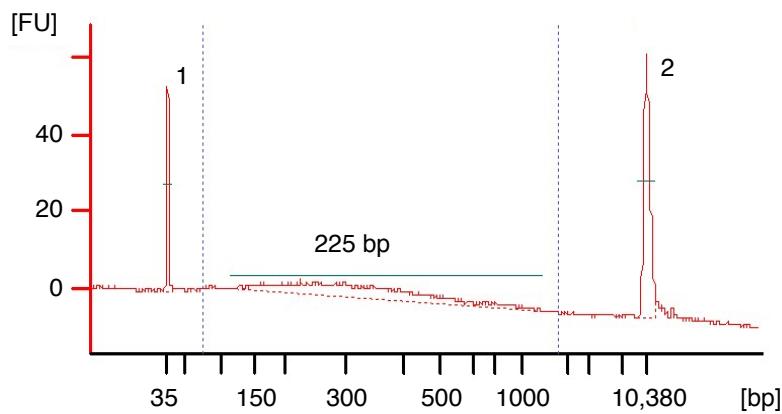
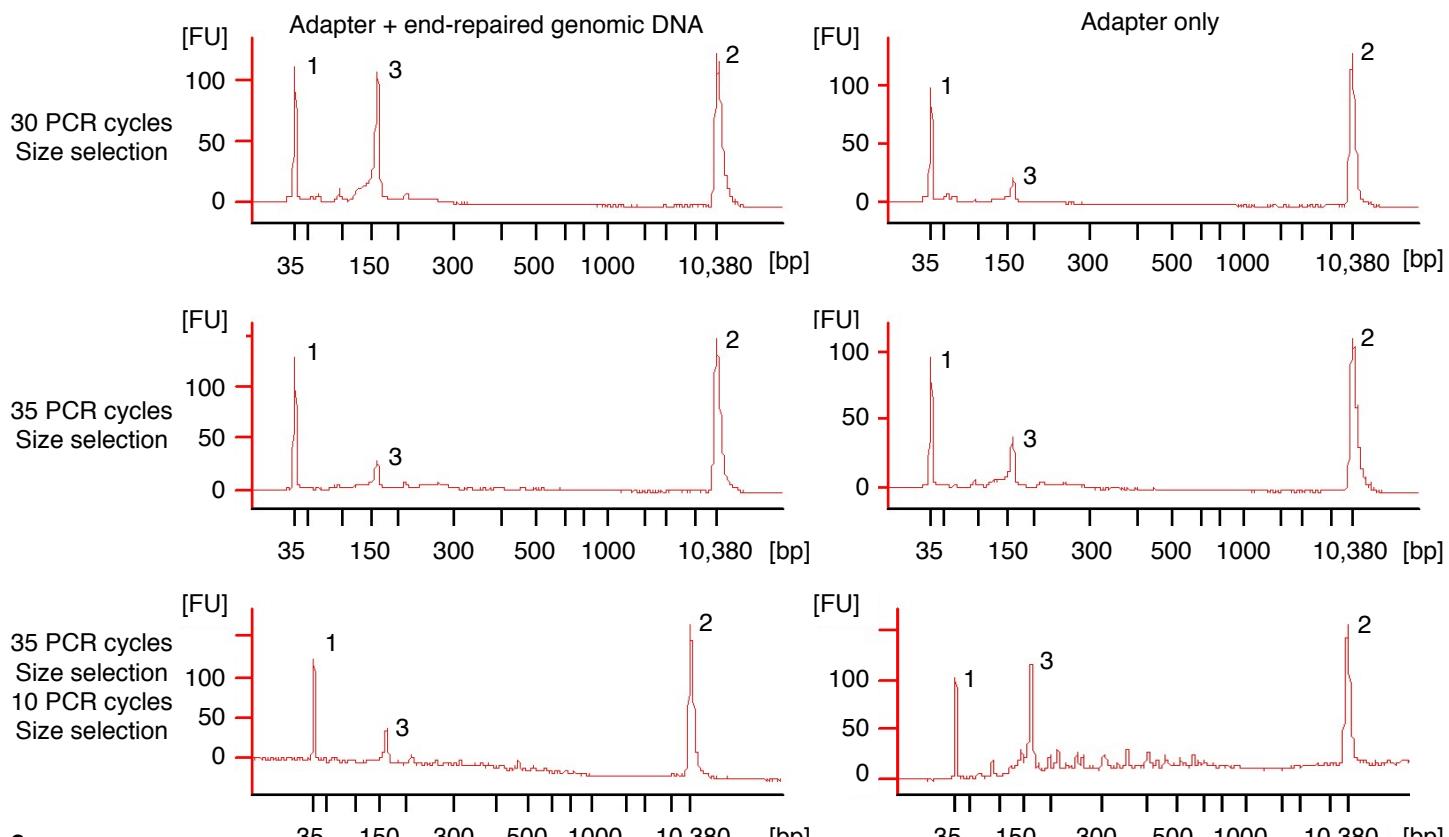
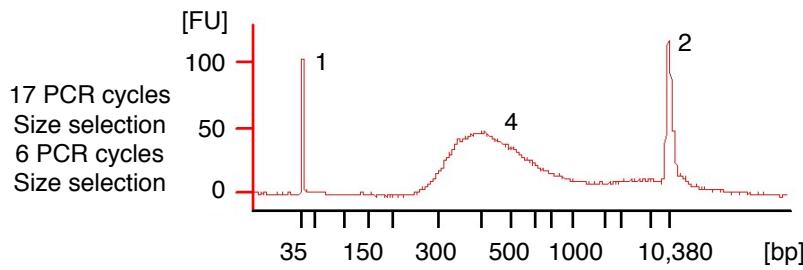


**a****b****c**

#### Supplemental Fig S1. Validation of library preparation.

**(a)** Bioanalyzer profile of sonicated *C. elegans* germ cell DNA (1: 35 bp marker; 2: 10,380 bp marker). **(b)** Bioanalyzer profiles of single MZ libraries prepared following the Schmitt et al. (2012) procedure with dA-tailing instead of dT-tailing (3: peak at expected size of adapter dimers; right column: adapter-only control). **(c)** Profile of a library created following our method. A peak is seen at expected size of adapter - genomic DNA ligation products (peak 4).