

Overview

Special issue on *Drosophila* spermatogenesis

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Drosophila melanogaster sperm are nearly 50 times longer than human sperm.¹ Their nuclei are exceptionally compact and have a distinctive needle-like morphology.² They develop in syncytial cysts where they are interconnected via intercellular bridges along their whole length, rather than being linked at a single point near their nuclei.³ Once transferred to the female, they are stored for ~2 weeks (i.e., for a whole generation time), rather than for days, as occurs in most mammals.⁴ These superficial differences obscure the fundamental similarity of spermatogenesis across distant phyla. Indeed, cellular processes and regulatory mechanisms discovered in *Drosophila* have proved to be conserved in other organisms, including humans.

Studies of *Drosophila* spermatogenesis have revealed important concepts and genes for each aspect of sperm behavior and function, from testis development through sperm storage and competition. These studies have taken advantage of comprehensive collections of mutants^{5,6} and RNAi-generating stocks,⁶⁻⁸ methods to drive knockdown or ectopic expression specifically in the germline or in their supporting cells⁹ or to create, within a testis, clones of cells of one genotype surrounded by cells of another (for example, see ref. 10); and markers for different cell types, stages and subcellular structures of spermatogenic cells (for example, see ref. 11). This special issue of *Spermatogenesis* reviews the latest findings from studies of *Drosophila* spermatogenesis, including cellular and molecular events that shape the testis and that drive male germ cells through a complex differentiation process to produce motile, fertile sperm.

Spermatogenesis proceeds in a gonad that is made of somatic cells, and to which primordial germ cells migrate. Interactions between the somatic gonad and the germ line are crucial for successful spermatogenesis. Many events in gonadogenesis, and some of the genes that underlie them, are conserved (see this issue's article by Whitworth et al.¹²). Moreover, in addition to contributing to our understanding of spermatogenesis, studies of *Drosophila* testis formation have illuminated aspects of sex determination, cell migration, cell determination, cell communication and cell-cell interaction.

The testis contains niches for both germline and somatic stem cells. Genetic analyses uniquely possible in *Drosophila* have

shown that these niches function through signaling pathways that ensure maintenance of stem cells throughout adult life; the use of these pathways is conserved for stem cell niches in testes of other animals (see articles in this issue by Matunis and colleagues,¹³ and by Zoller and Schulz¹⁴). Furthermore, cell signaling and cell-cell interactions in the niche and at later stages of gametogenesis serve as models for interactions that occur between different cell types in other tissues during development.

As with other cases of differentiation, spermatogenesis begins with a switch of gene expression program from that of a relatively undifferentiated state to one that results in formation of a specific cell type. In the case of spermatogenesis, the changes in gene expression promote the transition from a mitotic germline cell (the spermatogonium) to one that will form a functional gamete (see article in this issue by Lim et al.¹⁵). Some spermatogenesis genes are needed for meiosis and meiotic cytokinesis (for reviews of these processes, see this issue's articles by McKee and colleagues,¹⁶ and by Giansanti and colleagues,¹⁷ respectively). Others are required for formation of mature, motile sperm (described in this issue's article by Fabian and Brill¹⁸). Understanding this developmental program is relevant not only to spermatogenesis, but also gives insights into the process and regulation of cell cycle control and chromosome segregation (mitotic as well as meiotic), cell morphogenesis, chromatin condensation, organelle biogenesis and cell motility.

The resulting sperm contain a complex array of proteins (see article in this issue by Rettie and Dorus¹⁹). In addition to being of relevance to sperm form and function, identification and analysis of these proteins dissects the proteome of a single, pure cell type. Some sperm proteins are conserved across a wide range of species, indicating structures or functions integral to sperm in all organisms, whereas others are unique to *Drosophila*, providing a window into potential species-specific processes or functions.

After they are transferred to females, sperm are stored within the female reproductive tract (described in this issue's article by Schnakenberg et al.²⁰). During this time, they must be maintained in an active form, and must be released to fertilize eggs. In addition to allowing extended fertility after even a single mating, sperm storage provides an opportunity for sperm competition²¹

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and female sperm choice²² in situations where females mate multiple times (as occurs in *Drosophila*). Sperm competition and female sperm choice, as well as cross-species isolating mechanisms, are forms of selection that may drive dynamic changes in the sequences, abundance, or presence of some reproductive proteins, including those in the sperm proteome.

Drosophila has become a preeminent model for studying all of these processes, in part because *Drosophila* sperm are in many ways typical and in part because of the powerful genetics, conserved genes and well-characterized stages of spermatogenesis available in the fly system (previously reviewed in refs. 1, 23–25). The reviews in this issue offer an updated snapshot of this exciting field of research. We hope they will be of value to newcomers and experts alike.

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