



Marked inter-specific differences in the male olfactory system of honey bees (genus *Apis*)

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All honey bee species (genus *Apis*) display a striking mating behavior with the formation of male (drone) congregations, in which virgin queens mate with many drones. Bees' mating behavior relies on olfactory communication involving queen- but also drone pheromones. To explore the evolution of pheromonal sex communication in the *Apis* genus, we analyzed the neuroanatomical organization of the antennal lobe (primary olfactory center) in the drones of five species from the three main lineages (dwarf honey bees: *A. florea*, giant honey bees: *A. dorsata*, cavity-nesting honey bees: *A. mellifera*, *A. kochevnikovi* and *A. cerana*) and from three populations of *A. cerana* (Borneo, Thailand and Japan). In addition to differences in the overall number of functional units, the glomeruli, our data reveal marked differences in the number and position of macroglomeruli, enlarged functional units putatively dedicated to sex pheromone processing. Dwarf and giant honey bee species possess 2 macroglomeruli while cavity-nesting bees present 3 or 4 macroglomeruli, suggesting an increase in the complexity of sex communication during evolution in the genus *Apis*. The three *A. cerana* populations showed differing absolute numbers of glomeruli but the same three macroglomeruli. Overall, we identified 6 putatively homologous macroglomeruli in the genus *Apis*. One of these, which is dedicated to the detection of the major queen compound 9-ODA in *A. mellifera*, was conserved in all species. We will discuss the implications of these results for our understanding of sex communication in honey bees and propose a putative scenario of drone antennal lobe evolution in the *Apis* genus.