

## **Supporting Information**

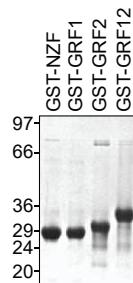
### **An autoinhibitory role for the GRF zinc finger domain of DNA glycosylase NEIL3**

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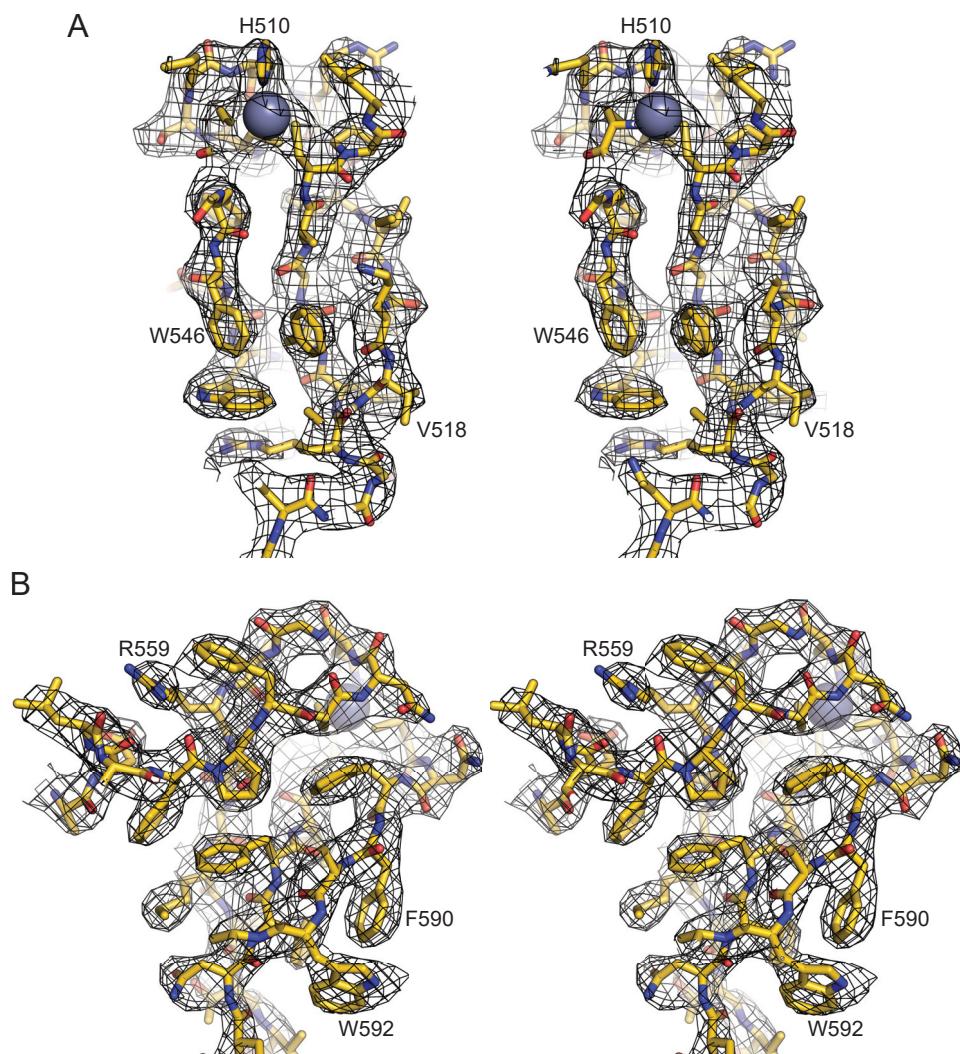
**Table S1.** DNA substrates composition from glycosylase assays

Substrate	Length (nt)	Sequence (5'-3')
<b>EMSA and fluorescence anisotropy</b>		
ssDNA	10	FAM-CTCAGGACTC
ssDNA	20	FAM-CTCAGGACTCAGTCGTCAG
ssDNA	30	FAM-CTCAGGACTCAGTCGTCAGCCCTTGACAG
ssDNA	40	FAM-CTCAGGACTCAGTCGTCAGCCCTTGACAGCGATGGAAGC
dsDNA	40	1: FAM-CTCAGGACTCAGTCGTCAGCCCTTGACAGCGATGGAAGC 2: GCTTCCATCGCTGTCAAGGGCTGACGAAC TGAGTCCTGAG
Fork	40	1: FAM-CTCAGGACTCAGTCGTCAG <u>CCCTGACAGCGATGGAAGC</u> 2: <u>CGAAGGTAGCGACAGTCCC</u> CTGACGAAC TGAGTCCTGAG
<b>Glycosylase assays</b>		
ssDNA	25	FAM-GACCACTACACC(DHT) <u>ATT</u> CCTTACAAC
Lead DHT fork	25	1: FAM-GACCACTACACC(DHT) <u>ATT</u> CCTTACAAC 2: <u>CAACATT</u> CCTTA <u>AG</u> GTGTAGTGGTC
Lag DHT fork	25	1: FAM-GACCACTACACC(DHT) <u>ATT</u> CCTTACAAC 2: GTTGTAAGGAATA <u>CC</u> ACATCACCAG
Lead AP-ICL fork	39	1: FAM-CCTTCAACCA <u>CC</u> ACCGCTCAACTC(AP)TG <u>A</u> ACTGCAGTCTGGTT 2: <u>CCGTT</u> CCGGTAGCGG <u>CA</u> AGAGTTGAGCGGTGGTTGAAGG
Lag AP-ICL fork	39	1: FAM-ATGGTCTGACGTGAATC(AP)TGCAACTGCCACCAACTTCC 2: GGAAGTTGGTGGCGAGTTGCAAG <u>T</u> GGGGATGGCCTTCCC
FAM, 6-carboxyfluorescein. Non-base-paired regions are shown in red. Damaged base pairs are underlined.		

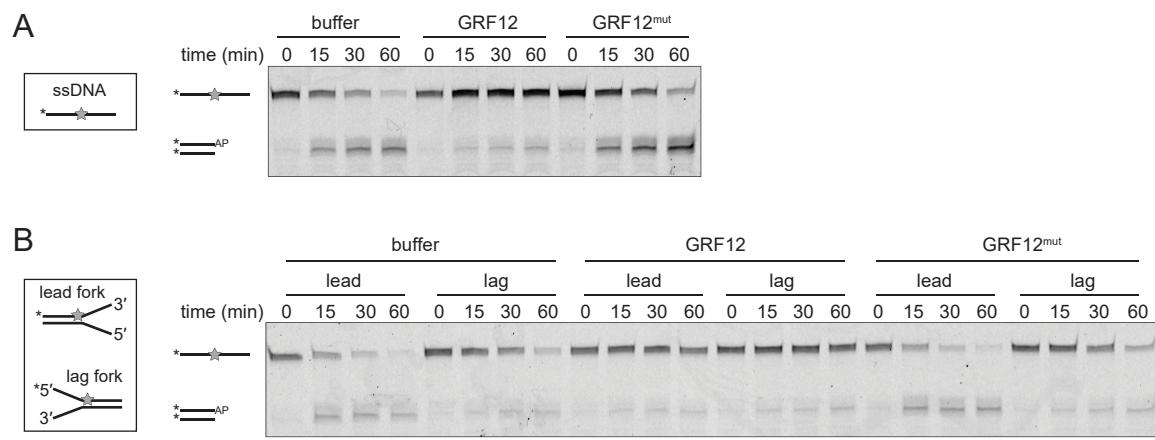
## Supplemental Figures



**Fig. S1.** Coomassie-stained SDS-PAGE gel of purified GST proteins used experiments shown in Fig. 1 and Fig. 2G.



**Fig. S2.** Stereoimages of representative regions of the refined model superimposed onto 2Fo-Fc electron density (contoured to  $1\sigma$ ). Labeled residues are for orientation purposes only. **A.** GRF1. **B.** GRF2.



**Fig. S3.** Representative gels used for data shown in Fig. 5. Glycosylase assays for GRF12 inhibition of 5,6-dihydrothymine in ssDNA (A) and forked substrates (B).