Week 5: 6. July 2015 - 10. July 2015

6. July 2015:

- 1) Digest of mmoX, mmoY, mmoZ, mmoD, mmoG, phi with Ndel/EcoRI
 - · Pipetting scheme according to the protocol Restriction Digest
 - First adding Ndel and incubate the Reaction at 37 °C for 1 hour, afterwards heat inactivation of Ndel at 65 °C for 2 minutes. Adding 1 ul of EcoRl and incubate the digest again for 1 hour at 37 °C
 - Verify 10 μl of the digestion on agarose gel

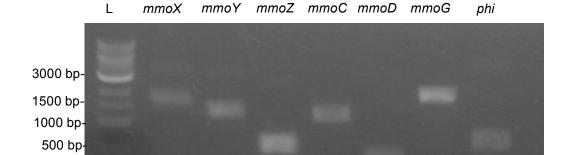


Figure 1: Digest of mmoX, mmoY, mmoZ, mmoD, mmoG and phi with Ndel/ EcoRl. 10 μl of digest were checked on 1% (w/v) agarose gel. Expected sizes: mmoX- 1605 bp, mmoY- 1191 bp, mmoZ- 546 bp, mmoC-1082 bp, mmoD- 333 bp, mmoG- 1702 bp and phi- 585 bp. As ladder (L) 1kB Ladder (NEB) was used.

- 2) Purification of digested mmoX, mmoY, mmoZ, mmoC, mmoD, mmoG, phi
 - Using Wizard® SV Gel and PCR Clean-Up System (Promega) and following the provided manual
- 3) Dephosphorylation of opened pET-30 using shrimp alkaline phosphatase
 - Pipetting scheme according to the protocol Shrimp Alkaline Phosphatase Treatment (Fermentas).
- 4) Ligation of mmoX, mmoY, mmoZ, mmoC, mmoD, mmoG, phi into pET-30
 - Pipetting scheme according to the Protocol Ligation with T4 DNA Ligase (NEB)
 - Use dephosphorylated pET-30
 - Control Reaction: pET-30 Selfligation
 - Incubate the reaction at 16 °C overnight

7. July 2015

- 1) Transformation of *E. coli* TOP10 cells with pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoD, pET-30+mmoG, pET-30+phi
 - Add 5 µl of ligation reaction to chemically competent *E. coli* TOP10 cells.
 - Following the protocol for transformation of chemical competent *E. coli* cells
 - Plate on LB+Kan [50 µg/ml] and incubate overnight at 37 °C
- 2) Invitro mutagenesis of hps to remove Xbal Restriction Site
 - Use QuickChange Site-Directed Mutagenesis Kit (Agilent) and follow the provided manual
 - Primer: hps mut1/mut2
 - Template: pCR4+hps
 - a) Dpnl Treatment
 - Add 1 µl of DpnI to the PCR Reaction and incubate at 37 °C for 1 hour
 - b) Transformation of E. coli TOP10 cells with DpnI treated PCR reaction
 - Add 1 µl of Dpnl treated PCR reaction to chemically competent E. coli TOP10 cells
 - Following the Protocol for Transformation of chemical competent E. coli cells
 - Plate on LB+Amp [100 µg/ml] and incubate overnight at 37 °C

8. July 2015

- The ligation was successful, control reaction with selfligated pET-30 showed less clones compared to the ligation reactions
- 1) Colony-PCR to screen for clones containing the pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoC, pET-30+mmoD, pET-30+mmoG, pET-30+phi
 - Constructs: pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoD, pET-30+mmoG, pET-30+phi
 - Pipetting scheme and PCR program according to PCR with Taq-DNA Polymerase Protocol
 - Primer: mmoX_E1/E2, mmoY_E1/E2, mmoZ_E1/E2, mmoC_E1/E2, mmoD_E1/E2, mmoG_E1/E2, phi_E1/E2
 - Check 6 Clones per Construct

- Positive Control: add 1 µl of pSC1B3+mmoX, pSC1B3+mmoY, pSC1B3+mmoZ, pCR4+mmoC, pSC1B3+mmoD, PCR fragment mmoG, pSC1B3+phi
- Negative Control: add 1 µl MilliQ Water

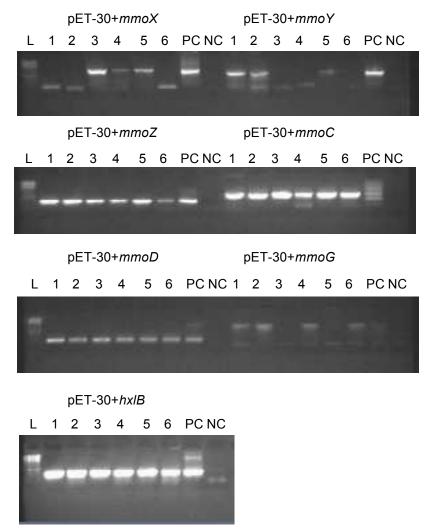


Figure 2: Colony-PCR to identify clones containing pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoD, pET-30+mmoD, pET-30+mmoG, pET-30+phi expression constructs. Numbers 1-6 determine the checked clone. As positive control (PC) 1 μl pSC1B3+mmoX, pSC1B3+mmoY, pSC1B3+mmoZ, pCR4+mmoC, pSC1B3+mmoD, PCR fragment mmoG, pSC1B3+phi was used. As negative control (NC) 1 μl MilliQ Water was added. 10 μl of PCR were analzyed on 1 % (w/v) agarose gel. Expected sizes: mmoX-453 bp, mmoY-1197 bp, mmoZ-552 bp, mmoC-1088 bp, mmoD-339 bp, mmoG-1708 bp, phi-591 bp. As ladder (L) 1 kB Ladder (NEB) was used.

- 2) <u>Inoculation of liquid culture for plasmid isolation of pET-30+mmoX</u>, <u>pET-30+mmoY</u>, <u>pET-30+mmoD</u>, <u>pET-30+mmoD</u>, <u>pET-30+mmoD</u>, <u>pET-30+phi</u>
 - Inoculate one clone with 5 ml LB+Kan [50 μg/ml] and incubate overnight at 37 °C shaking at 220 rpm.

- 3) Inoculation of liquid culture for plasmid isolation of pCR4+hps-mut Xbal
 - Inoculate one clone with 5 ml LB+Amp [100 μg/ml] and incubate overnight at 37 °C shaking at 220 rpm.
- 4) Amplification of mmoG, medh2 and phi for BioBrick System
 - Pipetting scheme and PCR program according to PCR with Phusion-HF DNA Polymerase Protocol
 - Primer: mmoG-BioBrick Fwd/Rev, medh2-BioBrick Fwd/Rev, phi-BioBrick Fwd/Rev
 - Template: PCR Fragment mmoG for TOPO cloning, pCR4+medh2, pCR4+phi

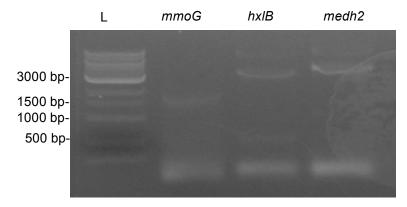


Figure 3: Amplification of *mmoG, phi* and *medh2* **for BioBrick System**. 20 μl of PCR were checked on 1% (w/v) agarose gel. Expected sizes: *mmoG*-1708 bp, *phi*-591 bp, *medh2*-1203 bp. As ladder (L) 1kB Ladder (NEB) was used.

9. July 2015

- 1) Plasmid Isolation of pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoD, pET-30+mmoG, pET-30+phi
 - Use the QIAprep Spin Miniprep Kit (Qiagen) and follow the provided manual.
 - Send the plasmids for sequencing with T7-FWD primer
 - Expression constructs seem to be correct
- 2) Plasmid Isolation of pCR4+hps-mut Xbal
 - Use the QIAprep Spin Miniprep Kit (Qiagen) and follow the provided manual.
 - Send for Sequencing
 - Mutagenesis was successful, Xbal restriction site was successfully mutat



Figure 4: Alignment of *hps* original sequence with sequencing results of *hps* invitro mutagensis to remove Xbal restriction site. Red colour indicates the Xbal restriction site in the hps sequence. Cyan labeled A in the *hps* sequence indicates the base we wanted to change into a C.

3) Transformation of *E. coli* BL21 cells with expression plasmids

- Expression Plasmids: pET-30+mmoX, pET-30+mmoY, pET-30+mmoZ, pET-30+mmoB, pET-30+mmoC, pET-30+mmoD, pET-30+mmoG, pET-30+phi
- Add 1 μl of Expression Plasmid to chemically competent E. coli BL21 cells.
- Following the protocol for transformation of chemical competent E. coli cells
- Plate on LB+Kan [50 μg/ml] and incubate overnight at 37 °C