# Sortase

## Week 9

Summarized below are the experiments conducted this week in chronological order. Click on the experiment name to view it. To go back to this summary, click **Summary** in the footer.

# Summary

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# 1 Digestion and Ligation of pSB1C3-Gb1-SortA

## Responsible

Aman Mebrahtu

#### Protocols used

- Digestion and Ligation

## Modifications and comments to protocols

The amount of enzyme used was adjusted to the concentration of the DNA sample.

# Experimental Set Up

The experimental set up for the digestions are stated in the tables below. A reaction volume of  $10 \mu l$  was used for both of the digestions.

Table 1: Concentrations of DNA to be digested

| Sample | Concentration $[ng/\mu l]$ |
|--------|----------------------------|
| pSB1C3 | 41.0                       |

Table 2: Digestion reaction of pSB1C3

| Reaction         | Volume $[\mu l]$ |
|------------------|------------------|
| DNA              | 5                |
| Buffer SH $10 X$ | 1.0              |
| Pst1             | 0.2              |
| EcoR1            | 0.2              |
| Sterile water    | 3.6              |

In this ligation, we decided to go forward with an insert:vector molar ratio of 10:1. The calculations shown on the tables below were adjusted so that each reaction contains 50 ng of backbone and the inserts are adjusted according to the desired molar ratio. Since the concentrations of the purified DNA fragments were relatively low, this resulted in an overload of reaction volume which is indicated by a minus sign for sterile water.

Table 3: Ligation Reaction of pSB1C3-Gb1-SortA

| 8                        |        |
|--------------------------|--------|
| Component                | Volume |
| Gb1-SortA                | 12.89  |
| pSB1C3                   | 4.17   |
| T4 DNA Ligase            | 0.5    |
| T4 DNA Ligase Buffer 10X | 2.5    |
| Sterile water            | 4.95   |



# Results and Conclusions

The result of the ligation is to be confirmed with transformation.



# 2 Transformation of pSB1C3-GB1-SortA

## Responsible

Reskandi Rudjito and Maren Maanja

## Protocols used

- Transformation

# Experimental Set Up

The ligation was performed overnight at a temperature of  $16^{\circ}\mathrm{C}$  was later transformed into both TOP 10 and BL21 cells.

Table 4: Re-Transformation of pSB1C3-T7-Lys

|                             | ı v                               |
|-----------------------------|-----------------------------------|
| Sample                      | Comment                           |
| pSB1C3-GB1-SortA Top10      | Transformed into Top10 cells      |
| pSB1C3-GB1-SortA BL21 (DE3) | Transformed into BL21 (DE3) cells |
| positive control Top 10     | No antibiotics                    |
| positive control BL21       | No antibiotics                    |

## **Results and Conclusions**

No colonies were observed indicating that the previous ligation procedure had been unsuccessful.

# 3 Digestion of Gb1-SortA

## Responsible

Reskandi Rudjito and Maren Maanja

#### Protocols used

- Digestion

# Experimental Set Up

Table 5: Concentrations of DNA to be digested

| Sample    | Concentration $[ng/\mu l]$ |
|-----------|----------------------------|
| Gb1-SortA | 21.9                       |

Table 6: Digestion of Gb1-SortA

| Table 0. Digestion of GSI Soluti |  |  |
|----------------------------------|--|--|
| Volume $[\mu l]$                 |  |  |
| 8                                |  |  |
| 2.0                              |  |  |
| 0.2                              |  |  |
| 0.2                              |  |  |
| 1.6                              |  |  |
|                                  |  |  |

The digested products were subjected to purification and to maximize the amount of DNA eluted, we performed re-elution of DNA from the purification columns.

## **Results and Conclusions**

The result of the digested Gb1-SortA is shown in the figure below (lane 5). The expected size of the digested fragment is 669 bp.

Table 7: Concentrations of digested products after purification

| Sample                  | Concentration $[ng/\mu l]$ |
|-------------------------|----------------------------|
| Gb1-SortA               | 4.0                        |
| Gb1-SortA (re-ellution) | 2.0                        |

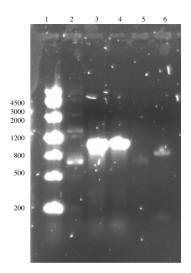


Figure 1: Digested pSB1C3. (1) Ladder (2)-(4) Not of interest (5) Digested GS (6) Not of interest

# 4 Ligation and Transformation of pSB1C3-Gb1-SortA

## Responsible

Reskandi Rudjito

## Protocols used

- Ligation
- Transformation

# Experimental Set Up

In order to perform the ligation, the two DNA fragments consisting of the Gb1-SortA and pSB1C3 backbone are digested and purified prior to ligation. The concentrations of each DNA fragment are stated in Table 8.

Table 8: Concentrations DNA fragments for Ligation

| Sample            | Concentration $[ng/\mu l]$ | Size [bp] |
|-------------------|----------------------------|-----------|
| Digested Gb-SortA | 4.0                        | 669       |
| Digested pSB1C3   | 5.9                        | 2200      |

In this ligation, we have decided to go forward with an insert:vector molar ratio of 10:1. The calculations shown on the tables below are adjusted so that each reaction contains 50 ng of backbone and the inserts are adjusted according to the desired molar ratio.

Table 9: Ligation Reaction of pSB1C3-Gb1-SortA

| Component                | Volume |
|--------------------------|--------|
| Gb1-SortA                | 17.76  |
| pSB1C3                   | 4.24   |
| T4 DNA Ligase            | 0.5    |
| T4 DNA Ligase Buffer 10X | 2.5    |
| Sterile water            | 0      |

The ligation was performed overnight at a temperature of 16  $^{\circ}\mathrm{C}$  was later transformed into both TOP 10 and BL21 cells.

Table 10: Re-Transformation of pSB1C3-T7-Lys

| Sample                      | Comment                           |  |
|-----------------------------|-----------------------------------|--|
| pSB1C3-GB1-SortA Top10      | Transformed into Top10 cells      |  |
| pSB1C3-GB1-SortA BL21 (DE3) | Transformed into BL21 (DE3) cells |  |
| positive control Top 10     | No antibiotics                    |  |
| positive control BL21       | No antibiotics                    |  |

## **Results and Conclusions**

The ligation product was subjected to gel electrophoresis, as shown in the figure below.



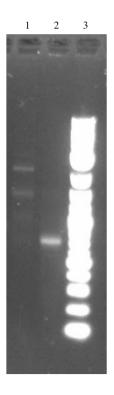


Figure 2: Digested pSB1C3. (1) Ligated pSB1-C3-Gb1-SortA (2) Digested pSB1C3 (3) Ladder

# Discussion and Troubleshoot

Lane 1 from Figure 2 shows that there are several bands is the upper part of gel which could possible indicate ligated plasmids.

# 5 Transformation of pSB1C3-Gb1-SortA

## Responsible

Aman Mebrahtu and Reskandi Rudjito

#### Protocols used

- Transformation

# Experimental Set Up

Transformation was done on both TOP 10 cells and BL21 cells to improve efficiency.

Table 11: Re-Transformation of pSB1C3-T7-Lys

| Sample                      | Comment                           |
|-----------------------------|-----------------------------------|
| pSB1C3-GB1-SortA Top10      | Transformed into Top10 cells      |
| pSB1C3-GB1-SortA BL21 (DE3) | Transformed into BL21 (DE3) cells |
| positive control Top 10     | No antibiotics                    |
| positive control BL21       | No antibiotics                    |

## **Results and Conclusions**

Transformation was unsuccessful.

## Discussions and Troubleshoot

We have been getting negative result in our transformation procedures. Thus, to come around this problem, we will try to purify the ligation product before transformation and also try electro-competent cells instead of chemically competent cells.

# 6 Re-Transformation of pSB1C3-Gb1-SortA using purified ligation product and electro-compotent cells

## Responsible

Reskandi Rudjito and Aman Mebrahtu

## Protocols used

- PCR Purification
- Transformation

# Experimental Set Up

In this experiment we tried both chemically competent and electro-competent cells with purified ligation products.

Table 12: Re-Transformation of pSB1C3-T7-Lys

| Sample                  | Comment   |
|-------------------------|---|
| pSB1C3-GB1-SortA        | Transformed into Top10 cells                        |
| pSB1C3-GB1-SortA        | Transformed into BL21 (DE3) cells                   |
| pSB1C3-GB1-SortA        | Transformed into electro-competent BL21 (DE3) cells |
| positive control Top 10 | No antibiotics                                      |
| positive control BL21   | No antibiotics                                      |

## Results and Conclusions

As a result we saw several positively transformed colonies.

Table 13: Result of Transformation

| Table 13. Result of Transformation |                    |  |
|------------------------------------|--------------------|--|
| Sample                             | Number of colonies |  |
| Chemical TOP 10 cells              | 0                  |  |
| Chemical BL21 (DE3) cells          | 2                  |  |
| Electro-competent BL21 (DE3) cells | 1                  |  |

# 7 Colony picking and Colony PCR of Transformed colonies containing pSB1C3-Gb1-SortA

## Responsible

Reskandi Rudjito and Aman Mebrahtu

#### Protocols used

- Colony picking
- PCR

# Experimental Set Up

The PCR was done with verification primers (VF2 and VR). Table 8 shows the composition used for the PCR and Table 9 shows the PCR conditions used in the thermocycler. The PCR reaction used for each sample was 25  $\mu$ l.

| Table 14: PCR Reaction |                          |                  |  |  |  |
|------------------------|--------------------------|------------------|--|--|--|
| Master Mix             | Component                | Volume $[\mu l]$ |  |  |  |
| 1                      | PCR Grade Nucleotide mix | 1                |  |  |  |
|                        | R_Primer (VR)            | 5                |  |  |  |
|                        | F_primer (VF2)           | 5                |  |  |  |
|                        | Template DNA             | 2                |  |  |  |
|                        | Sterile water            | 14               |  |  |  |
|                        | Total Volume             | 25               |  |  |  |
| 2                      | PCR Reaction buffer 10 x | 5                |  |  |  |
|                        | Taq DNA Polymerase       | 0.5              |  |  |  |
|                        | Sterile water            | 19.5             |  |  |  |
|                        | Total Volume             | 25               |  |  |  |

Table 15: PCR condition using Taq polymerase

| Step                 | Cycles       | Temperature {°C} | Time     |
|----------------------|--------------|------------------|----------|
| Initial Denaturation | 1            | 94               | 2 min    |
| Denaturation         | 30           | 94               | 30  secs |
| Annealing            |              | 52               | 1 min    |
| Extension            |              | 72               | 1 min    |
| Final Extension      | 1            | 72               | 7 min    |
| Hold                 | indefinitely | 4                | _        |

## **Results and Conclusions**

The result of the gel electrophoresis is shown below. The amplicon containing Gb1-SortA should be 941 bp in size.



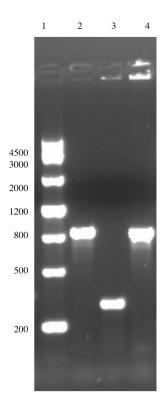


Figure 3: Colony PCR of Gb1-Sortase. (1) Ladder (2) Colony 1 (3) Colony 2 (4) Colony 3 - Electroporation

# Discussions and Troubleshoot

The result of the colony PCR shows that colony 1 and 3 contains the right insert. This is confirmed by the fact that the colony PCR show bands of above 800 bp and the expected size is around 941 bp. Colony 2 however most likely contained a contaminating sequence that was successfully cloned into the backbone.

# 8 Glycerol Stocks and Plasmid Extraction of pSB1C3-Gb1-SortA

## Responsible

Reskandi Rudjito and Aman Mebrahtu

## Protocols used

- Glycerol stocks
- Plasmid extraction

## Experimental Set Up

Colony 1 and 3 containing the pSB1C3-Gb1-SortA was cultured in LB containing chloram-phenicol. Samples of the two cultures were saved as glycerol stocks and stored in the -80°C freezer. The liquid culture was also used for plasmid extraction, where the extracted plasmid was subjected to sequence analysis.

## **Results and Conclusions**

The concentrations of the extracted plasmids are summarized below.

| Table 16: Concentration of Plasmid |                            |  |  |  |
|------------------------------------|----------------------------|--|--|--|
| Sample                             | Concentration $[ng/\mu l]$ |  |  |  |
| pSB1C3-Gb1-SortA (Colony 1)        | 322.1                      |  |  |  |
| pSB1C3-Gb1-SortA (Colony 3)        | 180.4                      |  |  |  |