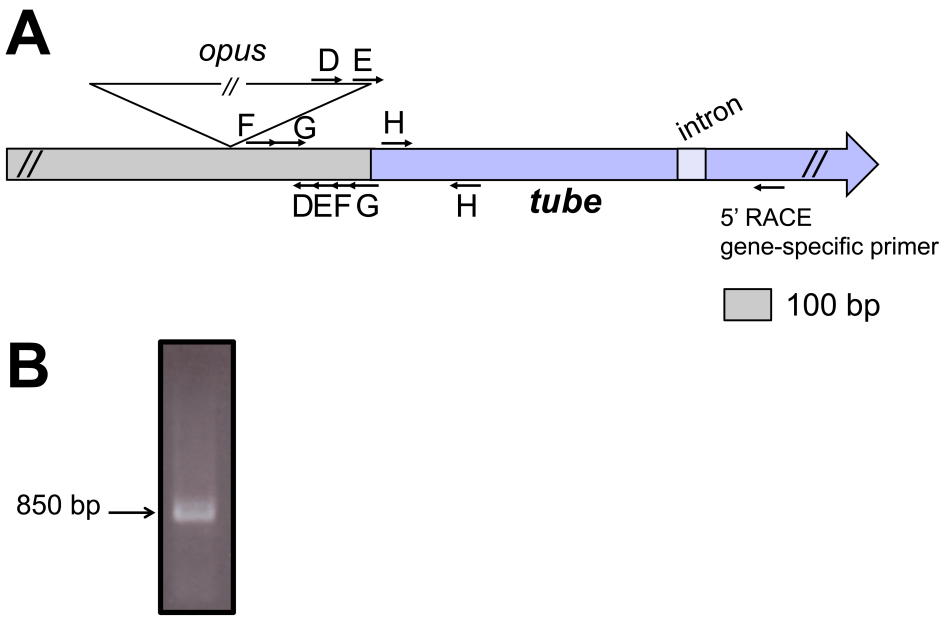

Figure S4. Downregulation of *tube* by *opus* insertion does not depend on an alternative transcription start site.

A. Schematic of primer locations for quantitative RT-PCR and 5' RACE to assay for use of an alternative transcription start site in embryos from tub^{ste}/tub^{null} females. Quantitative RT-PCR primer sets are labeled D-H. Products were detected from primer set H, but not from primer sets D-G.

B. Results from 5' RACE using RNA prepared from embryos from tub^{ste}/tub^{null} females. Expected size of *tube* transcript originating from wild-type transcription start size was 830 bp.