

Original Article

Barking up the wrong frog: global prevalence of misdirected amplexus in anuran amphibians

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ABSTRACT

Reproduction is an energy-expensive life process in many organisms, and accurate conspecific recognition is crucial to successful reproduction. Nonetheless, misdirected attempts at amplexus towards heterospecifics appear to be common in many anuran amphibians. Such reproductive interference can have significant ecological and evolutionary consequences, but its prevalence remains unknown. Here, we compile a global dataset of anuran misdirected amplexus and test how phylogenetic relatedness, ecological niche, breeding phenology, and geography influence the prevalence of anuran misdirected amplexus. We find misdirected amplexus to be significantly higher among more closely related heterospecific pairs than between random pairs, but still occurred between species of different families. Misdirected amplexus was also significantly more common in arboreal and terrestrial anurans, and among species pairs with similar body sizes or with similar microhabitat use. We also show that misdirected amplexus is significantly more common among explosive breeders compared with prolonged breeders and is more prevalent in temperate regions compared with the tropics. Overall, we show that misdirected amplexus among anuran amphibians is not a rare phenomenon and that its prevalence is influenced by evolutionary relatedness, microhabitat use, ecological similarity, and breeding phenology. These interactions have the potential to cause species decline, hence understanding them is crucial.

Keywords: interspecific amplexus; heterospecific mating; heterospecific amplexus; reproductive interference; niche overlap; breeding phenology; explosive breeding

INTRODUCTION

Reproduction is a fundamental aspect of every living organism. In sexually reproducing organisms, the existence of the species largely depends on successful mate recognition and reproduction (Ryan *et al.* 2003). Given that species also invest in other life processes, such as survival and growth (Chen *et al.* 2013), reproductive investment should be optimized by assessing the costs associated with resource allocation, survival, and fecundity (Williams 1966, Vitt *et al.* 2009). The energetic costs of reproduction specifically are threefold: (i) the production of eggs or sperm; (ii) investment towards secondary sexual characters; and (iii) investment towards reproductive behaviour (Wootton 1985). According to the 'costs of reproduction' hypothesis, reproductive strategies and life-history traits (e.g. reproductive age, clutch size) that maximize the sum of the present and future

reproductive successes would evolve (Williams 1966, Stearns 1989, Harshman and Zera 2007, Bourne *et al.* 2023). Although several studies have found support for the 'costs of reproduction' hypothesis (Harshman and Zera 2007), there are several instances where species indulge in erroneous interactions. These sexual interactions between animal species that decrease the fitness of at least one of the species involved are known as reproductive interferences (Cothran 2015). Seven types of reproductive interference have been described: signal jamming, heterospecific rivalry, misdirected courtship, heterospecific mating attempts, erroneous female choice, heterospecific mating, and hybridization (Gröning and Hochkirch 2008). Such misdirected heterospecific interactions during the different stages of mate acquisition can lead to reproductive interference and can affect the fitness of individuals (McLain and

Shure 1987, McLain and Pratt 1999, Thum 2007, Gröning and Hochkirch 2008, Maisonneuve *et al.* 2023). Although reproductive interference is widespread in many organisms (Gröning and Hochkirch 2008) and can affect the demography and structuring of ecological communities (Ting and Cutter 2018, Iritani and Noriyuki 2021), its ecological significance and prevalence in different groups remain understudied. Here, we aim to understand the prevalence of reproductive interferences through misdirected mating attempts in anuran amphibians.

Amphibians show a wide variety of reproductive strategies that range from aquatic egg laying to loss of larval stages or direct development to viviparity and parental care (Duellman and Trueb 1994). Based on combinations of reproductive traits, such as amplexus type, oviposition sites, clutch characteristics, development, and parental care, researchers have now identified >74 reproductive modes or strategies (Nunes-de-Almeida *et al.* 2021), making amphibian reproduction the most conspicuous feature of its biology (Duellman and Trueb 1994, Wells 2007). Amphibians are also one of the most vulnerable vertebrates to current climate change and other anthropogenic pressures (Carey and Alexander 2003), and many species are undergoing a drastic decline globally (Beebee and Griffiths 2005, Collins 2010, Luedtke *et al.* 2023). This has sparked interest in understanding how climate change, habitat alterations, and anthropogenic pressure can affect the breeding behaviour of anurans and influence their reproductive fitness. However, many species of amphibians frequently make mistakes in identifying conspecifics during mating attempts (Mollov *et al.* 2010, Serrano *et al.* 2022, Brischoux and Lorrain-Soligon 2023). Such interactions with heterospecifics can have ecological, evolutionary (Calabrese and Pfennig 2022, Chen *et al.* 2022), and behavioural (Schroeder and Podos 2023) implications that can lead to reproductive interference and also reduce fitness (Hettyey *et al.* 2014). Although misdirected amplexus is widespread and is known from >150 species of amphibians (Serrano *et al.* 2022), what drives the prevalence of such heterospecific interactions remains largely unexplored.

Here, we compile a dataset of published records of misdirected amplexus in amphibians and test specific hypotheses that drive the prevalence of such events. It has been shown that misdirected amplexus attempts are more frequent in a few amphibian families, such as Bufonidae, Ranidae, Hylidae, and Rhacophoridae (Serrano *et al.* 2022). Thus, we initially tested the prevalence of misdirected amplexus across different families. It has also been hypothesized that reproductive interference is more frequent among closely related species (Gröning and Hochkirch 2008). Given that the anatomies of species that produce sexual signals during mate acquisition are largely constrained by evolutionary relatedness (de Kort and ten Cate 2001), mistakes in recognizing such sexual signals are more likely among species that show low species divergence (de Kort and ten Cate 2001, Sosa-López *et al.* 2016). Thus, we tested the hypothesis that misdirected amplexus is more prevalent among closely related heterospecifics. Misdirected amplexus with heterospecifics can also occur when there is considerable overlap in mate recognition traits, such as body size, between species within the community (Okuzaki *et al.* 2010). Structurally complex environments can also affect the efficient transmission of mating signals (Goutte *et al.* 2018, Hardt and Benedict 2021), thereby increasing the chances of

heterospecific mating owing to signal interference (Wollerman 1999, Gordon and Uetz 2012). Thus, we also tested the hypotheses that misdirected amplexus occurs among species with considerable overlap between species traits and microhabitats, and that misdirected amplexus would be more prevalent in structurally complex microhabitats, such as arboreal and terrestrial environments. Breeding phenology can also influence the chances of misdirected amplexus. Intense competition for mates in species with short bursts of synchronous breeding activity can lead to males amplexing heterospecific males or even inanimate objects (e.g. stones, shoes) (Pearl *et al.* 2005, Wells 2007). Thus, we tested whether misdirected amplexus was more frequent in explosively breeding species in comparison to prolonged breeders. We also tested whether misdirected amplexus was more common in temperate regions, where seasonal climatic changes restrict breeding activity to short periods, in comparison to tropical regions, which experience more stable climatic conditions.

MATERIALS AND METHODS

Data collection

We compiled a dataset of available published records on anuran misdirected amplexus through an exhaustive literature search. We used the software PUBLISH OR PERISH v.8, October 2021 (<https://harzing.com/resources/publish-or-perish>; Harzing 2007) to search for published articles indexed by 'Google Scholar' about misdirected amplexus in amphibians using specific keywords. The keywords included '(frog OR anura OR anuran) AND (interspecific OR intergeneric OR interfamilial OR homosexual OR cross OR erroneous OR unnatural OR unusual OR anomalous) AND (amplexus OR breeding OR mating OR fertilisation)'. Given that most observations of misdirected amplexus in amphibians were published in herpetological journals, we searched specifically for articles in 31 well-known herpetological journals using the same keywords (see [Supporting Information, File S1](#)). Articles that we found manually during random searches were added to this list. Additionally, we used the same keywords to search for images on social media and citizen science platforms, such as Facebook, Instagram, Wikimedia Commons, and iNaturalist (see [Supporting Information, File S1](#)). Social media and citizen science observations were included only when we could confirm the identity of the species, the locality, and when the observations were made in natural conditions and not in captivity. Species identities for such observations were confirmed based on references available on Amphibia-Web Database (<https://amphibiaweb.org/>) and Amphibians of the World v.6.2 (<https://amphibiansoftheworld.amnh.org/>). Data collection was carried out until December 2022.

Data classification

From each article in our list, we extracted the amplexed species, i.e. the species that was on the bottom and mounted upon, and the amplexant species, i.e. the species that was on the top and mounted the other individual. We also noted the family of both the amplexed and amplexant species, the geo-coordinates (if available), the zoogeographical zone, and the habitat in which the observation was made. For species wherein the exact coordinates were unavailable, we extracted the approximate latitude

and longitude based on the broad location. We then classified these locations as tropical (between 23.5°S and 23.5°N) and temperate (outside the tropical belt) regions. The type of amplexus (i.e. inguinal, axillary, etc.) exhibited naturally by the amplexant and amplexed species was noted following Carvajal-Castro *et al.* (2020). The type of amplexus generally seen in the amplexant and amplexed species during conspecific mating was also noted using the paper by Carvajal-Castro *et al.* (2020) or other published literature. To this dataset, we added the data on misdirected amplexus compiled by Serrano *et al.* (2022) and removed any repeated observations to obtain our final dataset. In addition to these, we also collected body size data and microhabitat data that were available for both the amplexed and amplexant species. The average body size (snout–vent length) data were obtained for males of each species from the amphibian traits database (Huang *et al.* 2023). Microhabitat use was classified as arboreal, terrestrial, aquatic, or burrowing, and was mostly obtained from the study by Moen and Wiens (2017) and the International Union for Conservation of Nature (IUCN) Red List of Threatened Species. In some cases, we also obtained microhabitat data from other published sources (see Supporting Information, File S1). Species that were classified as being semi-arboreal, semi-terrestrial, semi-aquatic, or semi-burrowing were grouped as arboreal, terrestrial, aquatic, or burrowing, respectively. We also classified the breeding phenology of the amplexant species as being either ‘explosive’ or ‘prolonged’ breeders following Wells (1977, 2007), Bell *et al.* (2017), and other published sources (see Supporting Information, File S1).

Statistical analyses

We performed all analyses in R v.3.6.0 (R Core Team 2019). To test whether misdirected amplexus is more prevalent in closely related species, we classified the taxonomic relatedness of the amplexant species with the amplexed species as interspecific (within genus), intergeneric (between genera of the same family), interfamilial (between different families of the same order), interorder (between the three orders of amphibia), and non-amphibian (amplexus with other non-amphibian groups). We then used a χ^2 goodness-of-fit test to test for differences in the frequency of misdirected amplexus between groups against the null expectation and then carried out pairwise comparisons between groups using the R package RVAIDEMEMOIRE v.0.9-69 (Hervé 2023). We also tested whether the phylogenetic distance between species pairs engaging in misdirected amplexus was more closely related to each other than random. To do this, we used a dated phylogeny that included each unique species pair from our dataset. We obtained this phylogeny by pruning out species from a large-scale dated amphibian tree generated by Portik *et al.* (2023) using the R package PHYTOOLS v.1.0-3 (Revell 2012). We extracted the phylogenetic variance–covariance matrix for our species pairs using the cophenetic.phylo function in the PICANTE v.1.8.2 package (Kembel *et al.* 2010), then extracted the phylogenetic distance between each pair of species in our dataset using the extract.val function in PHYLIN v.2.0.2 (Tarroso *et al.* 2015) and calculated the average observed pairwise distance. We then generated a null distribution of the mean pairwise distance by randomly shuffling the amplexed species within each biogeographical zone using the R package

PERMUTE v.0.9-7 (Simpson 2022) and generating 1000 new species pairs. We performed these block permutations within each biogeographical zone to produce a more conservative estimate of the null distribution. We compared the average observed pairwise distance with the null distribution of mean pairwise distance obtained from the block permutation of amplexed species using a one-sample *t*-test.

To test whether misdirected amplexus is also more prevalent in sympatric species lacking niche segregation, we evaluated the overlap in body size, amplexus type, and microhabitat use. We first calculated the mean absolute difference in body size between amplexed and amplexant species in our dataset. We then performed a 1000 block permutation of the body-size values of the amplexed species within each biogeographical zone using the PERMUTE v.0.9-7 package (Simpson 2022) and calculated the absolute difference in body size between species pairs to generate a null distribution of body-size differences. We then tested whether the absolute difference in body size was lower than the random null expectation using a one-sample *t*-test. Additionally, we tested the prevalence of misdirected amplexus in species occupying different microhabitats (arboreal, terrestrial, aquatic, or burrowing) using a χ^2 goodness-of-fit test, then performing pairwise comparisons using RVAIDEMEMOIRE v.0.9-69 (Hervé 2023). We then scored each species pair in our dataset for microhabitat and amplexus type overlap and tested the prevalence of misdirected amplexus among species with and without microhabitat and amplexus type overlap using a χ^2 goodness-of-fit test. We also tested the prevalence of misdirected amplexus among species with explosive and prolonged breeding strategies and across climatic regimes, i.e. temperate vs. tropics, using a χ^2 goodness-of-fit test.

RESULTS

Our dataset included 482 observations, of which 392 (81.3%) observations were of misdirected amplexus between amphibians heterospecifics, 45 (9.3%) were incidences of necrophilia between conspecifics, three (0.62%) were misdirected amplexus between non-amphibian heterospecifics, and 42 (8.7%) were observations of misdirected amplexus with inanimate objects. Of these, we had 237 unique pairs of species involved in heterospecific amplexus involving 179 amplexant species and 181 amplexed species. Misdirected amplexus was recorded in 19 frog families, but ~76% of the species belonged to four families: Bufonidae (23.2%), Ranidae (21.5%), Hylidae (19%), and Rhacophoridae (11.4%) (Fig. 1A).

The average cophenetic distance between heterospecific amplexic interactions was 186.75 and was significantly lower than the null distribution of average cophenetic distance obtained by random block permutation of amplexed species ($t = 99.213$, d.f. = 1000, $P < .0001$; Fig. 1B). A χ^2 goodness-of-fit test on the number of misdirected amplexus indicated significant differences in the taxonomic relatedness of species interactions ($\chi^2 = 261.62$, d.f. = 4, $P < .0001$; Fig. 1C). About 40.5% of the observations of misdirected amplexus were intergeneric interactions within the same family and were significantly greater than all other interaction types ($P < .0001$) except for interfamilial interactions ($P = .36$), which accounted for

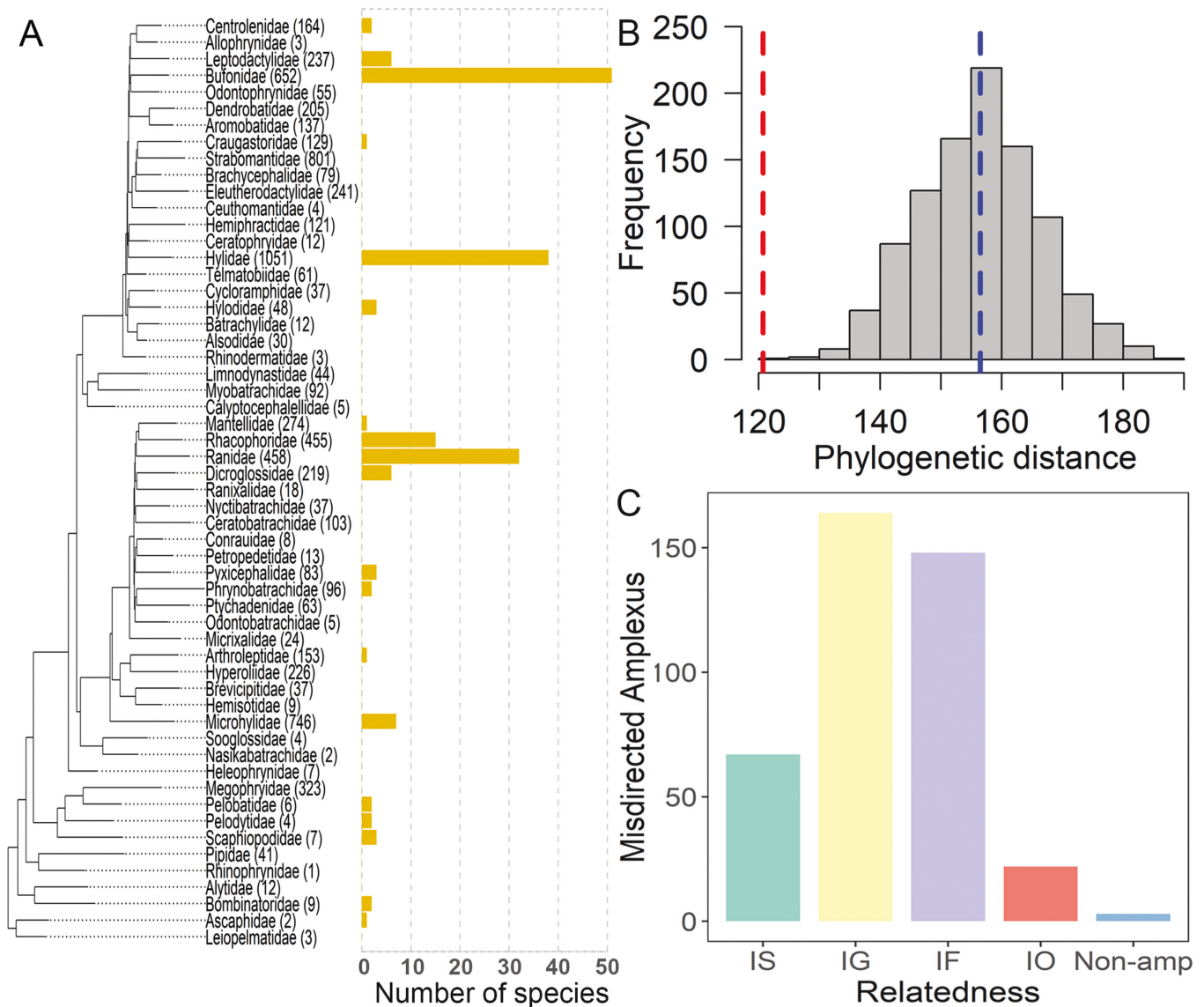


Figure 1. Phylogenetic distribution and relatedness among anuran species involved in misdirected amplexus. A, family-level phylogenetic tree of anurans adapted from Portik et al. (2023), with the number of species involved in misdirected amplexus. The numbers within parentheses represent the total number of species in each family. B, average pairwise phylogenetic distance between species pairs involved in misdirected amplexus (red vertical dashed line) compared with a random distribution of the mean phylogenetic distances. The blue vertical dashed line indicates the mean of the random distribution. C, frequency of misdirected amplexus observed across different levels of taxonomic organisation. Abbreviations: IF, interfamilial; IG, intergeneric; IO, interorder; IS, interspecific; Non-amp, with non-amphibian.

36.6% of the observations. Interspecific misdirected amplexus between species within the same genus accounted for 16.58% of the observations and was significantly greater than misdirected amplexus towards other amphibian orders ($P < .0001$) and other non-amphibians ($P < .0001$).

A one-sample t -test indicated that the mean absolute difference in body size was significantly lower compared with a null distribution of the mean body size differences obtained by random block permutation of amplexed species ($t = 52.798$, d.f. = 1000, $P < .0001$; Fig. 2A). We also found significant differences in misdirected amplexus across microhabitat preferences ($\chi^2 = 35.374$, d.f. = 3, $P < .0001$; Fig. 2B). Misdirected amplexus was significantly more frequent in terrestrial (36.12%) and arboreal (36.12%) species compared

with aquatic ($P = .009$) and burrowing species ($P < .0001$). Misdirected amplexus was also significantly more frequent in aquatic species (20%) compared with burrowing species (7.7%; $P = .007$). Instances of misdirected amplexus were also higher when there was an overlap in microhabitat preferences ($\chi^2 = 28.75$, d.f. = 1, $P < .0001$; Fig. 2C) and the amplexus type ($\chi^2 = 28.77$, d.f. = 1, $P < .0001$; Fig. 2D) of the amplexant and amplexed species.

Comparing the prevalence of misdirected amplexus among breeding strategies, we found significantly greater instances of misdirected amplexus in explosive breeders compared with prolonged breeders ($\chi^2 = 24.961$, d.f. = 1, $P < .0001$; Fig. 3A) and in temperate regions compared with the tropics ($\chi^2 = 49.005$, d.f. = 1, $P < .0001$; Fig. 3B).

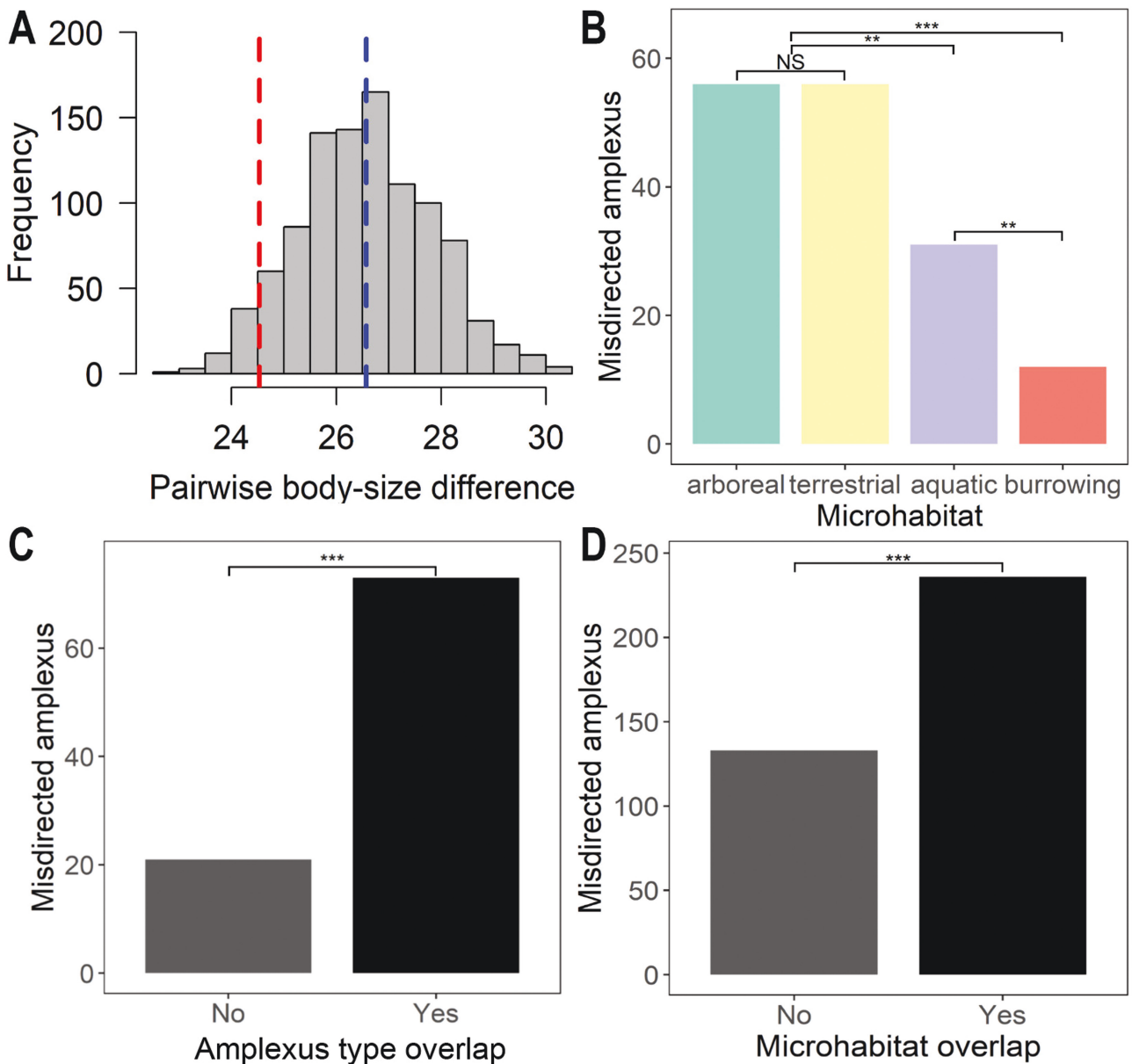


Figure 2. A, average difference in body size between species pairs involved in misdirected amplexus (red vertical dashed line) compared with a random distribution of body-size differences. The blue vertical dashed line indicates the mean of the random distribution. B, frequency of misdirected amplexus across different microhabitat usage. C, frequency of misdirected amplexus among species pairs with and without microhabitat overlap. D, frequency of misdirected amplexus among species pairs with and without similar types of amplexus. *** $P < .0001$; ** $P < .01$; NS, non-significant.

DISCUSSION

Our data, along with other recent studies, indicate that misdirected amplexus is not a rare phenomenon in anurans (Serrano *et al.* 2022, Brischoux and Lorrain-Soligon 2023). We collected 437 observations of true misdirected amplexic interactions (i.e. excluding instances of intraspecific necrophilia) spanning a period of about a century, from 1923 to 2023. Of the 437 records, there were 237 unique heterospecific amplexic interactions with 179 species involved in misdirected amplexus, suggesting that misdirected amplexus occurred more frequently (or was more frequently observed) in some species than in others. Misdirected amplexic interactions were unevenly distributed

across the anuran tree and were primarily from four anuran families: Bufonidae, Hylidae, Ranidae, and Rhacophoridae.

Phylogenetic relatedness

Signals from closely related species are generally more similar to each other than to those of distantly related species, owing to common descent (Sosa-López *et al.* 2016). Thus, it is thought that reproductive interference is more likely among closely related sympatric species that show low levels of species divergence (Gröning and Hochkirch 2008, Kyogoku 2015). Several studies examining conspecific recognition among several groups have found that the ability to discriminate heterospecifics decreases

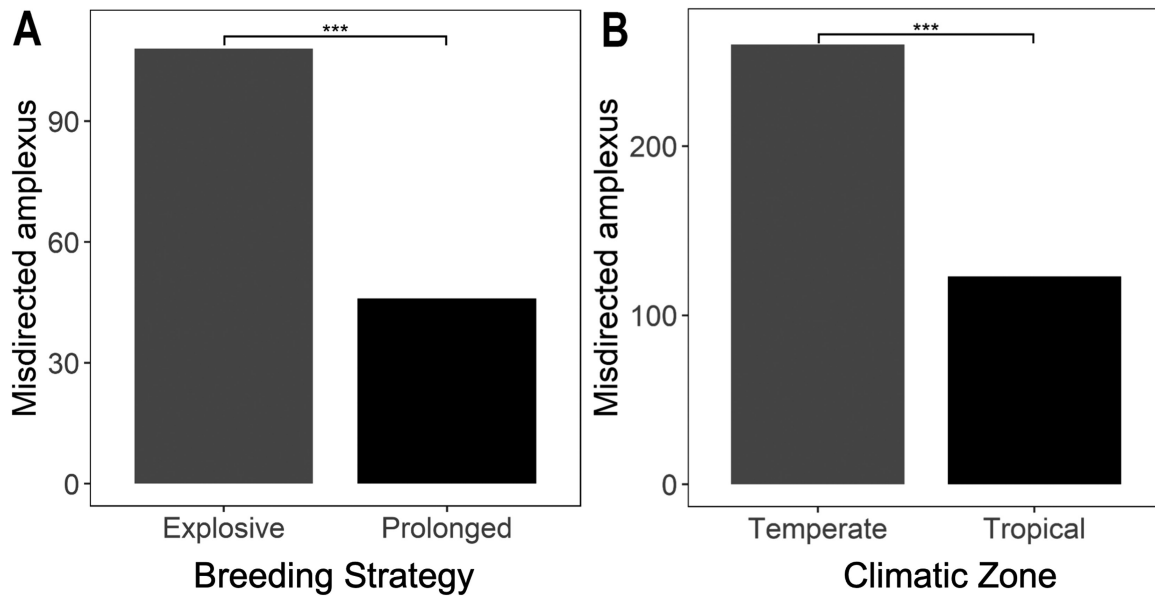


Figure 3. Frequency of misdirected amplexus between species with explosive and prolonged breeding strategies (A) and across geographical regions (B). *** $P < .0001$.

with evolutionary relatedness (de Kort and ten Cate 2001, Sosa-López et al. 2016). For instance, experiments have shown that in many species of birds and arthropods, males are unable to discriminate conspecifics from closely related heterospecifics, while discriminating distantly related heterospecifics (Verrell 1994, Clarke et al. 1996, de Kort and ten Cate 2001, Costa-Schmidt and Machado 2012, Saveer et al. 2014, Wolfenden et al. 2015, Wehi et al. 2017). Our results also indicated that the phylogenetic distance between species engaging in heterospecific amplexic interactions was significantly lower than random, lending support for the hypothesis that misdirected amplexus is greater among closely related heterospecifics. However, we also found that instances of intergeneric and interfamilial misdirected amplexus were significantly higher than misdirected amplexic events between species within genera. There were also some cases of misdirected amplexus with members of different amphibian orders, e.g. *Lithobates sphenoccephalus* (order Anura) amplexing on *Ambystoma maculatum* (order Caudata).

Many amphibians use pheromones for intraspecific sexual signalling during the breeding season (Woodley 2010). Some of these sexual pheromones are highly conserved across species and could lead to mistaken conspecific recognition (Clarke et al. 1996, Saveer et al. 2014). For example, mating secretions (termed amplexins) produced in the nuptial glands of the frog *Rana temporaria* during the breeding season were structurally similar to the sexual pheromones associated with courtship behaviours in plethodontid salamanders (Willaert et al. 2013). Additionally, some courtship pheromones, such as the sodefrin-precursor-like factor (Herrboldt et al. 2021), which enhances female receptiveness in plethodontid salamanders (Houck et al. 2008), have now been shown to be expressed in many anurans (Janssenswillen et al. 2015, Bossuyt et al. 2019).

Although misdirected amplexus might not be random and is more prevalent among closely related species, the conserved nature of many chemical pheromones across amphibians could still lead to mistaken conspecific recognition

(Rollmann et al. 2003) and misdirected amplexus among distantly related species.

Niche and microhabitat overlap

Habitat structure and composition can be a source for different levels of background noise and can significantly alter acoustic signals and affect effective communication (Forrest 1994, Boncoraglio and Saino 2007, Luther and Gentry 2013, Zaffaroni-Caorsi et al. 2023). Complex environments with dense vegetation cover can reflect sound, causing temporal signal distortions (Richards and Wiley 1980, Wells and Schwartz 1982, Forrest 1994, Blumenrath and Dabelsteen 2004). Such degradation or alteration of acoustic signals can lead to greater signal interference (Mathevon and Aubin 1997, Holland et al. 1998) and affect mating success (Gordon and Uetz 2012). Our analyses found that misdirected amplexus is more prevalent in arboreal and terrestrial species compared with aquatic and burrowing environments. Above-ground environments are structurally more complex than aquatic or burrowing environments (Gibert and Deharveng 2002, Grosberg et al. 2012), and thus, could lead to greater signal distortion and interference (Blumenrath and Dabelsteen 2004).

It has been suggested that the lack of sufficient niche segregation among sympatric species can lead to reproductive interference (Gröning and Hochkirch 2008). Our analyses showed that heterospecific reproductive interactions were more prevalent between species pairs that share similar ecological or reproductive traits. Heterospecific reproductive interactions were significantly more common in species with similar body sizes and that overlapped in their amplexus type and microhabitat use. Intense behavioural interference (including reproductive interference) between species generally leads to niche segregation (Grether et al. 2017) through character displacement (Laiolo 2012), niche partitioning (Friberg et al. 2013, García-Navas et al. 2023), or species exclusion that reduces interspecific competition (Kishi et al. 2009, Pastore et al. 2021). However, in some conditions,

habitat filtering for similar traits might still allow ecologically similar species to coexist even with reproductive interferences (Ruokolainen and Hanski 2016).

Breeding phenology

Many species of frogs breed in large aggregations, often triggered by rainfall and lasting for very short periods (Wells 1977, Ulloa *et al.* 2019). Such explosive breeding strategies result in synchronous emergence of the species, which, consequently, reduces the size disparity among tadpoles, leading to reduced cannibalism in tadpoles and eggs (Petranka and Thomas 1995). However, large-scale emergence of a species in a short period can have cascading effects on the assembly and functioning of a species community (Fouquet *et al.* 2020; Getman-Pickering *et al.* 2023), leading to intense scramble competition for resources (Wells 2007). Such intense competition for mates, along with the inability to recognize conspecifics owing to acoustic interference, can lead to males amplexing heterospecifics (Pearl *et al.* 2005, Wells 2007). Our results also support this hypothesis, in that misdirected amplexus was significantly more common in explosive breeders compared with prolonged breeders. However, we still found that ~30% of the misdirected amplexus were in species that showed prolonged breeding. Such patterns could be because of the simplistic dichotomous definitions of explosive and prolonged breeders. In a more real sense, these two categories represent separate ends of a continuum, and a species population might lie anywhere on the gradient (Wells 2007). Many species have prolonged breeding seasons that are divided into short breeding bouts. For example, in the Costa Rican gliding tree frog, *Agalychnis spurrelli*, the breeding period is spread out across 4–5 months, but each breeding event lasts only 1 or 2 days, with a large number of individuals aggregating around ponds (Güell and Warkentin 2023). Breeding phenology can vary across populations of the same species (Cayuela *et al.* 2016) or vary temporally (Klaus and Loughheed 2013). For instance, in the North American frog *Rana boylei*, northern populations breed explosively, with breeding activity lasting only 1 week (Rombough and Hayes 2007), whereas the southern populations show prolonged breeding activity lasting for 6–7 weeks (Wheeler and Welsh 2008). Further studies on the prevalence of misdirected amplexus across populations varying in breeding phenology might provide a better understanding of the costs associated with these mating strategies. Breeding phenology of anurans also varies with latitude (Jørgensen *et al.* 1986). The tropics provide a more stable climate for breeding conditions to persist year round, whereas the breeding window in temperate zones is shorter owing to high variability in climate (Janzen 1967, Deutsch *et al.* 2008, Gutiérrez-Pesquera *et al.* 2016, Shah *et al.* 2017). Such constrained breeding activity might lead to more species breeding explosively, resulting in scramble competition (Vitt *et al.* 2009) and a significantly greater prevalence of misdirected amplexus in the temperate zones than in the tropics.

Implications of misdirected amplexus

Studies have shown that reproductive interferences can potentially lead to population decline and community restructuring (Kyogoku and Wheatcroft 2020). Furthermore, climate change could alter anuran breeding strategies (Todd *et al.* 2010,

Walpole *et al.* 2012, Klaus and Loughheed 2013, Forti *et al.* 2022) and increase negative heterospecific interactions (Parejo 2016, Canestrelli *et al.* 2017). The presence of invasive species could also lead to negative heterospecific reproductive interactions with native species (D'Amore *et al.* 2009). For instance, in island endemic species, invasive species can cause decreased reproductive success and chronic physiological stress in native species (Narayan *et al.* 2015), leading to population decline. Although we shed some insights into the factors affecting prevalence of misdirected amplexus in anurans, further studies exploring the ecological impacts of such maladaptive behaviours are crucial. Such studies are particularly important given the current amphibian crisis (Geyle *et al.* 2021, Lötters *et al.* 2023, Luedtke *et al.* 2023) and the need for informed conservation strategies (McCoy *et al.* 2021).

CONCLUSION

Misdirected amplexus in anurans amphibians is not a rare phenomenon. Its prevalence is influenced by multiple phylogenetic, ecological, and geographical factors. Although close phylogenetic relatedness influences the prevalence of misdirected amplexus, it can still be observed among distantly related species. Misdirected amplexus is also not random and is more common among frogs with explosive breeding strategies and among species with similar ecological traits. Geographical region also influences the prevalence of misdirected amplexus, with temperate zones showing higher prevalence compared with the tropics. Our study sheds some insights into the factors affecting the prevalence of misdirected amplexus in anurans. However, future studies on how these interactions are maintained within populations will be crucial to understand the ecological consequences of such negative heterospecific interactions.

SUPPLEMENTARY DATA

Supplementary data is available at *Biological Journal of the Linnean Society* online.

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AUTHOR CONTRIBUTIONS

S.S. and V.P.C. conceived the study; S.S., V.A., P.J., and V.P.C. collected the data; V.P.C. performed the analyses and made figures; S.S. and V.P.C. drafted and critically revised the manuscript.

CONFLICT OF INTEREST

None declared.

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DATA AVAILABILITY

Data underlying this study are available online as [Supporting Information](#) for this article at the *Biological Journal of the Linnean Society*.

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