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## Sperm ion channels and transporters in male fertility and infertility

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### Abstract

Mammalian sperm cells must respond to cues originating from along the female reproductive tract and from the layers of the egg in order to complete their fertilization journey. Dynamic regulation of ion signalling is, therefore, essential for sperm cells to adapt to their constantly changing environment. Over the past 15 years, direct electrophysiological recordings together with genetically modified mouse models and human genetics have confirmed the importance of ion channels, including the principal Ca<sup>2+</sup>-selective plasma membrane ion channel CatSper, for sperm activity. Sperm ion channels and membrane receptors are attractive targets for both the development of contraceptives and infertility treatment drugs. Furthermore, in this era of assisted reproductive technologies, understanding the signalling processes implicated in defective sperm function, particularly those arising from genetic abnormalities, is of the utmost importance not only for the development of infertility treatments but also to assess the overall health of a patient and his children. Future studies to improve reproductive health care and overall health care as a function of the ability to reproduce should include identification and analyses of gene variants that underlie human infertility and research into fertility-related molecules.

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Regulation of ion balance is essential for sperm motility and fertility. In particular, ions pass through channels at least 1,000 times faster than through transporters<sup>1</sup>. Thus, ion channels enable sperm to rapidly respond to guidance cues in the female reproductive tract. Calcium (Ca<sup>2+</sup>) influx through Ca<sup>2+</sup>-permeable ion channels can affect cell signalling by altering local electrostatic fields and protein conformations<sup>2</sup>. The speed and effectiveness of Ca<sup>2+</sup> signalling is a consequence of a more than ~10,000-fold gradient maintained across the cell

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All authors researched data for the article, made substantial contributions to discussion of content, and wrote, reviewed and edited the manuscript before submission.

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plasma membrane; the intracellular  $\text{Ca}^{2+}$  concentration (10–100 nM free) is, therefore, low when compared with the extracellular concentration ( $\sim 1\text{--}2\text{ mM}$ )<sup>2,3</sup>.

$\text{Ca}^{2+}$  signalling is a conserved mechanism to modulate cell motility by increasing flagellar asymmetry.  $\text{Ca}^{2+}$  influx is required for altering flagellar waveform in *Chlamydomonas*<sup>4</sup>, the steering and turning of sperm from sea urchins<sup>5,6</sup> and fish<sup>7</sup>, and sperm hyperactivation in mammals<sup>8</sup>. The axoneme of motile cilia and flagella have  $\text{Ca}^{2+}$ -binding sites that regulate flagellar curvature by modulating the motor protein dynein ATPase<sup>9-12</sup>. Although various molecular mechanisms can chelate, compartmentalize or extrude  $\text{Ca}^{2+}$  upon its entry into the cytosol<sup>2</sup>, specialized  $\text{Ca}^{2+}$ -selective ion channels are the only  $\text{Ca}^{2+}$  entry sites in the sperm flagella<sup>13,14</sup>.

In mammals, flagellar  $\text{Ca}^{2+}$  entry is facilitated by the cation channel of sperm (CatSper), the sperm-specific  $\text{Ca}^{2+}$  channel complex<sup>13,15-18</sup> (FIG. 1). CatSper-dependent  $\text{Ca}^{2+}$  entry induces hyperactivated motility during sperm capacitation<sup>16,17</sup>. Hyperactivated motility is the swimming pattern observed in most sperm retrieved from the oviductal ampulla at the time of fertilization and is characterized by a deep and asymmetrical flagellar bend<sup>8,19,20</sup>. Sperm hyperactivation helps to free sperm cells from the oviductal epithelium<sup>21,22</sup>, to facilitate their upstream progression<sup>23</sup>, and to penetrate the zona pellucida to fuse with the egg<sup>15</sup>. Specific signals, such as progesterone and other secretion factors, which are present within the oviduct around ovulation, stimulate hyperactivation of bovine and human sperm<sup>24,25</sup>. Studies have suggested that human CatSper functions as a polymodal sensor that translates physical and chemical cues in the reproductive tract into a  $\text{Ca}^{2+}$  response<sup>23,26</sup>. Thus, elucidating the molecular regulatory mechanisms by which CatSper is regulated is fundamentally important to understanding sperm hyperactivation and mammalian fertilization.

Within the past two decades, the development of genetically modified mouse models and the application of direct electrophysiological recordings have improved our understanding of the molecular basis of mammalian sperm sensory signalling<sup>27</sup>. Species-specific variations in sperm signalling such as differences in overall receptor expression, molecular composition, and/or regulatory mode of primary ion channels have also been revealed<sup>28</sup>. Furthermore, application of state-of-the-art techniques such as super-resolution imaging<sup>29-32</sup>, imaging flow cytometry<sup>33-35</sup> and cryo-electron tomography<sup>36-38</sup> to sperm cells has begun to more directly inform the molecular and structural bases of sperm motility. Thus, the signal transduction pathways that lead to the mechanical transitions in the axoneme to regulate sperm motility have, at least partially, been revealed.

In this Review, we explore the literature regarding ion channel signalling in the context of regulating mammalian sperm motility and male fertility, focusing on sperm ion channels and membrane transporters for which genetic and/or electrophysiological evidence is available to support their roles in male fertility in mice and humans. We summarize important findings, current controversies and challenges, and provide insights into molecular mechanisms, future perspectives and clinical developments in the field.

## Sperm intracellular alkalinization

Mammalian sperm remain quiescent during maturation and storage in the acidic luminal environment of the epididymis (pH 6.6–6.8)<sup>39–41</sup>. Variations in pH along the female reproductive tract is one of the physiological cues that stimulates sperm capacitation, including hyperactivated motility and the acrosome reaction<sup>42</sup>. In humans, the pH of luminal fluid in the female reproductive tract increases gradually from the vagina (pH ~4.4) towards the cervix (pH 6.5–7.5); the pH of cervical mucus is in the range 5.4–8.5 and the pH of the uterus is >7 (REFS<sup>43,44</sup>). The luminal pH is normally highest in the fallopian tubes (7.3–7.7 in humans and ~7.9 in pigs and rabbits)<sup>43</sup>. Thus, spermatozoa encounter a drastic extracellular pH change during their journey through the female reproductive tract. Upon ejaculation, human spermatozoa are mixed with seminal plasma (pH 7.2–8.4), which alkalinizes the acidic vaginal environment<sup>45,46</sup>. The optimal pH for sperm motility is in the range 7.0–8.5 in bulls<sup>47</sup> and humans<sup>48</sup>, which is consistent with the pH of the oviductal fluid. Intracellular alkalinization of sperm cells during capacitation can be caused by HCO<sub>3</sub><sup>-</sup> uptake from the fluid in the female reproductive tract through HCO<sub>3</sub><sup>-</sup> transporters and extrusion of H<sup>+</sup> through proton carriers (FIG. 1 a).

### Proton carriers

Carrier-mediated mechanisms constitute the major route for proton transport across the plasma membrane in sperm<sup>39</sup>. Membrane transporters such as sodium–hydrogen exchangers (NHEs) and the voltage-gated proton-selective ion channel HV1 are the best-studied examples.

**Sodium–hydrogen exchangers.**—NHEs, encoded by the *Slc9* gene family, transport Na<sup>+</sup> into cells and H<sup>+</sup> out of cells<sup>49</sup>, thereby regulating intracellular pH (FIG. 1a). NHE1 (REF.<sup>50</sup>), NHE5 (REF.<sup>51</sup>) and NHE8 (REF.<sup>52</sup>) are expressed in multiple tissue types in mammals, including the testis. However, two NHEs — sNHE (encoded by *Slc9c1*)<sup>53,54</sup> and NHA1 (encoded by *Slc9b1*)<sup>54</sup> — are expressed specifically in sperm flagellum. Gene knockout studies in mice have demonstrated the importance of pH regulation by NHEs in sperm physiology (TABLE 1). In these studies, loss of sNHE<sup>53</sup> or NHA1 and NHA2 (encoded by *Slc9b2*)<sup>55</sup> together caused male infertility and knockout of NHE8 (encoded by *Slc9a8*) in male germ cells resulted in defects in acrosome formation and male infertility<sup>56</sup>. However, the extent of the contribution made by each NHE in controlling intracellular pH in mouse sperm is not clear. For example, impaired sperm motility in sNHE-deficient sperm was only partially rescued by artificial alkalinization but was completely rescued by administration of cyclic AMP (cAMP) analogues<sup>53</sup>. The intracellular cAMP levels and protein expression of soluble adenylyl cyclase (sAC) were attenuated in the absence of sNHE, NHA1 and NHA2 (REFS<sup>55,57</sup>), suggesting a functional redundancy in NHEs for sAC expression and/or their relationship with cAMP signalling pathways to facilitate sperm motility regulation. In sea urchins, sNHE can control intracellular pH homeostasis more rapidly than typical transporters by responding to hyperpolarization and cAMP<sup>58</sup>. As no other NHE has been reported in the sea urchin genome, sNHE might act as a solo voltage-dependent NHE. Whether other, as-yet-uncharacterized, NHEs are involved in pH regulation in mammalian sperm is unknown.

**Proton channels.**—The HV1 proton channel has been suggested to control human sperm intracellular pH<sup>59</sup>. HV1 is homologous with the voltage sensor domain (VSD) of voltage-gated channels and functions as a dimeric channel complex, but does not contain a separate pore domain<sup>60,61</sup>. Molecular dynamic simulations of HV1 homology models suggest that HV1 probably contains an internal water wire within the central crevice of the VSD for selective proton transfer<sup>62</sup>. Full-length HV1 and N-terminal cleaved HV1<sub>Sper</sub> have been detected in human sperm<sup>63</sup>. Both channels exhibit voltage-dependent activation, which requires a pH difference across the membrane ( $\Delta$  pH), and are inhibited by zinc<sup>59,63</sup>. The voltage dependence of heterologously expressed HV1<sub>Sper</sub> is also affected by simultaneous changes in intracellular and extracellular pH. Electrophysiological recordings of human sperm have not been able to distinguish HV1<sub>Sper</sub> from HV1 current<sup>63</sup>; thus, uncovering the molecular mechanisms by which HV1 and HV1<sub>Sper</sub> regulate the pH of human sperm requires further studies. Such studies would rely on human genetic evidence because HV1 is absent in mouse sperm<sup>59</sup>. Interestingly, a 2018 study found that the HV1 channels are arranged in bilateral lines along one side of the flagellar membrane<sup>32</sup>, which suggests that the combined effect of H<sup>+</sup> efflux through many HV1 channels alters the local intraflagellar pH and, in doing so, can regulate pH-sensitive molecules.

Intracellular alkalinization has been predicted to precede membrane potential hyperpolarization and Ca<sup>2+</sup> signalling in mouse and human spermatozoa after ejaculation<sup>13,64-67</sup>. Thus, understanding the molecular mechanism of sperm pH regulation and sensing is crucial. HV1 and sNHE are both confined to the principal piece of the flagella<sup>53,59</sup>. Interestingly, the ion channels CatSper and KSper (sperm-specific K<sup>+</sup> channel), which are both pH-sensitive, are also found in the principal piece (FIG. 1a), suggesting compartmentalized pH regulation and sensing machineries.

## Membrane potential hyperpolarization

Electrochemical gradients across cellular plasma membranes are generally maintained by Na<sup>+</sup>/K<sup>+</sup> ATPase<sup>68</sup>. Resting membrane potential is primarily set by K<sup>+</sup> channels and is typically about  $-70$  mV in somatic cells, including nerve fibres, in the absence of excitation<sup>69</sup>. By comparison, it is less polarized in mammalian spermatozoa (for example, about  $-40$  mV in mouse and human sperm)<sup>64,65,70</sup>. During capacitation, the sperm membrane hyperpolarizes — K<sup>+</sup> efflux through activated KSper is primarily responsible for this change in membrane potential<sup>64</sup> (FIG. 1a). In turn, hyperpolarization regulates various membrane proteins, including the voltage-gated proton channel HV1, ion exchangers and Ca<sup>2+</sup> channels. Abnormal depolarization of membrane potential might be associated with human male subfertility<sup>71-74</sup>; thus, understanding the molecular interactions and regulatory mechanisms of Na<sup>+</sup>/K<sup>+</sup> exchange and KSper is critical to understanding sperm physiology. Genetic evidence regarding which mutations in genes encoding Na<sup>+</sup>/K<sup>+</sup> ATPases and KSper are associated with fertility defects in men remains to be collected.

## Na<sup>+</sup>/K<sup>+</sup> transporters

Na<sup>+</sup>/K<sup>+</sup> ATPase transporters contribute to the regulation of membrane potential owing to unequal exchange of cytoplasmic Na<sup>+</sup> for extracellular K<sup>+</sup> (REF.<sup>68</sup>). The transporter consists

of a group of isozymes that contain  $\alpha$ -subunits and  $\beta$ -subunits. The catalytic  $\alpha$ -subunit of  $\text{Na}^+/\text{K}^+$  ATPase facilitates ion permeation of the plasma membrane<sup>68</sup>. The  $\alpha 1$  subunit is expressed in a variety of tissue and cell types in humans and rats, including sperm<sup>75,76</sup>; however, the  $\alpha 4$  subunit is specific to male germ cells<sup>76</sup>. When rat or human  $\alpha 4$  subunit is expressed in transgenic mice, it is particularly abundantly expressed in the sperm flagellum, which is consistent with its expression in wild-type rat and human sperm<sup>77,78</sup>. Treatment of rat sperm with a low concentration of ouabain (which selectively inhibits  $\alpha 4$ ) increased intracellular  $\text{Na}^+$ , depolarized the membrane potential, increased intracellular calcium and decreased intracellular pH<sup>79</sup>. Because  $\alpha 4$  does not directly transport  $\text{H}^+$ , it has been proposed that  $\alpha 4$  provides electrochemical energy that facilitates pH regulation by NHEs<sup>77,80</sup>. In particular, sNHE has a putative voltage sensor<sup>53</sup> and so could potentially be regulated by an  $\alpha 4$ -established membrane potential. Measuring intracellular pH under various transmembrane sodium gradients will help to test this possibility.

The potential contribution of the  $\alpha 1$  subunit to sperm function remains to be clarified, as  $\alpha 1$  subunit knockout is embryonically lethal in mice<sup>81</sup>. By contrast, disruption of *Atp1a4*, which encodes the  $\alpha 4$  subunit, in mice causes complete sterility<sup>82</sup> (TABLE 1), demonstrating that  $\alpha 4$ -mediated ion transport is critical for sperm function. Furthermore,  $\text{Na}^+/\text{K}^+$  ATPase  $\alpha 4$ -deficient sperm from the knockout mice are severely bent at the junction of the midpiece and the principal piece, exhibit increased intracellular  $\text{Na}^+$  levels and have depolarized membrane potentials, consistent with the phenotypes observed in mouse sperm with altered osmoregulation<sup>83</sup>. Contrastingly, transgenic mouse sperm that express functional rat *Atp1a4* exhibit increased membrane hyperpolarization and also demonstrate increased total sperm motility and hyperactivated motility<sup>77</sup>; these effects starkly contrast those observed in the *Atp1a4*-disrupted mice that are completely sterile<sup>82</sup>, demonstrating the importance of functional ATP1A4 in sperm motility and fertility. Expression of human *ATP1A4* in transgenic mice results in similar motility phenotypes but not in a significant change in sperm membrane potential<sup>78</sup>. The expression of rat or human  $\alpha 4$  does not affect the acrosome reaction in transgenic mouse sperm<sup>77,78</sup>. These studies demonstrate that  $\alpha 4$  activity primarily maintains sperm intracellular  $\text{Na}^+$  levels and contributes to setting membrane potential. As a result,  $\alpha 4$  affects several vital parameters, such as intracellular pH and  $\text{Ca}^{2+}$ , which are essential for sperm motility and hyperactivation.

### The KSper channel

KSper was first recorded in mouse sperm in 2007 (REF<sup>64</sup>). Upon intracellular alkalinization, KSper is activated and further hyperpolarizes the spermatozoan cellular membrane (FIG. 1a). Genetic disruption of *Kcnu1* demonstrated that SLO3, the pore-forming  $\alpha$ -subunit, mediates KSper in mouse sperm<sup>70,84</sup> (TABLE 1). Despite their normal morphology and motility, mouse spermatozoa lacking SLO3 are infertile owing to impaired hyperpolarization during capacitation. However, a residual  $\text{K}^+$  current has been observed in *Kcnu1*-knockout sperm at very positive potentials<sup>84</sup>, which suggests that another voltage-gated  $\text{K}^+$  current might exist in mouse sperm. However, a subsequent study clarified that all voltage-gated outward current is abolished in sperm of *Kcnu1/Catsper1*-double-knockout mice<sup>14</sup>, demonstrating that the residual  $\text{K}^+$  current observed in *Kcnu1*-knockout sperm is caused by  $\text{K}^+$  efflux through CatSper and provides evidence that KSper and CatSper are the sole

mediators of voltage-dependent  $K^+$  and  $Ca^{2+}$  currents, respectively, in uncapacitated mouse epididymal sperm in response to alkalization. Heterologous expression of leucine-rich repeat-containing protein 52 (LRRC52), an auxiliary subunit of SLO3, modulates SLO3 gating by shifting its voltage and pH dependence to more negative values, close to those of native KSper currents<sup>85,86</sup>. Consistently, upon deletion of *Lrrc52*, the activity of KSper at rest is attenuated and its activation requires increasingly positive voltages and higher pH, resulting in a more depolarized membrane potential<sup>86</sup>. Abrogation of LRRC52 results in male subfertility and reduced litter sizes<sup>86</sup> (TABLE 1), supporting the association of this abnormal depolarization of membrane potential with male subfertility in humans<sup>71-74</sup>.

Native human KSper (hKSper), unlike mouse KSper, is sensitive to both intracellular alkalization<sup>65,87</sup> and calcium<sup>65,88</sup>. It exhibits  $Ca^{2+}$  sensitivity in the same manner as another SLO family member SLO1 (also known as BK channel and encoded by *Kcnma1*), and pH sensitivity in the same manner as SLO3. These unusual characteristics — whereby hKSper displays hybrid characteristics of the mouse counterpart, SLO3, which is pH-sensitive, and SLO1, which is  $Ca^{2+}$ -sensitive — have resulted in controversy regarding the exact molecular composition of hKSper. However, studies have shown that heterologous expression of human SLO3 in *Xenopus* and 293T cells does give rise to currents that exhibit the properties of native hKSper in human sperm<sup>65,89,90</sup> — that is, activation by  $Ca^{2+}$  and alkalization, inhibition by progesterone, and a single-channel conductance of 70 pS (REF. 65). Moreover, SLO3 and LRRC52 were identified in human sperm by mass spectrometry<sup>65</sup>, further supporting the view that SLO3 and LRRC52 comprise hKSper. A 2017 study showed that a single-nucleotide polymorphism of human SLO3 (C382R) can endow the channel with enhanced pH and  $Ca^{2+}$  sensitivities<sup>89</sup>, suggesting that species-specific SLO3 variants could have acquired different  $Ca^{2+}$  sensitivity. KSper sensitivity to intracellular pH and  $Ca^{2+}$  might define its role in mouse and human sperm: in mouse sperm, the KSper-dependent, capacitation-associated membrane hyperpolarization is an upstream signalling event that increases the force driving  $Ca^{2+}$  influx through CatSper<sup>91</sup>; in humans, KSper probably functions downstream of CatSper because activation of hKSper requires an increase in cytosolic  $Ca^{2+}$  (REF<sup>92</sup>); KSper-induced hyperpolarization would further affect CatSper. An improved understanding of how sperm membrane potential is regulated during capacitation will provide insights into species-specific fertilization processes.

## Ca<sup>2+</sup> influx and signal transduction

The intracellular  $Ca^{2+}$  increase in sperm, which is required for hyperactivated motility and the acrosome reaction, originates primarily from influx of extracellular  $Ca^{2+}$ ; in these specialized cells, the organelles that are typically used for intracellular  $Ca^{2+}$  storage and release, such as the endoplasmic reticulum, are less developed than in somatic cells. Mouse spermatogenic cells possess T-type  $Ca_v$  channels<sup>93</sup>, but electrophysiological recordings have demonstrated that  $Ca_v$  currents gradually decrease during spermiogenesis and become undetectable in epididymal sperm<sup>94</sup>, suggesting that  $Ca_v$  currents are only required during spermatogenesis and not in mature sperm cells. So far, CatSper is the only  $Ca^{2+}$  channel in which genetic mutations have been reported to cause male infertility. Thus, the flagellar-specific  $Ca^{2+}$  channel CatSper is a predominant pathway by which  $Ca^{2+}$  enters mammalian sperm.

## Molecular organization of CatSper

CatSper was first identified in 2001 as a gene (*Catsper1*) encoding a putative sperm  $\text{Ca}^{2+}$  channel with a single repeat of six transmembrane domains in human and mouse testis<sup>15</sup>. The topology of CATSPER1 is unlike that of conventional voltage-gated  $\text{Ca}^{2+}$  channels, which are typically composed of four repeats of six transmembrane domains. Subsequently, *Catsper2* (REF<sup>17</sup>), *Catsper3* and *Catsper4* (REFS<sup>18,95</sup>) were identified. CatSper currents are absent in mouse sperm in which one subunit (*Catsper1*, *Catsper2*, *Catsper3* or *Catsper4*) has been knocked out<sup>18</sup>, indicating that the CatSper pore is a heterotetramer composed of CATSPER1–4 and that one of each is required for channel function. In addition, CatSper channels comprise at least six additional subunits that do not form the pore (FIG. 1b): four transmembrane proteins that are predicted to contain large extracellular domains (ECDs), denoted CATSPER $\beta$ <sup>96</sup>, CATSPER $\gamma$ <sup>97</sup>, CATSPER $\delta$ <sup>98</sup> and CATSPER $\epsilon$ <sup>29</sup>; and two small cytoplasmic proteins, CATSPER $\zeta$ <sup>29</sup> and EF-hand calcium-binding domain-containing protein 9 (EFCAB9)<sup>31</sup>. CatSper is the most complex ion channel known<sup>31,98</sup>. Knocking out any one of the four genes that encode CATSPER1–4 in mice renders males infertile<sup>15,18,95,99</sup> (TABLE 1) and *CATSPER* loss-of-function mutations have also been identified in infertile men<sup>100–104</sup>. Sperm cells in which *Catsper1*, *Catsper2*, *Catsper3* or *Catsper4* has been knocked out fail to develop hyperactivated motility<sup>16–18,99</sup>, which is consistent with the inability of *Catsper1*-null sperm to fertilize oocytes with an intact zona pellucida<sup>15</sup>. Direct whole-cell patch clamp studies of spermatozoa have demonstrated that CatSper is  $\text{Ca}^{2+}$ -selective channel activated by intracellular alkalinization<sup>13,67</sup>.

Deletion of *Catsper $\delta$* , which encodes one of the nonpore-forming transmembrane subunits, CATSPER $\delta$ , also abrogates CatSper current and hyperactivated motility, resulting in infertility in male mice<sup>98</sup> (TABLE 1). This phenotype arises from the loss of not only CATSPER $\delta$  but also each transmembrane domain-containing CATSPER protein in sperm cells of *Catsper $\delta$* -null males, and is similar to the all-or-nothing expression pattern of the CatSper pore-forming subunits<sup>29–31,98</sup>. By contrast, knocking out *Catsper $\zeta$*  and/or *Efcab9* does not completely eliminate the formation of functional CatSper channels and correspondingly results in male subfertility<sup>29,31</sup> (TABLE 1). Consistently, reduced CATSPER $\zeta$  protein expression has been reported in men with asthenozoospermia<sup>105</sup>, suggesting that these two non-transmembrane subunits, CATSPER $\zeta$  and EFCAB9, function as true auxiliary subunits that could modulate CatSper expression levels and/or channel kinetics. Interestingly, *Catsper $\zeta$*  orthologues have been found only in mammals, implying that its regulatory function might be specific to mammalian CatSper channels<sup>29</sup>. Direct analyses of the functions of the other non-pore-forming transmembrane subunits, CATSPER $\beta$ , CATSPER $\gamma$  and CATSPER $\epsilon$  await their investigation using genetic disruption in mice and/or the discovery of human genetic abnormalities related to these proteins.

## Spatial organization of CatSper

**Flagellar  $\text{Ca}^{2+}$  signalling nanodomains.**—Super-resolution microscopy has been used to demonstrate that the macromolecular CatSper channel forms four linear (also called quadrilateral)  $\text{Ca}^{2+}$  signalling nanodomains along the sperm tails in both mice and humans<sup>29–31</sup> (FIGS 1,2) and organizes a network of intracellular signalling molecules such as calmodulin-dependent protein kinase II (CaMKII) and calcineurin<sup>30</sup>. Compartmentalized

domains enable specific and fast-triggering downstream events and are common cellular adaptations for effective  $\text{Ca}^{2+}$  signalling in many biological systems<sup>2</sup>. A 2019 study in mice demonstrated that each of the four CatSper nanodomains is further resolved into two row structures<sup>31</sup> (FIG. 2b). CATSPER $\zeta$  and/or EFCAB9 deficiency disrupts not only the linearity of the CatSper nanodomains but also the two-row organization, renders the proximal sperm flagella rigid and alters sperm motility<sup>29,31</sup>. As a result, the presence and integrity of these nanodomains serves as an indicator of fertilizing capability. These studies also indicate that CATSPER $\zeta$  and EFCAB9 regulate the compartmentalization of  $\text{Ca}^{2+}$  signalling in mammalian sperm, and might, therefore, modulate the mechanism by which CatSper facilitates  $\text{Ca}^{2+}$  influx.

**CatSper assembly and trafficking to the flagellar membrane.**—Despite the absolute requirement of the CatSper channel for male fertility and the importance of CatSper nanodomain formation for sperm  $\text{Ca}^{2+}$  signalling, knowledge of how CatSper is assembled and organized into nanodomains is limited. This lack is largely due to an inability to heterologously reconstitute CatSper under laboratory conditions. Although the timing of translation of CatSper subunits is not known, mouse CATSPER $\delta$  has been suggested to assist the assembly of the CatSper channel in the endoplasmic reticulum or the Golgi, as CATSPER1 is degraded in its absence<sup>98</sup>. CATSPER $\delta$  and CATSPER $\epsilon$  might dimerize, as they have a region of high homology at their C terminus<sup>29</sup>. In 2018, a 6-bp in-frame deletion within *CATSPERE* in an infertile man was reported; the sperm cells of this patient did not respond to progesterone to elicit CatSper-mediated  $\text{Ca}^{2+}$  increase and in vitro fertilization (IVF) failed<sup>106,107</sup>. This alteration occurred within the region of CATSPER $\epsilon$  that exhibits high homology with CATSPER $\delta$ , suggesting that CATSPER $\epsilon$  might also have a role in the assembly of the large CatSper complex together with CATSPER $\delta$ . Studies in mice have suggested that only a correctly assembled CatSper channel, probably with all of its transmembrane subunits in place, is trafficked to the flagellar membrane<sup>98</sup>.

Questions remain regarding how CatSper traffics to the flagellar membrane and whether intraflagellar transport (IFT) machinery and adaptor proteins exist that specifically interact with CatSper or other sperm ion channels and transporters is unknown. Studies using genetically altered mouse models have shown that IFT is essential for mammalian spermiogenesis but that epididymal spermatozoa are devoid of IFT proteins<sup>108</sup>. For example, IFT25 and IFT27, which are dispensable for ciliogenesis in somatic cells, are required for flagellar formation as they participate in assembling and transporting structural components of the fibrous sheath that is unique to spermatozoa<sup>109,110</sup>. Identification of the CatSper interactome from spermatids, not mature sperm, and application of super-resolution microscopy could provide valuable information on the trafficking of flagellar membrane proteins and quadrilateral compartmentalization (FIGS 1,2) during spermiogenesis.

**Human versus mouse CatSper.**—Interestingly, two studies in infertile patients who lacked CatSper current demonstrated that CatSper assembly in humans is strikingly different from CatSper assembly in mice<sup>103,104</sup>. Infertile men who had lost one genomic copy of *CATSPER2* (owing to a copy number variation)<sup>104</sup> or both copies owing to contiguous deletion of a genomic region encompassing *CATSPER2* and *STRC* (15q15.1–15q15.3,



deletion of which leads to the rare deafness–infertility syndrome)<sup>103,111,112</sup> had otherwise normal semen parameters despite substantially reduced or absent expression of CATSPER2, respectively, in their spermatozoa. The protein expression levels of the other three CatSper pore-forming subunits in these unique human CatSper-deficient sperm cells were only marginally changed, or were not changed at all<sup>103,104</sup>, in contrast to the changes observed in mouse CatSper-knockout sperm. Moreover, the uniquely quadrilateral sub-flagellar compartmentalization was observed for CATSPER3 and CATSPER4 in the principal piece of human *CATSPER2*-deficient spermatozoa<sup>103</sup>. These results suggest that human CATSPER1, CATSPER3 and CATSPER4 proteins can still traffic to the flagella, but they cannot form functional channels without CATSPER2. Thus, the all-or-nothing expression pattern of the CatSper complex apparently does not exist in human sperm cells. Formation of trimeric complexes lacking CATSPER2 and their proper trafficking to the flagella is unlikely to happen because a tetramer of an ion channel pore is often assembled as a dimer of dimers. These findings illustrate the species-specific differences in the assembly and trafficking patterns of CatSper.

A more thorough understanding of the organization of CatSper into Ca<sup>2+</sup> signalling nanodomains requires further investigation. For example, probing the absence or presence and the sub-flagellar localizations of other CatSper auxiliary subunits in these human CATSPER2-deficient sperm will provide more insight into their roles in channel assembly and trafficking and domain organization. Additional unidentified channel subunits that specifically function in trafficking might also exist. This possibility was implied by a recent CatSper proteome study of mouse sperm cells, which showed several candidate CatSper-associated proteins, including one with a conserved domain involved in membrane trafficking<sup>31</sup>. The likely existence of additional unknown CatSper subunits highlights the molecular and regulatory complexity of this important channel.

### Modulation of CatSper activity

#### Molecular mechanisms of alkaline activation and Ca<sup>2+</sup> sensitivity of CatSper.

—Ca<sup>2+</sup> entry through CatSper requires channel activation by intracellular alkalization<sup>13,31,66,67</sup>. The pH sensitivity of CatSper was initially attributed to a conserved histidine-rich region in the N terminus of CatSper1 among mammals<sup>13,15</sup>, but, subsequently, the molecular basis by which CatSper can sense pH and Ca<sup>2+</sup> has been elucidated. The mechanism relies upon EFCAB9, a testis-specific protein that has co-evolved with other core members of the CatSper complex<sup>31</sup>. EFCAB9 was identified in a comparative proteomics screen for proteins differentially expressed in *Catsper1*-null sperm when compared with wild-type sperm<sup>31</sup>. EFCAB9 is the only CatSper subunit that contains known Ca<sup>2+</sup>- and calmodulin-binding motifs. Three EF-hand motifs were identified in EFCAB9 and it was shown to bind CATSPER $\zeta$  in a pH-dependent and Ca<sup>2+</sup>-dependent manner<sup>31</sup>. An increased Ca<sup>2+</sup> concentration facilitated EFCAB9–CATSPER $\zeta$  complex formation, whereas alkalization impeded the interaction<sup>31</sup>. That EFCAB9 and CATSPER $\zeta$  work as one functional unit is supported by mouse models in which either *Efcab9* or *Catsperz* or both has been knocked out. These three mouse models exhibit identical phenotypes and demonstrate that the expression of these proteins is interdependent: if *Efcab9* has been knocked out, CATSPER $\zeta$  is absent, and vice versa<sup>31</sup>. Although EFCAB9-deficient sperm exhibit reduced

CatSper expression, when intracellular pH is low, wild-type and *Efcab9*-null sperm show a similarly dense CatSper current ( $I_{\text{CatSper}}$ )<sup>31</sup>, indicating that EFCAB9–CATSPER $\zeta$  normally limits CatSper-mediated Ca<sup>2+</sup> entry. When intracellular pH rises, CatSper is activated in a Ca<sup>2+</sup>-dependent manner, but in the absence of EFCAB9, the channel is less sensitive to intracellular Ca<sup>2+</sup> changes and less responsive to alkalinization<sup>31</sup>. Thus, a working model was generated: before capacitation, the pH-sensing and Ca<sup>2+</sup>-binding EFCAB9–CATSPER $\zeta$  complex stabilizes the closed CatSper pore; upon alkalinization and Ca<sup>2+</sup> entry through CatSper, EFCAB9–CATSPER $\zeta$  undergoes structural rearrangements releasing its inhibition of the channel. Further opening of the channel is followed by Ca<sup>2+</sup> entry, which would be bound by EFCAB9, stabilizing its prolonged open state<sup>31</sup>.

**Steroid hormones in human CatSper regulation.**—Both mouse and human CatSper channels are activated by intracellular alkalinization; however, activation of CatSper by physiological ligands has only been reported in humans<sup>66,67</sup> and rhesus macaques<sup>113</sup>, suggesting that CatSper is regulated by species-specific mechanisms. For example, progesterone, a steroid hormone secreted by cumulus cells, can robustly evoke human<sup>66,67,102</sup>, but not mouse<sup>67</sup>, CatSper currents. Similarly, prostaglandin E<sub>1</sub> (PGE<sub>1</sub>), which is abundant in the seminal plasma and also secreted by cumulus cells, was found to activate human, but not mouse, CatSper<sup>67</sup>.  $\alpha/\beta$  hydrolase domain-containing protein 2 (ABHD2) has been identified as the non-genomic progesterone receptor in human sperm<sup>114</sup>. ABHD2 abolishes endocannabinoid 2-arachidonoylglycerol (2-AG) inhibition of CatSper as a progesterone-dependent lipid hydrolase, enabling CatSper activation<sup>114</sup>. Other steroid hormones, such as pregnenolone sulfate and testosterone, also modulate CatSper-mediated Ca<sup>2+</sup> influx into human sperm; they probably bind to the same sites as progesterone<sup>115</sup>. Whether testosterone and other steroids function as human CatSper agonists or antagonists remains controversial<sup>115-117</sup>, and the mechanisms by which these ligands bind to and modulate CatSper activity remain largely unknown.

Additional ligands and chemicals have been reported to activate human CatSper directly or indirectly<sup>115,118-120</sup>. These include structurally diverse endocrine-disrupting chemicals (EDCs) such as *p,p'*-dichlorodiphenyldichloroethylene, 4-methylbenzylidene camphor and triclosan<sup>119-121</sup>. Thus, the presence of any of these EDCs in the female reproductive tract could interfere with human sperm function by modulating CatSper activity.

The non-pore-forming CatSper transmembrane subunits have been suggested to bind factors that alter CatSper gating as they have large ECDs<sup>29,96-98</sup> (FIG. 1b). They could potentially bind ligands and function as sensory transducers in the polymodal Ca<sup>2+</sup> signalling exhibited by human sperm. Disrupting one of the genes, *CatSperd*, in mice results in an identical phenotype to that seen in mice with knockout of the pore-forming CatSper subunits at the organismal and cellular levels<sup>98</sup>, illustrating that CATSPER $\delta$  is essential to form a functional CatSper. In the future, biochemical and structural characterization of isolated ECDs from the non-pore-forming CatSper transmembrane subunits, in addition to characterization of the cytoplasmic CatSper subunits themselves, is a sensible approach to improving our understanding of the molecular mechanisms of CatSper.

## Signalling pathways in capacitation

Sperm capacitation involves a cascade of signalling pathways (FIG. 1a). Studies have indicated that different pathways directly or indirectly regulate CatSper, the central signalling hub in network with multiple signalling proteins.

### Signalling crosstalk and CatSper

During capacitation, the rise in intracellular pH activates not only CatSper but also KSper. Activation of KSper hyperpolarizes the membrane to further drive  $\text{Ca}^{2+}$  influx through CatSper in mouse sperm<sup>64,70,84</sup>.  $\text{HCO}_3^-$ ,  $\text{Ca}^{2+}$  and bovine serum albumin (BSA) have long been recognized as indispensable for sperm capacitation and fertilization in vitro<sup>122</sup>, engaging signalling pathways that regulate CatSper-mediated calcium signalling.

**$\text{HCO}_3^-$  uptake and cAMP-dependent PKA activation.**—At the molecular level, capacitation is initiated when spermatozoa are exposed to a high concentration of  $\text{HCO}_3^-$  in the luminal fluid of the female reproductive tract, which also has a higher  $\text{Ca}^{2+}$  concentration than the luminal fluid of the epididymis<sup>39</sup>.  $\text{HCO}_3^-$  enters the sperm through  $\text{HCO}_3^-$  transporters<sup>42</sup> and activates a unique sAC<sup>123</sup>, resulting in increased cAMP levels. The cAMP-dependent PKA pathway is important in the regulation of sperm motility: sAC-deficient male mice are infertile<sup>124</sup>, as are mice deficient in the catalytic subunit of PKA,  $\text{Ca}\alpha 2$  (REF.<sup>125</sup>).  $\text{HCO}_3^-$  also stimulates  $\text{Ca}^{2+}$  entry into sperm by raising intracellular pH (FIG. 1a). However, the role of cAMP in the regulation of  $\text{Ca}^{2+}$  influx is less clear. Membrane-permeable analogues of cyclic nucleotides stimulate  $\text{Ca}^{2+}$  entry in mouse<sup>15,96,126,127</sup> and human sperm<sup>26</sup>. However, a series of studies have also shown that an increase in intracellular cAMP (stimulated by  $\text{HCO}_3^-$ , 3-isobutyl-1-methylxanthine, uncaging of cAMP, or adenosine) fails to stimulate  $\text{Ca}^{2+}$  influx in mouse<sup>128</sup> and human sperm<sup>66</sup>. Furthermore, membrane-permeable analogues of cyclic nucleotides, but not physiological concentrations of cAMP or cGMP, activate human CatSper when applied extracellularly, but not intracellularly<sup>26,129</sup>. However, these data are inconsistent with the findings of studies in mouse sperm, in which cAMP directly applied intracellularly through a pipette activated CatSper<sup>130</sup>, but cyclic nucleotide analogues applied extracellularly to the bath solution had no effect<sup>13</sup>. Thus, cAMP and PKA regulation of CatSper-mediated  $\text{Ca}^{2+}$  influx is probably species-specific and needs to be further clarified.

**Calcium influx induced by loss of cholesterol.**—Together with high levels of  $\text{HCO}_3^-$  and  $\text{Ca}^{2+}$  concentration, serum albumin is a key component in mammalian sperm capacitation in vivo as well as in vitro<sup>131,132</sup>. Although BSA is known to induce  $\text{Ca}^{2+}$  influx in sperm in vitro, the molecular mechanism by which this occurs has not been fully elucidated. However, BSA-induced  $\text{Ca}^{2+}$  influx is obliterated in *Catsper1*-null spermatozoa<sup>133</sup>. As cholesterol release from the sperm plasma membrane by BSA is associated with activation of cAMP–PKA pathways during sperm capacitation in both mice<sup>134</sup> and humans<sup>135</sup>, lipid signalling by cholesterol efflux might participate in regulating CatSper-mediated  $\text{Ca}^{2+}$  signalling via the cAMP–PKA pathway. In this respect, caveolin 1 — a protein associated with cholesterol-rich lipid rafts — has been found in the CatSper  $\text{Ca}^{2+}$  signalling nanodomains in mouse sperm. The role of the CatSper channel as a domain

organizer is notable, as caveolin 1 localization is dysregulated in the absence of the CatSper channel but CatSper nanodomains remain intact in the absence of caveolin 1 (REF.<sup>30</sup>).

#### **Downstream signalling that leads to motility regulation at the axoneme.—**

Under capacitating conditions, cAMP-stimulated PKA activity leads to sperm motility changes within a minute<sup>136-138</sup>. By contrast, hyperactivated motility and a capacitation-associated increase in tyrosine phosphorylation (P-Tyr) occur much later in time<sup>139</sup>. The cAMP–PKA pathway is thought to control downstream P-Tyr because permeable cAMP analogues are able to induce P-Tyr, even in the absence of HCO<sub>3</sub><sup>-</sup> and calcium ions<sup>140</sup> and administration of PKA inhibitors completely blocks P-Tyr development<sup>141</sup>. Interestingly, earlier onset and increased P-Tyr was observed in *Catsper1*-null and *Catsperd*-null spermatozoa following incubation under capacitation conditions<sup>30</sup>, suggesting that PKA activity and P-Tyr are suppressed by a CatSper-mediated Ca<sup>2+</sup> signalling pathway. Super-resolution imaging has revealed a striking spatial confinement of P-Tyr to the axoneme in capacitated wild-type sperm (FIG. 3a); P-Tyr spreads and fills the extra-axonemal region in the absence of CatSper<sup>30</sup>. As P-Tyr requires PKA activation, the signal transduction to suppress P-Tyr in the periaxoneme has been suggested to involve active protein phosphatase 1 (PP1) and/or protein phosphatase 2A (PP2A) to limit PKA activity and protein tyrosine phosphatase<sup>30</sup>. In the same study, multiple tyrosine-phosphorylated proteins were identified from capacitated mouse sperm<sup>30</sup>, including axonemal proteins and a testis-specific tyrosine kinase, FER/FERT (FIG. 3b). Subsequently, a 2016 study revealed that the capacitation-associated P-Tyr increases are mostly eliminated in sperm from kinase-inactivating mutant (D743R) *Fer*<sup>DR/DR</sup> males<sup>142</sup>, demonstrating that FER/FERT is the master tyrosine kinase responsible for capacitation-associated P-Tyr. However, *Fer*<sup>DR/DR</sup> males are fertile<sup>143</sup>, although sperm from these mice do display reduced fertilizing ability in vitro<sup>142</sup>. Thus, P-Tyr is not essential for mouse fertilization in vivo but might have a functional role, such as timing motility regulation and/or determining the lifespan of sperm in the female oviduct.

A Ca<sup>2+</sup> signal can be directly translated into mechanical changes in the axoneme: transient treatment with the Ca<sup>2+</sup> ionophore A23187 can bypass the P-Tyr development in mouse sperm and the CatSper signalling network to induce hyperactivated motility in vitro<sup>144,145</sup>. Nevertheless, CatSper-mediated Ca<sup>2+</sup> signal transduction, originating from the linear CatSper nanodomains, is required for flagellar Ca<sup>2+</sup> regulation in vivo. Disruption of the nanodomain compartmentalization<sup>29,31</sup>, or loss of calcineurin<sup>30,146</sup> or EFCAB9 (REF.<sup>31</sup>), both of which are Ca<sup>2+</sup>-binding proteins from the nanodomains, leads to changes in the flagellar envelope and fertility defects in mice. Discovery of more Ca<sup>2+</sup>-dependent molecules associated with the CatSper nanodomains and/or the axoneme will further elucidate the downstream Ca<sup>2+</sup> signalling affecting dynamic motility regulation.

### **Other channels in sperm**

In addition to CatSper and KSper, HIV1 and sNHE, a number of other channels are involved in regulating sperm function.

## DSper and TRPV4

The presence of a channel that facilitates influx of  $\text{Na}^+$ , as is observed in neuronal excitation, has been hypothesized in sperm and named 'DSper'<sup>28</sup>. The DSper current was recorded in human sperm as a non-CatSper, non-selective cation conductance with outward rectification and pronounced temperature sensitivity<sup>147</sup>. In this study, DSper was potentiated during capacitation and was not diminished by either  $\text{Mg}^{2+}$  or NNC55-0396 administration, which block and inhibit CatSper, respectively. DSper is reversibly activated by warm temperatures (22–37°C;  $T_{1/2}$  34°C in uncapacitated human sperm,  $T_{1/2}$  31 °C in capacitated human sperm). Pharmacological screening of human sperm has suggested that TRPV4 mediates this temperature-dependent DSper current<sup>147</sup>. Capacitated rabbit and human sperm cells have been shown to move towards higher temperatures in vitro<sup>148</sup>, and subsequent study showed that capacitated human sperm swim up a temperature gradient by modulating hyperactivated motility<sup>149</sup>. However, in humans, sperm thermotaxis remains controversial because temperature differences within the female reproductive tract have not been reported and sperm capacitation is not linked to ovulation<sup>150</sup>. Regardless, direct recording of human spermatozoa by whole-sperm patch clamp<sup>147</sup> and mouse genetics studies<sup>151</sup> suggest that TRPV4 could be a molecular basis of temperature sensing at least in mouse and human sperm. However, there is a discrepancy regarding the protein distribution of TRPV4 in human sperm: data from two separate studies that used antibodies raised against the same epitope are conflicting. In one study, TRPV4 was reported in the post-acrosomal and neck regions, but not in the flagella<sup>152</sup>, whereas in the other study, TRPV4 was found in both the acrosome and the flagella<sup>147</sup>. This discrepancy highlights the limitation of immunolabelling in determining protein distribution in sperm cells without stringent controls. In mouse sperm, TRPV4 was detected along the tail and in the head and the signal was absent in *Trpv4*-knockout sperm<sup>151</sup>, validating this specific distribution in mice. Mouse genetics support the involvement of TRPV4 in sperm thermotaxis<sup>151</sup>. Unlike wild-type sperm, *Trpv4*-knockout sperm failed to respond to increasing temperature and exhibited delayed hyperactivated motility, despite still being capable of fertilization<sup>151</sup>. Specific DSper current via TRPV4 might have been undetectable in the previous mouse sperm recordings<sup>13</sup>, as in the absence of external  $\text{Ca}^{2+}$ , the  $\text{Na}^+$  moving through CatSper might have been masking the  $\text{Na}^+$  movement through DSper. Isolating the DSper current and confirming the existence of DSper-dependent thermotactic behaviour in mouse or human CatSper-deficient sperm will clarify whether and how TRPV4 contributes to the molecular basis of mammalian sperm thermotaxis.

## P2X2

Extracellular ATP has been reported to raise the intracellular concentration of  $\text{Ca}^{2+}$  and to stimulate the acrosome reaction in sperm<sup>153</sup>. Mouse spermatozoa have an ATP-gated current ( $I_{\text{ATP}}$ ), which is the only ion current detected from the midpiece of mammalian spermatozoa by patch-clamp recordings<sup>154</sup>.  $I_{\text{ATP}}$  is an intrinsically inwardly rectifying, cation-non-selective and divalent-permeable current, and is mediated by the homomeric P2X2 purinergic receptor. Consistent with the properties of P2X2 current in *Xenopus* oocytes<sup>155,156</sup>,  $I_{\text{ATP}}$  is also activated by  $\text{Zn}^{2+}$  and by an acidic extracellular pH in mouse sperm<sup>154</sup>. In light of these observations, P2X2 might function during sperm maturation in the epididymis where  $\text{Zn}^{2+}$  is abundant and the pH is more acidic than in the oviduct.

Neither oviductal fluid nor cumulus cells from ovulated mice evoked sperm  $I_{ATP}$ , suggesting that the female reproductive tract is not the primary site of sperm P2X2 function in mice<sup>154</sup>. The genetic disruption of *P2rx2* abolishes  $I_{ATP}$  in mouse sperm but does not affect sperm progressive motility, hyperactivation or the acrosome reaction<sup>154</sup>. However, frequent mating renders *P2rx2*-null male mice less fertile, suggesting that ATP-activated  $Ca^{2+}$  influx confers an advantage under high sexual demands<sup>154</sup>.

Subtle sperm fertility phenotypes in the ion-channel knockout mice discussed in this Review could be manifested more prominently in natural mating settings, in which sperm competition presumably exerts a stronger force than it does in a more controlled environment. In vitro, ATP supplementation rescues the immobility observed in metabolite starvation-induced sperm from both wild-type and *P2rx2*-null males<sup>154</sup>. Presumably,  $I_{ATP}$  can deliver  $Ca^{2+}$  to the mitochondria to drive ATP production. The function of  $I_{ATP}$  in human sperm is not clear, as ATP-gated current has not been found in human sperm<sup>26</sup>. The source of extracellular ATP for P2X2 activation also remains uncertain.

### Ca<sub>v</sub>2.3

The voltage-gated calcium channel Ca<sub>v</sub>2.3, encoded by *Cacnale*, mediates R-type  $Ca^{2+}$  currents in neurons<sup>157</sup>. Ca<sub>v</sub>2.3 was thought to have a function in sperm physiology based on its immunological detection in mouse sperm<sup>158,159</sup>. However, male mice lacking Ca<sub>v</sub>2.3 are only mildly subfertile<sup>160</sup> (TABLE 1). Non-capacitated Ca<sub>v</sub>2.3-lacking sperm swim more linearly and exhibit a lower rising rate of  $Ca^{2+}$  transients induced by BSA in their heads than wild-type sperm<sup>160</sup>. This subtle but interesting difference prompted the question as to whether Ca<sub>v</sub>2.3 functions in the acrosome reaction, and this was investigated in a study to determine whether membrane lipids can stimulate the acrosome reaction via modulation of Ca<sub>v</sub>2.3 during capacitation<sup>161</sup>. SNX-482, a Ca<sub>v</sub>2.3-specific blocker, reduced acrosome reactions induced by cholera toxin B or GM1 in sperm incubated under capacitating conditions, and *Cacnale*-null sperm exhibited significant reductions in the rates of the acrosome reaction and successful IVF<sup>161</sup>. Intriguingly,  $Ca^{2+}$  transients in the sperm head were associated with lipid modulation of the activity and localization of Ca<sub>v</sub>2.3 (REF.<sup>161</sup>). The calcium transients occurred only at the apical acrosome of wild-type sperm but in the equatorial segment of *Cacnale*-null sperm with faster kinetics. These Ca<sub>v</sub>2.3 data were not recorded using the current gold-standard method of patch clamping mouse sperm, implying that the resulting readout could be indirect<sup>14,162</sup>. Careful modification of the recording conditions required for recording from mouse sperm incubated under capacitated conditions and/or acrosome-reacted sperm might enable further biophysical characterization of the contribution of Ca<sub>v</sub>2.3 to the acrosome reaction. Using CATSPER-null sperm for Ca<sub>v</sub>2.3 patch clamp measurements will also help visualize the current mediated by Ca<sub>v</sub>2.3 by eliminating the major contribution of  $Ca^{2+}$  influx mediated by CatSper.

### PKD1, PKDREJ and PKD2

*PKD1* and *PKD2* encode polycystin 1 (PC1) and polycystin 2 (PC2 or TRPP2), respectively. PC1 is a putative transmembrane receptor, whereas PC2 can independently form a TRP-like ion channel<sup>163</sup>. Together PC1 and PC2 can form a heterotetrameric receptor channel complex and co-localize in renal cilia<sup>164-166</sup>. Mutations in either *PKD1* or *PKD2* can cause

autosomal dominant polycystic kidney disease (ADPKD), which results in the formation and expansion of collecting tubule-derived renal cysts<sup>164,167</sup>. Men with ADPKD also exhibit an increased rate of infertility and necrospermia<sup>168-170</sup>. Additionally, immotile sperm that lack the two central axoneme microtubules (9+0) have been identified in infertile men with PKD<sup>170</sup>.

A *PKDI* homologue, *Pkdrej*, has also been implicated in fertility and has been studied in some detail in mice<sup>171</sup>. The *PKDI* homologue *PKDREJ* (polycystic kidney disease and receptor for egg jelly) was found to be expressed specifically in the testicular tissues of humans<sup>172</sup> and mice<sup>173</sup> and localizes to the plasma membrane of the mouse sperm head<sup>173</sup>. However, no direct evidence supports a role for PKDREJ in acrosomal exocytosis. Mice that are homozygous for a disrupted *Pkdrej* allele (*Pkdrej*<sup>tm/tm</sup>) are still fertile (TABLE 1) but less fertile than wild-type mice when compared using sequential mating trials or artificial insemination assays<sup>171</sup>. *Pkdrej*<sup>tm/tm</sup> sperm navigate the female reproductive tract in vivo with reduced efficiency compared with wild-type sperm, as shown by fewer sperm reaching the cumulus matrix surrounding the egg over the same amount of time, and are slower to develop the ability to undergo a zona pellucida-induced acrosome reaction under capacitating conditions. However, as a comparable proportion of *Pkdrej*<sup>tm/tm</sup> sperm and wild-type sperm developed hyperactivated motility over the same time course in vitro, PKDREJ has been suggested to be a chronoregulator, not a master regulator, of capacitation<sup>171</sup>.

## CFTR and functionally related transporters

Cystic fibrosis transmembrane regulator (CFTR) is a Cl<sup>-</sup>-permeable and HCO<sub>3</sub><sup>-</sup>-permeable anion channel. HCO<sub>3</sub><sup>-</sup> conductance through CFTR is low compared with that of Cl<sup>-</sup><sup>174,175</sup>. Mutations in the gene encoding CFTR are the cause of cystic fibrosis, an autosomal recessive, monogenetic disease that results in severe phenotypes including progressive lung disease. Another cystic fibrosis phenotype is male infertility, which affects 97–98% of men with cystic fibrosis owing to congenital bilateral absence of the vas deferens (CBAVD)<sup>176</sup>. However, mutations in *CFTR* have also been found in 8.9% of otherwise healthy men with reduced sperm counts and/or with poor sperm quality; for example, sperm that exhibited reduced motility or abnormal morphology, or both<sup>177</sup>.

Xu and colleagues characterized the role of CFTR in sperm using a heterozygous (*Cftr*<sup>+/-</sup>) cystic fibrosis mouse model<sup>178</sup> — a heterozygous mouse was used because the homozygotes rarely survive past weaning<sup>179</sup>. When compared with wild-type sperm, fewer *Cftr*<sup>+/-</sup> sperm could achieve capacitation; they exhibited decreased membrane hyperpolarization and cAMP production in response to HCO<sub>3</sub><sup>-</sup>, decreased motility and reduced fertility in vitro and in vivo<sup>178</sup>. In accordance with these data, CFTR inhibitor-172, a CFTR channel blocker, inhibits the acrosome reaction, HCO<sub>3</sub><sup>-</sup>-dependent increases in intracellular pH and membrane hyperpolarization, and inhibits an HCO<sub>3</sub><sup>-</sup>-dependent increase in cAMP concentration in mouse sperm<sup>178,180</sup>. Likewise, another CFTR inhibitor, diphenylamine-2-carboxylic acid (DPC), also blocks capacitation-associated hyperpolarization (and seems to inhibit capacitation in general) as well as Cl<sup>-</sup> influx<sup>181</sup>. By contrast, genistein, which activates CFTR, induced hyperpolarization under noncapacitating conditions in mouse

sperm and resulted in  $\text{Cl}^-$  influx: intracellular  $\text{Cl}^-$  was measured using *N*-(ethoxycarbonylmethyl)-6-methoxyquinolinium bromide (MQAE), a fluorescent  $\text{Cl}^-$  probe<sup>181</sup>. External  $\text{Cl}^-$  was required for the genistein-induced hyperpolarization<sup>181</sup>. This external  $\text{Cl}^-$  dependence of capacitation was also observed in guinea pig sperm<sup>182</sup>, suggesting a role of CFTR in sperm capacitation in guinea pig. In the same study, the  $\text{Cl}^-/\text{HCO}_3^-$  exchanger known as solute carrier family 26, number 3 (SLC26A3) was suggested to work together with CFTR by pumping out  $\text{Cl}^-$  that had entered through CFTR<sup>182</sup>. In support of this interaction, mouse SLC26A3 and CFTR were co-immunoprecipitated, together with other members of solute carriers (SLC26A6 and SLC9A3R1)<sup>183</sup>. This CFTR-SLC interaction model was supported by a study showing that pharmacological inhibition or blocking SLC26A3 with an antibody could inhibit the acrosome reaction and hyperactivated motility<sup>182</sup>. Furthermore inhibition of CFTR and SLC26A3 was found to inhibit db-cAMP-induced  $\text{Cl}^-$  influx, capacitation-associated hyperpolarization and pH change, and  $\text{HCO}_3^-$  was found to induce hyperpolarization<sup>183</sup>.

Human mutations in *SLC26A3* can result in congenital chloride diarrhoea (CLD), and men with CLD often also have subfertility and oligoasthenoteratozoospermia<sup>184</sup>. Accordingly, *Slc26A3* knockout in mice also results in CLD and subfertility<sup>185</sup>. As both CFTR and SLC26A3 are expressed in the epithelial cells of the male reproductive tract as well as in the sperm cells themselves<sup>184,186</sup>, their respective roles in sperm versus other fertility-related processes can be difficult to determine. In the future, assessing SLC26A3 protein expression and function in heterozygous *Cftu*<sup>+/-</sup> mice could inform its role in CFTR regulation and sperm capacitation.

Direct whole-cell patch recordings from mouse testicular sperm demonstrated a  $\text{Cl}^-$  component to the membrane current that is ATP-dependent and is stimulated by cAMP, cGMP and genistein, and inhibited by CFTR inhibitor-172 and DPC<sup>187</sup>. However,  $\text{Cl}^-$  current could still be recorded from CFTR loss-of-function (F508) mouse testicular sperm, although in F508 mice this current is less sensitive to cAMP and CFTR inhibitor-172 (REF.<sup>187</sup>). In the same study,  $\text{Cl}^-$  current was also recorded from mouse wild-type epididymal sperm, but the effects of the CFTR-modulating compounds observed were not as substantial as seen in the testicular sperm. Thus, further work is required to clarify the extent to which CFTR conducts  $\text{Cl}^-$  current in mature sperm cells.

The precise localization of CFTR within sperm also requires clarification. Immunocytochemistry studies have shown that CFTR is localized in the equatorial segment of human<sup>178,180</sup>, mouse<sup>178</sup> and guinea pig sperm cells<sup>182</sup> but have also demonstrated its presence in the midpiece of human and mouse sperm<sup>181,183</sup>. CFTR has also been found simultaneously in both the equatorial segment of mouse sperm heads and the sperm midpiece<sup>188</sup>. Likewise, immunocytochemistry has demonstrated that SLC26A3 localizes within the heads of guinea pig sperm<sup>182</sup> but was shown to localize to the midpiece of mouse sperm in a separate study<sup>183</sup>. This discrepancy highlights the importance of genetic studies including the generation of knockout mice, for example, in sperm studies, as sperm cells have been repeatedly shown to be prone to non-specific antibody binding.



To date, the mouse and human sperm ion channels and membrane transporters that are clearly implicated in male fertility have been largely localized in the flagella (FIG. 1a; TABLE 1). Identifying and investigating channels located in other compartments of the sperm cells, such as acrosomal channels, requires further studies using current state-of-the-art techniques that provide specificity, sensitivity and high resolution in time and space, as well as evidence from genetic studies in both mice and humans.

## Sperm Ca<sup>2+</sup> homeostasis

To support dynamic cellular Ca<sup>2+</sup> signalling and to prevent Ca<sup>2+</sup> intoxication, Ca<sup>2+</sup> must be efficiently cleared from the cytosol after its entry through Ca<sup>2+</sup>-permeable channels.

### PMCA4

Plasma membrane Ca<sup>2+</sup>-ATPases (PMCAs) are highly conserved Ca<sup>2+</sup> extrusion pumps that maintain low basal levels of intracellular Ca<sup>2+</sup> (REF.<sup>189</sup>). Two isoforms, PMCA1 and PMCA4 (encoded by *Atp2b1* and *Atp2b4*, respectively), are expressed abundantly in testis and highly conserved across species<sup>190</sup>. Genetic knockout of PMCA4 in mice results in a >90% reduction in total PMCA expression<sup>191</sup>, indicating that PMCA4 is the primary PMCA in sperm cells. PMCA4-deficient male mice produce sperm cells with normal morphology but are totally infertile owing to impaired sperm motility and defective hyperactivated motility<sup>191-193</sup> (TABLE 1). Surprisingly, PMCA4-deficient mouse sperm are able to bind to zona pellucida and fertilize eggs in vitro<sup>192</sup>, suggesting that infertility in vivo is probably due to inefficient sperm navigation in the female reproductive tract. Interestingly, PMCA4 is primarily localized in the sperm principal piece but, unlike CatSper, does not exhibit a distinct pattern of distribution on the flagellar surface<sup>30</sup>. Ultrastructural analysis has revealed that mitochondria from PMCA4-deficient sperm are more condensed than wild-type sperm when incubated under capacitating conditions<sup>191</sup>; this characteristic is indicative of Ca<sup>2+</sup> overload. Whether PMCA4 participates in regulating CatSper-relevant Ca<sup>2+</sup> signalling and how sperm mitochondria are involved in Ca<sup>2+</sup> signalling during capacitation remains to be clarified.

### Mg<sup>2+</sup> transporter

A Mg<sup>2+</sup> transporter, CNNM4 (also known as ancient conserved domain-containing protein 4, ACDP4) is highly expressed in mature ameloblasts and intestinal epithelia<sup>194</sup>. Interestingly, CNNM4-deficient male mice are almost infertile<sup>195</sup>. *Cnnm4*-null spermatozoa exhibit rapid motility decreases and fail to develop hyperactivated motility<sup>195</sup>. Mg<sup>2+</sup> levels are significantly increased in CNNM4-deficient mouse sperm, whereas Ca<sup>2+</sup> levels are not affected and removal of Mg<sup>2+</sup> from the medium can rescue *Cnnm4*-null sperm motility<sup>195</sup>, suggesting that abnormally high levels of Mg<sup>2+</sup> are detrimental to sperm function. In addition, *Cnnm4*-null sperm exhibit excessive P-Tyr with impaired Ca<sup>2+</sup> influx<sup>195</sup>, similar to the phenotype of CatSper-deficient sperm<sup>30</sup>. These results suggest that CNNM4 might be involved in sperm Ca<sup>2+</sup> homeostasis and/or could be functionally linked to CatSper. Additional manipulation to a *Cnnm2* allele, which encodes another CNNM family Mg<sup>2+</sup> transporter, to make CNNM2 non-functional rendered the resulting *Cnnm2*<sup>+/-</sup>/*Cnnm4*<sup>-/-</sup> male mice completely infertile. The *Cnnm2*<sup>+/-</sup>/*Cnnm4*<sup>-/-</sup> sperm exhibited a more severe

reduction in motility and a loss of BSA-induced  $\text{Ca}^{2+}$  response, compared with *Cnnm4*<sup>-/-</sup> sperm, suggesting that CNNM2 works together with CNNM4 to regulate intracellular  $\text{Mg}^{2+}$  homeostasis and male fertility<sup>196</sup>. Whether the  $\text{Mg}^{2+}$  efflux activities of CNNMs are associated with  $\text{Ca}^{2+}$  homeostasis and how they can potentially regulate sperm  $\text{Ca}^{2+}$  signalling await further investigation.

## Clinical implications

The manipulation of ion channels in order to affect fertility could be leveraged for clinical application; for example, to produce male contraceptives or fertility treatments. Additionally, male fertility can reflect overall health as the dysregulated expression of ubiquitous proteins implicated in non-reproductive diseases might also result in altered sperm function.

## Novel non-hormonal male contraceptives

Most contraceptive strategies have been developed for use by women, but interest in generating a novel male contraceptive remains. One strategy is the use of hormones to inhibit endogenous testosterone production and, as a result, to block spermatogenesis<sup>197</sup>. To bypass potential adverse effects associated with hormone use, non-hormonal contraceptives are more desirable.

Ion channels in sperm are attractive targets for the development of contraceptives and, conversely, for drugs that could be used to treat infertility. Like G-protein-coupled receptors, ion channels are good drug targets as they are implicated in a variety of pathophysiologies and present druggable sites at cell surfaces<sup>198,199</sup>. In fact, ~15% of current drug targets are ion channels<sup>199,200</sup>.

The sperm-specific nature of the CatSper ion channel complex means that targeting it should result in few unintended off-target effects. Some compounds have been identified that inhibit CatSper, but they are non-specific and also inhibit KSper with comparable potency and so are probably not specific enough to be used as contraceptives<sup>201</sup>. RU1968, a ligand of steroidal sigma receptors, has been shown to suppress progesterone-stimulated  $\text{Ca}^{2+}$  signalling and prostaglandin-stimulated  $\text{Ca}^{2+}$  signalling in human sperm<sup>202</sup>. RU1968 has also been shown to inhibit human CatSper with ~15-fold higher potency than human KSper, and not to inhibit mouse KSper at all<sup>201</sup>, demonstrating the specificity for CatSper inhibition. Hopefully RU1968 and other CatSper inhibitors can be used as a template for the design of drugs that could be used in contraception.

Many sperm ion channels have yet to be thoroughly explored as therapeutic targets at least partially (as has been asserted for  $\text{K}^{+}$  channel pharmacology) owing to difficulties in establishing robust, high-capacity functional assays that could be used to interrogate large chemical libraries for potential drugs with activity against not only the targets themselves but also related targets to determine the specificity of screening hits<sup>199</sup>. This difficulty has been somewhat overcome in sperm with the development of a high-throughput, automated screening platform to assess the effect of small molecules on human sperm motility and ability to undergo the acrosome reaction<sup>203</sup>. The drug library (ReFRAME) used in the study comprises drugs that have either been approved for use or have undergone preclinical

profiling<sup>204</sup>; thus, a hit in this type of screen could accelerate the search for a marketable drug for use as a male contraceptive or to modify fertility.

### Fertility as a proxy for overall health in men

The relationship between subfertility and overall health in men is becoming increasingly apparent<sup>205,206</sup>. For example, as ~10% of the genome is implicated in fertility, subfertility and/or infertility, phenotypes can indicate abnormalities in other biological processes, such as those resulting in fibrosis<sup>205</sup>.

In addition to genetic fertility associations, subfertility and infertility are also associated with developmental, lifestyle, oncological and cardiovascular disorders<sup>205</sup>. Lifestyle factors associated with infertility in men include obesity<sup>207</sup>, tobacco abuse<sup>208</sup> and stress<sup>208</sup>. For example, a large meta-analysis of 21 studies including 13,077 men showed a J-shaped association between the risk of abnormal sperm concentration and BMI, in which increased BMI was negatively correlated with semen sperm concentration<sup>207</sup>. It is well known that cancer treatment can impair male fertility<sup>209</sup>, and studies have now also supported a link between male infertility and the risk of developing testicular cancer<sup>210,211</sup>, suggesting that male infertility might serve as a marker of oncological risk. Thus, assessing fertility and sperm functionality could offer unique insights into overall male health: each sperm cell is designed to function outside of the male body as a single cell and is, therefore, particularly amenable to in vitro analysis<sup>212</sup>. However, although sperm analysis can be a convenient and non-invasive tool, male reproductive capability is generally not evaluated until late in a man's life, usually after failure to reproduce<sup>206,213</sup>. Thus, it is recommended that men who exhibit health issues such as cystic fibrosis or altered fertility should consider genetic counselling to better understand other health problems to which they might be predisposed and the risk associated with passing their genes onto their children.

**Specific notable mutations.**—Mutations in *CFTR* have been identified not only in men who exhibit cystic fibrosis but also in those with CBAVD and/or sperm of reduced quality<sup>176,177</sup>. Although cystic fibrosis is a recessive disease, mutagenesis of one *CFTR* copy can result in altered sperm parameters in mice<sup>178</sup> and could be associated with CBAVD in men<sup>214</sup>. Abnormal mucociliary clearance is associated with cystic fibrosis, although this primarily results from the abnormal biophysical properties of the airway mucus and not ciliopathy<sup>215</sup>. However, a relationship is seen between fertility and ciliopathies<sup>216</sup>.

Mutations in the genes encoding PKD1 or PKD2, which both localize to primary cilia, can result in ADPKD, which is associated with increased rates of infertility<sup>164,167-170</sup>. As the motility of cilia and flagella are both conferred by an axoneme, it is tempting to speculate that similar axonemal defects could result in similar phenotypes.

Because the CATSPER complex is a sperm-specific ion channel, mutations in *CATSPER* genes are less likely to cause more widespread health problems than alterations in other ion channels such as CFTR. For example, a *CATSPER* in-frame 6-bp deletion results in normal sperm motility in humans but failure to fertilize in IVF owing to defective hyperactivation and lack of Ca<sup>2+</sup> response to progesterone<sup>106,107</sup>. In such cases, intracytoplasmic sperm injection can be used to achieve fertilization and can result in clinical

pregnancy. Although mutations in *CATSPER* are most likely to affect fertility, they can be associated with other health problems, such as deafness. Deafness–infertility syndrome is a very rare syndrome characterized by both deafness and male infertility and is associated with homozygous deletions of *STRC* (which is expressed in the inner ear) and *CATSPER2* on chromosome 15q15 (REFS<sup>111,112</sup>).

These examples illustrate the importance of diagnosis of the underlying cause of male infertility. Mutations giving rise to channelopathies that can affect male fertility can also have more wide-ranging effects on other body systems that men might wish to be aware of before beginning to consider assisted reproductive technology (ART).

## Current controversies

A large body of evidence exists regarding the role of ion channels in sperm, but limitations of the experimental approaches used in the field must be taken into account and whether the findings from in vitro and/or animal studies are physiologically relevant and not species-specific must be considered before these data can be extrapolated to humans.

## Interpreting data from indirect approaches

Determining protein locations in sperm cells solely based on antibody detection has generated controversies in the field and needs to be considered critically. Studies have produced inconsistent data, including the specific localizations of some ion channels and receptors, including CFTR, SLC26A3 and TRPV4, in sperm cells.

The identification of CatSper as a primary  $\text{Ca}^{2+}$  channel in sperm illustrates the importance of genetic and/or other direct evidence, such as direct electrophysiological recording from sperm cells. Before the discovery of CatSper, the N-type and R-type voltage-gated  $\text{Ca}^{2+}$  channels,  $\text{Ca}_v2.2$  and  $\text{Ca}_v2.3$ , were thought to be the  $\text{Ca}^{2+}$  entry channels in sperm, as they had been immunologically detected in mouse sperm<sup>159,217</sup>. T-type voltage-gated  $\text{Ca}^{2+}$  currents were also suggested to contribute to the  $\text{Ca}^{2+}$  influx in spermatozoa, as the corresponding currents were recorded in mouse testicular sperm<sup>93,218,219</sup>. However, whole-cell patch clamping of mouse epididymal sperm cells, combined with genetic analysis, clarified that CatSper is the primary facilitator of  $\text{Ca}^{2+}$  influx in mature sperm and is specifically localized in the principal piece of the sperm tail<sup>13,18</sup>. Furthermore, T-type currents were shown to be diminished in spermatids and were not detected in mature sperm<sup>13,94</sup>, highlighting the importance of a direct approach. Immunological methods such as immunostaining and immunoblotting have also been used to demonstrate the expression of many neurotransmitter receptors in mammalian sperm, including receptors for norepinephrine, aspartate, serotonin, acetylcholine, GABA and glycine<sup>220,221</sup>. However, no neurotransmitter-mediated currents were detected by whole-sperm cell electrophysiological recordings when functional expression of the corresponding receptors was tested<sup>154</sup>, suggesting either a non-functional presence of these receptors or non-specific detection.

Studies that solely relied on pharmacological interrogation also illustrate the unreliable nature of indirect approaches. Sperm cells are particularly prone to non-specific antibody binding and are particularly amenable to chemical inhibitors and/or activators, probably

owing to their small dimensions and relative lack of cytoplasm (that is, their low copy number of the channels or receptors and large lipid surface-to-volume ratio), which demonstrates the risk of detecting artefacts in sperm cells when genetic or other more direct evidence (for example, electrophysiology data) is lacking.

### Compounding implications from in vitro studies

Various capacitation-associated changes have been described, including activation of cAMP/PKA<sup>141</sup>, increases in protein P-Tyr<sup>139</sup>, a rise in intracellular pH<sup>222</sup> and Ca<sup>2+</sup> (REFS<sup>223-226</sup>), membrane hyperpolarization<sup>227,228</sup>, and modulation of the lipid content of the sperm plasma membrane<sup>132</sup>. In particular, a substantial body of work has documented the involvement of P-Tyr in regulating sperm motility and fertility during sperm capacitation<sup>229</sup>. As a result, P-Tyr has been used as a hallmark of sperm capacitation for decades. However, mounting evidence now indicates that P-Tyr is not actually required for hyperactivated motility<sup>144</sup> or fertility<sup>142</sup> in mice. This new interpretation has arisen because sperm capacitation is typically studied using in vitro analyses that reflect an average value for entire sperm populations at a given time. Inducing capacitation in mouse or human sperm in vitro using buffer containing HCO<sub>3</sub><sup>-</sup> and BS A results in a heterogeneous population of sperm, in which as few as ~15% of mouse sperm are hyperactivated<sup>230</sup>; 2–14% of human sperm are acrosome-reacted<sup>231</sup>. Similar to the fact that in vitro experiments do not necessarily reflect the in vivo processes, contact with the zona pellucida was believed to induce the acrosome reaction, especially after an observation that the mouse acrosome reaction can be induced by zona pellucida sperm-binding protein 3 (ZP3; one of four glycoproteins that make up the zona pellucida) in vitro<sup>232</sup>. Subsequently, mouse spermatozoa were found to begin undergoing acrosome reaction in the isthmus region of the fallopian tube before arriving at the ampulla<sup>233,234</sup> and reacted spermatozoa were able to penetrate the zona in vivo<sup>235</sup>; the small number of mouse spermatozoa that arrive at the lumen of the ampulla and cumulus oophorus are all fully capacitated and acrosome-reacted in vivo<sup>236</sup>. Even so, determining the exact site of the acrosome reaction in other mammalian species awaits further technological development and the physiological functions of the acrosome reaction and P-Tyr are currently being re-evaluated.

Currently available tools based on flow cytometry combined with sorting have been used to detect and/or separate the in vitro capacitated heterogeneous sperm populations based on various capacitation parameters: P-Tyr<sup>35,237</sup>, acrosome reaction status<sup>238,239</sup> and membrane hyperpolarization<sup>74,240</sup>. However, the dynamics of how these parameters change in each individual sperm and the extent to which these parameters reflect the fertilizing ability of each sperm has not been established. The direct linking of molecular information with motility and fertilizing capacity awaits a new approach to be developed.

### Critical appreciation of species-specific regulation

Conclusions from studies using different animal models should be carefully interpreted for two reasons. First, it is becoming apparent that some controversies in the field are partly due to a lack of understanding of marked species-specific regulation. Studies that exemplify this are those that have investigated the different molecular mechanisms of CatSper and KSper activation: CatSper is universally activated by intracellular alkalinization as demonstrated by

sperm cells from sea urchin<sup>241</sup>, mouse<sup>13</sup>, rat<sup>28</sup>, human<sup>67</sup>, cow<sup>242</sup> and horse<sup>243</sup>; however, only human and primate CatSper currents are robustly potentiated by hormones such as progesterone<sup>66,67,113</sup>. Similarly, human KSper is more sensitive to Ca<sup>2+</sup> than to alkalization compared with mouse KSper. These discrepancies illustrate the divergence of molecular mechanisms that regulate CatSper and KSper across different species. In the future, these phenomena should be kept in mind, as other sperm ion channels and membrane receptors might have also diverged in a species-specific manner.

Second, the mouse model is limited in its ability to reflect the sperm biology of other species such as humans and domestic animals, at least in part because epididymal, not ejaculated, mouse sperm are usually used in in vitro experiments<sup>28,244</sup>. Ejaculated sperm cells are mixed with secretory factors from male glands and ultimately with other molecules and ions in the female reproductive tract. Conclusions drawn from in vitro experiments using mouse sperm must consider the absence of these factors. Future studies should ensure that in vitro experiments are combined with genetic modifications and/or that in vitro systems are supplemented with the appropriate secretory factors from the male and female reproductive tracts in order to better inform our understanding of species-specific regulation of ion channels and membrane receptors including CatSper.

## Future of the field

In order to better understand the regulation and role of ion channels in sperm, future studies must seek more evidence from in vivo studies and genetics evidence.

## Seeking more in vivo context

Sperm analysis is currently performed using functionally heterogeneous sperm populations; these populations are typically capacitated in vitro and include degenerating cells<sup>30</sup>. Researchers must consider that the population mean of these analyses of the changes undergone by the sperm is often presented and cannot be used to faithfully recapitulate the time-dependent and space-dependent changes that sperm undergo in the oviduct<sup>245,246</sup>, nor the physiological and signalling state of the small number of spermatozoa that reach the vicinity of, and fertilize, the eggs<sup>247,248</sup>. Consistent with this notion, a greater number of sperm are required for IVF than the number of sperm present at the site of fertilization in vivo<sup>249</sup>. Studies that better emulate in vivo conditions and/or the development of new systems that can directly link the molecular changes of individual cells to motility and fertilizing ability are research priorities in the future. Examples of such studies include motility-correlative molecular imaging of sperm cells<sup>30</sup>, probing acrosome reaction states of sperm from genetically modified mice encoding fluorescence proteins in the acrosome along the female reproductive tracts<sup>233,236,250</sup>, and electrophysiological recordings taken from capacitated human sperm<sup>147</sup>. Such studies, in which an effort is made to understand sperm function in a more physiologically relevant manner, are anticipated in the future.

## Identifying gene variants in human fertility

Decades of mouse genetics studies have resulted in an extensive list of genes encoding spermatozoan ion channels and transporters implicated in male infertility (TABLE 1).

However, only a few of these genes, for example CatSper-encoding genes or *CFTR*, have been implicated in human infertility, which is surprising considering advances in tools such as whole-exome analysis, but illustrates the practical difficulties of studying the inheritance of infertility traits in humans. Damaging mutations in CatSper might have occurred and, therefore, have been detected, with increased frequency owing to the large number of genes required to form this multi-subunit channel complex. Targeted genomic analysis of large cohorts with a specific functional characteristic (for example, infertile men with normal semen parameters but with deficits in Ca<sup>2+</sup> signalling or abnormal membrane potential) and analyses of large families that exhibit Mendelian inheritance of these infertility traits might accelerate the discovery of human variants in these genes. As our understanding has grown that clinically assessing male reproductive health could provide unique insights into the general health of a patient and his children, identification of human infertility genetic variants could serve as a guideline for counselling general health, as well as candidate drug targets for screening for contraception, and could also provide insights into drug design and mechanisms of action.

In the current era of ARTs, in which natural barriers to egg fertilization are removed, defining the genetic defects underlying infertility is of the utmost importance, as genetic mutations that lead to male infertility are able to be passed to the next generation. Understanding the long-term effect of using ARTs needs to be prioritized in research.

## Conclusions

Technical advances have substantially improved our understanding of the role of ion channels and membrane receptors in sperm function during fertilization. Current knowledge of the ion channels and membrane transporters found in sperm have been greatly shaped by gene knockout studies in mice and genetic evidence from humans. In particular, our understanding of the CatSper channel — its molecular and spatial organization, the regulatory mechanism by which it is regulated, and the CatSper-based signalling pathways that are required to trigger hyperactivated motility — are now fairly well understood. However, the field is not without controversies and debates, in particular regarding the way in which in vitro data can be applied in vivo, and how data collected from one species can be applied to others; these areas need to be addressed. Many features of ion channels are unique to sperm as the sperm-specific channel isoforms have unique properties that are not found in other cell types. Thus, understanding the ion channel mechanisms in sperm cells will advance our knowledge of causes of male infertility and should inspire improvements in assisted reproduction and the development of new contraceptives, and improve diagnoses of infertility.

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## Glossary

**CatSper**

Sperm-specific calcium channel.

**NHEs**

Sodium–hydrogen exchangers.

**HV1**

Proton channel.

**KSper**

Native sperm-specific potassium current/channel.

**Na<sup>+</sup>/K<sup>+</sup> ATPase**

Sodium–potassium adenosine triphosphatase; also known as the sodium–potassium pump.

**SLO3**

The mediator of KSper, which is also used as the name of the protein or channel expressed heterologously.

**Quadrilateral compartmentalization**

The four linear Ca<sup>2+</sup> signalling nanodomains.

**DSper**

Depolarizing channel of sperm.

**TRPV4**

Transient receptor potential cation channel subfamily V member 4.

**P2X2**

P2X purinoceptor 2.

**Ca<sub>v</sub>2.3**

R type, voltage-dependent, calcium channel,  $\alpha$  1 E subunit.

**PKD**

Polycystin, transient receptor potential cation channel, autosomal dominant polycystic kidney disease protein.

**PKDREJ**

Polycystin family receptor for egg jelly.

**CFTR**

Cystic fibrosis transmembrane conductance regulator.

**PMCA4**

Plasma membrane calcium ATPase 4.



**CNNM4**

Cyclin and CBS domain divalent metal cation transport mediator 4.

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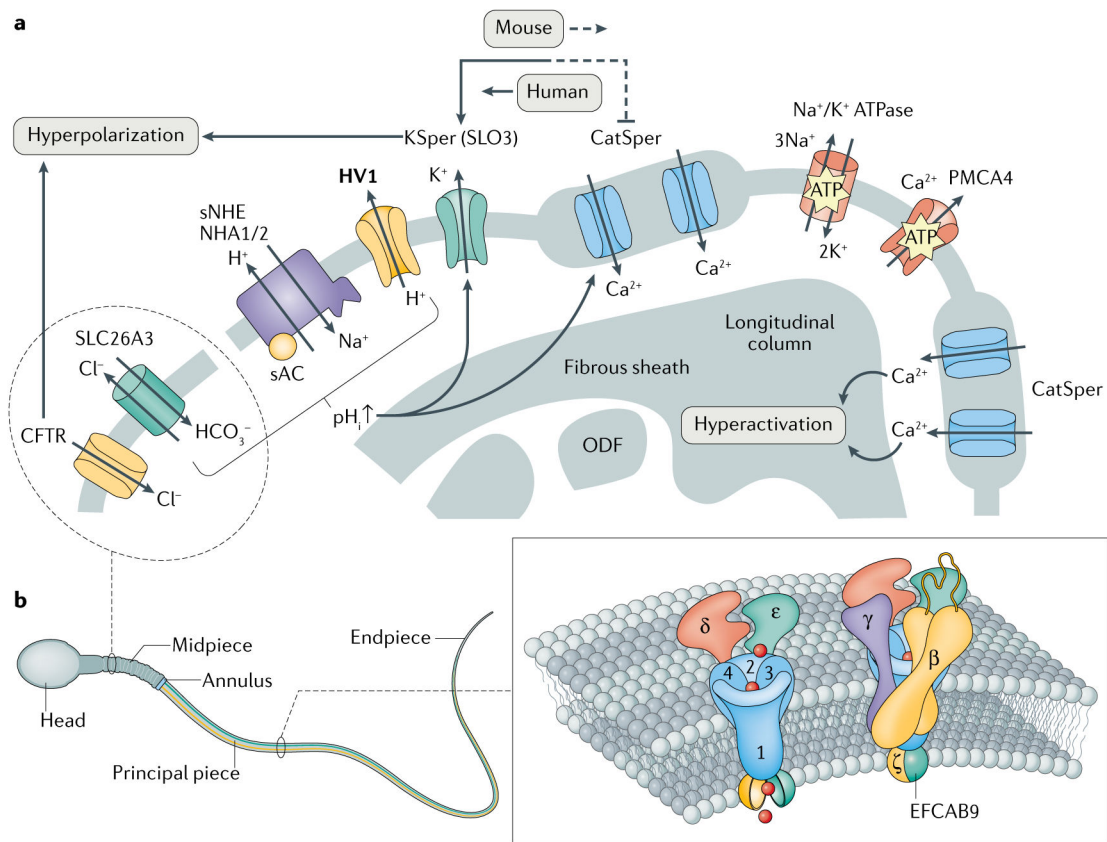
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### Key points

- Mammalian sperm cells undergo intracellular alkalization during their fertilization journey as they encounter a drastic extracellular pH change in the female reproductive tract.
- During capacitation, the sperm membrane potential hyperpolarizes, primarily via K<sub>Sper</sub> activation and K<sup>+</sup> efflux.
- Increases in intracellular Ca<sup>2+</sup> are required for inducing hyperactivated motility and acrosome reaction, two key physiological events essential for fertilization.
- CatSper, the multi-subunit Ca<sup>2+</sup> channel, is the predominant Ca<sup>2+</sup> entry pathway in sperm cells and organizes into linear Ca<sup>2+</sup> signalling nanodomains along the flagella.
- CatSper-mediated Ca<sup>2+</sup> signalling is integrated into other sperm capacitation signalling pathways including phosphorylation cascades.
- Improved understanding of spermatozoan ion channels and transporters will help elucidate the delicate and dynamic regulation of Ca<sup>2+</sup> homeostasis in sperm motility and fertility.



**Fig. 1 | Spermatozoan ion channels and membrane receptors.**

**a** | Partial cross-sectional view of the mammalian flagellar membrane that harbours ion channels and receptors. Intracellular alkalization is induced by sNHE and HV1 in mouse and human sperm, respectively. HCO<sub>3</sub><sup>-</sup> transporters such as SLC26A3, which interacts with the Cl<sup>-</sup>-permeable CFTR, can also carry HCO<sub>3</sub><sup>-</sup> into sperm cells from the female reproductive tract to trigger activation of soluble adenylyl cyclase (sAC). Consequently, in mouse sperm, the pH-sensitive channels K<sub>Sper</sub> and CatSper are activated and result in membrane hyperpolarization and Ca<sup>2+</sup> influx, respectively. The hyperpolarization contributed from both K<sub>Sper</sub> and CFTR during capacitation can inhibit further CatSper activation. By contrast, human K<sub>Sper</sub> exhibits less pH sensitivity and is robustly activated by Ca<sup>2+</sup> via CatSper. CatSper molecules are compartmentalized into four linear nanodomains along the principal piece of the tail; each nanodomain is apparently composed of two CatSper rows (two nanodomains are depicted in blue). Na<sup>+</sup>/K<sup>+</sup> ATPase α4 and PMCA4 function as transporters and sustain cytoplasmic ion homeostasis in sperm. As Na<sup>+</sup>/K<sup>+</sup> ATPase α4 contributes to membrane potential maintenance, both transporters are associated with Ca<sup>2+</sup> regulation, directly or indirectly. Unlike the other proteins that are localized in the principal piece, CFTR and SLC26A3 localize within the sperm head and/or midpiece (dashed line). HV1 expression is species-specific (in bold type). **b** | A CatSper channel complex linear nanodomain. CatSper is comprised of at least ten subunits including the pore-forming CATSPER1–4, the non-pore forming CATSPERβ, CATSPERγ, CATSPERδ and CATSPERε, and cytosolic CATSPERζ and EF-hand calcium-binding domain-containing protein 9 (EFCAB9). Note that large extracellular domains of the non-pore-

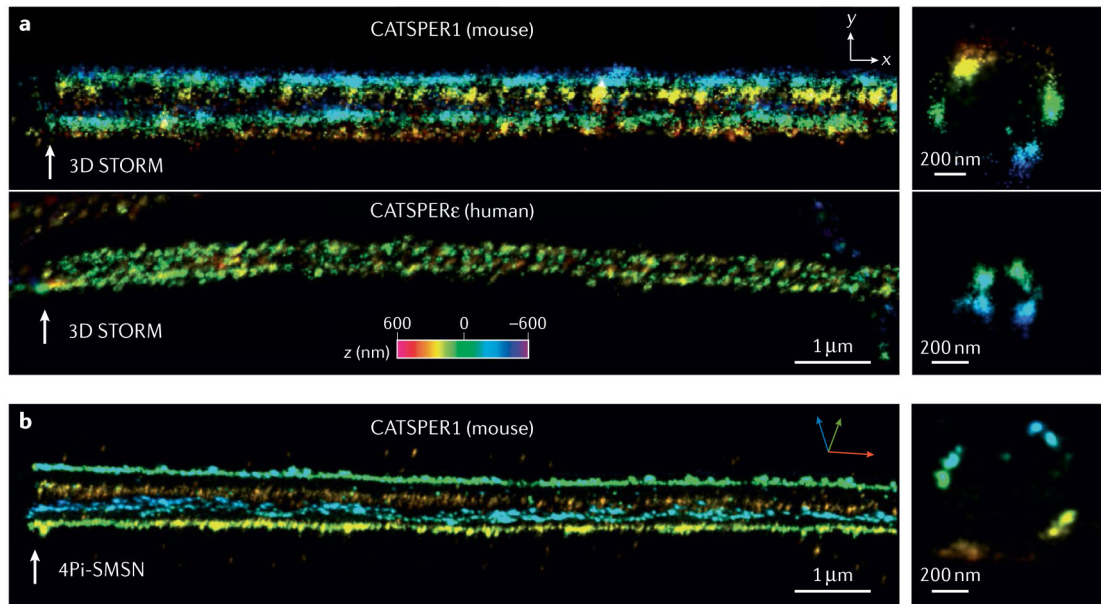
forming transmembrane subunits are predicted to surround the channel pore. ODF, outer dense fibre.

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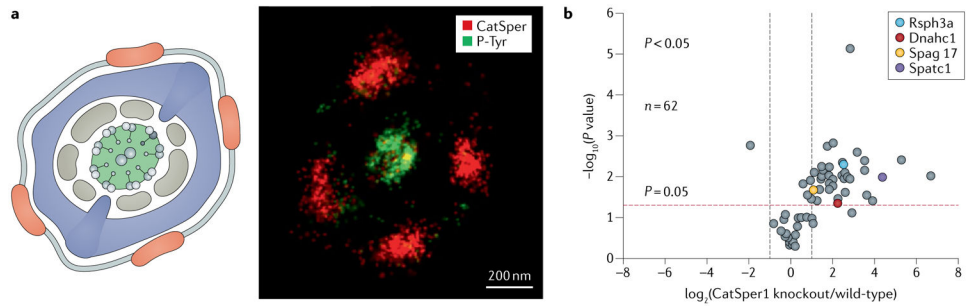
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**Fig. 2 | Quadrilinear CatSper  $\text{Ca}^{2+}$  signalling nanodomains in mammalian sperm.**  
**a** | 3D stochastic optical reconstruction microscopy (STORM) images. Distributions of CATSPER1 (mouse, upper) and CATSPER $\epsilon$  (human, lower) suggest conservation of the CatSper nanodomains in mammalian sperm. **b** | 3D 4Pi single-molecule switching nanoscopy (SMSN) images of CATSPER1 in mouse sperm. Note that each nanodomain is further resolved into two row structures. The colour encodes the relative distance from the focal plane along the  $z$  axis (color scale bar in  $x$ - $y$  projection). Adapted from (REF.<sup>29</sup>), CC BY 4.0 (<https://creativecommons.org/licenses/by/4.0/>).





**Fig. 3 | Capacitation-associated protein tyrosine phosphorylation.**

**a** | Tyrosine phosphorylation (P-Tyr) exhibits subflagellar localization. A cross-sectional view of the principal piece (left) shows CATSPER (red), fibrous sheath (blue), outer dense fibres (brown), and 9+2 axoneme (line structures in the centre). A two-colour 3D stochastic optical reconstruction microscopy (STORM) cross-sectional image of a capacitated wild-type spermatozoon (right) shows CATSPER1 (red) and P-Tyr (green). Note that P-Tyr is localized in the centre of the cross-section, corresponding to the axoneme. **b** | P-Tyr identified by quantitative whole-sperm proteome analysis from capacitated wild-type and *Catsper1*-null mice. Each protein is represented as a dot in a volcano plot of statistical significance ( $y$ -axis) against the average protein fold-change ( $x$ -axis) of *Catsper1*-null compared with wild-type spermatozoa. Four axonemal proteins (more than twofold change and  $P < 0.05$ ) are marked in colour.

Genetic studies of sperm ion channels and membrane transporters implicated in male fertility

Table 1 |

Gene	Tissue expression	Protein localization in sperm	Phenotypes of mutation	Species	Refs
<i>Ion channels</i>					
<i>Kcnul</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: impaired In vivo fertility: infertile	Knockout mouse	70,84
<i>Lrrc52</i>	Testis	ND	Spermatogenesis: ND Hyperactivated motility: ND In vitro fertilization: impaired In vivo fertility: subfertile	Knockout mouse	86
<i>Catsper1</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: impaired In vivo fertility: infertile	Knockout mouse	15,16
<i>CATSPER1</i>	Testis	Principal piece	Spermatogenesis: impaired Motility <sup>a</sup> : reduced In vitro fertilization: ND In vivo fertility: infertile	Human <sup>b</sup>	100 <sup>b</sup>
<i>Catsper2</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: ND In vivo fertility: infertile	Knockout mouse	17
<i>CATSPER2</i>	Testis	Principal piece	Spermatogenesis: impaired Motility <sup>a</sup> : reduced In vitro fertilization: ND In vivo fertility: infertile	Human <sup>c,d</sup>	102 <sup>c</sup> ; 103 <sup>c</sup> ; 104 <sup>d</sup> ; 111 <sup>c</sup> ; 216 <sup>c</sup>
<i>Catsper3</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: ND In vivo fertility: infertile	Knockout mouse	18,95
<i>Catsper4</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: ND In vivo fertility: infertile	Knockout mouse	18,95
<i>CATSPERE</i>	Testis	Principal piece <sup>e</sup>	Spermatogenesis: normal Motility <sup>a</sup> : normal In vitro fertilization: impaired In vivo fertility: infertile	Human <sup>f</sup>	29 <sup>e</sup> ; 106 <sup>f</sup>
<i>Catsperz</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired	Knockout mouse	29

Gene	Tissue expression	Protein localization in sperm	Phenotypes of mutation	Species	Refs
<i>Cacna1e</i>	Ubiquitous	Head	In vitro fertilization: impaired In vivo fertility: subfertile Spermatogenesis: ND Hyperactivated motility: normal In vitro fertilization: impaired In vivo fertility: mildly subfertile	Knockout mouse	160,161
<i>CFTR</i>	Multiple	Head/midpiece	Spermatogenesis: impaired Hyperactivated motility: ND In vitro fertilization: impaired In vivo fertility: impaired	Human, knockout mouse	178,180-183,188,251
<i>Pkdrej</i>	Testis	Head <sup>e</sup>	Spermatogenesis: normal Hyperactivated motility: normal but slow In vitro fertilization: ND In vivo fertility: fertile	Knockout mouse	171,172 <sup>e</sup> , 252
<i>Trpv4</i>	Ubiquitous	Head and tail	Spermatogenesis: normal Hyperactivated motility: normal but delayed In vitro fertilization: ND In vivo fertility: fertile	Knockout mouse	147,151,152,253
<b>Membrane transporters</b>					
<i>Atp1a4</i>	Testis	Principal piece	Spermatogenesis: normal Hyperactivated motility: reduced In vitro fertilization: impaired In vivo fertility: infertile	Knockout mouse	82
<i>Slc9a8</i>	Ubiquitous	Head (acrosome)	Spermatogenesis: impaired Motility <sup>a</sup> : reduced In vitro fertilization: ND In vivo fertility: infertile	Knockout mouse	56
<i>Slc9b1</i>	Testis	Principal piece	Spermatogenesis: normal Motility <sup>a</sup> : reduced In vitro fertilization: ND In vivo fertility: subfertile <sup>g</sup>	Knockout mouse	55
<i>Slc9b2</i>	Ubiquitous	Principal piece	Spermatogenesis: normal Motility <sup>a</sup> : reduced In vitro fertilization: ND In vivo fertility: subfertile <sup>g</sup>	Knockout mouse	55
<i>Slc9c1</i>	Testis	Principal piece	Spermatogenesis: normal Motility <sup>a</sup> : reduced In vitro fertilization: impaired In vivo fertility: infertile	Knockout mouse	53
<i>P2rx2</i>	Ubiquitous	Midpiece	Spermatogenesis: normal Hyperactivated motility: normal	Knockout mouse	154

Gene	Tissue expression	Protein localization in sperm	Phenotypes of mutation	Species	Refs
<i>Cnm4</i>	Ubiquitous	Principal piece	In vitro fertilization: normal In vivo fertility: mildly subfertile Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: impaired In vivo fertility: subfertile	Knockout mouse	195
<i>Ap2b4</i>	Ubiquitous	Principal piece	Spermatogenesis: normal Hyperactivated motility: impaired In vitro fertilization: normal In vivo fertility: infertile	Knockout mouse	191-193
<i>SLC26A3</i>	Multiple	Head/midpiece	Spermatogenesis: ND Hyperactivated motility: ND In vitro fertilization: ND In vivo fertility: subfertile	Human, knockout mouse	182,185,254

ND, Not determined.

<sup>a</sup>Motility was tested in the standard bath solution without capacitation components.

<sup>b</sup>Two separate insertion mutations in exon 1 (c.539–540insT and c.948–949insATGGC).

<sup>c</sup>Deaf infertility syndrome with a deletion encompassing *CATSPEP2* and *STRC* together.

<sup>d</sup>Copy number variation in the region of 43894500 to 43950000 in 15q15.3 encompassing a heterozygous deletion of *CATSPEP2*.

<sup>e</sup>Protein localization information only.

<sup>f</sup>Homozygous in-frame 6-bp deletion in exon 18 (c.2393\_2398delCTATGG).

<sup>g</sup>*Slc9b1/Slc9b2*-double-knockout mice are completely infertile.