

	Gene Symbol	Gene name	GenBank	Primer-Fw (5'-3')	Primer-Rv (5'-3')	Reference
Reference	<i>Lv</i> L40	Ubiquitin/ribosomal L40 fusion protein	KJ831563	GAGAATGTGAAGGCCAAGATC	TCAGAGAGAGTGC GACCATC	1
	<i>Lv</i> EF1 $\alpha$	Elongation factor 1-alpha	GU136229	TGGCTGTGAACAAGATGGACA	TTGTAGCCCACCTTCTTGACG	2
	<i>Lv</i> Actin	$\beta$ -actin	AF300705	CCACGAGACCACCTACAAC	AGCGAGGGCAGTGATTTC	3
Antimicrobial peptide	<i>Litvan</i> PEN1/2	Penaeidin 1/2	Y14925	GTCTGCCTGGTCTTCTTGG	GCAGCAATTGCGAGCATCTG	4
	<i>Litvan</i> PEN3	Penaeidin 3	AF390144	GTCTGCCTGGTCTTCTTGG	GCGCTTGTGAGAAGGAAATTC	4
	<i>Litvan</i> PEN4	Penaeidin 4	DQ211701	GTCTGCCTGGTCTTCTTGG	TTGCGGAGGGAATCCGTAG	4
	<i>Litvan</i> ALF-A	ALF Group A	EW713395	CTGATTGCTCTTGTGCCACG	TGACCCATGA ACTCCACCTC	5
	<i>Litvan</i> ALF-B	ALF Group B	GQ227486	GTGTCTCCGTGTTGACAAGC	ACAGCCCAACGATCTTGCTG	5
	<i>Litvan</i> ALF-C	ALF Group C	FE058235	ATGCGAGTGTCTGTCTCAG	TGAGTTTGTTCGCGATGGCC	5
	<i>Litvan</i> ALF-D	ALF Group D	FE151634	TGTGTTGGTTGTGGCACTGG	CAACGAGGTCAATGTCACCG	5
	<i>nLv</i> ALF1	new ALF-1	KJ000049	AAGCTCTCATTCTGGTTCGG	GGGTGTAACGAAGTACGTGC	this study
	Crustin	Crustin Type IIa	AY488496	CGAACCAGAGACACCTGTTG	CAGCACACTTGTAGTCGTTG	4
	c-type lys	conventional c-type lysozyme	AY170126	CTTCGAGCTCTATGTTCTAGA	TCAACCATTACAACGTCTGCA	4
	i-type lys	invertebrate i-type lysozyme	DQ398564	CGGTTGCGGAATTAGAAGAC	TCCACACACTTCTGACCATG	this study
Immune signaling pathway	<i>Lv</i> Toll1	Toll receptor 1	DQ923424	TCTGCTGACGAATACTCCCAAAC	GACTGCCACGCGATTTC	6
	<i>Lv</i> Toll2	Toll receptor 2	JN180637	CATGCCTGCAGGACTGTTTA	GGCCTGAGGGTAAGGTCTTC	7
	<i>Lv</i> Toll3	Toll receptor 3	JN180638	TCGTACAACCAGCTGACGAG	ATACTTCAGGTGGGCCACAG	7
	<i>Lv</i> Pelle	Pelle (IRAK4)	JN180645	TGACAGATTCCAGTGATCATGC	CTTCCAAGATGTGCGCTAACTGTG	8
	<i>Lv</i> TRAF6	TNFR-associated factor 6	KC346864	GTTTATGGCTACTCGGCAGATCC	CATGTGAGAGCTTTTGTGGGACTTC	4
	<i>Lv</i> Cactus	Cactus (I $\kappa$ B protein)	JX014314	AGTGACCTGAGAACCCTTCAC	CACACACTGACACTATCTCC	this study
	<i>Lv</i> Dorsal	Dorsal (Rel/NF- $\kappa$ B transcription factor)	FJ998202	TCACTGTTGACCCACCTTAC	GAGGGCATCTCAACATCAC	4
	<i>Lv</i> IMD	Immune deficiency homolog	FJ592176	AAGGTCGAGGTCAGCGAAC	GATCATCTGGGTGAGTCTGG	4
	<i>Lv</i> STAT	Signal transducer and activator of transcription	HQ228176	CATGGCTCTGGCAGATAAGC	GCATCTGCTGGTCTCTCTTC	4
	<i>Lv</i> Relish	Relish (Rel/NF- $\kappa$ B transcription factor)	EF432734	GAGGAGGCTTGTGCTATGC	CAGGTACAGGTCTGGTTAGG	4
	<i>Lv</i> GILT	$\gamma$ -interferon-inducible lysosomal thiol reductase	HQ317497	TCCACCTGTACTACGAGTCC	CGAATGGGAACATCTCCACC	4
<i>Lv</i> HMGBa	High-mobility group	HQ228174	GGATGAGAAGGTCAAGTACG	TCCTCGTCGTCTTCTCTC	4	
<i>Lv</i> DOME	JAK/STAT pathway receptor Domeless	KC346866	CTCAGGCTATGTTTCTCAGGATTCA	CACGGCAGTTCCTTATGGTCT	9	

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Microbial recognition	<i>Lv</i> LGBP	LPS and $\beta$ -1,3-glucan binding protein	EU102286	ACCGCAGCATCAGTTATACC	GTCATCGCCCTTCCAGTTG	3
	<i>Lv</i> $\beta$ GBP	HDL/1,3-beta-D-glucan-binding protein	AY249858	ACGAGAACGGACAAGAAGTG	TTCAGCATAGAAGCCATCAGG	this study
	<i>Lv</i> QM	QM protein	JX880087	TCGTGTGCTGGTGCTGATAGAT	GCCTCAATGACCTGCTCCTTGT	10
	<i>Lv</i> LT	C-type lectin	DQ871245	TCTGTTTCGAGGGAAAGTAG	GAACAAGAGGCATTTGTCCG	this study
	<i>Lv</i> Lectin-1	C-type lectin 1	HQ153047	CTTCTGGATCGGAGGGACTGA	TCGTCTTAGATTTGGCAGATAGCA	11
	<i>Lv</i> CTL3	C-type lectin 3	KF156943	ATGTTCTTCGTGCTCCTGCTGT	GCAGTGGTCGTAATGTTGTG	10
	<i>Lv</i> AV	Antiviral Protein (c-type lectin)	JX983205	AACGATCATGAGGACAGGTG	AATGAATCCCAGCGCATGAC	this study
	<i>Lv</i> Gal	Galectin	KF179103	AATGCTTTCACACGAGCAGC	GTCTTCAATCCAGACACAGG	this study
RNAi pathway	<i>Lv</i> Sid1	Sid-1	HM234688	GAAGCGATTGGCAGTCTATGAAC	TGGAAGCCTATCTCTGCAACTG	12
	<i>Lv</i> Dcr1	Dicer-1	EU676241	CCGAGATAGAACGGTTCAGTG	CGATAATTCCTCCCAACACCTG	13
	<i>Lv</i> Dcr2	Dicer-2	HQ541163	AGGAAATGCAATGTCGTGGTT	ACGAGCCCTCCCCCTAGATT	14
	<i>Lv</i> Ago1	Argonaute 1	HM234689	TGCGTCATTTGCCATCCAT	GCCATCTGGAGCGGAGAAG	12
	<i>Lv</i> Ago2	Argonaute 2	HM234690	GATGGCATGAAGTCTGCAGTTG	TGCGCACGACCATCACTAAG	12
	<i>Lv</i> TRBP1	(TAR) RNA-binding protein 1	HQ541157	CAAGAGTTGTGTATGCGCAG	AGCACCACCCTGGATGATG	4
proPO system	<i>Lv</i> proPO1	Prophenoloxidase 1	EU284136	CGGTGACAAAGTTCCTCTTC	GCAGGTCGCCGTAGTAAG	3
	<i>Lv</i> proPO2	Prophenoloxidase 2	EU373096	GCGGAATGACTTTGGTCTCG	AAGTGACGTCATCGGAGAGG	4
	<i>Lv</i> PPAE2	Prophenoloxidase-activating enzyme 2	FE090793	TCAGCAGGATTTGTGTTGCC	GTCTCAGTGAAGCCCCATC	4
	<i>Lv</i> $\alpha$ 2M-1	Alpha2-macroglobulin 1	EF073268	GGAGAGGAACGAGACCATC	TCAGCTTGGCGTCGAAGTG	4
	<i>Lv</i> $\alpha$ 2M-2	Alpha2-macroglobulin 2	DQ988330	TGCAGGTTCTAGTGTGGTAC	CACCAGGTAGTCGATGAC	4
Cytokine	<i>Lv</i> SVC1	single VWC domain protein 1	HQ541158	TGCCAACCTAGCTGAGCTTCA	CCATCAGGGCACACGTATTTG	14
	<i>Lv</i> AST	Astakine	HM473285	GACTGGGCGACTGTTCTTC	GGTACGCAAGATTGAGCTCC	4
	<i>Lv</i> Tetrasp	Tetraspanin D107	FE084500	GCTTGGCATCCTGAGAGAG	GCTGATTGGAATTGCCTTCG	4
	<i>Lv</i> Clq	Clq-binding protein	KC776604	TGAAGGAGAGCTCAAGAGTC	CCTCATACATGGTGAGTTCG	this study
	<i>Lv</i> Pxt	Peroxinectin	AY486425	CTGAAGAAAGGAGACCGATAC	TGGACGGCTTGATGTTATC	4
Redox system	<i>Lv</i> cMnSOD	Cytosolic manganese superoxide dismutase	AY486424	TCATGCTTTGCCACCTCTC	CCGCTTCAACCACTTCTTC	3
	<i>Lv</i> GST	Glutathione S-transferase (GST mu-class)	AY573381	CACCTACGAACACTACGAAC	GGTCTTGAAGCCGTCGAG	4
	<i>Lv</i> GPx	Glutathione peroxidase	AY973252	AGAGTTCGGCGACAAGCTG	TCGGTAGTGTTCTCCTGGTG	4
	<i>Lv</i> NOS	Nitric oxide synthase	GQ429217	GAGCAAGTTATTCGGCAAGGC	TCTCTCCAGTTTCTTGGCGT	15

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