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Regenerating the Cofactor

When used as a cofactor, tetrahydrobiopterin grabs molecular oxygen and attaches one oxygen atom to the target molecule, but it ends up having the other oxygen attached to itself in the process. Two enzymes are then needed to strip off this unwanted oxygen and restore the cofactor. The first, pterin-4a-carbinolamine dehydratase (PCD, PDB entry <u>ldcp</u>), strips off the oxygen and releases it as a water molecule. The second, dihydropteridine reductase (DHPR, PDB entry <u>ldrp</u>), adds a few hydrogen atoms and the cofactor is ready to go for the next reaction.



click on the above Jmol tab for an interactive visualization

Exploring the Structure

A recent crystallographic analysis revealed the structural basis for side effects caused by sulfa drugs. These drugs revolutionized medicine in the 1930s, being one of the first effective antibiotics for fighting bacterial infections. However, they show some side effects with the nervous system when used at high doses. A recent structure (PDB entry 4j7u), revealed that the drugs bind to sepiapterin reductase, blocking its action and ultimately blocking the production of neurotransmitters. To explore the structure of a sulfa drug bound to this enzyme, as well as the structure of the enzyme with its intended substrate (PDB entry <a>1sep__), click the image for an interactive JSmol.

Topics for further exploration

1. To see tetrahydrobiopterin in action, try searching for structures of phenylalanine hydroxylase, tyrosine hydroxylase and nitric oxide synthase. 2. More information about tetrahydrobiopterin is available at the Ligand Summary page.

References

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