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# Mammalian spermatozoa and cumulus cells bind to a 3D model generated by recombinant zona pellucida protein-coated beads

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The egg is a spherical cell encapsulated by the zona pellucida (ZP) which forms a filamentous matrix composed of several glycoproteins that mediate gamete recognition at fertilization. Studies on molecular mechanisms of sperm-egg binding are limited in many mammalian species by the scarcity of eggs, by ethical concerns in harvesting eggs, and by the high cost of producing genetically modified animals. To address these limitations, we have reproduced a three-dimensional (3D) model mimicking the oocyte's shape, by means of magnetic sepharose beads coated with recombinant ZP glycoproteins ( $B_{ZP}$ ) and cumulus cells. Three preparations composed of either ZP2 (C and N-termini;  $B_{ZP2}$ ), ZP3 ( $B_{ZP3}$ ) or ZP4 ( $B_{ZP4}$ ) were obtained and characterized by protein SDS-PAGE, immunoblot and imaging with confocal and electron microscopy. The functionality of the model was validated by adhesion of cumulus cells, the ability of the glycoprotein-beads to support spermatozoa binding and induce acrosome exocytosis. Thus, our findings document that ZP-beads provide a novel 3D tool to investigate the role of specific proteins on egg-sperm interactions becoming a relevant tool as a diagnostic predictor of mammalian sperm function once transferred to the industry.

Sexual reproduction is used by different organisms and involves the fusion of two highly specialized haploid cells, the egg and the spermatozoon. To ensure fertilization, sperm first must cross the cumulus cells surrounding the egg and then, contact, recognize and penetrate the extracellular egg coat (zona pellucida, ZP) to fuse with the plasma membrane. Initial sperm-egg recognition on the surface of the ZP is an essential step for fertilization and biochemical removal of the ZP matrix or its absence in gene-edited mice causes infertility<sup>1-3</sup>. Despite the crucial importance of fertilization, our knowledge about the molecular basis of this interaction is limited in humans and mice<sup>4,5</sup> and basically non-existent in other mammalian species.

A major challenge to obtaining conclusive results on specific molecules involved in sperm-ZP binding is due to limitations on obtaining and manipulating gametes. For example, in some mammalian species it is not feasible to get capacitated sperm and matured oocytes due to ethical concerns (i.e., humans) or restricted number of gametes (i.e., endangered species). There are also technical challenges in assaying and evaluating sperm-egg interactions which is a dynamic process involving multiple macromolecules. In recent years, three-dimensional (3D) systems have become a valuable tool to study and improve our understanding of different stages of reproduction<sup>6,7</sup>. A 3D system recreating the spherical shape oocytes and the biochemical characteristics of the ZP would be a powerful means to investigate unsolved issues related to *in vitro* fertilization<sup>8</sup>.

The mammalian ZP is composed of either three or four glycoproteins designated as ZP1, ZP2, ZP3 and ZP4. In mice, the zona matrix contains ZP1, ZP2 and ZP3. ZP4 is encoded by a pseudogene that does not express the cognate protein<sup>9</sup>. Although the zona matrices in pig<sup>10</sup>, cow<sup>11</sup> and dog<sup>9</sup> oocytes also are made up of 3 ZP proteins,

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these ZP matrices lack ZP1 (rather than ZP4) which is a pseudogene in these species<sup>9</sup>. The ZP matrices of other mammals including rat<sup>12</sup>, hamster<sup>13</sup>, bonnet monkey<sup>14</sup> and human<sup>15</sup> contain all four glycoproteins. Depending on the mammalian species, each ZP glycoprotein has been proposed as a ligand for sperm<sup>16–20</sup>. For example, in mouse and human, sperm bind to the N-terminus of ZP2<sup>20,21</sup> whereas in pigs and cows, ZP3 and/or ZP4 has been implicated in sperm-egg interaction<sup>18,19</sup>. This suggests that the role of individual ZP glycoproteins during fertilization may differ among mammals and needs to be investigated independently rather than extrapolating the findings of one species to another. Such investigations would be facilitated by *in vitro* model systems incorporating order-specific recombinant zona glycoproteins for validation of sperm-zona interactions in different mammals.

The contribution of individual ZP proteins to gamete recognition has been studied biochemically based on blocking potential sperm-ZP interactions with solubilized ZP<sup>22–24</sup>, purified native ZP proteins<sup>25</sup> and recombinant ZP proteins<sup>26–28</sup>. In addition, antibodies directed against specific epitopes have been used to evaluate the candidacy of particular zona proteins in gamete recognition<sup>29</sup>. In recent years, the ease of establishing gene-edited mice has opened the possibility of studying the role of ZP glycoproteins *in vivo* which has provided new insights into mouse and human fertilization<sup>20</sup>. Based on a ZP2-cleavage model of gamete recognition, it has been shown that the N-terminus of ZP2 attached agarose beads can decoy sperm and prevent fertilization *in vitro* and *in vivo*<sup>21</sup>. These same peptide beads can be used to select human sperm that appear to have enhanced ability to bind and penetrate a zona matrix in which human ZP2 replaces endogenous mouse ZP2 and may prove useful in assisted reproductive technologies<sup>30</sup>. Although the use of double and triple transgenic animals can potentially serve as models, the establishment of such animals is time-intensive, expensive and mostly restricted to mice which may not provide the best model for other mammals. Thus, there is considerable interest in developing new strategies to decode sperm-egg recognition<sup>31–33</sup> which will advance our understanding of *in vitro* fertilization and improve assisted human reproduction and livestock production. In this study, a new *in vitro* model is proposed and validated. The model is based on magnetic sepharose beads (B) coated with single recombinant ZP glycoproteins ( $B_{ZP}$ ) that mimic the 3D oocyte's shape. Recombinant porcine ZP2 (C and N-terminus), ZP3 and ZP4 glycoproteins were expressed with peptide tags to allow their identification and conjugation to magnetic sepharose beads. Beads, with individual zona glycoproteins were studied: 1) for their ability to support adhesion of *in vitro* matured porcine cumulus cells; 2) their potential to bind spermatozoa; 3) their ability to induce acrosome exocytosis; and 4) determine if these interactions were affected by the protocol used for sperm capacitation. In summary, this system recreates a 3D environment of ovulated eggs that is scalable and will offer insights into molecular mechanisms of gamete recognition in mammals.

## Results

### Secreted recombinant ZP glycoproteins are stably and uniformly conjugated to beads.

Expression plasmids encoding porcine ZP2C, ZP2N, ZP3 and ZP4 proteins (Fig. 1a, Supplementary Material Fig. S1) were expressed in Chinese Hamster Ovary (CHO) cells and secreted glycoproteins were successfully isolated. Each zona glycoprotein had the expected molecular mass<sup>10</sup>. ZP2C and ZP2N glycoproteins showed a molecular weight of 100 kDa, ZP3 reached 55 kDa, and ZP4 was 65 kDa on immunoblots probed with anti-Flag (ZP2C and ZP2N), anti-ZP3 (ZP3) and anti-V5 (ZP4) antibodies (Fig. 1b, Supplementary Material Fig. S1).

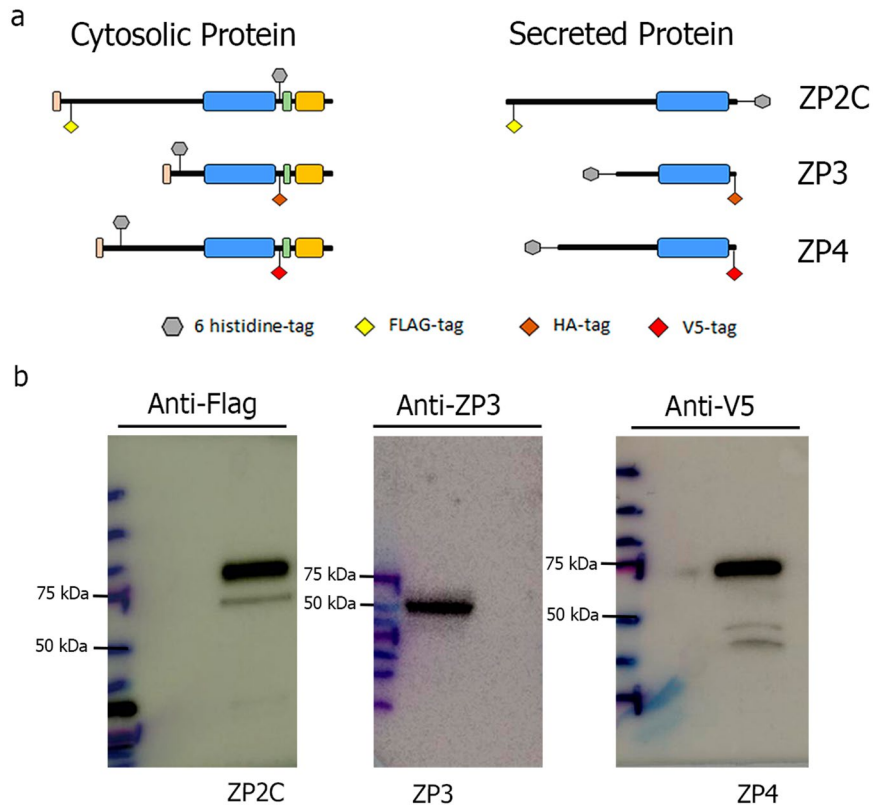
After incubation of beads with medium containing secretions from transfected CHO cells, all recombinant glycoproteins were successfully conjugated to beads (Fig. 2a). Electrophoresis and western blots confirmed their expected molecular weights (100 kDa for  $B_{ZP2}$ , 55 kDa for  $B_{ZP3}$  and 65 kDa for  $B_{ZP4}$ ) both in the secreted medium before bead conjugation (lane 1) and after elution from the beads (lane 2). The proteins were not detected in the medium (lane 3) confirming that all the glycoprotein secreted in the CHO medium was successfully bound to beads. Similar findings were observed for  $B_{ZP2N}$  (Supplementary Material Fig. S1c) and eluted glycoproteins were analysed by mass spectrometry to confirm their identities. All the validated peptides from mass spectrometry had a Scored Peak Intensity (SPI) higher than 70% and a peptide score greater than 5 (Supplementary Material Fig. S2). Protein concentrations in the eluted medium were 0.157, 0.125 and 0.172  $\mu\text{g}/\mu\text{l}$  for ZP2C, ZP3 and ZP4, respectively.

A thick and uniform coating of beads' surface by recombinant ZP glycoproteins was confirmed by immunofluorescent confocal microscopy and no signal was observed in the beads incubated with CHO-cell medium alone ( $B_{\text{CH}}$ ). Moreover, electrophoresis and western blots documented that adherence of recombinant glycoprotein to the sepharose beads was stable after storage for 0, 24, 48, 72 and 144 h (Fig. 2c and Supplementary Material Fig. S1c). Once both ZP2C and ZP2N were characterized subsequent assays used ZP2C because of greater abundance of ZP2C relatively to ZP2N which had an N-terminal histidine tag.

**ZP-coated beads form a 3D oocyte-like shape that supports sperm binding.** Recombinant ZP glycoproteins conjugated to magnetic sepharose beads ( $B_{ZP}$ ) form 3D spheres ( $\sim 120 \mu\text{m}$ ) that approximates the size of native oocytes (Fig. 3a). Scanning electron microscopy provides images of a sphere with a rough surface that approximates the zona matrix of native oocytes (Fig. 3a). The spherical shape of  $B_{ZP}$  is an advantage over flat surfaces because it allows analysis of contact and binding of spermatozoa in three dimensions when live-imaged by stereo-, optic- and scanning-electron microscopy (Fig. 3b,c). In addition, the  $B_{ZP}$  transformed the kinetic energy of spermatozoa into deformation of the proteinaceous surface when they bind (Fig. 3b). Sperm- $B_{ZP}$  binding assays were easily performed and recorded to evaluate sperm activity. The ability of ejaculated porcine sperm to bind to  $B_{ZP2}$ ,  $B_{ZP3}$  and  $B_{ZP4}$  varied among zona proteins over time (Fig. 3c). The number of  $B_{ZP}$  with at least one sperm bound ( $B_{ZP}\text{SB}$ ) increased with time and after 60 min, all preparations had more than 85% of the beads with at least one bound sperm.

### ZP-coated beads recapitulate the size and shape of native porcine cumulus-oocyte complexes (COC).

After 24 h of co-incubation, *in vitro* matured cumulus cells adhered to the surface of  $B_{ZP}$ . This recreated the size

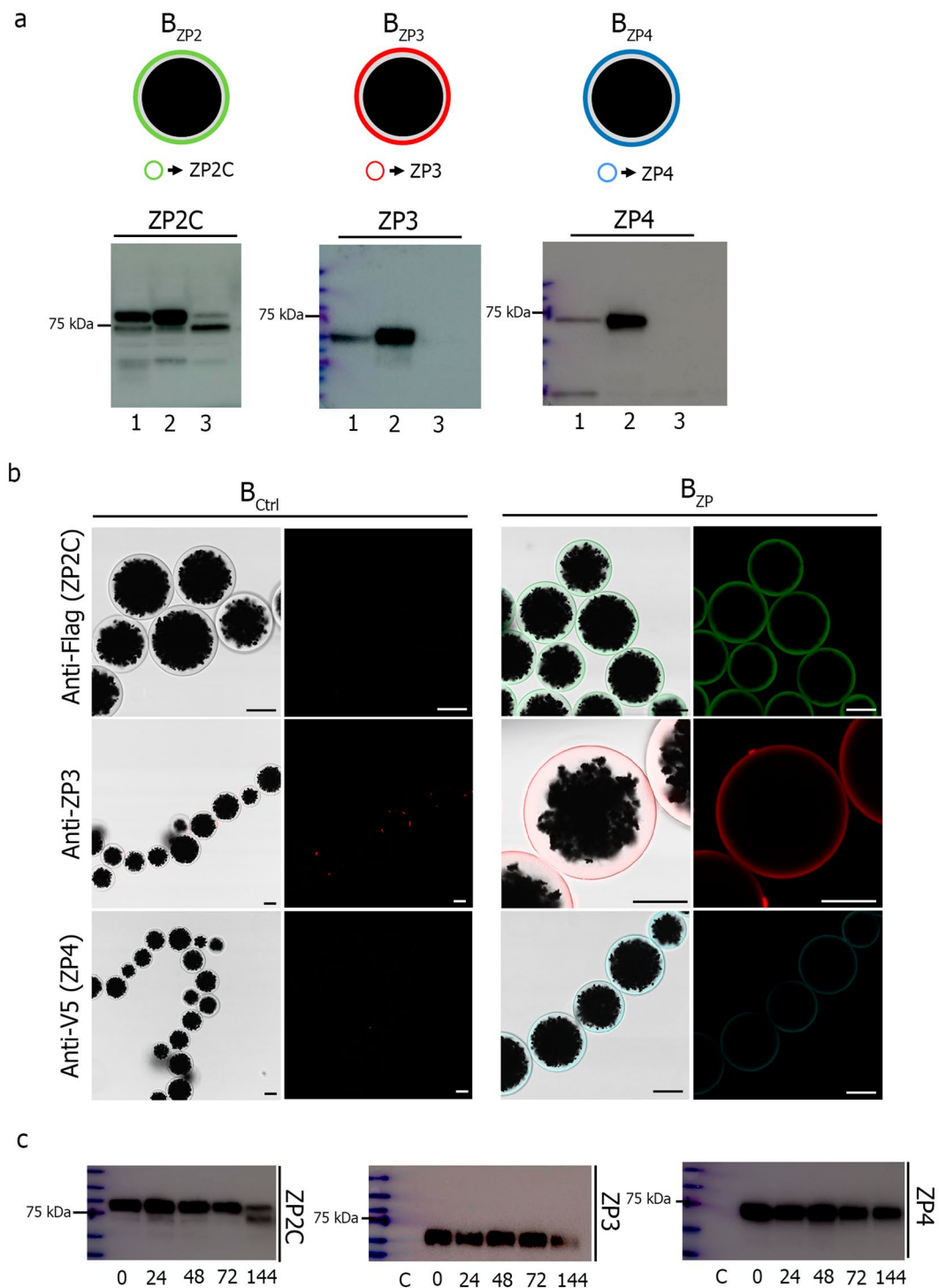


**Figure 1.** Design and expression of porcine recombinant ZP proteins. **(a)** Schematic representation of recombinant porcine ZP glycoproteins, ZP2C, ZP3 and ZP4. Signal peptide (pink), ZP domain (blue), processing region (green) and transmembrane domain (orange). **(b)** Proteins were expressed in CHO cells, separated by SDS-PAGE and analysed by western blot. ZP proteins were probed with anti-Flag antibodies for ZP2C, anti-ZP3 for ZP3 and V5 Epitope Tag antibody for ZP4. Molecular mass markers, left.

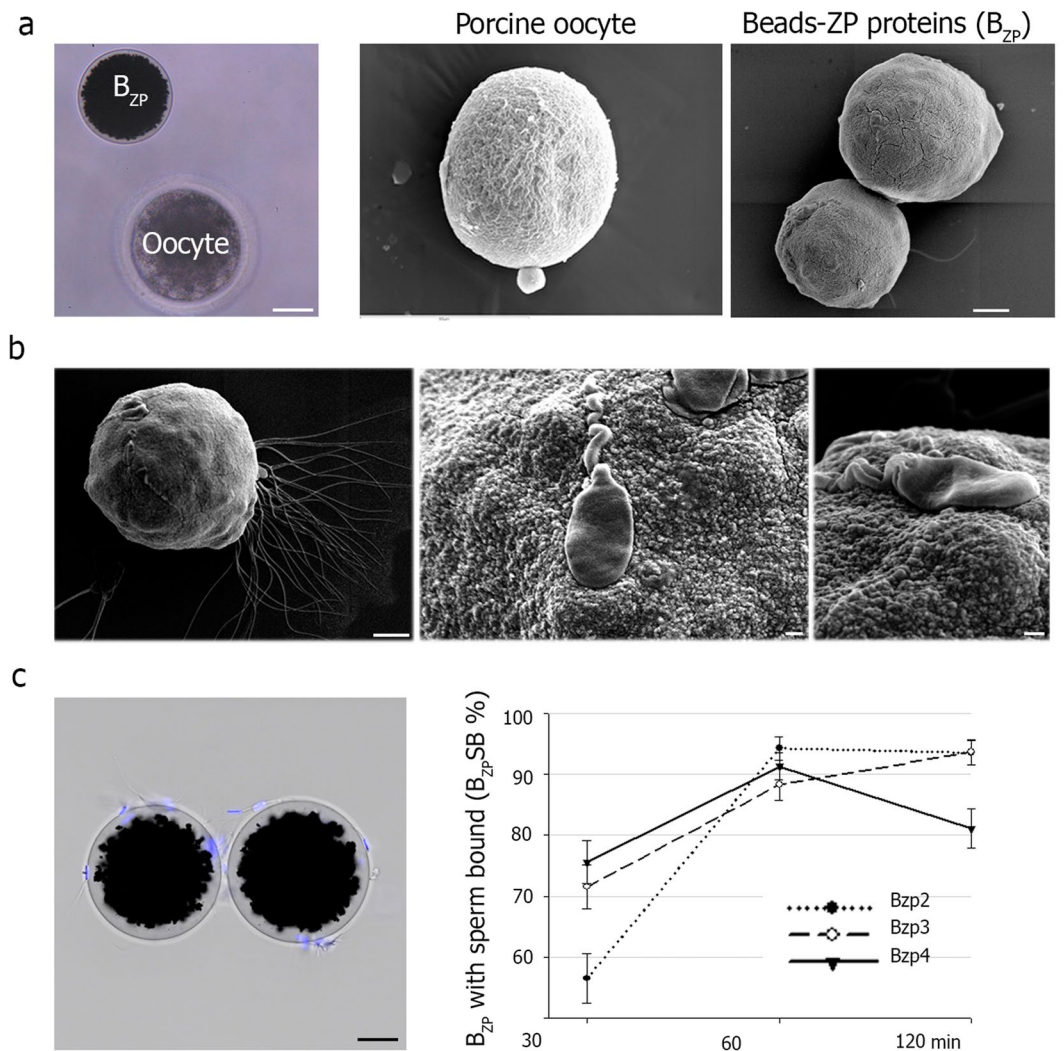
and shape of native porcine cumulus-oocyte complexes (COC) in which the cumulus cells and ZP glycoproteins form two layers external to the oocyte (Fig. 4a). The complex formed between cumulus cells and  $B_{ZP}$  ( $CB_{ZP}$ ) persisted after been mechanically washed (Fig. 4b) and no differences in adherence was observed among the three recombinant zona glycoproteins. In each case, more than 95% of  $B_{ZP}$  had at least one cumulus cell attached after washing (data not shown). Interestingly, the control beads with no ZP proteins on their surface (beads incubated with growth CHO-cell medium;  $B_{Ctrl}$ ) showed a lower number of cumulus cells per bead ( $5.99 \pm 0.45$ ,  $n = 201$ ) than  $B_{ZP3}$  ( $8.92 \pm 0.52$ ,  $n = 212$ ),  $B_{ZP4}$  ( $10.01 \pm 0.49$ ,  $n = 199$ ) ( $P < 0.001$ ) and  $B_{ZP2}$  ( $8.79 \pm 0.54$ ,  $n = 195$ ). No differences were found among the three recombinant zona glycoproteins (Fig. 4b).

**Acrosome integrity of sperm bound to ZP-coated beads.** The acrosomal status of the sperm bound to beads was verified by means of PNA-FITC staining and was confirmed at the ultrastructural level by scanning electron microscopy (Fig. 5a). Acrosome status was recorded for 93% of bound sperm at 30 min, 85% at 60 min and 79% at 120 min (Fig. 5a). Exocytosis was observed on the surface of the three models ( $B_{ZP2}$ ,  $B_{ZP3}$ ,  $B_{ZP4}$ ) (Fig. 5b) and exocytosis increased with time. Forty-five per cent of  $B_{ZP}$  showed acrosomal shrouds on their surface after 30 min incubation, 70% of  $B_{ZP}$  at 60 min and over 80% of  $B_{ZP}$  at 120 min.

**Sperm binding to ZP-coated beads reflect method of capacitation.**  $B_{ZP2C}$  were incubated for 0.5, 1, 2 and 18 h with spermatozoa capacitated by either swim-up or double centrifugation and the number of beads with at least one sperm bound ( $B_{ZP2SB}$ ) and the number of sperm bound per bead ( $S/B_{ZP2}$ ) was determined (Fig. 6). With either method of capacitation, the  $B_{ZP2SB}$  steadily increased for the first 2 h of co-incubation and then decreased after 18 h due to detachment of sperm from beads (Fig. 6 left panel). The temporal profile of  $S/B_{ZP2}$  was similar to  $B_{ZP2SB}$ . No differences in the  $S/B_{ZP2}$  were found between sperm capacitation protocols at time 0.5 h, but a higher  $S/B_{ZP2}$  after capacitation with double centrifugation was observed at 1 h ( $13.21 \pm 1.11$ ,  $n = 245$  vs.  $10.04 \pm 0.68$ ,  $n = 253$ ,  $P = 0.015$ ), 2 h ( $40.68 \pm 1.93$ ,  $n = 278$  vs.  $17.28 \pm 0.98$ ,  $n = 268$ ,  $P < 0.001$ ) and 20 h ( $27.02 \pm 2.14$ ,  $n = 204$  vs.  $10.42 \pm 0.74$ ,  $n = 234$ ,  $P < 0.001$ ) (Fig. 6 right panel).



**Figure 2.** Conjugation of ZP recombinant proteins to sepharose magnetic beads. **(a)** Schematic representation of ZP recombinant proteins coated beads ( $B_{ZP2}$ ,  $B_{ZP3}$  and  $B_{ZP4}$ ) (upper). SDS-PAGE and western blot of ZP2C, ZP3 and ZP4 proteins conjugated to magnetic beads (lower). Medium with secreted proteins before conjugation (lane 1), in the eluted fraction (lane 2), and media after conjugation (lane 3). Anti-Flag antibody was used for ZP2, anti-ZP3 for ZP3 and V5 Epitope Tag antibody for ZP4. **(b)** Confocal microscopy images of beads alone ( $B_{Ctrl}$ ) (left) and conjugated beads ( $B_{ZP}$ ) (right) showing uniform coating of beads with ZP proteins. Scale bar, 40  $\mu$ m. **(c)** SDS-PAGE and western blot of ZP2C, ZP3 and ZP4 proteins conjugated to magnetic beads after conjugation and storage for 0, 24, 48, 72 and 144 h. Lane C, beads incubated with CHO-cell growth medium without recombinant zona proteins.

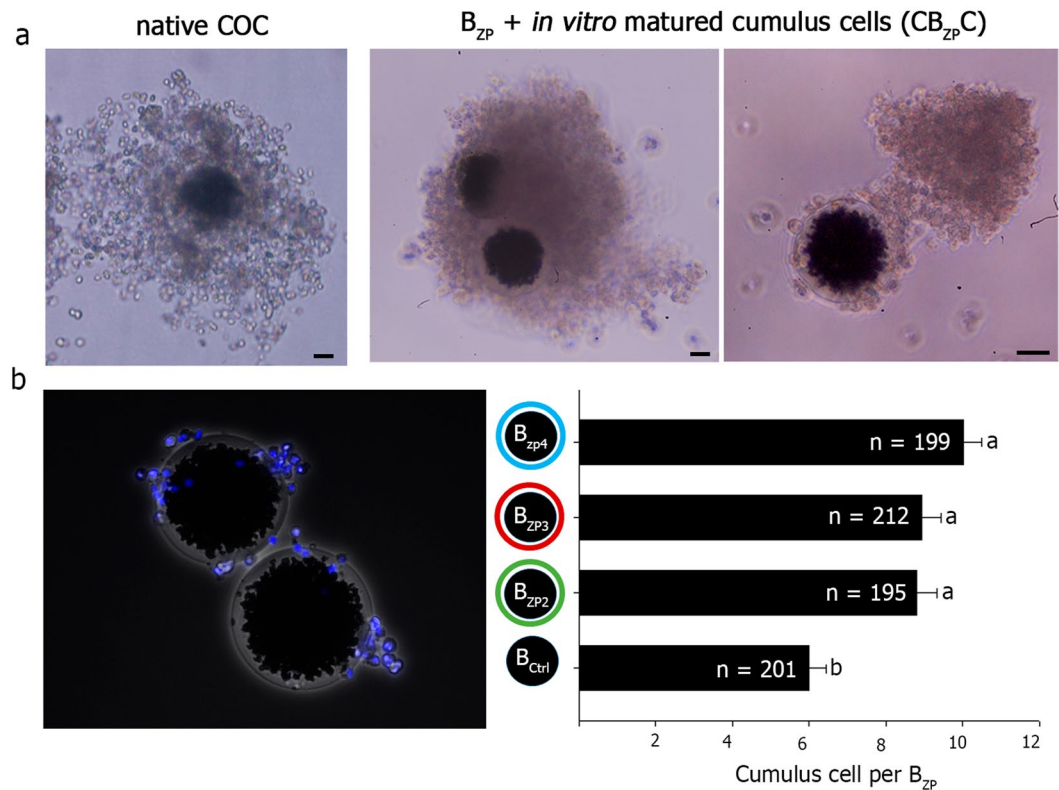


**Figure 3.** Sperm-B<sub>ZP</sub> binding characterization. **(a)** Magnetic bead conjugated to ZP proteins (B<sub>ZP</sub>) and native denuded porcine oocyte observed by stereomicroscopy. Scale bar, 30  $\mu$ m (left). Denuded porcine oocyte and beads conjugated with ZP proteins (B<sub>ZP</sub>) observed under scanning electron microscope. The typical sponge-like structure of the ZP is observed in the oocyte (middle) and is like the variegated surface of the recombinant zona protein-sepharose beads (right). Scale bar, 10  $\mu$ m. **(b)** Scanning electron microscopy of ZP-beads (B<sub>ZP</sub>) after co-incubation with porcine spermatozoa. Scale bar, 10  $\mu$ m (left). Higher magnification of sperm binding that documents small deformation mounds after interaction with the surface of a B<sub>ZP</sub> (middle, right). Scale bar, 1  $\mu$ m. **(c)** ZP coated beads (B<sub>ZP</sub>) with attached porcine spermatozoa after co-incubation with 200,000 sperm cells/mL and observed under stereomicroscope (sperm DNA appear in blue after being stained with bisbenzimidazole; scale bar, 30  $\mu$ m) (left). After co-incubation for 30, 60 and 120 min, B<sub>ZP</sub> were fixed (0.5% glutaraldehyde), stained (0.01 mM bisbenzimidazole) and evaluated under epifluorescence microscope (right). The percentage of B<sub>ZP</sub> with at least one sperm bound (B<sub>ZP</sub>SB) at each time was recorded.

## Discussion

The molecular basis of gamete recognition among mammals remains unclear. In gene-edited mice, Izumo<sup>34</sup>, Juno<sup>35</sup>, CD9<sup>36</sup> and ZP2<sup>20</sup> are required for sperm-egg recognition. However, these proteins have not been tested in other mammals, or the results have been controversial because of taxon-specific features of the fertilization. A significant impediment to obtaining unequivocal results has been the absence of a consistent assay for gamete recognition. Therefore, we have investigated zona glycoprotein-sepharose beads (B<sub>ZP</sub>) as a 3D model of mammalian gamete interactions that mimics the spherical shape of oocytes, easily translates among mammals and is scalable. Magnetic sepharose beads are conjugated to recombinant ZP glycoproteins to which cumulus cells and sperm bind and allow for assessment of acrosome status under varying experimental conditions. This technology represents a necessary first step in validating the role of individual egg proteins in sperm-egg recognition. As an example, we now report testing three different zona glycoproteins potentially involved in porcine gamete interactions.

The zona proteins are processed to remove a signal peptide at their N-terminus and are released from the C-terminus transmembrane domain so that an ectodomain can participate in the extracellular zona pellucida that surrounds ovulated eggs<sup>37–39</sup>. Therefore, care needs to be taken in the placement of epitope tags to ensure that they

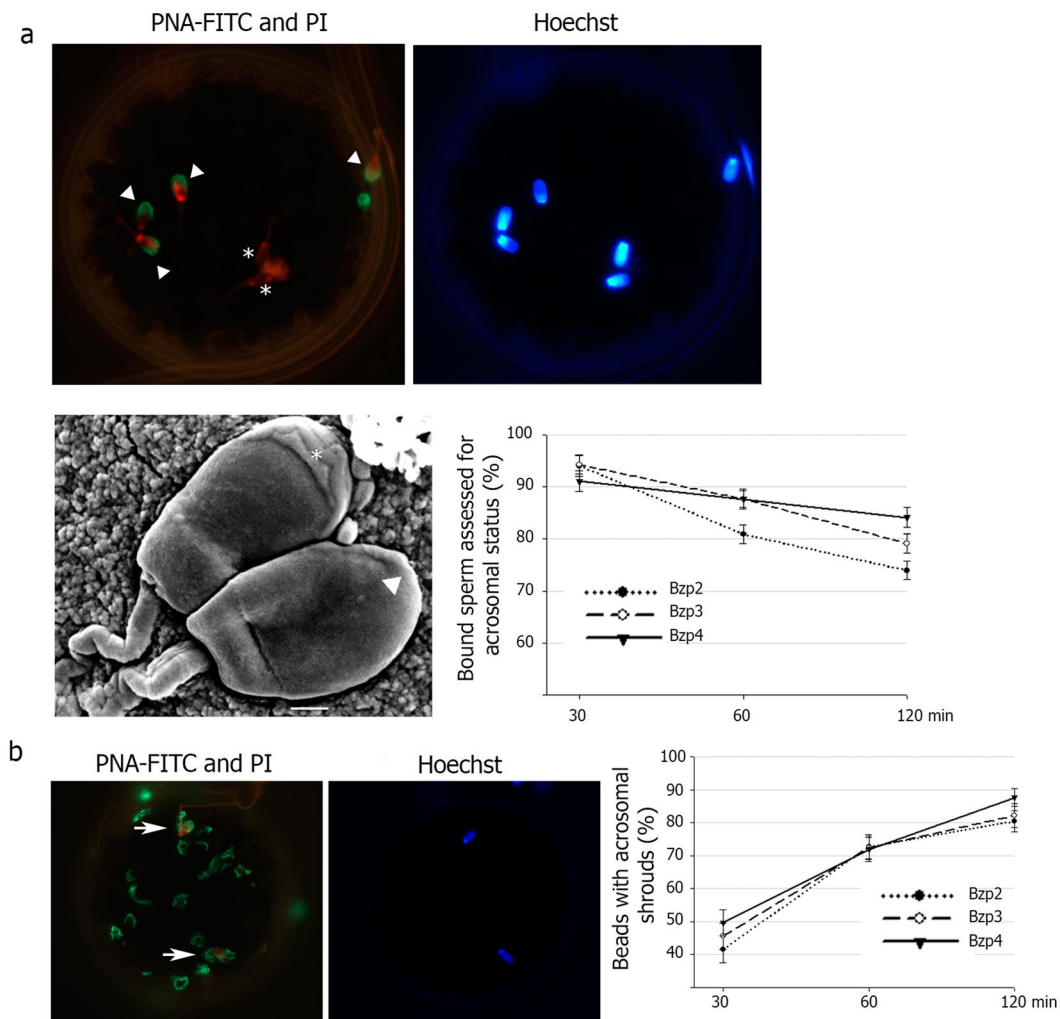


**Figure 4.** Mimicking the cumulus-oocyte complex (CB<sub>ZP</sub>C). (a) Expanded, native cumulus-oocyte complex (COC) after *in vitro* maturation. Scale bar, 30  $\mu$ m (left). ZP-beads after 24 h of incubation with isolated cumulus cells obtained from *in vitro* matured COCs, cumulus-ZP coated beads complex (CB<sub>ZP</sub>C). Scale bar, 30  $\mu$ m. The cells adhere to the bead surface resembling a native COC with two external coats, cumulus cells and ZP proteins (middle, right). (b) Quantification of the number of cells adhered to the ZP coated-bead (B<sub>ZP</sub>) stained with 0.01 mM bisbenzimidazole and mechanically washed (mean  $\pm$  SEM). No difference in the number of adherent cells was observed among the three zona proteins attached to sepharose beads (B<sub>ZP2</sub>, B<sub>ZP3</sub>, B<sub>ZP4</sub>). However, control beads without ZP proteins had fewer cells. Different letters (a,b) in the same column indicate differences between groups ( $P < 0.001$ ).

are present in recombinant zona glycoproteins. Because of glycosylation and multiple disulfide bonds, it is preferred that heterologous mammalian cells lines be used for production to ensure secretion of biologically active recombinant zona glycoproteins even the glycosylation patterns differ from endogenous ZP proteins. Designed recombinant glycoproteins of the ZP were successfully secreted, detected by commercial antibodies and conjugated to magnetic sepharose beads to provide a homogeneous coating over the entire surface. The conjugation of recombinant zona glycoprotein to the beads was stable, allowing a wide window of time in which to perform desired experiments.

Inert B<sub>ZP</sub> were selected because of their size, variegated surface and ease of handling. The native zona matrix porous matrix and its rough surface may be important for physiological responses when sperm bind to its surface<sup>40</sup>. Although individual recombinant ZP-proteins do not form native matrices, conjugated with sepharose beads provide a rough surface<sup>41</sup> with sufficient stiffness that responds to mechanical pressure of sperm-ZP contact. According to current models, the kinetic energy of the sperm is converted into elastic deformation of the zona matrix necessary for successful penetration<sup>42</sup>. The sepharose-glycoprotein texture allows this deformation which mimics, to some extent, the mechanical properties of the ZP. Furthermore, two sperm-oocyte binding patterns have been described when observed by SEM and Transmission Electron Microscopy (TEM) techniques. The most commonly pattern, a flat and tangential attachment of the sperm head to the surface of the ZP, which is then followed by intrusion into the zona in precisely this horizontal position<sup>43</sup> and a vertical sperm binding with ZP penetration by the tip of the head. This last pattern is found in oocytes where large, cluster-like numbers of bound spermatozoa are visible<sup>44</sup>. Both binding patterns are observed in the B<sub>ZP</sub> models. Sperm bound to the bead surface showing apical binding is observed in Fig. 3b (left image) and a sperm flattened on bead surface is observed in Fig. 3b (centre and right images). Besides, some artefact during samples processing and also that sperm could be artificially forced to have additional interactions with the bead material when compared to the physiological configuration of the zona pellucida, are not discarded, as a slightly “rolled” way in the mid-pieces, head deformations and aphysiological mid-piece bead interaction that are observed in SEM pictures.

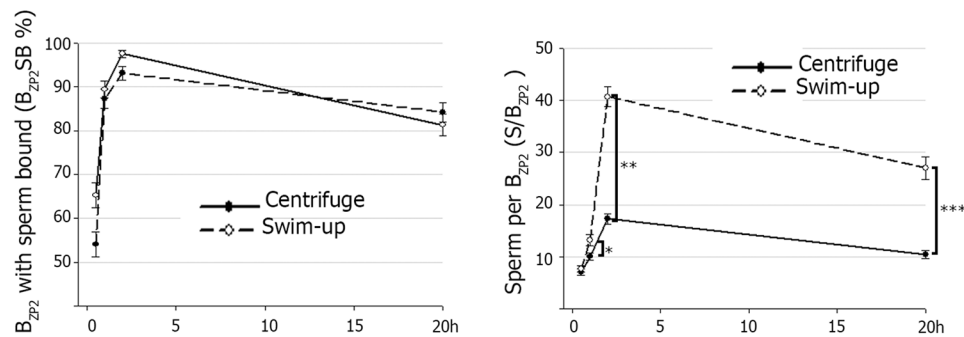
In most mammals, ovulated eggs are surrounded by a substantial layer of cumulus cells embedded in a hyaluronan matrix. The porcine cumulus cells-oocyte complex (COC) typically contains ~3000 cumulus cells<sup>45</sup> and forms a barrier that sperm must penetrate as they approach the ZP. The spherical B<sub>ZP</sub> and the characteristic of the



**Figure 5.** Acrosome status of sperm bound to the ZP-beads ( $B_{ZP}$ ). (a)  $B_{ZP}$  co-incubated with sperms were fixed (0.5% glutaraldehyde) and stained with 4  $\mu\text{g}/\text{mL}$  PNA-FITC (fluorescein isothiocyanate-conjugated peanut agglutinin), 20  $\mu\text{g}/\text{mL}$  PI (propidium iodide) and 0.01 mM bisbenzimidazole. Acrosome reacted ( $\Delta$ ) and non-reacted ( $*$ ) spermatozoa were observed on the bead surface. Images taken from epifluorescence microscopy (upper) and scanning electron microscopy (lower left). Scale bar, 1  $\mu\text{m}$ . Percentage of sperm bound to  $B_{ZP}$  assessed for acrosome status over time of co-incubation (lower right). (b) Acrosomal shrouds on the surface of  $B_{ZP}$ . Beads were fixed and stained with as in (a). Acrosomal shrouds (green) and spermatozoa heads (arrow) were observed by epifluorescence microscopy (left, middle). The number of  $B_{ZP}$  with acrosomal shrouds on the surface increased in a time-dependent manner ( $P < 0.05$ ) (right).

binding surface of recombinant zona glycoproteins result in *de novo* cumulus cells adhesion. This recreates the cumulus matrix appearance of native COCs in which cumulus cells adopt teardrop-like shapes and projections penetrate the zona matrix<sup>46</sup>. The presence of recombinant zona glycoproteins increased the number of cumulus cells tightly attached to the surface of the  $B_{ZP}$ . To the best of our knowledge, this is the first 3D model of gamete recognition that includes cumulus cells and provides a robust system to study the physiological activity of the fertilizing sperm.

To evaluate the role of individual zona proteins in gamete recognition, investigators have sought to assay sperm-binding activity and the ability to induce acrosome exocytosis. Two assays for sperm-ZP binding tests are the hemizona assay<sup>47</sup> and a competitive intact zona sperm binding test<sup>48</sup>, but reported results have been contradictory<sup>49</sup>. Individual recombinant zona proteins seemingly provide simpler, more reproducible and physiologically relevant sperm-ZP binding and acrosome reaction assays<sup>50</sup>. Taking advantage of recombinant zona glycoproteins coupled to sepharose beads, we have validated a sperm- $B_{ZP}$  binding assay using porcine gametes. Each  $B_{ZP}$  was treated as an individual oocyte using the same sperm:bead ratio commonly used in porcine IVF studies<sup>8</sup> and had time-dependent binding as is observed with native gametes<sup>51,52</sup>. The acrosome status of bound spermatozoa was readily determined by staining with PNA-FITC (acrosome reacted) and confirmed by the time-dependent presence of acrosomal shrouds on surface of the  $B_{ZP}$ <sup>52,53</sup>. While conducted using porcine gametes, the same design is easily translated to other species.



**Figure 6.** Sperm binding to ZP2C-coated beads ( $B_{ZP2}$ ) after capacitation by two different methods.  $B_{ZP2C}$  were incubated for 0.5, 1, 2 and 20 h with 200,000 porcine spermatozoa/mL capacitated by swim-up or double centrifugation method. After co-incubation,  $B_{ZP2C}$  were fixed (0.5% glutaraldehyde), stained (0.01 mM bisbenzimidazole) and evaluated by epifluorescence microscope to calculate the percent of  $B_{ZP2}$  with at least one sperm bound ( $B_{ZP2}$ -SB) (left) and to count the number of sperm bound per bead ( $S/B_{ZP2}$ ) (right). Higher number of sperm bound per  $B_{ZP2}$  after capacitation with double centrifugation was observed at 1 h (\* $P = 0.015$ ), 2 h (\*\* $P < 0.001$ ) and 20 h (\*\*\*) ( $P < 0.001$ ).

The method of sperm preparation affects gamete viability, motility, capacitation, ZP binding ability and acrosome exocytosis in both livestock<sup>54–56</sup> and humans<sup>57,58</sup>. Swim-up selection is a more physiological approach than washing by centrifugation and sperm cells selected by this method have lower progressive motility, slower capacitation speed, lower ROS content and DNA fragmentation levels than those sperm selected by centrifugation<sup>56,59,60</sup>. Using  $B_{ZP}$ , centrifuged sperm bound to the beads at a higher rate than swim-up sperm as is observed with native gametes<sup>56</sup>.

From a structural point of view, the  $B_{ZP}$  closely mimic native cumulus-oocyte complexes by maintaining a spherical shape, supporting cumulus cell adhesion and providing a glycoprotein-specific, variegated surface analogous to the native ZP. From a biological point of view, the  $B_{ZP}$  support sperm binding and recapitulates induction of acrosome exocytosis replete with the appearance of acrosomal shrouds and dependence of sperm binding on different methods of capacitation. From these observations, we conclude that the recombinant zona glycoprotein-coated beads will provide a valuable tool to explore the molecular basis of gamete recognition in a wide range of mammals. An additional movie file shows a graphical animation of the methodology developed (see Supplementary Movie).

## Methods

Unless otherwise stated, all chemicals were purchased from Sigma-Aldrich Química (Madrid, Spain). This study was carried out in strict accordance with the recommendations in the Guiding Principles for the Care and Use of Animals (DHEW Publication, NIH, 80–23). The protocol was approved by the Ethical Committee for Experimentation with Animals of the University of Murcia, Spain (Project Code: AGL2015-70159-P and AGL2015-66341-R).

**Recombinant zona pellucida (ZP) glycoproteins design and expression.** Expression plasmids (pcDNA3.1[+]) were designed and constructed to encode porcine ZP glycoproteins (ZP2: UniProt P42099; ZP3: UniProt P42098; and ZP4: UniProt Q07287) (GeneArt, Life Technology, Thermo Fisher Scientific). A histidine tag (6x) was added to the N-terminus of ZP2N (38–43 aa), ZP3 (41–46 aa) and ZP4 (22–27 aa), and to the C-terminus of ZP2C (644–649 aa). In addition, a FLAG-tag was added to ZP2C (38–45 aa) and ZP2N (642–649 aa); an HA tag was added to ZP3 (313–321 aa); and a V5 tag to ZP4 (469–482 aa). After verification by DNA sequence, ZP2C, ZP2N, ZP3 and ZP4 expression plasmids were amplified using Library Efficiency DH5 $\alpha$  Competent cells (Thermo Fisher Scientific) and purified with GenEluted Plasmid Kit.

Chinese Hamster Ovary cells (CHO cells, ECACC, The European Collection of Authenticated Cell Cultures) were grown (37 °C, 5% CO<sub>2</sub> and 95% humidity) for 48–72 h to 80–90% confluence using F-12 medium (Biowest, Nuaille, France) supplemented with 10% fetal bovine serum (Biowest, Nuaille, France) and 100 U/mL penicillin-streptomycin (GibcoBRL-Life Technologies, Gaithersburg, USA). Transient transfections were performed with X-tremeGene HP (Roche Applied Science, Indianapolis, USA) in accordance with the manufacturer's protocol. For each transfection, 4  $\mu$ L X-tremeGene HP transfection reagent were added to a final volume of 200  $\mu$ L Opti-MEM reduced-serum medium (Gibco-Invitrogen) pre-dissolved with 2  $\mu$ g template plasmid and incubated for 15 min at room temperature (RT). The complex was diluted by adding 2 ml Opti-MEM and overlaid on growing cells (37 °C, 5% CO<sub>2</sub> and 95% humidity). The medium containing the secreted glycoproteins was collected after 48 h, centrifuged (3,939  $\times$  g, 5 min, 4 °C) to remove cell debris and concentrated in Vivaspin Turbo 4 of 10,000 Da (Sartorius), obtaining a final volume of 200–300  $\mu$ L of concentrated glycoproteins in 20 mM sodium phosphate buffer, pH 7.4.

**Conjugation of recombinant glycoproteins to magnetic beads.** Magnetic sepharose beads (His Mag Sepharose Excel GE Healthcare) were handled with a magnetic rack (MagRack 6, GE Healthcare). The magnetic sepharose beads were homogeneously resuspended and 10  $\mu$ L of bead slurry was pipetted into a microcentrifuge



tube containing 500  $\mu\text{L}$  of 20 mM sodium phosphate buffer, pH 7.4. The beads were washed with 500  $\mu\text{L}$  of washing buffer (20 mM sodium phosphate, 0.5 M NaCl, 10 mM imidazole, pH 7.4) and finally with 500  $\mu\text{L}$  of binding buffer (20 mM sodium phosphate, 0.5 M NaCl, 0.1% Tween-20, pH 7.4). Concentrated recombinant ZP glycoproteins (ZP2C, ZP2N, ZP3 and ZP4) were incubated with washed magnetic beads (1:1 v/v) overnight at 4 °C with orbital agitation. After incubation, the beads coated with glycoproteins ( $B_{ZP2C}$ ,  $B_{ZP2N}$ ,  $B_{ZP3}$  and  $B_{ZP4}$ ) were washed twice with 20 mM sodium phosphate buffer (pH 7.4) to remove non-conjugated proteins, resuspended in 20 mM sodium phosphate buffer and stored at 4 °C until use. As a control, beads were incubated with non-transfected CHO cells media processed as though it contained recombinant protein ( $B_{\text{Ctrl}}$ ). The recombinant zona glycoproteins were eluted by adding 100  $\mu\text{L}$  of elution buffer (20 mM sodium phosphate, 0.5 M NaCl, 500 mM imidazole, pH 7.4) and incubated at 4 °C and 750 g in a thermomixer for 1 h. ZP2C, ZP3 and ZP4 glycoprotein concentrations were estimated by Bradford colorimetry at the Molecular Biology Service (SACE, University of Murcia, Spain). ZP2C, ZP3 and ZP4 were separated by SDS-PAGE and the bands corresponding to each glycoprotein were isolated and processed for proteomic analysis at the Molecular Biology Service (SACE, University of Murcia, Spain). The protein identity was determined by mass spectrometry, carried out by using a HPLC/MS system, in which an Agilent 1100 Series HPLC (Agilent Technologies, Santa Clara, CA, USA) is connected to an Agilent Ion Trap XCT Plus Mass Spectrometer (Agilent Technologies, Santa Clara, CA, USA) using an electrospray (ESI). Data processing was performed by the Data Analysis program for LC/MSD Trap Version 3.3 (Bruker Daltonik, GmbH, Germany) and Spectrum Mill MS Proteomics Workbench (Rev A.03.02.060B, Agilent Technologies, Santa Clara, California, USA).

**Western blot.** Cell growing medium containing concentrated glycoproteins was separated by SDS-PAGE and transferred to PVDF membranes which were probed with antibodies: anti-Flag for ZP2 (1:1000 v/v in TBST 1x, 1% BSA); anti-ZP3 kindly donated by Dr. Hedrick<sup>61</sup>, for ZP3 (1:2000 v/v in TBST 1x, 1% BSA); and V5 Epitope Tag Antibody for ZP4 (1:2000 v/v in TBST 1x, 1% BSA) (Thermo Fisher Scientific) prior to visualization by chemiluminescence (Pierce ECL-Plus, Thermo Fisher Scientific).

$B_{ZP}$  were analyzed by SDS-PAGE to verify that recombinant glycoproteins were successfully bound to the sepharose beads. To study the stability of the conjugation, after storage (4 °C, 20 mM sodium phosphate buffer, pH 7.4) for 0, 24, 48, 72 and 144 h, the  $B_{ZP}$  were washed twice with 20 mM sodium phosphate buffer, pH 7.4, resuspended in 20 mM sodium phosphate buffer, pH 7.4 and solubilized under reducing conditions in 4X SDS sample buffer (Millipore, USA). After 10 min at 100 °C, the supernatant was separated by SDS-PAGE, transferred to PVDF membranes and probed with the aforementioned ZP antibodies.

**Confocal microscopy.** After conjugation, the  $B_{ZP}$  were incubated (1 h, RT) with antibodies diluted in calcium- and magnesium-free PBS supplemented with 1% BSA. Antibody dilutions were: anti-FLAG for ZP2 (1:200 v/v); anti-ZP3 for ZP3 (1:200 v/v); and V5 Epitope Tag Antibody for ZP4 (1:200 v/v). The  $B_{ZP}$  then were incubated (1 h, RT) with secondary antibodies (anti-rabbit IgG-FITC conjugate (1:100 v/v) for ZP2 and ZP4; Alexa Fluor 488 Donkey anti-goat IgG (1:200 v/v) for ZP3), washed in PBS and fixed with 2% paraformaldehyde (Electron Microscopy Sciences, Hatfield, Philadelphia, USA). Stained  $B_{ZP}$  were placed on a chambered slide (25- $\mu\text{L}$  cavity) with Gene Frame (Advanced Biotechnologies, Leatherhead, UK) and covered with a coverslip. Samples were analysed with a Leica TCS SP8 confocal microscope and image analysis was performed using LAS X Core software analysis software (Leica Microsystems, Spain). FITC and Alexa Fluor 488 were excited at 488 nm.

**Incubation of ZP-conjugated beads with isolated *in vitro* matured cumulus cells.** To recreate an *in vitro* 3D model resembling a cumulus-oocyte complex, the models  $B_{ZP2C}$ ,  $B_{ZP3}$  and  $B_{ZP4}$  were incubated with cumulus cells derived from *in vitro* matured porcine cumulus oocyte complexes (COCs)<sup>52</sup>. COCs were isolated from ovaries obtained from 6 to 7-month-old animals slaughtered at an abattoir. After *in vitro* maturation, COCs were decumulated in PBS-1% BSA by pipetting, oocytes were discarded, and cumulus cells were centrifuged twice (1,200  $\times$  g, 10 min) in modified TALP medium<sup>62</sup>.  $B_{ZP}$  (50–55) were incubated (24 h, 38.5 °C, 5% CO<sub>2</sub>, 20% O<sub>2</sub> and saturated humidity) with 2,500 isolated cumulus cells/  $B_{ZP}$ <sup>63</sup> in 500  $\mu\text{L}$  TALP Adhesion of cumulus cells was confirmed microscopically by observing formation of cumulus mass plugs around coated beads ( $B_{ZP2C}$ ).  $B_{ZP2C}$  were then gently pipetted three times in PBS supplemented with 0.1% PVA to remove loosely adherent cumulus cells, fixed with 0.5% (v/v) glutaraldehyde, stained with 0.01 mM Hoechst 33342, mounted on cover slips and observed by epifluorescence microscope at 40 $\times$ . Three double-blind replicates were obtained to determine the number of cumulus cells remaining attached to the bead surface.

**Sperm preparation.** Ejaculated spermatozoa from boars proven to be fertile (Landrace  $\times$  Largewhite, 12–24 months old; CEFU, SA, Pliego, Murcia, Spain) from Artificial Insemination Center (AIC) were diluted (1:5 v/v) in Beltsville Thawing Solution (BTS). To validate sperm binding to the  $B_{ZP}$ , heterospermic samples were prepared either by double centrifugation<sup>64</sup> or swim-up<sup>56</sup>. For the double centrifugation, diluted sperm samples were centrifuged (300  $\times$  g, 3 min) after which the supernatant was collected and centrifuged again (800  $\times$  g, 5 min). The supernatant was discarded, and the sperm pellet was resuspended in modified TALP medium<sup>62</sup> supplemented with 1 mM sodium pyruvate, 0.6% BSA and 50  $\mu\text{g}/\text{mL}$  gentamycin. For the swim-up protocol, 1 ml diluted semen sample was laid below 1 ml NaturARTs PIG sperm swim-up medium (EmbryoCloud, Murcia, Spain) at the bottom of a conical tube. After 20 min at 37 °C (with the tube at a 45° angle), 800  $\mu\text{L}$  from the top of the tube were aspirated. Final sperm concentrations were adjusted to 200,000 sperm cells/mL with supplemented modified TALP media.

**Scanning electron microscopy.**  $B_{ZP}$  co-incubated (2 h) with double centrifuged sperm were fixed (1% glutaraldehyde in PBS (v/v), 30 min), washed three times in molecular biology water and transferred to dry on

a coverslip. Samples were then processed by the General Unit of Microscopy (University of Alicante, Spain). Finally, the beads were observed under an electron microscope JEOL JSM 840 SEM (Jeol Limited, London, United Kingdom). The shape and surface characteristics of the B<sub>ZP</sub> as well as the acrosome integrity of the bound sperm were recorded.

**Sperm-BZP binding assay.** B<sub>ZP</sub> (50–55) conjugated with ZP2C, ZP3, ZP4 were washed twice in TALP medium and co-incubated (38.5 °C, 5% CO<sub>2</sub>, 20% O<sub>2</sub> and saturated humidity) with double centrifuged spermatozoa (200,000 sperm/mL) in 500 µl medium. In addition, B<sub>ZP2C</sub> were inseminated with sperm prepared by swim-up (200,000 sperm/mL) to validate B<sub>ZP2C</sub> as a suitable model for sperm-ZP recognition independent of the method of sperm preparation. To assess the number of sperm bound and the integrity of their acrosomes after 0.5, 1, 2, and 20 h of co-incubation, an aliquot of B<sub>ZP2C</sub> was washed three times in PBS, supplemented with 0.1% PVA, fixed with 0.5% glutaraldehyde in PBS (v/v), and stained with 0.01 mM Hoechst 33342, 4 µg/mL fluorescein isothiocyanate-conjugated peanut agglutinin (PNA-FITC) and 20 µg/mL of propidium iodide (PI). B<sub>ZP</sub> were mounted on slides, and the number of sperms bound and acrosome status (acrosome from reacted sperm stains green), were recorded under an epifluorescence microscopy (Leica, DMLS, Barcelona, Spain). The percent of beads with at least one sperm bound (B<sub>ZP</sub>SB) and the mean number of sperm per bead (S/B<sub>ZP</sub>) were recorded and acrosome integrity was evaluated. Spermatozoa with damaged acrosome and acrosomal content were observed green. Three replicates were performed.

**Statistical analysis.** Data are presented as mean ± SEM and all percentages were modelled according to the binomial model of variables and arcsine transformation to achieve normal distribution. The percent of beads with at least one sperm bound, number of sperm per bead, beads with at least one cumulus cell attached and the mean number of cumulus cells per bead were analysed by one-way ANOVA. When ANOVA revealed a significant effect, values were compared by Tukey's test. A P value < 0.05 was considered statistically significant. The analysis was conducted using Systat v13.1 (Systat Software, Inc San Jose, CA, USA).

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### Author contributions

J.G.H., A.C., R.R. and M.J.M. performed the experiments. B.A., M.C.O. and M.J.M. designed the Z.P. expression plasmids. M.J.G.T. oversaw the electron microscopy studies. R.R. and M.J.M. conceived, planned and supervised the experiments. J.G.H., R.R. and M.J.M. wrote the original draft, reviewed and edited the manuscript. R.R. and M.J.M. obtained funding. All authors participated in final editing providing critical feedback and helped shape the research, analysis and manuscript.

### Competing interests

The authors declare no competing interests.

### Additional information

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