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Trade-offs between immunity and competitive ability in fighting ant males

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Fig1ab: fight_outcome: number of fights ending decided (one winner and one loser) vs undecided (both males dead or alive) for fights including one infected male or only two healthy males, and proportion of fights lost by infected males depending on infection stage

Columns:

"infection_stage": early-infected or late-infected; males fighting immediately after fungal spore exposure (early) or after establishment of the infection after an isolation period of 48h (late)

"dose": exposure dose of males; high, low

"colony": colony of origin

"fight_combination": combination of males per fight; healthy-healthy: two healthy males, healthy-infected: one healthy and one infected male

"fight_outcome": outcome of the fight; one dead (i.e. one winner and one loser), both alive, both dead

"male_ID_decided_fights": method used to differentiate between the two males in fights between a healthy and an infected male that had a winner and a loser; ddPCR: spore load determination with digital droplet PCR, colour-code: males were differentiated on the basis of the colour mark applied onto their abdomen

"infected_male_loser": fights in which the infected male died; yes: infected male loser; no: infected male winner, not possible: no differentiation between the males possible, both males: both males died

"male_pair_identification": numbers 1-208 refer to male pairs, for which scan sampling was performed (see Fig 1cdef); letters (E –early; L –late) with numbers (1-20 each) indicate male pairs, for which spore load was determined by ddPCR for both males after fight (see tab "cross-contamination")

Some of these data were also used to create Fig1 g,h.

Fig1cdef: behaviour: aggressive behaviour performed by the males towards their rivals (Fig 1c,d) and by the workers towards the males (Fig1 e,f)

Columns:

"infection_stage": early-infected or late-infected; males fighting immediately after fungal spore exposure (early) or after establishment of the infection after an isolation period of 48h (late)

"dose": exposure dose of males; high, low

"colony": colony of origin

"replicate": running number of male fighting pairs that were observed for their behavior (see also 1ab)

- "male_treatment": treatment of the male; healthy (sham-treated), infected (pathogenexposed)
- "fight_combination": combination of males per fight; healthy-healthy: two healthy males, healthy-infected: one healthy and one infected male
- "survival": mortality of the male; alive, dead
- "fight_outcome": outcome of the fight; one dead (i.e. one winner and one loser), both alive, both dead
- "male_biting": number of bites performed by the male towards the other male
- "male_holding": number of holdings (grabbing the rival with the mandibles) performed by the male towards the other male
- "male_besmearing": number of besmearings (male bending its abdominal tip to apply hindgut secretion onto the rival) performed by the male towards the other male
- "worker_biting": number of worker bites received by the male
- "worker carrying: number of worker carrying events received by the male
- "worker_dismembering": number of intensive worker attacks received by the male (sometimes body parts bitten-off)
- "worker_dragging": number of dragging events (male is held and pulled over the ground) received by the male by the workers

For the analysis, all male behaviours performed resp. worker behaviours received were summed and analysed as male-derived resp. worker-derived aggression.

Fig1gh: fight_costs: infected male mortality in the presence or absence of fighting: cost of fighting in dependence of infection stage

Columns

- "infection_stage": early-infected or late-infected; males fighting immediately after fungal spore exposure (early) or after establishment of the infection after an isolation period of 48h (late)
- "dose": exposure dose of males; high, low
- "colony": colony of origin
- "survival": mortality of male; alive, dead
- "male fighting": no: males without a rival (baseline mortality), yes: males fighting a rival

Some of these data (infected male mortality in the absence of fight) are repeated in tab S1_baseline_mortality, which also contained the mortality of healthy males in the absence of fight. See also tab 1a,b as we here show the mortality of all infected males either from decided fights (one male dead) or fights with both males dead.

FigS1: baseline_mortality: healthy and infected male mortality in the absence of fight *Columns:*

"infection_stage": early-infected or late-infected; males fighting immediately after fungal spore exposure (early) or after establishment of the infection after an isolation period of 48h (late)

"colony": colony of origin

Fig2 & S2: Immune_genes: expression of 3 immune genes and the housekeeping gene at 12h (Fig S2) and 24h (Fig 2) after exposure

Columns:

- "male_treatment": treatment of the male; sham: sham control treatment, low: low pathogen exposure, high: high pathogen exposure
- "timepoint": timepoint after treatment with the sham, low or high pathogen load, at which immune gene expression was measured; 12h, 24h
- "colony": colony of origin
- "male_ID": running number of the males used for immune gene expression (number not related to the main experiment)
- "immune_gene": immune gene analysed by ddPCR; Relish, PPOAF (Prophenoloxidase activating-factor), Defensin
- "immune_gene_expression_value": immune gene expression value, as quantified by ddPCR for the three genes (Relish, PPOAF, Defensin)
- "EF1_gene_expression_value": housekeeping gene (EF1) expression, as quantified by ddPCR "normalised_gene_expression": normalized gene expression; expression of immune gene / expression of EF1 (housekeeping gene)

cross_contamination: fungal load determination with ddPCR, inferred infection status and cross-contamination of the males in fights of healthy and infected males that ended with a clear winner and loser

Columns

- "infection_stage": early-infected or late-infected; males fighting immediately after fungal spore exposure (early) or after establishment of the infection after an isolation period of 48h (late)
- "replicate": running number of male fighting pairs that were analysed for their fungal load (see also 1ab)
- "survival": mortality of males; alive, dead
- "fungal_load": fungal load per male; fungal load/male (calculated by the quantified value of copies of the mRFP1 gene in the 20μl well of the ddPCR reaction, divided by the 2μl of extracted DNA used in the PCR reaction, times 50μl, which corresponds to the (final elution volume of DNA extraction), "below detection" threshold when less than 2 positive droplets in PCR.
- "inferred_male_type": infected, healthy; the male with the higher fungal load per pair determined to be the infected male, the male with either no spores or less spores than the other male in the fight determined as the healthy male
- "cross-contamination": cases in which both males had a detectable fungal load; yes: detectable fungal load in both males; no: detectable fungal load only in one male. Note that replicates in which none of the males had detectable spore load could not be included (which was the case for some of the late-infected pairings).

[&]quot;male_treatment": treatment of the male; sham: sham control treatment, low: low pathogen exposure, high: high pathogen exposure

[&]quot;survival": mortality of males after 24h without another male present; alive, dead