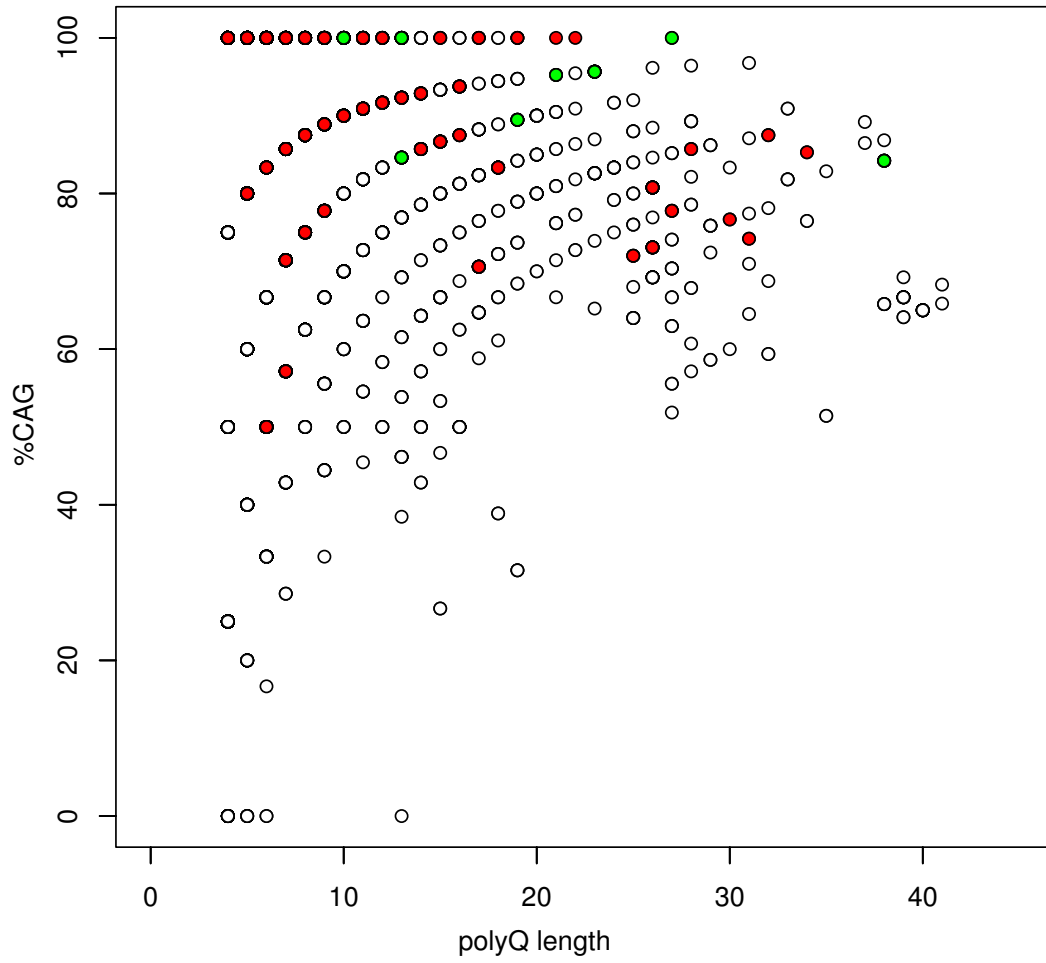


Supp.Fig1. Proteome-wide characterization of %CAG codon usage per polyQ length, in twelve primates (white). Showcased are the values for homorepeats related to polyQ-expansion diseases from human (in green) and nonhuman primates (in red).



Supp.File1. Set of human proteins for which itself or at least one of its orthologs has at least one region with four or more consecutive glutamines.

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Supp.File2. Set of human proteins related to polyQ-associated diseases.

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