

BBF RFC 92: The GoldenBricks assembly: A standardized one-shot cloning technique for complete cassette assembly

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1. Purpose

The RFC 92 propose a new standard assembly method for the PartsRegistry. The method makes one-shot cloning of a complete eukaryotic or prokaryotic cassette possible in one day while keeping the compatibility with the RFC10 BioBricks assembly.

2. Relation to other BBF RFCs

This RFC 92 is an upgrade of the RFC 10. It makes the BioBricks format compatible with the Golden Gate assembly method described previously for Yeast ORF in the RFC 88, and also in the RFC 28, RFC 53 and RFC 61.

3. Copyright Notice

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4. Introduction

The BioBricks are a collection of parts that can be assembled one by one in a standardized way. Golden Gate [1] is a technique that is capable of assembling several dozens of parts simultaneously. Taking inspiration from both techniques, we developed a new method we call GoldenBricks, merging the advantages of both technique. GoldenBricks make the assembly of a fully working eukaryotic or prokaryotic cassette possible from standard biological parts, while preserving the compatibility with the RFC 10 BioBricks format. The assembly process in itself takes a single day.

The GoldenBricks technique is a one-shot cassette construction using DNA parts coming either from a plasmid distribution or directly from PCR reaction, and engaging 5 or more different parts. GoldenBricks is designed for cloning both eukaryotic and prokaryotic cassette using the same prefixes, suffixes and protocols. If a non canonical part order is required (as for testing the strength of a terminator), it is still possible to assemble GoldenBricks sequentially with the standard BioBricks' technique.

GoldenBricks opens new perspectives for the PartsRegistry. It is faster, cheaper and safer and requires less sequencing runs than BioBrick RFC10 assembly. It is designed to be an efficient cloning techniques for researchers and companies but is also more suitable for less experimented users, such as iGEMers, high school iGEMers or the DIYbio community. The creation of large expression cassettes is up to now cheaper and faster than synthesizing the construct, which would guaranty the interest for a DNA databases over *de novo* synthesis for the years to come. Finally, the method is easily automatable.

5. Design principles

The GoldenBricks cloning technology relies on the type IIS restriction enzyme BsaI that cuts the DNA outside its recognition locus, no matter the sequence present. Therefore, several customized overhangs can be done simultaneously with a single enzyme. In GoldenBricks, the sequence of the overhangs is used to constrain the fragment to be assembled in the order wanted.

Once ligated, BsaI overhangs cannot be re-cut the ligation scar because the cutting site is different from the recognition site. Therefore, it is possible to digest and ligate at the same time, within the same time same tube, without DNA extraction or purification in between.

To keep the compatibility with the existing RFC 10 format, EcoRI, XbaI, SpeI and PstI are incorporated inside the GoldenBricks prefixes and suffixes. The GoldenBricks assembly reconstructs a BioBrick cassette with no illegal restriction site inside. The GoldenBricks parts can also be assembled using the usual BioBricks method even though the ligation scar is a little bit different.

Since the amount of DNA required for GoldenBricks' assembly is in the range of 50 ng, the quantity of DNA plasmid provided in the PartsRegistry distribution is sufficient to carry out several assemblies. The parts can be used out of the distribution plate and it is not necessary to do a preliminary DNA preparation anymore.

Finally, GoldenBricks incorporates an additional screening method. When two GoldenBricks overhangs ligate together, they all form a ligation scar that is digestible by the BbsI enzyme. The correct construction can therefore be checked using a single digestion that cuts between all the parts. Because GoldenBricks mutation rate is extremely low (no DNA modification or UV involved) and the number of clones to sequence can be significantly reduced.

5. Definition of the format

5.1 GoldenBricks for a prokaryotic cassette

A prokaryotic expression cassette is composed of a promoter, a repeat of several RBS (Ribosome Binding Site)-CDS (CoDing Sequence) units and a terminator cloned in a plasmid. The GoldenBrick format differentiates 5 different types of parts that use 10 different prefixes and suffixes. Constraining the assembly of the fragments in the canonical cassette order requires to have 5 types of overhangs (OH1 to OH5). To make the RBS-CDS unit repetition possible, the prefix of the RBS should match the suffix of the protein, that is to say that the overhang 2 is compatible with the overhang 4 (OH2=OH4). We will see later in the principle section how to control the assembly order.

The GoldenBrick prefixes and suffixes for a prokaryotic cassette are:

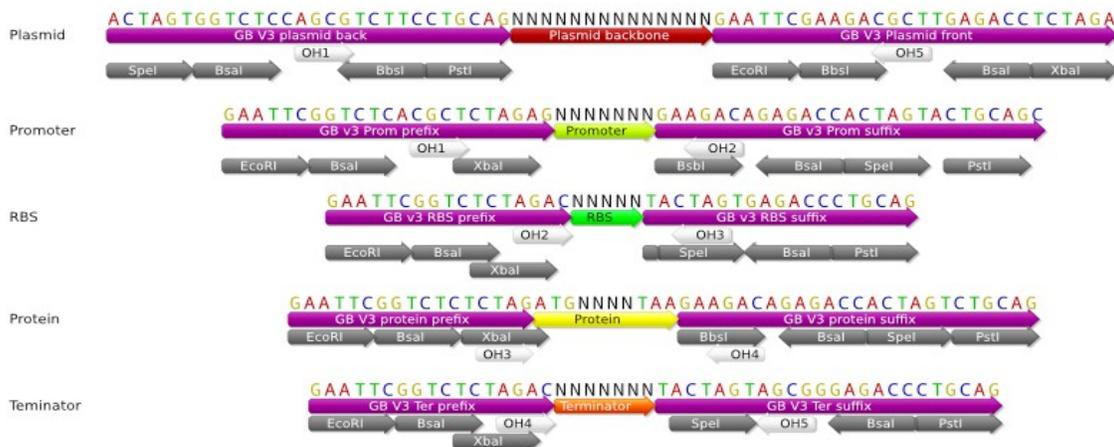


Fig 1: GoldenBrick prefixes and suffixes for a prokaryotic cassette [2]

5.2 GoldenBricks for an eukaryotic cassette

Eukaryotic cassettes are different from prokaryote cassettes, because the initiation of the translation is ensured by the Gpp RNA capping, induced by regulatory elements next to the promoter sequence. mRNAs are usually monocistronic in the genome, even-though the sequence of internal ribosome entry sites (IRES) has been identified in viral polycistronic mRNAs.

Considering that an eukaryotic promoter is equivalent to an prokaryotic promoter fused with a prokaryotic RBS, the GoldenBrick eukaryotic promoters use the prefix of a prokaryotic promoter and the suffix of a prokaryotic RBS. The IRES sequences are equivalent to a prokaryotic RBS. They are cloned between the same prefixes and suffixes.

before the BsaI cut, the equilibrium is driven toward the formation of the product by placing the mix alternatively between 37°C where BsaI cuts and 16°C where the T4 ligase ligates the fragments. Increasing the number of cycles increases the efficiency of the assembly but also increases the background level.

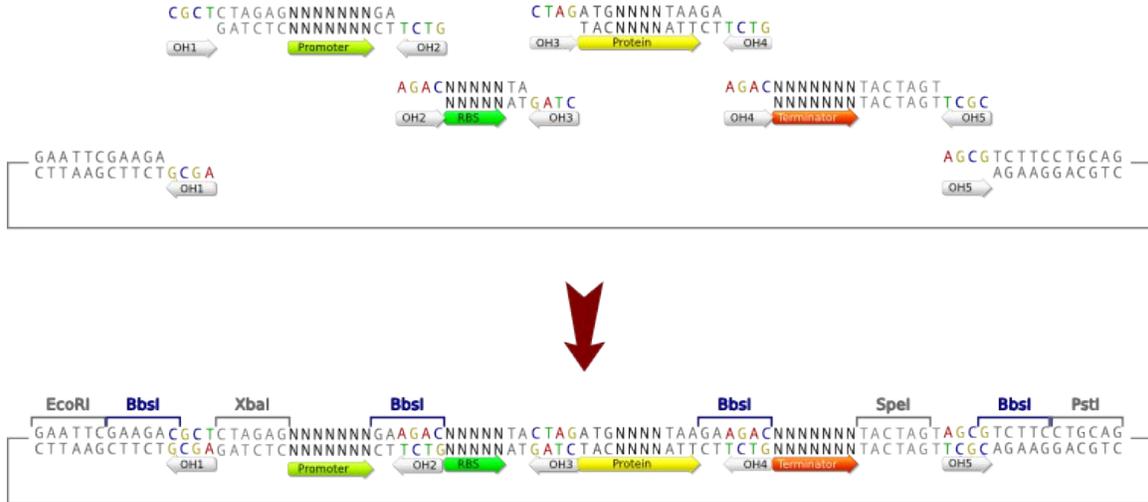


Fig 3: Assembly of the different DNA parts. The order of the parts in the construct is constrained by the overhangs' sequence.

The DNA mix is then transformed and plated with the antibiotic of the receiver plasmid.

6.2 Creation of a polycistron

Using the compatibility of overhangs 2 (OH2) and overhangs 4 (OH4) it is possible to assemble a polycistronic cassette using the GoldenBricks method. However, if one mixes all the RBSs and CDSs together at the same time, it is not possible to control which RBS ligates to which CDS, the order and the number of gene in the cassette. The sequential assembly protocol is designed to go around this limitation.

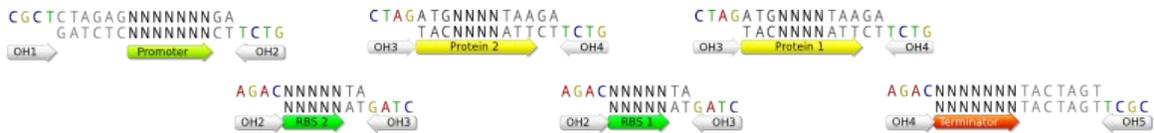


Fig 4: Assembly of a prokaryotic polycistronic mRNA

6.3 Control over the assembly order through sequential assembly.

In order to increase the probability of one peculiar RBS to match with one peculiar CDS, to diminish the formation of side products or in order to control RBS-CDS polymerization, we recommend the use of the sequential assembly. Sequential assembly consist in adding the DNA fragments one after the other in the mix rather than mixing them altogether at the beginning.

Two distinct approach wan be used:

- The sequential assembly, which consists in adding the terminator + CDS 1 + RBS 1 for a few cycles before adding the gene 2 + RBS 2, etc.. and then finally the promoter and the plasmid for a few more cycles. Sequential assembly significantly decrease the background level even in the case of one gene cassette.
- The parallel assembly which consists in assembling separately the RBS-CDS units and then pool the tube for a few cycles. The promoter and the plasmid are added in the end and the mix undergo a few more cycles.

6.4 Theoretical sides products to avoid

The advantage of GoldenBricks is that it is possible to make polycistronic mRNAs in a single cloning step. However, the compatibility of the overhangs 2 and 4 gives also the possibility for the promoter to ligate with a terminator.



Fig 5: Anticipated termination by the ligation of the promoter and the terminator.

The importance of this side-product can be significantly reduced by using a sequential assembly, adding the promoter and the plasmid later in the mix. This side-product can also be very-easily screened out by colony PCR.

6.6 Automatic assembly and automation

GoldenBricks standard is also fully automatable, because of the small DNA amount required and the absence of DNA purification step.

7. Control of the construction

7.0 Negative cloning control

GoldenBrick has a lower yield compared to standard assembly because it assembles more than two fragments at the time. This is compensated by the increase of speed and easiness in the screening process.

The original plasmid contains a red negative cloning control, containing on an RFP cassette J04450, visible with naked eyes. The last digestion and inactivation steps in the

protocol is often enough to eliminate the original receiver plasmid traces. Otherwise, the red clones it produces can be screened out by eyes.

7.1 Control using colony PCR

The GoldenBricks plasmids derives from the pSB1X3 backbone series (where X is the antibiotic resistance). It is therefore possible to do colony PCR using the VR and VF2 primer. Colony PCR helps to identify the brick that have the expected amount of fragment assembled in the cassette, without having to mini-prepare many clones.

7.2 Control using the generic BbsI digestion

After colony PCR screening, the candidates are inoculated and mini-prepped. The GoldenBricks format is designed with a single digestion screening tool to help finding the correct clone.

Goldenbricks prefixes and suffixes are designed so that once, ligated, the scar generated can be re-digested using the BbsI enzyme (see fig. 3). BbsI recognition sequence is quite common in CDSs in general. The enzyme cuts in between the assembled elements and also quite often also in the CDS, providing an informative digestion pattern.

7.3 Coupled control

To save time and mini-prepare, it is possible to apply a mixed screening strategy. Half of the colony PCR products are deposited on gels. The other half of the products that have the correct size are further digested with BbsI and run on gel for identification of expected clones.

8. Protocol

Simple assembly: A 60 fmol equimolar ratio of the different fragments were mixed with 15 units of T4 ligase and 5 units of BsaI in T4 ligase buffer, 20 μ L total. The tube was placed in a thermocycler that goes for 30 cycles 37°C for 2 min and 16°C for 5 min. In the end, a final digestion step of 5 min at 50°C and an a final inactivation of 5 min at 80°C was carried to eliminate the background before transformation.

Sequential assembly: The RBSs, CDS and the terminator were mixed in a 60 fmol ratio with 15 units of T4 ligase and 5 units of BsaI in (20 - promoter and plasmid volume) μ L of T4 ligase buffer. After 20 cycles (2 min at 37°C and 5 min at 16°C), the promoter and plasmid were added for another 10 cycles before proceeding to the final digestion, heat inactivation and transformation.

Semi-parallel assembly: The RBS1 with the CDS1 and terminator in one tube, and the RBS2 and CDS2 in a second one were mixed with 7 units of T4 ligase and 1.5 unit of

11. Availability

A few GoldenBrick parts are already available in the PartsRegistry for testing:

- [K812050](#): A GoldenBricked version of pSB1C3 with J04450
- [K812051](#): A GoldenBricked version of pSB1K3 with J04450
- [K812053](#): A GoldenBricked version of the strong RBS B0034
- [K812054](#): A GoldenBricked version of the RFP E1010
- [K812055](#): A GoldenBricked version of the terminator B0015
- [K812056](#): A GoldenBricked version of the pLac R0010 promoter
- [K812057](#): A GoldenBricked of an sfGFP protein

12. Additional information

Additional information can be found in the GoldenBricks page of our Team wiki (<http://2012.igem.org/Team:Evry/GB>).

13. Author's Contact Information

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Anna Młynarczyk, Aurore THELIE, Thomas Landrain, Andrew Tolonen, Nicolas Pollet, Jean-Loup Faulon and Alfonso Jaramillo are iGEM 2012 Evry team advisors and instructors.

14. References

[1] Engler C, Gruetzner R, Kandzia R, Marillonnet S (2009) Golden Gate Shuffling: A One-Pot DNA Shuffling Method Based on Type IIs Restriction Enzymes. PLoS ONE 4(5): e5553. doi:10.1371/journal.pone.0005553

[2] Image partially generated with the Geneious software version 5.4 created by Biomatters. Available from <http://www.geneious.com/>