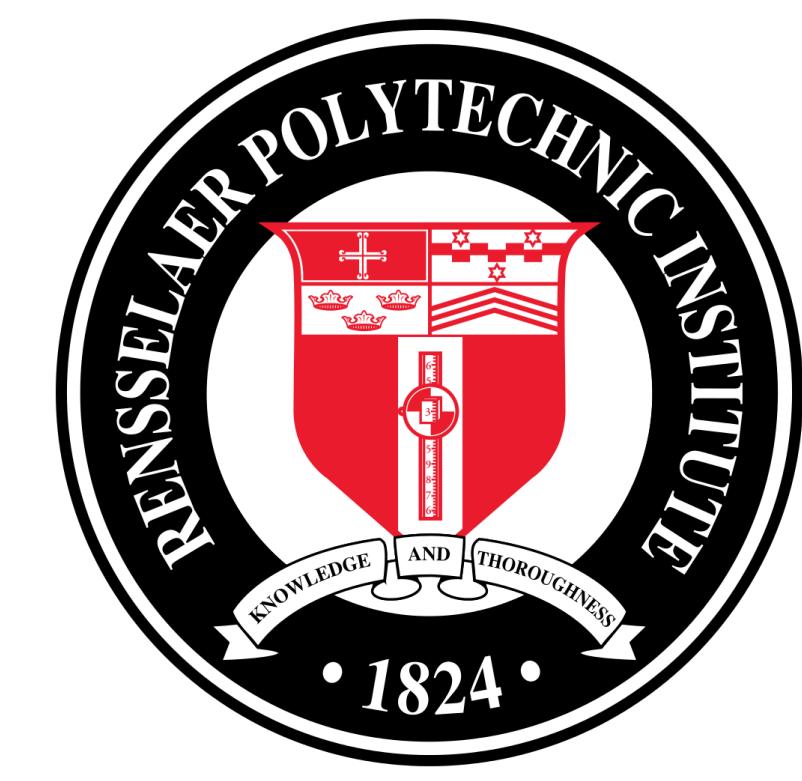


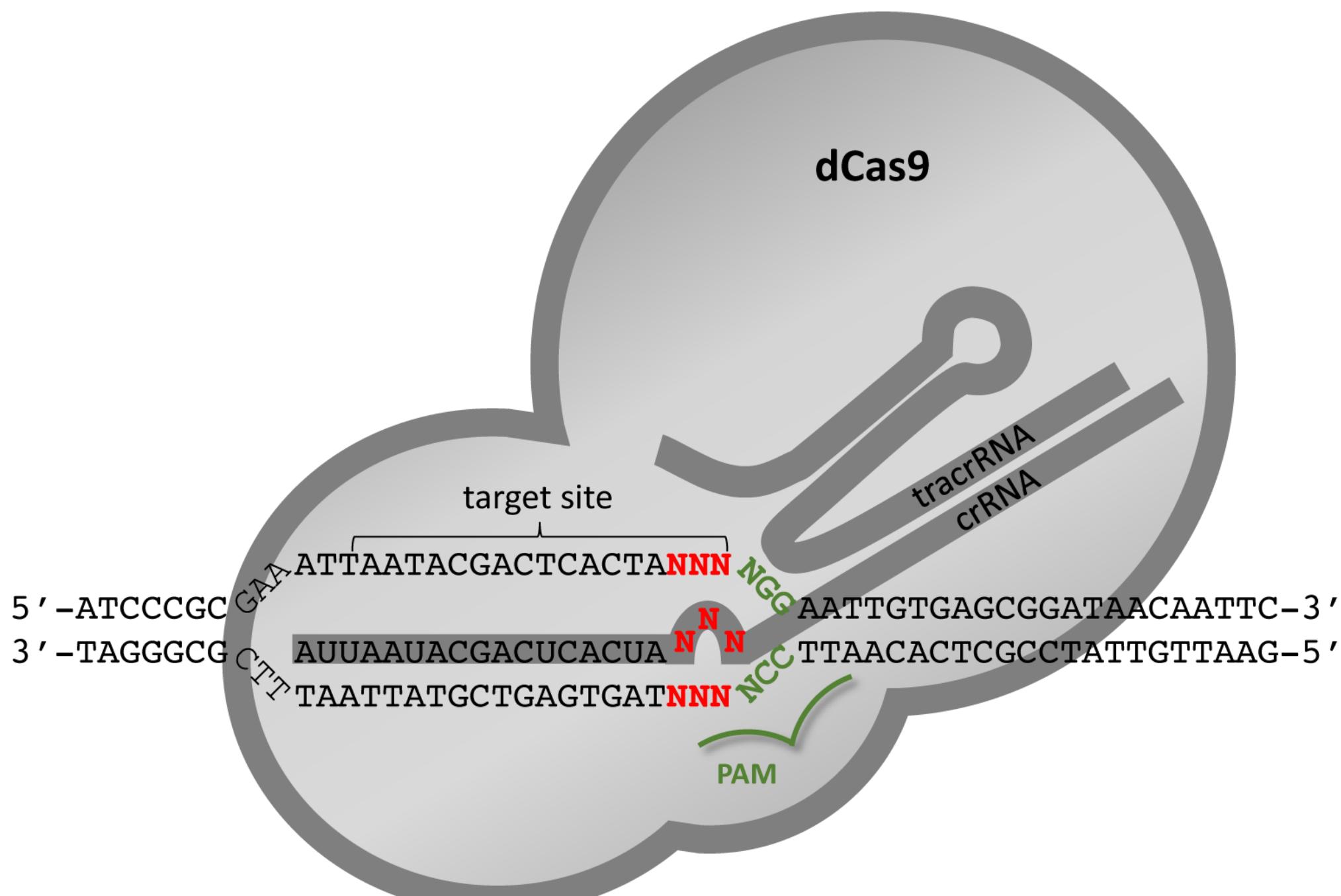
CRISPR Golden Gate Assembly Design Automation

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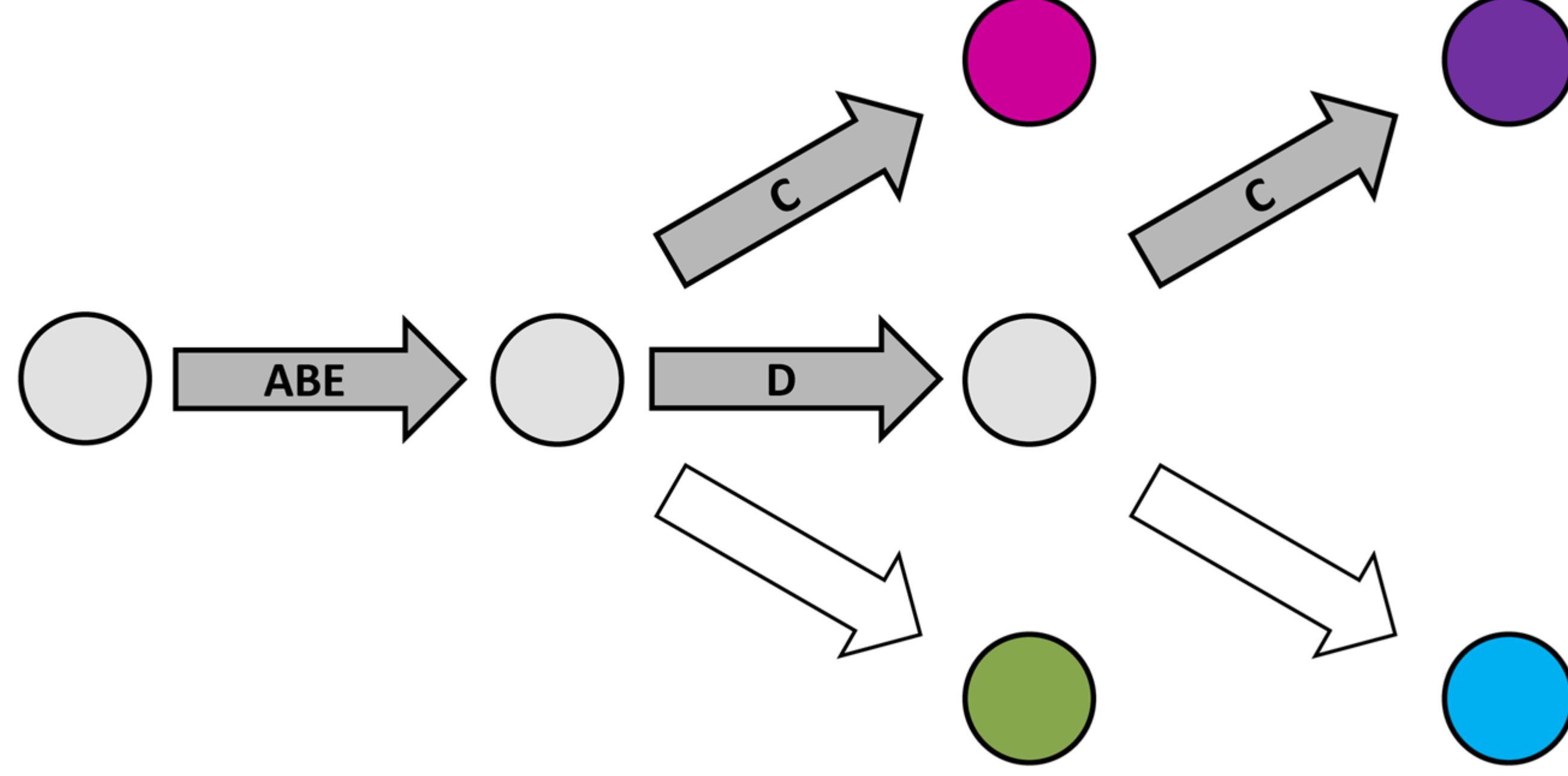
Background

A RNA-guided, DNA-binding protein known as dCas9 can be harnessed for metabolic engineering in *E. coli* by silencing genes that compete with the pathway of interest.

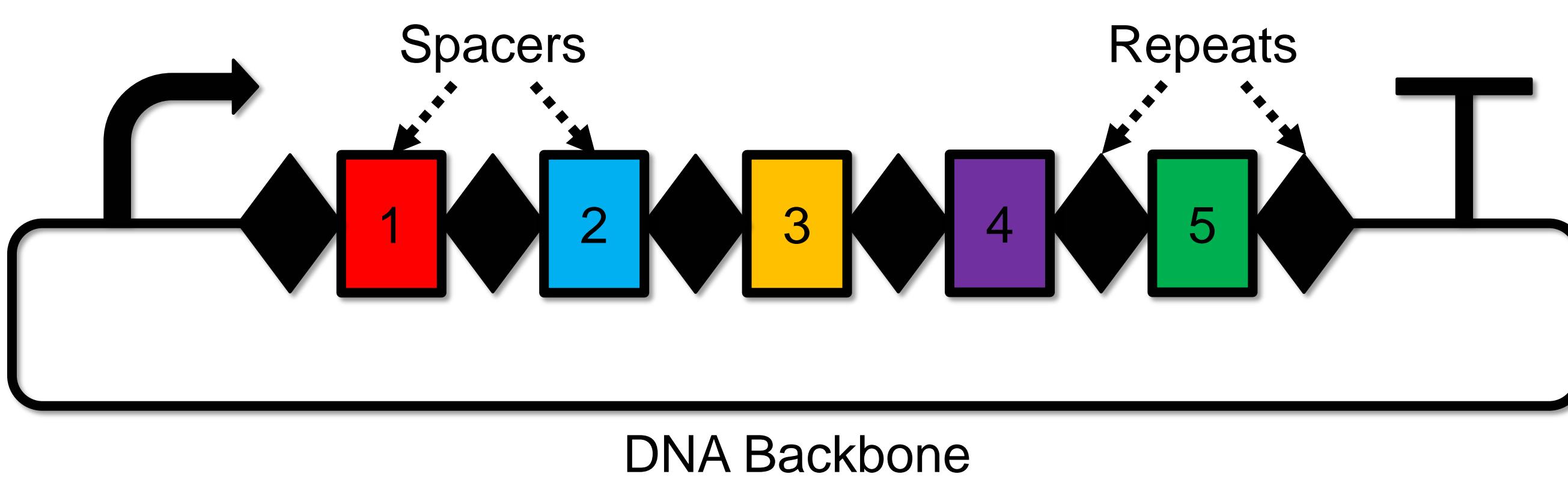


Ensemble of the violacein biosynthetic pathways and products. Major products include **prodeoxyviolacein**, **proviolacein**, **deoxyviolacein**, and **violacein**.

Pathway Ensemble

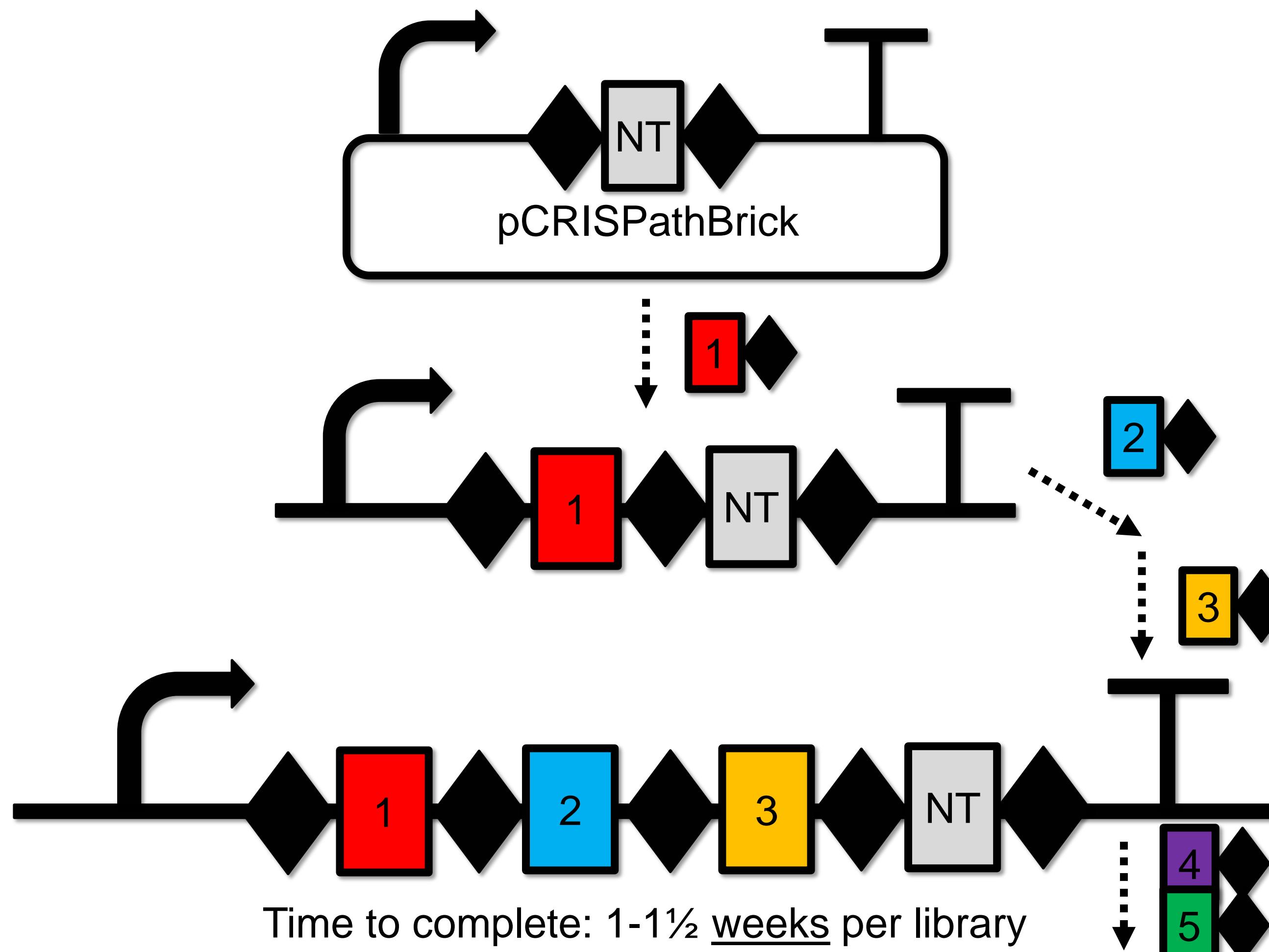


The instructions guiding dCas9 to appropriate binding locations are encoded in a DNA element known as a CRISPR array, composed of **spacers** which are regularly interspersed with short sequences known as **repeats**.

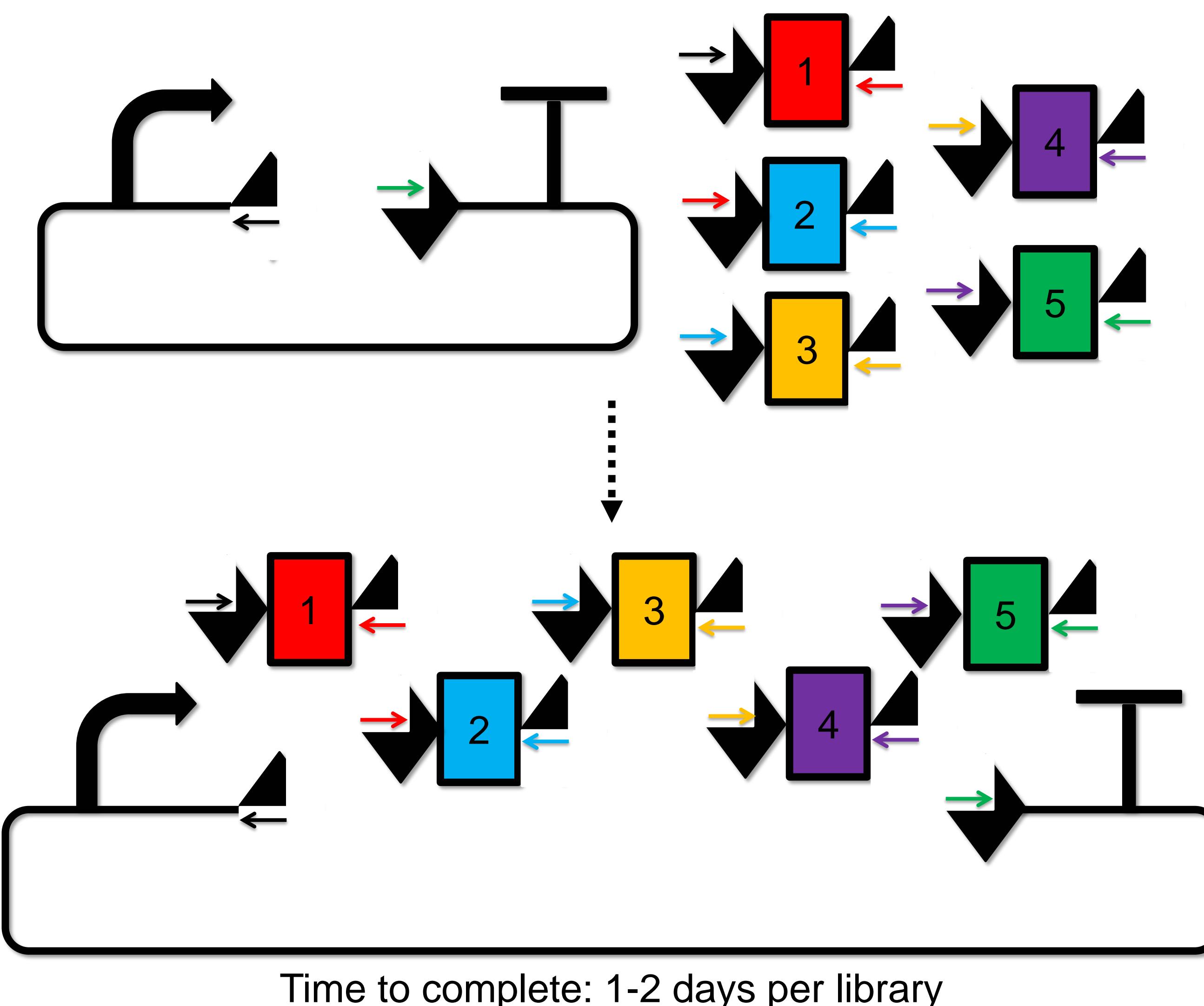


Constructing CRISPR Arrays

Construction of a CRISPR array has traditionally been a cumbersome task, while purchasing the arrays can be prohibitively expensive. Below is a typical array construction method, which requires a round of cloning for each spacer.



The Golden Gate assembly strategy allows researchers to construct libraries of CRISPR arrays in a one-pot reaction by the method shown below.



Automation and Implications

We have designed a MATLAB script which automates the selection of overhangs within the CRISPR repeat region. The algorithm has been successfully applied to construct Type II-A (Cas9) and Type V-A (Cpf1) CRISPR arrays.

DNA Prior to any cutting:

TCTAATTCTACGTTGTAGA [D30] CTAATTCTACGTTGTAGAT
AGATTAAGATGCAAACATCTA [D30] GATTAAGATGCAAACATCTA

DNA backbone after cutting out the dropout:

TCTAATTCTACGTTG
AGATTAAGATGCAAACATCTA
CTAA TTTCTACGTTGTAGAT
AAAGATGCAAACATCTA

Primer Orientation:

5'-GGCGAAGAC

TCTAATTCTACGTTG
AGATTAAGATGCAAACATCTA

1 TCTAATTCTACGTTGTAGAT
AGATTAAGATGCAAACATCTA
CAGAAGCCGG-5'

PCR Amplicon:

GGCGAAGACTGTAGAT
CCGGCTTCTGACATCTA

TCTAATTCTACGTTGGTCTCGGCC
AGATTAAGATGCAAACCAGAAGCCGG

Digested Amplicon:

TAGAT
A

TCTAATTCTA
AGATTAAGATGCAA

Primer Orientation:

5'-GGCGAAGAC

TCTAATTCTACGTTG
AGATTAAGATGCAAACATCTA

2 TCTAATTCTACGTTGTAGAT
AGATTAAGATGCAAACATCTA
CAGAAGCCGG-5'

PCR Amplicon:

GGCGAAGACTACGTTGTAGAT
CCGGCTTCTGATGCAAACATCTA

TCTAATTCTACGGTCTCGGCC
AGATTAAGATGCCAGAAGCCGG

Digested Amplicon:

CGTTTG
ACATCTA

TCTAATT
AGATTAAGAT

Primer Orientation:

5'-GGCGAAGAC

TCTAATTCTACGTTG
AGATTAAGATGCAAACATCTA

3 TCTAATTCTACGTTGTAGAT
AGATTAAGATGCAAACATCTA
CAGAAGCCGG-5'

PCR Amplicon:

GGCGAAGACTTCTACGTTGTAGAT
CCGGCTTCTGAAAGATGCAAACATCTA

TCTAATTGTCTCGGCC
AGATTAACAGAAGCCGG

Digested Amplicon:

TCTACGTTGTAGAT
GCAAACATCTA

T
AGATT

References

- Cress, B.F., Toparlak, O.D., Guleria, S., Lebovich, M., Stieglitz, J.T., Englaender, J.A., Jones, J.A., Linhardt, R.J. and Koffas, M.A., 2015. CRISPathBrick: modular combinatorial assembly of type II-A CRISPR arrays for dCas9-mediated multiplex transcriptional repression in *E. coli*. *ACS synthetic biology*, 4(9), pp.987-1000.
- Cress, B.F., Jones, J.A., Kim, D.C., Leitz, Q.D., Englaender, J.A., Collins, S.M., Linhardt, R.J. and Koffas, M.A., 2016. Rapid generation of CRISPR/dCas9-regulated, orthogonally repressible hybrid T7-lac promoters for modular, tuneable control of metabolic pathway fluxes in *Escherichia coli*. *Nucleic acids research*, p.gkw231.