## SUPPLEMENTAL INFORMATION

## Baffet\_Fig S1



Fig. S1. Expression of a wild-type dTBCB transgene fused to GFP rescues MT and polarity defects in a *dTBCB<sup>1</sup>* background

(A-B") Stage 10 egg chambers, expressing a *Ubi-\alpha-RFP-Tubulin* transgene and fixed to visualize the follicle cell MT network. Chambers are oriented with anterior to the left. (A) Wild-type. (B-B") *dTBCB<sup>1</sup>* mutant egg chamber expressing *Ubi-dTBCB-GFP* transgene. In (B") the rescued follicle cells show a normal MT network compared to the control (A).

(C-D") Stage 10 egg chambers, immunostained for Staufen (Stau) protein. (C) Wild-type. (D-D") *dTBCB*<sup>1</sup>mutant egg chamber expressing *Ubi-dTBCB-GFP* transgene. In (D"), the rescued oocyte shows a normal posterior Stau crescent, compared to the control (C).

(E-F") Stage 8 egg chambers, immunostained for aPKC. (E) Wild-type. (F-F")  $dTBCB^{l}$  mutant egg chamber expressing *Ubi-dTBCB-GFP* transgene. In (F"), the rescued follicle cells show a normal apical aPKC staining compare to the control (E). In rare circumstances dTBCB-GFP expression is incomplete (white bracket), in such cases the rescue of aPKC localization failed to occur.



Fig. S2. Overexpression of dTBCB-GFP in S2R+ *Drosophila* cells led to MT depolymerization

(A-B) Strong overexpression of the *Ubi-dTBCB-GFP* transgene (green) in S2R+ cells triggers MT network destabilization. MTs were detected with an  $\alpha$ -Tubulin antibody, red in (A) and white in (B).

## Baffet\_Fig S3 A

	Hatched eggs	n			
Wild-type	95%	250			
dTBCB <sup>1</sup> /+	96%	260			
α-Tub 84B⁵/+	75%	118			
dTBCB <sup>1</sup> /+ α-Tub 84B <sup>5</sup> /+	39%	123			



F

	Wild-type aPKC localisation	n
Wild-type	100%	35
dTBCB <sup>1</sup> /+	92%	65
α- <i>Tub 84B<sup>5</sup>/</i> +	90%	52
dTBCB <sup>1</sup> /+ α-Tub 84B <sup>5</sup> /+	30%	100

## Fig. S3. Genetic interaction between dTBCB and $\alpha\textsc{-}Tubulin\,84B$

(A) Trans heterozygous flies for  $dTBCB^{1}$  and  $\alpha$ -Tubulin  $84B^{5}$  mutations display a significant reduced fertility compared to wild-type and single heterozygous mutants.

(B-F) Follicle cells from stage 9 egg chambers, immunostained for aPKC. Trans heterozygous follicular cells for  $dTBCB^1$  and  $\alpha$ -Tubulin  $84B^5$  mutations can display a reduced apical aPKC distribution compared to wild-type and single heterozygous mutant cells (B-E). In F, percentages of chambers with localized aPKC are indicated for each genetic background.

Sup Table 1. Quantification of MT phenotype in germline cysts during oogenesis								
	Germarium	St2-St3	St4-St6	St7-St8	St9			
% of fluorescence diminution between <i>dTBCB<sup>1</sup></i> and wild-type germline cysts	43%	42%	43%	42%	53%			
Number of <i>dTBCB<sup>1</sup></i> cysts analyzed	46	15	17	13	4			
Number of wild-type cysts analyzed	53	10	10	14	4			