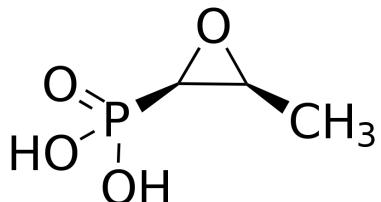


Fosfomycin

Fosfomycin

	
Systematic (IUPAC) name	
$[(2R,3S)-3\text{-methyloxiran}-2\text{-yl}] \text{phosphonic acid}$	
Identifiers	
CAS number	23155-02-4 ^[1] 78964-85-9 ^[2] (tromethamine)
ATC code	J01 XX01 ^[3]
PubChem	CID 446987 ^[4]
DrugBank	APRD00987 ^[5]
Chemical data	
Formula	$\text{C}_3\text{H}_7\text{O}_4\text{P}$
Mol. mass	138.059 g/mol
Pharmacokinetic data	
Bioavailability	30–37% (oral, fosfomycin tromethamine); varies with food intake
Protein binding	Nil
Metabolism	Nil
Half-life	5.7 hours (mean)
Excretion	Renal and fecal, unchanged
Therapeutic considerations	
Pregnancy cat.	B(US)
Legal status	℞-only (US)
Routes	Oral
✓ (what is this?) (verify) ^[6]	

Fosfomycin (also known as phosphomycin and phosphonomycin) is a broad-spectrum antibiotic^[7] produced by certain *Streptomyces* species.

Uses

Fosfomycin is indicated in the treatment of urinary tract infections, where it is usually administered as a single oral megadose.^[8]

The drug is well tolerated and has a low incidence of harmful side-effects.^[8] However, development of bacterial resistance under therapy is a frequent occurrence and makes fosfomycin unsuitable for sustained therapy of severe infections.

Additional uses have been proposed.^[9] The global problem of advancing antimicrobial resistance has led to a renewed interest in its use more recently.^[10]

Mechanism of action

Fosfomycin inhibits bacterial cell wall biogenesis by inactivating the enzyme UDP-*N*-acetylglucosamine-3-enolpyruvyltransferase, also known as MurA.^[11] This enzyme catalyzes the committed step in peptidoglycan biosynthesis, namely the ligation of phosphoenolpyruvate (PEP) to the 3'-hydroxyl group of UDP-*N*-acetylglucosamine. This pyruvate moiety provides the linker that bridges the glycan and peptide portion of peptidoglycan. Fosfomycin is a PEP analog that inhibits MurA by alkylating an active site cysteine residue (Cys 115 in the *Escherichia coli* enzyme).^[12]

Fosfomycin enters the bacterial cell through the glycerophosphate transporter.

Biosynthetic gene cluster

The complete fosfomycin biosynthetic gene cluster from *Streptomyces fradiae* has been cloned and sequenced and the heterologous production of fosfomycin in *Streptomyces lividans* has been achieved by Ryan Woodyer of the Huimin Zhao and Wilfred van der Donk research groups.^[13]

Resistance

Mutations that inactivate the non-essential glycerophosphate transporter render bacteria resistant to fosfomycin.^[14] [15]

Fosfomycin resistance enzymes

Enzymes conferring resistance to fosfomycin have also been identified and are encoded both chromosomally and on plasmids.^[16]

Glyoxalase superfamily enzymes

Three related but mechanistically distinct fosfomycin resistance enzymes (named, FosA, FosB and FosX) function by nucleophilic attack on carbon 1 of fosfomycin. This opens the epoxide ring and renders the drug ineffective. The enzymes differ by the identity of the nucleophile utilized in the reaction; glutathione for FosA, cysteine for FosB, and water for FosX.^[16]

FosC

FosC utilizes ATP and adds a phosphate group to fosfomycin, thus altering its properties and making the drug ineffective.^[17]

See also

- Fosmidomycin

External links

- Fosfomycin information at RxList^[18]

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