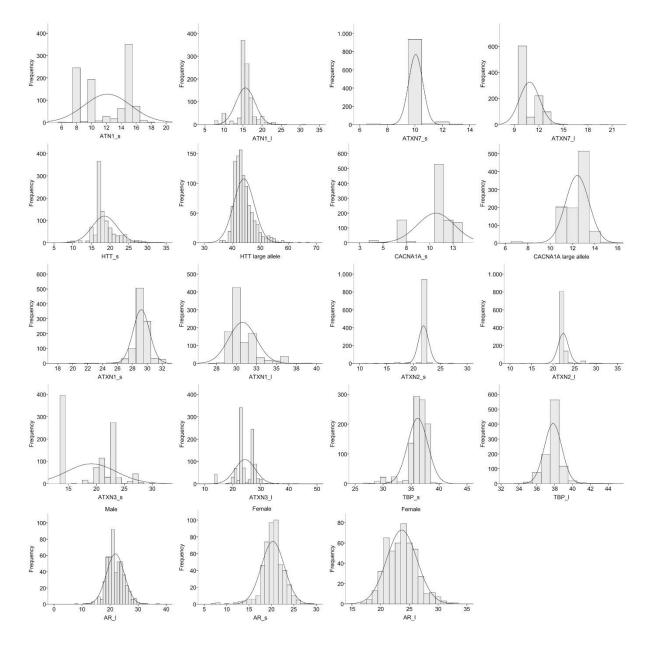
Allele		n	Mean ¹	Median	Mode	Range
HTT	long	996	44.1 ± 3.7	43	43	36-67
	short	998	18.5 ± 3.3	17	17	9-34
ATN1	long	997	15.6 ± 2.5	16	15	8-31
	short	997	12.1 ± 3.1	13	15	6-18
ATXN7	long	995	10.9 ± 1.2	10	10	9-20
	short	995	10.1 ± 0.5	10	10	7-13
CACNA1A	long	996	12.4 ± 1.0	13	13	7-16
	short	996	10.7 ± 2.0	11	11	4-13
ATXN1	long	996	30.8 ± 1.7	30	30	27-39
	short	996	29.2 ± 1.1	29	29	20-32
ATXN2	long	991	22.4 ± 1.2	22	22	13-31
	short	991	21.9 ± 0.9	22	22	11-27
ATXN3	long	991	24.4 ± 3.6	23	14	14-43
	short	991	19.1 ± 4.4	21	23	14-30
AR	long	990	22.8 ± 3.0	23	21	8-37
	short	990	21.2 ± 3.0	21	21	7-37
TBP	long	995	37.8 ± 1.0	38	38	34-44
	short	995	36.2 ± 1.8	36	36	27-40

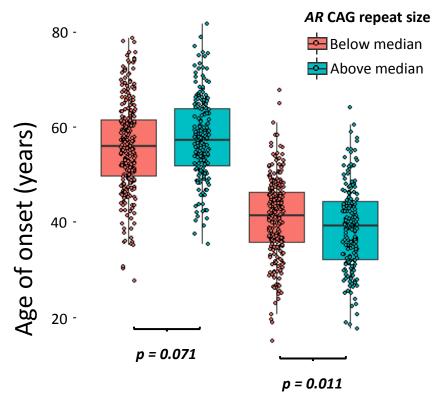
Supplementary Table 1: Distributions of the CAG repeat tracts in the PDAGs.

PDAG = polyglutamine disease-associated gene

Supplementary Figure 1: Distribution patterns of CAG repeat tracts in various polyglutamine diseaseassociated genes. Bars represent the frequency of a particular CAG repeat size in either the shorter (s) or the longer (l) allele of each gene. Curves represent the hypothetical normal distribution. For the *AR* gene histograms were produced for men and woman separately.



Supplementary Figure 2: Interaction between *AR* and mutant *HTT* **CAG repeat size.** Longer *AR* CAG repeat size tended to delay age of onset in HD patients with a below median number of CAG repeats in the expanded *HTT* allele (Mann-Whitney *U*-test p=0.071), while for patients with an above median expansion longer *AR* CAG repeats tended to advance age of onset (Mann-Whitney *U*-test p=0.011). Black horizontal lines represent medians, boxes display interquartile ranges and whiskers are 1.5 × interquartile range. Circles represent individual patient data with horizontally added jitter.

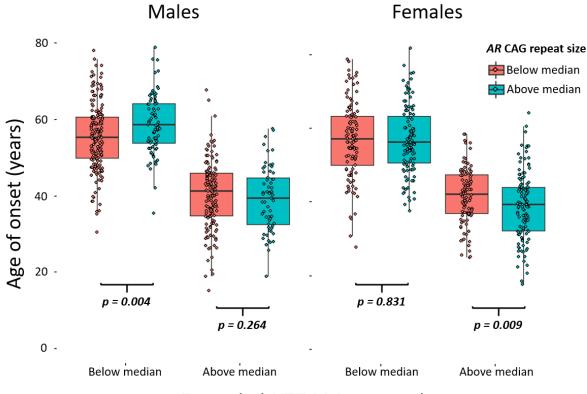


0 -

Below median Above median

Expanded HTT CAG repeat size

Supplementary Figure 3: Interaction between *AR* and mutant *HTT* CAG repeat size is sex-specific. In males a longer *AR* CAG repeat size delayed age of onset in subjects with a relatively low expanded *HTT* CAG repeat size (Mann-Whitney *U*-test p=0.004), while in females a longer AR allele resulted in an earlier age of onset in subjects with a relatively larger expanded HTT CAG repeat size (Mann-Whitney *U*-test p=0.004). Black horizontal lines represent medians, boxes display interquartile ranges and whiskers are 1.5 × interquartile range. Circles represent individual patient data with horizontally added jitter.



Expanded HTT CAG repeat size