

Arefin, Table S1

Gene	Primer name	Sequence	Annealing temperature
Rack1	Rack1 fw	CCC GTG ACA AGA CCC TGA T	50 °C
	Rack1 rev	TAG TTG CCA TCG GAG GAG AG	
rp49	rp49 fw	CTT CAT CCG CCA CCA GTC	50 °C
	rp49 rev	GGC GAC GCA CTC TGT TGT	
IM3	IM3 fw	TTG GGT CTG CTG GCT CTG	52 °C
	IM3 rev	TTC AAC TGG CAT CCT TCA TTC	
Diptericin B	DiptB fw	CTA TTC ATT GGA CTG GCT TGT G	54 °C
	DiptB rev	GTC CAT TGG GGC TCT GC	
Drosomycin	Drs fw	CCC TCT TCG CTG TCC TGA	52 °C
	Drs rev	TTA GCA TCC TTC GCA CCA G	
Attacin A	AttA fw	TGG TCA TGG TGC CTC TTT G	54 °C
	AttA rev	GAT TGT GTC TGC CAT TGT TGA	
PGRP-SB1	SB1 fw	GGG TGC GGT GTC TGC TC	52 °C
	SB1 rev	TCA CCG GCC ACG ATA AAG	
GNBP-like	CG13422 fw	CCA AGG CCA CCG TCA AG	52 °C
	CG13422 rev	TCG CTC AGA TCG TCC ATT T	
Tep1	Tep2 fw	GTC CTG CTC GCC CTT CTC	52 °C
	Tep2 rev	TCA AAT GCC AAA ACT CTA TGT CA	
Tep2	Tep1 fw	CGT TCT GCT GGC TTT CTT C	52 °C
	Tep1 rev	ATA CTG GTC GTC CGT CTT GTC	
Spds	Spds fw	GCG ATG GCC TGT GGT TT	52 °C
	Spds rev	CAG CGC GTG TTT CAT TAG C	
Glutactin	Glt fw	TAT CCG GTA CAA GAG CCA CAG	54 °C
	Glt rev	TTT CGG GGA GAT TTT CGT T	
CG7607	CG7607 fw	TAC ATA GCC GCC CAA AGA A	52 °C
	CG7607 rev	AGC CAT GCA AAA GAA AGT GAT	
PGRP-LF	LF fw	CGC CAT CAT GTT TCA CTC AA	52 °C
	LF rev	GAC CAA TAA AGG CCA CGA CTA	

Arefin, Table S1B

VDRC lines

Array genes (fig.4)

No	Gene name	CG no	Transformant ID	Type	Off target	Viability	Inserted Chromosome	Larval fatbody expression	Induction level (microarray)
1	Attacin A	CG10146	9287	GD		1 viable	3 yes		Strongly induced
2	Attacin A	CG10146	50319	GD		2 viable	3 yes		Strongly induced
3	Attacin A	CG10146	50320	GD		2 viable	2 yes		Strongly induced
4	DiptericinB	CG10794	102607	KK		0 viable	2 yes		Strongly induced
5	DiptericinB	CG10794	28736	GD		1 viable	3 yes		Strongly induced
6	Drosomycin	CG10810	2703	GD		5 viable	2 yes		Strongly induced
7	GNBP like 3	CG13422	107358	KK		0 viable	2 yes		Strongly induced
8	GNBP like 3	CG13422	7545	GD		0 viable	3 yes		Strongly induced
9	IM2	CG18106	43372	GD		3 viable	2 yes		Strongly induced
10	IM3	CG16844	29214	GD		2 viable	1 yes		Strongly induced
11	IM18	CG10332	49651	GD		0 viable	3 yes		Moderate
12	Metchnikowin	CG8175	109740	KK		1 viable	2 yes		Strongly induced
13	PGRP-SB1	CG9681	101298	KK		0 viable	2 yes		Strongly induced
14	SPH93	CG6639	104307	KK		0 viable	2 yes		Moderate
15	Akap200	CG13388	102374	KK		0 viable	2 yes		2 fold (LogFC)
16	Akap200	CG13388	109996	KK		0 viable	2 yes		2 fold (LogFC)
17	Akap200	CG13388	5647	GD		0 viable	2 yes		2 fold (LogFC)
18	Starry night	CG11895	107993	KK		0 viable	2 Not available		Moderate
19	Starry night	CG11895	1665	GD		0 viable	3 Not available		Moderate
20	Starry night	CG11895	51382	GD		0 viable	2 Not available		Moderate
21	Argonaute 2	CG7439	100356	KK		0 viable	2 yes		no
22	Argonaute 2	CG7439	49473	GD		0 viable	2 yes		no
23	Argonaute 2	CG7439	51521	GD		0 viable	2 yes		no
24	Larval transluc	CG32372	18977	GD		2 viable	3 yes		no
25	Wrinkled	CG5123	8269	GD		0 viable	3 no		no

Hemocytes enriched genes (fig. 6)

26	Lectin24DB	CG2958	105118	KK		0 viable	2		
27	Lectin24DB	CG2958	45294	GD		2 viable	3		
28	Lectin28C	CG7106	104201	KK		0 viable	2		
29	SPARC	CG6378	100566	KK		1 viable	2		
30	SPARC	CG6378	16677	GD		1 viable	2		
31	Eater	CG6124	4301	GD		0 viable	3		
32	dSR-C1	CG4099	49964	GD		0 viable	2		
33	Glutactin	CG9280	101918	KK		1 viable	2		
34	Glutactin	CG9280	15428	GD		1 viable	3		
35	Glutactin	CG9280	15429	GD		1 viable	2		
36	CG4250	CG4250	101949	KK		0 viable	2		

NIG lines (Fig.4)

37	GNBP like 3	CG13422	13422R-3	via	3
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Mutant lines (Fig.5)

Bloomington line: No	Gene name	CG no	Genotype	Stock no.	Type	Mutant type	Viability
38	Glutactin	CG9280		22539	Mutant	P-element ins	via

From Bruno Lemaitre lab (Fig.5)

39	GNBP-3	CG5008	GNBP-3/TM6c		Mutant		lethal
40	PGRP-SA	CG11709	y, w, DD, Seml (PGRP-SA		Mutant		via
41	PGRP-LC	CG4432	PGRP-LC ^{E12D3}		Mutant		via
42	PGRP-LE	CG8995	y ¹ w ^{67c23} PGRP-LE ¹¹²		Mutant		via
43	PGRP-LF	CG4437	PGRP-LF ²⁰⁰		Mutant		lethal

From Dominique Ferrandoni lab (Fig.5)

44	Tep2	CG7052	Tep2 (d11521)		Mutant		via
45	Tep3	CG7068	Tep3 (d03976)		Mutant		via
46	Tep4	CG10363	Tep4(EY04656)		Mutant		via
47	Tep2,3				Double mu	FRT medited	via

Reporter lines

48	Ddc-GFP					- 1.4 kb Ddc-GFP transgenic line was obtained from William McGinnis lab	
49	hml-GAL4;UAS-RFP					w1118 iso;+; hml-Gal4 UAS-RFP line was generated by recombining hml-Gal4 and UAS-RFP on 3rd chromosome	
50	vkg-GFP	CG16858		G00454	protein-trap	via	obtained from Cooley lab (Flytrap)

Arefin, Table S3

Gene Name	Abbreviation	Probe Set	Nematodes (log2 FC)	Wasps (Code/FC)	Injection of G+ and G- bacteria (FC)	Pathogenic G- Bacteria (FC)	Class
Immune induced molecule 3 elevated during infection	IM3	1622893 at	8.99	7.858119668	12.05	4.1	A - all infections
Immune induced molecule 4	IM4	1634366 at	8.76	13.28267974			C - shared EPN+wasps
Attacin-C	AttC	1626345 at	8.23	7.265109147	2.62		B - missing in pathogenic bacteria
Immune induced molecule 1	IM1	1641419 at	7.94	10.45256917	28.08	11.655	A - all infections
IM2-like	CG15065	1633053 at	7.87	2.870099541	26.73	2.685	A - all infections
Defensin	Def	1631660 at	7.81	14	9.4	2.365	A - all infections
Dipteracin B	DptB	1634271 at	7.81		2.86		F - shared EPN+bac. infection
GNBP-like	CG13422	1638235 at	7.29	4.04834199	9.270157347	9.95	A - all infections
Dmel_CG15067	CG15067	1627271 at	7.1	2	9.6		B - missing in pathogenic bacteria
Dmel_CG33470	CG33470	1625174 at	6.95	2	10.46		B - missing in pathogenic bacteria
Immune induced molecule 2	IM2	1639019 s at	6.93				D - EPN specific
Immune induced molecule 23	IM23	1640360 at	6.84	2.091029944	7.62		B - missing in pathogenic bacteria
Dmel_CG16836	CG16836	1629530 at	6.75	2	13.12		B - missing in pathogenic bacteria
Drosomyacin	Drs	1623669 at	6.59	2	7.259899432		B - missing in pathogenic bacteria
Immune induced molecule 10	IM10	1635189 at	6.02	3.318574591	11.7879722	5.48	A - all infections
Metchnikowin	Mtk	1626319 a at	5.95	14	23.24764177	2.62	A - all infections
Attacin-A	Atta	1627613 at	5.5	9.252935299	37.81	13.865	A - all infections
Attacin-A	Atta	1627551 s at	5.46	18.76399116	2.319343217		B - missing in pathogenic bacteria
Dmel_CG30026	CG30026	1625124 at	5.44	13.85899334	52.02091148	18.65	A - all infections
Peptidoglycan recognition protein SB1	PGRP-SB1	1631370 at	5.29				D - EPN specific
Dmel_CG15068	CG15068	1636490 at	5.27	10.20658452	3.54	5.44	A - all infections
Cecropin C	CecC	1635060 at	5.15	10.81605612			C - shared EPN+wasps
Dmel_CG4757	CG4757	1632719 at	5		1.69	1.98	F - missing in wasp infection
Dmel_CG33462	CG33462	1638021 at	4.66				D - EPN specific
Cecropin B	CecB	1640757 at	4.49	12	3.7		B - missing in pathogenic bacteria
Attacin-D	Attd	1626530 at	4.45	4.357954545	1.83	3.805	B - missing in pathogenic bacteria
Dmel_CG16775	CG16775	1631475 at	3.84		2.03		F - missing in wasp infection
Dmel_CG42335	CG42335	1623884 at	3.57				G - shared EPN+pathogenic G- bacteria
Dmel_CG18067	CG18067	1634477 at	3.53				G - shared EPN+pathogenic G- bacteria
Dmel_CG18107	CG18107	1640144 at	3.28	13	7.103805806	2.255	A - all infections
gp150-like	CG14762	1638772 at	2.95	3.338274045	2.63		B - missing in pathogenic bacteria
Dmel_CG4716	CG4716	1632809 at	2.95				D - EPN specific
Dmel_CG30029	CG30029	1629046 a at	2.94	8			C - shared EPN+wasps
accord	CG2217	1636492 at	2.92			1.68	G - shared EPN+pathogenic G- bacteria
Dmel_CG2217	CG2217	1634633 s at	2.92				D - EPN specific
Fibrinogen-like	CG5550	1636293 at	2.87	4.52221851			C - shared EPN+wasps
Dmel_CG4716	CG4716	1629201 at	2.81	2.364948282	2.32	13.7	A - all infections
Thioester-containing protein 1	Tep1	1628963 at	2.7		2.09	2.7	F - missing in wasp infection
Zn finger homeodomain 1	zfh1	1637734 at	2.66	12			F - missing in wasp infection
Dmel_CG6639	CG6639	1628262 s at	2.54				D - EPN specific
Dmel_CG42232	CG42232	1625698 at	2.47	12	4.81	2.16	A - all infections
Chronologically inappropriate morphogenesis	chinmo	1636205 at	2.47				D - EPN specific
Mucin 26B	Muc26B	1629484 s at	2.45				D - EPN specific
Mucin 96D	Muc96D	1632017 at	2.44		1.638986293	2.075	F - missing in wasp infection
polychaetoid	pyd	1636058 at	2.44	0.653075935			D - EPN specific
Mucin 96D	Muc96D	1637322 at	2.4				C - shared EPN+wasps
Insulin-like receptor	InR	1635733 x at	2.4				D - EPN specific
starry night	stan	1629141 at	2.35				D - EPN specific
Dmel_CG8160	CG8160	1625087 at	2.32				D - EPN specific
zye	CG5847	1638648 at	2.31	7.75063567	7.6	2.98	A - all infections
Beta amyloid protein precursor-like	Appl	1623710 at	2.31	1.162055336			C - shared EPN+wasps
Transposon.63	CG14695	1624033 at	2.31	0.797154497			C - shared EPN+wasps
Dmel_CG14695	CG14695	1632295 s at	2.31				D - EPN specific
Dmel_CG13482	CG13482	1639704 at	2.29	1.394326241			H - missing in bac. infection
NK7.1	CG8524	1636798 at	2.22		9.751731283	6.7	F - missing in wasp infection
anachronism	ana	1631303 s at	2.22				D - EPN specific
no receptor potential A	norpA	1630502 at	2.21				D - EPN specific
Dmel_CG11893	CG11893	1636576 s at	2.21				D - EPN specific
Chronologically inappropriate morphogenesis	chinmo	1635512 at	2.19				D - EPN specific
moody	CG4322	1628005 at	2.16				D - EPN specific
Dmel_CG12236	CG12236	1633112 at	2.14				D - EPN specific
DISCO Interacting Protein 1	DIP1	1624189 at	2.13				D - EPN specific
anon-fast-evolving-1H7	klu	1638708 s at	2.12				D - EPN specific
klumpfuss	Dmel_CG6279	1624215 s at	2.09				D - EPN specific
Dmel_CG13461	CG13461	1625672 s at	2.08				D - EPN specific
anon-fast-evolving-1H7	wech	1629347 at	2.08				D - EPN specific
Neurofibromin 1	Nf1	1638969 at	2.07				D - EPN specific
Mucin 68Ca	Muc68Ca	1635959 at	2.06		2.274807593		E - shared EPN+bac. infection
Muscle-specific protein 300	Msp-300	1640472 at	2.05		1.68	1.885	F - missing in wasp infection
Neurofibromin 1	Nf1	1639785 s at	2.04	0.8495051			C - shared EPN+wasps
Dmel_CG30069	CG30069	1625048 at	2.03				D - EPN specific
Odorant-binding protein 49a	Obp49a	1636834 s at	2.03				D - EPN specific
polyA-binding protein interacting protein 2	Paip2	1632945 at	2.02				D - EPN specific
Dmel_CG8388	CG8388	1634083 at	2.02				D - EPN specific
Thioester-containing protein 2	Tep2	1635730 at	2.02				D - EPN specific
SRPK	SRPK	1624932 at	2.01				D - EPN specific
IGF-II mRNA-binding protein	Imp	1628632 at	2.01				D - EPN specific
headcase	hdc	1627323 at	2	7.195665353	4.089814116		D - EPN specific
twin of eyeless	toy	1630067 a at	1.99				B - missing in pathogenic bacteria
Dmel_CG42335	CG42335	1632130 s at	1.98				D - EPN specific
Dmel_CG12071	CG12071	1623455 s at	1.97				D - EPN specific
Fascilin 1	Fas1	1629307 s at	1.96	0.594499033			C - shared EPN+wasps
Cenp-C	Cenp-C	1633512 at	1.96				D - EPN specific
Bruce	CG6303	1635601 at	1.95			1.54	G - shared EPN+pathogenic G- bacteria
Transposon.17	CG16772	1624143 a at	1.95				D - EPN specific
cabl	CG14478	1624183 a at	1.95				D - EPN specific
Dmel_CG16772	CG16772	1624183 a at	1.94				D - EPN specific
Dmel_CG14478	CG14478	1634454 at	1.93	0.954526709			C - shared EPN+wasps
Dmel_CG13323	CG13323	1624819 s at	1.93				D - EPN specific
grappa	spp	1635580 at	1.93				D - EPN specific
Mucin 68Ca	Muc68Ca	1631691 at	1.92		2.76		E - shared EPN+bac. infection
CG42342	CG42342	1638359 s at	1.92				D - EPN specific
pericentrin-like	cp309	1624587 at	1.91		7.212604808	2.095	F - missing in wasp infection
Kinesin heavy chain 73	Khc-73	1627796 s at	1.89				D - EPN specific
Elongase 68a	Elo68a	1631700 at	1.89				D - EPN specific
LD07388P/CAPRICIOUS	LD07388	1631471 at	1.88				D - EPN specific
Dmel_CG16711	CG16711	1634009 at	1.88				D - EPN specific
derailed	dri	1627136 at	1.87				D - EPN specific
Muscle-specific protein 300	Msp-300	1635229 at	1.87				D - EPN specific
Tie-like receptor tyrosine kinase	Tie	1634920 at	1.86	2.467849224			D - EPN specific
Dmel_CG10332/Immune induced molecule 18	CG10332/IM18	1624565 a at	1.85				C - shared EPN+wasps
Relish	Rel	1624297 at	1.85				D - EPN specific
leak	lea	1627250 at	1.85				D - EPN specific
Dmel_CG10911	CG10911	1629129 at	1.85				D - EPN specific
Peptidoglycan recognition protein LF	PGRP-LF	1622952 at	1.83				D - EPN specific
KP78b	KP78b	1627376 at	1.82	4	3.60014622	3.415	A - all infections
Dmel_CG34383	CG34383	1636905 at	1.82	0.943275096			C - shared EPN+wasps
I(3)neo38	CG10077	1624342 at	1.82			2.3	G - shared EPN+pathogenic G- bacteria
Dmel_CG10077	CG10077	1633145 at	1.8				D - EPN specific
armtage	armi	1639894 at	1.8				D - EPN specific
dumpy	dp	1639894 at	1.8				D - EPN specific
		1625511 at	1.79				D - EPN specific
		1640598 s at	1.78				D - EPN specific
		1638060 at	1.77				D - EPN specific
		1623907 at	1.76				D - EPN specific
		1634075 at	1.76				D - EPN specific

Cyclin T	CycT	1637897 at	1.75	0.716867073			C - shared EPN+wasp
Dmel_CG9821	CG9821	1634992 s at	1.74				D - EPN specific
Darkener of apricot	Doa	1640892 a at	1.73				D - EPN specific
Distal-less	Dll	1636088 at	1.71				D - EPN specific
Dmel_CG16743	CG16743	1630088 at	1.7	6.660108401	2.1		H - missing in bac. Injection
Dmel_CG6024	CG6024	1636848 at	1.7	0.949790795			C - shared EPN+wasp
ChT6	ChT6	1633303 at	1.69				D - EPN specific
anastral spindle 1	ana1	1637397 a at	1.69				D - EPN specific
tartan	trn	1639235 at	1.69				D - EPN specific
Thor	CG8846	1635900 at	1.67		1.71	3.44	F - missing in wasp infection
mutagen-sensitive 210	mus210	1632827 a at	1.67				D - EPN specific
nemo	nmo	1624300 s at	1.66	0.870678928			C - shared EPN+wasp
schnurri	shn	1625445 s at	1.66				D - EPN specific
trio	trio	1635047 s at	1.66				D - EPN specific
Myosin binding subunit	Mbs	1630456 at	1.65				D - EPN specific
Dmel_CG8771	CG8771	1635778 at	1.65				D - EPN specific
Dmel_CG3630	CG3630	1636461 at	1.64				D - EPN specific
Connectin	Con	1639913 at	1.63	0.912106008			C - shared EPN+wasp
telomere fusion	tefu	1630729 at	1.63				D - EPN specific
Dmel_CG4213	CG4213	1634084 at	1.63				D - EPN specific
unc-13	unc-13	1635684 a at	1.63				D - EPN specific
Dmel_CG5098	CG5098	1637079 at	1.63				D - EPN specific
Syncrip	Syp	1640760 at	1.62	0.27532097	2.416948355		B - missing in pathogenic bacteria
scabrous	sca	1636998 at	1.62	0.876908394			C - shared EPN+wasp
Prosap	Prosap	1625236 s at	1.62				D - EPN specific
Dmel_CG5524	CG5524	1639138 at	1.62				D - EPN specific
stumps	stumps	1634063 a at	1.61				D - EPN specific
Dmel_CG13003	CG13003	1630570 at	1.6				D - EPN specific
staufen	stau	1633016 a at	1.6				D - EPN specific
dim v-tubulin 5	dgt5	1641436 at	1.6				D - EPN specific
Dmel_CG32176	CG32176	1641704 at	1.6				D - EPN specific
enabled	ena	1627191 a at	1.59				D - EPN specific
CHKov1	CHKov1	1625997 s at	1.58				D - EPN specific
eukaryotic translation initiation factor 4G2	eIF4G2	1637987 at	1.55	0.862151836			C - shared EPN+wasp
Cullin-3	Cul-3	1624970 s at	1.55				D - EPN specific
golden goal	gogo	1625852 at	1.55				D - EPN specific
mRNA-like ncRNA in embryogenesis 2	MRE2	1629325 at	1.55				D - EPN specific
abrupt	ab	1629702 a at	1.55				D - EPN specific
naked cuticle	nkd	1630361 at	1.55				D - EPN specific
Myocardin-related transcription factor	Mrtf	1636679 at	1.55				D - EPN specific
anastral spindle 3	ana3	1640805 at	1.55				D - EPN specific
Dmel_CG14655	CG14655	1627852 at	1.54				D - EPN specific
serrano	sano	1627971 s at	1.54				D - EPN specific
Thd1	Thd1	1636521 at	1.54				D - EPN specific
Nutrient Amino Acid Transporter 1	NAAT1	1639528 at	1.54				D - EPN specific
Dmel_CG1146	CG1146	1637605 s at	1.53				D - EPN specific
Jumonji, AT rich interactive domain 2	Jarid2	1634092 at	1.52	0.858938891			C - shared EPN+wasp
enhanced adult sensory threshold	east	1624378 at	1.52				D - EPN specific
Dmel_CG16972	CG16972	1630207 at	1.52				D - EPN specific
couch potato	cpo	1632644 s at	1.52				D - EPN specific
Dmel_CG6945	CG6945	1630923 at	1.51				D - EPN specific
Dmel_CG42356	CG42356	1632958 a at	1.5			2.33	G - shared EPN+pathogenic G- bacteria
mustard	mtd	1626842 s at	1.5				D - EPN specific
Dmel_CG31340	CG31340	1633241 at	1.5				D - EPN specific
nervous fingers 1	nerfin-1	1634039 at	1.5				D - EPN specific
longitudinals lacking	lola	1635096 at	1.5				D - EPN specific
Dmel_CG42724	CG42724	1637665 at	1.5				D - EPN specific
onecut	onecut	1630376 at	1.49				D - EPN specific
bangles and beads	bnb	1632734 s at	1.49				D - EPN specific
CirI	CirI	1640640 at	1.49				D - EPN specific
Dmel_CG31780	CG31780	1624610 s at	1.48	7.268118978			C - shared EPN+wasp
Checkpoint suppressor homologue	CHES-1-like	1629295 at	1.48				D - EPN specific
Dmel_CG30091	CG30091	1632210 at	1.47				D - EPN specific
Not1	Not1	1638560 a at	1.47				D - EPN specific
Phosphodiesterase 11	Pde11	1639582 at	1.47				D - EPN specific
Dmel_CG9451	CG9451	1636526 at	1.46	2.029733094			C - shared EPN+wasp
Dmel_CG10383	CG10383	1629398 at	1.46			2.193194598	F - shared EPN+bac. Injection
APC-like	Apc	1628548 at	1.46			1.37	G - shared EPN+pathogenic G- bacteria
polyhomeotic proximal	ph-p	1623441 at	1.46				D - EPN specific
Myocyte enhancer factor 2	MeF2	1626392 s at	1.46				D - EPN specific
decapentaplegic	dpp	1630026 s at	1.46				D - EPN specific
Protein C kinase 98E	Pkc98E	1631059 at	1.46				D - EPN specific
Alhambra	Alh	1630415 at	1.45	1.02075416			C - shared EPN+wasp
kon-tiki	kon	1625687 at	1.45				D - EPN specific
Calmodulin-binding transcription activator	Camta	1626232 at	1.45				D - EPN specific
Dmel_CG42390	CG42390	1628867 s at	1.45				D - EPN specific
polo	polo	1636189 at	1.45				D - EPN specific
karst	kst	1637710 at	1.45				D - EPN specific
fat	ft	1624125 at	1.44				D - EPN specific
mushroom-body expressed	mub	1625921 at	1.44				D - EPN specific
Dmel_CG4612	CG4612	1627151 at	1.44				D - EPN specific
skuld	skd	1631516 s at	1.44				D - EPN specific
asense	ase	1635124 at	1.44				D - EPN specific
anterior open	aop	1627394 s at	1.43	1.00640539			C - shared EPN+wasp
Stromalin	SA	1626710 at	1.42				D - EPN specific
Meiotic central spindle	Meics	1629709 at	1.42				D - EPN specific
Dmel_CG6357	CG6357	1633059 at	1.42				D - EPN specific
Suppressor of zeste 2	Su(2)2	1633866 at	1.42				D - EPN specific
longitudinals lacking	lola	1640945 at	1.41	2.573891626			C - shared EPN+wasp
trithorax	trx	1624533 s at	1.41	0.936943992			C - shared EPN+wasp
ftz transcription factor 1	ftz-f1	1624520 a at	1.41				D - EPN specific
Dmel_CG9626	CG9626	1624725 at	1.41				D - EPN specific
Liprin-y	Liprin-y	1625011 at	1.41				D - EPN specific
Dmel_CG32369	CG32369	1638611 at	1.41				D - EPN specific
Dmel_CG7376	CG7376	1623124 at	1.4				D - EPN specific
Tis11 homolog	Tis11	1623863 a at	1.4				D - EPN specific
Dmel_CG5514	CG5514	1624644 a at	1.4				D - EPN specific
spalt major	salm	1627881 at	1.4				D - EPN specific
SoxNeuro	SoxN	1631408 at	1.4				D - EPN specific
Dp110	PI3K92E	1631594 s at	1.4				D - EPN specific
Dmel_CG32113	CG32113	1632287 at	1.4				D - EPN specific
daily-like	dlp	1636974 at	1.4				D - EPN specific
Dmel_CG13636	CG13636	1641738 a at	1.4				D - EPN specific
Dmel_CG12868	CG12868	1635030 at	1.39			3.062181652	F - missing in wasp infection
purity of essence	poe	1634654 at	1.39	0.854332193		8	C - shared EPN+wasp
Dmel_CG18769	CG18769	1633715 s at	1.39				D - EPN specific
B52	B52	1633821 at	1.39				D - EPN specific
rad50	rad50	1638887 a at	1.39				D - EPN specific
Dmel_CG8457	CG8457	1640747 s at	1.39				D - EPN specific
roundabout	robo	1626774 s at	1.38				D - EPN specific
gliolectin	glec	1628743 at	1.38				D - EPN specific
Dmel_CG7715	CG7715	1629765 at	1.38				D - EPN specific
midline fasciclin	mfas	1632298 s at	1.38				D - EPN specific
split ends	spen	1641518 a at	1.38				D - EPN specific
hibris	hbs	1637539 a at	1.37	0.74710544			C - shared EPN+wasp
myoblast city	mbc	1631013 at	1.37			2.135	G - shared EPN+pathogenic G- bacteria
Dmel_CG6700	CG6700	1627530 at	1.37				D - EPN specific
Dmel_CG11414	CG11414	1631406 at	1.37				D - EPN specific
Dmel_CG31357	CG31357	1632916 at	1.37				D - EPN specific
wings apart-like	wapl	1636092 a at	1.37				D - EPN specific
Sox21b	CG32139	1637750 at	1.36	0.459252157			C - shared EPN+wasp
Dmel_CG7294	CG7294	1631635 at	1.36			2.84	G - shared EPN+pathogenic G- bacteria
Dmel_CG43896	CG43896	1626206 at	1.36			1.54	G - shared EPN+pathogenic G- bacteria
CAP-D2 condensin subunit	CAP-D2	1625447 at	1.36				D - EPN specific
ariadne 2	ari-2	1625556 at	1.36				D - EPN specific
Dmel_CG2258	CG2258	1629104 at	1.36				D - EPN specific
Smad on X	Smox	1629290 at	1.35				D - EPN specific
Dmel_CG7142	CG7142	1635282 at	1.35				D - EPN specific
disconnected	disco	1639940 at	1.35				D - EPN specific
Dmel_CG8600	CG8600	1640510 at	1.35				D - EPN specific
Synaptotagmin 4	Syt4	1641475 at	1.35				D - EPN specific
Dmel_CG10289	CG10289	1641548 at	1.35				D - EPN specific

Dmel_CG6923	CG6923	1628420 s at	1.34			D - EPN specific	
par-1	par-1	1628849 at	1.34			D - EPN specific	
Tollo	Tollo	1632481 at	1.34			D - EPN specific	
hepsocat	hop	1639072 at	1.33			C - shared EPN+wasp	
insensitive	insv	1624375 at	1.33			D - EPN specific	
scribbler	sbb	1626352 at	1.33			D - EPN specific	
Dmel_CG42533	CG42533	1634804 at	1.33			D - EPN specific	
Dmel_CG34398	CG34398	1635127 at	1.33			D - EPN specific	
pou domain motif 3	pdm3	1631222 at	1.32	3.545454545		C - shared EPN+wasp	
gluon	glu	1635123 at	1.32	3.052313883		C - shared EPN+wasp	
Dmel_CG32767	CG32767	1622122 at	1.32			D - EPN specific	
mutagen-sensitive 312	mus312	1629113 a at	1.32			D - EPN specific	
pointed	pnt	1630010 a at	1.32			D - EPN specific	
chiffon	chif	1630995 at	1.32			D - EPN specific	
Interacts with the C terminus of ELL 1	Ice1	1631109 at	1.32			D - EPN specific	
Centrosomal protein 190kD	Cp190	1631940 s at	1.32			D - EPN specific	
Ankyrin 2	Ank2	1633313 at	1.32			D - EPN specific	
vielfaltig	vfi	1638370 s at	1.32			D - EPN specific	
pavarotti	pav	1623405 at	1.31			D - EPN specific	
Sin3A	Sin3A	1630165 s at	1.31			D - EPN specific	
Dmel_CG42863	CG42863	1634291 at	1.31			D - EPN specific	
Dmel_CG31122	CG31122	1637365 at	1.31			D - EPN specific	
Dmel_CG16896	CG16896	1638352 at	1.31			D - EPN specific	
longitudinals lacking	lola	1634495 s at	1.3	0.837751405		C - shared EPN+wasp	
optic ganglion reduced	ogre	1628323 s at	1.3		2.21608288	E - shared EPN+bac. injection	
Promyelocytic leukemia zinc finger ortholog	Plzf	1639534 at	1.3			D - EPN specific	
Leukocyte-antigen-related-like	Lar	1637537 at	1.29	0.92228936		C - shared EPN+wasp	
multiple ankyrin repeats single KH domain	mask	1641226 a at	1.29	0.875724028		C - shared EPN+wasp	
found in neurons	fne	1633852 at	1.29	0.717516202		C - shared EPN+wasp	
Glutactin	GLT	1630515 s at	1.29		1.06	G - shared EPN+pathogenic G- bacteria	
Dmel_CG9715	CG9715	1627694 at	1.29			D - EPN specific	
Dmel_CG5639	CG5639	1630064 at	1.29			D - EPN specific	
squeeze	sqz	1630772 at	1.29			D - EPN specific	
Dmel_CG12054	CG12054	1636061 at	1.29			D - EPN specific	
Megalyn	mgf	1641272 at	1.29			D - EPN specific	
Dmel_CG2225	CG2225	1630609 s at	1.28			D - EPN specific	
prospero	pros	1635500 a at	1.28			D - EPN specific	
female sterile (1) homeotic	fs(1)h	1625127 at	1.27	0.680376656		C - shared EPN+wasp	
nejire	nej	1622925 at	1.27			D - EPN specific	
roughest	rst	1625366 at	1.27			D - EPN specific	
Dmel_CG6181	Ge-1	1632251 s at	1.27			D - EPN specific	
Dmel_CG32982	CG32982	1634406 at	1.27			D - EPN specific	
Dmel_CG12105	CG12105	1640809 at	1.27			D - EPN specific	
monkey king	mkg	1622892 s at	1.26		1.21	G - shared EPN+pathogenic G- bacteria	
longitudinals lacking	lola	1625768 s at	1.26			D - EPN specific	
Dicer-1	Dcr-1	1627580 at	1.26			D - EPN specific	
BRWD3	BRWD3	1640098 at	1.26			D - EPN specific	
Dmel_CG13624	CG13624	1636321 s at	1.25		1.48	G - shared EPN+pathogenic G- bacteria	
debra	dbr	1628243 at	1.25			D - EPN specific	
Spc105-related	Spc105R	1628318 at	1.25			D - EPN specific	
Dmel_CG8449	CG8449	1633082 at	1.25			D - EPN specific	
Dmel_CG4419	CG4419	1624969 s at	1.24	1.0203125		C - shared EPN+wasp	
Serpin 28Dc	Spr28Dc	1636145 at	1.24		1.579484809	E - shared EPN+bac. injection	
growth arrest and DNA damage-inducible gene 45	Gadd45	1625139 at	1.24			G - shared EPN+pathogenic G- bacteria	
Dmel_CG10777	CG10777	1624719 at	1.24		1.16	D - EPN specific	
Brahma associated protein 170kD	Bap170	1628423 at	1.24			D - EPN specific	
Dmel_CG8232	CG8232	1636801 at	1.24			D - EPN specific	
homeodomain interacting protein kinase	hipk	1639306 s at	1.23	0.900560193		C - shared EPN+wasp	
Polycomblike	Pcl	1623525 at	1.23	0.87116529		C - shared EPN+wasp	
slender lobes	sle	1624252 s at	1.23		2.137019888	E - shared EPN+bac. injection	
asterless	ast	1625990 at	1.23			D - EPN specific	
Bub1-related kinase	BubR1	1641015 at	1.23			D - EPN specific	
Dmel_CG15784	CG15784	1640884 at	1.22	2.12216816		A - all infections	
necrotic	nec	1636653 at	1.22	2.349722029	3.943964525	2.285	C - shared EPN+wasp
Daxx-like protein	DLP	1630023 at	1.21			D - EPN specific	
Myosin heavy chain-like	Mhcl	1632231 a at	1.21			D - EPN specific	
mutagen-sensitive 201	mus201	1637542 s at	1.21			D - EPN specific	
Fasclin 2	Fas2	1638956 at	1.21			D - EPN specific	
Dmel_CG15744	CG15744	1639266 at	1.21			D - EPN specific	
Programmed cell death 4 ortholog	Pdcd4	1640020 at	1.21			D - EPN specific	
Hybrid male rescue	Hmr	1641024 at	1.21			D - EPN specific	
Dmel_CG2909	CG2909	1630975 at	1.2	2.771760155		H - missing in bac. Injection	
A kinase anchor protein 200	Akap200	1640838 s at	1.2		2.384756328	2.015	F - missing in wasp infection
Dmel_CG4022	CG4022	1635037 at	1.2	1.2772161		C - shared EPN+wasp	
dachsous	ds	1640627 at	1.2	1.049431232		C - shared EPN+wasp	
Dmel_CG9839	CG9839	1632554 at	1.2			D - EPN specific	
Dmel_CG10936	CG10936	1639703 s at	1.2			D - EPN specific	
Fancd2	Fancd2	1640096 at	1.2			D - EPN specific	
toutatis	tou	1640433 a at	1.2			D - EPN specific	
Cadherin 96Ca	Cad96Ca	1623418 at	1.19		0.399106685	E - shared EPN+bac. injection	
Tenascin accessory	Ten-a	1623755 at	1.19			D - EPN specific	
pr-set7	pr-set7	1624463 s at	1.19			D - EPN specific	
Breast cancer 2, early onset homolog	Brca2	1624941 at	1.19			D - EPN specific	
sister-of-Sex-lethal	ssx	1627150 at	1.19			D - EPN specific	
Ceramidase	CDase	1639396 s at	1.19			D - EPN specific	
mirror	mirr	1639798 at	1.19			D - EPN specific	
Toll	TI	1639321 s at	1.18	3	2.472136618	1.56	A - all infections
Imaginal disc growth factor 1	Idgf1	1632237 at	1.18	2.404311432		C - shared EPN+wasp	
tocan	toc	1641048 a at	1.18	1.033009352		C - shared EPN+wasp	
Transferrin 1	Tsf1	1632430 at	1.18			G - shared EPN+pathogenic G- bacteria	
<i>Na,K-ATPase Interacting</i>	<i>NKAIN</i>	1625992 s at	1.18			D - EPN specific	
odd paired	opa	1628865 at	1.18			D - EPN specific	
DNA ligase 1	DNA-lig1	1630390 at	1.18			D - EPN specific	
Wnk kinase	Wnk	1635378 at	1.18			D - EPN specific	
lethal (1) G0148	l(1)G0148	1639411 at	1.18			D - EPN specific	
male-specific lethal 1	msl-1	1639438 at	1.18			D - EPN specific	
Ankyrin 2	Ank2	1640337 a at	1.18			D - EPN specific	
Cadherin 87A	Cad87A	1641575 at	1.18			D - EPN specific	
Regulatory particle non-ATPase 5	Rpn5	1639091 at	1.17		2.037034174	3.24	E - shared EPN+bac. injection
Dmel_CG9568	CG9568	1639297 at	1.17			G - shared EPN+pathogenic G- bacteria	
Dmel_CG33225	CG33225	1627156 at	1.17			D - EPN specific	
cactus	cact	1629899 at	1.17			D - EPN specific	
retinal degeneration C	rdgC	1634061 a at	1.17			D - EPN specific	
Dmel_CG31140	CG31140	1634447 at	1.17			D - EPN specific	
Dmel_CG10948	CG10948	1636182 a at	1.17			D - EPN specific	
grappa	gpp	1637016 at	1.17			D - EPN specific	
Elongation factor 4a	CG9932	1638653 a at	1.17			D - EPN specific	
Dmel_CG1815	CG1815	1641089 s at	1.17			D - EPN specific	
Down syndrome cell adhesion molecule	Dscam	1637619 s at	1.16	0.818395743		C - shared EPN+wasp	
Smc5	Smc5	1623590 s at	1.16			D - EPN specific	
mini spindles	mmps	1628172 at	1.16			D - EPN specific	
abnormal oocyte	abo	1628418 at	1.16			D - EPN specific	
ATAC complex component 2	Atac2	1632308 at	1.16			D - EPN specific	
Dmel_CG30440	CG30440	1634125 at	1.16			D - EPN specific	
Sulfated	Sulf1	1635007 at	1.16			D - EPN specific	
Dmel_CG11247	CG11247	1636887 s at	1.16			D - EPN specific	
masquerade	mas	1639190 at	1.16			D - EPN specific	
knockout	ko	1641310 at	1.16			D - EPN specific	
SD27140	SD27140	1641365 s at	1.16			D - EPN specific	
Suppressor of cytokine signaling at 36E	Socs36E	1637703 a at	1.15	2.219902805		H - missing in bac. Injection	
expanded	ex	1625970 at	1.15	1.166760884		C - shared EPN+wasp	
terribly reduced optic lobes	trol	1640223 a at	1.15	0.582856592		C - shared EPN+wasp	
Inducer of meiosis 4	Ime4	1624263 at	1.15			D - EPN specific	
windei	wde	1626478 at	1.15			D - EPN specific	
MAN1	MAN1	1631696 s at	1.15			D - EPN specific	
tay bridge	tay	1633390 at	1.15			D - EPN specific	
centrosomin	cmn	1635619 a at	1.15			D - EPN specific	
Cht6	Cht6	1639149 at	1.15			D - EPN specific	
Dmel_CG15523	CG15523	1633846 at	1.14	2.299578059		C - shared EPN+wasp	
Dmel_CG9449	CG9449	1628150 a at	1.14		1.87	G - shared EPN+pathogenic G- bacteria	
Dmel_CG10462	CG10462	1625530 at	1.14			D - EPN specific	
Dmel_CG10376	CG10376	1626553 at	1.14			D - EPN specific	

lethal (3) L1231	I(3)L1231	1629496 at	1.14			D - EPN specific
Dmel_CG4951	CG4951	1631151 at	1.14			D - EPN specific
Dmel_CG5872	CG5872	1635168 at	1.14			D - EPN specific
alan shepard	shea	1637552 s at	1.14			D - EPN specific
enoki mushroom	enok	1639061 at	1.14			D - EPN specific
unknpt	unk	1633020 at	1.13	0.92385834		C - shared EPN+wasps
Ubiquitin-specific protease 64E	Ubp64E	1624727 s at	1.13			D - EPN specific
Dmel_CG11486	CG11486	1626332 s at	1.13			D - EPN specific
convoluted	conv	1629638 at	1.13			D - EPN specific
<i>Na,K-ATPase Interacting</i>	<i>NKAIN</i>	1633752 at	1.13			D - EPN specific
little imaginal discs	lid	1632855 s at	1.13			D - EPN specific
Dmel_CG14614	CG14614	1634430 at	1.13			D - EPN specific
Cadherin 74A	Cad74A	1635742 s at	1.13			D - EPN specific
Integrator 1	IntS1	1636297 at	1.13			D - EPN specific
Lap1	Lap1	1639189 at	1.13			D - EPN specific
lemming A	lmgA	1639494 at	1.13			D - EPN specific
RNA polymerase II 215kD subunit	RpII215	1640764 at	1.13			D - EPN specific
Dmel_CG7922	CG7922	1641352 at	1.13			D - EPN specific
LIM-kinase1	LIMK1	1641452 a at	1.13			D - EPN specific
glass	gl	1623923 s at	1.12			D - EPN specific
Kinesin-like protein at 61F	Klp61F	1624620 at	1.12			D - EPN specific
Dmel_CG13604	CG13604	1625603 at	1.12			D - EPN specific
pickled eggs	pigs	1626090 at	1.12			D - EPN specific
pyramus	pyr	1631552 at	1.12			D - EPN specific
SMC2	SMC2	1634149 at	1.12			D - EPN specific
Dmel_CG30020	CG30020	1634691 a at	1.12			D - EPN specific
Hairless	H	1638568 s at	1.12			D - EPN specific
relative of woc	row	1638839 at	1.12			D - EPN specific
ubiquitin-like protein-specific protease 1	Ulp1	1639435 s at	1.12			D - EPN specific
vestigial	vg	1641470 s at	1.12			D - EPN specific
Enhancer of decapping 3	Edc3	1641685 at	1.12			D - EPN specific
shifted	shf	1635403 at	1.11	2		C - shared EPN+wasps
Argonaute-1	AGO1	1632602 s at	1.11	0.828827339		C - shared EPN+wasps
Dmel_CG32486	CG32486	1624119 at	1.11			D - EPN specific
Transposon.32	SDU1615P	1624543 s at	1.11			D - EPN specific
nucampholin	ncm	1628866 at	1.11			D - EPN specific
combgap	cg	1630717 s at	1.11			D - EPN specific
quaking related 58E-1	qkr58E-1	1631768 at	1.11			D - EPN specific
Dmel_CG8290	CG8290	1636126 at	1.11			D - EPN specific
Activated Cdc42 kinase-like	Ack-like	1637111 a at	1.11			D - EPN specific
unc-104 ortholog	unc-104	1637684 at	1.11			D - EPN specific
prominin-like	CG7740	1641333 s at	1.11			D - EPN specific
Dmel_CG7192	CG7192	1641445 s at	1.11			D - EPN specific
Dmel_CG17836	Xrp1	1622243 a at	1.1		2.17	G - shared EPN+pathogenic G- bacteria
withered	whd	1626147 s at	1.1			D - EPN specific
ariadne	ari-1	1628016 s at	1.1			D - EPN specific
skywalker	sky	1631094 s at	1.1			D - EPN specific
Verprolin 1	Vrp1	1634562 s at	1.1			D - EPN specific
polychaetoid	pyd	1637428 a at	1.1			D - EPN specific
capsuleen	csul	1640862 a at	1.1			D - EPN specific
semaphorin 2a	Sema-2a	1629819 s at	1.09		2.452724324	E - shared EPN+bac. injection
Enhancer of bithorax	E(bx)	1622243 a at	1.09			D - EPN specific
Dmel_CG9004	CG9004	1625269 at	1.09			D - EPN specific
sloppy paired 2	slp2	1627053 at	1.09			D - EPN specific
egghead	egh	1631621 s at	1.09			D - EPN specific
Laminin B2	LanB2	1632666 at	1.09			D - EPN specific
Dmel_CG30389	CG30389	1637947 s at	1.09			D - EPN specific
sec24	sec24	1640083 at	1.09			D - EPN specific
Protein kinase related to protein kinase N	Pkn	1640417 a at	1.09			D - EPN specific
posterior sex combs	Psc	1631095 at	1.08	0.927774215		C - shared EPN+wasps
lethal (3) persistent salivary gland 2	(3)psg2	1621931 at	1.08			D - EPN specific
Dmel_CG42724	CG42724	1625578 at	1.08			D - EPN specific
Additional sex combs	Asx	1628901 at	1.08			D - EPN specific
Dmel_CG3363	CG3363	1631645 at	1.08			D - EPN specific
Transport and Golgi organization 6	Tango6	1636939 at	1.08			D - EPN specific
Bric-a-brac interacting protein 2	bip2	1639551 at	1.08			D - EPN specific
antiapoptosis clone 11	Aac11	1623100 at	1.07			D - EPN specific
Dorsal switch protein 1	Dsp1	1624304 s at	1.07			D - EPN specific
polymerase eta	DNAPol-η	1626494 at	1.07			D - EPN specific
Meltrin	Meltrin	1627649 at	1.07			D - EPN specific
crumbs	crb	1628146 at	1.07			D - EPN specific
Dmel_CG9754	CG9754	1628669 at	1.07			D - EPN specific
Dmel_CG5674	CG5674	1630550 a at	1.07			D - EPN specific
jim	jim	1636911 at	1.07			D - EPN specific
Dmel_CG12179	CG12179	1637361 a at	1.07			D - EPN specific
Myb-interacting protein 130	mip130	1639185 at	1.07			D - EPN specific
fat facets	faf	1640541 at	1.07			D - EPN specific
crooked legs	crol	1626018 s at	1.06			D - EPN specific
tonalli	tna	1625730 s at	1.06			D - EPN specific
Dmel_CG32206	CG32206	1628159 a at	1.06			D - EPN specific
Negative elongation factor A	Nelf-A	1631550 at	1.06			D - EPN specific
Reduction in Cnn dots 5	Rcd5	1632420 at	1.06			D - EPN specific
Dmel_CG9449	CG9449	1633200 at	1.06			D - EPN specific
Son of sevenless	Sos	1633335 at	1.06			D - EPN specific
Dmel_CG2519	CG2519	1633684 at	1.06			D - EPN specific
Dmel_CG6409	CG6409	1634278 at	1.06			D - EPN specific
capicua	cic	1635909 at	1.06			D - EPN specific
Dmel_CG30007	CG30007	1637712 at	1.06			D - EPN specific
Hexokinase A	Hex-A	1625638 a at	1.05		7.469895514 1.735	F - missing in wasp infection
Dmel_CG31496	CG31496	1628103 at	1.05			D - EPN specific
Furin 1	Fur1	1628952 s at	1.05			D - EPN specific
Distal-less	Dll	1630237 a at	1.05			D - EPN specific
Spinophilin	Spn	1633021 s at	1.05			D - EPN specific
Dmel_CG8389	CG8389	1637774 s at	1.05			D - EPN specific
lola like	lolal	1640311 s at	1.05			D - EPN specific
drongo	CG3365	1623693 a at	1.04		1.93	G - shared EPN+pathogenic G- bacteria
leonardo	14-3-3ζ	1625148 s at	1.04			D - EPN specific
Cyclin B3	CycB3	1626454 at	1.04			D - EPN specific
RhoGEF3	RhoGEF3	1626756 a at	1.04			D - EPN specific
Trithorax-like	Trl	1628275 at	1.04			D - EPN specific
<i>Prosap</i>	<i>Prosap</i>	1629160 s at	1.04			D - EPN specific
Dmel_CG12155	CG12155	1622471 at	1.04			D - EPN specific
bves	bves	1633387 at	1.04			D - EPN specific
wallenda	wnd	1633727 s at	1.04			D - EPN specific
Dmel_CG5604	CG5604	1634156 at	1.04			D - EPN specific
Dmel_CG8229	CG8229	1639048 a at	1.04			D - EPN specific
hephaestus	heph	1637478 s at	1.03	0.98616031 3.42258756		B - missing in pathogenic bacteria
worniu	wor	1631502 at	1.03	0.990381437		C - shared EPN+wasps
zipper	zip	1622840 a at	1.03	0.938871223		C - shared EPN+wasps
Suppressor of Cytokine Signaling at 16D	Socs16D	1636122 at	1.03	0.857330289		C - shared EPN+wasps
embargoed	emb	1622939 at	1.03	0.797073236		C - shared EPN+wasps
scalloped	sd	1625515 a at	1.03	0.755818539		C - shared EPN+wasps
echinoid	ed	1627506 at	1.03		0.318416425	E - shared EPN+bac. injection
Dmel_CG14562	CG14562	1623670 at	1.03			D - EPN specific
HERC2	HERC2	1629145 at	1.03			D - EPN specific
Kruppel homolog 1	Kr-h1	1631481 a at	1.03			D - EPN specific
Dmel_CG17078	CG17078	1631919 at	1.03			D - EPN specific
Nijmegen breakage syndrome	nbs	1632161 at	1.03			D - EPN specific
mastermind	mam	1632457 s at	1.03			D - EPN specific
Dmel_CG10508	CG10508	1623133 a at	1.02			D - EPN specific
Bre1	Bre1	1624301 at	1.02			D - EPN specific
Dmel_CG9702	CG9702	1634174 at	1.02			D - EPN specific
RNA-binding protein 1	Rbp1	1638486 at	1.02			D - EPN specific
Serpin 88Eb	Spn88Eb	1636946 at	1.01	12	3.468032833	B - missing in pathogenic bacteria
mitochondrial RNA polymerase	mtrNApol	1626738 at	1.01			D - EPN specific
Dmel_CG710	CG710	1633876 at	1.01			D - EPN specific
Dmel_CG14322	CG14322	1634309 at	1.01			D - EPN specific
domeless	dome	1632381 at	1	1.558303521		C - shared EPN+wasps
pebbled	peb	1622949 at	1	0.839048644		C - shared EPN+wasps
coronin	pod1	1641620 s at	1		2.009326839	E - shared EPN+bac. injection
Dmel_CG1233	CG1233	1625945 a at	1			D - EPN specific
scaf6	scaf6	1627184 at	1			D - EPN specific

ballchen	ball	1628342 s at	1			D - EPN specific
Dmel_CG13933	CG13933	1628791 at	1			D - EPN specific
Dmel_CG8366	CG8366	1632519 at	1			D - EPN specific
Dmel_CG15628	CG15628	1636248 at	1			D - EPN specific
piopio	plo	1638681 at	1			D - EPN specific
Zinc finger AT-hook protein	D19A	1640054 at	1			D - EPN specific
escargot	esg	1641639 at	1			D - EPN specific
Spermidine Synthase	Spd5	1630830 a at	-1	2.10757595		E - shared EPN+bac. injection
Dmel_CG13018	CG13018	1629541 at	-1			D - EPN specific
Dmel_CG33169	CG33169	1632606 a at	-1			D - EPN specific
Glutathione S transferase E3	GstE3	1637129 at	-1.01	6.271921203		E - shared EPN+bac. injection
Dmel_CG7607	CG7607	1625684 at	-1.01			D - EPN specific
Dmel_CG7231	CG7231	1633333 a at	-1.01			D - EPN specific
Dmel_CG14671	CG14671	1637379 at	-1.01			D - EPN specific
Dmel_CG33013	CG33013	1637916 at	-1.01			D - EPN specific
Dmel_CG2540	CG2540	1640452 at	-1.01			D - EPN specific
Bekka	Bka	1627349 at	-1.03			D - EPN specific
Dmel_CG34229	CG34229	1634245 at	-1.03			D - EPN specific
Dmel_CG31957	CG31957	1641624 at	-1.03			D - EPN specific
mitochondrial ribosomal protein L10	mRpl10	1624012 at	-1.04			D - EPN specific
Dmel_CG14210	CG14210	1624635 at	-1.04			D - EPN specific
Dmel_CG13516	CG13516	1638764 at	-1.04			D - EPN specific
Dmel_CG11875	CG11875	1627803 at	-1.05			D - EPN specific
Dmel_CG34053	CG34053	1626658 at	-1.06			D - EPN specific
Jonah 25Bii	Jon25Bii	1639196 at	-1.07	1.197022279		C - shared EPN+wasps
Dmel_CG2680	CG2680	1630821 at	-1.07			D - EPN specific
Dmel_CG8026	CG8026	1633840 a at	-1.07			D - EPN specific
cornichon related	cnrl	1638047 at	-1.07			D - EPN specific
Dmel_CG9117	CG9117	1632406 at	-1.08			D - EPN specific
Dmel_CG14057	CG14057	1633054 at	-1.08			D - EPN specific
Dmel_CG5835	CG5835	1627354 at	-1.09			D - EPN specific
Dmel_CG32262	CG32262	1633691 at	-1.09			D - EPN specific
Dmel_CG6041	CG6041	1627432 at	-1.11	1.282119657		C - shared EPN+wasps
Dmel_CG8021	CG8021	1624984 at	-1.11			D - EPN specific
Dmel_CG10778	CG10778	1630701 at	-1.11			D - EPN specific
mitochondrial ribosomal protein L36	mRpl36	1624765 at	-1.12			D - EPN specific
Dmel_CG14544	CG14544	1634023 at	-1.12			D - EPN specific
Dmel_CG9804	CG9804	1635849 at	-1.12			D - EPN specific
Dmel_CG15706	CG15706	1640917 at	-1.12			D - EPN specific
Dmel_CG13690	CG13690	1633171 at	-1.13	2.310195228		C - shared EPN+wasps
garnysstan	gny	1636630 s at	-1.13	1.093143152		C - shared EPN+wasps
technical knockout	tko	1626534 at	-1.13			D - EPN specific
Dmel_CG4186	CG4186	1629259 at	-1.13			D - EPN specific
Dmel_CG15985	CG15985	1629689 at	-1.13			D - EPN specific
Dmel_CG30010	CG30010	1637118 at	-1.14			D - EPN specific
Dmel_CG42235	CG42235	1627633 at	-1.15			D - EPN specific
Dmel_CG14463	CG14463	1640424 at	-1.15			D - EPN specific
Dmel_CG11781	CG11781	1630268 at	-1.16			D - EPN specific
Dmel_CG32448	CG32448	1638634 at	-1.16			D - EPN specific
Dmel_CG31229	CG31229	1630351 at	-1.17			D - EPN specific
Dmel_CG1488	CG1488	1630505 a at	-1.17			D - EPN specific
Dmel_CG15152	CG15152	1633949 at	-1.17			D - EPN specific
Dmel_CG31084	CG31084	1627438 at	-1.18			D - EPN specific
Dmel_CG33170	CG33170	1631473 at	-1.19			D - EPN specific
Dmel_CG7460	CG7460	1625149 at	-1.2			D - EPN specific
Dmel_CG31720	CG31720	1632145 a at	-1.21			D - EPN specific
Dmel_CG10300	CG10300	1636387 at	-1.21			D - EPN specific
mitochondrial ribosomal protein S18C	mRpS18C	1628621 at	-1.23			D - EPN specific
Dmel_CG7916	CG7916	1623817 at	-1.24			D - EPN specific
Dmel_CG31323	CG31323	1635371 at	-1.24			D - EPN specific
Dmel_CG10505	CG10505	1639363 at	-1.24			D - EPN specific
ZnT35C	CG3994	1639619 a at	-1.24			D - EPN specific
Dmel_CG13217	CG13217	1623464 at	-1.25			D - EPN specific
Dmel_CG32669	CG32669	1623556 at	-1.26			D - EPN specific
Dmel_CG13014	CG13014	1631684 at	-1.26			D - EPN specific
Allatostatin C	Ast-C	1634436 at	-1.26			D - EPN specific
Dmel_CG17036	CG17036	1628221 at	-1.27			D - EPN specific
Dmel_CG7006	CG7006	1633247 at	-1.27			D - EPN specific
Dmel_CG1134	CG1134	1633980 at	-1.27			D - EPN specific
Dmel_CG13066	CG13066	1631695 at	-1.29			D - EPN specific
fuseless	fusl	1640365 s at	-1.29			D - EPN specific
Dmel_CG15456	CG15456	1625166 at	-1.31	1.03422802		C - shared EPN+wasps
Dmel_CG13227	CG13227	1630218 at	-1.31			D - EPN specific
Niemann-Pick type C-2f	Npc2f	1634158 at	-1.31			D - EPN specific
Lcp65APsi	CR18775	1631002 at	-1.32	1.338640275		C - shared EPN+wasps
Dmel_CG18598	CG18598	1632319 at	-1.32			D - EPN specific
Dmel_CG15706	CG15706	1634197 at	-1.32			D - EPN specific
Dmel_CG34172	CG34172	1625901 s at	-1.33			D - EPN specific
Dmel_CG13024	CG13024	1633264 at	-1.34			D - EPN specific
Cyp9f3Psi	CG17875	1640584 at	-1.35			D - EPN specific
selenocysteine methyltransferase	CG10621	1626086 at	-1.36	3.322991509	0.77	B - missing in pathogenic bacteria
Dmel_CG14701	CG14701	1627723 at	-1.36			D - EPN specific
Dmel_CG9672	CG9672	1635525 at	-1.36			D - EPN specific
Tetraspanin 42E	Tsp42E	1634977 at	-1.37			G - shared EPN+pathogenic G- bacteria
Dmel_CG1532	CG1532	1623959 at	-1.38			D - EPN specific
Dmel_CG31789	CG31789	1630692 at	-1.43			D - EPN specific
Dmel_CG18643	CG18643	1638845 at	-1.43			D - EPN specific
Dmel_CG3690	CG3690	1628284 at	-1.45			D - EPN specific
Dmel_CG14572	CG14572	1630725 at	-1.45			D - EPN specific
Tak1-like 2	Takl2	1632359 at	-1.45			D - EPN specific
Glutathione S transferase E7	GstE7	1640065 at	-1.46			E - shared EPN+bac. injection
Dmel_CG14610	CG14610	1636683 at	-1.48	11		C - shared EPN+wasps
Juvenile hormone esterase	Jhe	1637833 at	-1.49			D - EPN specific
Ugt86Dj	Ugt86Dj	1634029 at	-1.51			D - EPN specific
Dmel_CG33282	CG33282	1625342 at	-1.53			D - EPN specific
Dmel_CG13085	CG13085	1631382 at	-1.55			D - EPN specific
Dmel_CG11825	CG11825	1635263 at	-1.56			G - shared EPN+pathogenic G- bacteria
Dmel_CG9521	CG9521	1626597 at	-1.58	1.36		D - EPN specific
Dmel_CG17571	CG17571	1635878 s at	-1.59			D - EPN specific
Insulin-like peptide 3	Ilp3	1627943 at	-1.61			D - EPN specific
Dmel_CG14141	CG14141	1641102 at	-1.61			D - EPN specific
Dmel_CG7130	CG7130	1628660 at	-1.63			D - EPN specific
Dmel_CG15336	CG15336	1632970 at	-1.65			D - EPN specific
Cuticular protein 67Fb	Cpr67Fb	1638742 at	-1.7	1.19722855		C - shared EPN+wasps
Jonah 65Ai	Jon65Ai	1636460 at	-1.75			G - shared EPN+pathogenic G- bacteria
Dmel_CG12490	CG12490	1636604 at	-1.75			D - EPN specific
Dmel_CG34112	CG34112	1630404 at	-1.77	0.409131603		C - shared EPN+wasps
Dmel_CG34267	CG34267	1626025 at	-1.78			D - EPN specific
Dmel_CG15554	CG15554	1634286 at	-1.78			D - EPN specific
Dmel_CG9394	CG9394	1636970 at	-1.83			D - EPN specific
Dmel_CG16771	CG16771	1630977 at	-1.84			D - EPN specific
Dmel_CG8299	CG8299	1633031 at	-1.84			D - EPN specific
Dmel_CG17681	CG17681	1633142 at	-1.85			D - EPN specific
Ecdysone-induced protein 63F 2	Eip63F-2	1639350 at	-1.85			D - EPN specific
Dmel_CG8560	CG8560	1639641 at	-1.87	1.215745914		H - missing in bac. Injection
Dmel_CG34451	CG34451	1635274 at	-1.89			D - EPN specific
Dmel_CG5883	CG5883	1637889 at	-1.95			D - EPN specific
Dmel_CG32379	CG32379	1628335 at	-2.15	0.463476071		C - shared EPN+wasps
Dmel_CG15155	CG15155	1623491 at	-2.15			D - EPN specific
Dmel_CG13012	CG13012	1625861 at	-2.17			D - EPN specific
Dmel_CG9826	CG9826	1625203 at	-2.22			D - EPN specific
Adipokinetic hormone	Akh	1631816 at	-2.26			D - EPN specific
Dmel_CG3344	CG3344	1632584 at	-2.31	1.177934444	0.865	H - missing in bac. Injection
Dmel_CG4363	CG4363	1630740 at	-2.33			D - EPN specific
Glutathione Synthetase	GS	1627321 x at	-2.35			D - EPN specific
Dmel_CG7881	CG7881	1623229 at	-2.4			D - EPN specific
Dmel_CG32750	CG32750	1634445 at	-2.41			D - EPN specific
Dmel_CG9903	CG9903	1624333 at	-2.55			D - EPN specific
Dmel_CG15773	CG15773	1637326 at	-2.85			D - EPN specific
Dmel_CG32483	CG32483	1628500 at	-3.08	1.163031446		C - shared EPN+wasps
Dmel_CG13135	CG13135	1625802 a at	-3.58			D - EPN specific

Gene identified by more transcripts in italics.

KEGG ID	KEGG path	FDR	P-value
dme04914	Progesterone-mediated oocyte maturation	2.42e-05	3.84e-07
dme04310	Wnt signaling pathway	0.00467	0.000148
dme04120	Ubiquitin mediated proteolysis	0.00988	0.000471
dme04630	Jak-STAT signaling pathway	0.0197	0.00156
dme04340	Hedgehog signaling pathway	0.0197	0.00153
dme04320	Dorso-ventral axis formation	0.0209	0.00199
dme04512	ECM-receptor interaction	0.0276	0.00306
dme03450	Non-homologous end-joining	0.0582	0.00739
dme03018	RNA degradation	0.0718	0.0103
dme00310	Lysine degradation	0.073	0.0116
dme03040	Spliceosome	0.0852	0.0149
dme03420	Nucleotide excision repair	0.104	0.0198
dme03440	Homologous recombination	0.112	0.0231
dme04013	MAPK signaling pathway - fly	0.161	0.0357
dme03013	RNA transport	0.179	0.0425

Arefin, Supplementary table 4

Arefin, Table S5

Name	Fold (compared to naive larval expression)	Function	Product
CG4250	no naive larval expression	circadian cycling in heads	LPS-induced tumor necrosis factor alpha type
Lectin24Db	no naive larval expression	galactose binding lectin	C-type lectin-like
Lectin28C	59.62	galactose binding lectin	C-type lectin-like
Eater	43.67	cell adhesion, recognition, phagocytosis,	bacterial cell surface binding protein
Scavenger receptor class C, type 1 (SR-C1)	25.44	scavenger receptor	Complement control module
Glutactin	19.67	extracellular matrix glycoprotein	Carboxylesterase type B
SPARC	13.75	ligand binding or carrier	SPARC/osteonectin-like

Arefin Table S6

		Gene Name	Abbreviation	MicroArrays (log2 FC)	qRT-PCR (log2 FC)	EPN Assay RNAi	EPN assay mutant	Detected in larval immune response
Candidates from our Microarray study	Recognition molecules	Immune induced molecule 3	IM3	8.99	9.59	ns		A - all infections
		Diptericin B	DptB	7.29	5.33	ns		A - all infections
		GNBP-like	CG13422	7.1	3.73	FB		B - missing in pathogenic bacteria
		Immune induced molecule 2	IM2	6.84		ns		B - missing in pathogenic bacteria
		Drosomycin	Drs	6.02	7.27	ns		A - all infections
		Metchnikowin	Mtk	5.5		ns		A - all infections
		Attacin-A	AttA	5.44	3.55	FB		A - all infections
		Peptidoglycan recognition protein SB1	PGRP-SB1	5.27	3.77	ns		A - all infections
		Thioester-containing protein 1	Tep1	2.66	2.82			C - shared EPN+wasps
		Dmel_CG6639	SPH93	2.47		ns		A - all infections
		starry night	stan	2.32		ns		D - EPN specific
		Immune induced molecule 18	CG10332/IM18	1.83		FB		D - EPN specific
		A kinase anchor protein 200	Akap200	1.2		ns		F - missing in wasp infection
		Dmel_CG7607	CG7607	1.01	3.78			D - EPN specific
		Argonaute 2	AGO2	0,79*		ns		E - shared EPN+bac. injection
		Spermidine Synthase	SpdS	-1	-1.23			E - shared EPN+bac. injection
		Peptidoglycan recognition protein LF	PGRP-LF	1.8	0,44*		sensitive	D - EPN specific
		Thioester-containing protein 2	Tep2	1.99	2.47		ns	B - missing in pathogenic bacteria
		Thioester-containing protein 4	Tep4	0,73*			ns	B - missing in pathogenic bacteria
		Thioester-containing protein 3	Tep3				sensitive	
		Gram-negative bacteria binding protein 3	GNBP3				ns	bac. injection
		Peptidoglycan recognition protein SA	PGRP-SA				ns	shared wasp+bac. injection
		Peptidoglycan recognition protein LC	PGRP-LC				ns	
		Peptidoglycan recognition protein LE	PGRP-LE				ns	
		Hemocytes enriched genes		Glutactin	Git	1.29	1.96	Hem
dSR-CI	CG4099					ns		wasp specific
Eater	CG6124					ns		
Lectin28C	CG7106					ns		
SPARC	CG6378					ns		
Lectin24DB	CG2958					ns		
Dmel_CG4250	CG4250					ns		wasp specific
larval translucida	Iti					ns		
Wrinkled	W					ns		

* - statistically significant but $|\log_2FC| < 1$

FB - significantly sensitive to EPN infection after knockdown in fat body (driver PPL-Gal4)

Hem - significantly sensitive to EPN infection after knockdown in hemocytes (driver He-Gal4)

ns - non-significant response to EPN

Table S1. Supplementary Material and Methods: A: Gene-specific primers used for confirmation of the array data by qPCR: see Figure 3 for results. B:

***Drosophila* stocks used for nematode infections (see Fig. 1, 4-6 for results)**

Table S2. The 100 most strongly regulated genes are enriched in immune genes.

Gene set enrichment analysis for the 100 most highly regulated genes in Fig 2 A was performed using Amigo (http://amigo.geneontology.org/cgi-bin/amigo/term_enrichment?session_id=).

Table S3. Complete set of transcripts that are regulated in different infection

models. The differentially expressed transcripts in *Drosophila* larvae after infection with common G- and G+ bacteria [3,4], pathogenic G- bacteria [4], wasps [5-7] and nematodes (this work) are shown.

Table S4. Significantly enriched KEGG pathways after gene set enrichment

analysis (GSEA). Enriched pathways after Fisher test for upregulated subset of significantly changed transcripts (logFC cutoff > 0.4 (TRUE), Q-VALUE cutoff 0.05:left column; and P-value at cutoff 0.05: right column) are shown.

Table S5: Hemocytes-enriched genes and their function. Genes were analyzed for enriched expression using previously published data [3].

Table S6: Comparison of functional analysis results and qPCR results with

microarray data. Complete set of genes used in our functional analysis and qPCR compared to microarray results.