

Figure S1: Water Surface Accessible Area for AA3 plotted against the Ramachandran angles  $\Phi$  and  $\Psi,\,nm^3$ 



Figure S2: Relative 'local' density of sodium **cations** (top) and **anions** (bottom) in the **first** (left) and **second** solvation shell of the **whole** tripeptide plotted against the molar concentration of ions. The concentrations are 0.20, 0.50, 1.00 and 2.00 M. Different lines correspond to different anions: blue solid line ( $\circ$ ) - F-; green dashed line ( $\Box$ ) -  $Cl^-$ ; black dotted line ( $\diamond$ ) -  $Br^-$ ; purple dash-dotted line ( $\bullet$ ) -  $I^-$ .



Figure S3: Relative 'local' density of sodium **cations** (top) and **anions** (bottom) in the **first** (left) and **second** solvation shell of the tripeptide **amino terminus** plotted against the concentration of ions. The concentrations are 0.20, 0.50, 1.00 and 2.00 M. Different lines correspond to different anions: blue solid line ( $\circ$ ) - F-; green dashed line ( $\Box$ ) -  $Cl^-$ ; black dotted line ( $\diamond$ ) -  $Br^-$ ; purple dash-dotted line ( $\bullet$ ) -  $I^-$ .



Figure S4: Relative 'local' density of sodium **cations** (top) and **anions** (bottom) in the **first** (left) and **second** solvation shell of the tripeptide **carboxyl terminus** group plotted against the concentration of ions. The concentrations are 0.20, 0.50, 1.00 and 2.00 M. Different lines correspond to different anions: blue solid line ( $\circ$ ) - F-; green dashed line ( $\Box$ ) -  $Cl^-$ ; black dotted line ( $\diamond$ ) -  $Br^-$ ; purple dash-dotted line ( $\bullet$ ) -  $I^-$ .



Figure S5: Relative 'local' density of sodium **cations** (top) and **anions** (bottom) in the **first** (left) and **second** solvation shell of the tripeptide **backbone** groups plotted against the concentration of ions. The concentrations are 0.20, 0.50, 1.00 and 2.00 M. Different lines correspond to different anions: blue solid line ( $\circ$ ) - F-; green dashed line ( $\Box$ ) -  $Cl^-$ ; black dotted line ( $\diamond$ ) -  $Br^-$ ; purple dash-dotted line ( $\bullet$ ) -  $I^-$ ..



Figure S6: Relative 'local' density of sodium **cations** (top) and **anions** (bottom) in the **first** (left) and **second** solvation shell of the tripeptide **side** groups plotted against the concentration of ions. The concentrations are 0.20, 0.50, 1.00 and 2.00 M. Different lines correspond to different anions: blue solid line ( $\circ$ ) - F-; green dashed line ( $\Box$ ) -  $Cl^-$ ; black dotted line ( $\diamond$ ) -  $Br^-$ ; purple dash-dotted line ( $\bullet$ ) -  $I^-$ .



Figure S7: Depth (left side) and position (right side) of the PMF minima between the **carboxyl terminus**,  $F^-$  anions (top) and  $Na^+$  cations (bottom) in NaF solution. The data are shown as functions of salt concentration. Different lines correspond to different PMFs: tripeptide group - water oxygen PMF corresponds to the solid blue line (thick,  $\circ$ ); tripeptide group - ion PMF corresponds to the green dashed line ( $\Box$ ); ion - water PMF corresponds to the black dotted line ( $\diamond$ ), ion - ion PMF corresponds to the purple dash-dotted line ( $\bullet$ ) -  $I^-$  and ion - counterion PMF corresponds to the red solid line (thin, crosses). The values for the minima depths are shown in  $k_BT$  units and the minima positions are shown in Angstroems.



Figure S8: The same data as on the Fig. 7 for the second side chain group and NaF solution.



Figure S9: The same data as on the Fig. 7 for the second side chain group and NaCl solution.



Figure S10: The same data as on the Fig. 7 for the second side chain group and NaBr solution.



Figure S11: The same data as on the Fig. 7 for the second side chain group and NaI solution.



Figure S12: The same data as on the Fig. 7 for the first backbone group and NaF solution.



Figure S13: The same data as on the Fig. 7 for the first backbone group and NaCl solution.



Figure S14: The same data as on the Fig. 7 for the first backbone group and NaBr solution.



Figure S15: The same data as on the Fig. 7 for the first backbone group and NaI solution.