Tissue-specific and repeat length-dependent somatic instability of the X-linked dystonia parkinsonism-

associated CCCTCT repeat

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Fig S1. Repeat-related outcomes in individuals exhibiting dystonia or Parkinson's symptoms at onset. a Inverse correlations between CCCTCT repeat length (in blood) with AAO in individuals expressing dystonia (n=194) or parkinsonism (n=43) at onset. Grey shaded areas show 95% confidence interval. b Distribution of repeat length (in blood) in individuals expressing dystonia (n=194) or parkinsonism (n=43). c Distribution of expansion indices (in blood) in individuals expressing dystonia (n=122) or parkinsonism (n=29). Box-whisker plots show median \pm interquartile range (IQR) and red dots show values in individual patient samples. b-c showed no statistical differences between groups.



Fig S2: Modal repeat length variation in brain regions. Of the 23 postmortem samples in which multiple brain regions were analyzed, 4 (17-012, 17-17, 19-017 and 21-031) exhibited variation by one repeat unit in the modal repeat length.



Fig S3: Examples of XDP CCCTCT repeat GeneMapper traces from different tissues. a-f

GeneMapper traces of blood and different brain tissues from individual 17-19. "x" axis shows base pair size and peak height respectively. Indicated on the "x"-axes are assigned repeat lengths based on standard repeat controls.



Fig. S4: Technical variability in expansion index values. Expansion indices were derived from 5 independent XDP CCCTCT repeat PCRs from samples 19-022 (Cereb) and 18-001 (Parietal Cx).



Fig S5: Examples of *LIPG* CCCTCT GeneMapper traces from different tissues.

a-f GeneMapper traces of different brain tissues from individual 17-19. "x" axis shows base pair size and peak height respectively. Indicated on the "x"-axes are assigned repeat lengths based on base pair size.