

S2 Table: List of PCR primers used in this study. The primers are used for qPCR.

Target gene	Sequence (5'-3')	Reference
REFERENCE GENES		
Actin	F: TGCCAACACTGTCCTTCTG R: AGAATTGACCCACCAATCCA	[1]
Enolase	F: GGTGATGAAGGTGGTTTGC R: GATGCAGCAACATCCATACC	[2]
GAPDH	F: GATGCACCCATGTTGTTG R: TTTGCAGAAGGTGCATCAAC	[2]
eIF3-S8	F: TGAGTGTCTGCTATGGATTGC R: TCGCGGCTCGTGGTA	[3]
MGST	F: TTGCTCTGTAAGGTTGGTTGC R: TGTCTGGTTAACTACAATCCTCCTG	[4]
RP49	F: CGTCATATGTTGCCAACT R: TTGAGCACGTTCAACAAAT	[5]
RPL8	F: TGGATGTTAACAGGGTTCAT R: CTGGTGGTGACGTATTGATAA	[6]
RPL13a	F: TGGCCATTACTGGTCGTT R: GAGCACGGAAATGAAATGGT	[2]
RPS5	F: AATTATTGGTCGCTGGAATT R: TAACGTCCAGCAGAACATGGTA	[6]
RPS18	F: GATTCCCAGTTGGTTTG R: CCCAATAATGACGCAAACCT	[2]
TBP-af	F: TTGGTTTCATTAGCTGCA R: ACTGCGGGAGTCAAATCT	[7]
Target genes		
AChE-1	F: AGTTGGCGAGATATGGTG R: GGAAAAATAGAGCGCGTGAG	This study
AChE-2	F: AACCGGCTTAAGGATTGAT R: CCCTCTCCCTTCAATCTTC	This study
CYP6AS1	F: GCGACCAATGCGAATGAAAC R: TCACGGCATTCCACCATTTC	This study
CYP6AS3	F: TCGAAAGGGACGAGGATATG R: AGTCATGGATGCCTACTGG	This study
CYP6AS4	F: GGCTGGATTGAAACGTAT R: CGCGTGGATTCTTCATT	This study
CYP6AS10	F: TTTCTAAATTGCCCCATC R: CGGGGACAATTCTGTTCTTA	This study
CYP9Q1	F: GTTCACGTCGAGCAAGATCA R: TCGTCAACACGCTTCAAC	This study
CYP9Q2	F: CCTGATCAAGAGCATCACGA R: GATCTGCTGAGGTGAAGG	This study
CYP9Q3	F: GTAGCCATTACCGCGTTCAC R: GTCTCGTCGATCTCCTGCTG	This study
CYP450	F: GGCGGGTAAAATGGTGTTC R: AGGATGGCAACCCATCACTG	This study
Abaecin	F: CAGCATTGCGATACGTACCA R: GACCAAGAAACGTTGAAAC	[6]
Apisimin	F: TGAGCAAAATCGTTGCTGTC R: AACGACATCCACGTTGATT	[6]
Defensin1	F: TGGCCTGCTAACTGTCTCAG R: AATGGCACTTAACCGAAACG	[6]
Defensin2	F: GCTGCTACCACTACGACATC R: CAACTACCGCCTTACGTCG	[6]

Vitellogenin	F: ACGTAATAAATGCCGCCAAG R: TGCATGTTGCTCTCCAACTC	This study
--------------	--	------------

GAPDH: glyceraldehyde 3-phosphate dehydrogenase; MGST: microsomal glutathione s-transferase; eIF: eukaryotic initiation factors; PRM: protamine; RP: ribosomal proteins; RPL: ribosomal proteins of the large subunit; RPS: ribosomal proteins of the small subunit; SRC-C: scavenger receptor class C; TARBP: TAR RNA-binding protein; TBP-af: TATA box binding protein - associated factor; TRBP: TAR RNA-binding protein; AChE: Acetylcholinesterase; CYP: cytochrome P

- [1] Cunha AD, Nascimento AM, Guidugli KR, Simoes ZLP, Bitondi MMG. (2005) Molecular cloning and expression of a hexamerin cDNA from the Honey bee, *Apis mellifera*. *J Insect Physiol* 51:1135–47.
- [2] Scharlaken B, de Graaf DC, Goossens K, Brunain M, Peelman LJ, Jacobs FJ (2008) Reference gene selection for insect expression studies using quantitative real-time PCR: The head of the Honey bee, *Apis mellifera*, after a bacterial challenge. *J Insect Science* 8: 33
- [3] Fisher P, Grozinger C.M. (2008) Pheromonal regulation of starvation resistance in Honey bee workers (*Apis mellifera*). *Naturwissenschaften* 95:723-729
- [4] Cornman RS, Tarpy DR, Chen Y, Jeffreys L, Lopez D, Pettis JS, Vanengelsdorp D, Evans JD (2012) Pathogen webs in collapsing Honey bee colonies. *PLoS One* 7:e43562
- [5] Liao Z, Jia Q, Li F, Han Z (2010) Identification of two piwi genes and their expression profile in Honey bee, *Apis mellifera*. *Arch Insect Biochem Physiol* 74:91–102.
- [6] Evans J (2006) Beepath: an ordered quantitative-PCR array for exploring Honey bee immunity and disease. *Journal of Invertebrate Pathology* 93:135-139.
- [7] Lourenço AP, Mackert A, Cristino AS, Simões ZLP (2008) Validation of reference genes for gene expression studies in the Honey bee, *Apis mellifera*, by quantitative real-time RT-PCR. *Apidologie* 39:372–385.