

 $C_{\rm K} = 0.01, C_{\rm Ph} = 0.1, \varepsilon_{KPh} = -0.5$ 



 $C_{\rm K} = 0.01, C_{\rm Ph} = 0.1, \varepsilon_{KPh} = -0.6$ 

**Figure S1.** Endpoint snapshots of simulations of a  $64^3$  lattice size and either  $t = 10^5$  MCS (left) or  $t = 10^6$  MCS (right). The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations and the lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) for each row of simulation snapshots is shown below. Visually, peptides in snapshots at  $t = 10^6$  are more aggregated than at  $t = 10^5$ , demonstrating that a total of  $10^5$  time steps is insufficient for system to reach an equilibrium.



 $C_{\rm K} = 0.02, C_{\rm Ph} = 0.1, \varepsilon_{KPh} = -0.5$ 



 $C_{\rm K} = 0.02, C_{\rm Ph} = 0.1, \varepsilon_{KPh} = -0.6$ 

**Figure S1 continued.** Endpoint snapshots of simulations of a 64<sup>3</sup> lattice size and either  $t = 10^5$  MCS (left) or  $t = 10^6$  MCS (right). The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations and the lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) for each row is shown on the left. Visually, peptides in snapshots at  $t = 10^6$  are more aggregated than at  $t = 10^5$ , demonstrating that a total of  $10^5$  time steps is insufficient for system to reach an equilibrium.



 $C_K=0.2,\ \mathcal{E}_{KPh}\ = -0.5$ 

 $C_K=0.3, \quad \mathcal{E}_{KPh}\ = -0.6$ 



 $C_K=0.5,\ \mathcal{E}_{KPh}\ =-0.7$ 

**Figure S2.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step  $t = 10^6$  MCS. Simulations do not contain phosphate. The lysine (C<sub>K</sub>) concentrations and lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) are shown below each snapshot.



**Figure S2 continued.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step t =  $10^6$  MCS. The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations were 0.01 and 0.1, respectively. The lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) is shown below each snapshot.



 $\varepsilon_{KPh} = -0.5$ 

 $\varepsilon_{KPh} = -0.6$ 



**Figure S2 continued.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step t =  $10^6$  MCS. The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations were 0.02 and 0.1, respectively. The lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) is shown below each snapshot.



 $\varepsilon_{KPh} = -0.5$ 

 $\varepsilon_{KPh} = -0.6$ 



 $\varepsilon_{KPh} = -0.7$ 

 $\varepsilon_{KPh} = -0.8$ 

**Figure S2 continued.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step t =  $10^6$  MCS. The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations were 0.03 and 0.1, respectively. The lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) is shown below each snapshot.



 $\varepsilon_{KPh} = -0.5$ 

 $\varepsilon_{KPh} = -0.6$ 



**Figure S2 continued.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step t =  $10^6$  MCS. The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations were 0.05 and 0.1, respectively. The lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) is shown below each snapshot.



 $\varepsilon_{KPh} = -0.5$ 

 $\varepsilon_{KPh} = -0.6$ 



**Figure S2 continued.** Snapshots of simulations of a  $64^3$  lattice size and at the end of time step t =  $10^6$  MCS. The lysine (C<sub>K</sub>) and phosphate (C<sub>Ph</sub>) concentrations were 0.07 and 0.1, respectively. The lysine-phosphate interaction strength ( $\varepsilon_{KPh}$ ) is shown below each snapshot.