



Figure S2. RT-PCR analysis of the endogenous *unc-60* mRNAs from synchronized L1 worms of N2 (lane 1), *asd-2* (*yb1540*) (lane 2), *asd-2* (*yb1540*); *ybIs1831*; *control* (*RNAi*) (lane 3) and *asd-2* (*yb1540*); *ybIs1831*; *asd-2* (*RNAi*) (lane 4). Splicing patterns of the mRNAs are schematically shown on the right. Triangles indicate positions and directions of the primers.