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Answer: The percentage of similarity in the protein is always higher than the percentage of similarity in the genes for each

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species because proteins only have 20 amino acids and are specific in function, shape, and size. Genes on the other hand are more flexible and can have 64 possible codon communications.

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Ap Biology Lab 21 Have A Blast

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Answers

BLAST Database is used to find similarities between biological sequences. The program compares nucleotide sequences and calculates the significance of matches. BLAST helps its users draw conclusions about evolutionary relationships among different species. One of, if not the most important pieces of evidence for evolution is DNA/proteins.

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Lab Answer: The percentage of similarity in the protein is always higher than the percentage of similarity in the genes for each

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

Blast Lab Answer Key - orrisrestaurant.com

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 *Drosophila Melanogaster* 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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Answers To Ap Blast Lab

Investigation 3: Undercovering Fossil Specimen using BLAST A. MORPHOLOGICAL OBSERVATION OF THE FOSSIL SPECIMEN AND FORMING THE HYPOTHESIS Make some general observations about the morphology (physical structure) of the fossil and record your observations in the space provided. The morphology of the fossil indicates that it is not a human.

Gotta Blast! - AP Biology Blog

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 Gene website

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

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AP Lab 3: Comparing DNA Sequences to Understand Evolutionary Relationships with BLAST. Background Information . Between 1990 – 2003, scientists working on an international research project known as the

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Human Genome Project were able to identify and map the 20,000 – 25,000 genes that define a human being.

AP Biology - Investigation 3 – DNA BLAST

Ap Biology Blast Lab Answers 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 Melangoblaster* F102063. Ap Biology Blast Lab Answers - ditkeerwel.nl this Ap Biology Blast Lab Answers, but stop in the works in harmful downloads. Rather

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