

1. Molecular Biotechnology

We used different molecular biological methods in our project. All used methods are listed below.

1. Genome preparation

In our project, we use the magnetotactic bacteria's (MSR-1) genome as template to amplify our target gene. Genome preparation is a method for isolating Genome from bacterial cell cultures. In this work the TIANamp Bacteria DNA Kit (Tiangen) was used. The Genome preparation was performed following the manufacturer's manual.

1. PCR

In addition to the common PCR for amplification of certain DNA fragments, several different types of PCR were used throughout our project. The purpose, procedure and generic use are listed in the table below.

Name	Purpose	Procedure	Notes
Common PCR	amplification of certain DNA fragments	Primers binding in the side of the target genes.	general reaction setup and procedure shown in tables below
colony/check PCR	Check on insert length/ correct integration of insert	primers binding in the middle of the insert or one in insert and one in vector are used	general reaction setup and procedure shown in tables below
SOE PCR	side directed mutagenesis	primers containing the desired base pair exchange/deletion/insertion are designed and used	Splicing by Overlapping Extension PCR
gradient PCR	Finding optimal conditions for our primers to bind and the PCRs in general	Several PCRs batches are run within the same thermocycler, differing in annealing temperature	Commonly used in preliminary experiments or appear nonspecific binding.

General reaction setup of a common PCR

component	volume	
Q5 Hot Start High-Fidelity 2X Master Mix (Purchased From NEB)	25µl	12.5µl
10µM Forward primer	2.5µl	1.25µl
10µM Reverse primer	2.5µl	1.25µl
Template	1µl (<1000ng)	0.5µl
ddH ₂ O	19µl	9.5µl

General reaction procedure of a Common PCR

pre-denaturation	98°C	30s	} 35 cycles
denature	98°C	10s	
anneal	T _m	30s	
elongate	72°C	20-30s per kb	
elongate	72°C	5min	
store	12°C	∞	

1. Prepare the reaction mixture at room temperature in the order indicated:

Component	Volume	
	Plasmid DNA	Unpurified PCR product
Water, nuclease-free	35 μ l	17 μ l
10X FastDigest® buffer or 10X FastDigest® Green buffer	5 μ l	2 μ l
DNA	5 μ l (up to 2 μ g)	10 μ l (~0.2 μ g)
FastDigest® enzyme1 (NEB)	2.5 μ l	1 μ l
FastDigest® enzyme2 (NEB)	2.5 μ l	1 μ l
Total volume	50 μ l	30 μ l

2. Mix gently and spin down.
3. Incubate at 37°C in a heat block or water thermostat for 30 min.
4. Inactivate the enzyme (optional).

6. Ligation

Ligation in the context of molecular biology is the enzymatic joining of previously restricted nucleic acid fragments by synthesis of new bonds with simultaneous breakdown of ATP.

1. Set up the following reaction in a microcentrifuge tube on ice.

(T4 DNA Ligase was added last. Note that the table shows a ligation using a molar ratio of 1:3 vector to insert for the indicated DNA sizes.) Use NEBioCalculator to calculate molar ratios.

COMPONENT	20 μ l REACTION
10X T4 DNA Ligase Buffer*	2 μ l
Vector DNA :Insert DNA	1:3
Nuclease-free water	to 20 μ l
T4 DNA Ligase(NEB)	1 μ l

2. Gently mix the reaction by pipetting up and down and microfuge briefly.
3. incubate at 16°C overnight.
4. Heat inactivate at 65°C for 10 minutes (optional) .
5. Chill on ice and transform 5-10 μ l of the reaction into 100 μ l competent cells.

7. Gibson Assembly

Gibson assembly is a novel method for the easy assembly of multiple linear DNA fragments. Regardless of fragment length or end compatibility, multiple overlapping DNA fragments can be joined in a single isothermal reaction. With the activities of three different enzymes, the product of a Gibson Assembly is a fully ligated double-stranded DNA molecule. In this work the *pEASY-Uni Seamless Cloning and Assembly Kit* (Trans Gene Biotech) was used.

1. Set up the following reaction in a microcentrifuge tube on ice.

2XAssmbly Mix	5 μ l
Linearized vector	X μ l
Inserts	Y μ l
ddH ₂ O	To 10 μ l

The molar ratio of X: Y is 1:2.

2. Gently mix the reaction by pipetting up and down and microfuge briefly.

3. Incubate at 50°C for 15min.
4. Chill on ice for several seconds and transform 5-10 µl of the reaction into 100 µl competent cells.

8. Transformation

The induction of competence of bacterial cells as well as the uptake of exogenous genetic material by these cells from their surroundings was done by using heat-shock transformation

Heat-Shock

1. Thaw cells on ice.
2. Add 1 µl of plasmid DNA.
3. Incubate on ice for 30 min.
4. Heat shock at 42°C for 60 s.
5. Incubate on ice for 3 min.
6. Add 350 µl of SOC media.
7. Incubate at 37°C for 1 h.
8. Plate 100µl on plates supplemented with the appropriate antibiotic.