

gene in red: gene with a real "Immunity" signature or strongly induced (>3) or a function related to resistance to pathogens  
 gene \* Genes found in several places

## Drosophila genes potentially involved in responses to microbial infection

Last update : June 2013

Expression pattern (Degregorio 2001,2002)

name Flybase link Symbol homology and biology Septic injury B.bassiana Key references (for useful tools with an emphasis on mutations)

### I. Family of receptors that could be involved Microbial recognition (classified by gene family)

#### CD36/croquemort (crq) family : 9 members

croquemort	<a href="#">CG4280</a>	crq	Genetic and molecular evidence exists for crq involvement in the phagocytosis of bacteria	-	-	<a href="#">Franc 1996</a> ; <a href="#">Franc 1999</a>
CG31741	<a href="#">CG31741</a>	crq-like	crq-like	-	-	
Santa-maria	<a href="#">CG12789</a>	santa-maria	crq-like	-	-	
peste	<a href="#">CG7228</a>	pes	Peste is required for entry of <i>Mycobacterium fortuitum</i> in S2 cells	-	-	<a href="#">Phillips 2005</a>
CG7227	<a href="#">CG7227</a>	crq-like	crq-like	-	-	
emp	<a href="#">CG2727</a>	emp	epithelial membrane protein; expressed in various embryonic tissues	-	-	
CG3829	<a href="#">CG3829</a>	emp-like	emp-like	-	-	
CG1887	<a href="#">CG1887</a>	emp-like	emp-like	-	-	
Sensory neuron membrane protein 1	<a href="#">CG7000</a>	Snmp1	more related to sensory neuron membrane protein- involved in pheromone signaling	-	-	
CG2736	<a href="#">CG2736</a>	emp-like	weakly related to emp	-	-	
BcDNA	<a href="#">CG10345</a>	CG10345	weakly related to emp	-	-	

#### Peptidoglycan recognition protein (PGRP) family : 13 members

Peptidoglycan recognition protein SA	<a href="#">CG11709</a>	PGRP-SA	secreted PGRP required for Toll activation by Gram-Positive bacteria in flies	9.5	-	<a href="#">Michel 2001</a>
PGRP-SC2	<a href="#">CG14745</a>	PGRP-SC2	PGRP with amidase activity-In vivo, may down-regulate the Imd pathway in Imd <sup>1</sup>	2.8	-	<a href="#">Bishoff 2006</a> ; <a href="#">Paredes 2011</a>
PGRP-SC1b	<a href="#">CG8577</a>	PGRP-SC1b	PGRP with amidase activity-In vivo, may down-regulate the Imd pathway in Imd <sup>1</sup>	-	-	<a href="#">Bishoff 2006</a> ; <a href="#">Paredes 2011</a> Costechareyre 2016
PGRP-SC1a	<a href="#">CG14746</a>	PGRP-SC1a	PGRP with amidase activity-In vivo, may down-regulate the Imd pathway in Imd <sup>1</sup>	-	-	<a href="#">Bishoff 2006</a> ; <a href="#">Paredes 2011</a> Costechareyre 2016
Peptidoglycan recognition protein LE	<a href="#">CG8995</a>	PGRP-LE	participates in sensing of Gram-negative bacteria upstream of the Imd pathway	-	-	<a href="#">Takehana 201</a> <a href="#">Takehana 201</a> <a href="#">Kaneko 2006</a> ; <a href="#">Yano 2008</a> ; <a href="#">Neven 2012</a> ; <a href="#">Bosco-Drayon 2012</a>
PGRP-SB1	<a href="#">CG9681</a>	PGRP-SB1	PGRP with amidase activity- Deletion of PGRP-SB1/SB2 has not phenotype	21.7	-	<a href="#">Zaidman-Remy 2011</a>
PGRP-SB2	<a href="#">CG9697</a>	PGRP-SB2	PGRP with amidase activity- Deletion of PGRP-SB1/SB2 has not phenotype	-	-	<a href="#">Zaidman-Remy 2011</a>
Peptidoglycan recognition protein LB	<a href="#">CG14704</a>	PGRP-LB	PGRP-LB down-regulate the Imd pathwayby scavenging peptidoglycan	3.7	-	<a href="#">Zaidman-Ren</a> <a href="#">Paredes 2011</a>
PGRP-SD	<a href="#">CG7496</a>	PGRP-SD	secreted PGRP required for Toll activation by DAP-type peptidoglycan	18.9	-	<a href="#">Bishchoff 2004</a>
Peptidoglycan recognition protein LC	<a href="#">CG4432</a>	PGRP-LC	Receptor of the Imd pathway, many isoform with distinct sensing and regulation	3.4	-	<a href="#">Choe 2002</a> ; <a href="#">Gottar 2002</a> ; <a href="#">Ramet 2002</a> ; <a href="#">Kaneko 2006</a> ; <a href="#">Neven 2012</a>
Peptidoglycan recognition protein LA	<a href="#">CG4384</a>	PGRP-LA	Modulate PGRP-LC sensing	-	-	<a href="#">Gendrin 2013</a>
Peptidoglycan recognition protein LF	<a href="#">CG4437</a>	PGRP-LF	Negative regulator of the Imd pathway	3.1	-	<a href="#">Maillet 2008</a>
Peptidoglycan recognition protein LD	<a href="#">CG3717</a>	PGRP-LD	related to prpg	-	-	

#### Drosophila scavenger receptor (dScR) family: 4 members

Scavenger receptor class C, type I	<a href="#">CG4099</a>	Sr-CI	dScR-I is expressed in embryonic hemocytes and binds to various ligands, including dsRNA	-	-	<a href="#">Pearson 199f</a> <a href="#">Ramet 2001</a> <a href="#">Ulvila, Parikka, et al.</a>
Scavenger receptor class C, type II	<a href="#">CG8856</a>	Sr-CII	-	-	-	
Scavenger receptor class C, type III	<a href="#">CG31962</a>	Sr-CIII	-	-	-	
Scavenger receptor class C, type IV	<a href="#">CG3212</a>	Sr-CIV	-	-	-	

#### Gram Negative Binding Protein (GNBP) family: 5 members

Gram-negative bacteria binding protein	<a href="#">CG5008</a>	GNBP3	required for Toll activation by Glucan	-	-	<a href="#">Gottar 2006</a>
Gram-negative bacteria binding protein	<a href="#">CG4144</a>	GNBP2	-	-	-	
Gram-negative bacteria binding protein	<a href="#">CG6895</a>	GNBP1	required for Toll activation by Gram positive bacteria	-	-	<a href="#">Gobert 2003</a> <a href="#">Pill-Floury 2004</a>
CG12780	<a href="#">CG12780</a>	GNBP-like	GNBP-like	2.8	-	
CG13422	<a href="#">CG13422</a>	GNBP-like	GNBP-like	13.3	5.1	

#### Nimrod/Eater/Draper family of phagocytosis receptors with EGF repeats

Nimrod A	<a href="#">CG42282</a>	NimA		-	-	<a href="#">Kunucz 2007</a>
Nimrod B1	<a href="#">CG33119</a>	NimB1		-	-	<a href="#">Kunucz 2007</a>
Nimrod B2	<a href="#">CG31839</a>	NimB2		-	-	<a href="#">Kunucz 2007</a>
Nimrod B3	<a href="#">CG34003</a>	NimB3		-	-	<a href="#">Kunucz 2007</a>
Nimrod B4	<a href="#">CG33115</a>	NimB4		-	-	<a href="#">Kunucz 2007</a>
Nimrod B5	<a href="#">CG16873</a>	NimB5		-	-	<a href="#">Kunucz 2007</a>
Nimrod C1	<a href="#">CG8942</a>	NimC1	shown to be required for phagocytosis of bacteria	-	-	<a href="#">Kunucz 2007</a>
Nimrod C2	<a href="#">CG18146</a>	NimC2		-	-	<a href="#">Kunucz 2007</a>
Nimrod C3	<a href="#">CG16880</a>	NimC3		-	-	<a href="#">Kunucz 2007</a>
Nimrod C4	<a href="#">CG16876</a>	NimC4	required for the uptake of apoptotic cells	-	-	<a href="#">Kunucz 2007</a> <a href="#">Kurant 2008</a>
draper	<a href="#">CG2086</a>	drpr	required for the uptake of apoptotic cells	-	-	<a href="#">Manaka 2004</a> <a href="#">McDonald 2006</a>
eater	<a href="#">CG6124</a>	eater	encodes a transmembrane receptor with EGF repeats; required for phagocytosis	-	-	<a href="#">Kocks 2005</a>

#### Putative chitin-binding lectins

Imaginal disc growth factor 1	<a href="#">CG4472</a>	ldgf1	chitin binding and growth factor activity in cell culture	4.1	-	
Imaginal disc growth factor 2	<a href="#">CG4475</a>	ldgf2	IDGF2 is a trophic factor promoting cellular and organismal survival	-	-	<a href="#">Broz 2017</a>
Imaginal disc growth factor 3	<a href="#">CG4559</a>	ldgf3		2.8	-	
Imaginal disc growth factor 4	<a href="#">CG1780</a>	ldgf4		-	-	

#### Galactins

Galactose-specific C-type lectin	<a href="#">CG9976</a>	Lectin-galC1	induced by infection in larvae	-	-	<a href="#">Tani et al., 2006</a>
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### II. Phagocytosis

### Hemocyte receptors that have been linked to phagocytosis

* <b>Nimrod C1</b>	<a href="#">CG8942</a>	<b>NimC1</b>	shown to be required for phagocytosis of beads and together with Eater of Gram positive bacteria			<a href="#">Kurucz 2007</a> <a href="#">Melcarne 2019</a>
* <b>draper</b>	<a href="#">CG2086</a>	<b>drpr</b>	required for the uptake of apoptotic cells			<a href="#">Manaka 2004</a>
* <b>eater</b>	<a href="#">CG6124</a>	<b>eater</b>	encodes a transmembrane receptor with EGF repeats; required for phagocytosis	-		<a href="#">Kocks 2005</a>
* <b>Scavenger receptor class C, type I</b>	<a href="#">CG4099</a>	<b>Sr-CI</b>	dsrC-I is expressed in embryonic hemocytes and binds to various ligands, including dsRNA	-		<a href="#">Pearson 1998</a> <a href="#">Ramet 2001</a> <a href="#">Ulvila, Parikka, et al.</a>
* <b>croquemort</b>	<a href="#">CG4280</a>	<b>crq</b>	Genetic and molecular evidence exists for crq involvement in the phagocytosis	-		<a href="#">Franc 1996</a> <a href="#">Franc 1999</a>
* <b>peste</b>	<a href="#">CG7228</a>	<b>pes</b>	Peste is required for entry of <i>Mycobacterium fortuitum</i> in S2 cells	-		<a href="#">Philips 2005</a>
* <b>Down syndrome cell adhesion molecule</b>	<a href="#">CG17800</a>	<b>Dscam</b>	The Dscam locus encodes a large repertoire of Immunoglobulin domain proteins, some being secreted; promotes			<a href="#">Watson FL Science 2005</a>

### Complement-like Proteins (Thiol Ester Proteins)

#### Secreted proteins that may play a role in the opsonization of microorganisms

<b>Thioester-containing protein 1</b>	<a href="#">CG18096</a>	<b>Tep1</b>	induced in larvae and adults	-		<a href="#">Lageux 2000</a> <a href="#">Bou Acoun 2010</a>
<b>Thioester-containing protein 2</b>	<a href="#">CG7052</a>	<b>Tep2</b>	induced in larvae and adults; required in cell culture for efficient phagocytosis	7.9	3.1	<a href="#">Lageux 2000</a> <a href="#">Bou Acoun 2010</a>
<b>Thioester-containing protein 3</b>	<a href="#">CG7088</a>	<b>Tep3</b>	required in cell culture for efficient phagocytosis of <i>S. aureus</i>	-		<a href="#">Lageux 2000</a> <a href="#">Bou Acoun 2010</a>
<b>Thioester-containing protein 4</b>	<a href="#">CG10363</a>	<b>Tep4</b>	induced in larvae and adults	3.4	2.5	<a href="#">Lageux 2000</a> <a href="#">Bou Acoun 2010</a>
<b>Thioester-containing protein 5</b>	<a href="#">CG13079</a>	<b>Tep5</b>				
<b>Macroglobulin complement-related</b>	<a href="#">CG7586</a>	<b>Mcr</b>	required in cell culture for efficient phagocytosis of <i>C. albicans</i>			<a href="#">Stroschein-Stevenson 2006</a>

### Complement-binding receptor

<b>Lipophorin receptor 2</b>	<a href="#">CG4823</a>	<b>LpR2</b>	putative alpha2M-receptor-like	4.5	2.1	
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### Other genes involved in phagocytosis

<b>Rac2</b>		<b>Rac2</b>	rac2 is required for efficient phagocytosis			<a href="#">Avel-Rochex, Williams 2005</a>
<b>pretaporter</b>	<a href="#">CG8556</a>	<b>prtp</b>	Involved in phagocytosis			<a href="#">Kuraishi et al., 2009</a>
<b>retinophillin</b>	<a href="#">CG1837</a>	<b>rtp</b>				<a href="#">Cuttelli 2008</a>
<b>pallbearer</b>	<a href="#">CG10233</a>	<b>pall</b>	Pallbearer functions in an SCF-dependent manner for ubiquitylation and proteasomal degradation in phagocytosis			<a href="#">Sylva 2007</a>
	<a href="#">CG3428</a>					

### Cytoskeleton component encoding genes

<b>TM9SF4</b>	<a href="#">CG7364</a>	<b>TM9SF4</b>	required for optimal phagocytosis of Gram-negative bacteria			<a href="#">Bergeret 2008</a>
<b>scab</b>	<a href="#">CG8095</a>	<b>scb</b>	A loss of alphaPS3 integrin decreased phagocytosis of <i>Staphylococcus aureus</i>			<a href="#">Nonaka 2013</a>
<b>Integrin betanu subunit</b>	<a href="#">CG1762</a>	<b>Itgbetanu</b>	loss of integrin beta reduced the phagocytosis of <i>S. aureus</i>			<a href="#">Shiratsuchi 2012</a>
<b>SCAR</b>	<a href="#">CG4636</a>	<b>SCAR</b>	required for phagocytosis			<a href="#">Nonaka 2013</a>

### Lysosomal enzymes

<b>Ectoderm-expressed 3</b>	<a href="#">CG3132</a>	<b>Ect3</b>	-	4.7	-	
<b>CG11459</b>	<a href="#">CG11459</a>	<b>CG11459</b>	encodes a Cathepsin L	6.5	3.7	
<b>phagocyte signaling impaired</b>	<a href="#">CG4845</a>	<b>psidin</b>	psidin encodes a lysosomal protein required in the blood cells for both degradation of engulfed bacteria and actin			<a href="#">Brennan 2007</a> <a href="#">Recruitment of sessile haemocytes upon immune challenge</a>
<b>Deoxyribonuclease II</b>	<a href="#">CG7780</a>	<b>DNaseII</b>	Dnase II is an enzyme involved in the degradation of DNA within phagolysosomes	4.9	2.2	

### III. Encapsulation: Genes linked to encapsulation

<b>lectin-24A</b>	<a href="#">CG3410</a>	<b>lectin-24A</b>	induced upon encapsulation			<a href="#">Keebaugh 2012</a>
<b>lectin-37Da</b>	<a href="#">CG33532</a>	<b>lectin-37Da</b>	lectin-galC1 like protein that enhance encapsulation	-		<a href="#">Ao, J 2007</a>
<b>lectin-37Db</b>	<a href="#">CG33533</a>	<b>lectin-37Db</b>	lectin-galC1 like protein that enhance encapsulation			<a href="#">Ao, J 2007</a>
<b>UDP-GlcNAc:alpha-3-D-mannoside-beta-1,2</b>	<a href="#">CG13431</a>	<b>Mgat1</b>	encodes a component of the N-Glycosylation machine involved in capsule formation			<a href="#">Mortimer 2012</a>
<b>Hemeses</b>	<a href="#">CG31770</a>	<b>He</b>	encodes a transmembrane glycoprotein-like that modulates the response to peptidoglycan			

### IV. Cytoskeleton genes involved in hemocytes function (From Fauvarques and Williams JCS 2011)

<b>Membrane proteins</b>						
<b>myspheroid</b>	<a href="#">CG1560</a>	<b>mys</b>	a beta-Integrin involved in adhesion and encapsulation			<a href="#">Irving et al., 2005</a>
<b>Neuroglian</b>	<a href="#">CG1634</a>	<b>Nrg</b>	a L1-CAM involved in adhesion and encapsulation			<a href="#">Nardi et al., 2005</a> <a href="#">Williams, 2009</a>
<b>TM9SF4</b>	<a href="#">CG7364</a>	<b>TM9SF4</b>	required for optimal phagocytosis of Gram-negative bacteria			<a href="#">Bergeret 2008</a>
<b>Integrin betanu subunit</b>	<a href="#">CG1762</a>	<b>Itgbetanu</b>	loss of integrin beta reduced the phagocytosis of <i>S. aureus</i>			<a href="#">Shiratsuchi 2012</a>
<b>scab</b>	<a href="#">CG8095</a>	<b>scb</b>	A loss of alphaPS3 integrin decreased phagocytosis of <i>Staphylococcus aureus</i>			<a href="#">Nonaka 2013</a>
<b>RhoGTPases</b>						
<b>Rac1</b>	<a href="#">CG2248</a>	<b>Rac1</b>	Rac1 and Bsk are both required for the proper encapsulation of eggs from the parasitoid wasp, <i>Recruitment of sessile haemocytes upon immune challenge</i>			<a href="#">Williams 2001</a> <a href="#">Williams 2007</a>
<b>Rac2</b>	<a href="#">CG8556</a>	<b>Rac2</b>	rac2 is required for efficient phagocytosis and encapsulation			<a href="#">Avel-Rochex, Williams 2005</a>
<b>Cdc42</b>	<a href="#">CG12530</a>	<b>Cdc42</b>	Cdc42r is required for efficient phagocytosis and encapsulation			<a href="#">Samson CJ 2003</a> <a href="#">Rogers 2003</a> <a href="#">Stramer 2005</a>
<b>Rho1</b>	<a href="#">CG8416</a>	<b>Rho1</b>	Tail retraction, Dynamic of cell cell contact			<a href="#">Stramer et al., 2005</a>
<b>Other cytoskeleton regulatory molecules</b>						
<b>SCAR</b>	<a href="#">CG4636</a>	<b>SCAR</b>	required for phagocytosis and lamellipodia			<a href="#">Pearson et al, Rogers et al., 2003</a>
<b>Zizimin-related</b>	<a href="#">CG11376</a>	<b>Zir</b>	zir is required for efficient phagocytosis and encapsulation			<a href="#">Samson CJ 2012</a>
<b>Actin-related protein 2</b>	<a href="#">CG9901</a>	<b>Arp2</b>	Phagocytosis, lamellipodia			<a href="#">Pearson 2000</a> <a href="#">Rogers 2003</a>
<b>Actin-related protein 3</b>	<a href="#">CG7558</a>	<b>Arp3</b>	Phagocytosis, lamellipodia			<a href="#">Pearson 2000</a> <a href="#">Rogers 2003</a>
<b>chickadee</b>	<a href="#">CG9553</a>	<b>chic</b>	A cofilin involved in lamellipodia, Spreading and phagocytosis			<a href="#">Pearson 2000</a> <a href="#">Rogers 2003</a>
<b>diaphanous</b>	<a href="#">CG1768</a>	<b>dia</b>	Filopodia, Rho-dependant signalling			<a href="#">Williams 2007</a>
<b>Signaling molecules</b>						
<b>Rho1</b>	<a href="#">CG8416</a>	<b>Rho1</b>	Rac1 activation, actin stress fibers			<a href="#">Williams 2007</a>
<b>slingshot</b>	<a href="#">CG6238</a>	<b>ssh</b>	Lamellipodia formation, F-actin distribution			<a href="#">Rogers 2003</a>

### V. Signaling

#### Genes encoding canonic components of the Toll signaling pathway

These genes encode components of the Toll pathway and are required for the expression of a subset of antimicrobial peptide genes (Drs, Cec, Def, AttA). Some of these genes (Toll, pII, tub, cact) are also required during hemocyte proliferation.

#### Cellular components

<b>spatzle</b>	<a href="#">CG6134</a>	<b>spz</b>	ligand of the Toll receptor	3.9	-	<a href="#">Lemaitre 1996</a>
<b>Toll</b>	<a href="#">CG5490</a>	<b>TI</b>	encodes a transmembrane receptor	3.1	-	<a href="#">Lemaitre 1996</a>
<b>Myd88</b>	<a href="#">CG2078</a>	<b>Myd88</b>	encodes an adaptor protein that functions downstream of Toll	-	-	<a href="#">Lemaitre 1996</a>
<b>tube</b>	<a href="#">CG10520</a>	<b>tub</b>	encodes an adaptor protein that functions downstream of Toll	-	-	<a href="#">Lemaitre 1996</a>
<b>pelle</b>	<a href="#">CG5974</a>	<b>pII</b>	encodes a kinase that functions downstream of Toll	2.9	-	<a href="#">Lemaitre 1996</a>
<b>cactus</b>	<a href="#">CG5848</a>	<b>cact</b>	encodes an IκB-like protein that functions downstream of Toll	7.8	2.4	<a href="#">Lemaitre 1996</a>
<b>Dorsal-related immunity factor</b>	<a href="#">CG6794</a>	<b>Dif</b>	DIF (Dorsal-related Immune Factor) encodes an NF-κB-like transcription factor	2.7	2	<a href="#">Rutschmann, Manfrulli 199 Meng 1999 Ip 1993 Matova 2006 Matova 2011 Pal 2008</a>
<b>dorsal</b>	<a href="#">CG6667</a>	<b>dl</b>	encodes an NF-κB-like transcription factor regulated by the Toll pathway	2.9	-	<a href="#">Lemaitre 1996 Han 1999 Pal 2008 Matova 2006 Matova 2011</a>

#### Pattern recognition receptor

* <b>Peptidoglycan recognition protein SA</b>	<a href="#">CG11709</a>	<b>PGRP-SA</b>		9.5	-	<a href="#">Michel 2001</a>
* <b>Gram-negative bacteria binding protein</b>	<a href="#">CG5008</a>	<b>GNBP3</b>	required for Toll activation by Glucan	-	-	<a href="#">Gottlar 2006</a>
* <b>Gram-negative bacteria binding protein</b>	<a href="#">CG6895</a>	<b>GNBP1</b>	required for Toll activation by Gram positive bacteria	-	-	<a href="#">Gobert 2003 Pili-Floury 2004</a>

#### extracellular components: the protease activity sensing branch

<b>Spatzle-Processing Enzyme</b>	<a href="#">CG16705</a>	<b>SPE</b>	In vivo RNAi and biochemical studies indicate a role of this Clip domain SP	4.8	3.3	<a href="#">Jang 2006 Kambris 2006</a>
<b>Serpin 42Dd</b>	<a href="#">CG9456</a>	<b>Spn42Dd</b>	Regulates the GNBPs-Dependent Toll Signaling Pathway	-	-	<a href="#">Fullaondo et al. 2011</a>
<b>modular serine protease</b>	<a href="#">CG31217</a>	<b>modSP</b>	ModSP is an apical serine protease that iterates signals from pattern-recognition receptors (PGRP-SA/GNBP1)	-	-	<a href="#">Buchon 2009</a>
<b>Gram-positive Specific Serine protease</b>	<a href="#">CG5896</a>	<b>grass</b>	A clip serine protease that functions upstream of SPE and downstream of ModSP	-	-	<a href="#">Kambris 2006 El Chamy 2008</a>
<b>nerotic</b>	<a href="#">CG1857</a>	<b>nec</b>	belongs to the <i>nerotic</i> cluster and encodes a protease that functions upstream	5.9	3.4	<a href="#">Levashina 1999</a>
<b>persephone</b>	<a href="#">CG6367</a>	<b>psh</b>	encodes a Clip domain serine protease required for Toll activation by entomopathogenic fungi	-	-	<a href="#">Liqoxyqakis 2002</a>
<b>Hayan</b>	<a href="#">CG6361</a>	<b>Hayan</b>	Hayan and Psh redundantly regulate SPE (Toll activation) and Melanization	-	-	<a href="#">Nam et al. 2011 Dudzic 2019</a>

#### Other Serine Proteases/Serpins potentially acting upstream of the Toll pathway

<b>Serine Protease Immune Response Inte spheroid</b>	<a href="#">CG2056</a>	<b>spirit</b>	An in vivo RNAi study indicates a role of this Clip-domain SP in the activatio	6.7	-	<a href="#">Kambris 2006</a>
	<a href="#">CG9675</a>	<b>spheroid</b>	An in vivo RNAi study indicates a role of this serine protease homolof (SPH) in the activation of Spatzle during	-	-	<a href="#">Kambris 2006</a>
	<a href="http://flybase.bio.indiana.edu/reports/FBc">http://flybase.bio.indiana.edu/reports/FBc</a>		An in vivo RNAi study indicates a role of this serine protease homolof (SPH) in the activation of Spatzle during	-	-	<a href="#">Kambris 2006</a>
* <b>Serpin 88Ea</b>	<a href="#">CG18525</a>	<b>Spn88Ea</b>	serpin-like that negatively regulate the Toll pathway and melanization	3.4	-	<a href="#">Ahmad 2005</a>

#### Genes encoding canonic components of the Imd pathway

These genes encode proteins required for the expression of antibacterial peptide genes after infection.

<b>immune deficiency</b>	<a href="#">CG5576</a>	<b>imd</b>	protein with homology to the mammalian RIP adaptor	2.1	-	<a href="#">Lemaitre 1996 Georgel 2001</a>
<b>TGF-beta activated kinase 1</b>	<a href="#">CG1388</a>	<b>Tak1</b>	encodes a homology of the mammalian MAPKKK TAK1	-	-	<a href="#">Vidal 2001</a>
<b>TAK1-associated binding protein 2</b>	<a href="#">CG7417</a>	<b>Tab2</b>	encodes a homology of the mammalian TAK1 binding protein	-	-	<a href="#">Kleino 2005 Gesellchen 2006 Zhuang 2006</a>
<b>Inhibitor of apoptosis 2</b>	<a href="#">CG8293</a>	<b>lap2</b>	encodes a homology of Inhibitor of apoptosis	-	-	<a href="#">Kleino 2005 Leulier 2006 Gesellchen 2005</a>
<b>Fas-associated death domain ortholog</b>	<a href="#">CG12297</a>	<b>Fadd</b>	homolog of mammalian FADD gene	-	-	<a href="#">Naitza 2002, Leulier 2002</a>
<b>Death related ced-3/Nedd2-like protein</b>	<a href="#">CG7486</a>	<b>Dredd</b>	encodes a homology of <b>Caspase-8</b>	-	-	<a href="#">Leulier 2000</a>
<b>kenny</b>	<a href="#">CG16910</a>	<b>key</b>	encodes a homology of mammalian ikkγ, structural component of the IKK complex	-	-	<a href="#">Rutschmann, Silverman and Maniatis, 2001</a>
<b>Immune response deficient 5</b>	<a href="#">CG4201</a>	<b>ird5</b>	encodes a homology of mammalian ikkβ, kinase component of the IKK complex	-	-	<a href="#">Lu et al., 2000 Silverman and Maniatis, 2001</a>
<b>Relish</b>	<a href="#">CG11992</a>	<b>Rel</b>	encodes a P105-like transcription factor	17.2	2.1	<a href="#">Henderson 1996 Dushay 1996</a>
* <b>Peptidoglycan recognition protein LC</b>	<a href="#">CG4432</a>	<b>PGRP-LC</b>	Receptor of the Imd pathway, many isoform with distinct sensing and regula	3.4	-	<a href="#">Choe 2002, Gottar 2002, Ramet 2002, Kaneko 2006</a>
* <b>Peptidoglycan recognition protein LA</b>	<a href="#">CG4384</a>	<b>PGRP-LA</b>	Modulate PGRP-LC sensing	-	-	<a href="#">Gendrin 2013</a>
* <b>Peptidoglycan recognition protein LE</b>	<a href="#">CG8995</a>	<b>PGRP-LE</b>	participates in sensing of Gram-negative bacteria upstream of the Imd pathw	-	-	<a href="#">Takehana 2006 Takehana 2006 Kaneko 2006 Yano 2008, Neven 2012, Bosco-Drayon 2012</a>
<b>negative regulator</b>				-	-	
<b>poor Imd response upon knock-in</b>	<a href="#">CG15678</a>	<b>pirk</b>	Negative regulator of the Imd pathway	3.8	-	<a href="#">Kleino 2008, Lhocine 2008 Aggarwal 2008</a>
<b>Peptidoglycan recognition protein LF</b>	<a href="#">CG4437</a>	<b>PGRP-LF</b>	Negative regulator of the Imd pathway	3.1	-	<a href="#">Maillet 2008</a>

#### Genes encoding proteins that affect the Toll and/or the Imd pathway(s)

<b>skpA</b>	<a href="#">CG16983</a>	<b>skpA</b>	This gene encodes a component of the SCF complex. High Diptericin expression is observed in a skpA mutant.	-	-	<a href="#">Khush 2002</a>
<b>bendless</b>	<a href="#">CG18319</a>	<b>ben</b>	RNAi in S2 cells indicates a role of Bendless in IKK activation / <b>Bendless mutant flies show reduced inducti</b>	-	-	<a href="#">Zhou 2005</a>
<b>Uev1A</b>	<a href="#">CG10640</a>	<b>Uev1A</b>	RNAi in S2 cells indicates a role of UEV1a in IKK activation	-	-	<a href="#">Zhou 2005</a>
<b>Helicase 89B</b>	<a href="#">CG4261</a>	<b>Hel89B</b>	required downstream of NF-κB for the activation of AMP genes	-	-	<a href="#">Yaci 2005</a>
<b>Plenty of SH3s</b>	<a href="#">CG4909</a>	<b>POSH</b>	Posh is a Ring-finger protein. JNK activation and Relish induction are delayed and sustained in POSH deficient	-	-	<a href="#">Tsuda 2005</a>
<b>caspar</b>	<a href="#">CG8400</a>	<b>casp</b>	encodes an homolog of the Fas-associating factor 1. Caspar represses the Imd pathway	-	-	<a href="#">Kim 2006</a>
<b>defense repressor 1</b>	<a href="#">CG12489</a>	<b>dnr1</b>	RING finger containing molecule. RNAi in S2 cells indicates an inhibitory role of DNR-1 on Dredd activity	-	-	<a href="#">Foley 2004</a>
<b>wnt inhibitor of Dorsal</b>	<a href="#">CG8458</a>	<b>wntD</b>	Negative regulator of the Toll pathway	-	-	<a href="#">Gordon et al., 2005</a>
<b>cylindromatosis</b>	<a href="#">CG5603</a>	<b>CYLD</b>	encodes a deubiquitylating enzyme that is a negative regulator of the NF-κB	-	-	<a href="#">Tschirritzis 2007</a>
<b>scrawny</b>	<a href="#">CG5505</a>	<b>scny</b>	negative regulator of the Imd pathway	-	-	<a href="#">Thevenon 2009</a>
<b>Ring and YY1 Binding Protein</b>	<a href="#">CG12190</a>	<b>RYBP</b>	<a href="#">dRYBP Contributes to the Negative Regulation of the Drosophila Imd Pathway.</a>	-	-	<a href="#">Aparicio 2013</a>
<b>Zn finger homeodomain 1</b>	<a href="#">CG1322</a>	<b>zfh1</b>	<a href="#">Transcription factor zfh1 downregulates Drosophila Imd pathway.</a>	-	-	<a href="#">Myllymäki H Wu 2007</a>
<b>immune response deficient 1</b>	<a href="#">CG9746</a>	<b>ird1</b>	<a href="#">Ird1 is a Vps15 homologue important for antibacterial immune responses in Drosophila</a>	-	-	<a href="#">Valanne 2010</a>
<b>G protein-coupled receptor kinase 2</b>	<a href="#">CG17998</a>	<b>Gprk2</b>	<a href="#">affect Toll signaling</a>	-	-	<a href="#">Haghayeghi Z Ji 2014</a>
<b>Pellino</b>	<a href="#">CG5212</a>	<b>Pli</b>	Pli interacts with the Pelle kinase - negative regulator of Toll signaling accord	-	-	<a href="#">Reard 2008 Kuitkenleuler 2009</a>
<b>Deformed epidermal autoregulatory fac</b>	<a href="#">CG9567</a>	<b>Def1</b>	<a href="#">regulates antimicrobial peptide gene expression</a>	-	-	<a href="#">Chiu 2005</a>
<b>lesswright</b>	<a href="#">CG3018</a>	<b>lwr</b>	Ubr9 negatively regulates the expression of the antifungal peptide gene drosomycin	-	-	<a href="#">Huang 2005</a>
<b>lesswright</b>	<a href="#">CG3018</a>	<b>lwr</b>	encodes an enzyme that conjugates a small ubiquitin-related modifier (SUMO). Mutations causes activation of T	-	-	<a href="#">Dilan 2014</a>
<b>trabid</b>	<a href="#">CG9448</a>	<b>trbd</b>	A deubiquitinase that interacts with TAK1 and negatively regulate the Imd pathway	-	-	

Transcription (Serpent, Caudal, Drifter...) and chromatin factors (Akirin) that pattern antimicrobial peptide genes in various tissues are not included (Moleskin)

Genes encoding components involved in the ecdysone response and vesicle trafficking (deep orange) are not included although they have a strong impact on Imd pathway and antimicrobial peptide expression

#### Genes encoding homologs of mammalian components of the IL1-R/TLR or TNF signalling pathways but has not yet been implicated in the immune response

##### Toll-like (TL) receptors

<b>Tehao</b>	<a href="#">CG7121</a>	<b>Tehao</b>	TL-5 expression induces the drosomycin antifungal peptide gene in cell cultur	-	-	
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Tollo	<a href="#">CG6890</a>	Tollo	oli-8/Tollo negatively regulates antimicrobial response in the Drosophila resp -	-	<a href="#">Akhouayri 2011</a>
Toll-6	<a href="#">CG7250</a>	Toll-6	-	-	
Toll-7	<a href="#">CG8595</a>	Toll-7	Toll-7 would function as a pattern recognition receptor for some virus	-	<a href="#">Nakamoto 2012</a>
18 wheeler	<a href="#">CG8896</a>	18w	No role of 18W in the Drosophila immune response	-	<a href="#">Iqoxyzgakis 2002</a>
Toll-4	<a href="#">CG18241</a>	Toll-4	-	-	
MstProx	<a href="#">CG1149</a>	MstProx	-	-	
Toll-9	<a href="#">CG5528</a>	Toll-9	Toll-9 is neither required to maintain a basal anti-microbial response nor to n-	-	<a href="#">Narbonne-Reveau 2011</a>

#### Others

IkappaB kinase-like 2	<a href="#">CG2615</a>	ik2	homolog of the mammalian TANK/IKKe gene	-	
cactin	<a href="#">CG1676</a>	cactin	encodes a conserved protein that interacts with cactus and modulates its act-	-	
ECSIT	<a href="#">CG10610</a>	ECSIT	homolog of the mammalian ECSIT gene	-	
TNF-receptor-associated factor 4	<a href="#">CG3048</a>	Traf4	homolog of the mammalian TRAF4 and 2 genes	-	<a href="#">Cha 2003</a>
CG4393	<a href="#">CG4393</a>	CG4393	weak homolog of the mammalian TRAF3 gene	-	
	<a href="#">CG10961</a>	0	homolog of the mammalian TRAF6 gene	-	<a href="#">Cha 2003</a>
NFAT homolog	<a href="#">CG11172</a>	NFAT	homology Rel domain transcription factor NFAT5?	-	

#### Genes encoding components of the JAK/STAT pathway

Some components of the Drosophila JAK/STAT pathway are involved in hemocyte proliferation and differentiation and in the expression of Turandot and Tep genes in the fat body.

This pathway is induced upon damage where it could play a role in the systemic adaptation to injury. It also participate in the gut repair response.

outstretched	<a href="#">CG5993</a>	os	ligand of Domeless, role in development and stem cell maintenance	-	<a href="#">Harrison 1998</a>
unpaired 2	<a href="#">CG5988</a>	upd2	ligand of Domeless, induced by infection in the gut	-	<a href="#">Agaïsse 2003</a> <a href="#">Osman 2012</a>
unpaired 3	<a href="#">CG33542</a>	upd3	ligand of Domeless. Produced by hemocyte and involved in the JAK-STAT activation in the fat-body.	-	<a href="#">Agaïsse 2003</a> <a href="#">Osman 2012</a>
domeless	<a href="#">CG14226</a>	dome	receptor of JAK-STAT Pathway	-	
Signal-transducer and activator of trans	<a href="#">CG4257</a>	Stat92E	homolog of the mammalian STAT gene	-	
hopscotch	<a href="#">CG1594</a>	hop	encodes the <i>Drosophila</i> homolog of mammalian JAK	-	
Suppressor of cytokine signaling at	<a href="#">CG15154</a>	Socs36E	Negative regulator of the JAK-STAT pathway	-	
Diedel	<a href="#">CG11501</a>	Diedel	putative negative regulator of the JAK-STAT pathway, found to be induced in S2	-	<a href="#">Boutros 2002</a> <a href="#">Muller 2005</a> <a href="#">Coste 2012</a>

#### Genes encoding components of the MAPK pathways

Some components of the MAPK pathways have been implicated in *Drosophila* antimicrobial responses.

Mekk1	<a href="#">CG7717</a>	Mekk1	regulates Turandot stress genes/regulate Duox activation	-	<a href="#">Inoue 2000</a> <a href="#">Brun 2006</a>
p38a	<a href="#">CG5475</a>	Mpk2	Overexpression of Mpk2 attenuates the expression of antimicrobial peptide g-	-	<a href="#">chen 2010</a>
p38b	<a href="#">CG7393</a>	p38b	participate in the stress response in the gut and phagocytic encapsulation	-	<a href="#">Shinzawa N ; chen 2010</a>
p38c	<a href="#">CG33338</a>	p38c	Regulate dopadecarboxylase at wound site	-	<a href="#">Davis2008</a>
MAP kinase kinase 4	<a href="#">CG9738</a>	Mkk4		-	<a href="#">Geuking 2010</a>
licorne	<a href="#">CG12244</a>	lic		-	<a href="#">Ha 2009</a>

#### Genes encoding components of the JNK pathway

The JNK pathway have been implicated in wound-healing of epidermis. It is activated downstream of the lmd pathway at the level of TAK1 and regulates proteins involved in cytoskeleton remodeling.

hemipterous	<a href="#">CG4353</a>	hep	Hemipterous encodes a MAPKKK required for JNK activation.	-	<a href="#">Boutros 2002</a> <a href="#">Delaney 2006</a> <a href="#">Kim 2006</a> <a href="#">Kallio 2005</a>
basket	<a href="#">CG5680</a>	bsk	basket encodes a MAPK activated by immune challenge in cultured cells.	-	
Jun-related antigen	<a href="#">CG2275</a>	Jra	-	3.8	
puckered	<a href="#">CG7850</a>	puc	negative regulator of the JNK pathway	2.5	

#### Genes encoding components of the Eiger/Wengen pathway

Eiger and Wengen are *Drosophila* proteins with similarities to mammalian TNF and TNF-R. There is contradictory result concerning the role of this pathway in immunity

eiger	<a href="#">CG12919</a>	egr	encodes a homologue of TNF. Enhances lethal effects of Salmonella infection	-	<a href="#">Schneider 201</a> <a href="#">Mabery 2010</a> <a href="#">Brandt 2004</a> <a href="#">Geuking 2009</a>
Grindelwald	<a href="#">CG10176</a>	grnd	encode the receptor of Eiger	-	<a href="#">Andersen 2005</a>

#### VI. Melanization

##### proPOphenoloxidase family: 3 members

PP01	<a href="#">CG42639</a>	proPO-A1	expressed in crystal cells	-	<a href="#">Rizki 1980</a> <a href="#">Bingqeli 2014</a>
PP02	<a href="#">CG8193</a>	proPO45	expressed in crystal cells	-	<a href="#">Bingqeli 2014</a>
PP03	CG42640	proPO59	expressed in lamellocytes	-	<a href="#">Nam 2008</a> <a href="#">Dudzic 2017</a>

##### Serine Proteases/Serpins

Serpin 27A	<a href="#">CG11331</a>	Spn27A	Serpin-like	6.5	2.1 <a href="#">Deagregorio 201</a> <a href="#">Iqoxyzgakis 2002</a>
Serine protease 7	<a href="#">CG3066</a>	Sp7	required for proPo activation in the hemolymph	5.5	<a href="#">Tang et al., 2011</a> <a href="#">Castillejo-Log Leclerc et al., 2006</a>
Melanization Protease 1	<a href="#">CG1102</a>	MP1	In vivo RNA indicates a role of MP1 for proPo activation in the hemolymph	3.9	<a href="#">Tang et al., 2006</a>
Serpin 28Dc	<a href="#">CG7219</a>	Spn28Dc	serpin-like	21.2	<a href="#">Scherfer 2008</a>
Serpin 77Ba	<a href="#">CG6680</a>	Spn77Ba	negatively regulate melanization in the trachea		<a href="#">Tang et al., 2008</a>
Hayan	<a href="#">CG6361</a>	Hayan	Terminal serine protease cleaving PPO in PO		<a href="#">Nam et al., 2012</a>

##### Other melanization factors

The list includes only the immune-inducibile genes encoding enzymes or structural components putatively involved in the melanization reaction.

Dopa decarboxylase	<a href="#">CG10697</a>	Ddc	Dopa decarboxylase	11.1	-
pale	<a href="#">CG10118</a>	ple	tyrosine hydroxylase	2.4	2.1
Punch	<a href="#">CG9441</a>	Pu	GTP cyclohydrolase	4.7	-
Dihydropteridine reductase	<a href="#">CG4665</a>	Dhpr	dihydropteridine reductase	3.2	-
yellow-f	<a href="#">CG18550</a>	yellow-f	dopachrome conversion enzyme	5.9	2
Chorion protein 19	<a href="#">CG6524</a>	Cp19	structural protein of the chorion	3.4	2.1
Multicopper oxidase-1	<a href="#">CG3759</a>	Mco1	laccase-like	2.2	-
Odorant-binding protein 28a	CG6641	Obp28a	Required for melanization		<a href="#">Benoit 2017</a>

## VII. Coagulation

<b>Hemolectin</b>	<a href="#">CG7002</a>	<b>Hml</b>	encodes a homolog of silkworm hemocytin and contains domains similar to mammalian von Willebrand factor. <a href="#">Goto 2001</a> <a href="#">Lesch 2007</a>		
<b>fondue</b>	<a href="#">CG15825</a>	<b>fon</b>	Silencing of Fondue by in vivo RNAi induces clotting defects in larvae.	4.4	- <a href="#">Scherfer 2006</a>
<b>Transglutaminase</b>	<a href="#">CG7356</a>	<b>Tg</b>	transglutaminase, similar to vertebrate clotting factor XIII and clotting factors in crustaceans and horseshoe crabs, involved in clot cross-linking		
<b>Edysone-induced gene 71Ee</b>	<a href="#">CG7604</a>	<b>Eig71Ee</b>	synonym I71-7; encodes a mucin required for bacteria entrapment in the clot		

### Putative coagulation factors

The list includes only the immune inducible genes encoding components putatively involved in the clotting reaction.

<b>CG5550</b>	<a href="#">CG5550</a>	<b>CG5550</b>	Fibrinogen-like	9.5	-
<b>Annexin B9</b>	<a href="#">CG5730</a>	<b>AnxB9</b>	Annexin V-like	2.5	-
<b>Retinoid- and fatty acid-binding glycoprotein</b>	<a href="#">CG11064</a>	<b>Rfabg</b>	Lipophorin; homologous to clotting proteins of other insects		
<b>Cuticular protein 49Ac</b>	<a href="#">CG8502</a>	<b>Cpr49Ac</b>	mucin domain & constituent of larval cuticle; <i>Drosophila</i> clot component identified by proteomics		
<b>CG11313</b>	<a href="#">CG11313</a>	<b>CG11313</b>	similar to PO-activating protease; <i>Drosophila</i> clot component identified by proteomics		
<b>Geisolin</b>	<a href="#">CG1106</a>	<b>Gel</b>	geisolin; structural cytoskeleton component; <i>Drosophila</i> clot component identified by proteomics		

## VIII. Hematopoiesis and cellular response (Partial)

### Hematopoiesis

<b>lozenge</b>	<a href="#">CG1689</a>	<b>Iz</b>	encodes an AML1 like transcription factor required for crystal cell differentiation	-	
<b>serpent</b>	<a href="#">CG3992</a>	<b>srp</b>	GATA factor that regulates embryonic blood cell development and cecropin I	-	
<b>glial cells missing</b>	<a href="#">CG12245</a>	<b>gcm</b>	encodes a DNA binding protein involved in plasmatocyte differentiation	-	
<b>gcm2</b>	<a href="#">CG3858</a>	<b>gcm2</b>	participates with gcm in hemocyte differentiation	-	
<b>u-shaped</b>	<a href="#">CG2762</a>	<b>ush</b>	U-shaped encodes a homolog of Friend of GATA that functions as a negative regulator of GATA	-	
<b>Notch</b>	<a href="#">CG3936</a>	<b>N</b>	involved in the specification of crystal cells	-	
<b>Serrate</b>	<a href="#">CG6127</a>	<b>Ser</b>	encodes a ligand of Notch; required for crystal cell specification	-	
<b>domino</b>	<a href="#">CG9696</a>	<b>dom</b>	domino is a chromatin factor required for cell proliferation. Homozygous mutant is a proliferation disrupter	-	
<b>proliferation disrupter</b>	<a href="#">CG18608</a>	<b>prod</b>	proliferation disrupter is a chromatin factor required for cell proliferation. Homozygous mutant is a proliferation disrupter	-	
<b>knot</b>	<a href="#">CG10197</a>	<b>kn</b>	encodes an homolog of EBF; required for lamellocyte differentiation	-	
<b>yantar</b>	<a href="#">CG18428</a>	<b>ytar</b>	mutation in Yantar induces defects in hemocyte proliferation and differentiation	-	
<b>PDGF- and VEGF-receptor related</b>	<a href="#">CG8222</a>	<b>Pvr</b>	encodes PDGF-receptor. Implicated in hemocyte proliferation	-	

Other genes causing blood cell proliferation are not listed here.

## IX. Antimicrobial agents: AMPs, ROS, NO, Iron sequestration and others

### Lysozyme (lys) family: 13 members including 1 gene encoding a protein with 4 lysozyme domains; probably involved in digestion

<b>Lysozyme X</b>	<a href="#">CG9120</a>	<b>LysX</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme B</b>	<a href="#">CG11179</a>	<b>LysB</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme D</b>	<a href="#">CG9118</a>	<b>LysD</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme S</b>	<a href="#">CG1165</a>	<b>LysS</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme E</b>	<a href="#">CG1180</a>	<b>LysE</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme P</b>	<a href="#">CG9116</a>	<b>LysP</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>Lysozyme C</b>	<a href="#">CG9111</a>	<b>LysC</b>	expressed in the midgut and may aid digestion.	-	-	<a href="#">Daffre 1994</a>
<b>CG7798</b>	<a href="#">CG7798</a>	<b>CG7798</b>	-	-	-	
<b>CG8492</b>	<a href="#">CG8492</a>	<b>CG8492</b>	encodes a large protein with 4 lysozyme domains	-	-	
<b>CG11159</b>	<a href="#">CG11159</a>	<b>CG11159</b>	-	-	-	
<b>CG16799</b>	<a href="#">CG16799</a>	<b>CG16799</b>	-	-	-	
<b>CG16756</b>	<a href="#">CG16756</a>	<b>CG16756</b>	-	-	-	
<b>CG30062</b>	<a href="#">CG30062</a>	<b>CG30062</b>	-	-	-	
<b>CG6429</b>	<a href="#">CG6429</a>	<b>CG6429</b>	<a href="#">Lysozyme (3.2.1.17)</a>	9.4	-	

### Inducible Antimicrobial peptides

#### Attacins: 4 members

<b>Attacin-A</b>	<a href="#">CG10146</a>	<b>AttA</b>	encodes an antibacterial peptide and is induced by infection	18.4	4.2	<a href="#">Asling 1995</a> <a href="#">Hendengren 2000</a>
<b>Attacin-B</b>	<a href="#">CG18372</a>	<b>AttB</b>	encodes an antibacterial peptide and is induced by infection	13.2	3.1	<a href="#">Hendengren 2000</a>
<b>Cecropin A1</b>	<a href="#">CG4740</a>	<b>CecA1</b>	encodes an antibacterial peptide and is induced by infection	31.7	2.5	<a href="#">Hendengren 2000</a>
<b>Attacin-D</b>	<a href="#">CG7629</a>	<b>AttD</b>	encodes an antibacterial peptide and is induced by infection	36.5	-	<a href="#">Hendengren 2000</a>

#### Diptericin (dpt) family: 2 members

<b>Diptericin</b>	<a href="#">CG12763</a>	<b>Dpt</b>	encodes an antibacterial peptide and is induced by infection	-	-	<a href="#">Wicker 1990</a> <a href="#">Reichert 1992</a>
<b>Diptericin B</b>	<a href="#">CG10794</a>	<b>DptB</b>	encodes an antibacterial peptide and is induced by infection	16.7	-	<a href="#">Hendengren 2000</a>

#### Cecropin (cec) family: 4 member

<b>Cecropin A2</b>	<a href="#">CG1367</a>	<b>CecA2</b>	encodes a peptide with both antibacterial and antifungal peptide and is induced by infection	25.8	-	<a href="#">Kylsten 1990</a>
<b>Cecropin A1</b>	<a href="#">CG1365</a>	<b>CecA1</b>	encodes a peptide with both antibacterial and antifungal peptide and is induced by infection	10.8	-	<a href="#">Kylsten 1991</a>
<b>Cecropin C</b>	<a href="#">CG1373</a>	<b>CecC</b>	encodes a peptide with both antibacterial and antifungal peptide and is induced by infection	11.0	-	<a href="#">Kylsten 1992</a>
<b>Cecropin B</b>	<a href="#">CG1878</a>	<b>CecB</b>	encodes a peptide with both antibacterial and antifungal peptide and is induced by infection	44.5	-	<a href="#">Kylsten 1990</a>

#### Drosomycin (drs) family: 6 members

<b>Drosomycin</b>	<a href="#">CG10810</a>	<b>Drs</b>	encodes an antifungal peptide and is induced by infection	8.3	7	<a href="#">Felbaum 1994</a>
<b>Drosomycin-like 2</b>	<a href="#">CG32279</a>	<b>Drs12</b>	drs-like	-	-	
<b>Drosomycin-like 3</b>	<a href="#">CG32283</a>	<b>Drs13</b>	drs-like	-	-	
<b>Drosomycin-like 4</b>	<a href="#">CG32282</a>	<b>Drs14</b>	drs-like	-	-	
<b>Drosomycin-like 5</b>	<a href="#">CG10812</a>	<b>Drs15</b>	drs-like	3.2	3.7	
<b>Drosomycin-like 6</b>	<a href="#">CG32268</a>	<b>Drs16</b>	drs-like	-	-	

## other AMPs

<b>Drosocin</b>	<a href="#">CG10816</a>	<b>Dro</b>	encodes an antibacterial peptide and is induced by infection	16.4	2.3	<a href="#">Bulet 1993</a>
<b>Metchnikowin</b>	<a href="#">CG8175</a>	<b>Mtk</b>	encodes a peptide with both antifungal and antibacterial activity and is induced	7.4	7.1	<a href="#">Levashina 1995</a>
<b>Defensin</b>	<a href="#">CG1385</a>	<b>Def</b>	encodes an antibacterial peptide and is induced by infection	27.6	3.2	<a href="#">Dimaro 1994</a>
<b>Andropin</b>	<a href="#">CG1361</a>	<b>Anp</b>	encodes an antibacterial peptide in the male genital tract	-	-	<a href="#">Samaklovic 1991</a>
<b>Listerinin</b>	<a href="#">CG9080</a>	<b>Listericin</b>	encodes an antimicrobial peptide (regulated by lmd in Degregorio 2002 or by	12.5	3.2	<a href="#">Goto et al., 2010</a>

## Iron sequestration

<b>Transferrin 1</b>	<a href="#">CG6186</a>	<b>Tsf1</b>	induced by infection, encodes a protein that mediates iron transport	2	2.3	
<b>Transferrin 3</b>	<a href="#">CG3666</a>	<b>Tsf3</b>	-	10.8	-	
<b>Zinc/iron regulated transporter-related 1</b>	<a href="#">CG6898</a>	<b>Zip3</b>	iron-zinc transporter	3.4	-	

## ROS Production by Duox in epit

<b>Dual oxidase</b>	<a href="#">CG3131</a>	<b>Duox</b>	In vivo RNAi indicates a role of duox in the production of ROS during oral bacterial infection.			<a href="#">Ha 2005</a>
<b>Immune-regulated catalase</b>	<a href="#">CG8913</a>	<b>Irc</b>	In vivo RNAi indicates a role of IRC in ROS homeostasis during oral bacteri	9	2.9	<a href="#">Ha 2005</a>

Duox is regulated at the transcriptional level by Mekk1, P38 and ATF2  
Duox activity is regulated by Gαq and PLCβ (Ha 2009)

## Nitric oxide synthase

<a href="#">CG6713</a>	<b>Nos</b>	Has been involved in immune signaling between gut and fat body. Th	-	-	<a href="#">Foley 2003</a>	<a href="#">Chakrabarti 2012</a>
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## Other

<b>Edin</b>	<a href="#">CG32185</a>		induced in S2 cell, RNAi files showed modestly impaired resistance to E. faecalis infection			<a href="#">Vanha-Aho 2</a>	<a href="#">Gordon 2008</a>
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## X. Antiviral defense

### RNAi pathway

#### Components of the RNAi pathway provides an antiviral defense

<b>Argonaute 2</b>	<a href="#">CG7439</a>	<b>AGO2</b>				
<b>Dicer-2</b>	<a href="#">CG6493</a>	<b>Dcr-2</b>				
<b>r2d2</b>	<a href="#">CG7138</a>	<b>r2d2</b>				
<b>Toll-7</b>	<a href="#">CG8595</a>	<b>Toll-7</b>	Toll-7 would function as a pattern recognition receptor for some virus	-	-	<a href="#">Nakamoto 2012</a>
<b>Methyltransferase 2</b>	<a href="#">CG10692</a>	<b>Mt2</b>	The RNA methyltransferase Dnm2 is required for efficient innate immune responses			<a href="#">Durdevic 2013</a>

## Other

<b>Ejaculatory bulb protein III</b>	<a href="#">CG11390</a>	<b>PebIII</b>	induced at the post-transcriptional level upon <i>Drosophila</i> C Virus infection			
<b>Pherokine 3</b>	<a href="#">CG9358</a>	<b>Phk-3</b>	induced at the post-transcriptional level upon <i>Drosophila</i> C Virus infection			
<b>virus-induced RNA 1</b>	<a href="#">CG31764</a>	<b>vir-1</b>	induced by the JAK-STAT pathway upon <i>Drosophila</i> C virus infection	3	-	
<b>refractory to sigma P</b>	<a href="#">CG10360</a>	<b>ref(2)P</b>	polymorphism at the ref(2)P locus affects Sigma virus replication			
<b>Vago</b>	<a href="#">CG2081</a>	<b>Vago</b>	Vago gene product controlled viral load in the fat body after infection with wit	3.4	-	<a href="#">Deddouche 2008</a>
<b>Adenosine deaminase acting on RN</b>	<a href="#">CG12598</a>	<b>Adar</b>	ADARs is an RNA editing enzyme that target double stranded RNA and may play a role in antiviral defense			<a href="#">Carpenter 2009</a>
<b>pastrel</b>	<a href="#">CG8588</a>	<b>pst</b>	the pastrel is associated with DCV-resistance by genome association studies and RNAi			<a href="#">Magwire 2012</a>

## XI. Miscellaneous

### Stress factors

<b>Turandot A</b>	<a href="#">CG31509</a>	<b>TotA</b>	induced after various stress conditions including infection	NT	NT	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Turandot B</b>	<a href="#">CG5609</a>	<b>TotB</b>	-	-	-	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Turandot C</b>	<a href="#">CG31508</a>	<b>TotC</b>	-	NT	NT	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Turandot F</b>	<a href="#">CG31891</a>	<b>TotF</b>	-	NT	NT	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Turandot M</b>	<a href="#">CG14027</a>	<b>TotM</b>	-	-	31.2	3.6	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>
<b>Turandot X</b>	<a href="#">CG31193</a>	<b>TotX</b>	-	NT	NT	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Turandot Z</b>	<a href="#">CG31507</a>	<b>TotZ</b>	-	NT	NT	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	
<b>Victoria</b>	<a href="#">CG33117</a>	<b>Victoria</b>	-	-	-	<a href="#">Ekengren 200</a>	<a href="#">Ekengren 2001</a>	

## Other

<b>Thor</b>	<a href="#">CG8846</a>	<b>Thor</b>	induced by infection; encodes a homolog of mammalian translation initiation	2.4	-	
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## XII. Other Genes induced by infection upon systemic infection (microarray)

This list includes all the other genes selected as *Drosophila* Immune Regulated Genes described in De Gregorio et al. 2001

### Other Induced serine proteases

<b>CG30087</b>	<a href="#">CG30087</a>	<b>CG30087</b>	serine protease-like	7	6	
<b>CG6639</b>	<a href="#">CG6639</a>	<b>CG6639</b>	serine protease-like	49.8	18.7	
<b>CG9649</b>	<a href="#">CG9649</a>	<b>CG9649</b>	serine protease-like	4.6	2.5	
<b>CG18563</b>	<a href="#">CG18563</a>	<b>CG18563</b>	serine protease-like	9.9	3.1	
<b>CG11842</b>	<a href="#">CG11842</a>	<b>CG11842</b>	serine protease-like	6.4	2.1	
<b>CG11841</b>	<a href="#">CG11841</a>	<b>CG11841</b>	serine protease-like	2.8	3.9	
<b>Hayan</b>	<a href="#">CG6361</a>	<b>Hayan</b>	serine protease-like	3.2	-	

CG9631	<a href="#">CG9631</a>	CG9631	serine protease-like	3.3	-
CG15046	<a href="#">CG15046</a>	CG15046	serine protease-like	5.6	-
Jonah 65Aiv	<a href="#">CG6467</a>	Jon65Aiv	serine protease-like	2.4	2.5
CG5909	<a href="#">CG5909</a>	CG5909	serine protease-like	5.2	2.8
CG31326	<a href="#">CG31326</a>	CG31326	serine protease-like	3.1	2.5
Ser7	<a href="#">CG2045</a>	Ser7	serine protease-like	2.7	2.6
CG9733	<a href="#">CG9733</a>	CG9733	close homolog of the proPAE gene	8.1	-
CG3505	<a href="#">CG3505</a>	CG3505	close homolog of the proPAE gene	5.9	2.5
<b>Other Induced serpins</b>					
Serpin 42Da	<a href="#">CG9453</a>	Spn42Da	serpin-like	2.3	-
Serpin 88Eb	<a href="#">CG6687</a>	Spn88Eb	serpin-like	11.5	4.4
Serpin 88Ea	<a href="#">CG18525</a>	Spn88Ea	serpin-like	3.4	-
<b>Induced Kunitz ser-protease inhi</b>					
CG16713	<a href="#">CG16713</a>	CG16713	Kunitz ser-protease inhibitors	7.2	2.7
CG3604	<a href="#">CG3604</a>	CG3604	Kunitz ser-protease inhibitors	4.5	-
<b>Bomanins</b>					
Immune induced molecule 1	<a href="#">CG18108</a>	IM1	Required for survival to Gram positive bacteria and fungi	11.3	7.6
CG18107		Bom	Bom		
Immune induced molecule 2	<a href="#">CG18106</a>	IM2	Bom	6.9	4.1
CG15065	<a href="#">CG15065</a>	CG15065	Bom	3.3	2.7
CG15068					
CG43202					
CG16836					
CG5778					
Immune induced molecule 23	<a href="#">CG15066</a>	IM23	-	10.3	10
CG15067	<a href="#">CG15067</a>	CG15067	-	3.7	3
CG5791	<a href="#">CG5791</a>	CG5791	-	7.7	7.5
<b>Induced small peptides (40-134a:</b>					
CG30080	<a href="#">CG30080</a>	CG30080	-	22.5	6.9
CG4250	<a href="#">CG4250</a>	CG4250	-	7.3	3.1
CG14957	<a href="#">CG14957</a>	CG14957	chitin-binding?	7.2	-
CG13324	<a href="#">CG13324</a>	CG13324	-	6.7	-
CG15126	<a href="#">CG15126</a>	CG15126	-	6.1	-
CG4269	<a href="#">CG4269</a>	CG4269	-	5.3	-
CG16836	<a href="#">CG16836</a>	CG16836	-	4.9	3.5
Acbp6	<a href="#">CG15829</a>	CG15829	Acyl-CoA binding protein 6	4.9	-
CG9928	<a href="#">CG9928</a>	CG9928	-	4.2	1.9
CG12494	<a href="#">CG12494</a>	CG12494	Withdrawn from Flybase	4.2	-
CG17107	<a href="#">CG17107</a>	CG17107	-	3.9	-
CG5773	<a href="#">CG5773</a>	CG5773	-	3.8	-
CG7738	<a href="#">CG7738</a>	CG7738	Withdrawn from Flybase	3.8	3.8
CG17278	<a href="#">CG17278</a>	CG17278	kazal serine protease inhibitor	3.4	-
CG13323	<a href="#">CG13323</a>	CG13323	-	3.1	-
CG8157	<a href="#">CG8157</a>	CG8157	-	2.9	-
CG9616	<a href="#">CG9616</a>	CG9616	-	2.8	-
Metallothionein C	<a href="#">CG5097</a>	MtnC	metallothionein	2.7	2.9
CG14599	<a href="#">CG14599</a>	CG14599	Withdrawn from Flybase	-	3.4
<b>Induced genes encoding a protein with an homolog in other species</b>					
Ninjurin A	<a href="#">CG6449</a>	NijA	Ninjurin family is inducing nonapoptotic cell death	11.7	-
Charged multivesicular body protein 2b	<a href="#">CG4618</a>	CHMP2B	4e-26 human BC2	5.5	-
CG1358	<a href="#">CG1358</a>	CG1358	1e-109 AF002196 ( <i>C. elegans</i> )	5.1	2
CG4199	<a href="#">CG4199</a>	CG4199	1e-102 Nfr1 ( <i>Xenopus laevis</i> ) Ferredoxin reductase like	5	-
Ady43A	<a href="#">CG1851</a>	Ady43A	1e-23 human adenosine kinase (U33936)	5	-
CG15829	<a href="#">CG15829</a>	CG15829	4e-08 mouse Acyl coA binding site (ACBP)	4.9	-
CG34349	<a href="#">CG11819</a>	CG34349	4e-40 human BAI1-associated 3	4.8	-
CG8791	<a href="#">CG8791</a>	CG8791	6e-30 mouse renal sodium-dependent phosphare transporter NTP1	4.8	2.7
CG10641	<a href="#">CG10641</a>	CG10641	5e-8 mouse caltractin CATR	4.4	-
CG5493	<a href="#">CG5493</a>	CG5493	5e-55 human cysteine dioxygenase pCDO	4.3	-
CG9701	<a href="#">CG9701</a>	CG9701	9e-68 mouse glycosyl hydrolase klotho	4.1	2.2
CG12116	<a href="#">CG12116</a>	CG12116	1e-10 mouse sepiapterin reductase	4	-
Urate oxidase	<a href="#">CG7171</a>	Uro	3e-65 mouse urate oxydase	3.9	2.8
CG32413	<a href="#">CG10487</a>	CG32413	9e-58 human glutaminyl peptide cyclotransferase precursor (QC)	3.8	-
Keap1	<a href="#">CG3962</a>	Keap1	1e-154 mouse Keap1	3.3	-
Gadd45	<a href="#">CG11086</a>	Gadd45	4e-09 human growth arrest and DNA-damage-inducible protein GADD45gar	3.1	-
astray	<a href="#">CG3705</a>	aay	1e-48 3-phosphoserine phosphatase ( <i>Arabidopsis thaliana</i> )	3	-
Death resistor Adh domain containing t	<a href="#">CG1600</a>	Drat	3e-53 <i>C. elegans</i> CAA98080	3	-
CG42797	<a href="#">CG3099</a>	CG42797	6e-89 mouse ubiquitin ligase NED4	3	2.3
Pabp2	<a href="#">CG2163</a>	Pabp2	PolyA binding protein	3	2.1
rhomboid-4	<a href="#">CG1697</a>	rho-4	3e-20 <i>Rattus norvegicus</i> Romboid related protein	2.9	-
lethal (1) G0232	<a href="#">CG3101</a>	I(1)G0232	2e-98 <i>Xenopus laevis</i> tyrosine phosphatases (A53978)	2.9	-

[Clemmons 2015](#)

[Broderick 2012](#)

Aspartyl-tRNA synthetase	<a href="#">CG3821</a>	Aats-asp	1e-144 yeast aspartyl tRNA synthetase	2.8	-
CG17119	<a href="#">CG17119</a>	CG17119	5e-76 human cystinosis	2.8	-
Pak3	<a href="#">CG14895</a>	Pak3	1e-99 human PAK3	2.8	-
CG5527	<a href="#">CG5527</a>	CG5527	2e-58 human KIAA0604	2.7	-
Cyp4p3	<a href="#">CG10843</a>	Cyp4p3	4e-37 human cytochromeP450 4B1	2.7	2.4
N-methyl-D-aspartate receptor-associat	<a href="#">CG3798</a>	Nmda1	4e-49 human protein KIAA0950	2.6	-
Arc2	<a href="#">CG13941</a>	Arc2	5e-06 mouse growth factor ARC	2.6	-
CG5188	<a href="#">CG5188</a>	CG5188	5e-57 human methionine aminopeptidase AMP1	2.5	-
CG8358	<a href="#">CG8358</a>	CG8358	1e-43 NEP mouse neprilysin	2.5	-
Mec2	<a href="#">CG7635</a>	Mec2	5e-76 human stomatin	2.5	2
glaikit	<a href="#">CG8826</a>	gkt	Tyrosyl-DNA phosphodiesterase 1	2.5	2.1
CG1607	<a href="#">CG1607</a>	CG1607	e-126 4F2/CD98 light chain [ <i>Mus musculus</i> ]	2.4	-
CG1607	<a href="#">CG1607</a>	CG1607	1e-126 mouse 4F2/CD98	2.4	-
Surfeit 4	<a href="#">CG6202</a>	Surf4	3e-88 mouse surfait protein SUR4	2.4	-
Cyp6w1	<a href="#">CG8345</a>	Cyp6w1	8e-66 human cytochrome P450 nifedipine oxidase	2.4	-
CG7033	<a href="#">CG7033</a>	CG7033	1e-169 human chaperonin containing t-complex CCT beta	2.3	-
yin	<a href="#">CG44402</a>	yin	1e-122 human oligopeptide transporter PET2	2.2	-
ergic53	<a href="#">CG6822</a>	ergic53	1e-104 human mannose specific lectin (U09716)	2.2	-
Purine-rich binding protein-alpha	<a href="#">CG1507</a>	Pur-alpha	3e-35 human transcriptional activator PUR-alpha	2.1	2.2
amnesiac	<a href="#">CG11937</a>	amn		2.1	2.2

### Induced unknown proteins

Some of the following genes encode proteins with a characterized domain.

Frost	<a href="#">CG9434</a>	Fst	involved in the cold response	13.2	2.6
CG13905	<a href="#">CG13905</a>	CG13905	-	13.1	-
CG14567	<a href="#">CG14567</a>	CG14567	-	12.7	2.8
CG13077	<a href="#">CG13077</a>	CG13077	-	10.3	-
CG13325	<a href="#">CG13325</a>	CG13325	-	7.9	-
Diuretic hormone 44	<a href="#">CG8348</a>	Dh44	-	5.8	-
CG15043	<a href="#">CG15043</a>	CG15043	-	5.7	2.4
CG14219	<a href="#">CG14219</a>	CG14219	-	5.7	-
CG16772	<a href="#">CG16772</a>	CG16772	-	5.6	4.7
Immune induced molecule 10	<a href="#">CG18279</a>	IM10	-	5.6	5
CG9989	<a href="#">CG9989</a>	CG9989	-	5.2	2.2
CG16718	<a href="#">CG16718</a>	CG16718	-	5.2	-
CG2217	<a href="#">CG2217</a>	CG2217	-	5	2.2
CG15745	<a href="#">CG15745</a>	CG15745	-	4.6	-
CG5778	<a href="#">CG5778</a>	CG5778	-	4.4	3
CG10420	<a href="#">CG10420</a>	CG10420	-	4.4	-
CG18067	<a href="#">CG18067</a>	CG18067	-	4.3	3.4
CG5118	<a href="#">CG5118</a>	CG5118	-	4	-
CG10912	<a href="#">CG10912</a>	CG10912	-	4	-
CG7296	<a href="#">CG7296</a>	CG7296	-	3.7	-
tamo	<a href="#">CG4057</a>	tamo	Tamo selectively modulates nuclear import in	3.6	<a href="#">Minakhina 2003</a>
CG7778	<a href="#">CG7778</a>	CG7778	-	3.6	-
CG10680	<a href="#">CG10680</a>	CG10680	-	3.4	-
Nimrod A	<a href="#">CG8855</a>	NimA	-	3.3	-
Activity-regulated cytoskeleton associa	<a href="#">CG12505</a>	Arc1	-	3.3	2.3
CG8965	<a href="#">CG8965</a>	CG8965	-	3.2	-
CG13311	<a href="#">CG13311</a>	CG13311	-	3.2	-
CG13618	<a href="#">CG13618</a>	CG13618	-	3.2	-
CG13641	<a href="#">CG13641</a>	CG13641	-	3.2	-
Cuticular protein 67Fb	<a href="#">CG18348</a>	Cpr67Fb	-	3.1	-
Sclp	<a href="#">CG2471</a>	Sclp	-	3.1	-
CG7016	<a href="#">CG7016</a>	CG7016	-	3	-
CG6426	<a href="#">CG6426</a>	CG6426	-	3	-
CG13795	<a href="#">CG13795</a>	CG13795	-	3	-
CG15293	<a href="#">CG15293</a>	CG15293	-	3	-
CG11413	<a href="#">CG11413</a>	CG11413	-	3	-
Nucleoporin 205kD	<a href="#">CG11943</a>	Nup205	-	3	2.2
CG33307	<a href="#">CG16887</a>	CG33307	-	2.9	-
Na <sup>+</sup> /H <sup>+</sup> hydrogen antiporter 2	<a href="#">CG4693</a>	Nha2	-	2.9	-
CG14762	<a href="#">CG14762</a>	CG14762	-	2.8	-
CG16743	<a href="#">CG16743</a>	CG16743	-	2.8	-
CG6357	<a href="#">CG6357</a>	CG6357	-	2.8	-
CG15784	<a href="#">CG15784</a>	CG15784	-	2.8	-
CG42807	<a href="#">CG6183</a>	CG42807	-	2.8	-
CG17560	<a href="#">CG17560</a>	CG17560	-	2.7	2.1
CG34370	<a href="#">CG13497</a>	CG34370	-	2.7	2.3
CG14841	<a href="#">CG14841</a>	CG14841	-	2.7	2
CG12502	<a href="#">CG12502</a>	CG12502	-	2.7	2.7
CG15282	<a href="#">CG15282</a>	CG15282	-	2.6	-
TRAM	<a href="#">CG11642</a>	TRAM	-	2.5	-
CG7695	<a href="#">CG7695</a>	CG7695	-	2.5	3.2
CG10383	<a href="#">CG10383</a>	CG10383	-	2.5	-
CG10910	<a href="#">CG10910</a>	CG10910	-	2.5	-
CG9186	<a href="#">CG9186</a>	CG9186	-	2.4	-



CG5059	<a href="#">CG5059</a>	CG5059	-	2.4	-
CG14661	<a href="#">CG14661</a>	CG14661	-	2.4	2.2
CG10332	<a href="#">CG10332</a>	CG10332	-	2.4	-
CG13193	<a href="#">CG13193</a>	CG13193	-	2.4	2
CG11417	<a href="#">CG11417</a>	CG11417	-	2.4	2.3
CG34347	<a href="#">CG15566</a>	CG34347	-	2.4	2.2
CG6672	<a href="#">CG6672</a>	CG6672	-	2.3	-
CG14907	<a href="#">CG14907</a>	CG14907	-	2.3	-
Mucin 55B	<a href="#">CG5765</a>	Mucin55B	-	2.3	-
CG18467	<a href="#">CG18467</a>	CG18467	-	2.3	2.4
CG2875	<a href="#">CG2875</a>	CG2875	-	2.3	-
DNApiol-iota	<a href="#">CG7602</a>	DNApiol-iota	-	2.2	-
CG14401	<a href="#">CG14401</a>	CG14401	-	2.2	-
CG6073	<a href="#">CG6073</a>	CG6073	-	2.2	2.1
CG17019	<a href="#">CG17019</a>	CG17019	-	2.2	-
CG14625	<a href="#">CG14625</a>	CG14625	-	2.1	2.3