

N°729 / OC / PC

TOPIC(s) : Biocatalytic cascade reactions / Enzyme discovery and engineering

Methylformamide Conversion by Formate Dehydrogenase to Fuel Reductive Aminase with NADPH and Amine in a Cascade Setup

AUTHORS

Artur MAIER / RUHR UNIVERSITY BOCHUM, UNIVERSITAETSSTRASSE 150, BOCHUM

Tanja KNAUS / UNIVERSITY OF AMSTERDAM, POSTBUS 94157, GD AMSTERDAM

Francesco MUTTI / UNIVERSITY OF AMSTERDAM, POSTBUS 94157, GD AMSTERDAM

Dirk TISCHLER / RUHR UNIVERSITY BOCHUM, UNIVERSITAETSSTRASSE 150, BOCHUM

PURPOSE OF THE ABSTRACT

Formate dehydrogenase (FDH) from *Candida boidinii* is well studied and applied in various setups as a NADH-regeneration system. Attempts were made to increase stability and due to its clear preference for NAD⁺, to shift the substrate specificity towards NADP⁺ through mutagenesis. Here, the impact of various known stability mutations [1,2] on the NADP⁺ accepting FDH (D195Q/Y196R/Q197N) [3], in the following FDH-QRN, and the bioinformatically predicted T130I mutation on wild type and mutants are elucidated. FDH (C23S/T130I) showed a 6-fold lower K_m value ($8.5 \pm 0.32 \mu\text{M}$ vs $51.0 \pm 1.0 \mu\text{M}$) and an approx. 3.5-fold increase in catalytic efficiency ($235.7 \text{ s}^{-1} \text{ mM}^{-1}$) compared to the wild type ($66.4 \text{ s}^{-1} \text{ mM}^{-1}$). FDH-QRN (C23S/T130I) showed a 2-fold lower K_m and increase of the catalytic efficiency to $18.37 \pm 1.4 \text{ s}^{-1} \text{ mM}^{-1}$ compared to the FDH-QRN ($16.74 \pm 1.55 \text{ s}^{-1} \text{ mM}^{-1}$). FDHs are known to accept formate derivatives such as methyl-, ethyl- and phenylformate, among others [4]. In silico docking and molecular dynamics simulations suggested that formamides might be suitable substrates for FDHs as well. All variants were screened for activity towards formamide (F), N-methylformamide (MF) and N,N-dimethylformamide (DMF), showing that all variants have the ability to accept formamides, with highest activities, with NAD⁺ as cofactor, achieved by FDH (T130I) for F with 83.1 mU/mg and for MF and DMF by FDH (C23S) with 68.3 mU/mg and 7.4 mU/mg, respectively. With NADP⁺ as cofactor, the FDH-QRN reached activities with F, MF and DMF of 6.2, 14.6 and 3.5 mU/mg. The NADP⁺-accepting variants were employed for NADPH regeneration in a cascade reaction for the reductive amination of cyclohexanone by reductive aminase from *Aspergillus oryzae* [5] with MF as the sole electron and amine donor, reaching conversion rates up to 63% in a whole cell approach, broadening the applicability of FDHs in biocatalysis.

FIGURES

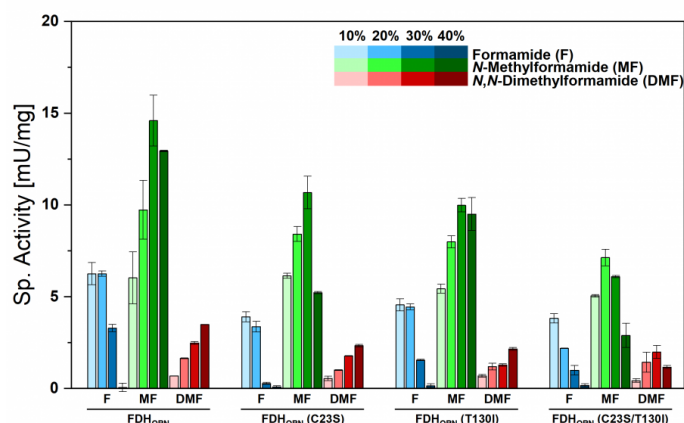


FIGURE 1

FDH Activity with formamide derivatives

Specific activity of the respective NADP⁺ accepting FDH variants with formamide (F), N-methylformamide (MF) and N,N-dimethylformamide (DMF) as the sole electron donor at concentrations ranging from 10-40% (v/v).

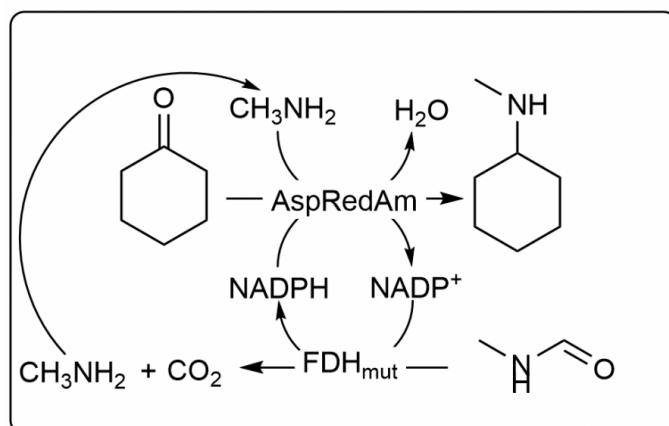


FIGURE 2

Reductive amination cascade

Scheme of the reductive amination of cyclohexanone catalyzed by AspRedAm coupled with a FDH mutant utilizing N-methylformamide for NADPH regeneration. N-methylformamide serves as the sole electron and methylamine donor.

KEYWORDS

NADPH regeneration | cascade | FDH | reductive amination

BIBLIOGRAPHY

- [1] Slusarczyk, H. et al., Eur. J. Biochem. (2000) 267, 1280-1289
- [2] Zheng, J. et al., Appl. Environ. Microbiol. (2016) 83(2):e02624-16
- [3] Wu, W. et al., J. Mol. Cat. (2009) 61, 157-161
- [4] Froehlich, P. et al., Org. Biomol. Chem. (2011) 26;9(22):7941-50
- [5] Aleku, G. et al. Nature Chem. (2017) 9, 961-969