

EFFECT OF POLYGLUTAMYLATION OF 5,10-METHYLENETETRAHYDROFOLATE ON THE BINDING OF 5-FLUORO-2'-DEOXYURIDYLATE TO THYMIDYLATE SYNTHASE PURIFIED FROM A HUMAN COLON ADENOCARCINOMA XENOGRAFT

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Abstract— $\text{CH}_2\text{-H}_4\text{PteGlu}$ and H_4PteGlu exist in human colon adenocarcinoma xenografts predominantly in the form of polyglutamate species at concentrations of $<3\text{ }\mu\text{M}$. The interaction of polyglutamates of $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}$ in the formation and stability of $[6\text{-}^3\text{H}]\text{FdUMP}$ –thymidylate synthase– $\text{CH}_2\text{-H}_4\text{PteGlu}_n$ ternary complexes has therefore been examined using enzyme purified from a human colon adenocarcinoma xenograft. Dissociation of these complexes was first-order and was dependent upon the concentration of folate. $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_{3-6}$ (0.9 to $1.6\text{ }\mu\text{M}$) were >200 -fold and $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_2$ ($18.2\text{ }\mu\text{M}$) was 18-fold more effective than $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_1$ ($335\text{ }\mu\text{M}$) at stabilizing ternary complexes for a $T_{1/2}$ for dissociation of 100 min. Polyglutamylation of $\text{CH}_2\text{-H}_4\text{PteGlu}$ also increased the affinity of binding of $[6\text{-}^3\text{H}]\text{FdUMP}$ to thymidylate synthase as determined by Scatchard analysis at folate concentrations of $10\text{ }\mu\text{M}$, where the K_d in the presence of $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_1$ was in the order of $4.0 \times 10^{-8}\text{ M}$, and for $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_{3-5}$ was between 3.7 and $5.5 \times 10^{-9}\text{ M}$. To examine whether this effect was due to differences in the rates at which $[6\text{-}^3\text{H}]\text{FdUMP}$ was bound (k_{on}) or dissociated (k_{off}) from the enzyme, the apparent rate of $[6\text{-}^3\text{H}]\text{FdUMP}$ binding was determined in the presence of $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_1$, $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_3$ and $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_4$. The k_{on} values were similar and were in the range of 1.7 to $2.3 \times 10^6\text{ M}^{-1}\text{ min}^{-1}$ for 10 or $20\text{ }\mu\text{M}$ folate concentrations. Differences in binding affinity determined for $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_1$ and longer polyglutamate forms of $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}$ were thus due to differences in k_{off} . The V_{max} for the initial velocity of $[6\text{-}^3\text{H}]\text{FdUMP}$ binding was achieved at $10\text{ }\mu\text{M}$ folate. Consequently, at concentrations of $\text{CH}_2\text{-H}_4\text{PteGlu}$ polyglutamates present in tumors, inhibition of thymidylate synthase by FdUMP *in vivo* would be expected to be transient, based upon the concentration of $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_n$ required for maximal formation and stability of the covalent ternary complex. It would be advantageous for modulation of $\text{CH}_2\text{-H}_4\text{PteGlu}_n$ pools to increase the concentrations of the longer polyglutamate species ($n \geq 3$) to maximize the interaction between FdUMP , thymidylate synthase and $\text{CH}_2\text{-H}_4\text{PteGlu}$.

Thymidylate synthase (EC 2.1.1.45; 5,10-methylenetetrahydrofolate:dUMP, C-methyltransferase) catalyzes the reductive methylation of dUMP to dTMP and is an important target in the treatment of adenocarcinoma of the colon by the anticancer agent FUra.† 5-Fluorouracil is metabolized to FdUMP which forms a covalent ternary complex with thymidylate synthase in the presence of $\text{CH}_2\text{-H}_4\text{PteGlu}$, resulting in the inhibition of dTMP biosynthesis. The

dissociation of this inhibitory complex is first-order, and dependent upon the concentration of unbound $\text{CH}_2\text{-H}_4\text{PteGlu}$ [1, 2]. Consequently, the duration of inhibition of thymidylate synthase *in vivo* may be dependent upon the endogenous concentration of $\text{CH}_2\text{-H}_4\text{PteGlu}$.

Intracellularly, reduced folates exist as polyglutamate forms with glutamate residues linked through the γ -carboxyl groups [3–8]. Polyglutamates of $\text{CH}_2\text{-H}_4\text{PteGlu}$ containing up to seven glutamate residues have increased affinity for thymidylate synthase [9–11]. However, the influence of polyglutamylation of the cofactor on the formation and stability of the ternary complex remains poorly characterized. Polyglutamylation of $\text{CH}_2\text{-H}_4\text{PteGlu}$ may be important for increasing the velocity of binding of $[6\text{-}^3\text{H}]\text{FdUMP}$ to thymidylate synthase [11, 12], for reducing the rate at which the nucleotide dissociates from the enzyme [11, 13], or for binding the folate at lower concentration [11]. In human colon adenocarcinoma xenografts, di-, tri-, tetra-, penta- and hexaglutamate forms of $\text{CH}_2\text{-H}_4\text{PteGlu}$ have been identified [14]. Due to the importance of thymidylate synthase as a target in this disease, the

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† Abbreviations: FUra, 5-fluorouracil; FdUMP, 5-fluoro-2'-deoxyuridylate; dUMP, deoxyuridylate; dTMP, thymidylate; $[6R]\text{CH}_2\text{-H}_4\text{PteGlu}_n$, the natural diastereoisomer of 5,10-methylenetetrahydrofolate where n = the number of glutamate residues; H_4PteGlu , tetrahydrofolate; PteGlu, folic acid; 5-CHO- H_4PteGlu , 5-formyltetrahydrofolate, leucovorin; DTT, dithiothreitol; BSA, bovine serum albumin; TMDG buffer, 0.1 M Tris-HCl, pH 7.5, containing 1.5 mM MgCl_2 , 2 mM DTT and 10% glycerol; and TCA, trichloroacetic acid.

effects of these different polyglutamate species on the stability of the FdUMP–thymidylate synthase–CH₂-H₄PteGlu_n complex, the affinity of binding of [6-³H]FdUMP, and the rate of ternary complex formation have been examined using thymidylate synthase purified from a human colon adenocarcinoma xenograft.

METHODS

Materials. [5-³H]dUMP (22 Ci/mmol) and [6-³H]FdUMP (15–20 Ci/mmol) were obtained from Moravsek Biochemicals, Brea, CA. Pteroylpolyglutamates containing from 2 to 7 glutamyl residues, prepared by solid phase synthesis, were purchased from American Radiochemicals Inc., St. Louis, MO. NCS tissue solubilizer and ACS and OCS liquid scintillation fluids were obtained from the Amersham Corp., Arlington Heights, IL. Folic acid (PteGlu) and all other reagents and chemicals were obtained from the Sigma Chemical Co., St. Louis, MO, or were of reagent grade.

Immune-deprivation of mice. Four-week-old CBA/CAJ mice were immune-deprived by thymectomy followed 3 weeks later by whole-body irradiation (950 rads at a dose rate of 170 rads/min) from a ¹³⁷Cs source, and reconstitution with 3 × 10⁶ syngeneic bone marrow cells injected i.v. [15]. Mice received s.c. grafts of neoplastic tissue after a further 8–16 days.

Tumor line. The human colon adenocarcinoma xenograft line HxVRC₅ has been described previously [16–20]. Briefly, tumors were established as xenografts from a poorly differentiated primary adenocarcinoma obtained from a previously untreated patient [15, 18]. The line has maintained both biological and biochemical characteristics observed within the original human specimen [15] and has been insensitive *in vivo* to treatment with 5-fluoropyrimidines [19]. This particular tumor was selected for investigation from a series of six colorectal adenocarcinoma xenograft lines, as it has demonstrated the highest thymidylate synthase activity [16, 17]. Tumors were used for enzyme purification upon reaching ≥1 g in weight.

Purification of thymidylate synthase and assay. The purification of thymidylate synthase >4000-fold from HxVRC₅ human colon adenocarcinoma xenografts was performed as described previously [21]. The activity of thymidylate synthase was determined from the catalytic release of [³H] from [5-³H]dUMP at 37°, 3 min [21]. Standard conditions used were 10 μM [5-³H]dUMP (sp. act. 2.7 Ci/mmol) and 80 μM [6RS]CH₂-H₄PteGlu.

Synthesis of [6R]CH₂-H₄PteGlu_n from PteGlu. The natural [6R] isomers of CH₂-H₄PteGlu polyglutamates were prepared by enzymatic synthesis from the corresponding pteroylpolyglutamates using dihydrofolate reductase purified from methotrexate-resistant *Lactobacillus casei* and an excess of formaldehyde as previously described [21]. Their purity and concentration were determined by HPLC analysis; storage was at –80° in long glass tubes flushed with argon [21]. [6RS]CH₂-H₄PteGlu was prepared as previously reported [21].

Stability of the [6-³H]FdUMP–thymidylate syn-

thetase–CH₂-H₄PteGlu_n ternary complex. Complex was formed with [6-³H]FdUMP (100 nM), [6R]CH₂-H₄PteGlu₁ (10 μM), [6R]CH₂-H₄PteGlu₂ (6 μM) or [6R]CH₂-H₄PteGlu_{3,6} (1.5 μM) and 0.5 to 1.5 ml enzyme (27–40 nM [6-³H]FdUMP binding sites) in a total volume of 1.0 to 2.0 ml TMDG buffer (0.1 M Tris–HCl, 1.5 mM MgCl₂, 2 mM DTT and 10% glycerol, pH 7.5) at 37° over a period of 45 min and placed on ice. An equal volume of ice-cold buffer was added subsequently. To determine the stability of the ternary complex formed with [6R]CH₂-H₄PteGlu₁ in the presence of different concentrations of [6R]CH₂-H₄PteGlu₁, four reaction mixtures (total volume 800 μl) each contained 400 μl ternary complex, TMDG buffer (pH 7.5) and [6R]CH₂-H₄PteGlu₁ at concentrations of 2.5, 25, 50 or 75 μM. Non-radiolabeled FdUMP (100 μM) was also added to start the exchange reaction with [6-³H]FdUMP according to the method of Lockshin and Danenberg [2]. Ternary complex and combined reagents were prewarmed separately at 37° for 3 min prior to mixing. Reaction mixtures were incubated at 37° for up to 45 min. At times 0, 15, 30 and 45 min, duplicate aliquots of 80 μl were placed on ice, and treated with 1 ml of an ice-cold 5% charcoal suspension (containing 1% BSA and 0.1% dextran in TMDG buffer, pH 7.5), to adsorb unbound FdUMP. After a further 20 min on ice, mixtures were centrifuged (12,000 g, 2°, 4 min), supernatant fractions were filtered, and dpm in 0.6 to 0.8 ml were determined by liquid scintillation counting. Data were analyzed by linear regression analysis to yield half-times (T_{1/2}) for dissociation of the ternary complex.

Alternatively, the stability of the ternary complex formed with [6R]CH₂-H₄PteGlu₂ was determined in the presence of increasing concentrations of [6R]CH₂-H₄PteGlu₂. Four reaction mixtures (total volume 700 μl) each contained 350 μl of ternary complex, TMDG buffer, [6R]CH₂-H₄PteGlu₂ at a concentration of 1.5, 4, 8 or 15 μM, and 100 μM FdUMP. Finally, the stability of ternary complex formed with tri-, tetra-, penta- and hexaglutamates of [6R]CH₂-H₄PteGlu was determined in the presence of 0.38, 0.88, 1.9 or 2.9 μM [6R]CH₂-H₄PteGlu_n using the method outlined for [6R]CH₂-H₄PteGlu₁.

Dissociation constant (K_d) for [6-³H]FdUMP in the presence of [6R]CH₂-H₄PteGlu_n. To determine relative differences in binding affinity between [6R]CH₂-H₄PteGlu₁ and [6R]CH₂-H₄PteGlu_n, enzyme (1 to 1.5 nM [6-³H]FdUMP binding sites) and [6R]CH₂-H₄PteGlu_n (10 μM) were incubated with various concentrations of [6-³H]FdUMP (0.2 to 200 nM) in a total volume of 250 μl Tris–HCl (25 mM, pH 7.4, containing 0.5% BSA, 1% β-mercaptoethanol and 10 mM sodium ascorbate), for 45 min at 37° by which time the reaction had reached equilibrium. The [6-³H]FdUMP–thymidylate synthase–CH₂-H₄PteGlu_n ternary complex was quantitated using TCA precipitation [22]. Briefly, duplicate reaction mixtures were treated with an equal volume of ice-cold 20% TCA to denature the protein. The mixtures were centrifuged for 5 min, 15,500 g, 2°, and the supernatant fractions were decanted. The precipitates were suspended subsequently in 1 ml TCA (10%), and the mixtures recentrifuged. This step was repeated twice, at which point the super-

natant fractions contained only background radioactivity. The precipitates were dissolved in 1 ml NCS at room temperature overnight, and radioactivity was determined following addition of 10 ml of OCS scintillant. Data were analyzed by the method of Munson and Rodbard [23] for a one-site binding model using the computer program "Ligand" and analyzed using an Apple IIe computer. Estimates for the association constant (K_a) were obtained using nonlinear least squares regression, where initial values were determined from the slopes of the Scatchard curves. The dissociation constant (K_d) was calculated as the inverse of K_a .

Influence of CH₂-H₄PteGlu_n on the apparent rate constant for FdUMP association (k_{on}) and dissociation (k_{off}). Measurement of the apparent rate constant for [6-³H]FdUMP association was essentially as described by Lockshin and Danenberg [2]. Reaction mixtures (1.9 ml) containing [6R]CH₂-H₄PteGlu_n (10 or 20 μM), enzyme (4.5 nM [6-³H]FdUMP binding sites), and buffer (25 mM Tris-HCl, pH 7.4, containing 1% BSA, 1% β-mercaptoethanol and 10 mM sodium ascorbate) were prewarmed at 37°; prewarmed [6-³H]FdUMP (10 or 25 nM) was added to start the reaction. At intervals of up to 2 min, aliquots (300 μl) were pipetted into an equal volume of ice-cold TCA (20%), and precipitated radiolabeled ternary complex was determined as described.

Under these conditions, the apparent bimolecular rate constant (k_{on}) for [6-³H]FdUMP association to thymidylate synthase was calculated from the equation

$$k_{on}t = \frac{1}{[E_0] - [FdUMP_0]} \times \ln \frac{[FdUMP_0]([E_0] - [X])}{[E_0]([FdUMP_0] - [X])}$$

where $[E_0]$ is the initial concentration of enzyme binding sites, $[FdUMP_0]$ is the initial concentration of [6-³H]FdUMP, and $[X]$ is the concentration of ternary complex at time t (min); k_{off} was calculated from the equation $K_d = k_{off}/k_{on}$ [2], where K_d and k_{on} had been determined experimentally.

Concentration of CH₂-H₄PteGlu_n at V_{max} for [6-³H]FdUMP binding. Reaction mixtures (100 μl), in duplicate, containing thymidylate synthase (3 nM [6-³H]FdUMP binding sites), [6-³H]FdUMP (25 nM), CH₂-H₄PteGlu_n (1–20 μM) and 25 mM Tris-HCl, pH 7.4 (containing 1% BSA, 1% β-mercaptoethanol and 10 mM sodium ascorbate), were incubated at 37° for 2 min. Ice-cold 20% TCA (100 μl) was added to terminate the reaction, and precipitated ternary complex was determined as described. K_m values were determined by the method of Cleland [24] using computer-assisted analysis.

RESULTS

The effect of polyglutamylation of CH₂-H₄PteGlu on the stability of the [6-³H]FdUMP–thymidylate synthase–CH₂-H₄PteGlu_n ternary complex was examined using enzyme purified from HxVRC₅ human colon adenocarcinoma xenografts. Results obtained for complex formed with [6R]CH₂-H₄PteGlu₄ and subsequently reincubated with [6R]CH₂-H₄PteGlu₄ are shown in Fig. 1 in comparison to dissociation of the ternary complex formed with [6R]CH₂-H₄PteGlu₁ in the presence of various concentrations of the monoglutamate. In both instances, dissociation of complexes was first-order, and the stability was dependent upon the concentration of the folate. At the lowest concentration of [6R]CH₂-H₄PteGlu₁ used (2.5 μM), the ternary complex dissociated with a $T_{1/2}$ of 32 min; for 75 μM [6R]CH₂-H₄PteGlu₁, the $T_{1/2}$ for dissociation was increased to 53 min. However, for [6R]CH₂-

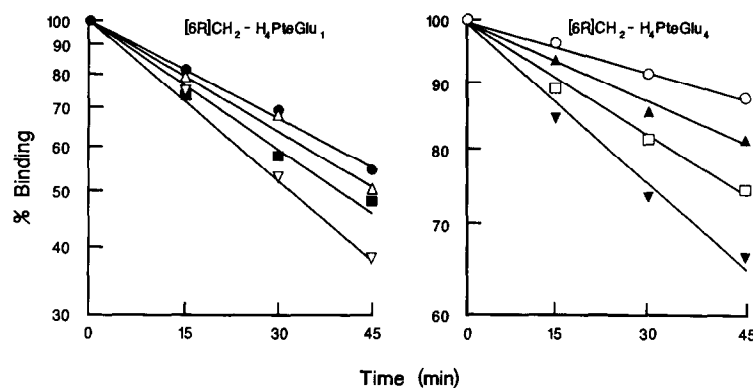


Fig. 1. Left panel: Stability of the [6-³H]FdUMP–thymidylate synthase–CH₂-H₄PteGlu₁ complex in the presence of increasing concentrations of [6R]CH₂-H₄PteGlu₁. Reaction mixtures containing radiolabeled ternary complex (6.8 nM), a 4000-fold excess of non-radiolabeled FdUMP (100 μM) and TMDG buffer, pH 7.5, in a total volume of 800 μl were incubated at 37° for up to 45 min in the presence of (▽) 2.5, (■) 25, (Δ) 50 or (●) 75 μM [6R]CH₂-H₄PteGlu₁. At 15-min intervals, 80-μl aliquots, in duplicate, were placed on ice and treated with 1 ml of an ice-cold 5% charcoal suspension. The amount of ternary complex remaining was determined as described in Methods. Right panel: Stability of the [6-³H]FdUMP–thymidylate synthase–CH₂-H₄PteGlu₄ complex in the presence of increasing concentrations of [6R]CH₂-H₄PteGlu₄. Reaction mixtures (800 μl) containing radiolabeled ternary complex (9.8 nM), FdUMP (100 μM), TMDG buffer, pH 7.5, and (▼) 0.38, (□) 0.88, (▲) 1.9 or (○) 2.9 μM [6R]CH₂-H₄PteGlu₄ were incubated at 37° for up to 45 min and processed as described to determine the amount of ternary complex remaining. All data were evaluated by linear regression analysis (left panel, $r^2 > 0.989$; right panel $r^2 > 0.963$).

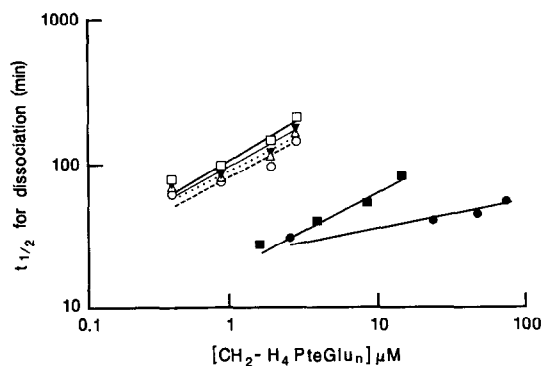


Fig. 2. Relationship between $T_{1/2}$ for dissociation and concentration of $[6R]CH_2-H_4PteGlu_n$. Data were derived from experiments described in the legend to Fig. 1 and were evaluated by linear regression analysis ($r^2 > 0.991$). Key: (●—●) $[6R]CH_2-H_4PteGlu_1$, (■—■) $[6R]CH_2-H_4PteGlu_2$, (△···△) $[6R]CH_2-H_4PteGlu_3$, (□—□) $[6R]CH_2-H_4PteGlu_4$, (○—○) $[6R]CH_2-H_4PteGlu_5$, and (▼—▼) $[6R]CH_2-H_4PteGlu_6$.

$H_4PteGlu_4$ at a concentration of $0.38 \mu M$, the $T_{1/2}$ for dissociation was 73 min, and it was increased to 215 min in the presence of $2.9 \mu M$ tetraglutamate.

The relationship between the $T_{1/2}$ for dissociation and concentration of $CH_2-H_4PteGlu_n$ was determined for $n = 1$ to 6 (Fig. 2). The magnitude of the difference between $[6R]CH_2-H_4PteGlu_1$ and $[6R]CH_2-H_4PteGlu_4$ was similar for $[6R]CH_2-H_4PteGlu_3$, $[6R]CH_2-H_4PteGlu_5$ and $[6R]CH_2-H_4PteGlu_6$; $[6R]CH_2-H_4PteGlu_2$ was somewhat more effective at stabilizing the ternary complex in comparison to $[6R]CH_2-H_4PteGlu_1$, but it was inferior to the longer polyglutamate chain length forms of the cofactor. Quantitation of the data from linear regression analyses yielded concentrations of $[6R]CH_2-H_4PteGlu_{3-6}$ (0.9 to $1.6 \mu M$) that were >200 -fold lower than required for $[6R]CH_2-H_4PteGlu_1$ ($335 \mu M$) to stabilize the complex with a $T_{1/2}$ for dissociation of 100 min (Table 1); for this degree of stabilization, $18 \mu M$ $[6R]CH_2-H_4PteGlu_2$ was required. When ternary complex was formed with $[6R]CH_2-H_4PteGlu_1$ and incubated in the presence of increasing concentrations of $[6R]CH_2-H_4PteGlu_4$, the relationship between $T_{1/2}$ for dissociation and folate concentration was identical to data derived from formation and stabilization of the

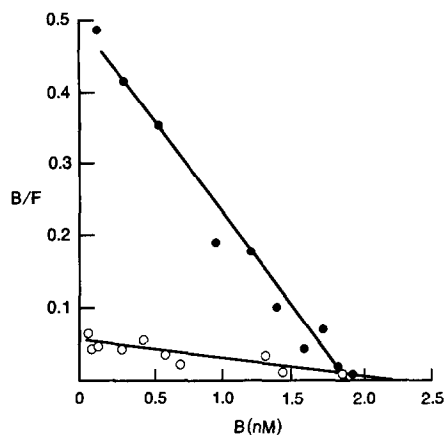


Fig. 3. Scatchard analysis for the binding of $[6-^3H]FdUMP$ to thymidylate synthase in the presence of (○) $[6R]CH_2-H_4PteGlu_1$ or (●) $[6R]CH_2-H_4PteGlu_4$. Reaction mixtures containing thymidylate synthase (1 to 1.5 nM), $[6R]CH_2-H_4PteGlu_n$ ($10 \mu M$), $[6-^3H]FdUMP$ (0.2 to 200 nM) and 25 mM Tris-HCl buffer, pH 7.4, containing 1% β -mercaptoethanol, 10 mM sodium ascorbate and 0.5% BSA in a total volume of $250 \mu l$, were incubated at 37° for 45 min. Ternary complex formed was determined by acid precipitation as described in Methods. Data were analyzed for a one-site binding model by the method of Munson and Rodbard [23].

complex with $[6R]CH_2-H_4PteGlu_4$ alone (data not shown).

To determine the effect of polyglutamylation of $CH_2-H_4PteGlu$ on the affinity of binding of $[6-^3H]FdUMP$ to thymidylate synthase, enzyme was incubated with $10 \mu M$ $[6R]CH_2-H_4PteGlu_n$ and various amounts of $[6-^3H]FdUMP$ at 37° for 45 min prior to measurement of bound $[6-^3H]FdUMP$ using TCA precipitation. Data plotted according to the Scatchard equation are shown in Fig. 3 for $[6R]CH_2-H_4PteGlu_1$ and $[6R]CH_2-H_4PteGlu_4$, where a large difference in slopes of the lines was evident. Using computer-assisted nonlinear least squares regression analysis [23], data fitted a one-site binding model; K_d values determined for $[6R]CH_2-H_4PteGlu_1$, $[6R]CH_2-H_4PteGlu_3$, $[6R]CH_2-H_4PteGlu_4$ and $[6R]CH_2-H_4PteGlu_5$ are shown in Table 2. From two separate experiments, the K_d for $[6-^3H]FdUMP$ binding in the presence of $[6R]CH_2-H_4PteGlu_1$ was in the order of 4×10^{-8} M. Addition of two glutamate residues decreased the binding constant by

Table 1. Concentrations of $CH_2-H_4PteGlu_n$ required to stabilize the ternary complex with a $T_{1/2}$ for dissociation of 100 min*

$CH_2-H_4PteGlu_n$ n	Concentration (μM)	Ratio $\frac{[CH_2-H_4PteGlu_1]}{[CH_2-H_4PteGlu_{2-6}]}$
1	335	1
2	18.2	18
3	1.3	268
4	0.9	372
5	1.6	216
6	1.2	291

* Calculated from linear regression analysis.

Table 2. Dissociation constants (K_d) for the binding of [6-³H]FdUMP to thymidylate synthase in the presence of CH₂-H₄PteGlu_n*

CH ₂ -H ₄ PteGlu _n n	K_d (M)	×-fold
1	4.2×10^{-8}	1
	3.8×10^{-8}	1
3	4.0×10^{-9}	10.1
	3.7×10^{-9}	10.8
4	3.8×10^{-9}	10.5
5	5.5×10^{-9}	7.3

* Reaction mixtures contained 1 to 1.5 nM enzyme, 10 μ M [6R]CH₂-H₄PteGlu_n, and varied concentrations of [6-³H]FdUMP (0.2 to 200 nM), as described in Methods.

Table 3. Rates of association (k_{on}) and dissociation (k_{off}) for the binding of [6-³H]FdUMP to thymidylate synthase purified from HxVRC₅ tumors

CH ₂ -H ₄ PteGlu _n n	k_{on} * (M ⁻¹ min ⁻¹)	k_{off} † (min ⁻¹)
1	2.0×10^6 (10) 1.9×10^6 (20)	8.0×10^{-2}
3	1.7×10^6 (20)	
4	2.3×10^6 (10) 2.1×10^6 (20)	8.7×10^{-3}

* Reaction mixtures contained 4.5 nM enzyme, 10 or 25 nM [6-³H]FdUMP and 10 or 20 μ M [6R]CH₂-H₄PteGlu_n, indicated in parentheses.

† Estimated from the equation $K_d = k_{off}/k_{on}$ [2] for 10 μ M [6R]CH₂-H₄PteGlu_n.

10-fold. The binding affinity was also 7- to 10-fold higher for [6R]CH₂-H₄PteGlu₄ and [6R]CH₂-H₄PteGlu₅ in comparison to [6R]CH₂-H₄PteGlu₁.

To determine whether the change in K_d for the binding of [6-³H]FdUMP to thymidylate synthase in the presence of [6R]CH₂-H₄PteGlu polyglutamates was due to differences in the rates at which FdUMP was bound (k_{on}) or dissociated (k_{off}), the apparent rate of [6-³H]FdUMP binding to HxVRC₅ thymidylate synthase in the presence of [6R]CH₂-H₄PteGlu₁, [6R]CH₂-H₄PteGlu₃ and [6R]CH₂-H₄PteGlu₄ (10 or 20 μ M) and two concentrations of [6-³H]FdUMP (10 or 25 nM) was determined (Table 3). Data for [6R]CH₂-H₄PteGlu₄ (20 μ M) are shown in Fig. 4. Irrespective of the number of glutamate residues added to CH₂-H₄PteGlu, the apparent k_{on} values for [6-³H]FdUMP binding were similar for concentrations of both 10 and 20 μ M, and were determined to be between 1.7 and 2.3×10^6 M⁻¹ min⁻¹. From the equation $K_d = k_{off}/k_{on}$, the k_{off} was calculated to be 8.0×10^{-2} min⁻¹ for [6R]CH₂-H₄PteGlu₁ (10 μ M) and 8.7×10^{-3} min⁻¹ for [6R]CH₂-H₄PteGlu₄ (10 μ M; Table 3).

The effect of the concentrations of [6R]CH₂-H₄PteGlu_n on the initial velocity of binding of [6-³H]FdUMP (25 nM) to thymidylate synthase was determined for [6R]CH₂-H₄PteGlu₁, [6R]CH₂-H₄PteGlu₃ and [6R]CH₂-H₄PteGlu₅ at 37° over a period of 2 min, during which time the reaction was linear (Fig. 5). The relationship between initial velocity and folate concentration was similar among the polyglutamate species. From computer-assisted analysis, the K_m values were determined to be 3.7 μ M for each polyglutamate form; at 10 μ M [6R]CH₂-H₄PteGlu_n, the rate of [6-³H]FdUMP binding had reached a maximum.

DISCUSSION

The existence in mammalian cells of folate species as polyglutamate forms is well established. For CH₂-H₄PteGlu, the presence of polyglutamates in L1210 cells was suggested when the covalent ternary complex isolated from these cells was found to have a higher molecular weight (1000 daltons) than the FdUMP-thymidylate synthase-CH₂-H₄PteGlu₁

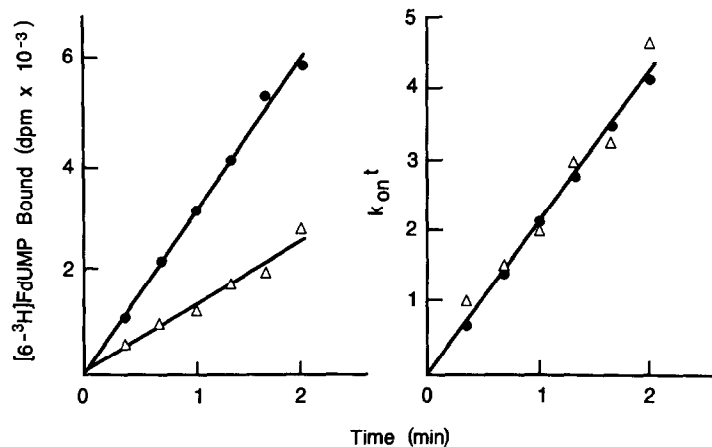


Fig. 4. Determination of the apparent k_{on} for [6-³H]FdUMP binding in the presence of [6R]CH₂-H₄PteGlu₄. Reaction mixtures contained thymidylate synthase (4.5 nM), [6R]CH₂-H₄PteGlu₄ (20 μ M), [6-³H]FdUMP [(Δ) 10 nM or (●) 25 nM] and 25 mM Tris-HCl buffer, pH 7.4, containing 1% β -mercaptoethanol, 10 mM sodium ascorbate and 1% BSA, in a total volume of 1.9 ml. Incubation was at 37° for up to 2 min. At timed intervals, aliquots (300 μ l) were pipetted into an equal volume of ice-cold 20% TCA, and the amount of ternary complex formed was determined as described in Methods.

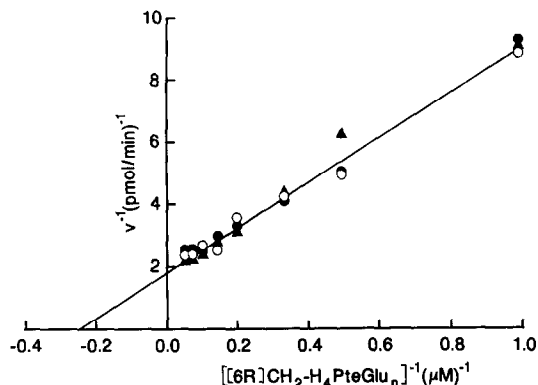


Fig. 5. Relationship between initial velocity (v) of ternary complex formation and concentration of (●) $[6R]CH_2-H_4PteGlu_1$, (▲) $[6R]CH_2-H_4PteGlu_3$ and (○) $[6R]CH_2-H_4PteGlu_5$. Reaction mixtures containing thymidylate synthase (3 nM), $[6-^3H]FdUMP$ (25 nM), Tris-HCl buffer (25 mM, containing 1% BSA, 1% β -mercaptoethanol and 10 mM sodium ascorbate, pH 7.4), and $[6R]CH_2-H_4PteGlu_n$ (1–20 μ M) were incubated at 37° for 2 min. Reactions were terminated and ternary complexes quantitated after acid precipitation as described in Methods.

complex when analyzed by gel electrophoresis [25]. More specifically, the predominant forms in mammalian liver and kidney have been determined to be the penta- and hexa-glutamates [26, 27], the relative distribution for liver dependent upon species. Similarly, in two human colon adenocarcinoma xenografts examined, $CH_2-H_4PteGlu_5$ constituted between 46 and 65% of the total $CH_2-H_4PteGlu_n$ pool, with $CH_2-H_4PteGlu_6$ ranging from 22 to 27% of the pool; di-, tri- and tetra-glutamates could also be detected, but in lower amounts [14]. Consequently, in attempting to understand the interaction between FdUMP, $CH_2-H_4PteGlu$ and thymidylate synthase in human colon adenocarcinoma xenografts, it was considered important to determine the way in which the various polyglutamate species of $CH_2-H_4PteGlu$ affected the formation and stability of the ternary complex.

The interaction of FdUMP, thymidylate synthase and $[6R]CH_2-H_4PteGlu_1$ has been well characterized for bacterial enzyme derived from *L. casei* [1] and mammalian thymidylate synthase using CCRF-CEM human leukemic cells as the enzyme source [2]. In both instances, an ordered mechanism of ligand binding was determined, the nucleotide binding prior to the folate, with dissociation of $[6R]CH_2-H_4PteGlu_1$ prior to FdUMP. Such an ordered mechanism resulted in the dependence of the stability of the complex upon the concentration of $[6R]CH_2-H_4PteGlu_1$, but not FdUMP. The influence of $[6R]CH_2-H_4PteGlu_5$ on the stability of the ternary complex has been reported for thymidylate synthase purified from MCF-7 human breast cancer cells [11], where increased stabilization was determined ($T_{1/2} \approx 187$ min) in comparison to $[6R]CH_2-H_4PteGlu_1$ ($T_{1/2} \approx 57$ min) at lower concentration (38 and 1 μ M respectively). For the colon tumor enzyme, approximately 120 μ M $[6R]CH_2-H_4PteGlu_1$ and 5 μ M $[6R]CH_2-H_4PteGlu_5$ would be required to

achieve similar levels of stability, data for $[6R]CH_2-H_4PteGlu_1$ being closer to the value determined for enzyme isolated from CCRF-CEM cells (≈ 180 μ M; [2]). In the current study, $[6R]CH_2-H_4PteGlu_{3-6}$ were similar in their capacities to stabilize the ternary complex and were considerably more effective than either $[6R]CH_2-H_4PteGlu_2$ or $[6R]CH_2-H_4PteGlu_1$. This effect for $[6R]CH_2-H_4PteGlu_{3-6}$ was accompanied by an increased binding affinity for $[6-^3H]FdUMP$ as determined by a 7- to 10-fold decrease in K_d determined by Scatchard analysis at 10 μ M $[6R]CH_2-H_4PteGlu_n$. The differences between $[6R]CH_2-H_4PteGlu_1$ and $[6R]CH_2-H_4PteGlu_5$ were in the range reported for thymidylate synthase purified from human breast cancer cells [11]. For enzyme from this source, a 2.4-fold increase in k_{on} and a 3-fold decrease in k_{off} were responsible for the difference in K_d reported for the polyglutamate species. However, in the current study, the increasing binding affinity for $[6R]CH_2-H_4PteGlu_n$ appeared to be due solely to a decrease in the rates of dissociation (k_{off}). Thus, current data demonstrating that the dissociation of FdUMP was a function of the concentration of the polyglutamate species of $CH_2-H_4PteGlu$ confirm the ordered mechanism of ligand binding as originally proposed [1, 2]. This is in contrast to suggestions by others [28] that the binding sequence observed with polyglutamates is in the reverse order from that seen with the monoglutamyl cofactor.

Concentrations of $CH_2-H_4PteGlu_n$ and $H_4PteGlu_n$ in cells have been estimated in the range of 3 μ M or less [29, 30]. In three human colon adenocarcinoma xenografts examined, the combined pools of $CH_2-H_4PteGlu_n$ and $H_4PteGlu_n$ were determined to be from 0.5 to 2.7 μ M, and for $CH_2-H_4PteGlu_n$ alone, in the order of 0.19 to 1.7 μ M [14]. Ullman *et al.* [31] found in L1210 cells that the concentrations of reduced folates required for optimal cell growth (100 nM) are suboptimal for achieving maximal cytotoxicity with FdUrd (500 nM). In addition, the presence of a high proportion of intracellular folates as polyglutamates (90%) has been considered important in enhancing the antithymidylate effect of FUra in cultured cells [32]. In human colon xenografts, however, there is no correlation between the distribution of polyglutamate species and the sensitivity of tumors to FUra [14]. It is also clear that concentrations of $CH_2-H_4PteGlu_1$ required to stabilize the covalent ternary complex (>300 μ M) would be considerably higher than those achievable physiologically in colon tumors, even with supplements of reduced folates such as 5-CHO- $H_4PteGlu$. Concentrations of $CH_2-H_4PteGlu$ polyglutamates also appear suboptimal for forming ternary complex at the maximal rate (10 μ M) or for stabilizing the complex to a level that would cause significant and prolonged inhibition of thymidylate synthase in human colon adenocarcinomas (several hours). In human colon adenocarcinoma xenografts [19], and in patients with colorectal adenocarcinoma [33, 34], responses to FUra are transient. More recently, the combination of FUra with the reduced folate 5-CHO- $H_4PteGlu$ (leucovorin) has increased the response rates by 3- to 5-fold for this disease from data generated in randomized clinical trials [35–37],

significantly prolonging the time to disease progression [35, 37] and overall survival [37]. Although no experimental studies have been reported to date, this is presumed to be due to an increase in the pools of CH₂-H₄PteGlu that, in turn, increase the formation and stability of the FdUMP-thymidylate synthase-CH₂-H₄PteGlu ternary complex and the degree and duration of thymidylate synthase inhibition. Data from our current study indicate that the choice of a schedule for leucovorin administration should be one that increases the concentrations of the longer polyglutamate chain length forms of CH₂-H₄PteGlu to optimize the interaction between FdUMP, thymidylate synthase and CH₂-H₄PteGlu.

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