

Group: \_\_\_\_\_

# Introduction

### Materials (per group)

Computer with an internet connection

## Procedure

- EXAMPLE:**
- |       |                   |
|-------|-------------------|
| DNA:  | T A C G G C T A G |
|       | ↓                 |
| mRNA: | A U G C C G A U C |

*Your DNA sequence:*

[illegible]

*mRNA sequence:*

[illegible]

3. Determine the codons.

EXAMPLE: mRNA:     A U G C C G A U C  
                                  ↓  
          Codons:     AUG   CCG   AUC

*Codons:*

\_\_\_\_\_

4. Translate the codon sequence into an amino sequence. Use the chart provided.

Codons:       AUG           CCG           AUC  
                                  ↓  
Amino Acids: Methionine   Proline     Isoleucine

*Amino Acid Sequence:*

\_\_\_\_\_  
\_\_\_\_\_  
\_\_\_\_\_

5. Write out the one-letter abbreviations for the amino acids in the sequence. Use the chart provided on the last page.

\_\_\_\_\_

5. Go to <http://www.ncbi.nlm.nih.gov/BLAST/> and choose Protein Blast Under the Basic Blast heading.

7. Using the table on the last page, enter the one-letter abbreviations for your amino acid sequence in the SEARCH box labeled: “Enter accession number(s), gi(s), or FASTA sequence(s)” – be sure to enter them in the correct order!

8. Scroll down and click on the “BLAST” button.



9. At the next page, click on the “FORMAT” button. It may take a few minutes to process your sequence.

10. At the next page, scroll down to the “Description” of proteins matched your sequence. Choose one that matches one on the list of possible proteins on the last page.

Descriptions

**Sequences producing significant alignments:**

Select: [All](#) [None](#) Selected:0

Alignments

Download

GenPept

Graphics

Distance tree of results

Multiple alignment

	Description	Max score	Total score	Query cover	E value	Ident	Accession
<input type="checkbox"/>	<a href="#">dystrophin [Homo sapiens]</a>	48.6	48.6	100%	1e-05	93%	<a href="#">AAL61554.1</a>

11. The protein our DNA sequence encodes is (should be in the list provided): \_\_\_\_\_

12. Scroll down to the “Alignments” section. Locate the alignment for Homo sapiens (Humans). Click on the Gene link under related information to the right. This link should provide information regarding the function or associated disease of the gene and protein. Additionally, conduct an internet search for the name of the protein. Look for information regarding the protein and associated disease.

## Alignments

Download ▾ [GenPept](#) [Graphics](#)

▼ Next ▲ Previous ▲ Descriptions

dystrophin [Homo sapiens]

Sequence ID: [gb|AAL61554.1|](#) Length: 57 Number of Matches: 1

Range 1: 5 to 19 [GenPept](#) [Graphics](#)

▼ Next Match ▲ Previous Match

Score	Expect	Identities	Positives	Gaps
48.6 bits(107)	1e-05	14/15(93%)	14/15(93%)	0/15(0%)

### Related Information

[Gene](#) - associated gene details

13. This protein is involved in the following disease: \_\_\_\_\_
14. Write a brief paragraph explaining the disease caused by this protein or a mutation in this protein.

## AMINO ACID CHARTS AND PROTEIN NAMES

Possible proteins
Presenilin 2
Synuclein
Laforin
Leptin
BRCA 2
Dystrophin
Apolipoprotein E

		SECOND NUCLEOTIDE									
		U		C		A		G			
FIRST NUCLEOTIDE	U	UUU	Phenylalanine (Phe)	UCU	Serine (Ser)	UAU	Tyrosine (Tyr)	UGU	Cysteine (Cys)	U	THIRD NUCLEOTIDE
		UUC		UCC		UAC		UGC		C	
		UUA	Leucine (Leu)	UCA		UAA	STOP	UGA	STOP	A	
		UUG		UCG		UAG		UGG	Tryptophan (Trp)	G	
	C	CUU	Leucine (Leu)	CCU	Proline (Pro)	CAU	Histidine (His)	CGU	Arginine (Arg)	U	THIRD NUCLEOTIDE
		CUC		CCC		CAC		CGC		C	
		CUA		CCA		CAA	Glutamine (Gln)	CGA		A	
		CUG		CCG		CAG		CGG		G	
	A	AUU	Isoleucine (Ile)	ACU	Threonine (Thr)	AAU	Asparagine (Asn)	AGU	Serine (Ser)	U	THIRD NUCLEOTIDE
		AUC		ACC		AAC		AGC		C	
		AUA		ACA		AAA	Lysine (Lys)	AGA	Arginine (Arg)	A	
		AUG	Methionine (Met) START	ACG		AAG		AGG		G	
	G	GUU	Valine (Val)	GCU	Alanine (Ala)	GAU	Aspartic Acid (Asp)	GGU	Glycine (Gly)	U	THIRD NUCLEOTIDE
		GUC		GCC		GAC		GGC		C	
		GUA		GCA		GAA	Glutamic Acid (Glu)	GGA		A	
		GUG		GCG		GAG		GGG		G	

BLAST Amino Acid Abbreviations	
AMINO ACID	abbreviation
Alanine	A
Arginine	R
Asparagine	N
Aspartic acid	D
Cysteine	C
Glutamine	Q
Glutamic acid	E
Glycine	G
Histidine	H
Isoleucine	I
Leucine	L
Lysine	K
Methionine	M
Phenylalanine	F
Proline	P
Serine	S
Threonine	T
Tryptophan	W
Tyrosine	Y
Valine	V