

Aminophosphonic And Aminophosphinic Acids Chemistry And Biological Activity

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13 nov 2024 i would like to use the uniprot api in python to get the uniprot accession given the sequence of amino acids but i can't seem to find a way to do so not from a sequence is there a

17 dec 2023 how can i create a heatmap visualization of amino acid attention scores with each row showing 30 amino acids and attention scores ask question asked 2 years 3 months ago modified

5 mar 2023 a mol object none on failure source of quote above since rdkit's managing of this case is logically reasonable you have to answer your question yourself how do i deal with unknown

26 apr 2021 i'd like to convert a column of three letter amino acids in excel to one letter and print out the one letter amino acids to each corresponding row in the excel file

13 apr 2018 thus a candidate protein string is derived by translating an open reading frame into amino acids until a stop codon is reached given a dna string s of length at most 1 kbp in fasta

having a fasta file containing protein sequences like these two showing below i would like to count how many times the amino acid a appears in each sequence sp p01920 dqb1 human hla class ii

20 feb 2018 i have the following dictionary which contains all the possible codons values triplets per amino acids keys letters this dictionary is also known as the dna codon table in bioinformatics

3 sep 2020 there's two ways to do this one way is to simply count the amino acid occurrence and use that as a predictor i hope i get the amino acids correct from school long time ago

9 sep 2022 i'm wondering if anyone knows any tools which allow me to count the frequency of amino acids at any specific position in a multiple sequence alignment for example if i had three

4 feb 2020 i am following this step to print the residues that i selected in pymol and i found that some of the residues got iterated more than 1 time does anyone know what the problem is ls iterate

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