

## THYMIDYLATE SYNTHASE IN ADVANCED GASTROINTESTINAL AND BREAST CANCERS

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**Knowledge of population thymidylate synthase (TS) levels in malignant tumors and normal tissues is essential for the use of TS as a predictor for 5-fluorouracil treatment. Tumor tissue TS levels in fresh frozen surgical biopsies from 136 patients with gastrointestinal or breast cancer, not previously subjected to chemotherapy, were analysed by [<sup>3</sup>H]FdUMP radioligand binding assay. TS levels were  $2.4 \pm 0.31$  pmol/g in liver metastases of colorectal cancer (n = 87),  $4.2 \pm 1.0$  pmol/g in primary colorectal cancer (n = 13),  $2.7 \pm 0.93$  pmol/g in gastric cancer (n = 13),  $3.1 \pm 1.7$  pmol/g in pancreatic cancer (n = 10),  $3.4 \pm 1.4$  pmol/g in breast cancer (n = 13) and  $0.58 \pm 0.075$  pmol/g in normal liver parenchyma (n = 24). TS levels were significantly higher in malignant tumor tissues compared to normal liver parenchyma.**

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For patients with advanced local growth or metastases of gastrointestinal or breast cancer the prognosis is poor. Surgery and radiotherapy may be beneficial for localized problems, such as intestinal, biliary, or urinary obstruction, painful osseal metastases, or neural compression, but generally these measures do not affect survival. Palliative chemotherapy, based on fluorinated pyrimidines, can prolong survival with an improved quality of life for a significant number of patients (1, 2). As it is difficult to predict the individual response to chemotherapy in these patients (3), an understanding of the biochemical strategy of cancer cells will be of importance for the design of enzyme pattern-directed anticancer chemotherapy (4). Inhibition of thymidylate synthase (TS, EC. 2.1.1.45) has been proposed as the major mechanism for the effect of 5-fluorouracil (5-FU) against tumor cells (5, 6).

TS catalyses de novo thymidine synthesis in eukaryotic cells, i.e. methylation of deoxyuridylate (dUMP) to thymi-

dylate (dTMP). The source of the methyl one-carbon group transferred by TS is polyglutamated 5,10-methylenetetrahydrofolate ( $\text{CH}_2\text{FH}_4$ ). The expression of the TS gene is cell cycle associated and high in the S-phase. High TS-levels are found in tissues with high cell turnover such as tumors. Surveys of normal tissues, recently reviewed by Spears (7), have shown relatively high levels in thymus, spleen, bone marrow, and testis, and low levels in visceral hollow organs. It has also been suggested that the expression of the TS gene is inhibited by its own end product and increased by dUMP in an autoregulatory manner (8-10). High TS enzyme levels prior to treatment, possibly due to gene amplification in tumors, may be responsible for innate 5-FU resistance (11). However, data on baseline levels of TS in normal human tissues and malignant tumors are scarce (12, 13).

The aim of the present study was to measure TS-levels in previously untreated human gastrointestinal cancer, female breast cancer and normal liver parenchyma, and to establish baseline levels of TS activity when measured with the ligand binding (ternary complex formation) assay technique.

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### Material and Methods

#### Patients

In 160 patients operated on at the Department of Surgery, Östra Sjukhuset, Göteborg, tissue specimens were

analysed for TS binding activity. The patients were operated on due to clinical routine indications, and all tissue specimens for TS analysis were taken from fresh tumor specimens or tumor biopsies for routine histological examinations. Normal liver biopsies were taken from 24 patients operated on for cholecystolithiasis. Histological examination of these showed normal liver parenchyma and these patients had no previous history of malignant disease. Eighty-seven patients had liver metastases of colorectal cancer, 13 had primary colorectal cancer, 13 gastric cancer, 10 advanced pancreatic cancer, and 13 female patients had breast cancer. The patients were followed postoperatively according to clinical routines, but there was no follow-up study protocol. Patient characteristics, i.e., age and sex, and tumor characteristics such as intratumor TS, degree of differentiation, and Dukes' classification of primary colorectal cancer were registered. Tissue  $\text{CH}_2\text{FH}_4$ -content was also measured in 38 tumor specimens and in liver parenchyma from 11 patients. Informed consent was obtained from all patients before the biopsy procedure.

#### Tissue preparation

The tumor biopsies were immediately frozen in liquid nitrogen and stored at  $-70^\circ\text{C}$  until the analyses of free TS and reduced folates were performed. Portions of frozen tumor tissue were then weighed and minced by means of a pair of scissors in a 10-fold excess of homogenisation buffer at  $4^\circ\text{C}$ . Homogenates were prepared by the action of rotating shearing blades (Turrax) for 30 s at  $4^\circ\text{C}$ .

#### Analysis of TS and methylenetetrahydrofolate

The  $[^3\text{H}]\text{FdUMP}$  ligand-binding assay was used for the analysis of free TS (14). One ml of the homogenate (duplicate samples) was centrifuged at  $4^\circ\text{C}$  and  $4\,000 \times g$  for 20 min. To 50  $\mu\text{l}$  of the cytosol, 25  $\mu\text{l}$  of homogenisation buffer, 50  $\mu\text{l}$   $[^3\text{H}]\text{FdUMP}$  (about 200,000 dpm) in  $\text{H}_2\text{O}$  and 25  $\mu\text{l}$  of 46,2  $\mu\text{mol}$   $\text{CH}_2\text{FH}_4$  was added. These procedures were carried out at  $2-4^\circ\text{C}$ . The tubes were incubated at  $37^\circ\text{C}$  for 10 min, and then 3 ml of 3% acid charcoal (rapidly stirred at  $4^\circ\text{C}$ ) was added followed by vortex mixing and centrifugation at  $4^\circ\text{C}$  and  $4\,000 \times g$  for 15 min to precipitate unbound radioactivity. Thereafter 0.8 ml of the supernatant was subjected to beta scintillation counting. TS concentrations were expressed as pmole bound FdUMP per gram tissue (wet weight).

Tumor tissue levels of  $\text{CH}_2\text{FH}_4$  and  $\text{FH}_4$  were assayed with some modifications according to the Priest methodology (15). To 50  $\mu\text{l}$  aliquots of cytosol or homogenisation buffer alone (blank) were added 50  $\mu\text{l}$  of the  $[^3\text{H}]\text{FdUMP}$  (about 800 000 dpm), 4 pmole of L. Casei TS in 25  $\mu\text{l}$  of homogenisation buffer (with 0,2% BSA). Buffer with or without formaldehyde was used for the estimation of

$\text{FH}_4 + \text{CH}_2\text{FH}_4$  and  $\text{CH}_2\text{FH}_4$  respectively. These procedures were carried out at  $2-4^\circ\text{C}$  and the tubes were then incubated at  $37^\circ\text{C}$  for 10 min, whereafter 1 ml ice-cold 3% acid charcoal was added for separation of protein-bound radioactivity. The samples were then centrifuged at  $4^\circ\text{C}$  and  $4\,000 \times g$  for 15 min, and scintillation counting performed. Folate concentrations were expressed as nmoles per gram tissue (wet weight).

#### Statistics

Differences between groups of patients were analyzed with a multifactorial ANOVA. Multiple regression analysis was performed for the analysis of variable correlation. All data are presented as mean  $\pm$  standard error of the mean (SEM).

#### Results

Baseline TS levels were  $2.4 \pm 0.31$  pmol/g in liver metastases of colorectal cancer ( $n = 87$ ),  $4.2 \pm 1.0$  pmol/g in primary colorectal cancer ( $n = 13$ ),  $2.7 \pm 0.93$  pmol/g in gastric cancer ( $n = 13$ ),  $3.1 \pm 1.7$  pmol/g in pancreatic cancer ( $n = 10$ ),  $3.4 \pm 1.4$  pmol/g in breast cancer ( $n = 13$ ) and  $0.58 \pm 0.08$  pmol/g in normal liver parenchyma ( $n = 24$ ). The distributions of TS levels were skewed in both gastrointestinal and breast cancers (Fig. 1), but close to normally distributed in healthy liver parenchyma. Baseline TS levels were significantly higher in malignant tissues than in normal livers, but the differences between the various types of malignancies were not statistically different. The TS level in primary colorectal cancers was higher than in hepatic metastases of colorectal tumors.

The degree of differentiation was correlated to tumor tissue TS levels (Fig. 2). The small number of primary

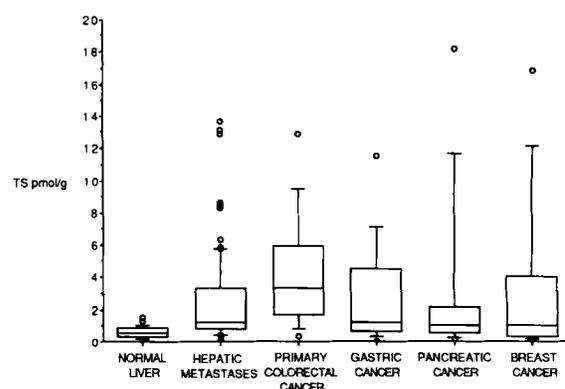


Fig. 1. TS levels in biopsies from normal liver parenchyma, ( $n = 24$ ), liver metastases of colorectal cancer, ( $n = 87$ ), primary colorectal cancer, ( $n = 13$ ), gastric cancer, ( $n = 13$ ), pancreatic cancer, ( $n = 10$ ), and breast cancer, ( $n = 13$ ). Lines and boxes represent the 10th, 25th, 50th, 75th and 90th percentile, circles represent individual extreme values below the 10th and above the 90th percentile. Baseline TS values in normal liver parenchyma are significantly lower than in the malignant tissues.

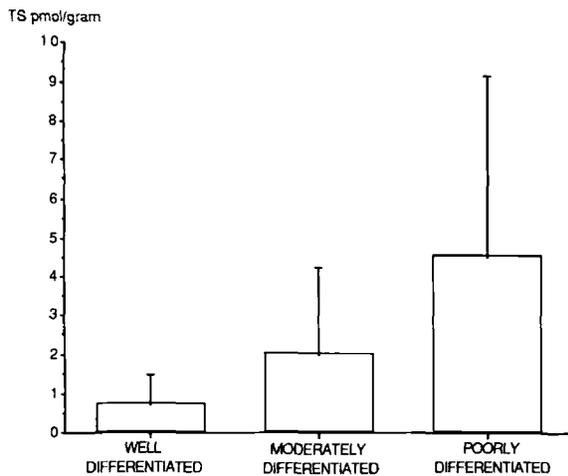


Fig. 2. Tumor tissue TS levels (mean  $\pm$  SEM) in 136 cancer patients in relation to differentiation grade, well differentiated ( $n = 21$ ), moderately differentiated ( $n = 66$ ) and poorly differentiated ( $n = 49$ ). The difference between poorly and well differentiated tumors was statistically significant ( $p < 0.05$ )

colorectal cancers did not allow reliable correlation analysis between Dukes' classification and intratumor TS levels.

In a limited number of samples, the levels of methylene tetrahydrofolates in tumor tissue were also measured but there was no significant correlation between folate and TS levels (Fig. 3). Mean  $\text{CH}_2\text{FH}_4$  levels were in normal livers  $2.2 \pm 0.52$  nmol/g ( $n = 11$ ), in liver metastases of colorectal cancers  $0.24 \pm 0.08$  nmol/g ( $n = 23$ ), in gastric cancers  $0.25 \pm 0.9$  nmol/g ( $n = 4$ ), in pancreatic cancers  $0.12 \pm 0.09$  nmol/g ( $n = 7$ ), and in breast cancers  $0.70 \pm 0.28$  nmol/g ( $n = 4$ ).  $\text{CH}_2\text{FH}_4$  was significantly lower in tumor tissue compared with normal liver parenchyma ( $p < 0.05$ ) but the differences between different tumor types were not statistically significant. From primary colorectal cancers we did not obtain enough biopsy material for the analysis of  $\text{CH}_2\text{FH}_4$ .

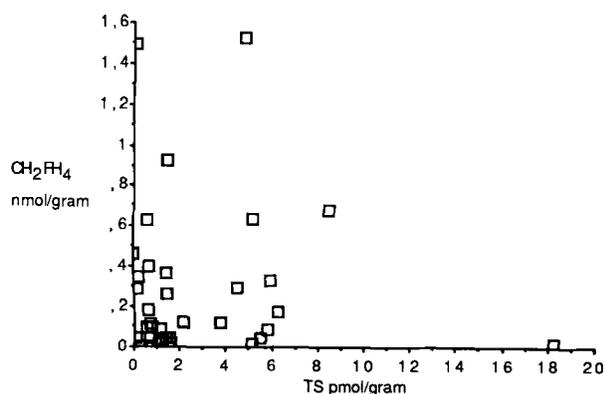


Fig. 3. Tumor tissue  $\text{CH}_2\text{FH}_4$  levels in 49 cancer patients in relation to tumor tissue TS levels. There was no statistically significant correlation between the two parameters.

## Discussion

The results in the present study reveal that TS levels are higher in malignant tissues than in normal liver parenchyma. Although no significant differences between tumor types were found, TS levels in malignant tumors were correlated to histological differentiation (16). Earlier reports, utilising different techniques, have also linked high TS levels to poor prognosis and metastatic spread in ovarian and colorectal cancer (12, 13, 17–19). In our previous studies of total and free TS levels after 5-FU therapy total TS in malignant tumors was significantly higher than in normal tissues (11, 20). However, in the earlier studies, free TS ( $\text{TS}_{\text{free}}$ ) was found to be higher in liver metastases than in primary colorectal tumors. On the other hand it has recently been reported that TS protein levels are equal in primary tumors and lymph node metastases of lung cancers (21).

Baseline TS levels were found to be higher in primary colorectal cancers than in liver metastases. However, the time course of tissue TS after i.v. 5-FU injections makes it difficult to estimate the true baseline, pre 5-FU TS, from TS after i.v. 5-FU injections. Earlier studies regarding TS levels after 5-FU i.v. push injections have shown a decline of inhibition starting 60 min after the injection (11) in accordance with a rebound increase of  $\text{TS}_{\text{free}}$  (20). In a recent study we have also found a divergence of  $\text{TS}_{\text{tot}}$  levels measured in tumor tissues and normal livers from one to five hours after i.v. 5-FU injections (Spears et al. unpublished study). Increasing  $\text{TS}_{\text{tot}}$  has also been observed in experimental tumors resistant to 5-FU, whereas decreasing  $\text{TS}_{\text{tot}}$  levels were found after i.v. 5-FU injections in the experimental tumors sensitive to 5-FU (22). This divergence may be related to if apoptosis is induced and further sustains the theory that 5-FU as other cell cycle specific agents do not kill cells by their biochemical actions per se but by dissociation of normally integrated cell cycle events, with induction of programmed cell death (7). 5,10- $\text{CH}_2\text{FH}_4$ -polyglutamate in combination with dUMP/FdUMP protects TS against proteolytic degradation which may be the key mechanism whereby the normal proteolytic mechanism of TS turnover is potentially slowed by TS protection within the TS-FdUMP-folate-polyglutamate ternary complex (23).

Measurements of TS in human tissues are associated with considerable problems due to low tissue levels of this enzyme. The samples in the present study had not been exposed to 5-FU and consequently FdUMP inhibition of enzyme posed no analytical problems. The advantages of the ligand binding assay of TS include high reproducibility and precision with a sensitivity requiring only 10 mg of tissue for duplicate assay (14, 24). The whole tissue is assayed and so the contribution of normal stromal elements is also included. If normal adjacent stromal cells produce a lot of thymidine, this could produce a bystander effect for contributing to DNA salvage pathway resistance

in the tumor cells, thus making whole tissue measurements important. The ligand binding assay is also conveniently performed in parallel with the folate assays. This helps to correct for tissue cellularity, in addition to describing the kinetic situation for optimal TS inhibition, predicting the response to 5-FU treatment and assessing the requirements for folate supplementation. Other available methods for TS analysis, such as the <sup>3</sup>H-displacement assay (25, 26), immunohistochemical labeling methods with monoclonal antibodies against TS protein (12), and reverse transcription polymerase chain reaction for the analysis of TS gene expression (27) may also be used for TS baseline studies. However, the optimal method for screening of large materials has not yet been determined.

Further prospective trials analysing tumor tissue TS and folate levels in relation to therapy response will be necessary in order to improve our tools for optimizing 5-FU + folate treatment in advanced cancer.

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