



Figure S5 Alignment of SUPR-1 homologs. The alignment was obtained by the ClustalW multiple-alignment program. Identical and similar residues are highlighted in black and gray, respectively. The *rpm-1(lf)* suppressing allele *ju1118* is shown in red above the sequences. Accession number for *C. elegans* SUPR-1 in UniProt is Q95XT1. Accession numbers for *C. briggsae* and *C. remanei* *supr-1* in Wormbase are CBG04186 and CRE30057, respectively.