Subject	Race (Gender)	Brain area	Age (years)	HD Vonsattel grade	Other pathology	Cause of death	PMI
Non-HD-1	Caucasian (F)	Frontal cortex	79	n.a.	Depression	Drug overdose	14 h
Non-HD-2	Caucasian (F)	Frontal cortex	77	n.a.	Lung disease/HBP/	ASCVD	8 h
					Arthritis/Breast cancer		
HD-1	Caucasian (F)	Frontal cortex	82	n.d.	Senile cerebral disease	Complications	6 h
HD-2	Caucasian (F)	Frontal cortex	78	III	Senile cerebral disease	Complications	2 h
Non-HD-3	Caucasian (M)	Caudate nucleus	58	n.a.	n.d.	n.d.	9 h
Non-HD-4	Caucasian (F)	Caudate nucleus	53	n.a.	HBP/Asthma	HASCVD	15 h
HD-3	Caucasian (M)	Caudate nucleus	43	n.d.	n.d.	Complications	10 h
HD-4	Caucasian (M)	Substantia nigra	58	n.d.	n.d.	ASCVD	17 h

Supplementary Table 1. Summary of demographic data, neuropathology and experimental results of human subjects

F, female; M, male; n.a., not applicable; n.d., not determined; PMI, post-mortem interval; HBP, high blood pressure; ASCVD, arteriosclerotic cardiovascular disease; HASCVD, hypertensive arteriosclerotic cardiovascular disease.

Supplementary Table 2. Relative mRNA levels of human and mouse HTT in brain cortex and the isolated brain capillaries (BCs) of R6/2 HD mice, the wild-type (WT) controls, and C57BL/6 (B6) mice.

	R6/2	WT	B6		
Mouse HTT					
BCs (n=2-3)	4.81±0.53	2.21±0.36	2.81±0.87		
Cortex (n=6-11)	6.40±0.69	6.56±0.50	6.42±0.88		
Human HTT					
BCs (n=2-3)	8.66±1.16	ND	ND		
Cortex (n=6)	4.32±0.60	ND	ND		

Methods for Supplementary Table 2. Brain capillaries were isolated from male and female R6/2 and wild-type (WT) mice of 12 weeks old or B6 mice of 6 weeks old (5-6 animals in each preparation) by the methods described in the article. The cDNA (1 μ L) was mixed with 7 μ L of DEPC-treated sterile deionized distilled water, 10 μ L of Power SYBR Green PCR Master Mix (Applied Biosystems, Warrington, UK) and 0.5 μ M forward and 0.5 μ M reverse primers: mouse HTT forward 5'-CTCAGAAGTGCAGGCCTTACCT-3' or reverse 5'-GATTCCTCCGGTCTTTTGCTT-3'; human HTT forward 5'-GCCGCTCAGGTTCTGCTTT-3' or reverse 5'-AAGGCCTTCATCAGCTTTTCC-3'. Quantitative RT-PCR was conducted as described in the text. Relative expression was calculated by $2^{-\Delta Ct} \times 10^4$. ΔCt is the relative quantity of Htt mRNA normalized to Gapdh mRNA. Data are given as mean±SEM of several independent preparations as indicated in the table. For BCs, each preparation was from 5-6 animals. ND, not detectable.



Supplementary Figure 1. Orthogonal views of the same imaging fields of the cortex (A) and striatum (B) shown in Figure 1. Images were taken from the z-section plane indicated by the blue line. The nuclear distribution of the p65 subunit of NF- κ B is identified using immunostaining of p65 (green), CD31 (red) positive brain capillary endothelial cells, and nuclei (blue). The images were acquired at 40× magnification on an LSM 880 confocal microscope (Zeiss, Jena, Germany). Scale bars indicate 50 μ m.



Supplementary Figure 2. mRNA levels of P-gp, Mrp2, and Bcrp in the cerebral cortex (A), liver (B), jejunum (C), and kidney (D) of male R6/2 HD mice at 12 weeks of age (white bars), compared to age-matched wild-type (WT) controls (black bars). Relative expression was calculated by $2^{-\Delta\Delta Ct}$. $\Delta\Delta Ct$ value was obtained by subtracting the mean ΔCt value of WT controls from the individual ΔCt value, the relative quantity of P-gp, Mrp2, or Bcrp mRNA normalized to Gapdh mRNA. The data are given as the mean±SEM of five to six animals. A fold change greater than 1 indicates up-regulation and a fold change less than 1 indicates down-regulation. (*, P < 0.05; **, P < 0.01 compared to WT mice).



Supplementary Figure 3. Western blotting and the quantification results of P-gp in the membrane fractions isolated from the whole brains of the WT (B6CBAFI/J) mice 2 hours after an intravenous injection of vehicle (2.5% aqueous dextrose solution; -TQD group) or 6 mg/kg tariquidar (TQD) (+TQD group). Tariquidar was freshly prepared in 2.5% aqueous dextrose solution. The quantitative results are presented as the mean±SEM of three mice.