

mutagenesis-PCR (for base exchange)

- Primer must be individually chosen according to plasmid-backbone and DNA-fragment
- Insert desired base exchange
- Primer- T_m should be between 55 – 65 °C
- Primers should preferably have a GC-rich end (for a better adhesion to the plasmid)
- Dilute Primer 1:10 to avoid primer-dimerization
- Dilute template 1:5

| mutagenesis-PCR approach | |
|--------------------------------|-------------------|
| | Volume in μ l |
| Template (1:5) | 15 |
| dNTP's each 100 nM | 3.8 |
| 10x Pfu-buffer with $MgSO_4$ | 1 |
| 2 primer each 10 μ M | 0.8 |
| Pfu-Polymerase | 0.5 – 4 |
| H ₂ O ad 50 μ l | 128.5 |

| PCR-program | | | |
|--------------|-------------------|-------------------|--------|
| | Temperature in °C | Duration | cycles |
| Denaturation | 95 | 1 min | |
| Denaturation | 95 | 0.5 min | |
| Annealing | Depends on primer | Depends on primer | |
| Elongation | Depends on primer | Depends on primer | |
| store | 4 | ----- | |
| | | | |
| | | | |

mutagenesis-PCR

- experiment date: _____; time: _____
- name of investigator: _____
- experiment date: _____; time: _____
- name of investigator: _____
- plasmid: name: _____ number: _____ production date: _____

| | |
|---|--|
| Primer 1(forw): Primer 1 – sequence: Primer1 – T _M : Primer 1 – binding position: | Primer 2 (rev): Primer 2 – sequence: Primer 2 – T _M : Primer 2 – binding position: |
|---|--|

| mutanegesis-PCR approach | |
|---------------------------------------|--------------|
| | Volume in µl |
| Template (1:5) | |
| dNTP's each 100 nM | |
| 10x Pfu-buffer with MgSO ₄ | |
| 2 primer each 10 µM | |
| Pfu-Polymerase | |
| H ₂ O ad 50 µl | |

| PCR-program | | | |
|--------------|-------------------|----------|--------|
| | Temperature in °C | Duration | cycles |
| Denaturation | | | |
| Denaturation | | | |
| Annealing | | | |
| Elongation | | | |
| store | | ----- | |