

Chapter 2

The Acrosomal Matrix

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2.1 Introduction

The acrosome is a single exocytotic vesicle overlying the nucleus of most animal sperm. While the sperm of some taxa (e.g., teleost fish) do not possess an acrosome, it is generally thought that mammalian sperm acrosomes may perform two principal exocytosis-dependent functions: (1) to serve as a site for the sequestration and release of proteins required for binding to and penetration of the zona pellucida and (2) to enable the sperm to fuse with the oolemma. For the past several decades, research has been largely guided by the concept that mammalian sperm acrosomes must be intact to bind to the zona pellucida prior to undergoing exocytosis, enabling the sperm to penetrate this vestment and fertilize the egg (Wassarman et al. 2004). Although this widely accepted dogma has been challenged on many levels, recent experiments utilizing modern, highly sophisticated imaging techniques have enabled investigators to track the acrosomal status of individual fertilizing spermatozoa. Using genetically engineered mouse strains that express green fluorescent protein in their acrosomes, these experiments have shown that almost all fertilizing mouse sperm have undergone acrosomal exocytosis prior to the time they penetrate the zona pellucida (Jin et al. 2011).

These contemporary experiments raise new questions of how the acrosome forms during spermatogenesis and functions in the fertilization process. Specific

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issues that need to be addressed include (1) the contribution of acrosomal biogenesis to the fertility of individual sperm, (2) the timing of acrosomal exocytosis *in vivo*, (3) the role of acrosomal compartmentalization (including structurally distinct domains within the vesicle and biochemical segregation of overlapping soluble materials and insoluble acrosomal matrix compartments), and (4) the functions of specific proteins in fertilization. While recent results challenge or even contradict experiments that were based upon previous reductionist paradigms, many of the older studies were carefully designed and yielded a solid body of evidence that should be reevaluated from a more holistic standpoint to see whether a reinterpretation sheds new light on the question of the physiological role of the acrosome in fertilization.

In a previous review, we focused on the structure and function of the sperm acrosomal matrix in fertilization, where we espoused three principal points: (1) the acrosome is not a “bag of enzymes,” simply comprised of soluble substances that are rapidly released when the outer acrosomal membrane fuses with the plasma membrane, (2) acrosomal exocytosis is not an “all-or-none” event, where the acrosome is either “intact” or “reacted” but instead incorporates intermediate steps involving the progressive loss of acrosomal matrix material, and (3) the acrosomal matrix is a molecular scaffold assembly that is dismantled by a self-regulated mechanism driven, in part, by proteolysis (Buffone et al. 2008). We will not recapitulate these points in this review, but we will discuss them in light of the many new findings from studies using genetically modified mice.

It is abundantly clear that fertilization is a complicated process that requires models to extend from a cluster of hundreds or thousands of sperm to the single-cell level. Given the heterogeneous nature in the structure and physiology in a population of sperm, it is not surprising that aspects of acrosomal structure, including the matrix, could depend on events during spermatogenesis and could impact function at several steps in the female reproductive tract or the cumulus–oocyte complex. Furthermore, the heterogeneity in the timing and location of acrosomal exocytosis in individual cells present within a population of spermatozoa is not yet fully understood. Thus, it is a challenge to sort out real population cell behaviors (e.g., disruption of cumulus cells) from irrelevant or deleterious cell behaviors (spontaneous exocytosis, cell demise). Perhaps the most important question is, what is the location and timing of these events in the one sperm that successfully fertilizes the egg? We must also recognize that there might be multiple means (pathways) to an end (fertilization). In light of the contemporary imaging studies indicating that most mouse sperm that fertilize mouse oocytes enclosed by cumulus masses have lost the green fluorescent protein marker of intact acrosomes (Jin et al. 2011; Hirohashi et al. 2011), we recognize that work in this area is still very active so it would be premature to attempt to develop a consensus model. Nevertheless, it is useful at this time to consider results utilizing another modern approach: gene knockouts in mice. Toward this goal, recently developed, genetically modified mouse models provide valuable new insights into the molecular mechanisms of mammalian fertilization involving the sperm acrosome.

2.2 Gene Knockout Studies in Mice: What They Do and Do Not Tell Us About Acrosomal Matrix Function

The knockout of a specific reproductive gene in the mouse is usually engineered with an explicit outcome (hypothesis) in mind: to demonstrate the importance of a particular protein by interfering with fertility. However, the resulting phenotypes are often different and/or are more complicated than expected. In many cases, knockouts of individual genes encoding acrosomal matrix proteins have not caused sterility; this outcome can be explained away with the understanding that sperm function and fertilization are critical for survival of a species, so it makes sense to have built-in redundancy. Nevertheless, closer inspection of some of these “fertile” knockouts has revealed important insights into the functions of acrosomal matrix and other fertilization proteins. Also, knockouts of multiple proteins can address questions of complementary proteins in a complex system. As the body of literature involving genetically engineered mice has grown, there have been many surprising new observations that force us to think more broadly and to ask different questions than had previously been considered. For example, we now know not to assume that acrosomal exocytosis will only occur on the ZP, and we also understand proteins thought to be involved in sperm–egg interactions can have other functions such as acrosome formation, transiting the uterus and oviduct, etc. Thus, we feel the literature is rich with data from gene knockouts, and this review is an attempt to bring both widely known and understated results to the reader’s attention with hope that new avenues for research will come to light (Table 2.1).

2.3 Acrosome Formation/Globozoospermia

The function of the sperm acrosome is dependent upon its structure. The sperm of each species has a characteristic configuration with defined acrosomal shapes that are determined, in part, by the underlying nuclear morphologies. However, each acrosome shares the common features of an inner acrosomal membrane (IAM) that is laminated to the nuclear foundation, a fusogenic outer acrosomal membrane (OAM) that is closely apposed to the plasma membrane over the acrosome, and an equatorial segment (ES), which forms the posterior acrosomal margin where the IAM and OAM meet. Acrosomal formation begins during late meiosis when small, proacrosomal vesicles form in pachytene spermatocytes (Bloom et al. 1975; Holstein and Roosen-Runge 1981; Anakwe and Gerton 1990). After the meiotic divisions, the resulting haploid spermatids inherit the proacrosomal vesicles and continue acrosome biogenesis in earnest. Proacrosomal vesicles eventually coalesce into a single acrosomal vesicle that then attaches to and begins to spread over the future anterior segment of the nucleus. The points of contact between the developing acrosome and spermatid nucleus define the prospective IAM. The acrosome and the nucleus continue developing throughout spermiogenesis until

Table 2.1 Mouse gene knockout models of acrosomal matrix and associated proteins

Mouse gene name	Common protein names	Proposed function	Observed KO phenotype	References
Acrosome formation/globozoospermia				
<i>Aggl</i>	Hrb	Acrosomal vesicle fusion	Infertile males; globozoospermia; proacrosomal vesicles do not fuse; acrosomal development stops at step 2	Kang-Decker et al. (2001) and Kierszenbaum et al. (2004)
<i>Csnk2a2</i>	Casein kinase II α' catalytic subunit	Head morphology, acrosome morphology	Infertile males; globozoospermia	Xu et al. (1999)
<i>Dpy19l2</i>		Attachment of acrosomal vesicle to nucleus/acroploxome	Infertile males; acrosomal vesicle fails to attach to nucleus, acrosome eliminated	Pierre et al. (2012)
<i>Galnt3</i>	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3	Coalescence of proacrosomal vesicles into acrosome	Infertile males	Miyazaki et al. (2012)
<i>Gmcll</i>	Germ cell-less (Drosophila)	Acrosome formation, chromatin condensation	Subfertile males; abnormal acrosome/head morphology	Kimura et al. (2003)
<i>Gopc</i>	Golgi-associated PDZ- and coiled-coil motif-containing protein	Head morphology and mitochondrion assembly	Infertile males; globozoospermia and mitochondrial defects	Yao et al. (2002)
<i>Ical1</i>	Islet cell autoantigen 1-like protein	Head morphology	Subfertile males; globozoospermia	He et al. (2015)
<i>Pafah1b1</i>	Platelet-activating factor acetylhydrolase, regulatory subunit	Acrosome/head formation	Infertile males; Abnormal acrosome morphology	Nayernia et al. (2003)
<i>Pick1</i>	Protein interacting with C kinase 1	Vesicle trafficking from Golgi to the acrosome	Infertile males; globozoospermia	Xiao et al. (2009)
<i>Pla2g3</i>	Phospholipase A2, group III	Acrosome and axoneme formation	Subfertile males; multiple abnormal head & tail morphologies	Sato et al. (2010)
<i>Sec23ip</i>	Sec23 interacting protein	Acrosomal vesicle fusion/formation	Subfertile males; globozoospermia, acrosomal vesicle does not form	Arimitsu et al. (2011)

<i>Spacal</i>	Sperm acrosome associated 1 protein	Acrosome attachment and lateral spreading over nucleus	Infertile males; globozoospermia, acrosome fails to form	Fujihara et al. (2012)
<i>Tbc1d20</i>	Tbc-1	Acrosomal vesicle formation	KO but no fertility info reported; gene mutation causes disorganized acrosomal granules, arrest at step 3	Sotomayor and Handel (1986) and Liegel et al. (2013)
Zona interacting proteins				
<i>Acr</i>	Proacrosin	Acrosomal matrix disassembly	Fertile males; slow acrosomal matrix dispersal	Baba et al. (1994) and Adham et al. (1997)
<i>Zan</i>	Zonadhesin	Species-specific sperm-egg recognition	Fertile but species-specific gamete recognition affected	Tardif et al. (2010)
<i>Zp3r</i>	Sp56, AM67	Zona binding?	Fertile males	Muro et al. (2012)
<i>Zpbp1</i>	Zona pellucida-binding protein 1	Acrosome formation, compaction	Infertile males; globozoospermia, acrosomal dysplasia	Lin et al. (2007)
<i>Zpbp2</i>	Zona pellucida binding protein 2	Acrosome formation	Subfertile males; abnormal head/acrosome morphology	Lin et al. (2007)
Oocyte cumulus complex and zona pellucida transit				
<i>Acr1</i>		ZP/cumulus transit	Fertile males with reduced litter size; IVF failure	Kawano et al. (2010)
<i>Prss21</i>				
<i>Acrbp</i>	SP32	Regulate proacrosin activity	No knockout published	Kanemori et al. (2013)
<i>Hyal5</i>	Hyaluronidase 5	Digest hyaluronic acid	Fertile males	Kimura et al. (2009)
<i>Npix2</i>	AM50, Narp/NP2	Acrosomal matrix structure	Fertile males; no analysis of sperm	Bjartmar et al. (2006)
<i>Pcsk4</i>	Proprotein convertase subtilisin 4/Kexin type 4	Processing/activation of sperm/acrosomal proteins, capacitation, acrosomal exocytosis, ZP binding	Infertile males; abnormal acrosomal morphology	Mbikay et al. (1997), Gyamera-Acheampong et al. (2006) and Tardif et al. (2012)
<i>Prss21</i>	TESP5, testisin	ZP/cumulus transit	Fertile males; in vitro assays showed motility, morphology, and IVF problems	Netzel-Arnett et al. (2009)
<i>Serpina5</i>	Protein C Inhibitor	Sertoli cell-germ cell interactions	Infertile males; premature release of spermatogenic cells, abnormal sperm motility morphology, IVF	Uhrin et al. (2000) and Uhrin et al. (2007)

(continued)

Table 2.1 (continued)

Mouse gene name	Common protein names	Proposed function	Observed KO phenotype	References
<i>Spam1</i>	PH-20	Cumulus transit	Fertile males	Baba et al. (2002)
<i>Spam1/Acr</i>		Work cooperatively for cumulus transit	Delayed cumulus transit	Zhou et al. (2012)
<i>Spam1/Prss21</i>		Work cooperatively for cumulus transit	Delayed cumulus transit	Zhou et al. (2012)
Inner acrosomal membrane proteins				
<i>Cd46</i>	Cellular differentiation marker 46	Stabilize acrosomal membranes, no role in complement protection	Fertile males; enhanced fertilization via rapid capacitation and spontaneous acrosomal exocytosis	Inoue et al. (2003)
<i>Spaca4</i>	SAMP14	Persists following acrosomal exocytosis	No knockout published	Shetty et al. (2003)
Equatorial segment proteins				
<i>Eqtn</i>	Equatorin	Fusion of outer acrosomal and plasma membranes anterior to equatorial segment	Males subfertile and have reduced litter size; sperm do not undergo acrosomal exocytosis	Wolkowicz et al. (2003) and Hao et al. (2014)
<i>Spesp1</i>	Sperm equatorial segment protein 1	Stability of equatorial segment membranes	Males subfertile; fusion of outer acrosomal and plasma membranes in equatorial segment	Fujihara et al. (2010)

they coordinately achieve structures similar to those found in ejaculated sperm. Final steps of metamorphosis occur during epididymal transit.

In recent years, the molecular mechanisms supporting the transmigration of a round spermatid acrosome to the form seen in mature testicular spermatozoa have been elucidated by many studies utilizing mutations or gene knockouts in the mouse. One of the earliest examples of the effect of acrosomal dysplasia is the blind-sterile mouse, whose round spermatids exhibit disorganized proacrosomal granules with no further development beyond step 3 of spermiogenesis (Sotomayor and Handel 1986). This phenotype is caused by mutations in the *Tbc1d20* gene, which encodes a GTPase-activating protein specific for Rab1 and Rab2 small GTPase families (Liegel et al. 2013). Another GTPase-activating protein, AGFG1 (also known as Hrb), specific in this case for ARF proteins, shows a similar phenotype (Kang-Decker et al. 2001; Kierszenbaum et al. 2004). Vesicle trafficking is key to formation of the acrosome since mutations affecting PICK1, the Golgi-associated PDZ- and coiled-coil motif-containing protein (GOPC), the casein kinase II α' catalytic subunit (CSNK2A2), and islet cell autoantigen 1-like protein (ICA1L) cause sterility or subfertility in males (Xu et al. 1999; Yao et al. 2002; Xiao et al. 2009; He et al. 2015). Mutations affecting SEC23IP, a phospholipase A1-like protein that interacts with Sec23, also cause male subfertility by interfering with acrosome biogenesis and leading to round-headed spermatozoa (Arimitsu et al. 2011). Spermatids of infertile male mice lacking GalNAc transferase 3 (GALNT3) produce proacrosomal vesicles of various sizes, which attach to the nuclear envelope but do not coalesce to form a single acrosomal vesicle, apparently causing oligoasthenoteratozoospermia through a severe reduction of mucin-type O-glycans, impaired acrosome formation, and increased apoptosis (Miyazaki et al. 2012). Acrosomal dysplasia does occur in cases where certain acrosomal matrix components such as zona pellucida-binding protein 1 (ZPBP) or 2 (ZPBP2) are missing (Lin et al. 2007).

Abnormalities in nuclear envelope structure can also interfere with the ability of the acrosome to form properly. Disruption of the mouse homologue (*Gmcl1*) of the *Drosophila melanogaster* germ cell-less gene causes insufficient chromatin condensation and abnormal acrosome structures in mutant sperm (Kimura et al. 2003). When the inner nuclear membrane protein DPY19L2 is absent, the nuclear dense lamina as well as the junction between the acroplaxome and the nuclear envelope are destabilized. As a result, the linking of the acrosome and the manchette to the nucleus is blocked; this disrupts vesicular trafficking, causes sperm nuclear shaping to fail, and leads to the elimination of the unbound acrosomal vesicle (Pierre et al. 2012). In elongating spermatids, abnormalities of anterior head shaping involving CSNK2A2 or the sperm acrosome-associated 1 protein (SPACA1) also lead to globozoospermia (Xu et al. 1999; Fujihara et al. 2012). Spermatids with mutant PFAH1B1 protein (the regulatory subunit of platelet-activating factor acetylhydrolase 1b) are sterile because the spermatids fail to form acrosomes properly and exhibit nuclei distorted in size and shape (Nayernia et al. 2003).

Final maturation of the sperm acrosome occurs within the epididymis. The classic case is the morphological alteration in the acrosome of guinea pig sperm

where the head changes drastically from a planar, tennis racket-shaped structure to a form reminiscent of a cupped hand or xistera, the handheld basket used in jai alai, as the sperm transit from the testis to the cauda epididymis (Fawcett and Hollenberg 1963). Another example is seen in the post-testicular development of spermatozoa of the tammar wallaby where the immature, uncondensed acrosome has a distinctive “thumbtack” appearance that is remodeled into a streamlined, condensed structure on the sperm head (Setiadi et al. 1997). Phospholipase A2, group III (PLA2G3), which is secreted in the proximal (caput) region of the mouse epididymis, is reported to have a role in epididymal maturation of sperm (Sato et al. 2010); however, the *Pla2g3* gene is also expressed in the testis, so it is difficult to attribute the subfertility of *Pla2g3*^{-/-} males to the lack of the *Pla2g3* protein in the epididymis rather than the testis. The molecular mechanisms for these structural alterations are not fully understood but could be due, in part, to changes in protein disulfide cross-links, oligosaccharide modifications, and/or repackaging of the acrosomal components (Anakwe et al. 1991; Setiadi et al. 1997; Fléchon 2015).

2.4 Zona Interacting Proteins

For many years, the prevailing notion was that sperm bound to the zona pellucida of unfertilized eggs through cell surface receptors or binding proteins present on the exterior of acrosome-intact sperm. Biochemical studies in the mouse led to the identification of ZP3R (originally called “sp56”) as the putative ZP3 receptor protein (Bleil and Wassarman 1990; Cheng et al. 1994; Bookbinder et al. 1995). Subsequent experiments established that ZP3R and its guinea pig homologue are actually components of the acrosomal matrix (Foster et al. 1997; Kim et al. 2001). ZP3R monomers are assembled into a multimeric complex in the particulate acrosomal matrix. During the course of spontaneous acrosomal exocytosis and coincident with release of ZP3R from the acrosome under capacitating conditions, the monomer is proteolytically processed from 67,000 M_r to 43,000 M_r (Buffone et al. 2009). The processing of ZP3R is dramatically reduced when sperm are incubated under non-capacitating conditions. The cleavage probably takes place in complement control protein domain (CCP) 6 or the bridge region between CCP6 and CCP7, which is not present in the guinea pig orthologue AM67. This cleavage has functional consequences since the cleaved form of ZP3R does not bind to unfertilized eggs. However, although ZP3R binds the zona pellucida, the elimination of ZP3R in the mouse by homologous recombination did not affect fertility in the nullizygous mice (Muro et al. 2012).

Other candidates proposed as proteins responsible for zona pellucida binding also turned out to be components of the acrosomal matrix. Zona-binding proteins include the homologues known as sp38 [boar (Mori et al. 1993)], IAM38 [bull (Yu et al. 2006)], ZPBP and ZPBP2 [mouse (Lin et al. 2007)]. In the pig, this protein competes with proacrosin for binding to the zona pellucida (Mori et al. 1993). As discussed previously, elimination of the *Zpbp* and/or *Zpbp2* genes

by homologous recombination results in infertility or subfertility in male mice due to multiple effects on acrosome biogenesis and other spermatogenic defects leading to dysfunctional spermatozoa (Lin et al. 2007). The serine zymogen proacrosin is another zona-binding protein (Howes et al. 2001; Furlong et al. 2005), but, as described below, elimination of the *Acr* gene by homologous recombination does not cause infertility in mice (Baba et al. 1994; Adham et al. 1997). Zonadhesin (ZAN) was originally identified biochemically as a binding protein of the porcine zona pellucida (Hardy and Garbers 1995) and then demonstrated to be associated with the acrosomal matrix and membranes (Bi et al. 2003). Homologues in other species were identified and a gene knockout of *Zan* was performed in the mouse, which does not affect fertility but, instead, enables the mouse sperm to bind more effectively to pig, cow, and rabbit ZP without enhancing attachment to the mouse ZP, essentially eliminating the filter for the species-specific interactions between the mutant mouse sperm and oocytes of other animals (Tardif et al. 2010). From the studies performed to date, it is clear that there are several zona pellucida-binding proteins present within the acrosomal matrix of sperm. Some of these are involved in essential steps of acrosome biogenesis such as ZPBP and ZPBP2. However, the elimination of other zona-binding proteins by homologous recombination does not affect fertility per se but may yield a subtler phenotype such as a loss of species-specific sperm–zona binding or the slower dissipation of the acrosomal matrix when exocytosis is initiated, leading to a delayed rate of fertilization. These traits appear to be redundant, yet they likely impart a competitive advantage to sperm from wild-type mice when one or more males mate with the same female.

2.5 Oocyte Cumulus Complex and Zona Pellucida Transit: Proteases, Regulatory Proteins, and Binding Proteins

Mammalian oocytes are surrounded by two formidable barriers to spermatozoa: the oocyte cumulus complex (OCC) that consists of multiple layers of cumulus cells with an extracellular matrix rich in hyaluronic acid and the ZP that is comprised of three or four glycoproteins (Litscher and Wassarman 2014). The sperm surface and acrosome contain several proteases that are purported to have a role in at least some limited digestion of the OCC extracellular matrix and ZP, but it remains unclear whether these proteases are essential for sperm to transit these barriers or whether mechanical forces produced by the sperm tail are sufficient (Bedford 1998; Litscher and Wassarman 2014). In addition, the acrosomal matrix contains numerous proteins that can bind to the ZP and presumably enable a spermatozoon to burrow into the ZP by the combined mechanical forces of the tail and limited enzymatic digestion by proteases. Gene knockouts of individual and multiple proteases have provided some important and unforeseen insights into this process. Indeed, the pattern of surprising knockout results was set by perhaps the first knockout in this field—the proacrosin (ACR) gene. For many years, proacrosin was purported to be

the main sperm protease for ZP penetration, but, as discussed previously, gene knockout experiments showed that proacrosin was not essential for zona penetration as mice were fertile and proacrosin-null sperm were able to transit the ZP (Baba et al. 1994). Further study showed that proacrosin-null sperm dispersed acrosomal contents more slowly than wild-type sperm, indicating that the role of acrosin was to disperse the acrosomal matrix (Yamagata et al. 1998). This also inspired research to identify novel acrosomal proteases involved in OCC and ZP transit.

PRSS21 (TESP5, testisin) is a candidate serine protease to complement acrosin; it is a GPI-linked protein on the surface of the sperm head, cytoplasmic droplet, and midpiece of the tail (Honda et al. 2002). PRSS21 knockout mice were fertile in long-term mating trials over several weeks, but *in vitro* analysis showed that the PRSS21-null sperm had decreased motility, abnormal morphology, and poor performance in fertilization assays (Netzel-Arnett et al. 2009).

Double knockout ACR/PRSS21 male mice were able to produce offspring *in vivo* but with significantly reduced litter size (Kawano et al. 2010). *In vitro*, ACR/PRSS21-null sperm failed to fertilize both cumulus-free and cumulus-intact oocytes, which suggests a role for these proteases in both cumulus penetration and ZP transit. Sperm penetration through the OCC was slower for ACR/PRSS21 sperm than wild-type sperm, and OCCs incubated with extracts of these sperm showed slower dispersal times than wild-type sperm extracts. Analysis of *in vitro* fertilization experiments showed numerous effects: the acrosomal contents of ACR/PRSS21 knockout sperm failed to disperse, the number of sperm bound to the ZP and the percentage of oocytes with fused sperm were very low, and the ratio of sperm bound to the ZP, percentage of ZP penetrated by sperm, number of sperm in the perivitelline space, and percent fertilized oocytes were zero or negligible. Scanning electron microscopy showed that oocytes incubated with ACR/PRSS21-null sperm did not have any sperm tracks, the putative paths of sperm through the ZP caused by limited enzymatic digestion. It is notable that the ACR/PRSS21 sperm appear to remain acrosome intact in these images, or perhaps they are acrosome reacted but retain a significant portion of undispersed acrosomal matrix material. The authors hypothesize that this may be due to the lack of PRSS21 activity rather than dispersal of the acrosomal matrix since ACR is known to activate the PRSS21 protease. They propose that ACR-null mice may have compensatory PRSS21 enzymatic activity for penetrating the ZP while PRSS21-null mice may have compensatory acrosomal matrix dispersal activity. ACR/PRSS21 double knockout mice lack both mechanisms and are subfertile due to the failure to fertilize all ovulated oocytes (Kawano et al. 2010). The authors propose that sperm lacking these two proteases are able to fertilize oocytes due to a compensatory mechanism in the uterine fluid; it is also tempting to speculate that at least some sperm lacking these proteases are able to fertilize oocytes due to simple mechanical forces from sperm motility. The authors suggest a role in cumulus dispersal and penetration, which is interesting since sperm lacking known hyaluronidases SPAM1 (Baba et al. 2002) and HYAL5 (Kimura et al. 2009) were able to fertilize oocytes. Together with double knockout experiments of SPAM1 and ACR or

SPAM1 and PRSS21 (Zhou et al. 2012), these results suggest that ACR and PRSS21 may work in concert with hyaluronidases for cumulus penetration. These results are also consistent with the notion that acrosomal exocytosis occurs before or during interaction with the cumulus (Jin et al. 2011).

A likely substrate of acrosin in the guinea pig acrosomal matrix is AM50, the orthologue of NPTX2 (neuronal pentraxin 2 or Narp) in mice. AM50, like other pentraxins, forms large disulfide-bonded pentameric or decameric complexes and is a major structural component of the acrosomal matrix in guinea pigs (Noland et al. 1994; Reid and Blobel 1994). When acrosin is activated during acrosomal exocytosis in guinea pig sperm or in isolated acrosomal matrices, AM50 is processed at acrosin-specific sites coincident with acrosomal matrix disassembly (Westbrook-Case et al. 1994; Kim et al. 2011). NPTX2 is present in the mouse and knockout of this protein yields fertile males (Bjartmar et al. 2006), although analysis of sperm performance has not been done. Curiously, pentraxin 3 (PTX3) is produced by mouse cumulus cells during cumulus expansion and localizes in the matrix; in vivo, *Ptx3*^{-/-} female mice have been reported to be infertile or subfertile due to severe abnormalities of the cumulus oophorus, although oocytes can be successfully fertilized in vitro (Varani et al. 2002; Salustri et al. 2004).

Protein C Inhibitor (PCI), a widely expressed serine protease inhibitor in seminal plasma, is present in the acrosome, has been shown to inhibit human sperm acrosin activity, and is detected on the surface of damaged or abnormal human sperm where it is assumed to prevent inappropriate acrosin activity (Moore et al. 1993; Hermans et al. 1994; Zheng et al. 1994; Uhrin et al. 2007). PCI has also been linked to fertility in humans as infertile patients had inactive PCI (He et al. 1999). PCI-null female mice are fertile but males are infertile and the primary defect appears to be with the Sertoli cell–spermatogenic cell junctions in the testis since spermatogenic cells were prematurely released into the seminiferous tubule lumen and immature and abnormal germ cells were found in the epididymis (Uhrin et al. 2000; Uhrin et al. 2007). Further analysis of PCI-null epididymal sperm showed 95 % abnormal sperm morphology (many with separated heads and tails and abnormally shaped heads), significantly reduced motility, and failure to bind either cumulus-intact or cumulus-free eggs (Uhrin et al. 2000; Uhrin et al. 2007). Whether the observed lack of cumulus and ZP binding and penetration in absence of PCI is due to problems of unregulated acrosin activity is a question to be investigated, but it is interesting to note that ACR/PRSS21-null mice also had this phenotype. Perhaps unregulated acrosin produces premature spontaneous acrosomal exocytosis and proteolysis of the acrosomal matrix such that sperm–ZP binding sites of the acrosomal matrix and/or ZP are prematurely processed and thus nonfunctional.

The pro-protein convertase PCSK4 is reported to be localized in the developing acrosome during spermatogenesis and, in mature sperm, it is present on the plasma membrane over the acrosome (Gyamera-Acheampong et al. 2006); whether there are two PCSK4 pools (acrosomal and cell surface) or it migrates from the acrosome to the sperm surface is unclear. PCSK4-null mice are infertile (Mbikay et al. 1997; Gyamera-Acheampong et al. 2006) and have abnormal acrosomal shapes (Tardif et al. 2012) that are likely caused by reduced ACRBP processing during

spermiogenesis that leads to improper packaging of proacrosin and perhaps other acrosomal matrix proteins into the acrosome (Kanemori et al. 2013). PCSK4-null mature sperm had accelerated tyrosine phosphorylation during capacitation, improper processing of the egg-binding protein ADAM2, increased spontaneous acrosomal exocytosis, and reduced ZP binding. Thus, PCSK4 has multiple functions in developing and mature sperm and identification of substrate proteins will help elucidate the mechanisms through which this protein functions.

2.6 Inner Acrosomal Membrane Proteins

As acrosomal exocytosis proceeds and acrosomal matrix proteins are released, some components of the IAM remain associated with the sperm and are in position to interact with egg vestments. The term “secondary binding” has been used to describe the interactions of these proteins with the ZP, particularly ZP2, but we envision these interactions to be less well defined and to involve multiple binding and release events. In this model, sperm would adhere to the ZP via multiple proteins that are then released as a result of limited proteolysis to produce repeated binding and release at the molecular level; the continual force of the tail would drive the head progressively further into the ZP. Numerous proteins have the ZP-binding and/or enzymatic properties that are consistent with this model.

The cellular differentiation marker CD46 is ubiquitously expressed on the surface of most mammalian cells and is involved in complement protection in many species, including humans, but not in rats, mice, and guinea pigs. In mice, it appears to be expressed solely in the testis, and, in sperm, CD46 is not on the PM as would be expected, but is located inside the acrosome on the IAM. It was thought that this might be to compensate for loss of sperm surface CD proteins that protect the cell from attack by complement, but sperm from CD46 knockout mice are not more susceptible to complement than control sperm, so the function of CD46 is not likely to be complement protection. Interestingly, CD46 knockout mouse sperm have *enhanced* fertility; the sperm show accelerated spontaneous acrosomal exocytosis and have an elevated litter size in one mouse strain (DBA/2N), and CD46-null sperm fertilized superovulated oocytes at a significantly higher rate than controls (Inoue et al. 2003). Thus, CD46 appears to be important in stabilizing the acrosomal membranes. How it does this from the inner acrosomal membrane is a mystery, but it is possible that, during acrosome formation in spermatogenesis, CD46 may interact with other proteins destined for the periacrosomal membranes. Consequently, CD46 might direct or influence sorting of proteins that stabilize these membranes so that, in the absence of CD46, these proteins are misdirected or do not function properly to stabilize the acrosomal membranes.

Several IAM proteins are known to have a role in sperm–ZP binding but do not have knockout models yet. Yu et al. (2006) also have extracted proteasomal activity from the IAM, but, again, there is no gene knockout data available with regard to its role in fertilization. SPACA4 (SAMP14) is present in the IAM, OAM, and ES of

human sperm; it is GPI-linked and is exposed after acrosomal exocytosis (Shetty et al. 2003). Antibodies to recombinant SPACA4 inhibit binding/fusion of human sperm to zona-free hamster eggs, but a knockout mouse line has not been generated.

2.7 Equatorial Segment Proteins

Equatorin (EQTN) localization is limited to the equatorial segment and the protein is resistant to fairly strong extraction conditions (high salt, 0.1 % Triton X-100, sonication), persists following acrosomal exocytosis on sperm in the perivitelline space, and is even present on sperm inside the egg (Manandhar and Toshimori 2001). Following acrosomal exocytosis, some EQTN is redistributed onto the PM over the ES (and some on the IAM), but the significance of this is not clear. Male EQTN-null mice are subfertile (~50 % of controls) and have reduced litter sizes (5.8 vs. 12.3 in control). EQTN-null sperm undergo normal acrosomal biogenesis (acrosomes are normal by immunofluorescence and by transmission electron microscopy), but do not undergo acrosomal exocytosis above control levels. Co-immunoprecipitation with SNAP25 and syntaxin 1a suggests a role in vesicle fusion during acrosomal exocytosis, but since the protein does not have a canonical transmembrane domain, it must do this via a membrane protein that interacts with the vesicle fusion proteins (Wolkowicz et al. 2003; Hao et al. 2014). Also, since the outer acrosomal and plasma membranes don't fuse in the ES, EQTN is not likely to be involved in fusion in that region. The reason why EQTN affects fusion of the OAM and PM in the more anterior zones is an area for more investigation.

Sperm equatorial segment protein 1 (SPESP1) is another ES-limited protein that is important for fertility in both expected and surprising ways. SPESP1-null and heterozygous males had delays in fertility and smaller litter sizes than wild-type males, and heterozygous and null sperm numbers in the oviduct were significantly reduced (Fujihara et al. 2010). Interestingly, *in vitro* assays showed that the number of sperm, percent progressively motile sperm, and motility patterns were the same for SPESP1-null, heterozygous, and wild-type mice, but the percent of fertilized oocytes and the percent fusibility with zona-free oocytes were significantly less in SPESP1-null and heterozygotes than wild type. Furthermore, in SPESP1-null sperm following acrosomal exocytosis, the plasma and outer acrosomal membranes were found to undergo fusion throughout the entire equatorial segment; since these membranes are typically nonfusogenic, these results show that SPESP1 is important in maintaining the stability of the ES which has an impact on fusibility with the egg plasma membrane (Fujihara et al. 2010).

2.8 Ten Questions

In light of recent developments in the field of fertilization using contemporary imaging techniques and molecular biology to create targeted mutations in genes considered to be important for fertilization, several issues remain that need to be addressed. These include, but are not limited to, the following ten questions: (1) Does acrosomal exocytosis need to occur in a specific region of the female reproductive tract or peri-oocyte environment for fertilization to take place? Current evidence suggests that most sperm have initiated acrosomal exocytosis prior to penetrating the zona pellucida of cumulus-enclosed oocytes. (2) Is binding of the zona pellucida by acrosomal matrix components exposed on the sperm surface required for fertilization or is the affinity of these proteins assistive but not essential for fertilization to take place? For example, ZP3R binds to the oocyte zona pellucida but is not required for fertilization. On the other hand, in the absence of ZAN, the species-specific filter that blocks heterospecific sperm from other phylogenetic classes from binding to mouse oocytes is lost; it would be interesting to determine if ZAN operates to filter the binding of sperm related at the genus or strain level. (3) Can we develop better tools to assess the integrity of the membranes overlying the acrosomal contents during various stages of capacitation and acrosomal exocytosis? (4) Do the components of the acrosome influence the interactions of sperm with the oviductal epithelium, either independently on a single-cell level or perhaps more generally through influencing the reciprocal connectivity between neighboring sperm and the female reproductive tract? (5) Do “acrosome-reacted” sperm that are experimentally recovered from the perivitelline space and are used successfully to fertilize other eggs still contain functional acrosomal matrix material? These experiments raise questions concerning the functions of the acrosomal contents in fertilization (they may be assistive but not required). (6) Are the different terms used to describe the acrosomal matrix important or are they describing the same properties from varied perspectives? Some reports refer to the acrosomal matrix as a particulate compartment of the acrosomal contents (Buffone et al. 2008). Others make reference to a crystalloid nature observed by transmission electron microscopy (Fléchon 2015). Still others classify the acrosomal matrix as an amyloid (Guyonnet et al. 2014). (7) What role does pH play in the intra-acrosomal dynamics and post-exocytotic dispersion of the contents? For example, the pH of the acrosome is acidic and gradually begins to alkalinize during the course of epididymal maturation; once exocytosis is initiated, the contents are rapidly exposed to the more alkaline pH of the milieu surrounding the sperm. (8) Does the change in pH occurring during these periods activate or inactivate acrosomal enzymes? This is true for some proteases such as the proacrosin zymogen activation, but there may be other enzymes exhibiting similar properties. (9) What are the substrates for acrosomal hydrolases? Initially, it was thought that acrosin acted upon the zona pellucida. However, it is clear that this serine protease is involved in the dispersion of the acrosomal matrix, but the actual substrates have yet to be elucidated. (10) Why are acrosomal proteins differentially compartmentalized?

The component segregation could be important for differential binding and/or release during exocytosis. Alternatively, the different domains might simply influence the shape of the acrosome. Clearly, there is still much to learn concerning the involvement of acrosomal matrix components in directing the structure of the acrosome during spermatogenesis and epididymal maturation as well as the function of the acrosome in the fertilization process.

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