

# PHYSICOCHEMICAL n-GRAMS TOOL (PnGT) TUTORIAL

## Introduction

**Physicochemical n-Grams Tool** is an open source standalone software for calculating physicochemical descriptors of protein developed using the Python scripting language.

The algorithms used in the development of this tool are based on the n-Grams curation techniques which are mainly used for natural language processing. Apart from language processing, now a days n-Grams modelling is also making a room in the computational biology and genomics.

The software currently calculates 33 physicochemical descriptors and the sequence length for given protein primary sequence. To find more specific information about our tool, please contact us.

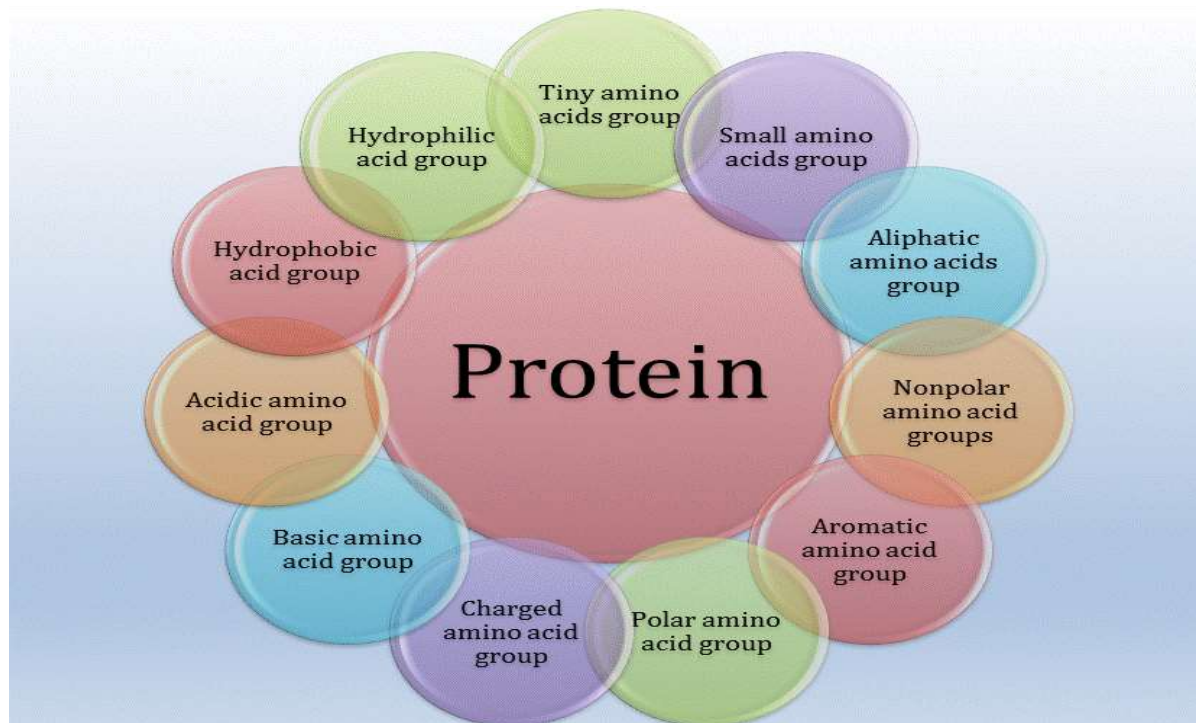


Figure.1 Physicochemical Amino Acids Property Groups

**Utility:** The tool is able to assess the **physicochemical properties** of the protein in an easier way with bi-gram, tri-Grams, tetra-Grams assessment of protein primary sequence.

**Assessment of the physicochemical n-Grams properties of the protein:** With the help of this PnGT tool, the analysis of proteomics data in the term of physicochemical n-Grams statistics is easy-to-do with the only primary sequence as feeding.

**Implementation in machine learning model development as feeding:** The outcomes in terms of physicochemical feature vectors from the Physicochemical n-Grams Tool can be directly implemented for the development of statistical (PCA, Regression Analysis) or machine learning model like SVM, kNN etc.

**Keywords:** Proteomics science, physicochemical parameter, n-Grams, machine learning, statistical modelling.

**Physicochemical n-Grams Descriptors and Calculations:** Conservation of multiple physicochemical groups along the sequence in a sliding window of length n residue. For example, physicochemical 2-grams: Aliphatic++; where n equals to 2

$$\{ \text{if aa} \}_i \in S^* \text{ AND } \{ \text{aa} \}_{(i+1)} \in S^*$$

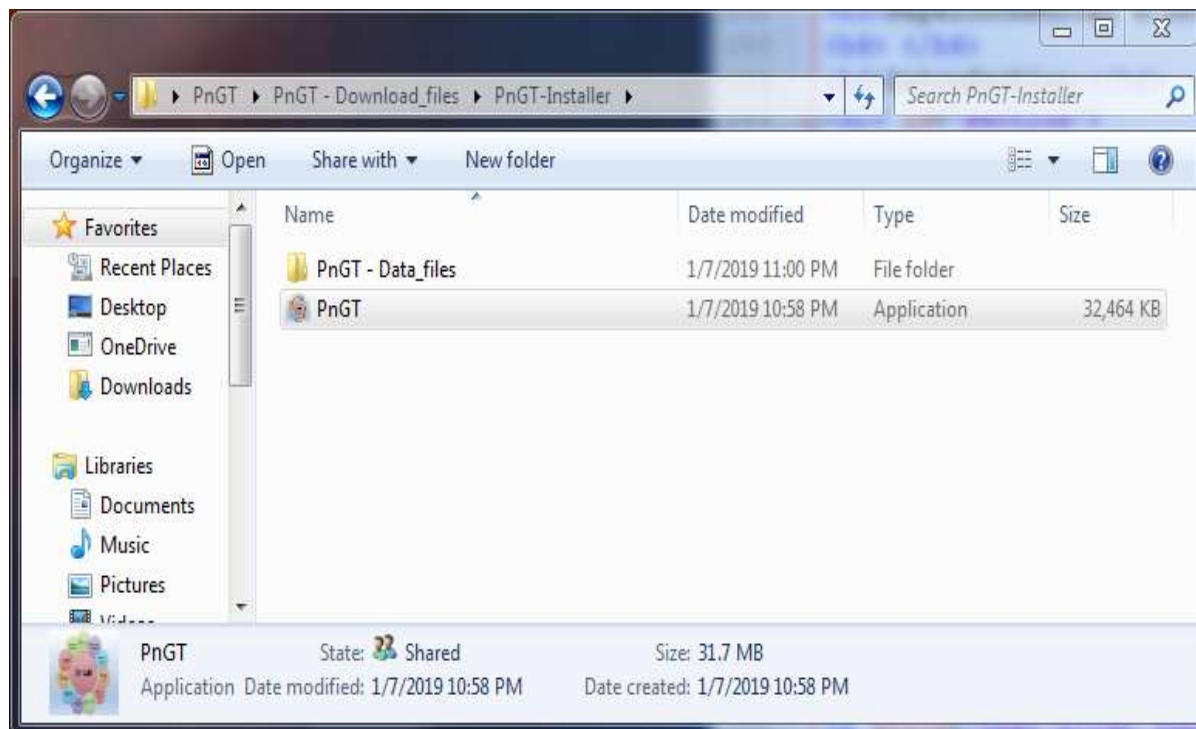
where, aai = ith residue of the protein

*Table 1. Physicochemical property groups of amino acids/*

Sr. no.	Name of amino acid property group	Amino acids in the specific group (S <sup>*</sup> )
1.	Tiny amino acids group	Ala, Cys, Gly, Ser, Thr
2.	Small amino acids group	Ala, Cys, Asp, Gly, Asn, Pro, Ser, Thr, Val
3.	Aliphatic amino acids group	Ile, Leu and Val
4.	Nonpolar amino acid groups	Ala, Cys, Phe, Gly, Ile, Leu, Met, Pro, Val, Trp, Tyr
5.	Aromatic amino acid group	Phe, His, Trp, Tyr
6.	Polar amino acid group	Asp, Glu, His, Lys, Asn, Gln, Arg, Ser, Thr
7.	Charged amino acid group	Asp, Glu, His, Arg, Lys
8.	Basic amino acid group	His, Lys, Arg
9.	Acidic amino acid group	Asp, Glu
10.	Hydrophobic acid group	Ala, Cys, Phe, Ile, Leu, Met, Val, Trp, Tyr
11.	Hydrophilic acid group	Asp, Glu, Lys, Asn, Gln

## Installation Steps for User

1. PnGT-Installer: To download “PnGT-Installer” go to [Download](#) tab and download PnGT-Installer to install Physicochemical n-Grams Tool.
2. Extract the zip file and open it, then locate “PnGT” as shown in Figure 2 below.



*Figure.2 PnGT Installer*

# Home Page

1. The PnGT will open as shown in Figure 3 below.

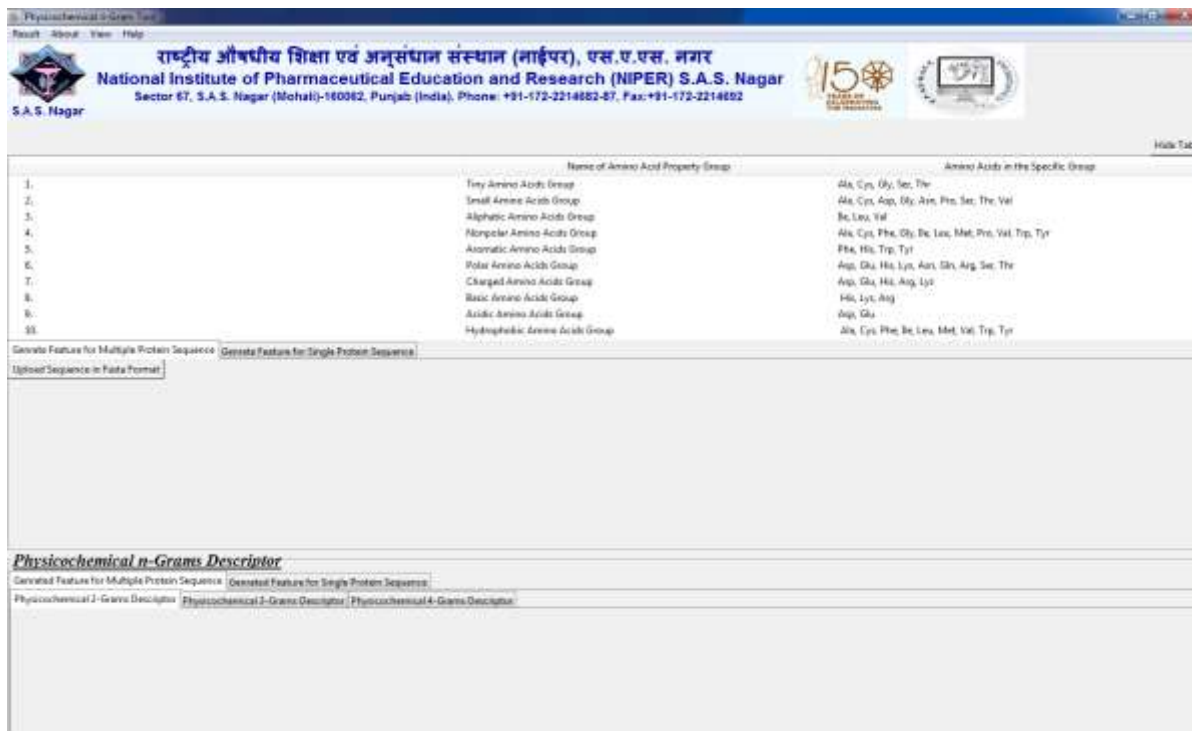


Figure.3 PnGT Home Page

## Physicochemical n-Grams Descriptors Calculation

- The tool is leveraged with two input method: FASTA format and simple sequence pasting method, if the user doesn't have the primary sequence in FASTA format.

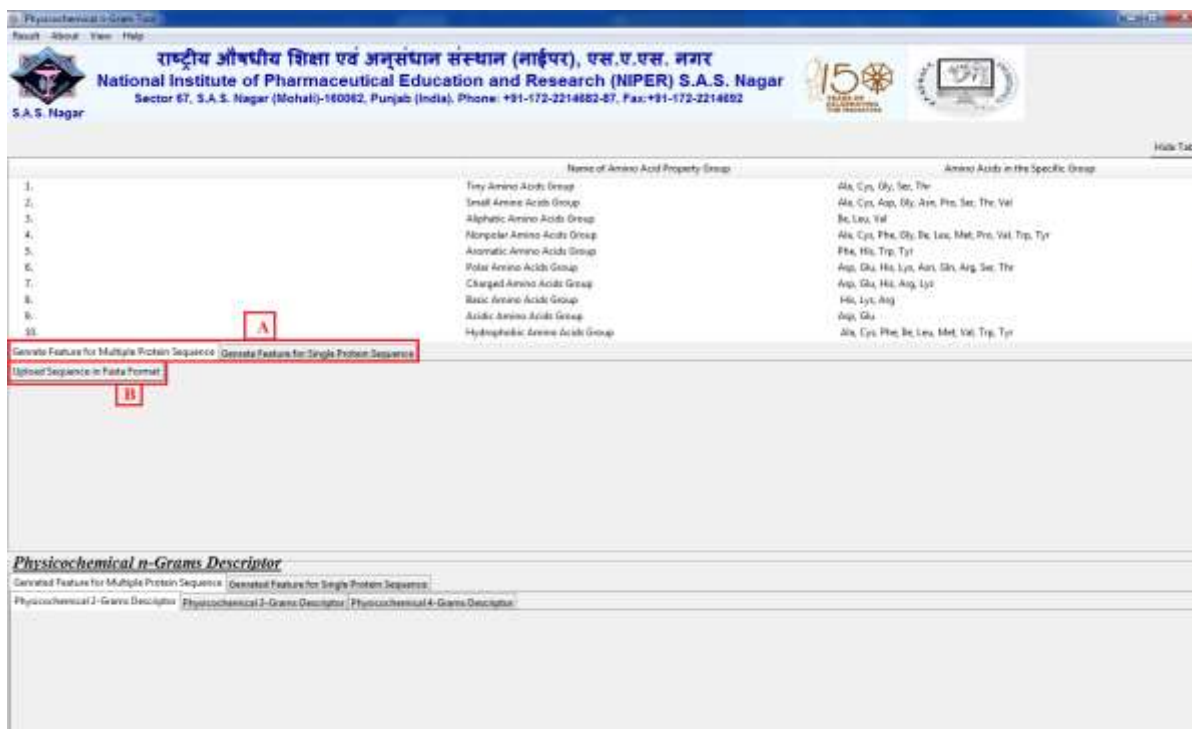




Figure.4A

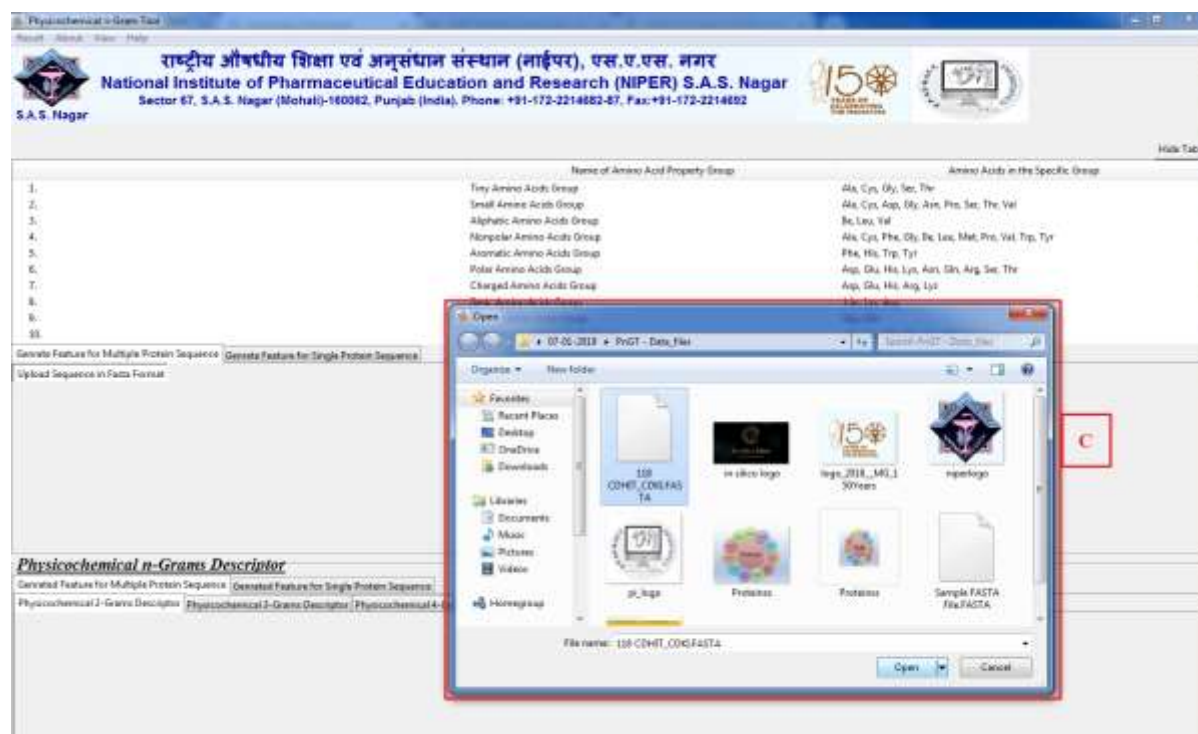


Figure.4B



Figure.4C



Physicochemicals Tool

File Edit Help


Name of Amino Acid Property Group

Amino Acids in the Specific Group

1. 2. 3. 4. 5. 6. 7. 8. 9. 10. 11. 12. 13. 14. 15. 16. 17. 18. 19. 20.

Upload File

Contact Us



Postal Address:  
Block A  
Department of Pharmacoinformatics,  
National Institute of Pharmaceutical Education and Research (NIPER)  
S.A.S. Nagar, Punjab-160062, India  
Please contact "indico10.meet@gmail.com" for any suggestions and queries

Amino Acids in the Specific Group

Ala, Cys, Gly, Ser, Thr  
Ala, Cys, Asp, Glu, Asn, Phe, Sec, Thr, Val  
Ser, Leu, Val  
Phe, Cys, Phe, Gly, Ile, Leu, Met, Pro, Val, Trp, Tyr  
Phe, His, Trp, Tyr  
Asp, Glu, His, Lys, Asn, Gly, Arg, Sec, Thr  
Asp, Glu, His, Arg, Lys  
His, Lys, Arg  
Asp, Glu

Amino Acid with One Letter and Three Letter Code

|                                 | Three Letter Code | One Letter Code |
|---------------------------------|-------------------|-----------------|
| Arginine                        | Arg               | R               |
| Asparagine                      | Asn               | N               |
| Aspartic Acid                   | Asp               | D               |
| Cysteine                        | Cys               | C               |
| Glutamic Acid                   | Glu               | E               |
| Glutamine                       | Gln               | Q               |
| Glycine                         | Gly               | G               |
| Histidine                       | His               | H               |
| Isoleucine                      | Ile               | I               |
| Lactamine                       | Leu               | L               |
| Lysine                          | Lys               | K               |
| Methionine                      | Met               | M               |
| Phenylalanine                   | Phe               | F               |
| Proline                         | Pro               | P               |
| Serine                          | Ser               | S               |
| Threonine                       | Thr               | T               |
| Tryptophan                      | Trp               | W               |
| Tyrosine                        | Tyr               | Y               |
| Valine                          | Val               | V               |
| 21st Amino Acid: Selenocysteine | Sec               | U               |
| 22nd Amino Acid: Pyrrolysine    | Pyl               | O               |

References

1. Livingstone, C.D. and Barton, G.J., 1993. Protein sequence alignments: a strategy for the hierarchical analysis of residue conservation. *Bioinformatics*, 9(6), pp.745-756.  
2. Ramana, J. and Gupta, D., 2010. Machine learning methods for prediction of CDK-inhibitors. *PLoS one*, 5(10), p.e13357.

Figure.6 The Snapshot showing widget of About Menu: Amino Acid and widgets of Help Menu: Contact Us & References.