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*Ap Biology Blast Lab
Answers*

AP Bio Investigation 3:

BLAST Lab ~~AP BLAST Lab~~

Comparing DNA Sequences **AP**

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*Lab 4: Plant Pigments and
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Answer: The percentage of similarity in the protein is always higher than the

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percentage of similarity in the genes for each species because proteins only have 20 amino acids and are specific in function, shape, and size. Genes on the other hand

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are more flexible and can have 64 possible codon communications.

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Cladogram generated from
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unknown sequence aligns

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BLAST Database is used to find similarities between biological sequences. The program compares nucleotide sequences and

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calculates the
significance of matches.
BLAST helps its users draw
conclusions about
evolutionary relationships
among different species.
One of, if not the most

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important pieces of
evidence for evolution is
DNA/proteins.

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Strong The Big BLAST Lab!
- AP Biology Lab Answer:
The percentage of
similarity in the protein
is always higher than the
percentage of similarity

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in the genes for each species because proteins only have 20 amino acids and are specific in function, shape, and size.

Blast Lab Answer Key -

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1) What Species in the BLAST result has the most similar gene sequence to the gene of interest ? -
The species in the BLAST lab is the Dorsophilla

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Melangoblaster F102063. 2)

Where is the species
located on the cladogram?

- The species is within
the 4th leaf under the
flies. 3) How similar is
that gene sequence?

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Investigation 3:

Undercovering Fossil

Specimen using BLAST A.

MORPHOLOGICAL OBSERVATION

OF THE FOSSIL SPECIMEN AND

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FORMING THE HYPOTHESIS

Make some general observations about the morphology (physical structure) of the fossil and record your observations in the space

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provided. The morphology of the fossil indicates that it is not a human.

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Once you have found the

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gene on the site, copy the gene sequence and input it into a BLAST query. Follow the instructions below to BLAST your own gene of interest: 1. On the Entrez Gene website search the

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term “keratin 18”. 2.
Click on the first link
that appears and scroll
down to “NCBI Reference
Sequences”. 3.

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In this laboratory
investigation, you will
use BLAST to compare

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several genes, and then use the information to construct a cladogram. A cladogram is treelike, with the endpoints of each branch representing a specific species. The

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closer two species are located to each other, the more recently they share a common ancestor.

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AP . . .

AP Lab 3: Comparing DNA Sequences to Understand Evolutionary Relationships with BLAST. Background Information . Between 1990-2003, scientists

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working on an international research project known as the Human Genome Project were able to identify and map the 20,000-25,000 genes that define a human being.

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AP Biology - Investigation 3 - DNA BLAST

Ap Biology Blast Lab
Answers 1) What Species in
the BLAST result has the
most similar gene sequence

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to the gene of interest ?
- The species in the BLAST
lab is the Dorsophilla
Melangoblaster F102063. Ap
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but stop in the works in
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Lab 4: Plant Pigments and
Photosynthesis AP Biology
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Answer: The percentage of
similarity in the protein

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is always higher than the percentage of similarity in the genes for each species because proteins only have 20 amino acids and are specific in function, shape, and size.

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Genes on the other hand are more flexible and can have 64 possible codon communications.

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BLAST Database is used to find similarities between biological sequences. The program compares

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nucleotide sequences and calculates the significance of matches. BLAST helps its users draw conclusions about evolutionary relationships among different species.

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One of, if not the most important pieces of evidence for evolution is DNA/proteins.

AP Biology BLAST Lab by
Emma Smith - Prezi

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NotebookBy: Stephanie
Strong The Big BLAST Lab!
- AP Biology Lab Answer:
The percentage of
similarity in the protein
is always higher than the

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percentage of similarity in the genes for each species because proteins only have 20 amino acids and are specific in function, shape, and size.

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1) What Species in the
BLAST result has the most
similar gene sequence to
the gene of interest ? -
The species in the BLAST

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lab is the Dorsophilla
Melangoblaster F102063. 2)
Where is the species
located on the cladogram?
- The species is within
the 4th leaf under the
flies. 3) How similar is

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that gene sequence?

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OF THE FOSSIL SPECIMEN AND
FORMING THE HYPOTHESIS

Make some general
observations about the
morphology (physical
structure) of the fossil
and record your

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observations in the space provided. The morphology of the fossil indicates that it is not a human.

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Once you have found the gene on the site, copy the gene sequence and input it into a BLAST query. Follow the instructions below to BLAST your own gene of interest: 1. On the Entrez

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Gene website search the term “keratin 18”. 2. Click on the first link that appears and scroll down to “NCBI Reference Sequences”. 3.

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use BLAST to compare several genes, and then use the information to construct a cladogram. A cladogram is treelike, with the endpoints of each branch representing a

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specific species. The closer two species are located to each other, the more recently they share a common ancestor.

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1990-2003, scientists working on an international research project known as the Human Genome Project were able to identify and map the 20,000-25,000 genes that

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define a human being.

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Answers 1) What Species in
the BLAST result has the

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most similar gene sequence
to the gene of interest ?
- The species in the BLAST
lab is the Dorsophilla
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