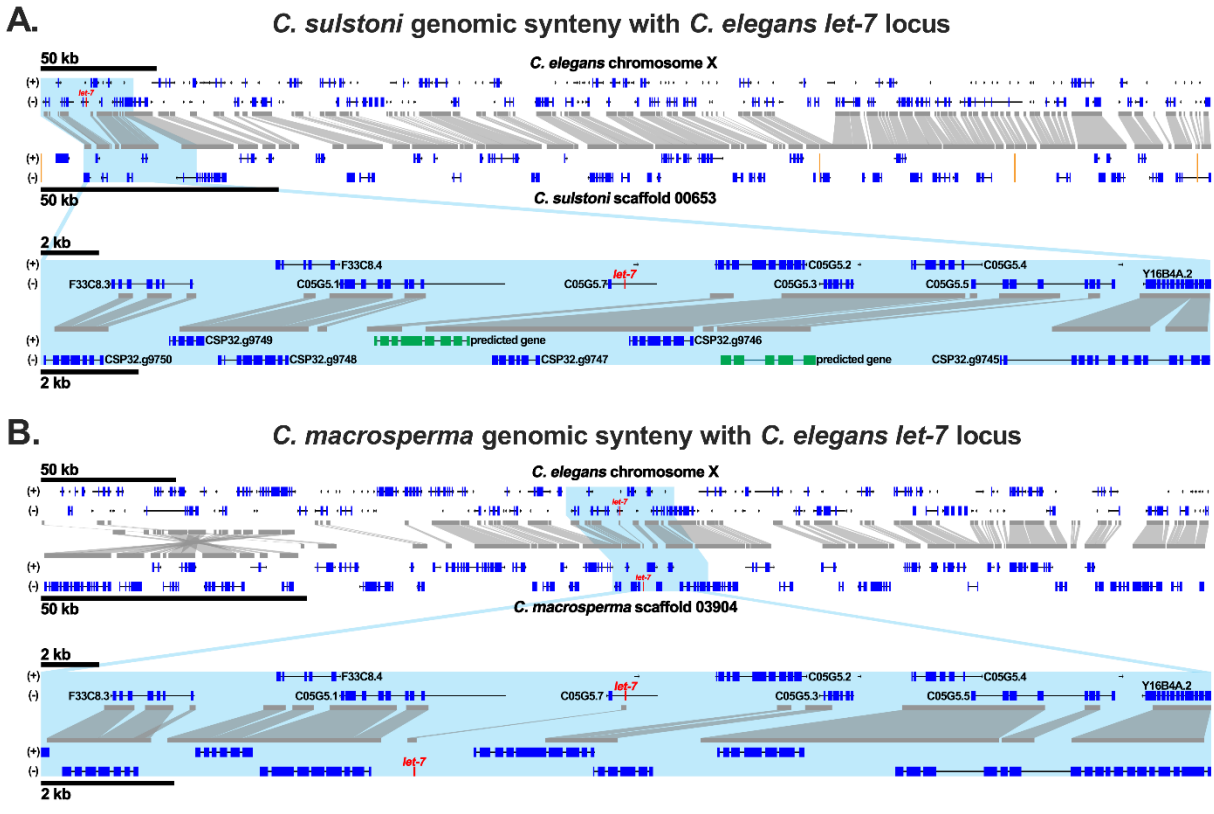

Figure S2 Expanded view: Synteny of a portion of *C. elegans* chromosome X containing the *let-7* sequence with *C. sulstoni* scaffold 00653 (A) and with *C. macrosperma* scaffold 03904 (B). Regions with sequence similarity are outlined in gray. Sequence gaps are shown in orange. *let-7* is shown in red. Annotated genes are shown in blue. Predicted genes are shown in green. Highlighted in light blue are zoomed in views (as shown in Fig. 1) of the *C. elegans let-7* locus and its respective syntenies with *C. sulstoni* (A) and *C. macrosperma* (B).