



Identification of a candidate queen cuticular hydrocarbon affecting fire ant social form



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Fire ant *Solenopsis invicta*



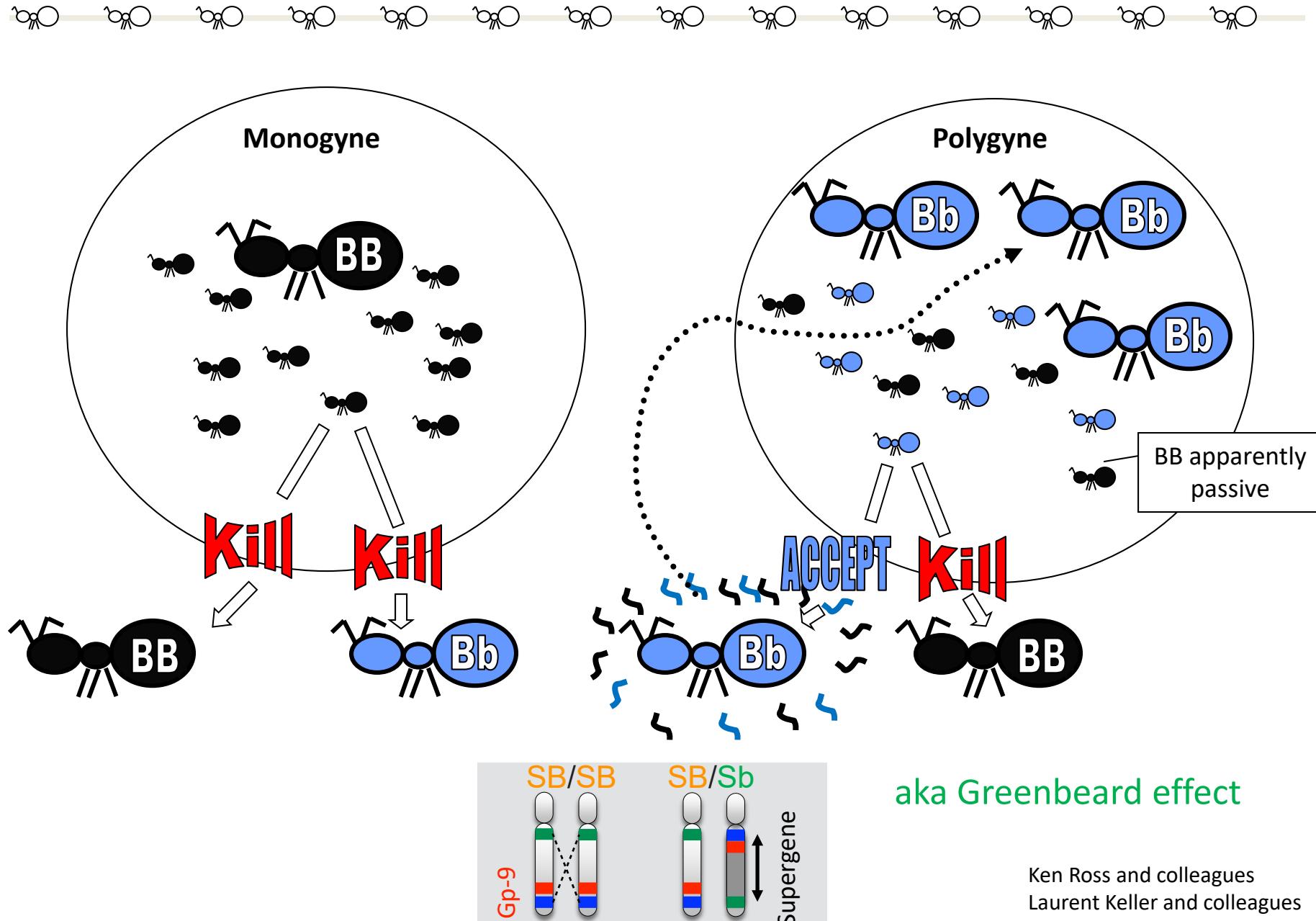
Monogyne

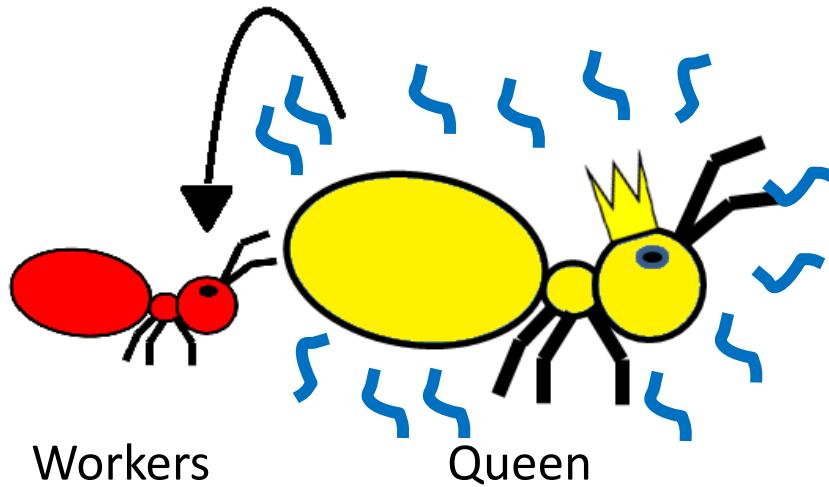


Polygyne

A selfish element drives colony form
(simultaneously a social supergene)

Fire ant social organization is linked to selfish Gp-9 locus





Can we determine:

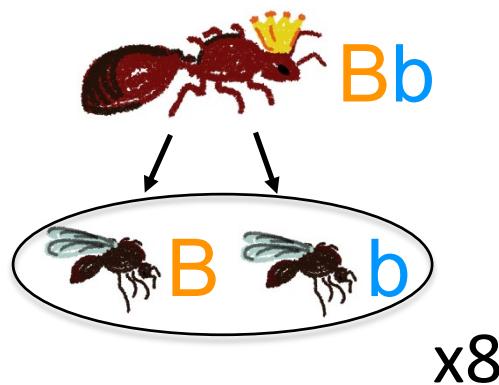
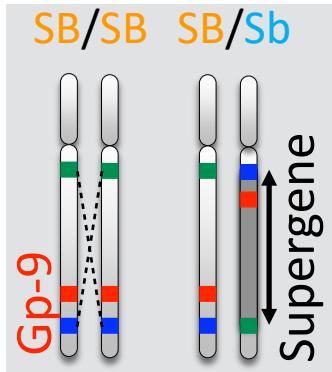
- Odor
- Receptor



Genomic
analysis

Chemical
ecology

Copy number variation analysis

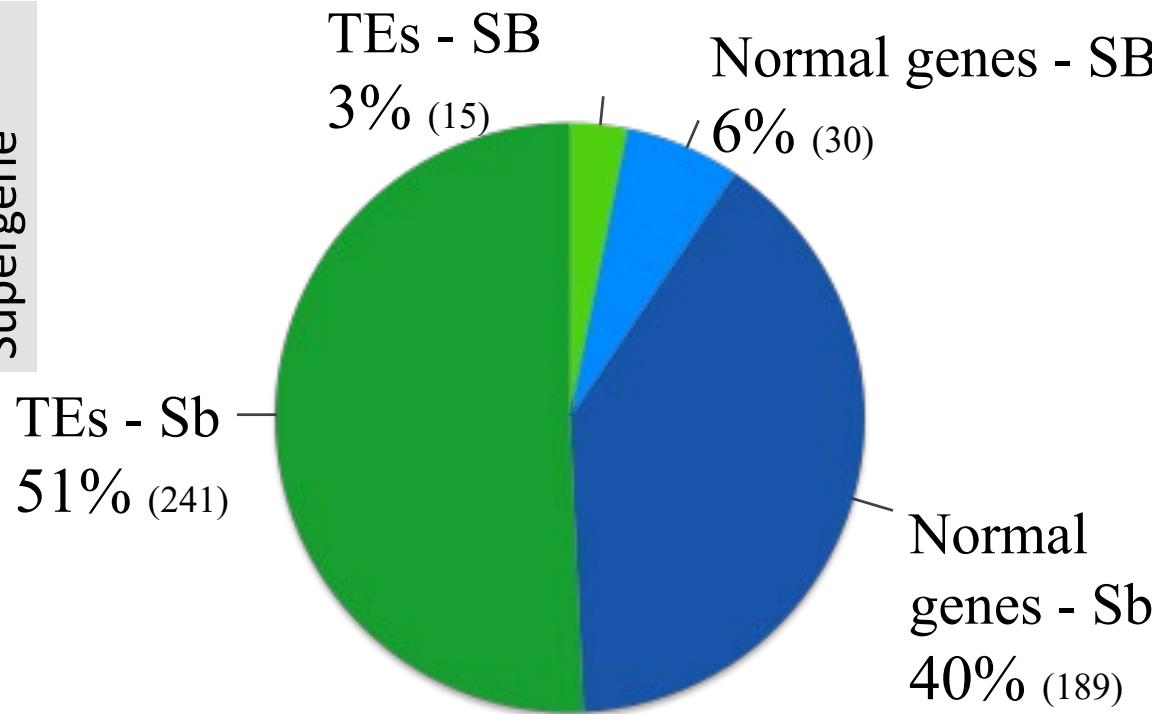
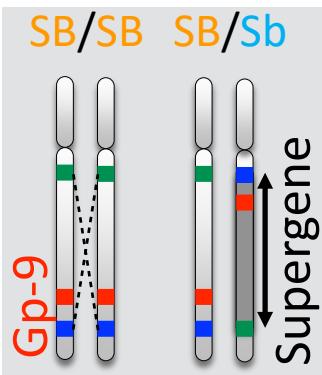


DNA
seq data →

map (BWA)
against
**reference
gene list**

→ mapped
reads
SB ≠ Sb?

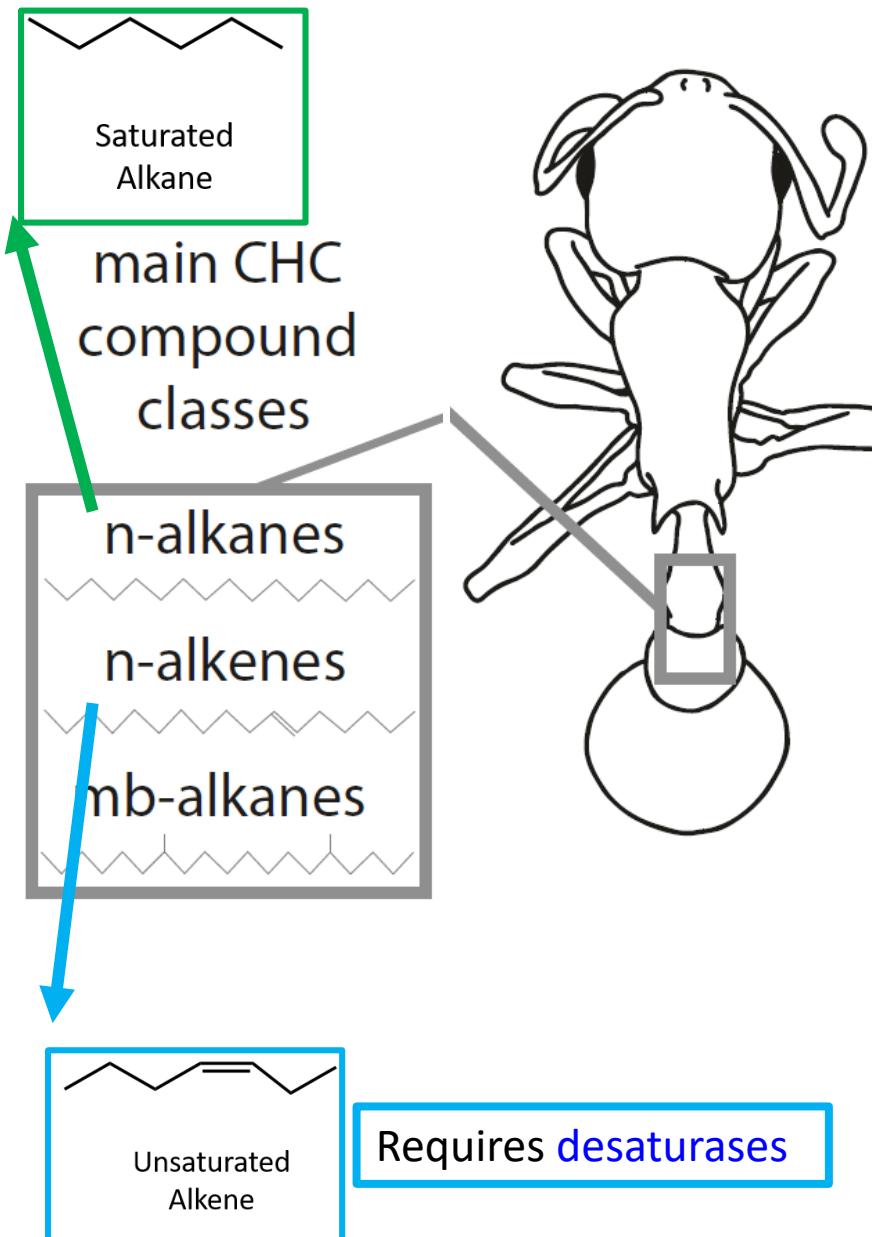
475 genes with CNV difference: Sb vs SB, [including a desaturase]



~50% are **transposable elements**

Surprisingly many “normal” genes

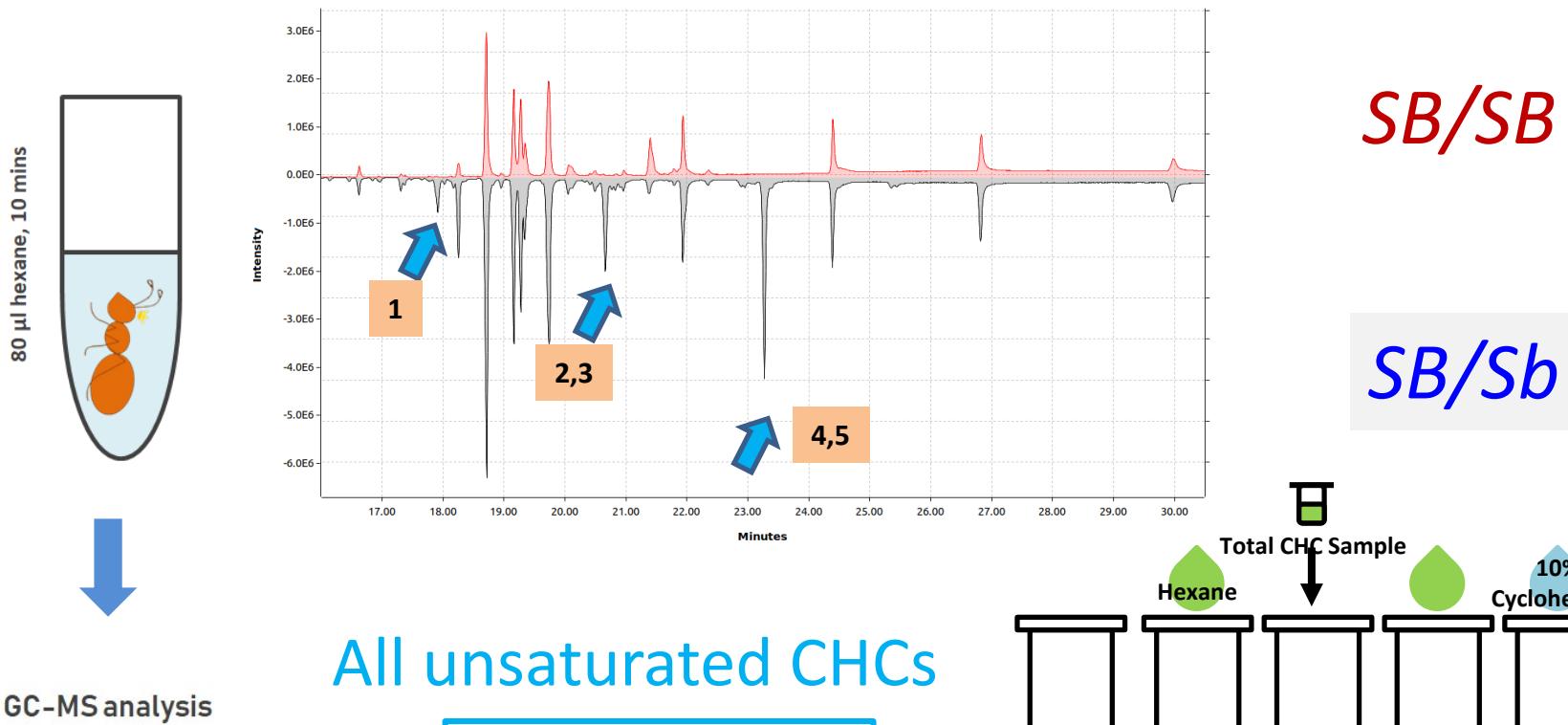
- Including 49 encoding putative **cuticular hydrocarbon synthesis enzymes**
- **Desaturase enzyme**



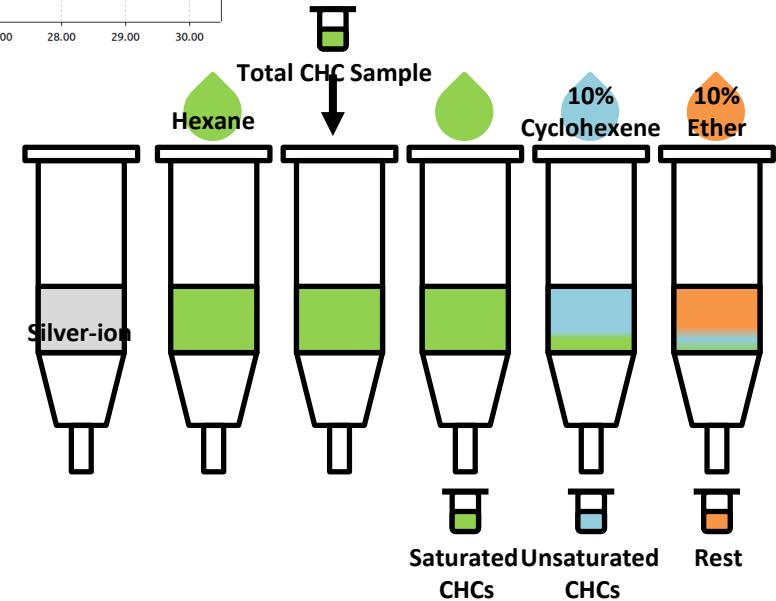
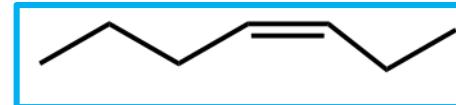
More unsaturated hydrocarbons in SB/Sb queens



Gas chromatography-mass spectrometer



GC-MS analysis

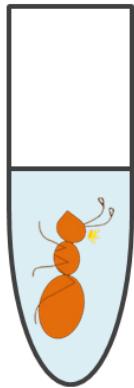


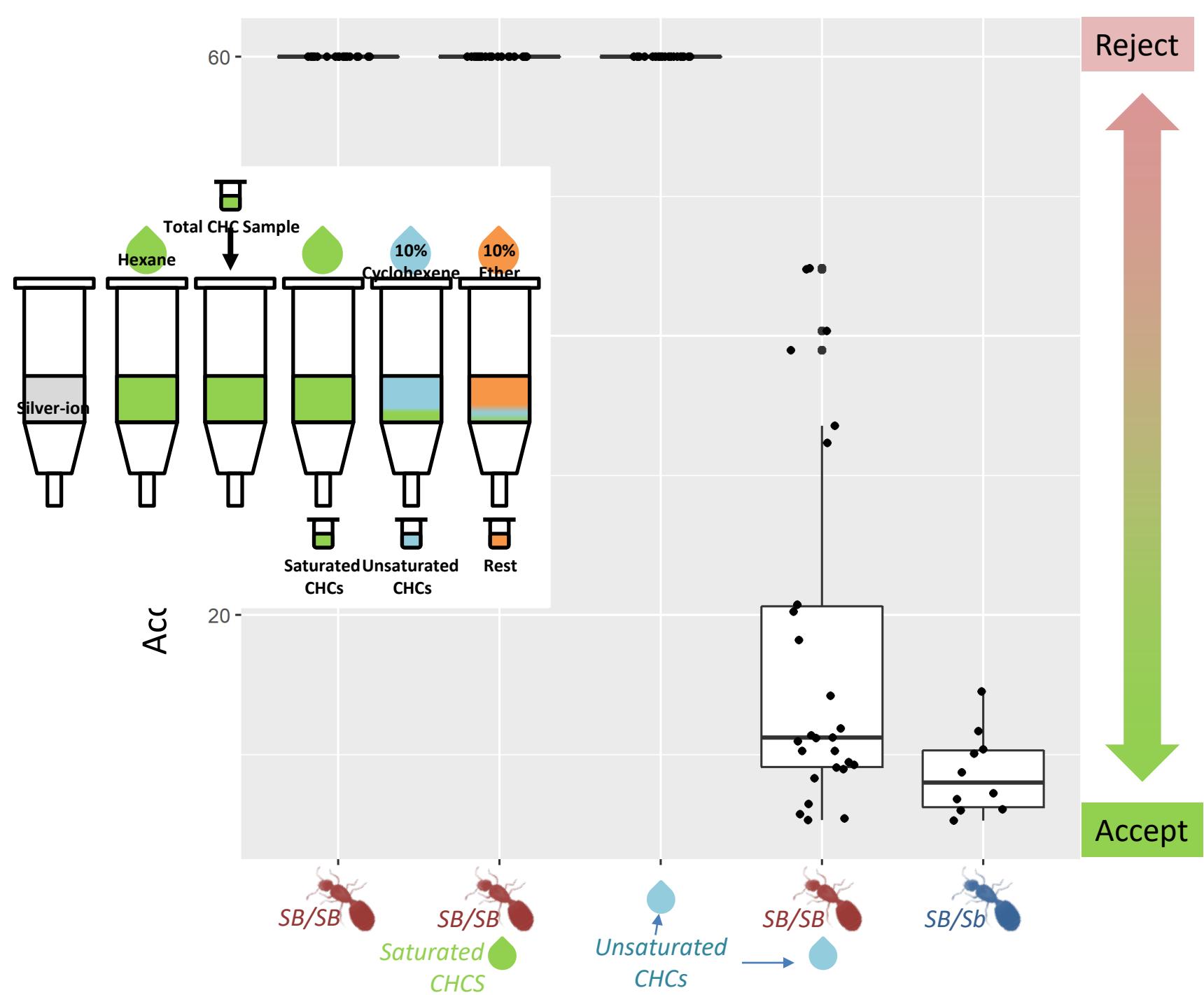
Hexane extract of queens → “paper surrogate”

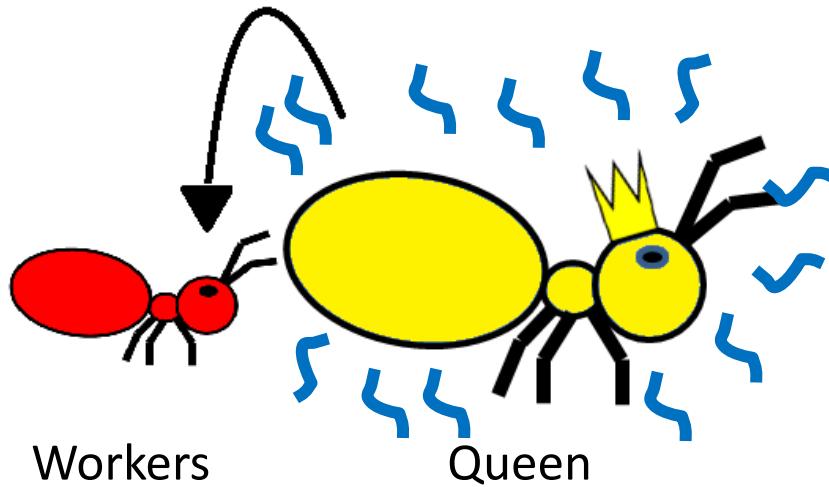


- *Bb* queen taken home
- *BB* queen tossed

80 µl hexane, 10 mins





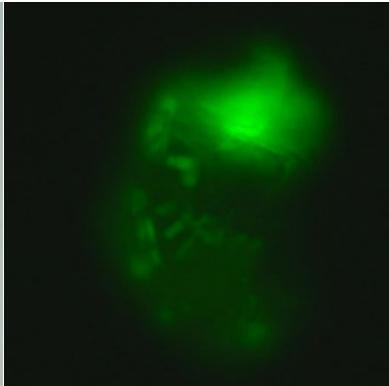


Narrowed it down to 1 unsaturated CHC
Trying to synthesize it

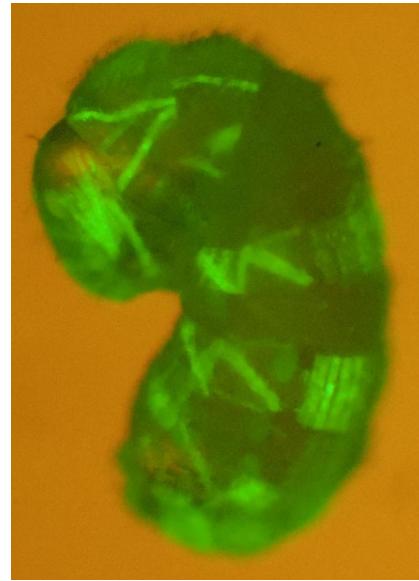
PiggyBac mediated transgenesis & CRISPR/Cas9 mutagenesis



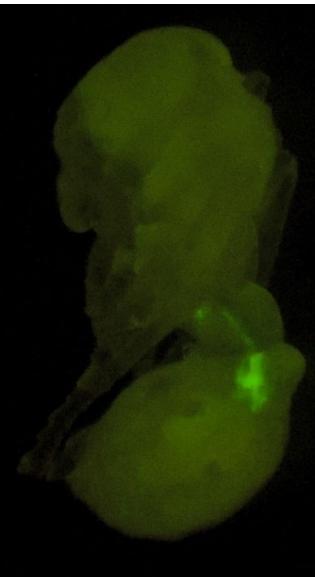
Embryo



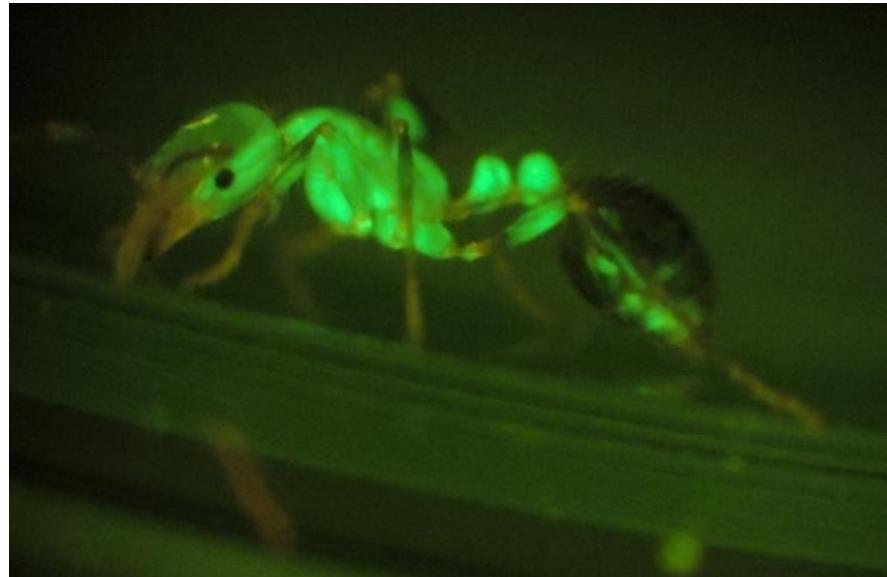
Larvae



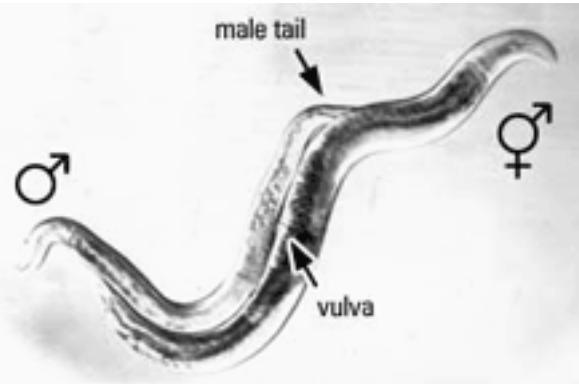
Pupae



Adult



Our interests



1. Social evolution
 - Social supergene

2. Sex determination
 - Novel non-coding gene?

1. Sex ratio bias in *Caenorhabditis*
 - Sperm competition?

2. Selfish gene in *C. briggsae*
 - Toxin-antidote

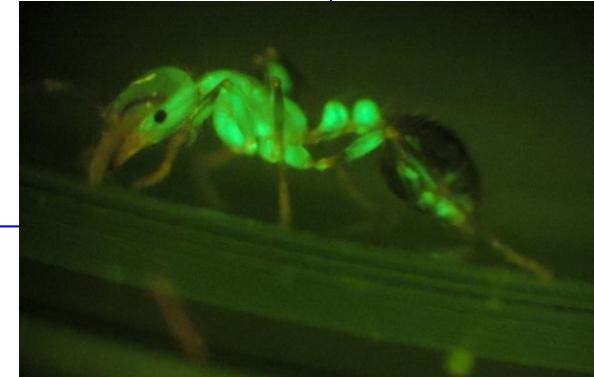
Collaborative possibilities



Can help:

in fire ants (ants/insects) and *Caenorhabditis* worms (nematodes):

- Genetic analysis
- Transgenesis
- CRISPR/Cas9 mutagenesis



Need help:

- Chemical ecology, chemistry
- Other monogyne/polygyne ants

