Figure S2

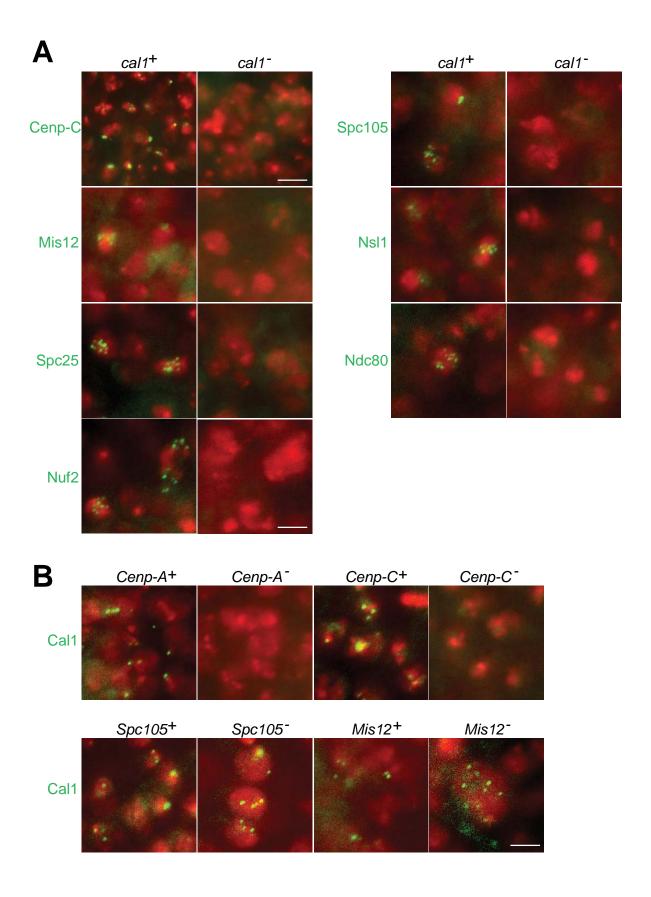


Figure S2. Call acts at the top of the kinetochore assembly pathway.

(A) Localization of EGFP-fusions of Cenp-C, Spc105, Mis12, Ns11, Spc25, Ndc80 and Nuf2 in homozygous *cal1*^{c03646} embryos (*cal1*^r) and in sibling control embryos (*cal1*⁺) within the CNS after germband retraction. Representative mitotic figures are shown with the kinetochore proteins in *green* and DNA staining in *red*. Magnification in the first two Cenp-C panels is indicated by the upper bar = 6 μm; magnification in all other panels by the lower bar = 3 μm.

(B) Localization of Cal1-EGFP in *cid*^{T12-1}/*cid*^{T22-4} (*Cenp-A*⁻), *Cenp-C*^{prl41} (*Cenp-C*), *Spc105*¹ (*Spc105*⁻) and *Mis12*^{f03756} (*Mis12*⁻) mutant embryos as well as in sibling control embryos (*Cenp-A*⁺, *Cenp-C*⁺, *Spc105*⁺ and *Mis12*⁺, respectively). Representative regions with Cal1-EGFP in *green* and DNA staining in *red* are shown at the stage where phenotypic abnormalities start in the mutant embryos, i.e. during mitosis 16 in *Cenp-C* and *Spc105* mutants and during the later mitotic divisions in the CNS in *Mis12* and *Cenp-A/cid* mutants. Bar = 3 μm.