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# 1 Quantification of the Effects of Hydrophobicity and 2 Mass Loading on the Effective Coverage of Surface- 3 Immobilized Elastin-like Peptides

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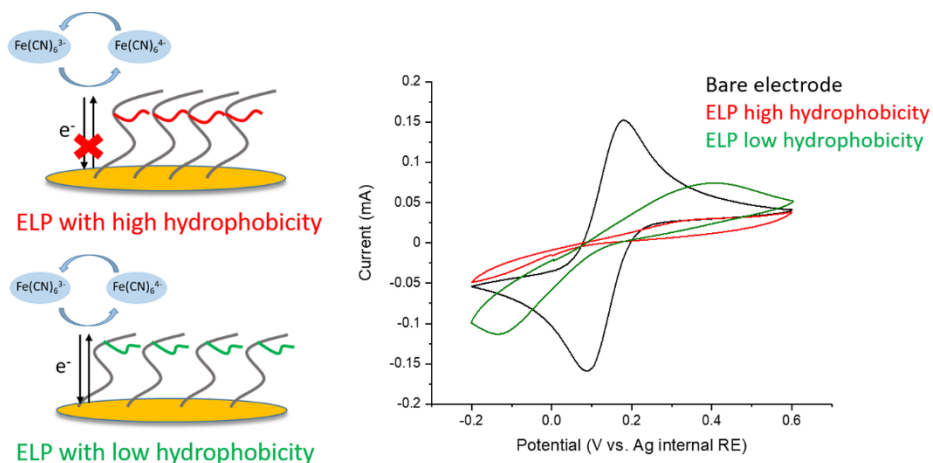
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## 10 **Abstract**

11 Elastin-like peptides (ELPs) immobilized to solid surfaces have recently gained attention for use  
12 in electrochemical applications in sensing as well as bioenabled electrode assembly. Key to the  
13 success of these applications is understanding how ELPs impact the access and electron transfer  
14 of reacting species to the solid surface (effective surface coverage). In this study, short ELPs with  
15 varying hydrophobicity and sequence length were designed for gold attachment, and the effect on  
16 the ability of a redox probe to access a gold surface was characterized by cyclic voltammetry. A  
17 quantitative model describing the relationship between ELP effective surface coverage as a  
18 function of mean hydrophobicity and mass loading was elucidated based on the results, showing

19 the ability to precisely control surface properties by tuning the ELP sequence. This model will be  
20 useful for the design of surface-bound ELP sequences that exhibit desired effective surface  
21 coverage for electrochemical as well as biomaterial applications.

## 22 Graphical abstract



23 Surface properties are controlled by tuning elastin-like peptide sequences

## 24 Keywords

25 Elastin-like polypeptides, electrode modification, cyclic voltammetry, biotechnology

## 26 1. Introduction

27 Recently, protein and polypeptide engineering have emerged as promising tools for  
28 electrochemical applications. Taking advantage of the ability to precisely define sequences and  
29 achieve multiple specific functions, protein and polypeptide-containing thin film electrode  
30 modifications have already been applied to biosensing, enzyme-based electrode design, and  
31 biomedical device manufacturing [1–3].

32 Elastin is a particularly attractive polypeptide platform for electrochemical applications.  
33 Engineered elastin-like polypeptide sequences (ELPs) are derived from the hydrophobic  
34 domains of tropoelastin, with a repetitive motif consisting of Val-Pro-Gly-Xaa-Gly, where  
35 X is a guest residue that can be substituted by any amino acid except proline [4]. ELPs are  
36 well-known as stimuli-responsive biopolymers, exhibiting reversible thermal-dependent  
37 lower solution critical temperature (LCST) behavior in aqueous solutions, where they are  
38 soluble below the transition temperature and insoluble above it. This transition temperature  
39 ( $T_i$ ) is dependent on the environmental conditions [5,6] as well as peptide concentration,  
40 chain length [7], and hydrophobicity [8]. With these unique sequence-defined  
41 characteristics, including stimulus-responsive and self-assembly behavior, ELPs have been  
42 designed for variety of applications, such as: protein purification, drug delivery, and tissue  
43 engineering [9]. The transition behavior of ELPs is also maintained in immobilized  
44 assemblies, leading to surface-bound “smart” applications [10].

45 Label-free electrochemical sensing platforms based on ELP transducers assembled on gold  
46 surfaces have recently been explored, where elastin in the insoluble state blocks electron  
47 transfer between a redox probe, specifically the common  $\text{Fe}(\text{CN})_6^{3-/4-}$  redox couple, and  
48 the gold electrode surface [11]. When the immobilized elastin is in the soluble state, the  
49 redox probe can access the electrode surface and electron transfer happens more readily. In  
50 essence, the current changes observed are indicative how exposed the gold surface is to the  
51 electrolyte when modified with molecular layers [12]. However, the impact of changing the  
52 assembled ELP guest residue content and mass loading on surface exposure, as measured  
53 via electron transfer performance between a redox probe and the gold electrode, is largely  
54 unexplored. The sensitivity of elastin-based electrochemical sensing platforms depends on

55 the differences in surface exposure, which may vary depending on the chosen ELP  
56 sequence. In addition, the application of ELPs has expanded to bioenabled electrode  
57 assembly, where surface access and chemical versatility is desired [13–17]. There has also  
58 been a particular recent interest in exploiting the responsive behavior of short elastin  
59 polypeptides on functionalized gold nanoparticles [18], which is useful in other non-  
60 electrochemical areas including drug delivery [19], imaging [20], and as active plasmonic  
61 waveguides [21]. Surface exposure is also important in these applications, as exposed  
62 substrates may be prone to biofouling [22].

63 In this study, we hypothesize that surface access (or effective surface coverage), as  
64 measured via electron transfer between a redox probe and the gold electrode, can be  
65 controlled by modifying the guest residue hydrophobicity and mass loading of assembled  
66 engineered short ELPs. Herein, we utilized cyclic voltammetry (CV) to investigate the  
67 effective surface coverage of different ELPs when assembled on gold electrodes with  
68 varying guest residues and length. A quartz crystal microbalance with dissipation (QCM-  
69 D) was utilized to estimate the mass loading and observe the layer formation process. The  
70 results show that effective surface coverage can be precisely and predictably tuned using  
71 assembled ELPs. The proposed model for effective surface coverage will serve as guidance  
72 for future ELP-based electrochemical sensing platforms and electrode design.

73

74

75

## 76 **2. Materials and methods**

### 77 **2.1 Materials**

78 Detailed materials information is provided in the Supplementary Data.

### 79 **2.2 Peptide design**

80 Peptide sequences utilized in this study consisted of an elastin-like motif, (VPGXG)<sub>n</sub>, where n =  
81 number of pentapeptide repeats. The N-terminus was modified with cysteine (C) for all peptides.  
82 The ends of the peptides were acylated and amidated. All sequences are shown in Table S1. All  
83 peptides were purchased from GenScript at purities above 95%.

### 84 **2.3 Electrode preparation**

85 Screen printed electrodes (Metrohm DropSens, DRP-220BT, L33 × W10 × H0.5 mm) with a gold  
86 working electrode (4 mm diameter), silver reference electrode, and platinum counter electrode  
87 were used in this study. Before electrode functionalization, all SPEs were prepared by cleaning in  
88 0.5 M H<sub>2</sub>SO<sub>4</sub> solution (60 μL to fully cover the SPE electrodes), performing cyclic voltammetry  
89 scans from -0.2 V to 1.3 V (versus internal silver reference electrode) at the scan rate of 100 mV  
90 s<sup>-1</sup>, with 9 scans accumulations [23]. After cleaning, SPEs were rinsed with deionized (DI) water  
91 and dried with N<sub>2</sub>. All SPEs were used once per test in this study, and not reused.

### 92 **2.4 Peptide incubation**

93 All peptides were prepared at 10 μg/mL in 0.01 M phosphate buffered saline (PBS, pH 7.4). For  
94 experiments on screen-printed electrodes, 60 μL of the peptide solution was deposited on the SPE  
95 electrode surface. The same amount of PBS solution (60 μL) was added for the bare electrode

96 experiments. To ensure the electrode surface was maximally saturated with peptide adsorption, all  
97 SPE samples were incubated for overnight at 4°C. Prior to electrochemical characterization,  
98 electrodes were gently rinsed with PBS solution and dried with N<sub>2</sub> gas. All peptide solutions were  
99 prepared freshly prior to experiments. In buffer concentration experiments, all solutions were made  
100 in 0.1 M PBS (pH=7.4) unless otherwise stated.

## 101 **2.5 Cyclic voltammetry**

102 All electrochemical experiments were performed on SPEs that connected to a Metrohm Autolab  
103 potentiostat (controlled by Nova 2.1 software). The redox buffer contained equimolar amounts of  
104 4 mM K<sub>3</sub>Fe(CN)<sub>6</sub> and 4 mM K<sub>4</sub>Fe(CN)<sub>6</sub> in 0.01 M PBS (pH 7.4) with 0.1 M KCl. Cyclic  
105 voltammetry (CV) measurements on SPEs were conducted at room temperature, at a range of -0.2  
106 V to 0.6 V and a scan rate of 100 mV s<sup>-1</sup> without any preconditional potential or accumulation time.  
107 For each sample, 60 μL redox solution was used to fully cover the SPE electrodes, and five scans  
108 were collected. In forward of CV scans, ferrocyanide is oxidized to ferricyanide, and in reverse  
109 scans, ferricyanide is reduced to ferrocyanide. Experiments with varying scan rate were collected  
110 using equimolar amounts of 5 mM K<sub>3</sub>Fe(CN)<sub>6</sub> and 5 mM K<sub>4</sub>Fe(CN)<sub>6</sub> in 0.01 M PBS (pH 7.4) with  
111 0.1 M KCl at a range of 10-100 mV s<sup>-1</sup>. Redox probe concentration experiments were collected  
112 with equimolar solutions from 0 to 100 mM in 0.01 M PBS with 0.1M KCl solution at 50 mV s<sup>-1</sup>.

## 113 **2.6 Quartz crystal microbalance with dissipation**

114 A quartz crystal microbalance with dissipation (QCM-D, QSense Explorer, controlled by QSoft  
115 software, Biolin Scientific) was used to investigate the peptide adsorption behavior and the relative  
116 mass loading of each peptide on gold surfaces. Frequency shifts and dissipation changes were

117 monitored simultaneously versus time. The details of these experiments can be found in our  
118 previous work [15,16], and in the Supplementary Data.

## 119 **2.7 Circular dichroism**

120 Circular dichroism (CD) spectroscopy was utilized to analyze the secondary structure of triple-  
121 repeat elastin-like peptides. Procedures can be found in our previous work [16], with details in the  
122 Supplementary Data.

## 123 **2.8 Atomic force microscopy**

124 Atomic force microscopy (AFM) was used to analyze the topography of surface-immobilized  
125 triple-repeat elastin-like peptides VKV and KVK. Procedures can be found in our previous  
126 works[15,16], with details in the Supplementary Data.

## 127 **2.9 Statistical analysis**

128 Data are represented as the mean  $\pm$  the standard deviation. Analysis of variance (ANOVA) was  
129 performed using Minitab to determine if a factor had a significant effect. Statistical groupings were  
130 determined by Tukey's *post hoc* test. Simple linear regression of average values was performed  
131 using Minitab. Best fit lines are obtained using the method of least squares. All statistical tests  
132 used  $\alpha = 0.05$ . For results in the main text,  $n = 3$  except in the case of single-repeated peptide where  
133  $X = E$  ( $n = 5$ ) and triple-repeated peptide where  $X = K_3$  ( $n=4$ ).

134

135

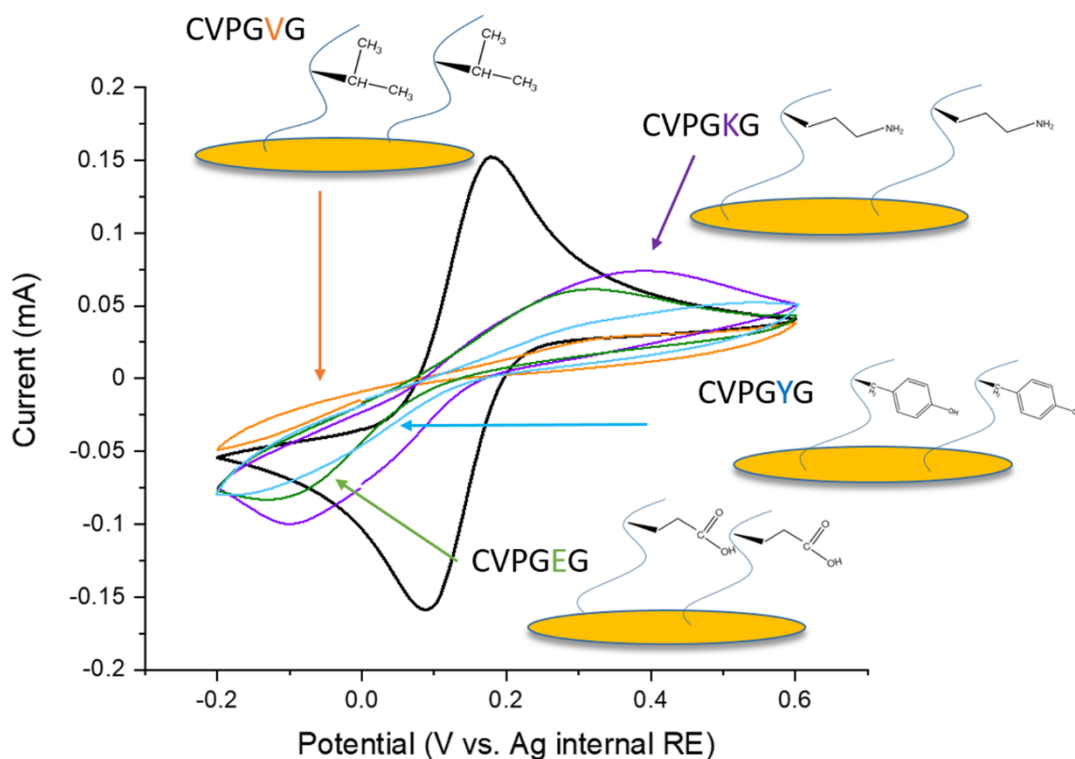
136



137 **3. Results**

138 **3.1 The influence of gold-immobilized single-repeat ELP guest residue on**  
139 **Fe(CN)<sub>6</sub><sup>3-/4-</sup> electron transfer**

140 Cyclic voltammetry (CV) was performed in the presence of a Fe(CN)<sub>6</sub><sup>3-/4-</sup> redox probe on  
141 gold electrodes with assembled ELPs. To investigate the effect of guest residue  
142 hydrophobicity, single-repeat elastin peptides with different guest residues were designed  
143 with the sequence Ac-CVPGXG-NH<sub>2</sub>, featuring an N-terminal cysteine (C) for rapid  
144 immobilization to the gold surface via a thiol bond. Table S1 contains all sequences  
145 explored in this study. All peptides were acetylated and amidated to ensure any charge  
146 effects were imparted only by the guest residues. Lysine (K) and glutamic acid (E), served  
147 respectively as positively and negatively charged guest residues. Valine (V) and tyrosine  
148 (Y) were selected as neutral guest residues of varying hydrophobicity. The single-repeat  
149 peptides were assembled on prepared screen printed electrodes (SPEs) and exposed to the  
150 redox solution. Each CV experiment consisted of five scans. Stable data indicated the  
151 assembled elastin did not change significantly during the scanning period (Figure S1).  
152 Representative CVs are shown in Figure 1.



153  
 154 **Figure 1.** Cyclic voltammograms obtained on ELP-modified SPEs with varying ELP guest  
 155 residue hydrophobicity demonstrate controlled electron transfer of a label-free redox probe.  
 156 Results were collected in 0.01 M PBS with 4 mM  $\text{Fe}(\text{CN})_6^{3-/4-}$  redox couple and 0.1 M KCl,  
 157 from -0.2 V to 0.6 V (vs. silver internal reference electrode) with a 0.1 V/s scan rate.  
 158 Representative data are shown, and include bare gold SPE electrode (black), V modified  
 159 electrode (orange), K modified electrode (purple), E modified electrode (green), and Y  
 160 modified electrode (blue).

161  
 162 On bare gold electrodes, as shown in Figure 1 (black line), a pair of redox peaks was  
 163 observed, with a peak-to-peak separation of ~60 mV which is expected for diffusion-

164 controlled, reversible redox reactions. In contrast, CVs taken on peptide-modified  
165 electrodes either had no discernible peaks, or had increased peak-to-peak separations,  
166 which is indicative of a quasi-reversible redox reaction. Plots of peak current versus the  
167 square root of scan rate for selected peptides in this study (Figure S2) confirm the quasi-  
168 reversible nature, with slight deviations from perfect linearity observed. In addition, the  
169 peak currents observed on peptide-modified electrodes were lower compared to peak  
170 currents observed on bare electrode. The results indicated that all ELPs were successfully  
171 immobilized on electrode surface, and different peptide layers were reducing the available  
172 electrode surface area or hindering the ability of electron transfer from the redox probe in  
173 solution to the underlying electrode.

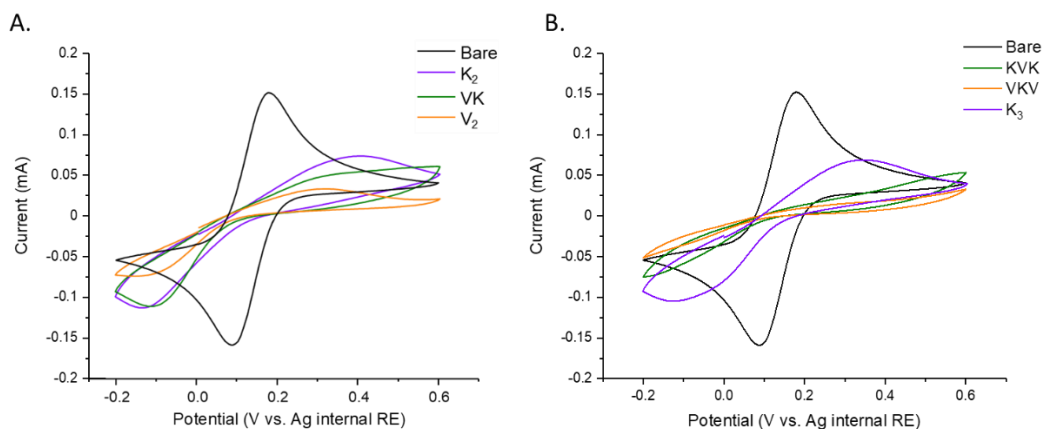
174 To quantify the extent to which the peptides block electron transfer, the total charge passed  
175 on the second CV scan of each peptide sample was normalized to the bare gold electrode  
176 sample. Thus, to compare samples we define effective surface coverage,  $f_e$ , as:

177 Equation 1:  $f_e = 1 - \frac{\text{total charge for sample } (C_i)}{\text{total charge for bare gold } (C_0)}$

178 Using this definition, we discovered  $f_e$  was guest residue dependent, being  $0.24 \pm 0.03$ ,  $0.37$   
179  $\pm 0.12$ ,  $0.43 \pm 0.11$ ,  $0.57 \pm 0.02$  for guest residue X = K, E, Y and V, respectively. We  
180 observed that in general  $f_e$  was positively related to hydrophobicity, with immobilized  
181 peptides containing guest residue X = V, Y and E in one statistical grouping, and X = Y,  
182 K, E being in another statistical grouping (see Table S2 for ANOVA and *post hoc* testing  
183 results), indicating that hydrophobicity of the guest residue has a significant impact on  
184 electron transfer.

185 **3.2 The influence of gold-immobilized double- and triple-repeat ELP guest residue**  
186 **on  $\text{Fe}(\text{CN})_6^{3-/4-}$  electron transfer**

187 To further investigate the effects of ELP guest residue hydrophobicity and length in  
188 controlling  $f_e$ , double- and triple-repeat ELPs were designed with varying ratios of  
189 positively charged guest residue (K) and neutral hydrophobic guest residue (V) (Table S1  
190 contains exact peptide sequences). Briefly, three double-repeat peptides were designed  
191 having the general form of  $\text{Ac-CVPGX}_1\text{GVPGX}_2\text{G-NH}_2$ , with guest residues occurring in  
192 the order  $X_1X_2 = \text{K}_2, \text{VK}$  and  $\text{V}_2$ . Three triple-repeat peptides were designed with the  
193 general form of  $\text{Ac-CVPGX}_1\text{GVPGX}_2\text{GVPGX}_3\text{G-NH}_2$ , with guest residues occurring in  
194 the order  $X_1X_2X_3 = \text{K}_3, \text{KVK}$  and  $\text{VKV}$ . Figure 2 shows the representative CV scans on  
195 double- and triple- repeat peptide functionalized electrodes as well as a bare gold electrode  
196 for comparison.



197  
198 **Figure 2.** Cyclic voltammograms demonstrate that the average guest residue hydrophobicity of  
199 double- and triple-repeat elastin peptides assembled on gold impact the electron transfer of a label-  
200 free redox probe. Results were collected in 0.01 M PBS with 4 mM  $\text{Fe}(\text{CN})_6^{3-/4-}$  redox couple and

201 0.1 M KCl, from -0.2 V to 0.6 V (vs. silver internal reference electrode) with a 0.1 V/s scan rate.  
202 Representative cyclic voltammograms are shown for (A) bare electrode (black), K<sub>2</sub> (purple), VK  
203 (green), and V<sub>2</sub> (orange); (B) bare electrode (black), K<sub>3</sub> (purple), KVK (green) and VKV (orange).

204  
205 The CVs obtained on electrodes functionalized with double- and triple-repeat ELPs  
206 exhibited a similar trend shown in Figure 1, where it was observed that as the fraction of V  
207 increases, the average hydrophobicity increases, and  $f_e$  increases, coinciding with an  
208 increase in peak-to-peak separation and decrease in peak current. For electrodes  
209 functionalized with double-repeat ELPs K<sub>2</sub>, VK, and V<sub>2</sub>,  $f_e$  was  $0.23 \pm 0.06$ ,  $0.43 \pm 0.10$ ,  
210 and  $0.57 \pm 0.03$ , respectively. On triple-repeat ELP-modified electrodes with K<sub>3</sub>, KVK and  
211 VKV,  $f_e$  was  $0.32 \pm 0.03$ ,  $0.55 \pm 0.09$ , and  $0.77 \pm 0.04$ , respectively. ANOVA and *post hoc*  
212 analysis (Tables S3 and S4) indicated that the average hydrophobicity of the guest residues  
213 had a significant impact on electron transfer for double- and triple repeat peptide samples.

214

### 215 **3.3 A quantitative model for effective surface coverage**

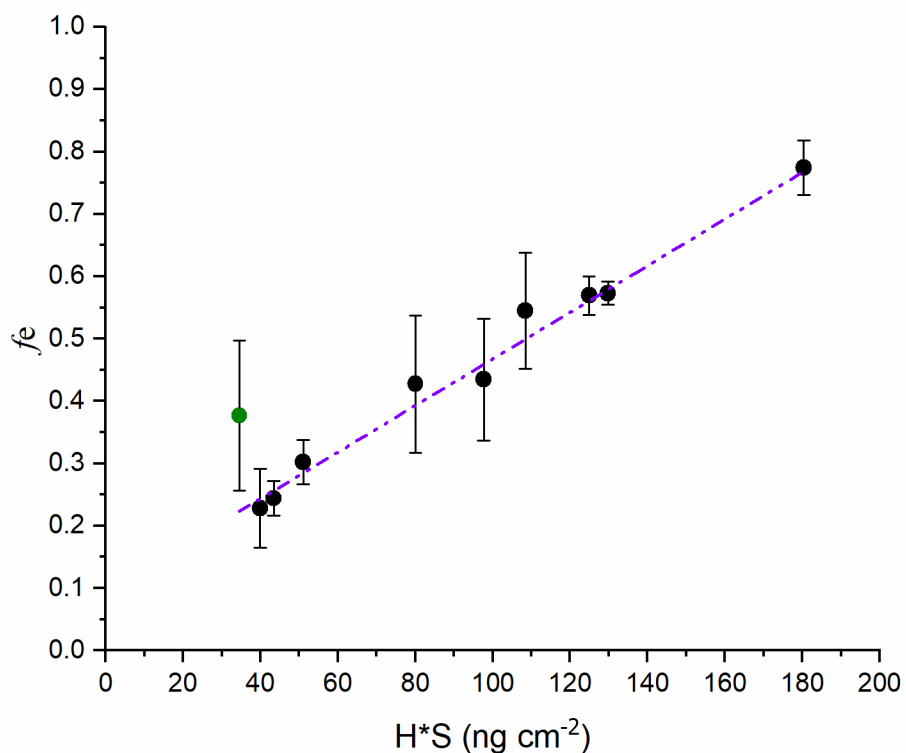
216 To compare the relative mass loading on gold electrodes of different ELPs, hydrated mass  
217 loading of ELPs were estimated using a quartz crystal microbalance with dissipation  
218 (QCM-D). Table S5 shows estimated relative hydrated mass loadings for all peptides in  
219 this study, and Figure S3 shows representative QCM-D runs of triple-repeat ELPs.

220 Given that relative mass loading and hydrophobicity were hypothesized to significantly  
221 affect the  $f_e$ , we proposed the following model:

222 Equation 2:  $f_e = k_e S (H) + f_{e,min}$

223 Where  $k_e$  is a parameter to describe the dependence of  $f_e$  on hydrophobicity and  
224 pentapeptide loading ( $\text{cm}^2 \text{ng}^{-1}$ ),  $S$  is the mass loading of elastin per unit area ( $\text{ng cm}^{-2}$ ), and  
225  $f_{e,min}$  is the minimal amount of coverage for a monolayer of elastin.  $H$  is the relative  
226 hydrophobicity normalized to valine such that  $H$  is the average hydrophobicity compatibility  
227 index of the guest residues divided by the hydrophobicity compatibility index of valine[24].  
228 Calculation examples are provided in Table S6.

229 Based on Equation 2, a plot of  $f_e$  versus  $H*S$  will yield a straight line, where the slope is  
230 equal to  $k_e$  and the intercept represents  $f_{e,min}$ . Figure 3 shows  $f_e$  values calculated from data  
231 represented in Figure 1 and 2 plotted as a function of  $H*S$ . Linear regression was performed  
232 on average values and the linear relationship between  $f_e$  and  $H*S$  was statistically  
233 significant (see Figure S4, and Table S7 for regression results).



234

235 **Figure 3.** Effective coverage,  $f_e$ , is linearly related to hydrophobicity compatibility with valine (H)  
 236 multiplied by the mass loading (S) of ELPs immobilized on gold. Each data point represents the  
 237 average  $\pm$  the standard deviation for at least  $n = 3$  independent trials. A simple linear regression  
 238 was performed using Minitab, and the purple line represents the best fit line. Results from the  
 239 linear regression can be found in the Supplementary Data. The data point in green represents an  
 240 outlier that is not included in the linear regression for the final model.

241

242 Conditions such as scan rate, redox probe concentration and PBS concentration were varied  
 243 to identify impact on the model. Experiments conducted at different scan rates are provided  
 244 in Supplementary Data Figure S5 and S6. Based on the results, scan rate needs to be sufficiently

245 high for the model to be valid. Redox probe concentration was varied, and data collected on VKV  
246 peptide-modified SPEs (Figure S7). When the redox concentration  $> 1\text{mM}$ , peak current is 100X  
247 greater or more compared to the control of 0 mM redox probe. Thus, the concentration was  
248 sufficiently high in experiments used to develop the model. In addition, Figure S7B shows that  
249 there is a linear relationship between peak current and concentration, indicating that the redox  
250 probe concentration does not significantly change the properties of the assembled peptide layer.  
251 The effect of higher PBS concentrations on the model is shown in Figure S8, where 10X PBS  
252 results in a linear relationship between  $fe$  and  $\text{H}^*\text{S}$ , but the slope is lower ( $\sim 0.0015$ ). This indicates  
253 that the model is valid at higher salt concentrations and sufficiently high scan rates, but the slope  
254 may be dependent on ionic strength.

255

256 An alternative model based on ELP length instead of mass loading is provided in the  
257 Supplementary Data (Figure S9). In the alternative model,  $fe$  is a linear function of  $\text{H}/\text{L}$ , where  
258 each length of peptide has its own distinct slope and  $\text{L}$  = the number of elastin repeats. As the  
259 peptide length increases, the slope increases, indicating the dependence of  $fe$  on  $\text{H}/\text{L}$  becomes  
260 stronger such that a small change in hydrophobicity for a long peptide will have a greater impact  
261 on  $fe$ . Previous studies have shown that more intermolecular aggregations and hydrophobic  
262 collapse happens when ELP length increases.[5,25] Overall, the alternative model based on length  
263 provides additional evidence that  $fe$  may be impacted by the number of hydrophobic interactions  
264 in an ELP layer, indicated by the model based on mass loading in Figure 3 (see Discussion section  
265 for more insight into this result).

266



### 267 3.4 The effect of guest residue charge on peptide assembly

268 The data gathered with the single repeat peptide where X = a negatively charged glutamic  
269 acid (E) was higher than expected (shown in green in Figure 3). Figure S3A shows that the  
270 data point lies outside of the 95% confidence interval for the linear relationship. Without  
271 the outlier, linear regression results in  $k_e = 0.0038 \pm 0.0004 \text{ cm}^2 \text{ ng}^{-1}$  and  $f_{e,min} = 0.093 \pm$   
272  $0.046$  (95% confidence intervals). To investigate if the electrostatic effect during peptide  
273 assembly, we exposed E and K peptides to acidic and basic environments, respectively, to  
274 neutralize the guest residues. CV experiments were performed after the assembly in  
275 neutralized conditions to quantify the impact on  $f_e$ . As demonstrated in Figure S10, peptides  
276 assembled under neutral conditions had the same CV results as when they were assembled  
277 in their charged states. This result suggests the electrostatic charge in single-repeat peptides  
278 has minimal influence on the peptide assembly on gold. Therefore, further investigation is  
279 needed to understand why the outlier exists in our model.

## 280 4. Discussion

281 When considering the physical implications of the model for  $f_e$  in Equation 2, it important to note  
282 that the model (linear fit) was not significant when using traditional hydrophobicity scales by  
283 Wimley and White [26], or the scale developed by Urry based on elastin [27]. The relationship  
284 manifested specifically as a function of relative hydrophobe compatibility with valine. Therefore,  
285 we speculate that likely number of hydrophobic interactions, which would be directly related to  
286 hydrophobe compatibility with valine (common to all pentapeptide repeats) and the mass loading  
287 of the peptides on the surface, is the main predictor of  $f_e$ . Similarly, the hydrophobicity of elastin  
288 guest residues has been shown to influence mechanical properties [28].

289

290 We also note that secondary structure features of the immobilized ELPs may play a role in the  
291 observed model, as they change as a function of hydrophobicity. It has been previously  
292 demonstrated that guest residue substitutions, specifically hydrophobicity, can alter the propensity  
293 of elastin to form  $\alpha$ -,  $\beta$ -, and  $\pi$ -turns in solution [29]. In a recent study,  $\beta$ -turn propensity alone was  
294 not a significant driver of ELP properties in solution, but when dimerization was considered, the  
295  $\beta$ -turn content altered ELP properties, specifically hydrophobic accessible surface area [30]. The  
296 short ELP surface-assembled structures were investigated in our previous work, where Au-binding  
297 was shown to only occur when the cysteine moiety was included in the sequence, and the  
298 characteristic  $\beta$ -turn structure was observed when the peptides were immobilized to gold.[16,17,31]  
299 In this current study, surface-immobilized triple-repeat ELPs, VKV and KVK, were investigated  
300 by atomic force microscopy (AFM), and results are provided in SI (Figure S11). The AFM reveals  
301 that the triple-repeat ELPs have distinct topological features when attached to a gold surface.  
302 Circular dichroism (CD) was utilized in our study to qualitatively analyze the characteristic  
303 features of triple-repeat ELPs in aqueous condition. The secondary structures were more distinct  
304 with the increasing fraction of valine (Figure S12), indicating the ELP structures in solution are at  
305 least correlated with the results in Figure 3, and surface-bound structure is worthy of future  
306 investigation.

307 The importance of the developed model is that it can be used to predict the effective  
308 coverage of different short ELPs when immobilized on surfaces with various combinations  
309 of guest residues. We note that while it is somewhat analogous to previous models which  
310 predict the transition temperature of higher molecular weight ELPs in solution [7,32], our

311 model is not describing LCST behavior, but instead the properties which impact the  
312 accessibility of the surface to aqueous components. For example, this model can be applied  
313 to designing peptide-functionalized surfaces to control electrode organization, where  
314 accessibility of the modified electrode surface is desired. We recently published studies  
315 where a single-repeat ELP sequence (X=E) enhanced the ionic transport of a thin ionomer  
316 layer, despite being adsorbed to the gold electrodes [15]. This current study shows that a  
317 single-repeat peptide where X=E would have relatively low  $f_e$ , thus corroborating the results  
318 of our previous study. Another area where the model will have impact is in the field of  
319 sensor design where ELPs are proposed as an active transducer [11]. It would be ideal to  
320 have the greatest difference in  $f_e$  between ELP in the soluble and insoluble states, and this  
321 study provides a basis on which future work optimizing ELP-based sensor sequences can  
322 be built. This study also provides direction for future studies where the impact of bound  
323 ELP on redox probe diffusion, electrochemically active surface area, and electron transfer  
324 rate will be fully elucidated and quantified.

## 325 **5. Conclusion**

326 This work elucidates the relationship between effective surface coverage, ELP guest  
327 residue, and mass loading for ELP sequences immobilized to gold. Specifically, there exists  
328 a linear relationship between effective coverage and the product of the ELP guest residue  
329 hydrophobe compatibility with valine and mass loading. This model demonstrates the  
330 potential for ELPs to be precisely designed for future electrochemical and biomaterial  
331 applications.

332

333 **Associated Content**

334 Supplementary data includes supporting experimental details as well as Figures S1–S12 and  
335 Tables S1–S7.

336

337 **Author Contributions**

338 Zihang Su: Conceptualization, methodology, formal analysis, investigation, resources, data  
339 curation, writing - original draft, visualization. ChulOong Kim: investigation, formal analysis,  
340 writing – review & editing. Julie N. Renner: Conceptualization, methodology, formal analysis,  
341 writing – review & editing, supervision, project administration, funding acquisition.

342

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