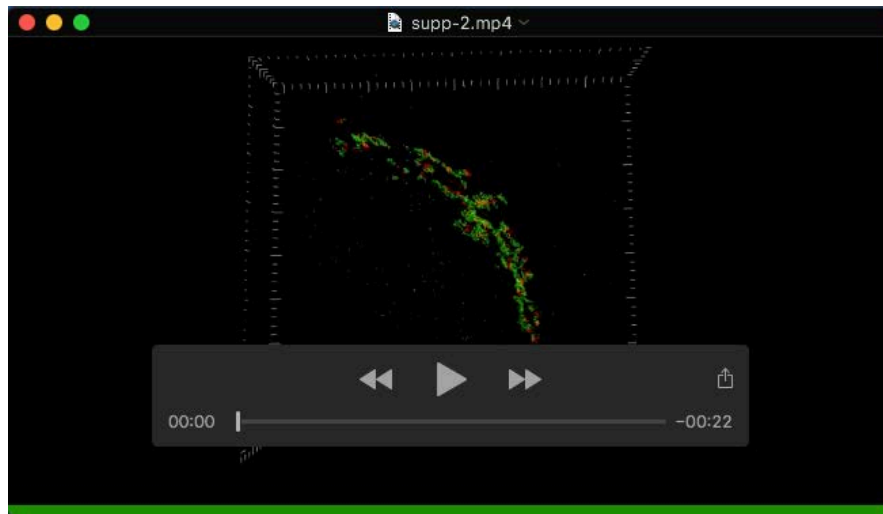


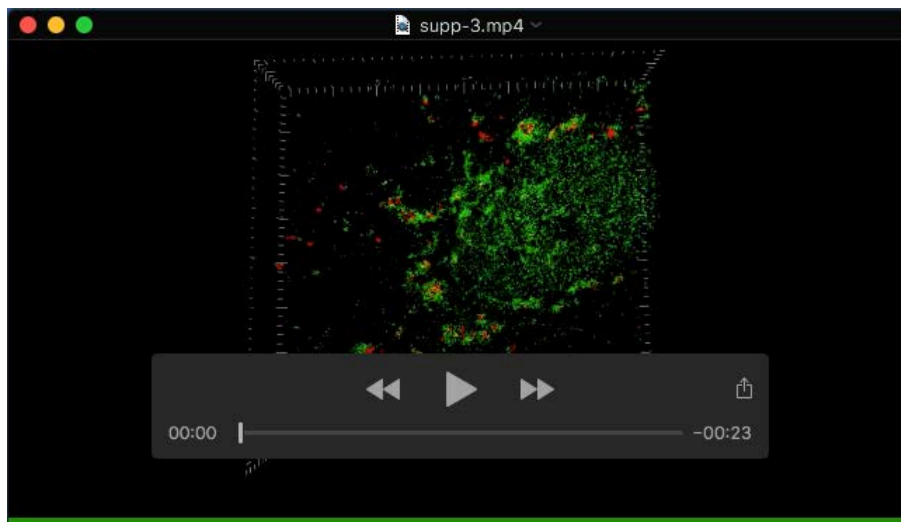
**Table S1**

| <b>Antibody</b>   | <b>SOURCE</b>             | <b>IDENTIFIER</b> | <b>IF</b> | <b>WB</b>        |
|-------------------|---------------------------|-------------------|-----------|------------------|
| SEPT1             | Santa Cruz                | SC-398586         |           | 1:200 /<br>1:500 |
| SEPT1             | This study                | N/A               | 1:200     | 1:500            |
| SEPT2             | Sigma-Aldrich             | HPA018481         | 1:200     | 1:5,000          |
| SEPT6             | This study                | N/A               | 1:50      |                  |
| SEPT6             | Sigma Aldrich             | HPA005665         |           | 1:1,000          |
| SEPT7             | This study                | N/A               | 1:100     |                  |
| SEPT7             | Santa Cruz                | SC-20620          |           | 1:500            |
| SEPT9             | (Diesenberg et al., 2015) | N/A               | 1:400     |                  |
| SEPT9             | abnova                    | H00010801         |           | 1:500            |
| GM130             | BD Bioscience             | 610822            | 1:200     | 1:100            |
| GM130             | Abcam                     | ab52649           | 1:200     |                  |
| $\gamma$ -tubulin | Abcam                     | ab11316           |           | 1:1,000          |
| $\alpha$ -tubulin | Sigma-Aldrich             | T5168             | 1:500     | 1:2,000          |
| EEA1              | BD Biosciences            | 610456            | 1:100     |                  |
| Ergic-53          | Santa Cruz                | sc-398893         | 1:100     |                  |
| COPI              | (Faulstich et al., 1996)  | N/A               | 1:50      |                  |
| API               | BD Bioscience             | 610386            | 1:200     |                  |
| GGA1              | Abnova                    | H00026088-B01     | 1:100     |                  |
| Vps26             | Abcam                     | 23892             | 1:100     |                  |
| CEP170            | Invitrogen                | 41-3220           | 1:100     | 1:100            |
| CEP170            | Abcam                     | ab72505           |           | 1:200            |
| LAMP-1            | BD pharmingen             | 555798            | 1:200     |                  |
| Golgin97          | Invitrogen                | A-21270           | 1:50      |                  |
| TGN46             | Abcam                     | ab50595           | 1:100     |                  |
| GAPDH             | Sigma-Aldrich             | G8795             |           | 1:20,000         |
| Golgin104         | Sigma-Aldrich             | HPA018019         |           | 1:400            |
| Giantin           | Biologend                 | 924302            | 1:400     | 1:500            |
| Talin             | Sigma-Aldrich             | T3287             |           | 1:2,000          |
| Myc               | Santa Cruz                | 9E10              | 1:200     | 1:1,000          |
| GFP               | Clontech                  | 632381            |           | 1:2,500          |



**Movie 1. SIM control GM130green\_golgin97red**

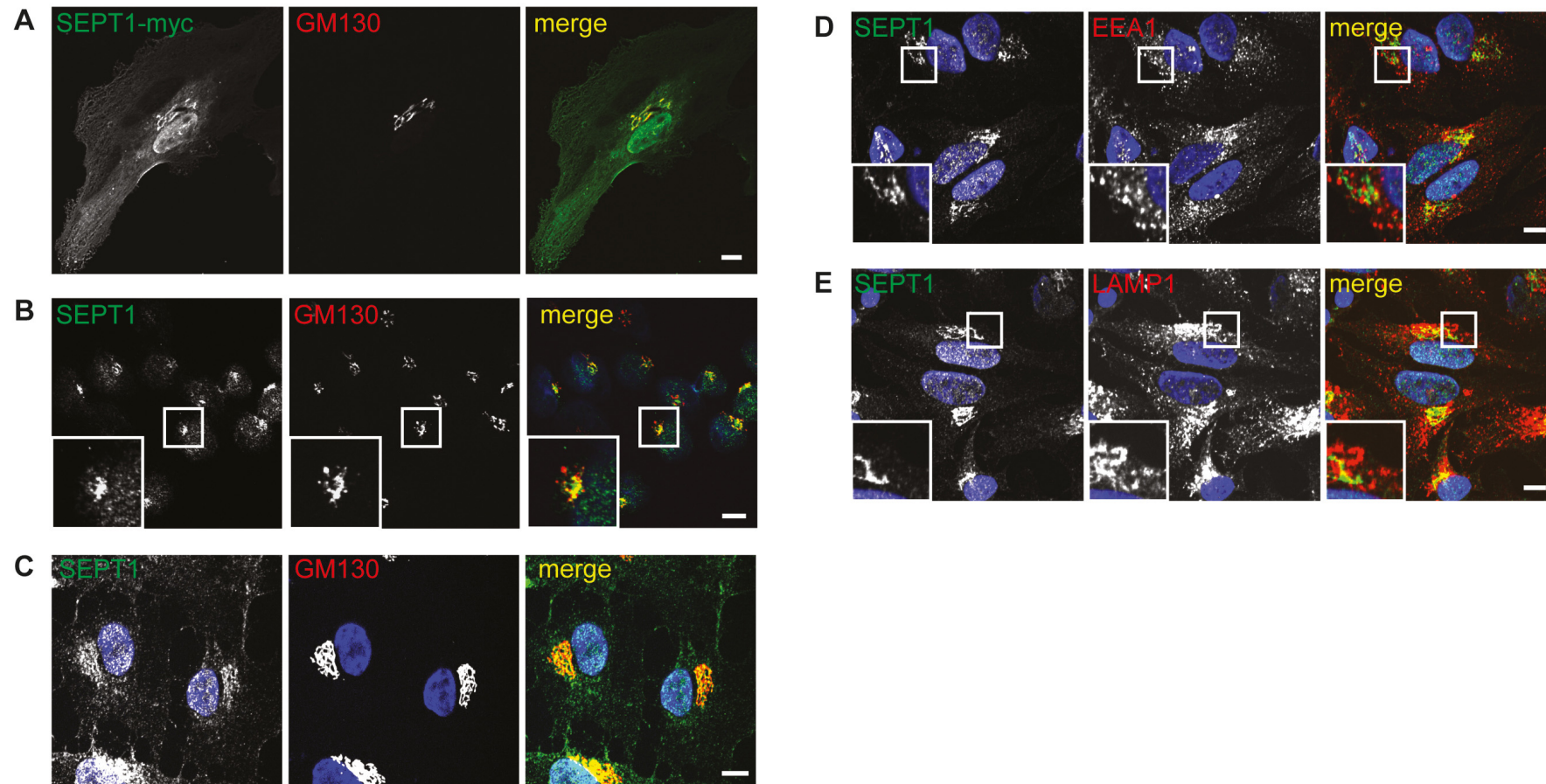
HeLa cells were treated with siCtrl, fixed, immunostained for GM130 (green) and golgin97 (red) and monitored by 3D-SIM microscopy.



**Movie 2. SIM siSEPT1 GM130green\_golgin97red**

HeLa cells were treated with siSEPT1#1, fixed, immunostained for GM130 (green) and golgin97 (red) and monitored by 3D-SIM microscopy.

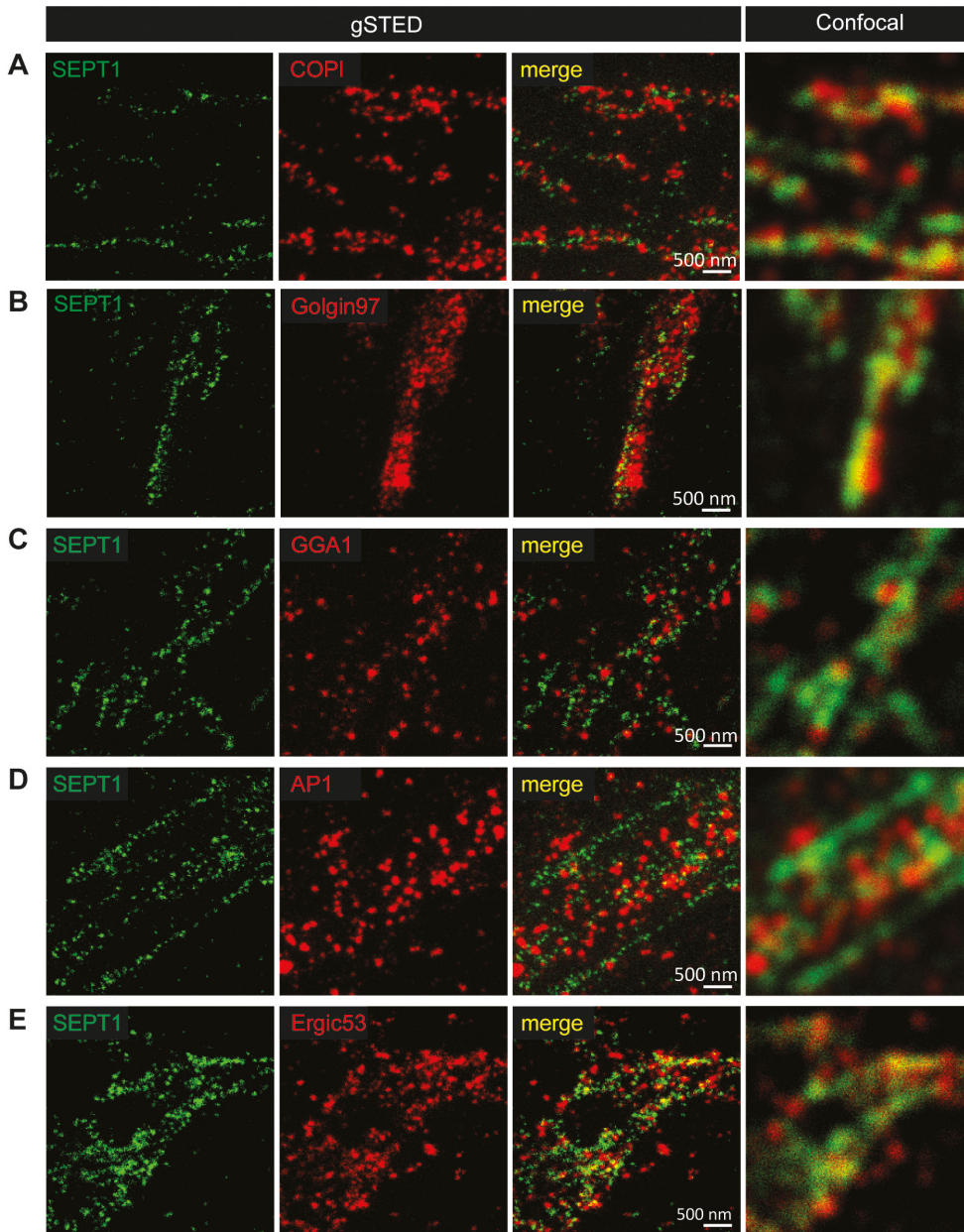
Figure S1



**SEPT1 localizes predominantly to the cis-Golgi.**

- (A) Localization of myc-tagged SEPT1 upon overexpression in HeLa cells, and immunostaining for myc and GM130.
- (B) Immunostaining of endogenous SEPT1 and GM130 in Jurkat cells.
- (C) Immunostaining of endogenous SEPT1 and GM130 in RPE1 cells.
- (D) Staining of SEPT1 and EEA1 (early endosome) in HeLa cells.
- (E) Staining of SEPT1 and LAMP1 (lysosome) in HeLa cells. All scale bars, 10  $\mu$ m.

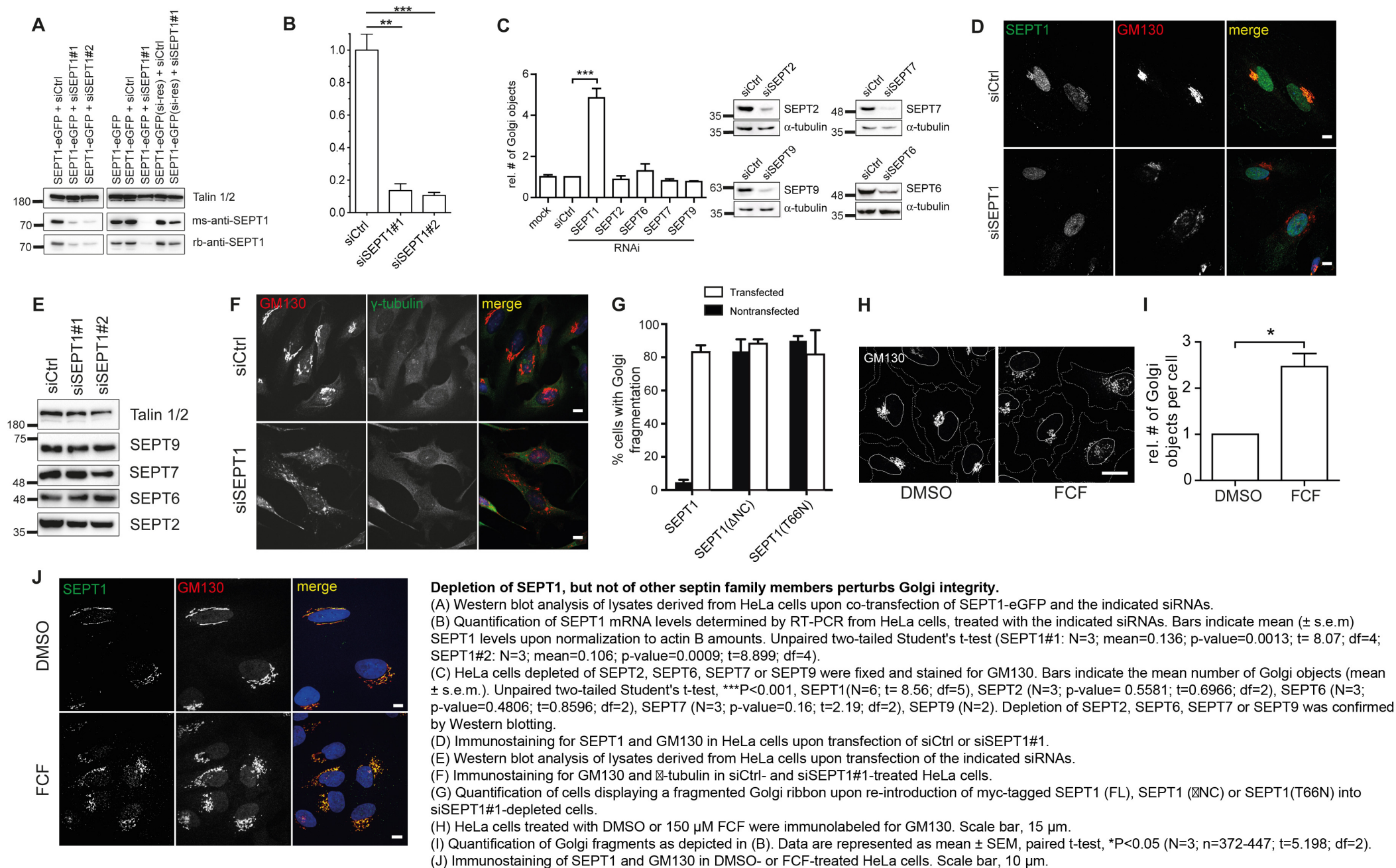
## Figure S2



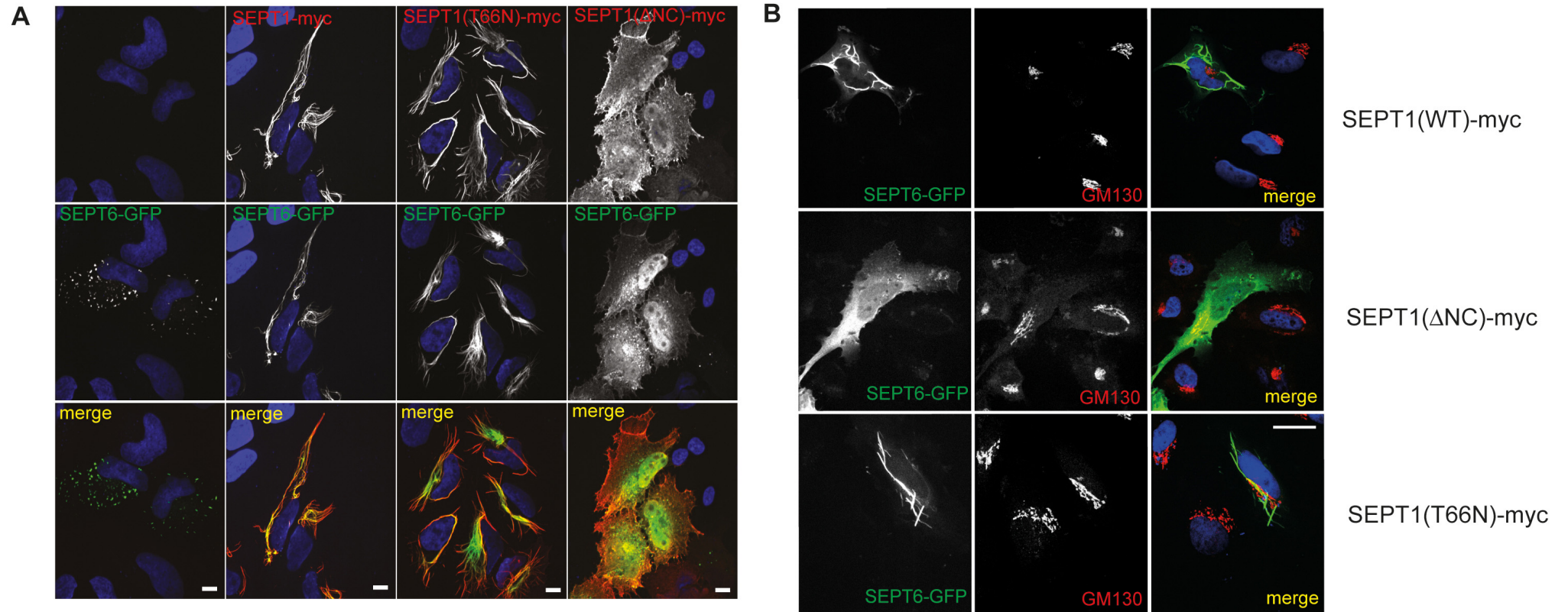
### **gSTED microscopy analysis of SEPT1 at Golgi membranes.**

(A) Analysis of HeLa cells co-immunolabeled with antibodies recognizing SEPT1 and the trafficking adaptor COPI (recruited to the cis-Golgi and to ER-Golgi-Intermediate-Compartment, ERGIC),  
(B) Golgin97 (trans-Golgi),  
(C) GGA1 (trans-Golgi trafficking adaptor),  
(D) AP1 (trans-Golgi adaptor), or  
(E) Ergic53. All scale bars, 500 nm.

Figure S3



## Figure S4

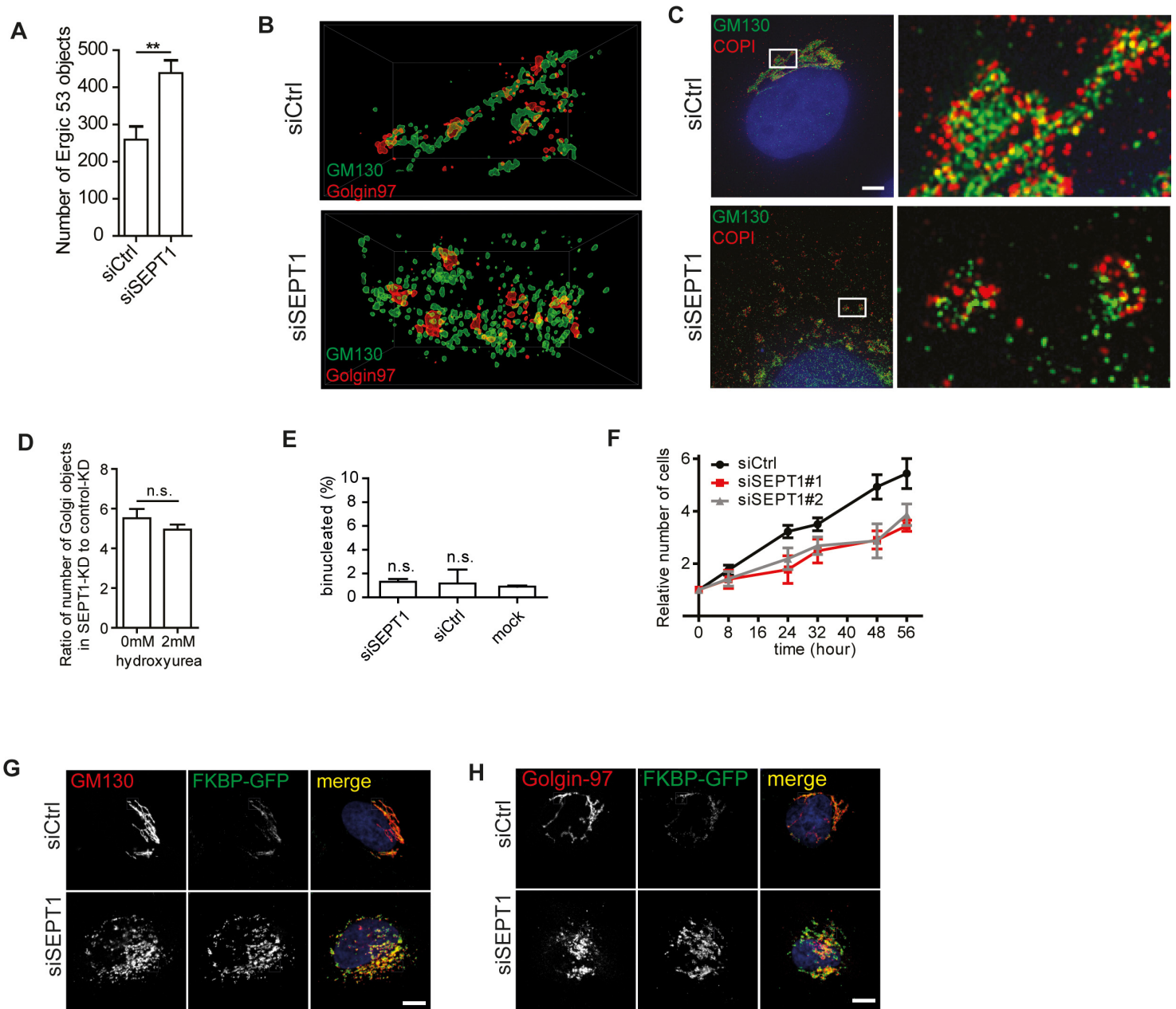


Analysis of SEPT1 variants used for rescue experiments.

(A) SEPT6-eGFP overexpressed alone, or co-expressed with SEPT1-myc (WT) or mutants. Co-expression with SEPT1(WT) or SEPT1(T66N) triggers formation of artificial septin filaments. SEPT1( $\Delta$ NC) cannot form filaments with SEPT6. Scale bar, 10  $\mu$ m.

(B) Immunostaining of GM130 upon co-transfection of HeLa cells with SEPT1-myc (WT,  $\Delta$ NC or T66N) and SEPT6-eGFP.

**Figure S5**



**Analysis of Golgi defects in SEPT1-depleted cells by 3D-SIM, and of the impact of SEPT1 on mitosis, proliferation and secretion.**

(A) Number of Ergic53-positive objects in control or SEPT1-depleted cells, as determined from 3D-SIM images. Data are represented as mean  $\pm$  SEM, unpaired two-tailed Student's t-test,  $**P < 0.01$  ( $n=9$ ;  $t=3.623$ ;  $df=16$ ).

(B) Surface representation of cis- (GM130) and trans- (Golgin97) Golgi objects derived from 3D-SIM images.

(C) 3D-SIM image of GM130 and COPI labeling in HeLa cells after control or SEPT1-depletion.

(D) Relative increase in the number of Golgi objects observed in cells treated with 0 mM or 2 mM hydroxyurea (mean  $\pm$  SEM). SEPT1 depletion increases the number of Golgi objects, and this defect is not cured by administration of hydroxyurea. Unpaired two-tailed Student's t-test,  $*P < 0.05$  ( $N=3$ ;  $n=224-355$ ;  $p\text{-value}=0.344$ ;  $t=1.072$ ;  $df=4$ ).

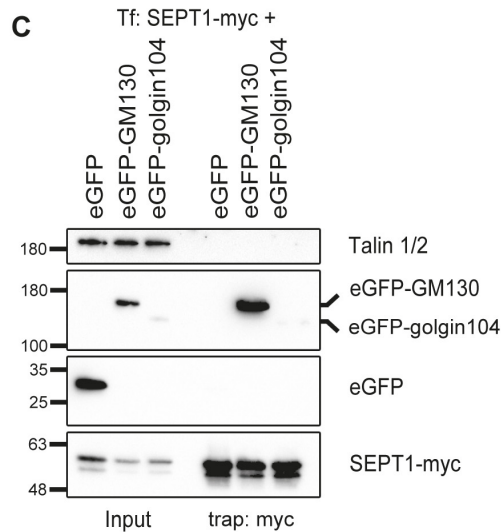
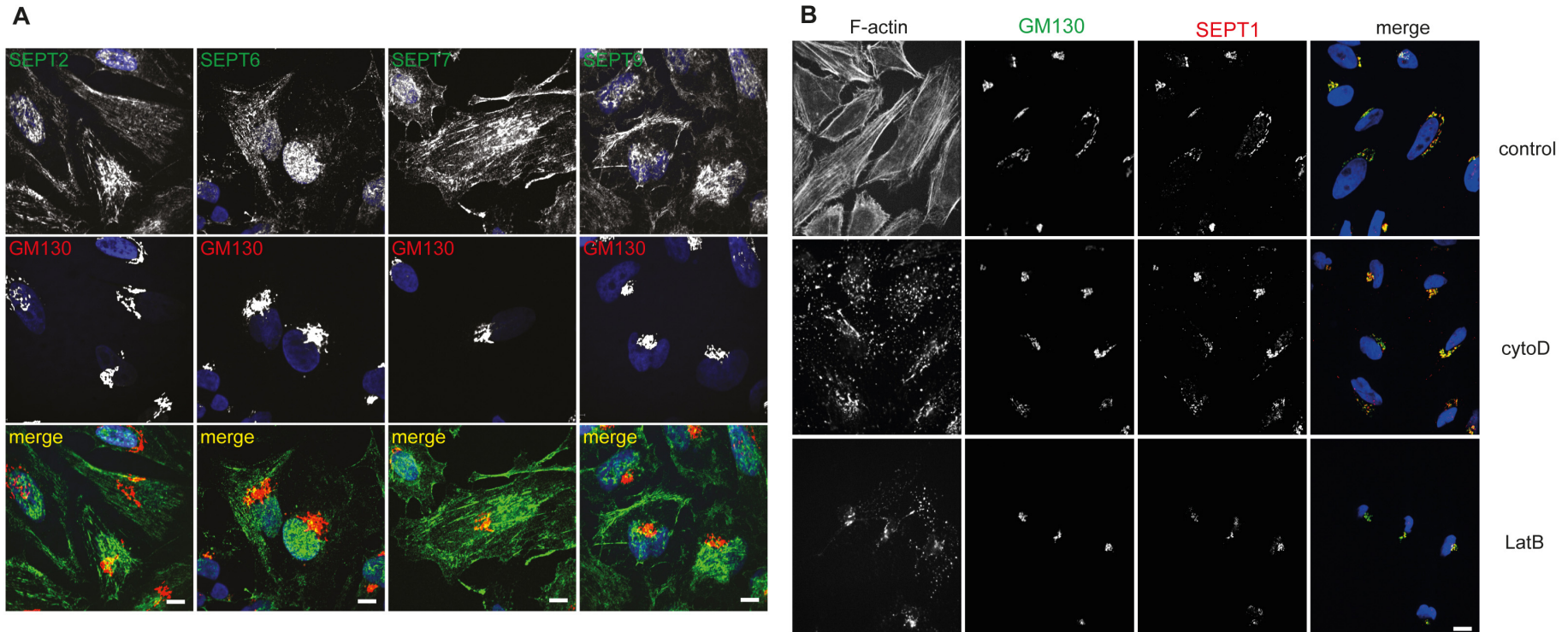
(E) Percentage of binucleated cells in control and SEPT1-depleted cells (mean  $\pm$  SEM), unpaired two-tailed Student's t-test, ns = not significant.

(F) Proliferation of HeLa cells transfected with control, siSEPT1#1 or siSEPT1#2 siRNAs and cells. Data indicate relative numbers of cells at different time points (mean  $\pm$  s.e.m;  $N=3$ ).

(G) Secretion of FKBP(F36M)-GFP reporter was induced for 30 min in control and SEPT1-depleted cells, and fixed cells were then immunostained for GM130 (cis-Golgi). Scale bar, 5  $\mu$ m.

(H) Cells as treated in (G) were immunostained with Golgin97 (trans-Golgi). Scale bar, 5  $\mu$ m.

**Figure S6**



**Localization of highly abundant septin family members, implication of F-actin on SEPT1 recruitment to Golgi membranes, and association of SEPT1 with GM130 and golgin104.**

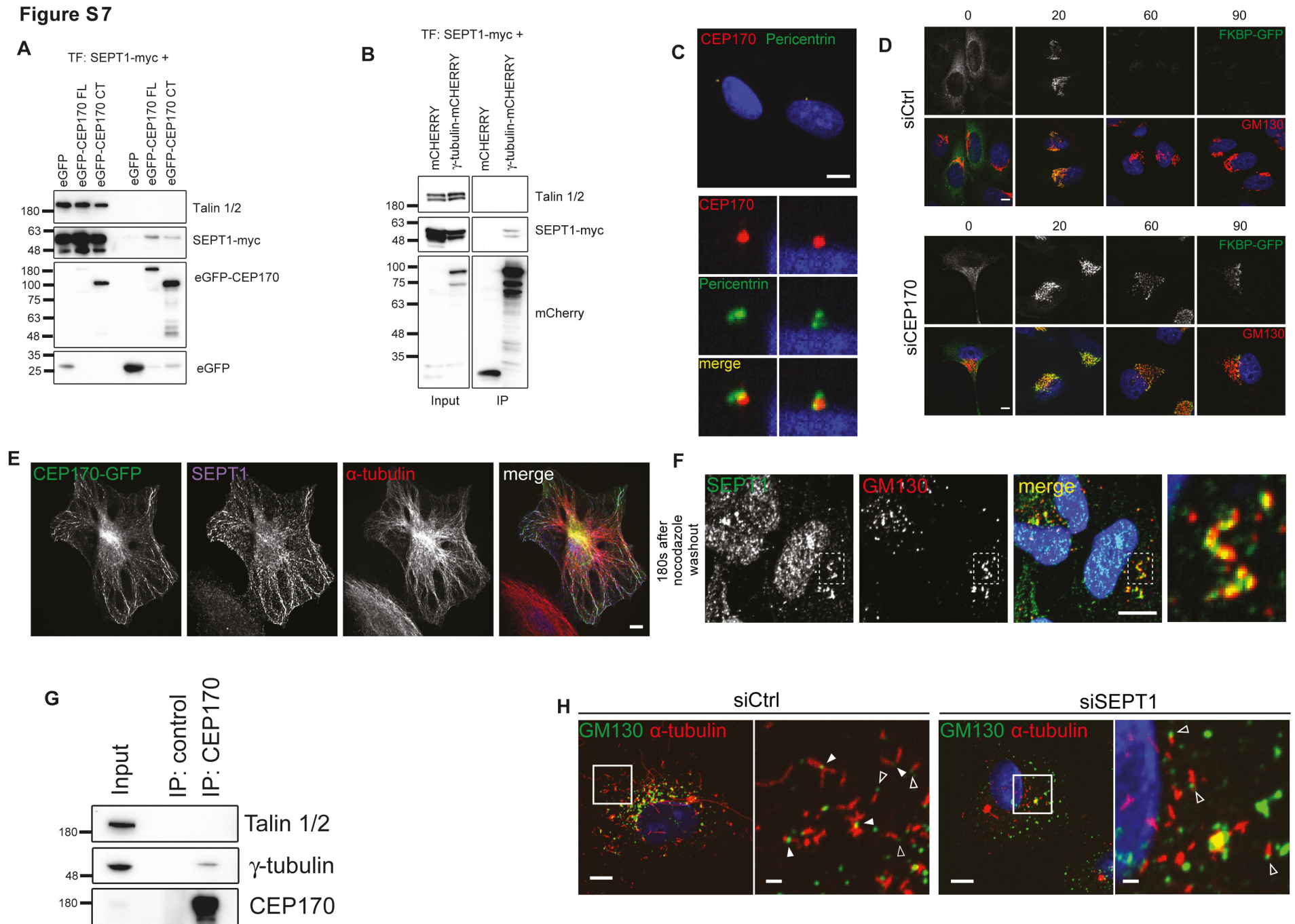
(A) HeLa cells were immunolabeled for GM130 and SEPT2, SEPT6, SEPT7 or SEPT9. None of these septin family members displays a prominent localization at the Golgi.

(B) HeLa cells were incubated with DMSO (control), 2  $\mu$ M cytochalasin D or 2  $\mu$ M latrunculin B for 30 min, fixed in 2% PFA and immunolabeled for F-actin (AlexaFluor568-labeled phalloidin), GM130 and SEPT1. All scale bars, 10  $\mu$ m.

(C) Co-immunoprecipitation of eGFP, eGFP-GM130 or eGFP-golgin104 with myc-tagged SEPT1 from co-transfected Hek293T cells. Samples were probed by Western blotting for myc (SEPT1), eGFP and talin 1/2.



## Figure S7



### SEPT1 associates with CEP170 and $\gamma$ -tubulin to connect cis-Golgi membranes to the microtubule nucleating machinery.

(A) SEPT1-myc co-purifies with eGFP-CEP170 full length, but not eGFP-CEP170 CT from co-transfected Hek293T cells. Samples were probed by Western blotting for myc (SEPT1), eGFP (eGFP), CEP170 (eGFP-CEP170 FL and CT) and talin 1/2.

(B)  $\gamma$ -tubulin-mCherry co-immunoprecipitates SEPT1-myc from transfected Hek293T cells.

(C) CEP170 localizes to the centrosome. Endogenous CEP170 and pericentrin were immunolabeled in methanol-fixed HeLa cells. Note that cold methanol disrupts septin filament integrity (not shown).

(D) Secretion of FKBP(F36M)-GFP reporter was induced for 30 min in control and CEP170-depleted HeLaM C1 cells, and fixed cells were then immunostained for GM130 (cis-Golgi) and monitored by confocal fluorescence microscopy.

(E) Upon overexpression in HeLa cells eGFP-CEP170 co-localizes with SEPT1 on microtubules.

(F) RPE1 cells were treated with nocodazole for 2 h, fixed in 2% PFA 3 min after wash-out, and immunolabeled for GM130 and SEPT1. Enlarged figure inset (right panel) depicts SEPT1 connecting neighboring GM130-positive Golgi objects.

(G) Endogenous  $\gamma$ -tubulin affinity-purifies together with CEP170 from Jurkat cell lysates. Endogenous CEP170 was immunoprecipitated, and the affinity-purified material was analyzed by Western blotting.

(H) Control and SEPT1-depleted RPE1 cells were fixed 90 s after nocodazole washout and stained for  $\alpha$ -tubulin and GM130. Open arrows: single microtubule growing from an individual Golgi object. Closed arrows: grouped microtubules growing from an individual Golgi object. All scale bars, 10  $\mu$ m.