# INTRODUCTION TO BIOINFORMATICS

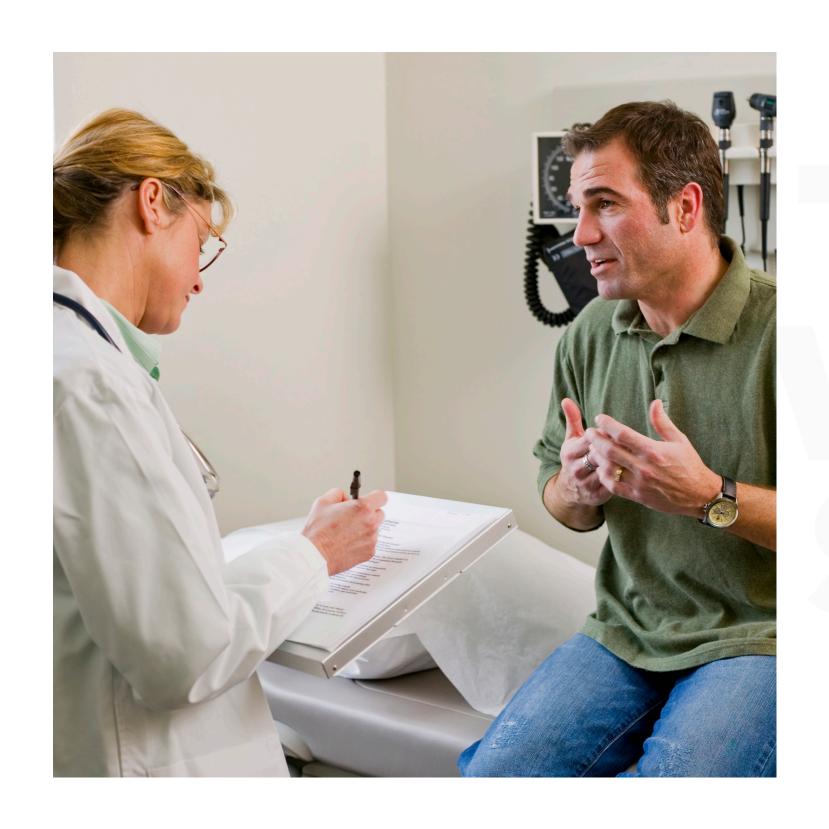




#### PICTURE THIS!

Many patients have been coming into your clinic over the last 3 weeks. In fact, the number appears to double each week. Help our doctors understand the disease and use your judgment to decide on a treatment.





A patient named Randy comes into the office with complaints of fever, headaches, and sore throat. You suspect a lung infection, but further tests are required.

## Start by reviewing Randy's info and filling in his Patient Chart.



PATIENT NAME: Randy Lynee

WEIGHT: 146

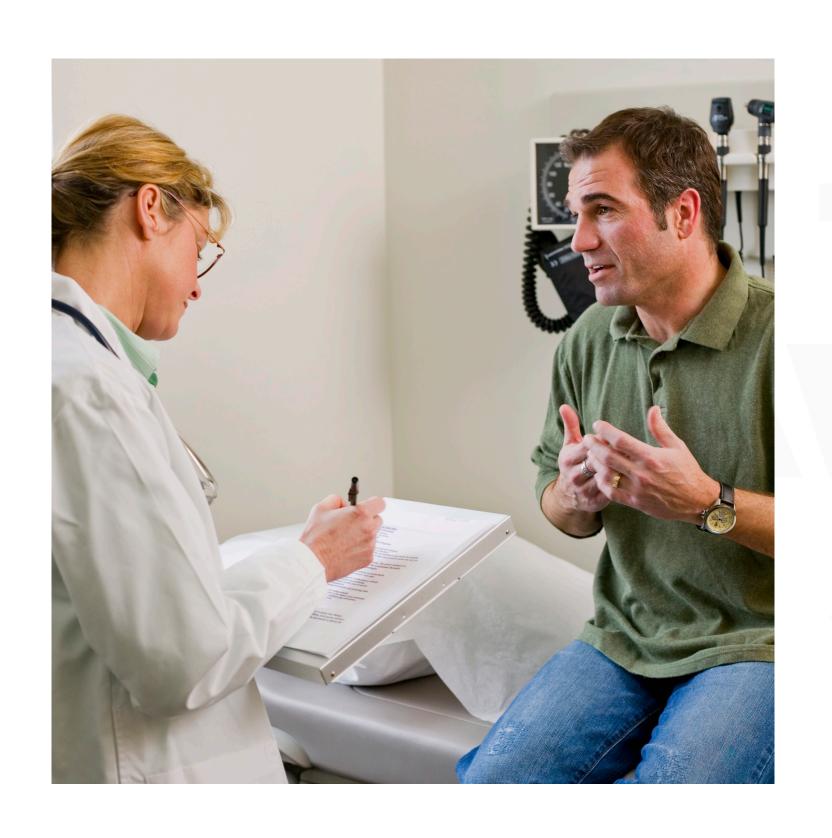
BLOOD PRESSURE: HEART RATE: 150 bpm

KEY SYMPTOMS:

-102 Dearce Temperature

- 102 Degree Temperature
- Sore throat
- Headache

INITIAL DIAGNOSIS: LUNG Infection



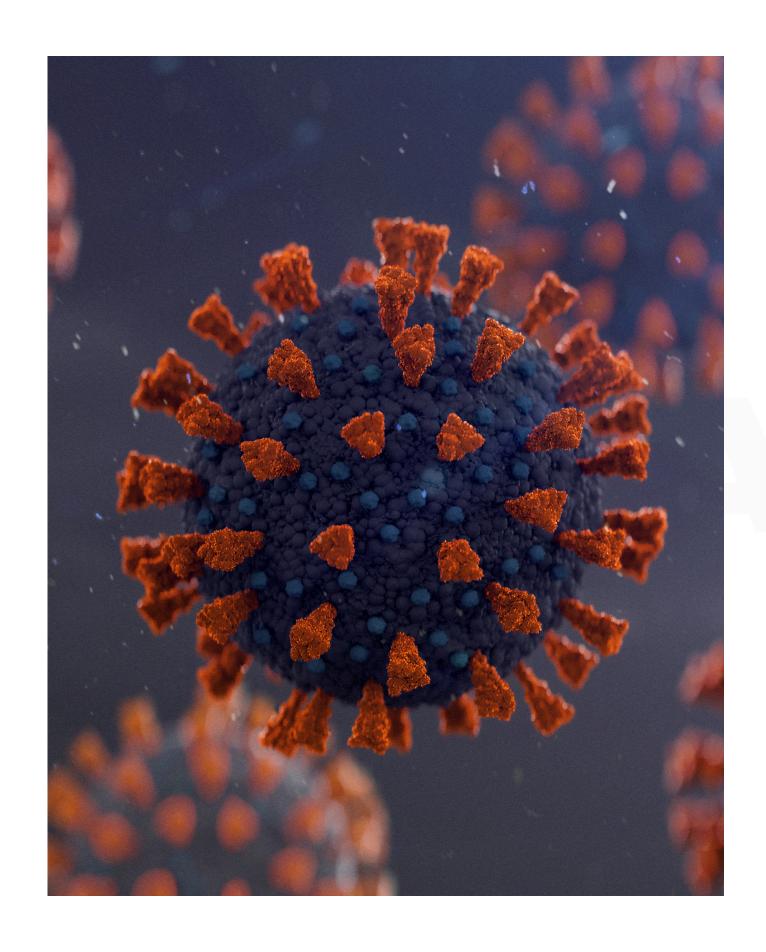
### ORDER TESTS

To diagnose Randy, you order a radiograph. You also administer antimicrobial treatment for three days.

#### TEST RESULTS

Antimicrobial Treatment: Irregular no reduction of symptoms in three days

Radiograph: Irregular evidence of pneumonia



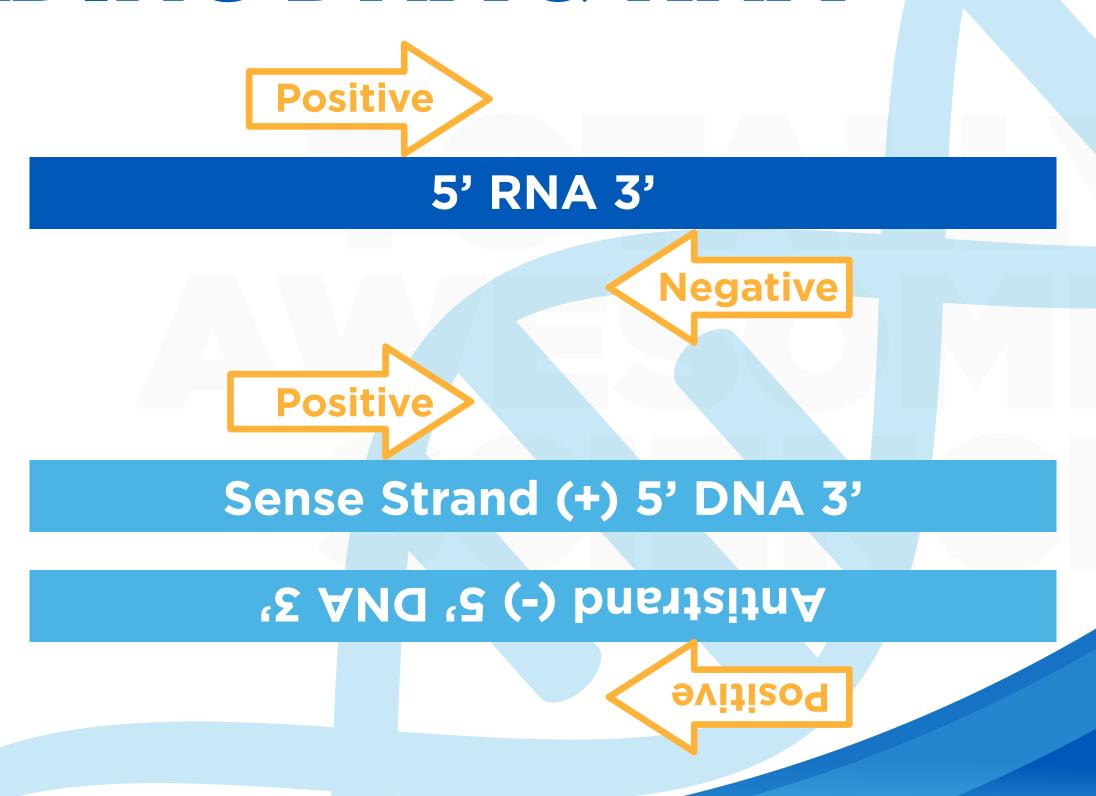
#### VIRUS REPORT!

#### Unidentified virus was

- RNA
- Single stranded
- Positive Sense
- Enveloped
- 29881 Base pair length (RNA bases)
- 9860 Amino
   Acid encoded

### READING DNA & RNA

Like words on a page are read from left to right, DNA is read in a specific direction.



#### READING DNA & RNA

#### Sense (Positive sense)

Words are usually read from left to right

#### Sense (Negative sense)

Thgir ot tfel morf dear yllausu era sdrow

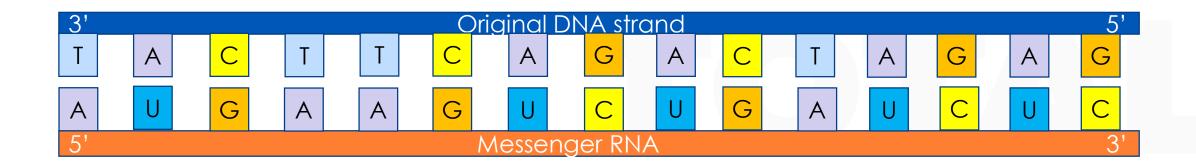


#### Sense strand

Words are usually read from left to right Morps era usually read from left to right with the Morps era usually read from left to right with the Morps erand

**Antisense strand** 

#### TRANSCRIPTION RULES



In DNA the base pairing rules include Adenine-Thymine; Guanine-Cytosine. RNA contains Uracil instead of Thymine.

## The pairing is as follows:

- Adenine → Uracil
- Thymine → Adenine
- Cytosine → Guanine
- Guanine → Cytosine

#### TRANSCRIPTION CAN TAKE MANY FORMS

- DNA TO RNA -



Polymerase

RNA

DNA

#### TRANSCRIPTION CAN TAKE MANY FORMS

- RNA TO DNA -

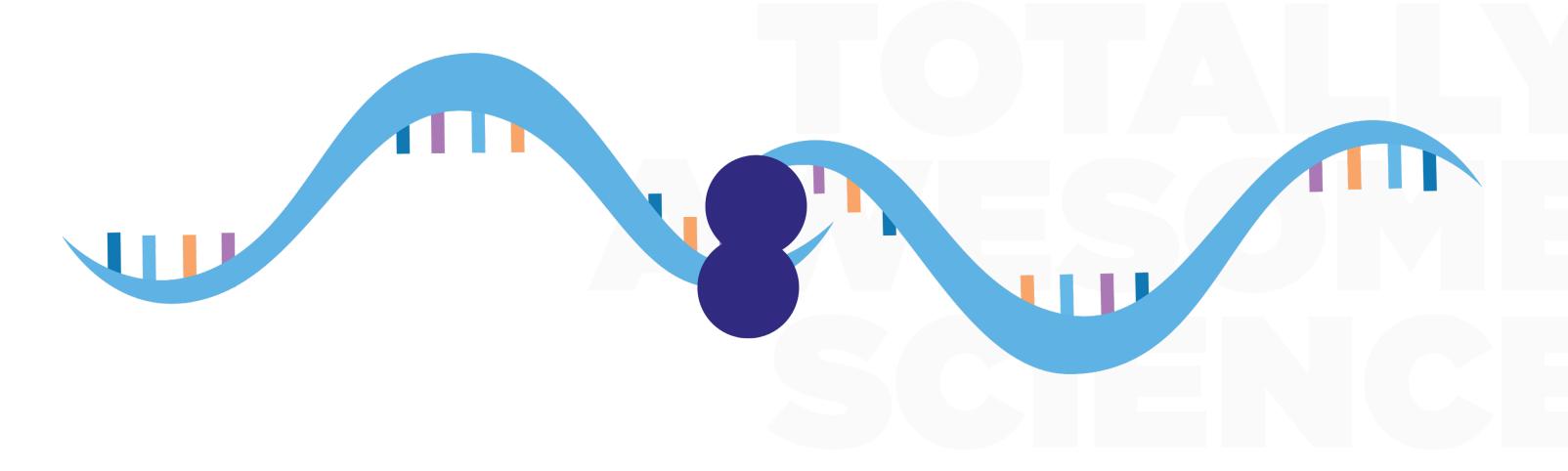






#### TRANSCRIPTION CAN TAKE MANY FORMS

- RNA TO RNA -







#### TRANSLATION

Segments of DNA that hold the code to make a protein are called genes.

Each gene can make an average of three proteins.

What human proteins do you know about?



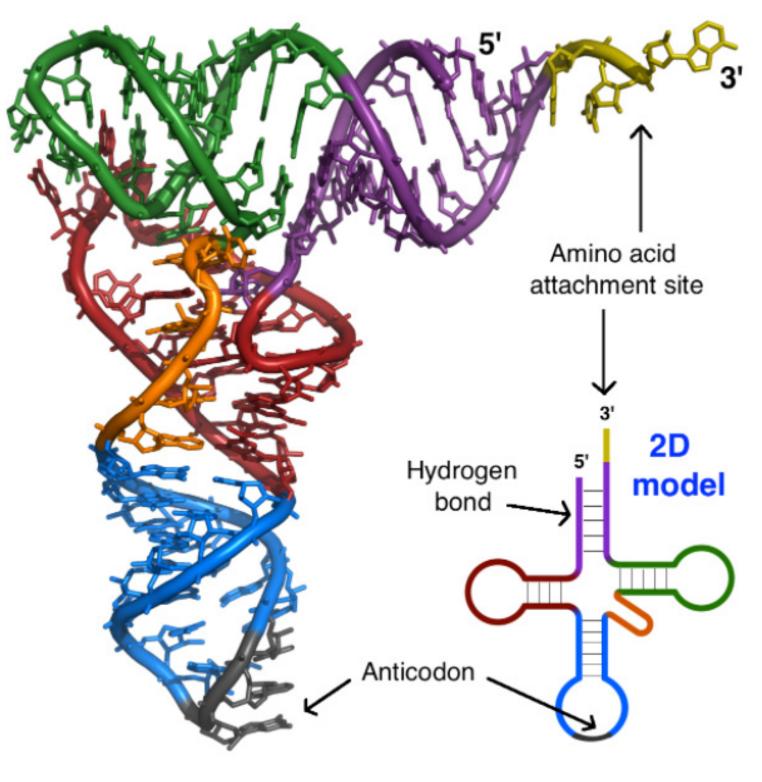




A protein with 4 chains that carries oxygen for your red blood cells

A protein that is found in skin cells that help protect your cells from UV radiation

An enzyme that helps your digestive system to break down lactose-the sugar found in milk



\_Image modified from "TRNA-Phe yeast," by Yikrazuul (CC BY-SA 3.0). The modified image is licensed under a CC BY-SA 3.0 license.\_

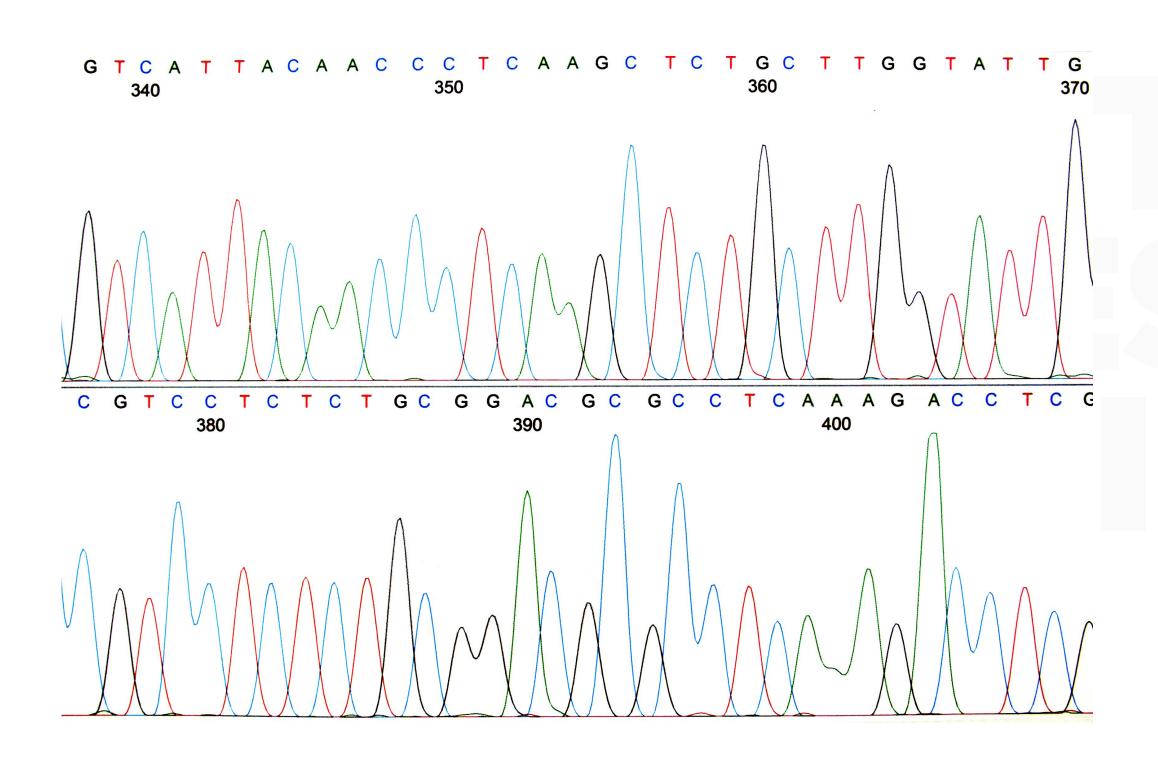
In order to change RNA code to amino acid sequence, a "translator" is needed. This is the function of transfer RNA.

tRNA contains an anticodon on one end and an amino acid on the other.

### WHAT IS NEW ABOUT THIS VIRUS?

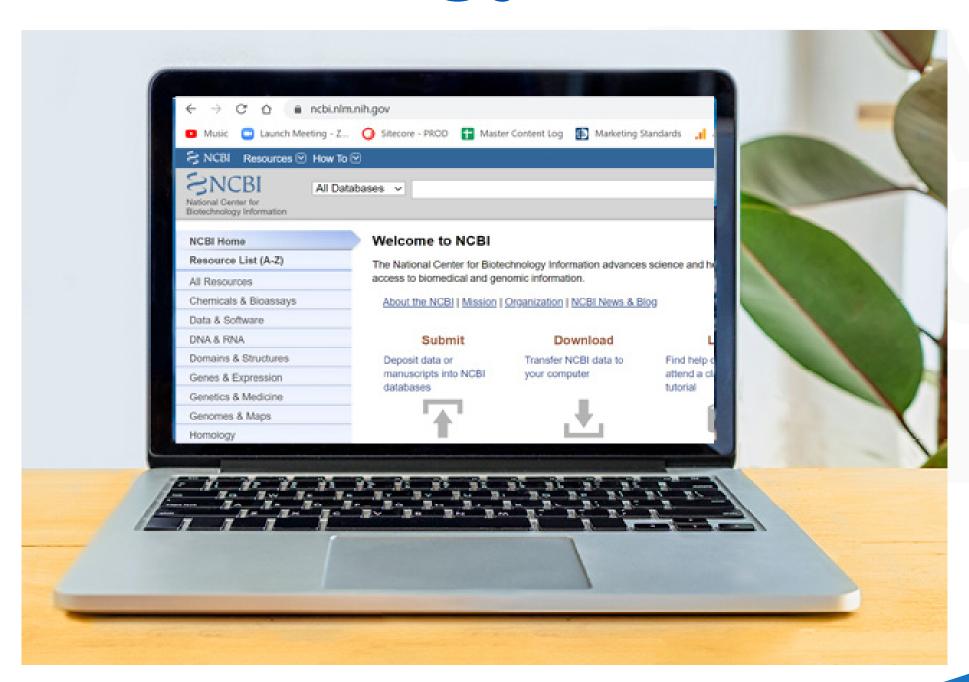


## VIRUS SEQUENCED!



The mysterious virus has been identified through a database search. It was found to be similar to SARS-Cov-2.

## National Center for Biotechnology Information



### WATCH THIS!



ne.	PROMI

#### **INVESTIGATING VIRAL PROTEIN**

#### BACKGROUND

1. What do scient

2. This virus does considered living

3. Where do new

#### **RESEARCH:**

SARS-CoV-2 produced and the following about the function below and collections.

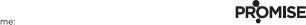
E ENVELOPE PR https://tinyurl.com/y

#### NSP1

https://tinyurl.com/y

- 1. My protein is
- 2. My protein has
- 3. My protein is
- 4. My protein app
- 5. What is the fu

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#### ANALYSIS OF CORONAVIRUS EVOLUTION

SANF RD

SANF: PRD

Coronaviruses (CoV), like Severe Acute Respiratory Syndrome (SARS-CoV), are positive sense RNA viruses that have been circulating from animal populations into human populations for many years. Researchers collected complete genome sequences from several populations of viruses and constructed a phylogenetic tree to represent their evolutionary relatedness (Figure 1).

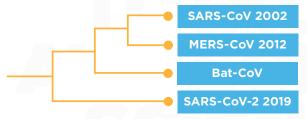


Figure 1. Phylogenetic tree representing evolutionary relatedness among corona virus strains based on whole viral genomes sequence comparison

A researcher studying adaptations in CoV sequenced the spike protein (S) in the human coronaviruses SARS, MERS, SARS-CoV-2, and Bat coronavirus. There are several substitutions found only in the SARS-CoV.

	Bat-CoV	MERS-CoV	SARS-CoV	SARS-CoV-2
Bat-CoV	-			
MERS-CoV	61.16%	-		
SARS-CoV	21.89%	58.76%	-	
SARS-CoV-2	2.12%	61.37%	22.53%	-

Table 1. Percentage of differences in the spike protein among CoV species

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# DISCOVER MORE ON VIRAL PROTEINS!

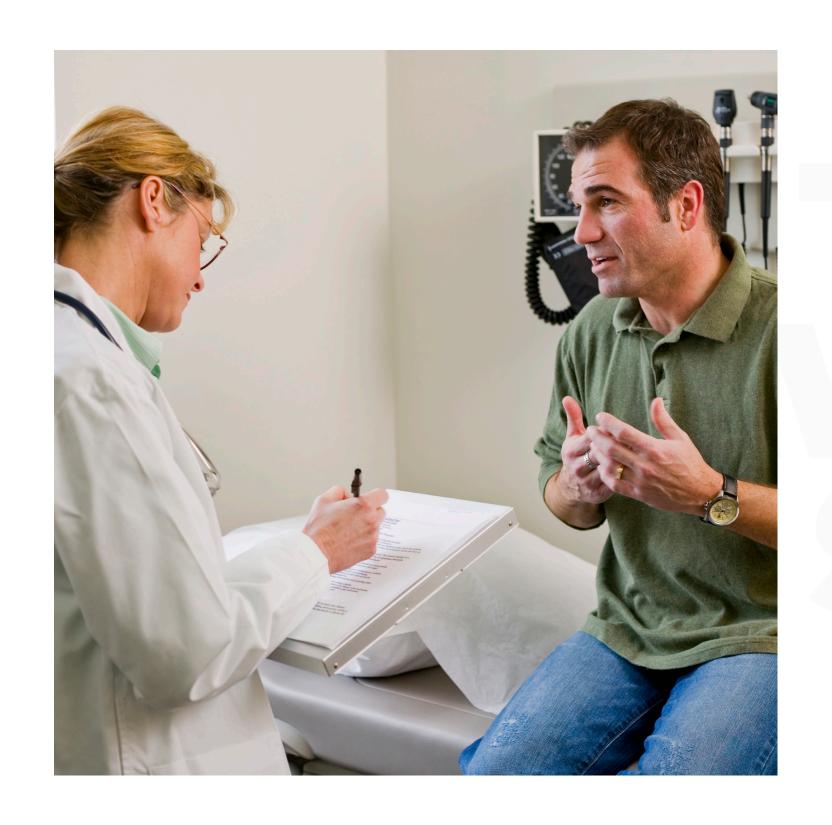
Complete the

Investigating Viral

Protein and the

Analysis of Coronavirus

Evolution handouts.



Now that you know about viral proteins, let's get back to Randy. You have diagnosed him with COVID 19. What is the best treatment plan?

## WATCH THIS!

