REGULATION OF ENZYMES BY LIGAND INDUCED CHANGE IN 211 POLYMERIZATION.

211 POLYMENIZATION. <u>Thomas W. Traut, Nancy Cheng, and Margaret M.</u> <u>Matthews</u>, University of North Carolina School of Medicine, Department of Biochemistry, Chapel Hill, North Carolina, USA Many enzymes readily change their aggregation state, by dissociation or association of subunits. This constitutes a mechanism for regulation, if the different molecular weight spe-cies vary in their specific activity, and if the equilibrium between these species is readily perturbed by physiological effectors. Our studies show that several enzymes involved in the synthesis or degradation of pyrimidines exhibit such regulathe synthesis or degradation of pyrimidines exhibit such regula-tion: The multifunctional protein UMP synthase (EC 2.4.2.10 + EC 4.1.1.23), and uridine kinase (EC 2.7.1.48) from Ehrlich ascites cells, and N-carbamoyl- β -alanine amidohydrolase (EC 3.5.1.6) from rat liver. Generally, substrates are positive effectors, while immediate or more distant end products are negative effectors.

UMP synthase shows only positive regulation: The 50,000 dalton monomer is inactive; effectors result in association to the active dimer. Uridine kinase shows negative regulation: the native enzyme exists as polymers of 4 or more subunits, and CTP or UTP produce dissociation to the 31,000 dalton subunit and loss of activity. NC β A amidohydrolase shows bimodal regulation: the native enzyme has an Mr of 235,000; the substrate produces association to 470,000 daltons; the product, β -alanine, causes dissociation to 120,000 daltons. Ligand induced changes in polymerization may be a general

mechanism for regulating enzyme activity.

THE MECHANISM OF INSULIN-INDUCED INCREASE OF THE RATE OF DE NOVO PURINE BIOSYNTHESIS 212 IN PRIMARY CULTURED RAT HEPATOCYTES

IN PRIMARY CULTURED RAT HEPATOCYTES Masami Tsuchiya, Mitsuo Itakura, Kamejiro Yamashita The University of Tsukuba, Institute of Clinical Medicine, Sakura-mura, Niihari-gun, Ibaraki 305, JAPAN The de novo purine biosynthesis was studied 8 hours after the administration of insulin. The rate of de novo purine biosynthesis assayed by [14C]formate incorporation started to increase at the insulin con-centration of 10⁻¹⁰M and reached the plateau at 10⁻⁷M or above with its level 2.8 folds higher than control. Serbosphoribesyl leveroposphate(PEPP) availability 5-phosphoribosyl 1-pyrophosphate (PRPP) availability assayed by $[1^{4}C]$ adenine incorporation increased steeply and reached the plateau with its level 2.3 folds higher than control at 10^{-9} M and stayed at this folds higher than control at 10 ⁻M and stayed at this level in the higher concentration. By the addition of Actinomycin D(lµg/ml) or cycloheximide(lµg/ml) increased rate of de novo purine biosynthesis and increased PRPP availability decreased to the control level. Amidophosphoribosyltransferase(ATase) activity per unit amount of protein or DNA increased 1.5 to 1.8 folds with increased Vmax and unchanged Km for PRPP. ATP concentration increased by 27% which was counterbalanced with decreased concentrations of AMP(73%), ADP(69%), GMP(73%) and GDP(79%) concentration. Re-sults of these studies suggest that insulin increases the rate of de novo purine biosynthesis by increasing ATase activity and PRPP availability.

213 THE EFFECTS OF URIDINE AND THYMIDINE ON THE DEGRADA-TION OF 5-FLUOROURACIL (5-FU), URACIL AND THYMINE BY RAT LIVER DIHYDROPYRIMIDINE DEHYDROGENASE (DPD).

Mendel Tuchman, Margaret L. Rammaraine, Robert F. O'Dea. University of Minnesota Medical School, University of Minnesota Hospitals, Departments of Pediatrics and Pharmacology, Minneapolis, Minnesota. U.S.A.

The effects of various nucleosides and nucleotides on the activity of DPD partially purified from rat liver was studied using 5-FU, uracil and thymine as substrates. 5-FU was found to have a Km similar to those of uracil and thymine. DPD showed the highest specific activity when utilizing 5-FU as substrate (Table). + .

+	5-FU (N)	Uracil (N)	Thymine (N)
Km	3.5 ± 0.4 (13)		2.2 ± 0.3 (10)
specific	0.82± 0.04 (18)	0.68± 0.05(20)	0.56± 0.04 (14)
activity		\$ §	t a p t

*calculated Km in µM, "nmols/min/mg pr ber of individual liver preparations. "nmols/min/mg protein, "mean ± S.E., num Uridine was found to be a potent non-competitive inhibitor of

DPD for all three substrates. Thymidine is much less potent as an inhibitor. Compared to 5-FU and uracil, thymine reduction is the most susceptible to inhibition by uridine and thymidine (Table). 1 t 1 E 111

Inhibitor	5-ru,	Uracii	Inymine
uridine	0.71	0.62	0.16
thymidine	24	41	8
*Ki expressed as	μM		

Uridine may enhance 5-FU cytotoxicity more than thymidine.

214 THYMIDINE INCORPORATION IN NUCLEOSIDE TRANSPORT DEFICIENT T LYMPHOMA CELLS. Buddy Uliman and Bruce Aronow. Univ. KY Medical Center, Dept. of Biochemistry, Lexington, KY, U.S.A. Nucleoside transport deficiency in mammalian cells is generally associated with an inability to transport all nucleosides and an inability to bind or be affected by p-nitrobenzylthioinosine (NBMPR), a potent inhibitor of nucleoside transport. Among nucleoside transport-deficient (NT-) S49 T lymphoma clones, however, the ability to take up and respond to the physiological effects of thymidine varied considerably. The NT- clone, AE₁, was almost as capable of incorporating thymidine into TTP as the wildtype parent. This incorporation of thymidine into TTP by AE₁ cells and a large percentage of thymidine incorporation into TTP by wildtype cells were unresponsive to NEMPR. Thus, thymidine uptake in NT- cells can be uncoupled genetically or pharmacologically in a specific manner from the uptake of other pharmacologically in a specific manner from the uptake of other nucleosides. Consequently, AE_1 cells were just as sensitive as wildtype cells to thymidine toxicity but were resistant to two thymidine analogs, BrdUrd and FdUrd. Moreover, the growth inhibitory effects of thymidine were ameliorated to a much greater extent in wildtype cells than in AE_1 cells by either NBMPR or deoxycytidine. Interestingly, however, AE_1 cells could not grow in hypoxanthine-amethopterin-thymidine entry indicated that AE_1 cells and NBMPR-treated wildtype cells transported thymidine at measurable rates. This residual thymidine transport capability can account for the phenotype of the NT⁻ cells toward thymidine.

215 REGULATION OF PURINE BIOSYNTHESIS IN G1 PHASE ARRESTED MAMMALIAN CELLS. Buddy Uliman and Bruce Aronow. Univ. KY Medical Center, Dept. of Biochemistry, Lexington, KY, U.S.A. The effects of G1 phase growth arrest on purine biosynthesis were studied in cultured S49 T lymphoma cells. Incubations of wildtype cells for 18 hr with dibutyryl cyclic AMP, which induces G1 arrest, reduced the rates of purine biosynthesis by 95%. Time course and concentration dependence studies indicated that the decrease in rates of purine biosynthesis correlated with the extent of G1 concentration dependence studies indicated that the decrease in rates of purine biosynthesis correlated with the extent of G1 phase arrest. Similar studies with somatic cell mutants deficient in some component of cyclic AMP action or metabolism indicated that the depression in purine synthetic rates required G1 arrest and did not result from cell death. Rates of RNA and DNA synthesis were also markedly diminished in the growth arrested cells. Measurements of purine synthesis in the presence of azaserine indicated that there was an early block in purine biosynthesis mean to the formation of presence of azaserine indicated that there was an early block in purine biosynthesis prior to the formation of phosphoribosylformylglycinamide. Additionally, the activities of adenylosuccinate synthetase and IMP dehydrogenase were diminished in Gl arrested cells. The levels of all controlling enzymes, substrates, and cofactors, however, were similar in cycling and Gl arrested cells. Despite diminished rates of purine biosynthesis, the levels of intracellular nucleotides in Gl cells were not reduced. These results suggest that perturbations in the consumption of nucleotides via inhibition of nucleic acid synthesis have profound effects on the purine pathway and indicate the importance of feedback inhibition by nucleotides in the regulation of purine synthesis in situ. nucleotides in the regulation of purine synthesis in situ.

MYOADENYLATE DEAMINASE DEFICIENCY: DIAGNOSIS BY

 $\begin{array}{c} 216 & \mbox{MYOADENYLATE DEAMINASE DEFICIENCY: DIAGNOSIS BY} \\ \text{FOREARM ISCHEMIC EXERCISE TESTING AND PLASMA PURINE} \\ \mbox{MEASUREMENTS. Peter A. Valen, Denny A. Nakayama,} \\ \mbox{Judith A. Veum, and Robert L. Wortmann. Medical College of Wisconsin, Medical College of Wisconsin Affiliated Hospitals, \\ \mbox{Department of Medicine, Milwaukee, WI USA.} \\ \mbox{Quantitation of lactate and ammonia (NH3) production after forearm ischemic exercise (FIE) has been proposed as a screening test for myoadenylate deaminase (MAD) deficiency. Plasma lactate, NH3 and purines (adenosine inosine, hypoxanthine, xanthine) were determined 1, 3, 5 and 10 minutes after 90 seconds of FIE and correlated with exercise performance (<math display="inline">\Sigma$ kg x sec), measured with a grip dynamometer, strain gauge and strip recorder. In 13 normals lactate and NH3 production correlated with exercise performance (r=0.76 and 0.72, respectively). Three patients with histochemically-defined MAD deficiency failed to generate NH3 despite increasing lactate and adequate performance. \\ \end{tabular}

patients with histochemically-defined MAD deficiency failed to generate NH3 despite increasing lactate and adequate performance. Retesting normals at submaximal performance produced lactate and NH3 generation patterns indistinguishable from those of the 3 patients. Normal subjects even with submaximal performance increased plasma purines from 10 to 38 µM after FIE. The MAD deficient patients raised their purine levels only 0-7 µM. These data demonstrate that submaximal exercise efforts by normals can give FIE test results indistinguisable from those of MAD deficient individuals if only lactate and NH₃ are assessed and plasma purine measurements after FIE testing are effective in identifying the deficiency state. These results support the hypothesis that disordered purine metabolism occurs in patients

hypothesis that disordered purine metabolism occurs in patients with MAD deficiency.