

# Cloning – Gateway BP-Reaction

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Version: 1.0 - [Print Version \(.PDF\)](#)

**ATTENTION:** This is expensive. One reaction as described here is ca. 7 Euro!  
If possible use Gateway BP-Reaction II because the enzyme is more stable, cheaper and available in smaller amounts.

1. Design PCR-Primers with attB1.1 and attB2.1 sites
2. Gel-purify you PCR product
3. Make sure you have a PCR product with attB1.1 and attB2.1 and one DONR™ clone
4. Measure the DNA concentration of both constructs
5. Calculate the amount in ng needed of PCR product (25 fmol each):  
**ng needed = (length of the PCR product in bp) x 0.0165**
6. Calculate the volume in µl needed of DONR™ plasmid (75 ng):  
**µl needed = 75 ng needed / (concentration in ng/µl)**
7. Calculate the volume in µl needed of DONR plasmid (75 ng):  
**µl needed = 75 ng needed / (concentration in ng/µl)**
8. Prepare in a new eppi the Gateway® reaction:

PCR-product	( ? ng)
pDONR™-vector	(75 ng)
add water to a total volume of	<b>3µl</b>
9. Remove the 5 x **BP-Clonase™** Reaction Buffer from -80°C
10. Pipett **1 µl** of this **BP-Clonase™** Reaction Buffer to the Gateway® reaction
11. Remove the 5 x **BP-Clonase™** enzyme mix from -80°C
12. Pipett **1 µl** of this **BP-Clonase™** enzyme mix to the Gateway® reaction
13. Store the enzyme mix and buffer immediately at -80°C !!!
14. Incubate at room temperature for **1 hour** or over night
15. Add **0,5 µl** of **ProteinaseK** solution and incubate for **10 min** at **37°C (IMPORTANT)**
16. Transform DH5α bacteria
17. Plate bacteria with proper antibiotic selection

## Materials needed:

BP-Clonase™ enzyme mix (# 11789-013) by [Invitrogen](#)

## **Commented Protocol:**

### **1. Design PCR-Primers with attB1.1 and attB2.1 sites**

Check out the protocol on primer design. In short - you should take care of some things. The Gateway® clones have a reading frame which should be kept. Design primers that the PCR product starts with a ATG and ends with a STOP-codon or the last aminoacid (if you want to make a fusion protein). [Primer3plus](#) is a powerful tool helping you to pick primers with the right annealing temperature which should be 60°C. Try to avoid self similarity and other things as usual, but because you are very limited in the position of the primers (its start and stop), I only care about annealing temperature and give it a try. Then just add to the primer which binds the start codon the attB1.1-sequence at his 5' End . To the primer which binds the stop codon or the last aminoacid add the attB2.1-sequence at his 5' End . The open reading frame is indicated and you should change the last two NN to code for an aminoacid of your choice. Good luck for the PCR! Because of the long 5' overhang and the restrictions on picking the primers, getting the PCR to work can be tricky.

There are improved and better att sites available:

attB1.1 GGG-GCA-ACT-TTg-tac-aaa-aaa-gtt-gNN

attB2.1 GGG-GCA-ACT-TTG-TAC-AAC-aaa-gtt-gNN

The original att sites:

attB1 GGGG-ACA-AGT-TTg-tac-aaa-aaa-gca-ggc-tNN

attB2 GGGG-ACC-ACT-TTG-TAC-AAG-aaa-gct-ggg-tNN

### **2. Gel-purify you PCR product**

Purification of the PCR-product is needed to get rid of smaller side-products, which remove primer-dimers which can result in false positive colonies. Remember that you want to clone DNA, so the cutting should be made on the weakest UV-light available and as fast as possible. And of course you **NEVER** make a picture of the gel before. Use the kit for gel-purification available in your lab.

### **3. Make sure you have a PCR product with attB1.1 and attB2.1 and one DONR™ clone Gateway®**

You need a PCR product with the attB1.1 and attB2.2 and the DONR™ vector MUST have attP1 and attP2 sites, or it will not work.

The amount of plasmids is not soo important as in a multiple Gateway® reaction, because it is more efficient. If you want to optimize you can calculate equimolar amounts of both plasmids as described in the How to measure DNA. Here we use double the amount of DEST-vector, because most of the ones we use are round and about double the size of the ENTR™ clones.

#### **4. Measure the DNA concentration of both constructs**

The amount of ENTR™ is not so important as in a multiple Gateway® reaction, because it is more efficient. If you want to optimize you can calculate equimolar amounts of both plasmids as described in the multiple Gateway® protocol. Here we use double the amount of DEST-vector, because most of the ones we use are round and about double the size of the ENTR™ clones.

#### **5. Calculate the amount in ng needed of PCR product (25 fmol each):**

$\text{ng needed} = (\text{length of the PCR product in bp}) \times 0.0165$

#### **6. Calculate the volume in $\mu\text{l}$ needed of DONR™ plasmid (75 ng):**

$\mu\text{l needed} = 75 \text{ ng needed} / (\text{concentration in ng}/\mu\text{l})$

The DONR-vector should be tested for low background colonies (due to a mutated ccdB-gene) when transferred in DH5alpha-bacteria.

#### **7. Calculate the volume in $\mu\text{l}$ needed of DONR plasmid (75 ng):**

$\mu\text{l needed} = 75 \text{ ng needed} / (\text{concentration in ng}/\mu\text{l})$

#### **8. Prepare in a new eppi the Gateway® reaction:**

PCR-product	( ? ng)
pDONR™-vector	(75 ng)
add water to a total volume of	<b>3<math>\mu\text{l}</math></b>

#### **9. Remove the 5 x BP-Clonase™ Reaction Buffer from -80°C**

It is most efficiently mixed by pipetting up and down, do not vortex.

#### **10. Pipett 1 $\mu\text{l}$ of this BP-Clonase™ Reaction Buffer to the Gateway® reaction**

#### **11. Remove the 5 x BP-Clonase™ enzyme mix from -80°C**

It is most efficiently mixed by pipetting up and down, do not vortex.

#### **12. Pipett 1 $\mu\text{l}$ of this BP-Clonase™ enzyme mix to the Gateway® reaction**

This is expensive stuff don't leave it to rot in the ice-bucket!

#### **13. Store the enzyme mix and buffer immediately at -80°C !!!**

The enzymes loses 50% activity after 15 freeze-thaw cycles. The advantage of BP-Clonase™II would be that it can be stored at -20 °C because it contains already the buffer. This is expensive stuff don't leave it to rot in the ice-bucket!

#### **14. Incubate at room temperature for 1 hour or over night**

Incubation over-night will enhance the reaction ca. 5-10 fold. This is especially important for PCR products over 5.000 bp.

#### **15. Add 0,5 µl of ProteinaseK solution and incubate for 10 min at 37°C (IMPORTANT)**

This step will enhance the reaction ca. **100 fold!!!!**. This is different to the LR-reactions which are only enhanced 2 fold by adding the proteinase K!!!!

#### **16. Transform DH5α bacteria**

For electro competent cells use 1-2 µl, for chemical competent all.

#### **17. Plate bacteria with proper antibiotic selection**

The resulting ENTR™-vectors are kanamycin resistant.

### **Known Issues:**

- The reaction is very efficient. You can obtain about 200 colonies of which about 95 % are correct.
- BP reactions work better with linear templates like PCR-products. If you want to use plasmids, linearize them first with a suitable restriction enzyme.
- The obtained plasmids are big. To check for correct clones digest with Sty I and in parallel with Eco RI and Hind III. Compare the pattern of bands with the predicted band size to find the correct clones.

### **References and Comments:**

The protocol is a evolution of the supplied informations, mainly because they were not clear enough and complex. It uses only half of the recommended amounts to be more cost effective. I did it as described before several times sucessfully.

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