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Bienzymatic ATP-dependent Bacterial System Fueled by Metaphosphate as Efficient Tool to Break the Chemical Stability of δ-Lactams

AUTHORS

Mélanie HALL / UNIVERSITY OF GRAZ, HEINRICHSTRASSE 28, GRAZ Sebastian ROTH / UNIVERSITY OF GRAZ, HEINRICHSTRASSE 28, GRAZ Somayyeh GANDOMKAR / UNIVERSITY OF GRAZ, HEINRICHSTRASSE 28, GRAZ Federico ROSSI / UNIVERSITY OF GRAZ, HEINRICHSTRASSE 28, GRAZ

PURPOSE OF THE ABSTRACT

Medium-sized 5- and 6-membered ring lactams are – in contrast to smaller 4-ring β -lactams – molecules with remarkable stability, a feature which appears connected to the increased resonance stabilization of the amide bond and the higher partial C-N double bond character in these lactams [1,2]. Chemical hydrolysis to the corresponding amino acids requires harsh reaction conditions (reflux and strong acid) and up to now, no enzyme active on monocyclic γ - and δ -lactams has been reported.

Our goal is to establish a biocatalytic platform for the ring opening of γ - and δ -lactams under mild reaction conditions. In this context, we explored the biocatalytic potential of bacterial ATP-dependent oxoprolinases, which function in pair (OpIA and OpIB) [3]. Strong activity in the presence of excess of ATP could be detected on δ -valerolactam and a range of derivatives. An ATP recycling system (Figure 1) based on cheap Graham's salt (sodium hexametaphosphate) and a polyphosphate kinase [4] allowed the use of catalytic amounts of ATP (1 mol%). Further improvements were obtained by co-expressing OpIA and OpIB using the pETDUET1 vector, a strategy which enhanced the soluble expression yield and the protein stability. Finally, a range of alternate phosphodonors were investigated in place of ATP. In some cases, activity was detected, providing hints on a possible mechanism, which was further studied by 31P-NMR.

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FIGURE 1

Biocatalytic metaphosphate-fueled ring-opening of delta-valerolactam derivatives by ATP-dependent-enzymes

FIGURE 2

KEYWORDS

valerolactam | ATP recycling | cascade | oxoprolinase

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