Cysteine Cathepsins Inhibition Affects Their Expression and Human Renal Cancer Cell Phenotype

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Figure S1. Detection of recombinant human Cts L and B activity through gelatin zymography assay. Recombinant purified human Cts L and B were loaded on the gelatin-gel zymogram. The Cts were activated through incubation of gel in an activating buffer (pH 4.8). Further staining with Coomassie brilliant blue G-250 showed the induction of proteolytic activity as bright white bands on dark background in correspondence with the bands of the enzymes.



Figure S2. Determination of the peptide inhibitory properties against human purified CtsB and L. The activities of human recombinant Cts B and L, were measured via fluorimetric analysis by exploiting the fluorogenic properties of the Triticain- α substrate Ac-PLVQ-AMC. In the assay, the 20 nM of recombinant human Cts L and Cts B were incubated with 50 μ M substrate without Cts inhibitors (red line) and with Ac-PLVE-FMK and Ac-VLPE-FMK (blue and green line, respectively) at a concentration of 2 μ M. Fluorescence was measured as relative as relative fluorescence unit (RFU).



Figure S3. Combined effect of PXT and the inhibitors on 769-P proliferation: 769-P cells were treated with 100 nM of PXT alone or in combination with 20 μ M of Ac-PLVE and Ac-VLPE and cell proliferation was measured after 24, 48, and 72 h via MTT assay. Data represent the mean (± S.D.) of three independent experiments, each performed in triplicate.



Figure S4. Spheroids size and circularity: (**a**,**b**) size and circularity of 769-P and (**c**,**d**) A498 cells upon treatment with Ac-PLVE-FMK and Ac-VLPE-FMK. Data represent the mean (± S.D.) of three independent experiments, each performed in triplicate.



Figure S5. Effect of Cts inhibitory peptides on cell stiffness: representative optical phase contrast image of cells (first row) and stiffness maps determined by indentation (Young's modulus) of 769-P cells (second row) with and without treatment with PLVE and VLPE. Calibration bars represent 25 μ m. The graph shows the values of Young's modulus of all analyzed samples, data are expressed as mean (± S.D.) and significance was calculated through one-way ANOVA followed by Dunnet's test. * = *p* < 0.05, ** = *p* < 0.01.



Figure S6. The effect of the peptide inhibitors on human 769-P and A498 renal cancer cell on lysosomes integrity. (**a**) 769-P and (**b**) A498 cells were treated with increasing doses of Ac-PLVE-FMK (red bars) and Ac-VLPE-FMK (blue bars) (2.5–250 μ M). NR uptake was measured after 24, 48, and 72 h via MTT assay. (**c**) LysoTracker red evaluation in 769-P and (**d**) A498 cells after 24, 48, and 72 h with the inhibitors (20 μ M). Data represent the mean (± S.D.) of at least three independent experiments, each performed in triplicate.



* The mean, normalized densitometry measurements for all samples (each protein) are provided in the main text of the paper.

72h













Figure S7. Uncropped western blots.