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The origin and evolution of queen and fertility signals in Corbiculate bees

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Abstract

Background: In social Hymenoptera (ants, bees and wasps), various chemical compounds present on the cuticle have been shown to act as fertility signals. In addition, specific queen-characteristic hydrocarbons have been implicated as sterility-inducing queen signals in ants, wasps and bumblebees. In Corbiculate bees, however, the chemical nature of queen-characteristic and fertility-linked compounds appears to be more diverse than in ants and wasps. Moreover, it remains unknown how queen signals evolved across this group and how they might have been co-opted from fertility signals in solitary ancestors.

Results: Here, we perform a phylogenetic analysis of fertility-linked compounds across 16 species of solitary and eusocial bee species, comprising both literature data as well as new primary data from a key solitary outgroup species, the oil-collecting bee *Centris analis*, and the highly eusocial stingless bee *Scaptotrigona depilis*. Our results demonstrate the presence of fertility-linked compounds belonging to 12 different chemical classes. In addition, we find that some classes of compounds (linear and branched alkanes, alkenes, esters and fatty acids) were already present as fertility-linked signals in the solitary ancestors of Corbiculate bees, while others appear to be specific to certain species.

Conclusion: Overall, our results suggest that queen signals in Corbiculate bees are likely derived from ancestral fertility-linked compounds present in solitary bees that lacked reproductive castes. These original fertility-linked cues or signals could have been produced either as a by-product of ovarian activation or could have served other communicative purposes, such as in mate recognition or the regulation of egg-laying.

Keywords: Fertility signals, Queen pheromones, Ancestral state reconstruction, Bees

Background

In order to be able to effectively organize themselves, insect societies require reliable information and communication systems [1]. In social insects, various communication mechanisms have evolved, of which chemical cues and signals are likely the most important [2]. Chemical substances present on the cuticle of individuals can serve various purposes, from the primary function of preventing desiccation [3] to having roles in nestmate and species recognition [4–6], courtship regulation [7–9] and signalling caste or reproductive status [10–19]. With respect to the latter, it has recently been shown in some hymenopteran species that

fertility-linked cuticular compounds help to regulate the reproductive division of labour between queens and workers in one of several distinct ways. For example, in the buff tail bumblebee, three ant species and the common wasp, it was shown that specific long-chain linear and methyl-branched alkanes act as sterility-inducing queen signals that stop the workers from reproducing [11, 12, 16, 19–23], reviewed in [24]. Moreover, in the common wasp, one of the main sterility-inducing pheromones was also shown to be used by the queen to mark her eggs, thereby enabling the workers to recognize and “police” eggs laid by other workers [25]. Finally, in two ants species, egg-laying workers were shown to produce specific alkenes that enable them to be recognized and policed via aggression by other workers [23, 26, 27].

The important role of cuticular hydrocarbons as queen pheromones in ants, wasps and the bumblebee contrasts

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with what is known from the honeybee, where most existing studies have indicated that mandibular gland compounds, including the keto acid 9-oxo-decenoic acid (9-ODA), act as the principal sterility-inducing queen signals [24]. Nevertheless, even there, it is likely that some cuticular fertility-linked compounds are also active as queen pheromones, since queens from which the mandibular gland was removed still inhibited worker reproduction [28, 29] and that extracts of the tergal glands, which produce specific compounds on the dorsal part of queen's cuticle [30, 31], also partially inhibited worker ovary development. In addition, in the stingless bee *Friesella schrottkyi*, it has recently been shown that specific linear and methyl branched alkanes were characteristic for the queen and that non-polar cuticular queen extracts inhibited worker reproduction [32].

In order to gain more insight into the origin and evolution of social insect queen pheromones, several studies have started to apply formal phylogenetic methods, and these show that there is remarkably high evolutionary conservation of queen signals [12, 16, 33]. For example, an ancestral state reconstruction has shown that structurally related saturated hydrocarbons were the most common chemical class of fertility-linked cues across more than 60 species of ants, bees, and wasps [12]. In addition, identical or structurally related saturated hydrocarbons were found to be bio-active across different *Lasius* ants [16] and even across several independently evolved social insect lineages, such as ants, wasps and bumblebees [12]. These results were interpreted as implying that queen pheromones likely evolved from pre-existing fertility signals in solitary ancestors, in which they may have had a different function [12, 24]. Furthermore, it has been suggested that the presence of honest signals of fertility in primitively eusocial species could lead to conditional helping strategies, whereby the most fecund individuals receive most help, thereby facilitate the transition to advanced eusociality with a pronounced reproductive division of labour [34]. Yet despite the potentially huge importance of fertility signals in the evolution and maintenance of eusociality, many questions remain. For example, it remains unknown why in bees the chemical nature of queen-characteristic and fertility-linked compounds is much more diverse than in ants and wasps [12], how queen signals evolved across this group and how they might have been co-opted from fertility signals in solitary ancestors. In fact, in the current most extensive phylogenetic analysis of fertility signals across over 60 social insect species, bees were the least represented, with only eight species. Furthermore, the published analysis [12] did not comprise any solitary outgroups, thereby limiting the power to infer how queen pheromones could have evolved from ancestral fertility-linked signals in solitary ancestors.

In the present study, we test the hypothesis that queen signals evolved from fertility-linked cues present in solitary ancestors [24] and perform a phylogenetic analysis of such compounds across 16 species of solitary and eusocial bee species. We use data from literature but also include new primary data from a key solitary outgroup species, the oil-collecting bee *Centris analis*, which is the sister taxon of all Corbiculate bees [35, 36], as well as data from the highly eusocial stingless bee *Scaptotrigona depilis*. Including *Centris* as an outgroup in our analysis allowed us to study character states present just before the transition from a solitary to a social lifestyle. In this way, we are able to obtain important insight with respect to what ancestral signals queen pheromones could have evolved from, and provide important clues as to what could be bioactive queen pheromones across different groups of bees.

Methods

Data collection

Chemical data on the identity of fertility signals across different species of solitary and social bees were collected from the literature through a systematic review of published studies, in which we compiled chemical data from fourteen different species (Additional file 1: Table S3), as well as through inclusion of new primary data of two more species (see below). Chemical compounds were classified as fertility linked whenever they were overproduced by queens in comparison to workers in eusocial species, or mature females versus non-egg laying virgin females in solitary species. These differences in chemical profiles could be either quantitative or qualitative. In order to standardize our data as much as possible, we mainly focused on studies that analysed apolar whole-body extracts, which recovers primarily cuticular compounds, but also smaller amounts of gland-derived compounds, such as 9-ODA in the honeybee [37, 38]. In addition, we obtained new data on the identity of fertility-linked signals in a key solitary outgroup, the solitary Centridini bee *Centris analis*, which is the closest extant relative of the Corbiculate bees [35, 36], as well as data on the queen signals produced by the highly eusocial stingless bee *Scaptotrigona depilis*. The oil collecting bee *Centris analis* was sampled using trap-nests for solitary bees [39]. Egg laying females ($n = 5$) were sampled when they visited their nest cavities whereas virgin females ($n = 8$) were collected on the day they emerged from trap-nests that were placed in the laboratory. Egg-laying queens ($n = 5$) and workers ($n = 15$) of the stingless bee *Scaptotrigona depilis* were sampled from the experimental meliponary from the University of São Paulo. Cuticular chemical data for the two bee species were obtained through gas chromatography and mass spectrometry analysis. Detailed sampling methods and details on the chemical analyses are provided in supplementary material.

Ancestral state reconstruction

Our total dataset comprised data on fertility-linked cuticular compounds across sixteen bee species, of which five were solitary and eleven eusocial (Additional file 1: Table S3). Compounds that were characteristic for either mature queens in social species or mature egg-laying females in solitary ones were grouped into different classes that correspond to known or presumed biosynthetic pathways (linear alkanes, branched alkanes, alkenes, alkadienes, aldehydes, alcohols, fatty acids, keto acids, esters, terpenes, terpene alcohols or lactones) [10, 40–44]. Subsequently, the presence or absence of at least one fertility-linked or queen characteristic compound belonging to one of each of these classes were coded as binary characters and ancestral states were reconstructed with Mesquite 3.01 [45] using maximum likelihood, based on the Markov k -state 1 parameter model, assuming equal rates of changes for both gains and losses. Phylogenetic relationships among species were based on the molecular phylogenies given in refs. [36, 46–48]. As branch lengths were not available in all cases, branch lengths were set equal to 1, which corresponded to assuming a punctuational mode of evolutionary change [49]. The likelihood threshold of 50% was used to consider that a given character state was likely present.

The advantage of our analysis being performed at the level of broad biosynthetic series is that in this way we were able to detect evolutionary conservation of fertility-linked compounds even if there would be small changes in the bioactivity of particular compounds across the study species. For example, overexpression of an elongase in queens of one particular lineage could result in an increase in chain length of fertility-linked linear or methyl-branched hydrocarbons [10, 50, 51] and would likely result in a change in the bioactivity of particular individuals compounds, but the use of this particular class of compounds as queen pheromones would still qualify as evolutionarily conserved. We should note, however, that the use of compounds belonging to a particular class as queen or fertility signals does not preclude other compounds of that same class also being used for other biological functions.

Ethical note

Data collection with live animals during this experiment was carried out at the University of São Paulo within the framework of a research project approved by the federal government (CNPq-Brazil 402661/2012-5). According to the country's law, research with bees is exempted from ethics committee approval when performed in research institutions.

Results

Our results show that, in comparison to ants and wasps [12], there is a strikingly high diversity of fertility-linked

chemical compounds in both solitary and eusocial bees. In particular, specific linear alkanes, alkenes, esters and fatty acids were already present with high likelihood (>50 %) as fertility-linked compounds in the ancestor of all solitary and social bees included in our analysis (Fig. 1, Table 1), whereas branched alkanes first appeared as fertility-linked compounds in the common ancestors of the Megachilidae and Apidae (Fig. 1, Table 1). In general, bees present odd long chain linear alkanes and alkenes ranging from heneicosane to hentriacontane, but other chain lengths including even chain length substances were also present in lower frequencies. Furthermore, specific terpenes also feature with high likelihood as fertility-linked signals in the common ancestor of *Centris* and the Corbiculate bees (Fig. 1, Table 1). Whereas both linear alkanes and alkenes are relatively conserved as fertility signals across all species analysed, fatty acids seem to have been lost in the common ancestor of stingless bees and bumblebees, and esters appear to have been gained and lost as fertility signals several times (Fig. 1). The use of keto acids, such as 9-ODA, as well as aldehydes, alcohols, alkadienes, terpene alcohols and lactones as queen or fertility signals appears to be restricted to specific lineages (Additional file 1: Figure S1).

Discussion

Overall, our results provide the first direct demonstration that many of the queen signals of Corbiculate bees are likely derived from fertility cues that were already present in solitary ancestors that lacked reproductive castes. In particular, our ancestral state reconstruction demonstrates that queen signals belonging to several distinct chemical classes, including linear alkanes, alkenes, ester and fatty acids, are likely derived from ancient fertility signals present in the common ancestor of all extant bee species, and originated at least ca. 120 Mya [36]. Similarly, we find that branched alkane and terpene queens signals are probably also derived from ancestral fertility signals, and started to be used before the origin of eusociality in Corbiculate bees, ca. 65 Mya [35, 52, 53], even though they had a more recent origin. By contrast, keto acids, such as 9-ODA, as well as aldehydes, alcohols, alkadienes, terpene alcohols and lactones are found to be used as queen or fertility signals only in very specific lineages. Overall, the diversity of fertility-linked compounds in both social and solitary bees was much higher than in other hymenopteran groups, including ants or wasps [12].

In general, our phylogenetic analysis adds credence to the idea that also in bees, compounds present on the cuticle, including both saturated and unsaturated hydrocarbons as well as other classes of compounds,

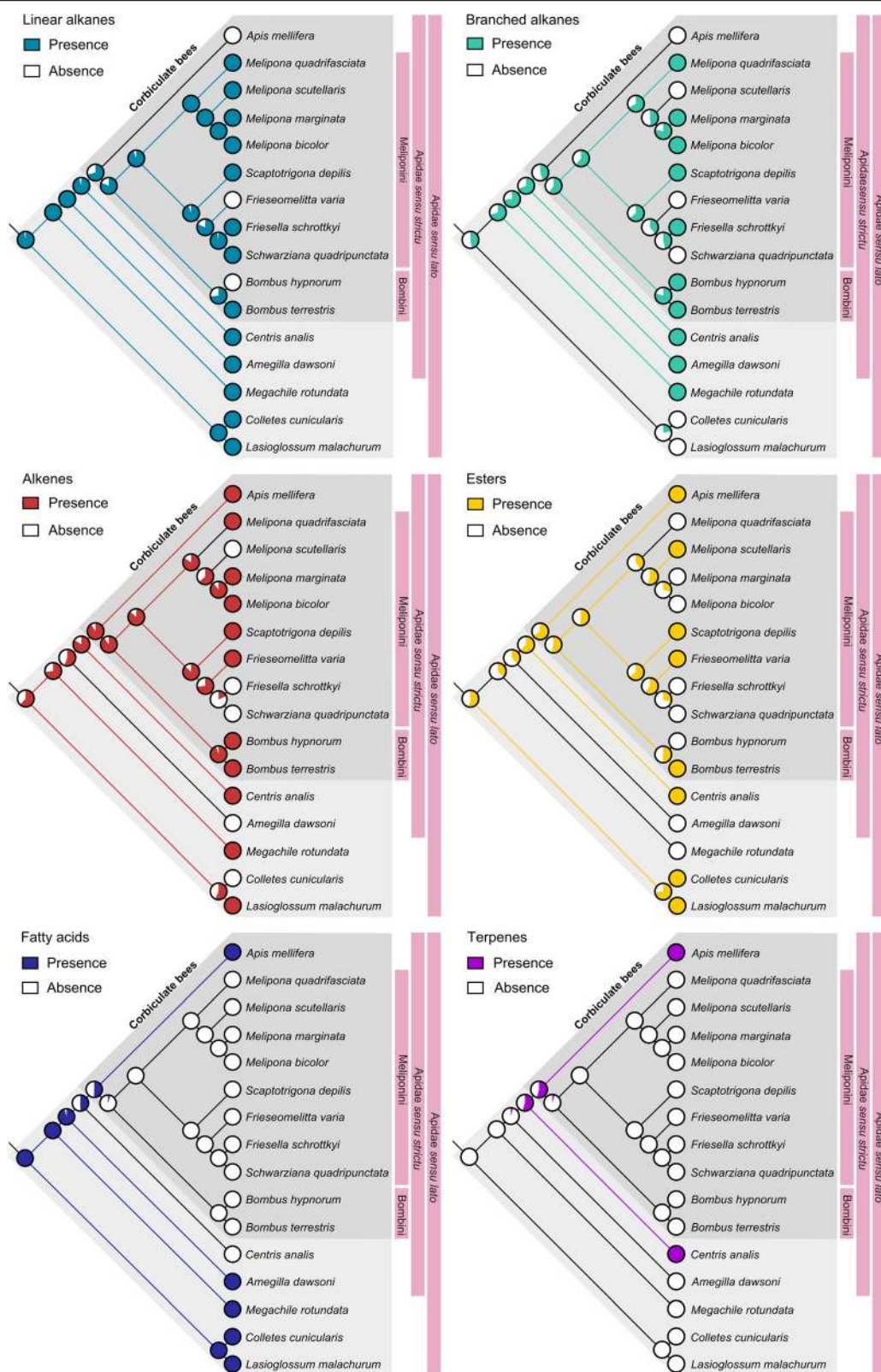


Fig. 1 Maximum likelihood ancestral state reconstruction of six major classes of fertility signals in bees. Pies show the likelihood in percentage of a given character being present in the node and coloured branches represent branches for which the likelihood was greater than 50 %

Table 1 Maximum likelihood, in percentage, of a given fertility signal being present as ancestral state of different bee clades.

Compound Class	Apidae <i>sensu lato</i>	Megachile + Apidae <i>sensu strictu</i>	Apidae <i>sensu strictu</i>	Centris + Corbiculate bees	Corbiculate bees	Meliponini	Bombini
Linear alkanes	96.9 %	99.0 %	98.8%	95.2 %	70.8 %	94.2 %	72.8 %
Branched alkanes	47.5 %	69.2 %	73.4 %	68.4 %	44.3 %	60.3 %	79.3 %
Alkenes	61.8 %	72.7 %	55.5 %	83.7 %	90.4 %	86.2 %	94.7 %
Esters	52.4 %	36.5 %	38.3 %	62.2 %	65.0 %	51.8 %	51.0 %
Fatty acids	99.0 %	99.3 %	95.7 %	49.7 %	50.3 %	0.4 %	0.4 %
Alcohols	1.4 %	1.4 %	9.3 %	0.8 %	0.1 %	0.0 %	0.0 %
Alkadienes	0.8 %	0.0 %	0.0 %	0.0 %	0.0 %	0.0 %	0.1 %
Aldehydes	0.8 %	0.2 %	0.8 %	9.3 %	1.4 %	0.2 %	8.7 %
Keto acids	0.0 %	0.0 %	0.0 %	0.0 %	3.7 %	0.0 %	0.0 %
Terpenes	1.1 %	0.0 %	4.6 %	54.1 %	54.1 %	0.0 %	0.0 %
Terpene alcohols	0.0 %	0.0 %	0.1 %	9.3 %	1.4 %	0.0 %	8.7 %
Lactones	3.7 %	0.0 %	0.0 %	0.0 %	0.0 %	0.0 %	0.0 %

In our analysis of 16 species, the Bombini (bumblebees) include the genus *Bombus*, Meliponini (stingless bees) include the genera *Melipona*, *Scaptotrigona*, *Friesella* and *Schwarziana*, Corbiculate bees correspond to all Bombini, Meliponini and *Apis mellifera* honeybees, Centris + Corbiculate bees to *Centris analis* and all Corbiculate bees, Apidae *sensu strictu* to *Amegilla dawsoni*, *Centris analis* and all Corbiculate bees, Megachile + Apidae *sensu strictu* to *Megachile rotundata* plus all Apidae s. s. and Apidae *sensu lato* to all species included in our analysis. Likelihoods higher than 50 % are shown in bold

such as ester and fatty acids, may act as conserved and ancient queen pheromones [12, 24]. In fact, direct evidence for cuticular compounds being used as sterility-inducing queen signals are now available for bumblebees, and strong suggestive evidence for their use has been provided also in honeybees and stingless bees. In particular, in the buff-tailed bumblebee *Bombus terrestris*, it has been shown that the linear alkane pentacosane inhibits worker ovary development [12, 21] but that some queen-characteristic esters did not suppress worker reproduction [12], even though they were still suggested to have a role in signalling fertility [20]. Similarly, in the stingless bees *Friesella schrottkyi*, queen cuticular extracts have been shown to inhibit worker ovary activation, and several hydrocarbons, including pentacosane, elicited an electroantennographic response, which suggests that cuticular compounds may act as queen pheromones in this species as well [32]. Finally, even in the well-studied honeybee, it is likely that some cuticular compounds complement the well-characterised mandibular gland compounds [54–56] in producing bio-active queen pheromone, since queens from which the mandibular gland was removed still inhibited worker reproduction [28, 29] and that extracts of the tergal glands, which produce specific compounds on the dorsal part of queen's cuticle [30, 31], also partially inhibited worker ovary development. Specific bioactive cuticular compounds have as yet not been identified, but queen-specific N-15 long-chain alkenes [31] or specific esters or fatty acids [30], such as hexadecanoic acid, (Z)-9-octadecenoic acid, hexadecanoic and octadecenoic acid methyl ester or decyl decanoate [57, 58], are amongst the possibilities.

If it is correct that some groups of social insect queen pheromones were co-opted from fertility signals that were already present in solitary ancestors, the question arises what function these fertility-linked compounds had in the solitary species. Previously, three main possibilities have been suggested, namely that they originally served as sex pheromones, that they were a by-product of ovarian activation, or that they were involved in regulating egg-laying, reviewed in [24]. Although these hypotheses are not mutually exclusive, for bees, especially this first hypothesis appears to be well supported, given that in *Apis* honeybees, the major queen pheromone component 9-ODA [54, 56, 59, 60] also doubles up as a male attractant sex pheromone [61–63], and that queen tergal gland secretion has been shown to enhance the effect of 9-ODA both in terms of inhibiting worker ovary activation [64] as well as in enhancing the queen's attractiveness to drones [65]. Moreover, in the eusocial Halictine bee *Lasioglossum malachurum*, a synthetic blend of the fertility-linked saturated lactones applied onto dummy bees was able to induce male inspection and pouncing behaviour (though not actual copulation behaviour, which appeared to be regulated by isopentenyl esters of unsaturated fatty acids) [66]. In addition, in the solitary bee *Colletes cunicularius*, several fertility-linked long-chain alkenes have also been shown to act as contact sex pheromones [67, 68] and in *Megachile rotundata*, alkene fraction extracts from virgin females, containing also the compounds linked to fertility, were shown to attract males in bioassays [69]. In all cases mentioned above, sex pheromones were characterised or presumed to be a blend of compounds functioning synergistically to attract males and elicit mating behaviour,

and could contain both fertility-linked compounds as well as compounds which were produced mainly by virgin females. Reduced production of non-fertility linked sex pheromones or increased production of compounds that inhibit male copulatory behaviour could explain the loss of the attractiveness of females after mating [70, 71], although change in context [72] or transfer of anti-aphrodisiac pheromones by the males themselves during mating has also been shown to be important in some cases (e.g. the transfer of anti-aphrodisiac s fatty acid, which are transferred as part of the male mating plug in bumblebees, [73], or the direct transfer of anti-aphrodisiac male cuticular compounds during mating in *Lasioglossum zephyrum*, *L. malachurum* and *Osmia rufa* [70, 74]. Either way, it is clear that a subset of fertility-linked sex pheromones could readily be co-opted as queen pheromones after the origin of eusociality [24]. Possibly, the fact that queen and fertility signals double up as sex pheromones in several bee species may also explain the relatively large diversity found in fertility-linked compounds, as sex pheromones are generally subject to diversifying selection [75, 76].

In the future, several of the hypotheses supported by our current dataset may be further validated by looking at the degree of evolutionary conservation in the expression levels of the actual underlying enzymes involved in the biosynthesis of different fertility-linked compounds [10, 40, 43], as well as by investigating conservation at the receptor level [77, 78]. Indeed, much ongoing research is targeted towards elucidating the roles of different enzymes in synthesizing different cuticular compounds in social insects, including those of different desaturases, elongases and decarboxylases involved in hydrocarbon biosynthesis [50, 79, 80], as well as those of lipophorins, involved in hydrocarbon transport [81, 82]. Recently, the enzymes involved in the production of the honeybee QMP blend were also described [83, 84], and the olfactory receptor AmOR11 was identified as the putative receptor for 9-ODA [85]. Together with targeted bioassays to test the bioactivity of particular fertility signals, such genomic studies may in the future yield unprecedented insight into the evolution of a key signalling system in insects, and allow us to uncover how social signalling systems were built on the groundplan of solitary ancestors [86, 87].

Conclusions

In a recent study, it was suggested that queen pheromones in different groups of social insects had been co-opted from ancient fertility signals present in common solitary ancestors [12]. Based on the analysis of fertility-linked compounds in different species of social bees and their solitary ancestors, our analysis provides direct support for this hypothesis. In particular, our results show that compounds belonging to four different chemical

classes (linear alkanes, alkenes, esters and fatty acids) are used as queen signals across several species of social bees, and that these same types of compounds were also present as fertility signals in their solitary ancestors. Other classes of compounds, however, only emerged later on in the evolutionary history of bees, or are specific to particular lineages.

Additional file

Additional file 1: Detailed data collection methodology, additional figure S1 with ancestral state reconstruction of the remaining six classes of chemical compounds and datasets used for the phylogenetic analysis. (DOCX 2511 kb)

Competing interests

The authors declare that they have no competing interests.

Authors' contributions

RCO and TW conceived the original idea, analysed the data and wrote the manuscript. CAO, MMCN, AVN, DAA, MCC and FN helped to collect both biological and chemical data and revised the manuscript. All authors read and approved the final manuscript.

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References

- Hölldobler B, Wilson EO. The ants. Cambridge: Harvard University Press; 1990.
- Wilson EO. The insect societies. The insect societies. Belknap Press; 1971.
- Jackson LL, Baker GL. Cuticular lipids of insects. *Lipids*. 1970;5(2):239–46.
- Lenoir A, Fresneau D, Errard C, Hefetz A. Individuality and colonial identity in ants: the emergence of the social representation concept. In: Detrain C, Deneubourg J, Pasteels J, editors. *Information Processing in Social Insects*. Birkhäuser Basel; 1999. p. 219–37.
- van Zweden J, d'Ettore P. Nestmate recognition in social insects and the role of hydrocarbons. In: *Insect hydrocarbons Biology, Biochemistry, and Chemical Ecology*. Edited by Blomquist GJ: Cambridge University Press; 2010: 222–243.
- Lucas C, Pho DB, Jallon JM, Fresneau D. Role of cuticular hydrocarbons in the chemical recognition between ant species in the *Pachycondyla villosa* species complex. *J Insect Physiol*. 2005;51(10):1148–57.
- Peschke K. Cuticular Hydrocarbons Regulate Mate Recognition, Male Aggression, and Female Choice of the Rove Beetle, *Aleochara Curtula*. *J Chem Ecol*. 1987;13(10):1993–2008.
- Ming QL, Lewis SM. Mate Recognition and Sex Differences in Cuticular Hydrocarbons of the Diurnal Firefly *Elychnia corrusca* (Coleoptera: Lampyridae). *Ann Entomol Soc Am*. 2010;103(1):128–33.

9. Zhang B, Xue HJ, Song KQ, Liu J, Li WZ, Nie RE, et al. Male mate recognition via cuticular hydrocarbons facilitates sexual isolation between sympatric leaf beetle sister species. *J Insect Physiol.* 2014;70:15–21.
10. Howard RW, Blomquist GJ. Ecological, behavioral, and biochemical aspects of insect hydrocarbons. In: *Annu Rev Entomol.* 2005;50:371–93.
11. Holman L, Jorgensen CG, Nielsen J, d'Etorre P. Identification of an ant queen pheromone regulating worker sterility. *Proceedings of the Royal Society B-Biological Sciences.* 2010;277(1701):3793–800.
12. Van Oystaeyen A, Oliveira RC, Holman L, van Zweden JS, Romero C, Oi CA, et al. Conserved class of queen pheromones stops social insect workers from reproducing. *Science.* 2014;343(6168):287–90.
13. Monnin T. Chemical recognition of reproductive status in social insects. *Ann Zool Fenn.* 2006;43(5-6):515–30.
14. Liebig J. Hydrocarbon profiles indicate fertility and dominance status in ant, bee, and wasp colonies. *Insect Hydrocarbons: Biology Biochemistry and Chemical Ecology.* 2010;254–281.
15. Peeters C, J L. Fertility signaling as a general mechanism of regulating reproductive division of labor in ants. In: *Organization of insect societies: from genome to socio-complexity.* Edited by Gadau J, J. F. Cambridge, MA: Harvard University Press; 2009: 220-242.
16. Holman L, Lanfear R, d'Etorre P. The evolution of queen pheromones in the ant genus *Lasius*. *J Evol Biol.* 2013;26(7):1549–58.
17. Holman L. Queen pheromones: The chemical crown governing insect social life. *Commun Integr Biol.* 2010;3(6):558–60.
18. Liebig J, Peeters C, Oldham NJ, Marktädter C, Hölldobler B. Are variations in cuticular hydrocarbons of queens and workers a reliable signal of fertility in the ant *Harpegnathos saltator*? *Proc Natl Acad Sci USA.* 2000;97(8):4124–31.
19. D'Etorre P, Heinze J, Schulz C, Francke W, Ayasse M. Does she smell like a queen? Chemoreception of a cuticular hydrocarbon signal in the ant *Pachycondyla inversa*. *J Exp Biol.* 2004;207(7):1085–91.
20. Sramkova A, Schulz C, Twele R, Francke W, Ayasse M. Fertility signals in the bumblebee *Bombus terrestris* (Hymenoptera: Apidae). *Naturwissenschaften.* 2008;95(6):515–22.
21. Holman L. Bumblebee size polymorphism and worker response to queen pheromone. *PeerJ.* 2014;2, e604.
22. Endler A, Liebig J, Schmitt T, Parker J, Jones G, Schreier P, et al. Surface hydrocarbons of queen eggs regulate worker reproduction in a social insect. *Proc Natl Acad Sci USA.* 2004;101:2945–50.
23. Smith AA, Millar JG, Hanks LM, Suarez AV. Experimental evidence that workers recognize reproductives through cuticular hydrocarbons in the ant *Odontomachus brunneus*. *Behav Ecol Sociobiol.* 2012;66(9):1267–76.
24. Oi CA, van Zweden JS, Oliveira RC, Van Oystaeyen A, Nascimento FS, Wenseleers T. The origin and evolution of social insect queen pheromones: Novel hypotheses and outstanding problems. *BioEssays.* 2015.
25. Oi Cintia A, Van Oystaeyen A, Oliveira RC, Millar JG, Verstrepen KJ, van Zweden JS, et al. Dual Effect of Wasp Queen Pheromone in Regulating Insect Sociality. *Curr Biol.* 2015;25(12):1638–40.
26. Smith AA, Millar JG, Suarez AV. A social insect fertility signal is dependent on chemical context. *Biol Lett.* 2015;11.
27. Smith AA, Holldobler B, Liebig J. Cuticular Hydrocarbons Reliably Identify Cheaters and Allow Enforcement of Altruism in a Social Insect. *Curr Biol.* 2009;19(1):78–81.
28. Velthuis H. Queen substances from the abdomen of the honey bee queen. *Z vergl Physiol.* 1970;70(2):210–21.
29. Maisonnasse A, Alaux C, Beslay D, Crauser D, Gines C, Plettner E, et al. New insights into honey bee (*Apis mellifera*) pheromone communication. Is the queen mandibular pheromone alone in colony regulation? *Frontiers in Zoology.* 2010;7(1):18.
30. Mucignat-Caretta C, Bortolotti L, Costa C. Chemical Communication in the Honey Bee Society. 2014.
31. Smith R-K, Taylor Jr OR. Unsaturated extracted hydrocarbon caste differences between European queen and worker honey bees, *Apis mellifera* L. (Hymenoptera: Apidae). *J Kans Entomol Soc.* 1990;63(3):369–74.
32. Nunes TM, Mateus S, Favaris AP, Amaral MF, von Zuben LG, Clososki GC, et al. Queen signals in a stingless bee: suppression of worker ovary activation and spatial distribution of active compounds. *Scientific Reports.* 2014;4:7449. doi: 10.1038/srep07449.
33. van Zweden JS, Bonckaert W, Wenseleers T, d'Etorre P. Queen Signaling in Social Wasps. *Evolution.* 2014;68(4):976–86.
34. Holman L. Conditional helping and evolutionary transitions to eusociality and cooperative breeding. *Behav Ecol.* 2014;25(5):1173–82.
35. Cardinal S, Danforth BN. The antiquity and evolutionary history of social behavior in bees. *PLoS One.* 2011;6(6):1–9.
36. Cardinal S, Danforth BN. Bees diversified in the age of eudicots. *Proceedings of the Royal Society of London B: Biological Sciences.* 2013;280(1755):20122686.
37. Engels W, Rosenkranz P, Adler A, Taghizadeh T, Lübke G, Francke W. Mandibular gland volatiles and their ontogenetic patterns in queen honey bees, *Apis mellifera carnica*. *J Insect Physiol.* 1997;43(4):307–13.
38. Moritz RFA, Crewe RM. The volatile emission of honeybee queens (*Apis mellifera* L.). *Apidologie.* 1991;22(3):205–12.
39. de Jesus BMV, Garofalo CA. Nesting behaviour of *Centris (Heterocentris) analis* (Fabricius) in southeastern Brazil (Hymenoptera, Apidae, Centridini). *Apidologie.* 2000;31(4):503–15.
40. Blomquist GJ, Bagnères A-G. *Insect Hydrocarbons Biology, Biochemistry, and Chemical Ecology.* Cambridge University Press. 2010.
41. Plettner E, Slessor KN, Winston ML, Oliver JE. Caste-selective pheromone biosynthesis in honeybees. *Science.* 1996;271(5257):1851.
42. Plettner E, Slessor KN, Winston ML. Biosynthesis of mandibular acids in honey bees (*Apis mellifera*): De novo synthesis, route of fatty acid hydroxylation and caste selective β -oxidation. *Insect Biochem Mol Biol.* 1998;28(1):31–42.
43. Morgan ED. *Biosynthesis in insects.* Royal Society of Chemistry; 2010
44. Kanehisa M. The KEGG database. *Silico Simulation of Biological Processes.* 2002;247:91–103.
45. Maddison WP, Maddison DR. Mesquite: a modular system for evolutionary analysis. In. 3.01 edn; 2014.
46. Ramírez SR, Nieh JC, Quental TB, Roubik DW, Imperatriz-Fonseca VL, Pierce NE. A molecular phylogeny of the stingless bee genus *Melipona* (Hymenoptera: Apidae). *Mol Phylogenet Evol.* 2010;56(2):519–25.
47. Rasmussen C, Cameron SA. Global stingless bee phylogeny supports ancient divergence, vicariance, and long distance dispersal. *Biol J Linn Soc.* 2010;99(1):206–32.
48. Hedtko SM, Patiny S, Danforth BN. The bee tree of life: a supermatrix approach to apoid phylogeny and biogeography. *BMC Evol Biol.* 2013;13:138.
49. Pagel M. Detecting Correlated Evolution on Phylogenies: A General Method for the Comparative Analysis of Discrete Characters. *Proceedings of the Royal Society of London B: Biological Sciences.* 1994;255(1342):37–45.
50. Falcón T, Ferreira-Caliman MJ, Nunes FMF, Tanaka ÉD, do Nascimento FS, Bitondi MMG. Exoskeleton formation in *Apis mellifera*: Cuticular hydrocarbons profiles and expression of desaturase and elongase genes during pupal and adult development. *Insect Biochem Mol Biol.* 2014;50:68–81.
51. Hashimoto K, Yoshizawa AC, Okuda S, Kuma K, Goto S, Kanehisa M. The repertoire of desaturases and elongases reveals fatty acid variations in 56 eukaryotic genomes. *J Lipid Res.* 2008;49(1):183–91.
52. Michener CD, Grimaldi DA. The oldest fossil bee: Apoid history, evolutionary stasis, and antiquity of social behavior. *Proc Natl Acad Sci U S A.* 1988;85(17):6424–6.
53. Michener CD. *The bees of the world.* 2nd ed. Johns Hopkins University Press. 2007.
54. Tan K, Liu X, Dong S, Wang C, Oldroyd BP. Pheromones affecting ovary activation and ovariole loss in the Asian honey bee *Apis cerana*. *J Insect Physiol.* 2015;74:25–9.
55. Butler C, Fairey EM. The role of the queen in preventing oogenesis in worker honeybees. *J Apic Res.* 1963;2(1):14–8.
56. Hoover SR, Keeling C, Winston M, Slessor K. The effect of queen pheromones on worker honey bee ovary development. *Naturwissenschaften.* 2003;90(10):477–80.
57. Wossler TC, Crewe RM. Mass spectral identification of the tergal gland secretions of female castes of two African honey bee races (*Apis mellifera*). *J Apic Res.* 1999;38(3-4):137–48.
58. Espelie KE, Butz VM, Dietz A. Decyl decanoate: a major component of the tergite glands of honeybee queens (*Apis mellifera* L.). *J Apic Res.* 1990;29(1):15–9.
59. Slessor KN, Winston ML, Le Conte Y. Pheromone communication in the honeybee (*Apis mellifera* L.). *J Chem Ecol.* 2005;31(11):2731–45.
60. Le Conte Y, Hefetz A. Primer pheromones in social hymenoptera. *Annu Rev Entomol.* 2008;53:523–42.
61. Law JH, Regnier FE. Pheromones. *Annu Rev Biochem.* 1971;40(1):533–48.

62. Brockmann A, Dietz D, Spaethe J, Tautz J. Beyond 9-ODA: sex pheromone communication in the European honey bee *Apis mellifera* L. *J Chem Ecol*. 2006;32(3):657–67.
63. Nagaraja N, Brockmann A. Drones of the dwarf honey bee *Apis florea* are attracted to (2E)-9-oxodecenoic acid and (2E)-10-hydroxydecenoic acid. *J Chem Ecol*. 2009;35(6):653–5.
64. Wossler TC, Crewe RM. Honeybee queen tergal gland secretion affects ovarian development in caged workers. *Apidologie*. 1999;30(4):311–20.
65. Renner M, Vierling G. Die rolle des taschendrüspheromons beim hochzeitsflug der bienenkönigin. *Behav Ecol Sociobiol*. 1977;2(3):329–38.
66. Ayasse M, Engels W, Lubke G, Taghizadeh T, Francke W. Mating expenditures reduced via female sex pheromone modulation in the primitively eusocial halictine bee, *Lasioglossum (Evyllaesus) malachurum* (Hymenoptera : Halictidae). *Behav Ecol Sociobiol*. 1999;45(2):95–106.
67. Mant J, Brändli C, Vereecken NJ, Schulz CM, Francke W, Schiestl FP. Cuticular hydrocarbons as sex pheromone of the bee *Colletes cunicularius* and the key to its mimicry by the sexually deceptive orchid, *Ophrys exaltata*. *J Chem Ecol*. 2005;31(8):1765–87.
68. Vereecken NJ, Mant J. Population differentiation in female sex pheromone and male preferences in a solitary bee. *Behav Ecol Sociobiol*. 2007;61:811–21.
69. Paulmier I, Bagnères A-G, Afonso CM, Dusticier G, Rivière G, Clément J-L. Alkenes as a sexual pheromone in the alfalfa leaf-cutter bee *Megachile rotundata*. *J Chem Ecol*. 1999;25(3):471–90.
70. Ayasse M, Paxton RJ, Tengö J. Mating behavior and chemical communication in the order Hymenoptera. *Annu Rev Entomol*. 2001;46:31–78.
71. Thomas ML. Detection of female mating status using chemical signals and cues. *Biol Rev*. 2011;86(1):1–13.
72. Engels W. Pheromones and reproduction in Brazilian stingless bees. *Mem Inst Oswaldo Cruz*. 1987;82:35–45.
73. Baer B, Morgan ED, Schmid-Hempel P. A nonspecific fatty acid within the bumblebee mating plug prevents females from remating. *Proc Natl Acad Sci USA*. 2001;98(7):3926–8.
74. Kukuk P. Evidence for an antiaphrodisiac in the sweat bee *Lasioglossum (Dialictus) zephyrum*. *Science*. 1985;227(4687):656–7.
75. Lofstedt C. Moth pheromone genetics and evolution. *Philosophical Transactions of the Royal Society B: Biological Sciences*. 1993;340(1292):167–77.
76. Maan ME, Seehausen O. Ecology, sexual selection and speciation. *Ecol Lett*. 2011;14(6):591–602.
77. Ozaki M, Wada-Katsumata A. Perception and olfaction of cuticular compounds. *Insect hydrocarbons; biology, biochemistry, and chemical ecology* 2010:207–221.
78. Ozaki M, Hefetz A. Neural mechanisms and information processing in recognition systems. *Insects*. 2014;5(4):722–41.
79. Helmkampf M, Cash E, Gadau J. Evolution of the insect desaturase gene family with an emphasis on social Hymenoptera. *Mol Biol Evol*. 2014;32(2):456–71.
80. Tsutsui ND. Dissecting ant recognition systems in the age of genomics. *Biol Lett*. 2013;9(6):20130416.
81. Fan Y, Schal C, Vargo EL, Bagnères A-G. Characterization of termite lipophorin and its involvement in hydrocarbon transport. *J Insect Physiol*. 2004;50(7):609–20.
82. Blomquist GJ, Nelson DR, De Renobales M. Chemistry, biochemistry, and physiology of insect cuticular lipids. *Arch Insect Biochem Physiol*. 1987;6(4):227–65.
83. Malka O, Karunker I, Yeheskel A, Morin S, Hefetz A. The gene road to royalty—differential expression of hydroxylating genes in the mandibular glands of the honeybee. *FEBS journal*. 2009;276(19):5481–90.
84. Malka O, Niño EL, Grozinger CM, Hefetz A. Genomic analysis of the interactions between social environment and social communication systems in honey bees (*Apis mellifera*). *Insect Biochem Mol Biol*. 2014;47:36–45.
85. Wanner KW, Nichols AS, Walden KK, Brockmann A, Luetje CW, Robertson HM. A honey bee odorant receptor for the queen substance 9-oxo-2-decenoic acid. *Proc Natl Acad Sci USA*. 2007;104(36):14383–8.
86. Page R, Linksvayer TA, Amdam G. Social life from solitary regulatory networks: a paradigm for insect sociality. *Organization of insect societies: from genomes to socio-complexity*. Cambridge: Harvard University Press; 2009. p. 357–76.
87. Johnson BR, Linksvayer TA. Deconstructing the superorganism: social physiology, groundplans, and sociogenomics. *The Quarterly Review of Biology*. 2010;85(1):57–79.

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