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An Ecology of Sperm: Sperm Diversification by Natural Selection

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postcopulatory selection, sexually transmitted disease, sperm aging, sperm competition, reactive oxygen species, genotype-by-environment interactions

Abstract

Using basic ecological concepts, we introduce sperm ecology as a framework to study sperm cells. First, we describe environmental effects on sperm and conclude that evolutionary and ecological research should not neglect the overwhelming evidence presented here (both in external and internal fertilizers and in terrestrial and aquatic habitats) that sperm function is altered by many environments, including the male environment. Second, we determine that the evidence for sperm phenotypic plasticity is overwhelming. Third, we find that genotype-by-environment interaction effects on sperm function exist, but their general adaptive significance (e.g., local adaptation) awaits further research. It remains unresolved whether sperm diversification occurs by natural selection acting on sperm function or by selection on male and female microenvironments that enable optimal plastic performance of sperm (sperm niches). Environmental effects reduce fitness predictability under sperm competition, predict species distributions under global change, explain adaptive behavior, and highlight the role of natural selection in behavioral ecology and reproductive medicine.

1. A FRAMEWORK OF SPERM ECOLOGY

In almost all species, only a tiny fraction of ejaculated sperm reaches an egg and interacts with it for fertilization. The function of these few sperm is central to the study of evolution because only those functioning sperm deliver genetic information to the next generation. Sperm function is also central to other biological areas. For example, reduced sperm function is one of the most important known causes of human infertility in the Western world (Hirsh 2003, Pizzol et al. 2014) and is central to assisted reproduction technologies. For other species on the planet, sperm function is the target of animal breeders to improve the reproductive capacity of livestock (Billard & Cosson 1992, Froman et al. 2006) and the target of geneticists to optimize conservation programs (Roldan & Gomendio 2009). It is under substantial scrutiny in ecotoxicology as a trait affected by environmental pollution (Hayes 2011, Tavares et al. 2013) and is used in a range of toxicity bioassays (Hoorstra et al. 2004, Rajkovic et al. 2006). A striking feature of sperm cells is their enormous evolutionary diversification, particularly in morphology (reviewed in Pitnick et al. 2009), which is currently attributed to sexual selection: Diversification in sperm form and function arises because the sperm of genetically different males compete, and the outcome of the competition varies within different female genotypes, thus leading to selection for competitive sperm (Birkhead et al. 2009, Manier et al. 2013).

Here we propose a framework for sperm evolution and diversification that incorporates the environmental and genetic components of sperm function. We start by briefly reviewing various ways in which the large number of environments affects many different sperm functions and their relative strengths. We then apply several simple ecological concepts to sperm biology to provide a more comprehensive view of its role in ecology, evolution, and medicine.

1.1. The Sperm Phenotype

Variation in sperm form and function—the cellular phenotype—comes from three sources and their interactions: the male nuclear genotype, the male mitochondria, and the environment (**Figure 1**). This definition extends previous ones that consider genetic variation in sperm form and function between males (Pizzari & Parker 2009), herein called the genotype effect. Research into sperm biology has been organized roughly into these three main sources of variation (**Figure 1**). In this review, we do not cover variation in the sperm phenotype that might arise from variation in the genetic makeup of sperm within an ejaculate and any possible resulting differences in haploid gene expression (Parker & Begon 1993). Although this issue is very interesting, we focus on environmental effects.

1.2. Male Nuclear Genetic Effects on Sperm Phenotype

By suggesting that “sperm phenotypes are predominantly determined by testicular gene expression and hence the diploid genome of the male,” Pitnick et al. (2009, p. 75) implied that environmental sources are not important in explaining the sperm phenotype and sperm diversification. This summary reflects four decades of intense research on sperm competition (Bernasconi et al. 2004; Birkhead et al. 1998, 2009; Parker 1970). This view is also implicit in the literature on animal breeding as well as in procedures within medicine that seek correlations between sperm function and genotype (so-called sperm function tests) (Aitken 2006, World Health Organization 2010). Although sperm competition is a successful research field, a need to extend this view is apparent from the fact that male genotype effects explain only a small to moderate proportion of variation in sperm function (Dowling et al. 2010, Simmons et al. 2014). For example, crosses of six male and female *Drosophila melanogaster* genotypes were carried out under highly controlled laboratory conditions (Clark et al. 1999), but genotype explained only 6–11% of the variation in paternity.

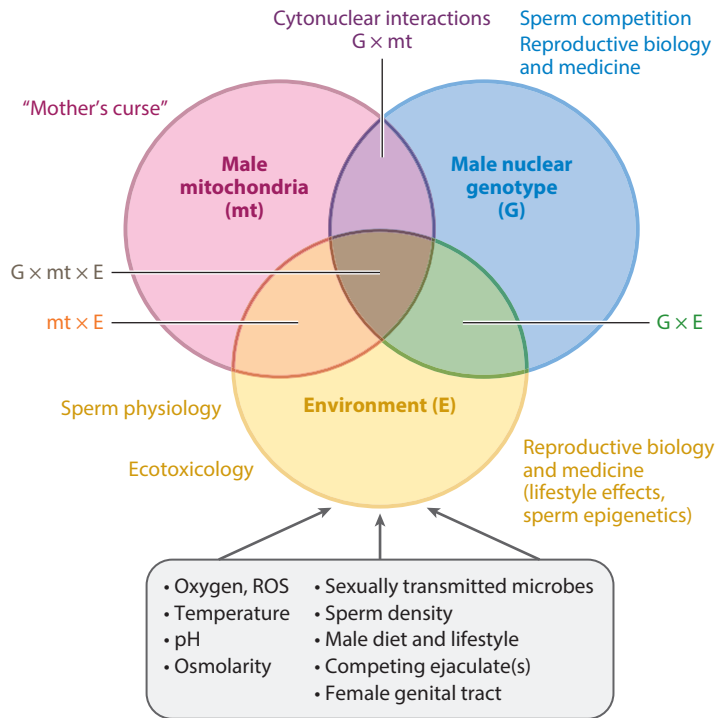


Figure 1

Three major influences on phenotypic sperm function and their corresponding research fields. The male nuclear genotypic contribution to sperm function is studied mainly in the context of sexual selection and male reproductive medicine (andrology) (*blue*). The male mitochondrial contribution to sperm function, and its interaction effects with the male nuclear genotype, is studied within evolutionary genetics in the context of “mother’s curse” and cytonuclear interactions (*magenta*). External environmental effects are studied in the context of sperm physiology, ecotoxicology, and reproductive medicine. Genotypically fixed aspects of competing ejaculates and female reproductive tracts are also important in studies of sperm competition. Sperm ecology considers environmental contributions to the sperm phenotype (*yellow*) and how they can lead to genotypic variation. Examples of major sperm environments are listed in the gray-shaded box (*bottom*). Abbreviation: ROS, reactive oxygen species.

Similarly, despite intense research efforts in reproductive medicine, approximately 10–15% of infertility cases are currently attributed to genetic factors in males (Pizzol et al. 2014).

1.3. Male Mitochondrial Effects on Sperm Phenotype

Mitochondria affect many aspects of the sperm phenotype (Aitken et al. 2009, Dowling et al. 2007, Froman & Kirby 2005, Innocenti et al. 2011, Yee et al. 2013, Zini & Al-Hathal 2011). In particular, mitochondrial genetic variation can have a substantial impact (but see Friberg & Dowling 2008): Variation in mitochondrial production of reactive oxygen species (ROS) explained 68% of the variation in sperm motility in humans (Koppers et al. 2008). Few studies, however, have experimentally manipulated the mitochondrial haplotype (but see Friberg & Dowling 2008, Yee et al. 2013) and, therefore, separate mitochondrial and mitochondrial-genotype (mt × G) effects on sperm function. Fruit fly sperm carrying mitochondrial haplotypes combined with a foreign

Reactive oxygen species (ROS):
a by-product of normal metabolism; these chemically reactive molecules can cause damage to membranes and DNA and also act as cellular signals

Sperm trait: a sperm character explained by male nuclear and mitochondrial genotype

Genotype-by-environment ($G \times E$) interaction: different genotypes respond differently to various environmental conditions

nuclear background have had, on average, a 30% lower sperm competitive ability than they do when expressed with their coevolved background (Yee et al. 2013).

Two aspects of mitochondrial effects on the sperm phenotype relate to sperm diversification. First, the exclusive maternal inheritance of mitochondria (in almost all species) reduces the possibility that sperm functions can evolve via sperm competition if mitochondria govern these sperm functions. Selective advantages may occur through local mitochondrial adaptations in females (Rand 2001). However, mutations with a negative effect on sperm function can accumulate if these mutations have only small, or positive, effects on females (Innocenti et al. 2011, Yee et al. 2013), a process known as “mother’s curse.” Second, molecular signaling from mitochondria to the nucleus, such as variable ROS production, can differ between environments (Murphy 2009, Wallace et al. 2011). This provides an opportunity for mitochondrial-environmental ($mt \times E$) interactions and perhaps for local adaptations of mitochondria (Dowling 2014, Wolff et al. 2014).

1.4. Environmental Effects on Sperm Phenotype

In the eighteenth century, Spallanzani observed that snow-chilled sperm recover their motility in warmer temperatures (Mann 1964). Mann (1964) added that the nineteenth century “abounds in studies on the effect of changes in the medium on sperm motility and survival” (p. 54). Despite this history, and in contrast to recent extensive research on genotype and mitochondrial effects, current evolutionary and ecological research has largely ignored environmental effects on the sperm phenotype and sperm diversification (but see Blanckenhorn & Hellriegel 2002; for ecotype effects on sperm cells in plants, see Delph et al. 1997). Instead, several currently unconnected research fields deal with environmental effects on the sperm phenotype, i.e., effects that go beyond a mere variation in sperm number: (a) the substantial medical literature of lifestyle effects on sperm function (Aitken et al. 2014, Fraga et al. 1996, Yauk et al. 2008); (b) the literature on fertilization biology in marine systems (Adriaenssens et al. 2012; Jensen et al. 2014; Levitan 1995, 2000; Schlegel et al. 2014); (c) ecotoxicological research on the effects of environmental pollutants and endocrine disruptors on sperm function across a wide range of taxa (Hayes 2011, Lewis & Ford 2012, Tavares et al. 2013); (d) applied research on storage, transport, and long-term cryostorage of sperm (Leahy & Gadella 2011, Mann 1964); and (e) sperm aging, which encompasses the successive or collective accumulation of damage across all the environments through which a sperm cell has passed (e.g., Pizzari et al. 2008, Reinhardt 2007, Siva-Jothy 2000, Tarin et al. 2000). Environmental effects on the sperm phenotype can also be deduced from the fact that intramale variation (Pitnick et al. 2009) and intraejaculate variation for sperm traits are abundantly reported. Finally, sperm epigenetics (offspring variation based on environmental alterations of sperm cells), just like sperm aging, describes collective and cumulative environmental effects, often without specifying the underlying molecular mechanism. This is an emerging field, but most effects concern epigenetic alteration at the spermatid stage (Dada et al. 2012, Jenkins & Carrell 2012, Johnson et al. 2011) rather than of mature sperm (but see Marshall 2015).

1.5. Introducing the Research Field of Sperm Ecology

By applying basic individual-level approaches, which have been successful in developing whole-organism ecology, sperm ecology aims to characterize interactions between sperm cells and their environment and to examine the consequences of this interaction. This aim requires a consideration of the nuclear genotypic, mitochondrial, and environmental components of the sperm phenotype and their interrelations, such as genotype-by-environment ($G \times E$) and mitochondria-by-environment ($mt \times E$) interaction effects (**Figure 1**). By using the concept of the sperm phenotype, sperm ecology extends existing research areas by combining the focus on additive genetic

effects in research on sperm competition (Simmons & Moore 2009) with the environmental effects that ecotoxicology, reproductive medicine (i.e., lifestyle effects on sperm function), and other fields outlined in Section 1 have identified. In addition, specifying the environments that sperm cells encounter in different female genotypes may provide a useful route to characterize the outcome of reproductive interactions (see, e.g., Aranha et al. 2008, Rosengrave et al. 2009, Yeung et al. 2006). The fact that sperm function in individual males is not always highly repeatable (Birkhead & Fletcher 1995, Garcia-Tomas et al. 2006, Peters et al. 2004; but see Gage et al. 2004) suggests a role for environmental effects in explaining fitness variation in nature.

Environmental effects on sperm may be apparent as a temporal variation in sperm function. In contrast to the concept of sperm competition, for which the evolutionary outcome is important (i.e., only the end points of competition), sperm ecology takes a longitudinal, cellular-lifetime approach (**Figure 2**, **Supplemental Figure 1**; for all **Supplemental Material**, follow the corresponding link from the Annual Reviews home page at <http://www.annualreviews.org>), yielding several advantages. First, sperm may be in competition for a variable amount of time; hence, temporal variation will help to predict the end points of sperm competition in these cases while simultaneously considering the universal cellular trade-off between energy expenditure and life span (**Figure 2**) (for various examples of cellular trade-offs in sperm, see Burness et al. 2004, Gage et al. 2004, Hughes & Davey 1969, Levitan 2000, Reinhardt & Otti 2012, Ribou & Reinhardt 2012).

Second, the longitudinal approach incorporates delayed environmental impacts on sperm (see below) including the view that differences in offspring phenotype or quality arise because fertilizing sperm have different exposure histories or durations (e.g., Burrue et al. 2013, Ghaleno et al. 2014, Immler et al. 2014, Lane et al. 2014, Marshall 2015). Third, a lifetime view allows one to consider sperm physiological changes as phenotypic plasticity at the cellular level. Despite Spallanzani's early observations, and despite the central position of phenotypic plasticity in whole-organism biology, the issue of sperm phenotypic plasticity has not been addressed by sperm competition and has been addressed only rarely in other areas of evolutionary and ecological research (Crean et al. 2013, Jensen et al. 2014, Poland et al. 2011, Purchase et al. 2010). We further argue that male and female reproductive traits evolve to accelerate sperm function via sperm phenotypic plasticity. This process is similar to niche construction in whole-organism ecology, and we refer to such created sperm environments as sperm niches.

Finally, by testing for adaptive $G \times E$ or $mt \times E$ interactions, sperm ecology aims to describe whether, and how, natural selection favors specific sperm phenotypes in specific environments. This approach may also include sexual selection if male and female genotypes are regarded as specific environments for sperm. Therefore, sperm ecology contributes to explaining sperm diversification. Importantly, sperm ecology does not necessarily require competition between genotypes to produce evolutionary changes (**Figure 2**) and, as such, is a parsimonious concept. In summary, by integrating sperm biology with four basic ecological concepts, e.g., environmental variation, phenotypic plasticity, niche construction, and $G \times E$ interactions (local adaptation), sperm ecology may contribute to explaining phenotypic adaptations as well as sperm diversification in three important ways: (a) characterizing and quantifying the effects of environmental variation on sperm function, (b) assessing the role of natural selection in sperm diversification, and (c) suggesting a pathway for the evolution of male and female reproductive traits.

2. CHARACTERIZING AND QUANTIFYING THE EFFECTS OF ENVIRONMENTAL VARIATION ON THE SPERM PHENOTYPE


Environmental variation can act on sperm in several ways. First, an external environment (such as temperature for ectothermic animals; water pressure, UV radiation, and salinity for

Phenotypic

plasticity: the ability of a genotype to vary phenotypically under different environmental conditions

Local adaptation:

a form of genotype-by-environment interaction where the fitness of a genotype is highest in the environment in which that genotype evolved

 **Supplemental Material**

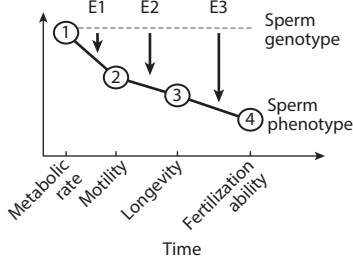
Sperm ecology

Environmental and temporal effects on sperm phenotype
Origin of, and diversification by, G x E interactions

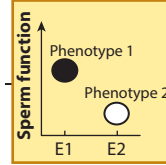
Unit of study



Overall E effect on sperm phenotype



E effect over time (e.g. fitness episodes 1 to 4)

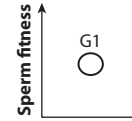


General observation

Sperm competition

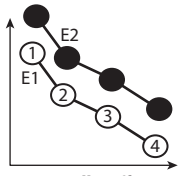
Sperm phenotype = male genotype
Diversification via G x G interactions

Total sperm fitness of sperm genotype considered; temporal and environmental variation not relevant

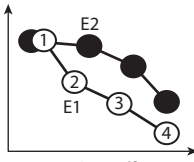


Origin of sperm phenotypic variation

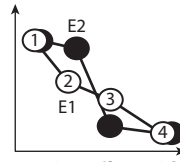
Environmental: Different environments cause different sperm functions, different temporal variation (polyandry or monogamy):



Overall E effect



E x time effect



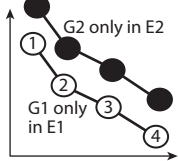
E x time effect, with trade-offs

Genotypic (mutations)

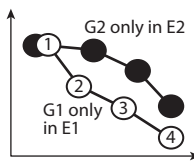


Evolution of sperm genotypes

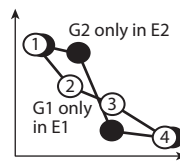
Divergent natural selection of different genotypes in different environments:



Overall E effect



E x time effect



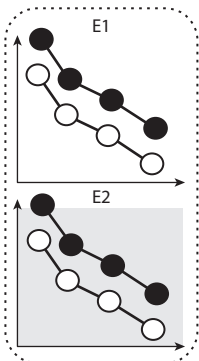
E x time effect, with trade-offs

Competitor sperm mutant favored under polyandry



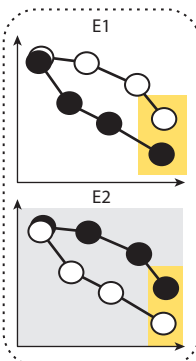
Maintenance of sperm genotypic diversity

Only environmental effect



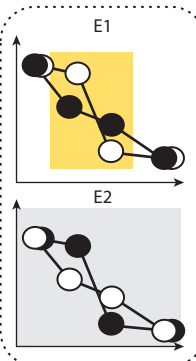
Diversity not maintained

G x E interaction effect on sperm phenotype



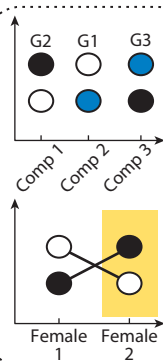
Diversity maintained

G x E x time interaction effect on sperm phenotype



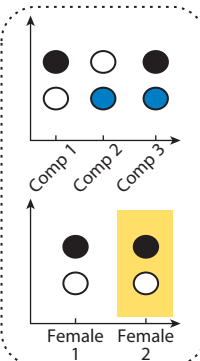
Diversity maintained

G x G interaction Sperm competition nontransitive



Diversity maintained

Only genotypic effect Sperm competition transitive (good genes)



Diversity not maintained

broadcast-spawning organisms; or in vitro treatments for laboratory cases) directly impacts sperm cells. Second, environments can act on males and females and generate microenvironments for sperm cells that are different from the external environment (**Supplemental Figure 1**). For example, smoking results in increased systemic levels of ROS, including in reproductive compartments; food items may affect the pH in seminal fluid; or increased temperature may alter ion or enzyme concentrations. Third, longitudinal variation in sperm function may vary across different environments, such as reduced metabolism under hypoxia.

▶ **Supplemental Material**

We use two literature search methods to describe environmental effects on sperm: (a) a random selection of relevant articles and (b) a directed search by specific environments on specific sperm functions (see **Supplemental Table 2**). Our literature search on environmental effects on sperm yielded 27,514 articles, or 8,042 if restricted to articles published in 2000–2014. Of the latter, 900 articles were randomly chosen, of which 178 articles (19.8%) (**Supplemental Table 1**) matched our criteria (**Supplemental Table 2**). These articles provide an estimate of the significance of environmental effects on sperm within ecology and evolution.

2.1. Many Environmental Factors Affect Sperm Function

Phenotypic sperm function is affected by various environments, including temperature, pH, osmolarity and concentration of specific ions, oxygen concentration, oxygen radicals and antioxidants, diet (male, maternal, and paternal as well as its amount and composition), larval or adult population density, photoperiod, UV radiation, sexually transmitted microbes, viruses, exposure to airborne or food-borne chemicals (male, maternal, and paternal), external nucleic acids, and sperm density (**Supplemental Table 2**). The data may be biased, as results that show no environmental effect on sperm were published less frequently (but for a discussion of the substantial number of studies reporting the absence of environmental effects, see Section 2.2). Yet, it seems most environments tested show some effect on sperm.

2.2. The Literature on Environmental Effects on Sperm Phenotype Is Large and Largely Neglected by Ecology and Evolution Research

After applying our search criteria (**Supplemental Table 2**), accounting for the fact that different environments were studied to a varying extent (**Supplemental Table 1**) and excluding 597 articles that appeared under more than one environment (e.g., two environments examined by one paper), we suggest that between the years 2000 and 2014 an estimated 1,293–2,180 (mean = 1,736) articles explored environmental effects on sperm (**Supplemental Table 1**). The 178 articles we studied in detail examined a total of 458 environment–sperm function combinations (2.57 per study). Of these, 356 combinations (78%) reported in 163 studies (91.6%) showed at least one environmental effect on at least one function, whereas 64 studies (35.9%) showed no effect on at least one function. Projecting to the other articles, we find the abundance of environmental

Figure 2

Origin, evolution, and maintenance of genetic variation in sperm function as envisaged by the concept of sperm ecology presented here and compared with the concept of sperm competition. Sperm ecology considers temporal variation in sperm function, i.e., over various fitness episodes, noting how different environments increase deviation in sperm phenotype from the sperm genotype. This allows for the consideration of delayed environmental effects on sperm (for details, see **Figure 3**) and cellular trade-offs. Natural selection favors certain sperm phenotypes in some environments, without requiring a sperm competitive advantage. $G \times E$ interactions increase sperm function diversity (for details, see **Figure 4**). Abbreviations: Comp, competition; E, environment; G, male nuclear genotype.

effects is remarkable: Of the projected 1,736 (1,293–2,180) relevant articles, 1,590 articles may have explored 4,086 environment–sperm function combinations, with 3,187 environment–sperm function combinations potentially showing environmental effects on sperm function.

Although the assertion that environment shapes the sperm phenotype (as proposed in **Figure 1**) may not seem surprising, it is noteworthy that only very few of these articles originate from evolution and ecology research. Broadly defining “ecology and evolution journals” (**Supplemental Table 2**), only 8 (4.5%) articles from our random search concerned ecology and evolution. A similar number resulted from our directed search: only 40 (5.4%) out of 7,445 articles, including 18 that examined consequences of predicted global change on sperm function, were found.

Supplemental Material

2.3. Magnitude and Shape of Environmental Effects on Sperm Phenotype

Our summary (**Supplemental Table 2**) revealed that some environments such as brief high altitude visits (Okumura et al. 2003), brief pollution events, and brief temperature elevations (Paul et al. 2008) affect the sperm phenotype after even a short transient impact, whereas others were found after a sustained period of action. Some environmental effects became apparent immediately, whereas others, including those found in offspring, appeared much longer after the environmental impact. Many environmental effects, such as DNA damage, membrane damage, and sperm mortality, are irreversible and hence permanent at the cellular level or in males. However, examples also show that other environmental effects are reversible at those levels (Aitken et al. 2012, Bencic et al. 2000, Le Comber et al. 2004, Okumura et al. 2003, Villegas et al. 2003).

Among the fitness-related aspects of sperm function, none appeared so canalized as to be consistently inert to environmental effects. Across species, sperm morphology, metabolism, motility, longevity, fertilization ability, epigenetic signatures on the nuclear genome, and offspring health were all affected. Within species, these characters were not equally affected, and sometimes the effects were not positively correlated with each other (**Supplemental Table 2**). The magnitude of effects was so variable as to prevent any generalization. Compared with controls, environmental effects on sperm populations or sperm cells ranged from no, or minute, effects to substantial reductions in sperm function, including complete failures. Even natural variation in environmental conditions (such as temperatures $>37^{\circ}\text{C}$ or changes in pH or osmolality) generated substantial variation in sperm function (**Supplemental Table 2**).

2.4. Phenotypic Plasticity in Sperm Function

Almost all studies incorporated in our literature search assessed environmental effects against control sperm from the same male, the same genotype, or the same population. In other words, almost all 397 environment–sperm function combinations represent phenotypic plasticity at either the genotype or male population level. Some studies even demonstrated plasticity at the level of the individual sperm cell or the ejaculate, for example, by showing that in vitro effects were reversible (Le Comber et al. 2004, Otti et al. 2013) or that human sperm repeatedly bind and unbind to the epithelium (Pacey et al. 1995) or move in and out of a hyperactivated state (Mortimer & Swann 1995). It may be noteworthy that even sperm morphology can be plastically (but not necessarily reversibly) affected by the environment. For example, compared with ejaculated sperm, the female sperm storage organ of several insect species shows substantial membrane alteration (Renieri & Vegni Talluri 1974, Riemann & Thorson 1971; for a review of phenotypic plastic, environmental effects on sperm size, also see Marshall 2015).

Physiological responses in sperm, equivalent to cellular phenotypic plasticity, are not unexpected, given the diverse chemistry of male and female genital tracts that sperm cells have to

master. However, explicitly spelling out the existence of sperm phenotypic plasticity may help to formalize predictions of when such plasticity would be adaptive. Adaptive plasticity depends on how often a given environment is encountered, the duration of the encounter, and the reliability of information (for a general framework, see Pfennig et al. 2010). Interestingly, these predictor variables of sperm phenotypic plasticity can now be linked to the substantial body of cell biology studies that examine the considerable ability of individual sperm cells to respond to chemical, surface, or other conditions encountered in situ (Alvarez et al. 2012, 2014; Babcock et al. 2014; Bahat & Eisenbach 2006; Friedrich & Jülicher 2007).

2.5. The Relative Size of Genotypic, Mitochondrial, and Environmental Effects on Sperm Phenotype

Surprisingly few studies have estimated the relative size of genotype and environmental effects simultaneously in the same system (and possibly none have separated genotype, mitochondrial, and environmental effects). One study used female gene expression after sperm receipt as a parameter of sperm fitness and varied sperm age (environmental effect) within three different sperm genotypes (populations) (Otti et al. 2015). Approximately 16 times as many female genes were differentially expressed in response to environmental effects (79 genes) as opposed to genotype effects (5 genes). When this comparison was restricted to genes with substantial differences, 5 times as many genes were still expressed in response to environmental than to genotype effects (Otti et al. 2015). However, even though quantifying the number of differentially expressed genes in females may be useful, this number does not necessarily translate directly into differences in sperm fitness. The statistical table in the *Drosophila melanogaster* study by Clark et al. (1999) suggests less than 1% of the variation in paternity was explained by environmental (laboratory) effects, compared with 6–11% explained by male genotype effects. Both studies have the limitation that they include seminal fluid effects. More closely related to sperm function are the careful analyses by Purchase & Moreau (2012) and Purchase et al. (2010) on sperm swimming speed in fish across a pH and temperature gradient. These studies showed that genotype explained 3 times as much variance as did pH, whereas temperature explained 1.3 times as much variance as did genotype.

We conclude that environmental effects on sperm function are ubiquitous, take many forms, and may be as large as genotype effects, or even larger. Given our randomized literature search, it is not tenable to assume that male genotype effects almost exclusively shape the sperm phenotype. Environmental effects can be direct, indirect, or phenotypically plastic. They can be caused by sustained action or brief impact and are permanent or reversible. Evolutionary and ecological research should not ignore environmental effects when examining variation in reproductive success.

3. CONSEQUENCES OF ENVIRONMENTAL EFFECTS ON SPERM PHENOTYPE

3.1. Reduced Significance and Hampered Predictability of Sperm Competition

The observations that sperm functions vary between environments and that sperm can accumulate damage and information during their passage through an environment have important consequences. First and foremost, in many species the sperm genotype can occupy a very large phenotypic space (**Figure 3**). This fact severely hampers our ability to predict sperm function on the basis of genotype alone, in both the absence and the presence of sperm competition. Whenever sperm of two males compete, their sperm phenotypes have “stored” an environmental component, and this history may be decisive in their competition, even if both compete in the same environment.

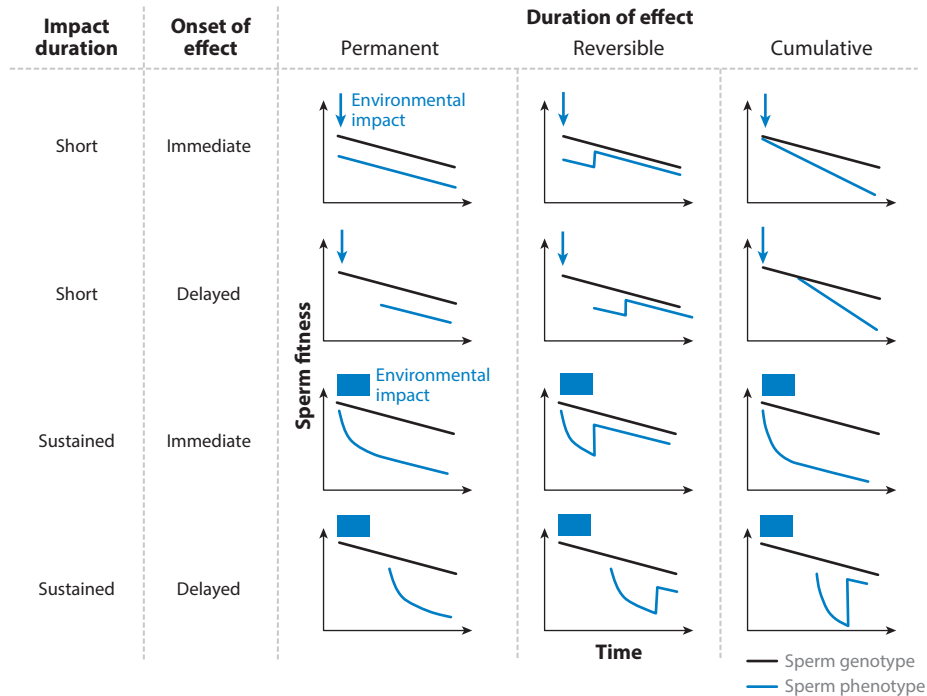


Figure 3

Summary of environmental impacts on the sperm phenotype (*blue lines*) generating deviations from the sperm genotype (*black lines*). The sperm genotype contains an a priori aging component, hence a decline over time. Impacts can directly affect the sperm cell and act immediately or in a delayed fashion, or they can indirectly affect the male, in which case they are inherently delayed. The impact can be a short burst or sustained, leading to permanent, reversible, or cumulative effects. In sum, the sperm phenotype can occupy almost any value under the sperm genotypic curve.

Supplemental Material

Importantly, this history will not give consistent differences between two competing males unless the sperm function follows a linear decline over time (cf. Reinhardt 2007, figure 1). Therefore, the loaded raffle in sperm competition (Parker 2009) cannot be expressed by a loading coefficient that is independent of time and environment. In nature, there will hardly ever be two males with identical lifestyles, habitat utilization, or age at the time of mating (when their sperm compete); hence, they will not have sperm with an identical environmental component (**Supplemental Figure 1**). We therefore predict that such carryover environmental effects on sperm competition or fertility are a universal feature in the animal kingdom. However, its testing is severely hampered by the paucity of studies addressing the impact of environmental effects on postejaculatory reproductive success (Almbro et al. 2011, Breckels & Neff 2014, Gasparini & Evans 2014, Mehliis & Bakker 2014, Vasudeva et al. 2014).

3.2. Sperm Function May Determine Species Range

In certain cases, sperm velocity was not affected by environmental variation, such as increased temperature and decreased water pH in several marine invertebrates (Byrne et al. 2010), the Atlantic cod (Frommel et al. 2010), or the oyster (Havenhand & Schlegel 2009). But in other

cases, males within a population were affected differently by environmental variation such that consistent variation between populations did not emerge. Examples of the latter include responses of sperm motility to ocean acidification in a polychaete (Schlegel et al. 2014) and a sea urchin (Schlegel et al. 2012). These examples show that sperm phenotypic plasticity (in males or at the population level) can buffer environmental variation and enable population persistence.

However, there are also dozens of studies showing that some sperm functions have optima at certain intermediate states that are close to current environmental conditions. In these cases, environmental effects on sperm function may limit a population's range or its ability to cope with altered environmental conditions. This has been suggested for some species under predicted global climate change: For example, reduced sperm velocity is expected under higher UV radiation doses [stickleback (Rick et al. 2014), sea urchins (Lu & Wu 2005, Nahon et al. 2009)], increased CO₂ concentration in the water [sea star (Uthicke et al. 2013), mussels (Vihtakari et al. 2013), oyster (Barros et al. 2013), coral and sea cucumber (Morita et al. 2010)], and increased water temperature [guppy (Breckels & Neff 2013)]. Sperm longevity may be reduced under increased water temperatures [sea urchin (Binet & Doyle 2013)] or may undergo altered trade-offs with sperm velocity [sea urchin (Caldwell et al. 2011), stickleback (Mehlis & Bakker 2014)]. Similar effects may occur under decreased water pH [sea urchin (Caldwell et al. 2011)]. Acclimatization by males to higher temperatures, possibly representing indirect environmental effects, may not shift thermal critical limits of sperm velocity (Adriaenssens et al. 2012).

3.3. Intraejaculate Heterogeneity

As males pass through different environments but continue to produce sperm, ejaculates become heterogeneous in terms of sperm age (Reinhardt & Siva-Jothy 2005), epigenetic marks (Aoki et al. 2006), and other characters (Dorado et al. 2013, Satake et al. 2006). Immler et al. (2014) found that sperm from the same ejaculate produce different offspring phenotypes when sperm are exposed to different treatments. An important consequence is that different environments may cause different patterns of ejaculate heterogeneity and so contribute to phenotypic and genetic divergence of populations that live in different habitats.

3.4. The Brynhild Effect

Some sexual selection models have suggested that females benefit from creating barriers to sperm that only the best sperm can pass to fertilize eggs (Birkhead et al. 1993). The observation that sperm cells experience, or even accumulate, environmental damage is not entirely consistent with those models. Instead, sperm ecology predicts the adaptive evolution of filter mechanisms against damaged sperm regardless of their genotype (see also Siva-Jothy 2000, Reinhardt 2007). Additionally, stronger female barriers that represent harsh environments for sperm are predicted to cause greater damage to sperm. Termed the Brynhild effect, this situation is similar to that of the eponymous female character in the Nordic epic saga *Nibelungenlied* (Song of the Nibelungs) who resided inside a ring of fire. Noble men aiming to get across to marry Brynhild either died or were injured. The man who finally succeeded needed magical powers to cross the fire.

3.5. Sperm Viability Is Not a Good Fitness Indicator

Some environments may be so stressful that sperm apoptosis is initiated, for example, during attacks by retroviruses that inject foreign RNA or DNA into sperm DNA. Aitken & Baker (2013) pointed out that apoptosis may then be selectively advantageous: "Selective deletion of damaged germ cells

is clearly a critical component of the mechanisms used to safeguard the genome of a given species” (p. 265). Though expressed in a group selectionist way, this remark illustrates how sperm viability or apoptotic activity per ejaculate is not necessarily a sign of low male quality but may be adaptive for a male if apoptosis prevents damaged sperm that would result in lower-fitness offspring from outcompeting his own genetically undamaged sperm (Aitken & Koppers 2011, Aitken et al. 2013). Aitken & Baker (2013) also highlighted another benefit of apoptosis: “By engaging in regulated cell death exhibiting many features of apoptosis, moribund spermatozoa ensure they can be efficiently removed from the male or female reproductive tract without provoking a damaging inflammatory response” (p. 266).

As a consequence, the widely used proportion of dead sperm per ejaculate (or proportion of live sperm/sperm viability) may be an indicator of the environmental history of a male or a positive indicator of the ability of a genotype to respond to its environment ($G \times E$), rather than exclusively reflecting a negative genotype.

3.6. Variance Effects in Numerical Sperm Competition

Sperm competition predicts a numerical advantage for males delivering more sperm. Because of the environmental component of the sperm phenotype, sperm ecology also specifies this prediction of male advantage. Continuous, and constant, sperm production automatically results in ejaculates with more sperm that also contain more recently produced sperm, i.e., sperm that have been exposed to an environment for shorter periods (Reinhardt 2007). The general numerical advantage seen in sperm competition may therefore be due to the fact that in larger ejaculates more sperm are present in the fresh cohort.

3.7. Mean Ejaculate Traits May Be Noninformative

In most species, only a few sperm reach an egg. As selection acts to maximize sperm functions, mean ejaculate values of sperm motility or longevity may be less informative in predicting paternity than are some maximum values (Holt & van Look 2004, Mossman et al. 2010, Reinhardt & Otti 2012). We suggest that current medical diagnostics of infertility (World Health Organ. 2011) may benefit from considering this notion.

3.8. Trade-Offs in Sperm Function Can Hamper Comparability

Sperm function can decline within seconds of activation (see examples in Levitan 1995, Purchase et al. 2010, Reinhardt & Otti 2012). Trade-offs in sperm function associated with such rapid decline can severely hamper the comparison of individuals and lead to false conclusions (as illustrated in Reinhardt & Otti 2012, figure 2).

3.9. Adaptive Habitat Choice

Given the environmental effects on sperm function and that altered sperm function translates into reproductive success, we predict that males and females are under selection to choose specific environmental conditions that positively affect sperm function.

3.10. Selection for Sperm Niches

Alternatively to adaptive habitat choice, environment-dependent sperm function can select for male and female traits that create environments in which sperm function is improved. These

so-called sperm niches may, for example, allow an organism to colonize new habitats. This has been suggested by Elofsson et al. (2003), who argue that ovarian fluid chemistry allowed sticklebacks to overcome the osmotic constraints on sperm imposed by a new freshwater habitat. The most obvious sperm niche is seminal fluid, an evolutionarily diverse character (Avila et al. 2011, Poiani 2006). Seminal fluid fulfills niche functions by buffering the pH for sperm, reducing oxidative stress to sperm, supporting motility, extending longevity, and improving offspring development across a variety of taxa (e.g., Aitken & Clarkson 1988, Bromfield et al. 2014, Heise et al. 2010, Kang et al. 2008, Rickard et al. 2014, Scaggiante et al. 1999, Shaliutina-Kolesova et al. 2014). Specific examples include seminal antioxidants (Avila et al. 2011, Poiani 2006) or antimicrobial properties (Otti et al. 2009, 2013; Poiani 2006). Other niches may transiently reduce sperm motility and save cellular energy resources by containing sperm in bundles, via spermatophores, or through additional sperm types (Reinhardt 2007).

Many female traits also serve as niches by plastically improving sperm function. Reinhardt (2007), Holt & Lloyd (2010), and Heifetz & Rivlin (2010) review female traits that reduce sperm metabolism and sperm oxidative stress, thereby extending sperm longevity during prefertilization storage. These traits include hypothermia at the sites of sperm storage, reduction of sperm motility by binding sperm to epithelia, as well as packing sperm tightly or organizing them in bundles. Sperm niche functions are also found in immunological and antioxidant protection, interference with sperm metabolism, reduction in the number or size of mitochondria, or delays of sperm activation. Recent work on insects suggests that the process by which sperm metabolism is reduced is adaptive to females (i.e., it delays infertility) (Reinhardt & Ribou 2013, Ribou & Reinhardt 2012). Both Reinhardt & Ribou (2013) and Ribou & Reinhardt (2012) supported the idea that this effect is specifically directed toward sperm phenotypes, and not sperm genotypes, because the sperm genotype could not be predicted on the basis of sperm metabolism.

3.11. Selection May Be Directed Against Environmental, not Genotypic, Components of the Sperm Phenotype

If environmental effects on sperm are often damaging, then male and female traits that discriminate against sperm on the basis of the sperm phenotype unrelated to the sperm genotype are predicted to evolve. In males, such traits include those that specifically disfavor aged sperm phenotypes, for example, repeated mating with the same female (though functional for sexual selection, mating with different females also gives rise to this trait), sperm transfer to other males, sperm discard without copulation, continuous sperm production, and the many ways of bringing reproductive events forward in time, i.e., closer to sperm production (reviewed in Reinhardt 2007). In females, repeated mating to the same male reduces representation of aged or environmentally damaged sperm phenotypes in the fertilization set. Sperm dumping by females also has this effect if it is related to sperm storage time. If sperm stratify in males by age or quality cohorts, the behavior of mate copying by females would automatically increase the representation of higher-quality sperm (reviewed in Reinhardt 2007).

However, though these traits automatically alter ejaculate variability in terms of environmental effects, relatively few empirical tests exist. For example, in a cricket species, nonused sperm were expelled by males while younger sperm were more successful in reaching the female sperm storage organ (Reinhardt & Siva-Jothy 2005). In bedbugs, Otti et al. (2013) demonstrated that antibiotic activity in the seminal fluid transferred during one mating was sufficient to reduce sperm mortality caused by simultaneously transferred bacteria. Finally, because experimental scrotum insulation results in reduced sperm motility or DNA fragmentation (Banks et al. 2005, Brito et al. 2003), we may conclude that the scrotum evolved to reduce environmental effects on sperm.

In summary, environmental effects on the sperm phenotype yield substantial conceptual and methodological consequences for ecological and evolutionary research. Even though some predicted consequences remain untested, they substantially alter our understanding of variation in fitness and reproductive success. We suggest that the consideration of environmental effects on fitness is a worthwhile scientific enterprise.

4. NATURAL SELECTION AND DIVERSIFICATION IN SPERM PHENOTYPE

4.1. A Model for Phenotypic Sperm Evolution

Sperm phenotypic diversification is possible along two principal paths: via sperm environments enabling sperm phenotypic plasticity and via sperm traits. Both ways can lead to adaptive population genetic changes and contribute to evolutionary change. Selection can operate on increased male reproductive success by directly favoring sperm traits (**Figure 4**). Sperm phenotypic plasticity opens a second route for selection: Male traits may be favored that create sperm niches in which phenotypically plastic sperm function better than they would in the absence of such niches (**Figure 4**), making niches adaptive paternal effects. An obvious example of the evolution of a sperm niche is seminal fluid. Sperm cells experiencing a newly evolved sperm niche may then allow further sperm diversification by providing an arena in which novel sperm traits evolve (**Figure 4**) [cf. genetic assimilation (West-Eberhard 2003)].

Alternatively, sperm niches may reduce the opportunity for sperm evolution because sperm experience a stable environment without selection pressure. An imaginary example is a sperm phenotype that has benefitted through a male mutation that led to increased sugar content in the seminal fluid (**Figure 4**). This sperm phenotype may now benefit from a male mutation that

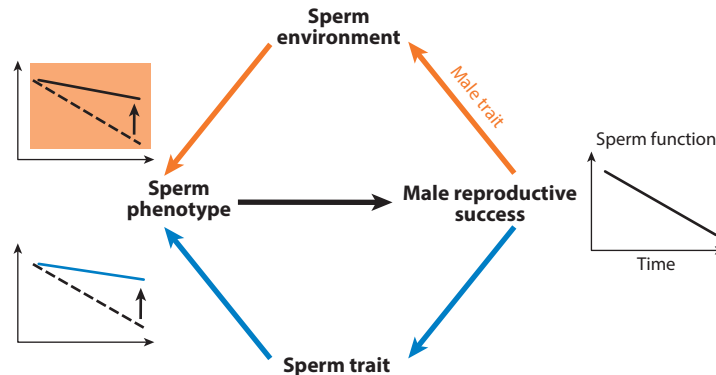


Figure 4

Model of sperm phenotypic variation leading to genotypic variation. Increased male reproductive success (with or without competition) may be generated if male traits are favored that directly code for sperm traits (*blue arrows*). Alternatively, male traits are favored that indirectly affect sperm function by creating environments that improve sperm function, but without altering sperm traits (*orange arrows*). Under this simple model, sperm environments can either shield sperm from external influences or generate niches that allow an increase in phenotypic space. These new sperm environments may then canalize specific phenotypes (*blue arrows*) and allow for selection of new genotypes or further extension of the sperm niche. The most obvious candidate for this model is the evolution of seminal fluid, but it also accommodates male-female coevolution.


generates, say, thermal stability in the sperm environment (**Figure 4**) and so optimizes energy expenditure (thus, no change in sperm traits). Alternatively, this sperm phenotype may benefit from a mutation that increases the permeability of the sperm membrane for sugars (**Figure 4**). These kinds of successive changes can explain environment-mediated sperm diversification if the male environment changes (e.g., generally increased sugar availability to the male), but they do not require such changes (e.g., mutations that increase male sugar uptake or allocation to seminal fluid).

Although this verbal model is exceedingly simple, we propose it may, for example, explain the evolution of species-specific ion channels or surface proteins in sperm in response to altered internal conditions (Lishko et al. 2012) as well as the evolution of seminal fluid complexity (Avila et al. 2011, Poiani 2006). Alternatively, it may provide a plausible mechanism for examples of ecological speciation in which divergence in diet specialization translated into postmating reproductive isolation (Nosil 2012). This model may also help to explain how males may adaptively alter the fertilization ability of their sperm such that sperm, or ejaculates, function best under their paternal environments (Marshall 2015). And it can incorporate coevolutionary interactions such that male traits that induce females to create sperm niches evolve (such as male seminal substances that manipulate the sperm storage ability of females; see Avila et al. 2011). Whereas genotype effects encompass the sperm-activating ability by seminal fluids of other males, even of another species (den Boer et al. 2010, Morrell et al. 2014, Usinger 1966), our model can generate coevolution-like dynamics between sperm and males, where males evolve niches to harness their own sperm.

4.2. Genotype-by-Environment Interaction Effects on Sperm Phenotype

In general, $G \times E$ interaction effects on the sperm phenotype have rarely been addressed. For example, of the 178 articles in our literature research that contained relevant data, 6 articles (3%) presented the data separately for two, or more, breeds or separate populations (broadly equivalent to genotypes). This suggests that very few of the thousands of articles reporting effects on sperm cells caused by diet, parental smoking habits, caffeine consumption, pH or temperature, salinity, antioxidants or their addition or blocking, vaginal lubricants, traditional medicinal herbs, etc., examined the generality of the effect beyond one genotype. None of these studies reported a formal $G \times E$ analysis.

We have extended our literature search in a second step and assessed the title and abstracts of all 7,445 articles without search terms (manually) for evidence of $G \times E$ interactions and of local adaptation. This resulted in a number of additional articles. Whereas no significant $G \times E$ interaction effect was reported for the sperm motility decline in two trout populations over decreasing pH (Purchase & Moreau 2012) or in two very different environments for sperm motility of Atlantic cod (Beirao et al. 2014), most other studies seem to indicate the presence of $G \times E$ interaction effects (**Table 1**). The common existence of species-specific maxima of sperm motility at different temperatures, osmolarities, or pH values that correlate to habitat conditions (**Supplemental Table 2**) (Alavi & Cosson 2005) suggests that local adaptation in sperm is common.

 [Supplemental Material](#)

5. CONCLUSIONS

$G \times E$ interaction effects are likely relatively common. However, their adaptive significance and magnitude await further quantification, especially in the light of widespread sperm phenotypic plasticity. This is a major task for sperm ecology.

The coevolution-like dynamics between sperm and internal environments of males or females have evolutionary and medical consequences that can be summarized in one sentence: Do not take sperm out of context. The lack of a coevolved sperm niche for sperm tested in medical sperm

Table 1 Examples of studies examining genotype-by-environment ($G \times E$) interaction effects or adaptive phenotypic plasticity on the sperm phenotype

Type of evidence	Level of comparison	Group	Sperm function and environments examined	Reference(s)
$G \times E$	Interspecific	Sea urchin	Fertilization ability, but not motility, of sperm decreased with increasing CO ₂ water concentration in the water in only one species.	Sung et al. 2014
$G \times E$	Interspecific	Sea urchin	Species differed in their sperm motility response to temperature.	Rahman et al. 2009
$G \times E$	Interspecific	Bird	Species differed in their sperm motility response to cooling protocols.	Blanco et al. 2000
$G \times E$	Interspecific	Cichlid fish	Consistent differences in sperm motility pattern existed between mouthbrooding and externally spawning species.	Reinhardt & Otti 2012
$G \times E$	Single nucleotide polymorphism	Ram	Temperature treatment increased DNA fragmentation in sperm only in bearers of one allele but not another.	Ramon et al. 2014
$G \times E$	Karyotype	Mice	After irradiation, sperm defragmentation was larger in Y-bearing than in X-bearing sperm, leading to reduced egg-binding ability.	Kumar et al. 2013
$G \times E$	Population	Cattle	Altitude and season produced changes in motility that differed between two breeds of cattle.	Barros et al. 2006, Chacur et al. 2013
$G \times E$	Population	Chicken	Sperm of four chicken breeds differed in their susceptibility to different freezing methods.	Schramm 2008
$G \times E$, phenotypic plasticity	Population	Guppies	Populations responded to experimental evolution under altered temperatures with an increase in sperm length, but not sperm motility. Populations also displayed phenotypic plasticity in sperm length and sperm motility.	Breckels & Neff 2014
$G \times E$	Genotype	Humans	Exposure of males to certain environments caused altered morphology or chromatin integrity in only some male genotypes (review).	Axelsson et al. 2010
$G \times E$	Genotype	Fruit fly	Larval-rearing density affected the sperm length of only some genotypes.	Morrow et al. 2008
$G \times E$	Isogenic line	Fruit fly	Laboratory \times genotype interaction effects accounted for 12–19% of the variation in paternity.	Clark et al. 1999
$G \times E$	Genotype	Flour beetle	Different genotypes varied in their sperm defense ability (P1) in relation to nutritional manipulation.	Lewis et al. 2012
$G \times E$	Genotype	Honeybee	Colonies showed an age \times genotype interaction effect on sperm viability.	Stürup et al. 2013
$G \times E$	Genotype	Cod	Significant, but likely small, $G \times E$ interaction effect was found on sperm velocity.	Purchase et al. 2010
$G \times E$	Genotype	Bedbug	Twice as many female genes were differentially expressed in response to $G \times E$ interaction effects as opposed to genotype effects.	Otti et al. 2015

(Continued)

Table 1 (Continued)

Type of evidence	Level of comparison	Group	Sperm function and environments examined	Reference(s)
Local adaptation	Interspecific	Sea urchin	Sperm motility, metabolism, and temperature-dependent motility differed across four closely related species such that sperm function was highest under conditions resembling the native habitat.	Mita et al. 2002
Local adaptation	Interspecific	Fish	Sperm motility, metabolism, and temperature-dependent motility differed across four closely related species such that sperm function was highest under conditions resembling the native habitat.	Lindroth 1947
Local adaptation	Population	Fruit fly	Local adaptation in male fertility to temperature was found.	Rohmer et al. 2004
Local adaptation	Population	Cichlid fish	Populations from two different habitats differed in their activation threshold of sperm motility, and the threshold reflected the ionic concentrations in these habitats.	Morita et al. 2010
Local adaptation	Population	Stickleback fish	Sperm from males of a saltwater population were motile in saltwater, but sperm from freshwater or brackish water populations were not.	Elofsson et al. 2003
Local adaptation	Individual	Bees	Ejaculates contained much more live sperm when males were reared at natural temperatures in a hive versus when reared at lower or higher temperatures.	Bienkowska et al. 2011
Adaptive phenotypic plasticity?	Individual	Bedbug, cricket	Adaptive variation in sperm metabolic rates was found between male and female sperm stores.	Reinhardt & Ribou 2013, Ribou & Reinhardt 2012
Adaptive phenotypic plasticity	Individual	Sea squirt	Fertilization ability of sperm varied adaptively with population density.	Crean & Marshall 2008, Crean et al. 2013
Adaptive phenotypic plasticity	Individual	Tubeworm	Sperm kept under low salinity produced offspring that survived better under low salinity.	Ritchie & Marshall 2013

function tests may provide an explanation for the poor predictive power of sperm function tests for conception and paternity (Aitken 2006; but see Birkhead et al. 1999, Froman & Feltmann 1998). Also consistent with this assertion, the predictive power of sperm function tests would likely be further improved if sperm function were measured after sperm have had contact with the female reproductive tract (Glazener et al. 2000; also Holman & Snook 2008).

SUMMARY POINTS

1. Sperm cells have been suggested to be the morphologically most diverse cell type to have evolved via sperm competition. Here we add that sperm cells show substantial phenotypic diversity caused by environmental effects.
2. Direct and indirect environments can act immediately or in a delayed way and shape sperm function in a decisive manner.

3. Male and female environments (indirect sperm environmental effect) can differ between or—as a result of ecological specialization—within populations. In this way, natural selection may substantially contribute to the evolution of sperm diversity via local adaptation of sperm.
4. Regardless of whether postejaculatory sexual selection, i.e., sperm competition or female sperm choice, augments the explanation of sperm diversification, sexual selection will benefit from the consideration of environmental influences on sperm.
5. Applying ecological concepts may help formalize descriptions of sperm biology. Such application may also help to identify the (currently largely lacking) mechanistic basis of competition between sperm.

FUTURE ISSUES

1. What is the relative significance of sperm phenotype evolution via sperm niches (environmental effects) and via sperm traits (genotype effects)?
2. How frequently do reproductive traits evolve that act on sperm phenotypic variation that is not related to sperm genotypic characters?
3. To what extent is the outcome of sperm competition between two males repeatable across environments?
4. Are large environmental effects on the sperm phenotype the reason why sperm competition ability has low heritability (Simmons & Moore 2009), and are they the reason why genotype explains only a low proportion of reproductive success (Pischedda & Rice 2012)?
5. What are the extent and mechanisms of sperm epigenetic alterations and their effect on offspring characters?
6. Is there support for Marshall's (2015) idea that external fertilizers are more likely to show adaptive sperm phenotypic plasticity than are internal fertilizers?

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