

Supplementary information for:

The Essential Role of Prolines and Their Conformation in Allosteric Regulation of Kaiso Zinc Finger DNA-Binding Activity by the Adjacent C-Terminal Loop

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Supplementary Table S1. Oligonucleotides used for cloning and EMSA. Protein binding sites are shown in bold.

1.1. Oligonucleotides used for cloning

Kaiso_P523Q_d	GTATCAGTGCCGTTACTGT
Kaiso_P523Q_r	ACAGTAACGGCACTGATAC
Kaiso_P577A_d	GATGCTCTGGGGACTCA
Kaiso_P577A_r	TGAGTCCCCAGAAGCATC
Kaiso_P588A_d	CATGCATGCAGGTCTTAC
Kaiso_P588A_r	GTAAAGACCTGCATGCATG
Kaiso_P588V_d	CATGTATGCAGGTCTTAC
Kaiso_P588V_r	GTAAAGACCTGCATACATG
Kaiso_579_r	TTGTCGACTTACCCAGAAGGGATCTTGACTA
Kaiso_593_r	TTGTCGACTTATTGAAAGACCTGCATGGAT
Kaiso_604_r	TTGTCGACTTATGATCTATCGGAAAGATATGC
Kaiso_472_d	TTGAATTGCAAACAAACGTATGAAAGTA

1.2. DNA probes for EMSA

Kaiso BSx3

AACAGCTATGACCATGATTACGCCAAGCTCGAAATTAAACCTCACTAAAGGGAACAAAAGCTGGAGCTCCACCG
CGGTGGCGGCCGCTCTAGAACTAGTGCT**TCTGCC**ATAACGGTGC**TCTGCC**ATAACGGTGC**TCTGCC**
ATAAGCTTATCGATACCGTCGACCTCGAGGGGGGGCCGGTACCCAATTGCCCTATAGTGAGTCGTATTACAA
TTCACTGGCGTCGTTTAC

Pita BSx5 as nonspecific DNA

GTAAAACGACGGCCAGTGAGCGCGCGTAATACGACTCACTATAGGGCGAATTGGGTACCGGGCCCCCTCGA
GGTCGACGGTATCGATAAGCTTGATCTTAG**CCAAGACGCGAACCCGAATCCGAAACTTAGCCAAGACGCGAACCCGAATCCG**
ACCCGAATCCGAAACTTAGCCAAGACGCGAACCCGAATCCGAAAGATCTAATATCGAATTCTGCAGCCGGGGATCCACTAG
AAACTTAGCCAAGACGCGAACCCGAATCCGAAAGATCTAATATCGAATTCTGCAGCCGGGGATCCACTAG
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TAATCATGGTCATAGCTGTT

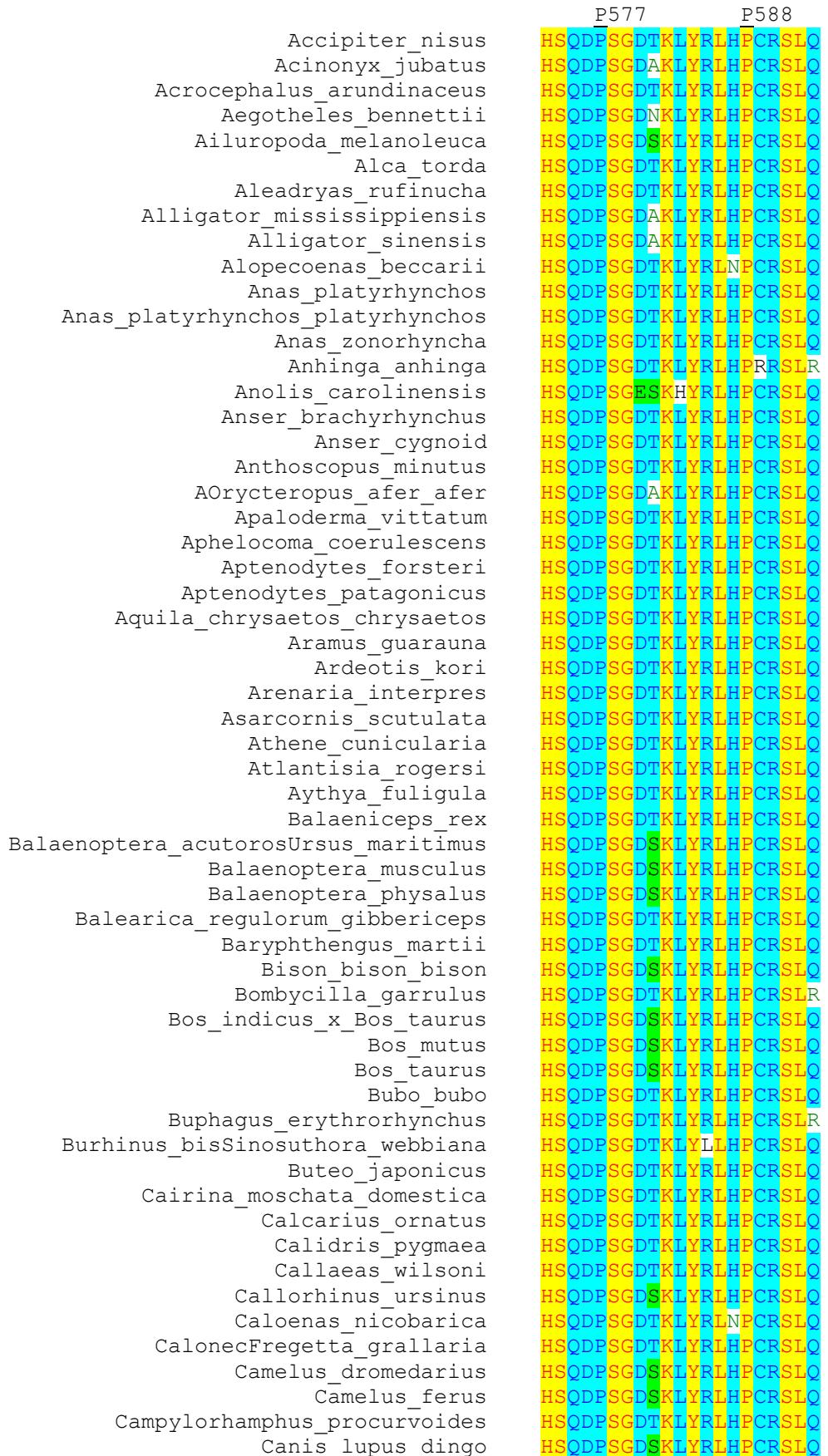
Supplementary Table S2. The presence of protein–DNA hydrogen bonds along last 90 ns of 100 ns molecular dynamics trajectories of Kaiso complexes with Kaiso binding sequence DNA. Only bonds with over 20% presence in wild-type protein are shown. Grey fill indicates H-bonds weakened 5x times or more compared to wild-type protein. Light grey fill indicates H-bonds weakened 2-5x times compared to wild-type protein.

donor	acceptor	wt	P523Q	P577A	P577 ^{cis}	P588A	P588V	P588 ^{cis}
TYR550-Side	DA30-PO4	85.41%	81.04%	81.72%	85.11%	85.91%	88.22%	68.06%
TYR562-Side	DG31-PO4	77.72%	74.75%	76.12%	67.03%	73.83%	74.15%	77.54%
THR538-Side	DG28-PO4	72.43%	72.46%	69.73%	61.04%	55.14%	53.09%	16.27%
TYR584-Side	DA30-PO4	71.73%	62.67%	68.73%	73.73%	68.73%	72.65%	66.77%
ARG549-Side	DC29-PO4	67.93%	85.83%	67.03%	60.24%	87.81%	68.06%	83.43%
TYR536-Side	DC8-PO4	66.63%	42.61%	17.98%	2.00%	0.90%	42.22%	2.40%
SER508-Side	DG10-PO4	63.84%	57.78%	62.14%	61.94%	66.33%	71.46%	41.82%
TYR503-Side	DT9-PO4	54.65%	69.16%	63.54%	66.33%	69.63%	65.17%	66.97%
ARG511-Side	DG10-Base	45.65%	60.98%	61.14%	60.84%	62.54%	49.50%	45.51%
ALA598-Main	DA14-Base	40.36%	41.22%	36.96%	38.16%	43.86%	43.01%	3.69%
ARG510-Side	DT25-PO4	39.56%	34.03%	8.49%	6.59%	16.08%	0.20%	0.90%
TYR522-Side	DG27-PO4	36.76%	44.41%	46.95%	36.06%	33.67%	10.18%	38.12%
TYR597-Side	DG28-Base	32.17%	40.72%	35.16%	32.87%	0.60%	45.11%	10.08%
ARG475-Side	DC18-PO4	31.77%	40.22%	25.57%	9.89%	6.59%	2.99%	6.69%
SER567-Side	DC4-Base	30.27%	60.68%	56.84%	0.00%	11.89%	20.46%	0.00%
ARG501-Side	DT9-PO4	25.37%	34.03%	20.58%	22.88%	25.37%	24.35%	24.45%
ARG595-Side	DG27-Base	24.98%	24.35%	21.98%	27.57%	11.09%	27.15%	2.30%
DC29-Base	GLU535-Side	22.08%	39.92%	39.66%	42.86%	42.66%	19.16%	44.71%
LYS477-Side	DC18-PO4	20.78%	16.57%	0.10%	0.40%	8.89%	0.10%	27.84%

Supplementary Table S3. The presence of protein–DNA hydrogen bonds along last 90 ns of 100 ns molecular dynamics trajectories of Kaiso complexes with CG-methylated DNA. Only bonds with over 15% presence in wild-type protein (and the bond E535-5MC8, which was previously found to be important for specific methyl-DNA recognition) are shown. Designations are the same as in Supplementary Table 2.

donor	acceptor	wt	P577A	P577^{cis}	P588A	P588V	P588^{cis}
ARG549-Side	5MC30-PO4	100.00%	100.00 %	92.71%	100.00 %	100.00%	100.00 %
ARG511-Side	DG9-Base	100.00%	56.74%	43.66%	90.11%	70.83%	33.57%
TYR503-Side	5MC8- PO4	83.62%	79.82%	65.13%	80.02%	70.73%	82.52%
THR538-Side	DG29-PO4	73.83%	31.37%	56.44%	71.13%	68.13%	4.30%
TYR584-Side	DG31-PO4	73.13%	79.82%	63.74%	77.72%	76.72%	76.52%
TYR562-Side	DG32- PO4	71.23%	70.93%	58.14%	64.94%	69.53%	65.73%
TYR550-Side	DG31-Side	70.63%	68.93%	65.43%	69.03%	69.03%	60.04%
TYR522-Side	5MC28- PO4	63.44%	28.57%	39.76%	64.24%	43.96%	7.69%
SER508-Side	5MC8- PO4	37.36%	7.89%	0.0%	9.49%	0.10%	0.0%
VAL504-Main	DG9- PO4	28.57%	52.05%	45.65%	47.35%	51.15%	61.64%
ALA598-Main	DC13- PO4	24.48%	19.18%	25.97%	31.27%	14.69%	10.09%
5MC30-Side	GLU535-Side	24.48%	3.80%	10.89%	10.59%	17.88%	1.10%
ARG595-Side	DG29-PO4	20.88%	1.90%	1.00%	9.09%	0.20%	8.09%
ARG595-Side	5MC28-Base	19.68%	28.17%	29.57%	26.57%	28.77%	17.58%
SER578-Side	DG32- PO4	17.98%	1.70%	2.90%	1.40%	5.39%	0.60%
TYR597-Side	DG29-Base	16.48%	16.58%	15.88%	26.17%	18.18%	25.47%
CYS505-Main	DG9-PO4	15.98%	1.40%	2.60%	4.40%	1.00%	0.0%
GLN563-Side	DT33-Base	14.79%	3.50%	5.39%	4.60%	5.59%	2.00%
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5MC8-Side	GLU535-Side	1.70%	6.09%	6.19%	9.99%	1.50%	1.20%

Supplementary Figure S1. Multiple sequence alignment of C-terminal extensions of Kaiso proteins from 400 species of Vertebrates.



Canis_lupus_familiaris	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Canis_lupus_familiaris_0	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Canis_lupus_familiaris_1	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Capra_hircus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cardinalis_cardinalis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cariama_cristata	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Catagonus_wagneri	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Catharus_fuscescens	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Catharus_ustulatus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cavia_porcellus_	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cephalopterus_ornatus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cephus_grylle	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Certhia_brachydactyla	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Certhia_familiaris	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cettia_cetti	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chaetorhynchus_papuensis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Charadrius_vociferus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chauna_torquata	HSQDP	SGDA	KLYR	LHPC	CRS	IQ
Chelonia_mydas	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chelonoidis_abingdonii	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chelydra_serpentina	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chinchilla_lanigera	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Chlamydotis_macqueenii	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chloebia_gouldiae	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chloropsis_cyanopogon	HSQDP	SGNT	KLYR	LHPC	CRS	IQ
Chloropsis_hardwickii	HSQDP	SGNT	KLYR	LHPC	CRS	IQ
Chordeiles_acutipennis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chroicocephalus_maculipennis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chrysemys_picta_bellii	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Chunga_burmeisteri	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Ciccaba_nigrolineata	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Ciconia_maguari	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cinclus_mexicanus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Circaetus_pectoralis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cisticola_juncidis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Climacteris_rufus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cnemophilus_loriae	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cochlearius_cochlearius	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Columbina_picui	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Copsychus_sechellarum	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Corvus_brachyrhynchos	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Corvus_monedulaoides	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Corythaeola_cristata	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Corythaixoides_concolor	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Crocodylus_porosus	HSQDP	SGDA	KLYR	LHPC	CRS	IQ
Crocuta_crocuta	HSQDP	SGDA	KLYR	LHPC	CRS	IQ
Cyanistes_caeruleus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cyanoderma_ruficeps	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Cyanoderma_ruficeps_0	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Daphoenositta_chrysoptera	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Dasyornis_broadbenti	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Dicaeum_eximium	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Dicrurus_megarhynchus	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Donacobius_aVidua_chalybeata	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Dromas_ardeola	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Drymodess_brunneopygia	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Dryoscopus_gambensis	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Edolisoma_coeruleuscens	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Egretta_garzetta	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Elachura_formosa	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Emberiza_fucata	HSQDP	SGDT	KLYR	LHPC	CRS	IQ
Enhydra_luDelphinapterus_leucas	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Equus_asinus_asinus	HSQDP	SGDS	KLYR	LHPC	CRS	IQ
Eriothacus_rubecula	HSQE	PSGD	TKLYR	LHPC	CRS	IQ

Erpornis_zantholeuca	HSQDP	SGDTKLYRILHPCRSI
Erythrocercus_mccallii	HSQDP	SGDTKLYRILHPCRSI
Eschrichtius_robustus	HSQDP	SGDSKLYRILHPCRSI
Eudyptes_chrysocome	HSQDP	SGDTKLYRILHPCRSI
Eudyptes_filholi	HSQDP	SGDTKLYRILHPCRSI
Eudyptes_robustus	HSQDP	SGDTKLYRILHPCRSI
Eudyptes_sclateri	HSQDP	SGDTKLYRILHPCRSI
Eudyptula_minor	HSQDP	SGDTKLYRILHPCRSI
Eulacestoma_nigropectus	HSQDP	SGDTKLYRILHPCRSI
Eurypyga_helias	HSQDP	SGDTKLYRILHPCRSI
Eurystomus_gularis	HSQDP	SGDTKLYRILHPCRSI
Falco_tinnunculus	HSQDP	SGDTKLYRILHPCRSI
Falcunculus_frontatus	HSQDP	SGDSKLYRILHPCRSI
Felis_catus	HSQDP	SGDTKLYRILHPCRSI
Ficedula_albicollis	HSQDP	SGDTKLYRILHPCRSI
Formicarius_rufipectus	HSQDP	SGDTKLYRILHPCRSI
Fregata_magnificens	HSQDP	SGDTKLYRILHPCRSI
Fukomys_damarensis	HSQDP	SGDSKLYRILHPCRSI
Fulmarus_glacialis	HSQDP	SGDTKLYRILHPCRSI
Furnarius_figulus	HSQDP	SGDTKLYRILHPCRSI
Gavia_stellata	HSQDP	SGDTKLYRILHPCRSI
GeoGeoMyotis_myotis	HSQDP	SGDSKLYRILHPCRSI
Geospiza_fortis	HSQDP	SGDTKLYRILHPCRSI
Geospiza_parvula	HSQDP	SGDTKLYRILHPCRSI
Glareola_pratincola	HSQDP	SGDTKLYRILHPCRSI
Glaucidium_brasiliandum	HSQDP	SGDTKLYRILHPCRSI
Gopherus_agassizii	HSQDP	SGDTKLYRILHPCRSI
Gopherus_evgoodei	HSQDP	SGDTKLYRILHPCRSI
Gorilla_gorilla_gorilla	HSQDP	SGDSKLYRILHPCRSI
Grallaria_varia	HSQDP	SGDTKLYRILHPCRSI
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Herpetotheres_cachinnans	HSQDP	SGDTKLYRILHPCRSI
Himantopus_himantopus	HSQDP	SGDTKLYRILHPCRSI
Hippolais_icterina	HSQDP	SGDTKLYRILHPCRSI
Hipposideros_armiger	HSQDP	SGESKLYRILHPCRSI
Hirundo_rustica_rustica	HSQDP	SGDTKLYRILHPCRSI
Homo_sapiens	HSQDP	SGDSKLYRILHPCRSI
Horornis_vulcanius	HSQDP	SGDTKLYRILHPCRSI
Hypotaenidia_okinawae	HSQDP	SGDTKLYRILHPCRSI
Ibidorhyncha_sthryothorus_ludovicianus	HSQDP	SGETKMYRILHPCRSI
Ifrita_kowaldi	HSQDP	SGDTKLYRILHPCRSI
Irena_cyanogasteranas_sempalmata	HSQDP	SGDTKLYRILHPCRSI
Jacana_jacana	HSQDP	SGDTKLYRILHPCRSI
Junco_hyemalis	HSQDT	SGDTKLYRILHPCRSI
Lamprotornis_superbus	HSQDP	SGDTKLYRILHPCRSI
Lanius_ludovicianus	HSQDP	SGDTKLYRILHPCRSI
Larus_smithsonianus	HSQDP	SGDTKLYRILHPCRSI
Leiothrix_lutea	HSQDP	SGDTKLYRILHPCRSI
Lepidothrix_coronata	HSQDP	SGDTKLYRILHPCRSI
Leptocoma_aspasia	HSQDP	SGDTKLYRILHPCRSI
Leptonychotes_weddellii	HSQDP	SGDSKLYRILHPCRSI
Leptosomus_discolor	HSQDP	SGDTKLYRILHPCRSI
Leucopsar_rothschildi	HSQDP	SGDTKLYRILHPCRSI
Limosa_lapponica_baueri	HSQDP	SGDTKLYRILHPCRSI
Lipotes_vexillifer	HSQDP	SGDSKLYRILHPCRSI
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Lonchura_striata_domestica	HSQDP	SGDTKLYRILHPCRSI

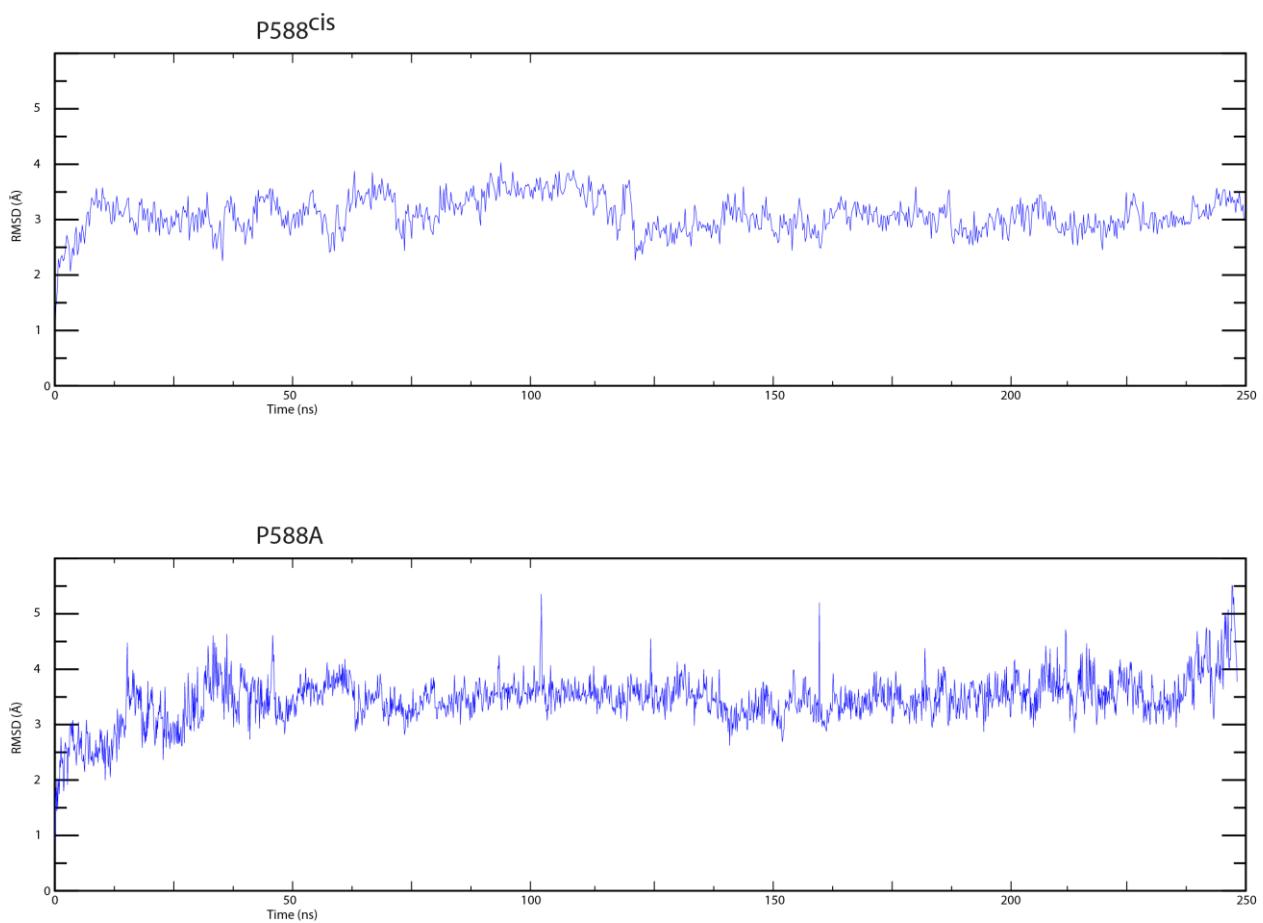
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Malurus_cyaneus_samueli
Malurus_elegans
Marmota_monax
Marmota_monax_0
Melospiza_melodia
Mesembrinibis_cayennensis
Mesitornis_unicolor
Microcaecilia_unicolor
Mionectes_macconnelli
Molossus_molossus
Molothrus_aterrimus
Monodon_monoceros
Moschus_moschiferus
Motacilla_alba
Muntiacus_muntjak
Muntiacus_reevesi
Mus_caroli
Mus_musculus
Mus_musculus_0
Mus_spicilegus
Myiagra_hebetior
Myotis_brandtii
Myotis_davidii
Myotis_lucifugus
Mystacornis_crossleyi
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Neomonachus_schauinslandi
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Neopipo_cinnamomea
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Nipponia_nippon
Notiomystis_cincta
Nyctereutes_procyonoides
Nyctibius_bracteatus
Nyctibius_grandis
Nycticryphes_semicollaris
Nyctiprogne_leucopyga
Oceanites_oceanicus
Oceanodroma_tethys
Octodon_degus
Odobenus_rosmarus_divergens
Oenanthe_oenanthe
Onychorhynchus_coronatus
Opisthomus_hoazin
Oreocharis_arfaki
Origma_solitaria
Oriolus Oriolus
Ornithorhynchus_anatinus
Otus_sunia
Ovis_aries
Ovis_aries_0
Oxyrhynchus_madagascariensis
Oxyruncus_cristatus
Pachycephala_philippinensis
Pachyramphus_minor
Pan_paniscus

HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDAK	I	Y	R	L	H	P	C	R	S	I	Q	
HS	QDP	S	GDAK	I	Y	R	L	H	P	C	R	S	I	Q	
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	K	S	I	Q
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDSK	I	Y	R	L	H	P	C	K	S	I	Q	
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
HS	QDP	S	GDT	K	I	Y	R	L	H	P	C	R	S	I	Q
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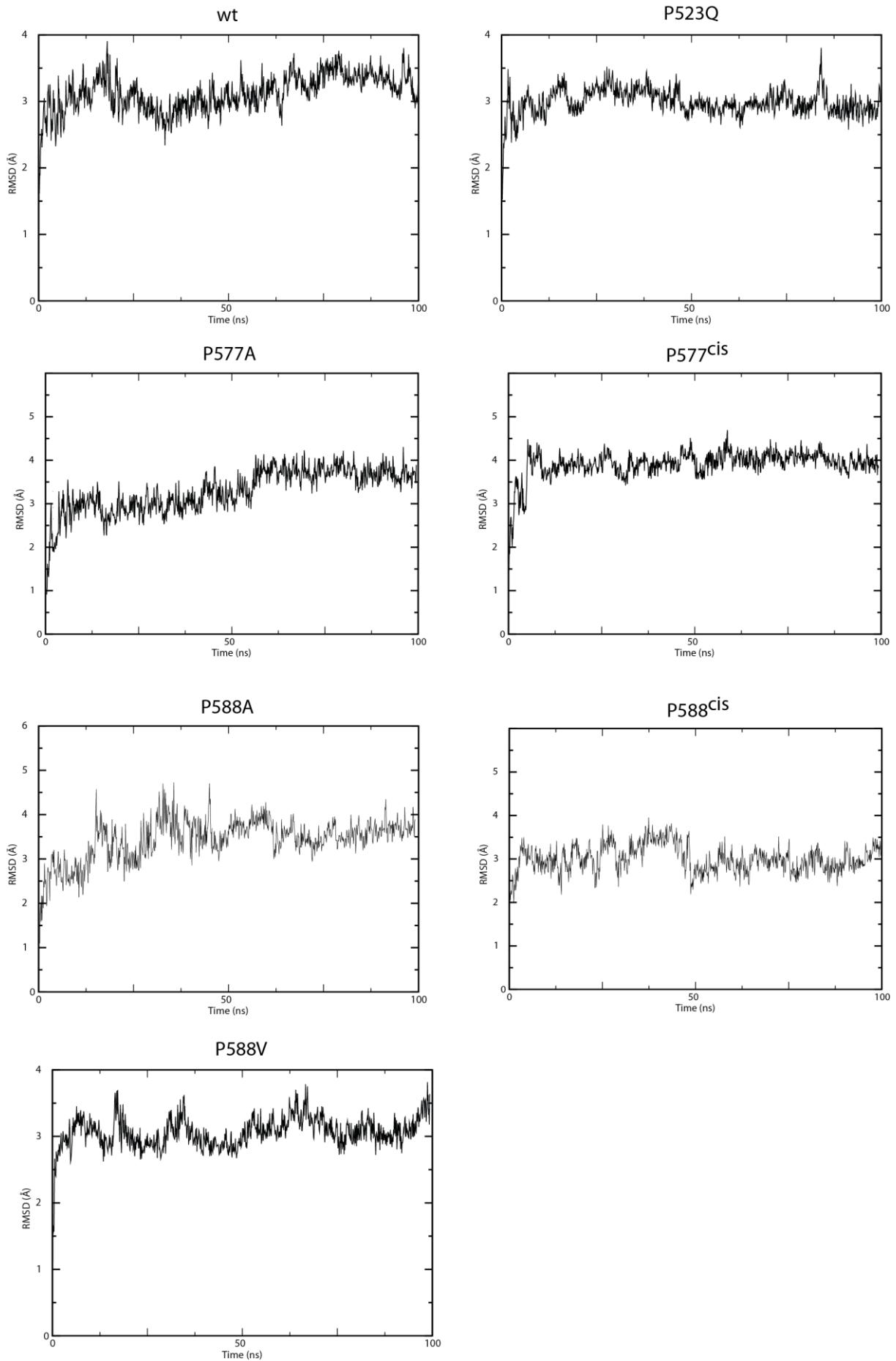
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	Serinus_canaria	HSQDP	SGDT	KLY	R	LH	PCRSI	Q		
	Setophaga_kirtlandii	HSQDP	SGDT	KLY	R	LH	PCRSI	Q		
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	Sylvia_aChionis_minor	HSQDP	SGDT	KLY	R	LH	PCRSI	Q		
	Syrrhaptes_paradoxus	HSQDP	SGDT	KLY	R	LH	PCRSI	Q		
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		Terrapene_carolina_Bos_mutus	HSQDP	SGD	S	KLY	R	LH	PCRSI	Q
		Thalassarche_chlororhynchos	HSQDP	SGDT	KLY	R	LH	PCRSI	Q	
		Thinocorus_orbignyanus	HSQDP	SGDT	KLY	R	LH	PCRSI	Q	
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		Todus_mexicanus	HSQDP	SGDT	KLY	R	LH	PCRSI	Q	
		Toxostoma_redivivum	HSQDP	SGDT	KLY	R	LH	PCRSI	R	
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		Xiphorhynchus_elegans	HSQDP	SGDT	KLY	R	LH	PCRSI	Q	
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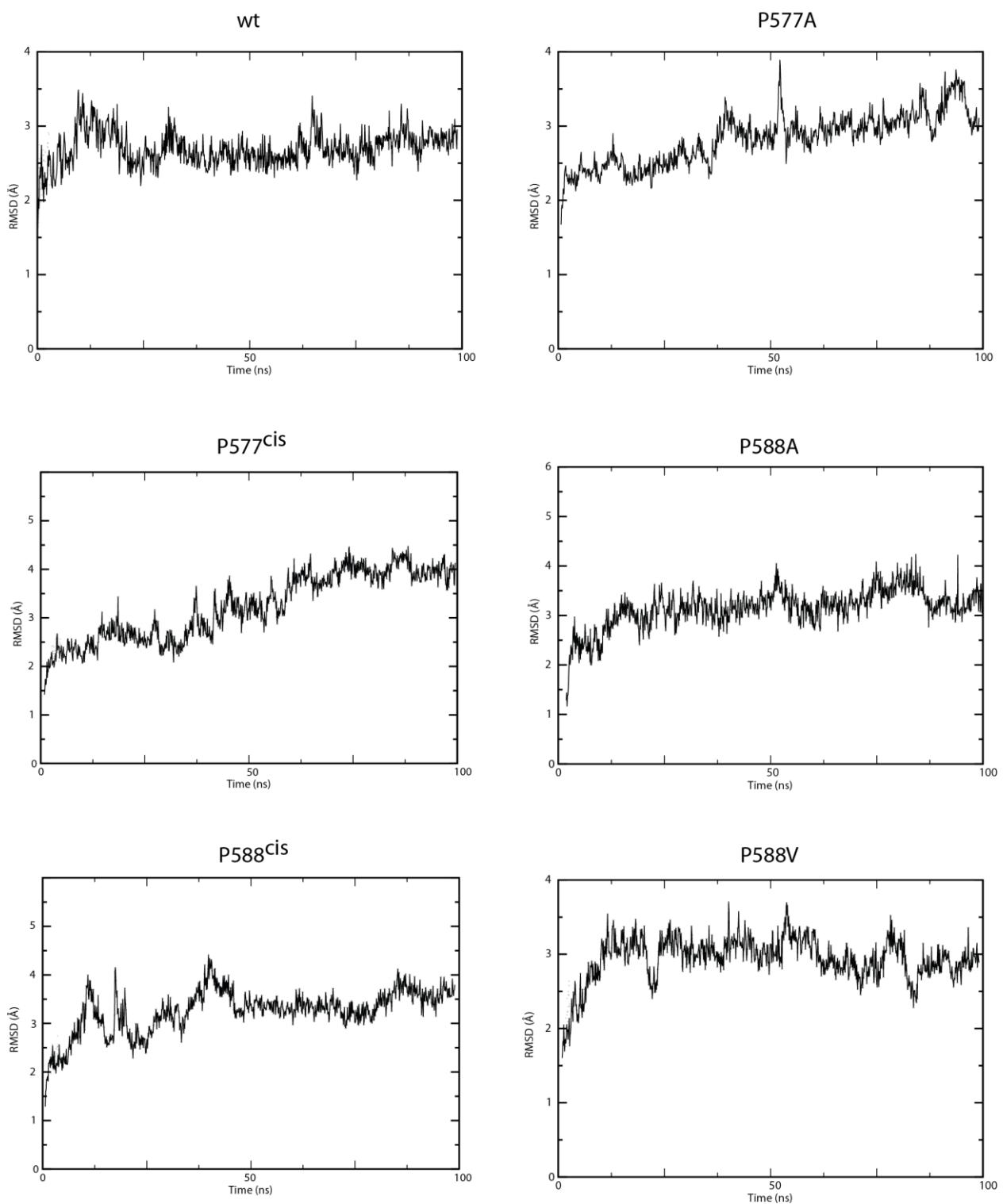
Supplementary Figure S2. RMSD of 250 ns molecular dynamics simulation trajectories of Kaiso P588A and P588^{cis} complexes with KBS DNA.



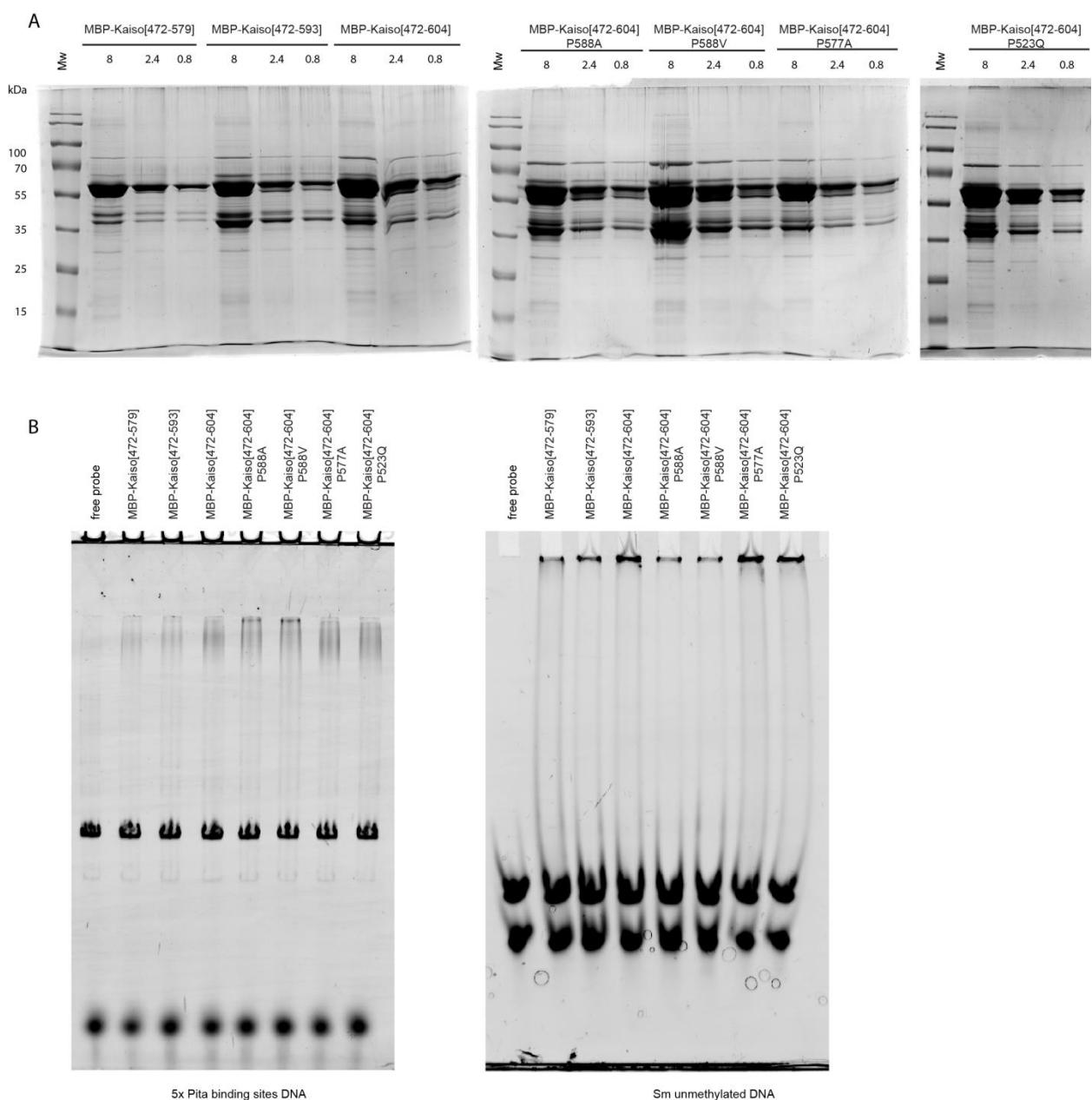
Supplementary Figure S3. RMSD of molecular dynamics simulation trajectories of Kaiso complex with KBS DNA.



Supplementary Figure S4. RMSD of molecular dynamics simulation trajectories of Kaiso complex with methylated DNA.



Supplementary Figure S5. (a) Protein preparations used for EMSA. (b) Non-specific binding controls at 10 μ M.



Supplementary Figure S6. Uncropped gels of EMSA.

