

Supplemental Information

Cytosolic carboxypeptidase 1 is involved in processing of α -tubulin and β -tubulin

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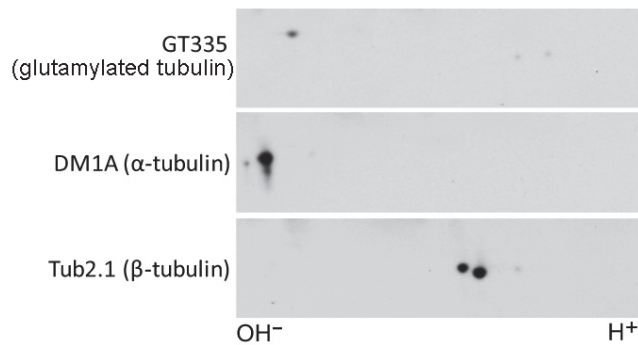
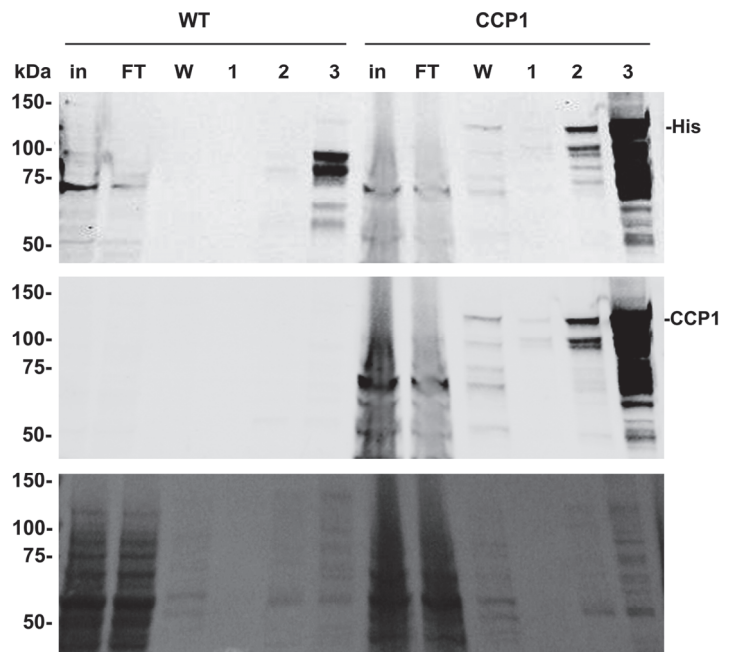


Figure S1. Levels of polyE tubulins in HEK293T cells. Cell lysates of HEK293T cells were subjected to high-resolution 2DE. There are almost no poly-glutamylated tubulins.

Figure S2. Purification of recombinant CCP1. Immunoblot analysis of fractions obtained during purification. Sf9 cells were infected with WT or His-tagged CCP1 recombinant baculovirus. Cell extracts (input, in) were loaded into the Talon column and non-bound fraction (flow-through, FT), wash (W), elute with 10 mM imidazole (1), elute with 20 mM imidazole (2), and elute with 80 mM imidazole (3) were collected. Aliquots of fractions (0.1% of in and FT, 1% of other fractions) were separated by SDS-PAGE and either probed with tetra-His antibody (upper panel), CCP1 antisera (middle panel) or stained with ponceau S solution (lower panel). The theoretical MW of CCP1 is 130 kDa.



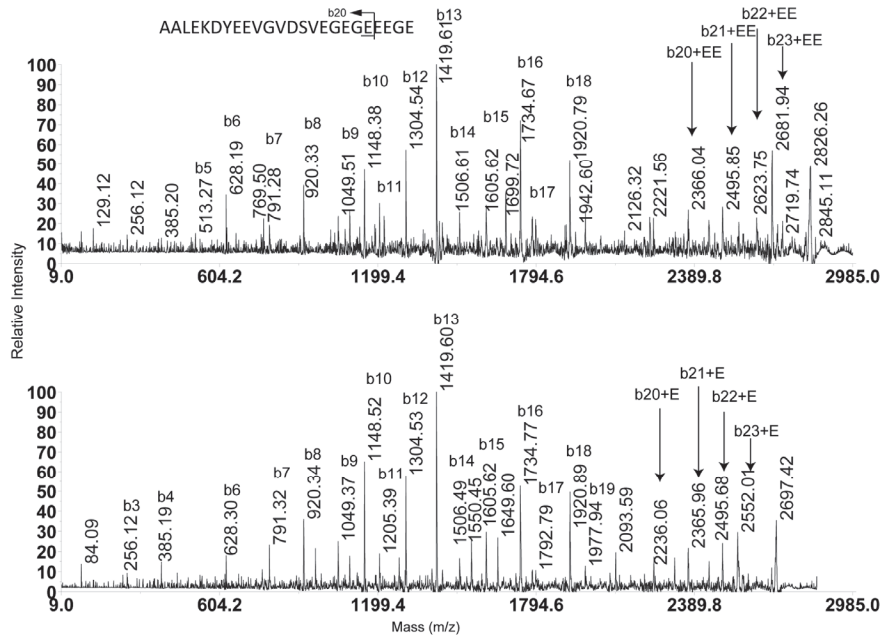


Figure S3A. Fragmentation profiles of the C-terminal representative peptides from α -tubulin. (Top) Positive MALDI TOF-TOF MS/MS spectrum of the precursor 2696.2 of $\alpha 1a/\alpha 1b/\alpha 3$. (Bottom) Positive MALDI TOF-TOF MS/MS spectrum of the precursor 2825.3 of $\alpha 1a/\alpha 1b/\alpha 3$. Each of the sequence is shown in the figures. Mainly b ions are labelled in the mass spectra. The polyglutamylated site of both peptides is underlined.

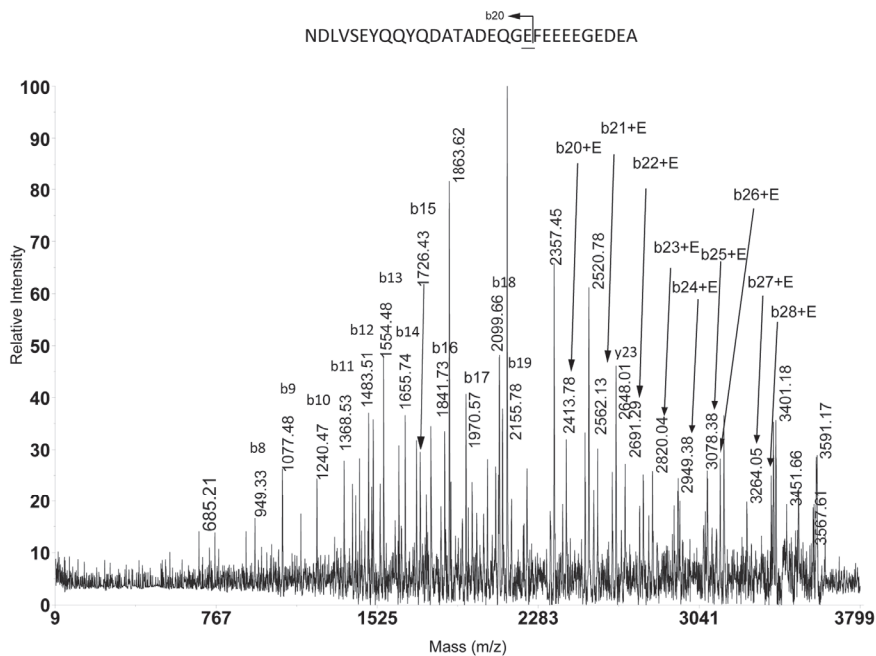


Figure S3B. Fragmentation profiles of the C-terminal representative peptides from β -tubulin. Positive MALDI TOF-TOF MS/MS spectrum of the precursor 3595.8 of $\beta 2b$. Each of the sequence is shown in the figures. Mainly b ions are labelled in the mass spectra. The polyglutamylated site of both peptides is underlined.

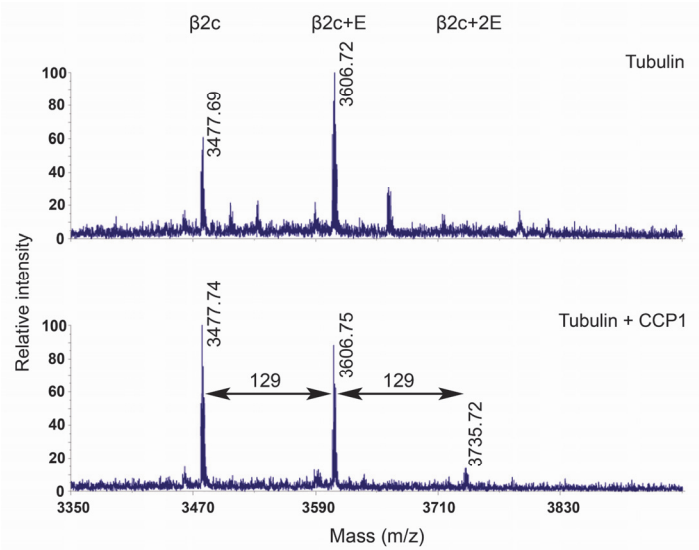


Figure S4. β 2c tubulin isoform processing by purified CCP1. Representative spectra of β 2c form of porcine β -tubulin. After incubation, tubulin was isolated on polyacrylamide gel, digested with CNBr, and analyzed as described. The form shown is the C-terminal region of β 2c. **Top panel:** spectrum of β 2c brain tubulin alone. **Bottom panel:** spectrum of β 2c brain tubulin after incubation with purified CCP1. 129 is the molecular weight of Glu residue in Da.

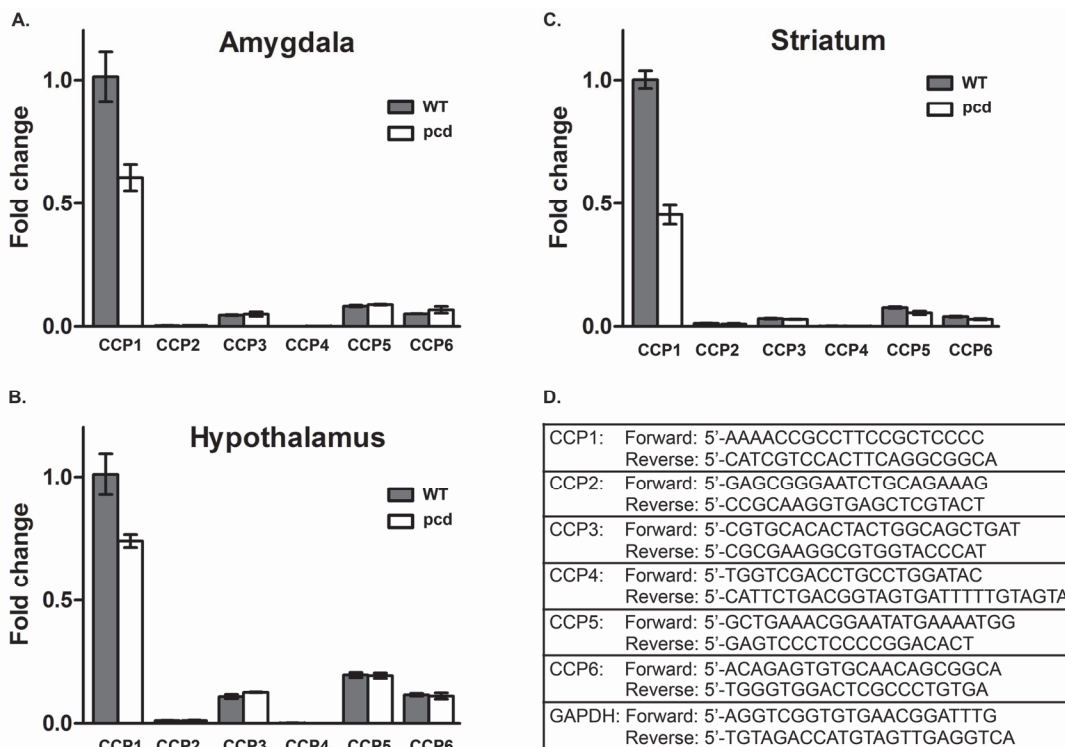


Figure S5. Relative levels of CCP mRNAs in the brain of wild-type and *pcd* mice. Quantitative real-time PCR was performed to determine mRNA levels of CCP1 through 6 in amygdala, hypothalamus, and striatum (panels A, B, and C, respectively) of adult wild-type and mutant mice. CCP1 mRNA is the most abundant among other CCP mRNAs in wild-type mice. The levels of CCP1 mRNA are lower in *pcd* mice, but the levels of other CCP transcripts are not affected by CCP1 mutation. **D:** Mouse primers used for quantitative real-time PCR. Fold change in expression was calculated using the DDCT method. GAPDH was used as an internal control. n = 4 (wild-type), n = 2 (*pcd*).

Table S1. Primers used for Quantitative real-time PCR of human cell lines.

CCP1	Forward: 5'-GCAGTGAAGCGTTTACCCT Reverse: 5'-GCTGGGGCGATATGGCTC
CCP2	Forward: 5'-GGGAGGCAAACGAGGAAT Reverse: 5'-CTGCCCATCATCCGTGT
CCP3	Forward: 5'-GCTAGGGAGATGGGTGCC Reverse: 5'-AGGGGTTCCGTTTCTAAGCC
CCP4	Forward: 5'-GATATTCGGAGGCTCATCCA Reverse: 5'-CTGCATACCGCTCACTTTGA
CCP5	Forward: 5'-GAGCACAGCAGCCTTACT Reverse: 5'-GCAACCCATTGTGACTTT
CCP6	Forward: 5'-TTCCTATTGCTACACCCT Reverse: 5'-CACCTTTTCAACCACG
GAPDH	Forward: 5'-GAAGGTGAAGGTCGGAGT Reverse: 5'-GAAGATGGTGATGGGATTTC

Movie S1. Athletic performance of *pcd/ΔTtll1* mouse.

The *pcd/ΔTtll1* mouse was made to grasp a cord with the body on the downside of the cord. The mouse climbed onto the cord within a few seconds.

Movie S2. Athletic performance of *pcd* mouse.

The *pcd* mouse was made to grasp a cord with the body on the downside of the cord. The mouse failed to climb onto the cord within the allotted time of 1 minute.

Movie S3. Comparison of athletic performance of *pcd* mouse with *pcd/ΔTtll1* and wild-type mice.

The movie shows the severe ataxia of *pcd* mouse compared with *pcd/ΔTtll1* and wild-type mice.