

Zinc phosphodiesterase – a new member of the metallo beta-lactamase superfamily

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The *elaC* gene of *Escherichia coli* encodes a binuclear zinc phosphodiesterase (ZiPD) [1]. ZiPD belongs to the metallo- β -lactamase superfamily of proteins. Proteins sharing this fold catalyze a wide variety of different reactions. Hydrolytic metallo- β -lactamase proteins mostly bind zinc, recombinant glyoxalase II (a thiolesterase) is isolated with iron, manganese and zinc, while redox-active rubredoxin-oxygen-oxidoreductase requires a bi-iron site. In general, the metallo- β -lactamase fold allows for remarkably different catalytic activities and metal selectivities. ZiPD homologues are found in all domains of life and the homolog gene in *B. subtilis* is essential for viability [2]. In order to characterize the function of *E. coli* gene and the gene product:

- (a) the metal binding residues were identified by mutational exchange and kinetic as well as X-ray absorption spectroscopic analysis of the mutant proteins;
- (b) the enigmatic physiological function of *E. coli* ZiPD was investigated by the characterization of the *E. coli* *elaC* deletion mutant;
- (c) the metal specificity of different members of the metallo beta lactamase superfamily was addressed by a comparison of the metal binding affinities [3-5] and
- (d) the function of conserved sequence pattern in the *ElaC* genes [6].

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