

1 **Supporting Information for**

2 **Investigating the correlation between the protein adhesion**  
3 **simulation and the biocompatibility of polymeric substrate for**  
4 **skin tissue engineering applications**

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## 2 Geometry optimization of the systems

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*Table S1. The geometry optimization task carried out throughout the simulations.*

Algorithm	Smart
Energy (kcal/mol)	$10^{-3}$
Force (kcal/mol/Å)	0.5

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5 **Table S1** shows the criterias selected for the Geometry optimizaion task in Materials Studio's  
6 Forcite module. This task helps to change and optimize the geometry of the structre  
7 according to the chosen criterias, energy and force in this case. This process continues until  
8 either the final iteretion or the criterias are reached. Hence, a successful task result in a  
9 substrate with the minimum potential energy.

## 10 Polymer surface and bulk substrate annealing

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*Table S2. The anneal task carried out on the unit polymerics surface.*

Annealing cycles	2
Initial temperatures (K)	298
Mid-cycle temperature (K)	450
Ensemble	NVT
Thermostat	Berendsen
Time step (fs)	1
Total simulation time (ns)	2

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*Table S3. The anneal task carried out on the bulk polymerics surface.*

Annealing cycles	2
Initial temperatures (K)	298
Mid-cycle temperature (K)	450
Ensemble	NVT
Thermostat	Berendsen

Time step ( $fS$ )	1
Total simulation time ( $nS$ )	3.5

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2 **Table S2 and S3** shows the criteria for unit and bulk polymeric cell, respectively. Since  
3 using the Geometry optimization task might cause the structure to be trapped in a local  
4 energy minimum and finishing the simulation prematurely, utilizing the Anneal task will  
5 search for the minimum energy of the structure in a much wider range by increasing and then  
6 decreasing the temperature of the substrate controlled via a selected thermostat.

### 7 **Dynamics simulation parameters carried out on the protein-polymer system**

8 *Table S4. The final molecular dynamics run carried out on the protein-polymer surface.*

Ensemble	NVT
Thermostat	Berendsen
Temperature ( $K$ )	310
Time step ( $fS$ )	1
Total simulation time ( $nS$ )	2.5

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10 After placing the protein near the relaxed polymeric surface, a Dynamics task will be carried out  
11 **Table S4**. In this task the structures will move according to the computed forces and factors like  
12 temperature controlled via a thermostat and the selected ensemble.

### 13 **Lattice parameters of Polymer surface and bulk substrate**

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15 *Table S5. the lattice parameters of constructed unit and bulk polymer surfaces.*

Samples	Unit models	Bulk models
Lattice dimensions	$43.092\text{\AA} \times 43.092\text{\AA} \times 20\text{\AA}$	$129.275\text{\AA} \times 129.275\text{\AA} \times 20\text{\AA}$

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1 The lattice parameters of unit cells with periodic boundary condition constructed by assembling the  
 2 polymer oligomers together. Furthermore, the bulk models were created by assembling the fabricated  
 3 the unit cells next to each other in the  $x - y$  plane.

#### 4 Berendsen thermostat

5 The Berendsen method utilizes an external bath with constant temperature or pressure with adjustable  
 6 time constants. Here, the gradual heat transfer between the selected structure and the environment  
 7 which is governed by the temperature scaling factor ( $\lambda$ ) (Eq. S1).

$$\lambda = \sqrt{1 - \frac{\Delta t}{\tau} \left( \frac{T_{instant} - T_0}{T_{instant}} \right)} \quad (\text{Eq. S1})$$

8 Where  $\Delta t$ ,  $\tau$ ,  $T_{instant}$ , and  $T_0$  corresponds to time step, relaxation time, instantaneous temperature, and  
 9 target temperature. This method allows for an accurate constant-temperature ensemble and can be  
 10 expanded to polyatomic molecules [1].

#### 11 Total energy of the simulated protein-polymer systems

12 *Table S6. The total energy of the simulated protein-polymer systems*

	$E_{\text{non-bonded}}$ ( $\frac{\text{Kcal}}{\text{mol}}$ )		$E_{\text{H-bond}}$ ( $\frac{\text{Kcal}}{\text{mol}}$ )		$E_{\text{van der Waals}}$ ( $\frac{\text{Kcal}}{\text{mol}}$ )	
	Collagen	Fibronectin	Collagen	Fibronectin	Collagen	Fibronectin
<b>CH/G-1:3</b>	387.193	180.900	-11.118	-7.372	376.075	173.528
<b>CH/G-1:2</b>	365.438	182.775	-3.500	-17.244	361.938	165.531
<b>CH/G-1:1</b>	347.983	151.122	-21.230	-17.345	326.753	133.777
<b>CH/G-3:1</b>	428.634	226.194	-10.372	-34.229	418.262	191.965
<b>CH/G-2:1</b>	410.120	237.332	-10.281	-18.342	399.839	218.99

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15 **References**

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2 1. Berendsen, H.J., et al., *Molecular dynamics with coupling to an external bath*. The Journal of  
3 chemical physics, 1984. **81**(8): p. 3684-3690.

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