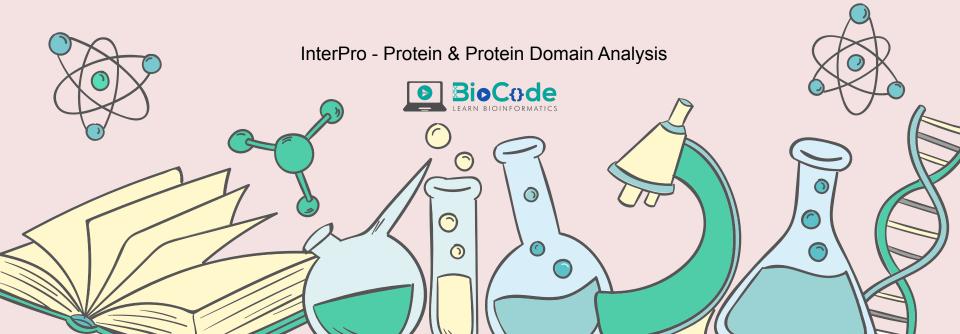
# InterPro Database

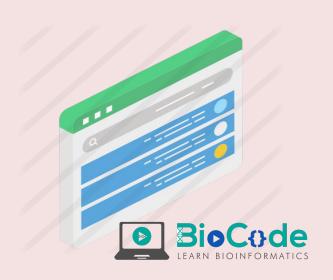




### Content

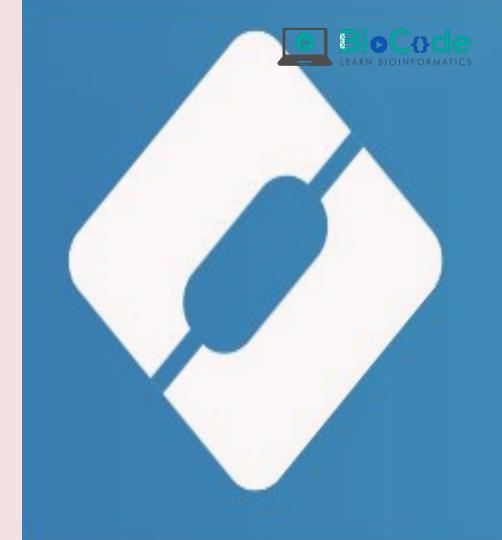


- Introduction to InterPro
- Benefits for Research
- Constitutes
- Input & Output
- Analysis
- Conclusion



# <sup>o</sup> Introduction

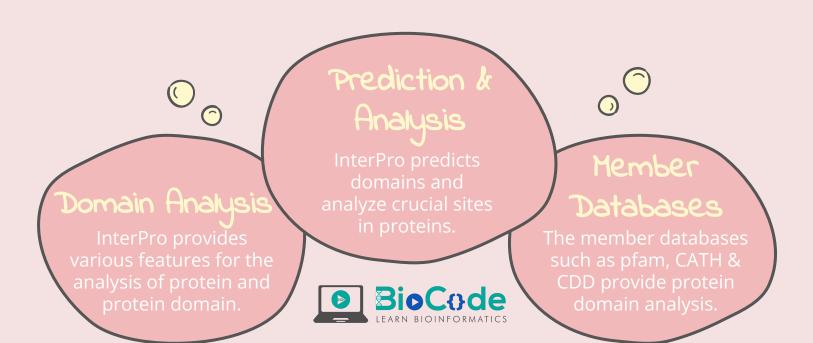
InterPro is a resource that provides functional analysis of protein sequences and provides protein family, domain and functionality of protein information. The member databases also provide such information.





InterPro combines signatures from multiple, diverse databases into a single searchable resource, reducing redundancy and helping users interpret their sequence analysis results.

#### Constitutes



## Input & Output

### Input

InterPro allows to search the query by sequence (FASTA), text search (protein names) or domain architecture.

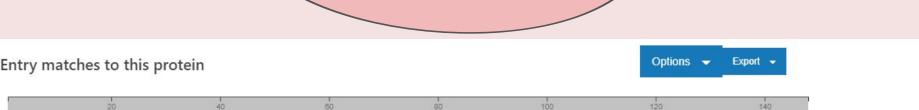




### Output

InterPro fetches search results itself and from member databases and provides list of the latest entries.





BioC@de

 Family ▼ Predictions

SIGNAL\_PEPTIDE (1)

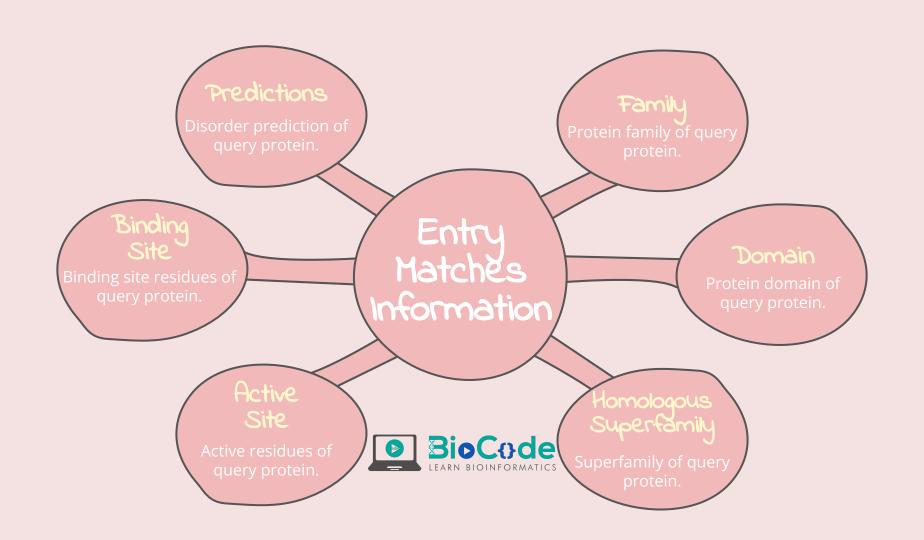
SignalP-TM (5) SignalP-noTM (6) SignalP-noTM (7)

SIGNAL\_PEPTIDE\_H\_REGION (2) SIGNAL\_PEPTIDE\_C\_REGION (3) SIGNAL\_PEPTIDE\_N\_REGION (4)

IPR038843 PTHR35523







### Query Search





Query

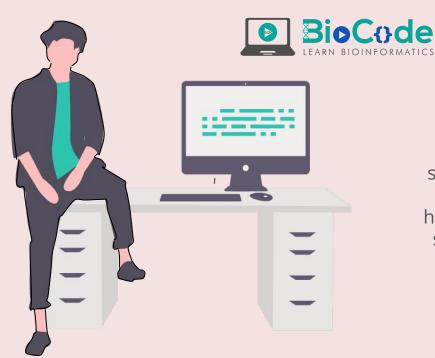
InterPro query can be search by FASTA sequence or by text as protein, domain names, keywords or GO terms.



Results

InterPro provides graphical representation, GO terms information and hyperlinks to member databases.

#### Conclusion



InterPro combines signatures from multiple, diverse databases into a single searchable resource, reducing redundancy and helping users interpret their sequence analysis results. InterPro provides links to integrated databases for protein & protein domain analysis.