

# Body size shapes average behaviour, social impact, and social responsiveness in agonistic interactions

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## Abstract

Social interactions shape behavioural phenotypes and fitness, and therefore represent major drivers of evolution. Increasingly, it is recognised that plasticity in response to social partners (“social plasticity”) operates through two causal pathways: an individual’s *impact on* and *responsiveness to* partner phenotypes. Functional links among an individual’s average behaviour, social impact, and social responsiveness may have unanticipated evolutionary consequences. Body size is likely to be a key driver of such links, as size often determines the outcome of social interactions. This is particularly true in competitive contexts, where larger contestants generally have an advantage and individuals are expected to decide whether to escalate a contest or retreat based on the difference between their own body size and that of their rivals. We quantified the extent to which the three components, average behaviour, social impact and responsiveness, vary as a function of body size in competitive interactions, using southern field crickets (*Gryllus bimaculatus*) as a study system. We found that smaller animals were on average less aggressive, elicited more aggression in their opponents, and were more responsive by reducing their aggression more strongly when interacting against large opponents. Thus, all

three social behavioural components—average aggression, social impact and social responsiveness—were size-dependent. If heritable, such associations in traits mediating aggression may either constrain or facilitate evolutionary responses to selection in ways not previously recognised. Our findings provide the first empirical evidence for hidden relationships among the key components of social traits, highlighting the need for a multivariate perspective to better predict the evolution of social behaviour.

**Keywords:** Aggression, animal contests, animal personality, resource competition, *Gryllus bimaculatus*, individual-by-environment interaction, indirect genetic effects, reaction norm, social evolution, social plasticity.

## Introduction

Social interactions affect the behavioural decisions, and fitness, of individuals and their social partners (Westneat & Fox, 2010). Social interactions are ubiquitous because all animals interact with conspecifics to forage, compete, and reproduce (Trivers, 1985; Frank, 2007). Two fields of evolutionary biology have extensively studied social interactions. First, behavioural ecology has focussed on predicting the behaviour that individuals should express to maximize fitness in situations where their phenotype is affecting, and affected by, the phenotype of conspecifics. To achieve this aim, behavioural ecologists have used game theory, for example, to predict behaviour expressed during cooperative or competitive interactions (Parker, 1974; Maynard Smith, 1982; McNamara & Leimar, 2020). Second, quantitative genetics has developed theoretical models to predict how selection on traits expressed during social interactions translates into micro-evolutionary change (Moore et al., 1997; McGlothlin et al., 2010; Bijma, 2014). Both fields emphasize the key role that social interactions play as drivers of variation, selection, and evolution.

Individual phenotypes expressed during social interactions (called “interacting phenotypes”) arise from three key components: (1) the individual’s average phenotype, (2) its partner’s social impact, and (3) its social responsiveness (Figure 1) (Araya-Ajoy et al., 2020; de Groot et al., 2023; Wijnhorst et al., 2025). An individual’s average phenotype is determined by environmental and genetic factors that permanently influence its phenotype, known as direct environmental (DEEs) and direct genetic effects (DGEs), respectively (Falconer & Mackay, 1996; Lynch & Walsh, 1998). Variation among individuals in these factors contributes to repeatable differences in average phenotype, which can give rise to broader suites of correlated traits, including behavioural syndromes and animal personalities (Dall et al. 2004; Sih et al. 2004; Réale et al. 2007, 2010). Social impact describes how interacting individuals elicit phenotypic change in their social partners (Araya-Ajoy et al., 2020; de Groot et al., 2023). When this influence has a genetic basis, it is called an indirect genetic effect (IGE; Moore et al., 1997; Wolf

et al., 1998; McAdam et al., 2014). Social responsiveness, in contrast, is a form of phenotypic plasticity defined as the slope of the reaction norm, and captures how an individual's phenotype changes as a function of the phenotype of social partners (Dingemanse & Araya-Ajoy, 2015). Although conceptually related, social impact and social responsiveness arise from distinct causal pathways (Araya-Ajoy et al., 2020; de Groot et al., 2023; Wijnhorst et al., 2025). Previous research has primarily focused on covariance parameters that encapsulate effects of both social impact and social responsiveness, such as those called “responsiveness” (André & Day, 2007; André, 2015), “response coefficient” (Akçay et al., 2009; Akçay & Van Cleve, 2012; Van Cleve & Akçay, 2014), or the “interaction coefficient  $\psi$ ” (Moore et al., 1997; Bleakley et al., 2010; McGlothlin et al., 2010). Consequently variation in both social responsiveness and social impact has so far mostly been disregarded.

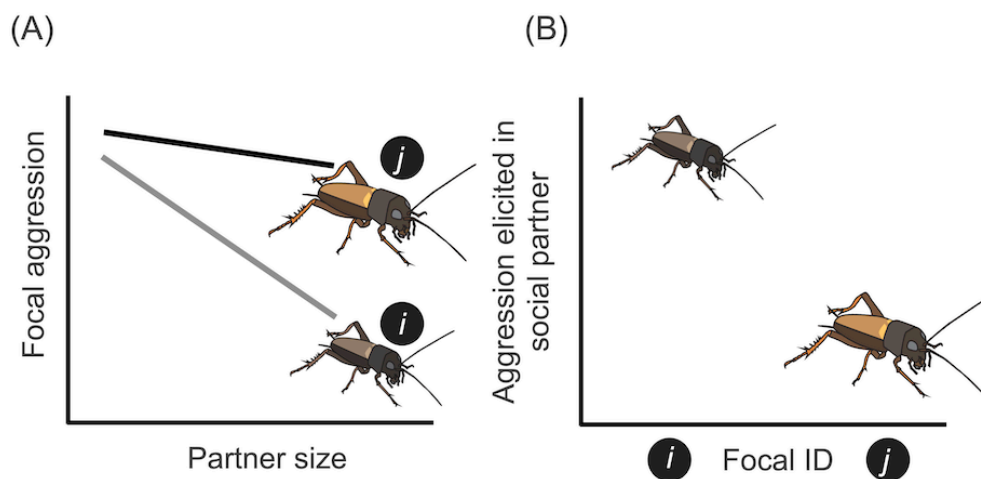
Quantitative genetics theory predicts that the evolutionary response to selection on interacting phenotypes depends on the additive genetic variance in, and covariance between, direct and indirect genetic effects. Direct genetic effects (DGEs) influence an individual's average phenotype, while indirect genetic effects (IGEs) capture the impact of an individual's genotype on the phenotype of its social partners. Positive genetic correlations between DGEs and IGEs are expected to amplify the response to selection, whereas negative correlations should constrain it (Moore et al., 1997; McGlothlin et al., 2010; Bijma, 2014). In addition, genetic correlations between the average phenotype and the level of social responsiveness—the extent to which individuals adjust their phenotype in response to others—should also shape evolutionary trajectories (Bailey et al. 2021). Such correlations imply that the amount of variance explained by DGEs may change depending on the phenotype of social partners, potentially enhancing or dampening evolutionary responses depending on the correlation sign. If interacting phenotypes are indeed shaped by three components—average phenotype, social impact, and social responsiveness—then accurate predictions of micro-evolutionary change require estimates of genetic variance in, and covariance among, *all* three. This raises two key questions: (1) do

populations harbour heritable variation in each component of interacting phenotypes, and (2) are these components genetically correlated?

Heritable variation in average trait values is already well established for interacting phenotypes, as demonstrated by studies reporting non-zero variance attributable to direct genetic effects (DGEs; Stirling et al., 2002; Dochtermann et al., 2015, 2019; Santostefano et al., 2025). Indeed, heritable variation in aggression has been documented across a wide diversity of taxa, including mammals (Kessler et al., 1977; Benus et al., 1991), birds (Verbeek et al., 1996; Duckworth & Kruuk, 2009), and insects (Edwards et al., 2006; Santostefano et al., 2017b). Likewise, heritable variation in social impact is evidenced by studies reporting non-zero variance attributable to indirect genetic effects (IGEs; Bailey & Desjonquères, 2022; Santostefano et al., 2025). Meanwhile, evidence for heritable variation in social responsiveness comes from studies reporting genetic differences in plastic responses to social partners (*e.g.*, Kent et al., 2008; Bleakley & Brodie, 2009; Chenoweth et al., 2010; Bailey & Zuk, 2012). Correlations between DGE and IGE have also been reported (*e.g.*, McAdam & Boutin, 2004; Wilson et al., 2009, 2011), indicating genetic covariance between average phenotype and social impact. Genetic correlations between average phenotype and social responsiveness may also exist, as suggested by among-individual correlations between these components observed across a range of species (*e.g.*, Benus et al., 1990; Natarajan et al., 2009; Strickland et al., 2021). Nevertheless, to our knowledge, no study has yet attempted to infer variation or covariation across all three components of interacting phenotypes.

This study aims to address this gap by testing whether average behaviour, social responsiveness, and social impact (co)vary across individuals due to variation in body size. We designed an experiment to quantify, first, the three components of interacting phenotypes (average trait expression, social impact, and social responsiveness) and, second, test whether body size mediates their expression. We used agonistic interactions in male southern field crickets (*Gryllus bimaculatus*) as model system. Our analysis indirectly addresses patterns of

(co)variance that—in line with Cheverud (1988)’s conjecture—are predictive of genetic (co)variance in this species (Santostefano et al., 2017a,b; Han et al., 2018). This is because aggression and aggression elicited in conspecifics are both repeatable and heritable, and are negatively correlated at the individual and additive genetic level (Santostefano et al., 2017b). We hypothesised that functional links between average aggression, social impact, and social responsiveness would arise due to variation in a non-behavioural trait: body size. In many animals, including field crickets, body size is a critical determinant of fighting success and influences aggression during contests (Gammell & Hardy 2003; Briffa, 2015). Larger individuals are more likely to win fights (Simmons, 1986; Shackleton et al., 2005; Reaney et al., 2011) and tend to be more aggressive (Dixon & Cade, 1986; Simmons, 1986). We thus predicted that (1) larger males to be more aggressive and that (2) males should in general reduce their aggression when faced with large opponents. We also expected that larger, more aggressive males are less responsive to opponent size, thereby linking average aggression and responsiveness (Figure 1A). Lastly, larger males should reduce their opponent’s aggression, thereby linking body size and social impact, and average aggression and impact (Figure 1B).



**Figure 1.** Illustration of size-mediated differences in social responsiveness (A) and social impact (B). In this example, individual *i* is smaller, more responsive to social partners’ phenotype, and elicits more aggression in social partners than individual *j*.

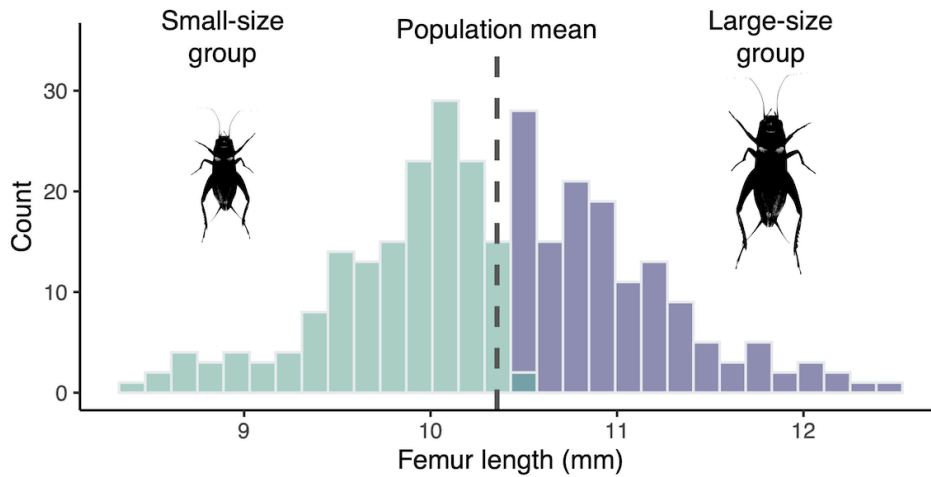
## Materials and methods

### *Origin and maintenance of the crickets*

All crickets were derived from our stock population that originated from Sevilla (Spain) and was maintained under controlled conditions at LMU Munich for four generations (Ratz et al. 2024). The stock was kept under a 12L:12D photoperiod at 28°C and 60% humidity, and provided with commercial dry cat food, fish flakes, and water. We separated a total of 340 males as juveniles in their last nymphal stage and housed them in individual containers (10 × 10 × 9 cm) furnished with a plastic tube section for shelter, and supplied with ad libitum water and dry cat food.

### *Experimental design*

Our aim was to evaluate the relationships among body size, average aggression, social impact, and social responsiveness to partner body size. We thus designed an experiment where each individual was first categorized as “small” or “large” (Figure 2); we then created dyads of all combinations based on these size groups (focal-opponent: small-small, small-large, large-small, large-large). While doing so, we ensured that each individual interacted (repeatedly) with “small” and with “large” opponents. To this end, we allocated males that had reached sexual maturity (*i.e.*, ≥10 days after adult moulting) to one of two size groups based on the population-average body size. As a proxy for body size, we used the length of the rear femur measured using digital callipers (Mousseau & Roff, 1989; Nosil, 2002). To purge measurement error in our analysis, we used the mean of three measurements of femur length taken from each individual (Arnqvist & Martensson, 1998; Fruciano, 2016). We assigned males below the population mean to the “small” group, and males above the population mean to the “large” group measured on approximately 80 experimental males.



**Figure 2.** Distribution of the rear femur size among the 340 experimental males. As part of our experiment, males below the population mean were assigned to the “small” group, while males above the population mean were assigned to the “large” group.

### *Aggression assays*

Aggression was assayed repeatedly using a well-established protocol (Santostefano et al., 2016, 2017a,b; Han et al., 2018; Jäger et al., 2019). Briefly, males were first marked with non-toxic acrylic paint one day prior to their first aggression assay. To begin an assay, we placed a focal male and an opponent male (arbitrarily assigned) in the centre of an arena ( $14.5 \times 15.5 \times 9 \text{ cm}^3$ ). We allowed approximately 1 min for acclimation, and recorded the interaction for five minutes using a high-resolution digital video camera (Basler GenICam, Germany) at a distance of 43 cm above the arena, and set at 25 frames/s and  $1600 \times 1200$  pixels resolution. Males generally rapidly engage in agonistic displays that often escalate into fights (Adamo & Hoy, 1995; Hofmann & Stevenson, 2000). After each test, males were returned to their original individual containers.

Over the course of the experiment, each male was assayed 8 times with a “small” opponent and 8 times with a “large” opponent. While our goal was to test each male in a total of 16 times (realized mean = 12.5 times), individuals that died or lost two or more legs were



removed from the experiment before completing all contests. All videos were scored using the EthoVision software (Noldus EthoVision XT 10, Noldus Information Technology). To prevent carry-over effects across assays caused by winner or loser effects, we allowed a minimum of one day before conducting repeat aggression assays on the same individuals (Han & Dingemanse, 2017; Santostefano et al., 2017b; Han et al., 2018). This interval is conservative as such effects generally disappear within 6 h after fighting (Khazraie & Campan, 1999; Rillich & Stevenson, 2011). As a measure of aggressiveness, we used the total movement toward the opponent, which is computed in EthoVision as the total distance focal males moved relative to their opponents calculated across all video frames (Santostefano et al., 2016, 2017b). We recorded a total of 1822 dyadic interactions involving 289 males.

### ***Statistical analysis***

All analyses were performed using R (v.4.1.2; R Development Core Team, 2021). The following packages were used for general analysis and plotting: *car* (v.3.0; Fox et al., 2019), *MASS* (v.7.3; Venables & Ripley, 2002), *rstan* (v. 2.21.5; Stan Development Team, 2022), *shinystan* (v.2.6.0; Gabry & Veen, 2022), *bayesplot* (v.1.9.0; Gabry & Mahr, 2024), *ggplot2* (v.3.4.0; Wickham, 2016), and *cowplot* (v.1.1.1; Wilke, 2020).

Our aim was to estimate the main effects of focal body size (*i.e.*, size-related average behaviour), opponent body size (*i.e.*, size-related social impact) and the interaction between focal and opponent body size (*i.e.*, size-related social plasticity). Aggression was standardized, and log-transformed to approach normality. The model was fitted in *rstan* using common regularizing priors for all model parameters, assuming normal distributions of mean 0 and standard deviation of 1. Priors for standard deviations and variances followed a half-normal probability function. All correlations were assigned a Lewandowski-Kurowicka-Joe (LKJ) prior with shape parameter 1, which specifies a uniform probability density function bounded between -1 and 1 (Lewandowski et al., 2009). The model was fitted with 4 chains, a warmup of 2000

iterations, 4000 sampling iterations, and a thinning interval set to one. Visual inspection of plots of posterior distributions versus fitted values using the *shinystan* interface revealed satisfactory fit.

The model was fitted with focal ( $A_{D,i}$ ), opponent ( $A_{S,j}$ ) and dyad ( $Dyad_{i,j}$ ) identity as random effects (Equation 1), and random slopes for focal identity ( $\psi_i$ ) as a function of opponent size group  $Z_j$  (small or large). For measurement  $k$  of focal individual  $i$  interacting with individual  $j$ , focal aggression  $Y$  was modelled as:

$$Y_{ijk} = \mu + A_{D,i} + \beta_1 Z_i + (\bar{\psi} + \psi_i) Z_j + \beta_2 Z_i Z_j + A_{S,j} + Dyad_{i,j} + e_{ijk} \quad (1)$$

Where  $\mu$  represents the population average aggression,  $\beta_1$  the fixed effect of focal size group  $Z_i$  (coded as small or large),  $\bar{\psi}$  the population-level social responsiveness to social partner size,  $\beta_2$  the fixed interaction effect between focal and partner size, and  $e_{ijk}$  the residual error. Prior model checks revealed that the residual variance varied as a function of opponent size (see Table S2), and we thus modelled heteroskedasticity by allowing the residual variance to depend on opponent size:  $e_{ijk} \sim \mathcal{N}(0, \sqrt{e^{\mu_e + \beta_e Z_j}})$ , where  $\mu_e$  is the residual variance in the average social environment, and  $\beta_e$  the difference in between-social-environment variance across the two size groups.

## Results

### *Size-driven average behaviour, social responsiveness, and social impact*

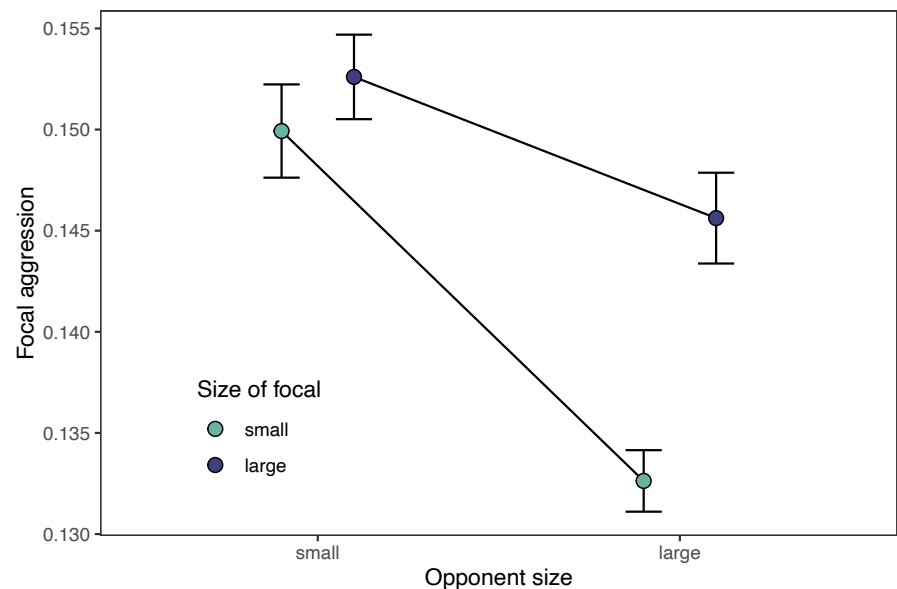
We detected strong evidence for size effects on average level of aggression, social responsiveness, and social impact (Table 1; Fig. 3). Small and large animals differed in both their average aggression and level of social responsiveness: large individuals were on average more aggressive (positive effect of “focal size”; Table 1, Fig. 3) and reduced their aggression less strongly with opponent size (positive effect of focal  $\times$  opponent size; Table 1, Fig. 3). Thus, body size mediated the magnitude of social responsiveness. Meanwhile there was a negative main

effect of opponent size on focal aggression (Table 1; Fig. 3), indicating that larger opponents overall suppressed aggression in focal individuals. Thus, body size also mediated the magnitude of social impact.

**Table 1:** Median estimates ( $\pm$  95% CI) of the main effects on focal aggression.

	Estimate (log)	Estimate (back-transformed)
Fixed effects ( $\beta$ ) <sup>1</sup>		
Intercept ( $\mu$ )	0.008 (−0.043, 0.062)	0.141 (0.130, 0.143)
Focal size ( $\beta_1$ )	0.117 (0.046, 0.189)	0.004 (0.001, 0.006)
Opponent size ( $\bar{\psi}$ )	−0.253 (−0.322, −0.185)	−0.008 (−0.010, −0.006)
Focal $\times$ opponent size ( $\beta_2$ )	0.130 (0.004, 0.255)	—

<sup>1</sup>Focal size measures size-related average behaviour, opponent size measures size-related impact, focal  $\times$  opponent size measures size-related social responsiveness.



**Figure 3.** Large males were on average more aggressive, more socially impactful, and less responsive to opponent body size. Circles and error bars indicate means  $\pm$  standard errors. Data are based on 1822 dyadic interactions involving 289 males.

## Discussion

We have shown that body size affects all three major components of contest behaviour: average aggression, social impact, and social responsiveness. Specifically, small individuals were less aggressive, less impactful, but more responsive to size differences in opponents compared to large individuals. Overall, our findings indicate that body size can constrain the expression, and plausibly evolution, of flexible social behaviours.

Our findings that focal body size mediates all three components (*i.e.* average aggression, social impact, and social responsiveness) highlight the plurality of roles that size can play in social interactions. The level of aggression and the magnitude of the impact on conspecifics were greater in large males, whereas responsiveness was greater in small males. The association between size and average aggression is expected given that larger males, which are typically stronger competitors (Shackleton et al., 2005; Reaney et al., 2011), should benefit from maintaining high levels of aggression to assert dominance (Dixon & Cade, 1986; Simmons, 1986). Likewise, the association between size and impact is expected given that large males should suppress aggression in their competitors, as these competitors should retreat and avoid escalating a contest when the likely to win is relatively low (Rillich et al. 2007). Importantly, this association between size and impact implies that variance attributable to differences among social partners, and potentially to indirect genetic effects (IGEs), can be shaped by its body size. Yet body size in insects is subject to developmental plasticity, being largely determined by resource acquisition during juvenile development (Edgar, 2006; Mirth et al., 2007; Chown & Gaston, 2010). Thus, genes influencing resource acquisition and allocation during juvenile development might play a predominant role in driving indirect genetic effects during social interactions. To our knowledge, this perspective is yet to be considered, and we encourage future work testing possible genetic links between juvenile growth and social impact in interacting adults.

Large focal individuals showed a weaker response to variation in their opponents' size, and were thus less socially responsive than small individuals. This finding demonstrates that body size not only influences aggression in agonistic interactions but also shapes how interacting individuals adjust their own behaviour to others. It further raises the question as to why body size influences the level of responsiveness in social interactions. One reason to expect responsiveness to decrease with body size is because males larger than the population mean are statistically more likely to encounter smaller rivals and win fights (Simmons, 1986; Shackleton et al., 2005; Reaney et al., 2011). Given their higher probability of success, large males may benefit from maintaining high aggression toward any rivals, as predicted by game theory (e.g., Phillips et al., 2018; Fouilloux et al., 2022). In contrast, small males, who are more likely to lose contests, may gain by finely tuning their behaviour to the phenotype of their opponents, avoiding the costs of escalated fights with stronger competitors while still exploiting opportunities to win against weaker ones. More generally, the coexistence of socially responsive and unresponsive strategies within a population is predicted by game theory (Wolf et al., 2008, 2011) and reflects commonly observed patterns in natural populations (e.g., Araya-Ajoy & Dingemanse, 2017; Montiglio et al., 2017; Perez et al., 2019). While responsiveness might be maintained as a mechanism to adaptively adjust behaviour in aggressive disputes (Tello-Ramos et al., 2019; Wolf et al., 2008), behavioural consistency can be an alternative successful strategy. This can be the case, for example, when individuals gain information about conspecifics by monitoring their interactions with others and where behavioural consistency could therefore reduce the occurrence of extended fights (Dall et al., 2004; Schuett et al., 2010).

To our knowledge, only a handful of studies have examined such links between an average social behaviour and its associated *impact on* and *responsiveness to* others. In mice, highly aggressive males are less responsive, reducing less their aggression toward females compared with less aggressive males (Benus et al., 1990; Caramaschi et al., 2008; Natarajan et al., 2009). Meanwhile, in agamid lizards, individuals that are more responsive to changes in

population density are more likely to form social associations (Strickland et al., 2021). From an evolutionary perspective, our findings suggest that variation in key traits mediating social interactions, such as body size, could help maintain multiple aggression–impact–responsiveness strategies in populations, provided that alternative strategies yield comparable selective advantages. Such size dependence of social impact and social responsiveness also implies that indirect genetic effects (IGEs) could exert stronger or weaker evolutionary effects, either accelerating or constraining responses to selection (Wolf et al., 1998; McAdam et al., 2014), depending on the phenotypic composition of the group. For example, IGEs may have a stronger influence on the expression and evolution of aggression in populations composed of predominantly smaller individuals. As with other environmentally dependent traits (Postma & van Noordwijk, 2005; Nussey et al., 2007), these findings suggest that group composition could alter heritability, the ratio of genetic to phenotypic variance (Falconer & Mackay, 1996), thereby increasing or reducing the evolutionary potential of interacting phenotypes (Bijma & Wade, 2008). There is now a pressing need for empirical research to test these evolutionary implications. Quantitative genetic studies adopting a multivariate perspective on interacting phenotypes are particularly promising to tackle this gap as they allow quantifying the genetic (co)variances underlying social traits. Such work will be crucial to identify genetic associations that may constrain or promote social evolution (Figure S1).

It is becoming clear that two distinct mechanisms, social impact and social responsiveness, cause plastic responses during social interactions, and that underlying correlations have the potential to shape the evolutionary trajectory of interacting phenotypes (Araya-Ajoy et al., 2020; de Groot et al., 2023). Using a reaction norm framework, we show that these components covary with body size, a key morphological trait mediating social interactions. These findings imply that social effects, and IGEs, can themselves be plastic and shaped by the social environment. This is an important and unsuspected property of social interactions and their evolutionary consequences which merits further investigation. If the association between body

size and the social behaviour has a genetic basis, selection on one will also drive evolutionary change in the other, as expected for any pair of genetically correlated traits (Lande, 1979; Lande & Arnold, 1983; Walsh & Blows, 2009). Social interactions are central to many processes that drive the evolution of both social and non-social traits. Our study highlights the importance of considering the multivariate nature of traits mediating social interactions. We suggest that the reaction norm approach provides a useful framework to explore phenotypic plasticity in the social context, opening up novel and exciting directions in the study of social evolution.

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# **Data and code availability**

<https://osf.io/jgrez>

# **Conflict of interest**

We declare no conflicts of interest.

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