

POLY U-DEPENDENT RIBOSOMAL BINDING OF Lys-tRNA^{Lys}

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Received 11 May 1970

1. Introduction

There is some evidence that U-U base pairs can be formed at the polynucleotide level. Thus, enhanced hypochromicity of poly U at 0° [1] suggests that this polynucleotide acquires some ordered structure at this temperature and infrared studies [2] indicate that hydrogen bonding is involved in this transition. Moreover, polyamines were found to stabilize poly U ordered structure for, in the presence of one spermine equivalent, poly U exhibits a cooperative melting profile with 50% hypochromicity and T_m equal 29° [3]. A similar though much more stable complex is formed with the 5-methyl homolog of poly U [4]. The random coil-spermine complex transition is accompanied by abrupt changes in ORD spectrum, e.g., a four-fold increase of the long wavelength Cotton effect and a violet shift of the first trough [5]. The compactness of the resulting structure is evidenced by the marked increase (from 4.5 S to 9.5 S) of the sedimentation coefficient; the $S_{20,w}$ value becomes relatively independent of ionic strength in sharp contrast to the behavior of the random coil poly U [6]. Ordered poly U does not bind to ribosomes unless "melted out" [7]. The kinetics of formation of the poly U ordered state is independent of poly U concentration within a fairly wide range (10^{-5} to 10^{-3} M [3]) suggesting that it is an intramolecular phenomenon. Since poly U as well as oligo U's do not appear to show non-cooperative stacking interactions [6, 8, 9] it may tentatively be assumed that the ordered state of poly U corresponds to a hairpin-like helical molecule with hydrogen-bonded U-U base pairs. This is supported by the observation that poly

N-methyl U does not form any kind of ordered structure [18].

The above considerations and the fact that the stability of the trinucleotide codon-ribosomes-mRNA complex is higher than that of the corresponding trinucleotide-complementary polynucleotide complex [10] suggested the possibility of detecting U-U base in codon-anticodon interaction, such as occurs in the mRNA-directed ribosomal binding of aminoacyl-tRNAs. In looking for U-U interaction the possibility of competition from A-U base pairing should clearly be avoided. The experimental approach was to look for a poly U-dependent ribosomal binding of Lys-tRNA^{Lys} (lysine codons, AAA) in the absence of tRNA^{Phe}. Two lysine-accepting tRNA species are known in *E. coli* [11]; they would be expected [12] to have the anticodons (3'-5') UUC or UUU (UUI would appear to be excluded as anticodon for it would recognize asparagine codons AAU). Evidence for the occurrence of poly U-directed ribosomal binding of Lys-tRNA^{Lys} is presented in this communication.

2. Materials and methods

Pseudomonas sp. 412 was grown and ribosomes and ribosomal subunits prepared as previously described [13]. Ribosomes from this strain, a wild type psychrophile, are particularly suitable for assays near 0°. The ribosomes were washed twice with 1.0 M ammonium chloride [13]. Three samples of *E. coli* tRNA were used. Sample A was commercial (Schwarz Bioresearch) unfractionated *E. coli* tRNA. When charged with ¹⁴C-

lysine it accepted 37 pmoles/ A_{260} unit. Sample B was a tRNA^{Lys}-enriched fraction from *E. coli* obtained by chromatography on DEAE-Sephadex [14]. It accepted 270 pmoles of lysine and 3.6 pmoles of phenylalanine/ A_{260} unit (Lys: Phe = 75). Sample C was a highly purified preparation of tRNA^{Lys} from *E. coli* B, obtained by rechromatography of sample B on hydroxyapatite [15]. It accepted 1115 pmoles of lysine and 0.8 pmole of phenylalanine (Lys:Phe = 1400). tRNA^{Phe} (90% phenylalanine accepting capacity) was provided by the Oak Ridge National Laboratory.

Binding assay samples (0.05 ml) contained, tris-acetate buffer, pH 7.2, 2.5 μ moles; NH_4Cl , 5 μ moles; magnesium acetate, 0.5 μ moles; B-mercaptoethanol, 1 μ mole; NH_4Cl -washed *Pseudomonas* sp. 412 ribosomes, 2.5 A_{260} units; poly U (Miles), 10 μ g; Lys (^{14}C)-tRNA, 19–26 pmoles (^{14}C -Lysine specific radioactivity, 271 μ Ci/ μ mole; 1 pmole = 535 cpm in the Packard Tricarb Scintillation Counter). Samples with 30 S ribosomes contained in addition 0.2 μ mole of spermidine. The samples were incubated for 5 min at 30° (to form the poly U-ribosomal complex) followed by chilling to 0° prior to the addition of Lys-tRNA and further incubated for 60 min at 0°. Binding was measured by the procedure of Nirenberg and Leder [16].

3. Results and discussion

As seen in table 1, there was poly U-directed binding of ^{14}C -Lys-tRNA both with highly purified tRNA^{Lys} and with partially purified preparations. Binding was observed with either 30 S or 70 S ribosomes although net binding values were, as usual higher (about three-fold) with the latter. In contrast, no poly U-directed binding was observed when unfractionated tRNA was the source of Lys-tRNA. This was true irrespective of the experimental conditions (0° to 37°; 5 to 30 mM Mg^{2+}). In this case, non-specific binding in the absence of poly U was always 25–35% higher than in its presence, and observation made earlier by Nirenberg and Leder [6] in their original study of the binding system.

The effects of Mg^{2+} concentration, temperature, and tRNA^{Phe} on the binding of highly purified Lys-tRNA^{Lys} are shown in fig. 1. Increasing the Mg^{2+} concentration from 5 to 20 mM increased non-specific binding but augmented the poly U-dependent binding only moderately (fig. 1A), whereas increasing tempera-

Table 1
Poly U-dependent binding of Lys-tRNA to ribosomes.
(pmoles/assay, each assay in duplicate).

	Lys-tRNA preparations					
	A		B		C	
	minus poly U	plus poly U	minus poly U	plus poly U	minus poly U	plus poly U
<i>Binding to 70 S ribosomes</i>						
Expt. 1	0.72	0.51	0.63	0.94	0.55	0.97
Expt. 2	0.78	0.48	0.68	1.04	0.47	0.95
<i>Binding to 30 S ribosomes</i>						
Expt. 1	0.28	0.18	0.26	0.38	0.32	0.47
Expt. 2	—	—	0.22	0.35	0.33	0.51

ture (fig. 1B) above 10° markedly decreased poly U-directed binding without affecting the non-specific reaction. This shows that this binding is relatively weak. Fig. 1C shows that binding is prevented by tRNA^{Phe} (ratio tRNA^{Phe}: Lys-tRNA^{Lys}, 1:6). On the other hand, when tRNA^{Phe} is added after binding of Lys-tRNA^{Lys} has taken place, addition of tRNA^{Phe} has virtually no effect between 0° and 10° whereas at higher temperatures the more stable complex with tRNA^{Phe} is apparently formed at the expense of the less stable one with Lys-tRNA^{Lys}.

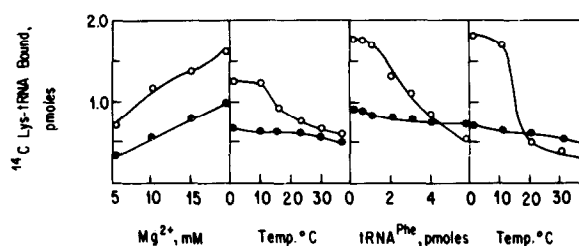


Fig. 1. Effect of Mg^{2+} , temperature and tRNA^{Phe} on poly U-dependent binding of Lys-tRNA^{Lys} "1100" to 70 S ribosomes.

- Effect of Mg^{2+} concentration, incubation for 60 min at 0°.
- Effect of temperature, 20 mM Mg^{2+} .
- Effect of tRNA^{Phe} added at 0 time. Incubation for 60 min at 0°, 20 mM Mg^{2+} .
- As in C., but tRNA^{Phe} (6 pmoles) was added after 60 min at 0° and samples were incubated for another 30 min at temperatures indicated. ●—●, without poly U; ○—○, with poly U. Other conditions as in table 1.

Addition of polyamines (spermine, spermidine) to samples with 70 S ribosomes appeared not to increase poly U-dependent binding of Lys-tRNA^{Lys} although it augmented non-specific binding in the absence of the polymer. However, with 30 S ribosomes no binding was detected in the absence of polyamines.

The results described provide further support for the idea that weak U-U base pairs can be formed under appropriate conditions. The codon-anticodon interaction leading to poly U-dependent ribosomal binding of Lys-tRNA^{Lys} would appear to be of the same nature as that responsible for the assumption of ordered structure by poly U. Weakness of the poly U-tRNA^{Lys} interaction is evidenced by the easy dissociation of the complex by tRNA^{Phe} and by elevated temperature.

Some relaxation of base pairing specificity in the presence of the ribosome at the 3'-terminus of the codon, or in the case of tRNA^{Met} at the 5'-terminus, is known but the involvement of the ribosome, if any, is not understood. Uhlenbeck, Baller and Doty [17] have recently shown that in the case of tRNA^{Met} "wobble" may be demonstrated in the absence of the ribosome. The formation of a Lys-tRNA-poly U ribosomal complex likewise implies little or no involvement on the part of the ribosome in affecting U-U base pairing specificity.

Acknowledgements

We are indebted to Dr. S.Ochoa for his advice and critical reading of the manuscript. We wish to thank

Mrs. S.Konar for technical assistance. Acid by American Cancer Society grants E 542 and PRA61 (to W.S.); E102 and P530 (to P.S.); U.S.P.H.S. grant CA00558 (to P.S.). This is publication No. 1374 of the Cancer Commission of Harvard University.

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