

iGEM 2015 – Microbiology – BMB – SDU

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| Project type: Cloning | | Creation date: |
| Project title: Control with Leucin Zipper | | Written by: ADK, TBA and JSP |
| | | Performed by: ADK, TBA, EMT, AC and JSP |

1. SOPs in use

iGEM2014_SOP0010_v01_Phusion PCR

iGEM2014_SOP0017_v01_ Fast digest

iGEM2014_SOP015_v02_ligation

iGEM2013_SOP0021_v01_ Colony PCR with MyTaq

iGEM2013_SOP0009_v01_TSB transformation

2. Purpose

To conjugate a leucin zipper to the T18 and T25 domain as used in the bacterial two-hybrid system and to furthermore verify that the bacterial two-hybrid system works.

3. Overview

| Date (DD.MM.YY) | Person(s) (initials) | Experiments | SOPs |
|----------------------------|---------------------------------|--|----------------------------------|
| 28.07.15 | JSP | Phusion PCR, amplification of Leucine Zipper gene. | iGEM2014_SOP0010_v01_Phusion PCR |
| 29.07.15 | JSP | FastDigest | iGEM2014_SOP0017_v01_Fast digest |
| 29.07.15 | ADK | Phusion PCR, amplification of Leucine Zipper gene. | iGEM2014_SOP0010_v01_Phusion PCR |

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| 30.07.15 | JSP | FastDigest (Leucin Zipper) | iGEM2014_SOP0017_v01_ Fast digest |
| 30.07.15 | JSP | FastDigest (backbones) | iGEM2014_SOP0017_v01_ Fast digest |
| 31.07.15 | ADK | Miniprep of (Top10) pSB1C3- UT18-LeuZ, and (Top10) pSB1K3-KT25-LeuZ | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 31.07.15 | ADK | Fast digest of pcr product leuZ (G23) | iGEM2014_SOP0010_v01_ Phusion PCR |
| 31.07.15 | ADK | Fast digest of R63 (pSB1C3- UT18-LeuZ) and R64 (pSB1K3-KT25-LeuZ), with BamH1 and Pst1 to cut of the wrong LeuZ | iGEM2014_SOP0010_v01_ Phusion PCR |
| 31.07.15 | TBA, JSP | Ligation of Leucine zipper into backbones: pSB1K3-RFP-T25 pSB1C3-RFP-T18 pSB1C3- T18 pSB1K3-T25 | iGEM2014_SOP015_v02 _ligation |
| 31.07.15 | JSP | Phusion PCR, amplification of Leucine Zipper gene. | iGEM2014_SOP0010_v01_ Phusion PCR |
| 01.08.15 | ADK | Transformation of pSB1K3- RFP-T25(Y57) pSB1C3-RFP-T18 (Y58) pSB1C3-T18 (Y59), pSB1K3- T25 (Y60), ligated with leuzine zipper (Y63) into <i>E. coli</i> Top10 | iGEM2013_SOP0009_v0 1_TSB transformation |
| 03.08.15 | ADK | Colony PCR with MyTaq of first draw of Y59 and Y58 | iGEM2013_SOP0021_v0 1_ Colony PCR with MyTaq |
| 04.08.15 | ADK | Colony PCR with MyTaq of first draw of Y59 and Y58 | iGEM2013_SOP0021_v0 1_ Colony PCR with |

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| | | | MyTaq |
| 04.08.15 | ADK | Transformation of pSB1K3 into Top10 | iGEM2013_SOP0009_v01_TSB transformation |
| 05.08.15 | ADK | Cheeked if our Freeze stocks, contains the right plasmids | |
| 07.08.15 | ADK | Miniprep of: #6: MG1655/ pSB1C3-KT25 #7: MG1655/pSB1K3-UT18C #22: pSB1C3-KT25-LeuZ-RFP #23: pSB1K3-UT18-LeuZ-RFP | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 08.08.15 | ADK | Miniprep and freeze stock of pSB1K3 (R71) and pSB1C3 (R72) | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 08.08.15 | ADK, TBA | Fast digest of: R67: MG1655/ pSB1C3-KT25 R68: MG1655/pSB1K3-UT18C R69: pSB1C3-KT25-LeuZ-RFP R70: pSB1K3-UT18-LeuZ-RFP With BamH1 and Pst1 | iGEM2014_SOP0017_v01_Fast digest |
| 08.08.2015 | ADK | Ligation of cut R67, R68, R69, R70 with leuZ (Y63) | iGEM2014_SOP015_v02_ligation |
| 09.08.2015 | ADK | Transformation of R67, R68, R69, R70 with leuZ (Y63) into <i>E. coli</i> MG1655 | iGEM2013_SOP0009_v01_TSB transformation |
| 10.08.2015 | JSP | Transformation of R54 & R55 (suspect we used backbones with error in sequences) | iGEM2013_SOP0009_v01_TSB transformation |

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| 11.08.2015 | ADK | Miniprep of transformations from yesterday, stored as R77 and R78 respectively. | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 12.08.2015 | ADK, TBA | Fast digest of R77, R78, R63 and R64, using BamH1 and Pst1. PCR of G24 (true leucin zipper), stored as G43. | iGEM2014_SOP0017_v01_Fast digest |
| 13.08.2015 | ADK | PCR of G24, stored as G44 and G45 Fast Digest of G45 using BamH1 and Pst1 | iGEM2014_SOP0010_v01_Phusion PCR iGEM2014_SOP0017_v01_Fast digest |
| 13.08.15 | JSP | Ligation of Leucine zipper into backbones: pSB1K3-RFP-T25 pSB1C3-RFP-T18 pSB1C3-T18 pSB1K3-T25 | iGEM2014_SOP015_v02_ligation |
| 14.08.15 | ADK | Transformation of yesterday's ligations into MG1655 | iGEM2013_SOP0009_v01_TSB transformation |
| 15.08.2015 | JSP | Colony PCR on pSB1K3-RFP-T25 and pSB1C3-RFP-T18 (Y76+Y77) ligated with leucin zipper (Y63) | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 15.08.15 | JSP | Ligation of Leucine zipper into backbones: pSB1C3-T18 pSB1K3-T25 (Y74+Y75) | iGEM2014_SOP015_v02_ligation |
| 16.08.2015 | JSP | Transformation of ligation from 15.08.2015 (Y63+Y74/Y75) | iGEM2013_SOP0009_v01_TSB transformation |
| 17.08.2015 | TBA, JSP. | Fast Digest of R16, R17, R18 | iGEM2014_SOP0017_v01_ |

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| | TBA, ADK | & R19 with BamH1 + Pst1. Stored as Y86, Y87, Y89 & Y88 respectively. Ligation of Y86, Y87, Y88 & Y89 with Y63 (Leucinzipper) | Fast digest iGEM2014_SOP015_v02 _ligation |
| 18.08.2015 | ADK | Transformation of yesterdays ligations | iGEM2013_SOP0009_v0 1_TSB transformation |
| 20.08.2015 | JSP | FastDigest of pSB1C3-T25 (R17) and pSB1K3-T18 (R18), and Leucine Zipper (G46). Ligation of pSB1C3-T25 (R17) and pSB1K3-T18 (R18), and Leucine Zipper (G46). | iGEM2014_SOP0017_v01_ Fast digest iGEM2014_SOP015_v02 _ligation |
| 21.08.2015 | JSP | Transformation of ligation from 20.08.2015 | iGEM2013_SOP0009_v0 1_TSB transformation |
| 21.08.2015 | JSP | Mini-prep on pSB1C3- T18+T25 and pSB1K3- T18+T25 from freeze-stock #5- #8 - stored as R89-R92 | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 22.08.2015 | AC | Colony-PCR on transformation from 21.08.2015 | iGEM2013_SOP0021_v0 1_Colony PCR with MyTaq |
| 25.08.2015 | JSP | FastDigest on R32 (PcstA- RFP) and R89-92 (pSB1C3- T18+T25 and pSB1K3- T18+T25) Ligation of the above digested | iGEM2014_SOP0017_v01_ Fast digest iGEM2014_SOP015_v02 _ligation |

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| | | products. | |
| 26.08.2015 | TBA | Transformation of ligations from 25th. | |
| 27.08.2015 | JSP | Colony-PCR on 4 selected colonies from each transformation. | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
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| 30.08.2015 | TBA ETM | Miniprep from #52 and #53, stored as R110 and R111 Transformation (test of the two-hybrid system) | iGEM2013_SOP0009_v01_TSB transformation |
| 31.08.2015 | | Colony PCR | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 01.09.2015 | EMT, JSP | PCR on plasmids | iGEM2014_SOP0010_v01_Phusion PCR |
| 02.09.2015 | TBA | Transformation (test of the two-hybrid system) into Δ cyaA | iGEM2013_SOP0009_v01_TSB transformation |
| 04.09.2015 | JSP | Transformation (test of the two-hybrid system) into Δ cyaA | iGEM2013_SOP0009_v01_TSB transformation |
| 05.09.2015 | JSP | Colony-PCR with specific primers and on wt Δ cyaA + BTH101 | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 06.09.2015 | JSP | Transformation (test of the two-hybrid system) into Δ cyaA | iGEM2013_SOP0009_v01_TSB transformation |
| 06.09.2015 | JSP | Colony-PCR on transformation from 05.09.2015 | iGEM2013_SOP0021_v01_Colony PCR with |

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5. Experiment history

| Date (DD.MM.YY) | Person(s) (initials) | Alterations to SOPs and remarks to experiments | SOPs |
|--------------------|-------------------------|---|--------------------------------------|
| 28.07.15 | JSP | <p>Primers in use:</p> <p>Zip_F - 019 cgctTCTAGaGGGATCCgaaaa tttgatatttcaatctggtatgaaacagctgg aagacaaagttga</p> <p>Zip_R - 020 atatCTGCAGCggccgctACTA GTaacgttcaccaaccagtttttcaga</p> <p>Two PCR was done; one with normal volume of primers (2,5 µl) and one with half amount (1,25 µl). HF buffer used in both occations.</p> | iGEM2014_SOP0010_v01_ Phusion PCR |
| 29.07.15 | JSP | <p>Restriction enzymes used: BamHI and PstI</p> <p>Incubated for 30 min + 15 min in ice afterwards.</p> <p>10 µl of PCR product used (7,4 ng/µl)</p> | iGEM2014_SOP0017_v01_ Fast digest |
| 29.07.15 | ADK | <p>Pcr produkt of G20 (leuZ) used HF buffer and CG buffer, 0,5µl primers: Zip_F - 019 and Zip_R - 020.</p> | iGEM2014_SOP0010_v01_ Phusion PCR |
| 30.07.15 | JSP, TBA | <p>For the FastDigest 7 µl (approx. 0,21 µg) PCR product</p> | iGEM2014_SOP0017_v01_ Fast digest |

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| | | was used (G23) | |
| 30.07.15 | JSP, TBA | <p>FastDigest was done on the following backbones:</p> <p>pSB1K3-RFP-KT25-LeuZ (R54): 10 µl</p> <p>pSB1C3-RFP-UT18-LeuZ (R55): 48 µl</p> <p>pSB1K3-KT25-LeuZ (R62): 30 µl</p> <p>pSB1C3-UT18-LeuZ (R61): 30 µl</p> <p>For R55, R61, R62: 6 µl 10x FastDigest buffer, 2 µl enzyme (of both FasAP and FD enzymes) (in a total volume of 60 µl)</p> | iGEM2014_SOP0017_v01_ Fast digest |
| 31.07.2015 | ADK | Miniprep of (Top10) pSB1C3-UT18-LeuZ, and (Top10) pSB1K3-KT25-LeuZ - Done according to the SOP | GE healthcare kit |
| 31.07.2015 | ADK | (Attempt 1) Fast digest of pcr product leuZ (G23): 2µl 10x FD buffer, 1µl BamH1, 1µlPst1, 16µl G23 | iGEM2014_SOP0017_v01_ Fast digest |
| 31.07.15 | ADK | Fast digest of R63 (pSB1C3-UT18-LeuZ) and R64 (pSB1K3-KT25-LeuZ), with BamH1 and Pst1. Made two mix: 8µl R63/10µl R64, 3µl 10x FD buffer, 1µl BamH1, 1µl Pst1, 1µl phosphate FD, and micro H ₂ O to a total volume of | iGEM2014_SOP0010_v01_ Phusion PCR |

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| | | <p>30µl.</p> <p>(Attempt 2) Fast digest of pcr product leuZ, used the last of leuZ product from G21, G22 (approx. 70µl in total), 2µl 10x FD buffer, 1µl BamH1, 1µl Pst1</p> | |
| 31.07.15 | JSP | <p>PCR on leucin zipper from R27. Both HF and GC was tested. Half amount of primers used (1,25 µl). A sample with no template was also tested. PCR was made with temperature gradient.</p> | iGEM2014_SOP0010_v01_Phusion PCR |
| 31.07.15 | TBA, JSP | <p>For ligation approx. 0, 20 and 50 fmol was used of digested leucin zipper(Y63) (0, 0,2 µl + 0,4 µl)</p> <p>Digested backbones used for ligation: Y57, Y58, Y59 and Y60</p> <p>Ligation was set to overnight.</p> | iGEM2014_SOP015_v02_ligation |
| 01.08.2015 | ADK | <p>Transformation of pSB1K3-RFP-T25(Y57)</p> <p>pSB1C3-RFP-T18 (Y58)</p> <p>pSB1C3-T18 (Y59), pSB1K3-T25 (Y60), ligated with leuzine zipper (Y63) into <i>E. coli</i> Top10</p> <p>- Done according to the SOP - plated out on 25µCML and 25µ, according to the plasmid resistance</p> | iGEM2013_SOP0009_v01_TSB transformation |
| 03.08.2015 | ADK | <p>Colony PCR with MyTaq of</p> | iGEM2013_SOP0021_v0 |

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| | | first draw of Y59 and Y58. Annealing temp. 55 ⁰ | 1_ Colony PCR with MyTaq |
| 04.08.15 | ADK | Colony PCR with MyTaq of first draw of Y59 and Y58. Annealing temp. 60 ⁰ . 5µl MyTaq, 1µl VF primer, 1µl VR primer, 3µl water pr. reaction and 0,5µl boiled culture (boiled in 50µl water) | iGEM2013_SOP0021_v0 1_ Colony PCR with MyTaq |
| 04.08.15 | ADK | Transformation of pSB1K3 into Top10 - because we have a suspicion that we might have swapped up the two plasmids - plated out on 25γ kan plates | iGEM2013_SOP0009_v0 1_TSB transformation |
| 05.08.15 | ADK | Checked if our freeze stocks contains the right plasmids according to our freeze stock book, by plaiting them out on plates according to their resistance, and to see if those containing RFP turned red. We checked the following: #5: MG1655/pSB1C3-UT18C #6: MG1655/ pSB1C3-KT25 #7: MG1655/pSB1K3-UT18C #8: MG1655/pSB1K3-KT25 #17: TOP10/pSB1C3-UT18C-LeuZ #20: Top10/pSB1K3-KT25-LeuZ #22: pSB1C3-KT25-LeuZ-RFP #23: pSB1K3-UT18-LeuZ-RFP #33: TOP10/pSB1K3-RFP- | |

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| | | KT25-LeuZ | |
| 07.08.15 | ADK | Miniprep of #6, #7, #22, #23 - done according to the SOP in the Kit | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 08.08.15 | ADK | Miniprep and freeze stock of pSB1K3 (R71) and pSB1C3 (R72) Freezestock 700µl overnight culture, and 300µl glycerol | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| | | Fast digest of: R67: MG1655/ pSB1C3-KT25 R68: MG1655/pSB1K3-UT18C R69: pSB1C3-KT25-LeuZ-RFP R70: pSB1K3-UT18-LeuZ-RFP With BamH1 and Pst1 - done according to SOP | iGEM2014_SOP0017_v01_Fast digest |
| 08.08.2015 | ADK | Ligation of cut R67, R68, R69, R70 with leuZ (Y63), overnight. Done according to the SOP, inserted 0Fmol, 20Fmol, 50Fmol. | iGEM2014_SOP015_v02_ligation |
| 09.08.2015 | ADK | Transformation of R67, R68, R69, R70 with leuZ (Y63) into <i>E. coli</i> MG1655. Done according to the SOP, and plated out on CML and KAN LA-plates according the backbones resistance | iGEM2013_SOP0009_v01_TSB transformation |

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| 10.08.2015 | JSP | Transformation of R54 and R55 | iGEM2013_SOP0009_v01_TSB transformation |
| 11.08.2015 | ADK | Miniprep of transformations from yesterday, stored as R77 and R78 | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 12.08.2015 | ADK, TBA ADK | Fastdigest of R63, R64, R77, R78. stored Y74, Y75, Y76 and Y77 respectively PCR of leuZ G24 - two attempts. Second attempt added 4µl MgCl ₂ | iGEM2014_SOP0017_v01_Fast digest iGEM2014_SOP0010_v01_Phusion PCR |
| 13.08.2015 | ADK | PCR of leuZ G24, didn't add MgCl ₂ , used half amount of primers , HF buffer. stored as G45, G44 FastDigest of G45 using BamH1 and Pst1. Left for ~ 45 min. Ligation of Y74, Y75, Y76 and Y77 with digested G45 | iGEM2014_SOP0010_v01_Phusion PCR |
| 13.08.2015 | JSP | For ligation approx. 0, 20 and 50 fmol was used of digested leucin zipper(Y78) (0, 0,4 µl + 0,9 µl) Digested backbones used for ligation: Y74, Y75, Y76 and Y77 - approx. 10 µl backbone was used to each ligation. Ligation was set to overnight. | iGEM2014_SOP015_v02_ligation |

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| 14.08.15 | ADK | Transformation of yesterday's ligations into MG1655 - Done according to SOP | iGEM2013_SOP0009_v01_TSB transformation |
| 15.08.2015 | JSP | Colony PCR on Y76 and Y77 ligated with Y63. No real alterations to SOP. Colonies were streaked out on agar plates and added directly to MyTaq solutions. | |
| 15.08.2015 | JSP | For ligation approx. 0, 20 and 50 fmol was used of digested leucin zipper(Y63) (0, 0,25 μ l + 0,5 μ l) Digested backbones used for ligation: Y74 and Y75- approx. 10 μ l backbone was used to each ligation. Ligation was set to overnight. | iGEM2014_SOP015_v02_ligation |
| 16.08.2015 | JSP | Transformation of ligation from 15.08.2015 (Y63+Y74/Y75): no alterations to SOP | iGEM2013_SOP0009_v01_TSB transformation |
| 17.08.2015 | TBA, JSP | Fast Digest of R16 (pSB1C3-UT18), R17 (pSB1C3-KT25), R18 (pSB1K3-UT18) and R19 (pSB1K3-KT25) with BamH1 + Pst1. By error, 2 μ L of each restriction enzymes was added instead of 1 μ L. Incubated for approx. 50 min. Digested R16 stored as Y86, digested R17 stored as Y87, digested R18 stored as Y89 and | iGEM2014_SOP0017_v01_Fast digest |

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| | TBA, ADK | <p>digested R19 stored as Y88.</p> <p>Ligation of Y86 - Y89 with Y63 (digested Leucin zipper). Possible contamination of Y63. Ligations left overnight at 16 degrees.</p> | iGEM2014_SOP015_v02_ligation |
| 18.08.2015 | ADK | Transformation of yesterdays ligations | iGEM2013_SOP0009_v01_TSB transformation |
| 20.08.2015 | JSP | <p>FastDigest of pSB1C3-T25 (R17) and pSB1K3-T18 (R18), and Leucine Zipper (G46). All cut with BamHI and PstI - digestions was given approx. 2 hours.</p> <p>Ligation of pSB1C3-T25 (R17) and pSB1K3-T18 (R18), and Leucine Zipper (G46). Instead of using 30 and 50 fmol inserts, 60 and 90 fmol inserts were used. Ligations were left overnight.</p> | <p>iGEM2014_SOP0017_v01_Fast digest</p> <p>iGEM2014_SOP015_v02_ligation</p> |
| 21.08.2015 | JSP | <p>Transformation of ligation from 20.08.2015</p> <p>pSB1C3-T25 + Leucine Zip (Y91+Y90)</p> <p>pSB1K3-T18 + Leucine Zip (Y92+Y90)</p> <p>Strain used: MG1655</p> <p>When streaked out on agar plates, Y91 was added to</p> | iGEM2013_SOP0009_v01_TSB transformation |


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| | | kanamycin plates, and Y92 chloramphenicol plates. This should have been the other way around(!). As 10 µl (max.) was still left in the eppendorf tubes, these left overs was streaked out on the appropriate agar plates. | |
| 21.08.2015 | JSP | <p>Mini-prep on pSB1C3-T18+T25 and pSB1K3-T18+T25 from freeze-stock #5-#8 - stored as R89-R92</p> <ul style="list-style-type: none"> - only 1,5 mL of liquid culture used for miniprep (for higher concentration, next time use all 5 mL) - No alteration to protocol. | Bio-Rad Quantum Prep: plasmid miniprep Kit |
| 22.08.2015 | AC | Colony-PCR on transformation from 21.08.2015 | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 25.08.2015 | JSP | <p>FastDigest on R32 (PcstA-RFP) and R89-92 (pSB1C3-T18+T25 and pSB1K3-T18+T25)</p> <p>R32 cut with SpeI+PstI R89-92 cut with XbaI+PstI</p> <p>Stored as Y93(PcstA-RFP), Y94(T18), Y95 (T25)</p> <p>Ligation of the above digested products. No alteration to SOP.</p> | <p>iGEM2014_SOP0017_v01_Fast digest</p> <p>iGEM2014_SOP015_v02_ligation</p> |

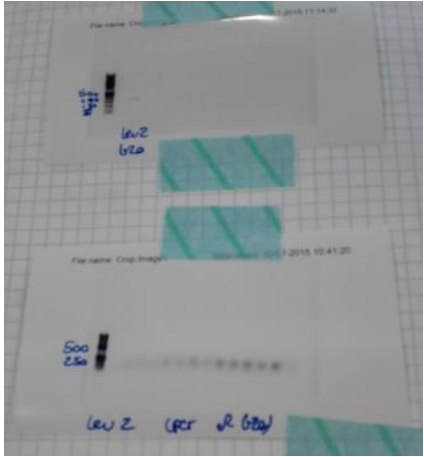
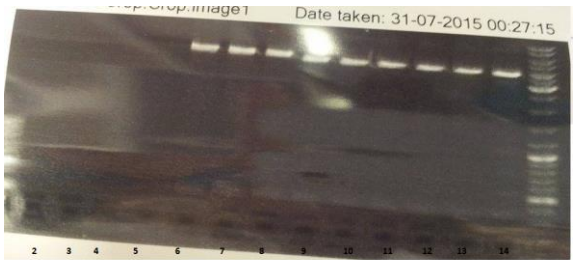
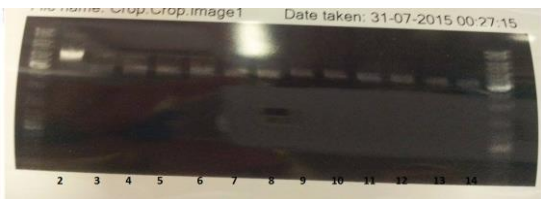
| | | | |
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| 26.08.2015 | TBA | Transformation of yesterdays overnight ligations into MG1655. | |
| 27.08.2015 | JSP | Colony-PCR. No alteration to SOP. | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
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| 30.08.2015 | TBA EMT | Miniprep of overnight cultures of #52 and #53. stored as R110 and R111, respectively. Transformation of 8 plasmid combinations to test the two-hybrid system | iGEM2013_SOP0009_v01_TSB transformation |
| 31.08.2015 | | Results inconclusive. colony PCR on colonies from selected plates. | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 01.09.2015 | EMT, JSP | PCR of plasmids in use to check if lenght of content matches what they supposedly contain | iGEM2014_SOP0010_v01_Phusion PCR |
| 02.09.2015 | TBA | Transformation of different Rxx+Rxx, Rxx+Rxx | iGEM2013_SOP0009_v01_TSB transformation |
| 04.09.2015 | JSP | Transformation (test of the two-hybrid system) into Δ cyaA. The following combinations was used: R111+R113 (T18-Zip+25-Zip) (R113 transformed into R111 | iGEM2013_SOP0009_v01_TSB transformation |

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| | | <p>containing ΔcyaA strain.) R111+R104 (T18-Zip+T25) (R104 transformed into R111 containing ΔcyaA strain.) R107+R113 (T18+T25-Zip) (R113 transformed into R107 containing ΔcyaA strain.) R108+R103 (T18+T25) (R103 transformed into R108 containing ΔcyaA strain.) R110 R111 R108 R107</p> | |
| 05.09.2015 | JSP | <p>Colony-PCR on transformation from <i>04.09.2015</i> - No changes to SOP</p> | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 06.09.2015 | JSP | Transformation (test of the two-hybrid system) into Δ cyaA | iGEM2013_SOP0009_v01_TSB transformation |
| 06.09.2015 | JSP | Colony-PCR with specific primers and on wt Δ cyaA + BTH101 | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 06.09.2015 | JSP | Transformation (preparation for test of the two-hybrid system) into BTH101: R103, R104, R107, R108, R112 and R113 were transformed. | iGEM2013_SOP0009_v01_TSB transformation |
| 07.09.2015 | JSP | Transformation (test of the two-hybrid system) into | iGEM2013_SOP0009_v01_TSB transformation |

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| | | <p>BTH101. R102+R98 were transformed into BTH101 strain containing R103. R112 was transformed into BTH101 containing R104. R113 was transformed into BTH101 containing R111. These transformation were also made: R97+R98 (double-plasmid transformation) R97 R98 R112 R113</p> | |
| 08.09.2015 | JSP | Colony-PCR on transformation from <i>06.09.2015</i> | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |
| 08.09.2015 | JSP | Colony-PCR on transformation from <i>07.09.2015</i> | iGEM2013_SOP0021_v01_Colony PCR with MyTaq |

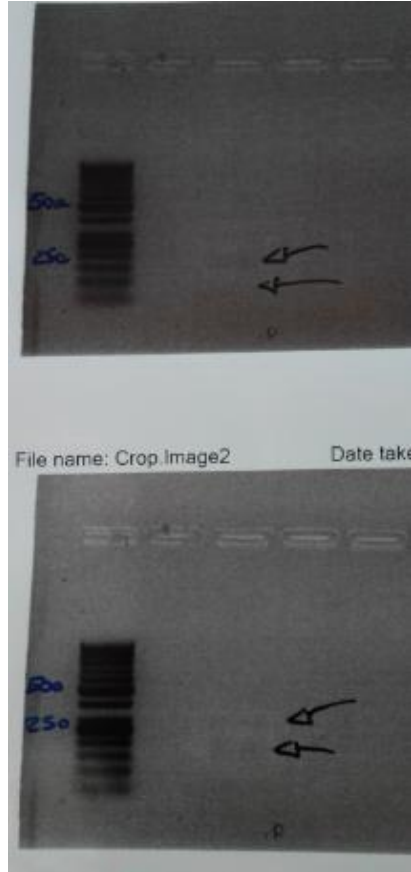
6. Results

| Date (DD.MM.YY) | Picture | Comments |
|--------------------|---|---|
| 28.08.15 |  | PCR on leucin-zipper from R49: Strong bands at approx. 100-200 bp (difficult to distinguish between the bands in the ladder) Conc. from |

| | | |
|----------|---|--|
| | | purification: 7,6 ng/μl (labeled G20) |
| 29.07.15 | no picture | No bands on gel after FastDigest. |
| 29.07.15 |  | Pcr of G20: Both buffers worked, got to concentrations: G21=12,88ng/μl and G22=10,79ng/μl. Did also purified G20: 20μl G20(per product of leuZ), 300μl binding buffer, 300μl isopropanol - ran it on the column and washed with washing buffer. Got the concentration G23=30,18ng/μl |
| 30.07.15 | <p>Row 1</p>  <p>Row 2</p>  | <p><u>Row 1:</u> 2-5: G23 6-8: R54 9-14: R55</p> <p><u>Row 2:</u> 2: R55 3-8: R61 9-14: R62</p> <p>R54 + R55 was expected to be around 4000 bp. This is consistent with the bands shown.</p> <p>R61+R62 was expected to be around 3000 bp. This is also consistent with the bands shown. G23 was</p> |

| | | |
|------------|--|--|
| | | expected to around 160 bp. No bands shown can be explained by a low concentration. |
| 31.07.15 | | Got following concentration: R63 (pSB1C3-UT18-LeuZ)=121,57ng/μl, and R64 (pSB1K3-KT25-LeuZ)=83,86ng/μl |
| 31.07.15 | | Fast digested leuZ: Got no bands on the gel |
| 31.07.2015 | | Fast digest of pSB1C3-UT18-LeuZ (R63) and pSB1K3-KT25-LeuZ (R64), with BamH1 and Pst1 - Got the following concentration: pSB1C3-UT18-LeuZ (Y61):7,96ng/μl, and pSB1K3-KT25-LeuZ (Y62): 5,81ng/μl |

31.07.2015



(Attempt 2) Fast digest of pcr product leuZ. Got two very unclear bands - Measured the concentration of the digested leuZ (Y63) on the nanodrop, Y63=13,8ng/ μ l

31.07.15

Row 1



Row 2



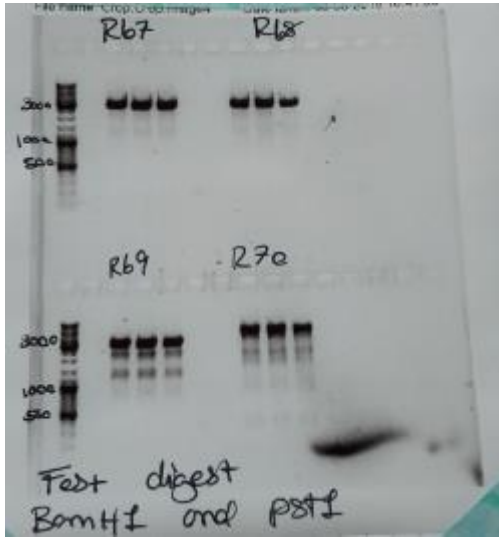
Row 1

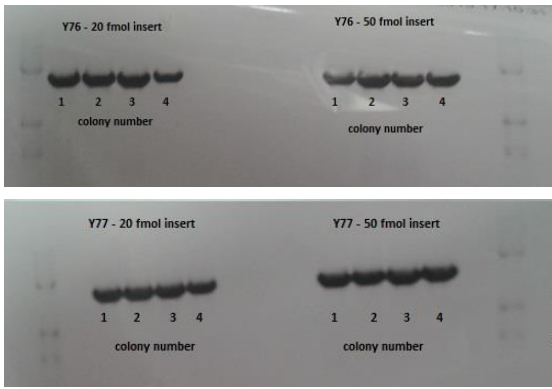
2-11: PCR with GC buffer
12-15: PCR with HF buffer

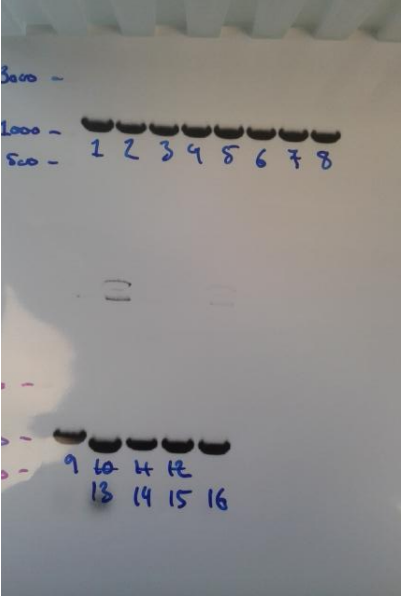
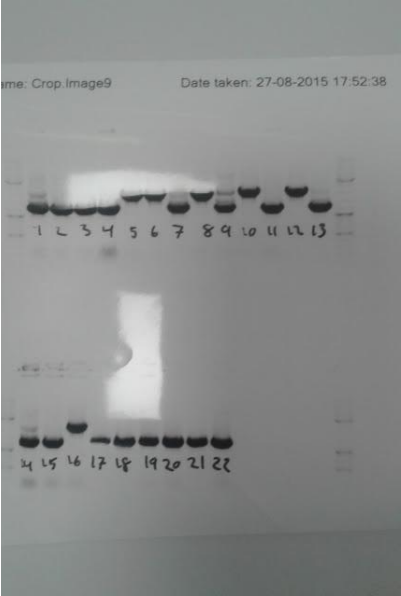
Row 2

2-7: PCR with HF buffer
8-12: No template
For those PCR reactions with DNA template, the bands were as expected around 160 bp (with some exceptions: row 1, #14 and row 2, #5). The bands from the PCR reactions with no template were as expected much lower: approx. 50 bp.

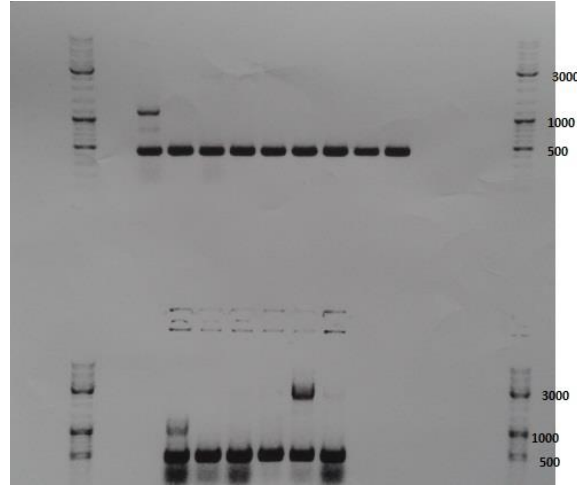
| | | |
|------------|--|--|
| 03.08.2015 | | Transformation: got colonies of the Top10 with Y58(pSB1C3-RFP-T18-leuZ) and Y59(pSB1C3-T18-leuZ) on the CML plates, but unfortunately, we didn't got any colonies on the Kan plates with Y57 (pSB1K3-RFP-T25), and Y60 (pSB1K3-T25) - We try to plate these out on Cml plates, which gave colonies |
| 03.08.2015 | | Colony PCR failed, no bands - will try again tomorrow, with 10µl MyTaq, 1µl VF primer, 1µl VR primer, 8µl water pr. reaktion. I |
| 05.08.2015 | | Transformation failed |
| 06.08.2015 | | Check of freeze stock: It looks like all sample contains the right plasmids. We made overnight culture of #6, #7, #22, #23. |
| 07.08.2015 | | Miniprep: - R67 (#6: MG1655/pSB1C3-KT25)=96,8ng/µl -R68 (#7 MG1655/pSB1K3- |

| | | |
|-------------|--|--|
| | | <p>UT18C)=58,7ng/μl - R69 (#22 pSB1C3- KT25-LeuZ- RFP)=212,9ng/μl -R70 (#23: pSB1K3- UT18-LeuZ-RFP) =116,7ng/μl</p> |
| 08.08.15 | | <p>Miniprep: pSB1K3 (R71)=103,4ng/μl pSB1C3 (R72)=198,04ng/μl</p> |
| 08.08.15 |  | <p>Fastdigest: we are a little uncertain of why we get so many bands. Have decided to purified the clear bands, and ligate these backbones eith Y63 (leuZ cut with BamH1 and Pst1)</p> |
| 09.08.2015 | | |
| 10.08.2015 | | <p>Yesterdays transformations failed. all of them</p> |
| 11. 08.2015 | | <p>Transformations with correct plasmid backbones from the 10th succeeded</p> |
| 12.08-2015 | | <p>First PCR G24 (true leucin zipper) attempt</p> |

| | | |
|------------|--|--|
| | | <p>with both MyTaq and phusion (with extra MgCl₂) polymerase failed due to wrong primers.... second attempt using MyTaq yielded only poor concentrations.</p> <p>Fast Digests was succesful, but due to insuccicient amount of G24, ligation was postponed.</p> |
| 13.08.2015 | | <p>Phusion PCR of G24 without extra MgCl₂ yielded better concentrations.</p> <p>Fast digest of G45</p> |
| 14.08.15 | | <p>Transformation. Colonies of MG1655/Y76 and MG1655/Y77 - MG1655/Y74 and MG1655/Y75 failed</p> |
| 15.08.2015 |  <p>The figure displays four agarose gel images arranged in a 2x2 grid. The top row shows results for strain Y76: the left gel is labeled 'Y76 - 20 fmol insert' and the right gel is 'Y76 - 50 fmol insert'. The bottom row shows results for strain Y77: the left gel is 'Y77 - 20 fmol insert' and the right gel is 'Y77 - 50 fmol insert'. Each gel has four lanes numbered 1 to 4, representing different colony numbers. To the right of the gels, molecular weight markers are indicated at 3000, 1000, and 500 nucleotides (nt). In all cases, bands are visible in lanes 1, 2, 3, and 4, with the bands for Y76 appearing slightly lower than those for Y77.</p> | <p>Colony PCR showed bands at approx. 2000-2500 nt. For Y76 the band should be around 2200 nt and for Y77 the band should be around 2300 nt.</p> |
| 17.08.2015 | | <p>Transformation from</p> |

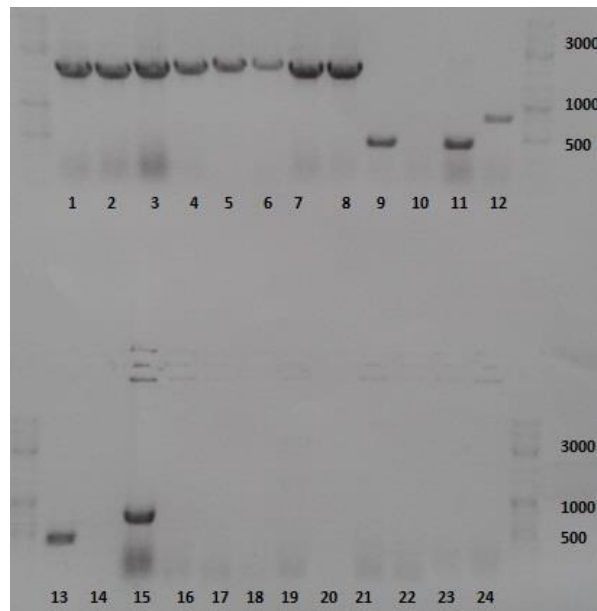
| | | |
|------------|---|--|
| | | 16.08.2015 failed. No colonies. |
| 22.08.2015 |  | <p>Transformation was successful, even though only 10 μl instead of approx. 100 μl was streaked out. The results from colony-PCR confirmed the successful ligation+transformation (the bands should be approx. 1200-1300 nt)</p> <p>1-8: Y91 9-16: Y92</p> |
| 27.08.2015 |  | <p>Transformation was successful (#5, #6, #8, #10, #12, #16). Bands were expected to around 2100-2200 nt long.</p> <p>1-4: Y94-20 (PcstA-RFP-T18) 5-8: Y94-50 (PcstA-RFP-T18) 9-12: Y95-20 (PcstA-RFP-T25) 13-16: Y95-50 (PcstA-RFP-T25)</p> |

05.09.2015




Not succesful. We would have expected Two bands in each of them (one at approx. 1000 bp and one at 2000 bp.) Instead we got a band at approx. 450 bp. Probably a contaminant.

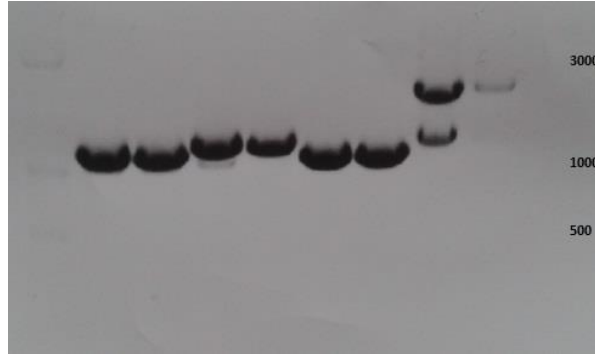
06.09.2015



What we expected:
1+2: R107 (approx. 2100)
3+4: R108 (approx. 2100)
5+6: R110 (approx. 2100)
7+8: R111 (approx. 2100)
9+10: R111 (approx. 2100 (VF/VR-primers) or 750 bp (001+003 primers))
11+12:R107 (approx. 2100 (VF/VR-primers) or 750 bp (001+003 primers))
13+14:R108 (approx. 2100 (VF/VR-primers) or 850 bp (001+003 primers))
15+16: R108+R103 (two bands at approx. 750 and 850 bp 001+002/3 primers used)
17+18:R111+R104 (two bands at approx. 750 and 850 bp 001+002/3 primers used)
19+20:R107+R113 (two

| | | |
|------------|--|---|
| | | <p>bands at approx. 750 and 850 bp 001+002/3 primers used)</p> <p>21+22:R113+R111 (two bands at approx. 750 and 850 bp 001+002/3 primers used)</p> <p>23: 'clean' ΔcyaA: no band</p> <p>24: 'clean' BTH101: no band</p> <p>Conclusion: From these results it can be concluded that only colony 1-8+23+24 were correct (and partially 12)</p> |
| 07.09.2015 |  | <p>What was expected:</p> <p>1+2: R103 (1050 bp)</p> <p>3+4: R104 (1150 bp)</p> <p>5+6: R107 (approx 2100 bp)</p> <p>7+8: R108 (approx 2100 bp)</p> <p>9+10:R111 (approx 2100 bp)</p> <p>Conclusion:</p> <p>Transformations were succesful</p> |

08.09.2015



What was expected:

1+2: R103+R102 (1050-1150 bp)

3+4: R103+R98 (1050-1300 bp)

5+6: R104+R112 (approx 1100 bp)

7+8: R111+R113 (1300+2100 bp)

Conclusion:

Transformations were successful

This was also confirmed by the presence of blue colonies at the plate with colonies transformed with R111+R113 (T18-zip+T25-zip)

7. Appendices