Cortactin depletion results in short tubulobulbar complexes and spermiation failure in rat testes

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Summary
Tubulobulbar complexes are actin-related endocytic structures that form at sites of intercellular attachment in the seminiferous epithelium and are proposed to internalize intact junctions. In this study, we test the prediction that altering the structure/function of tubulobulbar complexes results in failure to release mature spermatids from Sertoli cells. We used an in vivo knockdown strategy to target cortactin, a component of tubulobulbar complexes, in Sprague Dawley rats. In each animal, one testis was surgically injected with cortactin siRNA reagents and the other testis was injected with non-targeting siRNA. After three days, experimental and control testes were processed for immunoblotting, electron microscopy or immunofluorescence microscopy. In testis sections immunostained for cortactin or labeled for filamentous actin, fluorescence microscopy revealed that tubulobulbar complexes were shorter in siRNA-treated testes relative to controls. Significantly, in the knockdown testes, spermiation was delayed in some tubules and had failed in others. When evaluated by electron microscopy, adhesion complexes (ectoplasmic specializations) remained associated with mature spermatids that failed to be released from Sertoli cells. Immunoblots both of whole testis lysates and of isolated seminiferous epithelial lysates confirmed that cortactin expression was knocked-down in experimental testes and in the seminiferous epithelium respectively, relative to controls. Moreover, in testes injected with siRNA reagents with a dye modification on one of the four targeting siRNA sequences, dye clusters were detected at the base of the epithelium confirming that the reagents entered Sertoli cells. Our results are consistent with the hypothesis that tubulobulbar complexes internalize intercellular junctions and that they are a significant component of the sperm release mechanism.

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Key words: Tubulobulbar complexes, Cortactin, Spermatogenesis, Testis, Spermiation, Actin

Introduction
The disassembly of massive intercellular adhesion junctions between Sertoli cells and mature spermatids is fundamental to the process of sperm release (spermiation) from the seminiferous epithelium. During this disassembly, unique structures termed “tubulobulbar complexes” form at the intercellular attachment sites (Russell and Clermont, 1976), and have been proposed to be responsible for internalizing ‘intact’ junctions into Sertoli cells (Guttman et al., 2004b; Young et al., 2009b; Young et al., 2012).

We previously have shown that tubulobulbar complexes contain adhesion proteins (α6β1 integrin, nectin2 [Sertoli cell] and nectin3 [spermatid]) known to be present at ectoplasmic specializations (Guttman et al., 2004b; Young et al., 2009b). Moreover, regions of the complexes enriched in junction proteins also are associated with the endosomal markers Rab5 (Young et al., 2012) and EEA1 (Young et al., 2009b), as well as with LAMP1, SGP1 and PKCζ (Guttman et al., 2004b). Significantly, the absence of tubulobulbar complexes in estradiol treated rats (D’Souza et al., 2009) and in amphiphysin knockout animals (Kusumi et al., 2007) is correlated with a failure to release mature spermatids from the seminiferous epithelium. In addition, spermiation failure is one of the first phenotypes observed in men undergoing hormonal methods of contraception (McLachlan et al., 2002). Together, these observations are consistent with our working hypothesis that tubulobulbar complexes that form between Sertoli cells and spermatids are subcellular machines that internalize intact intercellular adhesion junctions and that the complexes are a key element of the sperm release mechanism.

Although tubulobulbar complexes are unique to the seminiferous epithelium they bear a striking resemblance to podosomes, particularly those in osteoclasts (Ochoa et al., 2000), to membrane tubules reconstituted in cell-free systems (Wu et al., 2010), and to models of clathrin-based endocytosis generally in cells (Taylor et al., 2011). Podosomes in osteoclasts, reconstituted membrane tubules, the necks of mature clathrin coated pits, and tubulobulbar complexes all consist of elongate cores of tubular membrane surrounded by networks of actin filaments and related proteins. However, unlike in the other systems where the tubular central core consists of a single membrane, the core of tubulobulbar complexes consists of a...
‘double-membrane’ tube formed by the attached plasma membranes of two adjacent cells. Among key components present in many of these systems are Arp2/3, N-WASP, amphiphysin, dynamin, and cortactin (Kusumi et al., 2007; Ochoa et al., 2000; Taylor et al., 2011; Vaid et al., 2007; Wu et al., 2010; Young et al., 2009a). Knockdown of cortactin in osteoclasts prevents podosome formation even though actin-related structures in other parts of cells are unaffected (Tehrani et al., 2006).

If tubulobulbar complexes are a significant part of the sperm release mechanism, then perturbation of tubulobulbar complex structure/function should lead to spermiation failure. Because there are no culture models of sperm release, the approach we currently are taking to verify this prediction is to deplete cortactin, a known component of tubulobulbar complexes in rats, by intratesticular injection of siRNA reagents, determine if the knockdown has an effect on tubulobulbar complexes, and then determine whether there is a spermiation defect.

In this study we report the results of cortactin knockdown experiments. Although tubulobulbar complexes were present in siRNA treated rat testes, they were shorter three days after injection than those in contralateral control testes injected with a scrambled sequence. Moreover, spermiation delay and failure were detected in the knockdown testes, and ectoplasmic specializations (tissue specific adhesion junctions) remained associated with those spermatids retained by the epithelium. We conclude that apical tubulobulbar complexes are a significant part of the normal process of sperm release and that interfering with their structure/function results in spermiation failure.

Results
Normal arrangement and structure of tubulobulbar complexes in rat
Tubulobulbar complexes that form in apical regions of the seminiferous epithelium are elongate tubular extensions of the attached plasma membranes of the spermatid and Sertoli cell that project into the Sertoli cell in regions previously occupied by unique adhesion junctions known as ectoplasmic specializations. In rats, as many as twenty-four complexes, each ranging from 2–3 μm in length develop in two parallel rows associated with the concave face of the “hook shaped” spermatid heads (Russell and Clermont, 1976; Russell, 1979c) (Fig. 1). The “double-membrane” core of each complex is cuffed by a dendritic actin network (Young et al., 2009a) (Fig. 2A,B). In addition to actin, other molecular components of tubulobulbar complexes include Arp2/3 (Vaid et al., 2007), N-WASp (Young et al., 2009a), cortactin (Young et al., 2009a), dynamin (Vaid et al., 2007), amphiphysin (Kusumi et al., 2007), and coflin (Guttman et al., 2004a). Formation of each complex is initiated in the Sertoli cell by a clathrin-coated pit (Young et al., 2009a), which retains an attachment to the neighboring germ cell plasma membrane as the complex elongates and matures (Russell and Clermont, 1976) (Fig. 2A). In regions of the coated pit where the germ cell plasma membrane remains attached to the Sertoli cell, fine filamentous connections occur between the two cells, and the germ cell membrane contains a distinct sub-membrane density (Russell and Clermont, 1976) (Fig. 2D,E). As the complex matures a swelling develops in the distal third of the structure just proximal to the coated pit (Fig. 2C). This bulbar region is not associated with actin. Rather, it is closely associated with a cistern of endoplasmic reticulum. The bulbar region eventually buds from the complex and is internalized by the Sertoli cell (Russell and Clermont, 1976; Russell, 1979c).

Cortactin knockdown by intratesticular siRNA
In the first set of experiments on five animals, cortactin bands on Western blots of whole testis lysates from testes injected with cortactin (Cttn) targeted siRNAs were clearly weaker relative to those of similar lysates from contralateral control testes injected with non-targeting siRNAs (Fig. 3A). We concluded that cortactin in the testis was successfully knocked down by intratesticular injection of siRNA.

Tubulobulbar complexes are shorter in cortactin knockdown testes
Tubulobulbar complexes in sections of stage VII seminiferous epithelium appeared qualitatively shorter in cortactin knockdown testes relative to control testes when double labeled for cortactin (Fig. 3B,C) and actin (Fig. 3D).

As an index of tubulobulbar complex length we quantified the numbers of pixels above a set threshold occupied by the total cluster of tubulobulbar complexes adjacent to the concave surfaces of hook-shaped spermatid heads at the apex of the epithelium. Bar = 10 μm. (B) Schematic illustration of tubulobulbar arrangement adjacent to mature spermatids.

Fig. 1. Position and arrangement of tubulobulbar complexes associated with Sertoli/spermatid attachment sites in rat seminiferous epithelium. (A) 1 μm thick section at stage VII of spermatogenesis showing clusters of tubulobulbar complexes adjacent to the concave surfaces of hook-shaped spermatid heads at the apex of the epithelium. Bar = 10 μm. (B) Schematic illustration of tubulobulbar arrangement adjacent to mature spermatids.
shorter in experimental than in control testes. We conclude that cortactin knockdown resulted in shorter tubulobulbar complexes.

Spermatids retained in the epithelium

Mature spermatids are released as spermatozoa into the lumen of seminiferous tubules in Stage VIII of spermatogenesis (Leblond and Clermont, 1952). During this spermiation process adhesion junctions (ectoplasmic specializations) between Sertoli cells and spermatids disappear, Sertoli cell cytoplasm is gradually withdrawn from spermatid heads, spermatids detach from the apex of the epithelium, and residual lobes from the released spermatids are retained by Sertoli cells. In light micrographs of the seminiferous epithelium from control testes at Stage VIII, spermatids were being released in the normal fashion (Fig. 6A,B). In the epithelium at stage VIII of experimental animals retraction of the Sertoli cell cytoplasm away from spermatids appeared delayed in some tubules (arrows in Fig. 6C). Significantly, after most spermatids in controls had been released (Fig. 7A,B) some spermatids in experimental testes appeared not to be released and were retained by the epithelium (arrow marked by asterisk in Fig. 6C, arrowheads in Fig. 7C–F). When evaluated ultrastructurally, ectoplasmic specializations were absent from Sertoli cell regions adjacent to spermatids being released from control epithelia (arrowheads in Fig. 7G,H), but were still present in association with spermatids retained at the apex of experimental testes (Fig. 7I,J). We conclude that there is evidence of a delay and a failure of spermiation in experimental testes, and that adhesion junctions (ectoplasmic specializations) persisted at intercellular attachment sites where spermatids were retained by the epithelium.

siRNA reagents enter Sertoli cells and result in cortactin knockdown in isolated seminiferous epithelia

To verify that the siRNA administered by intratesticular injection actually entered Sertoli cells and resulted in cortactin knockdown in the seminiferous epithelium, we ran additional experiments using the same pool of cortactin targeting and non-targeting duplexes but with one nucleotide from each pool modified with DY547 on the 5’ end of the sense strand. This nucleic acid tag can be visualized by confocal microscopy using an appropriate laser. The dye was detected in the seminiferous epithelium of all animals that received the modified siRNA and non-targeting reagents (Fig. 8B).

Fig. 2. Ultrastructural features of tubulobulbar complexes associated with Sertoli/spermatid attachment sites at the apex of rat seminiferous epithelium. (A) Longitudinal section of a tubulobulbar complex. Bar = 200 nm. (B) Cross section through proximal tubular region. Bar = 100 nm. (C) Cross section through distal tubular region and coated-pit. Notice the density (asterisk) associated with the spermatid plasma membrane and the connections (arrowhead) with the Sertoli cell plasma membrane of the coated pit. Bar = 100 nm. (D) Longitudinal section through distal tubular region and coated-pit illustrating fine filamentous attachments between the plasma membranes of the Sertoli cell and the spermatid (arrowheads), and the density associated with the spermatid plasma membrane (asterisk). Bar = 100 nm.

To confirm that cortactin was indeed knocked-down in experimental testes, Western blots were run on whole testis lysates from animals using the same reagents as those used for the dye morphological studies. Cortactin bands were weaker in experimental lysates than in contralateral control testis lysates (Fig. 8A). To confirm that cortactin knockdown occurred in the seminiferous epithelium, seminiferous epithelium was isolated from two of the above animals and then analysed by immunoblotting. In both animals, successful depletion of cortactin was detected (Fig. 8C). As in our previous experiments, the number of positive pixels above threshold was significantly less adjacent to mature spermatids in experimental animals than in controls when sections were labeled for cortactin (Fig. 8D). We conclude that siRNA reagents entered Sertoli cells and that the reagents targeting cortactin resulted in lower overall levels of the protein in the seminiferous epithelium than in controls. These results confirm the successful delivery of the siRNA into the seminiferous epithelium.

Discussion

There is a growing body of evidence that tubulobulbar complexes that develop adjacent to mature spermatids are subcellular machines that internalize intact intercellular junctions (Guttmann et al., 2004b; Young et al., 2009b; Young et al., 2012). In this study we use intratesticular injection of siRNA reagents to target a known component of tubulobulbar complexes, alter tubulobulbar complex structure and correlate this altered structure with a fundamental biological event during spermatogenesis – sperm release.

Apical tubulobulbar complexes each consist of a tubular process of the spermatid plasma membrane that extends into a corresponding invagination of the neighboring Sertoli cell (Russell and Clermont, 1976). A clathrin-coated pit in the Sertoli cell initiates the formation of the structure and remains attached to the tip of the complex as it lengthens. A network of actin filaments cuffs the double-membrane tubular core and as the complex matures, a large swelling or bulb forms in distal regions (Russell, 1979a). The bulb buds from the complex and enters endosomal compartments of the Sertoli cell (Guttmann et al., 2004b; Young et al., 2012).

The location, timing and unique structure of tubulobulbar complexes have led to a number of suggested functions. Among these are that they eliminate cytoplasm from spermatids (Russell,
anchor elongate spermatids to the epithelium (Russell and Clermont, 1976), facilitate shaping of spermatid heads (Kierszenbaum and Tres, 2004), and remove excess acrosomal contents from spermatids (Tanii et al., 1999). The observation that tubulobulbar complexes also occur at the base of the seminiferous epithelium in association with junction complexes between neighboring Sertoli cells indicates that the primary function of tubulobulbar complexes is probably not related specifically to spermatids per se, but is more related to a cellular process or event common to both apical and to basal sites. At the apex of the epithelium, ectoplasmic specializations located between Sertoli cells and spermatids must ‘disassemble’ to enable sperm release. At the base of the epithelium, similar junctions located between Sertoli cells need to remodel as the next generation of spermatogenic cells translocate through the junctions to complete meiosis and then to develop into sperm cells. A unifying function for tubulobulbar complexes in the seminiferous epithelium is that they internalize intercellular junctions (Guttman et al., 2004b; Russell, 1979b).

There now is substantial evidence that tubulobulbar complexes do internalize adhesion junctions that occur between Sertoli cells and spermatids. First, tubulobulbar complexes develop in areas previously occupied by testis specific adhesion junctions known as ectoplasmic specializations (Tani et al., 1999). The observation that tubulobulbar complexes also occur at the base of the seminiferous epithelium in association with junction complexes between neighboring Sertoli cells indicates that the primary function of tubulobulbar complexes is probably not related specifically to spermatids per se, but is more related to a cellular process or event common both to apical and to basal sites. At the apex of the epithelium, ectoplasmic specializations located between Sertoli cells and spermatids must ‘disassemble’ to enable sperm release. At the base of the epithelium, similar junctions located between Sertoli cells need to remodel as the next generation of spermatogenic cells translocate through the junctions to complete meiosis and then to develop into sperm cells. A unifying function for tubulobulbar complexes in the seminiferous epithelium is that they internalize intercellular junctions (Guttman et al., 2004b; Russell, 1979b).

There now is substantial evidence that tubulobulbar complexes do internalize adhesion junctions that occur between Sertoli cells and spermatids. First, tubulobulbar complexes develop in areas previously occupied by testis specific adhesion junctions known as ectoplasmic specializations, and the structures develop prior to release of spermatids from the epithelium. Second, integral membrane junction proteins have been localized in tubulobulbar complexes and are concentrated at their ends (Guttman et al., 2004b; Young et al., 2012). Third, these proteins colocalize with early

Fig. 3. Evidence of cortactin knockdown. (A) Western blots of seminiferous epithelial lysate from five animals indicate weaker cortactin bands from experimental testes in comparison to control. Calnexin was used as a loading control. (B) Immunofluorescence of stage VII spermatids labeled for cortactin (green) and DAPI (blue). Cortactin staining outlines individual tubulobulbar complexes from experimental and control testes and reveals a difference in length. Bar = 5 μm. (C) Quantitation of knockdown was determined by probing cryosections from control and experimental tissue labeled for cortactin. Mature spermatids were randomly selected and set to a predetermined threshold using ImageJ software. The number of positive pixels was determined using the “analyze particles” function. The number of pixels was considered to be an index of tubulobulbar complex length in each cluster. Experimental tubulobulbar complexes were shown to be significantly different from control tubulobulbar complexes (P=0.0028) (D) Quantitation of knockdown determined by probing cryosections from control and experimental tissue labeled for filamentous actin. Experimental tubulobulbar complexes were shown to be significantly different from those in controls (P=0.0004).
endocytic markers Rab5 and EEA1 (Young et al., 2009b; Young et al., 2012). Significantly, nectin 3, one of the adhesion molecules present in the spermatid plasma membrane, is concentrated at the ends of tubulobulbar complexes and is absent from spermatozoa after release from the epithelium. These data, together with the observation that the double membrane bulks bud from the structures, are consistent with the conclusion that apical tubulobulbar complexes are subcellular endocytic machines that internalize ‘intact’ intercellular adhesion junctions into Sertoli cells.

If tubulobulbar complexes associated with spermatids do in fact internalize junctions, then perturbing their structure/function should prevent junction turnover ultimately delaying or preventing sperm release from the epithelium. In this study we used an in vivo siRNA approach to knockdown cortactin, a protein concentrated at tubulobulbar complexes in the seminiferous epithelium (Young et al., 2009a), and a component of the actin cuff surrounding the central double-membrane core of each structure.

Cortactin is known to coordinate membrane dynamics and cytoskeletal remodeling in other systems (Schafer, 2002) – two features that also make it a key player during the process of clathrin-mediated endocytosis (Merrifield et al., 2005). Two of cortactin’s five functional domains allow it to simultaneously bind F-actin and Arp2/3. Therefore, cortactin not only promotes nucleation and branching but also acts to stabilize newly formed branches (Weaver et al., 2001). These features, paired with cortactin’s affinity to bind newly incorporated actin molecules (Bryce et al., 2005) make it a key component in the formation of an actin network.
Temporarily silencing the expression of cortactin leads to unstable actin-based structures in a number of systems. In the absence of cortactin, lamellipodial protrusions retract due to instability (Bryce et al., 2005). Cortactin knockdown in osteoclasts causes a loss of podosome formation (Tehrani et al., 2006). Significantly, podosomes somewhat resemble tubulobulbar complexes (Young et al., 2009b) in that they are tubular structures with a central membrane core cuffed by a dendritic actin network.

Here we show that when administered to rats by intratesticular injection, siRNA reagents enter Sertoli cells and caused a knockdown of cortactin in the seminiferous epithelium. As observed in other systems (Tehrani et al., 2006) where knockdown of cortactin prevented the formation of some structures without altering other cortactin containing features, reduced cortactin levels in the seminiferous epithelium altered the phenotype of tubulobulbar complexes without appearing to effect actin bundles in ectoplasmic specializations where cortactin also has been localized (Kai et al., 2004; Young et al., 2009b).

When labeled for cortactin and/or actin, tubulobulbar complexes were clearly shorter in epithelia from testes injected with targeting siRNA reagents relative to similar epithelia from contralateral testes injected with non-targeting siRNA reagents. This result was confirmed quantitatively using a threshold method to evaluate the numbers of positive pixels generated by labeling tubulobulbar clusters adjacent to spermatid heads. Using this method, the pixels above threshold remaining in each image were counted and considered an index of tubulobulbar complex length. Numbers of pixels present in tubulobulbar complexes from knockdown animals were significantly different from controls, both for actin and cortactin stained material, supporting our conclusion that cortactin knockdown reduces tubulobulbar complex length.

Significantly, in knockdown epithelium we detected a delay in retraction of Sertoli cell away from mature spermatids and spermiation failure. Moreover, in cases where spermatids remained attached to the epithelium, adhesion junctions (ectoplasmic specializations) were also present, an observation consistent with a failure of the normal junction removal mechanism.

**Fig. 6. Evidence of delayed spermiation.** Light micrographs of Stage VIII seminiferous tubules from control (A,B) and cortactin targeted siRNA (C) injected testes. Notice that in panel (C) Sertoli cell cytoplasm (arrows) still engulfs the spermatid heads (arrowheads) at the apex of the epithelium whereas this cytoplasm has been withdrawn in controls. Also notice that there is one spermatid head (arrow marked with asterisk) that is positioned deep in the epithelium in (C). Bar = 10 μm.

**Fig. 7. Evidence of spermiation failure and persistence of adhesion junctions in cortactin siRNA testes.** (A–F) 1 μm thick plastic sections stained with toluidine blue of stages VIII (late)/IX of spermatogenesis. In sections from control testes (A,B), spermatids are in the final stages of release (arrowhead in (A)) or have been released as spermatozoa from the epithelium (B). In comparable sections from cortactin siRNA testes (C–F), numerous spermatids (arrowheads) are observed at the apex of the epithelium, some still surrounded by the apical process of the Sertoli cell (arrowhead in (C)). Others are positioned deeper in the epithelium (arrowheads in D,E,F). Bar = 10 μm. When viewed at the ultrastructural level, spermatids being released from the epithelium at late stage VIII in sections of control testes lack ectoplasmic specializations in Sertoli cell regions adjacent to spermatid heads (arrowheads in (G,H) and apical regions of the Sertoli cell are being withdrawn from the heads. In sections at stages VIII (late)/IX of cortactin siRNA testes (I,J), ectoplasmic specializations are present adjacent to spermatid heads that are positioned in the epithelium similar to those shown in panels D,E,F. Bar = 200 nm.
We propose that short tubulobulbar complexes in the knockdown epithelia cause a failure of the structures to completely internalize intercellular adhesion junctions between Sertoli cells and spermatids and that this results in a delay in and possibly eventual failure of sperm release. Because we only evaluated one time point (3 days) after siRNA injections, we cannot distinguish between (1) an initial complete disassembly of tubulobulbar complexes followed by partial recovery over the 3-day time period, and (2) a continuous prevention of tubulobulbar complexes to develop to their full length. Collins and coworkers (Collins et al., 2011), on the basis of a high resolution ultrastructural study, conclude that one of the actin filament network’s primary roles during clathrin-mediated endocytosis is to elongate the bud neck in order to drive the endocytosed vesicles away from the plasma membrane. This is consistent with our observation of short tubulobulbar complexes after cortactin knockdown. Due to a lack of stability in the actin network caused by a depletion of cortactin, tubulobulbar complexes may not be able to elongate and acquire or maintain normal length and thereby fail to internalize the same amount of junctional membrane.

The time period (3 days or roughly 72 hrs) from injection of the siRNA reagents until evaluation of phenotype in our study was sufficient for us to observe effects on tubulobulbar complexes and on sperm release. In the rat, stage VII of spermatogenesis, the stage when tubulobulbar complexes are subcellular machines that internalize intact intercellular adhesion junctions and that their function is essential for normal sperm release. Significantly, these findings introduce a new paradigm for junction internalization generally in cells and link the mechanism to a biologically significant event – sperm release.

Interestingly, cortactin homozygous knockout mice appear phenotypically normal and live a normal life span (Schnoor et al., 2011); however, their fertility and testicular morphology have not been reported. Similarly, the anti-androgen flutamide is reported to reduce testicular cortactin levels and to cause phenotypic changes in ectoplasmic specializations (Anahara et al., 2006), but any effects on tubulobulbar complex morphology or on spermatogenesis have not been evaluated.

Our data presented here are consistent with the hypothesis that tubulobulbar complexes are subcellular machines that internalize intercellular adhesion junctions and that their function is essential for normal sperm release. Significantly, these findings introduce a new paradigm for junction internalization generally in cells and link the mechanism to a biologically significant event – sperm release.

Materials and Methods
Reagents
The paraformaldehyde was obtained from Fisher Scientific (Ottawa, ON, CAN). The secondary antibodies and phallotoxins conjugated to Alexa fluorochromes were obtained from Invitrogen (Burlington, ON, CAN) and those conjugated to horseradish peroxidase were purchased from Jackson ImmunoResearch Laboratories, Inc. (Westgrove, USA). Unless otherwise indicated, all other reagents used in the studies were obtained from Sigma–Aldrich (Oakville, ON, CAN).

siRNAs
An ON-TARGETplus pool of four siRNAs synthesized by Dharmacon (Lafayette, CO) was used to target rat cortactin (Cttn). An ON-TARGETplus non-targeting pool – a pool of four non-targeting siRNAs – was used as a negative control.

For dye modification studies, the same pool of four oligonucleotides was used with one of the four duplexes having a DYS474 modification on the 5’ end of the sense strand.

PEI-mediated siRNA delivery
siRNAs were mixed with in vivo-jetPEI (Polyplus, Illkirch, France) at an N/P ratio of 1/8 at room temperature. For intratesticular injection administration, 50 μl of the Cttn targeting siRNA mixture was injected into right testis and 50 μl of the non-targeting control was injected into the left testis.
Animals

Animals used in this study were reproductively active male Sprague-Dawley rats. They were obtained from Charles River Animal Research Laboratories. The rats were maintained according to guidelines established by the Canadian Council on Animal Care and using protocols approved by the UBC and SFU animal care committees. All experiments were done at least in duplicate using tissue from different animals.

Animal surgeries

Handling

All animals were handled daily for a minimum of one week before surgical treatment to decrease animal anxiety around investigators and staff. Handling included weighing and a brief physical inspection. A food ‘treat’ was given after each handling session.

Surgical preparation

Animals were weighed and anesthesia was induced with 5% isoflurane. Anesthesia was maintained with 2–3% isoflurane for the remainder of the surgical protocol. Surgical sites were shaved and then washed 3 times with surgical soap followed by 70% ethanol. Lacerlube was applied to protect the eyes, and a lactate ringer’s solution (LRS) was injected subcutaneously to maintain hydration.

Pre-emptive pain management consisted of a single dose each of buprenorphine at 0.05 mg/kg and of ketoprophen at 5.0 mg/kg injected subcutaneously. In addition, a local analgesia, bupivacaine, was injected subcutaneously along each incision site.

Surgical procedure

After surgical preparation, animals were transferred to a surgical table equipped with a heat source. Heart rate and oxygen saturation were monitored with a pulse oximeter equipped with a foot clip sensor. Deep anesthesia was ensured by a lack of withdraw reflex upon toe pinch. Using sterile technique, an incision of approximately 1–1.5 cm was made through the right side of the scrotum and skin was opened to expose the spermatic fascia enclosing the testis. An additional incision was made through the fascia to expose the surface of the testis. The testis was positioned to avoid major blood vessels and 32-gauge needle attached to a Hamilton syringe system was inserted through the capsule and into the testis. A total volume of 50 μL of Cttn targeting siRNA and PEI mixture was injected into the testis. The needle was withdrawn and the spermatheca fascia sutured closed with two or three discontinuous sutures. Subeuticular sutures were used to close the skin incision. Using the same surgical procedure, 50 μL of a non-targeting siRNA and PEI mixture were injected into the left testis as a control.

Post-surgical care and processing

After surgery, rats were placed in a cage under an infrared heater and allowed to recover. The cage contained clean, dry bedding, hydrogel and bacon softies. In most cases, animals were up and eating within 30 minutes. Animals were then transferred into their home cages with their original cage mates that also had the same surgery on the same day. Animals were monitored closely for the remainder of the experiment.

After three days, and in each experiment, animals were deeply anesthetized and the testes removed for processing. The testes of two animals were fixed for electron microscopy, those of another two animals were fixed and cryo-sectioned for immunofluorescence analysis and those of a third pair were processed for electron microscopy, those of another two animals were fixed and cryo-sectioned into the left testis as a control.

Immunofluorescence

Tissue preparation

Isolated testes were removed from anesthetized animals and perfused briefly with warm PBS to clear blood and then perfused with warm fixative (PBS, 3% paraformaldehyde, pH 7.3) for 30 min.

Frozen sections

Fixed testes were frozen onto aluminum stubs with OCT compound (Sakura Finetek USA, Torrence, CA) by immersion in liquid nitrogen. Sections of 5 μm were cut using a cryo-microtome, attached to poly-L-lysine coated glass slides, immediately plunged into ~20°C acetone for 5 min, air dried, and then processed for immunofluorescence.

Immunostaining

Slides with attached cryosections were rehydrated and blocked with 5% normal goat serum (NGS) in TBPS-BSA (PBS containing 0.5% Tween-20 and 0.1% bovine serum albumin) for 20 min at room temperature. The primary antibody used to probe cryosections was rabbit anti-cortactin (KE-20) (Sigma–Aldrich, St. Louis, MO, USA). The primary antibody added to the slides was made up in TBPS-BSA with 1% NGS, and incubated overnight at 4°C in a humidity chamber. The material was washed extensively with the TBPS-BSA (wash buffer), then incubated for 60 minutes at 37°C with secondary antibody conjugated to a fluorochrome (goat anti-rabbit AlexaFluor 568 (Invitrogen)). The slides were again washed and then coverslips were mounted using Vectashield (Vector Labs, Burlington, ON). The tissue was visualized using a Zeiss Axioskop microscope fitted with appropriate filter sets for detecting fluorescence and with appropriate optics for phase microscopy and an inverted Zeiss Axiovert LSM 5 confocal microscope equipped with epi-fluorescence and LSM 5 Pascal software. Image acquisition was done with the LSM 5 Pascal software.

Filamentous actin was labeled by making up AlexaFluor 568 phalloidin in TBPS-BSA and staining slides for 20 minutes at room temperature. Slides were then extensively washed in TBPS-BSA.

Immunoblotting

Western blots were performed on either whole testis alone, or on whole testis and seminiferous epithelium lysates both from experimental and control (non-targeting siRNA) testes. Blots were probed first with cortactin antibody, and then the blots were stripped and stained with an antibody to calnexin as a loading control. For whole testis lysates, testes were removed from rats under deep anesthesia and decapsulated. The whole testis tissue was extensively homogenized in RIPA lysis buffer (150 mM NaCl, 50 mM Tris, pH 7.4, 5 mM EDTA, 1% Nonidet P-40, 1% deoxycholic acid [sodium salt], 10% SDS) with complete, mini-EDTA-free protease inhibitor (Roche, Mississauga, ON, CAN).

For blots run on both whole testis and seminiferous epithelium, half of each testis was prepared as described above and the other half was cut into small pieces in ice-cold PEM/250 buffer (0.2 M Peps, 0.1M EGTA, 0.1M MgCl2, Sucrose). The small pieces of seminiferous tubules were collected, placed in fresh PEM/250 buffer and epithelium was separated from tubule walls as described elsewhere (Guttmann et al., 2007). The isolated sheets of epithelium were collected and pelleted by centrifugation. Pellets were resuspended and extensively homogenized in RIPA buffer.

Lysates were loaded into wells of 1-mm-thick 10% SDS-PAGE gels at equal concentrations as determined by protein assay and run according to standard protocols (Laemmli, 1970). Proteins were transblotted onto Immobilon-P transfer membrane (Millipore, Billerica, MA) and then washed for 5 minutes at room temperature with TBST (500 mM NaCl, 50 mM Tris, 0.1% Tween-20). The blots were blocked for 12 hours at 4°C using 4–5% non-fat milk (Blotto, Santa Cruz Biotechnology, Santa Cruz, CA). Membranes were washed 3 times, 5 minutes each wash, and then incubated for 1 hour at room temperature with rabbit-anti-cortactin (Sigma–Aldrich, Oakville, ON, CAN) at a concentration of 1:10,000. Blots were washed 3 times; 5 minutes each wash with TBST and then incubated for 1 hour at room temperature with goat anti-rabbit horseradish peroxidase-conjugated antibody. Blots were washed 3 times; 5 minutes each with TBST and then reacted with ECL (GE Lifesciences, Pittsburgh, PA, USA) and bands were visualized with Bioflex EC film (Interscience, Markham, ON, CAN).

Controls for loading were performed by stripping membranes with methanol for 10 minutes at room temperature. After stripping, blots were washed in TBST 3 times for 5 minutes each wash. The blots were blocked for 12 hours at 4°C using 4–5% non-fat milk. Membranes were washed 3 times, 5 minutes each wash, and then incubated for 1 hour at room temperature with rabbit anti-calnexin (Sigma–Aldrich, Oakville, ON, CAN) at a concentration of 1:80,000. Blots were then washed 3 times; 5 minutes each wash with TBST and then incubated for 1 hour at room temperature with goat anti-rabbit horseradish peroxidase-conjugated antibody. Blots were washed 3 times, 5 minutes each with TBST and then reacted with ECL (GE Lifesciences, Pittsburgh, PA, USA) and bands were visualized with Bioflex EC film (Interscience, Markham, ON, CAN).
Relative quantitation and statistical analysis of actin and cortactin fluorescence as an indication of tubulobulbar complex length
Quantitation of knockdown was determined by probing cryosections of tissue from control and experimental tissue labeled for cortactin or for actin. Ten mature spermatids with associated labeled clusters of tubulobulbar complexes at stage VII of spermatogenesis from each control testis and ten cells from each experimental testis were randomly selected and each cell was set to a predetermined threshold using ImageJ software (Abramoff et al., 2004). When set to a common threshold, the number of positive pixels was determined using the “analyze particles” function. The number of pixels was considered to be an index of tubulobulbar length in each cluster.

Statistics
To determine if there were any significant differences between the mean values from experimental and then control animals, a one-way analysis of variance was used (ANOVA). No significant differences were found (P>0.05) and mean values were combined. Mean values ± SD are indicated in figure legends (Fig. 3C,D, Fig. 8D); one-tailed, unpaired t test (with Welch’s correction) for statistical significance.

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Competing Interests
The authors have no competing interests to declare.

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