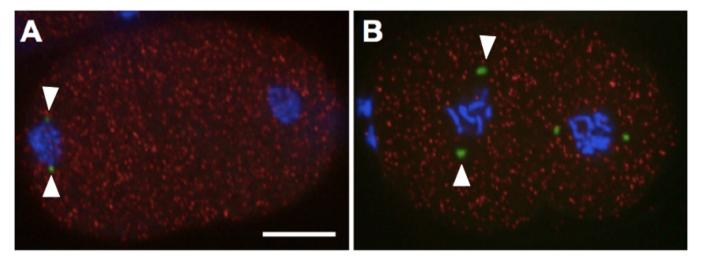


Localization of *tbg-1* mRNAs and GFP::TBG-1 protein in Early *C. elegans* Embryos

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Description

<u>tbg-1</u> encodes gamma-tubulin, a ubiquitous and highly conserved component of centrosomes in eukaryotic cells (Strome *et al*, 2001). Using smFISH we determined the localization of <u>tbg-1</u> transcripts (red). <u>tbg-1</u> transcripts are detected within distinct foci throughout the cytoplasm during both the first (A) and second (B) mitosis. <u>tbg-1</u> transcripts are not enriched at centrosomes or either blastomere. In contrast, GFP tagged <u>TBG-1</u>proteins (green signal; arrowheads) localize at centrosomes, as previously shown (Strome *et al*, 2001). Shown are projections from selected focal planes. Bar=10μm.

New Findings: The first observation of *tbq-1* mRNA localization in early *C. elegans e*mbryos.

Reagents

RNA probes targeting <u>tbg-1</u> mRNAs (Quasar 670; red) were designed using Stellaris Probe Designer (Biosearch Technologies). smFISH was performed as described previously (Osborne-Nishimura et al., 2015; Shaffer et al., 2013). For hybridization, embryos were incubated with <u>tbg-1</u> RNA probes (Quasar 670) at 39°C for four hours in the dark. Following hybridization, the embryos were washed and mounted with DAPI containing (blue) medium. To visualize centrosome-associated <u>TBG-1</u> protein, we used transgenic strain that expresses GFP::<u>TBG-1</u> (TH27; Hannak et al., (2002), green).

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