

Supplementary material

Virulence of *Melissococcus plutonius* and secondary invaders associated with European foulbrood disease of the honey bee

Oleg Lewkowski, Silvio Erler

Table S1: Sample overview (number of individuals) per colony for each assay of experiment 1 – ‘Impact of host background’: survival, larval weight on day 7, relative *M. plutonius* load per larvae; and CFUs used for infection for each treatment group (mean ± SE).

Colony	Treatment	Survival	Larval weight	<i>M. plutonius</i> load	CFUs per ml	CFU per larvae
A	control	70	65	8	0	0
	49.3	70	59	12	$4.8 \times 10^6 \pm 1.6 \times 10^6$	1206 ± 398
	119	66	57	12	$3.2 \times 10^6 \pm 1.5 \times 10^6$	797 ± 380
	4-127	71	65	12	$7.0 \times 10^7 \pm 3.4 \times 10^7$	17512 ± 8424
	total	277	246	44		
B	control	47	44	8	0	0
	49.3	84	76	12	$4.8 \times 10^6 \pm 1.6 \times 10^6$	1206 ± 398
	119	42	38	8	$3.2 \times 10^6 \pm 1.5 \times 10^6$	797 ± 380
	4-127	42	39	8	$7.0 \times 10^7 \pm 3.4 \times 10^7$	17512 ± 8424
	total	215	197	36		

Table S2: Sample overview (number of individuals) for each assay of experiment 2 – ‘Impact of co-infection’: survival, larval weight on day 8; and CFUs used for infection for each treatment group (mean ± SE).

Treatment	Survival	Larval weight	CFUs per ml	CFU per larvae
control	73	83	0	0
<i>M. p.</i>	59	62	$1.4 \times 10^8 \pm 6.3 \times 10^7$	34605 ± 15654
<i>E. f.</i> *	65	40	$1.1 \times 10^9 \pm 5.2 \times 10^8$	$2.7 \times 10^5 \pm 1.3 \times 10^5$
<i>P. a.</i> *	69	42	$3.4 \times 10^7 \pm 1.4 \times 10^7$	8387 ± 3474
total	266	227		

* *M. p.* treatment on day 1

Table S3: Larval weight (mean \pm SD) on day 8 (day 7 post-infection) from the preliminary infection experiment. Controls and *M. plutonius* (strain LMG 20360, ST1, CC13) inoculated larvae.

Colony	Treatment	Weight (mg)	N
A	control	139.6 \pm 23.2	23
	LMG 20360	148.1 \pm 20.3	66
	total		89
B	control	124.4 \pm 33.3	23
	LMG 20360	134.7 \pm 21.0	64
	total		87

Fig. S1. Larval survival in preliminary infection experiment (*M. plutonius* strain LMG 20360, ST1, CC13) with colony A and colony B (cox regression model: $P > 0.05$).

