An Examination of Peptide-Tether Binding to Gold and Glass Surfaces MASSACHUSETTS INSTITUTE

by

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Submitted to the Department of Mechanical Engineering In partial fulfillment of the requirements for the degree of

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Abstract

For many biological applications, it is beneficial to know that a peptide will bind to a surface. In this thesis, a tether, bead, peptide complex is constructed using the gold binding peptide (GBP) of sequence (CGGVSGSSPDS). Several assays and assay modifications are developed and tested to attempt to attach the GBP first to a gold slide and then to gold nanoparticles. Four peptides: FO₂, K1, K2, K3, known to bind to sapphire were attached to glass to see if it was possible to modify the sapphire assay to work with glass.

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Background

I. Peptide Binding

Knowing that a peptide from a natural amino acid binds to a surface provides information for biological applications, showing that protein and cell deposition can occur without additional chemical reactions [4]. Specific binding is also useful for producing biosensors and other self assembling components **[6,1].** Peptide display is a common method to determine whether or not a peptide can attach to a surface. Large groups of peptides are displayed on bacteria, bacteriophage, or beads, so that a single sequence of the amino acid can be picked for its binding capabilities. This process can be accomplished **by** creating a few new phages with different genes inserted into their genomes. The most desirable phage is selected for and amino acids are added to its library. Large amounts **(10x)** of the phage with the added amino acids are exposed to the binding surface and the ones that adhere are kept and analyzed **[2,3].** There were six metabolically stable gold binding peptides found **by** Brown, the peptide that contributed to the greatest fraction bound has the following sequence: (alPRGvYKIDSNmh)g.

Ionic bonds, hydrogen bonds, and hydrophobic interactions primarily contribute to peptidesurface binding **[1].** Studies with sapphire show that ionic bonding, and bonding between hydroxyl groups and metal ions (divalent metal ion bridging), allow attachment to occur. The strength of the peptide surface bond is determined **by** three factors: the number of charge interactions, the strength of the charge interaction, and the orientation of the charge groups relative to the surface [4]. The way peptides bind to the surface allows for binding specificity. **A** gold binding peptide (GBP) engineered to attach to inorganic surfaces, specifically gold, was shown to attach more readily to gold than sapphire. The streptavidin on the gold quantum dots bound to biotin (on GBP1) **[5].**

Peptides were attached to three different surfaces in this thesis: gold coated slides **(1"** x **3"** x 0.04" Cr/Au), gold nanoparticles, and glass. The gold binding peptide of sequence **(CGGVSGSSPDS)** was selected because of its strong affinity for gold. The peptide contains four serines that all have a hydroxyl groups on their side chains. Hydroxyl groups have been found

to bind well to gold surfaces **[3].** Four peptides that were proven to bind to sapphire were tested to see if they could also bind to glass **[1].** The four peptides chosen to test were F0² (KRHKQKTSRMGK), **K1 (GK) 6,** K2 (GGKK) ³, K3 (GGGKKK) ².

II. Construction of Tethers

It is important to choose a tether that will not interact with the surface and where one end of the tether only binds to the peptide and the other only to a bead. **DNA** was used as the tether in all experiments performed in this thesis. **DNA** is beneficial to use because it is easy to make the molecule any desired length and concentration using PCR. Problems sometimes arise when using **DNA** because the **DNA** backbone is negatively charged. **If** the peptide binds to a surface that is positively charged, the **DNA** will bind to the surface, eliminating its usefulness **[1].**

The beads attached to the tether were 0.8pm polystyrene beads that could easily be seen through a microscope. The goal of the each of the assays was to produce single molecule tethers. Two important considerations were taken into account when creating the assays. Primarily, a peptide attached to a tether should bind to the surface and a control tether without a peptide attached should not bind to the surface. Also, as the concentration of peptide tethers increased, the number of beads attached to the surface should increase as well **[1]. If** both of these conditions are met, the assay should be successful in creating a peptide/tether surface attachment.

Experimental Procedure - Gold Experimentation

I. Gold Slide Setup

Gold slides **1"** x **3"** X .040" Cr/Au were purchased from EMF. Four pieces of 3M Scotch **⁶⁶⁵** removable double sided tape was placed on the gold slide and a glass coverslip was placed over the tape (with the longer side perpendicular to the longer side of the gold slide), creating three channels. The gold slides could be used again after soaking them in soapy water for the night. The tape and coverslide were easy to remove after being soaked but the gold slides scratched easily. Care had to be taken at all times not to scratch the gold or the washes during the assay would not be as successful.

II. Tether Development

DNA tethers were constructed via PCR. **A 1010bp** dsDNA tether was made with a biotin molecule **(300** nm) on one of its **5'** ends and a digoxineginin on the other. The PCR was completed using a biotin conjugated forward primer **(5' -** biotin **- AAT CCG** CTT **TGC** TTC **TGA CT - 3')** and an amine conjugated reverse primer **(5' - NH2 - TTG AAA TAC CGA CCG TGT GA - 3')** on the M13mp18 plasmid. **All** of the PCR ingredients were added to a PCR tube and the appropriate program was run on the **DNA** engine. The PCR products were cleaned with the QiaQuick purification kit and the **DNA** concentration was measured using the Nanodrop.

The next process in the gold binding assay was to attach the **DNA** tether to the peptide through Sulfo-SMCC conjugation. The Sulfo-SMCC was attached to the **DNA** through the terminal amine group. After adding the **DNA** to the linker, the two were incubated for 2 hours at 4 degrees Celsius. The **DNA-SMCC** was cleaned using a Bio-Rad column and the product was spun at **¹⁰⁰⁰**x g for 4 minutes. Once the **DNA** and **SMCC** were attached, the peptide could be added because the peptide attaches to the **SMCC.** The gold peptide has a N-terminal cysteine which provides the necessary sulfhydryl group needed to attach the peptide to the **SMCC [1].** The DNA-SMCC-Peptide mixture was incubated overnight at 4 degrees Celsius. In the morning, the product was cleaned using a Bio-Rad Micro Bio Spin **30** column and the tether construction was complete.

Figure **2.** Tethered peptide attached to a surface.

Ill. Assay Development

The first goal of the assay development was to see if the beads alone attached to the gold slides. It is not desirable to have the beads attach to the slide, so the assay should be constructed in such a way that only the peptide attaches to the surface of the slide. Streptavidin coated 0.8pm beads were flowed through three channels. In the first channel, beads stored in PBST were flowed through. In the second channel, beads incubated in Casein, a blocking protein, were added. Beads coated with **200bp DNA** were flowed through the third

channel. It was beneficial to have all channels, including the control, located on the same slide because it made it easy to guarantee a comparison was done on the same surface of the slide. The microscope could be moved from one channel to the next over the tape without changing the position.

As expected, the channel with the beads in PBST had the greatest number of beads bound to the slide. The beads covered in **DNA** to prevent non specific adhesion had less beads bound to the slide, but not the least. Channel two, the beads covered with Casein, had the least number of beads bound to gold. The exact number of beads in each channel was impossible to determine because there were too many beads to count. This experiment provided good insight into the formation of the gold binding assay. Beads should be covered in Casein before flowing them through the channels.

Figure 3. Flow channel with three different types of beads

The gold binding assay was tried on a double channel gold slide. One of the channels was loaded with the gold peptide at a concentration of **30** ng/A. The other channel was loaded with the control **DNA, NH2-3500 bp.** The slides were incubated at room temperature in a humidity chamber for one hour. While the slides were incubating, the beads were prepared. $0.8\mu m$ streptavidin coated beads were spun **6** minutes at **10k** RPM with PBST. The process was repeated four times and each time the beads were washed in PBST, except for the final time, when they were suspended in casein. CBPST (Casein in PBST) was made and flowed through the cells to prevent the beads from binding to the surface. Another **30** minute incubation

followed the adding of the Casein. The beads were sonicated before they were loaded into the channels to prevent clumping. **A 30** minute incubation took place after the beads were added, and when time was up, a final wash through with CPBST was executed. Once washed, the slides were ready to examine.

IV. Results of the Gold Binding Assay

The first attempt at the gold binding assay was not successful. The channel with the gold peptide at a concentration of **30** ng/A contained a large amount of stuck beads. Hardly any moving beads were present at all. Because the beads were coated with casein, only a few stuck to the surface. The major contribution to the amount of beads present was due to the **DNA** tethers. The **DNA** ended up sticking to the slide and not the peptide, which made it look like the beads were stuck to the surface. The control looked similar to the gold peptide channel but had a larger quantity of stuck beads.

The assay was tried two more times. The first time the concentration of the gold peptide was reduced to 3 ng/ λ and the second time the concentration was reduced to 0.3 ng/ λ . In both cases, it was hard to tell which surface of the slide was being viewed while looking in the microscope, and the large presence of stuck beads still covered the slides in both the sample and control channels. The slides were also close to opaque making any observations hard to acquire. **A** final change was made to the assay to try to reduce the number of stuck beads. Instead of coating the beads in Casein, **PEG** was used to attempt to reduce the number of **DNA** tethers attached to the beads.

V. Gold Binding **Assay Modification**

PEG, polyethylene glycol, was used to coat the beads so that the desired amount of **DNA** would cover the beads. Coating the beads in **PEG** makes them much more resistant to **DNA** absorption allowing only a few tethers to attach to the beads as opposed to having the beads

completely covered **by** tethers. To cover the beads in **PEG,** the amine coated polystyrene beads were first spun at 6K rpm in water. The beads were then resuspended in **PEG** and incubated overnight. Before using the beads they were washed in water and spun for **10** minutes at 3K rpm (this process was repeated two more times).

Figure 4. DNA tethers attached to coated beads

To test the PEG assay, the beads and **PEG** were loaded onto a gold slide. Minimal improvements were accomplished with this test. There were still many beads stuck to the slide in all channels. Due to the fact that a promising gold slide assay was not yet developed and the difficulty surrounding the viewing of the slides with the microscope, further testing was abandoned. Another method for testing the gold binding peptide needed to be found. The next attempt was made using gold nanoparticles.

VI. Gold Nanoparticles

The motivation for using gold nanoparticles was to make the results more visible. Instead of using an almost opaque gold slide, a glass slide with small gold particles would be easy to view. The same gold binding peptide was added to attach to the gold. The gold nanoparticles came as a liquid and were stuck on the glass slides by non specific adhesion.

Two channels were constructed on the slide. **A** mixture of **5pL** of 0.8pm nanoparticles, 5pm of 30 ng/ λ gold peptide-DNA, using the same DNA tether as before, and 20_{PL} of PBST was added to the first channel. **5pL** of 0.8mm beads and **25pL** of PBST was added to the second. The slide was incubated for one hour and then **5pL** of 60mg/pL of **200bp DNA** was added to both channels, which acted as a blocking agent. After the **DNA** was added the slide was incubated for an additional **30** minutes. During incubation the beads were prepared **by** spinning and resuspending them in PBST. The beads were added after the completion of the incubation and then the slide was put back in the humidity chamber for an additional **30** minutes. Once complete, the final step was the wash with **50pL** of PBST.

When examining the slide after the assay was complete, the peptide did not appear to attach to the gold nanoparticles. The gold nanoparticles were visible with the microscope, but nothing else was visible on the slide. After testing with both the gold slides and the gold nanoparticles, it was not evident if the gold binding peptide ever attached to the gold. In case something went wrong in the construction of the tether, the **DNA** tethers were remade and one of the gold slide tests was completed again. The peptide chosen had attached to gold in several other studies, but even when the tether was rebuilt, the results were the same. The only objects that appeared on the slides were a large number of stuck beads. Modifications need to be made to the gold assay to make the beads less sticky. It was later found that the polymer coating on the gold nanoparticles blocked the binding of the peptide. Further testing could be done to see if the polymer coating could be removed or nanoparticles without the coating could be found. Due to the time constraints of this thesis, the direction was shifted to **look** at glass biding peptides.

Experimental Procedure - Glass Experimentation

I. Glass Binding **Peptides**

There were several peptides engineered **by** Krauland et. *al.* that bind to sapphire. These peptides, K1, K2, and K3 have sequences shown in table one **[1].** The goal of this

experimentation was to see if these sapphire binding peptides, along with F02 isolated using a yeast surface display, will also bind to glass.

Table 1. Peptides chosen to test on glass

II. Glass Assay

A standard glass slide with three flow cell channels was constructed **by** attaching an etched glass coverslip to a glass slide using double sided tape. Three samples were added to the slide: the control **(1** ng/A **NH2),** sample one **(10** ng/X TAMRA **F02),** and sample two **(1** ng/A TAMRA $FO₂$). Another slide with two channels was created for the remaining two samples: sample three (10 ng/ λ FO₂), and sample four (1 ng/ λ FO₂). All the samples contained the 3500 bp DNA tether, already attached to the peptide prior to loading. TAMRA is a protein dye that reacts with the amine terminus of proteins to form protein-dye conjugates. Once the samples were loaded, the slides were incubated at room temperature for two hours. 100μ L of 1 mg/ml Casein was next added to the slides and the slides were left for **30** minutes in a humidity chamber at room temperature. While the slides were incubating, **0.8** pm Streptavidin coated beads were prepared as they were in the gold binding assay. The final resuspension was done in 1 mg/ml Casein PBST right before the beads were loaded onto the slide. The beads were also sonicated at **30** seconds at **30%** prior to loading. The slides containing the beads were again left to incubate for **30** minutes in the humidity chamber. The final wash through was with Casein.

The same assay was used to create the slides with the K1, K2 and K3 samples. The 1 ng/ λ NH₂ control was used, and this time there were six other samples: **K1** at 1 ng/A and **10** ng/A, **K2** at 1 ng/A and **10** ng/A, and **K3** at 1 ng/A and **10** ng/A. The **3500 bp DNA** tether was once again attached to all the samples and the NH₂ before loading the samples onto the slide. The procedure for attachment can be found in Section II: Tether Development.

III. Results of **Glass Assay**

Each of the 12 slides were examined using a microscope and the number of tethers were counted **by** picking **10** random locations, counting the number of beads in the view **of** the microscope, and then taking an average. The height of the bars below corresponds to the number of tethers seen in the view of the microscope at one time using the camera view. The error bars were generated using a **95** percent confidence interval. The labels on the horizontal axis of the graph name the peptide.

Figure 5. Number of tethered peptides bound to glass

Overall, the **FO***2* slides contained several more tethers than the K1, K2, K3 slides. The TAMRA slides contained less tethers than the slides without TAMRA, showing that the dye somewhat restricted the binding of the peptide to the surface. The **K1,** K2, K3 peptides have a weaker binding affinity to glass, but because the peptides did bind to glass, further experimentation should be done to see what concentration and conditions would increase the number of peptides bound.

Conclusion

Similar assays were used to bind the GBP to gold and the five sapphire binding peptides to glass. Although the gold assay was not successful, it was useful to see the versatility of the assay, which was successful in binding the peptides to glass. **A** few more changes might make it possible to bind the GBP to gold nanoparticles. Future work could also be done on the glass peptides using the optical trap to pull on the tethers to get a distribution of pull forces.

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