

Supporting Information for

Metal Ion Interactions in the Active Site of Human

Topoisomerase II α

Joseph E. Deweese, F. Peter, Guengerich, Alex B. Burgin, and Neil Osheroff

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FIGURE S1. DynaFit script and results files for fitting of pre-equilibrium k vs. Mg^{2+} ion concentration using variable values for the metal ion dissociation, DNA cleavage, and ligation steps and a fixed value for the metal ion binding. See Fig. 3 of main text for plot. A, the reaction scheme with variable values for the metal ion dissociation (k_{-1}), DNA cleavage (k_2), and ligation (k_{-2}) steps using a fixed value for the metal ion binding. B, same as in A with different starting values for k_{-1} , k_2 , and k_{-2} .

A

SCRIPT

```
;DYNAFIT script file    ./test/
;Basic mechanism for TOPO
; all units in nM, sec
;Trial 8

[task]

data  =  velocities
task   =  fit

[mechanism]

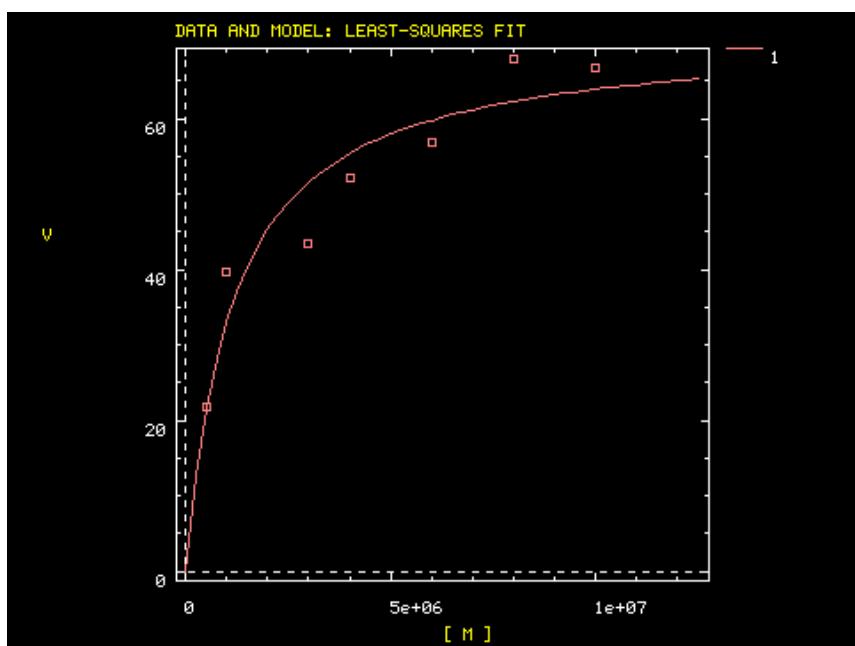
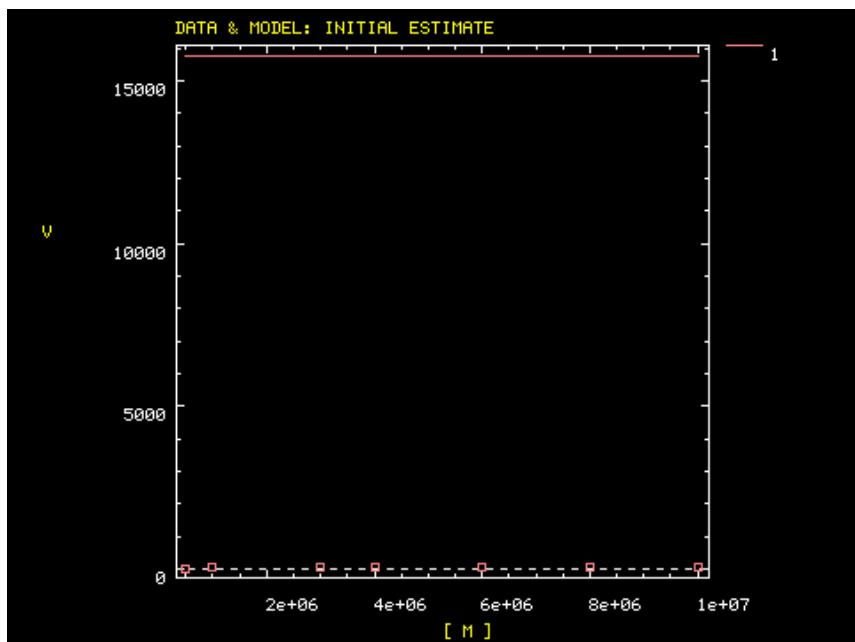
ED      +      M      <==>  EDM      :      k1      k-1
EDM     <==>  EPM     :      k2      k-2

[constants]
k1 = 1, k-1 = 200?
k2 = 200?, k-2 = 40?

[sweep]
[velocity]
variable M
file ./scripts/TopoIIrun.txt
[concentrations]
ED = 100

[responses]
EPM = 1
[progress]
delay = 0.001
[output]
directory ./projects/data/output/x
[end]
```

RESULTS



SCRIPT FILE

TASK

Fit of initial velocities

DATA

```
file :scripts:TopoIIrun.txt
```

OUTPUT

LEAST-SQUARES FIT

mean square	23.2506
standard deviation	4.82189
log(determinant)	-5.82
log(condition number)	-6.72
Marquardt parameter	1.64e+04
execution time (min)	0.05611
datapoints	7
parameters	3
iterations	20
subiterations	39
function evaluations	63
error status	0

PARAMETERS & STANDARD ERRORS

Set	Parameter	Initial	Fitted	Error	%Error
	k-1	200	1.161e+06	310000	27
	k2	200	8.056	290	3600
	k-2	40	2418	36000	1500

COVARIANCE MATRIX

Set	Parameter	Covariances		
	k-1	a	a	
	k2	b	76	b
	k-2	c	76	100

EIGENVECTORS AND EIGENVALUES

		Eigenvectors		
	Eigenvalues	0.00	0.37	2.63
	log(C)	7.03	0.85	0.00
Set	Parameter	1	2	3
	k-1	1	0	84
	k2	2	-70	37
	k-2	3	-70	-37
				52
				-60
				60

CONDITION INDICES

Set	Parameter	Index	7	1	0
	k-1	0	72	27	
	k2	50	13	36	
	k-2	49	14	36	

PLOTS

Best-fit plot: :projects:data:output:x:tab:fit_01.tab

B

SCRIPT

```
;DYNAFIT script file  ./test/
;Basic mechanism for TOPO
; all units in nM, sec
;Trial 10

[task]

data  =  velocities
task  =  fit

[mechanism]

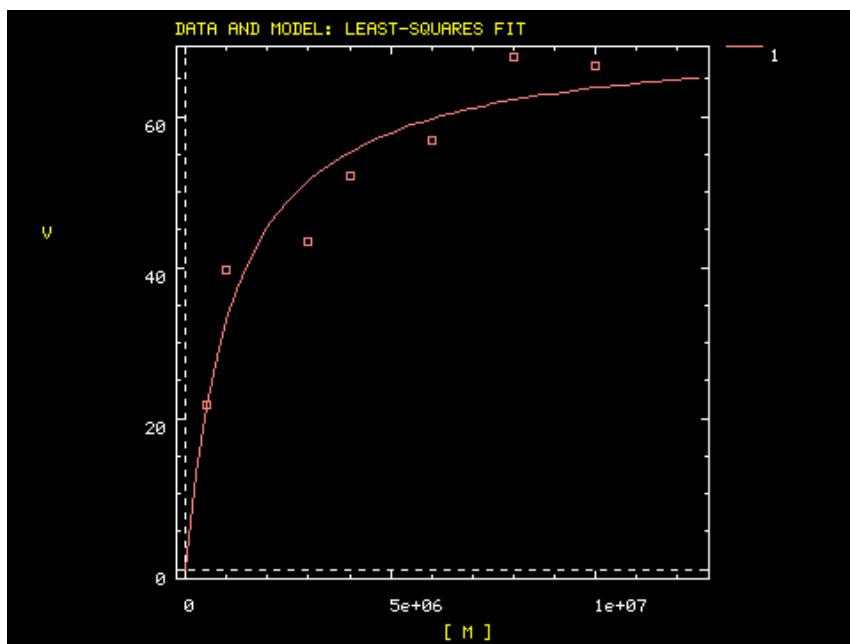
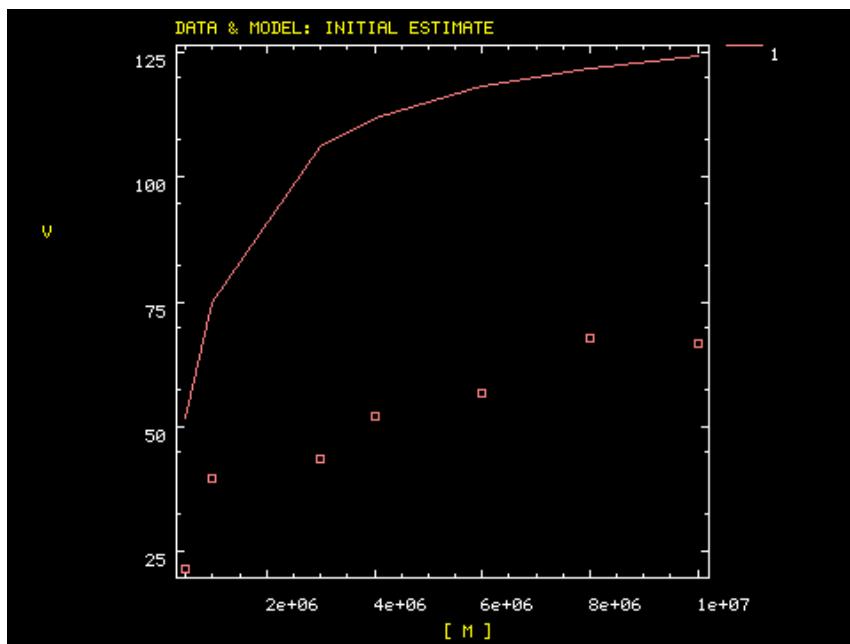
ED      +      M      <==>   EDM    :      k1      k-1
EDM    <==>  EPM    :      k2      k-2

[constants]
k1 = 0.01, k-1 = 10000?
k2  = 10?, k-2 = 2000?

[sweep]
[velocity]
    variable  M
    file    ./scripts/TopoIIrun.txt
[concentrations]
    ED  =  100

[responses]
    EPM = 1
[progress]
    delay = 0.001
[output]
    directory ./projects/data/output/x
[end]
```

RESULTS



SCRIPT FILE

TASK

Fit of initial velocities

DATA

file :scripts:TopoIIrun.txt

OUTPUT

LEAST-SQUARES FIT

mean square	23.2609
standard deviation	4.82295
log(determinant)	-2.78
log(condition number)	-8.92
execution time (min)	0.01333
datapoints	7
parameters	3
iterations	10
function evaluations	11
error status	1

PARAMETERS & STANDARD ERRORS

Set	Parameter	Initial	Fitted	Error	%Error
	k-1	10000	13860	470000	3400
	k2	10	7.218	3100	43000
	k-2	2000	2311	430000	19000

COVARIANCE MATRIX

Set	Parameter	Covariances
	k-1	a a
	k2	b 100 b
	k-2	c 100 100

EIGENVECTORS AND EIGENVALUES

Set	Parameter	Eigenvectors		
		Eigenvalues		
		0.00	0.21	2.79
	log(C)	8.90	1.11	0.00
		1	2	3
	k-1	1	-1	83 55
	k2	2	-71	37 -58
	k-2	3	-70	-40 58

CONDITION INDICES

Set	Parameter	Index	9	1	0
			0	69 30	
	k-1		50	14 34	
	k2		49	16 34	

PLOTS

Best-fit plot: :projects:data:output:x:tab:fit_01.tab

FIGURE S2. DynaFit script and results files for fitting of pre-equilibrium k vs. Mg²⁺ ion concentration using variable values for the DNA cleavage and ligation steps and fixed values for the metal ion binding and dissociation. See Fig. 3 of main text for plot. A, the

reaction scheme with variable values for the cleavage (k_2) and ligation (k_{-2}) steps using fixed values for the metal ion binding (k_1) and dissociation (k_{-1}). B , same as A using different starting values for k_2 and k_{-2} .

A

SCRIPT

```
;DYNAFIT script file  ./test/
;Basic mechanism for TOPO
; all units in nM, sec
;Trial 12

[task]

data  =  velocities
task  =  fit

[mechanism]

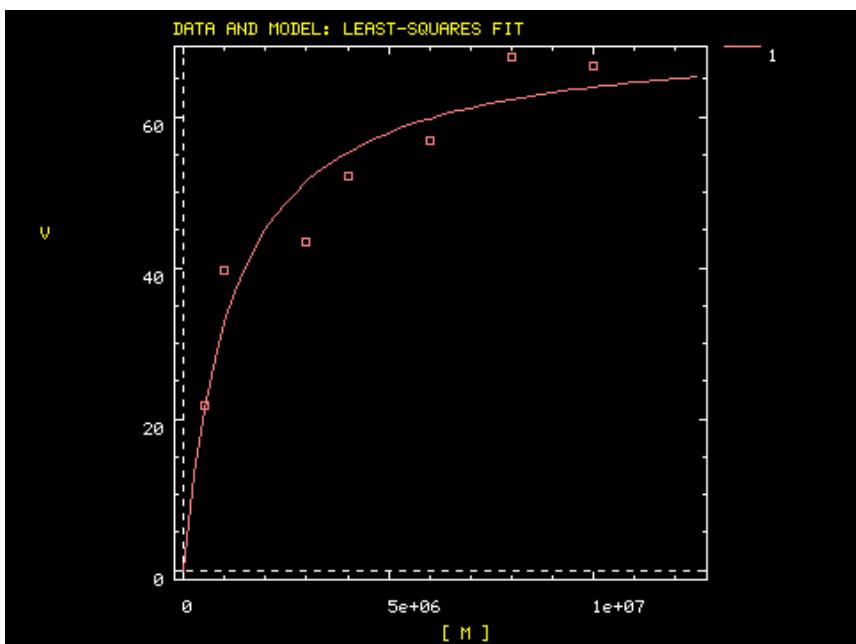
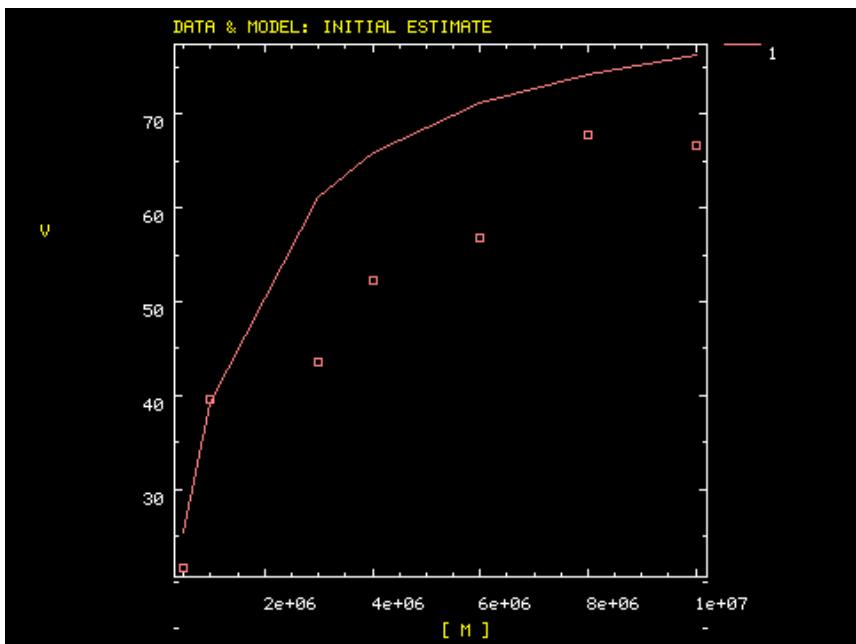
ED      +      M      <==>   EDM    :      k1      k-1
EDM    <==>  EPM    :      k2      k-2

[constants]
k1 = 0.1, k-1 = 120000
k2  =  7?, k-2 = 2100?

[sweep]
[velocity]
    variable  M
    file    ./scripts/TopoIIrun.txt
[concentrations]
    ED    =  100

[responses]
    EPM = 1
[progress]
    delay = 0.001
[output]
    directory ./projects/data/output/x
[end]
```

RESULTS



SCRIPT FILE

TASK

Fit of initial velocities

DATA

```
file :scripts:TopoIIrun.txt
```

OUTPUT

LEAST-SQUARES FIT

mean square	23.2593
standard deviation	4.82279
log(determinant)	-1.23
log(condition number)	-1.98
Marquardt parameter	32
execution time (min)	0.03778
datapoints	7
parameters	2
iterations	23
subiterations	19
function evaluations	45
error status	0

PARAMETERS & STANDARD ERRORS

Set	Parameter	Initial	Fitted	Error	%Error
	k2	7	6.472	1	16
	k-2	2100	2198	170	7.9

COVARIANCE MATRIX

Set	Parameter	Covariances
	k2	a a
	k-2	b 98

EIGENVECTORS AND EIGENVALUES

		Eigenvectors	
Eigenvalues		0.02	1.98
log(C)		1.98	0.00
Set	Parameter	1	2
	k2	1 -70	-70
	k-2	2 -70	70

CONDITION INDICES

Set	Parameter	Index	2	0
		k2	49	49
		k-2	49	49

PLOTS

```
Best-fit plot: :projects:data:output:x:tab:fit_01.tab
```

B

SCRIPT

```
;DYNAFIT script file  ./test/
;Basic mechanism for TOPO
; all units in nM, sec
;Trial 16

[task]

data  =  velocities
task  =  fit

[mechanism]

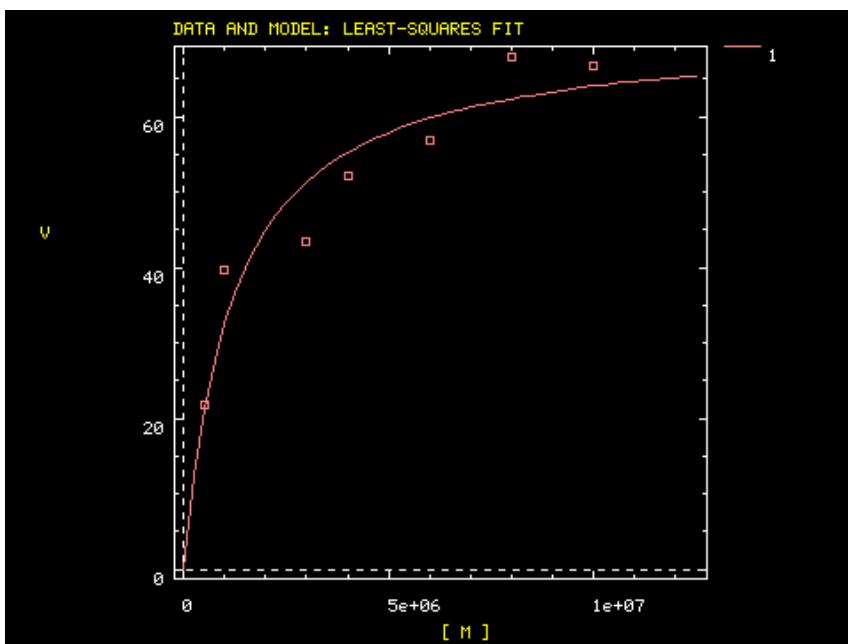
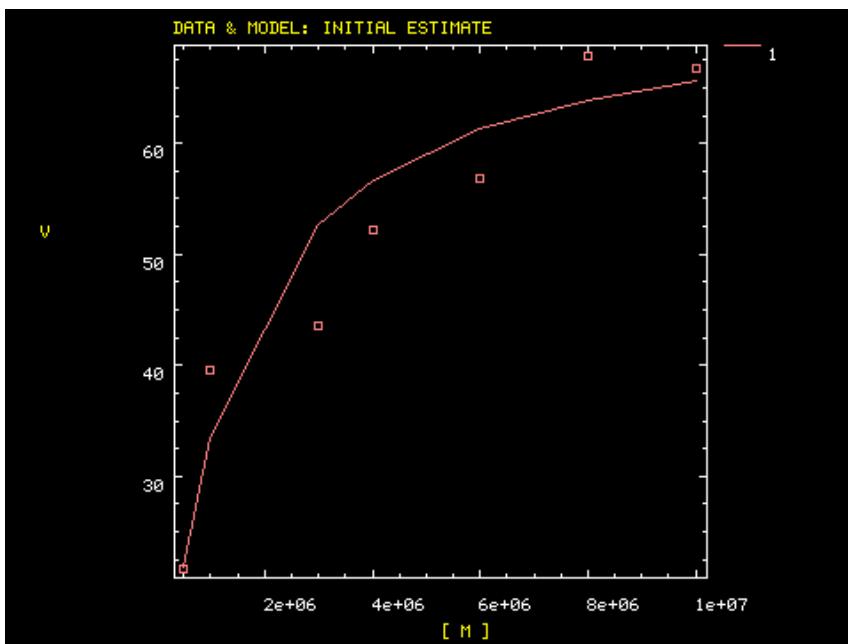
ED      +      M      <==>  EDM      :      k1      k-1
EDM    <==>  EPM      :      k2      k-2

[constants]
k1 = 0.1, k-1 = 120000
k2  =  2?, k-2 = 1000?

[sweep]
[velocity]
    variable  M
    file    ./scripts/TopoIIrun.txt
[concentrations]
    ED   =  100

[responses]
    EPM = 1
[progress]
    delay = 0.001
[output]
    directory ./projects/data/output/x
[end]
```

RESULTS



SCRIPT FILE

TASK

Fit of initial velocities

DATA

file :scripts:TopoIIrun.txt

OUTPUT

LEAST-SQUARES FIT

mean square	23.2976
standard deviation	4.82676
log(determinant)	-2.89
log(condition number)	-3.88
Marquardt parameter	1.31e+05
execution time (min)	0.03777
datapoints	7
parameters	2
iterations	11
subiterations	40
function evaluations	55
error status	0

PARAMETERS & STANDARD ERRORS

Set	Parameter	Initial	Fitted	Error	%Error
	k2	2	1.969	3.1	160
	k-2	1000	1009	1500	150

COVARIANCE MATRIX

Set	Parameter	Covariances
	k2	a a
	k-2	b 100

EIGENVECTORS AND EIGENVALUES

	Eigenvectors		
Eigenvalues	0.00	2.00	
log(C)	3.88	0.00	
Set	Parameter	1	2
	k2	1 -70	-70
	k-2	2 -70	70

CONDITION INDICES

Set	Parameter	Index	4	0
	k2	50	50	
	k-2	50	50	

PLOTS

Best-fit plot: :projects:data:output:x:tab:fit_01.tab