

21.07.2020

Transformation see protocol

- Thawed 2 Eppis each with 200 µl overnight bacterial culture from E.coli BL21 strain
- Prepare 2 approaches on the flame:
 - Eppi 1: 1 µl sterile H₂O
 - Eppi 2: 1 µl self-prep marLac

22.07.2020

Prepare 2YT medium: on three liter

- 48 g Trypton (16 g per L)
- 30 g Yeast (10 g per L)
- 15 g NaCl (5 g per L)

overnight culture:

- in 5 ml LB medium for glycerine
- in 45 ml LB medium for production
- overnight at 37°C 170rpm

23.07.2020

glycerine stock:

- OD of overnight culture: 0,417
- 2 ml of preculture for 2 min and 13.000 rpm in the centrifuge
- resuspend cell pellet in 400 µl LB and 2µl ampicillin
- add 600 µl sterilem glycerine, invert, incubate 15 min on ice
- storage cells at -80°C

production of marLac, 30°C, 3h:

- OD of overnight culture 1:10 dilution: 0,708
- 4 5 L sterile Erlenmeyer flesks fill with 1 L of sterile 2YT medium
 - o 1 ml Amp (c = 1 µl/ml of 100 µM Stock)
 - o 0,1 ml CuSO₄ (c = 0,1 mM of 1 M Stock)
 - o add 10 ml of overnight culture
- Start OD:

| 1 | 2 | 3 | 4 |
|-------|------|-------|-------|
| 0,028 | 0,03 | 0,026 | 0,028 |

- put it in the shaker at 95 rpm and 37°C, until the culture reaches a OD of 0,5
 - o let it grow for 2 hours

| 1 | 2 | 3 | 4 |
|-------|-------|-------|-------|
| 0,561 | 0,814 | 0,787 | 0,872 |

- remove pre-induction sample (VI)
- add 500 µl IPTG (12:00 Uhr)
- After 3h (15:00 Uhr) with 30°C remove after induction sample (induction)

| | | | | |
|----------|-------|-------|-------|-------|
| dilution | 1 | 2 | 3 | 4 |
| 1:10 | 0,255 | 0,282 | 0,309 | 0,315 |
| - | 2,55 | 2,82 | 3,09 | 3,15 |

- fill it in 1 L centrifuge beaker and balance it
 - in the centrifuge for 15 min, 9000 rpm and 4°C
 - discard the supernatant and put the pellet in a Falcon (all 4 pellets in one Falcon)
 - o do everything on ice
 - weigh the pellet and store it on -20°C
- ➔ m_{Pellet}: 11,53 g

27.07.2020

Cell disruption and first affinity chromatography.

- See protocol
- dilution of not induced sample and induced sample:

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 |
|-----------------------|---------------|-----------------|-------------|---------------|---------------------|--------------------|-----------|-----------|-----------|-----------|
| Sample | Marker | not ind. | ind. | Lysate | flow through | Wash eluate | E1 | E2 | E4 | E6 |
| Loading volume | 5 µl | 10 µl | 10 µl | 10 µl | 10 µl | 10 µl | 15 µl | 15 µl | 15 µl | 15 µl |

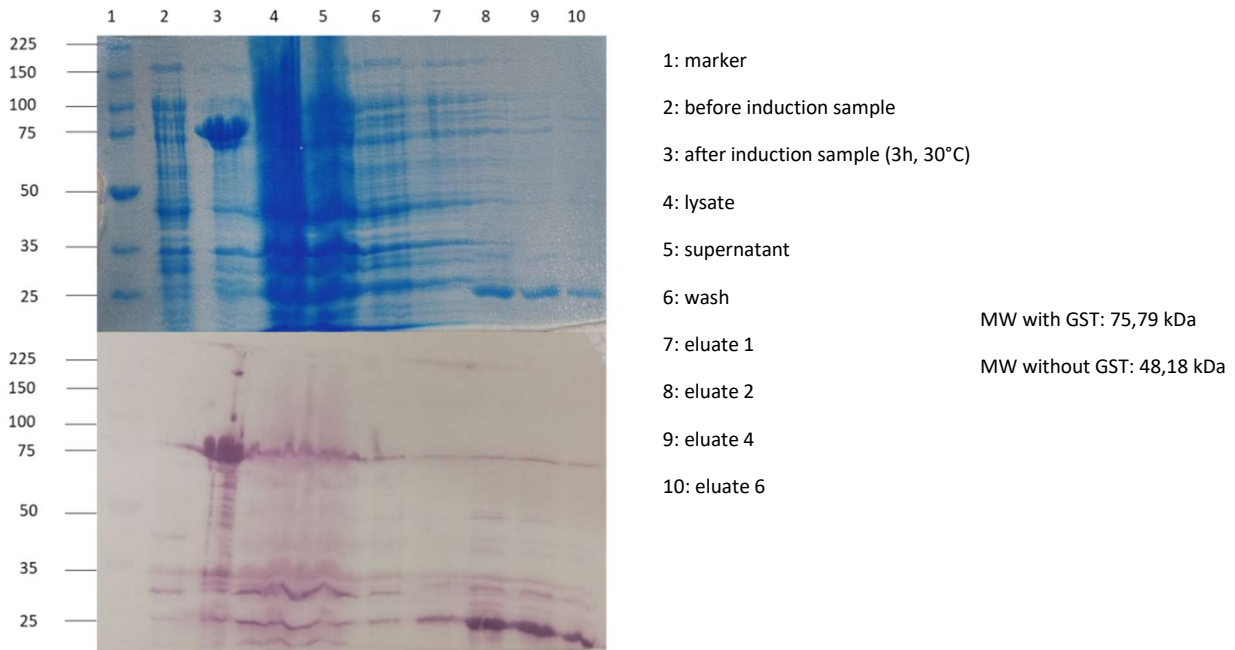
| | OD | water | SDS |
|-------------|-------|-------|------|
| Not induced | 0,561 | 10,51 | 2,6 |
| induced | 2,55 | 143,4 | 35,8 |

- then run the SDS Gel for 50min and 200V

Westernblot

- see protocol

E.coli BL21 (DE3) mar pGEX-6P-1 with CuSO₄, medium: 2YT; 3h; 30°C



Dialysis of the eluate:

- cut off a section of a dialysis tube, approx 10 cm
- attach with a clamp the lower end of the tube
- pipette the eluate in the dialysis tube
- add 20 µl Precision protease
- attach the top end with a clamp too and fix it on a swimmer
- in 1 L (100-times amount of buffer) overnight for 4°C let the dialysis happen while stirring gently

28.07.2020

buffer: see protocol

- TBS-T (pH=7,4) 1 L
- TBS (pH=7,4) 500 ml
- Tris-HCl 50 mM

2YT medium: for 4 L

- 64 g Trypton
- 40 g Yeast
- 20 g NaCl

2. affinity chromatography:

- see protocol
- put the eluate again in a dialysis tube and dialysis it in pH7 for 1,5h:
 - o buffer: phosphatbuffer pH 7, see protocol

concentration:

- see protocol
- ➔ Mar: $c = 28,45 \mu\text{M}$; $V = 800 \mu\text{l}$

Sample preparation and loading scheme.

| | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |
|-----------------------|--|---|------------------|------------------|------------------|------------------|--------------------------------------|-----------------|-------------------|
| Sample | elution fraction (before digestion) | Elution fraction (after digestion) | D1 | D2 | D3 | D4 | Elution fraction (eluted GST) | Marker | Protease |
| Volume | 60 μl | 60 μl | 60 μl | 60 μl | 60 μl | 60 μl | 60 μl | 5 μl | 10 μl |
| Sample buffer | 20 μl | 20 μl | 20 μl | 20 μl | 20 μl | 20 μl | 20 μl | - | 2,5 μl |
| Boil time | 5 min | 5 min | 5 min | 5 min | 5 min | 5 min | 5 min | - | 5 min |
| | Centrifuge all samples 5 min at 11.000g before loading | | | | | | | | |
| Loading volume | 15 μl | 15 μl | 15 μl | 15 μl | 15 μl | 15 μl | 15 μl | 5 μl | 5 μl |

- run the SDS gel for 50min 200 V

Westernblot:

- see protocol

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29.07.2020

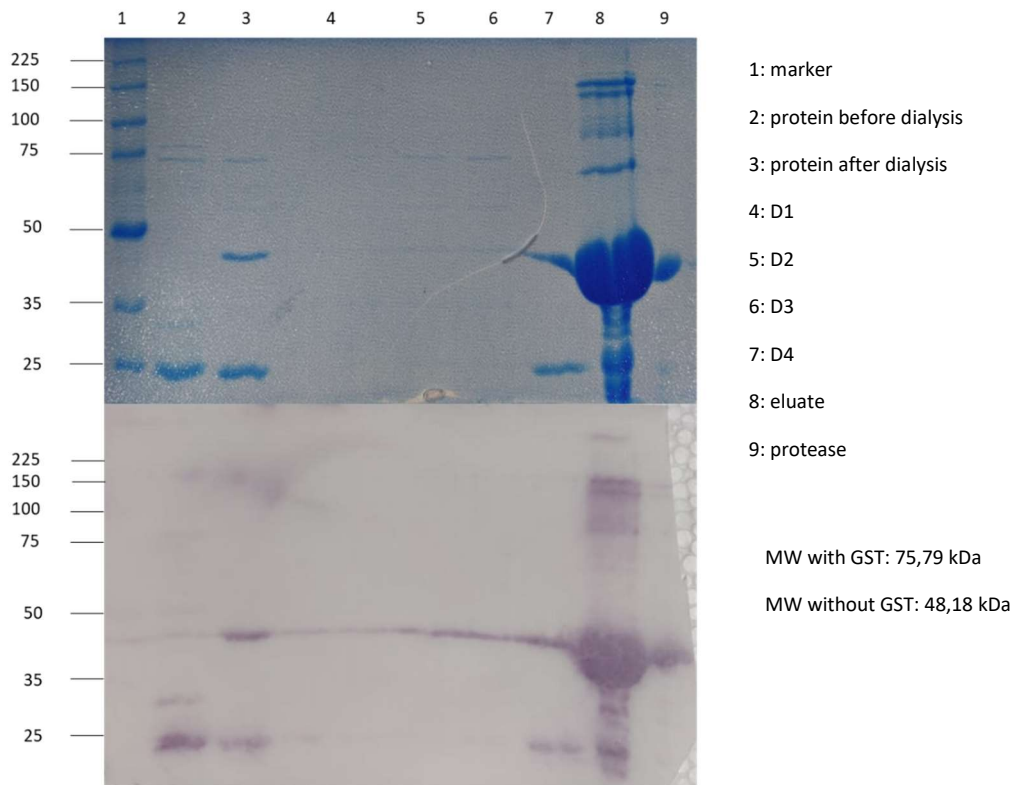
pour LB-Agar-plates

- see protocol

detect the Westernblot

- see protocol

E.coli BL21 (DE3) mar pGEX-6P-1 with CuSO₄, medium: 2YT; 3h; 30°C



overnight culture:

- see protocol
- for the BaLac-production on the 30.07.2020
 - o 50 ml in 500 ml flask without chicane
 - o 50 µl ampicillin
 - o inoculate with the tranformants of the 28.07.2020
- for BL 21 BaLac glycerine stocks:
 - o 5 ml LB in a test tube
 - o 5 µl ampicillin
 - o inoculate with the tranformants of the 28.07.2020