PEPTIDE CHAIN INITIATION IN A SPECIES OF NOSTOC AND IN CHLOROPLASTS OF EUGLENA GRACILIS

Francesco SALA, Silvana SENSI and Bruno PARISI Institutes of Genetics and Botany, University of Pavia, 27100 Pavia, Italy

Received 4 August 1970

1. Introduction

Peptide chain initiation in Escherichia coli requires the coordinate interaction of at least three protein factors (initiation factors). Through a sequence of partially understood reaction, N-formyl-methionyltRNA (fMet-tRNA), selected by the initiator codon AUG, is bound to the ribosome and becomes available for the formation of the first peptide bond [1-5]. A number of reports have demonstrated the presence of fMet-tRNA in other bacteria [6], in mitochondria [7-8] and in chloroplasts [9]. Furthermore, mitochondria [8] and chloroplasts [10] incorporate N-formyl-methionine into the N-terminal position of polypeptides. These findings raise the question of whether the reactions for peptide chain initiation are similar in all prokaryotic organisms and in organelles endowed with bacterial type ribosome (70 S), as already demonstrated in the case of the reactions for peptide chain elongation [11]. The similarity in the initiation reactions may be ascertained in vitro by testing the functional interchangeability of ribosomes and initiation factors prepared from different sources. This approach has already shown that ribosomes prepared from Bacillus subtilis and from mitochondria of Neurospora crassa may carry out the same initiation reactions catalyzed by E. coli ribosomes. In addition E. coli initiation factors are active on ribosomes from B. subtilis and from mitochondria of N. crassa [12].

In the present investigation we provide evidence that this is the case also for a species of a blue-green alga (Nostac sp.), and for chloroplasts of Euglena gracilis, both containing ribosomes of the bacterial type [13, 14].

2. Methods

E. gracilis strain Z was grown in the light at 27° in an heterotrophic acidic medium [15]. The cells were collected by centrifugation, washed twice with water and used immediately. Chloroplasts were prepared according to Eisenstadt and Brawerman [16]. "Unwashed" chloroplast ribosomes were prepared by resuspending the chloroplasts in a buffer containing 0.01 M tris-HCl (pH 7.8), 0.01 M MgCl₂, 0.05 M NH₄Cl, 0.006 M β -mercaptoethanol and 2% Triton-X-100 (BDH-Chemicals). For the preparation of "washed" chloroplast ribosomes, the concentration of NH₄Cl was raised to 0.5 M. After clarification at 30,000 g for 30 min, the ribosomes were pelleted by centrifugation at 105,000 g for 3 hr and resuspended in the low NH₄Cl buffer without Triton-X-100,

Nostoc sp., a strain isolated in our laboratory, was grown in the light at 27° in an autotrophic chelated medium [17]. Cells were harvested during the growth phase by centrifugation, washed twice with water and frozen at -70° until used. "Washed" ribosomes were prepared as described for *E. gracilis*, except that cell disruption was achieved by grinding in a mortar with sand (BDH-Chemicals; 40-100 mesh).

All ribosomal preparations were kept at -70° and used only once after thawing.

The preparation of *E. coli* ribosomes, initiation factors, and *N*-formyl-¹⁴C-methionyl-tRNA (f-¹⁴C-Met-tRNA) as well as the analytical methods and the assay for fMet-puromycin are described elsewhere [12]. *E. coli* ¹⁴C-phenylalanyl-tRNA (¹⁴C-Phe-tRNA) was prepared as described by Kaji, Kaji and Novelli [18].

3. Results and discussion

E. coli ribosomal preparations contain initiation factors, which can be washed off by treatment with high concentrations of NH₄Cl [19]. The results of table 1 show that this appears to be the case also for ribosomes prepared from chloroplasts of E. gracilis. Indeed, "unwashed" ribosomal preparations may carry out the synthesis of fMet-puromycin while "washed" preparations are almost totally dependent on added E. coli factors. We were unable to perform the same experiments on "unwashed" Nostoc ribosomes, due to difficulties encountered in the preparation of active ribosomes from this organism. However, washed ribosomes of Nostoc sp., as in the case of "washed" chloroplast ribosomes, are able to catalyze the synthesis of fMet-puromycin in heter-

Table 1
Synthesis of fMet-puromycin by "washed" and "unwashed" ribosomal preparations from Euglena gracilis chloroplasts,

Nostoc sp. and Escherichia coli.

D.:h		fMet-puromycin (pmoles per mg ribosome)	
Ribosomes from		-Initiation factors	+E. coli initiation factors
E. coli	"washed"	0.4	81.1
	"unwashed"	31.6	40.0
E. gracil	is "washed"	0.9	49.4
	"unwashed"	15.3	23.7
Nostoc sp."washed"		8.0	44.6

The 50 μ l assay mixtures contained 100 μ moles/ml tris-HCl pH 7.2, 50 μ moles/ml KCl, 5 μ moles/ml magnesium acetate, 1 μ mole/ml GTP, 10 μ moles/ml β -mercaptoethanol, 0.8 absorbance units/ml AUG and 422 pmoles/ml f-¹⁴C-Met-tRNA (sp. act. 233 μ Ci/ μ mole). Where indicated, the reaction mixtures contained 1.75 mg/ml of E. coli "washed" ribosomes, 1.40 mg/ml of chloroplast "washed" ribosomes, 1.55 mg/ml of chloroplast "unwashed" ribosomes, 1.55 mg/ml of chloroplast "unwashed" ribosomes, 2.0 mg/ml "washed" Nostoc sp. ribosomes. When present, 0.50 mg/ml of E. coli initiation factors were added. The extent of synthesis of fMet-puromycin was assayed with the method of Leder and Bursztyn [20] after incubation for 15 min at 24° followed by a 20 min incubation at the same temperature in the presence of 1 μ mole of puromycin.

Table 2
Characteristics of fMet-puromycin synthesis by "washed" ribosomes prepared from Nostoc sp., chloroplasts of Euglena gracilis and Escherichia coli.

	% Activity of ribosomes from		
Reaction mixtures	Nostoc sp.	Chloroplasts of E. gracilis	E. coli
Complete	100	100	100
-E. coli factors	10	11	2.2
-AUG	10	19	18
-GTP	32	44	23
-fMet-tRNA, +Phe-tRNA	1.5	0.7	0.6

The reaction mixtures and the assay conditions were as described in table 1. The complete mixtures containing ribosomes of *Nostoc* sp., of chloroplasts and of *E. coli* synthesized 44.6, 25.4 and 62.8 pmoles fMet-puromycin per mg ribosomes, respectively.

ologous mixtures containing *E. coli* initiation factors. In every case the properties of the reaction, i.e. dependence on the initiator codon AUG, GTP and fMet-tRNA are similar to those determined for the initiation reactions in *E. coli* (table 2).

No attempt was made to isolate initiation factors from the NH₄Cl-wash of chloroplast and *Nostoc* sp. ribosomes, owing to quantitative limitations in the material under study.

The above results give further support to the idea that in prokaryotic organisms (bacteria and blue-green algae) and in cellular organelles (chloroplasts and mitochondria) protein synthesis initiates by a mechanism which is similar if not identical to that outlined for E. coli. Such a concept is strengthened by this and previous reports that E. coli initiation factors and E. coli fMet-tRNA are both utilized for pentide chain initiation by the ribosomes prepared from B. subtilis [12], N. crassa mitochondria [12], Nostoc sp. and E. gracilis chloroplasts. It should be added that while such results argue in favor of an exchangeability of initiation factors F₁ and F₂, nothing may be concluded in the case of factor F₃, since such a factor is required for peptide chain initiation in vitro with natural mRNA, but not with the trinucleotide AUG [1, 3]. The striking structural and functional similarities in the complex

mechanism for peptide chain initiation gives further support to the hypothesis of a close phylogenetic relationship between bacteria, blue-green algae, chloroplasts and mitochondria.

Acknowledgements

The authors are grateful to Dr. O.Ciferri for critical discussion, to Dr. O.Tiboni for the isolation of the *Nostoc* strain and to Mr. G.Di Pasquale for skillful technical assistance. This work was supported by a grant from the Consiglio Nazionale delle Ricerche.

References

- [1] M.Revel, H.Herzberg and H.Greenshpan, Cold Spring Harbor Symp. Quant. Biol. 35 (1969) 261.
- [2] R.E.Thach, J.W.B.Hershey, D.Kolakofsky, K.F.Dewey and E.Renold-O'Donnel, Cold Spring Harbor Symp. Quant. Biol. 35 (1969) 277.
- [3] A.J.Wahba, Y.B.Chae, K.Iwasaki, R.Mazunder, M.J.Miller, S.Sabol and M.A.G.Sillero, Cold Spring Harbor Symp. Quant. Biol. 35 (1969) 285.
- [4] A.J.Wahba, K.Iwasaki, M.J.Miller, S.Sabol, M.A.G.Sillero and C.Vazquez, Cold Spring Harbor Symp. Quant. Biol. 35 (1969) 291.

- [5] G.Brawerman, Cold Spring Harbor Symp. Quant. Biol. 35 (1969) 307.
- [6] K.S.Seligman and L.R.Finch, FEBS Letters 8 (1970) 9.
- [7] A.E.Smith and K.A.Marcher, J. Mol. Biol. 38 (1968) 241.
- [8] H.Küntzel and F.Sala, Z. Physiol. Chem. 350 (1969) 1158.
- [9] G.Burckard, B.Eclancher and J.H.Weil, FEBS Letters 4 (1969) 285.
- [10] H.Bachmayer, Biochim. Biophys, Acta 209 (1970) 584.
- [11] O.Ciferri and B.Parisi, in: Progress in Nucleic Acid Research and Molecular Biology, Vol. 10, eds. J.N. Davidson and W.E.Cohn (Academic Press, New York, 1970) p. 121.
- [12] F.Sala and H.Küntzel, European J. Biochem. 15 (1970) 280.
- [13] B.Parisi, A.Perani, O.Tiboni and O.Ciferri, Giorn. Bot. It., in press.
- [14] G.Brawerman, Biochim. Biophys. Acta 72 (1963) 317.
- [15] S.H.Hutner, A.C.Zehalsky, S.Aaronson, H.Baker and O.Frank, in: Methods in Cell Physiology, Vol. 2, ed. D.M.Prescott (Academic Press, New York, 1966) p. 217.
- [16] J.M.Eisenstadt and G.Brawerman, in: Methods in Enzymology, Vol. 12 A, eds. L.Grossman and K.Moldave (Academic Press, New York, 1967) p. 476.
- [17] N.Lazaroff and W.Vishniac, J. Gen. Microbiol. 25 (1961) 365.
- [18] A.Kaji, H.Kaji and G.D.Novelli, J. Biol. Chem. 240 (1965) 1185.
- [19] W.M.Stanley, M.Salas, A.J.Wahba and S.Ochoa, Proc. Natl. Acad. Sci. U.S. 56 (1966) 290.
- [20] P.Leder and H.Bursztyn, Biochem. Biophys. Res. Commun. 25 (1966) 233.