

Introduction:

Bee local: A comparison of productivity and pathogen load
in local vs. Claifornia re-queened colonies

Andre Burnham, Fiona McLaughlin, **P. Alexander Burnham** &
Herman Lehman



The
UNIVERSITY
of **VERMONT**

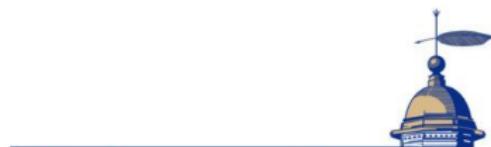
Acknowledgments

Co-Authors:

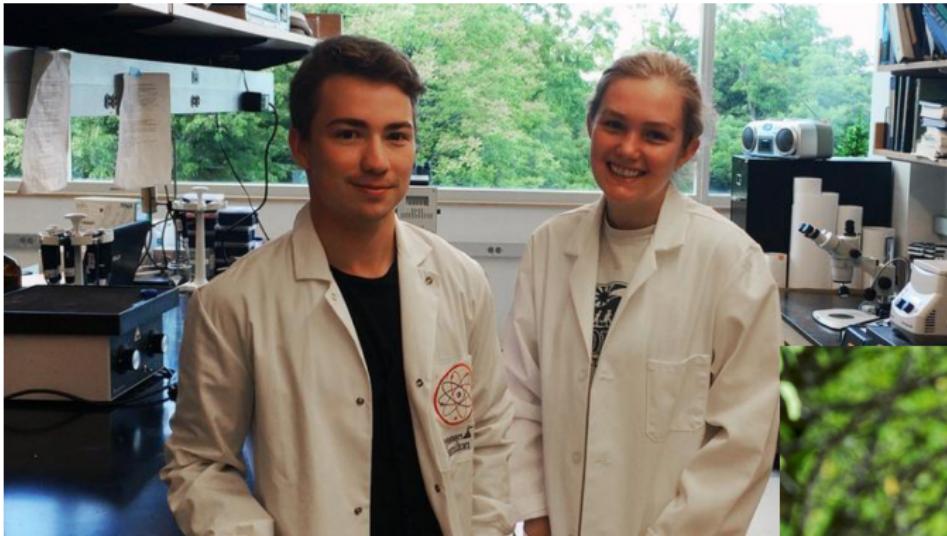
- ▶ Andre Burnham
- ▶ Fiona McLaughlin
- ▶ Dr. Herman Lehman

Thank you to:

- ▶ The Casstevens Family
- ▶ Nancy Thompson
- ▶ Samantha Alger



The Bee Team

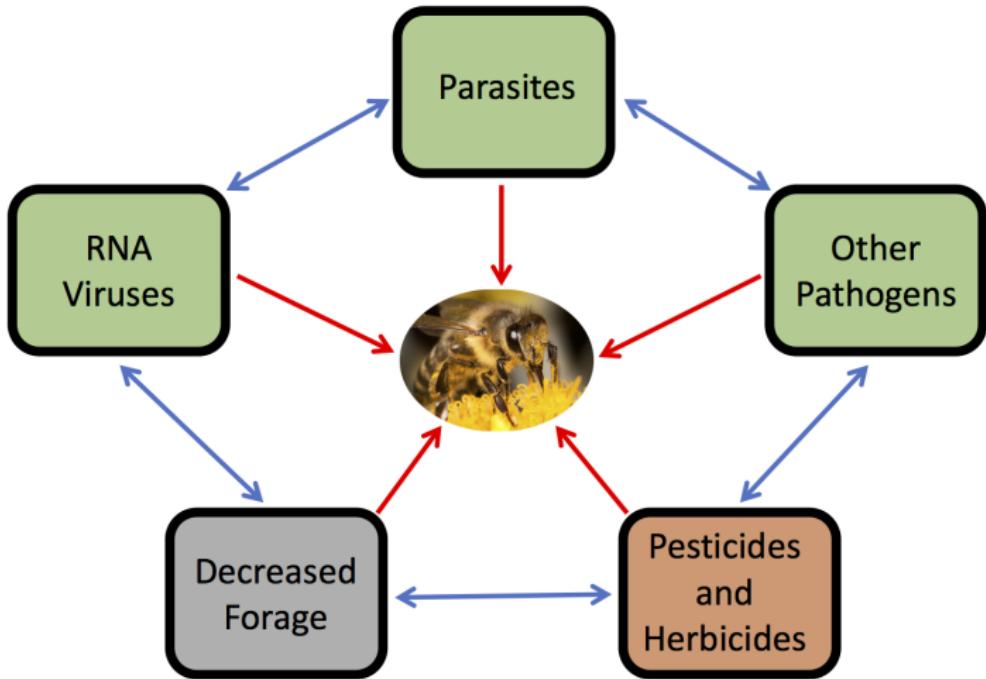


Honey Bees are Important

- ▶ 30% of the world's food is derived from pollination (Aizen et al., 2009)
- ▶ Pollinators are responsible for between \$235-577 billion (Gallai et al., 2009)
- ▶ Honeybees are responsible for \$14 Billion in the USA (Morse & Calderone, 2000)



Threats to Bees:



Honey Bee Pathogens

VIRUSES:

- ▶ Deformed Wing
- ▶ Black Queen Cell
- ▶ Israeli Acute Paralysis



Deformed wing Virus
University of Florida,
Entomology Dept.

PARASITES:

- ▶ Nosema (ceranae/apis)
- ▶ Varroa Mite



Varroa destructor
North Carolina State University,
Cooperative Extension



American Foulbrood
Bee Informed Partnership

Troubles for Beekeepers (re-queening)



The basic premises behind this study

- ▶ Imported VS Local
- ▶ Local Adapation



The basic premises behind this study

- ▶ Mass-Produced VS Handmade
- ▶ Selection by the Breeder



The question:

“Are locally-bred queens more successful than imported queens?”

Experimental Design

- ▶ 20 colonies re-queened with Californian-bred queens
- ▶ 20 colonies re-queened local-bred (Vermont) queens
- ▶ 2 sites, 10 Local and 10 California for each
- ▶ Sampled for pathogens and productivity measures
- ▶ Sampled at different time points for 3 months

Pictures of the Yards



What we sampled

- ▶ Growth:
 - ▶ Colony Mass
 - ▶ Brood Production
- ▶ Foraging:
 - ▶ Pollen Production
- ▶ Pathogens:
 - ▶ Varroa
 - ▶ Nosema spp.
 - ▶ RNA Viruses

Data Analysis

```
aov.out <- aov(Nosema ~ Origin * Time + Error(FieldID) ,  
                  data=QueenDF)  
  
summary(aov.out)
```

Repeated Measures ANOVA output

```
Error: FieldID
          Df     Sum Sq   Mean Sq F value Pr(>F)
Origin      1 3.156e+13 3.156e+13    7.972 0.00779 ***
Time        1 2.589e+12 2.589e+12    0.654 0.42413
Origin:Time 1 9.223e+11 9.223e+11    0.233 0.63234
Residuals   35 1.386e+14 3.959e+12
---
Error: Within
          Df     Sum Sq   Mean Sq F value Pr(>F)
Time        1 5.620e+10 5.620e+10    0.017 0.896098
Origin:Time 1 4.934e+13 4.934e+13  15.111 0.000275 ***
---
```

Graphics

```
NosemaSummary <- ddply(QueenDF, c("Origin", "NosemaDay"),  
                      summarise,  
                      n = length(Nosema),  
                      mean = mean(Nosema, na.rm = TRUE),  
                      sd = sd(Nosema, na.rm = TRUE),  
                      se = sd / sqrt(n))
```

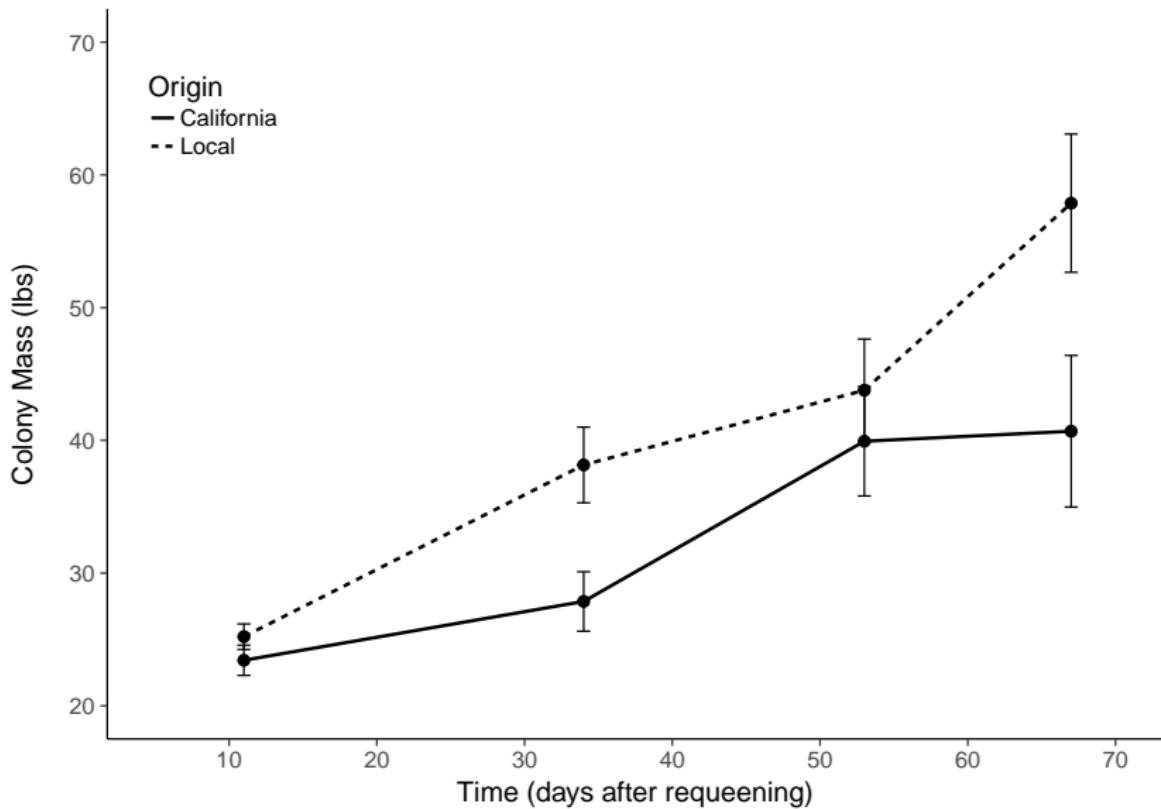
Graphics

```
nosema <- ggplot(data = NosemaSummary,
                    aes(x = NosemaDay,
                        y = mean,
                        group = Origin)
) + geom_point(size=3)
+ scale_colour_manual(values = c("black", "black"))
+ labs(x = "Time (days after requeening)",
      y = "Nosema Load (spores/bee)")
+ coord_cartesian(ylim = c(0, 4300000),
                  xlim = c(10,70))
+ geom_errorbar(aes(ymin = mean - se,
                     ymax = mean + se, width = 0.9))
+ geom_line(aes(linetype=Origin), size=1)
+ scale_fill_brewer(palette = "Paired")
+ theme_classic(base_size = 17)
+ theme(legend.position=c(.15, .85)
+ labs(linetype="Queen Origin")
```

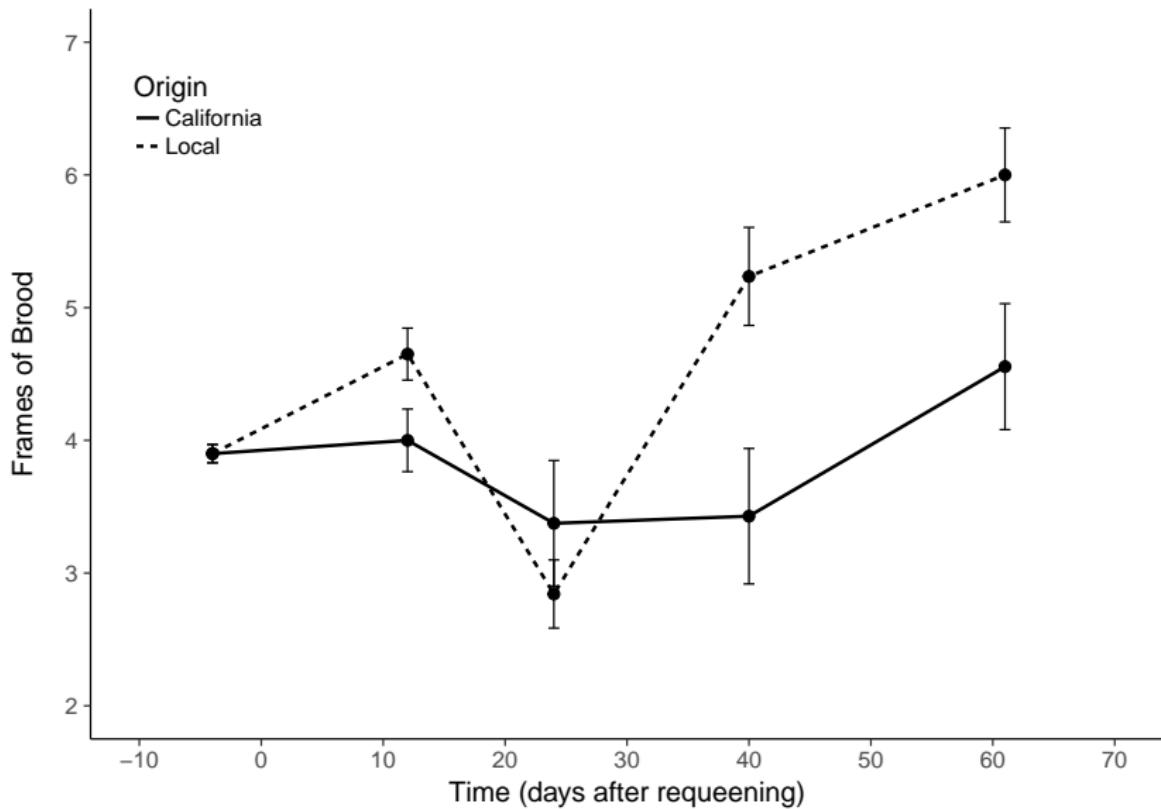
Our Predictions

- ▶ Local queens (colonies) will have higher growth through the season
- ▶ Local queens will be better foragers
- ▶ Local queens (colonies) will have lower pathogen loads

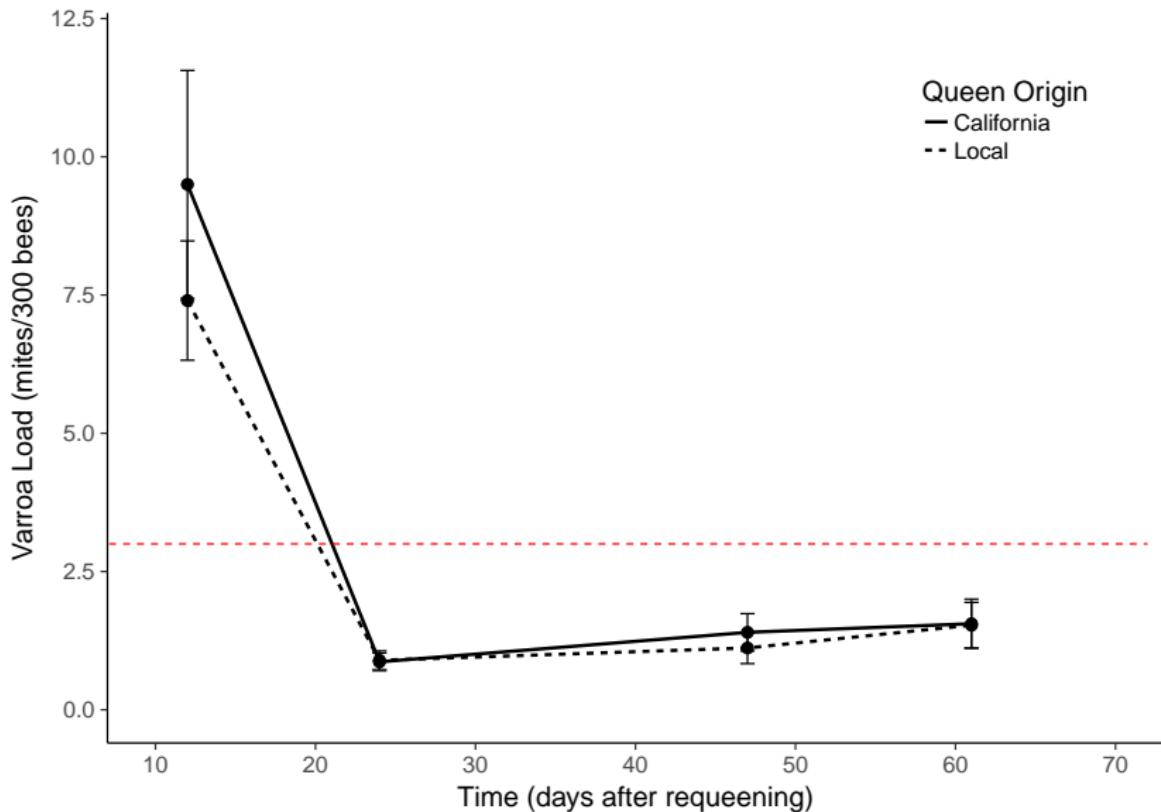
Colony Mass (growth)



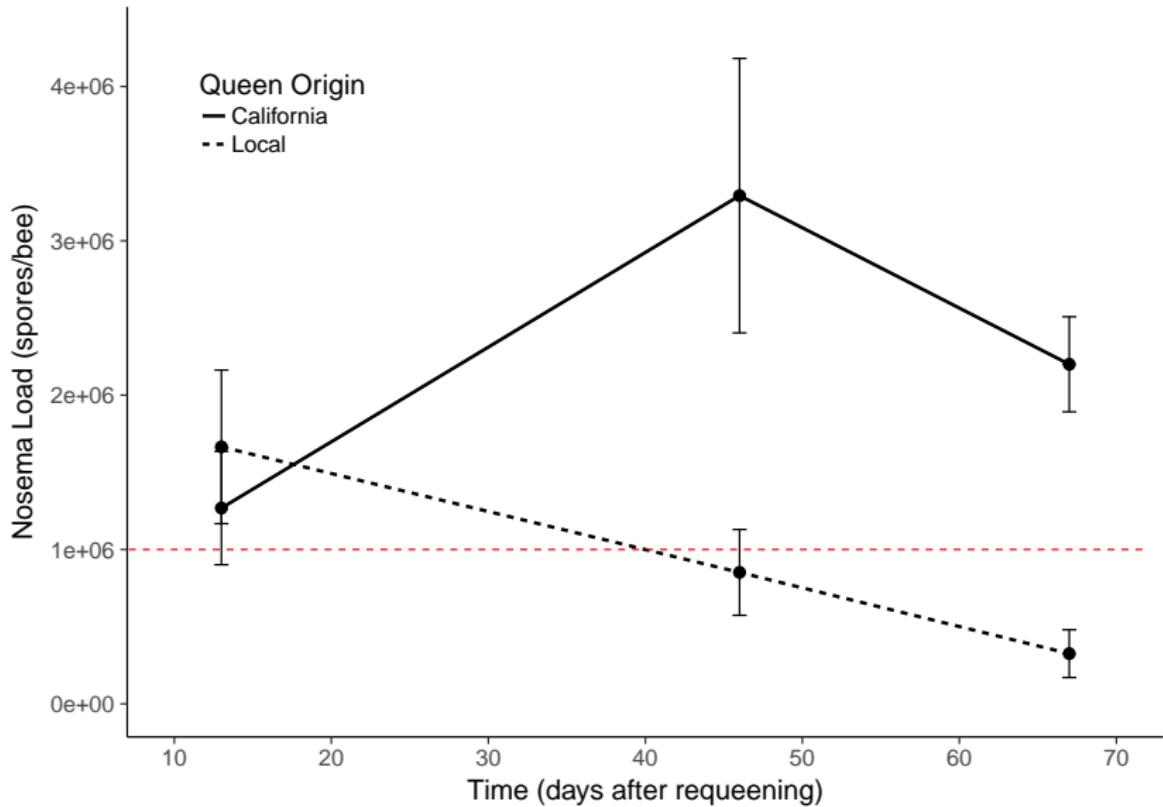
Frames of Brood (growth)



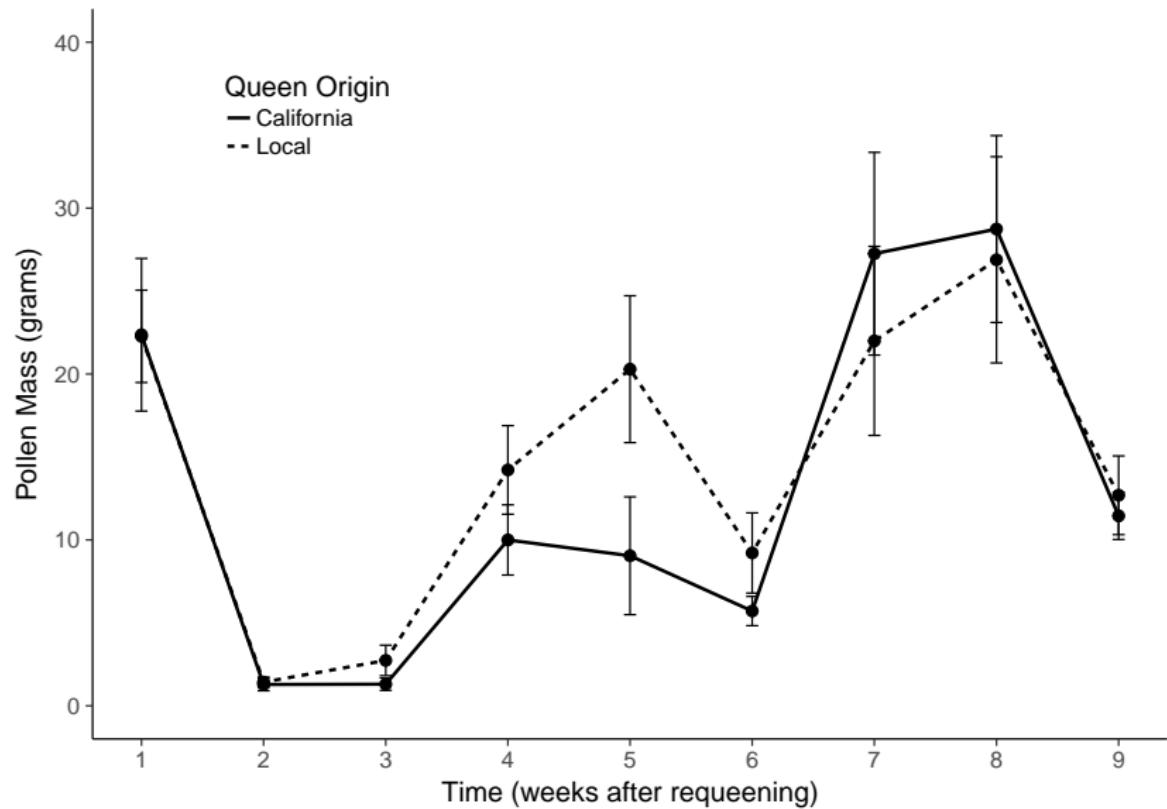
Pollen Collection (foraging)



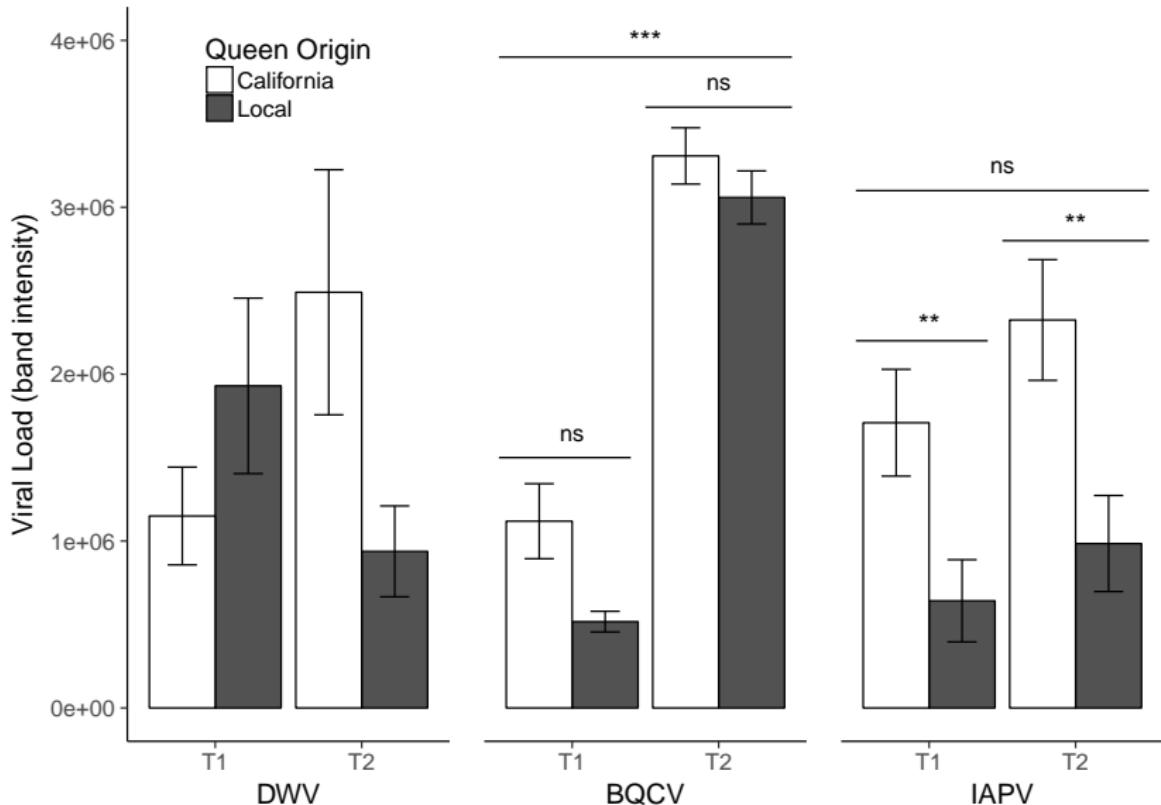
Varroa Load (pathogens)



Nosema Load (pathogens)



Viral Load (pathogens)



In Summary

- ▶ Colony Mass = **Higher in Local**
- ▶ Amount of Brood = **Higher in Local**
- ▶ Pollen Collection = **No Difference**
- ▶ Varroa Load = **No Difference**
- ▶ Nosema Load = **Lower in Local**
- ▶ RNA Viruses = **Mixed Results**

In Summary

- ▶ Overall, colonies re-queened with locally raised queens had higher growth
- ▶ Pollen collection did not seem to be involved in this growth
- ▶ Some pathogens seemed to have less of an effect on local honeybees and others had similar effects across both groups

Implications

- ▶ Locally-raised queens outperform mass-produced, California queens in their northern environment.
- ▶ This could be evidence for the importance of care in breeding stocks (mass produced vs handmade)
- ▶ **And/Or** This could be evidence for local (genetic) adaptation (imported vs. local)

Thank you!



Questions?

