



האוניברסיטה העברית בירושלים THE HEBREW UNIVERSITY OF JERUSALEM



COREMI COST action Training School « Molecular and Morphological Identification of Mites Associated with the Red Poultry Mite"

Overall objectives

- Understanding the utility of morpho-molecular identification of mites
- Getting trained in mite morphological identification (theoretical and practical knowledge)
- Getting able to estimate what molecular analyses can bring to the study of mite species and communities

Framework of TS WG3: <u>Genetic structure</u> in a changing world

Leader: Lise Roy

- A methodological WG for an integrative approach
 - DNA markers
 - Molecular epidemiology of PRM (spread routes...), composition of associated communities (natural enemies, pathogenic microorganisms, ...)
 - Study of mechanisms underlying physiological features (immunological interactions with hen, resistances against acaricides...)
- Cross-disciplinary interests
 - direct links with all three other WGs



An ecosystem-level integrative approach

Why is PRM breaking up in some farms, whilst it keeps +/- stable in others?





Starting flock

Mid-flock



L. Roy, first conference of COREMI in Foggia, Italy, May 28th, 2015



Red mites: 2 different mite families, 2 different ecologies



- Eggs laid and molt performed on host
- as headlice



Eggs Mite feces

Poultry Red Mite lays eggs and molts in the environment (nest in wild avifauna, litter, dried droppings, cracks crevices in farms)

The Poultry Red Mite (*Dermanyssus gallinae*) = a micropredator



A pest mite typically dwelling in the same microhabitats as predatory mites ...

An example of morpho-molecular investigation on genus *Dermanyssus*

- Previous works highlighted important shortcomings in *Dermanyssus* species diagnostics due to highly variable morphological characters
 - ⇒ A feature common to many parasitic mites
 - ⇒ Use of molecular tools to contribute clarifying the issue: my PhD
- My PhD work as a case in point to illustrate general methods
 1) Main striking results with info on methods : phylog

2) Then, basic knowledge with punctual references to my work



An example of morpho-molecular investigation Inventory of *Dermanyssus* mites and species delineation phylog morphology

- Wild vs domestic birds:
- Different species of *Dermanyssus* found in wild birds
 - 5 species recorded in wild birds in France (9 different bird orders sampled): 416, D. apodis n. sp
 - <u>D. gallinae</u>, <u>D. carpathicus</u> <u>D. apodis</u>, <u>D. hirundinis</u>, <u>D. longipes</u>
- A single species in poultry farms:
 - <u>D. gallinae</u>

D. gallinae = the only generalist species and the only species found in farm contexts

Strongly supported monophyly of populations GOn and MAR along with several subtle morphological characters separate these populations from other species of the gallinae-group. These popuations constitute an entity which appears to be of specific statu and which we describe here under the name D. apode emale (holotype) (Fig. 6A-B): hosoma. Length of setae: anterior pair of hyp

long (47-65 µm), lateral pair 26 µm long (24-31 µm), posterior pair 26 µm (26-28 µm). Capitulum 96 µm (96-137 µm) long (from its basis to apex of palp coxae), 151 um(143-166 um) wide basally and 117 μm (104–130 μm) wide distally, (i.e. between lateral margins of palp coxae' apex).

field sampling

Setae all of palp genu lanceolate. Anti air wider than other gnathosomal setae. Idiosoma: 840 μm (735–1050 μm) long and 693 um) wide Domum: dorsal shield length 714 um (646







An example of morpho-molecular investigation Most spectrum within *Dermanyssus*



Whilst most of *Dermanyssus* species were recurrently recorded from a large diversity of birds, only *D. gallinae* was confirmed with a large host spectrum

Other species: either specific to a bird genus or to 1-2 bird families

phylog morphology field sampling bioassays

Roy et al. 2009b

An example of morpho-molecular investigation Host spectrum within *Dermanyssus*



An example of morpho-molecular investigation Spread routes consistent with trade routes

- High gene flow within European countries and Brazil
 - Same populations 1000 km apart within countries
 - Patterns consistent with known trade routes
- Populations from several different
 European countries (and Japan) related
 to each another

- excluding Norway (consistent with drastic isolation from an economic perspective)





Roy & Buronfosse 2011 (+ Oines & Brännström 2011, Chu et al. 2016)







Roy & Buronfosse 2011

An example of morpho-molecular investigation Spread routes: ways and vectors

In France, mite transfer between farms at least during spent hen removal



<u>No</u> transfer between wild bird and domestic fowl

At international scale, gene flow identified, but specific modes of mite transfer to be further explored



Persistance of PRM after empty period

> genpop morphology field sampling bioassays

Roy & Buronfosse 2011

An example of morpho-molecular investigation Genetic structure of *D. gallinae* s. l.

D. gallinae s. str. most likely results from recurrent hybridisation events between <u>incipient</u> species





Roy & Buronfosse 2011