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.