

GENETYCZNE ASPEKTY WIĄZANIA N₂ BAKTERII Z RODZAJU *AZOSPIRILLUM*

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Wpłynęło w listopadzie 2003

1. Wstęp. 2. Metabolizm wiązania azotu. 3. Pozyskiwanie energii. 4. Podsumowanie

Genetical aspects of nitrogen fixation by bacteria *Azospirillum* species

Abstract: Biological nitrogen fixation is catalysed by the nitrogen enzyme complex, which includes dinitrogenase (MoFe protein, *nifDK* gene products), containing the active site of dinitrogen reduction, and dinitrogenase reductase (Fe protein, *nifH* gene product) supplies the reducing power to the dinitrogenase. Nitrogenase is subject to elaborate control at the trans-criptional and post-translational levels by the concentrations of intracellular nitrogen and oxygen. Nitrogenase activity is also regulated at a post-translational level by two mechanisms. One of the best understood post-transcriptional regulatory mechanisms is reversible ADP-ribosylation, which has been found in some phototrophs and the *Azospirillum* genus. The dinitrogenase reductase ADP-ribosyltransferase/dinitrogenase reductase-activating glycohydrolase (DRAT/DRAG) systems, which involves reversible nitro-genase inactivation *via* ADP ribosylation in response to micromolar concentrations of ammonia. DRAT and DRAG are encoded by the *draTG* operon, which is not co-regulated with the *nif* genes but which is found near the *nifHDK* operon in *R. rubrum*, *A. lipofrum* and *A. brasilense*. Both DRAT and DRAG activities are regulated *in vivo*, but the mechanisms for their regulation are unknown.

1. Introduction. 2. Metabolism nitrogen fixation. 3. Energy supply. 4. Summary

Słowa kluczowe: *Azospirillum*, wiązanie azotu

Key words: *Azospirillum*, nitrogen fixation