

US EPA ARCHIVE DOCUMENT

Importance of Basic and Translational Approaches in Bed Bug Research



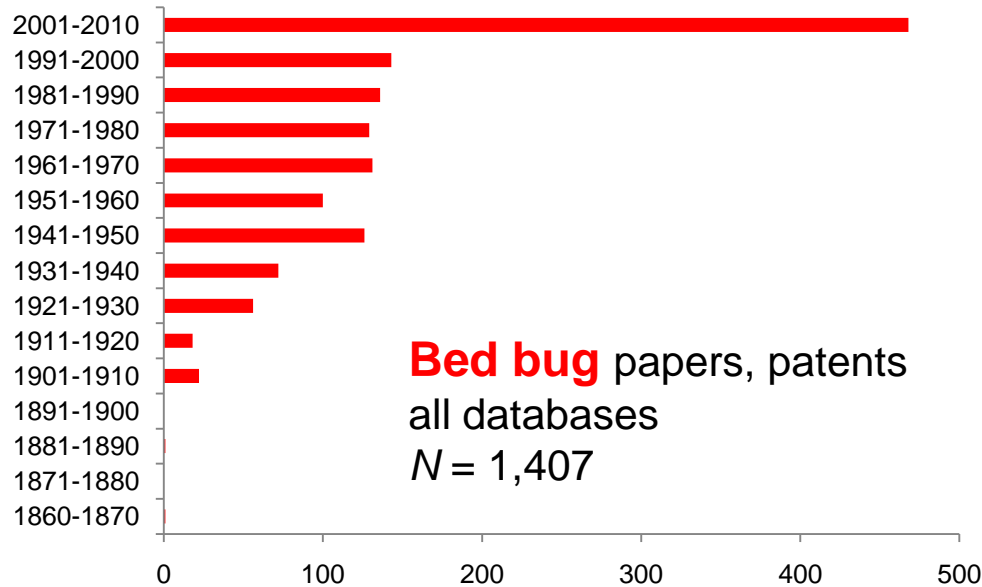
Coby Schal <coby_schal@ncsu.edu>

**Department of Entomology
W. M. Keck Center for Behavioral Biology
North Carolina State University**

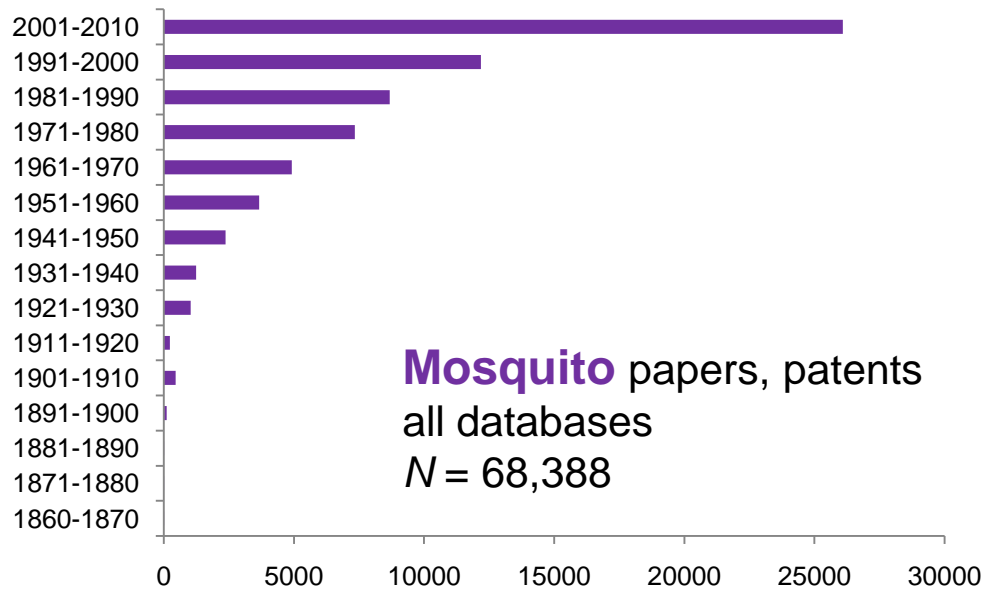
Presentation Outline

- **Trends of scientific publishing on BB**
- **Research approaches**
- **3 Examples of translational research**
 1. Chemical ecology: Pheromones, other semiochemicals
 2. Population genetics
 3. Gene expression & transcriptomes
- **Perspective: Fundamental & translational research**

Literature trends

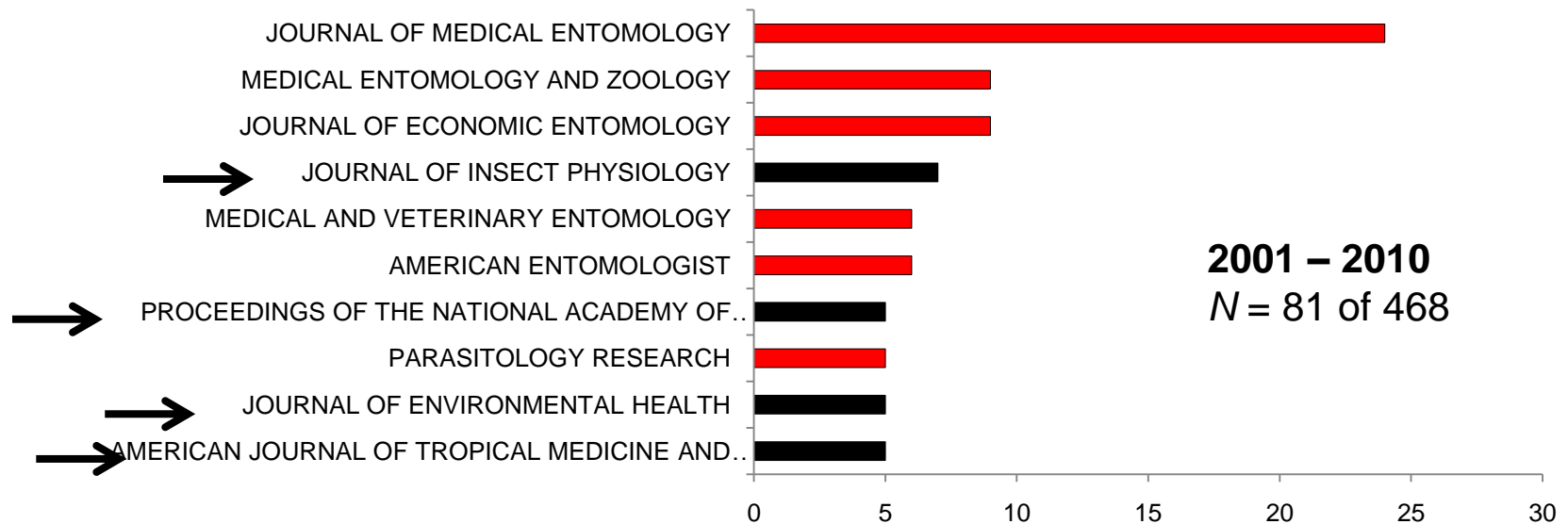
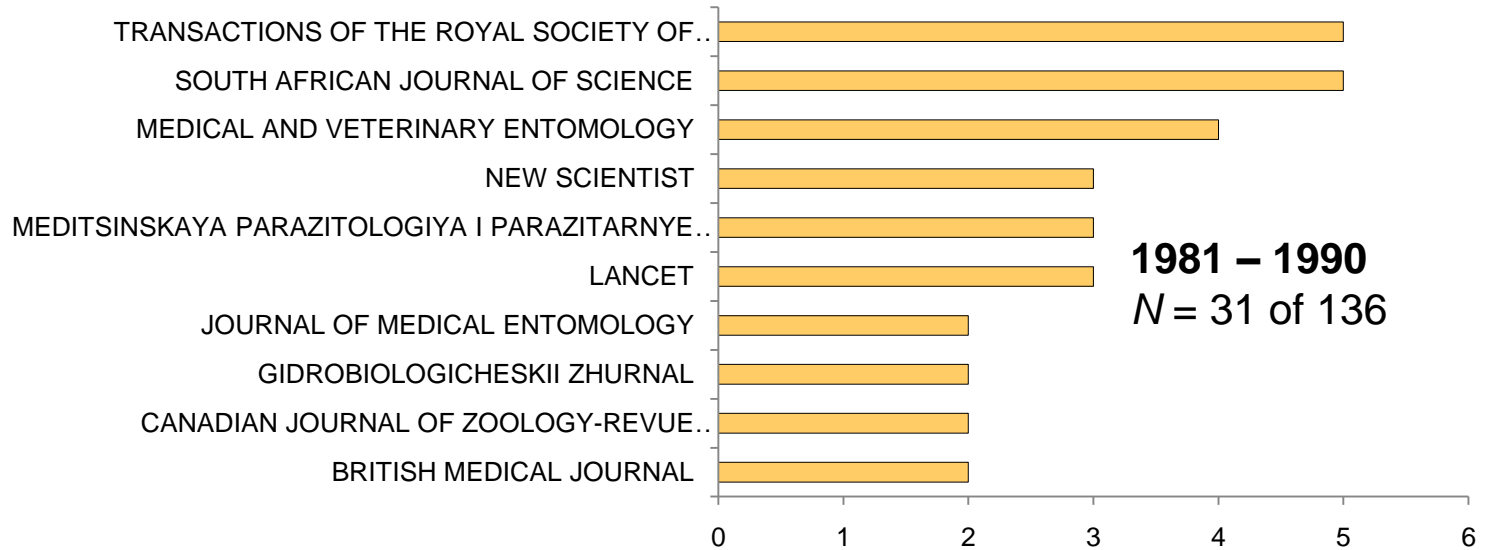


- “flat” productivity for 60 yrs – 1940 – 2000
- upsurge in information in the last decade



- mosquito : BB = 50 :1

Bed bugs: all databases, top 10 journals



Linear Model

Society benefits with information and cool technologies and sometimes...information for practitioners and decision makers

Applied research and development... stem from basic research spin-offs

Basic research, inspired by curiosity,...adds to 'pool of knowledge'

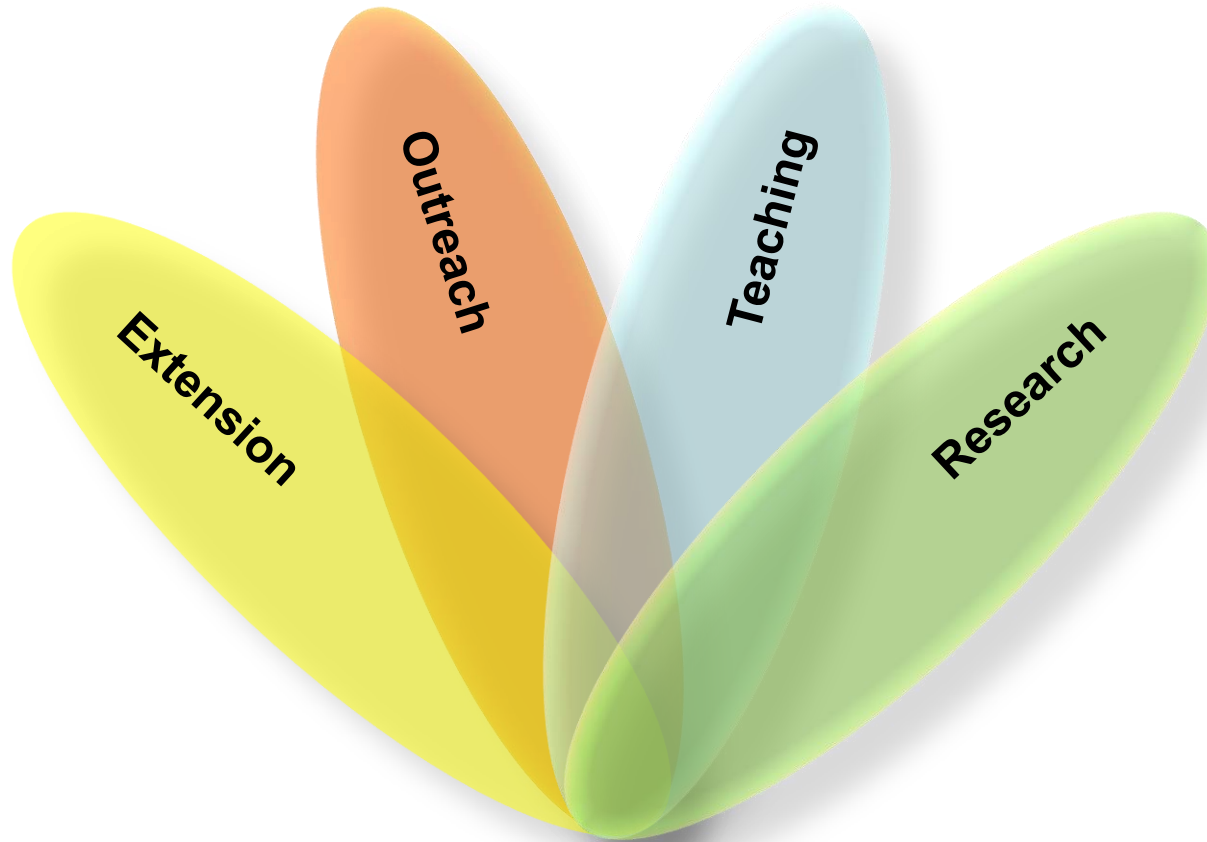
Government invests in "hands-off"
Basic research

- Applied research poorly funded
- Compartmentalization: Different teams engaged in basic & applied
- Basic researcher may be off on another project before application

Long time, often decades

21st Century Approach:

Integrating Research, Extension, Outreach and Teaching



Translational research

mission-oriented, multidisciplinary, incentivized
shorter time-frame to practical applications

A (partial) 'Laundry' List of Needs

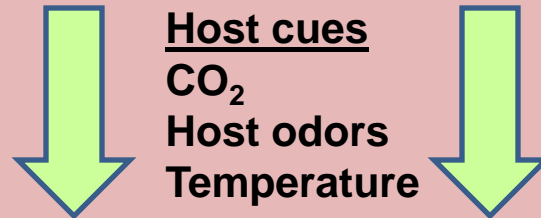
- Disease vectors, bite allergies, asthma
 - Abiotic interactions: temp, humidity
 - Biological control agents: viruses, fungi, bacteria
 - New pesticides, modes of action, resistance
 - Novel non-chemical targets: cuticle, midgut, ovaries
1. BB behavior, communication, aggregation
 2. Population biology
 3. Unique genes and interactions of gene products, BB genome
- More topics later..... Dr. Kells

Example 1: Host finding, chemicals

Host seeking



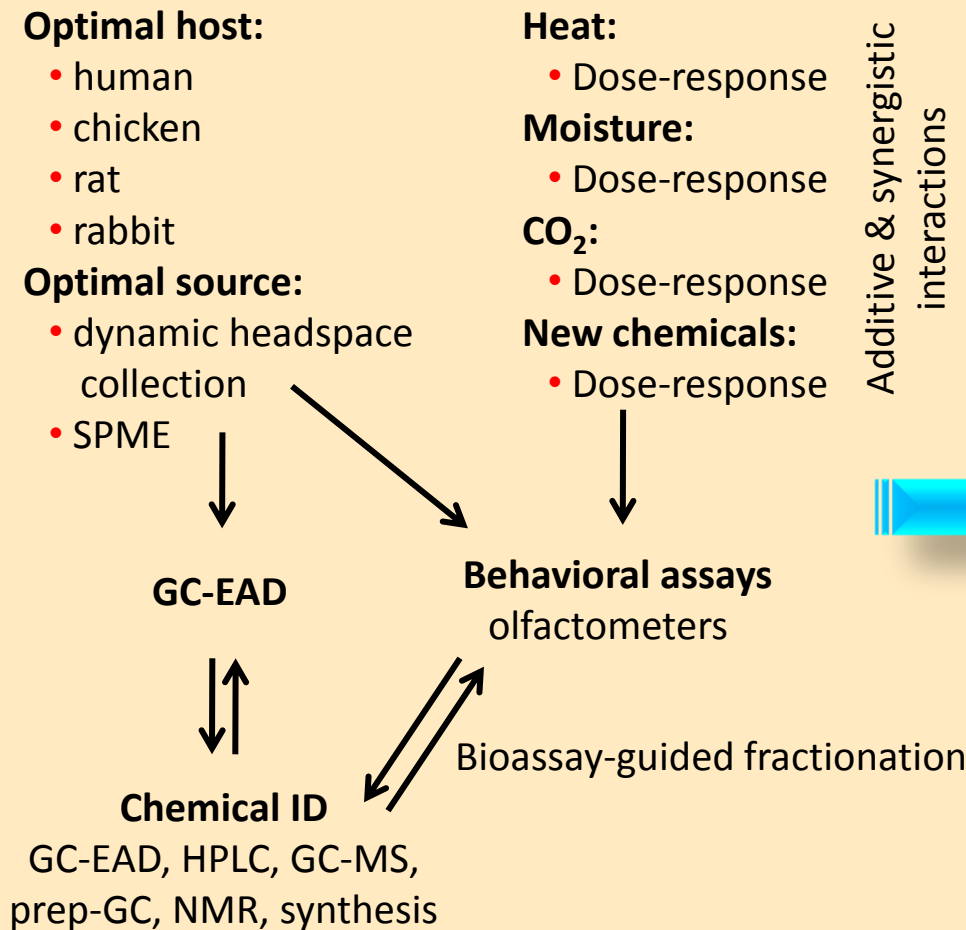
Host location



Host rejection/acceptance; engorgement

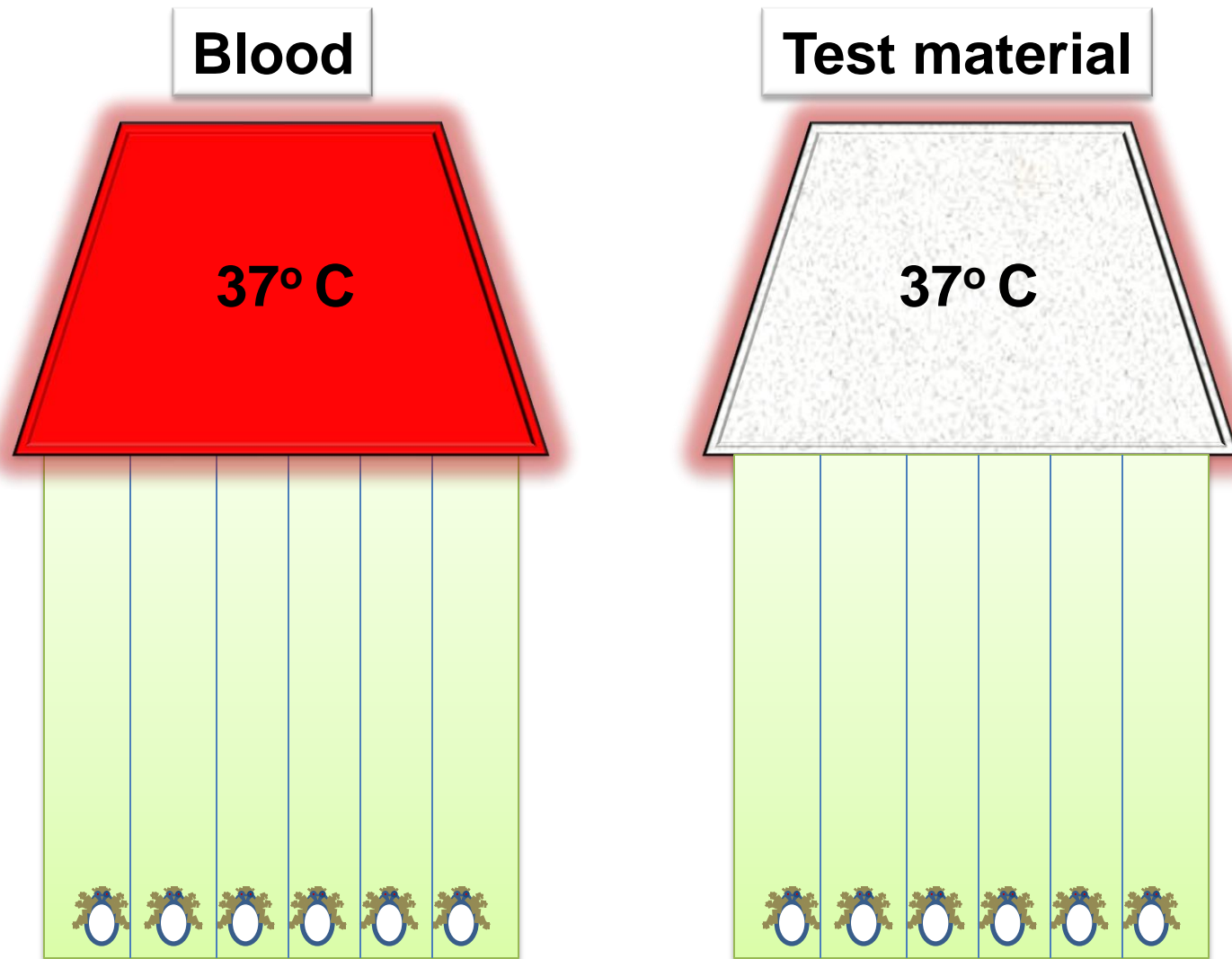


Sensory ecology associated with host-seeking behavior

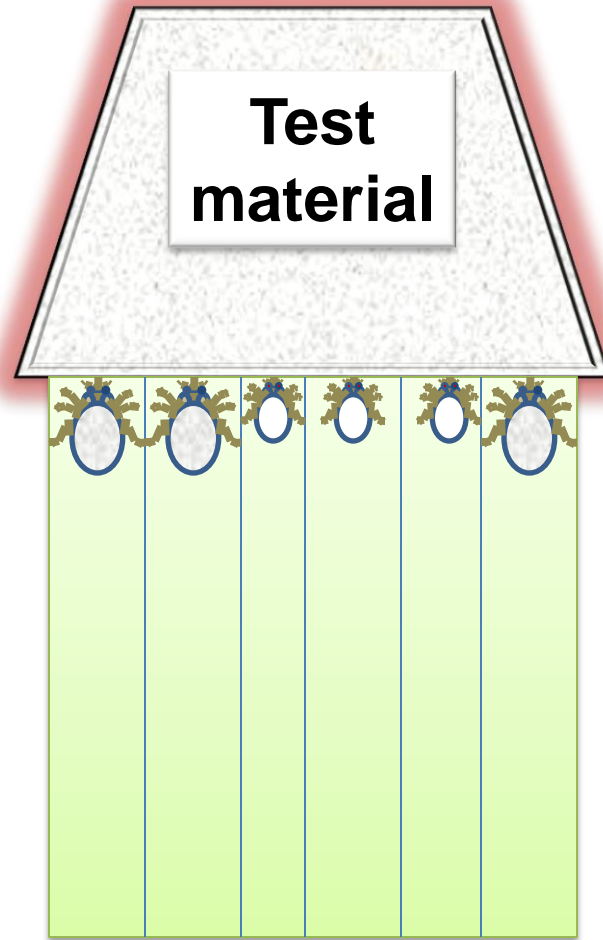
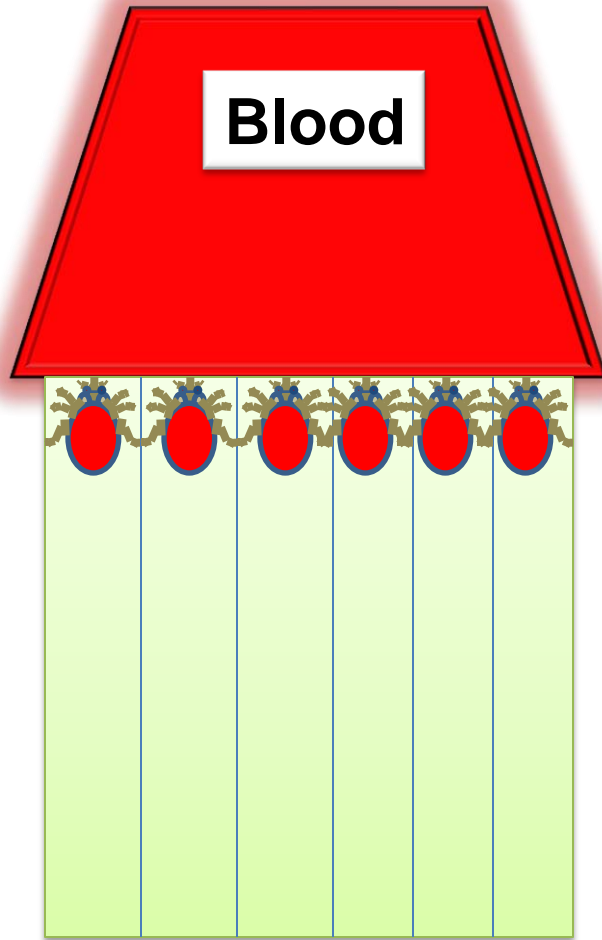


- Host-attractants for traps
- Sex pheromones
- Aggregation pheromones
 - ✓ Detection
 - ✓ Monitoring
 - ✓ Dissemination of biological control agents
 - ✓ Attract-and-Kill
- Repellents
- Attractants for baits?
- Phagostimulants
- **More..... Dr. Feldlaufer**

What's in blood that bed bugs like?

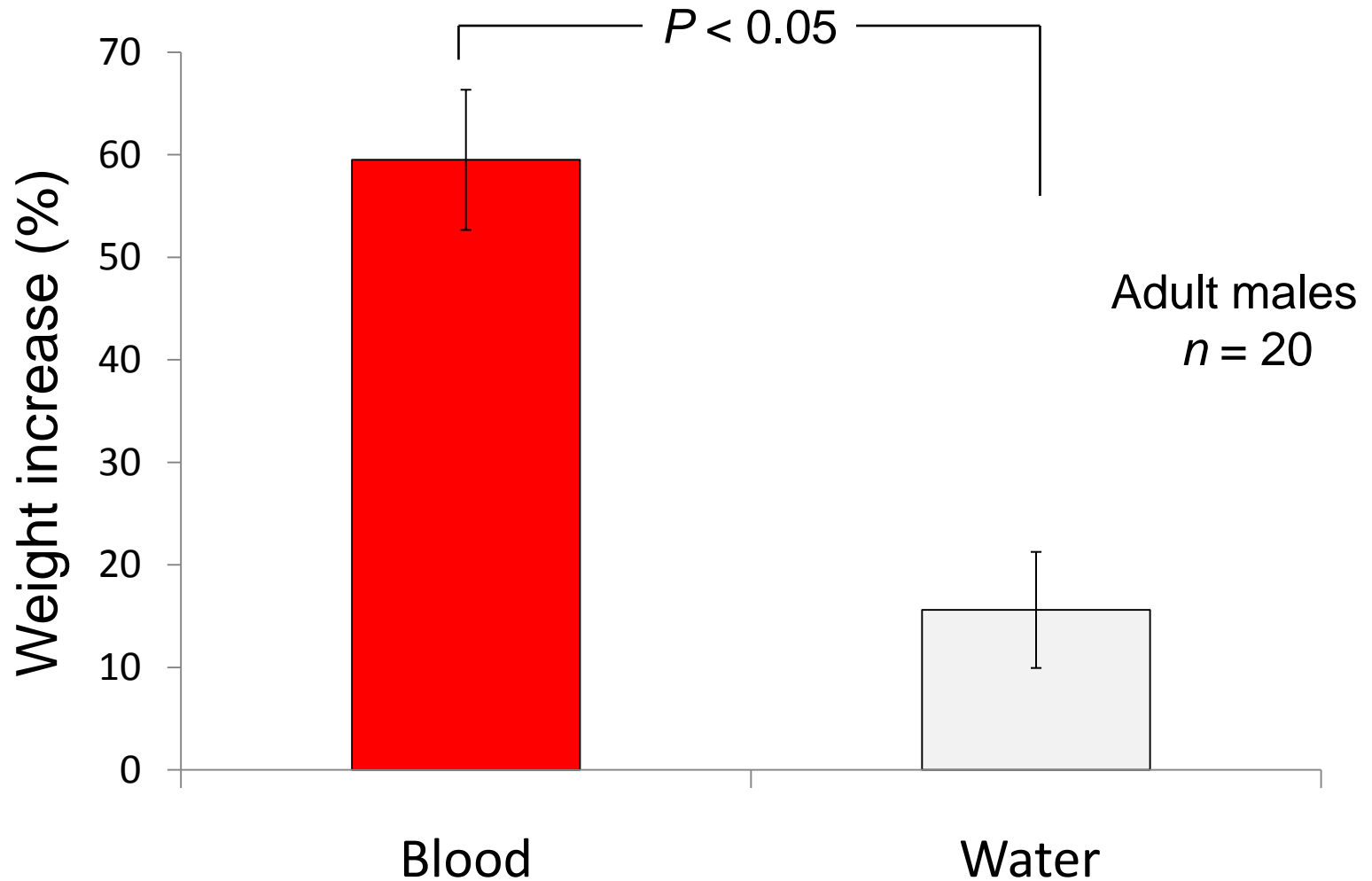


15 min feeding period

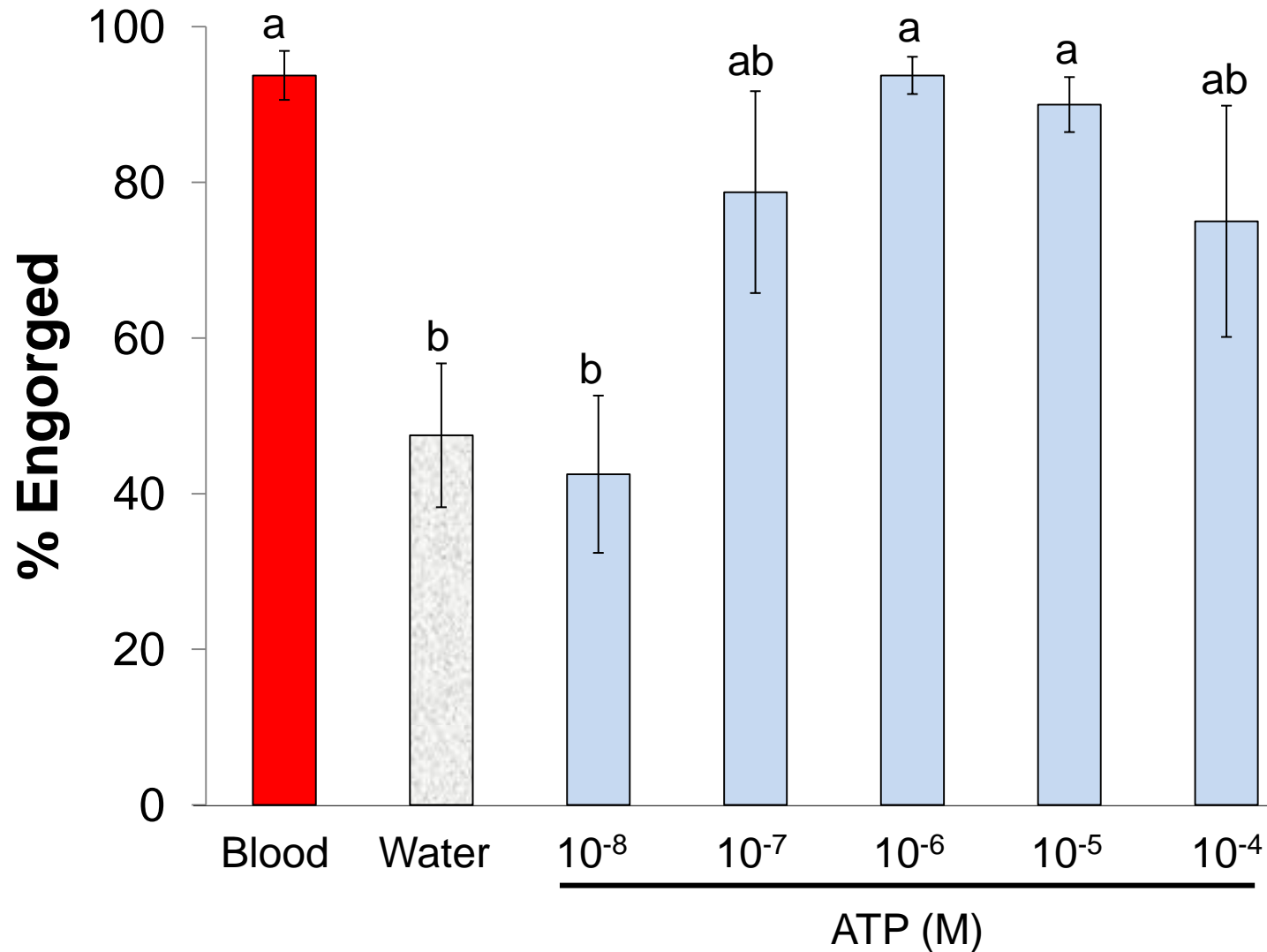
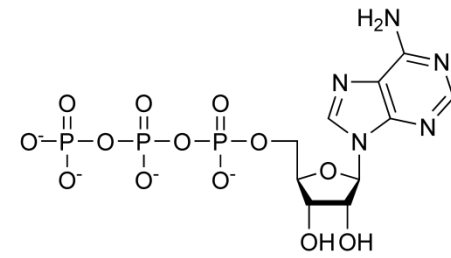


Individual bugs weighed before and after feeding

Blood contains phagostimulants



Water + ATP = Blood!



- Blood contains phagostimulants
- ATP is a highly effective phagostimulant
- All life stages displayed similar gorging responses to ATP
- ATP is a more effective phagostimulant than other adenosine nucleotides
ATP > ADP > AMP
- Other phagostimulants?



Richard Naylor



Alvaro Romero

Example 2: Population genetics

Where have bed bugs resurged from?

3 Major hypotheses:

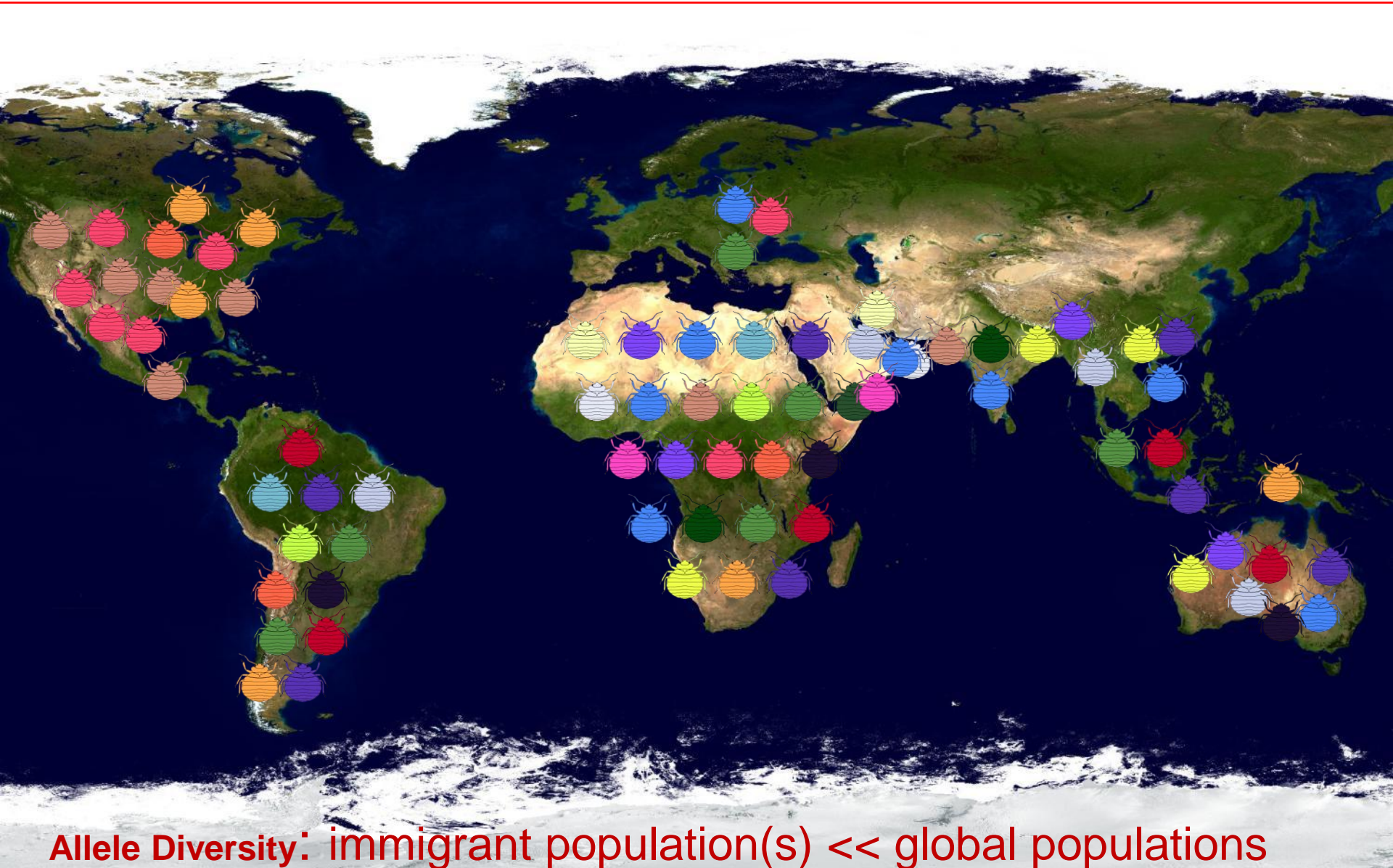
Local “home grown” bed bug populations

1. Residential, urban reservoirs
2. Agricultural (e.g., poultry farms) reservoirs

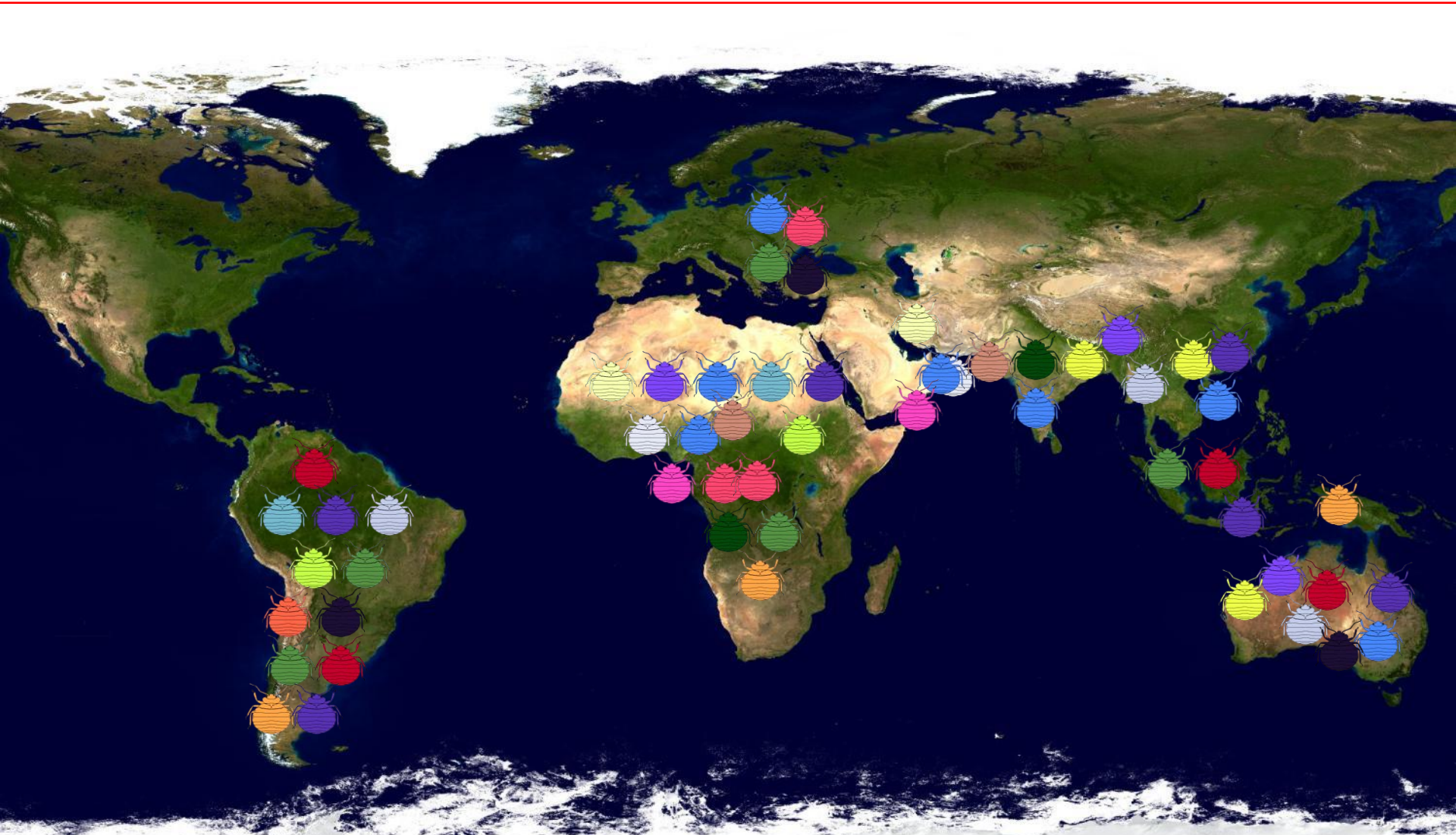
Global “immigrant” bed bugs

3. Outside USA: travel, globalization

A single, or few, invasion(s): Low diversity



Multiple, independent invasions: High diversity



Allele Diversity: immigrant population S ~ source population S

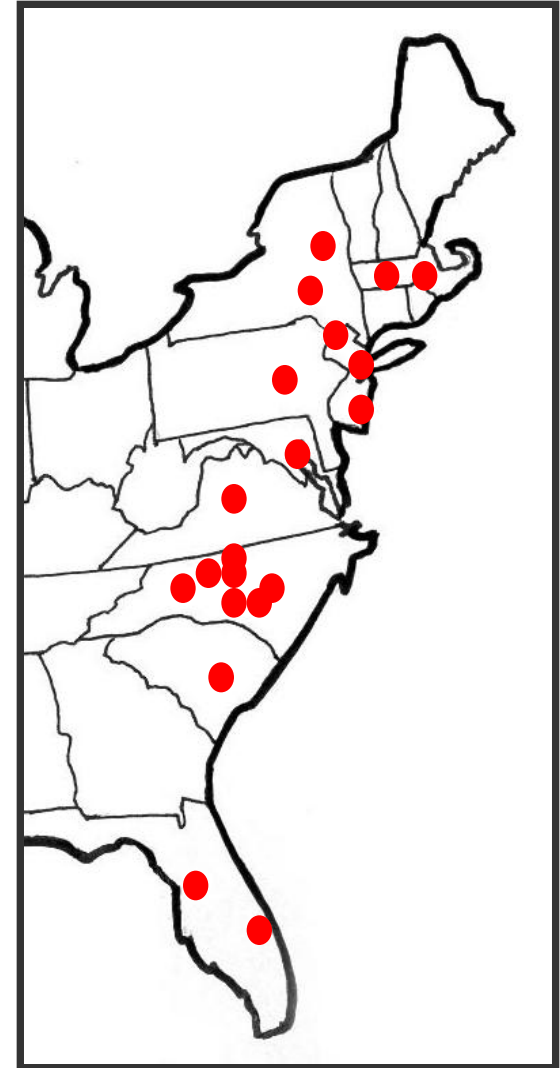
Broad-scale BB population genetics

(Vargo, Booth, Saenz, Schal)

- Microsatellite markers:
 - ✓ Genetic diversity
 - ✓ Genetic differentiation
- 21 populations from 9 states



Virna Saenz



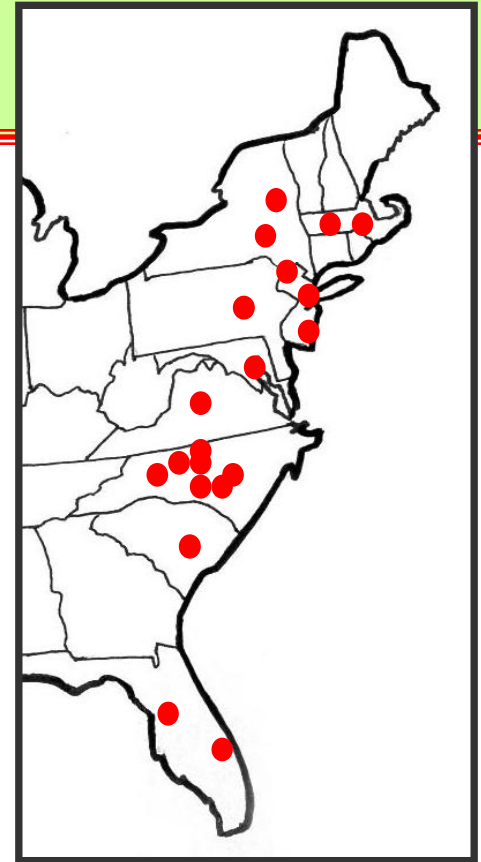
Genetic diversity

Across populations

- **Very high genetic diversity**
 - 3 to 17 alleles/locus (mean = 8.4)
 - But, need comparisons to “candidate” source populations
 - Suggests multiple sources

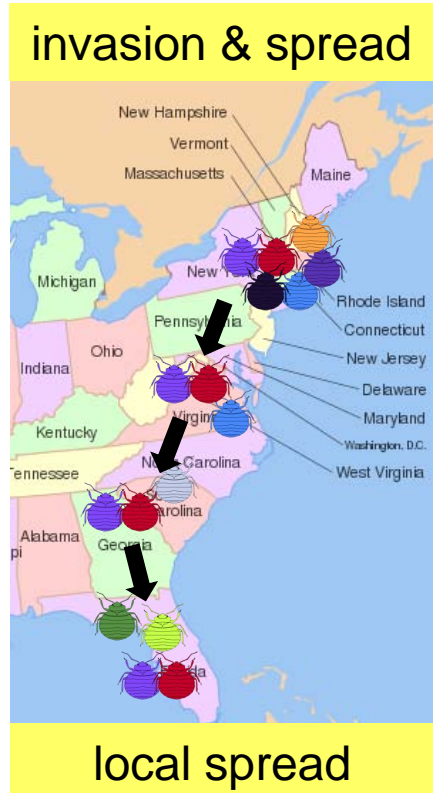
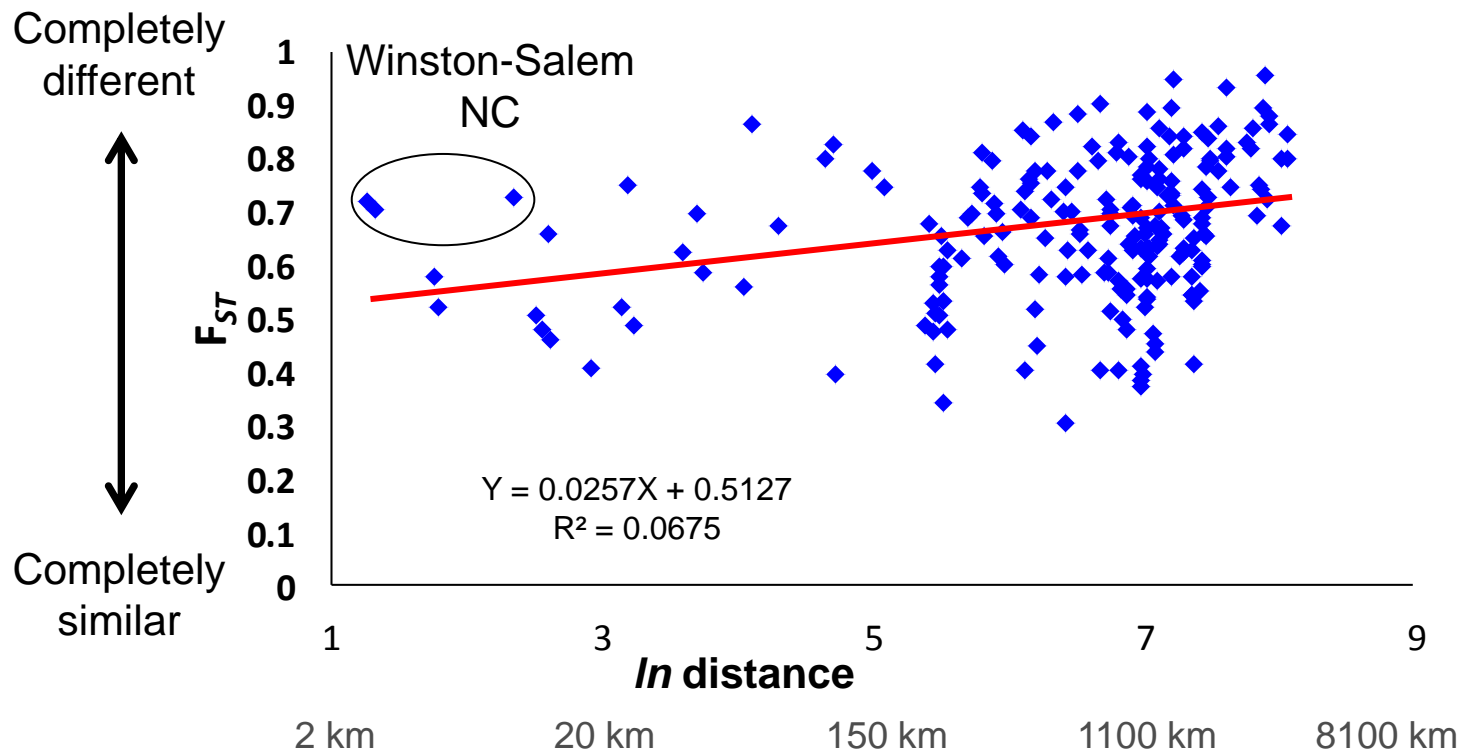
Within populations

- **Very low genetic diversity**
 - 1 to 4 alleles/locus
 - Mean relatedness of individuals
 - $r = 0.75$ (0.5 = sibs)
 - Very high levels of inbreeding
 - Suggests single/few sources per infestation



Genetic differentiation of populations

strong genetic differentiation!
but, no “Isolation-by-Distance”



Broad-scale BB population genetics

(Vargo, Booth, Saenz, Schal)

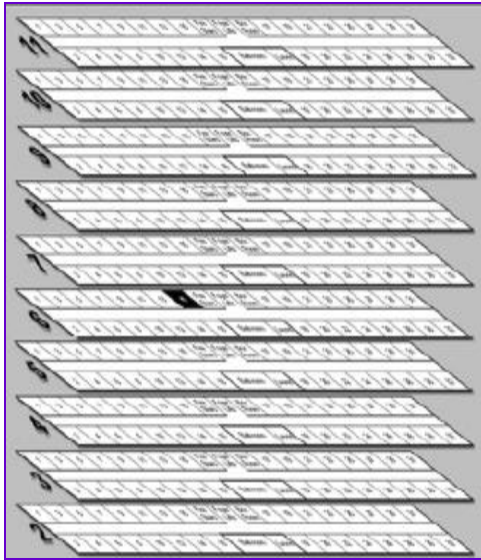
- **Within** populations: Very low genetic diversity & high inbreeding
 - Populations appear to be founded by a singly mated female

yet,

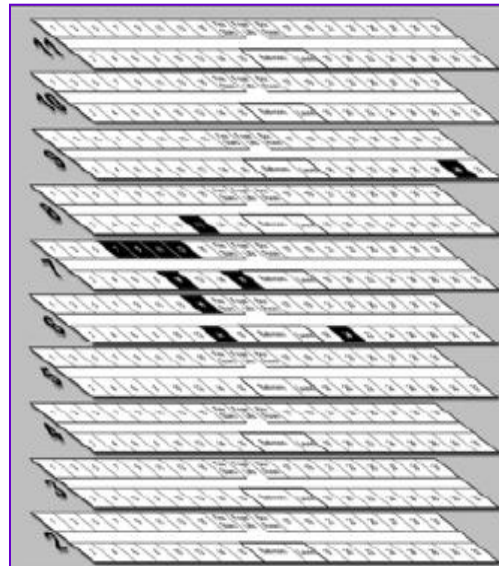
- **Across** populations: Very high genetic diversity & high genetic differentiation
 - Multiple introductions
 - “Jump”, human-mediated transport

How do bed bugs spread within a building?

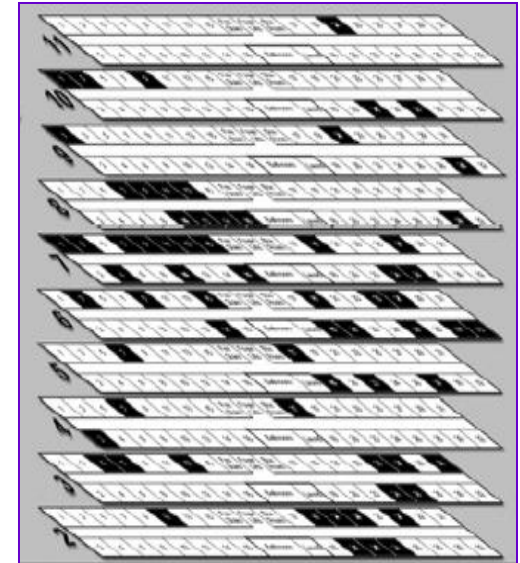
May 2003



May 2004

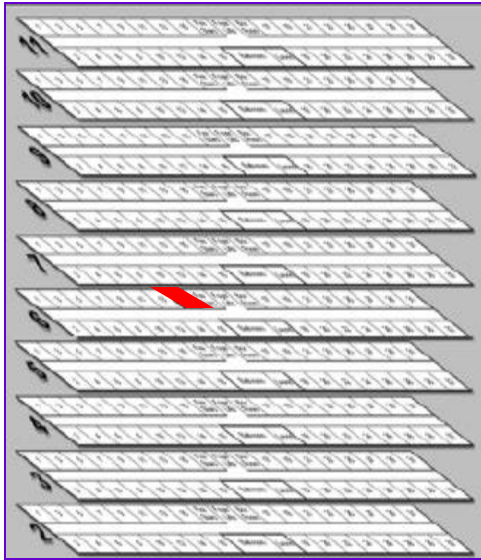


June 2005

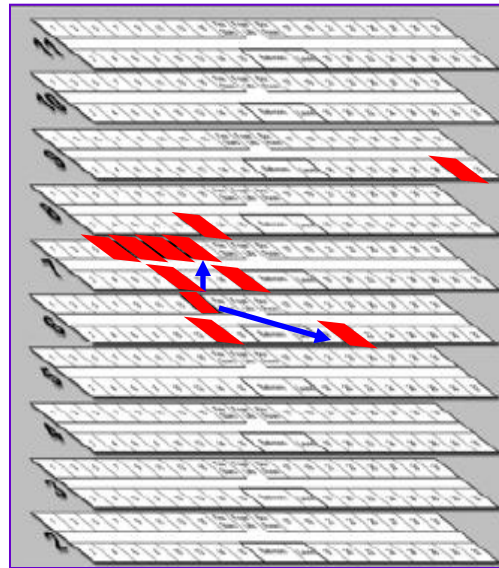


How do bed bugs spread within a building?

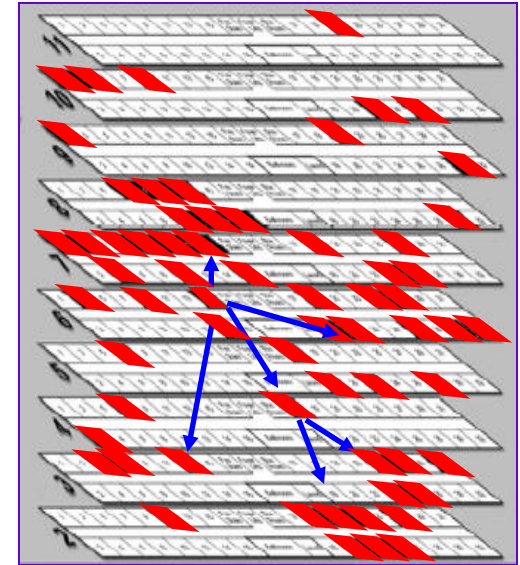
May 2003



May 2004



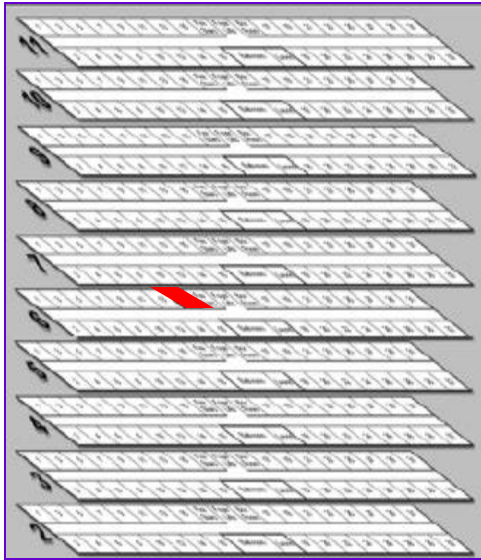
June 2005



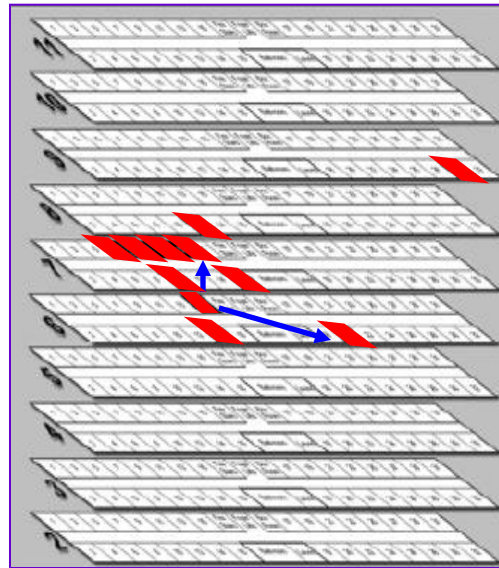
Single introduction?

How do bed bugs spread within a building?

May 2003

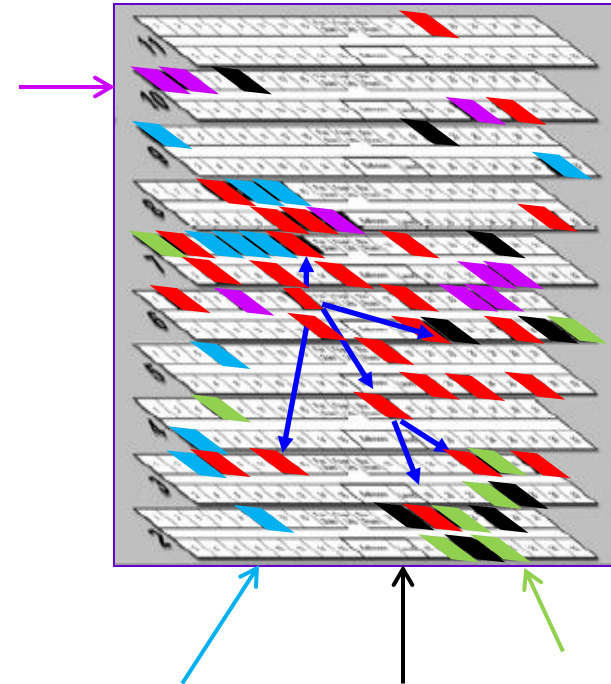


May 2004



Single
introduction

June 2005



Multiple
introductions?

Empirical test: BB spread within a building

How many individuals start an infestation?

Within apartments

- Low genetic diversity
 - ✓ 1 to 4 alleles/locus → Few!
- High relatedness
 - ✓ $r = 0.54$

How many introductions?

Among apartments

- Low genetic diversity
 - ✓ 2 to 5 alleles/locus → Single!

Raleigh, NC
Isolated building, recent infestation

Floors	9 th	916	915	914	913		912	911	910		909	908	907
													906
		917	918	919	920		901		902		903	904	905
	8 th	816	815	814	813		812	811	810		809	808	807
													806
		817	818	819	820		801		802		803	804	805
	7 th	716	715	714	713		712	711	710		709	708	707
													706
		717	718	719	720		701		702		703	704	705
	6 th	616	615	614	613		612	611	610		609	608	607
													606
		617	618	619	620		601		602		603	604	605
5 th		516	515	514	513		512	511	510		509	508	507
													506
		517	518	519	520		501		502		503	504	505
4 th		416	415	414	413		412	411	410		409	408	407
													406
		417	418	419	420		401		402		403	404	405
3 rd		316	315	314	313		312	311	310		309	308	307
													306
		317	318	319	320		301		302		303	304	305

Empirical test: BB spread within a building

How many individuals start an infestation?

Within apartments

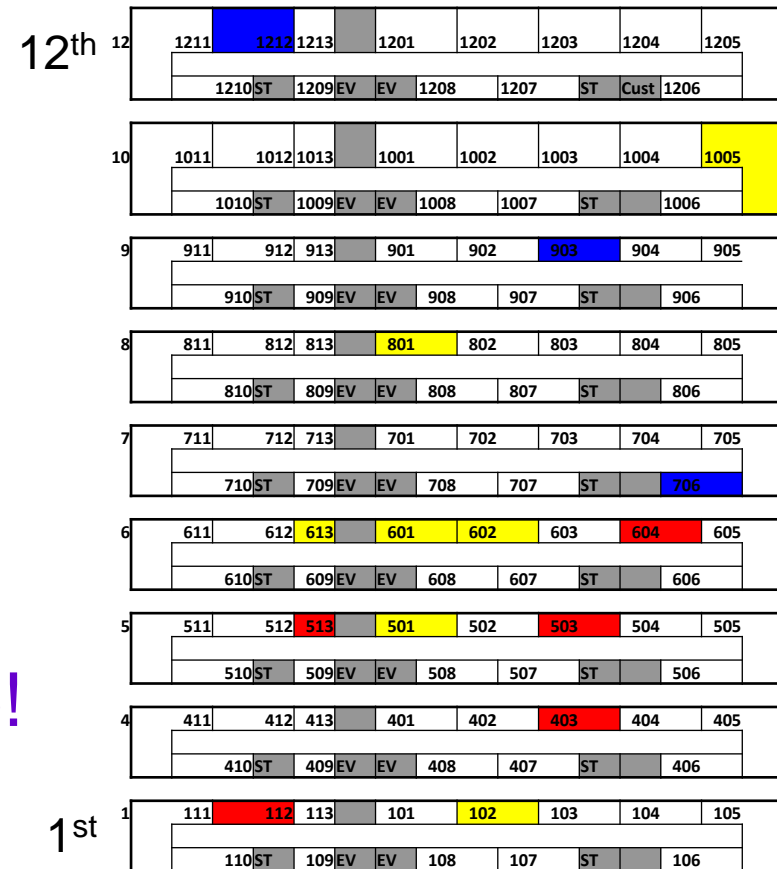
- Low genetic diversity
 - ✓ 1 to 4 alleles/locus → Few!
- High relatedness
 - ✓ $r = 0.65$

How many introductions?

Among apartments

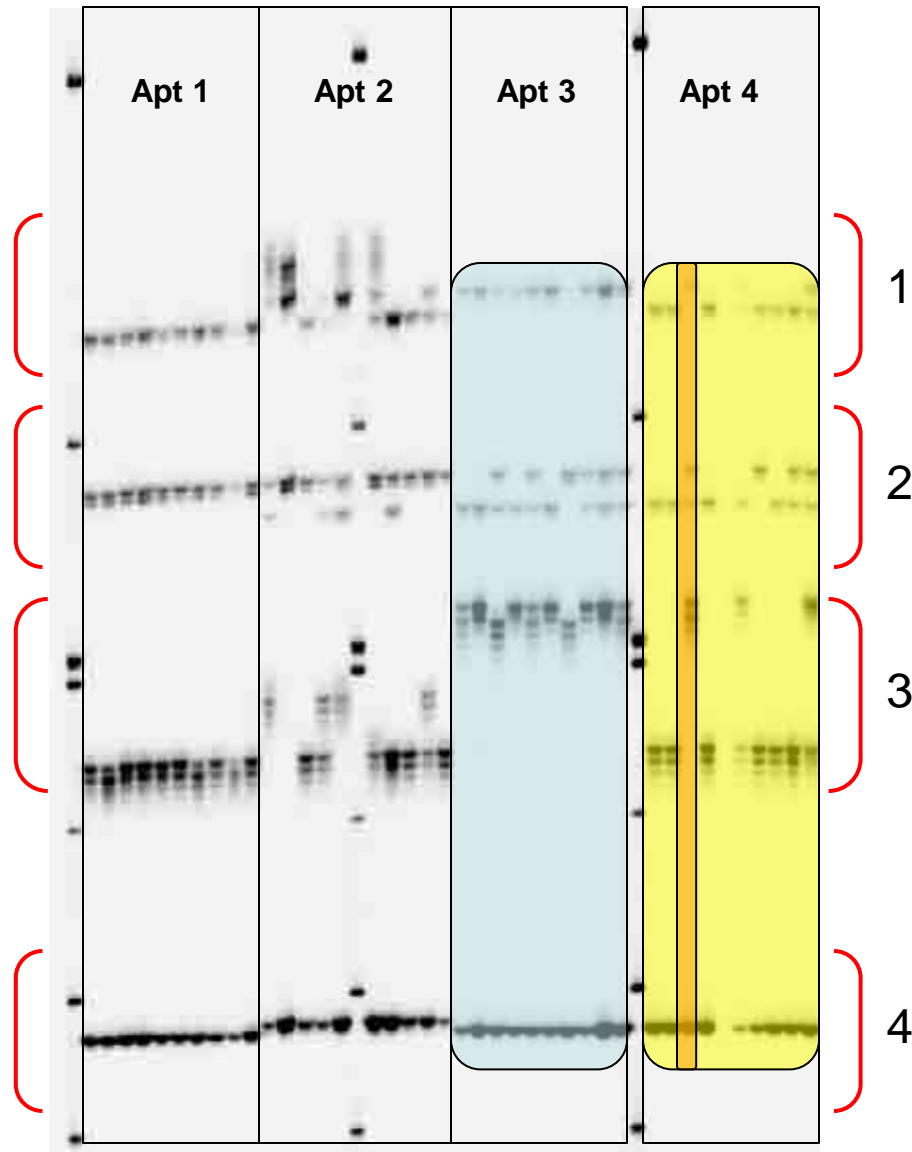
- High genetic diversity
 - ✓ 2 to 11 alleles/locus → Multiple!

Jersey City, NJ
Building complex, old infestation



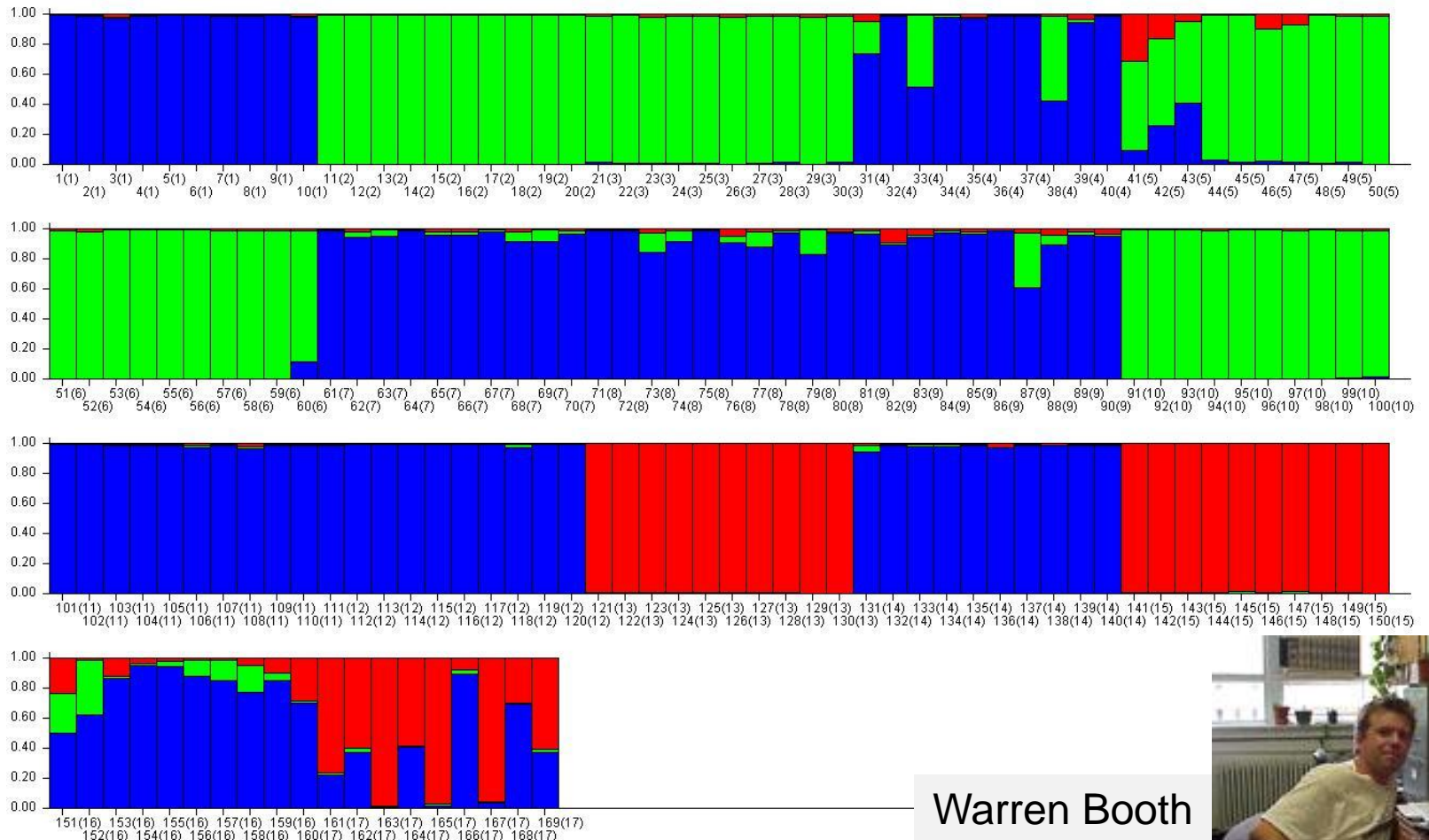
collections by Changlu Wang

Empirical test: BB spread within a building



Empirical test: BB spread within a building

Jersey City, NJ: 17 populations
3 genetic clusters



Warren Booth

software: STRUCTURE



Implications of Population Genetics

Large scale (continental, East Coast)

- Bed bug populations come from multiple geographic sources
- Extensive human-mediated movement of bed bugs across large geographic areas

Interception important at transportation hubs

Fine scale (within buildings)

- Infestations often start with a single introduction, and then spread; but also multiple introductions

Early detection critical

Forensic applications: sources of bed bug infestations

Example 3: Genome, Transcriptome

- Genes common to BB and other insects?
- Genes common to hematophagous arthropods
- Genes and gene products unique to BB
- Why do BB not vector diseases?
- When and where genes are turned on or off
(transcripts specific to tissue-, time-, instar-,
feeding-, resistance-, etc.)
- Gene networks:
immunity, heat/cold response, feeding behavior,
sexual behavior, host-finding.....

Example 3: Genome, Transcriptome

OPEN ACCESS Freely available online

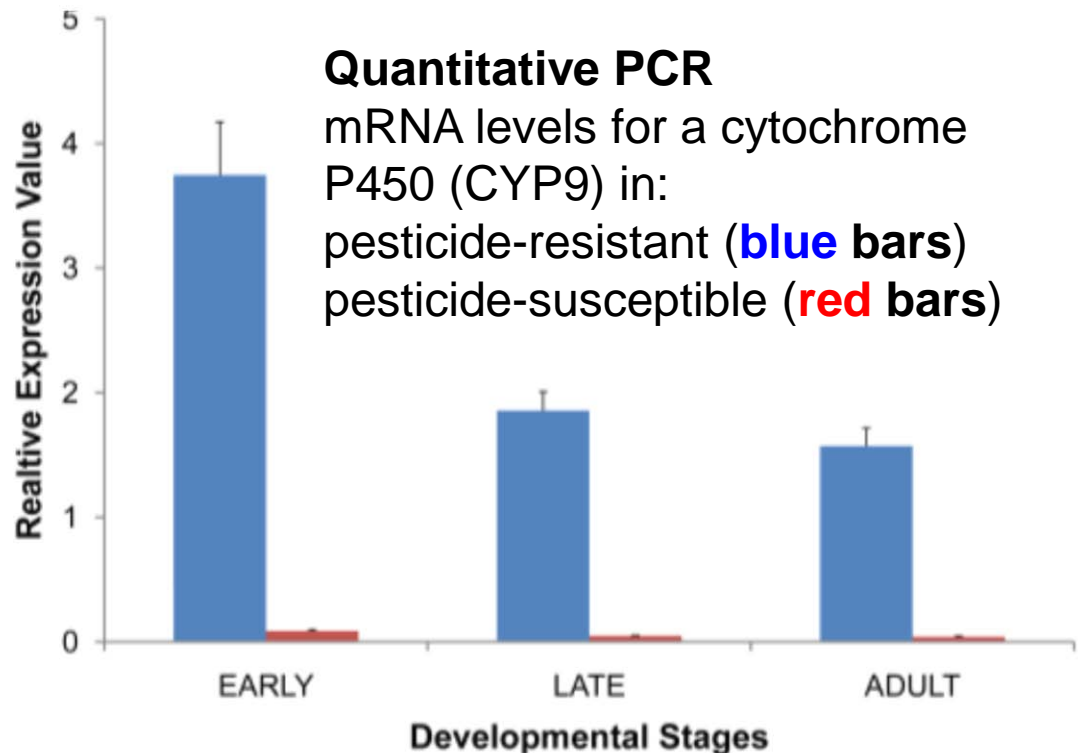


Transcriptomics of the Bed Bug (*Cimex lectularius*)

Xiaodong Bai¹*, Praveen Mamidala¹*, Swapna P. Rajarapu¹, Susan C. Jones², Omprakash Mittapalli^{1*}

¹ Department of Entomology, Ohio Agricultural and Research Development Center, The Ohio State University, Wooster, Ohio, United States of America, ² Department of Entomology, The Ohio State University, Columbus, Ohio, United States of America

“Nearly 85.9% of the *C. lectularius* sequences showed similarity to insect sequences, but **44.8%** of the deduced proteins of *C. lectularius* did not show similarity with sequences in the GenBank”



Perspective: Fundamental and Translational research

- **Government & private sector investments**
- **Innovative, medium- and long-term**
- **Accountable to stakeholder input**
- **Spawns ideas for novel tactics, products**
- **Research \rightleftharpoons Pest control**
- **Apply discoveries from research in the lab to “clinical” studies**
- **Break down compartmentalization**

Collaborators & co-workers: Ed Vargo

**Warren Booth, Alvaro Romero, Virna Saenz, Rick Santangelo
Changlu Wang**

Blanton J. Whitmire Endowment

NC State University



USDA-NIFA-AFRI



**NSF Postdoctoral Research
Fellowship in Biology**



Pest management companies:

Orkin

Cooper

Rentokil/JC Ehrlich

Terminix

Life cycle

Egg to Adult = 45 days at 26° C



bedbuginfo.dondodd.com/



sciencedude.ocregister.com

