

Genome Editing and Engineering

Course No: BT-637



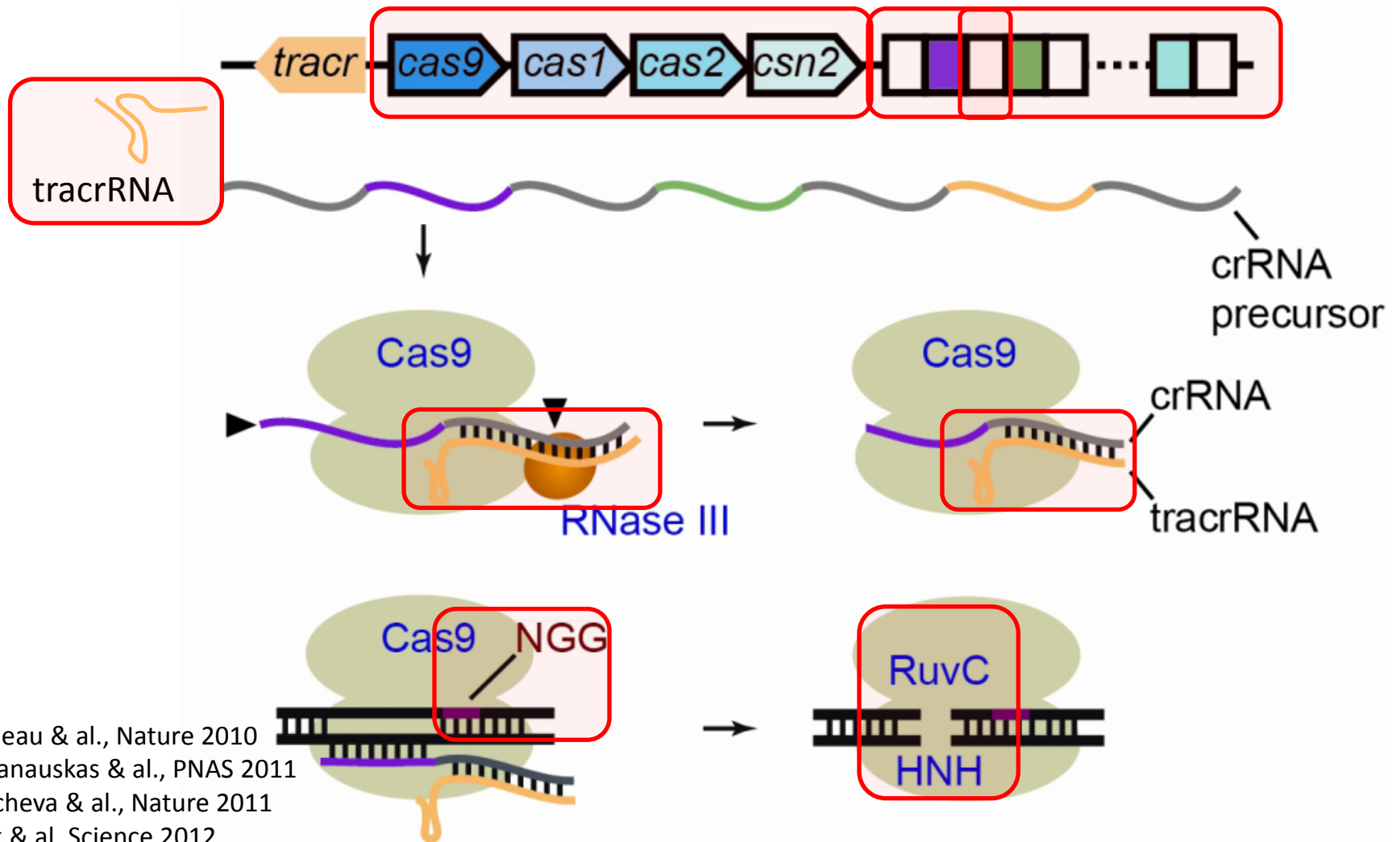
LECTURE-17

Dr. Kusum K. Singh

Department of Biosciences and Bioengineering

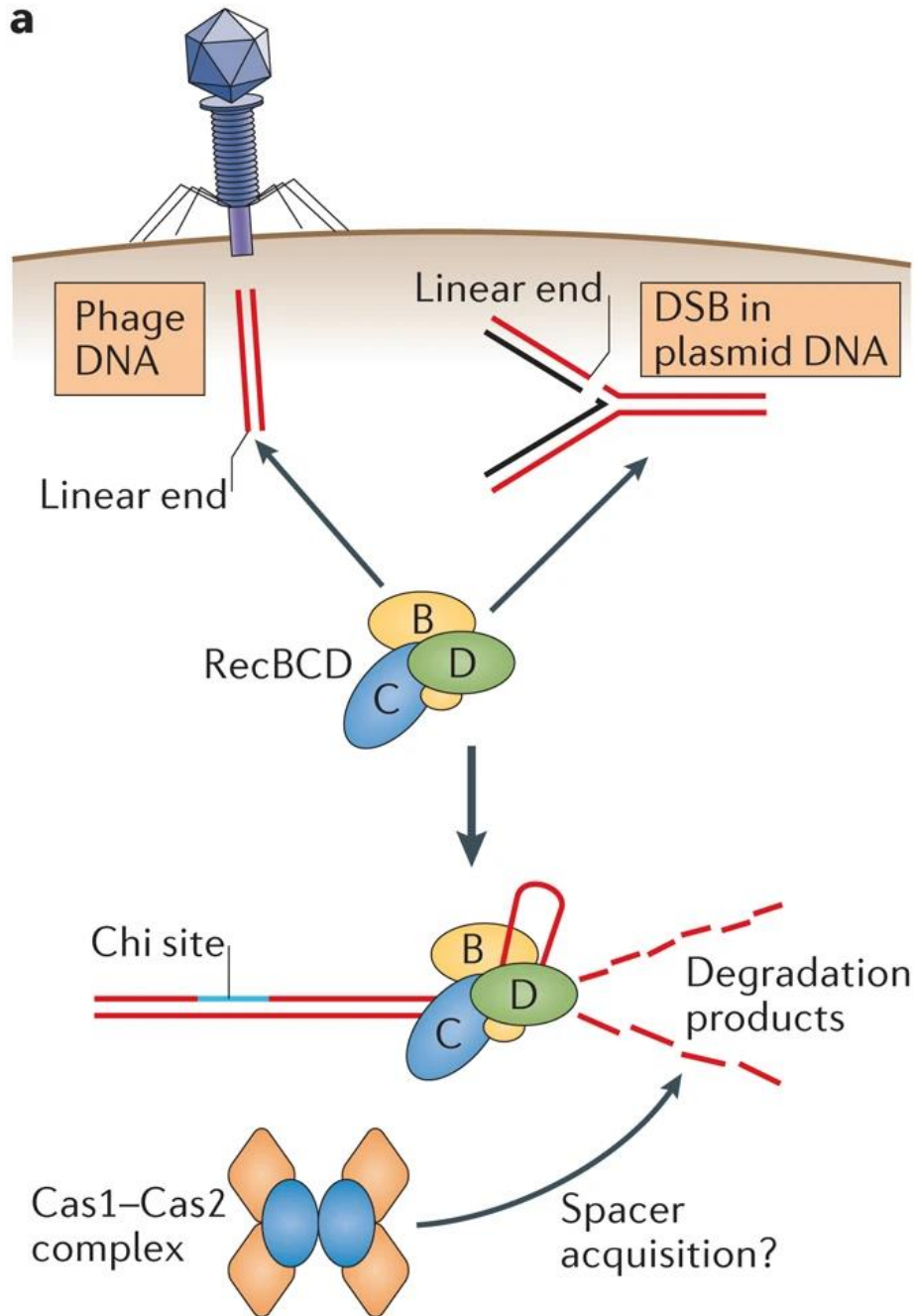
Indian Institute of Technology Guwahati

S.pyogenes type II-A system

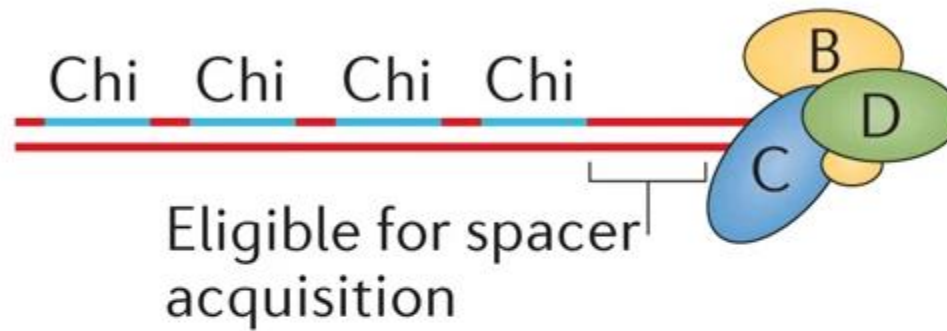


Garneau & al., Nature 2010
Sapranaukas & al., PNAS 2011
Deltcheva & al., Nature 2011
Jinek & al, Science 2012
Gasiunas & al., PNAS 2012
Sternberg & al., Nature 2014

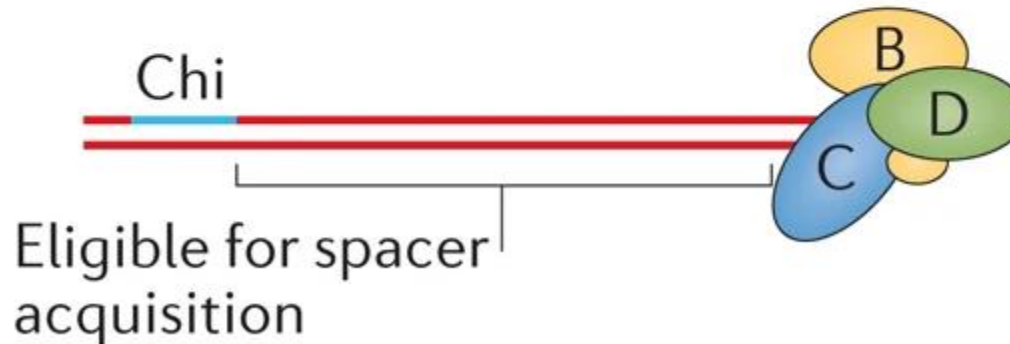
a

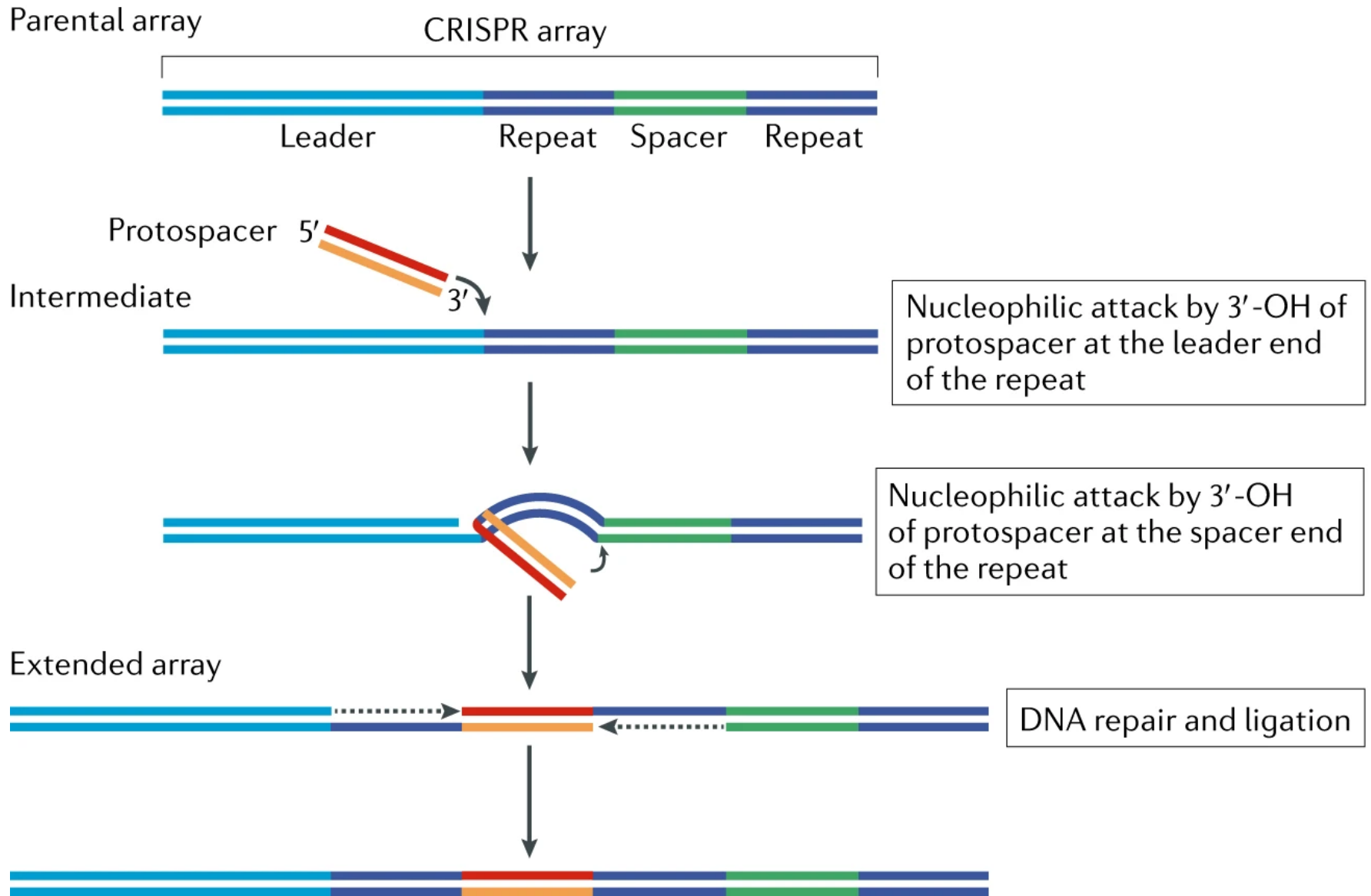


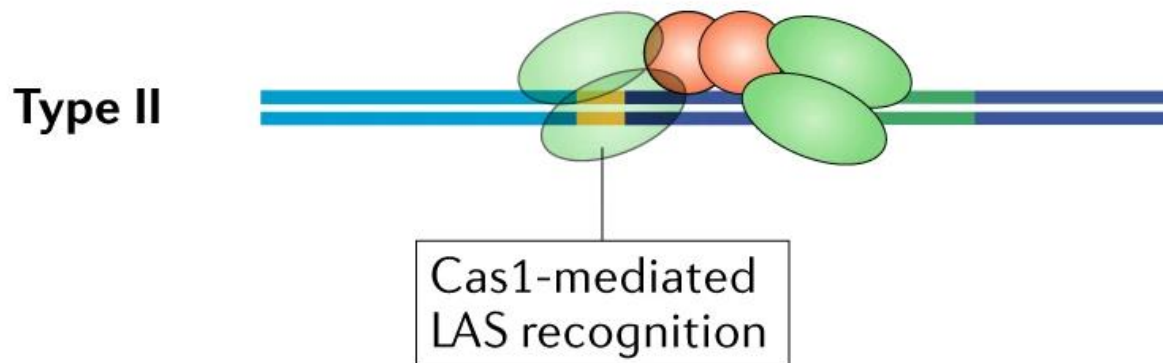
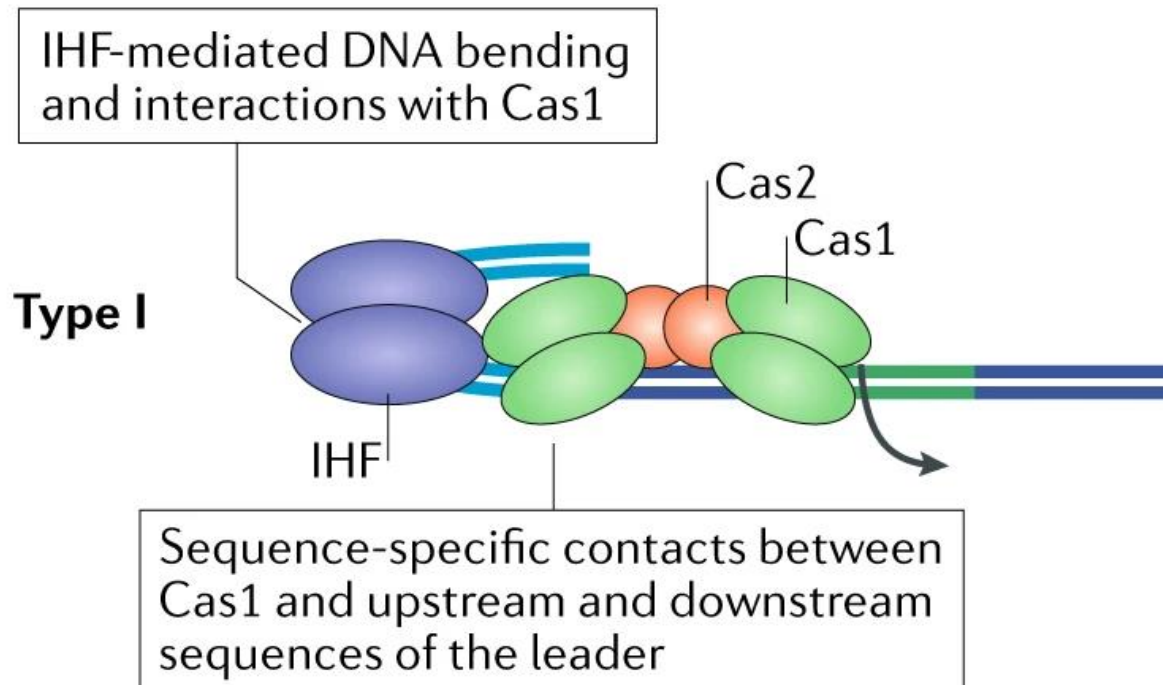
Self DNA

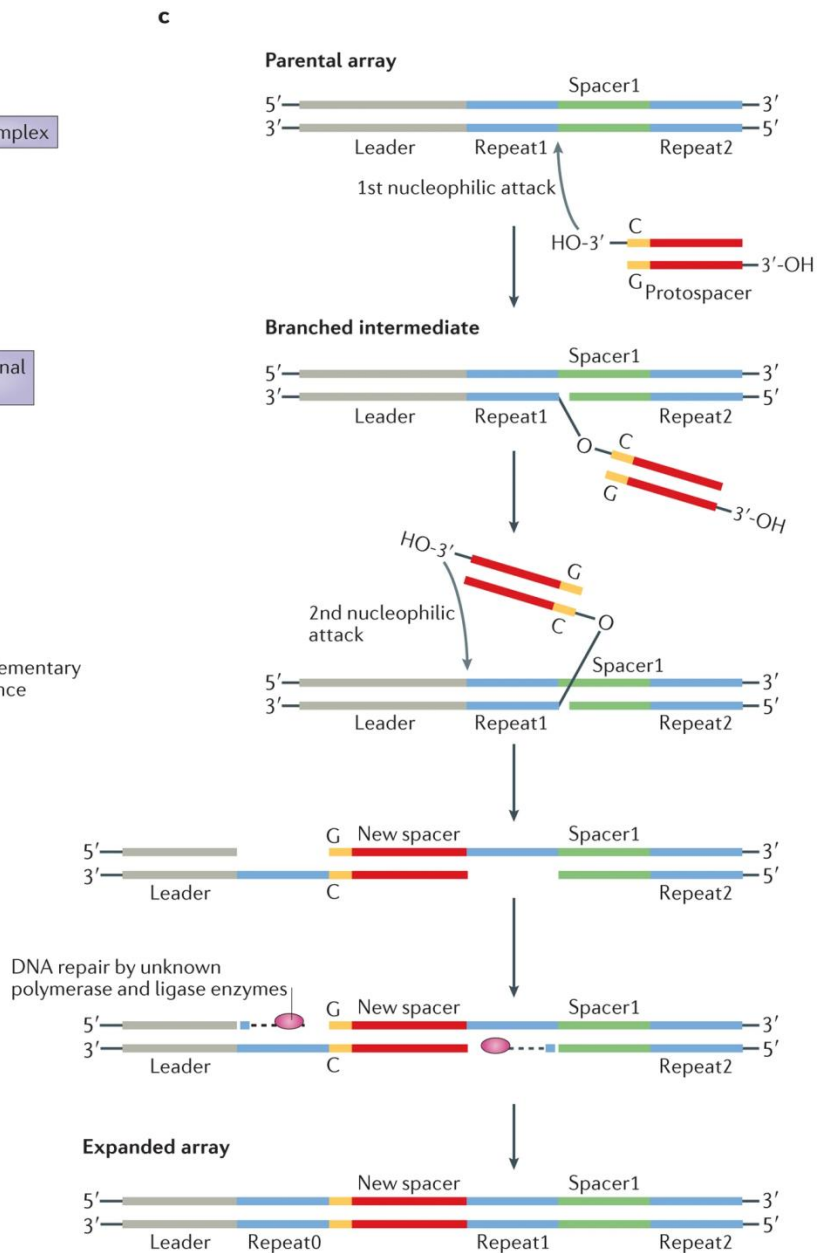
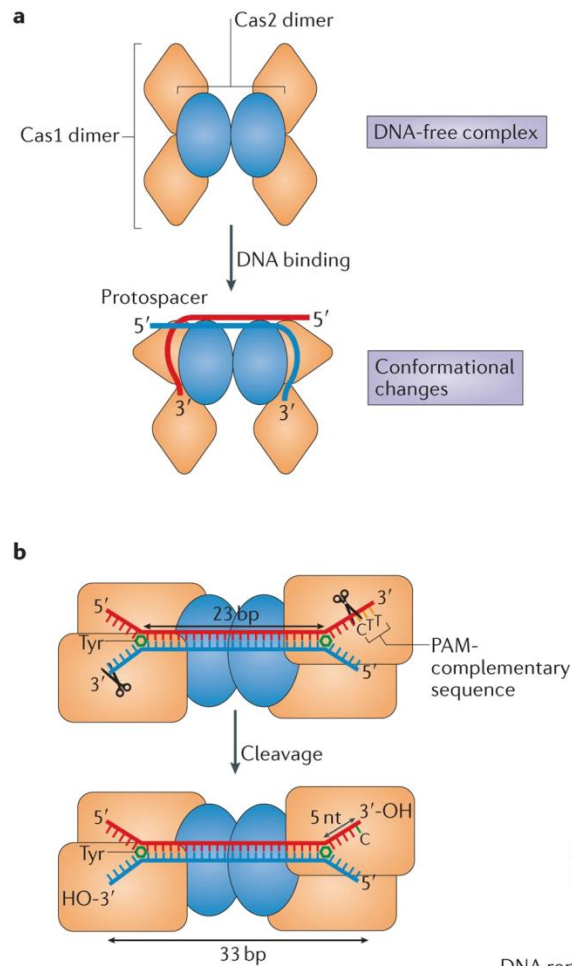


Foreign DNA

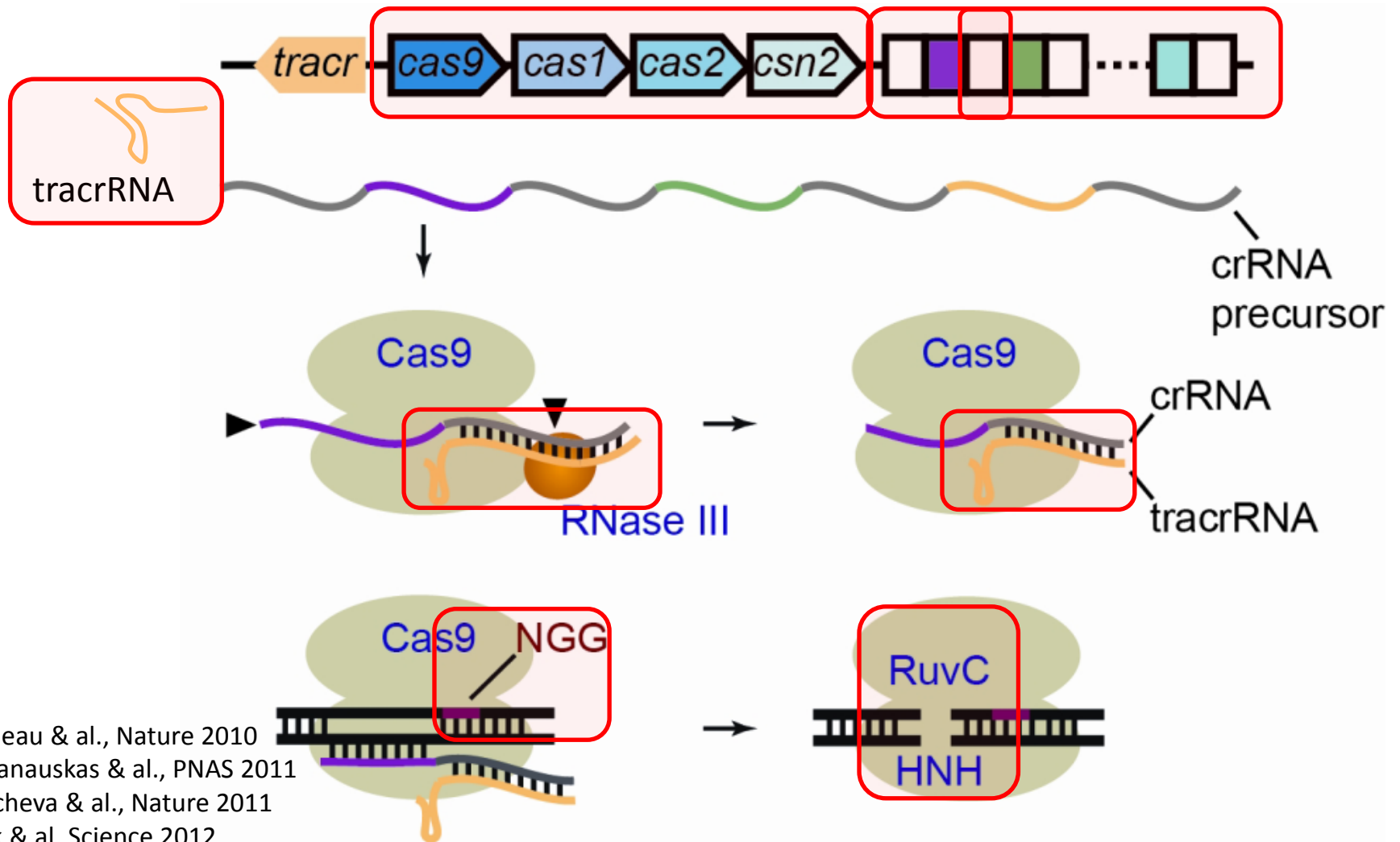






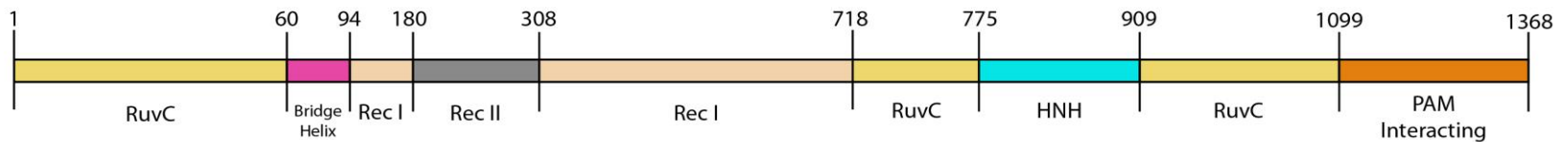
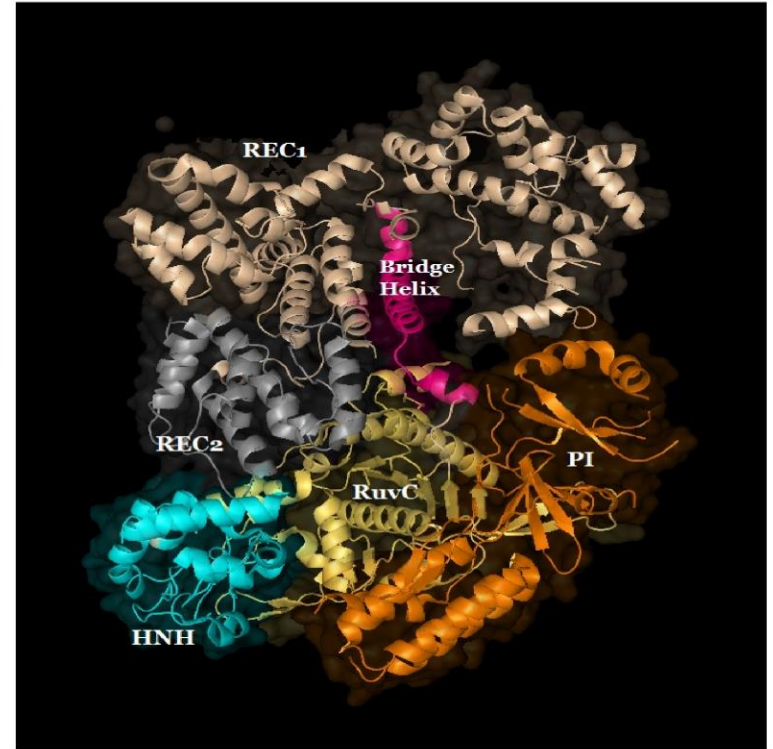
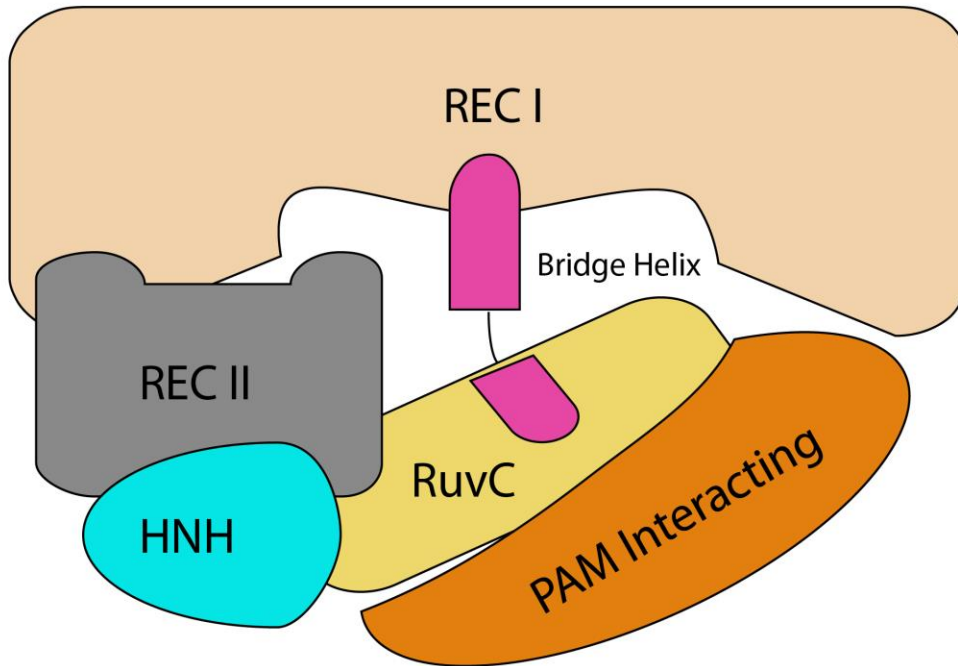


S.pyogenes type II-A system

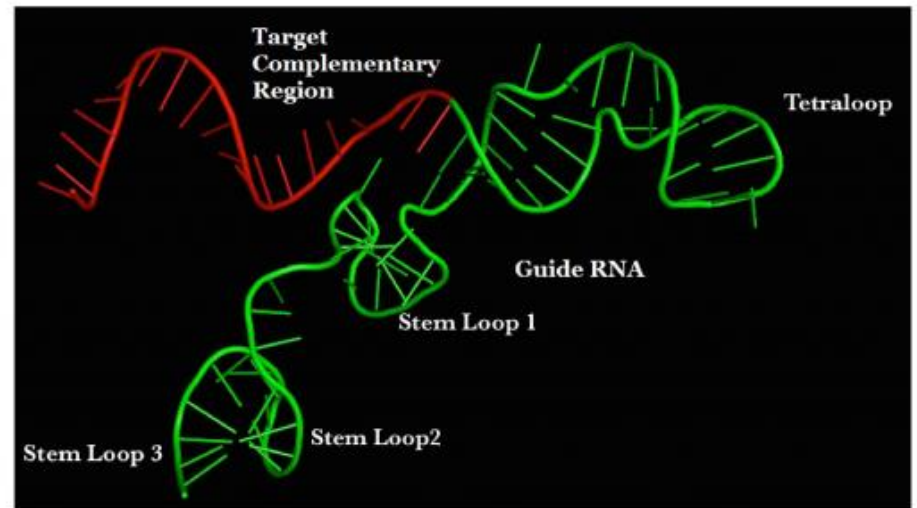
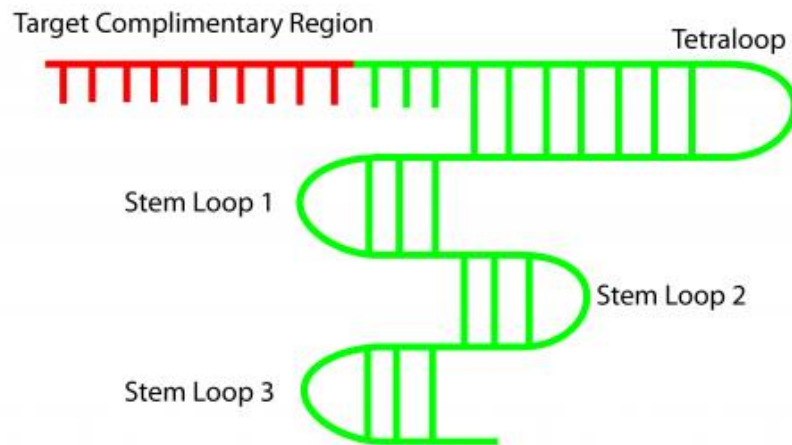


Garneau & al., Nature 2010
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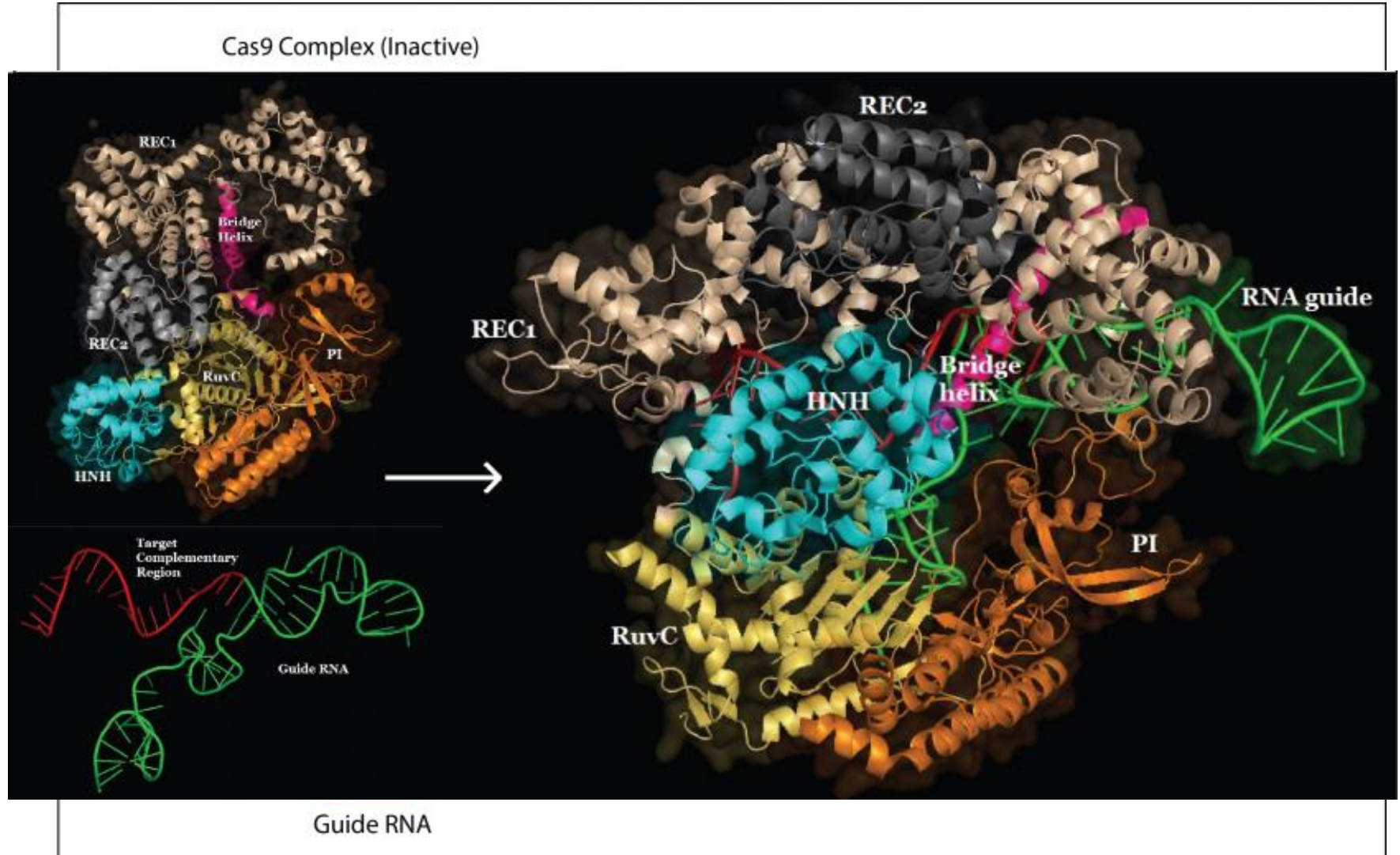
S.pyogenes type II-A system



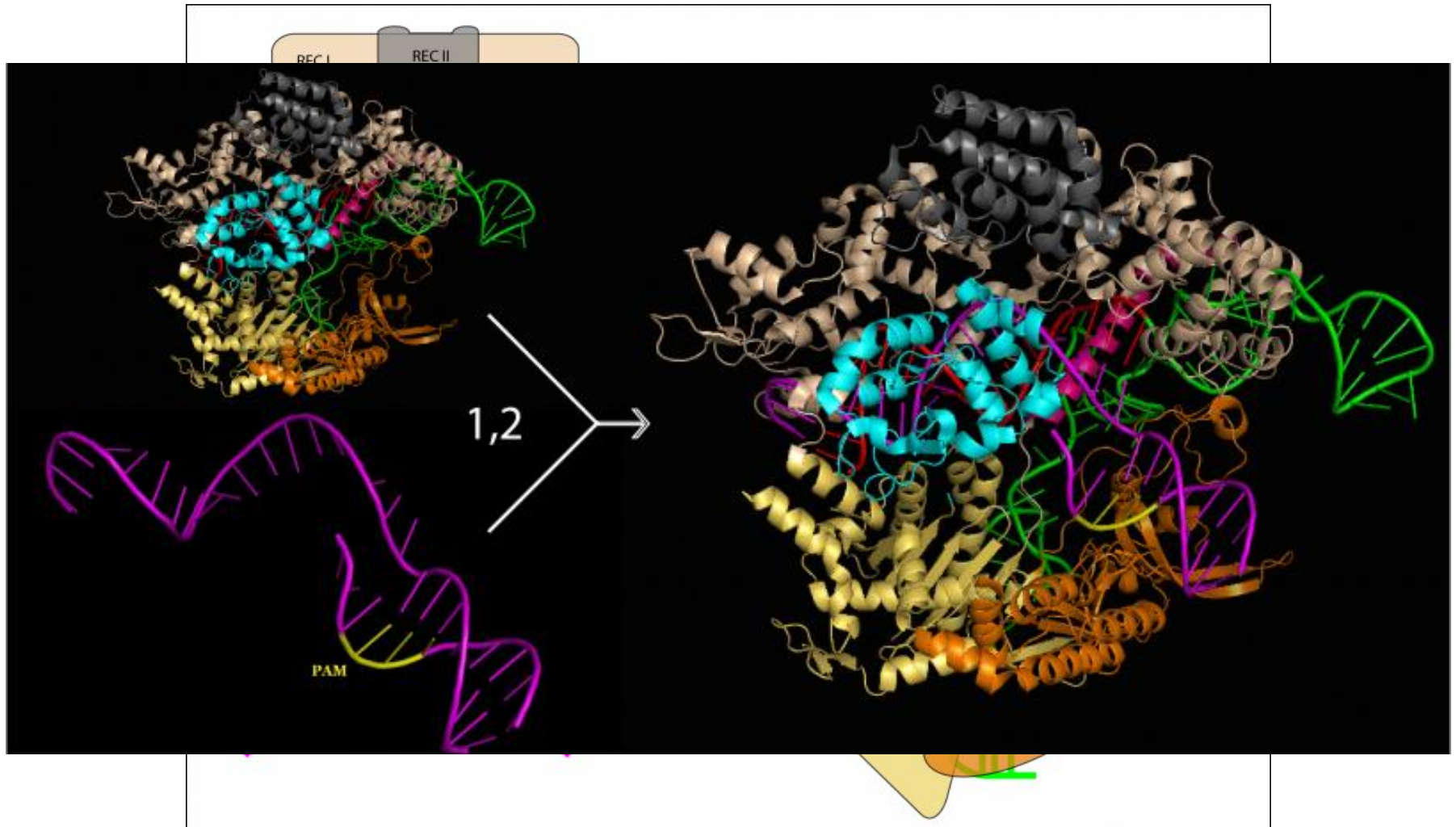
S.pyogenes type II-A system



S.pyogenes type II-A system



S.pyogenes type II-A system

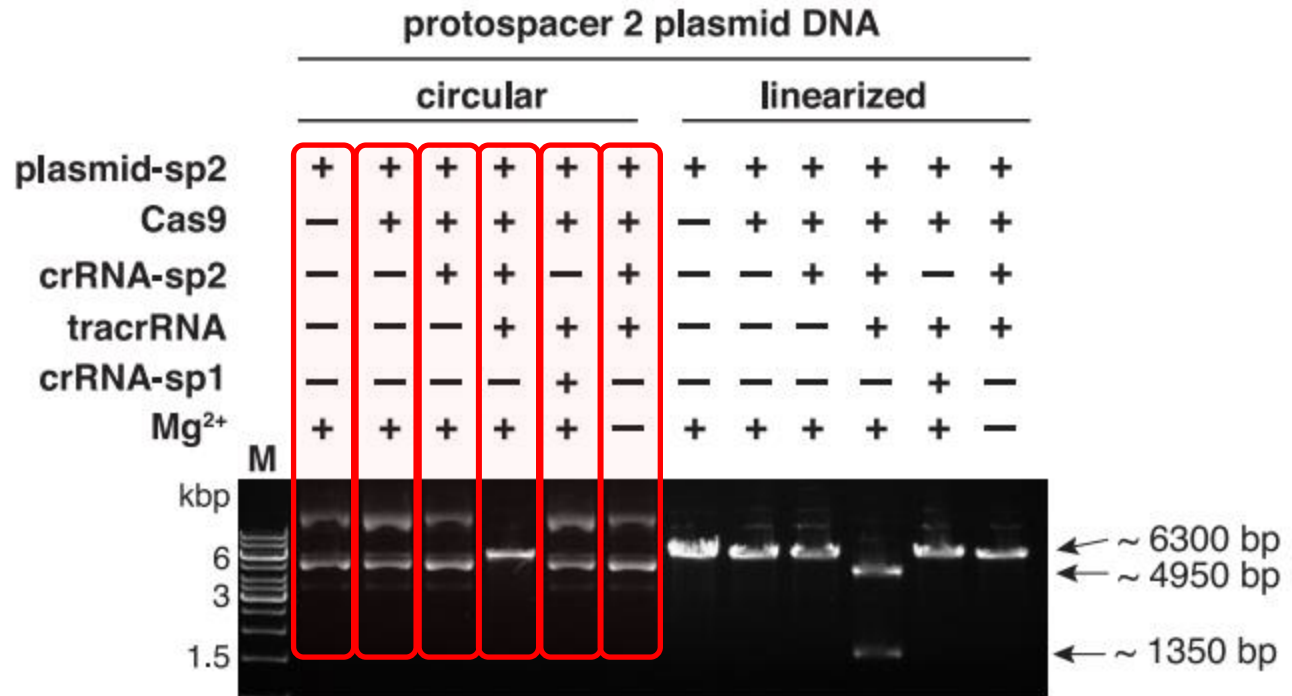
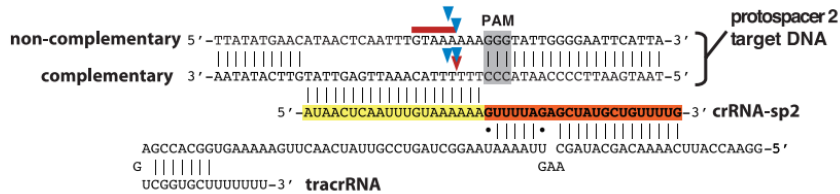


S.pyogenes type II-A system

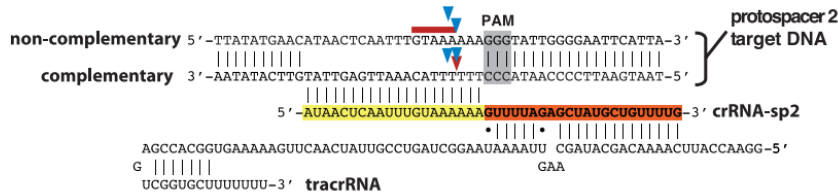
A Programmable Dual-RNA–Guided DNA Endonuclease in Adaptive Bacterial Immunity

Martin Jinek,^{1,2*} Krzysztof Chylinski,^{3,4*} Ines Fonfara,⁴ Michael Hauer,^{2†}
Jennifer A. Doudna,^{1,2,5,6‡} Emmanuelle Charpentier^{4‡}

Cas9 DNA endonuclease guided by two RNA

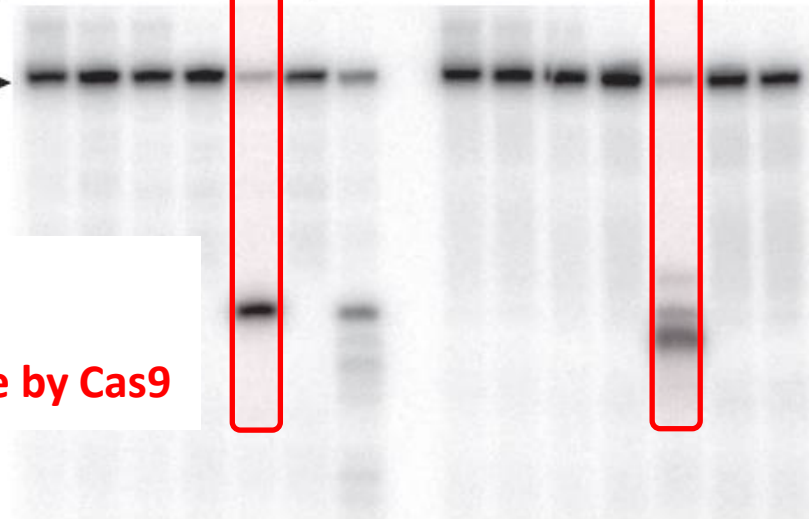


Cas9 DNA endonuclease guided by two RNA



	protospacer 2 oligonucleotide DNA											
	complementary						non-complementary					
labeled strand	-	+	+	+	+	-	-	+	+	+	+	-
partner strand	-	+	+	+	+	-	-	+	+	+	+	-
Cas9	-	-	+	+	+	+	-	-	+	+	+	+
crRNA-sp2	-	-	-	+	+	-	-	-	-	+	+	-
tracrRNA	-	-	-	-	+	+	-	-	-	-	+	+
crRNA-sp1	-	-	-	-	-	+	-	-	-	-	-	+

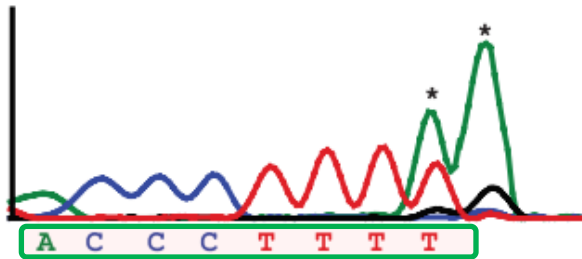
50 nt →



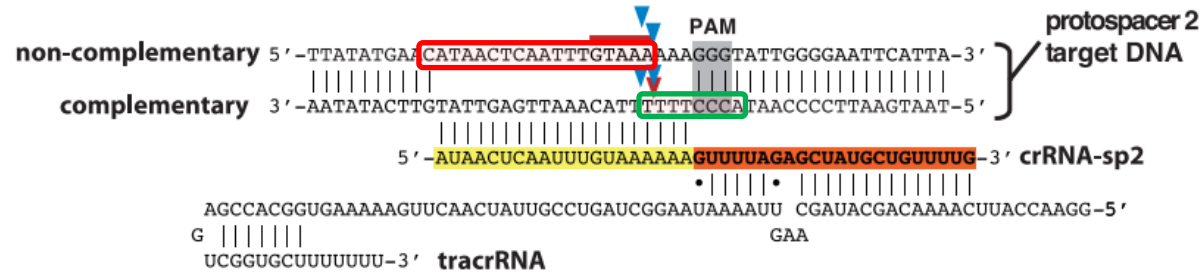
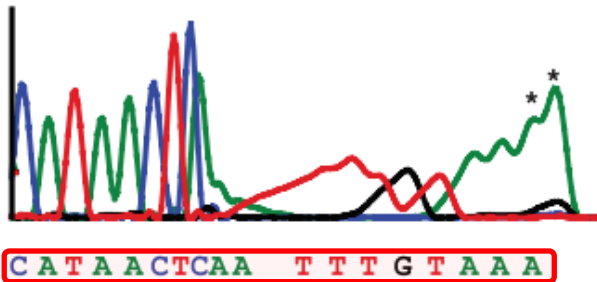
- pre-crRNA processing by RNase III
- activating crRNA-guided DNA cleavage by Cas9

Cas9 DNA endonuclease guided by two RNA

non-complementary strand binding primer

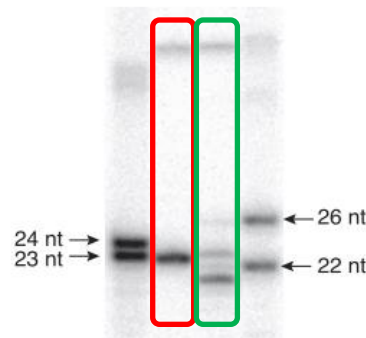


complementary strand binding primer



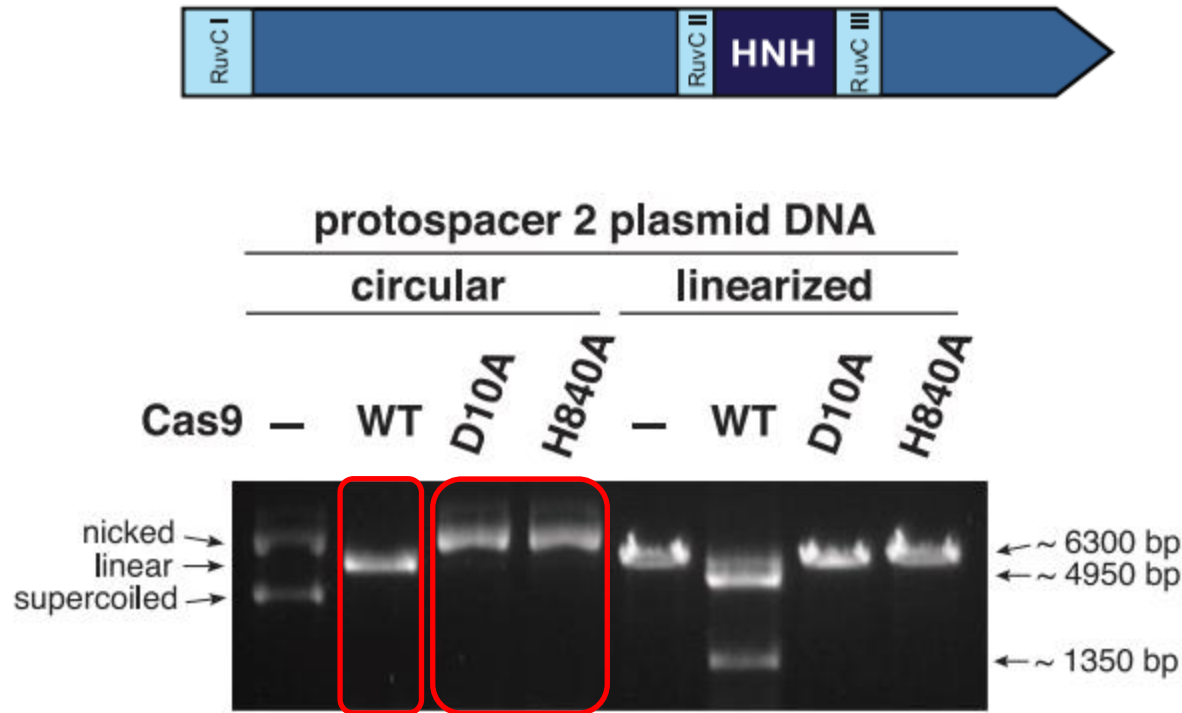
complementary strand
non-complementary strand

M P P M



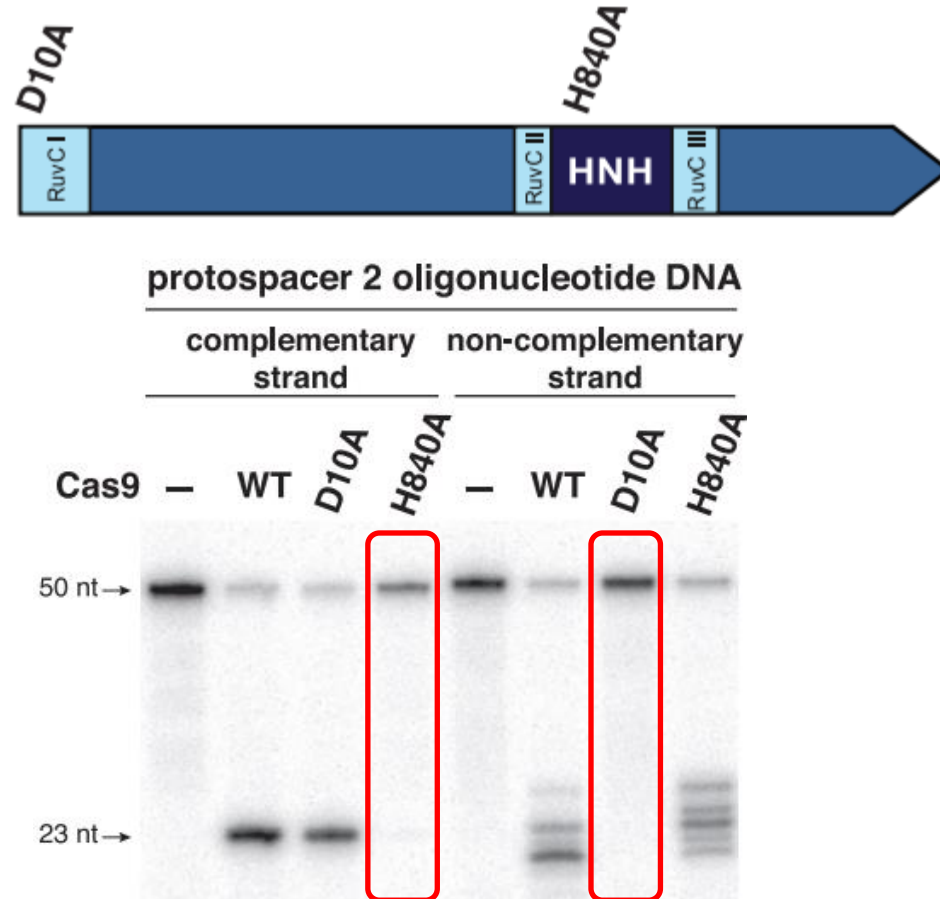
- Non complementary = 1st cleaved endonuclease
- 3'-5' exonuclease

Each Cas9 nuclease domain cleaves one DNA strand



- HNH = one plasmid strand
- RuvC = one plasmid strand

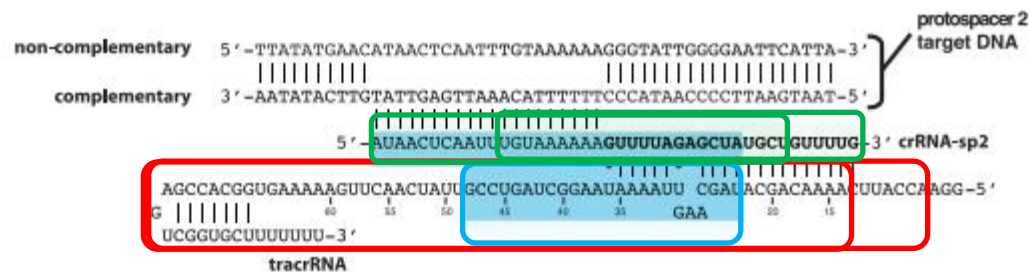
Each Cas9 nuclease domain cleaves one DNA strand



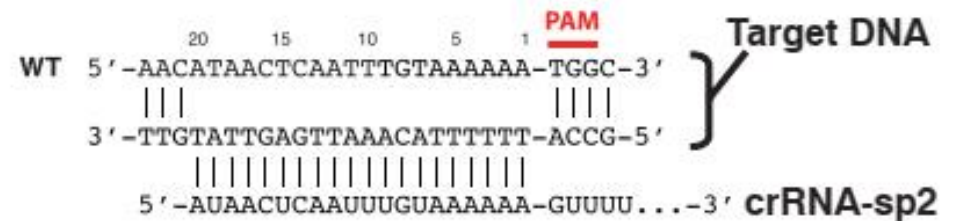
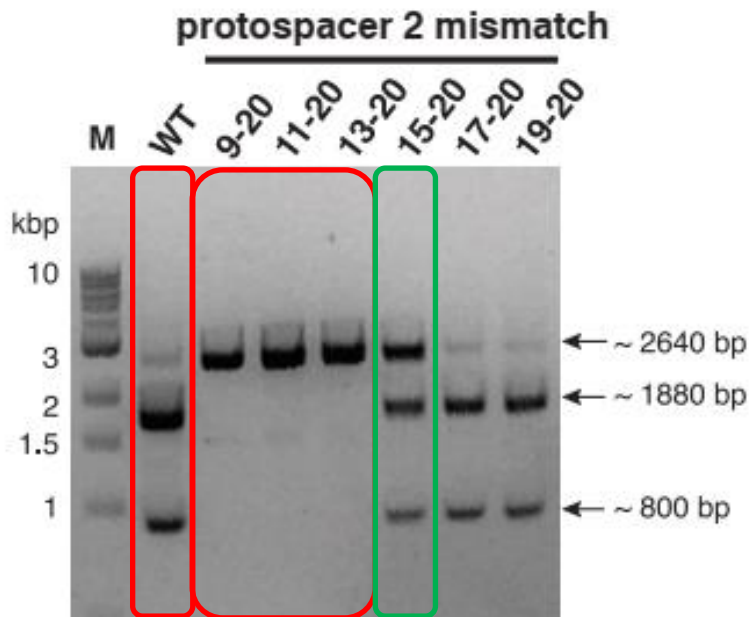
- HNH = complementary strand
- RuvC = non complementary strand

Dual RNA requirements

C



Protospacers sequence



mismatched targets

9-20 5' -**TATTGAGTTAA**GTAAAAAA-3'

11-20 5' -**TATTGAGTTA**TTGTAAAAAA-3'

13-20 5' -**TATTGAGT**ATTTGTAAAAAA-3'

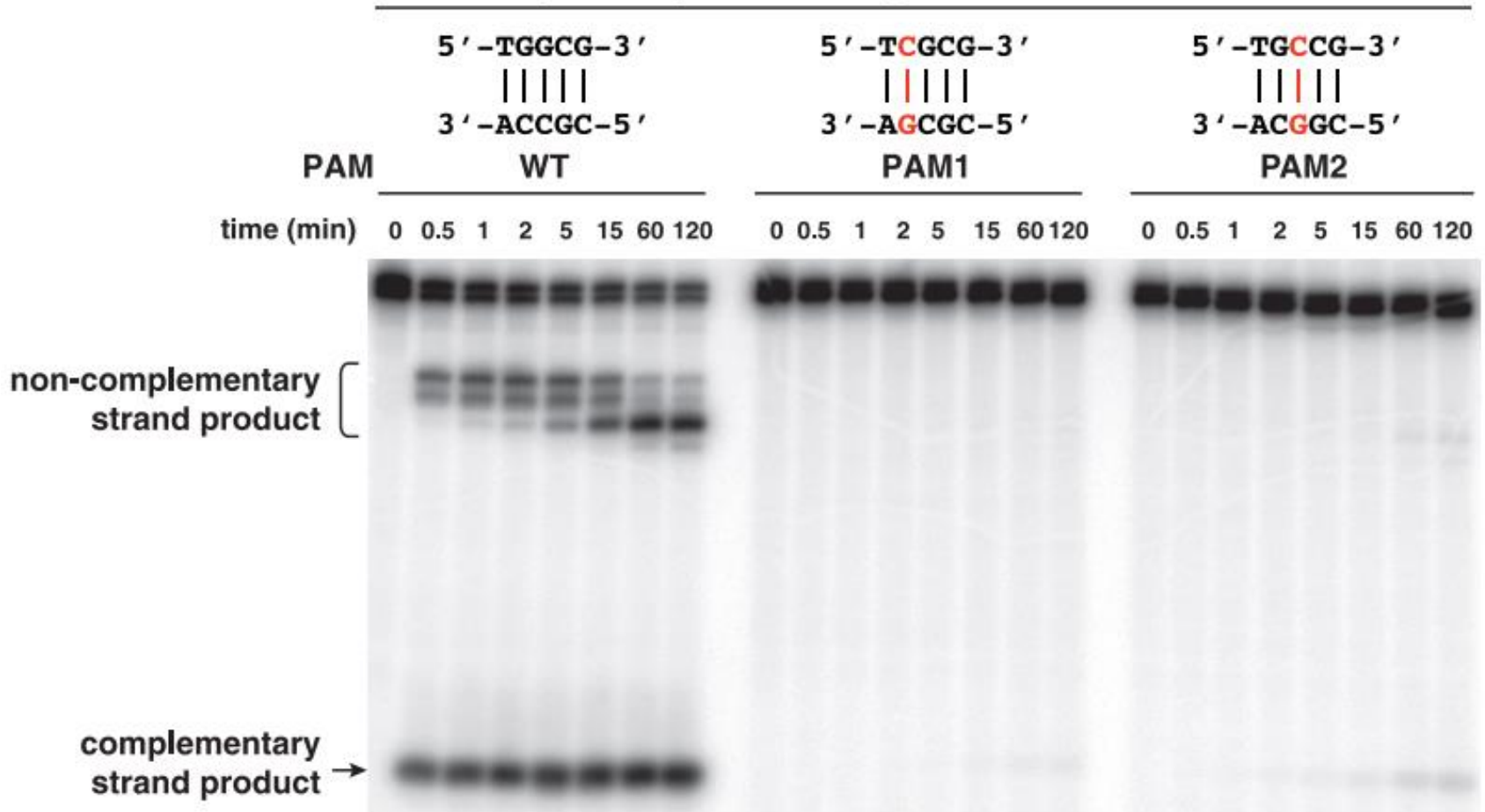
15-20 5' -**TATTGA**CAATTTGTAAAAAA-3'

17-20 5' -**TATT**CTCAATTTGTAAAAAA-3'

19-20 5' -**TA**AACTCAATTTGTAAAAAA-3'

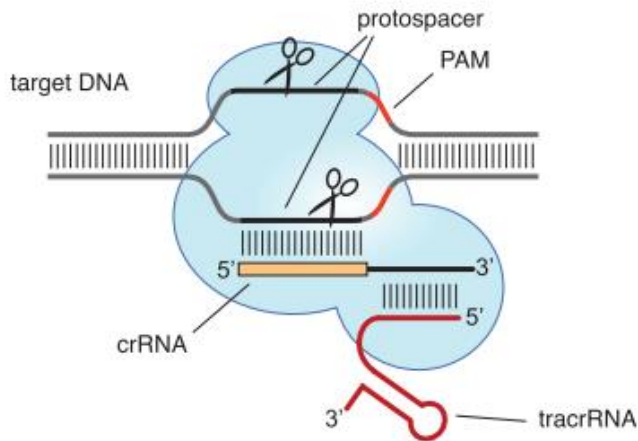
PAM sequence

protospacer 4 oligonucleotide DNA

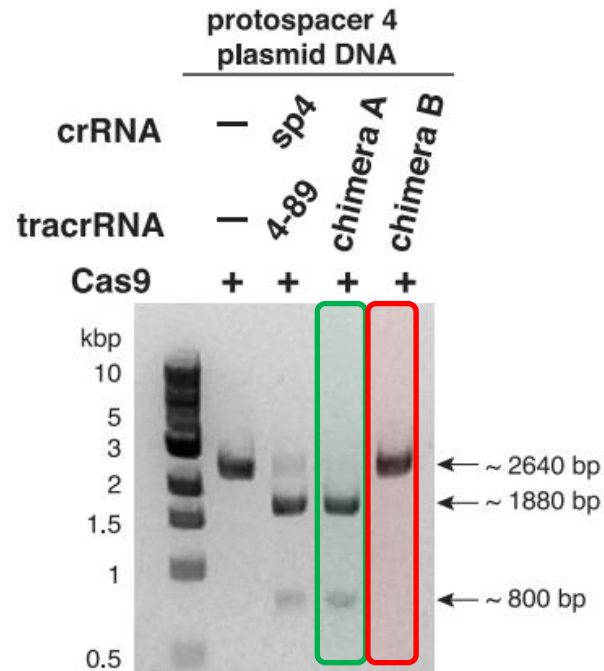
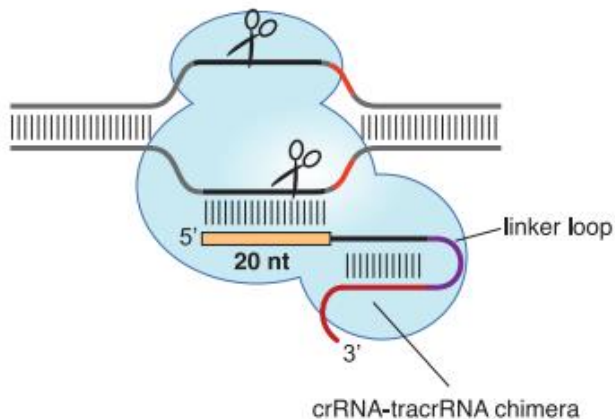


Single chimeric RNA

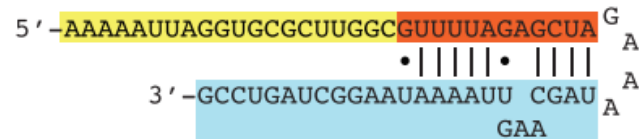
Cas9 programmed by crRNA:tracrRNA duplex



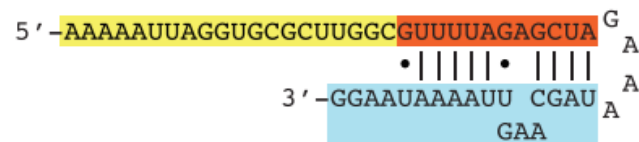
Cas9 programmed by single chimeric RNA



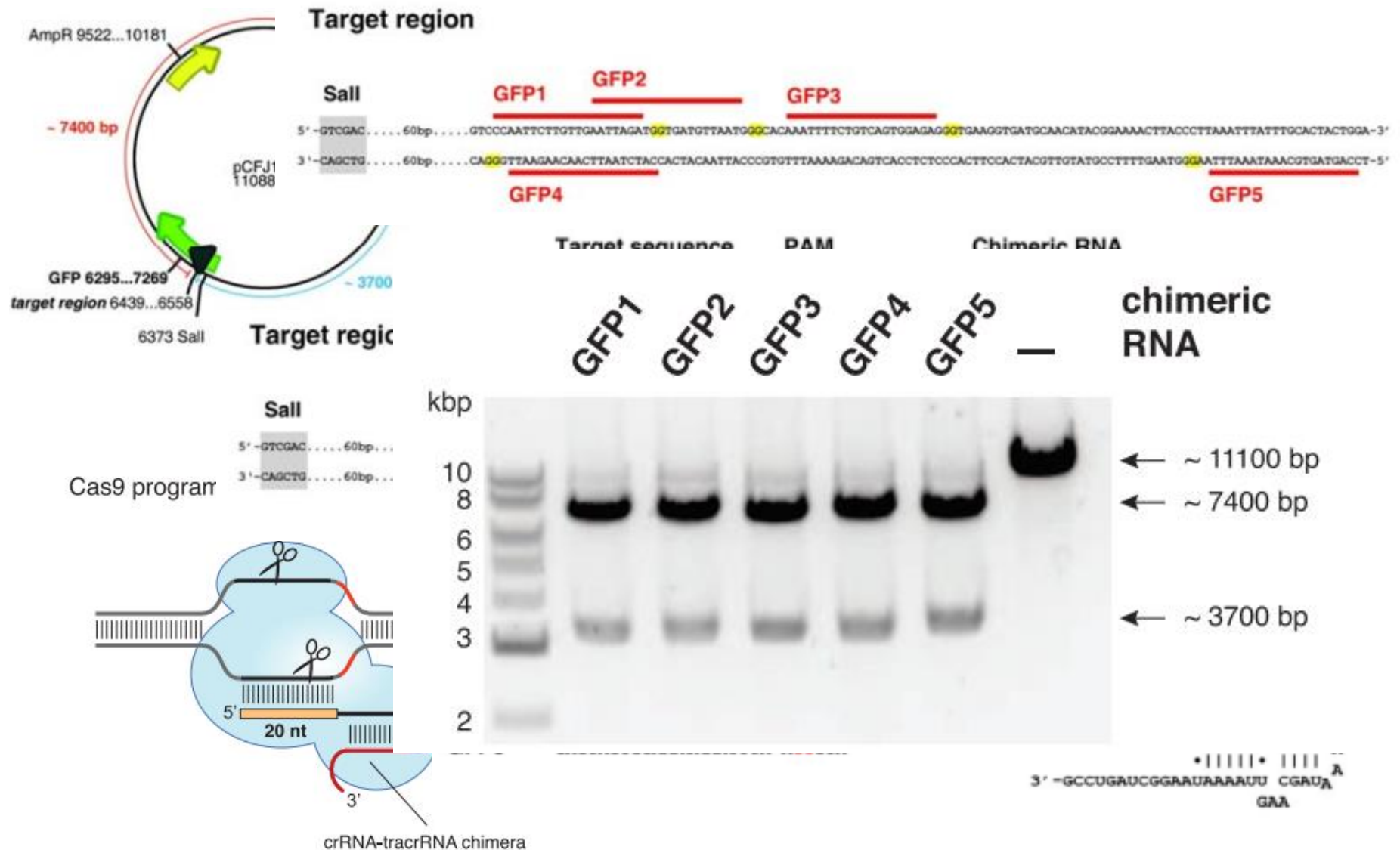
chimera A



chimera B



Engineered site specific guide RNA



Conclusions of Lecture-17

- Type IIA CRISPR-Cas System
- Molecular requirements (Cas9, tracrRNA, crRNA, PAM in target DNA)
- Engineered Single guide RNA
- System = simple, efficient, versatile and programmable
- RNA-programmed Cas9 = gene-targeting and genome-editing

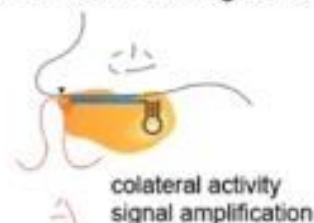
Abudayyeh et al., *Science* (2016)
 East-Seletsky et al., *Nature* (2016)
 Cox et al., *Science* (2017)
 Abudayyeh et al., *Nature* (2017)
 Smargon et al., *Mol. Cell* (2017)
 Konermann et al., *Cell* (2018)
 Yan et al., *Mol. Cell* (2018)

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RNA knockdown



Virus detection/degradation



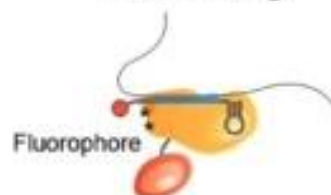
Gootenberg et al., *Science* (2017)
 Gootenberg et al., *Science* (2018)
 Joung et al., (2020)

A-to-I editing



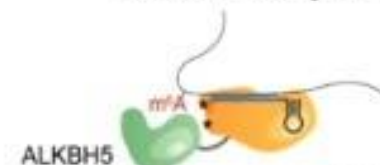
Cox et al., *Science* (2017)

RNA tracing



Wang et al., *Science* (2019)
 Yang et al., *Mol. Cell* (2019)

RNA demethylation



Li et al., *NAR* (2020)

Alternative splicing



Konermann et al., *Cell* (2018)
 Jillette et al., *bioRxiv* (2019)

Alternative polyadenylation



Anderson et al., *bioRxiv* (2019)

Proximity labeling



Zhang et al., *NAR* (2020)
 Han et al., *bioRxiv* (2020)
 Lin et al., *bioRxiv* (2020)
 Li et al., *bioRxiv* (2020)

Questions??