Centrobin is essential for C-tubule assembly and flagellum development in *Drosophila melanogaster* spermatogenesis

Jose Reina¹*, Marco Gottardo²*, Maria G. Riparbelli², Salud Llamazares¹, Giuliano Callaini², and Cayetano Gonzalez¹,³

Centrobin homologues identified in different species localize on daughter centrioles. In *Drosophila melanogaster* sensory neurons, Centrobin (referred to as CNB in *Drosophila*) inhibits basal body function. These data open the question of CNB's role in spermatocytes, where daughter and mother centrioles become basal bodies. In this study, we report that in these cells, CNB localizes equally to mother and daughter centrioles and is essential for C-tubules to attain the right position and remain attached to B-tubules as well as for centrioles to grow in length. CNB appears to be dispensable for meiosis, but flagellum development is severely compromised in Cnb mutant males. Remarkably, three N-terminal POLO phosphorylation sites that are critical for CNB function in neuroblasts are dispensable for spermatogenesis. Our results underpin the multifunctional nature of CNB that plays different roles in different cell types in *Drosophila*, and they identify CNB as an essential component for C-tubule assembly and flagellum development in *Drosophila* spermatogenesis.

Introduction

Centrobin was initially identified in humans through a yeast two-hybrid screen for proteins that interact with the tumor suppressor BRCA2 (Zou et al., 2005). Like human Centrobin (referred to as CNTROB in humans), its homologues in other species are components of daughter centrioles in different cell types (Zou et al., 2005; Januschke et al., 2011, 2013; Gottardo et al., 2015). In mammals, CNTROB has been reported to be required for centriole duplication and elongation as well as for microtubule nucleation and stability through its binding to tubulin and its effect in stabilizing centrosomal P4.1-associated protein (CPAP; Jeong et al., 2007; Jeffery et al., 2010, 2013; Gudi et al., 2011, 2014, 2015; Shin et al., 2015).

*Centrobin* homologues identified in different species localize on daughter centrioles. In *Drosophila* primary spermatocytes (Singh et al., 2014). In *Drosophila* type I sensory neurons, CNB functions as a negative regulator of ciliogenesis: CNB depletion enables daughter centrioles as functional basal bodies that template ectopic cilia, and mother centrioles modified to carry CNB cannot function as basal bodies (Gottardo et al., 2015).

The function of a daughter centriole protein like CNB as a negative regulator of ciliogenesis opens the question of CNB’s role in *Drosophila* primary spermatocytes where after centrosome duplication all four centrioles, mothers and daughters alike become basal bodies that assemble axonemal-like structures, which are the precursors of sperm flagella (Tates, 1971; Riparbelli et al., 2012, 2013; Gottardo et al., 2013). To investigate this issue, we have studied CNB localization and function in spermatogenesis.

Results and discussion

**CNB localizes equally to mother and daughter centrioles in *Drosophila* primary spermatocytes**

As in other cell types in *Drosophila* and other species, CNB localization in spermatogonial cells was asymmetric, restricted to only one of the two centrioles that were labeled with anti-SAS4 antibodies (Fig. 1A, red and green, respectively; n = 86 cells from

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1. Institute for Research in Biomedicine (IRB Barcelona), The Barcelona Institute of Science and Technology, Barcelona, Spain;
2. Department of Life Sciences, University of Siena, Siena, Italy;
3. Institución Catalana de Recerca i Estudis Avançats, Barcelona, Spain.

*Jose Reina and M. Gottardo contributed equally to this paper. Correspondence to Cayetano Gonzalez: gonzalez@irbbarcelona.org; M. Gottardo’s present address is Center for Molecular Medicine, University of Cologne, Cologne, Germany.*

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Centrobin is required for C-tubule assembly

CNB is required in primary spermatocytes for basal bodies to achieve normal length and for the proper assembly of the C-tubule

To determine whether CNB has a function in the basal body–CLR complex, we examined trans-heterozygous PBac(RB) Cnb[e00267]/Df(3L)ED4284 (henceforth referred to as Cnb mutant) spermatocytes. We found that the length of YFP-Asterless (ASL) signal (a proxy for centriole length) was significantly shorter ($P < 0.001$) in Cnb mutant (0.97 ± 0.28 µm) than in WT males (1.55 ± 0.11 µm; Fig. 2, A and C, gray bars). Longitudinal EM sections from centriole pairs in which both mother and daughter were cut along their entire length confirmed this observation (Fig. 2, B, D, and E) and revealed that daughter basal bodies were shorter than their mothers in Cnb mutant spermatocytes (0.62 ± 0.07 µm and 0.78 ± 0.04 µm, respectively; $n = 5$ centriole pairs; Fig. 2 E and Table S2). This was not the case in WT spermatocytes, where as previously reported (Riparbelli et al., 2012), we found that mother and daughter basal bodies were of equal length (1.02 ± 0.04 µm and 1.00 ± 0.05 µm, respectively; $n = 5$ centriole pairs; Fig. 2 E and Table S2). EM sections also revealed that 10 out of 16 mutant axonemes sectioned longitudinally were irregularly...
shaped and had abnormal axonemes with doublets that were aberrantly interrupted halfway through the CLR (Fig. 2 D).

We then examined the ultrastructural details revealed by transversal EM sections. As reported previously (Gottardo et al., 2015), unlike basal bodies in sensory neurons that are made of doublets, basal bodies in WT spermatocytes are made of triplets (Fig. 3, A and A’). At the transition zone, the outermost tubule (C-tubule) progressively lost protofilaments (Fig. 3, B and B’), hence becoming an open longitudinal sheet that extended beyond the transition zone along the axoneme (Fig. 3, C and C’, red arrow). At the point where C-tubule remnants and B-tubules intersect, short radial projections could be observed (Fig. 3, C and C’, blue arrow; Riparbelli et al., 2012).

In addition to this basic layout of triplets, doublets, and open sheaths, about half of basal body–CLR complexes in Drosophila WT spermatocytes present a luminal singlet microtubule independently of the Z position of the cross-section (Tates, 1971; Carvalho-Santos et al., 2012; Riparbelli et al., 2012). The presence of this luminal singlet is as erratic in Cnb mutant as it is in WT males and does not represent a Cnb mutant phenotype.

We serially sectioned 11 basal body–CLR complexes from two Cnb mutant males. In all 11 samples, C-tubules were either misplaced outwards, away from the nearly straight line defined by the A/B doublet (Fig. 3, D and D’, blue), or lost, leaving only duplets behind (Fig. 3, D, D’, G, and G’, green; Table S1). These abnormalities extended distally along the axoneme, where C-tubule–derived longitudinal sheets were often lacking in Cnb mutant spermatocytes (Fig. 3, E–F’ and H–I’, red arrow). The short radial projections observed in WT axonemes (Fig. 3, C and C’, blue arrow) also appeared to be lacking in Cnb mutant CLR’s (Fig. 3, F, F’, I, and I’, blue arrow), but this observation must be interpreted with caution because of the technical difficulties of detecting such small structures that are located in regions of electron-dense material.

These observations reveal that CNB is required in Drosophila spermatocytes for centrioles to properly position and stabilize C-tubules, grow in length up to full-sized basal bodies, and template normal axonemes. Our results identify CNB as one of the very few proteins known so far to be required for C-tubule assembly, including Δ-tubulin in Paramecium tetraurelia (Garreau de Loubresse et al., 2001) and Chlamydomonas reinhardtii (Dutcher and Trabuco, 1998) along with UNC and SAS4 in Drosophila (Gottardo et al., 2013; Zheng et al., 2016). Notably, as in Cnb mutant spermatocytes, basal bodies are also shorter than normal in Sas4f122a and Unc mutant spermatocytes (Gottardo et al., 2013; Zheng et al., 2016), hence suggesting that short centrioles and lacking or mispositioned C-tubules may be causally linked.

Drosophila SAS4 and its human homologue CPAP are closely functionally related to the corresponding Centrobin homologues CNB/CNTRB. Mutants in the N-terminal helical motif of the PN2-3 domain of CPAP and SAS4 cause overelongation of newly formed centriolar/ciliary microtubules, and so does CNTRB overexpression by stabilizing and driving the centriolar localization of CPAP (Gudi et al., 2011, 2014, 2015). Moreover, the mutants in PN2-3 domain of CPAP referred to above result in biciliated cells, and so does Cnb loss of function in Drosophila sensory neurons.
that presented a uniform nuclear size (Fig. S1 B), which is a very sensitive readout of faithful meiotic chromosome segregation (González et al., 1989). These results strongly suggest that shortened centrioles and lacking C-tubules have no major observable consequences on meiosis progression.

However, soon after meiosis, when the basal body attaches to the nucleus and the axoneme starts to grow, pushing the old CLR caudally, Cnb mutant spermatids presented CLRs that were much longer than those from WT cells: 5.6 ± 1.4 μm (n = 7) and 1.5 ± 0.8 μm (n = 6), respectively (Fig. 4, A and B). Abnormally long CLRs have also been reported in Unc mutant early spermatids (Gottardo et al., 2013). In WT cells at this stage, CNB signal was strongest on the basal body proximal to UNC and was also detectable over the perinuclear plasm (Tates, 1971), a cap over the nuclear hemisphere where the basal body is engaged (Fig. S1 C).

Cnb mutant phenotypes were also conspicuous at later stages of elongation. Transversal sections through the tails of WT elongating spermatids showed two mitochondrial derivatives flanking the axoneme, which was almost completely surrounded by a double membrane (Fig. 4 C). Among the sample of 32 Cnb mutant elongating spermatids that we analyzed in detail (belonging to three cysts from two different males), we found a wide range of phenotypic variability (Fig. 4, D–G). Five presented a fairly WT 9 + 2 configuration, whereas the other 27 included cells that had no axoneme (n = 5; Fig. 4 D, NA) or axonemes with missing or highly disarranged duplets and in which the surrounding membrane was widely open (n = 22; Fig. 4, E and F). Among the latter, six had some duplets that carried C remnants, and 16 did not. Interestingly, not all duplet-bound open sheaths observed in Cnb mutant cells were C remnants, such as for instance the one shown in Fig. 4 F that originated from an A-tubule and was oriented in the wrong direction. We also found four axonemes that presented one to three triplets with an attached open sheath (Fig. S2 A, arrows). No such structures have been reported in WT cells. The presence of a fraction of Cnb mutant spermatids with axonemes that presented C remnants suggests that a certain fraction of Cnb mutant primary spermatocytes retain C-tubules.

In addition to lacking and disarranged axonemes, Cnb mutant elongating spermatids also presented ectopic nuclei at sections where only tails were present in WT cysts (Fig. 4 G, Nu, and Fig. S2 B). Scattered nuclei have also been reported in other Drosophila mutants like Unc and Sas4 (Baker et al., 2004; Riparbelli and Callaini, 2011). Head-to-tail attachment defects have been documented in the sperm of a rat CNTROB mutant strain (hd rats) that was male sterile (Liška et al., 2013).

Our results reveal that CNB depletion brings about a spectrum of mutant phenotypes at different stages of spermatogenesis that includes shorter centrioles, mispositioned or absent C-tubules, lack of C-tubule derivatives, abnormal or absent axonemes, and scattered nuclei along elongating spermatid bundles. We cannot discard that CNB may have independent functions accounting for each of these phenotypes. However, the apparent pleiotropic effect of CNB loss may simply reflect the downstream consequences of centrioles that are shorter than normal and present C-tubule defects.

**Amino acids T4, T9, and S82, which are critical for centriole asymmetry in Drosophila neuroblasts, are dispensable for CNB function in spermatogenesis**

Expression of YFP-CNBWT, a fusion protein between YFP and WT CNB, rescued all the phenotypes that we observed in Cnb mutant males. These included (A) the ultrastructural abnormalities, which affected premeiotic basal body–CLR complexes and postmeiotic axonemes (Fig. 5, A–D), (B) centriole (YFP-ASL) length (Fig. 5 I), which was rescued from 0.97 ± 0.28 μm (n = 27) in Cnb mutant males to 1.47 ± 0.16 μm (n = 20), very close to WT length (1.55 ± 0.11 μm, n = 21), and (C) fertility (Fig. 5 J), which was rescued to 81 ± 7, which is indistinguishable from WT levels (87 ± 4; offspring per male in single-pair mating tests; n = 15). These results confirm that all these phenotypes are indeed brought about by Cnb loss of function and are not caused by other unrelated mutations.

Remarkably, all these Cnb mutant traits were also rescued by the YFP-CNB4A-T9A-S82A transgene (Januschke et al., 2013).
Basal bodies, CLRs, and spermatids recovered WT ultrastructure including properly positioned C-tubules and C-tubule remnants (Fig. 5, E–H); centriole (YFP-ASL) length was recovered to 1.26 ± 0.14 µm (n = 17); and, notably, the amount of offspring per male in single-pair mating experiments was also fully recovered to WT levels (82 ± 7; n = 15; Fig. 5 J).

YFP-CNBT4A-T9A-S82A carries a mutant version of CNB in which three consensus POLO phosphorylation sites have been substituted by alanine. In larval neuroblasts, YFP-CNBT4A-T9A-S82A retains specific daughter centriole binding, but unlike YFP-CNBT4A-T9A-S82A, it does not rescue the loss of PCM and microtubule aster during interphase caused by Cnb loss of function (Januschke et al., 2013). The efficient rescue of Cnb mutant traits in spermatogenesis by YFP-CNBT4A-T9A-S82A strongly suggests that CNB phosphorylation by POLO in these three critical sites is not necessary for CNB function during spermatogenesis, hence underpinning the diversity of cell type–specific molecular functions of CNB.

The different, and in some aspects opposite, roles of CNB in basal body structure and ciliogenesis in type I sensory neurons and spermiogenesis are remarkable. In neurons, CNB inhibits basal body function on the centriole to which it binds in WT cells is only the daughter centriole (Gottardo et al., 2015). In primary spermatocytes, however, CNB localizes to both mother and daughter centrioles, which in these cells are equally able basal bodies, and acts as a positive regulator of ciliogenesis (Tates, 1971; Fritz-Niggli and Suda, 1972; Gottardo et al., 2013). One conspicuous difference between these two cell types is that basal bodies in spermatocytes contain C-tubules, whose assembly, positioning, and stability requires CNB. In vertebrates, where centrioles also have C-tubules, a recent study has found that in addition to its primary localization to daughter centrioles, CNTROB associates with mother centrioles at the onset of ciliogenesis, and CNTROB loss causes defective axonemal extension (Ogungbenro et al., 2018). These results suggest that the different effect that CNB depletion has on ciliogenesis in Drosophila neurons and spermatocytes may reflect the doublet versus triplet structure of the corresponding basal bodies.
Materials and methods
Fly stocks and husbandry
The following fly stocks were used in this study: PBac(RB) Cnbe00267 (Exelexis); Df(3L)ED4284 (Bloomington Drosophila Stock Center); pUbqYFP-ASL (Rebollo et al., 2007); ANAI-GFP (from T. Avidor-Reiss; Blachon et al., 2008); pUbq YFP-CNBP and pUbq YFP-CNBP ASLΔT4A.T3R2A (Januschke et al., 2013); and UNC-GFP (Baker et al., 2004). Throughout the study, Cnb mutant refers to PBac(RB) Cnb e00267/Df(3L)ED4284. This allelic combination causes the loss of interphase asters in larval neuroblasts as effectively as Cnb-RNAi driven from ubiquitous promoters. However, unlike RNAi-driven CNB depletion, this allelic combination does not have any significant effect on adult viability (Januschke et al., 2013). All crosses were performed at 25°C.

Fertility tests
Fertility tests were performed by scoring the offspring from each of 15 males individually mated to three w1118 females.

Immunocytochemistry
Testes from pupae were dissected in PBS and placed in a small drop of 5% glycerol in PBS on a glass slide, squashed under a small coverslip, and frozen in liquid nitrogen. After removal of the coverslip, the samples were immersed in methanol for 10 min at −20°C followed by 15 min in PBS and 1 h in PBS containing 0.1% BSA (PBS-BSA). Samples were then incubated with primary antibodies in a humid chamber either for 1 h at room temperature or overnight at 4°C, washed in PBS-BSA three times for 10 min, incubated for 1 h at room temperature with Alexa Fluor-conjugated secondary antibodies (Invitrogen), and then washed again as before. Immunostained preparations were mounted in small drops of Vectashield (Vector Laboratories). Images were taken either on an Axiolab was purchased using a 63× 1.47 NA objective (Leica Microsystems). Acquired images were processed using ImageJ (1.48s; National Institutes of Health). Grayscale digital images were collected separately and then pseudocolored and merged using Photoshop (7.0; Adobe). Final figures were prepared using Illustrator (Adobe). First antibodies: rabbit anti–DSas-4 (from J. Gopalakrishnan; Gopalakrishnan et al., 2011), DM1a anti–α-TUB (1:500; T9026; Sigma-Aldrich), rabbit anti-Asl (1:500; Januschke et al., 2013), mouse antiacetylated Tub (1:500; T6793; Sigma-Aldrich), chicken anti-PLP (from A. Rodrigues-Martins; Rodrigues-Martins et al., 2007), and rabbit anti-CNB (1:500). Alexa Fluor conjugates were used at 1:1,000 dilution. DNA was stained with DAPI.

Production of anti-CNB antibody
A PCR fragment encoding amino acids 235–549 of CNB was cloned into the pHAT vector (European Molecular Biology Laboratory). The resulting 6×His protein fusion protein was purified using TALON resin (Takara Bio Inc.) according to the manufacturer’s instructions and was used to generate rabbit antibodies at Harlan (ENVIGO).

Fluorescence-based estimation of centriole length
As a proxy for centriole length, we measured the distance between the distal end of the YFP-ASL signal and the point at which both centrioles intersect on the internal part of the V shape. Because emitted light scatters over an area that is larger than the actual fluorescent particle, measurements based on fluorescent labels overestimate the actual particle’s size. EM-based measurements provided much more precise data.

Phase-contrast microscopy
Phase-contrast microscopy of nonfixed testes was performed as previously described (Glover and González, 1993). In brief, testes were dissected and placed in a small drop of PBS on a siliconized slide, cut at about one third from its apical end with tungsten needles, covered with a coverslip, and gently squashed by removing the PBS with a small piece of blotting paper. We used a 40× dry phase-contrast objective. Images were acquired with an AxioCam MRm camera (ZEISS).

Transmission EM
Testes isolated from larvae and pupae were prefixed in 2.5% glutaraldehyde buffered in PBS overnight at 4°C. After prefixation, the material was carefully rinsed in PBS and postfixied in 1% osmium tetroxide in PBS for 2 h at 4°C. Samples were then washed in the same buffer, dehydrated in a graded series of ethanol, embedded in a mixture of epon-Araldite, and polymerized at 60°C for 48 h. 50–60-nm-thick sections were obtained with an Ultracut E ultramicrotome (Reichert) equipped with a diamond knife, mounted on copper grills, and stained with uranyl acetate and lead citrate. Images were taken with either a CM 10 transmission electron microscope (Philips) operating at an accelerating voltage of 80 kV or a Tecnai Spirit transmission electron microscope (FEI) operating at 100 kV equipped with a Morada charge-coupled device camera (Olympus).

Statistical analysis
We performed one-way ANOVA tests using Prism (GraphPad Software). Error bars represent SEM.

Online supplemental material
Fig. S1 shows EM sections showing that basal body–CLR complexes in Cnb mutant early spermatids retain the ultrastructural defects observed in early spermatoocytes, immunofluorescence and phase-contrast microscopy documenting normal meiosis in Cnb mutant males, and immunofluorescence staining of CNB in WT spermatids. Fig. S2 shows ultrastructural details of a Cnb mutant axoneme that presents triplets with open sheaths as well as immunofluorescence showing scattered nuclei over the tails in Cnb mutant spermatids. Table S1 shows the number of C-tubules observed in 11 Cnb mutant basal bodies. Table S2 shows the mother and daughter basal body length in WT and Cnb mutant males.

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