

Supplemental Material

The Kinetic Mechanism for DNA Unwinding by Multiple Molecules of Dda Helicase Aligned on DNA[†]

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Running title: DNA unwinding by multiple helicases aligned on DNA

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Abbreviations: RQF – Rapid Quench Flow; NLLS – non-linear least squares.

Mechanisms for data fitting using the program Kinetic Global Kinetic Explorer.

Mechanisms were written using the Model Editor of the program according to the following definitions:

- D1: DNA substrate
- D2: DNA substrate after one kinetic step for unwinding
- D3: DNA substrate after two kinetic steps for unwinding
- D4: DNA substrate after three kinetic steps for unwinding
- P: DNA substrate after sufficient number of kinetic steps for ssDNA to form
- E: One molecule of Dda
- EE: Two molecules of Dda
- EEE: Three molecules of Dda
- EEEE: Four molecules of Dda
- A: The trailing molecule of Dda after dissociation of the leading molecule (see Fig. 2 for explanation of the trailing molecule)
- B: The trailing molecule of Dda after one kinetic step for translocation

DNA unwinding of the 14T16bp substrate in 3 kinetic steps with three Dda molecules.

ED1 = ED2 = ED3 = EP
ED1 = E + D1
ED2 = E + D2
ED3 = E + D3
EED1 = EED2 = EED3 = EEP
EED1 = ED1
EED2 = ED2
EED3 = ED3
EED1 = E + AD1
EED2 = E + AD2
EED3 = E + AD3
AD1 = BD1
AD2 = BD2
AD3 = BD3
BD1 = ED1
BD2 = ED2
BD3 = ED3
AD1 = E + D1
AD2 = E + D2
AD3 = E + D3
BD1 = E + D1
BD2 = E + D2
BD3 = E + D3

Chi² = 0.01377
Chi²/DoF = 0.001059
Sigma = 0.0325

DNA unwinding of the 14T20bp substrate in 4 kinetic steps with two Dda molecules.

ED1 = ED2 = ED3 = ED4 = EP

ED1 = E + D1

ED2 = E + D2

ED3 = E + D3

ED4 = E + D4

EED1 = EED2 = EED3 = EED4 = EEP

EED1 = ED1

EED2 = ED2

EED3 = ED3

EED4 = ED4

EED1 = E + AD1

EED2 = E + AD2

EED3 = E + AD3

EED4 = E + AD4

AD1 = BD1

AD2 = BD2

AD3 = BD3

AD4 = BD4

BD1 = ED1

BD2 = ED2

BD3 = ED3

BD4 = ED4

AD1 = E + D1

AD2 = E + D2

AD3 = E + D3

AD4 = E + D4

BD1 = E + D1

BD2 = E + D2

BD3 = E + D3

BD4 = E + D4

$\text{Chi}^2 = 0.00350236$

$\text{Chi}^2/\text{DoF} = 0.000269412$

Sigma = 0.0164

DNA unwinding of the 21T16bp substrate in 3 kinetic steps with three Dda molecules.

ED1 = ED2 = ED3 = EP
ED1 = E + D1
ED2 = E + D2
ED3 = E + D3
EED1 = EED2 = EED3 = EEP
EED1 = ED1
EED2 = ED2
EED3 = ED3
EEED1 = EEED2 = EED3 = EEEP
EEED1 = E + EED1
EEED2 = E + EED2
EEED3 = E + EED3
EEED1 = EAD1
EEED2 = EAD2
EEDE3 = EAD3
EED1 = E + AD1
EED2 = E + AD2
EED3 = E + AD3
EAD1 = EED1
EAD2 = EED2
EAD3 = EED3
AD1 = BD1
AD2 = BD2
AD3 = BD3
BD1 = ED1
BD2 = ED2
BD3 = ED3
AD1 = E + D1
AD2 = E + D2
AD3 = E + D3
BD1 = E + D1
BD2 = E + D2
BD3 = E + D3

Chi² = 0.0221537
Ch²/DoF = 0.00170413
Sigma = 0.0412811

DNA unwinding of the 21T20bp substrate in 3 kinetic steps with three Dda molecules.

ED1 = ED2 = ED3 = ED4 = EP
ED1 = E + D1
ED2 = E + D2
ED3 = E + D3
ED4 = E + D4
EED1 = EED2 = EED3 = EED4 = EEP
EED1 = E + EED1
EED2 = E + EED2
EED3 = E + EED3
EED4 = E + EED4
EEED1 = EEED2 = EEED3 = EEED4 = EEEP
EEED1 = E + EEED1
EEED2 = E + EEED2
EEED3 = E + EEED3
EEED4 = E + EEED4
EEED1 = EAD1
EEED2 = EAD2
EEED3 = EAD3
EEED4 = EAD4
EED1 = E + AD1
EED2 = E + AD2
EED3 = E + AD3
EED4 = E + AD4
EAD1 = EBD1 = EED1
EAD2 = EBD2 = EED2
EAD3 = EBD3 = EED3
EAD4 = EBD4 = EED4
AD1 = BD1 = ED1
AD2 = BD2 = ED2
AD3 = BD3 = ED3
AD4 = BD4 = ED4
AD1 = E + D1
AD2 = E + D2
AD3 = E + D3
AD4 = E + D4
BD1 = E + D1
BD2 = E + D2
BD3 = E + D3
BD4 = E + D4

$\text{Chi}^2 = 0.0121836$
 $\text{Ch}^2/\text{DoF} = 0.000937204$
 $\text{Sigma} = 0.0306138$

DNA unwinding of the 28T16bp substrate in 3 kinetic steps with four Dda molecules.

ED1 = ED2 = ED3 = EP
ED1 = E + D1
ED2 = E + D2
ED3 = E + D3
EED1 = EED2 = EED3 = EEP
EED1 = ED1
EED2 = ED2
EED3 = ED3
EEEED1 = EEEE2 = EEEE3 = EEEEP
EEEED1 = E + EED1
EEEED2 = E + EED2
EEEED3 = E + EED3
EEEED1 = EAD1
EEEED2 = EAD2
EEEED3 = EAD3
EEEED1 = EAED1
EEEED2 = EAED2
EEEED3 = EAED3
EEAD1 = EEBD1 = EED1
EEAD2 = EEBD2 = EED2
EEAD3 = EEBD3 = EED3
EED1 = E + AD1
EED2 = E + AD2
EED3 = E + AD3
EAD1 = EBD1 = EED1
EAD2 = EBD2 = EED2
EAD3 = EBD3 = EED3
AD1 = BD1
AD2 = BD2
AD3 = BD3
BD1 = ED1
BD2 = ED2
BD3 = ED3
AD1 = E + D1
AD2 = E + D2
AD3 = E + D3
BD1 = E + D1
BD2 = E + D2
BD3 = E + D3

$\text{Chi}^2 = 0.0263758$
 $\text{Ch}^2/\text{DoF} = 0.00219798$
 $\text{Sigma} = 0.0468826$

Mechanism for DNA unwinding of the 28T16bp substrate in 3 kinetic steps.

ED1 = ED2 = ED3 = ED4 = EP
ED1 = E + D1
ED2 = E + D2
ED3 = E + D3
ED4 = E + D4
EED1 = EED2 = EED3 = EED4 = EEP
EED1 = ED1
EED2 = ED2
EED3 = ED3
EED4 = ED4
EEEED1 = EEEED2 = EEEED3 = EEEED4 = EEEEP
EEEED1 = E + EED1
EEEED2 = E + EED2
EEEED3 = E + EED3
EEEED4 = E + EED4
EEEED1 = EEEAD1
EEEED2 = EEEAD2
EEEED3 = EEEAD3
EEEED4 = EEEAD4
EEAD1 = EEBD1 = EED1
EEAD2 = EEBD2 = EED2
EEAD3 = EEBD3 = EED3
EAD1 = EBD1 = EED1
EAD2 = EBD2 = EED2
EAD3 = EBD3 = EED3
AD1 = BD1
AD2 = BD2
AD3 = BD3
BD1 = ED1
BD2 = ED2
BD3 = ED3
AD1 = E + D1
AD2 = E + D2
AD3 = E + D3
BD1 = E + D1
BD2 = E + D2
BD3 = E + D3

Chi² = 0.00912995
Ch²/DoF = 0.000702303
Sigma = 0.026501