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Title: Interspecific interactions and premating reproductive isolation

Year: 2008

Version: Published version

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Please cite the original version:

Tynkkynen, K., Kotiaho, J. S., & Svensson, E. I. (2008). Interspecific interactions and premating reproductive isolation. In A. Gordoba-Aguilar (Ed.), *Dragonflies: Model Organisms for Ecological and Evolutionary Research* (pp. 139-152).

Interspecific interactions and premating reproductive isolation

Katja Tynkkynen, Janne S. Kotiaho, and Erik I. Svensson

Overview

Interspecific interactions have several evolutionary consequences: for example, two species may compete, hybridize, or behave aggressively towards each other, or there may be predator–prey interactions. One consequence of these interactions is the evolution of premating reproductive isolation between the two species. The most obvious interspecific interaction, which has an effect on reproductive isolation, is the avoidance of hybridization, or, in other words, the reinforcement process. The theory of reinforcement states that when hybridization is maladaptive, selection pressure causes a divergence in female mate preference and/or in male secondary sexual characters. It is often assumed that, ultimately, females are responsible for hybridization because they are the choosier sex and their co-operation is needed for successful copulations. Despite this view, it is possible that males, rather than females, are responsible for hybridization, especially in species in which males can force copulations. In addition to the avoidance of maladaptive hybridization, other interspecific interactions, such as aggression or predation, may also have an effect on premating reproductive isolation. For example, if interspecific aggression is directed towards males with the most exaggerated sexual characters because of the similarity of these characters between the two species, natural selection for sexual character divergence may arise. Just like the reinforcement process, this process may lead to a strengthening of premating reproductive isolation. In this chapter, we will review the effect of interspecific interactions on premating reproductive isolation. As examples we will mostly use the studies conducted on *Calopteryx* damselflies.

11.1 Introduction

In nature, there are several kinds of interaction between species. Species can be involved in predator–prey interactions (including parasites; Zuk and Kolluru 1998; Kotiaho 2001; Svensson and Friberg 2007), they may compete or defend territories against each other (e.g. Nomakuchi and Higashi 1996; Genner *et al.* 1999; Adams 2004; Tynkkynen *et al.* 2006), and two species can hybridize (see Mallet 2005). Interactions between species can have evolutionary consequences, influencing reproductive isolation (Box 11.1) and speciation events. For example, competition over resources can cause divergent selection on ecological or behavioural

traits, strengthening premating isolation between the two incipient species (see Coyne and Orr 2004 for a review of speciation).

Although there are several factors which may affect reproductive isolation, we will here concentrate on two questions. First, how can interspecific interactions, especially hybridization, aggression, and predation, affect premating reproductive isolation? Second, is reproductive isolation a direct target of selection or does it evolve as a correlated response to selection on other traits? We will mainly use *Calopteryx* damselflies as a model species, which have been under intensive study concerning our study questions.

Box 11.1 Reproductive isolation

Reproductive isolation can arise from several reproductive barriers, which may work before or after mating. Pre-mating barriers include factors that prevent the occurrence of mating, whereas post-mating reproductive barriers include factors that, after mating, prevent the formation of offspring, or that cause the hybrid offspring to be unviable or sterile (Coyne and Orr 2004).

In this chapter, we concentrate on pre-mating reproductive isolation. Pre-mating reproductive isolation can be classified as caused by several factors (see Coyne and Orr 2004 for more

detailed description). First, there may be sexual isolation (also called behavioural isolation), in which reproductive isolation is based on behavioural differences between two species, or mate-recognition ability of individuals (for an example case, see Box 11.2). Second is ecological isolation, which means that gene flow between species is impeded because of a divergence in habitat or timing of reproduction. Finally, mechanical isolation occurs when reproductive structures are incompatible, thus preventing normal copulation (Coyne and Orr 2004).

11.2 Causes and evolutionary consequences of hybridization in animals

Hybridization means crossing of genetically distinguishable groups or taxa, leading to the production of viable hybrids (*sensu* Mallet 2005). Hybridization is a surprisingly prevalent phenomenon in nature: at least 25% of plant species and 10% of animal species hybridize (Mallet 2005). In animals, hybridization is mostly observed in conspicuous and intensively studied taxa such as birds (Grant and Grant 1992), mammals (see Coyne and Orr 2004), *Drosophila* (Coyne and Orr 1989; Mallet 2005), and butterflies (Mallet 2005). In odonates, some suspected and verified hybrids have been observed (e.g. Asahina 1974; Bick and Bick 1981; Corbet 1999; Monetti *et al.* 2002; Sánchez-Guillén *et al.* 2005). However, odonate hybrids may be more common than reported, since precopulative tandems and matings between heterospecifics are commonly observed in nature (Corbet 1999).

11.2.1 Why do animals hybridize?

A pair of species can hybridize only if they occur in sympatry (at the same place), and if they have incomplete pre- and postmating isolation barriers. In animals, the ultimate causes of hybridization can roughly be divided into two groups: either there is incomplete mate-recognition ability (e.g. Sætre *et al.*

1997a; Randler 2002) or hybridization is the result of adaptive decision-making (Nuechterlein and Buitron 1998; Wirtz 1999; Veen *et al.* 2001).

Hybridization can be caused by incomplete mate-recognition ability, if, for example, behavioural isolation between the two species is incomplete. Such is the case for instance, when the two species have occurred in allopatry (at separate places), and then subsequently become sympatric. Here, the mate-recognition ability that evolved in allopatry may be poorly adapted to discriminating heterospecifics. It is also possible that two or more closely related species occur in sympatry without hybridization, but after some environmental change, they start to hybridize (Rhymer and Simberloff 1996; Seehausen *et al.* 1997a; Lamont *et al.* 2003; Box 11.2).

Hybridization is not always maladaptive. For example, when an individual is unable to find a conspecific mate, it may do better by breeding with a heterospecific than not at all. This adaptive explanation for hybridization requires that the hybridizing individual gains some fitness return, meaning that hybrids can have reduced, but non-zero fitness (Grant and Grant 1992; Nuechterlein and Buitron 1998; Sætre *et al.* 1999; Wirtz 1999; Veen *et al.* 2001; Randler 2002). However, the two origins for hybridization, incomplete mate-recognition ability and adaptive decision-making, are not mutually exclusive, since both may occur at the same time within a sympatric population (Sætre *et al.* 1997a; Veen *et al.* 2001).

Box 11.2 An example of the relaxation of pre-mating reproductive isolation

In Lake Victoria are at least 500 endemic haplochromine cichlid fish species (Seehausen *et al.* 1997b). In these cichlid species, male coloration is an important factor in sexual selection, and works as a pre-mating isolating barrier between species (Seehausen *et al.* 1997a; Seehausen and van Alphen 1998). Females of the cichlid species are able to recognize conspecific males based on their coloration in clear water conditions. However, recent anthropogenic

disturbance has increased the water turbidity and the coloration of the males is not easy for the females to observe. This has caused matings between heterospecifics that result in viable and fertile hybrids (Seehausen *et al.* 1997a). Thus, the relaxation of pre-mating reproductive isolation due to anthropogenic environmental change has resulted in formation of hybrid swarms, and may have led to species extinctions through introgression (Seehausen *et al.* 1997a).

11.2.2 Consequences of hybridization

Hybridization leads to different outcomes depending on the viability and fertility of the hybrid individuals. If hybrid fitness is not reduced, or if it is even higher than that of parents (a phenomenon known as heterosis), the mechanisms to avoid hybridization are not strong or are missing, and the gene pool of the two species may eventually fuse (see, for example, Coyne and Orr 2004; Taylor *et al.* 2006). In such a case, if one species is less abundant than the other, the fusion of the gene pools may lead to extinction of the less-abundant species (or the sympatric population) (e.g. Liou and Price 1994; Rhymer and Simberloff 1996; Levin 2004). In the example of the Lake Victoria cichlid fishes (Box 11.2), relaxation of premating reproductive isolation had resulted in extinctions through introgression (Seehausen *et al.* 1997a).

When species hybridize there is often a catastrophic reduction in fitness, and thus selection to avoid interspecific matings is likely to be very strong. As a consequence, the mate preferences and the sexual characters of a pair of potentially hybridizing species may diverge. This process is referred to as reinforcement because it should reinforce pre-mating reproductive isolation of the species (e.g. Dobzhansky 1951; Howard 1993; Sætre *et al.* 1997a; Rundle and Schluter 1998; Noor 1999; Higgie *et al.* 2000). As a result of reinforcement, the probability of copulation between heterospecifics, and thus the occurrence of hybrids in the wild, is reduced. The divergence of reproductively isolating traits

between allopatric and sympatric populations of two species is referred to as reproductive character displacement (*sensu* Howard 1993; see Butlin 1987 for a different definition of the concepts). The pattern may also be observed across sympatric populations, in which the divergence of isolation traits depends on the relative abundance of the sympatric species (Howard 1993; Noor 1999; Tynkynen *et al.* 2004). It should also be taken into account, that in addition to hybridization, matings between heterospecifics without production of hybrids may cause selection for reproductive isolation, if mating or courtship itself has a negative effect on fitness (see Chapman *et al.* 1998; Kotiaho *et al.* 1998; Kotiaho 2001).

When pre-mating isolation has been reinforced such that matings between heterospecifics are not prevalent, the selection to further enhance the pre-mating isolation may relax. It is possible, however, that sexual selection develops along with the reinforcement process, and that its role increases when reinforcement proceeds (Liou and Price 1994; Coyne and Orr 2004; Mallet 2005). When sexual traits or female mate preferences are shaped by reinforcement (i.e. by natural selection), a genetic correlation will be formed between male sex traits and female mate preference (Lande 1981; see also Liou and Price 1994; Coyne and Orr 2004). Similar patterns to reinforcement may also be caused by forces other than avoidance of maladaptive hybridization. For example, ecological factors and the associated selection pressures may cause pre-mating reproductive

isolation, which arises as a byproduct to divergent natural selection on different populations (e.g. Noor 1999; Coyne and Orr 2004; Mullen and Andrés 2007; see also Servedio 2001). One such ecological factor is interspecific aggression; an example of which is described later in this chapter.

11.2.3 Male coercion behaviour: can it lead to hybridization in odonates?

In general, females are thought to be choosy and it seems that female cooperation is frequently needed for a successful copulation. Therefore, it has been proposed that it is the females who are mainly responsible for hybridization (Sætre *et al.* 1997b; Wirtz 1999; Randler 2002). However, in some cases, males can coerce females to copulation, and thus male choosiness should not be neglected (e.g. Cordero 1999; Cordero and Andrés 2002). If females are indeed the choosier sex, it may be too costly for males to evolve to be choosy, since mating opportunities with conspecific females may also be missed (Parker 1983; Sætre *et al.* 1997b; Parker and Partridge 1998; Wirtz 1999). Nevertheless, it is likely that males will also be selected to display some discrimination of conspecifics, especially if the costs of reproduction activities are high (Wirtz 1999).

In some species of odonates, males are able to grasp a female and form the precopulatory tandem, and the females may have little option but to mate (Oppenheimer and Waage 1987; Cordero and Andrés 2002). This is the case in *Calopteryx* damselflies: when the anal appendages of the male grab the female neck to produce the tandem position, the pair usually proceeds with copulation (Oppenheimer and Waage 1987). Moreover, in most odonates it is unclear whether female mate choice even occurs (see Fincke 1997; Fincke *et al.* 1997; but see Córdoba-Aguilar 2006). The role of female mate choice seems to be more clear in calopterygid damselflies, in which males have conspicuously pigmented wings, and in which males court females by performing a flight display (Fincke *et al.* 1997; Siva-Jothy 1999; Córdoba-Aguilar 2002). However, since even in these species males can force females to mate (Cordero 1999; Cordero and Andrés 2002), it is thus still unclear who is, in fact, in control of the onset of mating.

Interestingly, in his classical studies, Waage (1975, 1979) suggested that character displacement observed in wing pigmentation of *Calopteryx maculata* and *Calopteryx aquabilis* is caused by avoidance of maladaptive hybridization by males rather than females. Although hybrids were not found, Waage (1975) observed copulations between heterospecifics followed by oviposition. Males of *C. maculata* had better mate-discrimination ability in sympatric populations than in allopatric ones, indicating occurrence of reinforcement. The selection for better mate discrimination may also have caused reproductive character displacement on female wing transparency; that is, on the trait in which males base their species recognition (Waage 1975, 1979). It is particularly interesting that in this case reproductive character displacement was observed in female characteristics and in male mate discrimination (Waage 1975, 1979), because in most other studies of reproductive character displacement it is the male secondary sexual characters or female mate-discrimination ability that has been investigated (e.g. Howard 1993; Sætre *et al.* 1997a; Rundle and Schluter 1998; Noor 1999; Higgie *et al.* 2000). It should be noted, however, that the suggestion of reinforcement made by Waage (1975) was challenged recently by the suggestion that reproductive character displacement may have arisen from selection caused by interference between mate-recognition signals rather than reinforcement (Mullen and Andrés 2007).

The role of females and males in reproductive isolation has been studied in detail in *Calopteryx splendens* and *Calopteryx virgo*. Hybrids between these two species occur in nature in low prevalence (less than 1% of adult individuals), but matings between heterospecifics are frequent (Svensson *et al.* 2007; K. Tynkkynen, J.S. Kotiaho, M. Luojumäki, and J. Suhonen, unpublished results). The discrepancy between number of observed matings and hybrids, and the pronounced genetic distance between the two species (Misof *et al.* 2000; Weekers *et al.* 2001), suggests that hybridization is likely to be maladaptive.

Females of these species mate assortatively and use male wing coloration as a visual cue in species recognition (Svensson *et al.* 2007). *C. splendens* males have melanized wing spots in their wings covering

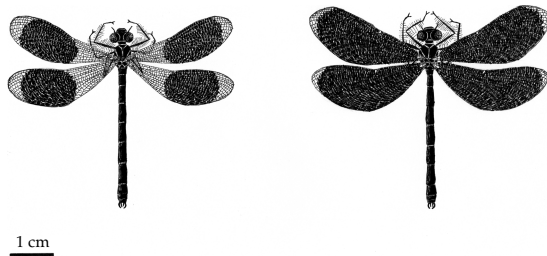


Figure 11.1 Males of *Calopteryx splendens* (left) and *Calopteryx virgo* (right). Artwork provided kindly by the artist Kaisa J. Raatikainen.

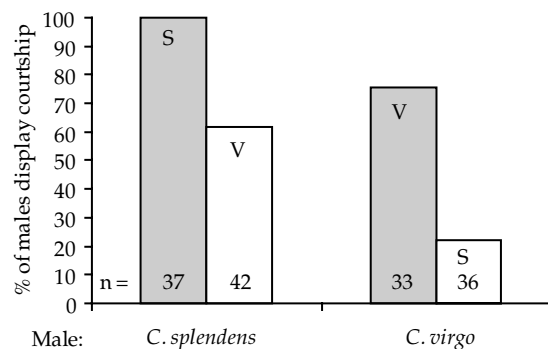


Figure 11.2 Percentage of males displaying courtship to conspecific (grey bars) or heterospecific (white bars) females. S indicates *C. splendens* and V means *C. virgo* females (Tynkkynen *et al.*, 2008). Reproduced with permission from Elsevier.

about 30–70% of wing area, and *C. virgo* have almost completely pigmented wings (Figure 11.1). As the extent of the wing pigmentation of *C. splendens* males was manipulated to resemble that of *C. virgo*, *C. virgo* females became more willing to copulate with *C. splendens* males, and at the same time the willingness of *C. splendens* females decreased (Svensson *et al.* 2007). Thus, there is strong sexual premating reproductive isolation between the two species (Svensson *et al.* 2007).

There is also a difference between *C. splendens* and *C. virgo* males in courting activity towards conspecific and heterospecific females (Tynkkynen *et al.* 2008; Figure 11.2). In central Finland, it was found that when heterospecific females were presented to territorial males of both species in field experiments, *C. splendens* courted as many as 62%

of the females, whereas *C. virgo* courted only 22% of the females (Figure 11.2). This suggests, that in contrast to females, among which the sexual premating reproductive isolation was strong (Svensson *et al.* 2007), the premating isolation in *C. splendens* males was incomplete. The same asymmetrical discrimination between males of the two species was verified independently in a study in southern Sweden, where it was found that *C. splendens* males copulate with heterospecific females in the field more often than do *C. virgo* males (Svensson *et al.* 2007). In contrast, *C. virgo* males were apparently more discriminatory towards heterospecific members of the opposite sex than females in both species and *C. splendens* males (Svensson *et al.* 2007). Thus, in the genus *Calopteryx*, a consideration of male mating behaviours and male mate choice, not just female mate choice, is crucial to understand the causes of hybridization. It is possible that eagerly courting *C. splendens* males are able to force *C. virgo* females to copulate (see Cordero 1999; Cordero and Andrés 2002), and *C. virgo* females yield to mating since it may be costly for them to resist persistent harassment from *C. splendens* males. The elevated species-recognition ability and choosiness of *C. virgo* males (Svensson *et al.* 2007; Tynkkynen *et al.* 2008) may be due to the sex- and species-specific costs of matings: *C. virgo* males have an elevated predation risk from birds, presumably because of their more conspicuous, entirely dark wings (Svensson and Friberg 2007).

11.3 Interspecific aggression

11.3.1 Causes for interspecific aggression

There are at least two causes of interspecific aggression: it can result from interspecific interference competition over resources or from mistaken species recognition. The first is probably more common, as documented by published studies (e.g. Nomakuchi and Higashi 1996; Genner *et al.* 1999; Adams 2004).

Aggressive behaviour related to territorial defence can mistakenly be directed towards heterospecific individuals due to their phenotypic similarity (Murray 1981; Nishikawa 1987; Alatalo *et al.* 1994; Tynkkynen *et al.* 2004). For example, interspecific

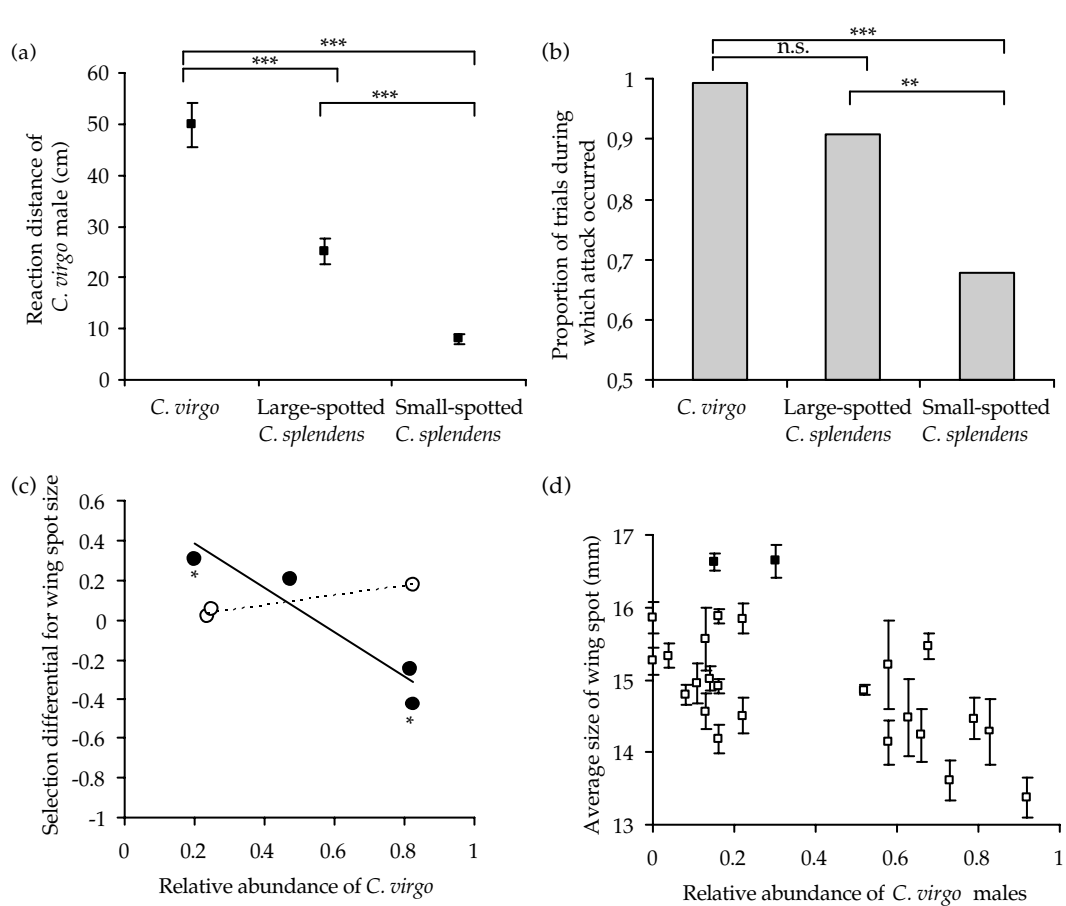


Figure 11.3 Interspecific aggression and character displacement in *C. splendens* and *C. virgo* males. (a, b) Results from the experiment in which aggressiveness of territorial *C. virgo* males was determined towards *C. virgo*, and *C. splendens* males with large and small wing spots. Reaction distance (a) and aggressiveness of reactions (b) were observed. Aggressiveness of reactions is indicated by proportions of trials in which attack occurred (Tynkkynen *et al.* 2004). NS, not significant; ** $P<0.01$, *** $P<0.001$. (c) Standardized selection differentials (measures of directional selection) on wing-spot size of *C. splendens* males in relation to relative abundance of *C. virgo* males. Filled circles indicate natural populations (controls), and open circles populations in which the relative abundance of *C. virgo* is reduced. There is significant interaction between the control and treatment (ANCOVA, $F_{1,3}=17.05$, $P=0.026$; Tynkkynen *et al.* 2005). For individual selection differentials, * $P<0.05$. (d) Character displacement in wing-spot size of *C. splendens* males. The spot size decreases as relative abundance of *C. virgo* males (number of *C. virgo* males divided by the total number of both species) increases. The two populations indicated by filled circles are geographically distant, eastern populations (Tynkkynen *et al.* 2004). Reproduced with permission from Blackwell Publishing.

aggression occurs between males of *C. virgo* and *C. splendens*, and the aggression seems to be at least partially based on mistaken species recognition (although there may also be a component of interference competition over territories or oviposition sites) (Tynkkynen *et al.* 2004, 2006). *C. virgo* males react from greater distance to large-spotted *C. splendens* males, but they are also more aggressive towards large-spotted than small-spotted *C. splendens*

(Figures 11.1 and 11.3a and b). Thus, it seems that, at least partially, interspecific aggression between *C. virgo* and *C. splendens* is based on mistaken species recognition. This is because if the behaviour was due to interference competition, *C. virgo* males should be equally aggressive towards large-spotted and small-spotted *C. splendens* males, whereas reaction distance could still vary, if the large-spotted *C. splendens* males are more detectable.

11.3.2 Consequences of interspecific aggression on premating reproductive isolation

Interspecific aggression may have an effect on premating reproductive isolation of closely related sympatric species at least in two ways. First, interspecific interference competition may cause ecological reproductive barriers to emerge. In other words, there may be shifts in habitat use or behavioural adaptations to avoid contacts with the heterospecific individuals. Such avoidance may lead into strengthening premating reproductive isolation (see Coyne and Orr 2004; Adams 2004). Second, interspecific aggression due to mistaken species recognition may have an effect on male sexual characters, which in turn may affect premating reproductive isolation through female mate choice or through interspecific male–male competition (e.g. Tynkkynen *et al.* 2004, 2005). Regardless of the origin of the aggression, interspecific aggression can reduce the fitness of the target. For example, interspecific aggression may force individuals of subdominant species to less preferred habitats or territories (Alatalo *et al.* 1994; Nomakuchi and Higashi 1996; Martin and Martin 2001; Melville 2002), it may reduce attractiveness of males to females when males are the targets of excessive interspecific harassment, or it may reduce the survival of individuals through injuries or depletion of energy reserves (e.g. Eccard and Ylönen 2002; Tynkkynen *et al.* 2005; Figure 11.3c). In addition, energetically demanding fighting with heterospecifics may reduce territory-holding potential and thus decrease territorial life span, which in turn can reduce reproductive success (Tynkkynen *et al.* 2006).

As an example, interspecific aggression from *C. virgo* towards large-spotted *C. splendens* males could have caused character displacement to the wing spot of *C. splendens* males such that wing-spot size decreases with increasing relative abundance of *C. virgo* males (Tynkkynen *et al.* 2004; Figure 11.3d). If the similarity between the males of the two species decreases, it may facilitate female mate recognition. In addition, it may cause a genetic correlation between a male sexual character and female mate preference (see Lande 1981). A genetic correlation may arise if males with a certain wing-spot size

avoid interspecific aggression, but have sufficiently large spots to appeal to females, giving these males an advantage in both natural and sexual selection. If this process occurs, it may enhance premating reproductive isolation between the species. Thus, it seems likely that interspecific aggression has a role in creating premating reproductive isolation between the two *Calopteryx* species (Mullen and Andrés 2007; Tynkkynen *et al.* 2008).

11.4 A role for predation in the evolution of premating isolation

An influential review on laboratory selection experiments in *Drosophila* concluded that reproductive isolation between populations may often evolve as a correlated response to divergent natural selection between different environments (Rice and Hostert 1993). If two populations are inhabiting different environments and have different ecologies, they will become reproductively isolated from each other, given sufficient time, even if some gene flow connects the populations (Rice and Hostert 1993). This is the so-called byproduct model of speciation, which has been advocated by many workers since Ernst Mayr and in which there is no particular need to invoke any direct or indirect fitness benefits to males or for females to choose their own conspecifics as mates (Coyne and Orr 2004). There is very strong empirical support for this byproduct model of speciation, which is valid for both allopatric and sympatric scenarios (Endler 1977; Hendry *et al.* 2000; Hendry 2001; Svensson *et al.* 2006).

Given the strong evidence for the byproduct model of speciation, it follows that investigators should focus on ecologically important traits between species or incipient species, and estimate the form and direction of natural or sexual selection on the same traits. Interspecific differences in wing coloration in *Calopteryx* damselflies are particularly interesting in this context since they function as a species-recognition mechanisms in both males and in females (see above). If natural or sexual selection on such wing colour differs between populations, divergent selection could cause sexual isolation and speciation as a correlated response. Interestingly, the extent of wing melanization and darkness of the wing patches

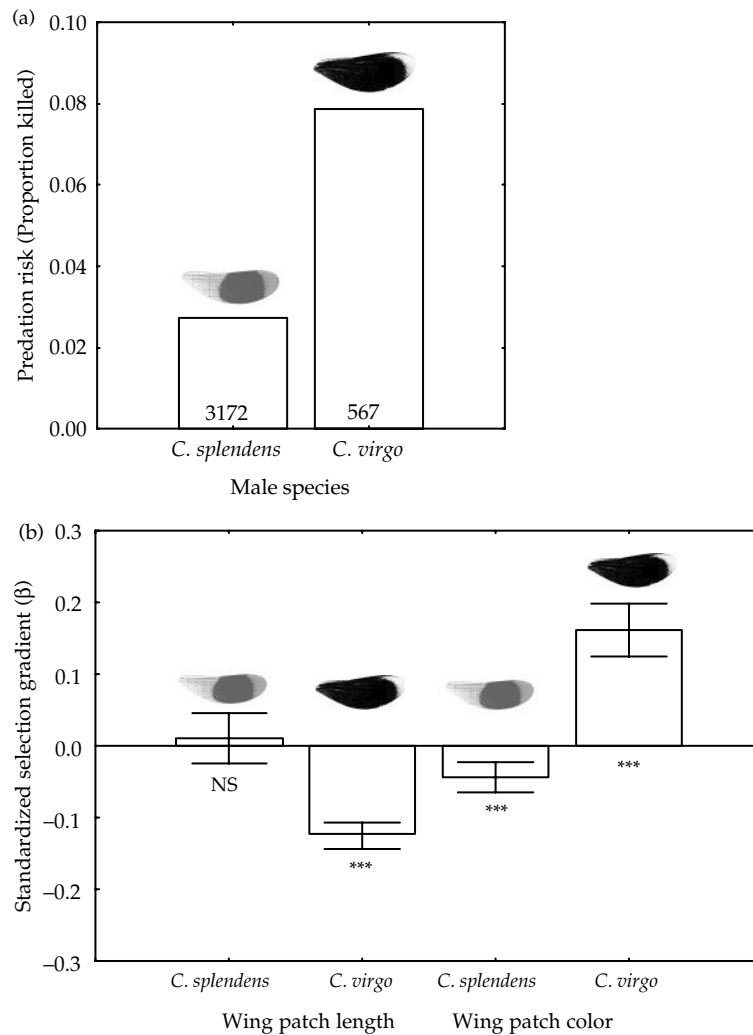


Figure 11.4 (a) Species-specific predation by *Motacilla alba* on *C. splendens* and *C. virgo* males. Male *C. virgo* suffered a 2.9-times-higher predation risk from wagtails than did male *C. splendens* and this difference was highly significant (99.9% confidence interval, 1.54–4.82; $P < 0.001$). Sample sizes (total no. of live and dead males) for each species are also provided in graph. Wings of the two different species inserted in graph for comparison. Note the almost entirely melanized wings in *C. virgo*, compared with *C. splendens* males. (b) Directional selection gradients (β) on wing coloration of *C. splendens* and *C. virgo*. Selection gradients on wings were estimated from morphological measurements of predated individuals and these wings were compared with measurements of live males captured at the same time and at the same locality (see Svensson *et al.* 2007). The selection gradients for all traits differed significantly between species ($P < 0.001$, in all cases). Estimated means \pm 95% confidence intervals are shown in figure. NS, not significant; *** $P < 0.001$. Reproduced with permission from Chicago University Press.

has a strong effect on predation risk by white wagtails (*Motacilla alba*) at a sympatric locality of *C. splendens* and *C. virgo* in southern Sweden (Figure 11.4) (Svensson and Friberg 2007). *C. virgo*, which has entirely dark wings (Figure 11.1), and is

thus presumably more visible to avian predators, had an almost three-times-higher predation risk than sympatric *C. splendens* males (Figure 11.4a). Moreover, both wing-spot size and the intensity of wing-spot darkness were subject to natural

selection, although the selection pressures differed between the two species (Figure 11.4b). Predation selected for a reduced extent of wing melanization but darker spots in *C. virgo*, whereas selection on the smaller wing spot in *C. splendens* was not as pronounced (Figure 11.4b). Taken together, these data indicate that predator-mediated selection acts on wing melanization and hence such selection could potentially interfere with both the character-displacement process between these two species and may indirectly also affect the development of reproductive isolation between populations or species. More generally, the evolution of reproductive isolation between incipient species is thought to be facilitated if the species-recognition characters are subject to direct selection, as in this case, rather than if the species-recognition characters are only 'arbitrary' signalling traits that are only subject to indirect selection (Felsenstein 1981; Dieckmann and Doebeli 1999).

11.5 Sexual isolation as a result of direct or indirect selection on female mate preferences, or a correlated response to divergent sexual selection?

Natural selection on the melanized wing spots in the genus *Calopteryx* could have caused sexual isolation (see Box 11.1) in this genus, irrespective of whether natural selection was caused by male interspecific interactions (Tynkkynen *et al.* 2004, 2005, 2006) or by avian predation (Svensson and Friberg 2007). However, studies on sexual isolation should preferably also be performed at lower taxonomic levels than species; that is, between conspecific populations. Such studies are still rare, relative to the numerous laboratory studies on *Drosophila* (Rice and Hostert 1993; Coyne and Orr 2004). Ideally, such studies should be performed not only between allopatric populations experiencing little or no gene flow, but between parapatric populations with varying degree of gene flow (Hendry *et al.* 2000; Hendry 2001), combined with simultaneous estimation of natural and/or sexual selection regimes in the different populations (Svensson *et al.* 2006). Sexual isolation could evolve between such parapatric populations, even in the presence of gene flow, provided that natural and/or sexual selection

is strong enough to overcome the homogenizing effects of gene flow (Smith *et al.* 1997; Hendry *et al.* 2002; Svensson *et al.* 2006). Most previous studies in this area have focused on the role of divergent natural selection and the associated effects on the development of reproductive isolation (Nosil *et al.* 2002, 2003; Nosil 2004). However, divergent sexual selection can also cause sexual isolation as a correlated response, although the empirical evidence for this is much more limited (Lande 1981; McPeck and Brown 2000; Svensson *et al.* 2006).

Svensson *et al.* (2004) have studied divergent sexual selection and sexual isolation within conspecific populations of *C. splendens* in southern Sweden. These populations are connected by varying degrees of gene flow, as revealed by analysis of molecular population divergence using amplified-fragment length polymorphism (AFLP) markers (Svensson *et al.* 2004). The average neutral molecular population divergence between 12 populations (pairwise F_{st}) ranges from close to zero to approximately 0.13 (Svensson *et al.* 2004), implying strong to moderate gene flow and varying degrees of parapatry between these populations. In spite of ongoing gene flow, sexual isolation has apparently evolved since in six out of seven populations, experimental 'immigrant' males presented to local females in the field have lower courtship success than local males (Figure 11.5a). We also confirmed, in a reciprocal transplant experiment between two of our most intensively studied populations experiencing more or less full gene flow ($F_{st} \approx 0$), that local females to a large extent preferred their own local males, rather than immigrant males (Figure 11.5b). The average courtship success of immigrant males was about 0.85 compared with local males; that is, a selection coefficient of -0.15 against immigrant males (Svensson *et al.* 2006). The average courtship success of male categories in this study was estimated from the average female responses to tethered males, measured on an 11-degree scale that takes into account all the distinct precopulatory behaviours in *Calopteryx* (Svensson *et al.* 2006). Such a strong selection coefficient against immigrant males could be the result of either strong direct selection against females mating with immigrant males; for example, because of physical injury or increased predation risk, a possibility that we consider unlikely.

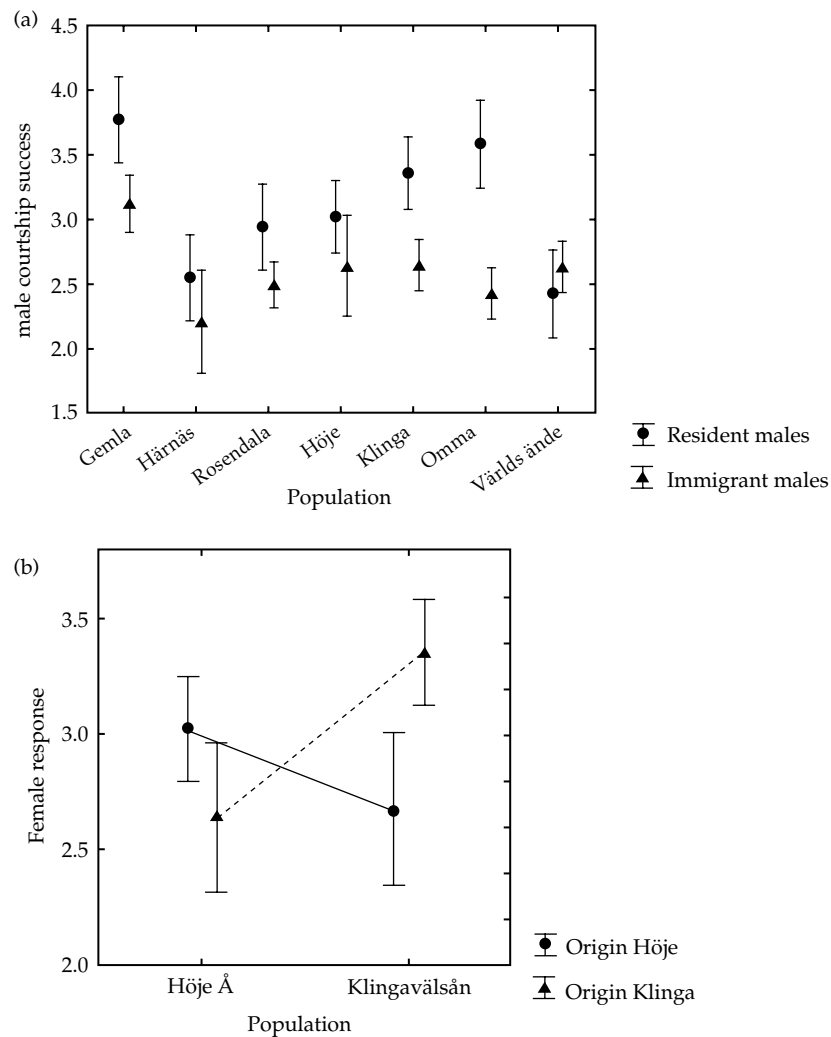


Figure 11.5 (a) Differences (means \pm 95% confidence interval) in male courtship success (measured by monitoring female response; y axis) between resident males and experimental 'immigrant' males in seven different populations of *C. splendens*. Female courtship response towards residents and immigrants was quantified on a nominal scale using well-described and distinct precopulatory behaviors (Svensson *et al.* 2006). Each male was presented to several different females in tethering experiments in the field, and the average female response was used as a measure of male courtship success. (b) A sexually selected fitness trade-off between two damselfly populations differing in morphology and ecology. Resident males have higher courtship success towards females than experimental immigrant males, resulting in a significant phenotype \times origin population interaction (Svensson *et al.* 2006). Reproduced with permission from Blackwell Publishing.

Another possibly more likely explanation is that sexual isolation has arisen as a correlated response to strong divergent selection between these populations. Sexual selection is clearly divergent between these populations (Svensson *et al.* 2006), so there is a clear potential for premating isolation to develop as a purely correlated response, exactly

as predicted by the byproduct models of speciation (Coyne and Orr 2004). Such byproduct models of speciation predict that sexual isolation will arise as a result of either divergent natural selection (Rice and Hostert 1993) or divergent sexual selection (Svensson *et al.* 2006), both which may play a role in odonate divergence (Svensson and Friberg

2007; Svensson *et al.* 2006). Another possibility may of course be that females are rejecting immigrant males because of indirect fitness costs, for example if offspring from such interpopulation matings have lower survival. However, we consider this possibility unlikely given the fairly pronounced gene flow between these populations (Svensson *et al.* 2004), and indirect fitness costs and benefits are also considered to be a relatively weak force in the evolution of mate preferences (Kirkpatrick and Barton 1997; Møller and Alatalo 1999; Arnqvist and Kirkpatrick 2005; Orteiza *et al.* 2005). The reason why premating isolation between the populations on the female side has not more efficiently reduced gene flow between the same populations could be due to male coercive mating behaviour (see above). If immigrant males are able to overcome discriminating local females by obtaining enforced copulations, such male mating behaviour would constrain population divergence by increasing gene flow in spite of emergent sexual isolation among females (see Parker and Partridge 1998).

11.6 Conclusions and suggestions for future research

The possibility that males, rather than females, are primarily responsible for hybridization, opens up new perspectives in future studies of hybridization and reproductive character displacement. For example, almost all studies on reinforcement and reproductive character displacement assume intrinsically that females are performing active mate choice and thus are responsible for hybridization. However, if males are driving the hybridization, it may lead to unexpected patterns. For example, there may be character displacement in secondary sexual characters of the males, but it may be caused by factors other than reinforcement of female mate preferences.

One alternative mechanism for reinforcement and patterns of reproductive character displacement is interference of the mate-recognition signals of the two species. For example, a female may have problems to distinguishing signals of her own species, increasing the costs of mate searching. This may cause similar selection on female mate preference and male sexual characters as does avoidance

of maladaptive hybridization (see Noor 1999; Coyne and Orr 2004; Mullen and Andrés 2007). We suggest that interspecific aggression originating from mistaken species recognition may form an unrecognized form of interspecific interference, which may explain reinforcement of premating isolation (see also Mullen and Andrés 2007). In contrast to increasing the costs for females, mistaken species recognition between males may cause selection on males for better species recognition or divergence in their sexual characters. If male sexual characters and female mate preference are genetically correlated (Lande 1981), interspecific aggression may drive population divergence in male traits, cause a correlated response in female preferences and, as a result, produce a pattern similar to the one predicted by classical reinforcement theory (see Alatalo *et al.* 1994; Tynkkynen *et al.* 2004, 2005, 2006).

The melanized wings in *Calopteryx* are unique in their multifarious and pleiotropic effects on suites of adaptive functions such as their links to male immunological condition (Rantala *et al.* 2000; Siva-Jothy 2000), their effect on predation risk (Svensson and Friberg 2007), their importance in male–male interspecific interactions (Tynkkynen 2004, 2005, 2006), and their role in female choice and intraspecific sexual selection (Siva-Jothy 1999; Svensson *et al.* 2004). In addition to all these documented links to adaptive functions, wing pigmentation also functions as a sexual isolation mechanism between species (Svensson *et al.* 2007). The adaptive and multiple functions of melanized wings in combination with their role in mediating sexual isolation make these an excellent illustration of so-called ‘magig-trait’ models in speciation research (Gavrilets 2004). Future work in the genus *Calopteryx* should aim to clarify whether and how these wing pigmentations have influenced speciation rates in a broader phylogenetic context and the relative importance of natural and sexual selection in speciation processes in this and other odonate groups (Svensson *et al.* 2006). The relative simplicity by which these wing pigmentations can be measured and experimentally manipulated in natural populations should make *Calopteryx* damselflies excellent model organisms in future speciation research.

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