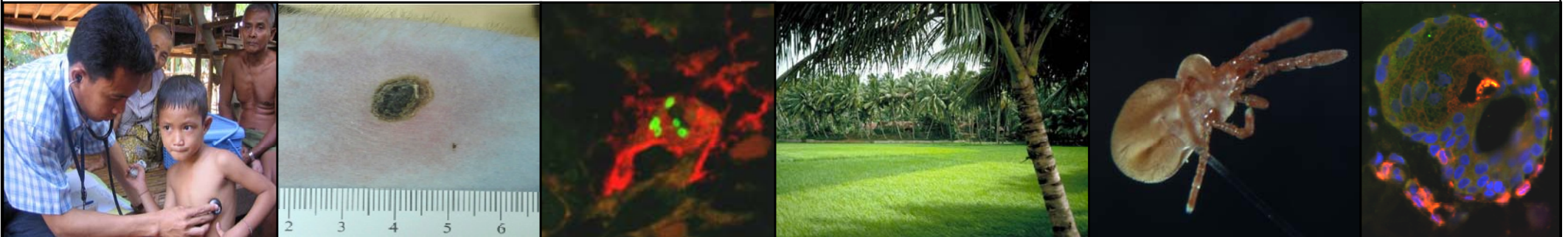


# Unresolved Entomological issues related to Scrub Typhus

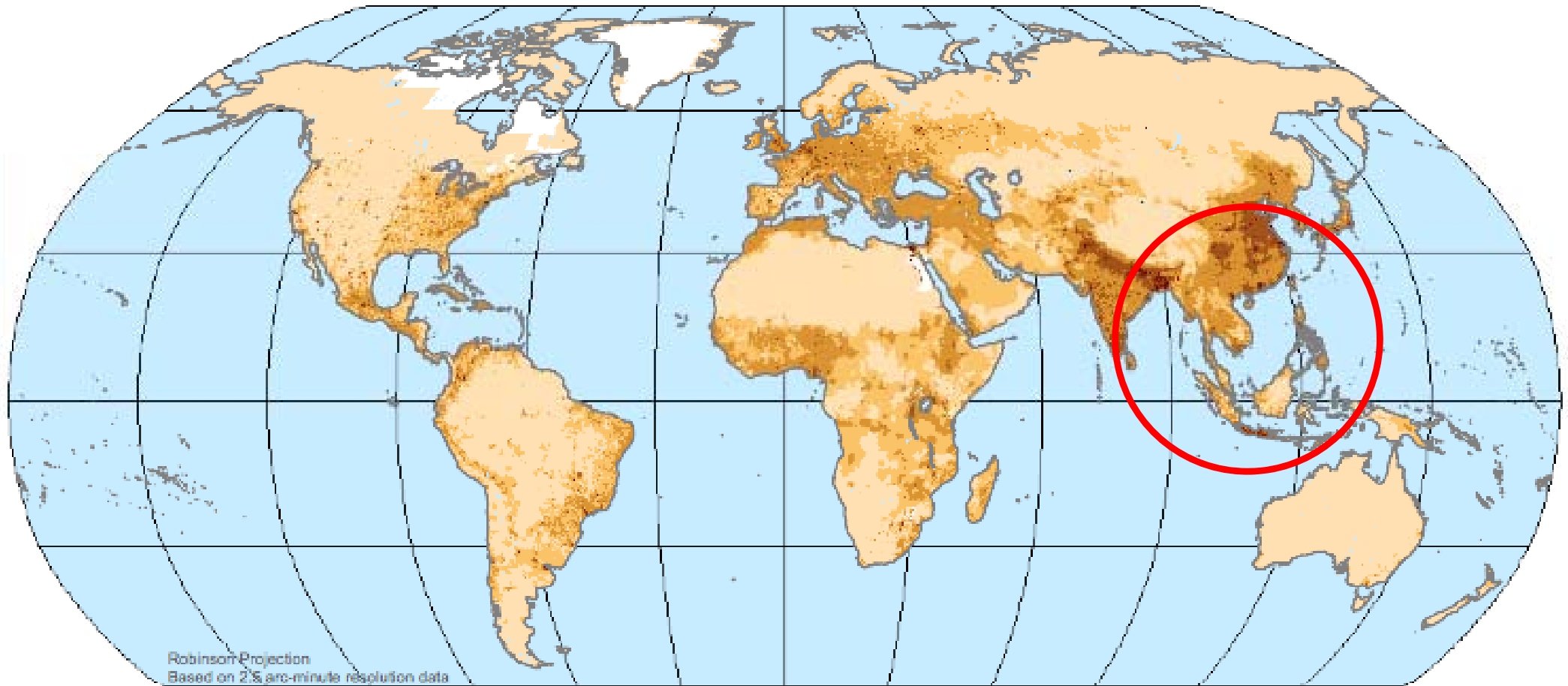
Daniel H. Paris

MORU, Bangkok, Thailand

JITMM, November 2013

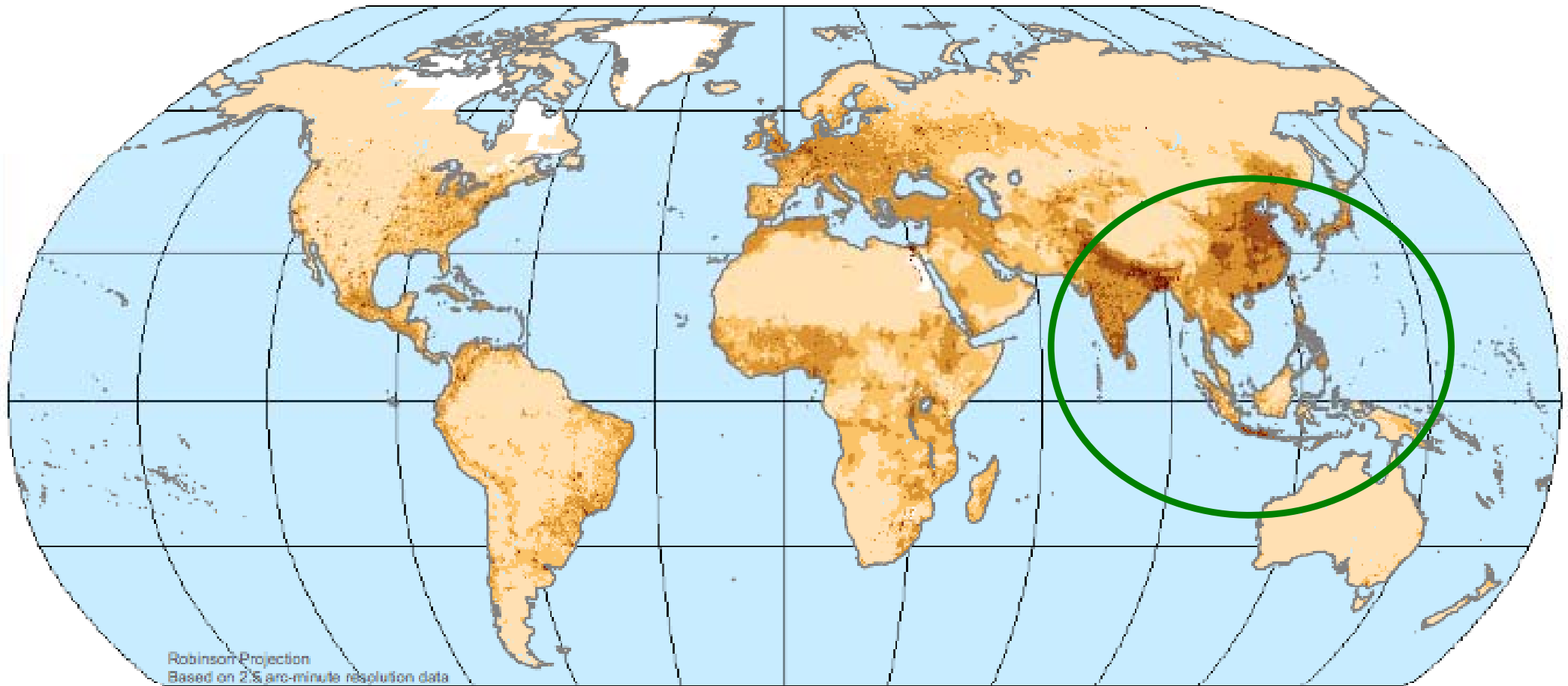


## The importance of Asia ...



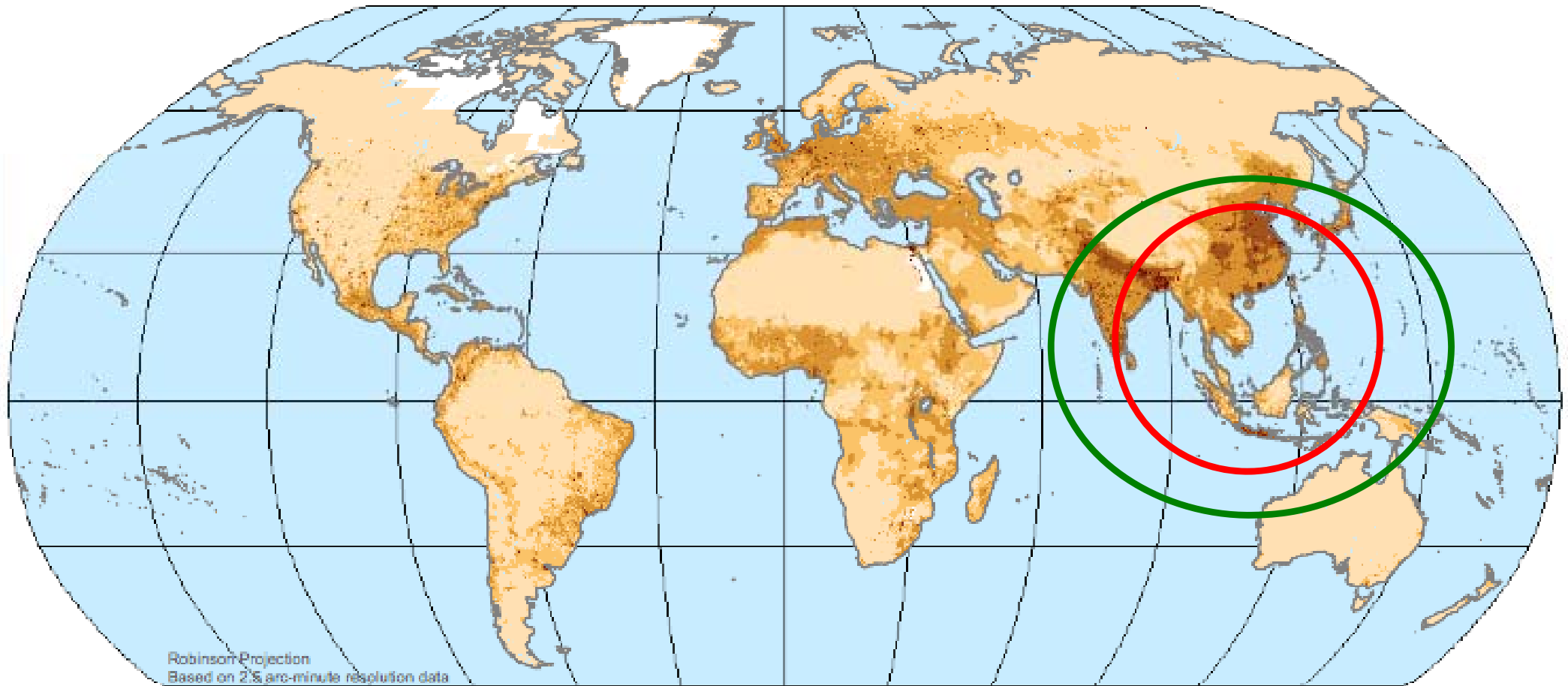
***Half of the world's population within 2000 miles of Bangkok !!!***

## The importance of Asia ...



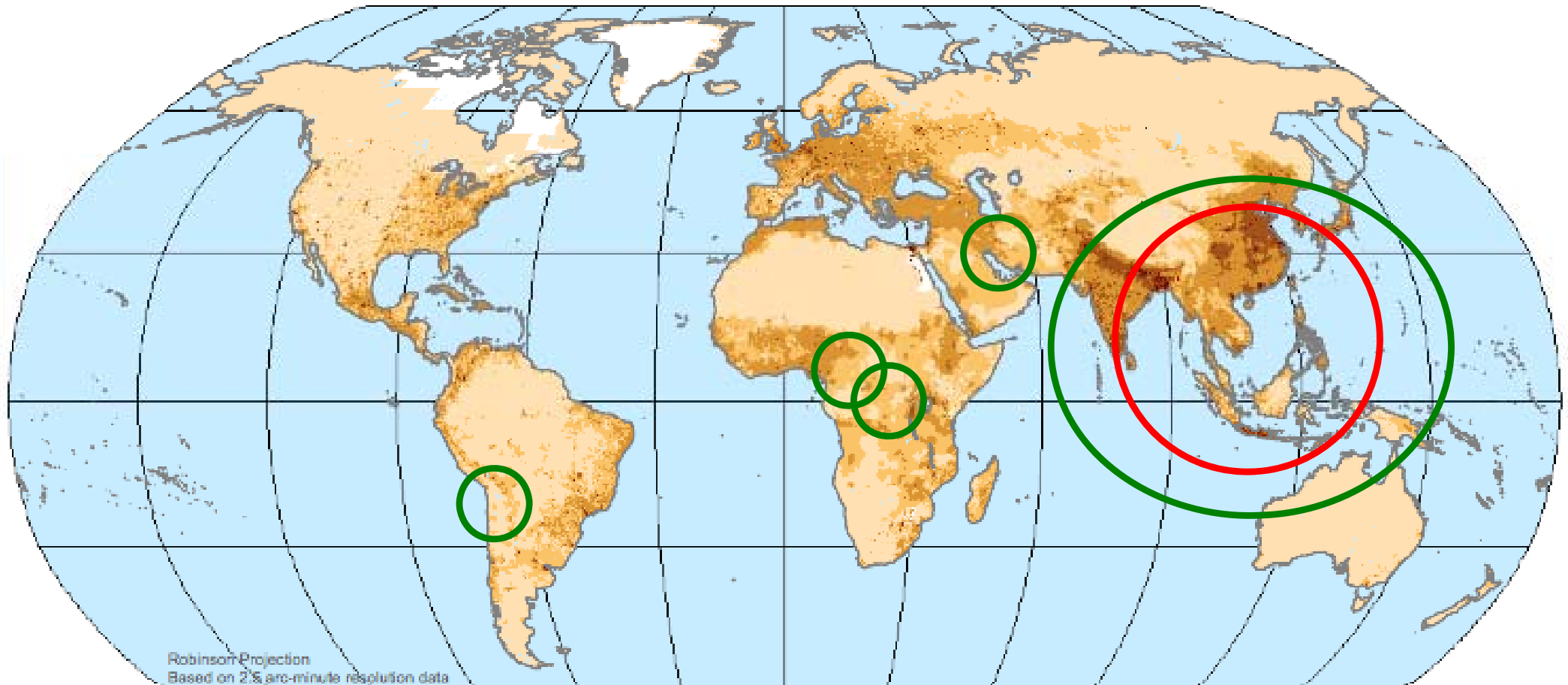
**>90 % of reported scrub typhus cases within 3000 miles of Bangkok !!!**

# The importance of Asia ...



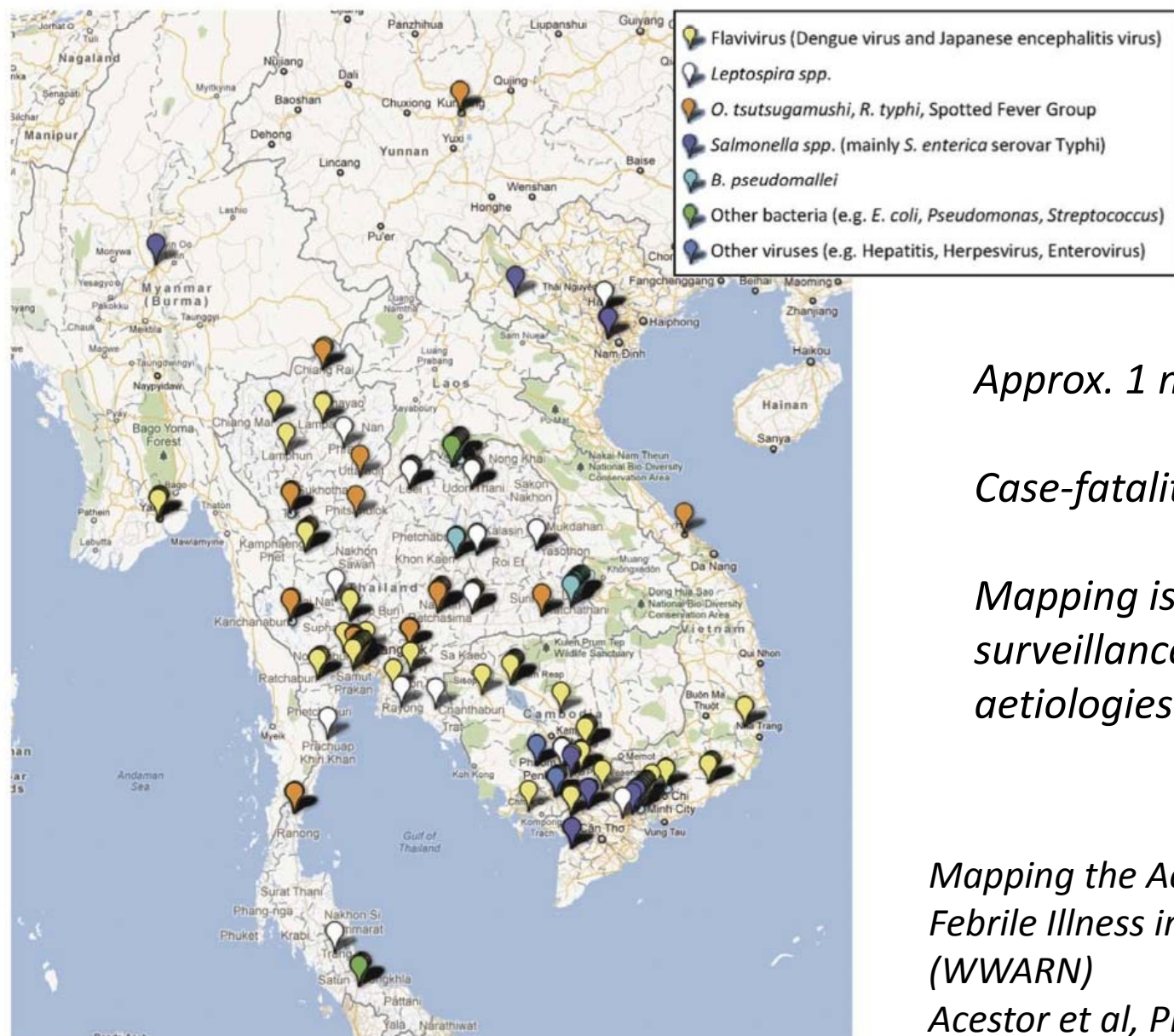
*... What about the rest of the world ... ???!*

# The importance of Asia ...



*... potential global tropical / subtropical distribution ... ?!*

# How important is scrub typhus?



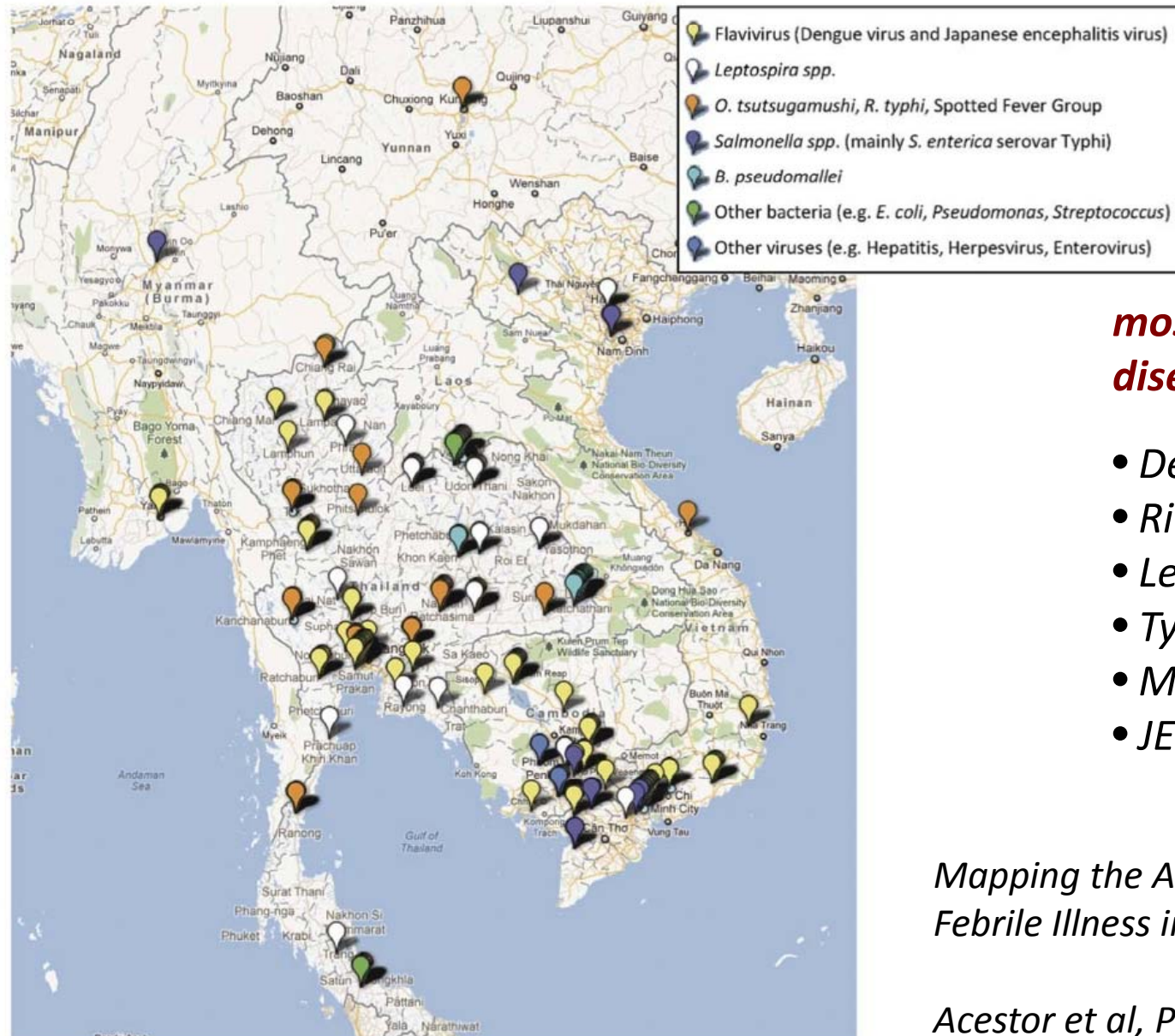
Approx. 1 mio cases estimated

Case-fatality rate 3-7%

Mapping is an excellent surveillance tool to identify aetiologies and localise 'gaps'

Mapping the Aetiology of Non-Malarial Febrile Illness in Southeast Asia 2012 (WWARN)  
Acestor et al, PLoS ONE, 2012

# How important is scrub typhus?



**most frequent reported diseases/pathogens:**

- Dengue
- Rickettsial infections
- Leptospirosis
- Typhoid
- Melioidosis
- JEV

*Mapping the Aetiology of Non-Malarial Febrile Illness in Southeast Asia 2012*

*Acestor et al, PLoS ONE, 2012*

# Typhus in SEA

Thailand  
Malaysia  
Laos  
Vietnam  
Cambodia  
Myanmar  
Bangladesh  
India  
... ?! etc.

*“Dengue-like illness”*

*“Typhus-like illness”*

- *burden 15-25% ST, 10-15% MT*
- *estimated 1 mio cases per year*
- *seroprevalence 30-50%*

***ST and MT (and SFG) combined = leading cause of treatable undifferentiated febrile illnesses***

***“Scrub typhus is probably one of the most underdiagnosed and underreported febrile illnesses requiring hospitalization in the region ”***



*Fringe habitats - rural*



*Markets - urban*



# Unresolved issues

- *Ecology and Epidemiology*
- ***Entomology***
- *Diagnosis and Diagnostics*
- *Scrub Typhus Pathogenesis / Immunity*
- *Developing a Vaccine against Orientia*



*Concerted efforts of  
multiple specialities ...*



# Entomology – open questions

- *Are Leptotrombidia the only species transmitting Orientia?*
- *What are the risk factors for getting bitten?*
- *What are the true hosts of Orientia?*
- *How do Orientia modulate the sex ratio of infected mites?*
- *What selective processes influence Orientia evolution?*
- *What do multiple strain infections (mites/humans) tell us?*
- *Mite bite site processes? What is the role of mite saliva?*

# Are *Leptotrombidia* the only species transmitting *Orientia*?

*The established vectors are all species of the genus Leptotrombidium*

*L. deliense, L. scutellare, L. imphalum, L. arenicola, L. umbricola*

*L. akamushi, L. deliense, L. pallidum ...*

*(Thailand, Malaysia, Japan, Korea, India)*

*Reports of non-Leptotrombidium mites assoc. w/ Orientia*

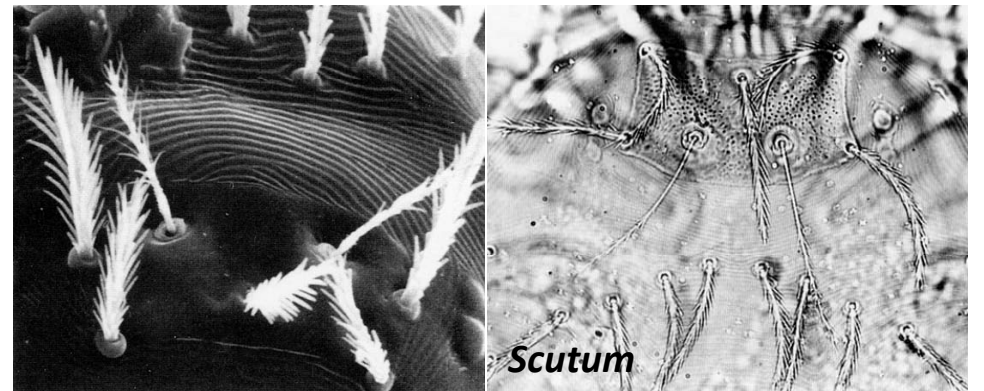
- *Blankaartia* sp. (Thailand)
- *Eutrombicula* sp. (Thailand)
- *Schoengastia* sp. (Thailand)
- *Gahrliepia* sp. (Japan, India)
- *Neotrombicula* sp. (Korea)
- *Cheladonta* sp. (Korea)

**Challenge**

- *Morpholgy* (scutum, hairs etc.)
- *Genetic characterisation* (COX genes etc.)
- *other ...*



<http://livingwithinsects.wordpress.com/>



# What are the true hosts of *Orientia*?

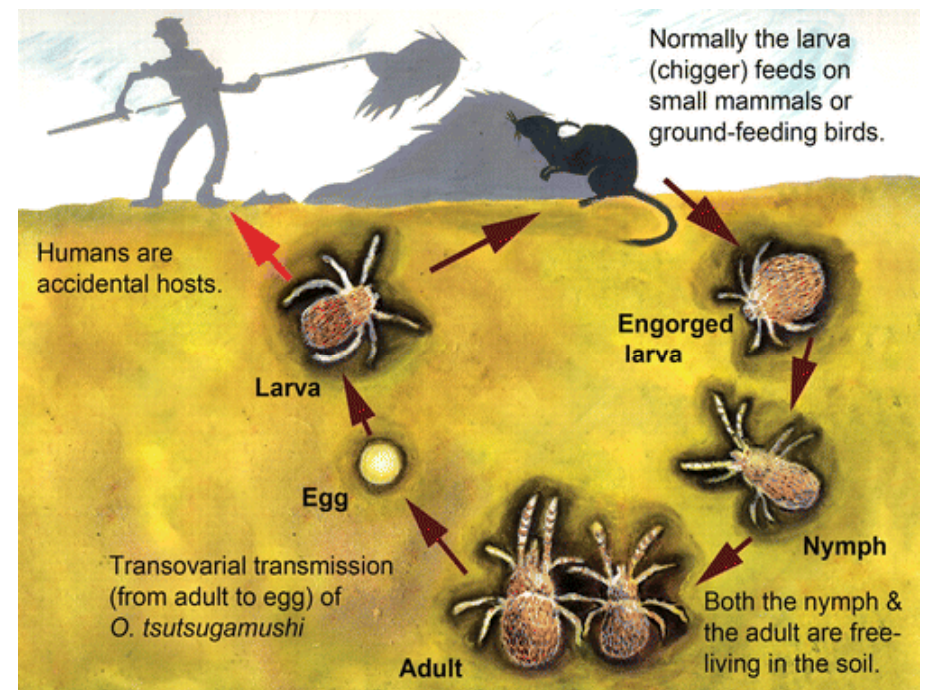
- Different mite species within genus have different infection rates
- Transmission of *Orientia*? Role of rodents? Humans?

- **Transmission rates**

- Rodent carriage of *Orientia* (range 14% to 33%)
- Rodent-mite transmission (range from <1% to max. 9%)
- Establishment of infection or vertical transmission rates post rodent feeding in nature remains unknown
- Co-feeding transmission rates?

- **Transmission experiments**

- Vertical transmission of mites infected via feeding on mice is low (<1 in 10,000)
- Basically the rodent seems to be a dead-end host as well?
- “*chicken and egg*” question ... ?!



Takahasi 1990  
Tanskul 1998  
Lerdthusnee 2003  
Jeong 2007

# What is the basis of “(re-)emergence”?

- *De novo* emergence vs. Re-emergence?
- Endemic disease – increased awareness?
- Other factors for maintenance of *Orientia* in nature? (other than vertical transmission that can explain antigenic diversity)
- Are there *Orientia* species other than *O. tsutsugamushi*?
- Actual geographic distribution?

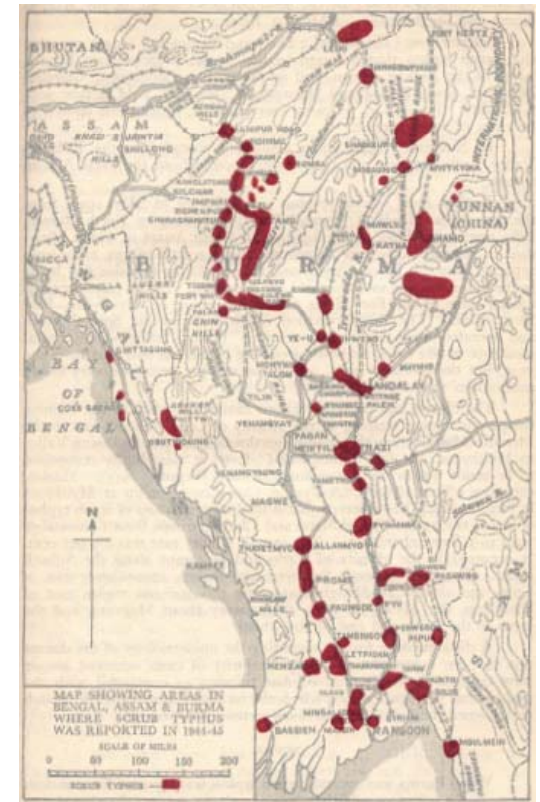


## Highly efficient vertical transmission

- Over 5 generations infection rates almost 100%
- Progeny is female in 95-100%, normal ratio is 2:1 (f:m)
- Variation per mite species  
*lower female rates in F1-F3 generations*

## Entity termed “mite islands”

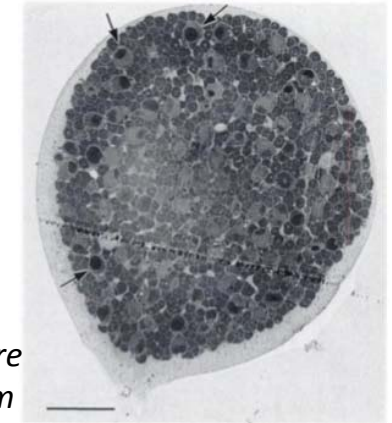
- Defined areas with high presence of infected chiggers
- Clonal characteristics of *Orientia*
- Chigger ‘re-attach’ to another rodent if incomplete feed, co-feeding issues



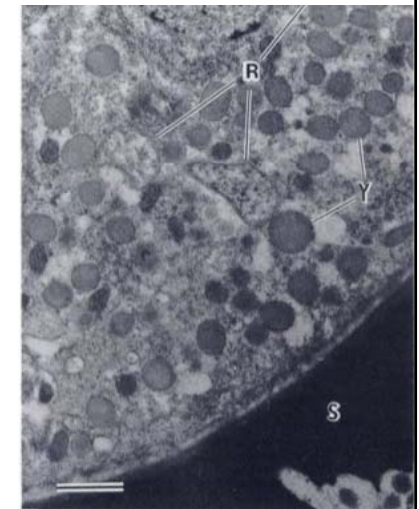
# How do *Orientia* modulate the sex ratio?

## Why are not all mites infected in nature?

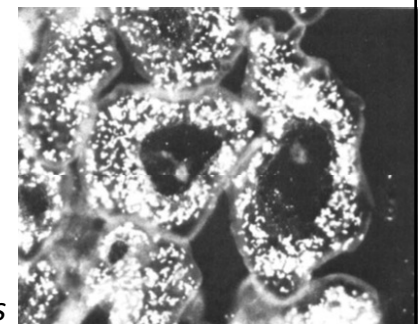
- Data shows around 1-3% are infected (Japan, SEA)
  - Population-survival advantage  
*(Fewer eggs, longer development time)*
  - Uninfected larvae acquire *Orientia* very rarely via rodent feeding  
*(infection rates <1%, vertical transmission almost absent)*
  - Proportion of transmission not vertically transmitted  
*(lose ability of female infected offspring)*



Spermatophore  
bar 5 um



Oocytes  
bar 1 um



Ovary  
Oocytes

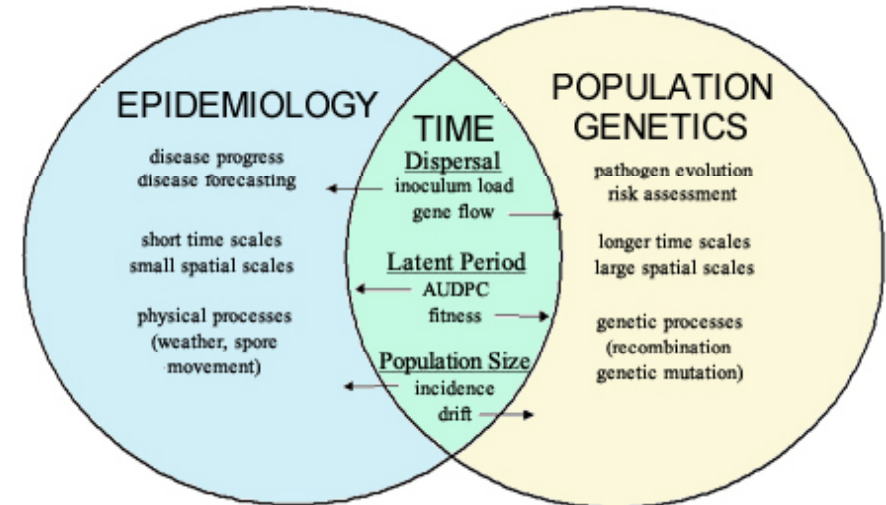
## Association with oocytes / ovaries and not male spermatophores

- Cytoplasmic inheritance factor (?)
- Parthenogenesis (?)

# What selective processes influence *Orientia* evolution?

## Some considerations

- Clonal organization of *Orientia*
- DNA polymorphism, genomic plasticity
- Small core genome, highly streamlined
- Gene transfer tools available
- Rodents less important than assumed
- *Role of co-feeding?*



- Which *Orientia* undergo transstadial and transovarial transmission?
- What are the characteristics of these endosymbionts?  
= *usually mutually beneficial interactions*
- *Infected mites - longer development times and decreased fecundity*  
= *more time for reproduction?*

## What tools to use?!

- *56kDa*: HV regions / immunogenic protein / selective pressure
- *MLST*: one mite – one strain – one ST (seq. type)
- *Whole genome sequencing*: population genetics

Cho 2007  
Nakayama 2010  
Sonthayanon 2010  
Duong 2010  
Takhampunya 2013

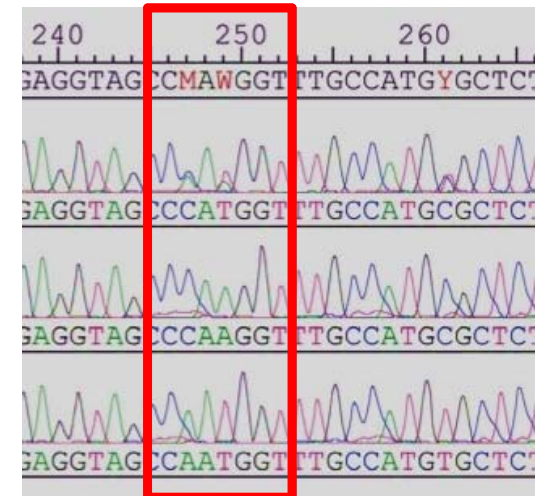
# What do multiple strain infections tell us?

## Evidence for multiple infections

- In humans: Thailand 25% with multiple infections
- In rodents: multiple genotypes described
- In mites: laboratory-reared and naturally-infected mites

## Heterogeneity / strain diversity

- Human pathogenic
- Human non-pathogenic = strains causing sub-clinical disease (*unknown*)
- Strains causing varying levels of disease severity
  - virulence factors (??) affected by background immunity



## Questions – Issues

- *Is recombination (and other mechanisms for increasing genetic diversity) occurring in rodents and mites ?*
- *If these ‘maintenance hosts’ are producing recombinant forms what are the underlying mechanisms?*



# Mite Bite Site

*Why is the mite bite and resulting eschar painless?*

Stylostome formation and different depths?

- Affects dissemination pathways?

Early host defenses?

- Neutrophils, monocytes influx

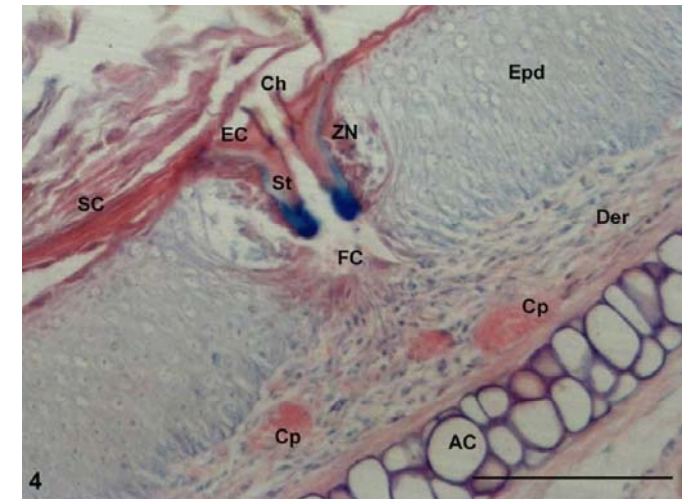
Immuno modulation – saliva/pathogen?

- IL-10, TGF $\beta$ , ag-presentation etc.

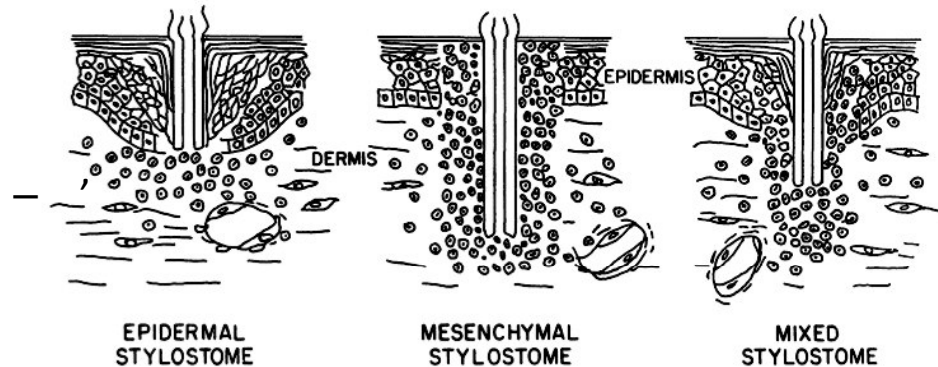
Local replication vs. early dissemination

- Haematogenic / lymphatic / 'monocyte-shuttles'
- Role of bacterial virulence?

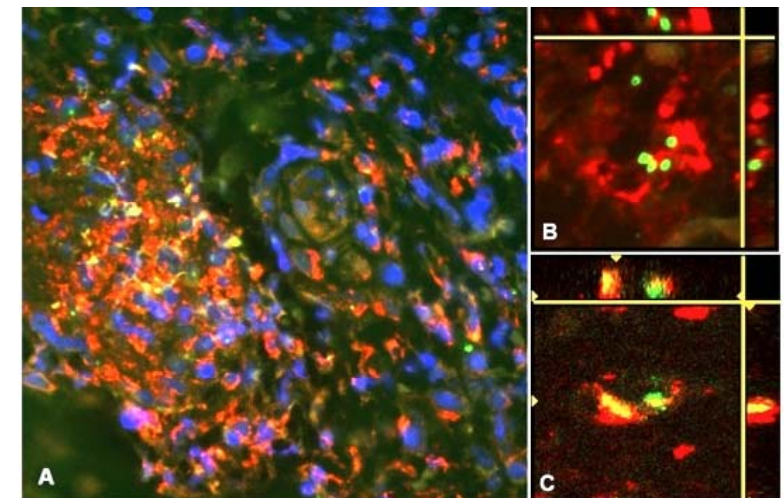
Inoculation bacterial load?



Shatrov, 2009



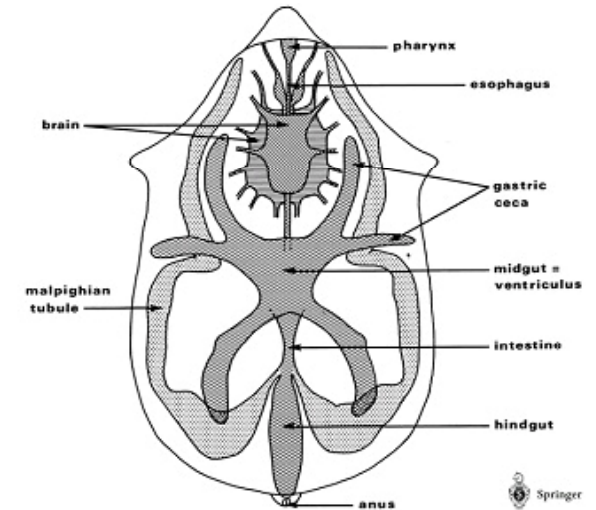
Hase, 1978



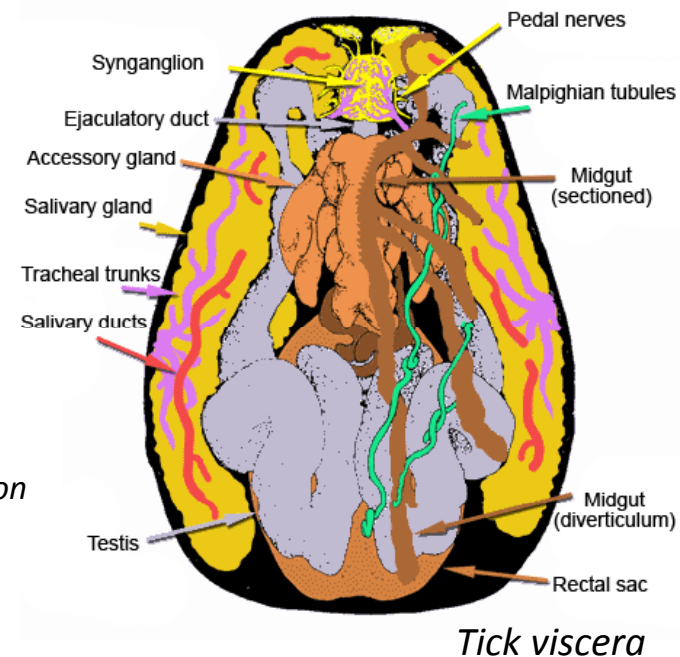
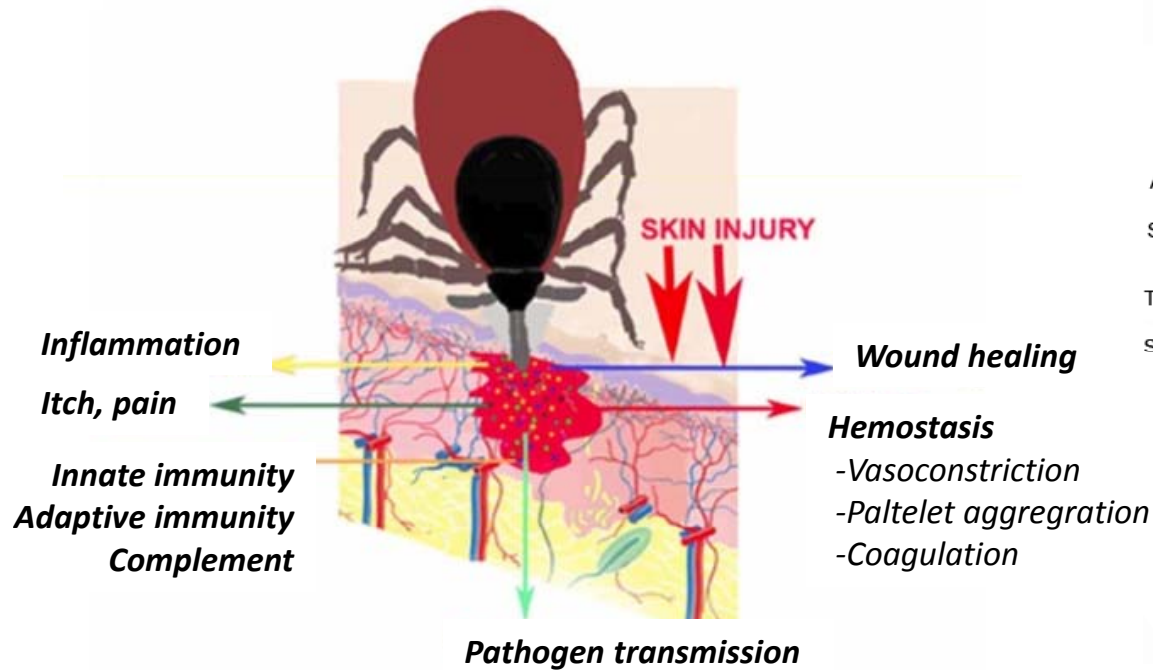
Paris, 2012

# What is the role of mite saliva?

- **Totally unexplored ... !**
- Parallels to tick saliva?
  - No blood feeding, lymphatic fluids
  - Inflammation, analgesia, immunomodulation, complement activation etc.
  - Hemostasis, vasoconstriction ...



Melhorn 2001



Tick viscera

# *Thank you !*

**MORU**

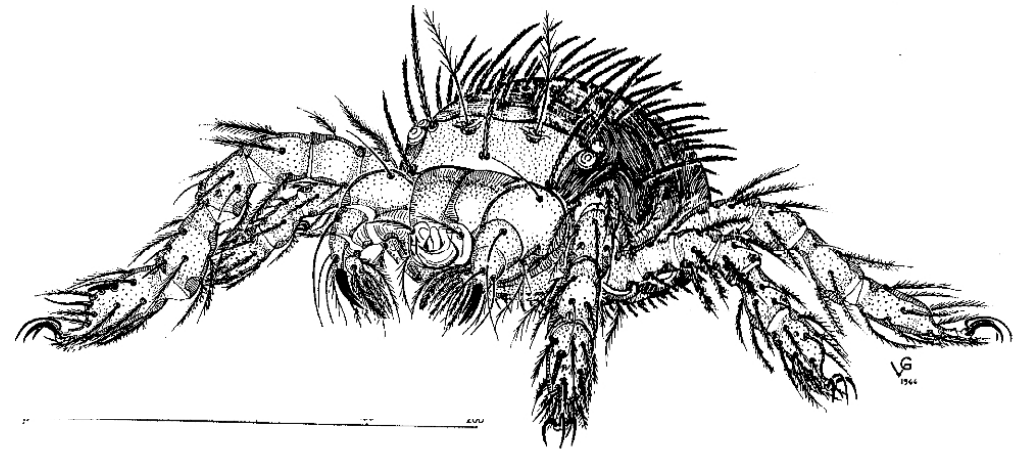
*Mahidol University*

*Oxford University*

**AFRIMS**

**NMRC**

*... and many more ...*



*Red mites and Typhus, R. J. Audy, 1965*

**Wellcome Trust**  
**NIH / NIAID**

