iGEM Munich 2017 Protocols

## Polymerase chain reaction

Adapted from: https://www.neb.com/protocols/2012/08/29/protocol-for-q5-high-fidelity-2x-master-mix-m0492. Use NEB Tm Calculator to calculate the Tm of the primers.

## Aim of the experiment

This experiment can be used for exponential amplification of a DNA of interest. There are different subspecies of the reaction in existence, which can be used for special functions (i.e. addition of certain short sequences at 3 or 5 point end, insertion of point mutation etc.)

#### **Materials**

- Q5-High-Fidelity 2x Master Mix
- Forward primer
- Reverse primer
- template DNA

### **Procedure**

1. To a PCR tube add following reagents:

Table 1: PCR-Mix

Concentration	Chemicals
1x	Q5-High-Fidelity 2x Master Mix
$0.5~\mu\mathrm{M}$	Forward primer
$0.5~\mu\mathrm{M}$	Reverse primer
up to 1000 ng	Template DNA
fill up to $50\mu\mathrm{l}$	$H_2O$

iGEM Munich 2017 Protocols

2. Transfer tube to a Thermocycler and run following program:

Table 2: Thermocycling conditions

Step	Temperature (°C)	Time (s)
Initial denaturation	98	30
25-35 cycles	98	30
	98	10
	annealing temperature	20
	72	30/kb
Final extension	72	300
Hold	4	for ever

# Possible follow-up protocols

The following protocols are the next steps of a possible cloning cycle after a Polymerase Chain Reaction (PCR):

- 1. Restriction digest
- 2. Agarose-Gel-electrophoresis
- 3. PCR clean-up