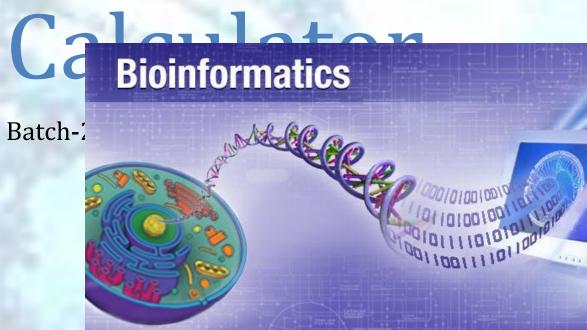
Name-Arabinda Chand

# Project-Amino Acid Calculator



Computer science 2/10/2022

#### Certificate

This is to certify that Arabinda Chand of class 12 of poddar brio international school has successfully completed his/her project work for the subject 'Computer science' for class XII project of the Central Board of Secondary Education in the year 2021-2022.

#### Acknowledgement

I would like to thank my teacher Miss. Pooja Nimroth(PGT computer science) who gave me this opportunity to work on this project. I got to learn a lot from this project about How I can take first step towards integrating computer science with biology. I would also like to thank our school principal Miss. Rashmi Singh.

At last, I would like to extend my heartfelt thanks to my parents because without their help this project would not have been successful. Finally, I would like to thank my dear friends who have been with me all the time.



- Introduction of sql, python & project
- Objectives
- Software & Hardware requirements
- Advantages & Disadvantages of the project, python & sql
- Coding
- Output screenshots
- Future through this project
- Bibliography

## Introduction To sq.

Structured query language provides codes to define data structure, manipulate data in a database & retrieving data in a database

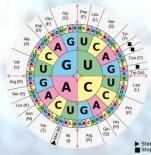
#### To python

Python is High level programming language. It was initially designed by Guido Van Rossum in 1991 in Centrum Wiskunde de Informatica,

Netherlands

#### To project

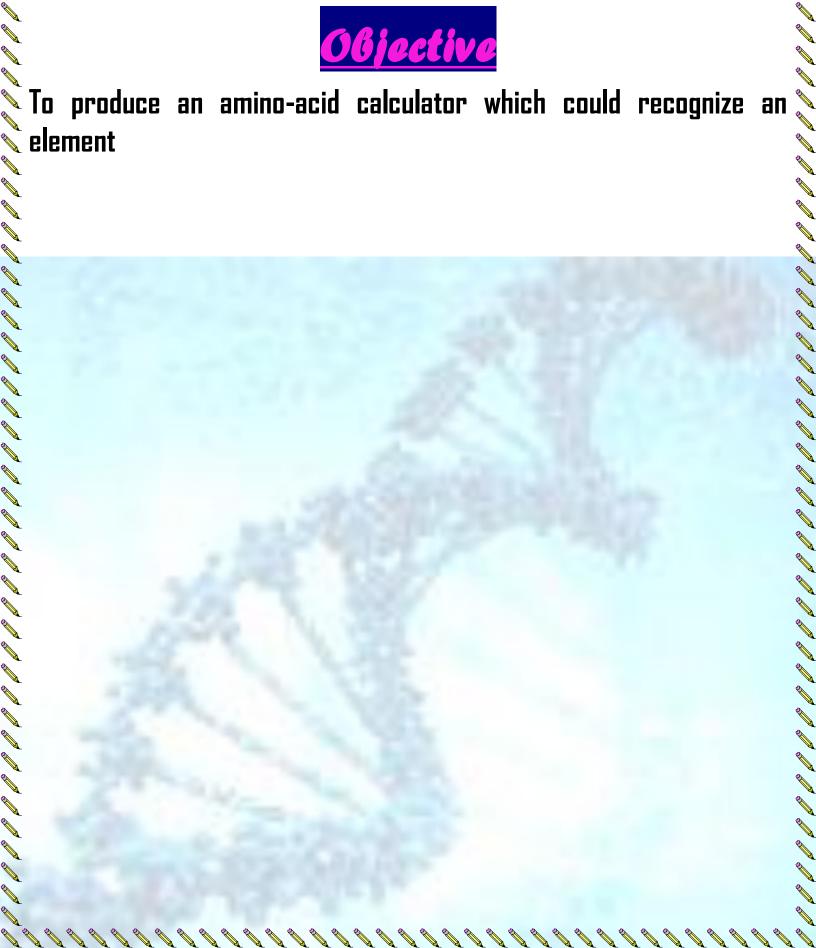
Introducing you to the future of calculators for biological application. Launching with the glory of genetics welcome to the world of dna with amino acid calculator. Which can tell you efficiently the name of any triplet (nucleotide base) codon corresponding to any of its amino acid. You can use it instead of memorizing the stack of 64 codon table of amino acids. Which can be problematic for many & it was also the first step to achieve the target of integrating biology with computer science with using practicality.











#### Hardware & software requirement

Hardware requirement-windows 7 laptop, a printer Software requirement-python, pip connector which can connect python mysql,mysql,word(2007)

2		
80 80 80°	Advantages of	Disadvantages of
	890	mysqe
	Sql is portable	More features implemented in
St.	It is of high speed	proprietor way
5 S S	Easy to learn	
200 m	Relational database	
	Not case-sensitive	Difficult to interface
2	Data consistency	
200	Object based	
SA SA	Client/server	
200	Programming & interactive	
2/102	language	
200	Advantages of	Disadvantages of
	python	python
30	Easy to learn & use	It is not the fastest
200		language(slower than compiled
200	757	ones)
800 AD	General purpose programming	Lesser libraries than c, java & perl
200	language	
5 S S	Interpreted(pre-compiled) &	Not strong on type binding(gives
200	Interactive	runtime error)
	Portable	Not easily convertible

P. J.

A			
	Python library, easy & short	Primitive Database access	S)
	syntax, highlighting & error		SV SV
	detection	Weak for mobile development	₹ Ş
Da.	Graphical User Interface		Ø.
	Extensible/extendable language	Memory consumption	S)
	Free & open source		Sy Sy
S.	Object oriented		8



- It has average 0.5 secs runtime and it has 121 bits
- It can be used instead of memorizing the table
- It is very fast & non tedious
- It is 99% accurate

- It focuses on use of computer science in biology
- It is originally made in india
- It can also tell you about start or stop codons

#### **Limitations**

- It can't be used for any other purpose
- Some people may prefer to rely more on memorizing for brainstorming for independence from the calculator



code in python-

import mysql.connector

econ=mysql.connector.connect(host='localhost',

password='12345678',

user='root',

database='amino\_acid\_calculator')

cursor=con.cursor()

code=input('enter your triplet')

sql='select\*from covid\_warrior where codon=%s'

dna=(code,)

cursor.execute(sql,dna)

output=cursor.fetchall()

for x in output:

print(x)

con.close()

code in mysql-

CREATE DATABASE amino\_acid\_calculator;

**USE** amino\_acid\_calculator;



```
CREATE TABLE covid_warrior (codon varchar(250) NOT NULL PRIMARY KEY,amino_acid varchar(250) NOT NULL);
INSERT INTO covid_warrior (codon,amino_acid)
VALUES ('UUU', 'Phenylalanine'),
('UUC', 'Phenylalanine'),
('UUA','Leucine'),
('UUG','Leucine'),
('CUU','Leucine'),
('CUC','Leucine'),
('CUA','Leucine'),
('CUG','Leucine'),
('AUU','Isoleucine'),
('AUC','Isoleucine'),
('AUA','Isoleucine'),
('AUG','Methionine(start)'),
('GUU','Valine'),
('GUC','Valine'),
('GUA','Valine'),
('GUG','Valine'),
('UCU','Serine'),
```

```
('UCC','Serine'),
('UCA','Serine'),
('UCG','Serine'),
('CCU','Proline'),
('CCC','Proline'),
('CCA','Proline'),
('CCG','Proline'),
('ACU','Threonine'),
('ACC','Threonine'),
('ACA','Threonine'),
('ACG','Threonine'),
('GCU','Alanine'),
('GCC','Alanine'),
('GCA','Alanine'),
('GCG','Alanine'),
('UAU','Tryptophan'),
('UAC','Tryptophan'),
('UAA','Stop(ochre)'),
('UAG','Stop(amber)'),
('CAU','Histidine'),
```

```
('CAC','Histidine'),
('CAA','Glutamine'),
('CAG','Glutamine'),
('AAU','Asparigine'),
('AAC','Asparigine'),
('AAA','Lysine'),
('AAG','Lysine'),
('GAU','Aspartic acid'),
('GAC','Aspartic acid'),
('GAA','Glutamic acid'),
('GAG','Glutamic acid'),
('UGU','Cysteine'),
('UGC','Cysteine'),
('UGA','Stop(opal)'),
('UGG','Tryptophan'),
('CGU','Arginine'),
('CGC','Arginine'),
('CGA','Arginine'),
('CGG','Arginine'),
('AGU','Serine'),
```

('AGC','Serine'), ('AGA','Arginine'), ('AGA', Argillile'), ('AGG','Arginine'), ('GGU','Glycine'), ('GGC','Glycine'), ('GGA','Glycine'), ('GGG','Glycine');

#### Future through this project

Bioinformatics is an emerging area where computers make old biology efficient whose era is yet to boom because people are getting interested for the speed, efficiency & accuracy of computer in biology.

### Bibliography

All In One Computer science
Preeti Arora Computer science
Rachna Sagar Computer science
Sumita Arora Computer science