

DNA homologies of ribosomal RNA genes of *Neurospora* species.

D. K. Mukhopadhyay

R. Mimiko

S. K. Dutta

Follow this and additional works at: <https://newprairiepress.org/fgr>



This work is licensed under a [Creative Commons Attribution-Share Alike 4.0 License](https://creativecommons.org/licenses/by-sa/4.0/).

Recommended Citation

Mukhopadhyay, D. K., R. Mimiko, and S.K. Dutta (1980) "DNA homologies of ribosomal RNA genes of *Neurospora* species.," *Fungal Genetics Reports*: Vol. 27, Article 8. <https://doi.org/10.4148/1941-4765.1671>

This Research Note is brought to you for free and open access by New Prairie Press. It has been accepted for inclusion in *Fungal Genetics Reports* by an authorized administrator of New Prairie Press. For more information, please contact cads@k-state.edu.

DNA homologies of ribosomal RNA genes of *Neurospora* species.

Abstract

DNA homologies of ribosomal RNA genes of *Neurospora* species.

Mukhopadhyay, D. K., R. Mniko, and S. K. Dutta.

DNA homologies of ribosomal RNA genes of
Neurospora species.

probably variable sequences. Extensive electron microscopic studies (Schibler et al. 1975 J. Molec. Biol. 94:503) of 45S precursor rRNA of several cold and warm blooded animals confirm that spacer regions vary extensively from species to species.

It was desirable to know whether such differences in rDNA sequences exist between Neurospora species. Any such difference should be detectable using standard procedures for DNA homology studies (Dutta 1976 Mycologia 68:388). rDNA sequences were isolated from N. crassa mycelial cells using the procedure described previously (Chattopadhyay et al. 1972 Proc. Natl. Acad. Sci. 69:3256). The purified rDNA was ³H-labeled (by nick translation) and reassociated with total DNA isolated from the heterothallic species N. crassa and from three homothallic

Ribosomal RNA genes (rDNAs) of Neurospora crassa contain DNA sequences which code for 17S, 5.8S, and 26S rRNAs, in addition to internal and external spacers (Free, Rice, and Metznerberg 1979 J. Bacte. 137:1219). As has been reported for many eukaryotes, the DNA sequences which code for 17S, 5.8S, and 26S rRNAs in Neurospora species are probably conserved while the internal and external spacer regions are

species: N. dodgei, N. lineolata, and N. africana. In addition, ³²P-labeled total DNA of N. crassa was re-annealed with unlabeled bulk DNA from N. crassa, N. dodgei, and N. lineolata.

TABLE 1

Summary of DNA:DNA Reassociation of N. crassa total ³²P-DNA and of ³H-rDNA with total DNA of Neurospora species

Unlabeled DNA Fragments	with ³ H-rDNA		with ³² P-total DNA	
	Percent Reassociation (Normalized)	T _{e50} in °C	Percent Reassociation (Normalized)	T _{e50} in °C
Heterothallic				
<u>N. crassa</u> 74A (FGSC #987)	100	88	100	86
Homothallic				
<u>dodgei</u> (FGSC #1692)	97	86	65	82
<u>N. lineolata</u> (FGSC #1910)	95		63	81
<u>N. africana</u> (FGSC #1740)	92	86	64	82

The purified ³H-rDNAs (nick translated) of N. crassa, sheared to 400 nucleotide fragments had 1 x 10⁶ cpm (counts per minute) per microgram of DNA. The ³H-rDNA C₀t used for these reactions was 2 x 10⁻³, at which there was no detectable self reaction. The ³²P-DNA of N. crassa DNA, sheared to 400 nucleotides had 20,000 cpm/μg DNA. ³²P-DNA C₀t used was 0.05 and the 1-2 percent reaction obtained with ³²P-total was routinely deducted from the total DNA:DNA reassociation. In all reactions unlabeled DNA C₀t was at least 700. T_{e50} (50% dissociation) was determined from thermal stability curves.

The results of various DNA:DNA and rDNA:DNA reactions are summarized in Table 1. With total ³²P-DNA of N. crassa, it was impossible to detect DNA sequence differences among the three homothallic species, although differences between heterothallic and homothallic species were obvious. However, 2 to 5 percent differences in nucleotide sequence were observed when purified rDNA of N. crassa was reacted with the three homothallic species. These observations suggest the existence of non-identical rDNA sequences among different species of Neurospora. Whether these differences are in the spacer regions is now being investigated. (Supported in part by the U. S. Department of Energy). - Department of Botany and the Cancer Research Center, Howard University, Washington, D. C. 20059.