

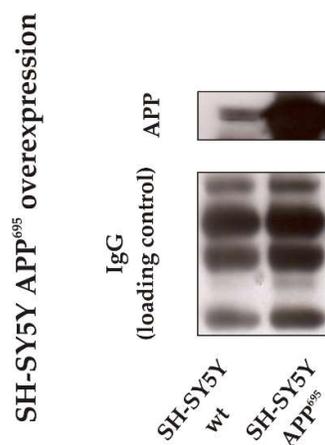
## Supplemental material

Table 1a. Cytotoxicity by LDH release

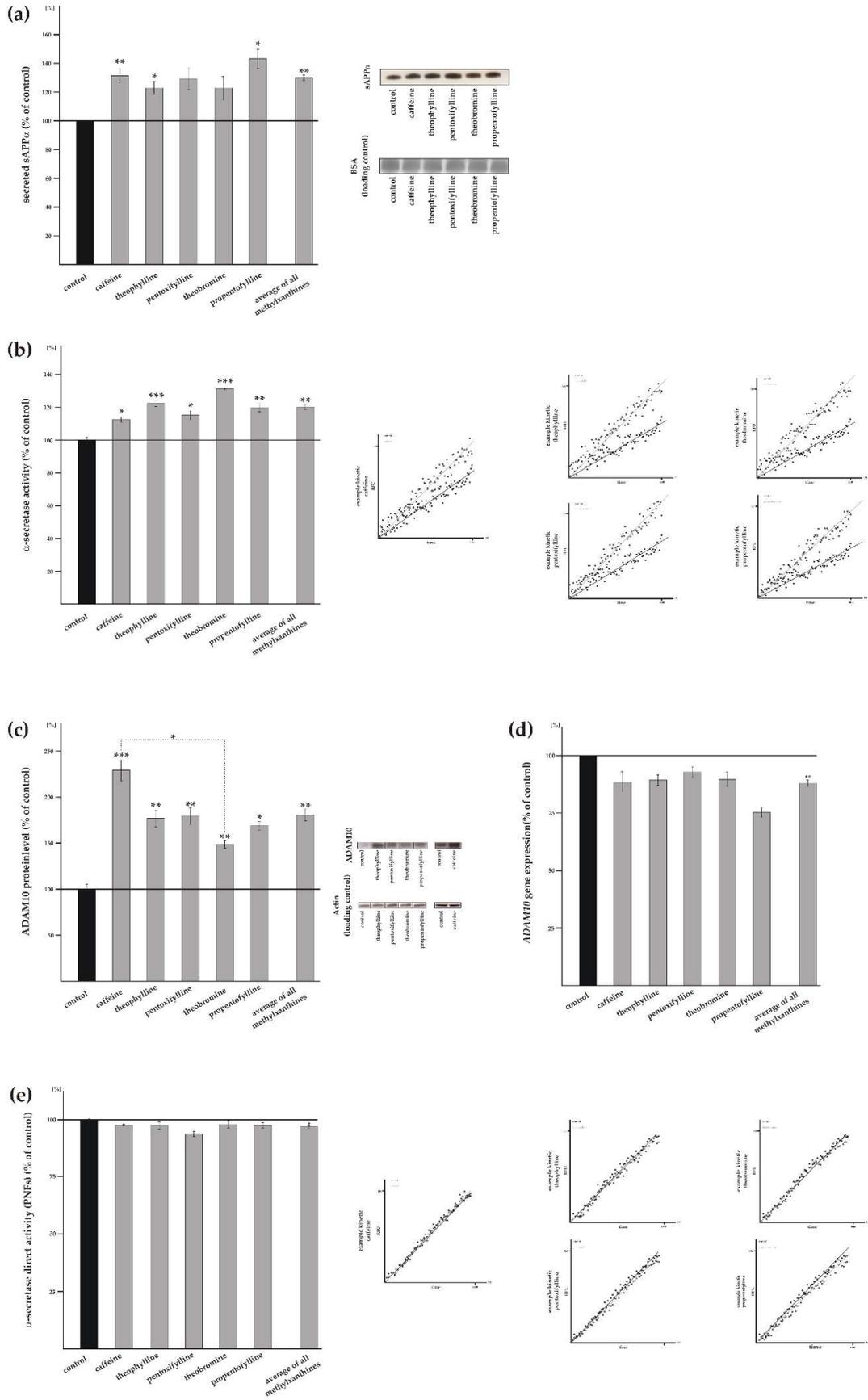
MTX	mean difference to control [%]
caffeine	-0.54
theophylline	-0.28
pentoxifylline	-0.06
theobromine	1.4
propentofylline	3.78

Table 1b. Cell proliferation by XTT release

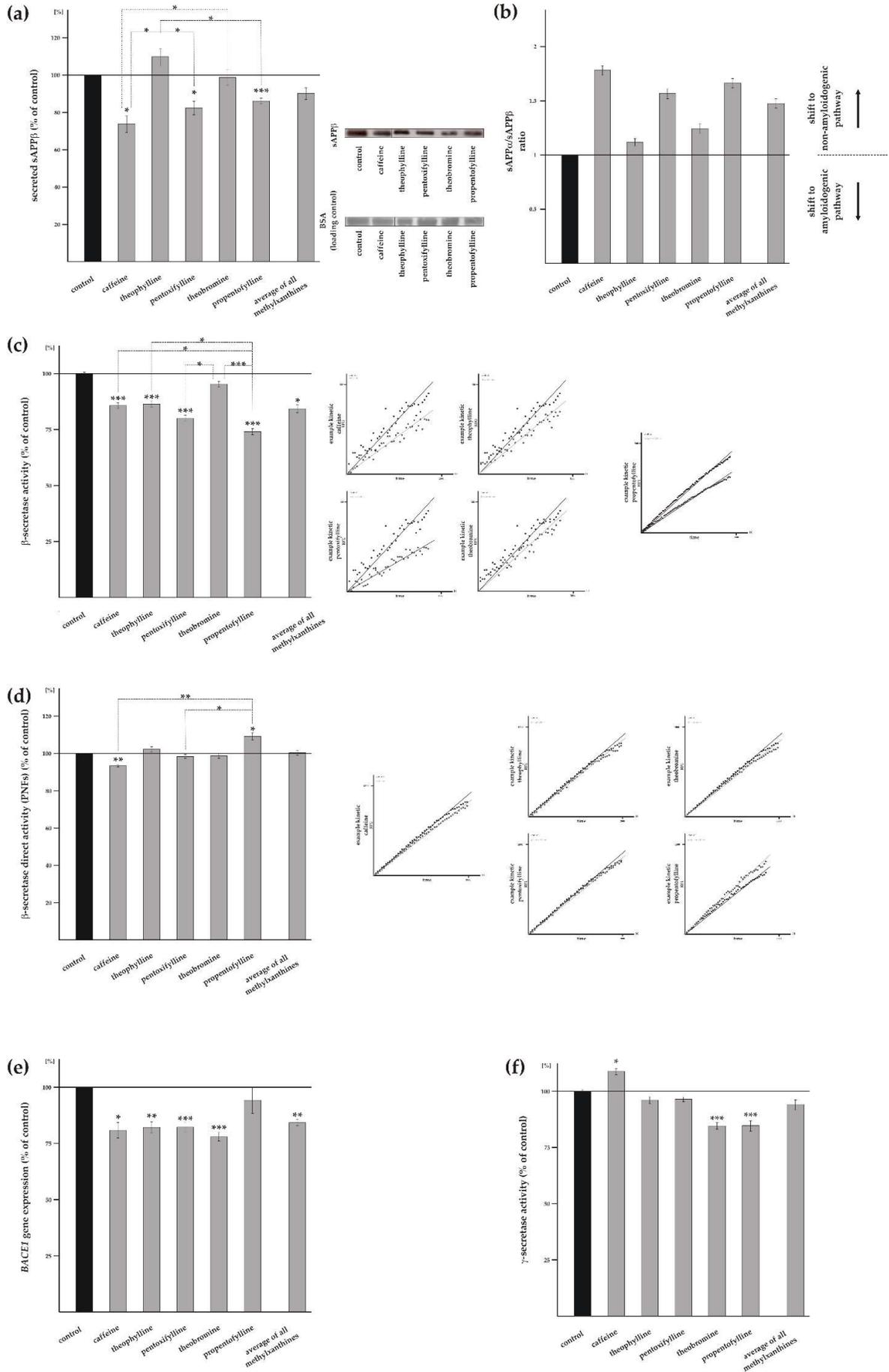
MTX	mean difference to control [%]
caffeine	3.8
theophylline	-3.27
pentoxifylline	2.34
theobromine	0.83
propentofylline	-1.91



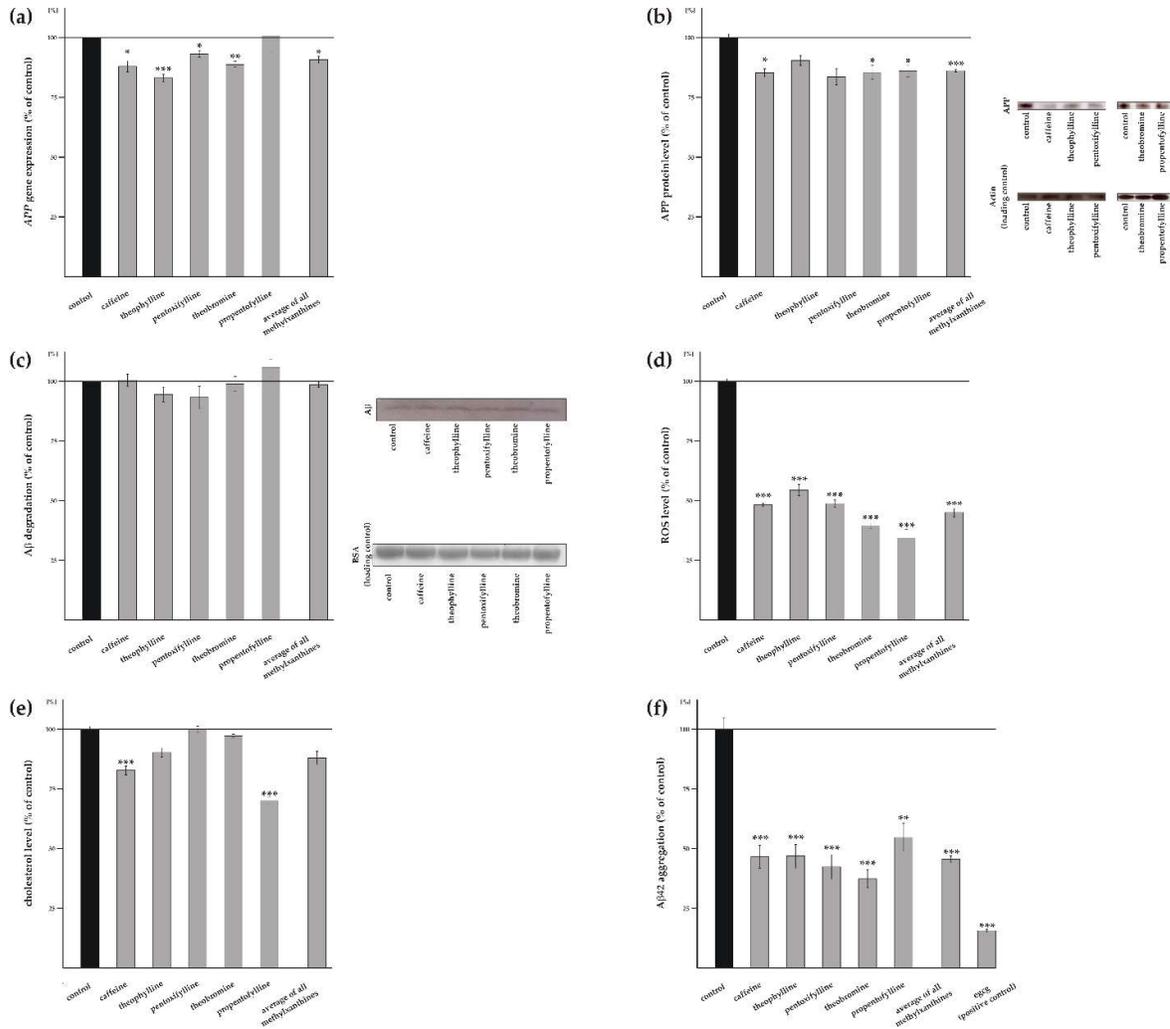
Supplemental Figure 1. Expression of APP<sup>695</sup> in SH-SY5Y wt and APP<sup>695</sup> transfected cells



**Supplemental Figure 2. Effects of caffeine, theophylline, pentoxifylline, theobromine and propentofylline on  $\alpha$ -secretase.** (a) Protein amounts of secreted soluble sAPP $\alpha$  of treated SH-SY5Y cells compared to untreated control cells ( $n \geq 9$ ). Representative WBs including load control are shown on the right. (b) Activity of  $\alpha$ -secretase in living SH-SH5Y cells after MTXs treatment ( $n \geq 4$ ). Representative kinetics are shown on the right side. (c) Protein level of ADAM10 ( $n \geq 4$ ). Representative WBs including load control are shown on the right. (d) *ADAM10* gene expression ( $n \geq 10$ ). (e)  $\alpha$ -secretase activity in post nuclear fractions of incubated SH-SY5Y cells ( $n \geq 4$ ) and resulting, representative kinetics after MTXs incubation. Error bars represent the standard error of the mean. Asterisks show the statistical significance calculated by unpaired Student's t test (\*  $p \leq .05$ ; \*\*  $p \leq .01$ ; \*\*\*  $p \leq .001$ ).



**Supplemental Figure 3. Influence of MTXs on amyloidogenic APP processing.** (a) Protein level of sAPP $\beta$  of treated SH-SY5Y APP<sup>695</sup> cells (n  $\geq$  13). Representative WBs including load control are shown on the right. (b) sAPP $\alpha$ /sAPP $\beta$  ratios. (c)  $\beta$ -secretase activity (n  $\geq$  19). Representative kinetics for each analyzed MTX are shown on the right. (d) Analysis of the activity of  $\beta$ -secretase in post nuclear fractions from SH-SY5Y cells (n  $\geq$  6). Example kinetics are illustrated on the right side. (e) Level of *BACE1* mRNA (n  $\geq$  10). (f) Activity of the  $\gamma$ -secretase (n  $\geq$  25). Error bars represent the standard error of the mean. Asterisks show the statistical significance calculated by unpaired Student's t test (\* p  $\leq$  .05; \*\* p  $\leq$  .01; \*\*\* p  $\leq$  .001).



**Supplemental Figure 4. Influence of MTXs on APP level, A $\beta$  catabolism and aggregation, reactive oxygen species and cholesterol.** (a) Gene expression of *APP* (n  $\geq$  10). (b) APP protein level in lysates of treated SH-SY5Y cells (n  $\geq$  6). Representative WBs including load control are shown on the right. (c) Influence of MTXs on ROS level in human neuroblastoma cells (n  $\geq$  8). (d) A $\beta$  degradation in the presence of MTXs (n  $\geq$  11). Representative WBs including load control are shown on the right. (e) Effect of MTXs on cholesterol level (n  $\geq$  4). (f) Influence of MTXs on A $\beta$ 42 aggregation (n  $\geq$  20). Epigallocatechin gallate (EGCG) served as positive control. Error bars represent the standard error of the mean. Asterisks show the statistical significance calculated by unpaired Student's t test (\* p  $\leq$  .05; \*\* p  $\leq$  .01; \*\*\* p  $\leq$  .001).