

Supplementary figures for: “Candidate genes involved in cuticular hydrocarbon differentiation between cryptic, parabiocotic ant species”

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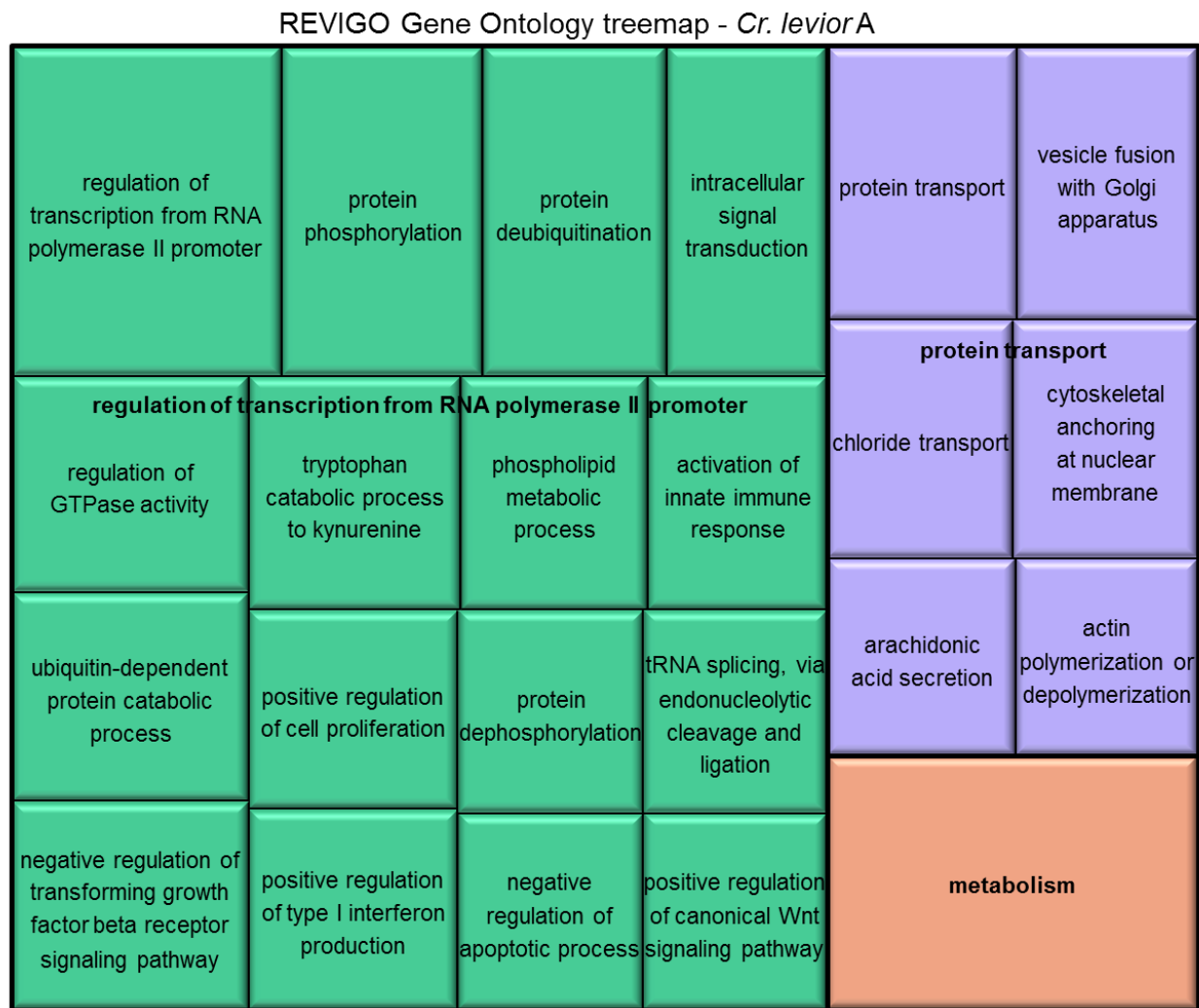


Figure S1: REVIGO treemap illustrating significantly enriched GO terms in *Crematogaster levior* A. Significantly enriched GO terms (Fisher-test: $p < 0.05$) of genes upregulated in *Cr. levior* A compared to B. Relative square sizes are scaled to match the degree of overexpression (Supek *et al.* 2011).

REVIGO Gene Ontology treemap - *Cr. levior* B

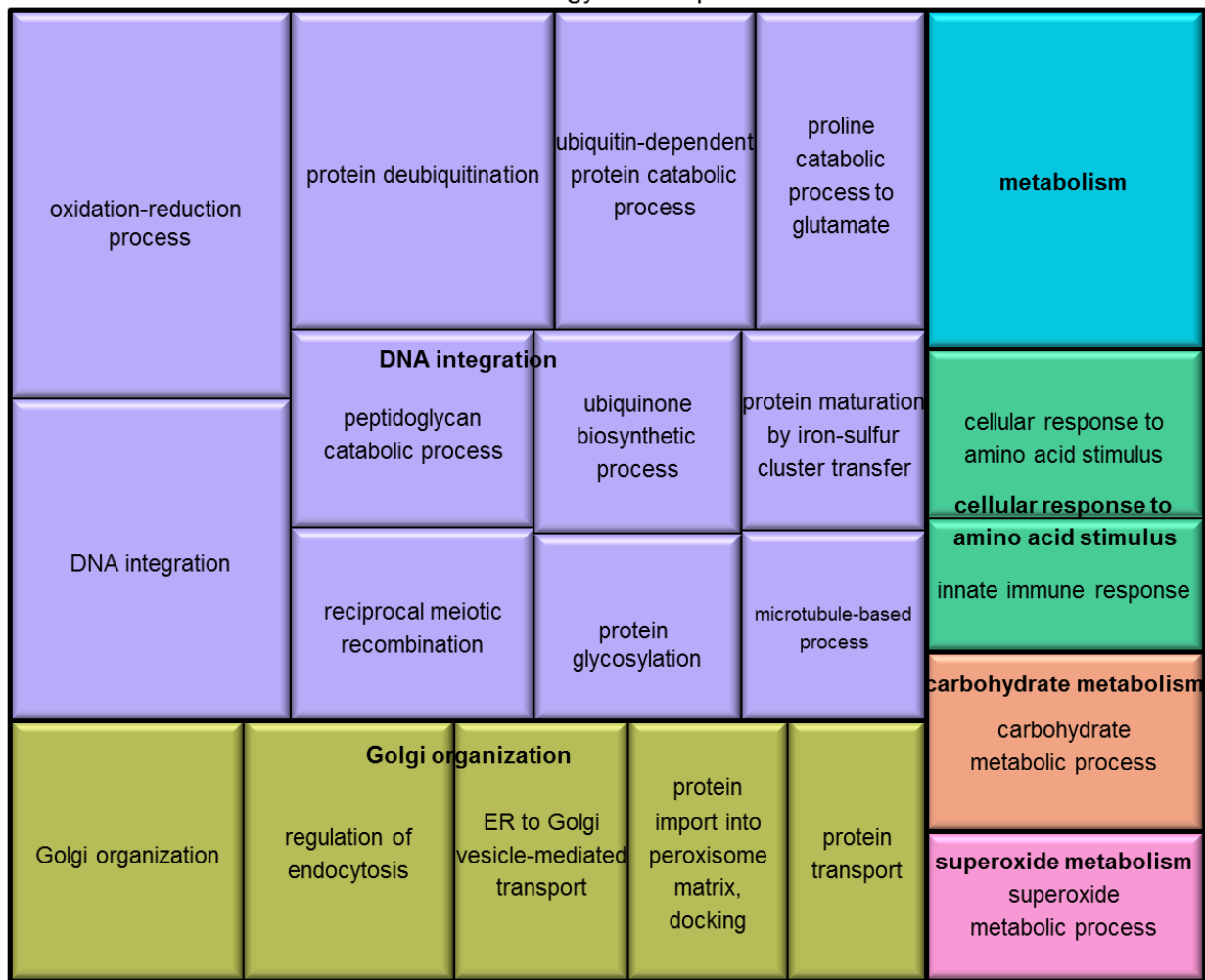


Figure S2: REVIGO treemap illustrating significantly enriched GO terms in *Crematogaster levior* B. Significantly enriched GO terms (Fisher-test: $p < 0.05$) of genes upregulated in *Cr. levior* B compared to A. Relative square sizes are scaled to match the degree of overexpression (Supek *et al.* 2011).

REVIGO Gene Ontology treemap - *Ca. femoratus* PAT

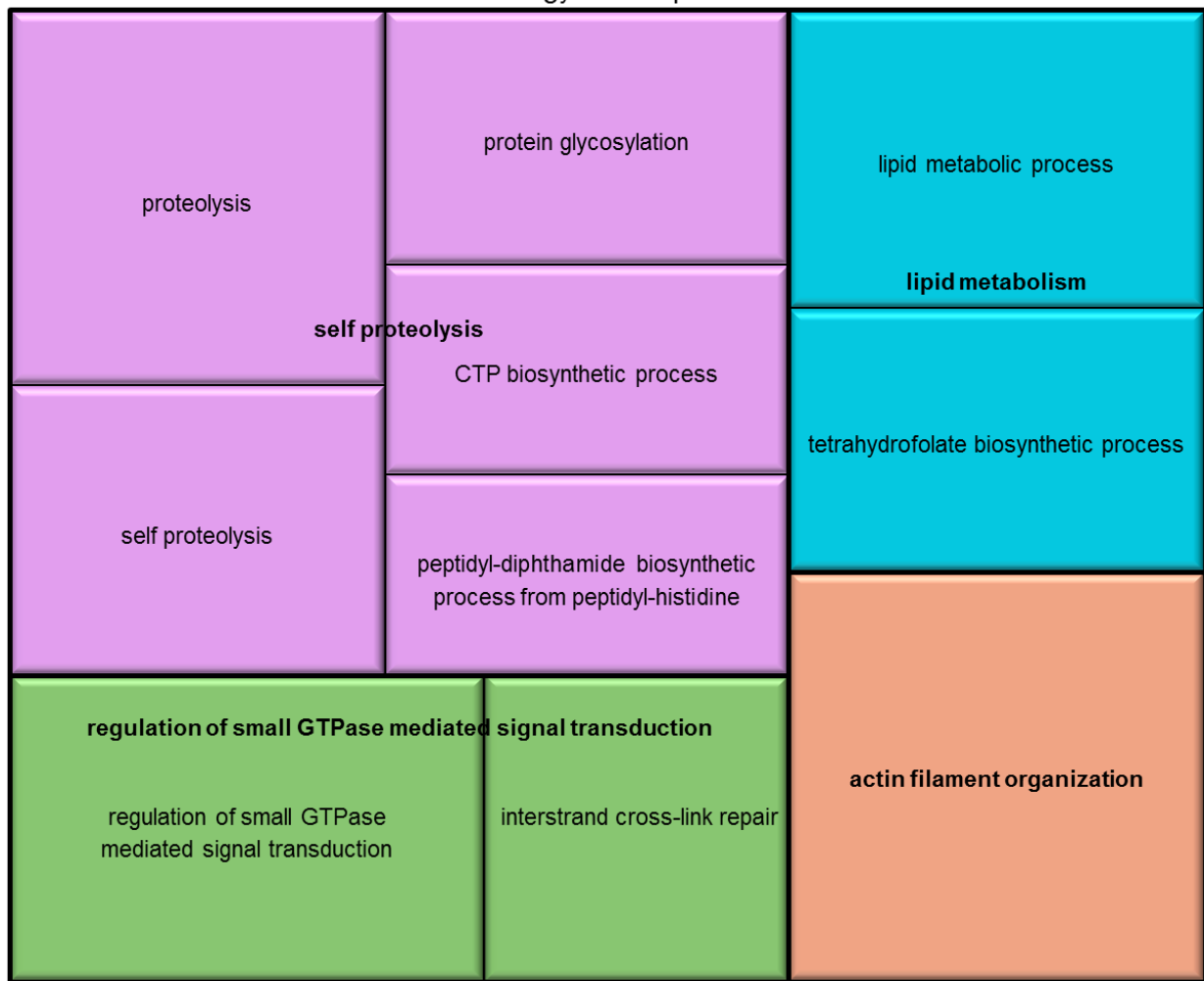


Figure S3: REVIGO treemap illustrating significantly enriched GO terms in *Camponotus femoratus* PAT. Significantly enriched GO terms (Fisher-test: $p < 0.05$) of genes upregulated in *Ca. femoratus* PAT compared to PS. Relative square sizes are scaled to match the degree of overexpression (Supek *et al.* 2011).

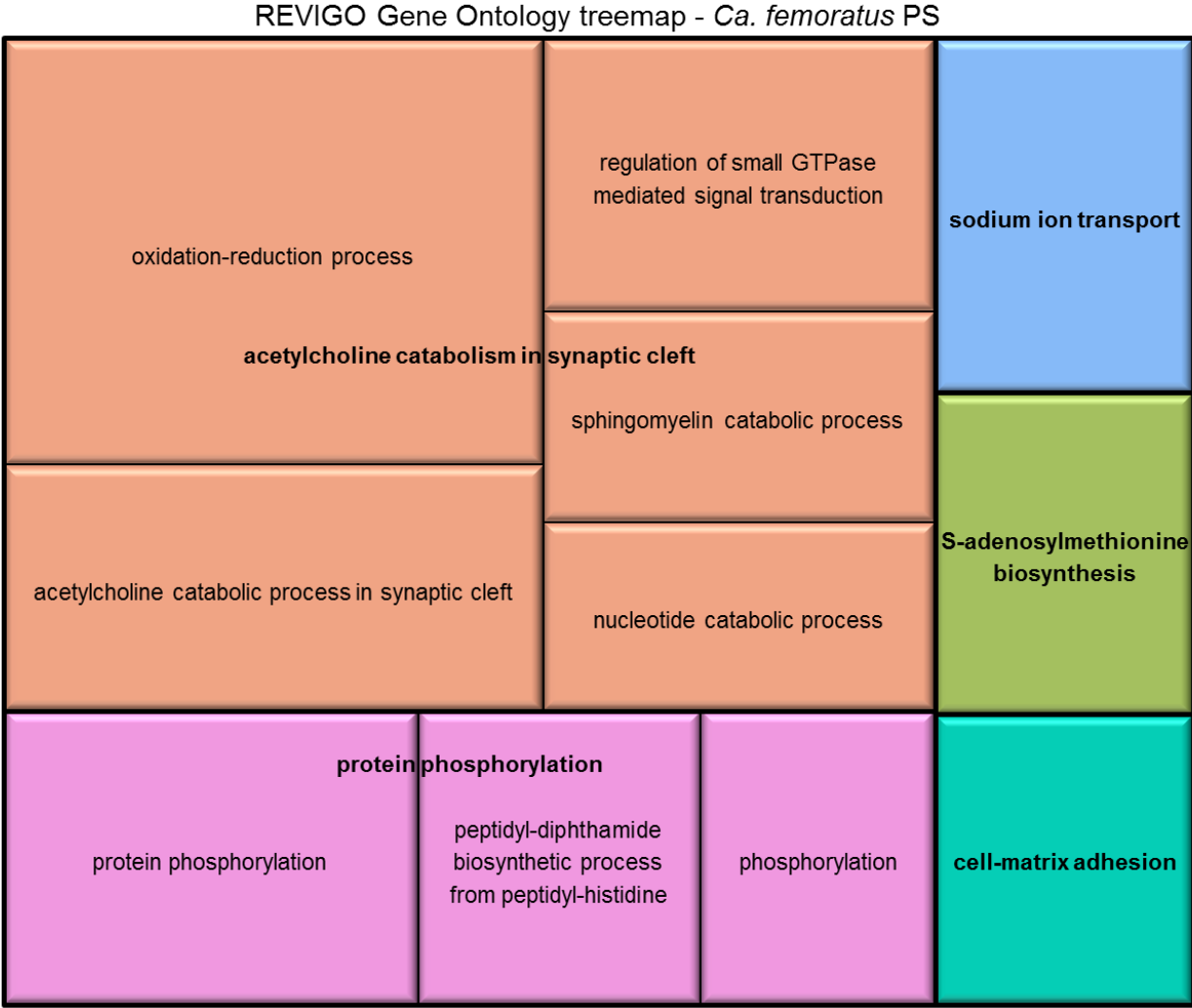


Figure S4: REVIGO treemap illustrating significantly enriched GO terms in *Camponotus femoratus* PS. Significantly enriched GO terms (Fisher-test: $p < 0.05$) of genes upregulated in *Ca. femoratus* PS compared to PAT. Relative square sizes are scaled to match the degree of overexpression (Supek *et al.* 2011).