



Figure S3. CJD1₍₆₀₋₁₅₁₎ homology model. The model is based on an alignment between CJD1₍₆₀₋₁₅₁₎ and the J domain of the *C. elegans* DnaJ homologue, dnj-2. PyMol software was utilized to create the images. N-termini are displayed in blue while C-termini are colored red.