FIGURE S1.—Sequencing of existing cDNAs reveals multiple *egl-9* mRNA isoforms. Boxes represent exons and lines represent introns. MYND and prolyl hydroxylase domains are filled with black. Nineteen cDNAs from the National Institute of Genetics, Mishima, Japan were sequenced in full, and the number cDNAs of each *egl-9* isoform in that collection is shown in the righthand column. The cDNA sequences have been submitted to www.wormbase.org.