Supplementary Information

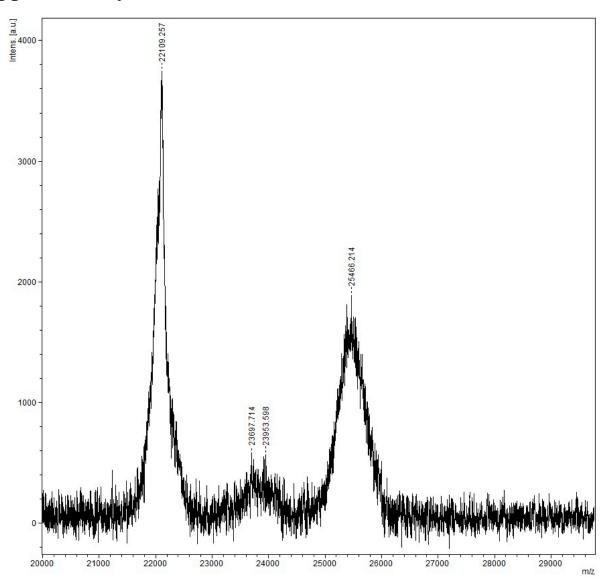


Figure S1. Mass spectrometric analysis of P1G10. The proteolytic fraction was desalted by passage on C4 HPLC column (Vidac). The absorbing fraction from HPLC was mixed with α -cyano-4-hydroxy-cinnamic acid (matrix) dissolved in 0.1 g/100 mL trifluoroacetic acid and processed in a UltrafleXtrem MALDI-TOF mass spectrometer (Bruker Daltonics, Leipzig, Germany).