle	Cut
<i>CG2909</i>	<i>CG2909</i>	100165-KK	V+	-----	CG
<i>CG2914</i>	<i>Ets21C</i>	106153-KK	+	DNA binding transcription factor	D
<i>CG3074</i>	<i>Swim</i>	BL-36591	+	Wnt-protein binding	CS
<i>CG3081</i>	<i>CG3081</i>	102507-KK	sF//+	-----	CG
<i>CG31332</i>	<i>Unc-115b</i>	BL-40839	+	Actin binding	Cyt
<i>CG31875</i>	<i>CG31875</i>	104638-KK	+	Chromatin binding	D
<i>CG32475</i>	<i>mth18</i>	4071-GD	+	G-protein coupled receptor signaling	CS
<i>CG32581</i>	<i>CG32581</i>	49728-GD	EPL//S-P	Zinc finger, RING-type	CGh
<i>CG32625</i>	<i>CG32625</i>	103216-KK	+	CHHC zinc finger	CGh
<i>CG32788</i>	<i>Crg-1</i>	BL-46971	+	Sequence-specific DNA binding	D
<i>CG33048</i>	<i>Mocs1</i>	-----	-----	Mo-molybdopterin cofactor biosynthetic	M
<i>CG33653</i>	<i>Caps</i>	BL-31984	+	Phospholipid binding	Neu
<i>CG3397</i>	<i>CG3397</i>	BL-25818	+	Oxidoreductase	RedOx
<i>CG34002</i>	<i>CG34002</i>	-----	-----	V5/Tpx-1-related	CGh
<i>CG3448</i>	<i>CG3448</i>	26198-GD	+	DNA repair protein XRCC4	D
<i>CG3568</i>	<i>CG3568</i>	18447-GD	+	-----	CG
<i>CG40300</i>	<i>ago-03</i>	35232-GD	+	mRNA poly(A) tail shortening	R
<i>CG40452</i>	<i>Snap25</i>	BL-34377	+	Target SNARE	Neu
<i>CG4174</i>	<i>CG4174</i>	BL-36845	+	Procollagen-proline 4-dioxygenase	P
<i>CG42516</i>	<i>CG42516</i>	-----	-----	Transcription factor TFIIC	D

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Age 24h/48h					
CG42644	CG42644	-----	-----	-----	CG
CG43144	CG43144	-----	-----	Non protein coding gene	CR
CG43646	CG43646	-----	-----	-----	CG
CG43676	CG43676	-----	-----	-----	CG
CG4594	CG4594	100448-KK	+	Dodecenoyl-CoA delta-isomerase	M
CG4920	<i>ea</i>	BL-41961	+	Serine-type endopeptidase	P
CG5022	CG5022	BL-33359	+	Ezrin/radixin/moesin like	Cyt
CG5096	CG5096	100951-KK	+	Leucine-rich repeat	CGh
CG5229	<i>chm</i>	BL-36869	wS	H4 histone acetyltransferase	D
CG5966	CG5966	BL-33932	+	Triglyceride lipase	M
CG6658	<i>Ugt86Di</i>	104262-KK	+	Glucuronosyltransferase	M
CG6665	CG6665	104625-KK	+	NICE-3	CGh
CG6794	<i>Dif</i>	30579-GD	+	DNA binding transcription factor	D
CG6912	CG6912	102806-KK	+	-----	CG
CG7035	<i>Cbp80</i>	BL-33648	+	RNA interference	R
CG7590	<i>scyl</i>	25506-GD	+	Negative regulation of growth; response to DNA damage	CD
CG7787	CG7787	BL-32843	+	Guanyl-nucleotide exchange factor	Tra
CG8573	<i>su(Hw)</i>	BL-33906	+	Negative regulation of chromatin silencing	D
CG8620	CG8620	BL-42638	+	-----	CG
CG8799	<i>l(2)03659</i>	100105-KK	+	ATPase, transporter	Tra
CG9214	<i>Tob</i>	108824-KK	+	Anti-proliferative	CS
Age 48h					
CG10160	<i>ImpL3</i>	10160R-1	S-P//S-P	Glycolysis	M
CG10245	<i>Cyp6a20</i>	3313-GD	+	Electron carrier	RedOx
CG1155	<i>Osi14</i>	102979-KK	+	DUF1676	CGh
CG11592	<i>Amnionless</i>	BL-41956	+	Amnionless	CGh

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Age 48h					
CG12242	<i>GstD5</i>	BL-42842	+	Glutathione transferase	RedOx
CG12505	<i>Arc1</i>	48130-GD	S/V	Nucleic acid binding	D
CG12846	<i>Tsp42Ed</i>	105847-KK	+	Tetraspanin	CGh
CG12908	<i>Ndg</i>	109625-KK	+	EGF-like calcium-binding	CA
CG1299	<i>CG1299</i>	108705-KK	EPL//wV+	Serine-type endopeptidase	P
CG13060	<i>CG13060</i>	49608-GD	+	-----	CG
CG13941	<i>Arc2</i>	3464-GD	S/Bs	-----	CG
CG14059	<i>Ilp8</i>	102604-KK	+	Insulin	CS
CG14196	<i>CG14196</i>	-----	-----	Monocarboxylate transmembrane transporter	Tra
CG14304	<i>CG14304</i>	106221-KK	+	Chitin	Cut
CG14545	<i>CG14545</i>	BL-35625	+	-----	CG
CG14695	<i>CG14695</i>	14695R-1	+	-----	CG
CG15209	<i>CG15209</i>	5534-GD	+	-----	CG
CG15347	<i>CG15347</i>	107479-KK	+	DUF753	CGh
CG15623	<i>c-cup</i>	106603-KK	+	Cytochrome b5-like heme/steroid binding domain	Mit
CG18859	<i>Or19a</i>	BL-31892	+	Olfactory receptor	Neu
CG2065	<i>CG2065</i>	BL-34098	+	Glucose/ribitol dehydrogenase	M
CG2885	<i>RabX2</i>	103311-KK	+	Small GTPase mediated signal transduction	CS
CG30196	<i>CG30196</i>	30196R-3	+	LPS-induced tumor necrosis factor alpha factor.	CD
CG31062	<i>side</i>	1283-GD	+	CD80-like, immunoglobulin C2-set; Fibronectin, type III	Neu
CG31279	<i>CG31279</i>	109930-KK	V+	-----	CG
CG32825	<i>Or19b</i>	109050-KK	sF//+	Olfactory receptor	Neu
CG34031	<i>CG34031</i>	-----	-----	-----	CG
CG34165	<i>CG34165</i>	109149-KK	sF//+	-----	CG
CG3767	<i>Jhl-26</i>	36566-GD	+	CHK kinase	CS
CG4181	<i>GstD2</i>	109123-KK	sF//+	Glutathione transferase	RedOx

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Age 48h					
<i>CG42255</i>	<i>CG42255</i>	100425-KK	F/sS	Metalloendopeptidase	P
<i>CG42352</i>	<i>Ir40a</i>	100912-KK	+	Ion channel	Neu
<i>CG42363</i>	<i>CG42363</i>	-----	-----	-----	CG
<i>CG4373</i>	<i>Cyp6d2</i>	101829-KK	+	Electron carrier	RedOx
<i>CG4423</i>	<i>GstD6</i>	105805-KK	F/S//Bs	Glutathione transferase	RedOx
<i>CG5955</i>	<i>CG5955</i>	102443-KK	sS/F	UDP-glucose 4-epimerase	M
<i>CG6042</i>	<i>Cyp12a4</i>	107088-KK	+	Electron carrier	RedOx
<i>CG6142</i>	<i>CG6142</i>	100131-KK	+	Glucose-methanol-choline oxidoreductase	M
<i>CG7080</i>	<i>CG7080</i>	104928-KK	+	-----	CG
<i>CG7356</i>	<i>Tg</i>	103601-KK	Bs/S	Protein-glutamine gamma-glutamyltransferase	P
<i>CG8927</i>	<i>CG8927</i>	14167-GD	+	Cuticle	Cut
<i>CG9521</i>	<i>CG9521</i>	106710-KK	+	Oxidation-reduction	RedOx
<i>CG9877</i>	<i>CG9877</i>	-----	-----	-----	CG
Age 24h					
<i>CG11486</i>	<i>CG11486</i>	BL-42492	+	Protein kinase, catalytic domain	R
<i>CG13316</i>	<i>Mnt</i>	BL-42558	EPL/nW//+	Sequence-specific DNA binding	D
<i>CG30059</i>	<i>CG30059</i>	BL-28520	+	N-acetylglucosamine-6-sulfatase	M
<i>CG32019</i>	<i>bt</i>	BL-31546	+	Fibronectin, type III; Immunoglobulin I-set	CA
<i>CG45050</i>	<i>CG43674</i>	BL-31375	+	DNA binding	D
<i>CG6544</i>	<i>fau</i>	106560-KK	+	-----	CG
<i>CG7178</i>	<i>wupA</i>	BL-31893	EPL	Tropomyosin binding; actin binding	Cyt
<i>CG8663</i>	<i>nrv3</i>	BL-29431	L	Potassium ion transport; sodium ion transport	Tra

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				Df//UAS	
CG10102	CR10102/Arc3	-----	-----	Non protein coding gene	CR
CG10916	CG10916	107518-KK	PL/nW//wS-P	Zinc ion binding	D
CG1303	<i>agt</i>	104331-KK	+	DNA methylation	D
CG14121	<i>l(3)S147910/ver</i>	14121R-1	+	Telomere capping	D
CG15009	<i>ImpL2</i>	15009R-3	+	Insulin-like growth facgtor binding	CS
CG16928	<i>mre11</i>	30476-GD	+	Nucelase activity	D
CG18213	CG18213	18213R-2	+	Zinc finger	CGh
CG1925	<i>mus205</i>	103775-KK	sS/Bs//S	DNA binding	D
CG2999	<i>unc-13</i>	33609-GD	+	Intracellular signal transduction	CS
CG3008	CG3008	43732-GD	S-P	Protein kinase activity	P
CG31705	CG31705	7681-GD	+	-----	CG
CG32021	CG32021	102286-KK	V-/S	-----	CG
CG5202	<i>escl</i>	49982-GD	+	Histone methyltransferase activity	D
CG5247	<i>lrbp</i>	5247R-2	sS//S-P	DNA binding	D
CG6272	CG6272	101871-KK	EPL	Sequence-specific DNA binding	D

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Age 24h/48h					
CG10344	CG10344	48026-GD	+	Cysteine S-palmitoleyltransferase	P
CG10539	S6k	10539-R3	wS	Ribosomal protein S6 kinase	CS
CG10541	Tektin-C	100094-KK	EPL//S-P	Microtubule binding	Cyt
CG10576	CG10576	BL-34388	+	Aminopeptidase	P
CG11253	CG11253	105377-KK	+	Sensory perception of sound	CGh
CG11263	CG11263	31112-GD	+	Ribonuclease H-like	R
CG12182	CG12182	12182R-1	S-V//S-P	-----	CG
CG12275	RpS10a	104132-KK	EPL//sS-P	Structural constituent of ribosome	R
CG12843	Tsp42Ei	101044-KK	sF//+	EC2 domain	CGh
CG13083	CG13083	106400-KK	PL/nW//wS-P	-----	CG
CG1342	Spn100A	1342R-1	+	Serpin domain	CGh
CG13890	CG13890	108862-KK	Cell Pig./+	Dodecenoyl-CoA delta-isomerase	M
CG13897	CG13897	13897R-1	+	DNA binding	D
CG14394	NijC	39307-GD	Bs/S/V+	Ninjurin	CA
CG14485	swi2	BL-28530	+	Response to ecdysone	CG
CG14715	CG14715	104124-KK	+	Peptidyl-prolyl cis-trans isomerase	P
CG15236	CG15236	15236R-1	S/Bs	-----	CG
CG15707	krimp	BL-37511	+	Tudor domain	R
CG17077	pnt	105390-KK	+	Protein binding	CS
CG2161	Rga	20826-GD	S // S/CD	G2/M transition DNA damage checkpoint	D
CG2849	Rala	43622-GD	S-P/nW//S-P	Ras	CS
CG30104	CG30104	106640-KK	sF//+	5'-nucleotidase	D
CG30488	CG30488	100513-KK	wV+/wBs	Allergen V5/Tpx-1-related	CGh
CG31262	CG31262	52605-GD	+	Sodium/solute symporter	Tra
CG31436	CG31436	21344-GD	S/V-	CHK kinase-like	CGh
CG31809	CG31809	BL-29424	+	Steroid dehydrogenase	M

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Age 24h/48h					
<i>CG31810</i>	<i>CG31810</i>	BL-29424	+	Steroid dehydrogenase	M
<i>CG32364</i>	<i>CG32364</i>	26045-GD	+	RNA recognition motif domain	R
<i>CG33140</i>	<i>CG33140</i>	18688-GD	EPL//S-P	-----	CG
<i>CG33302</i>	<i>Cpr31A</i>	45310-GD	S/Bs	Cuticle	Cut
<i>CG34215</i>	<i>CG34215</i>	-----	-----	-----	CG
<i>CG3649</i>	<i>CG3649</i>	3649R-1	sS-P//S-P	Phosphate:sodium symporter	Tra
<i>CG40178</i>	<i>CG40178</i>	110089-KK	sF//+	Redox homeostasis	Red-Ox
<i>CG4086</i>	<i>Su(P)</i>	BL-34955	+	Glutathione S-transferase	Red-Ox
<i>CG41320</i>	<i>CG41320</i>	109162-KK	sF//+	Thioredoxin-like fold	Red-Ox
<i>CG42330</i>	<i>Dscam4</i>	42882-KK	+	Cell adhesion	CA
<i>CG43079</i>	<i>nrm</i>	104295-KK	+	CD80-like, immunoglobulin C2-set	CA
<i>CG43103</i>	<i>CG43103</i>	-----	-----	-----	CG
<i>CG43117</i>	<i>CG43117</i>	103091-KK	+	-----	CG
<i>CG4570</i>	<i>CG4570</i>	109328-KK	sS	Transposase, Tc1-like	D
<i>CG5671</i>	<i>Pten</i>	35731-GD	LS	Phosphoprotein phosphatase activity	CS
<i>CG6013</i>	<i>CG6013</i>	6013R-2	+	HMG box domain	D
<i>CG7160</i>	<i>Cpr78E</i>	7160R-3	sS	Cuticle	Cut
<i>CG7354</i>	<i>mRpS26</i>	100067-KK	EPL//S-P	Ribosomal protein	R
<i>CG7478</i>	<i>Act79B</i>	BL-36857	+	Cytoskeleton	Cyt
<i>CG7532</i>	<i>l(2)34Fc</i>	7532R-1	+	Serine-type endopeptidase	P
<i>CG7607</i>	<i>CG7607</i>	9208-GD	+	Immunoglobulin-like domain	CA
<i>CG7916</i>	<i>CG7916</i>	101195-KK	sF//+	Haemolymph juvenile hormone binding	CGh
<i>CG8023</i>	<i>eIF4E-3</i>	BL-42804	+	Eukaryotic initiation factor 4G binding	R
<i>CG8768</i>	<i>CG8768</i>	8768R-2	sS	NAD-dependent epimerase/dehydratase	M
<i>CG9333</i>	<i>Oseg5</i>	100020-KK	S/V+	WD40-repeat	CGh

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CG11051	<i>Nplp2</i>	15305-GD	+	Neuropeptide hormone	CS
CG14332	CG14332	41165-GD	+	-----	CG
CG17876	<i>Amy-d</i>	109082-KK	+	Alpha-amylase	M
CG18363	<i>dic-04</i>	8416-GD	+	Mitochondrial carrier	Tra
CG33126	<i>NLaz</i>	101321-KK	sS/V+	Lipocalin/cytosolic fatty-acid binding	M
CG42798	CG42798	-----	-----	-----	CG
CG42834	CG42834	-----	-----	-----	CG
CG9486	CG9486	102802-KK	S/Bs	Acyl-CoA N-acyltransferase	M
Age 24h					
CG13053	CG13053	9227-GD	+	-----	CG
CG17134	CG17134	BL-42868	+	Aspartic-type endopeptidase	P
CG2360	<i>Ccp84Aa</i>	49694-GD	+	Chitin	Cut
CG3200	<i>Reg-2</i>	3200R-3	+	Phosphoglycolate phosphatase	M
CG32444	CG32444	101230-KK	+	Aldose 1-/Glucose-6-phosphate 1-epimerase	M
CG43114	CG43114	-----	-----	-----	CG
CG6579	<i>atilla</i>	101803-KK	sF//+	-----	CG
CG8800	CG8800	107835-KK	sF//+	Leucine rich repeat 4	CGh
Df//UAS					
CG10041	CG10041	110063-KK	+	Serine-type endopeptidase activity	P
CG10225	<i>RanBP3</i>	104432-KK	F//+	Negative regulation of Wnt signaling pathway	CS
CG10382	<i>wrapper</i>	10382R-3	+	Immunoglobulin subtype 2	CGh
CG11212	<i>Ptr</i>	BL-28973	+	Hedgehog receptor activity	CS
CG11357	CG11357	108467-KK	L/Bs	Transferring glycosyl groups	P
CG11607	<i>H2.0</i>	15876-GD	L//sS-P	Sequence-specific DNA binding	D
CG11797	<i>Obp56a</i>	100850-KK	+	Odorant binding	CS

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CG11883	CG11883	38590/GD	S-P//S-P	5'-nucleotidase activity	D
CG12052	<i>lola</i>	BL-35721	L/F	Sequence-specific Dna binding transcription factor activity	D
CG12256	CG12256	12256R-4	+	Serine-type endopeptidase activity	P
CG12287	<i>pdm2</i>	102126-KK	Bs	DNA binding	D
CG1273	CG1273	4496-GD	+	-----	CG
CG12952	<i>sage</i>	BL-25980	+	Sequence-specific DNA binding transcription factor activity	D
CG13618	CG13618	14714-GD	+	Haemolymph juvenile hormone binding	CGh
CG13848	<i>pinta</i>	13848R-4	+	Retinoid binding	M
CG14590	CG14590	41536-GD	+	Histone deacetylase binding	D
CG14869	CG6107	110157-KK	+	Zinc ion binding	P
CG15369	CG15369	15369R-2	+	Cysteine-type endopeptidase inhibitor activity	P
CG15739	CG15739	106016-KK	F//S-P	Phosphatase activity	P
CG16705	<i>SPE</i>	30972-GD	L//S-P	Serine-type endopeptidase activity	P
CG16712	CG16712	109238-KK	+	Serine-type endopeptidase inhibitor activity	P
CG16756	CG16756	102908-KK	S	Lysozyme activity	P
CG16947	CG16947	108103-KK	+	Zinc finger, RING-type	CGh
CG1725	<i>dlg1</i>	1725-R2	L//sS-P	Epidermal growth factor receptor binding	CA
CG1763	<i>nod</i>	48150-GD	+	DNA binding	D
CG1780	<i>ldgf4</i>	1780R-4	+	Imaginal disc growth factor receptor binding	CS
CG18024	<i>SoxN</i>	BL-25996	+	Chromatin DAN binding	D
CG18255	<i>Strn-Mlck</i>	-----	-----	Protein serine/threonine kinase activity	P
CG18410	<i>Ude</i>	18410R-2	+	DNA binding	D
CG18657	<i>NetA</i>	18657R-2	+	Netrin domain	CGh
CG1869	<i>Cht7</i>	42877-GD	sS-P//sS-P	Chitinase activity	Cut
CG2056	<i>spirit</i>	3285-GD	+	Serine-type endopeptidase activity	P
CG2488	<i>phr6-4</i>	2488R-3	+	DNA (6-4) photolyase activity	D

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CG2706	<i>fs(1)Yb</i>	110056-KK	+	Tudor domain	D
CG30069	CG30069	25562-GD	S/Bs	-----	CG
CG3097	CG3097	-----	-----	Zinc ion binding	P
CG3100	<i>b6</i>	14526-GD	+	Neuronal pentraxin receptor activity	CS
CG31098	CG10634	21300-GD	V+	Transferase activity	P
CG31721	<i>Trim9</i>	100767-KK	+	Zinc finger, B-box	CGh
CG31764	<i>vir-1</i>	102534-KK	+	-----	CG
CG31839	<i>nimB2</i>	105925-KK	+	Insulin-like growth factor binding protein	CS
CG31884	<i>Trx-2</i>	36297-GD	+	Disulfide oxidoreductase activity	D
CG32029	<i>Cpr66D</i>	112709-KK	S/Bs	Structural constituent of chitin-based cuticle	Cut
CG32055	CG32055	6335-GD	wS	Insulin-like growth factor binding	CS
CG32261	<i>Gr64a</i>	103342-KK	S/F	Taste receptor activity	CS
CG3240	<i>Rad1</i>	12676-GD	+	Exonuclease activity	D
CG32491	<i>mod(mdg4)</i>	BL-32995	F	DNA binding	D
CG3289	<i>Ptpa</i>	41913-GD	+	Protein tyrosine phosphatase activator activity	CS
CG33197	<i>mbl</i>	BL-29585	Bs/P	DNA binding	D
CG33960	<i>Sema-2b</i>	BL-28932	+	Sema domain	CS
CG33970	CG33970	33090-GD	+	Transporter activity	Tr
CG34406	CG44002	43767-GD	+	Zinc ion binding	D
CG42614	<i>scrib</i>	BL29552	PL/nW/sS-P/CD	Protein binding	CA
CG4322	<i>moody</i>	4322R-1	S-P//S-P	Melatonin receptor activity	CS
CG4379	<i>Pka-C1</i>	101524-KK	S-P	Protein binding	CS
CG4398	CG4398	4398R-1	+	-----	CG
CG4432	<i>PGRP-LC</i>	4432R-1	+	Protein dimerization activity	D
CG4565	CG4565	BL-32893	+	Histone-lysine N-methyltransferase activity	D
CG5249	<i>Blimp-1</i>	5249R-2	S-P/F	Nucleic acid binding	D

Spalt -

Name	Symbol	RNAi strain	Phenotype	Molecular	Molecular
				Df//UAS	
CG6134	<i>spz</i>	BL-28538	+	Protein homodimerization activity	CS
CG6713	<i>Nos</i>	BL-28792	+	Nitric-oxide synthase activity	D
CG7201	<i>CG7201</i>	101198-KK	sF//+	-----	CG
CG7420	<i>CG7420</i>	BL-32510	L	Regulator of chromosome condensation, RCC1	D
CG7577	<i>ppk20</i>	36659-GD	S	Sodium channel activity	Tr
CG7734	<i>shn</i>	105643-KK	sS-P//sS-P	RNA polymerase II activating transcription factor binding	D
CG7897	<i>gp210</i>	-----	-----	Protein import into nucleus	CS
CG7906	<i>CG7906</i>	BL-30519	+	Kazal domain	P
CG8084	<i>ana</i>	BL-27515	S/V+//S-P	Growth factor activity	CS
CG8589	<i>tej</i>	103264-KK	+	Tudor domain	D
CG8675	<i>CG8675</i>	8675R-1	S-P/nW//S-P	-----	CG
CG8773	<i>CG8773</i>	10203-GD	+	Metallopeptidase activity	P
CG8780	<i>tey</i>	106065-KK	Bs/S	Negative regulation of transcription	D
CG8945	<i>CG8945</i>	46925-GD	F	Zinc ion binding	P
CG9355	<i>dy</i>	9355R-1	L//S-P	Structural constituent of chitin-based cuticle	Cut
CG9460	<i>Spn42De</i>	9460R-3	+	Serine-type endopeptidase inhibitor activity	P
CG9850	<i>CG11169</i>	107199-KK	sF//+	Metalloendopeptidase activity	P