

Stupar Lab Meeting

Multiplex CRISPR Design

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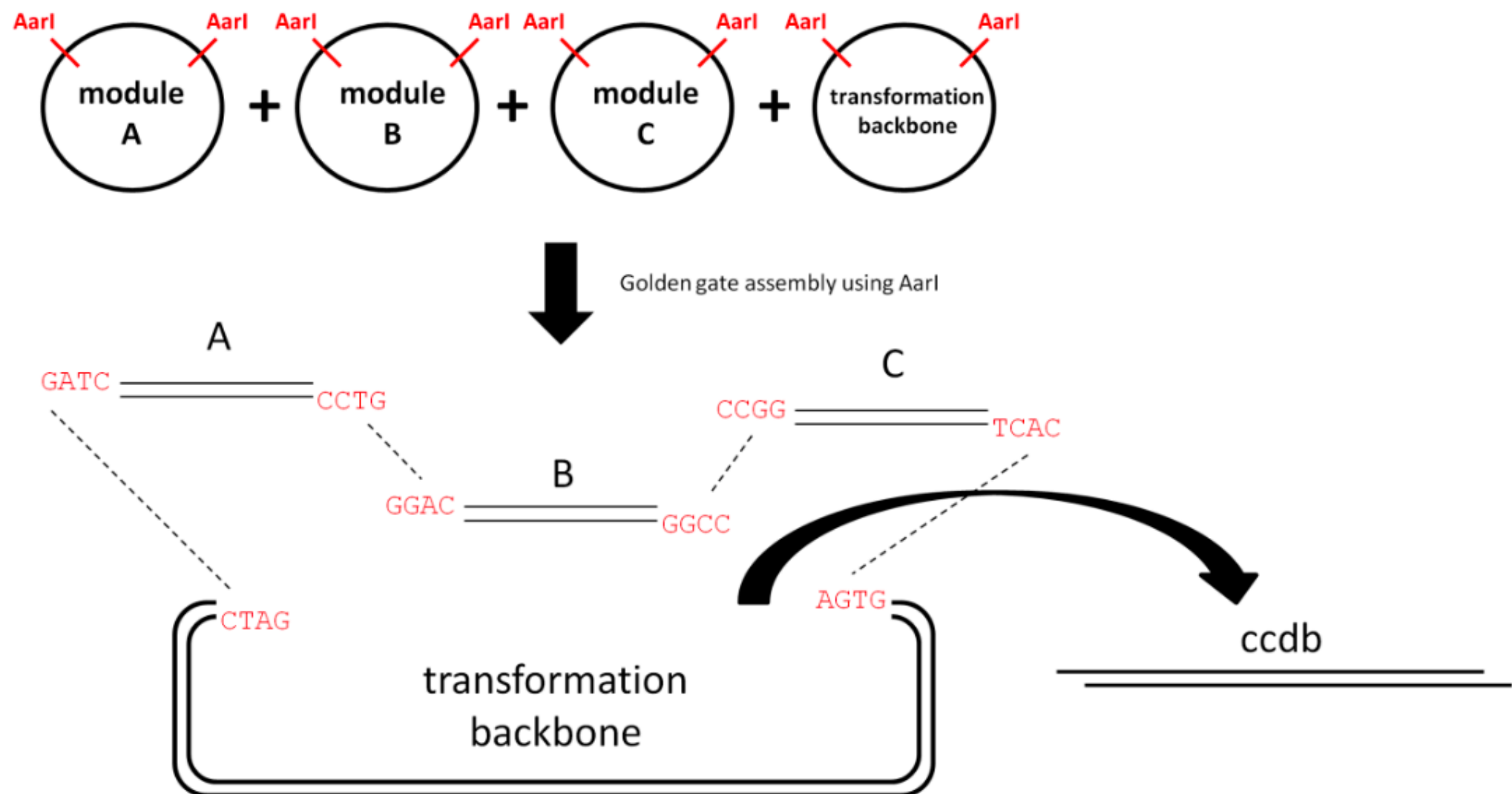
Slides: https://z.umn.edu/crispr_slides

Publication in *The Plant Cell*

- Čermák et al. (2017). A Multi-purpose toolkit to enable advanced genome engineering in plants. *The Plant Cell*.
- DOI: <https://doi.org/10.1105/tpc.16.00922>
- See the paper for a description of the protocols, and the details of the methods
- Website: <http://z.umn.edu/crisprmultiplex>
Note: Requires Javascript

Reagent Toolkit and Protocol

- A modular set of vectors that are built separately, then combined with a Golden Gate protocol



Reagent Toolkit and Protocol

Module	Number of Vectors	Description
Module A	61	Ready-to-use, with Cas9 or GFP cassettes
Module B	22	Add additional gRNAs or TALEN monomer
Module C	22	Add additional gRNAs, donor template, or expression cassettes
Transformation Backbone	31	Will be transformed into plant

Reagent Toolkit and Protocol

- Set of five protocols, with variants depending on what is being assembled (gRNAs, or TALENs)

Module B

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Plasmid ID	Gene	Promoter	Terminator	Protocols
pMOD_B0000	None	None	None	5
pMOD_B2000	TALEN_2 backbone with Esp3I ccdB cassette for repeat cloning	None (begins with P2A to be fused to TALEN 1)	HSP	1A , 5
pMOD_B2101	SapI ccdB cassette for cloning multiple gRNA protospacers with Csy4 spacers	35S	35S	3A , 3S2 , 5
pMOD_B2103	SapI ccdB cassette for cloning multiple gRNA protospacers with Csy4 spacers	CmYLCV	35S	3A , 3S2 , 5
pMOD_B2103b	SapI ccdB cassette (promoter in the backbone, not in the assembly) for cloning multiple gRNA protospacers with Csy4 spacers	CmYLCV	35S	3A , 3S1 , 3S2 , 5

Vector Selection

- Use drop-down menus to choose your vector
- Menus automatically update, and it will serve a vector map as a GenBank Flat File

Vector Selection

Enter your design with the drop-down menus.

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Step 1:	<div>Module A</div>
Step 2:	<div>AtCas9_H840A (Nickase)</div>
Step 3:	<div>35S</div>
Step 4:	<div>Please Submit Below</div>
Step 5:	<div></div>
	<div>Submit</div>

Primer and Map Construction

- Accepts FASTA file of target sequences
- Select target vector, promoter, restriction enzyme, and splicing system

Multi-gRNA Array Assembly - Primer Design and Map Construction

Please enter the following design parameters into the form. Note that not every vector from the Vector Selection page is available in this construction tool.

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Targets (FASTA):	<input type="button" value="Choose File"/> favorite targets.fasta
Target Vector:	<input type="text" value="pMOD_B2103"/>
Promoter System:	<input type="text" value="35S"/>
Restriction Enzyme:	<input type="text" value="BsaI"/>
Splicing System:	<input type="text" value="Csy4"/>
	<input type="button" value="Submit"/>

Primer and Map Construction

Uploaded! Your file had 6 target sequences.

You are using the 35S promoter, the BsaI restriction enzyme, and the Csy4 splicing system. You have chosen pMOD_B2103 as your target vector.

Download
GenBank File

Primer Designs

PCR Reaction 1

```
>o35S
TGCTCTTCGCGCATGGAGTCAAAGATTCAA
>CSY_gRNA11
TGGTCTCCTGGATCTATCATCTGCCTATACGGCAGTGAAC
```

PCR Reaction 2

```
>REP_gRNA11
TGGTCTCATCCAGATGTTCCGTTTTAGAGCTAGAAATAGC
>CSY_gRNA13
TGGTCTCCGAAGAAGAAGAACTGCCTATACGGCAGTGAAC
```

PCR Reaction 3

```
>REP_gRNA13
TGGTCTCACTTCAGACACGAGTTTTAGAGCTAGAAATAGC
>CSY_gRNA16
TGGTCTCCATATAATGCACCCTGCCTATACGGCAGTGAAC
```

Vector Map

[\[Download\]](#)

```
LOCUS      pMOD_B2103_favorite_targets_fasta
DEFINITION MODULE B with CmYLCV:SapI cddb cassette for cl
            spacers - Csy4 .
ACCESSION  urn
VERSION    urn
KEYWORDS   .
SOURCE     .
            ORGANISM .
            .
FEATURES             Location/Qualifiers
     rep_origin        1..857
                        /modified_by="User"
                        /label="High Copy Ori"
     promoter          878..1414
                        /note="35S Promoter"
     misc_feature       1422..1441
                        /note="Csy4"
     misc               1442..1461
                        /note="gRNA11"
     misc_feature       1462..1537
                        /note="gRNA Repeat"
     misc_feature       1538..1557
                        /note="Csy4"
     misc               1558..1577
                        /note="gRNA13"
     misc_feature       1578..1653
                        /note="gRNA Repeat"
```