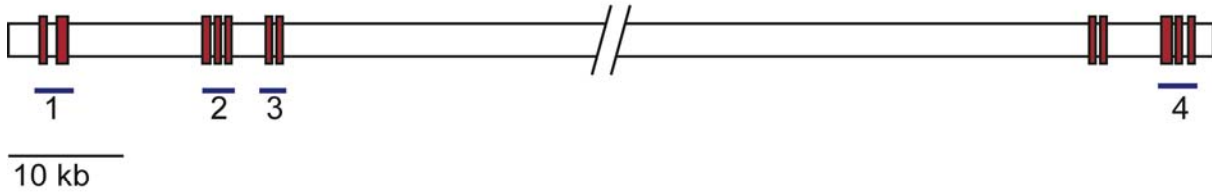
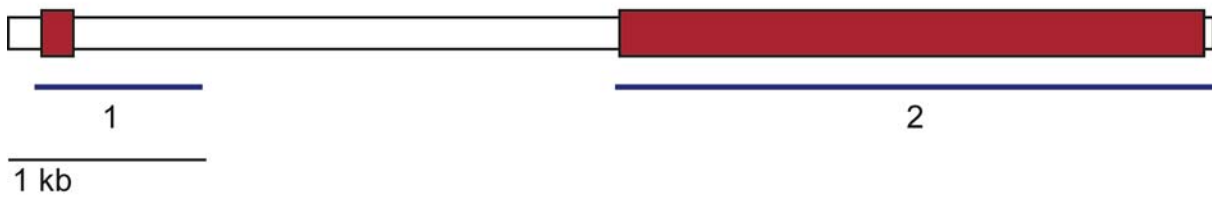


A. *pumilio*



B. *stonewall*



C. *cyclin A*

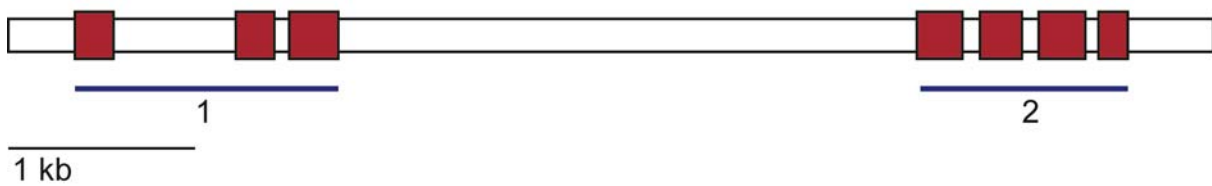


Figure S1 Sequenced fragments of *pumilio*, *stonewall*, and *cyclin A*. (A) *pum* sequencing. The *pum* locus spans approximately 160 kb, so four different regions of *pum*, labeled 1-4, that include 10 of 12 exons in were individually sequenced. The diagram corresponds to the *pum-A* isoform. The center hashes denote where internal sequence (~70 kb) was removed to allow for ease of viewing. (B) *stwl* sequencing. Two fragments of *stwl* were amplified, labeled 1-2. (C) *cycA* sequencing. Two fragments of *cycA* were amplified, labeled 1-2. The blue lines denote the amplified fragments. Red boxes denote exons.