

Potent inhibition of human folylpolyglutamate synthetase by a phosphinic acid mimic of the tetrahedral reaction intermediate

John J. McGuire^{a,*}, William H. Haile^a, Nadya Valiaeva^b, David Bartley^b,
 Jianxia Guo^a, James K. Coward^b

^aGrace Cancer Drug Center, Roswell Park Cancer Institute, Elm and Carlton Streets, Buffalo, NY 14263, USA

^bDepartments of Medicinal Chemistry and Chemistry, University of Michigan, Ann Arbor, MI 48109, USA

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Abstract

A phosphorous-containing pseudopeptide folate analog (Valiaeva *et al.*, *J Org Chem* 2001;66:5146–54) was designed to mimic the tetrahedral intermediate formed in the ATP-dependent reaction catalyzed by folylpolyglutamate synthetase (FPGS). This analog, methotrexate-phosphinate (MTX-phosphinate; 4-amino-4-deoxy-10-methylpteroyl-L-Glu- γ -[Ψ {P(O)(OH)-CH₂}]-glutarate), is a highly potent ($K_{i,s}$, 3.1 ± 0.5 nM), competitive inhibitor of recombinant human cytosolic FPGS. Within experimental limits, FPGS inhibition was not time-dependent, and preincubation of FPGS, inhibitor, and ATP did not potentiate the inhibition. These results suggest that slow phosphorylation to produce a more potent inhibitor form is not involved. MTX-phosphinate was not growth inhibitory to human CCRF-CEM leukemia cells at 1 μ M (70-fold above the concentration of MTX giving 50% growth inhibition), probably because of poor transport. Because of its exceedingly high potency as an FPGS inhibitor, MTX-phosphinate represents a lead structure from which cell-permeable analogs may be developed to test the hypothesis that FPGS inhibition is therapeutically efficacious.

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1. Introduction

FPGS, the enzyme that synthesizes folate and antifolate poly(γ -glutamate) metabolites (reaction 1), is a target for antifolate drug design.



Cellular folate pools occur exclusively as reduced folylpoly(γ -glutamates) ($\text{H}_4\text{PteGlu}_n$) [2]. $\text{H}_4\text{PteGlu}_n$ primarily function to: (a) enhance retention of intracellular folates; and (b) act as kinetically preferred intracellular 1-carbon carriers. $\text{H}_4\text{PteGlu}_n$ are essential since mutational inactivation of FPGS induces auxotrophy for glycine, a purine, thymidine, and methionine [2].

FPGS-specific inhibition should induce the FPGS[−] phenotype and be lethal. Importantly for cancer chemotherapy, FPGS inhibition may be tumor selective [reviewed in Ref. [3]] since: (a) dietary folate deficiency selectively decreases tumor growth; and (b) tumor and normal cells may express different FPGS isozymes. FPGS inhibition might be useful in overcoming one clinical antifolate resistance mechanism, decreased FPGS activity, since such tumors are collaterally sensitive to FPGS inhibition [4].

Several groups (e.g. [5–7]) have attempted to develop folate analog FPGS inhibitors. Modification of the Glu moiety may result in FPGS inhibition, whereas changes in

* Corresponding author. Tel.: +1-716-845-8249; fax: +1-716-845-8857.

E-mail address: John.McGuire@RoswellPark.edu (J.J. McGuire).

Abbreviations: AMT, aminopterin (4-amino-4-deoxy-pteroylglutamic acid); DHFR, dihydrofolate reductase (5,6,7,8-tetrahydrofolate:NADP⁺ oxidoreductase, EC 1.5.1.3); EC₅₀, drug concentration effective in inhibiting growth of cells by 50% relative to solvent-treated control cultures; FPGS, folylpolyglutamate synthetase (tetrahydrofolate:L-glutamate γ -ligase (ADP-forming), EC 6.3.2.17); rhcFPGS, recombinant human cytosolic FPGS; Glu, glutamic acid; $\text{H}_4\text{PteGlu}_n$, tetrahydrofolate poly(γ -glutamates) containing n total glutamate residues; IC₅₀, drug concentration effective in inhibiting enzyme activity by 50% relative to a solvent-containing control; MOI, multiplicity of infection; MTX, methotrexate (4-amino-4-deoxy-10-methylpteroylglutamic acid); MTX-phosphinate, 4-amino-4-deoxy-10-methylpteroyl-L-Glu- γ -[Ψ {P(O)(OH)-CH₂}]-glutarate; MTX-phosphonate, 4-amino-4-deoxy-10-methyl-pteroyl-L-Glu- γ -[Ψ {P(O)(OH)-O}]-glutarate; Orn, ornithine; and the use of Ψ as a symbol for peptide surrogates is as described by the IUPAC-IUB Joint Commission on Biochemical Nomenclature [1].

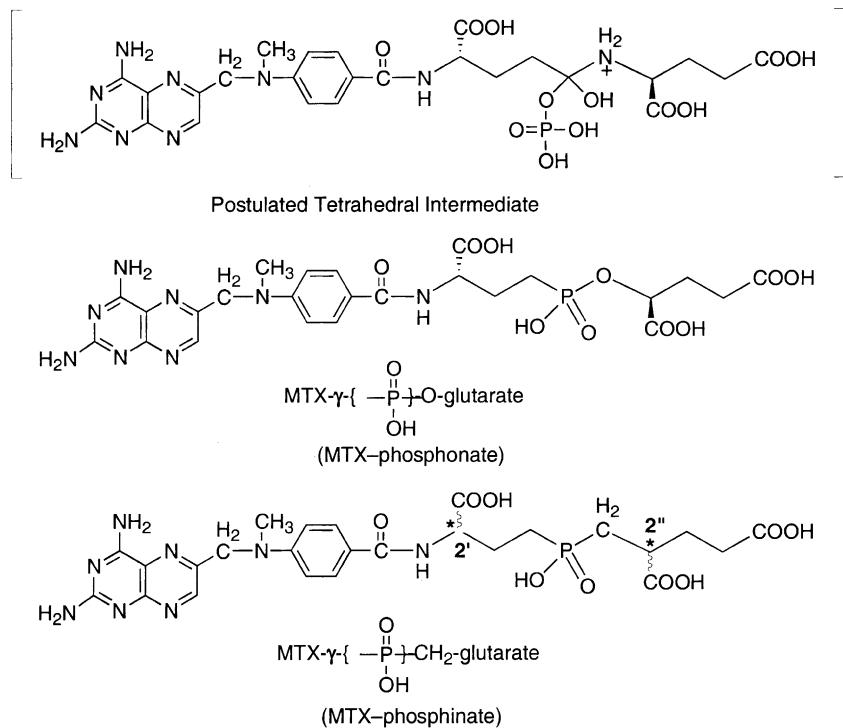


Fig. 1. Structures of the tetrahedral intermediate of the FPGS reaction and of MTX-phosphonate and MTX-phosphinate. Note that there are two chiral centers (*) at the 2' and 2'' positions in MTX-phosphinate, and the synthetic method [13] yielded all four diastereomers.

the pteroyl moiety lead to increased inhibitory potency and/or specificity [3]. Shane and colleagues showed that substitution of Orn for Glu in PteGlu (i.e. PteOrn) leads to potent FPGS inhibition [6], but Orn substitution leads to poor cellular uptake because of the positively charged δ -amino group [8].

Another approach is the design of mechanism-based FPGS inhibitors. FPGS-catalyzed ligation of at least the first two γ -Glu residues proceeds through acyl-phosphate intermediates at the γ -COOH of the polyglutamate chain [9]. Attack by the amino group of the incoming Glu leads to transient formation of a tetrahedral intermediate at the γ -carbon (Fig. 1); such intermediates may be mimicked by stable phosphorous-containing structures (phosphonamides, phosphonates, and phosphinates; discussed in Ref. [10]). Since the stability of phosphonamides is a concern [11], our synthetic effort has focussed on MTX-phosphonate (Fig. 1), a potent competitive inhibitor of human FPGS (K_i , 46 nM) [12], and on MTX-phosphinate. MTX-phosphinate has been synthesized as a mixture of two racemic diastereomers (Fig. 1) [13]; its characterization as an inhibitor of human FPGS is presented below.

2. Materials and methods

2.1. Materials

Common chemicals were reagent grade or higher. MTX was a gift of Immunex. AMT was from the Sigma Chemical

Co. MTX-phosphinate was synthesized as described previously [13]. Literature extinction coefficients [14] were used (that of MTX-phosphinate was assumed to be the same as that of MTX).

2.2. Enzymes and assays

A BaculoGoldTM Transfection Kit (Pharmingen) was used to co-transfect *Sf9* insect cells with pVL1392/cFPGS shuttle vector DNA (a gift of Dr. B. Shane), and BaculoGoldTM DNA according to the instructions of the manufacturer to yield recombinant rhcFPGS-encoding baculovirus. After titering, recombinant virus was used to infect (MOI = 4) *Sf9* monolayers, and cells were harvested at 72 hr. FPGS activity was extracted [15] and purified by $(\text{NH}_4)_2\text{SO}_4$ fractionation and BioGel A-0.5M chromatography (specific activity 6.4×10^6 pmol [^3H]Glu/hr/mg protein). K_m values for AMT and MTX as substrates are the same for this rhcFPGS as those reported for CCRF-CEM FPGS [12]. Inhibition constants K_{ii} and K_{is} [16] for rhcFPGS were determined [12] using MTX as the variable substrate and 2 and 8 nM MTX-phosphinate, levels yielding about 25 and 50% inhibition, respectively. Activity of and inhibitory potency (IC_{50}) against partially purified CCRF-CEM DHFR were assayed as described previously [15].

2.3. Cell culture

The human T-lymphoblastic leukemia cell line CCRF-CEM and its MTX transport-deficient subline R2, their

culture, and inhibition of growth under continuous (120-hr) drug exposure have been described previously [17]. Cell lines were negative for Mycoplasma (Mycoplasma Plus PCR primers).

3. Results and discussion

3.1. Inhibition of *rhcFPGS*

rhcFPGS did not use MTX-phosphinate as a substrate at 1 μ M, the highest level tested (data not shown). When measured at a fixed pteroyl substrate (MTX) and enzyme concentration, MTX-phosphinate (IC_{50} , 8 \pm 1 nM; average \pm range, N = 2) was an even more potent FPGS inhibitor than MTX-phosphonate (IC_{50} , 120 nM) [12]. Kinetic constants for *rhcFPGS* inhibition using MTX as the variable substrate were: K_{is} = 3.1 \pm 0.5 nM and K_{ii} = 56 \pm 5 nM (average \pm range; N = 2). Since $K_{is} \ll K_{ii}$, inhibition was essentially competitive; this is consistent with the Lineweaver-Burk replot of the kinetic data (not shown). Thus, MTX-phosphinate is the most potent FPGS inhibitor based on the MTX heterocycle that has been described to date. This high potency is even more impressive considering that binding of an antifolate FPGS inhibitor containing a Glu substitution (e.g. Orn) generally tracks the efficiency of the Glu-containing homolog to serve as an FPGS substrate [18], and MTX and MTX- γ -Glu are poor FPGS substrates [12]. If this relationship holds in the phosphinate series, it should be possible to substantially increase potency by replacing the 4-amino-4-deoxy-10-methylpteroyl heterocycle with one that, as a Glu-containing substrate, has a low K_m (e.g. DDATHF, K_m ca. 1 μ M for human FPGS [19]).

Phosphinate inhibitors of ATP-dependent reactions may exhibit slow formation of a phosphorylated inhibitor that has much tighter binding (e.g. [20]). However, FPGS inhibition does not appear to be time-dependent, since at fixed levels of MTX-phosphinate (3 or 8 nM), FPGS activity was linear with respect to time for up to 2 hr. Also experiments in which MTX-phosphinate (10 or 20 nM) bound to FPGS was preincubated with ATP and L-Glu in the absence of pteroyl substrate (i.e. MTX) for 30 min showed no increase in inhibition relative to a control in which inhibitor and MTX were added simultaneously. However, the time period over which these reactions were observed (1–2 hr) might preclude kinetic observation of a phosphorylated inhibitor if it is synthesized over a few minutes [20].

MTX-phosphinate was synthesized as a diastereomeric mixture [13] to test the hypothesis that the phosphinate substructure imparts potent FPGS inhibition. Based on the known stereospecificity of FPGS [2], it is likely that only the 2'S, 2''S MTX-phosphinate diastereomer (Fig. 1) is highly active, and thus the actual K_i of the inhibitory species may be as low as 0.75 nM. A stereospecific synth-

esis of the 2'S, 2''S diastereomer is being pursued to explore this possibility.

3.2. Inhibition of human DHFR

Inhibitory potency against human DHFR (CCRF-CEM cells) was measured. MTX-phosphinate was slightly less potent than MTX (IC_{50} , 2.1 \pm 0.1 vs 0.6 \pm 0.1 nM, respectively; average \pm range, N = 2) as a human DHFR inhibitor. The slightly lower potency of MTX-phosphinate may result from the presence of isomers (above) since the analog of MTX in which D-Glu replaces L-Glu is a much less potent DHFR inhibitor [21].

3.3. Cell growth inhibition

MTX-phosphinate and MTX were compared as growth inhibitors of CCRF-CEM human leukemia cells and a MTX transport-deficient subline, R2. MTX-phosphinate (cell growth 96.5 \pm 0.5% of control at 1000 nM; average \pm range; N = 2) was \geq 70-fold less growth inhibitory to CCRF-CEM cells than was MTX (EC_{50} , 14.5 \pm 0.5 nM; average \pm range; N = 2). The subline R2 (EC_{50} , 2550 \pm 50 nM; average \pm range; N = 2) was 175-fold resistant to MTX and also did not respond to 1000 nM MTX-phosphinate (cell growth 104 \pm 2% of control; average \pm range; N = 2), the highest level tested. Despite the fact that MTX is a relatively poor (high K_m) mammalian FPGS substrate (e.g. [12]), we employed the 4-amino-4-deoxy-10-methyl heterocycle because it allows a simple way to separate uptake from FPGS inhibition [12]. If an analog containing this heterocycle is taken up, the potent inhibition of DHFR (above) will allow immediate and potent growth inhibition (FPGS inhibition causes a delayed effect); growth inhibition is thus a surrogate for uptake. Since MTX-phosphinate is not growth inhibitory, it is not taken up efficiently into cells. This is not surprising because of its negative charge and dipeptide structure [12]. Since we have shown that the phosphinate substructure imparts more potent FPGS inhibition than the phosphonate, our efforts will now turn to designing phosphapeptide prodrugs that are active in whole cells.

A potential complication to the development of therapeutically effective FPGS inhibitors has been noted. Human FPGS exists in cytosolic and mitochondrial isoforms [22–24]. It is unknown whether both FPGS isoforms must be inhibited to induce cell death. Inhibition of mitochondrial FPGS could be difficult because mitochondria are impermeable to many antifolates (e.g. [25]). However, since purine and thymidylate syntheses occur in the cytosol, cytosolic FPGS inhibition alone may be sufficient to inhibit these two processes that are critical for DNA synthesis and repair. The FPGS inhibitors now under development should allow us to answer this important question.

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