Structural Basis for Left-Handed Z-DNA Binding by Human dsRNA Specific Adenosine Deaminase ADAR1

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Thomas Schwartz

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1. Gutachter : Prof. Dr. Udo Heinemann
2. Gutachter : Prof. Dr. Volker A. Erdmann

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II. Abbreviations

Å	Ångstrøm, 1 Å = 10^{-10} m
a.u.	asymmetric unit
aa	amino acid
acc	accession number
ADAR1	dsRNA dependent adenosine deaminase
β-ME	2-mercaptoethanol
bp	base pair
CD	circular dichroic
cDNA	complementary DNA
DNA	deoxyribonucleic acid
ds	double-stranded
DTT	dithiothreitol
E. coli	*Escherichia coli*
EDTA	ethylenediaminetetraacetic acid
EMSA	electrophoretic mobility shift assay
FPLC	fast performance liquid chromatography
HEPES	N-[2-Hydroxyethl]piperazine-N'-[4-butanesulfonic acid]
HTH	helix-turn-helix
kD	kiloDalton, 1 kD = 1.66018 • 10^{-21} g
M	molarity, [mol • l^{-1}]
MALDI-TOF	matrix assisted laser desorption ionization – time of flight
MOPS	3-[N-Morpholino]propanesulfonic acid
mRNA	messenger RNA
NCS	non-crystallographic symmetry
PAGE	polyacrylamide gel electrophoresis
PCR	polymerase chain reaction
PDB	Protein Data Bank
PMSF	phenylmethylsulfonyl flouride
rmsd	root mean square deviation
RNA	ribonucleic acid
SDS	sodium dodecyl sulfate
SIRAS	single isomorphous replacement including anomalous scattering
Tris	tris[hydroxymethyl]aminomethane
UV	ultra violet

V8	endoproteinase Glu-C
Zα	Z-DNA binding domain of ADAR1
Zβ	Zα-related domain in ADAR1
Zab	bipartite domain in ADAR1, consisting of Zα and Zβ

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