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PASTEURELLA MULTOCIDA THYMIDINE KINASE 1 EFFICIENTLY ACTIVATES PYRIMIDINE NUCLEOSIDE ANALOGS

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□ In the *Pasteurella multocida* genome only one putative deoxyribonucleoside kinase encoding gene, for thymidine kinase 1 (PmTK1), was identified. The PmTK1 gene was sub-cloned into *Escherichia coli* KY895 and it sensitized the host towards 2,2-difluoro-deoxycytidine (gemcitabine, dFdC), 3-azido-thymidine (AZT) and 5-fluoro-deoxyuridine (5F-dU). PmTK1 was over-expressed and purified with two different tags. Apparently, deoxyuridine (dU), and not thymidine (dT), is the preferred substrate. We suggest that PmTK1s could be employed as a species-specific activator of uracil-based nucleoside antibiotics.

Keywords Thymidine kinase; nucleosides; *Pasteurella multocida*; antibiotics

INTRODUCTION

Pasteurella multocida is a Gram-negative bacterium, which can cause zoonotic infections in humans. Infections can be treated with beta-lactam antibiotics, fluoroquinolones or tetracycline, however the increased use of these antibiotics has already promoted the appearance of resistance strains. The enzymes involved in the metabolism of nucleic acid precursors could represent putative targets and activators for new antibacterial drugs. Thymidine kinase 1 (TK1) catalyzes the reaction: thymidine (dT) + ATP → thymidine monophosphate + ADP, and is almost ubiquitous in all bacteria.^[1] TK1s could be employed as successful activators of nucleoside analogs.^[1,2] In this study, we identified a putative TK1 encoding gene, PmTK1, in the *P. multocida* genome. We describe here that if the PmTK1 gene is expressed in *E. coli*, the host becomes sensitive towards a few pyrimidine-based nucleoside analogs. We show that recombinant PmTK phosphorylates preferably deoxyuridine

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(dU) over dT, and that the obtained kinetic parameters do not necessarily reflects the nucleoside phosphorylating potential in the living cell.

MATERIALS AND METHODS

Cloning and Purification

The PmTK1 encoding gene, with accession number DCQ384602, has previously been cloned into the pGEX-2T expression vector from GE Healthcare (P744).^[2] PmTK1 was additionally subcloned into the TOPO pET101 vector (Invitrogen) giving the plasmid P993. P744 was over-expressed and purified without cleaving off the glutathione-S-transferase (GST) tag, as reported elsewhere.^[2] P993 was over-expressed, giving a C-terminus His-tagged enzyme, and purified as reported elsewhere.^[3] Protein purity was determined with SDS-page and the total protein concentration was determined with the Bradford method using BSA as the standard.

Minimal Inhibitor Concentration (MIC)

The in vivo tests using a kinase deficient *E. coli* strain, KY895, transformed with P744 on the nucleoside analog containing medium were done as reported previously.^[2] The nonlabeled nucleosides were from Sigma-Aldrich (St. Louis, MO, USA).

Enzyme Assay

Radioactive labeled thymidine was from GE-Healthcare, other labeled substrates were from Moravek, California. The enzyme assays have been described elsewhere.^[3]

RESULTS AND DISCUSSION

Pasteurella multocida TK1 is closely related to other TK1s from Gram-negative bacteria, such as *E. coli*.^[2] So far, Gram-negative TK1s have only been poorly studied. The PmTK1 gene has been introduced into the TK1-deficient *E. coli* KY895 and the host was screened on a variety of nucleoside analogs. The MIC values were very low on 2',2'-difluoro-deoxycytidine (gemcitabine, dFdC), 5-fluoro-deoxyuridine (5F-dU) and 3'-azido-thymidine (AZT). The host sensitivity towards AZT was at least 3.000-fold higher than the control, for both dFdC and 5F-dU it was 32-fold higher (Table 1). The observed PmTK1 activation ability was then studied on the recombinant PmTK1 carrying two different tags, a C-terminal linked 6xHis and an N-terminal linked GST. We determined the enzyme activity and kinetic parameters with dT, deoxyuridine (dU), deoxyadenosine (dA), deoxyguanosine (dG), deoxycytidine (dC), dFdC, 5F-dU, and AZT.

TABLE 1 MIC values for three nucleoside analogs. *E. coli* KY895 was transformed with the pGEX-2T vector, either without any insert or with the PmTK1 gene. Data are adapted from ref.^[2] LD100 is defined as the lowest concentration of the tested analog that completely inhibits the bacterial growth

<i>E. coli</i> transformed with	LD100(dFdC)/ μ M	LD100 (5F-dU)/ μ M	LD100(AZT)/ μ M
pGEX-2T only	31.6	0.316	>100
pGEX-2T-PmTK1 (P744)	1.0	<0.01	0.0316

PmTK1 phosphorylated, in the presence of ATP as the donor, dT, dU, 5F-dU, and AZT, but it did not accept as a substrate dA, dG or dC. Surprisingly, also dFdC was not a substrate for either the 6xHis tagged or the GST-tagged PmTK1. For PmTK-6xHis, the $K_{0.5}$ value for dU was 33 μ M, what is 5-fold lower than with dT. The V_{\max} value for dU was 0.703 U/mg and for dT it was 1.5-fold higher, giving the k_{cat}/K_m values which were 3-fold higher for dU comparing with dT (Table 2). For GST-PmTK, $K_{0.5}$ for dU was 82 μ M, what was 9-fold lower than for dT. However, the V_{\max} values were 6-fold lower for dU compared to dT, resulting in the similar k_{cat}/K_m values for both substrates. It is evident from these data that the GST-tag and 6xHis tag influenced the kinetic parameters of recombinant PmTK1. For GST-PmTK, the kinetics for dT and AZT exhibited positive cooperativity with a Hill coefficient of 1.6 and 1.8, respectively. In contrast, PmTK-6xHis followed the Michaelis-Menten kinetics with dU and dT. GST-PmTK phosphorylated AZT with $K_{0.5}$ of 430 μ M and V_{\max} of 0.814 U/mg. For 5F-dU the K_m , V_{\max} and Hill coefficient values were of 3157 μ M, 2.619 U/mg and 1.9, respectively. Apparently, both analogs were under in vitro conditions not very good substrates, and dFdC was not a substrate at all. However, under in vivo conditions all three tested analogs were successfully phosphorylated (Table 1).

PmTK1 is so far the only TK1, which prefers dU over dT, but the K_m values are high (Table 2). So far, only one more Gram-negative TK1 (SeTK1), from *Salmonella enterica*, has had their kinetic parameters determined. SeTK1

TABLE 2 Kinetic data for recombinant PmTK1 tagged with either a C-terminal 6xHis or an N-terminal GST-tag. Both recombinant TK1s had lower K_m values for dU than for dT. Both tags may have impaired the kinetic parameters. U: μ mol min⁻¹

	Substrate	$K_{0.5}/\mu$ M	$V_{\max}/$ U/mg	Hill coefficient	$k_{\text{cat}}/$ s ⁻¹	$k_{\text{cat}}/K_m/$ s ⁻¹ ·M ⁻¹ ·10 ³
PmTK-6xHis	dU	33 \pm 9	0.703 \pm 185	1	0.30	9.1
PmTK-6xHis	dT	164 \pm 40	1.074 \pm 158	1	0.46	2.8
GST-PmTK	dU	82 \pm 22	0.179 \pm 17	1	0.14	1.7
GST-PmTK	dT	717 \pm 62	1.122 \pm 1	1.8 \pm 0.4	0.90	1.3
GST-PmTK	AZT	430 \pm 28	0.814 \pm 44	1.6 \pm 0.2	0.66	1.5
GST-PmTK	5F-dU	2298 \pm 508	2.619 \pm 684	1.9 \pm 0.1	1.12	0.5

phosphorylates dT with K_m of 34 μM . Two Gram-positive TKIs, BcTK1 (from *Bacillus cereus*),^[2] BaTK1 (from *Bacillus anthracis*),^[4] and human TK1^[5] exhibit low K_m values for dT, 0.5–0.6 μM , while AZT K_m values are of 0.6–4 μM . SeTK1 is, like PmTK1, not very efficient with AZT having K_m of 73 μM . However, PmTK1 and SeTK1 efficiently sensitize *E. coli* KY895 giving MIC values for AZT of only 0.01–0.1 μM ^[2]. Apparently, Gram-negative TKIs could behave in a different and much more efficient way under *in vivo* conditions. Therefore, the *in vitro* kinetic parameters obtained with recombinant Gram-negative TKIs only poorly reflect the capacity of these TKIs in the living bacterial cell. In addition, our studies demonstrate that nucleoside analogs could be used as species-specific antibiotics.

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