

New molecular method to detect and quantify three microsporidia infecting bees, Vairimorpha (Nosema) apis\*, Vairimorpha (Nosema) ceranae\* and Vairimorpha (Nosema) bombi\* Aurélie Babin, Frank Schurr, Marie-Pierre Rivière, Marie-Pierre Chauzat, Eric Dubois

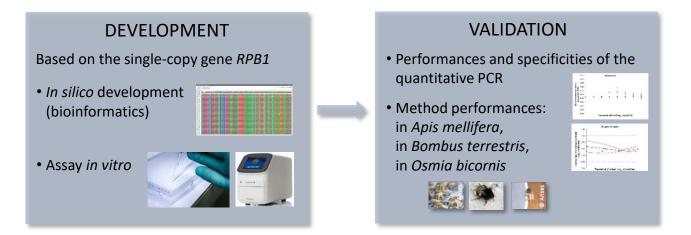
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\*Nosema apis, N. ceranae and N. bombi have been reassigned in 2020 from the genus Nosema to the genus Vairimorpha

- These three microsporidia are **intracellular parasites of bees**: *V. apis* and *V. ceranae* are honey bee parasites, *V. bombi* is a bumble bee parasite.
  - Current molecular methods do not allow the quantification of the three Vairimorpha species.

## OBJECTIVE

To develop and validate a **harmonised molecular method** (quantitative PCR) enabling the specific detection and quantification of these three parasites in honey bees, bumble bees and mason bees.



## PERSPECTIVES

- Study accurately *Vairimorpha* (co)infections, especially those devoid of clear clinical signs
- Study the microsporidia spillover between bee species
- Study the dynamics of coinfection/synergism with other pathogens and parasites
- Transfer to high-throughput methods for the analysis of big sample sets

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