oxy]acetanilide, 103789-06-6; 3-(acetylamino)-2,4-pentanedione, 5440-23-3; 4-acetyl-2,5-dimethoxazole, 23000-12-6; 3-(4-nitrophenoxy)propionic acid, 10572-16-4; α -aminopropiophenone hydrochloride, 16735-19-6; α -[[3-(4-nitrophenoxy)propionyl]-amino]propiophenone, 141399-55-5; 4-[2-(4-methyl-5-phenyl-2-oxazolyl)ethoxy]nitrobenzene, 107325-00-8; 4-fluorobenzonitrile, 1194-02-1; 4-[2-(5-methyl-2-phenyl-4-oxazolyl)ethoxy]benzonitrile, 103789-44-2; 4-hydroxybenzaldehyde, 123-08-0; 4-(2-phenyl-4-

thiazolylmethoxy)benzaldehyde, 103789-67-9; β -methyl L-aspartate hydrochloride, 16856-13-6; β -methyl N-benzoyl-L-aspartate, 39741-26-9; methyl 3-(benzoxylamino)-4-oxovalerate, 54819-26-0; methyl 2-(5-methyl-2-phenyl-4-oxazolyl)acetate, 103788-64-3; ethyl 4-chloroacetoacetate, 638-07-3; benzamide, 55-21-0; ethyl 2-(2-phenyl-4-oxazolyl)acetate, 84446-03-7; ethyl 2-(2-cyclohexyl-4-thiazolyl)acetate, 24087-96-5; cyclohexanethiocarboxamide, 7390-42-3.

N-[4-[[(3,4-Dihydro-4-oxo-1,2,3-benzotriazin-6-yl)methyl]amino]benzoyl]-L-glutamic Acid, a Novel A-Ring Analogue of 2-Desamino-5,8-dideazafolic Acid¹

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N-[4-[[(3,4-Dihydro-4-oxo-1,2,3-benzotriazin-6-yl)methyl]amino]benzoyl]-L-glutamic acid ("2-aza-2-desamino-5,8-dideazafolic acid", ADDF) was synthesized from 2-amino-5-methylbenzamide via a four-step sequence consisting of diazotization, benzylic bromination, condensation with dimethyl N-(4-aminobenzoyl)-L-glutamate, and ester hydrolysis. ADDF was an inhibitor of recombinant mouse thymidylate synthase; inhibition was competitive with 5,10-methylenetetrahydrofolate as variable substrate ($K_{\rm i}=2.3~\mu{\rm M}$). It was a substrate for murine follypolyglutamate synthetase with kinetic characteristics ($K_{\rm m}=28~\mu{\rm M}$) comparable to those of aminopterin, and it inhibited the growth of L1210 cells in culture (IC₅₀ = 0.52 $\mu{\rm M}$). The structural modification of the A-ring embodied in ADDF appears to offer a novel, heretofore unexplored approach to the design of TS inhibitors.

The potential therapeutic significance of folic acid analogues targeted against thymidylate synthase (TS) as opposed to dihydrofolate reductase (DHFR) was predicted more than 20 years ago by Borsa and Whitmore.² Shortly thereafter, Bird and co-workers3 reported the potent biological activity of 5,8-dideazafolic acid (1).4 This led to an extensive program of synthesis of quinazoline analogues substituted at the N^{10} position,⁵⁻⁷ on the phenyl ring,⁸ and in ring B,^{9,10} and to selection of N^{10} -propargyl-5,8-dideazafolic acid (CB3717) as a suitable candidate for biochemical¹¹⁻¹³ and clinical^{14,15} evaluation. While CB3717 had many desirable pharmacological properties, such as the ability to enter cells by a transport mechanism distinct from that of reduced folates and the classical DHFR inhibitor methotrexate (MTX), its clinical usefulness was hampered by low solubility at physiological pH, which gave rise to hepatic and renal toxicity.16

COOH

$$HN$$
 R
 CH_2NH
 CH_2NH
 $COOH$
 $COOH$

The toxicity encountered during clinical trials with CB3717 prompted a vigorous search for more soluble congeners, culminating in the discovery of a family of second-generation TS inhibitors that included 2-desamino-5,8-dideazafolic acid (2), 2-desamino-2-methyl-5,8-dideazafolic acid (3), and the corresponding N^{10} -methyl and N^{10} -propargyl analogues. Piochemical studies n^{20-22} revealed that, in general, replacement of the NH₂ group

at C² by H or Me resulted in weaker binding to purified TS but increased inhibition of the growth of cultured cells.

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This higher than expected biological potency was rationalized on the basis that (i) substrate activity for folylpolyglutamate synthetase (FPGS) is minimally affected by replacement of the 2-NH₂ group with H or Me, (ii) polyglutamylation increases TS binding by as much as 100-fold, (iii) in contrast to CB3717, 2-desamino and 2desamino-2-methyl analogues are accepted by the active transport mechanism used by MTX and reduced folates, and (iv) noneffluxing long-chain polyglutamates of these compounds are formed efficiently inside the cell. A large number of analogues of 1-3 modified at the C² position, in the benzoyl moiety, 23,24 and in the C9-C10 bridge re-

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Scheme I^a

$$HOOC \longrightarrow CH_3$$

$$X = NH_2$$

$$(iii) \longrightarrow G: X = NO_2$$

$$(iiii) \longrightarrow 7: X = NH_2$$

$$(iv) \longrightarrow N$$

$$X = Br$$

$$(v) \longrightarrow B: X = H$$

$$13,14: X = Br, OAc$$

$$COOR$$

$$CH_2NH \longrightarrow CONHCH$$

$$(CH_2)_2$$

$$COOR$$

$$(vii) \longrightarrow 15: R = Me$$

$$4 (ADDF): R = H$$

^a Reagents: (i) SOCl₂; (ii) NH₃; (iii) H₂/Pd-C; (iv), NaNO₂/HCl; (v) N-bromosuccinimide/AcOH/hv; (vi) 4-H₂NC₆H₄CONHCH-(COOMe)CH2CH2COOMe; (vii) NaOH.

gion^{25,26} were synthesized to permit detailed structure activity analysis. Out of these studies emerged the new TS inhibitor ICI D1694, 27 an exciting third-generation analogue with Me groups at C^2 and N^{10} , and a thiophene ring as the aryl moiety.28

The absence of a 2-NH₂ group in second- and thirdgeneration analogues of 1 is significant in terms of their ability to interact with TS. According to a recent X-ray crystallographic study of the ternary complex of CB3717 and 2'-deoxyuridylate (dUMP) with TS from Escherichia coli,29 the quinazoline ligand can bind to two different sites in the same domain of the enzyme, and when bound can have two different orientations vis-à-vis the nucleotide ligand. In one orientation, involving the "major" site, the 2-NH₂ group forms an H-bond to the C=O of the penultimate C-terminal residue (position 263, E. coli numbering) and to an ordered water molecule which is also H-bonded to N¹; in the other orientation, involving the

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"minor" site, the 2-NH2 group forms an H-bond to a serine OH (Ser180), and N¹ forms an H-bond to a glutamate COOH (Glu58) which is also H-bonded through an ordered water molecule to Ser180. According to another X-ray crystallographic analysis,30 this one involving a ternary complex between E. coli TS and 5-fluoro-2'-deoxyuridylate (FdUMP) instead of dUMP, the 2-NH2 group of the quinazoline forms an H-bond to the amide C=O of a residue (Ala263) in the C-terminal region, and N¹ is bound to an ordered water molecule (Wat430) that also interacts with the C=O group of Ala263 and with a highly-conserved arginine residue (Arg21) in the phosphate binding site. The active-site residues with which the 2-NH2 group of CB3717 interacts in mammalian TS have not yet been described, and may not be exactly the same as those in the E. coli enzyme. Moreover, the nature of these interactions may depend on the nucleotide ligand as well as the N¹⁰ substituent in the quinazoline. However, the evidence reported to date suggests that it is reasonable to suppose that the 2-NH₂ group binds to TS from all species via a network of H-bonds that also involves molecules of ordered water, and that these interactions are probably important not only in correctly aligning the folate cofactor and nucleotide substrate in the active site but also in sequestering them from bulk solvent. When the 2-NH₂ group is replaced by H or Me, binding to the enzyme is diminished because the only H-bonds possible between the quinazoline A-ring and the active site are those involving N¹ as an H-bond acceptor and N³ as an H-bond donor. The reason the desamino compounds actively inhibit TS in intact cells despite their weaker binding to purified enzyme appears to be that efficient uptake and polyglutamylation compensate for the partial loss of binding caused by removal of the 2-NH₂ group.

In considering possible approaches to improving the solubility of quinazoline inhibitors of TS while preserving their ability to form polyglutamates, we became interested in 2-desamino-4(3H)-oxoquinazoline analogues in which C^2 is replaced by a nitrogen atom with its lone pair of electrons. Attractive features of such 2-aza-2-desamino compounds were (i) that removal of H or Me might enable the molecule to extend more deeply into the active-site pocket, and (ii) that N² might interact with active-site residues by H-bonding, acting as an acceptor either directly or via a molecule of ordered water. Both of these properties ought to favor TS binding. Furthermore, in vivo activity might be anticipated if these compounds are good FPGS substrates and if the polyglutamates show the same increase (100-fold or more) in TS binding as is observed in 2-desamino-5,8-dideazafolates. In this report we describe the synthesis of N-[4-[(3,4-dihydro-4-oxo-1,2,3-dihydro-4benzotriazin-6-yl)methyllamino|benzoyl]-L-glutamic acid (4, "2-aza-2-desamino-5,8-dideazafolic acid", ADDF), which we believe is the first known example of a 2-azafolate analogue. We also present data showing that ADDF binds to TS, is a FPGS substrate, and inhibits the growth of tumor cells in culture.

Chemistry

As shown in Scheme I, sequential reactions of 2-nitro-5-methylbenzoic acid (5) with $SOCl_2$ and NH_3 afforded 2-nitro-5-methylbenzamide (6, 87%), which on catalytic hydrogenation yielded 2-amino-5-methylbenzamide (7, 96%). Treatment of 7 with $NaNO_2/HCl$ at low tem-

perature³² then gave 6-methyl-1,2,3-benzotriazin-4(3H)-one (8, 97%). Confirmation of ring closure to a benzotriazinone came from the ¹H-NMR spectrum, in which all three aromatic proton signals were markedly shifted downfield (δ 7.8–8.0), in agreement with the powerful electronwithdrawing character of the triazinone moiety. Several attempts were made to enhance the solubility of 8 by N³-acylation, with the expectation that this would be needed if subsequent benzylic bromination were done in CCl4, the traditional solvent for such reactions. Treatment of the Na salt of 8 with (pivaloyloxy)methyl chloride in DMF afforded a single product which appeared to be the desired N³-derivative 9. However, in contrast to the facile preparation of the analogous blocked compound in the quinazoline series, 17 all efforts to recrystallize 9 led to loss of the blocking group. We also tried to prepare the N3acetyl derivative 10 by reaction of the Ag salt of 8 with acetyl chloride, as has been described for the analogue without a 6-methyl substituent.³³ To our surprise, the reaction of 8 yielded a product whose ¹H-NMR spectrum contained two Me groups, but whose UV spectrum differed from that expected on the basis of the literature.33 Moreover, efforts to purify the product by silica gel chromatography led only to deacylation. Although definitive proof of identity was not obtained, this material was tentatively assigned the O-acetyl structure 11. Finally we tried to prepare 10 from the Na salt of 8 and Ac₂O in DMF, but no reaction occurred at room temperature, and the only product identified after heating was the benzoxazinone 12.34

Bromination of the benzotriazinone 8 without protection at N³ proved feasible through the use of hot AcOH as the solvent. The presence of the desired product 13 was evident in the ¹H-NMR spectrum, which showed a downfield singlet at δ 4.63 as compared with δ 3.30 for the Me group in 8. Also present were two other singlets, which we believe correspond to 14, presumably formed by solvolysis of 13. Heating the crude bromination product (estimated from the ¹H-NMR spectrum to contain roughly 40 mol % of 13) directly with dimethyl N-(4-aminobenzovl)-L-glutamate and NaHCO3 in DMF for 3 days afforded the protected diester 15 (45% crude yield), and further treatment of 15 for a few minutes with NaOH in aqueous MeOH afforded the diacid 4 (ADDF, 81%). The UV absorption spectrum of 4 [λ_{max} (0.1 M HCl) 203, 224, 290 nm; λ_{max} (pH 7.4) 293 nm; λ_{max} (0.1 M NaOH) 298 nm] showed a small bathochromic shift in going from acid to neutral to alkaline pH. IR spectra of both 4 and 15 measured in KBr disks showed strong absorption at 1690 cm⁻¹, indicating that the lactam tautomer 4A may be present in the solid state. The ¹H-NMR was consistent with the benzotriazinone structure, with all three aromatic B-ring protons deshielded (δ 8.10) relative to the 3',5'-protons (δ 6.60) and even the 2',6'-

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protons (δ 7.57) on the phenyl ring. However, a more lyglutamate metabolites that efflux from the cell at a slow to nonexistent rate. Nevertheless, it seems reasonable to conclude that ADDF is likely to form noneffluxing polyglutamates, and that the long-chain derivatives of ADDF would be likely to have K_i values in the low nanomolar Continuous exposure of mouse L1210 cells to ADDF for 48 h resulted in complete inhibition of growth at high concentrations and 50% inhibition of growth at a concentration of $0.52 \pm 0.08 \,\mu\text{M}$ (n = 3). Thus, the potency of ADDF was greater than that reported for 1 (2.7 μ M) and

interesting feature of the ¹H-NMR spectra of diester 15 and diacid 4 was the presence of upfield signals, at δ 1.7 for 15 (in CDCl₃ solution) and at δ 2.1 for 4 (in d_8 -DMSO solution), which we believe arise from the lactim tautomeric form 4B. Since the peak areas for these upfield signals in 4 and 15 closely approximated one proton, it appears that very little of the lactam 4A was present under aprotic conditions. The position of the lactam-lactim equilibrium presumably depends not only on the dielectric constant of the solvent, but also on its protic versus aprotic nature. Whether 4 is in the lactam (4A) or lactim (4B) tautomeric form when it binds to the TS active site will obviously be worthwhile to examine, since it could be the basis for a fundamental mechanistic difference between 2desamino-5,8-dideazafolates and 2-aza-2-desamino-5,8dideazafolates.

Biological Activity

ADDF was studied as an inhibitor of mouse TS, as an inhibitor of the growth of L1210 cells, and as a substrate for mouse liver FPGS. Affinity column chromatography on 10-formyl-5,8-dideazafolyl-Sepharose was used to purify recombinant mouse fibroblast TS expressed in E. coli.34 Using a spectrophotometric assay, this enzyme had a $K_{\rm m}$ of 11.3 \pm 2.7 μ M (n = 4) for (6R)-5,10-methylenetetrahydrofolate in excellent agreement with the $K_{\rm m}$ of 22–27 μ M measured using (6R,S) cofactor with either recombinant or naturally occurring enzyme from LU-37 mouse fibroblasts.34 ADDF was an inhibitor of this recombinant mouse TS with competitive kinetics against variable cofactor (Figure 1 contained in supplementary material); the K_i for this interaction was $2.3 \pm 0.3 \mu M$ (n = 3), as compared with published values of 0.067 μ M for 1^{27} and 2.0 μ M for 2.20 Thus, replacement of C² and the attached NH₂ group in 1 by a nitrogen atom with a lone pair of electrons had a similar effect on binding to TS as replacement of the 2-NH2 group by hydrogen. This lends support to our hypothesis that the structural changes embodied in 2aza-2-desamino antifolates such as ADDF would be a new approach to inhibitors of this enzyme.

The polyglutamate derivatives of CB3717, the 2desamino analogue of CB3717, and ICI D1694 have been shown to be at least 100-fold more potent inhibitors of TS than the monoglutamate forms of these compounds.^{27,28} Because of this clear indication that the potential for polyglutamylation is a major factor in the design of folate analogues inhibitors to TS, the ability of ADDF to serve as a substrate for mouse liver FPGS was studied in vitro. ADDF was a substrate for this enzyme with a $K_{\rm m}$ of 28 \pm 4.3 μ M and a $V_{\rm max}$ 1.6 \pm 0.1 times that of folic acid (n=4). These kinetic values are equivalent to those previously observed for aminopterin with this same source of enzyme. and would rank ADDF as a better substrate than CB3717 $(K_{\rm m}$ = 40 μ M) but a less efficiently used substrate than either 1 ($K_{\rm m} = 6.4 \,\mu{\rm M}$) or 2 ($K_{\rm m} = 4.8 \,\mu{\rm M}$). The fact that the first-order rate constant for the reaction with ADDF as the substrate $(V_{\text{max}}/K_{\text{m}} = 6.3 \pm 0.1 \text{ relative to folic acid})$ was ca. 5-fold less than has been reported for 1 and 2 suggested that the 2-aza-2-desamino compound may be utilized somewhat less efficiently than either the 2-amino or 2-desamino analogues. It should be noted that this in vitro assay measures the conversion of a folyl monoglutamate to a diglutamate, and not to tri- or higher posimilar to that of 2 $(0.43 \mu M)$.²⁰ An interesting feature of the N¹⁰-unsubstituted quinazolines 1 and 2 that sets them apart from pure inhibitors of TS is that their inhibitory effect on the growth of L1210 cells is restored to normal levels by a combination of thymidine (dThd) and hypoxanthine (Hx). This observation has been explained on the basis that these quinazolines have a second site of action in the folate pathway, resulting in inhibition of both thymidylate synthesis and purine synthesis. This second site has been identified as DHFR.20,27 Similar cell culture reversal experiments demonstrated a very interesting pattern with ADDF: when L1210 cells were growth in the presence of 5.6 μ M dThd, the IC₅₀ of ADDF increased to $0.89 \pm 0.09 \mu M$ (n = 3)whereas, in the presence of 32 μ M Hx, the IC₅₀ of ADDF was $0.97 \pm 0.18 \,\mu\text{M}$. On the other hand, when both dThd and Hx were present, ADDF was not inhibitory at concentrations as high as 30 μ M. Hence, ADDF appeared to inhibit at least two folate-dependent reactions, one of which was TS. The potential for ADDF inhibition of glycinamide ribonucleotide transferase (GARFT) was directly assessed with enzyme purified from mouse L1210 cells: no inhibition was observed at concentrations as high as 30 μ M.

In summary, the activity of ADDF against tumor cells. its ability to bind to TS and FPGS, its novel molecular structure, and its relatively easy synthesis all suggest that this compound is a promising lead to the design of cytotoxic antifolates. Manipulations of the structure of ADDF that would result in increased FPGS substrate activity seem particularly interesting at this time.

Experimental Section

IR spectra were obtained on a Perkin-Elmer Model 781 double-beam recording spectrophotometer; only peaks above 1200 cm⁻¹ are reported. UV spectra were obtained on a Varian Model 210 instrument. ¹H-NMR spectra were obtained on a Varian EM360L spectrometer with Me₄Si or Me₃Si(CH₂)₃SO₃Na as the reference. TLC analyses were done on fluorescent Eastman 13181 silica gel sheets or Eastman 13254 cellulose sheets. Spots were visualized under 254-nm UV illumination. Column chromatography was done on Baker 3405 (60-200 mesh) silica gel or Whatman DE-52 preswollen DEAE-cellulose. Solvents in moisture-sensitive reactions were dried over Linde 4A molecular sieves (Fisher, Boston, MA). HPLC was done on Waters C₁₈ radial compression cartridges (analytical: 5 μ m particle size, 5 × 100 mm; preparative: 15 μ m particle size, 25 × 100 mm). Melting points were determined in Pyrex capillary tubes in a Mel-Temp apparatus (Cambridge Laboratory Devices, Cambridge, MA) and are not corrected. Microanalyses were performed by Quantitative Technologies, Inc., Bound Brook, NJ.

6-Methyl-1,2,3-benzotriazin-4(3H)-one (8). A mixture of 5-methyl-2-nitrobenzoic acid (5) (36.2 g, 0.2 mol) and SOCl₂ (50 mL) was heated under reflux for 20 min, during which a homogeneous solution was obtained. After removal of the excess SOCl2 with the aid of a water aspirator, the residue was dissolved in dry THF (40 mL) and the solution was added dropwise with stirring to an ice-cold solution of NaOH (8 g, 0.2 mol) in 28% NH₄OH (300 mL). The precipitate was collected, washed with H₂O, and recrystallized from EtOH-H₂O to obtain 5-methyl-2-nitrobenzamide (6) (31.5 g, 87%) as a white solid: mp 176-178 °C (lit.31

mp 176-177 °C); IR (KBr) ν 1655 cm⁻¹ (amide C=O). A solution of 6 (31.5 g, 0.175 mol) in MeOH (250 mL) was shaken with 5% Pd-C (0.5 g) under 3 atm of H₂ for 24 h in a Parr apparatus. A solid formed after the initial heat of reaction subsided. The mixture was heated to boiling to redissolve the product, and it was filtered while hot. The filtrate was evaporated under reduced pressure, and the residue was dried at 90 °C for 1 h (caution: some sublimation may occur) to obtain 2-amino-5-methylbenzamide (7) as a white solid (25.3 g, 96%): mp 173-175 °C (lit.31 mp 179 °C); ¹H-NMR (CDCl₃) δ 6.60 (d, 1 H, J = 8 Hz, C₃-H). To an ice-cold suspension of 7 (25.3 g, 0.168 mol) in 3.6 N HCl (260 mL) was added dropwise over 25 min a solution of NaNO₂ (12.75 g, 0.185 mol) in H₂O (100 mL), while keeping the internal temperature below 5 °C. After another 20 min of stirring at this temperature, 10 N NaOH (100 mL) was added, causing all the solid to dissolve. The solution was acidified to pH 2 with 12 N HCl and chilled. The solid was filtered and recrystallized from EtOH to obtain off-white needles (26.2 g, 97%): mp 217-218 °C (dec. gas evolution) (lit. 32 mp 219–220 °C); IR (KBr) ν 1680 (lactam C=O); ¹H-NMR (d_6 -DMSO + D₂O) δ 3.30 (s, 3 H, 6-Me), 8.00 (m, 3 H, aryl); UV (95% EtOH) λ_{max} 208, 225, 282 nm.

Dimethyl N-[4-[[(3,4-Dihydro-4-oxo-1,2,3-benzotriazin-6yl)methyl]amino]benzoyl]-L-glutamate (15). N-Bromosuccinimide (1.37 g, 7.7 mol) was added in a single portion to a solution of 8 (1.13 g, 7 mmol) in glacial AcOH (70 mL) in an oil bath at 60 °C. The resulting solution was heated at 70 °C with illumination from a 150-W floodlamp for 2 h. After evaporation of the solvent under reduced pressure, the residue was partitioned between CHCl₃ and H₂O. The CHCl₃ layer was washed with 5% NaHCO₃, rinsed with H₂O, and evaporated to obtain a product (0.946 g) which was estimated to consist of a 2:3 mixture of 6-(bromomethyl)-1,2,3-benzotriazin-4(3H)-one (13) [1H-NMR (CDCl₃) δ 4.63] and unchanged 8 [1 H-NMR (CDCl₃) δ 3.30]. The entire mixture of 13 and 8 from the bromination reaction (estimated to contain 1.97 mmol of 13 from the δ 4.63/ δ 3.30 ratio) was added in a single portion to a solution of dimethyl N-(4aminobenzoyl)-L-glutamate (0.588 g, 2 mmol) in dry DMF (10 mL). Then, solid NaHCO₃ (0.168 g, 2 mmol) was added, and the mixture was kept in an oil bath at 65 °C for 3 days. The solvent was evaporated under reduced pressure, and the residue was partitioned between CHCl₃ and H₂O. The CHCl₃ layer (TLC: R_f 0.70, 0.65, 0.4, 0.0; silica gel, 19:1 CHCl₃-MeOH) was evaporated, and the residue was chromatographed on a silica gel column (45 g, 2.5 × 40 cm) with 20:1 CHCl₃-MeOH as the eluent. Pooled fractions containing the spot with $R_f = 0.4$ were evaporated in two batches, and the residues were dried separately in vacuo (P2O5, 65 °C). The first batch (0.254 g, 28%) was TLC-homogeneous, whereas the other (0.154 g, 17%) contained a small impurity. Rechromatography of the TLC-homogeneous batch afforded a beige powder (0.22 g, 87% recovery): mp 94-102 °C; IR (KBr) ν 3430, 3040, 2960, 1740, 1690, 1640, 1615, 1580, 1515, 1445, 1420, 1335, 1285, 1265 cm⁻¹; NMR (CDCl₃) δ 1.71 (br s, 1 H, lactim OH), 2.43 (m, 4 H, CH_2CH_2), 3.63 (s, 3 H, γ -COOMe), 3.75 (s, 3 H, α -COOMe), 4.65 (m, 4 H, CH₂N, NH, α -CH), 6.58 (d, J = 9 Hz, 2 H, 3'- and 5'-H), 7.68 (d, J = 9 Hz, 2 H, 2'- and 6'-H), 7.8-8.4 (m, 3 H, aryl). Anal. $(C_{22}H_{23}N_5O_6\cdot 0.6H_2O)$ C, H, N.

N-[4-[[(3,4-Dihydro-4-oxo-1,2,3-benzotriazin-6-yl)methyl]amino]benzoyl]-L-glutamic Acid (4, ADDF). A stirred cloudy solution of 15 (195 mg, 0.42 mmol) in MeOH (5 mL) was treated dropwise with 1 N NaOH (5 mL) over 1 min. The solution quickly became homogeneous, and after another 5 min the pH was adjusted to neutrality with HCl. The MeOH was evaporated under reduced pressure, and the product was purified by preparative HPLC (8% MeCN in 0.05 M NH4OAc, pH 6.9). Pooled pure fractions were concentrated by rotary evaporation and then freeze-dried. The residue was redissolved in H₂O (15 mL), and the solution was lyophilized again to obtain a pale-yellow solid (162 mg, 81%): mp ca. 150 °C (gas evolution after softening at 140-145 °C); IR (KBr) v 3420, 3150, 3040, 2980, 1690, 1615, 1580, 1555, 1520, 1445, 1405, 1330, 1285, 1270 cm⁻¹; NMR (d_6 -DMSO) δ 2.10 (m, 5 H, CH₂CH₂, lactim OH), 4.15 (m, 1 H, α -CH), 4.57 $(m, 2 H, CH_2N), 6.60 (d, J = 8 Hz, 2 H, 3'- and 5'-H), 7.02 (m, 2 H, 2 H, 3'- and 5'-H), 7.02 (m, 3 H, 2 H, 3'- and 5'-H), 7.02 (m, 3 H, 3'- and 5'-H), 7.02 (m, 3'- and 5'-H), 7.02 (m, 3'- and 5'-H), 7.02 (m, 3'- and$ 1 H, NH), 7.57 (d, J = 8 Hz, 2 H, 2'- and 6'-H, with overlapped m, 1 H, NH), 8.10 (m, 3 H, aryl); UV (0.1 M HCl) λ_{max} 203 nm (ε 27700), 224 (27600), 290 (12300); UV (pH 7.4 phosphate buffer) λ_{max} 216-218 nm (plateau, ϵ 32,200), 293 (24,300); UV (0.1 M

NaOH) $\lambda_{\rm max}$ 298 nm (ϵ 25,300). Anal. ($C_{20}H_{19}N_5O_6$ -0.75NH $_3$ -2H $_2$ O) C, H, N.

Enzyme Assays. A. Thymidylate Synthase (TS). The plasmid pETSM which allowed expression of mouse fibroblast TS at very high levels was obtained from Dr. Lee F. Johnson (Departments of Biochemistry and Molecular Genetics, Ohio State University). E. coli transformed with pETSM were grown in liquid medium, and expression of mouse TS was induced as previously described.35 Mouse TS extracted from the bacteria was purified to near homogeneity by affinity chromatography on a column of 10-formyl-5,8-dideazafolyl-Sepharose as described by Rode and co-workers³⁶ with minor modifications.³⁷ (6S)-Tetrahydrofolate was prepared and stored in 1-µmol aliquots as previously described. 38 Immediately prior to enzyme assays, an ampoule of tetrahydrofolate was dissolved in 1% 2-mercaptoethanol (2-ME), and the concentration was determined spectrophotometrically. A solution of 2 mM (6S)-tetrahydrofolate was made up in 1% 2-ME containing 25 mM formaldehyde and allowed to stand at room temperature for 10 min to allow formation of (6R)-5,10methylenetetrahydrofolate (CH₂THF). The CH₂THF, along with ADDF and 6 mIU of TS, was added to a cuvette in 950 μ L of 3-morpholinopropanesulfonic acid buffer, pH 6.8, and the cuvette was preincubated at 37 °C for 3 min. Both the buffer and the cuvette chamber were preequilibrated at 37 °C. The reaction was initiated by addition of dUMP, in a volume of 50 μ L, to the other reactants. Initial rates of reaction were calculated from the change in absorbance at 340 nm. Final concentrations of the reaction components were: CH2THF, 2.5-20 µM; buffer, 25 mM; formaldehyde, 2.5 mM; 2-ME, 15 mM; dUMP, 100 μ M.

B. Folylpolyglutamate Synthetase (FPGS) and Glycinamide Ribonucleotide Formyltransferase (GARFT). FPGS from mouse liver and GARFT from cultured L1210 cells were obtained and assayed as previously described. 37,39

Cell Growth Inhibition. The origin of the L1210 murine leukemia cells used to measure the effect of ADDF on cell growth, as well as the assay conditions, were described earlier.⁴⁰ Three separate experiments using replicate cultures were carried out, and the results were averaged.

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Supplementary Material Available: A Lineweaver-Burk plot for the FPGS-catalyzed addition of glutamic acid to compound 4 (1 page). Ordering information is given on any current masthead page.

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