

ELECTRONIC SUPPLEMENTARY MATERIAL

To be submitted to: Oecologia

Hyperparasitoids exploit herbivore-induced plant volatiles during host location to assess host quality and non-host identity

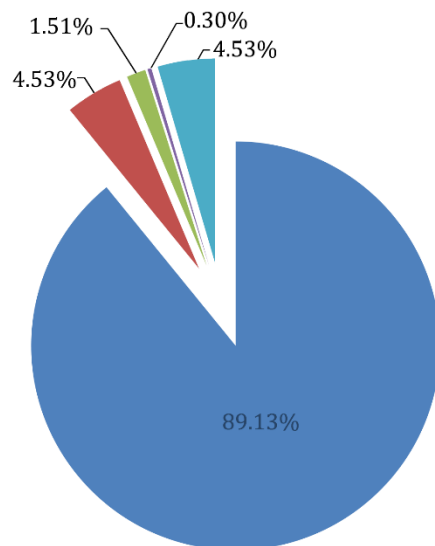
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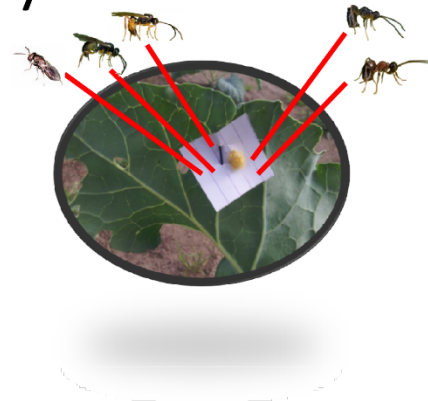
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Online Resource 1. Proportion of each species of hyperparasitoid collected in the common garden experiments from *Cotesia glomerata* cocoon clutches attached to *Brassica oleracea* “Kimmeridge” plants subjected to different induction treatments. Colors indicate different hyperparasitoid species: *Lysibia nana* (blue); *Acrolyta nens* (red); *Gelis agilis* (green); *Gelis* sp. (purple); *Pteromalus* sp. (light blue).

Hyperparasitoid community



total number of hyperparasitoids=331



- *Lysibia nana*
- *Acrolyta nens*
- *Gelis agilis*
- *Gelis* sp.
- *Pteromalus* sp.

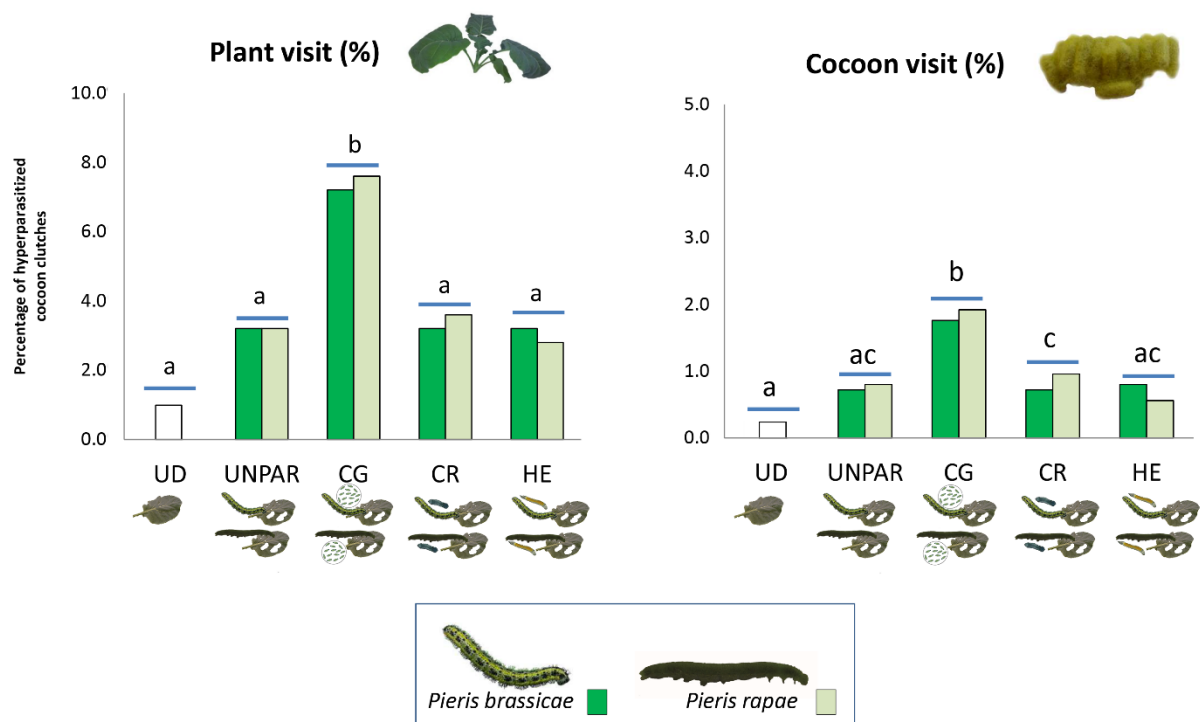
Online Resource 2. The effect of plant induction treatment on the overall hyperparasitism rates achieved on *Cotesia glomerata* cocoons at the clutch level. Hyperparasitism was modeled as the number of clutches yielding hyperparasitoids out of the fixed totals of five cocoon clutches attached to the plant (see main text for information about statistical analyses).

Model factor	Deviance	Degrees of freedom	P value
Overall (cocoon level)	427.74	499	
Factor			
Induction treatment (1)	43.604	8	<0.001
Replicate (2)	55.847	4	<0.001
(1) × (2)	14.251	32	0.9971

Online Resource 3. The effect of the herbivore species (*Pieris brassicae*, *P. rapae*) and parasitism (*Cotesia glomerata*, *C. rubecula*, *Hyposoter ebeninus*, none) on the overall hyperparasitism rates achieved on *C. glomerata* cocoons at the clutch level. Hyperparasitism was modeled as the number of clutches yielding hyperparasitoids out of the fixed totals of five cocoon clutches attached to the plant.

Model factor	Deviance	Degrees of freedom	P value
Overall (cocoon level)	366.17	399	
Factor			
Herbivore species (1)	0.086	1	0.7695
Parasitism (2)	21.875	3	<0.001
Replicate (3)	50.353	4	<0.001
(1) × (2)	2.012	3	0.5700
(1) × (3)	0.512	4	0.9724
(2) × (3)	4.376	12	0.9757
(1) × (2) × (3)	7.326	12	0.8

Online Resource 4. Percentage of *Cotesia glomerata* cocoon clutches that yielded *Lysibia nana* hyperparasitoids in the field trials either at the plant level (i.e. at least one cocoon clutch out of the five clutches attached to the plant contained *L. nana* hyperparasitoids) (left) or at the individual clutch level (right). The cocoons were collected from plants that were either left untreated (UD), infested with unparasitized *Pieris* caterpillars (UNPAR) or parasitized by *C. glomerata* (CG), *C. rubecula* (CR) or *Hyposoter ebeninus* (HE). Dark green bars indicate plant treatments with *P. brassicae*; light green bars indicate treatments with *P. rapae*, white bars indicate undamaged plants). Letters indicate significant differences between treatment groups (GLM, $P < 0.05$).



Online Resource 5. The effect of plant induction treatment on the hyperparasitism rates achieved by *Lysibia nana* on *Cotesia glomerata* cocoons at the plant level. Hyperparasitism was modeled as a binomial occurrence of *L. nana* hyperparasitoids (presence=1, absence=0) emerging from the five cocoon clutches attached per plant

Model factor	Deviance	Degrees of freedom	P value
Overall (plant level)	474.41	499	
Factor			
Induction treatment	36.577	8	<0.001
Replicate	46.770	4	<0.001
(1) × (2)	16.995	32	0.9865

Online Resource 6 The effect of plant induction treatment on the hyperparasitism rates achieved by *Lysibia nana* on *Cotesia glomerata* cocoons at the clutch level. Hyperparasitism was modeled as the number of clutches yielding *L. nana* hyperparasitoids out of the fixed totals of five cocoon clutches attached to the plant (see main text for information about statistical analyses).

Model factor	Deviance	Degrees of freedom	P value
Overall (cocoon level)	405.21	499	
Factor			
Induction treatment	40.061	8	<0.001
Replicate	52.702	4	<0.001
(1) × (2)	14.670	32	0.9963

Online Resource 7. The effect of herbivore (*Pieris brassicae*, *P. rapae*) and parasitism (*Cotesia glomerata*, *C. rubecula*, *Hyposoter ebeninus*, none) on the hyperparasitism rates achieved by *Lysibia nana* on *C. glomerata* cocoons at the plant level. Hyperparasitism was modeled as a binomial occurrence of *L. nana* hyperparasitoids (presence=1, absence=0) emerging from the five cocoon clutches attached per plant

Model factor	Deviance	Degrees of freedom	P value
Overall (plant level)	416.41	399	
Factor			
Herbivore species (1)	0.059	1	0.8077
Parasitism (2)	17.862	3	<0.001
Replicate (3)	42.737	4	<0.001
(1) × (2)	0.359	3	0.9486
(1) × (3)	3.112	4	0.5393
(2) × (3)	6.005	12	0.9158
(1) × (2) × (3)	6.088	12	0.9116

Online Resource 8. The effect of the herbivore species (*Pieris brassicae*, *P. rapae*) and parasitism (*Cotesia glomerata*, *C. rubecula*, *Hyposoter ebeninus*, none) on the hyperparasitism rates achieved by *Lysibia nana* on *C. glomerata* cocoons at the clutch level. Hyperparasitism was modeled as the number of clutches yielding *L. nana* hyperparasitoids out of the fixed totals of five cocoon clutches attached to the plant.

Model factor	Deviance	Degrees of freedom	P value
Overall (cocoon level)	347.83	399	
Factor			
Herbivore species (1)	0.165	1	0.6842
Parasitism (2)	19.407	3	<0.001
Replicate (3)	48.158	4	<0.001
(1) × (2)	1.392	3	0.7073
(1) × (3)	3.208	4	0.5236
(2) × (3)	5.383	12	0.9439
(1) × (2) × (3)	4.420	12	0.97