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Structural basis for error-free replication of oxidatively ...

For example, in phage T7 Pol, C is preferred over A because a lysine (Lys536) in the fingers domain sterically and/or electrostatically interferes with 8-oxoG when it is rotated from the anti to the syn conformation (Brieba et al., 2004; Brieba et al., 2005).

The Fpg/Nei Family of DNA Glycosylases: Substrates ...

Overall, the structures of the Fpg and Nei proteins are similar, with a distinct 2-domain architecture connected by a flexible hinge region (Fig. 1A and B using EcoFpg and EcoNei as examples).55, 56, 72 In general, the N-terminal region is predominantly  $\beta$ -sheet-rich and is composed of a  $\beta$ -sandwich flanked by  $\alpha$ -helices.

Evolution of the Genetic Code by Incorporation of Amino ...

Valine. Val is the simplest of the highly hydrophobic amino acids and has a  $\beta$ -branched aliphatic side chain. As expected, it is a component of TM protein domains and the cores of globular proteins.

Differing Conformational Pathways Before and After ...

To elucidate how human DNA polymerase ( $\text{pol } \beta$ ) discriminates dATP from dCTP when processing 8-oxoguanine (8-oxoG), we analyze a series of dynamics simulations before and after the chemical step with dATP and dCTP opposite an 8-oxoG template started from partially open complexes of  $\text{pol } \beta$ .

Structural basis for DNA-hairpin promoter recognition by ...

Coliphage N4 virion-encapsidated RNA polymerase (vRNAP) is a member of the phage T7-like single-subunit RNA polymerase (RNAP) family. Its central domain (mini-vRNAP) contains all RNAP functions of the full-length vRNAP, which recognizes a five- to seven-base pair stem and three-nucleotide loop hairpin DNA promoter.

Identification of enolase as a plasminogen-binding protein ...

[Show full abstract] crystal structure of alpha-enolase from *Streptococcus pneumoniae* at 2.0Å resolution, the first structure both of a plasminogen-binding and of an octameric alpha-enolase. While

Crystallography and Site-directed Mutagenesis of Yeast ...

Except if there is any hope of completely understanding the catalytic action of any enzyme, surely the best candidate for the protein is triosephosphate isomerase

(TIM).

WO1997048722A1 - Pyridine nucleotide-dependent enzymes ...

The invention discloses a bisubstrate analogue effective at inhibiting pyridine nucleotide-dependent enzyme(s).

Additionally, disclosed is a method for manufacture of said binding substrate analogue and a complex and drug relating to same. The analogue comprises Diazaborine or part thereof for use in the rational design of bisubstrate enzyme