

Figure S1. Sites of KLK13 hydrolysis in libraries used for primed site profiling. (A) Library with general formula ABZ-Val-Arg-Phe-Arg-Ser-X₂'-X₃'-Tyr(3-NO₂); (B) library shown in panel A after incubation with KLK13; (C) library with general formula ABZ-Val-Arg-Phe-Arg-Ser-Thr-X₃'-Tyr(3-NO₂); (D) library shown in panel C after incubation with KLK13; (E) peptide ABZ-Val-Arg-Phe-Arg-Ser-Thr-Gln-Tyr(3-NO₂); (F) peptide shown in panel E after incubation with KLK13. ABZ fluorescence was monitored at the excitation and emission wavelengths of 320 and 450 nm, respectively. In panels C, D and F—retention time and MS identification of the major product are given above the chromatogram. Note that all tested substrates generated the same ABZ-Val-Arg-Phe-Arg-OH (t_R 12.56 min) product signifying KLK13 catalyze hydrolysis at Arg-Ser peptide bond, regardless the identity of the variable regions. (G) MS spectra of the libraries and optimized substrate 2 were recorded and confirmed the identity of the investigated compounds.

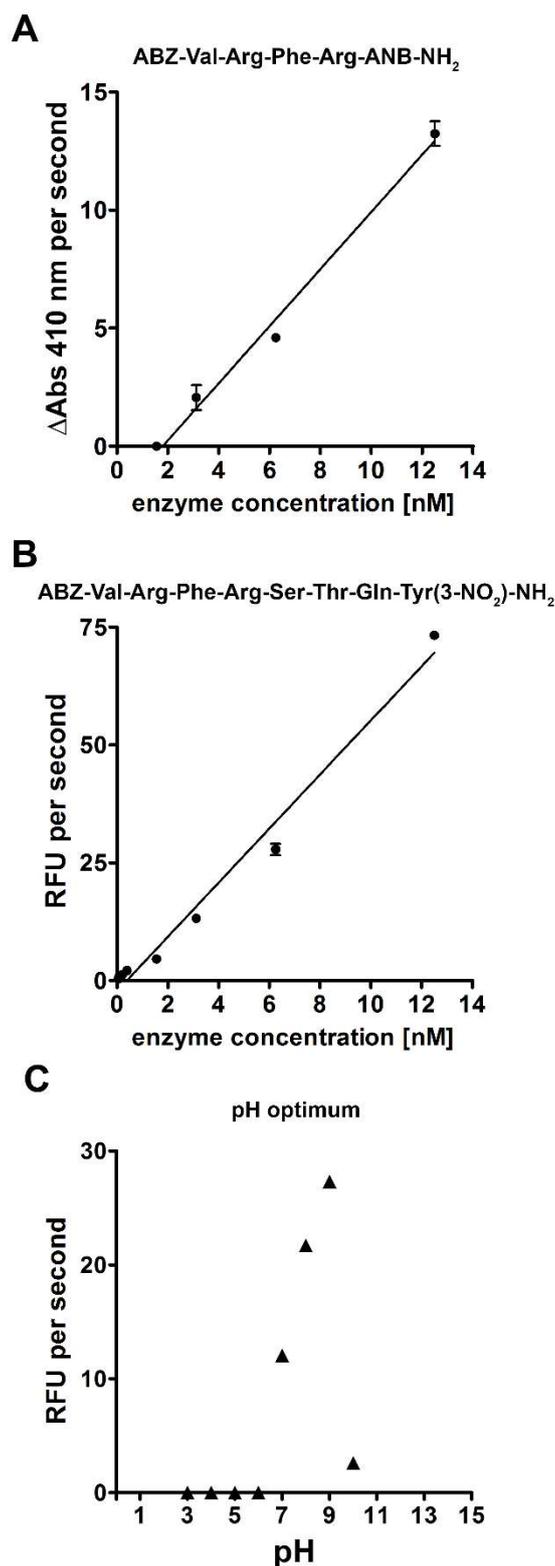


Figure S2. Detection limit and pH optimum of KLK13 as determined for the optimized substrates. Substrates **1** (A) and **2** (B) were titrated with decreasing amounts of KLK13 in 50 mM Tris-Cl (pH 7.5) buffer containing 1 mM EDTA and 0.5 μ M heparin at 37 °C. (C) pH dependence of KLK13. Processing of ABZ-Val-Arg-Phe-Arg-Ser-Thr-Gln-Tyr(3-NO₂)-NH₂ by KLK13 was monitored at pH range of 3 to 10 (citric acid/sodium citrate, sodium acetate/acetic acid, MES, MOPS, HEPES and TRIS buffers). Reaction progress was monitored by fluorescence detection at the excitation and emission wavelength 310 and 450 nm, respectively.