

## Supporting Information for:

### Critical Domain Sizes of Heterogeneous Nanopattern Surfaces with Optimal Protein Resistance

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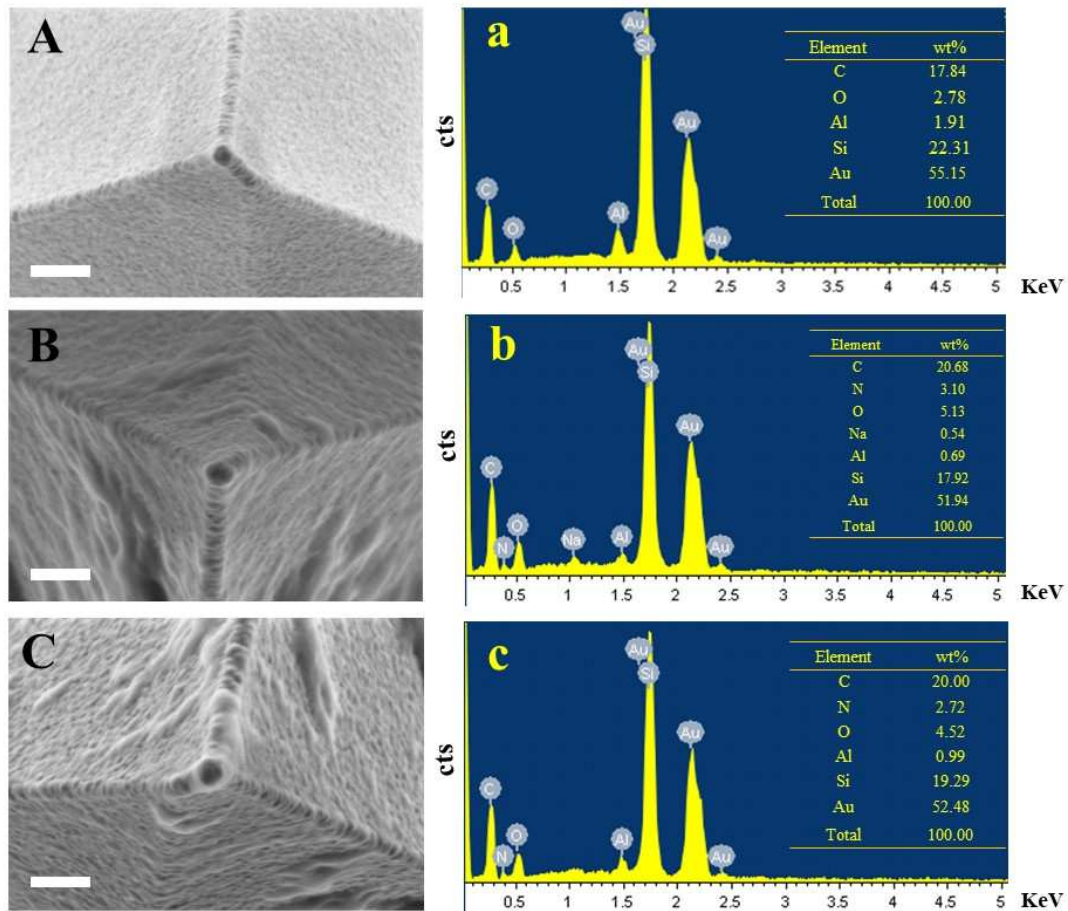
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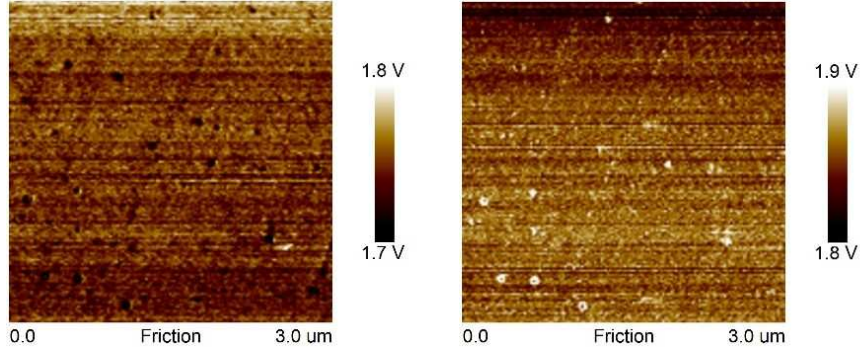
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**Figure S1.** Representative SEM and EDS results of gold coated tip (A, a), proteins functionalized tip (B, b), and proteins functionalized tip after being used in force measurement (C, c). Scale bar: 200 nm.



**Figure S2.** The representative lateral force trace (left) and retrace (right) maps of mPEG<sub>110</sub>-V-PMMA<sub>82</sub>-*b*-PFMA<sub>5</sub> V-shaped brush.

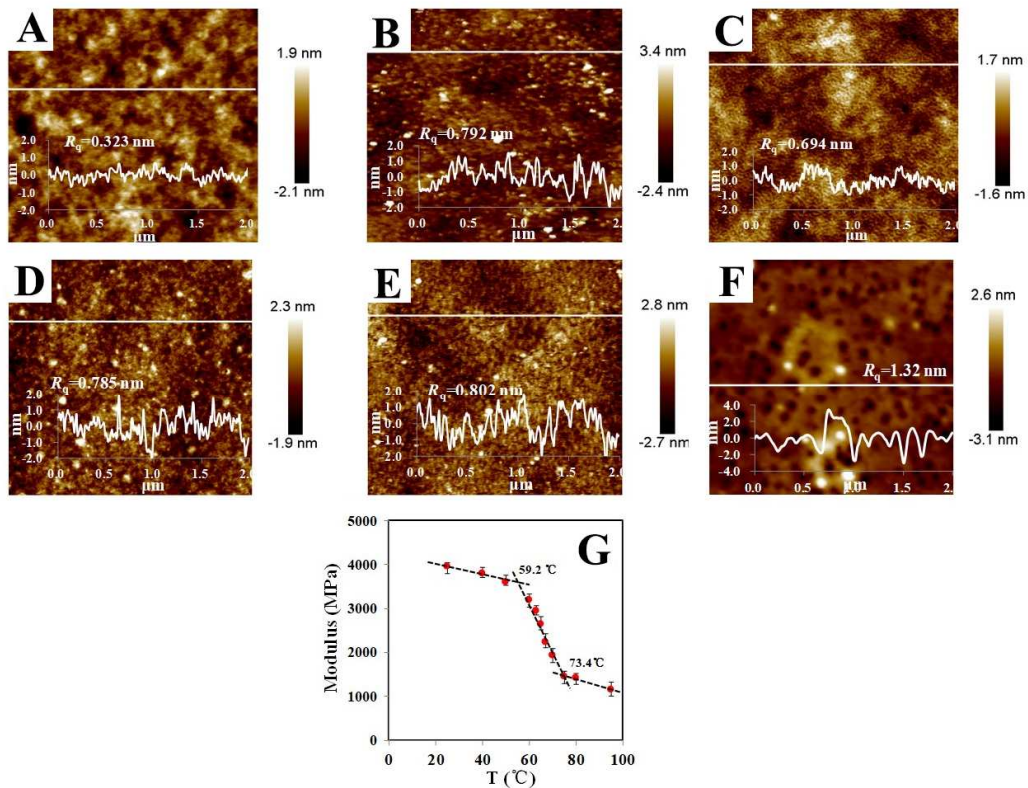
As we all know, friction coefficient ( $\mu$ ) can be calculated as the specific value of the friction force ( $F_{friction}$ ) and load force ( $F_{load}$ ). In the measurement of AFM,  $F_{friction}$  and  $F_{load}$  are converted from voltage, and  $\mu$  can be obtained through the relations as follows:

$$\mu = \frac{F_{friction}}{F_{load}} = \frac{k_1 V_{friction}}{k_2 V_{load}}$$

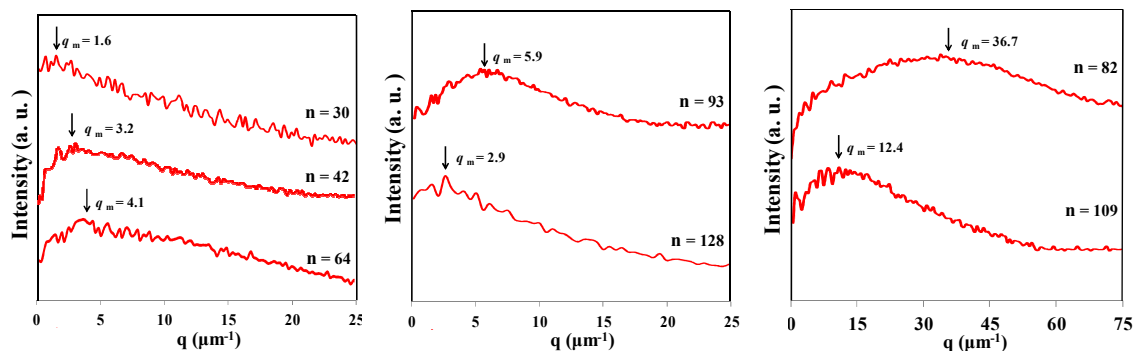
Where  $V_{friction}$  is friction induced voltage,  $V_{load}$  is load voltage (2 V was used in our work),  $k_1$  and  $k_2$  are two constants, which are related to lateral sensitivity and lateral force constant, normal sensitivity and normal force constant, respectively. Because Bruker MultiMode-8 instrument does not provide a way to calibrate lateral sensitivity and lateral force constant of a cantilever, the absolute value of friction coefficient can't be obtained. And the relative friction coefficient is calculated through the equation as follows:

$$\mu_{relative} = \frac{k_2}{k_1} \mu = \frac{V_{friction}}{V_{load}}$$

$V_{friction}$  is calculated as half of the difference value between the two friction images ( $V_{friction} = (V_{trace} - V_{retrace})/2$ ), as shown in [Figure S2](#). (According to the data of [Figure S2](#),  $V_{friction} = (1.885 - 1.705)/2 = 0.09$ ,  $\mu_{relative} = 0.045$ )



**Figure S3.** AFM height images (2.0 μm × 2.0 μm) of mPEG<sub>110</sub>-V-PMMA<sub>n</sub>-b-PFMA<sub>5</sub> brushes. (A-F) n = 30, 64, 82, 93, 109 and 128, respectively. (G) Modulus vs temperature for bright domain obtained by AFM PeakForce QNM mode.



**Figure S4.** Power spectrum obtained by circular average of FT AFM images of V-shaped brushes mPEG<sub>110</sub>-V-PMMA<sub>n</sub>-b-PFMA<sub>5</sub> in Figure 2.

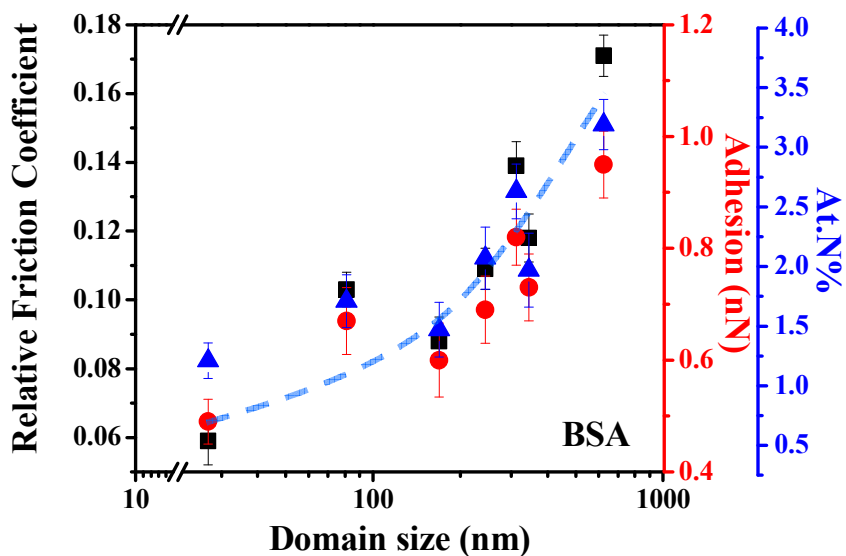
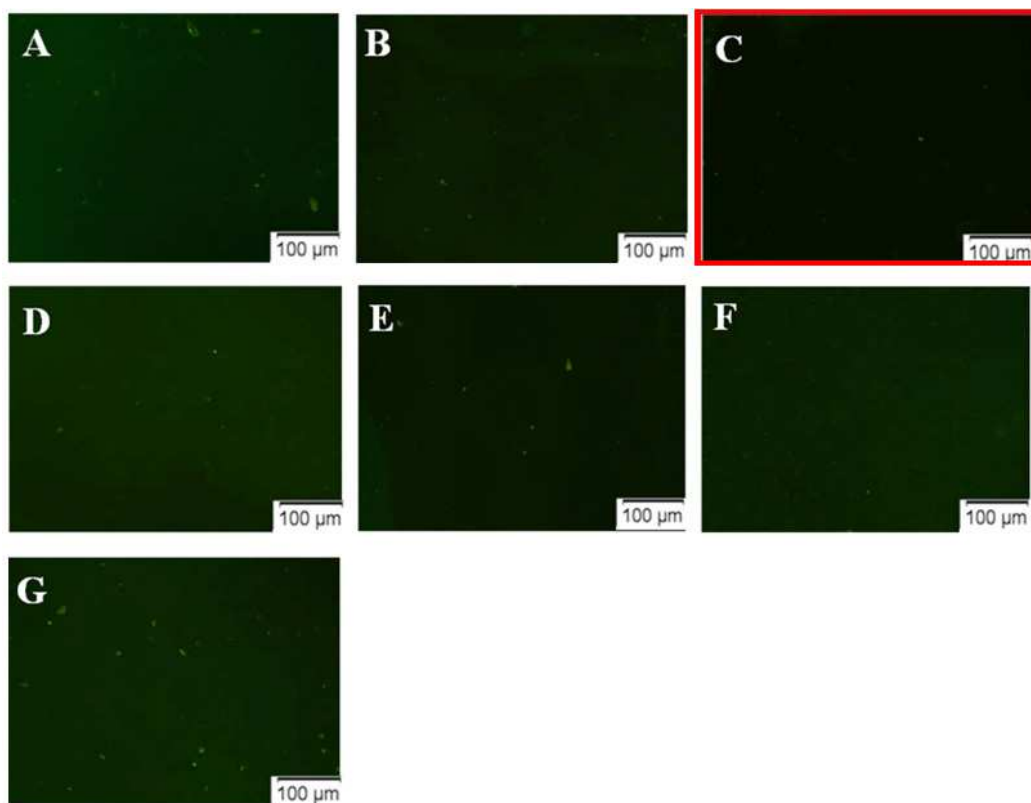
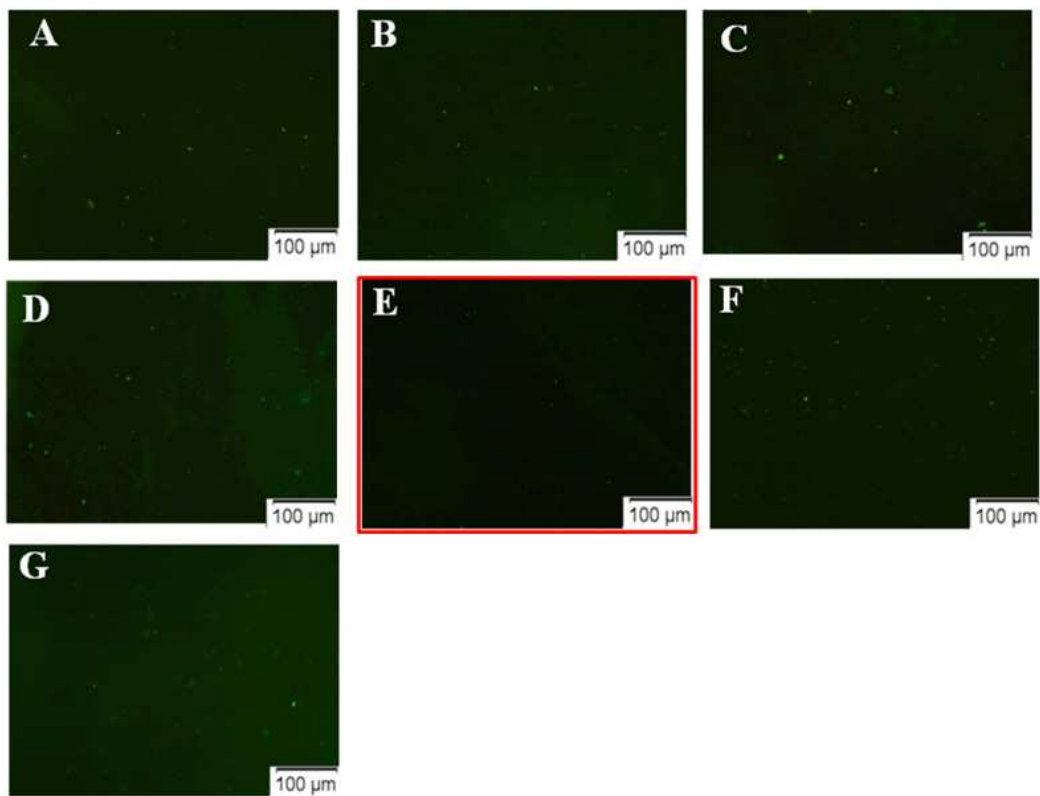


Figure S5. Atomic %  $N_{1s}$  of adsorbed protein, average measured adhesion force and the relative friction coefficient of BSA versus domain sizes in  $mPEG_{110}$ -V- $PMMA_n$ - $b$ - $PFMA_5$  brushes.



**Figure S6.** Fluorescence microscopy images of V-shaped polymer brushes with two arms of mPEG and PS after FITC-BSA adsorption. (A)  $mPEG_{42}$ -V- $PS_{43}$ , (B)  $mPEG_{42}$ -V- $PS_{78}$ , (C)  $mPEG_{42}$ -V- $PS_{128}$ , (D)  $mPEG_{110}$ -V- $PS_{61}$ , (E)  $mPEG_{110}$ -V- $PS_{119}$ , (F)  $mPEG_{110}$ -V- $PS_{169}$ , (G)  $mPEG_{110}$ -V- $PS_{227}$ .



**Figure S7.** Fluorescence microscopy images of V-shaped polymer brushes with two arms of mPEG and PS after FITC-HFg adsorption. (A) mPEG<sub>42</sub>-V-PS<sub>43</sub>, (B) mPEG<sub>42</sub>-V-PS<sub>78</sub>, (C) mPEG<sub>42</sub>-V-PS<sub>128</sub>, (D) mPEG<sub>110</sub>-V-PS<sub>61</sub>, (E) mPEG<sub>110</sub>-V-PS<sub>119</sub>, (F) mPEG<sub>110</sub>-V-PS<sub>169</sub>, (G) mPEG<sub>110</sub>-V-PS<sub>227</sub>.

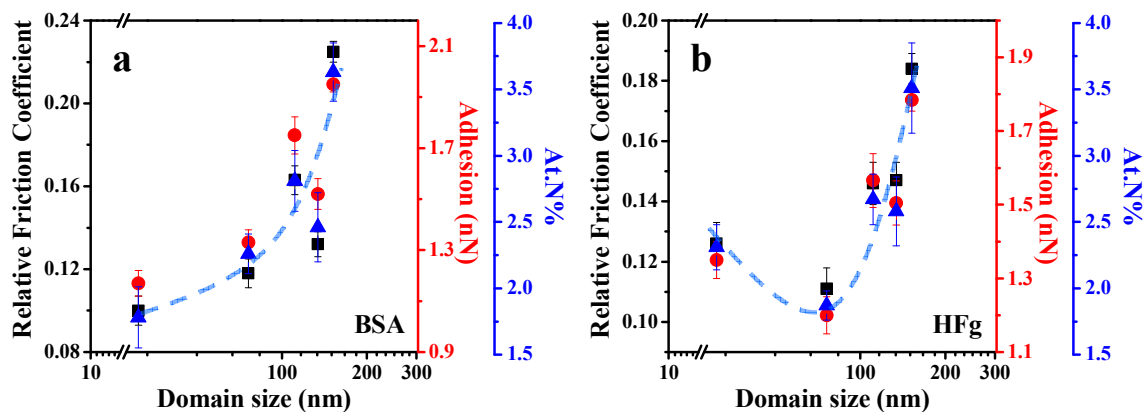
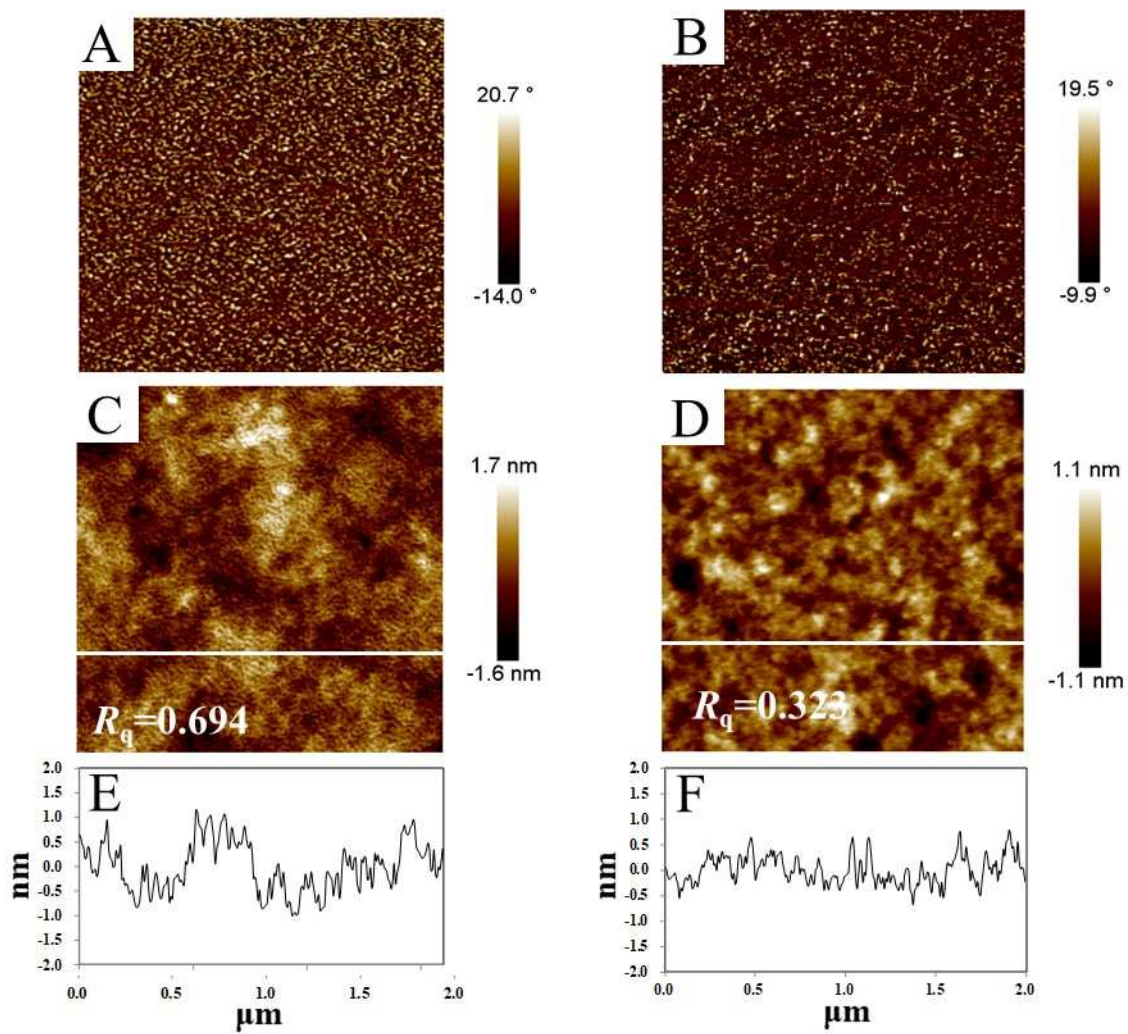
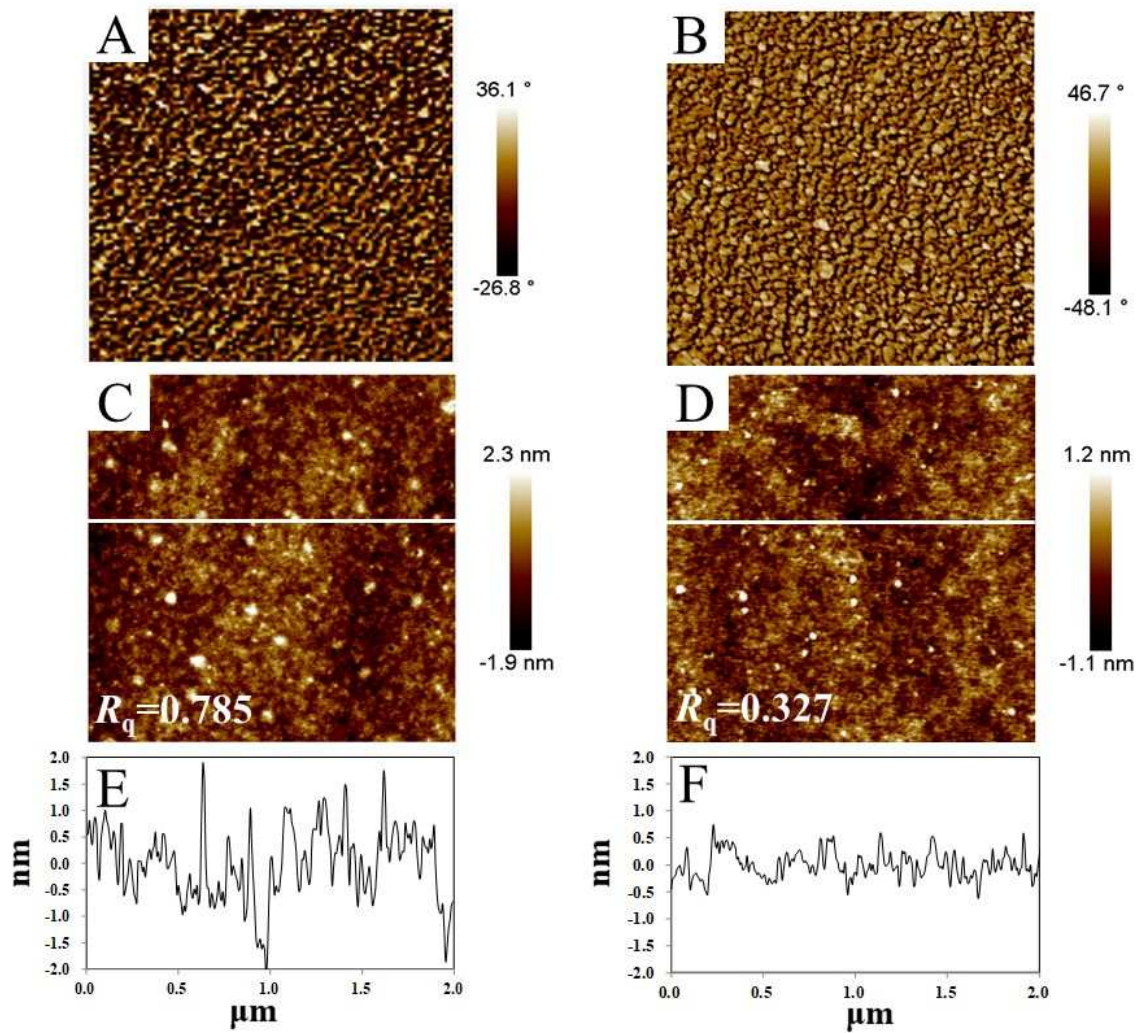


Figure S8. Atomic % N<sub>1s</sub> of adsorbed protein, average measured adhesion force and the relative friction coefficient of BSA (a) and HFg (b) versus domain sizes in mPEG-V-PS brushes.



**Figure S9.** AFM phase images, height images and height profiles of mPEG<sub>110</sub>-V-PMMA<sub>82</sub>-*b*-PFMA<sub>5</sub> brushes in air (A, C, E) and in PBS solutions (B, D, F), respectively.





**Figure S10.** AFM phase images, height images and height profiles of mPEG<sub>110</sub>-V-PMMA<sub>109</sub>-b-PFMA<sub>5</sub> brushes in air (A, C, E) and in PBS solutions (B, D, F), respectively.