

Interactomics: Protein Arrays and Label-Free Biosensors.

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Lecture-7.

Protein Immobilization for Protein-Protein Interaction Studies.

Welcome to MOOC NPTEL course on interactomics, today we will talk about protein-protein interactions study, immobilization of the ligand on sensor chip. Surface Plasmon resonance has transformed the study of bio-molecular interactions by delivering a platform that does not require the ligand or analyte to be labeled. SPR measures the interaction between a ligand which is immobilized on the sensor chip surface and an analyte which is passed in the solution. This measurement takes place in real time label-free environment providing kinetic, equilibrium and concentration data.

In today's SPR immobilization experiment, the ligand anti beta 2 microglobulin will be covalently immobilized to the surface of sensor chip using amine coupling chemistry. The direct immobilization of the ligand is known as direct coupled. Another way of immobilizing a bio-molecule is by using capturing chemistry. In this case, the ligand is not covalently immobilized to the sensor chip surface but is captured through the electrostatic interactions.

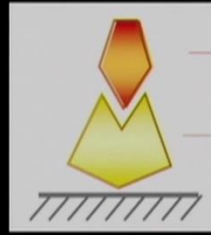
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Lecture Outline

- Immobilization of anti- β 2 microglobulin antibody on a sensor chip surface.
 - Activation— preparing for amine coupling
 - Immobilization
 - Deactivation
- Data analysis for immobilization.



Ligand and Analyte



- The analyte is the interaction partner that passes in solution over the surface.
- The ligand is the interaction partner attached to the surface.



The major steps involved in the immobilization of anti beta 2 microglobulin antibody will involve activation preparing for amine coupling immobilization and deactivation. So, let us have, the lab experimental session now. Let us learn a little about the basics of immobilization and SPR assay. In the molecular interaction study using surface plasmon resonance we will immobilize one of the interacting molecule that is anti-beta two microglobulin on the gold sensor chip surface while the protein beta two microglobulin will be passed over that surface in solution. Here, the ligand refers to the immobilized component and the interacting partner in the sample injected over the surface is referred to as the analyte.

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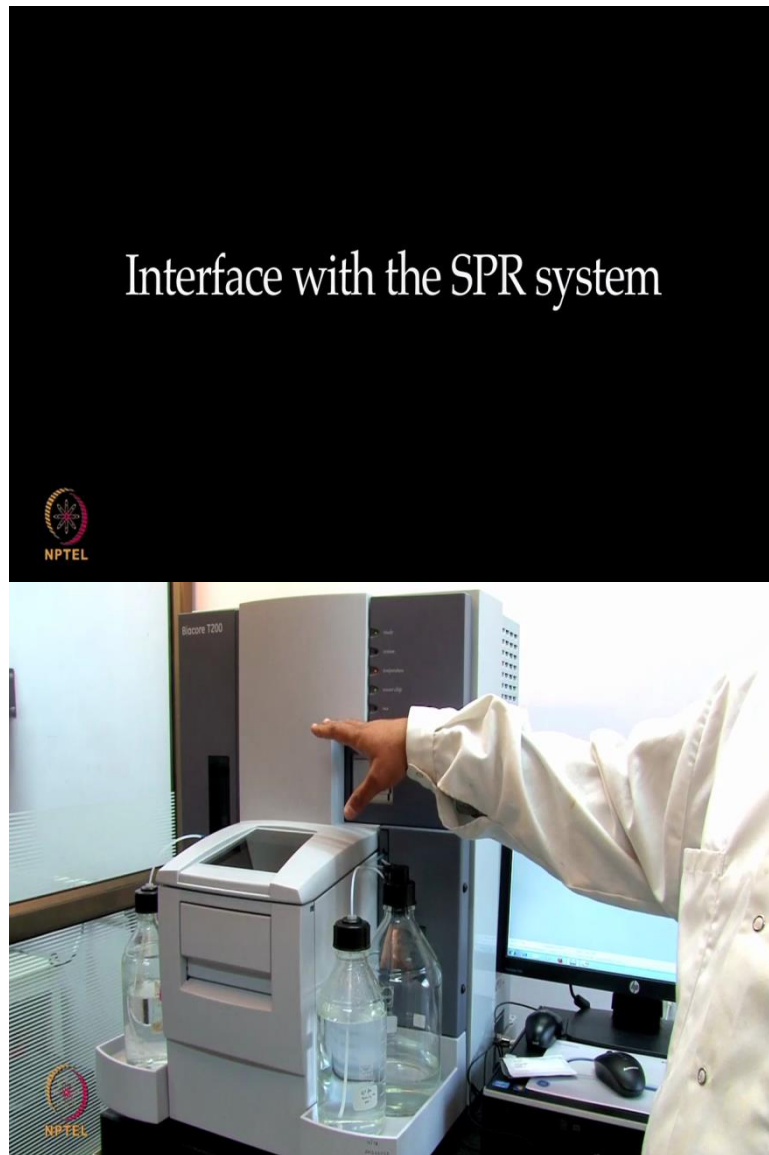
General Steps in a biacore SPR assay

Surface preparation

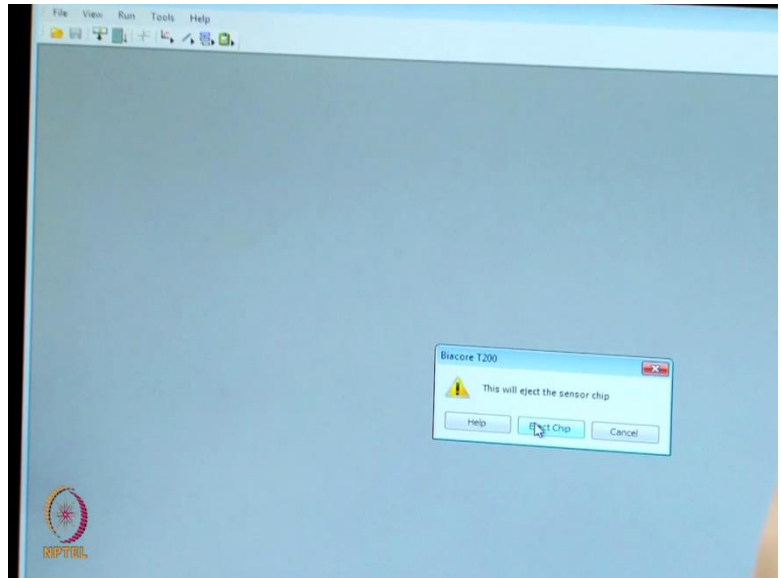


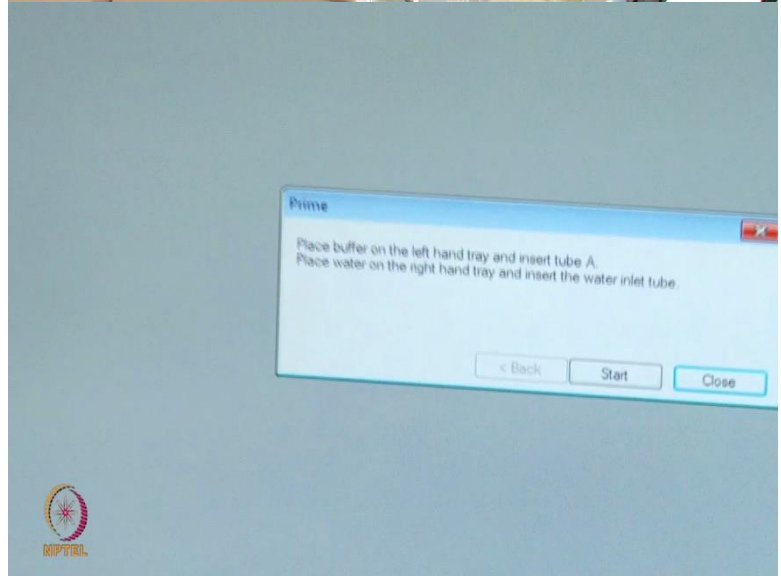
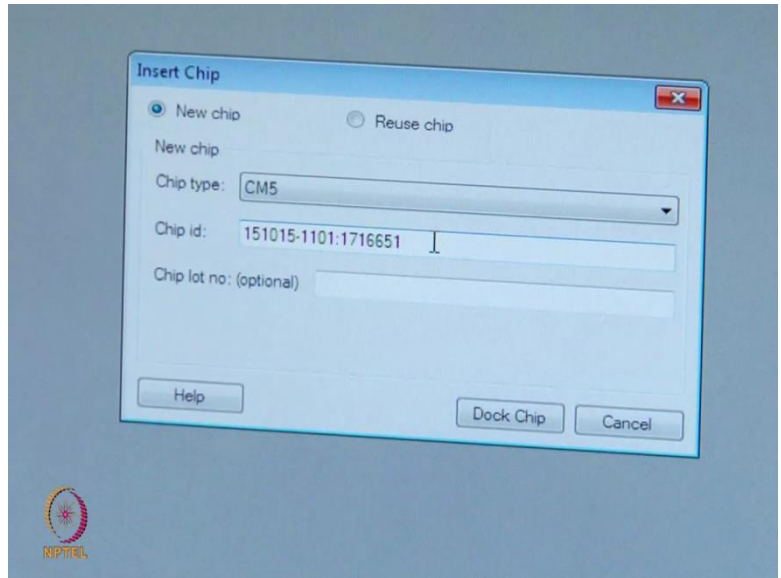
The three major steps in a biacore SPR assay involved; immobilization, the process by which ligand is attached to the sensor chip surface, interaction analysis where the analyte is injected over the sensor chip surface and the interaction between the analyte and immobilized ligand is monitored. And regeneration, the process of removing bound analyte from the ligand on the surface.

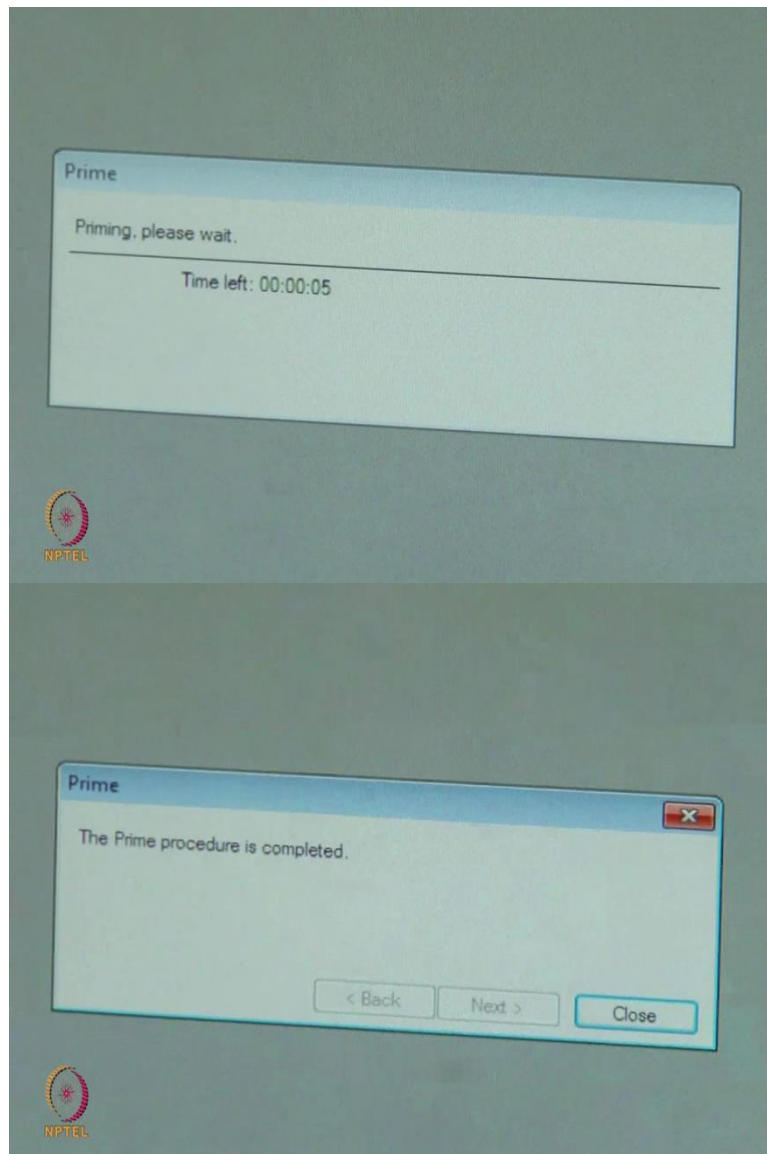
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So, this is a surface Plasmon resonance and it is a biacore T200. So, this machine has the various parts like the running buffer is connected here and any of the biological buffers can be connected as a running buffer. There could be have any pH from low to high. So, the regular buffers used in a biacore experiment are HBS EP-HBSN PBS for small molecule experiments.

Coming here, we have a water reservoir. The water reservoir is useful for cleaning needles and syringe and we have a waste reservoir here and this waste reservoir collects all the waste. The samples are actually sent in this compartment, chip is docked in this compartment. So, the experiment starts by picking up samples from this compartment and transferring them here at the interaction side and then the experiment is recorded on the screen. Here, some

kind of indications are provided when the machine is ready or the system is calibrating temperature, when the new chip is docked and the run is actually happening.

So, we will now, connect a new running buffer here and prime the system before our immobilization experiment. So, here the new HBS-EP plus buffer is connected and the new chip will be docked now. So, let us look at the chip now. So, this is a new chip, the chips are generally provided in these cassettes and then new CM5 chip will look like this. This chip will be inserted in chip docking area from the control software, we will eject to the sensor chip that is connected.

So, now the old chip will be removed and we will insert a new biacore chip CM5 here and the insertion or the way orientation of the chip is shown on the chip in arrows and we will close the compartment door and identify the chip from the chip type here and chip will be given a new id and sometimes it is very essential to add the name for the chip and also, the lot number and save dock chip.

Now, the chip is getting docked. So, we have connected a new buffer, we will prime the system. Priming is the process of sending buffer through the IRC and equilibrate the system before our experiment. Generally, in any biacore experiments, buffer should be connected and equilibrated or if there is no time, a minimum of three hours of equilibration is essential, otherwise when to start a new experiment do at least six primes on the system.

Now, that the chip is docked we will prime the system. Primly, we have connected the buffer. So, we just say start and it takes six minutes for the system to prime. Now, the prime procedure is complete, we will do an immobilization of a ligand. In this case today, we are actually immobilizing anti-beta two M antibody for that immobilization. Let us prepare a wizard, before we setup an immobilization protocol, let us understand a little about immobilization levels.

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Immobilization levels

- Binding capacity of the surface will depend on the levels of immobilized ligand.
- The term R_{max} refers to the binding capacity of the surface in terms of the response at saturation.

$$R_{max} = \frac{\text{analyte MW}}{\text{ligand MW}} \times R_L \times S_m$$

R_L : Immobilization level
 S_m : Stoichiometric ratio

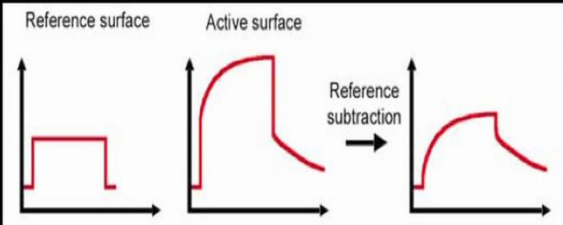
- Different applications require different immobilization levels.



The binding capacity of the chip surface will depend on the levels of immobilized ligand. The term maximum response refer to as R_{max} is described as the binding capacity of the surface in terms of the response at saturation. A theoretical R_{max} value can be calculated using the formula shown below where R_L is the immobilization level and the S_m is the stoichiometric ratio. A theoretical calculated R_{max} is often higher that the experimentally derived R_{max} for the same interaction. This could be because of several reasons such as the ligand is not fully active or that there is steric hindrance in the interaction.


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Reference surface



The bulk contribution can be subtracted by using a reference surface.

- Should be placed upstream of the active surface.
- Flow cells optimized for use in pairs (Fc1 + Fc2, Fc3 + Fc4)



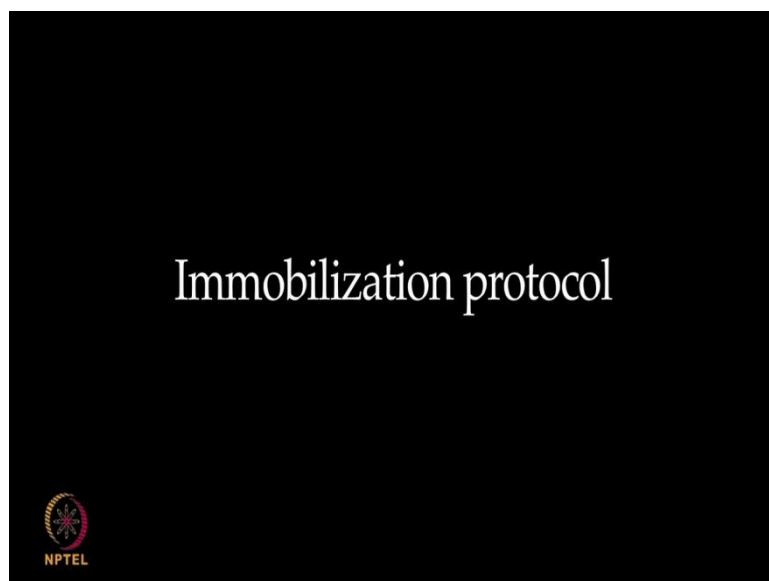
Different applications may require different binding capacities and thus, different immobilization levels. A low R_{max} is often beneficial in kinetics analysis while higher

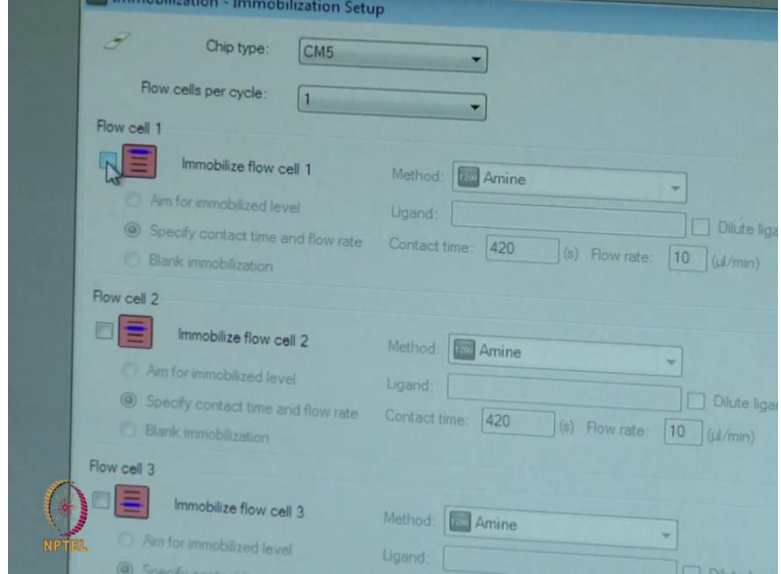
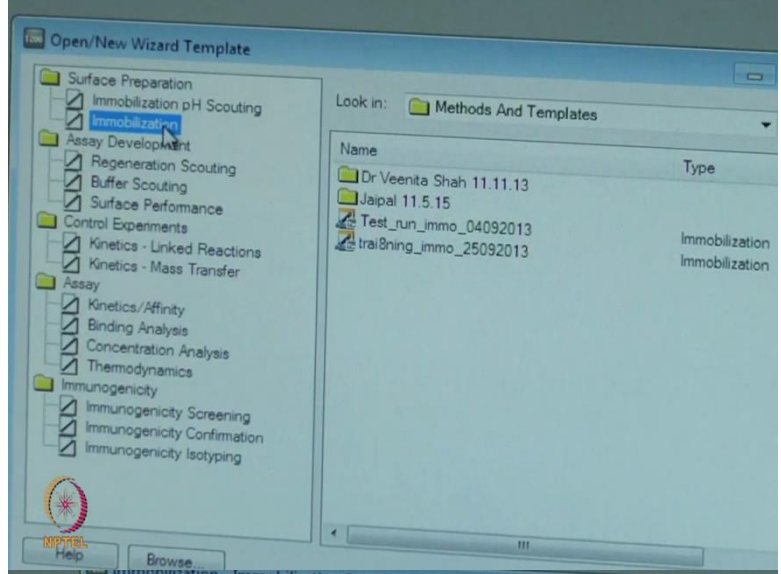
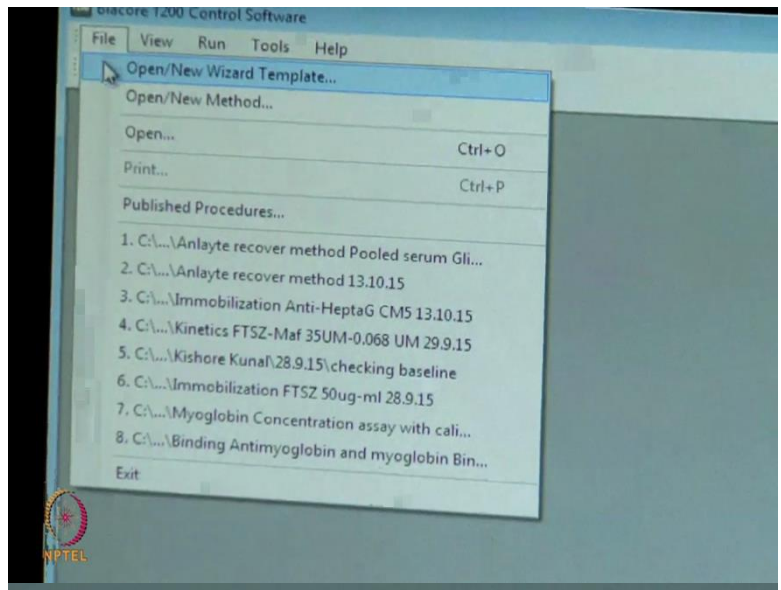
immobilization levels are advantageous in binding analysis and concentration measurements. Today, we are going to immobilize antibody on a CM5 chip using amine coupling chemistry and the figure here, shows a typical immobilization sensor gram using amine coupling.

The three major steps involved here are activation of the surface esters using EDC and NHS. Covalent coupling of the ligand on the sensor chip using amine groups of the ligand and deactivation of free esters with ethanol amine, we will analyse the results of anti-beta two microglobulin immobilization later in the lecture. Reference subtraction is particularly important for assays where measurement is taken during the sample injection.

The bulk contribution due to any difference in the sample matrix and running buffer can be subtracted by using a reference surface. This reference surface is typically placed upstream of the active surface. The flow cells on the chip surface are optimized accordingly for use in pairs that is flow cell one with flow cell two and flow cell three with flow cell four.

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Adjust contact time and flow rate
 Contact time: 420 (s) Flow rate: 10 (μl/min)

Immobilize flow cell 3
 Method: Amine
 Ligand: Dilute ligand
 Contact time: 420 (s) Flow rate: 10 (μl/min)

Immobilize flow cell 4
 Method: Amine
 Ligand: Dilute ligand
 Contact time: 420 (s) Flow rate: 10 (μl/min)



Immobilization - System Preparations

Prime before run
 Normalize detector

Temperature settings
 Analysis temperature: 25 (°C)
 Sample compartment temperature: 25 (°C)



Immobilization - Rack Positions

Position	Volume (μl)	Content
R2 B1	89	EDC
R2 B2	89	NHS
R2 B3	Empty	EDC/NHS, min. capacity 124μl
R2 B4	129	Ethanolamine
R2 C1	89	EDC
R2 C2	89	NHS
R2 C3	Empty	EDC/NHS, min. capacity 124μl
R2 C4	129	Ethanolamine
R2 C5	98	antib2m
R2 F6	120	BIA normalizing solution 70% (w/w) glycerol





File, open new wizard template, identify immobilization from surface preparation say new. So, we will immobilize for binding, flow cell one and two for binding and three and four for kinetics. Prime before run; normalize detector save next and the rack positions are displayed with the number of vials and the volume and the positions.

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Reagents Required

1. Ligand: Anti β 2microglobulin 30 μ g/ml (working concentration)
Stock concentration of ligand: 1mg/ml
2. Immobilization buffer: 10mM sodium acetate pH 5.0
3. Running buffer: HEPES-EP+ (10 mM HEPES, 150 mM NaCl, 3 mM EDTA, 0.05% (v/v) P20, pH 7.4)
4. N-ethyl-N-(dimethylaminopropyl)-carbodiimide (EDC) and N-hydroxysuccinimide (NHS) (1:1)
5. 1M Ethanolamine-HCl, pH 8.5

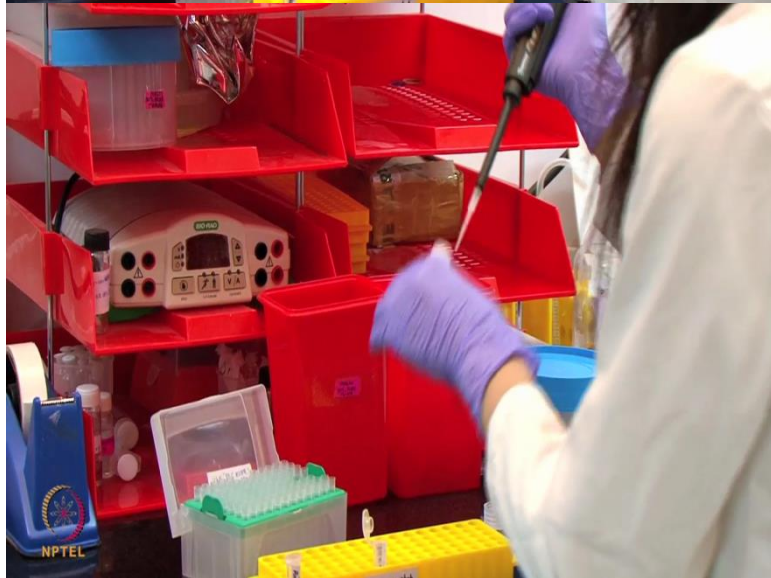


As was listed on the table, we will now have a closer look on the buffers and reagents required for immobilization of anti-beta two microglobulin . The reagents include a stock concentration of anti-beta two microglobulin from which a working concentration of 30 microgram per ml will be made using an immobilization buffer of 10 milimolar sodium acetate pH 5, we need HEPES-EP plus pH 7.4 which will include 10 milimolar HEPES, 150 milimolar NaCl, 3 milimolar EDTA and 0.05 percent P20. This will be used as the running buffer which is already connected to the system followed by priming of the system.

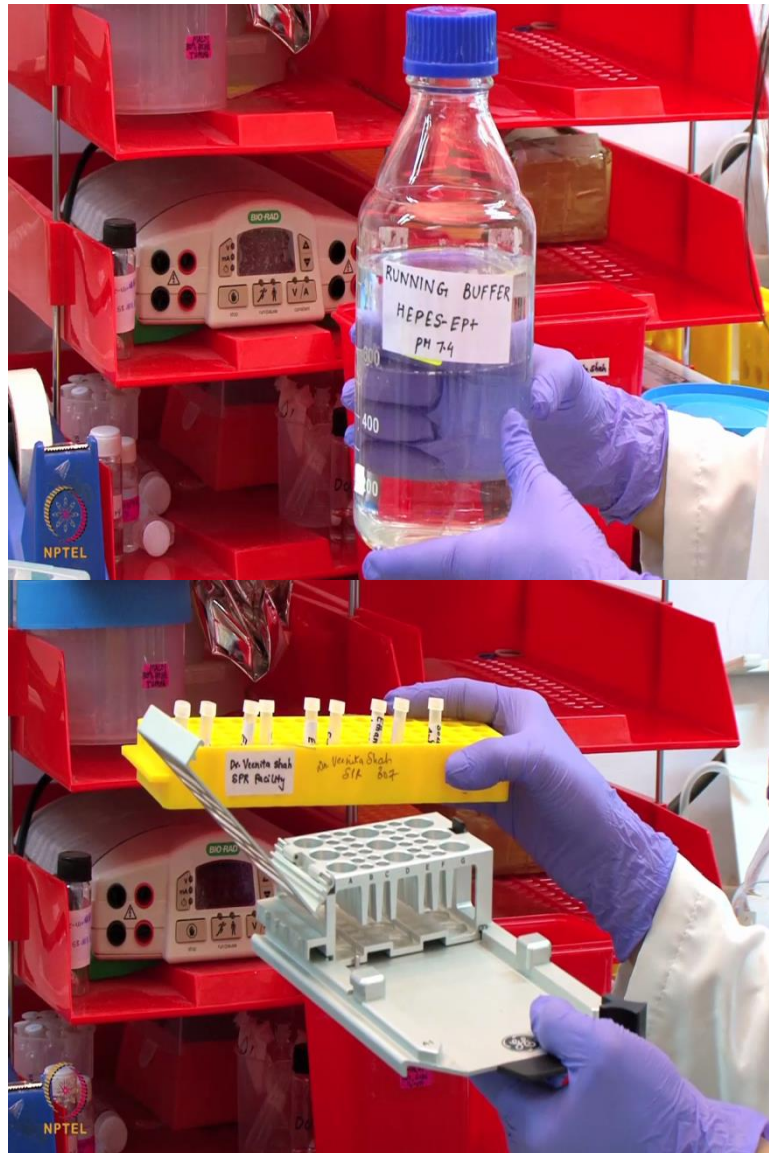
EDC and NHS in the amine coupling kit are used in one is to one ratio for surface activation. Lastly, we also require one molar ethanol amine HCl pH 8.5 for blocking the pre-ester groups on the surface. We shall now proceed to use the above mentioned reagents for our immobilization experiment. We will now work on the reagents required the immobilization of anti-beta two microglobulin on cellular chip surface.

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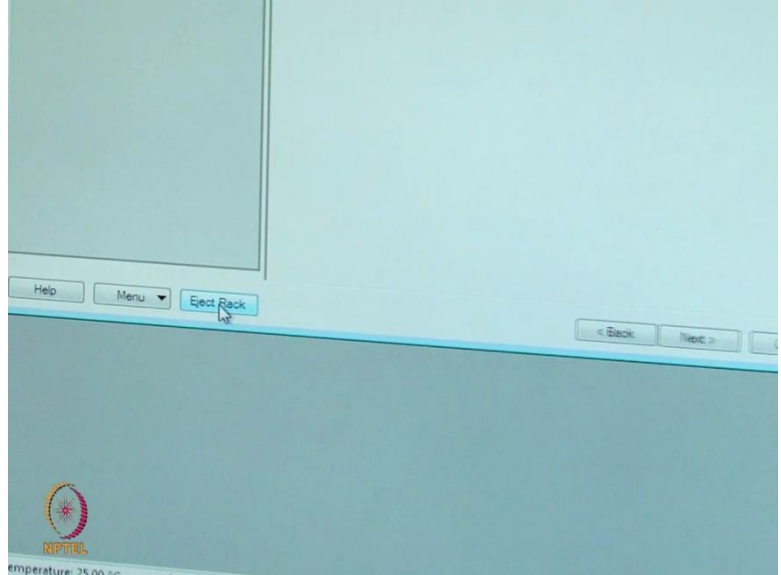
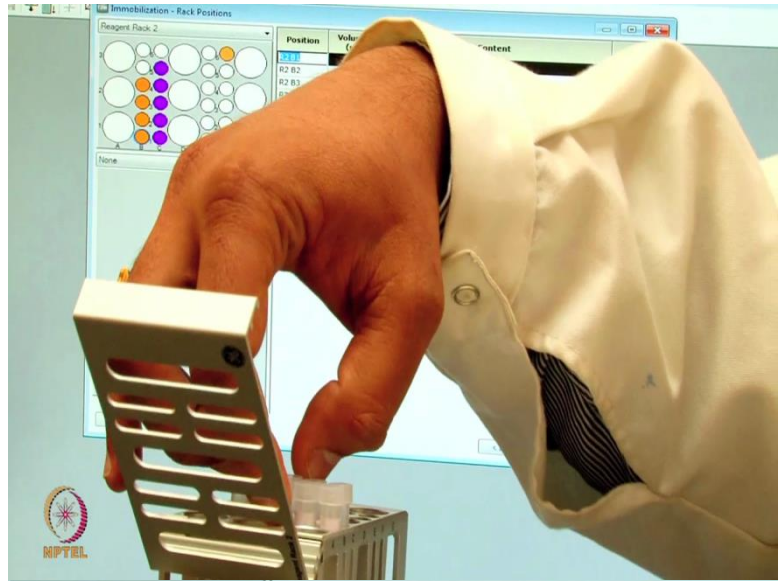


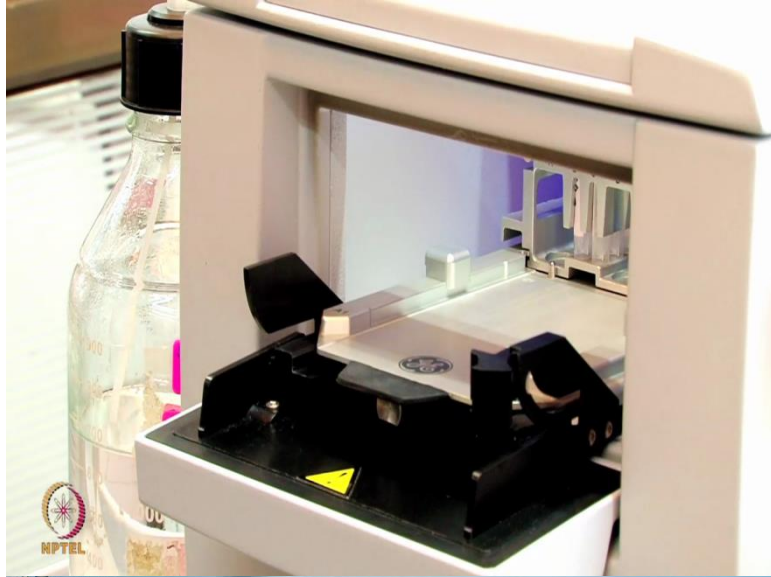




We will first tell you the stock concentration of the antibody which is 1 mg per ml of anti-beta two microglobulin . In 10 milimolar sodium acetate pH 5 to make a working antibody solution of 30 microgram per ml for this we will take 6 microliter of the ligand stock and mix it in 194 microliter of sodium acetate pH5. The choice of the correct immobilization buffer is an important parameter to consider and the pH scouting feature of the system will help in choosing the correct pH of the immobilization buffer. This is our 30 microgram per ml of ligand pH.

(())(17:47) of NHS, EDC and ethanol amine are prepared and transferred to this specialized tubes used for the system. We have now transferred all the reagents into this specialized tube. So, we have two NHS tube, Two EDC, two empty tubes for longer stability of EDC, NHS which will the makes inside the system two ethanol tube, one for the blank flow cell and the other for the active flow channel and one ligand solution. We will be using HEPES-EP plus





Immobilization - Rack Positions

Reagent Rack 2

Position	Volume (μl)	Content	Type
R2 B1	89	EDC	
R2 B2	89	NHS	Immob Fc 1
R2 B3	Empty	EDC/NHS, min. capacity 124μl	Immob Fc 1
R2 B4	129	Ethanolamine	Immob Fc 1
R2 C1	89	EDC	Immob Fc 2
R2 C2	89	NHS	Immob Fc 2
R2 C3	Empty	EDC/NHS, min. capacity 124μl	Immob Fc 2
R2 C4	129	Ethanolamine	Immob Fc 2
R2 C5	98	antiB2m	Immob Fc 2
R2 E6	130	antimalaria solution 70% (w/w) glycerol	Normalize

Eject Rack Tray

Rack Tray Ejected

Click OK to return the rack tray to the sample compartment.

Help OK

Time to auto close: 22

Help Menu Eject Rack

< Back Next > Close

R2 B2	89	NHS	Immob Fc 1
R2 B3	Empty	EDC/NHS, min. capacity 124µl	Immob Fc 1
R2 B4	129	Ethanolamine	Immob Fc 1
R2 C1	89	EDC	Immob Fc 1
R2 C2	89	NHS	Immob Fc 2
R2 C3	Empty	EDC/NHS, min. capacity 124µl	Immob Fc 2
R2 C4	129	Ethanolamine	Immob Fc 2
R2 C5	98	antib2m	Immob Fc 2
R2 F6	120	BIA normalizing solution 70% (w/w) glycerol	Normalize

Biacore T200 Control Software

Inserting rack: 7

- Make sure the correct sensor chip is docked
- Make sure all samples & reagents are loaded in the rack and microplate according to the Rack Positions setup. (Vials should be sealed with rubber caps and microplate with adhesive foil.)
- Place the buffer(s) on the left hand tray and insert the correct tubing(s). see below.
- Note! Standby after run will use buffer A.
- Make sure there is fresh water in the water bottle on the right hand tray.
- If necessary, empty the waste bottle before start of the run.

Estimated run time: 1 h 3 min (excluding standby)

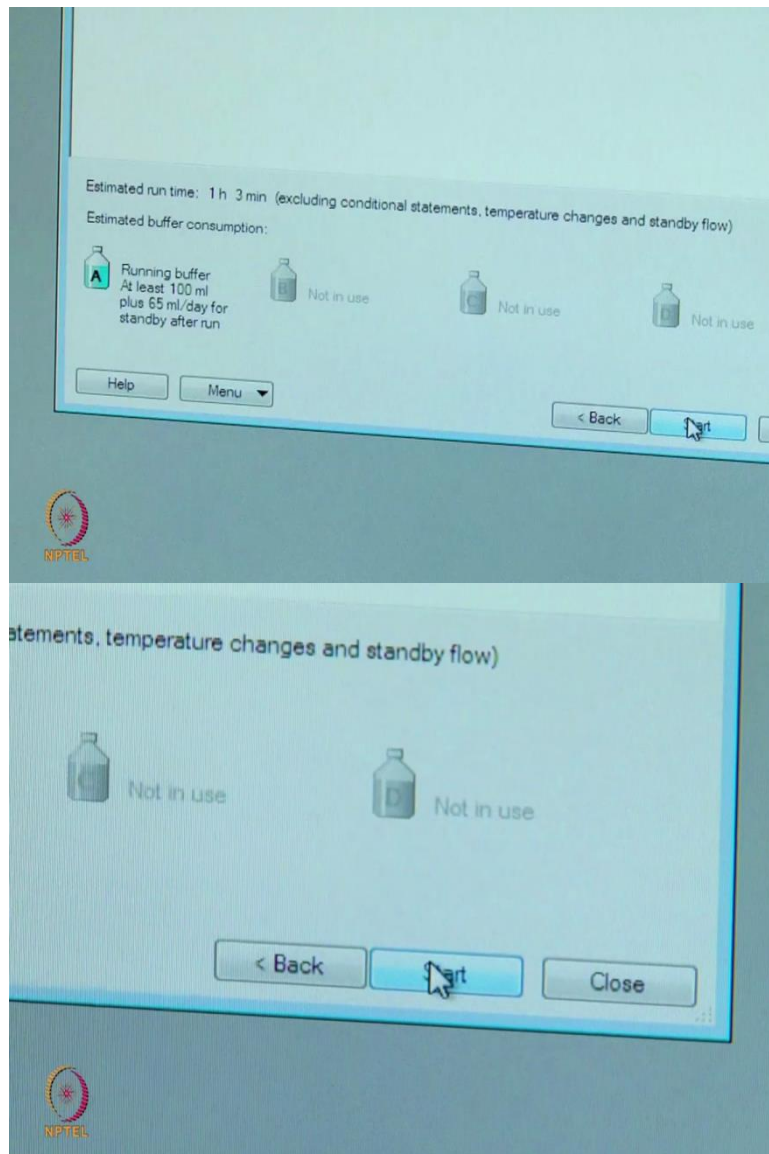
Estimated buffer consumption:

A Running buffer
At least 100 ml
plus 85 ml/day for
standby after run

Tahoma 10 **B I U**

Prepare Run Protocol

- Make sure the correct sensor chip is docked
- Make sure all samples & reagents are loaded in the rack and microplate according to the Rack Positions setup. (Vials should be sealed with rubber caps and microplate with adhesive foil.)
- Place the buffer(s) on the left hand tray and insert the correct tubing(s). see below.
- Note! Standby after run will use buffer A.
- Make sure there is fresh water in the water bottle on the right hand tray.
- If necessary, empty the waste bottle before start of the run.



We will now proceed to insert these tubes into the appropriate rack and then into this system for immobilization of anti-beta two microglobulin on the sensor chip surface. Now, the we made the template, we will insert the vial in the right position, we just fitted them in the required volumes of different chemicals to start with the EDC is positioned here, the NHS is positioned here, the ethanol amine is positioned here and there is empty vial , in a similar way another row is also made. So, one for the reference surface and other for the active surface to normalize the chip, normalization solution for the machine, to normalize the discrepancies on the RU responses on four different close channel.

We close the door this way and go on the screen here, we eject the rack, we will insert the rack into the sample compartment this way and need to lock it inside, lock it inside and then go on the screen, close the compartment. We go to the next tab. So here there some points we need to take a before start of the run. Make sure the correct sensor chip is docked. Make


sure all sample reagents are loaded in the rack and micro plate according to the rack position setup, vials should be sealed with caps, place the buffer, place water and make sure there is sufficient amount of water and buffer.

So, once we go to all these and everything is set in the machine, it shows the estimated run time as 1 hour 3 minutes and we have connected the running buffer. Now, we will start the experiment. This will prepare the chip for the immobilization process and immobilize the antibody or the ligand of our choice today the anti-beta two M. We need to save this as and now, the immobilization process is running.


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Points to ponder

- Ligand is the interacting partner attached to the surface and analyte is the one that passes in solution over the surface.
- R_{max} refers to the binding capacity of the surface in terms of the response at saturation.
- Different applications require different binding capacities on the surface, and thus different immobilization levels.
- Reference subtraction is very critical for an accurate data interpretation.



Data analysis for immobilization
of anti- β 2 microglobulin



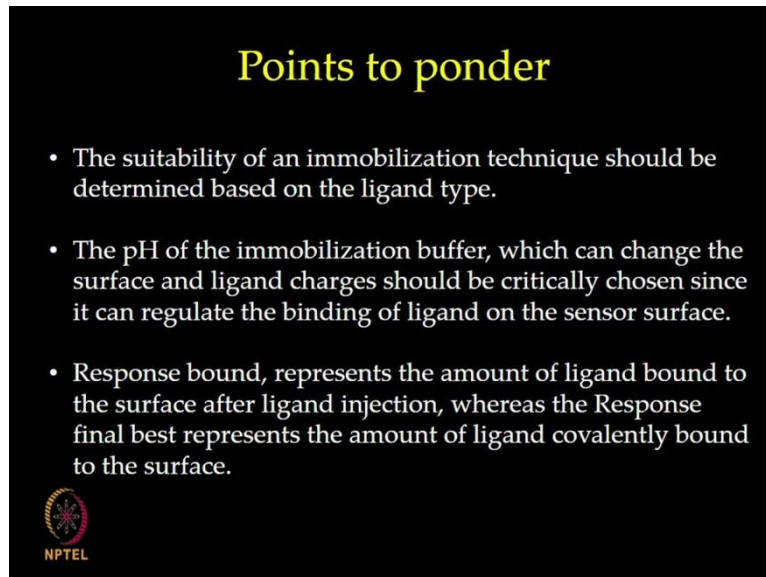


We will now look at the results of our anti-beta two microglobulin immobilization performed on CM5 chip. Looking at the immobilization results dialog box, there are two response levels calculated from the sensor gram. The response bound which represents, the amount of ligand bound to the surface after ligand injection whereas the response final test represents the amount of ligand covalently bound to the surface.

So, here as we observe 12894 RU of anti-beta two microglobulin is immobilized on the surface of the chip. Now, analyzing the immobilization sensor gram, we observe the baseline here followed by EDC-NHS activation of the dextran matrix which is again followed by baseline after activation of the surface. This is followed by covalent coupling of the ligand to the dextran matrix. The buffer washes away the loosely associated ligand molecules. Deactivation and further washing away of loosely associated ligand happens and the


difference in response between these two points reflect the amount of anti-beta two microglobulin immobilized.

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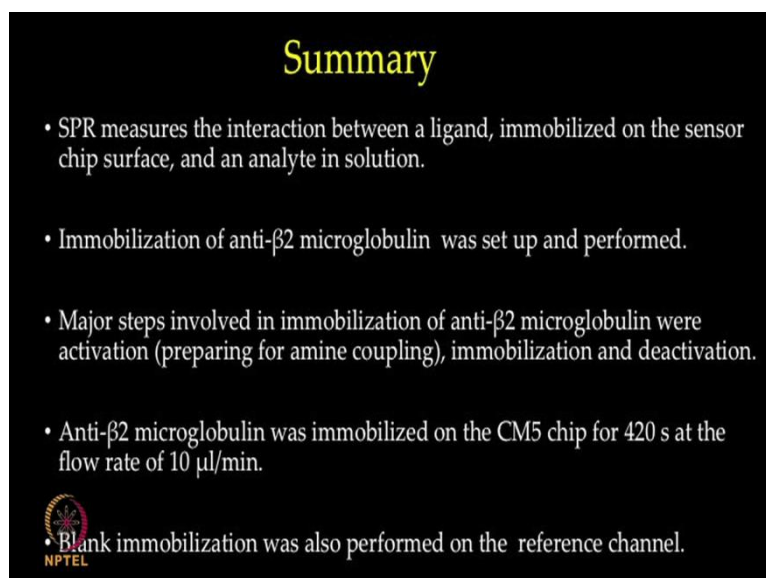
Points to ponder

- The suitability of an immobilization technique should be determined based on the ligand type.
- The pH of the immobilization buffer, which can change the surface and ligand charges should be critically chosen since it can regulate the binding of ligand on the sensor surface.
- Response bound, represents the amount of ligand bound to the surface after ligand injection, whereas the Response final best represents the amount of ligand covalently bound to the surface.




So, as we observed we have successfully immobilized anti-beta two microglobulin and we will now proceed for our binding experiment in our next lecture. Performing an interaction analysis on active and stable ligand surface is key to generate robust data set. We have witnessed this procedure for immobilizing anti-beta two microglobulin antibody on CM5 sensor chip surface. In next lecture, we will talk about protein-protein interactions study binding analysis. Thank you.

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Summary

- SPR measures the interaction between a ligand, immobilized on the sensor chip surface, and an analyte in solution.
- Immobilization of anti- β 2 microglobulin was set up and performed.
- Major steps involved in immobilization of anti- β 2 microglobulin were activation (preparing for amine coupling), immobilization and deactivation.
- Anti- β 2 microglobulin was immobilized on the CM5 chip for 420 s at the flow rate of 10 μ l/min.



- Blank immobilization was also performed on the reference channel.

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