

Nicotinamide N-methyltransferase: from disease to 3D structure



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The enzyme nicotinamide N-methyltransferase (NNMT) catalyses the N-methylation of nicotinamide, pyridine and structurally related compounds, using S-adenosyl-L-methionine (SAM) as methyl donor. NNMT has been found to be overexpressed in several malignancies, including clear cell renal cell carcinoma (ccRCC), oral squamous cell carcinoma (OSCC), bladder urothelial carcinoma (BUC), and non-small cell lung cancer (NSCLC). Moreover, NNMT upregulation has been recently reported in metabolic syndrome and in cancer stem cell (CSC)-enriched populations obtained from human cancer cell lines.

Results obtained from studies carried out to investigate the functional significance of alterations in enzyme expression associated with disease, and the role of NNMT in cancer cell metabolism, demonstrated that NNMT knockdown led to a significant decrease of anchorage-independent cell growth and cell proliferation, both *in vitro* and *in vivo*, in several human cancer cell lines.

The crystal structure of human NNMT as a ternary complex bound to both the demethylated donor S-adenosyl-L-homocysteine (SAH) and the acceptor substrate nicotinamide has been determined. These studies revealed the structural basis for nicotinamide binding and highlight several residues in the active site which may play a fundamental role in nicotinamide recognition and NNMT catalysis. The functional importance of these residues was probed by site-directed mutagenesis and enzyme kinetics analysis.

References

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