

# Genome Editing and Engineering

Course No: BT-637



## LECTURE-6

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# Introduction

- FokI (Flavobacterium okeanokoites)
- Biochemical and crystal structure
- Single catalytic domain
- Rearrange in higher order structure
- Cleave both strands of DNA

*Proc. Natl. Acad. Sci. USA*  
Vol. 95, pp. 10564–10569, September 1998  
Biophysics

## **Structure of *FokI* has implications for DNA cleavage**

DAVID A. WAH\*, JURATE BITINAITE†, IRA SCHILDKRAUT†, AND ANEEL K. AGGARWAL\*‡

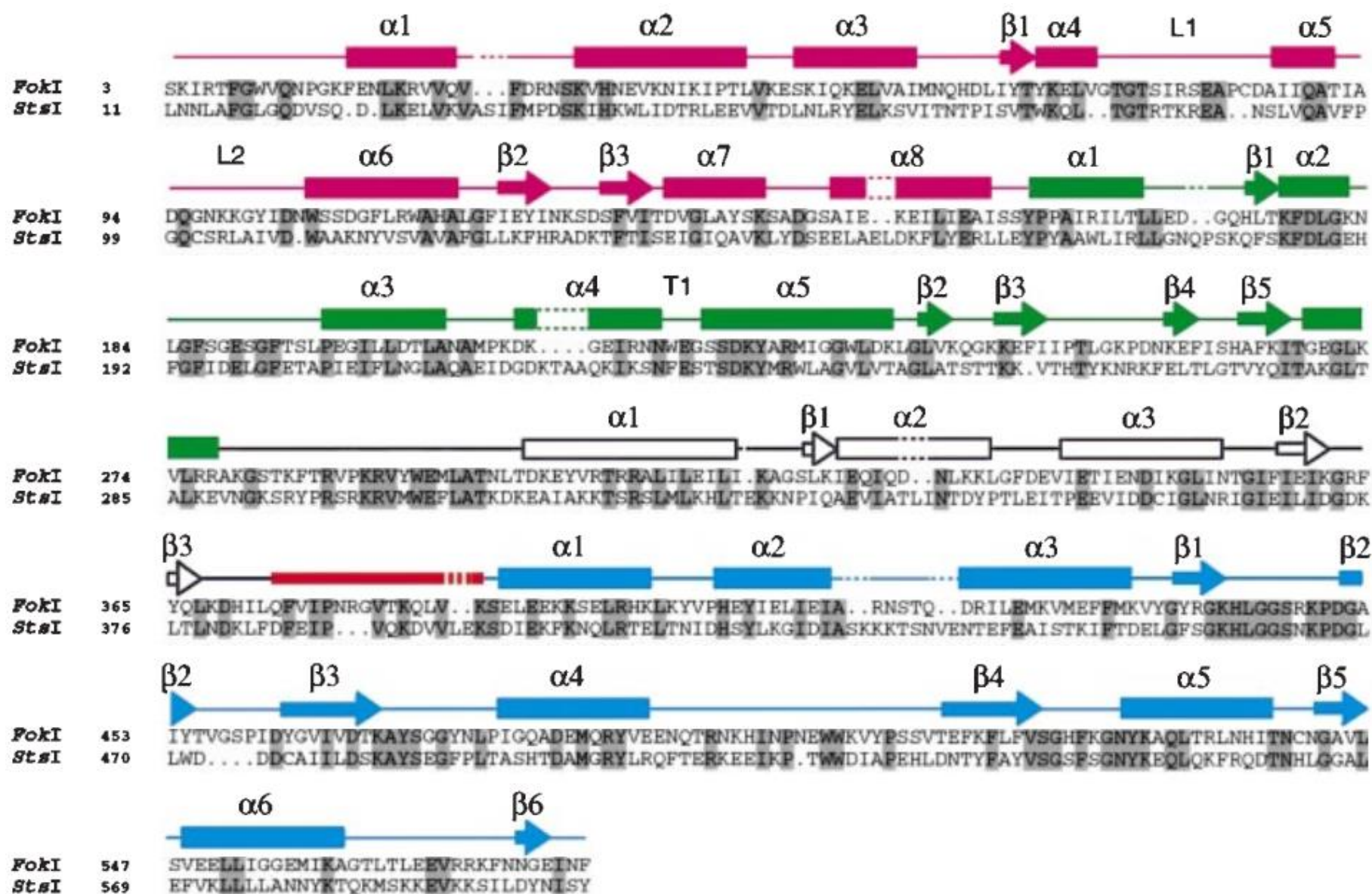
\*Structural Biology Program, Department of Physiology and Biophysics, Box 1677, 1425 Madison Avenue, Mount Sinai School of Medicine, New York, NY 10029; and †New England Biolabs, 32 Tozer Road, Beverly, MA 01915

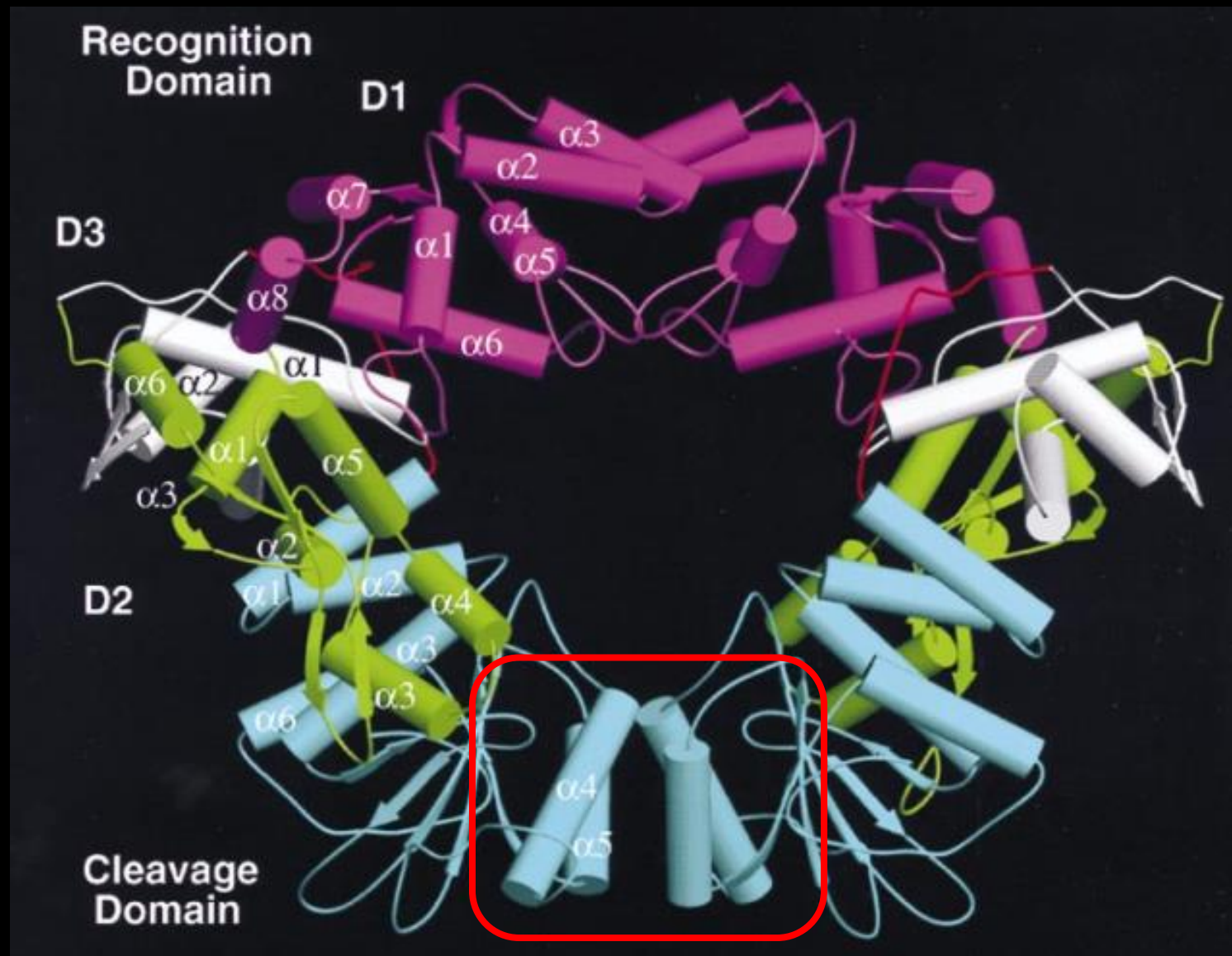
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## ***FokI* dimerization is required for DNA cleavage**

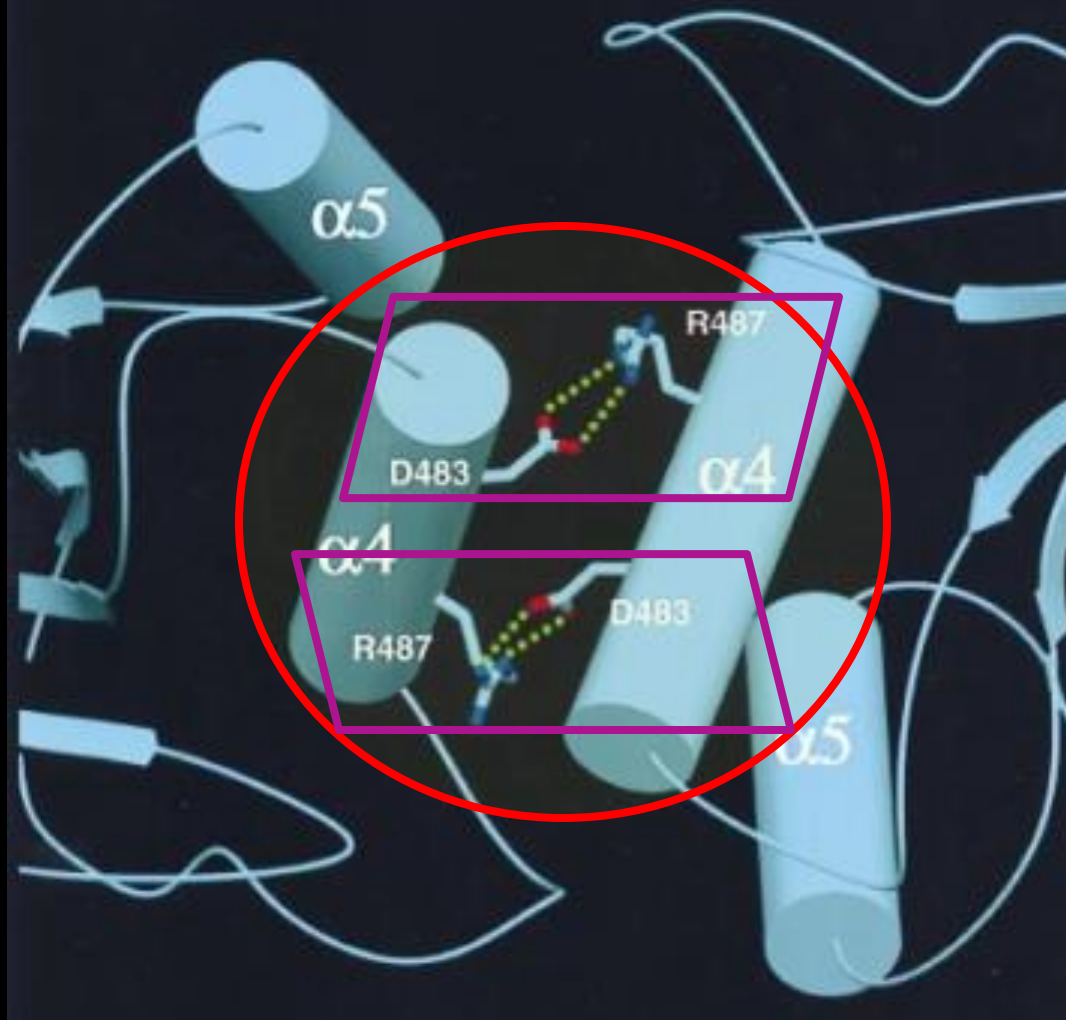
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## Dimer Interface





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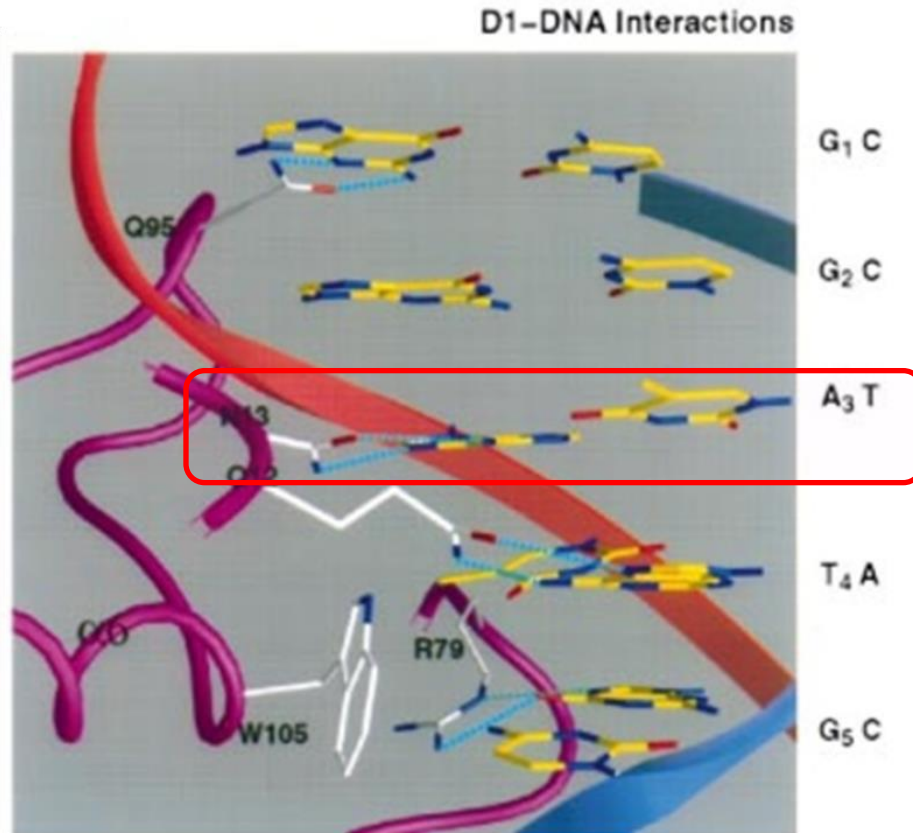
## *FokI* dimerization is required for DNA cleavage

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# Construction of FokI mutants

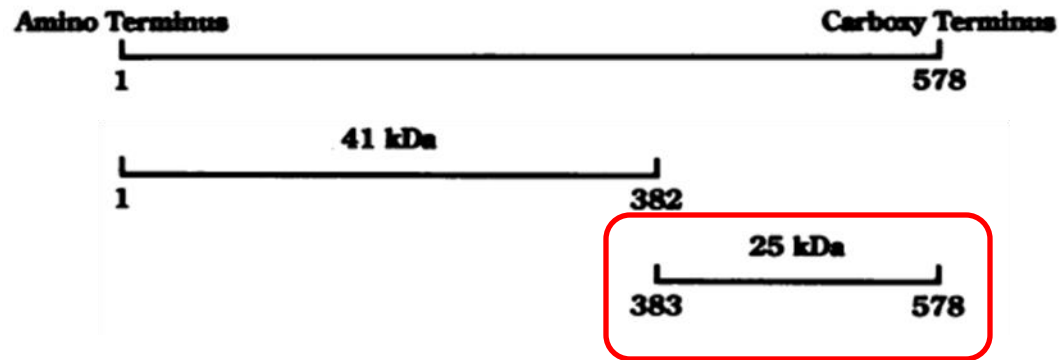
- Fok N13Y





# Construction of FokI mutants

- Fok N13Y
- Fok CD
- 196 a.a.
- 25 kDa



# Construction of FokI mutants

- Fok N13Y

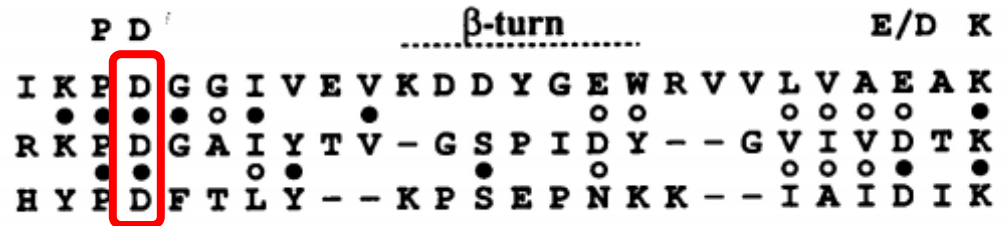
- Fok CD

Consensus

*Eco* RI (88-113)

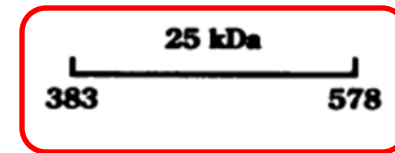
*Fok* I (447-469)

*Eco* RV (71-92)

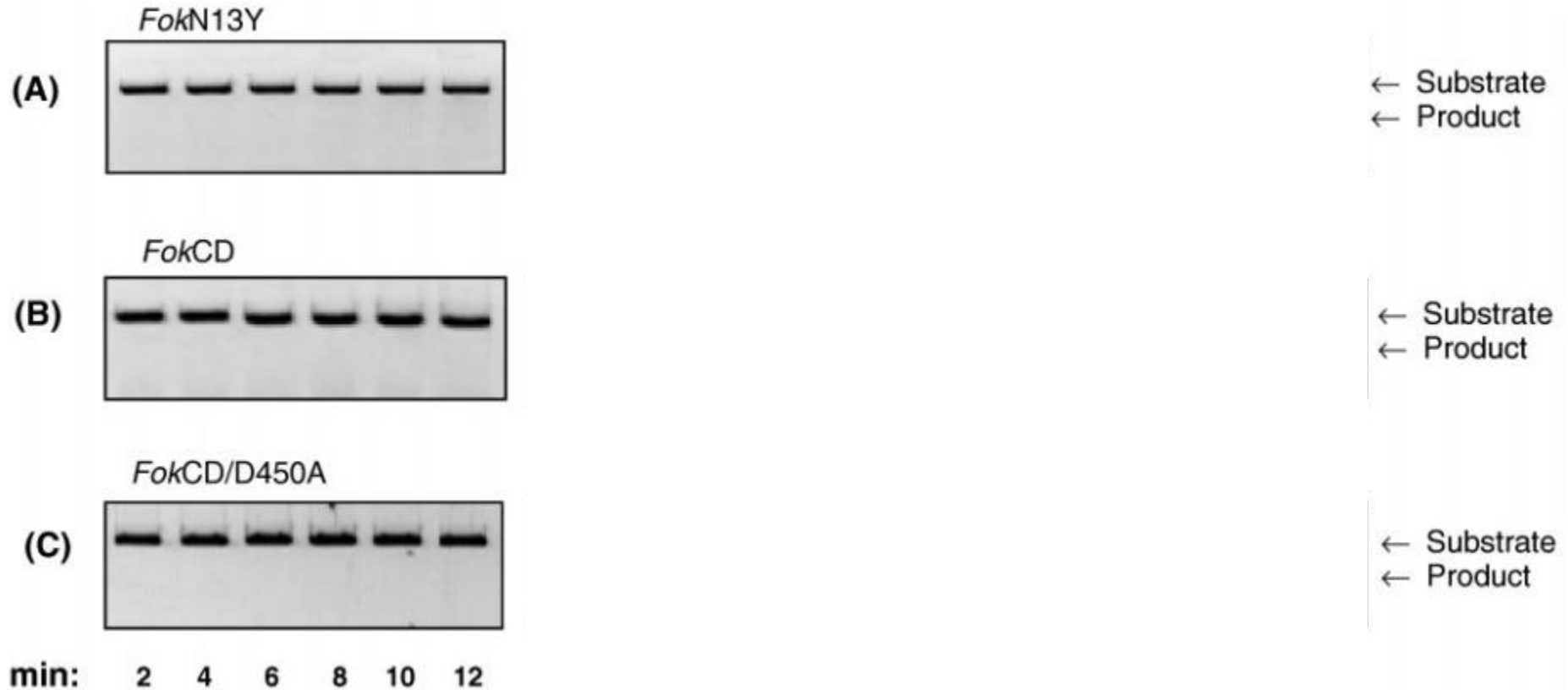


- Fok CD/D450A

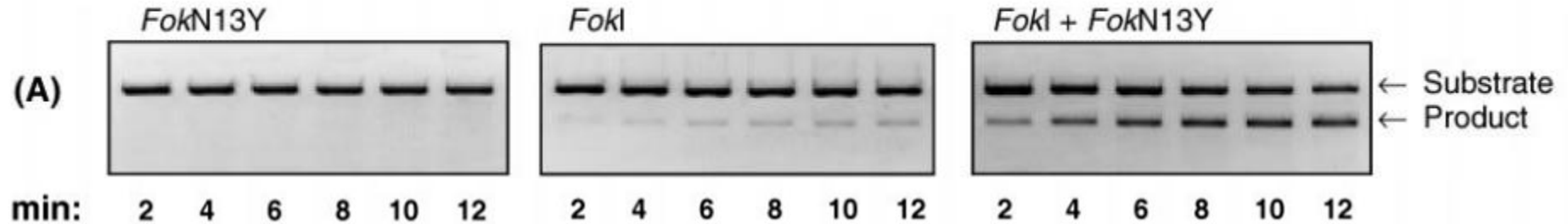
D450A;



# Construction of FokI mutants

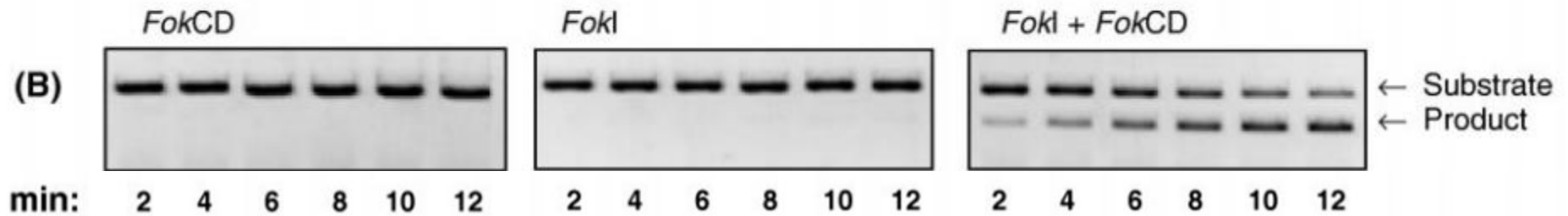


# Construction of FokI mutants



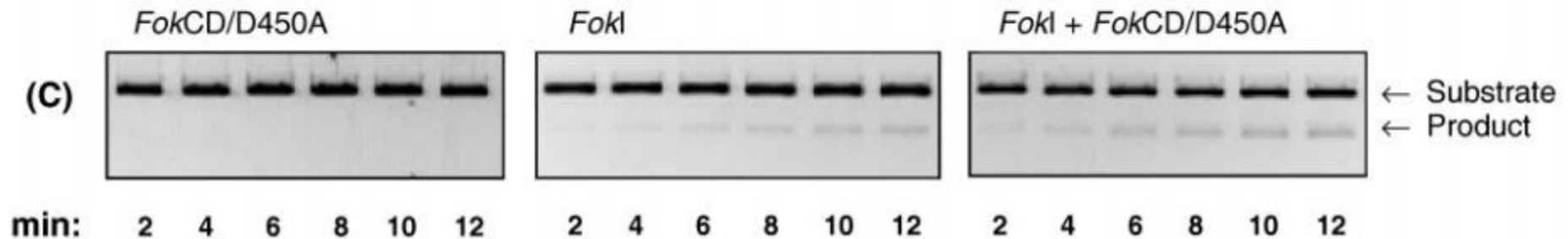
Wt. FokI interacts with Fok N13Y

# Construction of FokI mutants



Wt. FokI interacts with Fok CD

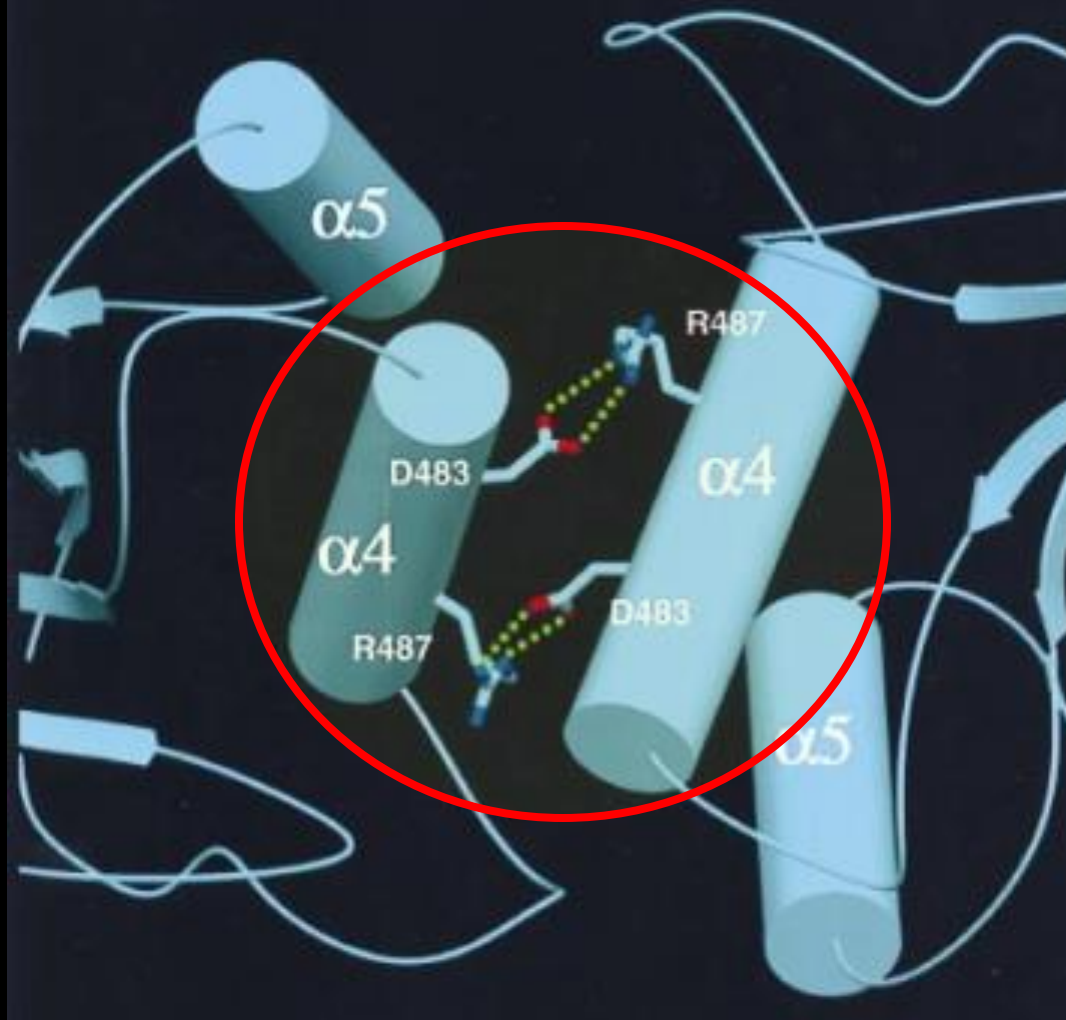
# Construction of FokI mutants



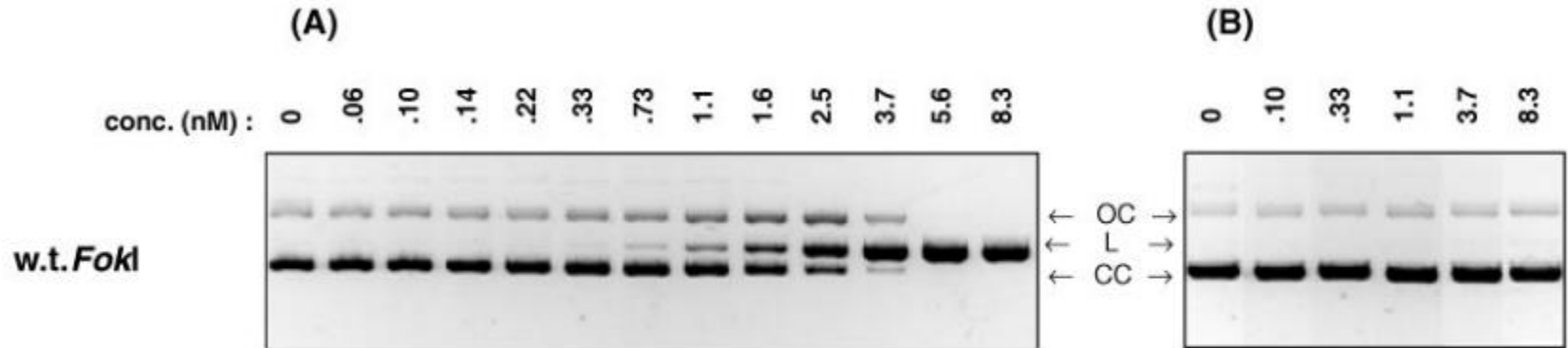
Wt. FokI does not interact with Fok CD/D450A



## Dimer Interface



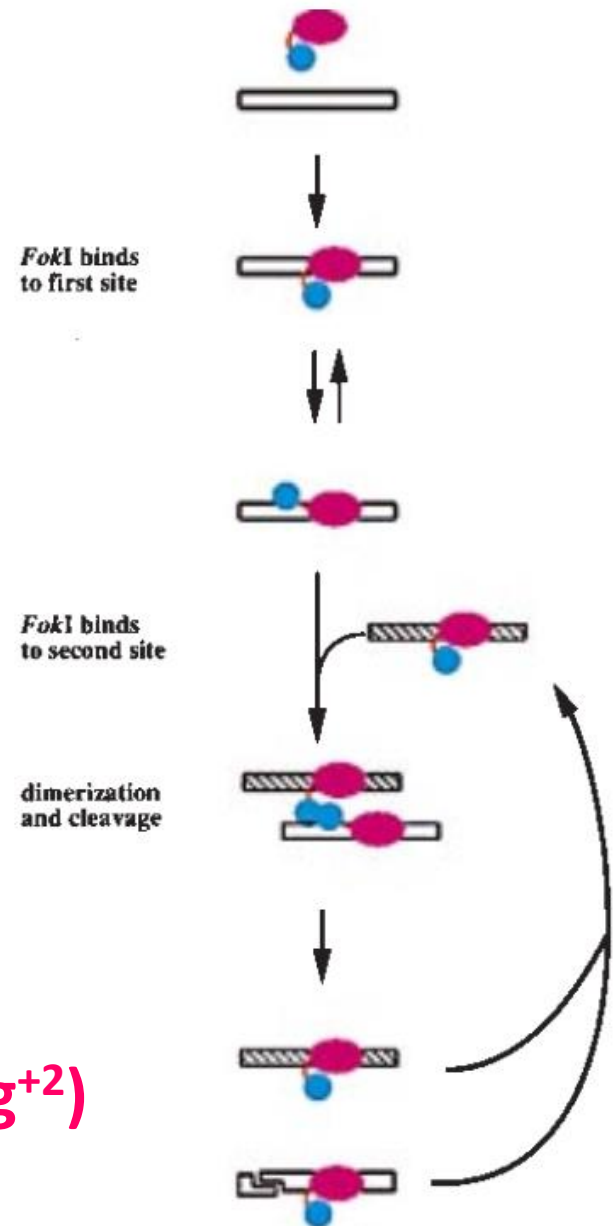
# Interface D483A; R487A mutants



Fok D483A, R487A = Reduced cleavage activity

# Dimer model

- FokI first binds to DNA as monomer
- The complex is inactive
- A second FokI monomer arrives:
  - i) when DNA is scanned until
  - ii) the monomer collides
- Correct orientation = dimerize
- Cleavage of ds DNA= FokI dimers ( $Mg^{+2}$ )



# Conclusions of Lecture-6

- Cleavage of both strand= after FOKI dimerization (Mg)
- Dimerization model offers (two control)
  - i) The release of CD depends on seq.-spe. binding
  - ii) Dimerization of CD = phosphodiester bond cleavage

**Thank You!**