

**iGEM TU/e 2014**

Biomedical Engineering

Eindhoven University of Technology

Room: Ceres 0.04

Den Dolech 2, 5612 AZ Eindhoven

The Netherlands

Tel. no. +31 50 247 55 59

[2014.igem.org/Team:TU\\_Eindhoven](http://2014.igem.org/Team:TU_Eindhoven)

## Insert + vector ligation

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# 1 Insert + vector ligation

- First determine the amount of vector you will need.
  - Aim for a 100 ng of vector:
 
$$\frac{100 \text{ ng}}{[\text{Vector}] \frac{\text{ng}}{\mu\text{L}}} = x \mu\text{L vector needed}$$
- Now determine the amount of insert you will need:
  - We will need molar equality for the ligation to work.
  - Find the average number of bases in the insert (432bp for the repeat samples):  $N_a$
  - Find the average number of bases in the vector:  $N_b$
  - Molar equality is found by:

$$\frac{100 \text{ ng}}{N_b} \times N_a = X \text{ ng of insert needed}$$

$$N_A = (6,022 \ 141 \ 29 \pm 0,000 \ 000 \ 27) \times 10^{23} \text{ mol}^{-1}$$

- Instead of aiming for molar equality (to increase success rate) use 5 times X ng of the insert.

$$\frac{5 * X \text{ ng}}{[\text{Insert}] \frac{\text{ng}}{\mu\text{L}}} = Y \mu\text{L of insert needed.}$$

Component	Quantity/mass/final concentration	Volume (uL)
H <sub>2</sub> O		Fill up to 20 μL
T4 Ligase Buffer 10x		2 μL
Insert DNA	X ng (stock ... ng/μL)	Y μL
Vector DNA	100 ng (stock ... ng/μL)	X μL
T4 DNA ligase	0.01 μL (400 U/μL stock)	1 μL
<i>Total</i>		20 μL

**Important:** add the ligase at the very end!!

- Gently mix the solution by pipetting up and down
- Incubate at 25°C for 2 hours in a PCR machine
- Heat inactivate for 10 min at 65°C