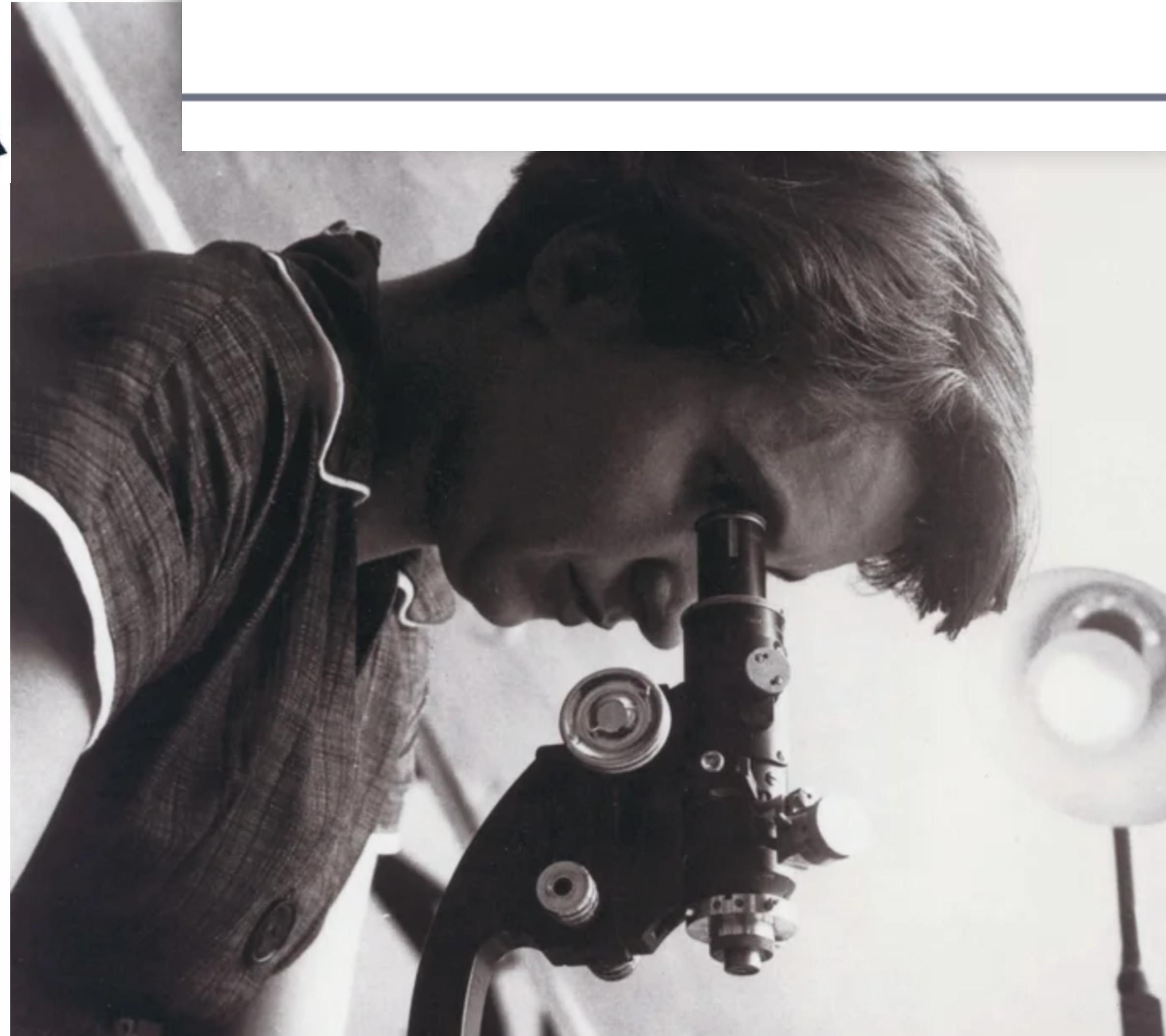




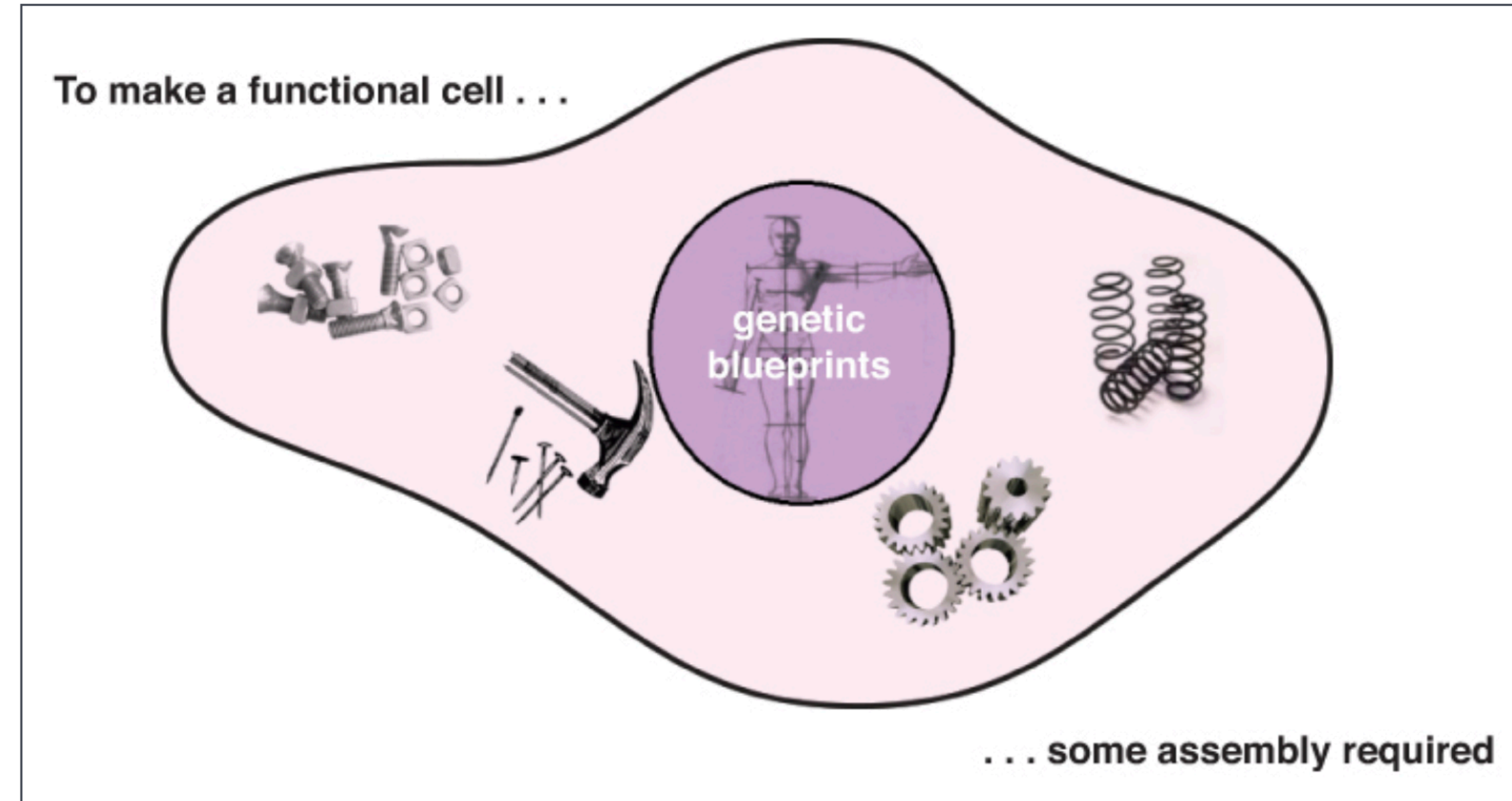
Team#1- Mit, Ryan, Nicole, Costa



Chapter 2: Central Dogma

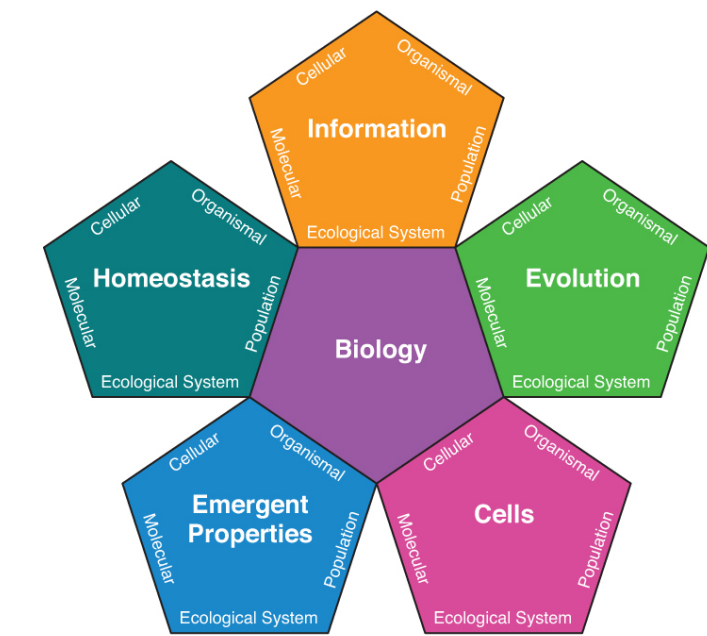
DNA contains instructions but each cell must construct its own components to perform a function. Original art.

What makes cells function as productive members of your body? How do cells control which genes are activated at different times or in different organs of your body? As you learned in Chapter 1, DNA is the molecular information passed down from generation to generation. However, DNA does not perform cellular functions, it only provides a molecular blueprint for cells. Proteins perform most cellular functions. Although the structure and function of DNA were understood by the late 1950's, researchers still did not understand how the different forms of RNA functioned. In this chapter, you will follow the path of researchers who made many ground-breaking discoveries about how cells produce proteins. This path will lead you to the fundamental concept of "central dogma," which explains how cells process molecular information from DNA to RNA to protein. It may surprise you to know that the path of discovery has not ended. Biologists of your generation will continue to make many important discoveries, building on the foundation of ideas you will learn in Chapter 2. The four Sections of Chapter 2 focus on information at the cellular level.



you are here		Big Ideas of biology				
		Information	Evolution	Cells	Homeostasis	Emergent Properties
levels of the biological hierarchy	molecules	1	4	7	10	13
	cells	2	5	8	11	14
	organisms I	3	6	9	12	15
	organisms II	16	19	22	28	25
	populations	17	20	23	29	26
	ecological systems	18	21	24	30	27

Integrating Concepts in Biology



Chapter 2: Central Dogma

2.1 How does DNA communicate information to the cell?

by A. Malcolm Campbell, Laurie J. Heyer, &
Christopher Paradise

Biology Learning Objective

- Describe the three major types of RNA and their functions.

What does RNA do for cells?

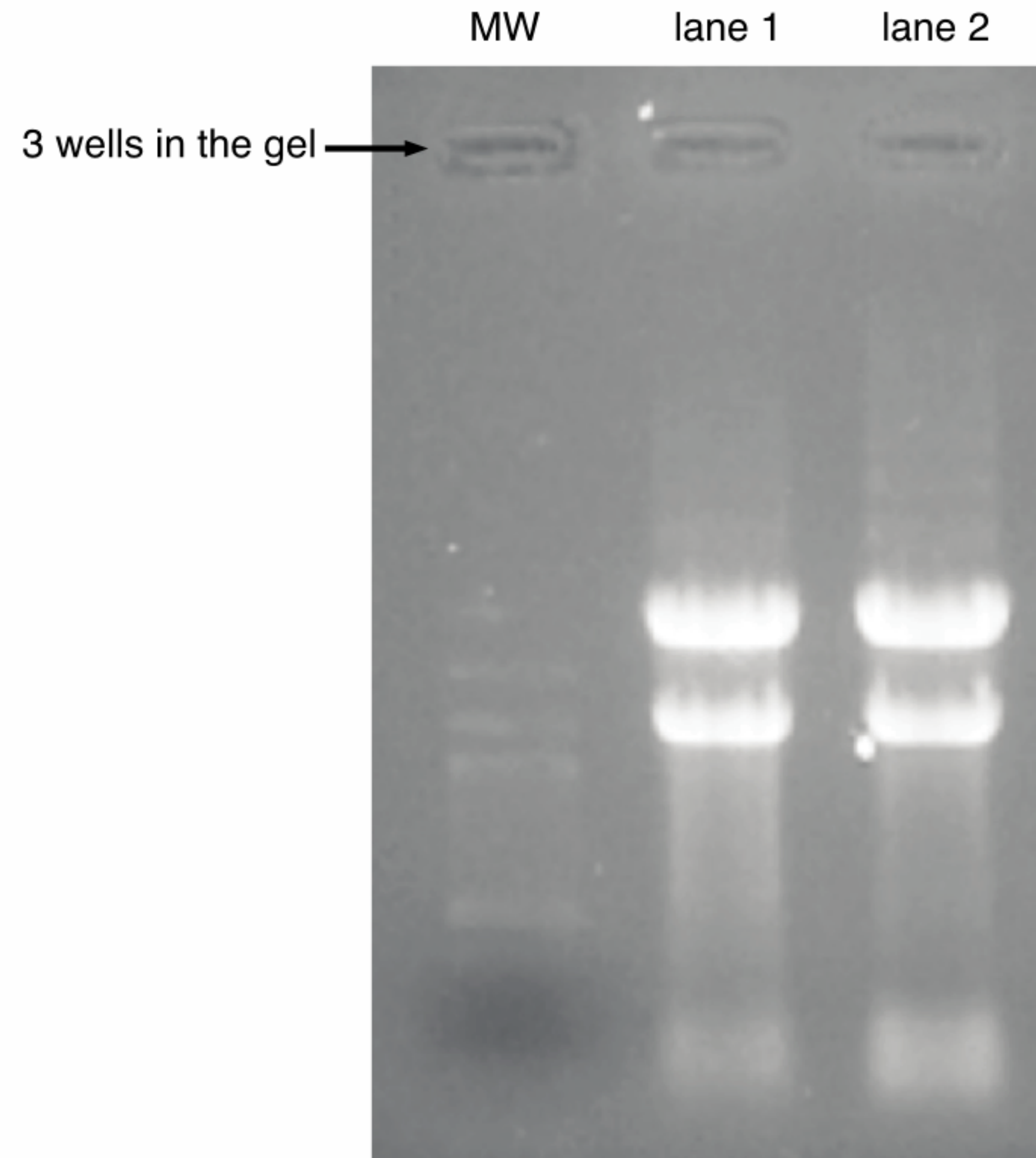


Fig. 2.3

Yeast RNA Separated by Size

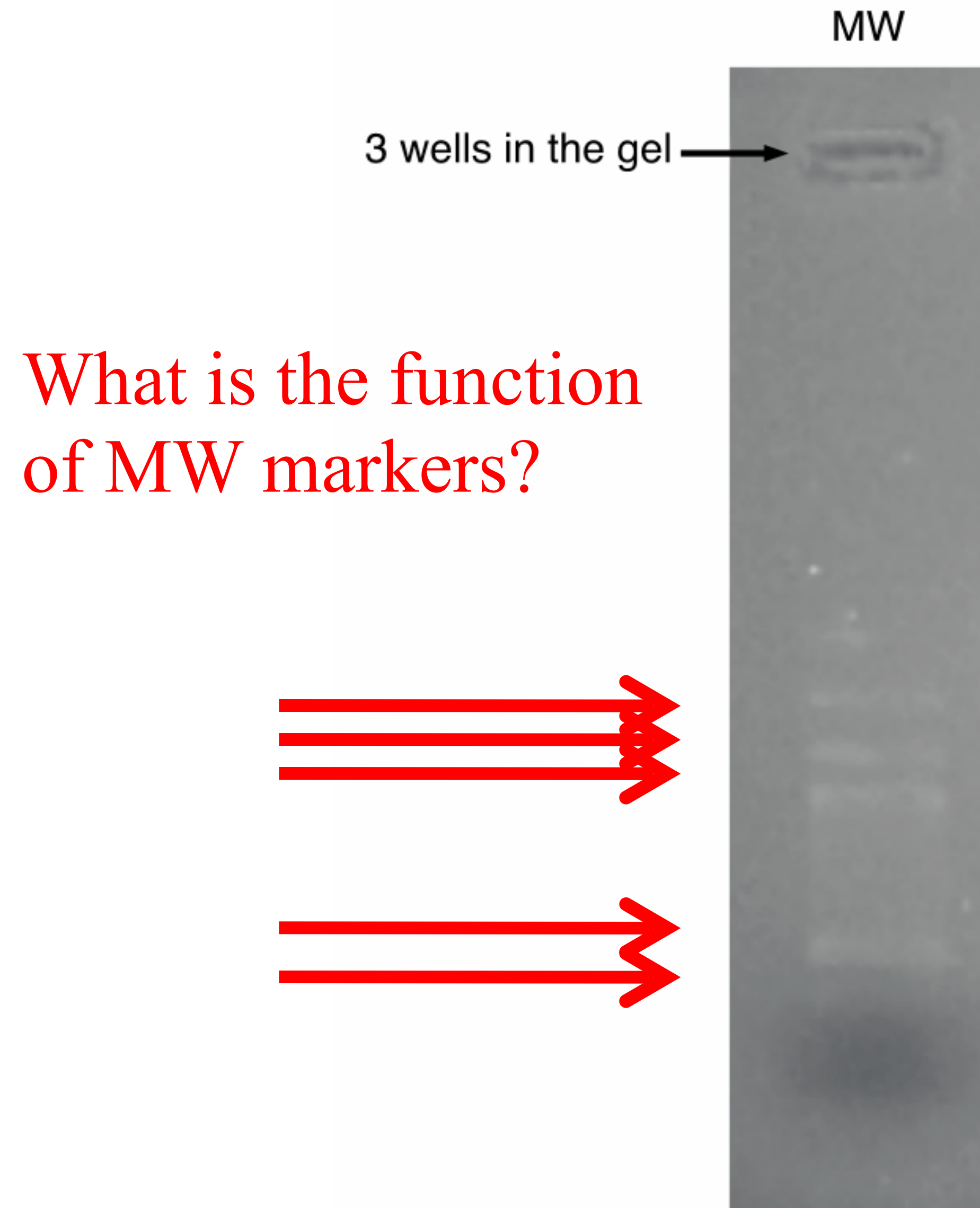


Fig. 2.3

Yeast RNA Separated by Size

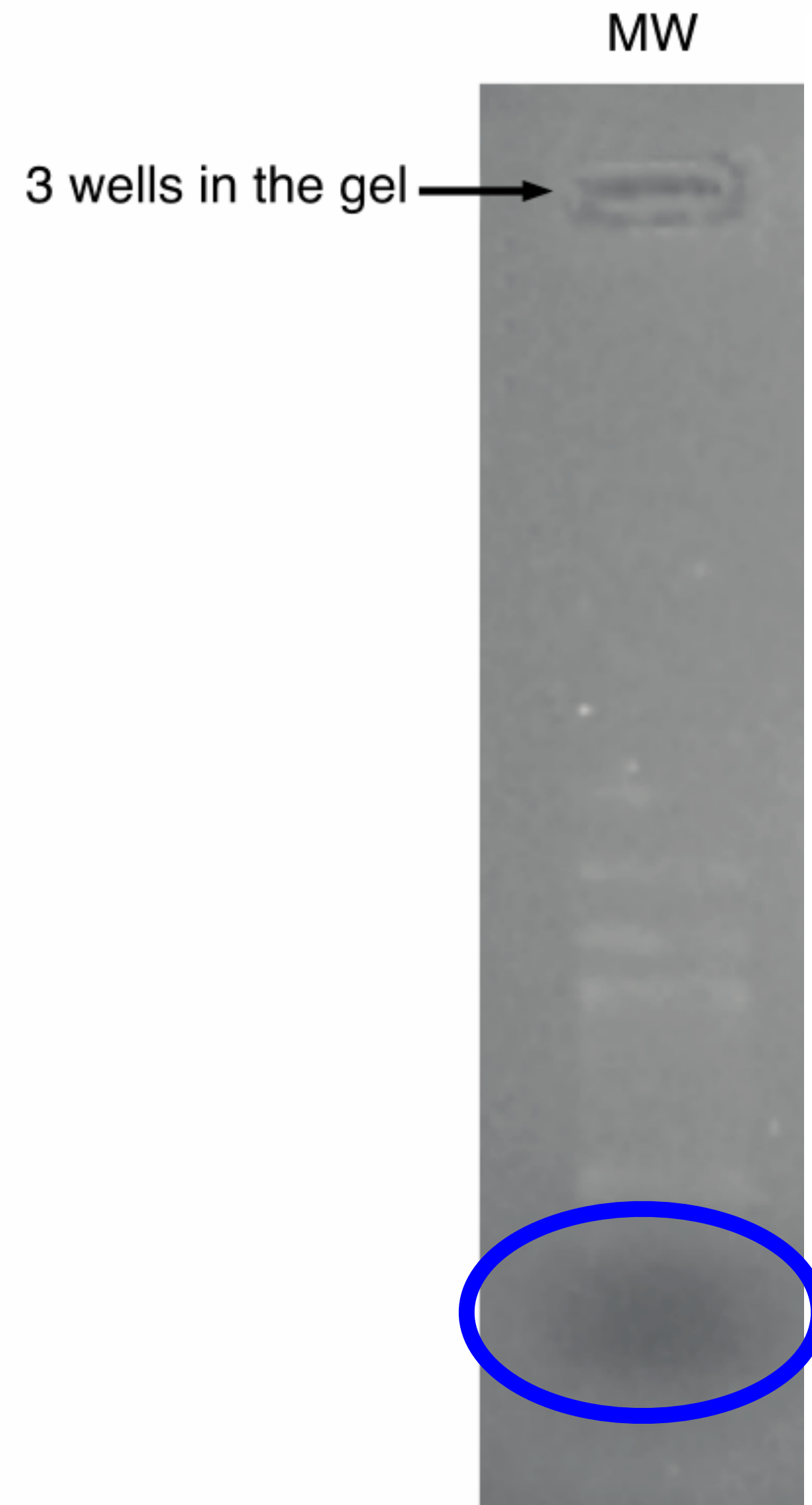


Fig. 2.3

Yeast RNA Separated by Size

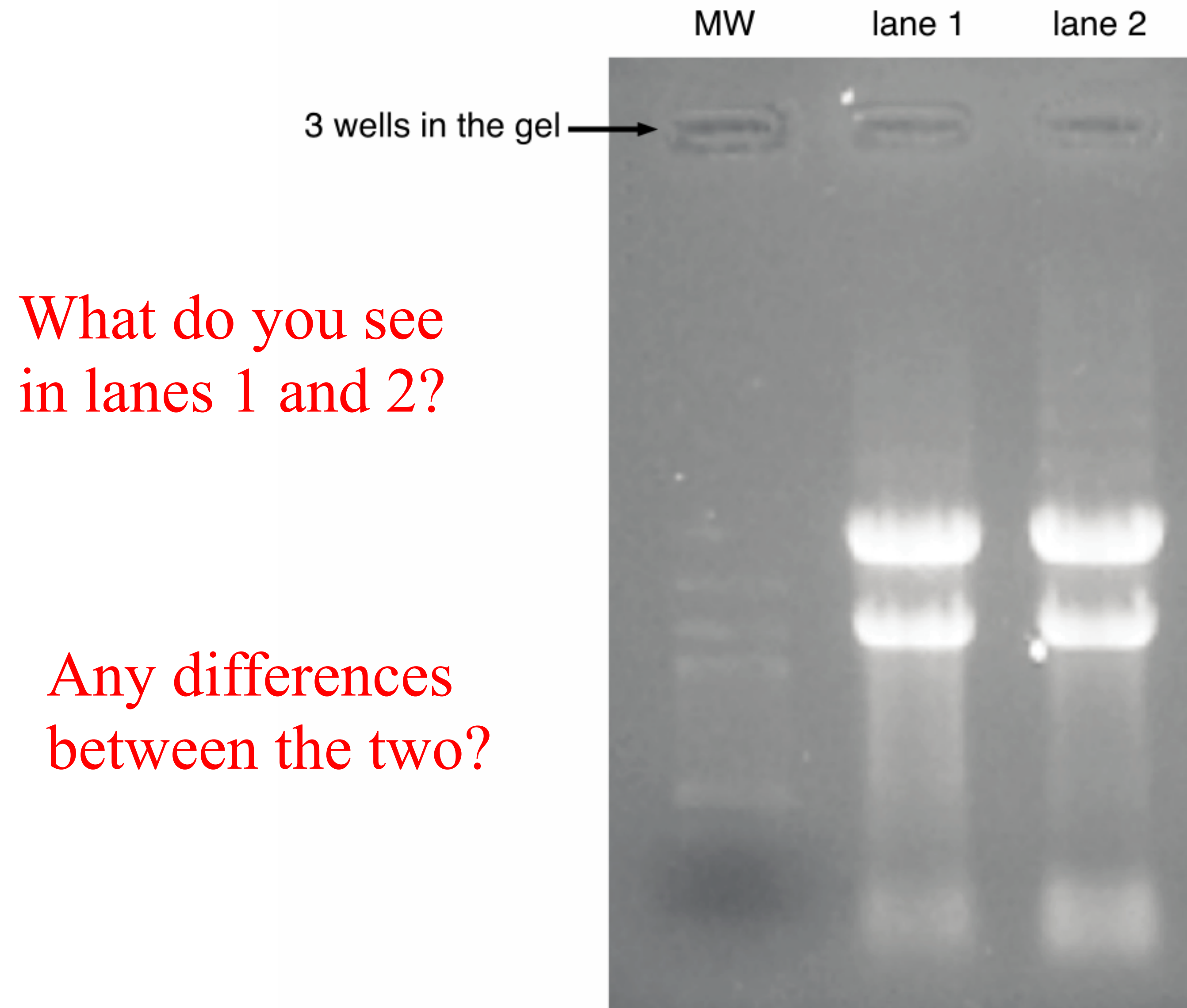


Fig. 2.3

Yeast RNA Separated by Size



Fig. 2.3

Yeast RNA Separated by Size

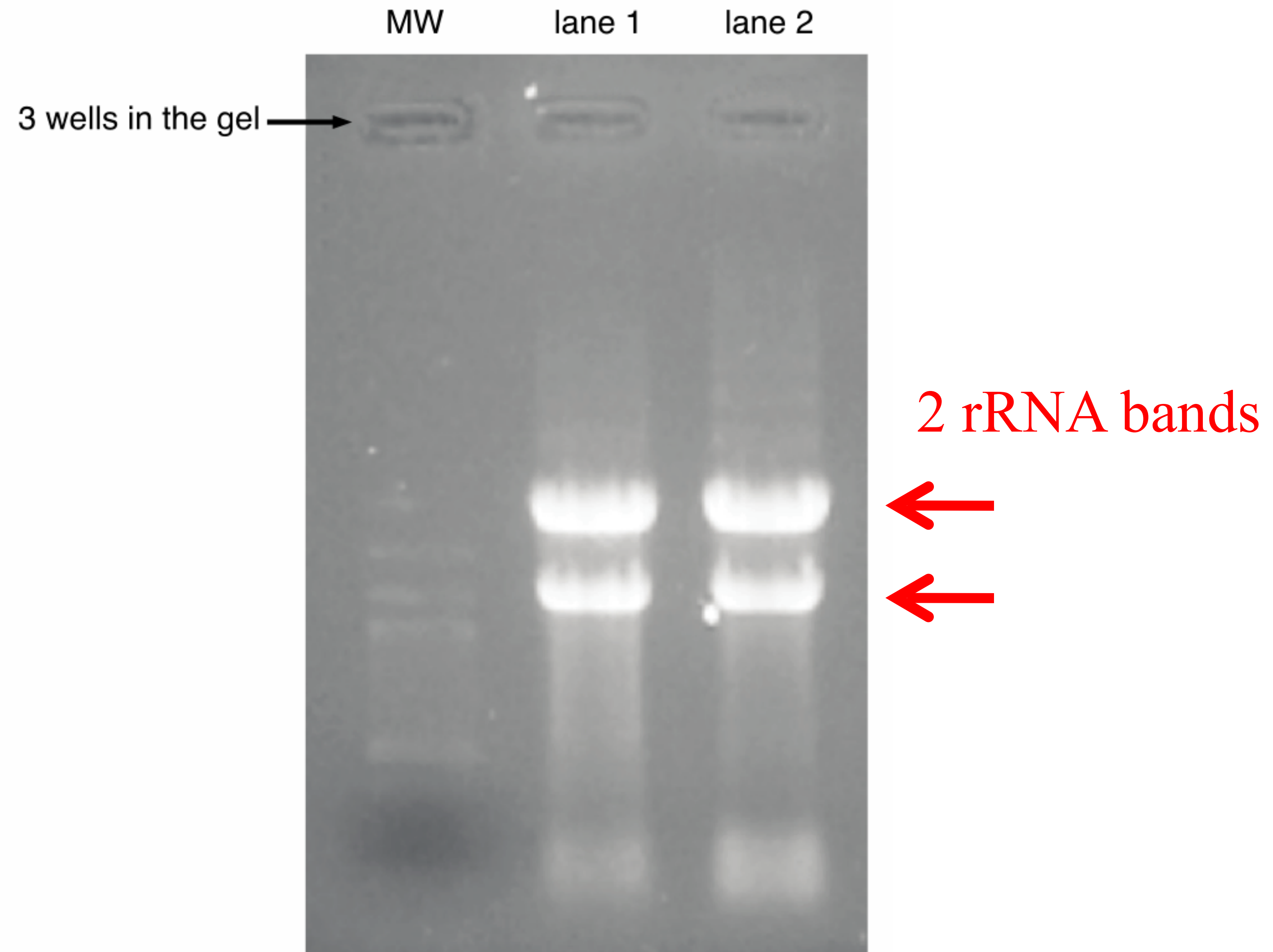


Fig. 2.3

Yeast RNA Separated by Size

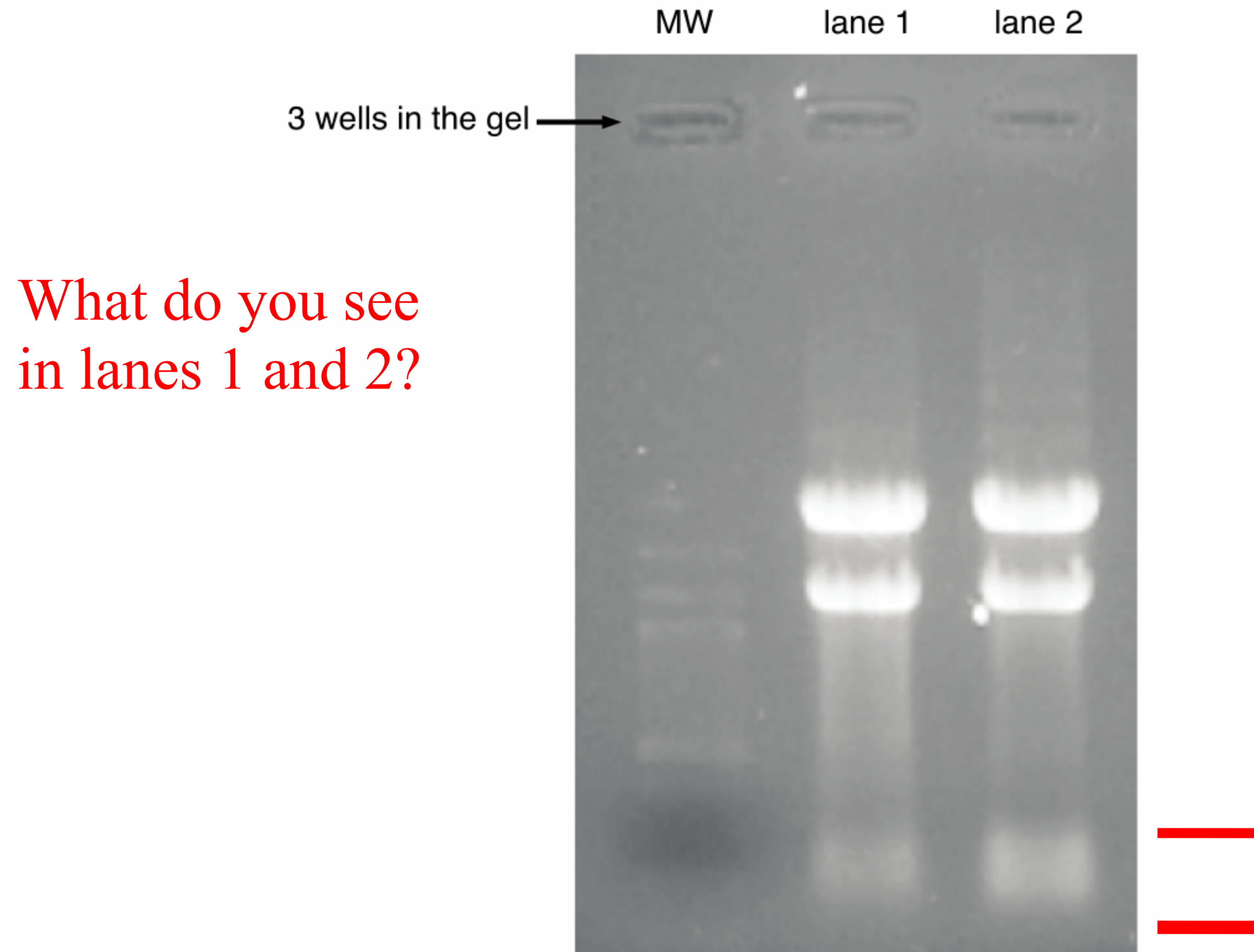


Fig. 2.3

Using logic and the existing data, it seemed clear to Brenner, Jacob, and Meselson that proteins could not be produced directly on DNA because proteins are produced in the cytoplasm and DNA is located inside the nucleus. rRNA is the most abundant form of RNA and accounts for about 80% of a cell's total RNA. Furthermore, the two sizes of rRNA molecules visible in Figure 2.3 lack the size and nucleotide sequence diversity required to produce the wide assortment of proteins found in any given cell. Data published by others indicated that when a cell is infected by a virus, the protein content of the cell changes within 10 minutes—such that nearly all of the new proteins in the infected cell are encoded by the virus genome and not the host genome. Therefore, protein-coding RNA needs to change very quickly, but rRNA in cells persists for a long time. In addition to the paired

rRNA molecules, you should see a category is visible as a cloud of RNA is spread nearly the entire length of the gel as a very faint smear, indicating the sizes range from very big to almost as small as the cloud at the bottom.

To further understand how DNA information is converted into proteins, you need to understand the second category of RNA. Look at the cloud of very small RNA molecules near the bottom of the gel in Figure 2.3. This band is **transfer RNA (tRNA)** and is critical for protein production. Accounting for about 18% of all RNA, tRNA was named for its function of *transferring* amino acids in the process of protein production. In 1964, a pair of investigators discovered the function of tRNA through experimentation (Figure 2.5). The investigators purified all of the RNA from some actively growing pea plants and incubated the RNA with millions of copies of radioactive amino acid leucine. They separated all of the RNA by size and measured the presence of RNA by ultraviolet (UV) absorption. Later, they detected the location of the radioactive leucine.

Trifecta: purpose, methods, findings

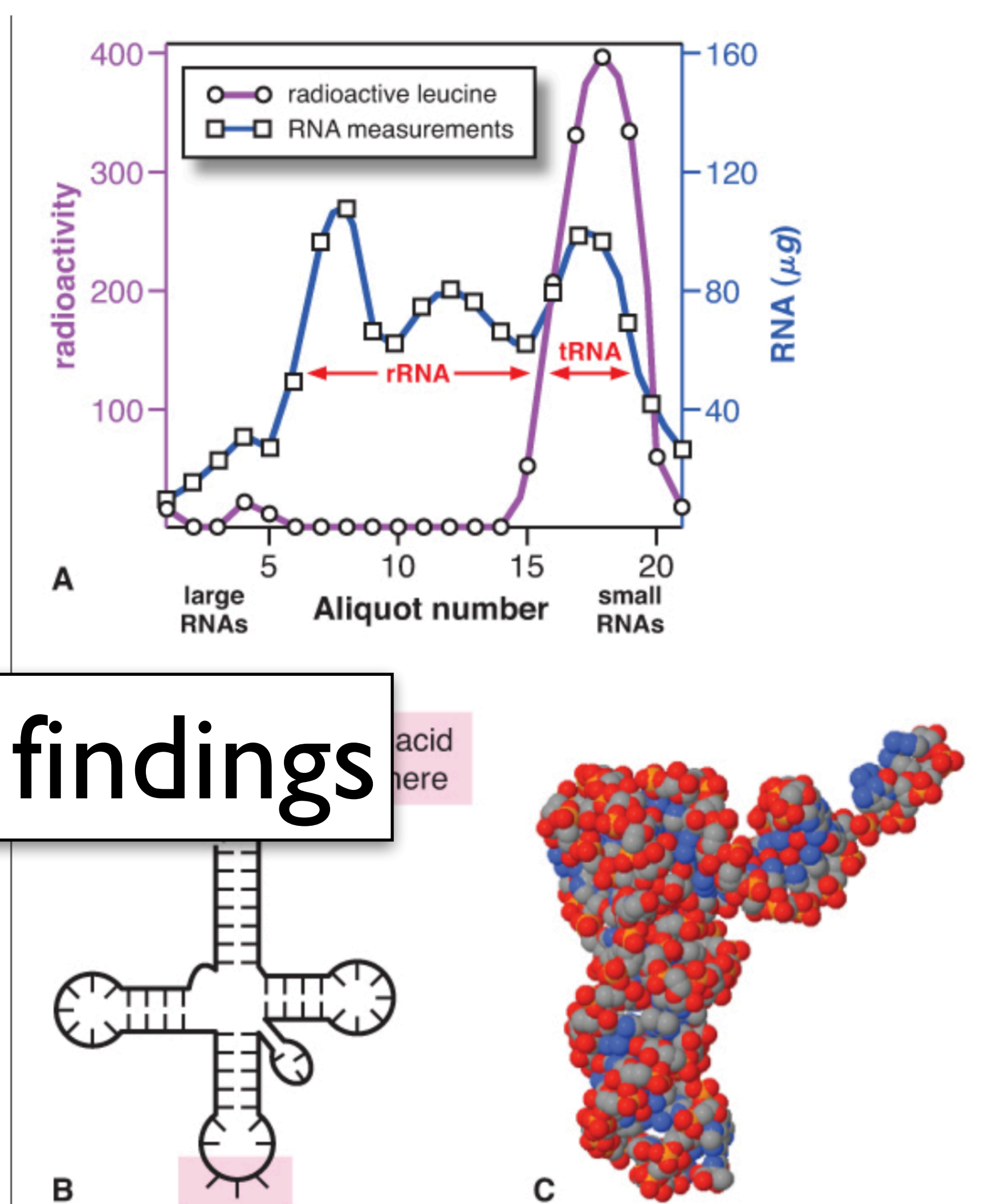


Figure 2.5 Structure and function of tRNA. **A**, RNA molecules interact with radioactive leucine. Total RNA from peas was extracted and incubated with radioactive leucine in the presence of cytoplasm from the peas. RNA molecules were separated by size. Radioactive leucine and non-radioactive RNA were detected separately as shown. **B**, Two-dimensional representation of tRNA as determined by base pairing within the molecule. **C**, Space filling three-dimensional image of tRNA; red, oxygen; blue, nitrogen; orange, phosphorus; and grey, carbon. Interactive

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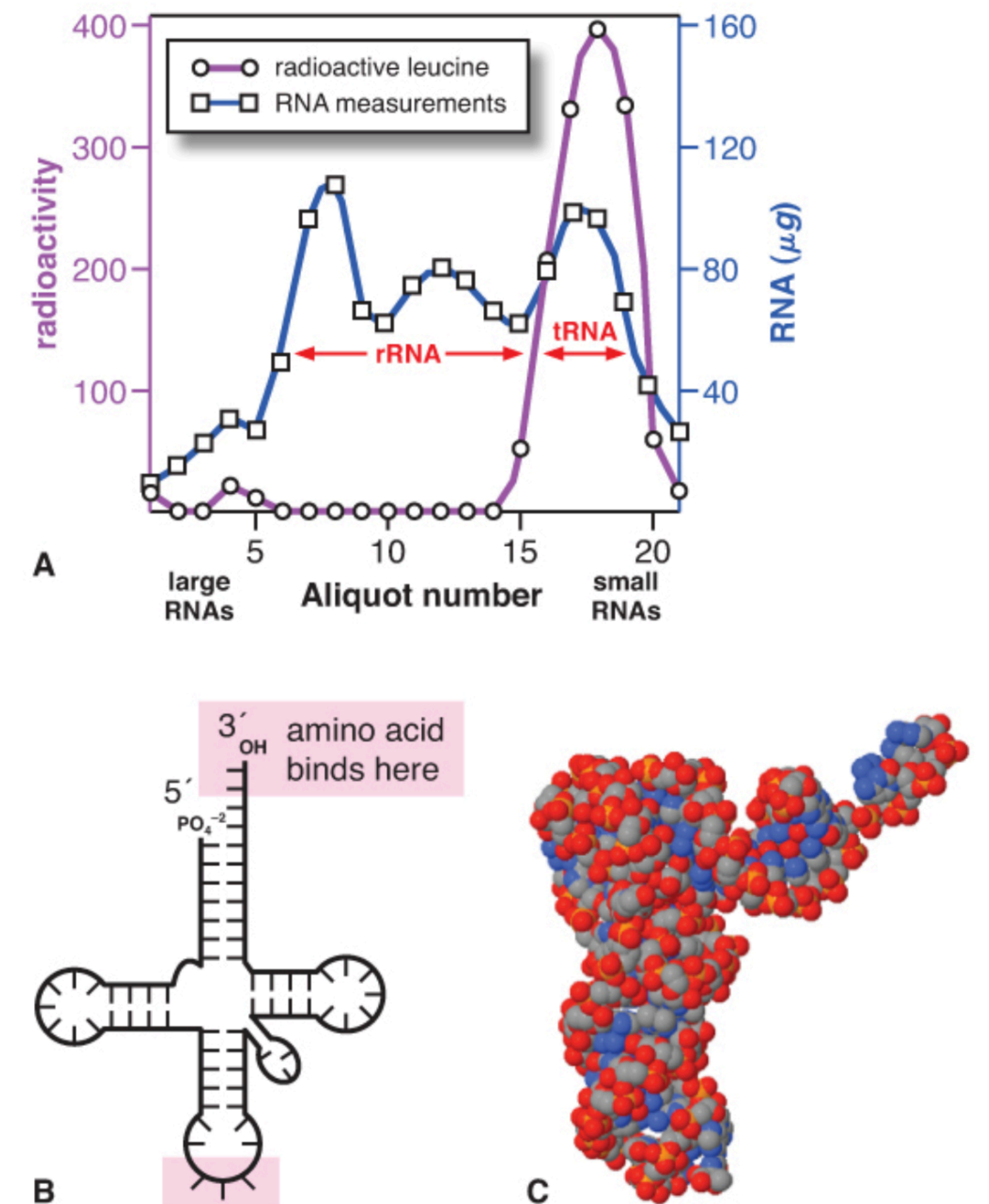


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05:00

Prepare to

Explain:

Purpose,

Methods,

Findings

(trifecta)

Trifecta?

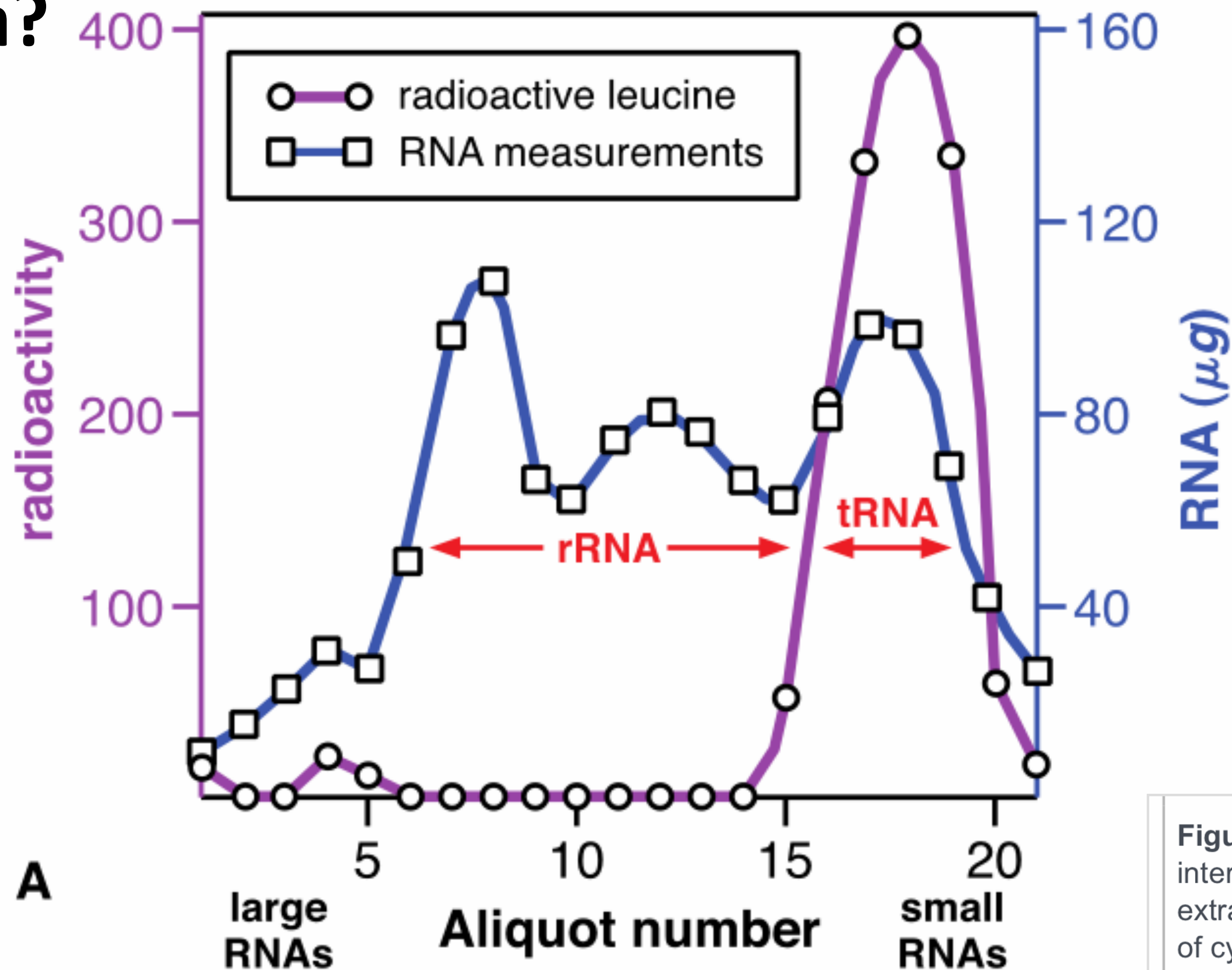


FIG. 2.—Size distribution of nuclear RNA and localization of transfer RNA on a sucrose density gradient. Sucrose density gradient (20 ml) of 20–5% sucrose in .01 M tris (pH 7.1) and 1% phenol was set up according to Bolton *et al.*¹⁷ and overlaid with an inverse gradient of approximately 1 mg RNA in a sucrose gradient of 4–0%. A covering layer of paraffin oil prevented tubes from collapsing and rendered gradients more stable for handling. The tubes were centrifuged 12–14 hr in the Spinco rotor No. 25 at 24,000 rpm, then pierced and 1 ml fractions collected. Each fraction was precipitated with ethanol/acetate (66/2%), thoroughly washed to remove the phenol, and then tested for C¹⁴-leucine incorporation as described in Table 1.

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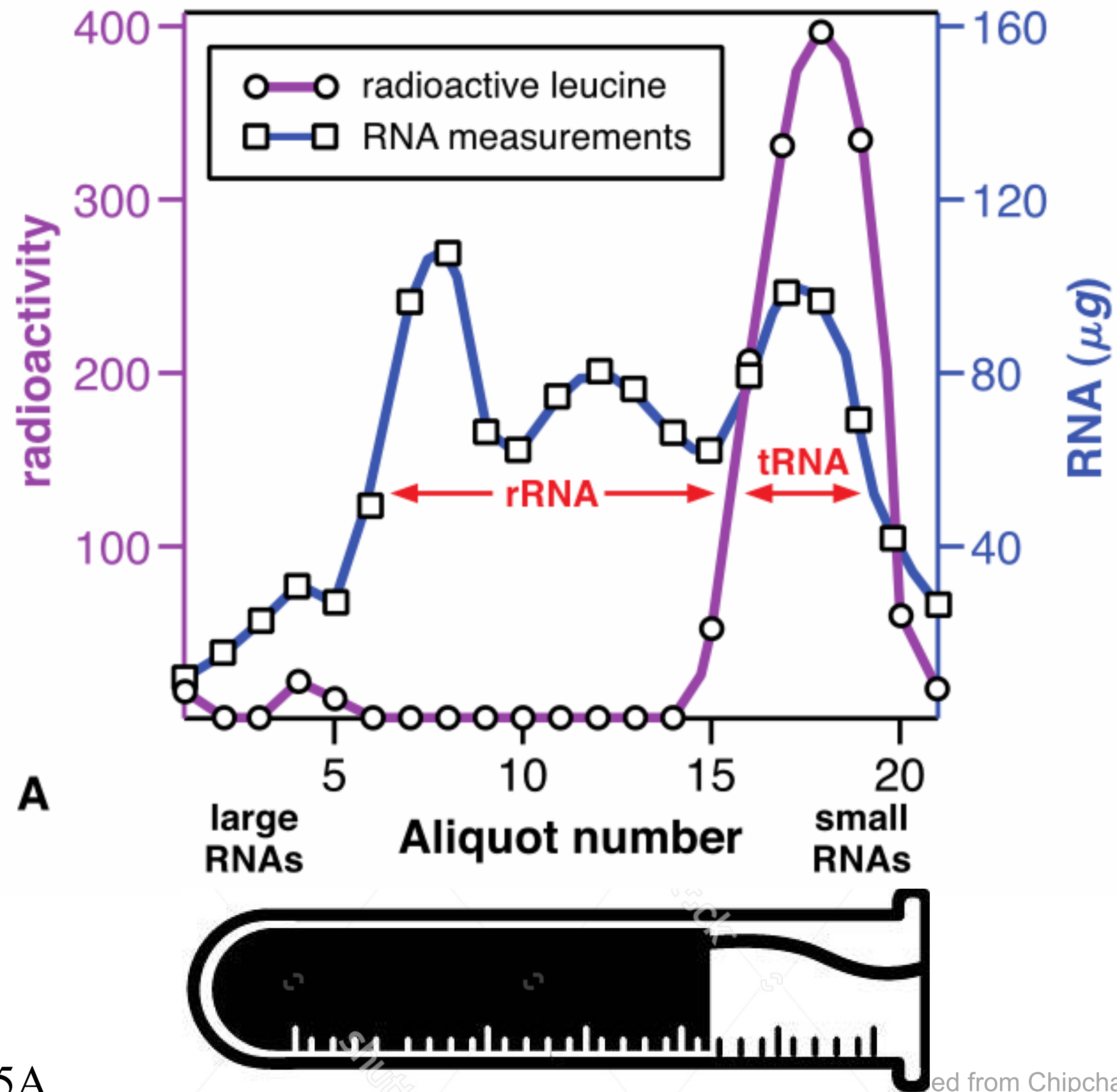


Figure 2.5A

SYNTHESIS OF TRANSFER RNA BY ISOLATED NUCLEI*

BY MARGARET I. H. CHIPCHASE AND MAX L. BIRNSTIEL

DIVISION OF BIOLOGY, CALIFORNIA INSTITUTE OF TECHNOLOGY

Communicated by James Bonner, March 15, 1963

In the course of earlier experiments on the incorporation of labeled nucleosides into RNA by isolated nuclei¹ we observed that much of the newly synthesized RNA is soluble in 1 *M* NaCl, as is transfer RNA.²⁻⁴ Sirlin has reported the incorporation of pseudo-uridine into nuclei, allegedly into transfer RNA,^{5, 6} and the presence of amino-acyl RNA in thymus nuclei has been shown by Hopkins.⁷ Since the pea nuclei with which we work are capable of protein synthesis,⁸ they might therefore be suspected of containing transfer RNA. It will be shown below that isolated pea nuclei not only contain, but possess the ability to synthesize, transfer RNA.

Materials and Methods.—Analytical reagent grade chemicals were used throughout. ATP, CTP, GTP, UTP, UMP, uridine, phosphocreatine, and crystalline DNase were obtained from Sigma. Creatine phosphokinase was obtained from the California Corp. for Biochemical Research. Sodium penicillin-G was a gift of Chas. Pfizer and Co., New York. 2-hydroxy-3-naphthoic

phosphate system.¹³ Alternatively, an ammonium sulfate fractionation was used. Nine volumes of ice-cold 2.5 *M* ammonium sulfate, pH 5, were added to the aqueous solution to yield a final concentration of 2.25 *M*.¹⁴ This solution was kept at 0°C for 10 min and the fine precipitate then centrifuged down at 35,000 × *g* for 15 min. This procedure precipitates approximately 90% of the dye-bound RNA¹⁴ leaving the dye-non-bound amino-acyl RNA in solution. The precipitate of dye-bound RNA was washed once with 5% TCA, and then twice with 70% ethanol + 0.5% sodium acetate pH 5 and once with absolute ethanol. The dye-non-bound RNA was precipitated from the dialyzed sulfate solution by the addition of TCA to give a final concentration of 5%, the mixture kept at 0°C for 10 min, and the precipitate sedimented at 35,000 × *g* in the Servall for 15 min. The precipitate was washed as above. Each precipitate was dissolved in distilled water, 560 mμ and 260 mμ absorptions determined, and an aliquot counted.

Experimental Results.—That nuclei contain an active transfer RNA is shown by the following experiment: isolated pea stem nuclei were incubated and extracted as described in the legend of Figure 1. The data of Figure 1 show that labeled

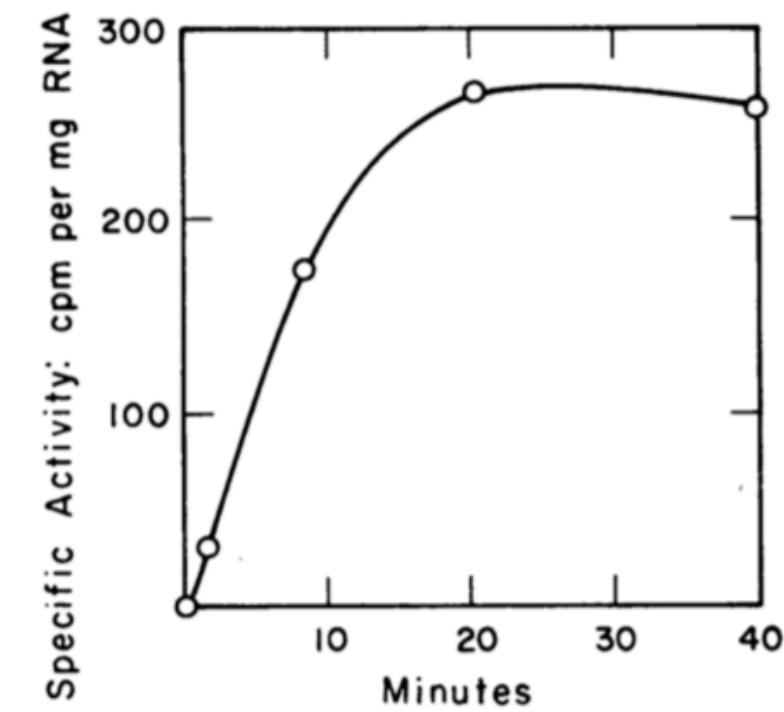


FIG. 1.—Formation of amino-acyl RNA in isolated nuclei. Incubation mixture: ATP, CTP, GTP, and UTP, .0001 *M* each; tris .02 *M*; phosphocreatine .02 *M*; creatine phosphokinase 100 μg/ml; CaCl₂ .003 *M*; MgCl₂ .0001 *M*; C¹⁴-protein hydrolysate 2 μc/ml; final pH 7.0, incubation at 37°C. Aliquots were precipitated at intervals by addition of an equal volume of phenol. RNA was extracted⁹ and washed as described in Table 1.

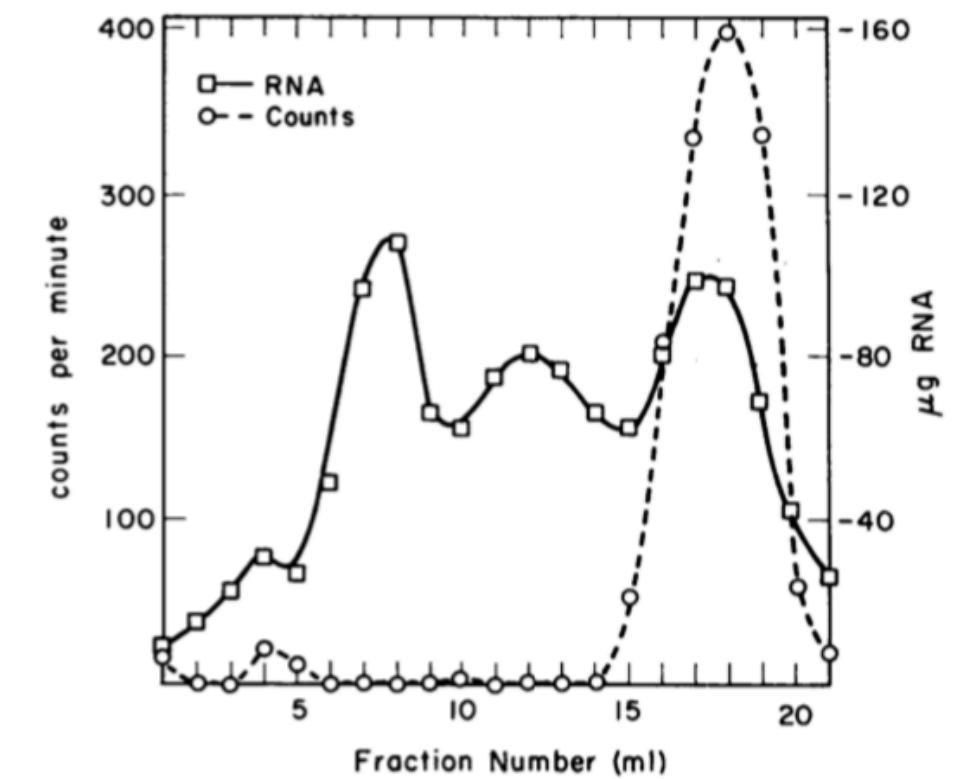


FIG. 2.—Size distribution of nuclear RNA and localization of transfer RNA on a sucrose density gradient. Sucrose density gradient (20 ml) of 20–5% sucrose in .01 *M* tris (pH 7.1) and 1% phenol was set up according to Bolton *et al.*¹⁷ and overlaid with an inverse gradient of approximately 1 mg RNA in a sucrose gradient of 4–0%. A covering layer of paraffin oil prevented tubes from collapsing and rendered gradients more stable for handling. The tubes were centrifuged 12–14 hr in the Spinco rotor No. 25 at 24,000 rpm, then pierced and 1 ml fractions collected. Each fraction was precipitated with ethanol/acetate (66/2%), thoroughly washed to remove the phenol, and then tested for C¹⁴-leucine incorporation as described in Table 1.

amino acids are incorporated into nuclear amino-acyl RNA.

That nuclear RNA can also bind amino acids *in vitro* is shown by the results of

Which RNA tells ribosomes what to make?

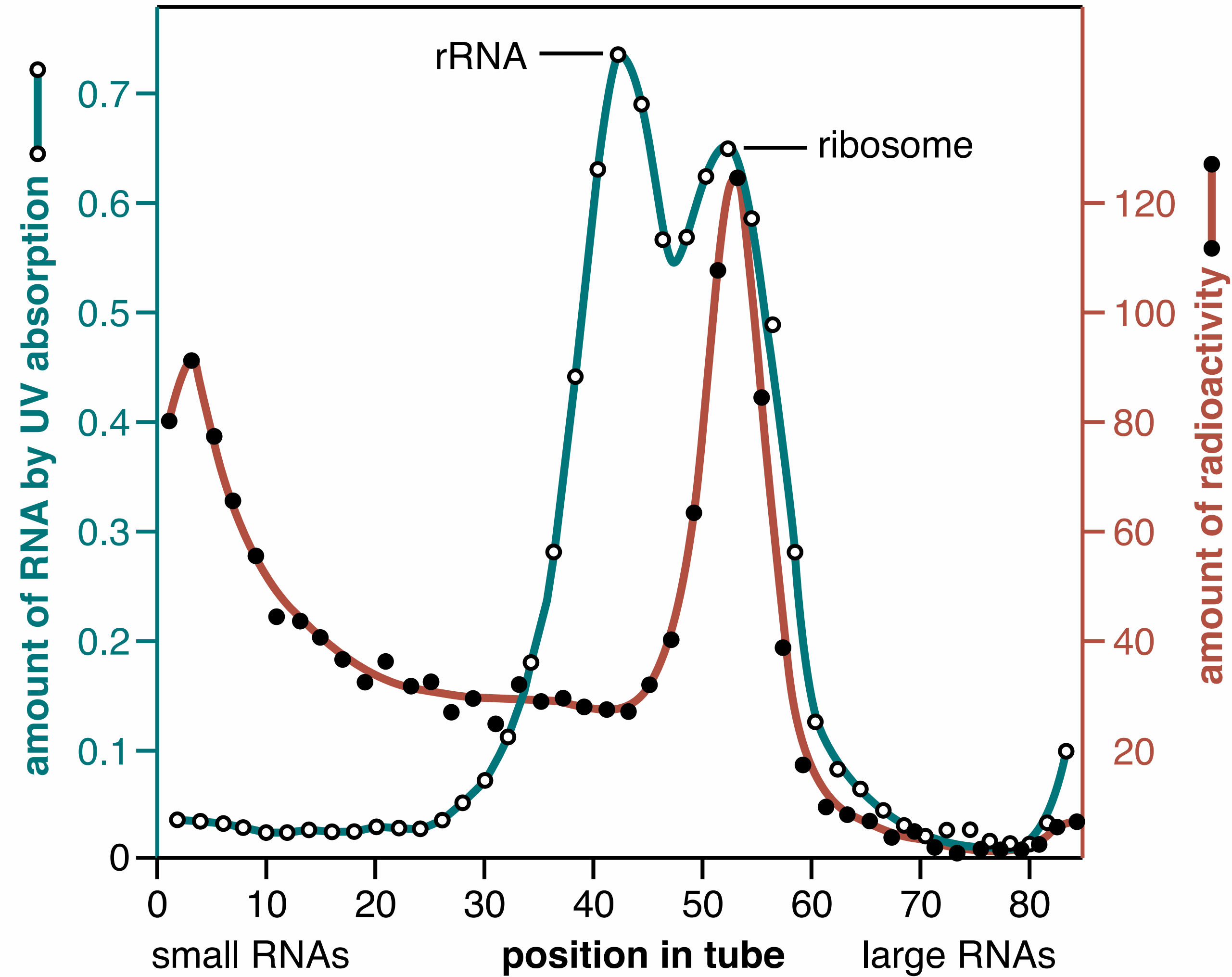


Fig. 2.6



Team#2- Annamaria, Paul, Michael, Ally





Team#1- Mit, Ryan, Nicole, Costa

VOTE

Team#2- Annamaria, Paul, Michael, Ally