

Thelytokous Parthenogenesis In Unmated Queen Honey Bees (*Apis mellifera capensis*): Central Fusion and High Recombination Rates

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ABSTRACT

The subspecies of honey bee indigenous to the Cape region of South Africa, *Apis mellifera capensis*, is unique because a high proportion of unmated workers can lay eggs that develop into females via thelytokous parthenogenesis involving central fusion of meiotic products. This ability allows pseudo-clonal lineages of workers to establish, which are presently widespread as reproductive parasites within the honey bee populations of South Africa. Successful long-term propagation of a parthenogen requires the maintenance of heterozygosity at the sex locus, which in honey bees must be heterozygous for the expression of female traits. Thus, in successful lineages of parasitic workers, recombination events are reduced by an order of magnitude relative to meiosis in queens of other honey bee subspecies. Here we show that in unmated *A. m. capensis* queens treated to induce oviposition, no such reduction in recombination occurs, indicating that thelytoky and reduced recombination are not controlled by the same gene. Our virgin queens were able to lay both arrhenotokous male-producing haploid eggs and thelytokous female-producing diploid eggs at the same time, with evidence that they have some voluntary control over which kind of egg was laid. If so, they are able to influence the kind of second division meiosis that occurs in their eggs *post partum*.

In the honey bee, *Apis mellifera*, unfertilised eggs normally develop into haploid males by arrhenotokous parthenogenesis. Unfertilized eggs are produced by queens for the production of males, and also by unmated queenless workers whose eggs also produce functional males (DZIERZON 1845). Very occasionally, however, a worker will lay an egg in which meiosis II is modified so that an unfertilised egg is able to restore diploidy and become female (MACKENSEN 1943; TUCKER 1958), in a form of parthenogenesis known as thelytoky. Thelytoky is ubiquitous in workers of the South African subspecies *A. m. capensis* (hereafter Cape) (ANDERSON 1963; ONIONS 1912), and is thought to be controlled by a single gene, *Th*, which a mapping study has suggested may be homologous to *Grainy Head* of *Drosophila melanogaster* (LATTORFF *et al.* 2007; LATTORFF *et al.* 2005). In Cape workers, two haploid pronuclei of second division meiosis fuse and produce a diploid zygote, which usually gives rise to a female that may be reared as a worker or a queen (JORDAN *et al.* 2008; MORITZ *et al.* 1996). Some Cape workers use this ability to produce female offspring and reproductively parasitize other colonies (ALLSOPP 1993; BAUDRY *et al.* 2004; DIETEMANN *et al.* 2006; JORDAN *et al.* 2008; NEUMANN *et al.* 2001).

During this form of automictic (meiotic) thelytokous parthenogenesis there is a normal reduction division, bivalent formation and formation of chiasmata during meiosis I (VERMA and RUTTNER 1983). If a locus is distant from the centromere there will be multiple recombination events between the locus and the centromere, and the two pairs of alleles will become randomly placed on the four chromatids. Thus thelytokous parthenogenesis involving recombination means that for any locus heterozygous in the mother, there is a 1/3 chance that the offspring will be homozygous, whichever way the pronuclei combine (Table 1, PEARCY *et al.* 2006).

This ratio arises because if we choose any one chromatid at random, two of the three remaining chromatids will carry the alternate allele.

If there is interference to recombination or if loci are positioned close to the centromere and cannot recombine, the way in which the chromatids fuse determines what happens to the zygosity of offspring. During thelytokous parthenogenesis the products of meiosis II can fuse in one of three ways (PEARCY *et al.* 2006; SUOMALAINEN *et al.* 1987). Let us assume that the four haploid pronuclei of meiosis II are aligned in a row as in: $A_1A_2B_1B_2$. A_1 and A_2 were derived from nucleus A of meiosis I and B_1 and B_2 were derived from nucleus B. Under terminal fusion, terminal pro-nuclei fuse (i.e. A_1 with A_2 or B_1 with B_2). Under central fusion A_2 fuses with B_1 , and in random fusion any of the pronuclei may fuse. Under terminal fusion without recombination, a locus heterozygous in the mother will become homozygous in the offspring. Under central fusion without recombination, a locus will remain heterozygous (Table 1). For completeness, other less likely scenarios for the fusion of gametes are given in Table 1.

When unmated Cape queens are stimulated to produce unfertilised eggs by exposure to carbon dioxide (MACKENSEN 1947), they too can produce diploid female offspring via thelytoky like queenless unmated workers (CREWE and ALLSOPP 1994). In contrast, when mated Cape queens lay unfertilised eggs they produce males via arrhenotoky (JORDAN *et al.* 2008). This indicates an extraordinary ability of Cape queens to manipulate the kind of parthenogenesis that occurs when they lay unfertilised eggs – thelytoky and arrhenotoky in unmated queens and arrhenotoky in

mated queens. It also suggests that mated Cape queens could potentially produce daughters both sexually and asexually (JORDAN *et al.* 2008).

In honey bees sex is determined by the combination of paternal and maternal alleles at a single locus, the Complimentary Sex Determiner (*csd*) locus (BEYE *et al.* 2003). If the individual is heterozygous at the *csd* it is female. If the individual is homozygous at the *csd* a diploid male develops, but these are removed by workers at the first larval instar and are therefore inviable (WOYKE 1963). If the individual is haploid and therefore hemizygous at the *csd*, it is male. The *csd* encodes an *SR*-type protein, which is enormously polymorphic (HASSELMANN and BEYE 2004) due to diversifying selection (BEYE *et al.* 2003).

Sex determination via a single complimentary sex locus has important consequences. In sexually producing populations we predict that selection will act to increase recombination rates at the *csd* because recombination increases the probability of heterozygosity at the *csd*. As expected, the region around *csd* shows a seven-fold increase in recombination rate relative to other parts of the genome (HASSELMANN and BEYE 2006), presumably as a mechanism for maintaining heterozygosity. But what is expected in thelytokous populations? In Table 1 we list the various kinds of gamete fusion that are possible, and the consequences of the different forms on the probability that a locus heterozygous in the mother will become homozygous in diploid offspring. Table 1 shows that, in the absence of recombination, central fusion is favoured over random fusion because heterozygosity is maintained. However under all of terminal, random and central fusion we expect a 1/3 reduction in heterozygosity at the sex locus if recombination occurs. Thus, in thelytokous

populations with central fusion we expect reduced levels of recombination to evolve, at least on linkage group 3, which contains the sex locus. Studies of recombination rates in Cape workers show that they are at least an order of magnitude lower than in arrhenotokous queen meiosis (BAUDRY *et al.* 2004; MORITZ and HABERL 1994), strongly suggesting that selection for reduced recombination has indeed occurred in thelytokous Cape workers.

Alternative means of parthenogenesis within the same species, and indeed the same individual, raise interesting questions concerning the mechanisms of gametogenesis in Cape queens. Gametogenesis in Cape queens is as yet undescribed, but is well understood for arrhenotokous populations. Queens in non-Cape populations store new eggs in their lateral oviducts (DADE 1977) with the maternal pronucleus arrested in metaphase I (SASAKI and OBARU 2002). Second division meiosis occurs only after oviposition (SASAKI and OBARU 2002) when the diploid products of meiosis I align perpendicularly to the egg axis and undergo the second meiotic division. A single central nucleus becomes the maternal pro-nucleus, whereas the other three nuclei degenerate and become polar bodies (NACHTSHEIM 1913; PETRUNKEWITSCH 1901; YU and OMHOLT 1999). If the egg has been fertilized, one of the 6-10 sperm pronuclei present in the egg will fuse with the maternal pronucleus to produce a zygote and eventually a diploid female. If the egg has not been fertilised the maternal nucleus continues to divide mitotically and will produce a haploid male by arrhenotokous parthenogenesis.

A detailed cytological description of thelytokous parthenogenesis in Cape worker-laid eggs is also available (VERMA and RUTTNER 1983). In thelytokous parthenogenesis

by Cape workers the central (rather than the terminal or random) pronuclei fuse to produce the restored diploid nucleus, as if one of the central maternal pronuclei takes the place of a sperm pronucleus. A linkage study (BAUDRY *et al.* 2004) has confirmed the cytological evidence of central fusion.

Here we examine the sex, recombination rates, and the mode of gamete fusion in offspring of virgin queens of *A. m. capensis* that had been treated with carbon dioxide to induce oviposition (MACKENSEN 1947). Jordan *et al.* (2008) showed that thelytokous reproduction is rare or absent in mated Cape honey bee queens, whereas it is normal in queens of the ant *Catoglyphos cursor* (PEARCY *et al.* 2004; PEARCY *et al.* 2006). This investigation provides insights into the evolution of the widespread occurrence of thelytoky in the Cape worker, and demonstrates that thelytoky is possible in the queen caste.

MATERIALS AND METHODS

Thelytokous and arrhenotokous reproduction in unmated queens

In September 2006 in Stellenbosch South Africa we reared *A. m. capensis* queen pupae using standard methods (HARBO 1986; LAIDLAW and PAGE 1997). Mature queen pupae were allowed to eclose in an incubator at 35° C, and the virgins were then matured in the incubator in individual vials for 7 days, while being fed *ad libitum* on diluted honey. The queens were then anaesthetized for 10 min with carbon dioxide to induce oviposition (MACKENSEN 1947) and introduced into nucleus colonies (HARBO 1986; LAIDLAW and PAGE 1997) populated with *A. m. scutellata* workers and

brood. To prevent mating, we clipped the wings of the queens, retaining the clippings for later genotyping. To limit the amount of worker reproduction in the nucleus colonies, and to aid the establishment of the virgin Cape queens, we used *A. m. scutellata* workers instead of *A. m. capensis* workers in the nucleus colonies. We anaesthetised the queens at least once more 2 days after introduction, and a third time if eggs had not been seen. Until oviposition was observed, the queens were prevented from leaving their host colony by a grid of queen excluder material tacked over the entrance. Induction of oviposition in virgin queen honey bees from populations other than the Cape honey bee does not induce thelytokous parthenogenesis, but arrhenotokous parthenogenesis (MACKENSEN 1947).

As the first virgin-queen brood approached maturity, we collected brood from both worker and drone cells. To determine whether virgin Cape queens can produce both thelytokous and arrhenotokous progeny simultaneously, pupae were first sexed morphologically, and a sample of drone and worker progeny were then genotyped at microsatellite loci *Am* 059, *Am* 014, *Am* 107 and *Am* 061 (SOLIGNAC *et al.* 2003) to determine if they were sons of host workers, daughters of Cape workers foreign to the host colonies, or sons and daughters of the resident virgin queen. The queen genotype was obtained from tissue from the clipped wing of the queen. A progeny was rejected as being the offspring of the queen if it did not share at least one allele with the laying virgin queen at all four loci analysed.

Recombination rates during thelytokous reproduction

For this question we focussed genotyping effort on the offspring of queen 1. This queen produced large numbers of worker brood in worker cells, and no drone progeny. Pupae from worker cells or newly emerged callow workers ($n = 44$) were genotyped at 28 microsatellite loci on linkage groups 1 and 3 (which contains the *csd*). These loci were all heterozygous in the queen. Microsatellite loci and PCR primers were obtained from the microsatellite-based map 'Solignac_3' generated from 2008 microsatellite and other PCR-based markers segregating in the worker progeny of two hybrid queens (SOLIGNAC *et al.* 2007). This level of coverage provides accurate estimation of map distances between marker loci.

Under thelytokous parthenogenesis with central fusion, the expected recombination rate between a locus and the centromere is $1/3$ (see above and Table 1). Exceptions will occur when loci are situated less than 100 cM from the centromere, if distortions are caused by lethal allelic combinations at the sex locus, or by any other distorter of fair meiosis. The recombination fraction between a locus and its centromere, θ , can be estimated as the proportion of offspring that are homozygous in offspring (assuming the locus is heterozygous in the mother) (BAUDRY *et al.* 2004). Assuming no distortions to fair meiosis, the map distance, D , between a locus and the centromere can be calculated from $D = -\frac{2}{3} \ln(1 - 3\theta)$ (BAUDRY *et al.* 2004; RIZET and ENGELMANN 1949). This relationship assumes that the probability of a chiasmata forming is Poisson distributed, and corrects for the occurrence of double crossovers. Rizett and Engelmann's equation can also be used to calculate the map distance between any two pairs of loci, in which case θ is the proportion of individuals that are heterozygous at one locus and homozygous at the second (BAUDRY *et al.* 2004).

Similarly, the inverse $\theta = \frac{1}{3}(1 - e^{-\frac{3}{2}D})$ can be used to convert map distances from the ‘Solignac_3’ map to the expected recombination fraction between two loci or a locus and the centromere in thelytokously-produced progeny under the assumption of fair meiosis. We used these equations to determine if patterns of recombination observed in the progeny of our queen differed from expectations under a model of thelytokous parthenogenesis with central fusion, or if they were more compatible with alternative modes of gamete fusion given in Table 1. We also used them to compare recombination rates in thelytokous reproduction observed here with recombination rates reported in (BAUDRY *et al.* 2004).

DNA extraction and microsatellite genotyping

Tissue was obtained from the hind legs of worker and drone pupae and newly emerged callows, and from the clipped wings of the virgin queens. DNA was extracted by grinding tissue in 500 μ l of 5% Chelex solution followed by 10 min boiling (WALSH *et al.* 1991). Standard PCR conditions (ESTOUP *et al.* 1994) were used to amplify microsatellite loci (SOLIGNAC *et al.* 2003). PCR products (1.2 μ L) from each multiplex reaction were added to 10 μ L formamide and 100 nL LIZ DNA size standard (Applied Biosystems). Samples were run on a 3130xl Genetic Analyser (Applied Biosystems), with capillary length 36 cm and injection time of 15 s at 1200 V, for 41 minutes. Resultant data files were analysed using Genemapper software (Applied Biosystems) and genotypes for each individual constructed.

Results

Thelytokous and arrhenotokous reproduction in the same queens

Both drone and worker brood were observed in all four colonies (Table 2). Workers were active contributors to egg laying in most colonies, reducing the number of queen laid progeny we sampled. Nonetheless we were able to confirm thelytokous reproduction by queens in all four colonies. In three colonies queens laid both arrhenotokous and thelytokous offspring (Table 2). There is also evidence that queens preferentially laid eggs in the correct cell size, with thelytokous workers mostly reared in worker-sized cells and arrhenotokous drones in drone-sized cells. In all, 185 queen-laid individuals were retrieved from the correct cells and 10 from incorrect cells. This deviates significantly from random ($\chi^2_1 = 157.0, P < 0.001$). These samples were taken as soon as the first queen progeny began to emerge, and are therefore not thelytokous granddaughters of the virgin queens.

Mode of thelytokous reproduction in a virgin Cape queen

In the absence of centromeric interference, expected recombination fractions between all pairs of loci, θ_{exp} , calculated from the map distances from the ‘Solignac_3’ map using the Rizet and Engelmann (1949) correction are universally 0.33. The observed recombination fractions, θ_{obs} , between pairs of loci are given in Figure 1.

On linkage group 1, loci *Am* 103, 210 and 491 were expected to show reduced recombination rates because they lie within or close to the centromeric region (BAUDRY *et al.* 2004) (Figure 1). We confirm a reduced recombination rate between

loci *Am* 103 and *Am* 210, but the region between *Am* 210 and *Am* 491 showed a θ_{obs} of 0.35 (Figure 1), suggesting that the centromere is more than 100 cM distant from *Am* 491. On the other hand, the region between *Am* 076 and *Am* 103 showed a θ_{obs} of only 0.23 suggesting that the centromere of linkage group 1 may be slightly more telomeric than suggested by Baudry *et al.* Excluding loci *Am* 076, 103 and 210, the average number of recombinants workers per locus on linkage group 1, $\bar{\theta}_1$, was 0.33 (s.e. ± 0.014). This is not significantly different from the expected 0.33 on the assumption of automictic parthenogenesis with either central, random or terminal fusion of gametes ($P > 0.05$, one-sample *t*-tests with 10 degrees of freedom). However $\bar{\theta}_1$ deviated significantly ($P < 0.001$) from 0 homozygotes expected under apomixis and 100% homozygotes expected under gamete duplication.

On linkage group 3, loci *Am* 009, *Am* 317, *Am* 194 and K0333b are within 100 cM of the terminal centromere, and were expected to show reduced recombination rates (BAUDRY *et al.* 2004; Figure 1). As expected, these loci showed lower recombination rates than most other loci on this chromosome and no recombinants at all were seen at locus *Am* 009. Excluding these four loci, $\bar{\theta}_3 = 0.29 \pm 0.02$, which is marginally significantly different from 0.33 % ($t_9 = 1.9$, $P = 0.05$). This reduction in expected homozygosity is expected due to the effects of *csd*, which may have cause selection inviability of some homozygotes especially near locus K0338 (Figure 2). None the less, the proportion of homozygous individuals observed for non-centromeric loci on linkage group 3 differed significantly ($P < 0.001$) from that expected under both gamete duplication (100% homozygosity) and apomixis (no homozygotes) based on one sample *t*-tests with 9 degrees of freedom.

The non-centromeric loci provide strong evidence that automictic parthenogenesis is more likely than gamete duplication or apomixis. The centromeric loci, where there is a reduction in recombination, can be used to determine whether random, central, or terminal fusion of gametes during automixis is more likely. Under central fusion with recombination we expect an increase in the proportion of individuals that are homozygous away from the centromere towards the chromosomal arms. Under terminal fusion we expect the reverse polarity (BAUDRY *et al.* 2004). In Figure 2 we have plotted the proportion of individuals that are homozygous against the map distances obtained from the Solignac_3 genetic maps. Linkage group 1 shows two gradients of increasing homozygosity away from the metacentric centromere. Linkage group 3 shows a gradient of increasing homozygosity away from the terminal centromere. These patterns (especially that on linkage group 1) are consistent with central fusion and are inconsistent with terminal fusion, random fusion or gamete duplication.

Baudry *et al.* (2004) calculated the map distance between loci *Am* 062 and *Am* 031 in the sexually-produced progeny of an *A. m. capensis* queen and between *Am* 062 and *Am* 109 in the progeny of an arrhenotokous *A. m. mellifera* worker. This allows us to make a direct comparison of recombination rates in a normal *A. m. capensis* queen meiosis, an *A. m. mellifera* arrhenotokous worker meiosis, and a thelytokous *A. m. capensis* queen meiosis (Table 3). Our calculated map distances (Table 3) are larger than those estimated from other progenies, including that of normal queen meiosis in *A. m. mellifera*, suggesting that there is no reduction in recombination rates in the thelytokous meiosis of the *A. m. capensis* queen. Further more, we can directly compare the recombination frequency between loci *Am* 031 and *Am* 062 in

thelytokous workers from Figure 3 of BAUDRY *et al.* (2004), and compare this directly to the recombination rate between these same loci in the thelytokous progeny of a virgin queen (this study). In the BAUDRY *et al.* study, 3 of 108 individuals showed recombination between these two loci, whereas in our study 10 of 42 individuals were recombinant showing that there is a highly significant reduction in worker thelytokous parthenogenesis compared to that observed in the virgin queen ($\chi^2_1 = 16.9, P < 0.001$).

DISCUSSION

Warmelo (1912) remarked that ‘...it would seem contrary to all the laws of nature that the African worker bee produce her progeny in a wholly different manner from the queen which is essentially a worker bee with fully developed reproductive organs’. Our study has shown that Warmelo was only half right with respect to thelytokous parthenogenesis in Cape queens and workers. In both castes, it appears that diploidy is restored by central fusion rather than terminal fusion of meiotic products, or other possible mechanisms of gamete fusion listed in Table 1. However, the massively reduced rates of recombination observed in thelytokous parthenogenesis of the Cape worker (BAUDRY *et al.* 2004; MORITZ and HABERL 1994) are apparently absent when a virgin Cape queen reproduces thelytokously.

Reduced rates of recombination are essential for the maintenance of genetic diversity in a parthenogen propagating thelytokously with central fusion (BAUDRY *et al.* 2004; BELSHAW and QUICKE 2003). In the case of honey bees where there is a single sex-determining locus which must be heterozygous for the expression of the female sex,

maintenance of heterozygosity is essential, at least at the *csd*. Absence of reduced rates of recombination in the queen suggests that reduced recombination in the worker is under separate genotypic control to the control of thelytoky itself. Lattorff *et al.* (2007; 2005) showed that in the Cape worker, thelytoky is controlled by a single locus. This locus also influences two other traits related to worker reproduction pleiotropically: ovary activation and the production of a queen-like pheromonal bouquet (LATTORFF *et al.* 2007). However our results suggest that this locus may not be responsible for reduced rates of recombination, which is likely to be under separate genetic control. Clonal worker lineages (of which there are probably many JORDAN *et al.* 2008) that do not successfully evolve reduced rates of recombination will be at a strong selective disadvantage against lineages that can do so, and are likely to go extinct due to increasing homozygosity at the *csd*.

We have confirmed genetically the remarkable ability of unmated Cape queens to produce both thelytokous and arrhenotokous eggs during the same period (CREWE and ALLSOPP 1994). Our data suggest that virgin Cape queens have at least partial control over which kind of meiosis their eggs undergo. Where queens produced both male and female offspring, these were mostly (but not always) found in the correct cells. This suggests that Cape queens can to a large extent choose the ploidy of their eggs. An alternative explanation is that virgin queens lay thelytokous and arrhenotokous eggs at random in worker and drone cells, but that the workers selectively rear only those eggs that are laid in the appropriate cells. However, as workers readily rear larvae of any sex in both drone cells and worker cells without selection (CALDERONE and KUENEN 2001), it seems much more likely that virgin Capensis queens can

influence whether they lay diploid or haploid eggs rather than workers removing the larvae which are located in the wrong cell type.

How could the ability to lay arrhenotokous or thelytokous eggs be advantageous to Cape queens? The ability to lay thelytokous eggs allows queens to effectively clone themselves. It has been argued that such an ability should be at a selective advantage during reproductive swarming, as the queen need not share the genome of her gyne offspring with her mating partners (FOURNIER *et al.* 2005; JORDAN *et al.* 2008; PEARCY *et al.* 2004). When producing workers, Cape queens can produce haploid eggs and fertilize them with their stored sperm. As workers are mostly sterile, the queen pays little or no fitness cost by sharing her genome with her mating partners (PEARCY *et al.* 2004), and may increase her fitness by generating a genetically variable worker progeny (JONES *et al.* 2004; MATTILA and SEELEY 2007; OLDROYD and FEWELL 2007; SEELEY and TARPY 2007). The ability to produce males that will potentially mate with other queens is also advantageous. The optimal strategy, if it is biologically possible, is to do all these things.

Jordan *et al.* (2008) reported that when Cape colonies undergo reproductive swarming, queens occasionally lay eggs in queen cells that are parthenogenetic offspring of themselves, suggesting that indeed, mated queens may have the ability to produce clonal queen offspring during reproductive swarming. Intriguingly, however, although two of these three individuals were shown by either morphological or genetic means to be female, they were homozygous at multiple loci that were heterozygous in their mother. Thus these offspring were presumably not produced by the same kind of thelytokous reproduction as observed here. The degree of

homozygosity would suggest that these offspring were the products of the terminal fusion of two pronuclei, or perhaps the mothers of these eggs had some kind of ability to eliminate sperm pronuclei, yet maintain heterozygosity at the sex locus. Although such a mechanism seems unlikely, a reciprocal situation is known to occur in the little fire ant *Wasmannia auropunctata*, where the maternal genome is eliminated in eggs destined to be queens, thus allowing the male mating partners of queens to be genetically reincarnated as queens (FOURNIER *et al.* 2005).

The mechanism by which a queen might choose (or at least influence) the ploidy of her unfertilised eggs is difficult to envisage. In arrhenotokous populations queens have complete voluntary control over whether or not a particular egg they lay is fertilized (RATNIEKS and KELLER 1998). If the queen encounters a drone-sized cell, (which she measures with her front tarsi, KOENIGER 1970), she refrains from releasing sperm onto the egg as it is laid. The egg then develops arrhenotokously as a male (WINSTON 1987). If she encounters a worker-sized cell she releases a minute amount of sperm onto the surface of the egg as it is laid, and these eggs develop as females (HARBO 1979). The process is remarkably accurate, and queens rarely make mistakes (RATNIEKS and KELLER 1998). No such mechanical option is available to Cape queens. In order to be able to choose the ploidy of her egg a queen must be able to influence the second division meiosis that occurs in her egg *after* it has been laid, presumably by some signal encoded as the egg is laid. If she desires to lay a female-producing egg in a worker cell (or a queen cell) she must cause the two central pronuclei to fuse *post partum*. If she desires to lay a male-producing egg in a drone cell she must cause all but one of the 4 pronuclei to degenerate, while the remaining pro-nucleus begins to divide mitotically, again *post partum*. And she must be able to

switch between the two kinds of parthenogenesis depending on the kind of cell she is laying in.

In many insect species mitotic division of the zygote is stimulated by the presence of sperm in the cytoplasm of the egg (SANDER 1985). In the Hymenoptera, however, an alternative mechanism is required because unfertilised eggs can develop by arrhenotokous parthenogenesis. In honey bee queens this stimulus is the physical squeezing of the egg as it is laid (SASAKI and OBARU 2002), so division occurs whether the egg is fertilized or not. Perhaps the queen goes through the same physical motion as she would to release sperm onto the egg, and this somehow stimulates the central pro-nuclei to fuse rather than to die, perhaps by a secretion from the accessory gland of the spermatheca.

Pearcy *et al.* (2006) explored the population genetics of thelytoky in the ant *C. cursor*. They showed that as with *A. m. capensis*, thelytoky is achieved by central fusion of automictic products. In *C. cursor*, colonies are established by parthenogenetic daughters of queens, which are generally highly inbred (PEARCY *et al.* 2004).

Workers, in contrast, are produced sexually, and workers may become the mothers of a replacement queen if their queen dies (PEARCY *et al.* 2004). This ant system differs from that of the Cape bee where queens almost always produce both daughter queens and workers sexually (JORDAN *et al.* 2008). A parthenogenetic queen lineage will eventually have a high rate of homozygosity, even if recombination is constrained, and will be uncompetitive with more heterozygous queens laid by workers. This may explain why asexual reproduction, which we have shown is possible in Cape honey bee queens, is rarely used for the production of daughter queens. Perhaps the method

of sex determination differs between *C. cursor* and the honey bee, so that inbreeding is less of a problem in the ant.

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TABLE 1

Predicted effects of recombination events and different kinds of gamete fusion during thelytokous parthenogenesis on the probability, r , that a locus heterozygous in the mother will be homozygous in the offspring, and the consequences if the locus is the complementary sex determiner (CSD)

<u>Mode of parthenogenesis</u>	<u>Recombination</u>			
		Absent		Present
	r	CSD	r	CSD
Terminal fusion	1	Inviabile	1/3	1/3 inviable
Central fusion	0	Viable	1/3	1/3 inviable
Random fusion	1/3	1/3 inviable	1/3	1/3 inviable
Gamete duplication	1	Inviabile	1	Inviabile
Apomictic	0	Viable	0	Viable

TABLE 2

Numbers of thelytokous and arrhenotokous progeny produced by unmated Cape queens and the number of queen-laid progeny found in incorrect cells.

Queen	Drones			Workers		
	Number genotyped	Number produced arrhenotokously by the virgin queen	Number laid incorrectly in worker cells by the virgin queen	Number genotyped	Number produced thelytokously by the virgin queen	Number laid incorrectly in drone cells by the virgin queen
1	47	0	0	99	99	0
2	76	17	10	4	4	0
3	48	48	0	4	4	0
4	19	19	0	32	4	0
Total	190	84	10	139	111	0

Genotypes of individual bees used to compile this table are given in on line supplementary material at.....

TABLE 3

**Linkage distances, *D*, in cM estimated between two pairs of loci on linkage group
1 from various progeny**

<i>D</i> estimated from progeny of:	Locus pair	
	Am 062-031	Am 062-109
<i>A. m. capensis</i> queen (normal meiosis) ¹	22.4	-
<i>A. m. capensis</i> queen (thelytokous parthenogenesis) ²	56.5	32.4
<i>A. m. mellifera</i> worker (arrhentokous parthenogenesis) ¹	-	6.5
<i>A. m. mellifera</i> queen (normal meiosis) ³	17.4	4.2

¹ Table 2 of Baudry *et al.* (2004) using the Haldane correction

² This study, calculated using the Rizet and Engelmann (1949) correction from data in Figure 1.

³ Derived from ‘Solignac-3’ map (SOLIGNAC *et al.* 2007)

Figure legends

FIGURE 1—Representation of Linkage groups 1 and 3 of the honey bee derived from the genetic map Solignac_3 (SOLIGNAC *et al.* 2007) showing the loci studied here. The cumulative map distance from one telomere is given after the name of each locus. The observed recombination fraction between pairs of loci, θ_{obs} , estimated as the proportion of bees homozygous at one locus and heterozygous at the second is given on the left hand side of each linkage group. The expected recombination fraction between all loci is 0.33 after the Rizet and Engelmann correction. The shaded areas on the chromosomes are thought to encompass the centromeres (BAUDRY *et al.* 2004). The location of the *complimentary sex determiner* (*csd*) locus is indicated on linkage group 3.

FIGURE 2—Proportion of individuals homozygous at a locus plotted against the ‘Solignac_3’ (SOLIGNAC *et al.* 2007) genetic maps for linkage groups 1 and 3. Centromeric regions determined by Baudry *et al.*(2004) are indicated by the bars. The location of the *complimentary sex determiner* locus is indicated on linkage group 3.

Fig 1

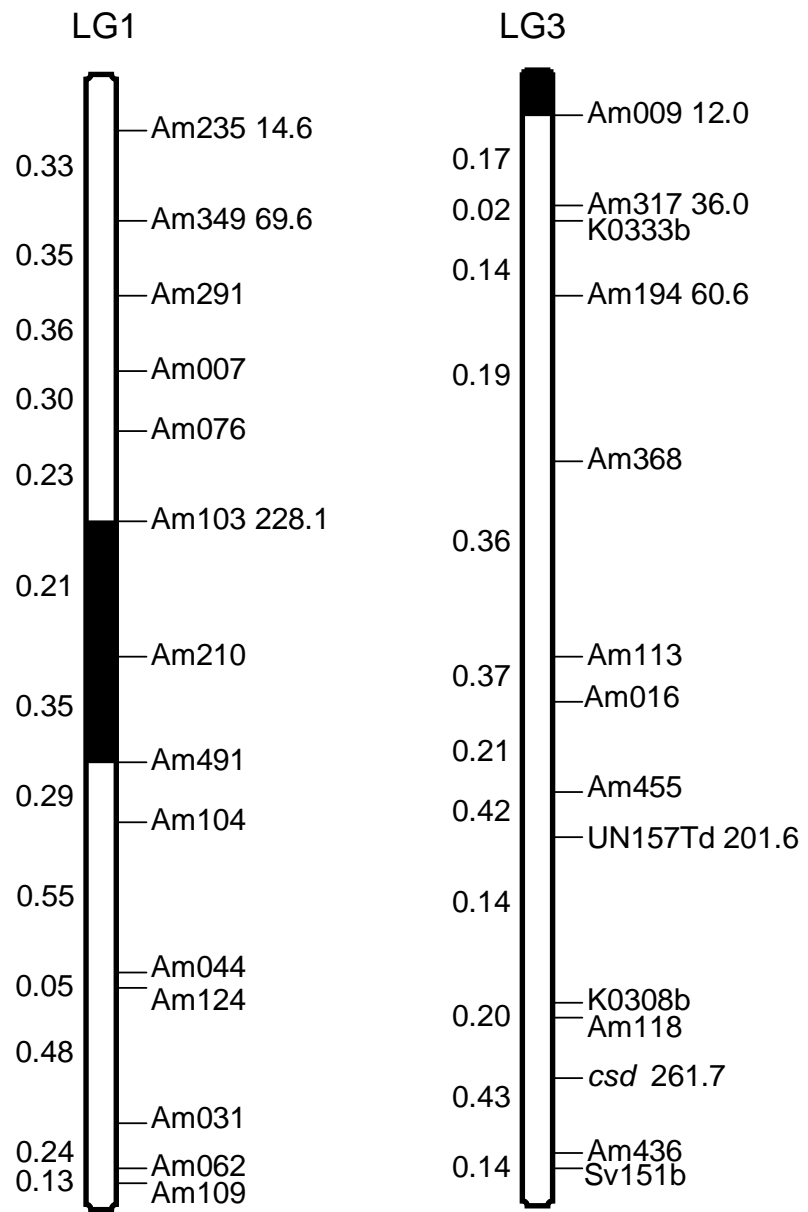


Fig 2

