REPRESSION OF TRYPTOPHAN OPERON RNA SYNTHESIS BY trp REPRESSOR IN AN IN VITRO COUPLED TRANSCRIPTION—TRANSLATION SYSTEM

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1. Introduction

The expression of the tryptophan (trp) operon in E. coli is negatively controlled by the interactions of at least three factors: the product of the regulatory gene (trp R), the intact trp operator and excess L-tryptophan [1-5]. Recently we have purified the protein product of the trp R gene and have characterized it in a purified in vitro transcription system directed by DNAs from trp transducing phages [6]. These in vitro studies were in agreement with the classical model of repression of the trp operon [1, 7]. In this model, the product of the regulatory gene is an inactive aporepressor, which by interaction with L-tryptophan is converted to the active repressor. The activated trp repressor binds to the DNA at the trp operator to block transcription initiating at the trp promoter. Similar conclusions from in vitro studies have been reported by Rose et al. [8]. However, the possible contribution of coupling between transcription and translation in the regulation of the trp operon [9-12] remains to be elucidated.

A DNA-directed RNA-dependent in vitro protein synthesizing system has been developed in which the faithful expression of the genetic information coded on ΦΧ174 replicative form DNA and T7 DNA has been observed [13, 14]. In this paper, we have applied this cell-free system to study the bacterial trp operon. Trp specific RNA was transcribed from the correct strand of template DNA from a trp transducing phage and was furnished to the translational machinery. Addition of the isolated trp repressor into this coupled transcription—translation system specifi-

cally inhibited the expression of the *trp* operon at the transcriptional level. The *trp* repression depended upon the concentration of L-tryptophan.

2. Materials and methods

2.1. Coupled system extracts were prepared from E. coli D24 (RNase Γ , λ , Γ , met) cells according to the method of Bryan et al. [13].

2.2. Synthesis of RNA and protein in the coupled system were measured by the method of Bryan et al. [13]. The standard reaction mixture (0.13 ml) contained 1.65 μ moles Tris-HCl (pH 7.8), 0.1 μ mole NaCl, 1.5 μ moles Mg(COOCH₃)₂, 0.9 μ mole 2-mercaptoethanol, 0.6 µmole phosphoenol pyruvate, 0.25 μ mole ATP, 0.05 μ mole each of α -32 P-GTP (42 Ci/ mole), CTP and UTP, about 4 nmoles each of 20 amino acids (14 C-leucine, 263 Ci/mole), 5 µg pyruvate kinase, 5 μg λpt60-3 DNA, 5 μg RNA polymerase, 5 A 260 units ribosome fraction, and 0.75 A_{260} unit soluble fraction. When appropriate, 20 μ l of partially purified trp repressor were added to the reaction mixture, which was then kept for 5 min at 0°C prior to the addition of the RNA polymerase. After incubation at 33°C for the desired time intervals, radioactivities in the cold 5% CCl₃COOH (TCA)precipitable materials and the hot 7% HClO₄ (PCA)precipitable materials were measured as the amounts of RNA and protein synthesized, respectively.

2.3. trp Repressor was partially purified from E. coli W3110 Ilv, leu, pro, trp A₉₈₆₅ trp R rec A J₂₇₃/KLFH (thr, leu, pro, trp R) [15] cells as

described previously [6]. The DNA-cellulose fraction (1 mg protein/ml) was used for the present experiments.

2.4. Other experimental methods including the preparation of phage stocks, the extraction of phage DNAs, the separation of DNA strands, and the purification of DNA-dependent RNA polymerase have been described elsewhere [16].

3. Results and discussion

Fig. 1 shows the kinetics of RNA and protein synthesis in the in vitro coupled system using λ pt60-3 DNA, which carries an intact *trp* operon [12], as the template. RNA synthesis proceeded linearly for 20 min then reached a plateau level. After a lag of about 4 min protein synthesis continued for at least 45 min, indicating that translation closely follows the transcriptional process. Very little RNA and protein was synthesized in the absence of the template DNA. The

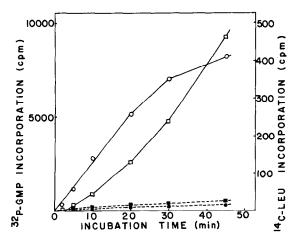


Fig. 1. Kinetics of RNA and protein synthesis in the coupled system under standard conditions. At the designated intervals, $20~\mu$ l samples were withdrawn from the standard reaction mixture (see Materials and methods) and pipetted into 2 ml of ice-cold 1% casamino acid solution to terminate the reaction. Radioactivities in the cold 5% TCA precipitable materials and the hot 7% PCA precipitable materials were measured, respectively. [32 P] GMP incorporation per 10 μ l with ($^{\circ}$ $^{\circ}$ $^{\circ}$ or without ($^{\circ}$ $^{\circ}$ $^{\circ}$) $^{\circ}$ $^{\circ}$ $^{\circ}$ $^{\circ}$) or without ($^{\circ}$ $^{\circ}$ $^{\circ}$ $^{\circ}$) or without ($^{\circ}$ $^{\circ}$ $^{\circ}$ $^{\circ}$ $^{\circ}$) or without ($^{\circ}$ $^{\circ}$ $^{\circ}$ $^{\circ}$) $^{\circ}$ $^{\circ}$

standard concentration of L-tryptophan in the coupled system is 0.0248 mM which is adequate for de novo protein synthesis in vitro [13, 14] but is insufficient to repress the trp operon in vivo [17, 18]. Since all the experiments described here were carried out with bacterial extracts prepared from a strain of E. coli carrying the intact trp R gene it is expected that the RNA and protein synthesis would be repressed by the action of endogenous trp repressor when the L-tryptophan concentration was increased to 0.509 mM, which is a sufficient level to cause repression in vivo and in vitro [3, 6, 12, 16]. However, no obvious inhibition of RNA and protein synthesis was observed when the concentration of L-tryptophan was increased (upper columns in table 1). We believe that some trp repressor molecules were removed from the soluble fraction of the cell extracts and/or were inactivated during preparation of the coupled system. Thus, most of the template DNA molecules, unhindered by trp repressor molecules, would be accessible for the synthesis of trp RNA and

Table 1

RNA and protein synthesis in the coupled system under difdifferent conditions

		Time	[³² P]GMP o incorporated		
Addition of trp			L-Trp conc.		
•	Synthesis		0.0248 mM	0.509 mM	b/a
		5'	1088 ^(a)	1055 ^(b)	_ 0.97
None	RNA	10'	2226	2309	1.04
		30'	6109	6303	1.03
		5'	32	31	0.97
None	protein	10'	74	71	0.96
	_	30'	196	206	1.05
		5′	885	726	0.82
Added	RNA	10'	1715	1489	0.87
		30'	6211	5442	0.88
		5'	27	22	0.79
Added	protein	10'	66	56	0.85
	•	30'	180	167	0.93

Reaction mixtures (0.13 ml) were set up as described in Materials and methods with L-tryptophan added to the final concentrations indicated. At the designated intervals, 20 µl samples were withdrawn from each reaction mixture to measure the net RNA and protein synthesis.

Table 2
Repression of trp RNA synthesis

Addition Time		trp ED RNA (% of input) L-Trp conc.		trp CBA RNA (% of input) L-Trp conc.	
None	5'	8.4	7.8	2.6	2.1
	10'	8.9	8.0	6.3	5.9
Added	5' 10'	5.3 6.5	1.2	2.1 2.4	0.87 1.1

[32P] GMP labeled RNAs were synthesized under various conditions as described in table 1 and the RNAs were purified from the reaction mixtures by phenol extraction [16]. 10 µl aliquots of the purified RNA samples dissolved in 110 μl of 0.2% SDS were separately hybridized to excess amounts (1.0 μg) of separated l-strands of Φ80ptED, Φ80ptCBA and Φ80 wild DNAs. Hybridizations were carried out in 0.15 ml of 0.30 M NaCl-0.03 M sodium citrate, pH 7.4 (2 × SSC) at 65°C for 4 hr. After RNase treatment (6 μg/ml of RNase A, Worthington Biochem. Co., and 3 units/ml of RNase T₁, Sankyo Co.) at 25°C for 30 min, the RNase-resistant RNA-DNA complex was collected on a presoaked membrane filter (Schleicher and Schuell Co., Bac-T-Flex, Type B6, 27 mm diam.) and washed with 50 ml of cold 2 × SSC and the radioactivity on a filter was counted. Hybridization efficiency was 95%. The difference in hybridization values between $\Phi80pt$ and $\Phi 80$ wild type was taken as a measure of the trp operon specific RNA, and is represented by percentage of input counts.

protein. When purified *trp* repressor together with a high concentration of L-tryptophan were added to the coupled system, RNA and protein synthesis was significantly inhibited (lower columns in table 1).

In order to know whether or not the reduction of RNA synthesis is due to the specific repression of transcription of the *trp* operon, we measured the amounts of *trp* RNA in the total RNA synthesized under various conditions. The results of DNA-RNA hybridization experiments are summarized in table 2. Without addition of *trp* repressor about 8 to 9% of the total RNA is *trp* RNA which is complementary to operator proximal *trp* E-D genes, regardless of the concentration of L-tryptophan. When *trp* repressor was added to the coupled system *trp* ED RNA synthesis was remarkably repressed by the higher concentration of L-tryptophan. Significant repression of *trp*

ED RNA synthesis was also observed at the lower concentration of L-tryptophan (about 40% repression at 5 min). The concentration of L-tryptophan that causes 50% repression in our coupled system is calculated to be about 0.03 mM, which seems to be several fold higher than that in other in vitro systems not involving translation [8, 19], our unpublished results [0.014 mM]).

During the initial 5 min of synthesis, there is much less RNA which is complementary to trp C-B-A genes than that complementary to trp E-D genes, indicating that transcription of the trp operon occurs sequentially from E gene to A gene as observed in vivo [20]. It is noteworthy that synthesis of significant amounts of trp RNA was observed even under fully repressed conditions (in the presence of both trp repressor and a high concentration of L-tryptophan). This could be explained by the following possibilities. (a) Some transcription initiated at the λN gene promoter proceeds into the trp operon even in the presence of active trp repressor on the trp operator [6, 12, 16]. (b) Transcription on the trp operon initiates mainly at the principal trp promoter but some transcription occurs at second minor promoter located between the trp D and C genes [21, 22], and the latter is not subject to the action of trp repressor [23].

In summary, we have demonstrated the repression of *trp* specific RNA synthesis in a coupled transcription—translation system. Repression was dependent upon the action of the *trp* repressor in the presence of excess L-tryptophan. This system would be useful for the further study of regulation of the *trp* operon.

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References

- [1] Cohen, G.N. and Jacob, F. (1959) C. R. Acad. Sci. 248, 3490-3492.
- [2] Yanofsky, C. (1971) J. Amer. Med. Assoc. 218, 1026– 1035.
- [3] Imamoto, F. (1973) Progr. Nucl. Acid. Res. and Mol. Biol. (Davidson, J.N. and Cohn, W.R., ed.) Vol. 13, p. 340-407.
- [4] Hiraga, S. (1969) J. Mol. Biol. 39, 159-179.
- [5] Morse, D.E. and Yanofsky, C. (1969) J. Mol. Biol. 44, 185-193.
- [6] Shimizu, Y., Shimizu, N. and Hayashi, M. (1973) Proc. Natl. Acad. Sci. U.S. 70, 1990-1994.
- [7] Jacob, F. and Monod, J. (1961) J. Mol. Biol. 3, 318–356.
- [8] Rose, J.K., Squires, C.L., Yanofsky, C., Yang, H.L. and Zubay, G. (1973) Nature New Biol. 245, 133-137.
- [9] Morse, D.E. and Yanofsky, C. (1969) J. Mol. Biol. 41, 317-328.
- [10] Rose, J.K. and Yanofsky, C. (1972) J. Mol. Biol. 69, 103-118.
- [11] Imamoto, F. (1973) J. Mol. Biol. 74, 113-136.

- [12] Imamoto, F. and Tani, S. (1972) Nature New Biol. 240, 172-175.
- [13] Bryan, R.N., Sugiura, M. and Hayashi, M. (1969) Proc. Natl. Acad. Sci. U.S. 62, 483-489.
- [14] Gelfand, D.H. and Hayashi, M. (1970) Nature 228, 1162-1165.
- [15] Morse, D.E. and Yanofsky, C. (1969) J. Mol. Biol. 44, 185-193.
- [16] Shimizu, N. and Hayashi, M. (1973) submitted.
- [17] Pouwels, P.H. and Stevens, W.F. (1973) Molec. Gen. Genet. 120, 55-68.
- [18] Doolittle, W.F. and Yanofsky, C. (1968) J. Bact. 95, 1283-1294.
- [19] McGeoch, D., McGeoch, J. and Morse, D.E. (1973) Nature New Biol. 245, 137-140.
- [20] Imamoto, F. (1969) Mol. Gen. Genet. 105, 298-305.
- [21] Jackson, E.N. and Yanofsky, C. (1972) J. Mol. Biol. 69, 307-313.
- [22] Shimizu, N., Shimizu, Y. and Hayashi, M. (1973) submitted.
- [23] Morse, D.E. and Yanofsky, C. (1968) J. Mol. Biol. 38, 447-451.