

Pakistan Institute of Engineering and Applied Sciences Nilore, Islamabad, 45650 Pakistan

THESIS PROJECT PROPOSAL PROFORMA

Project Title	Large Margin Prediction of Host-Pathogen Protein Interactions		
Nature	Computational		
Prerequisites	 Knowledge of the following will be helpful: Machine Learning / Computational Intelligence / Pattern Classification / Data mining Python Programming Bioinformatics Interested students will be required to take Machine Learning and Bioinformatics courses if they have not already done so. 		
Field	Scientific Computing and Computational Intelligence		
Expected Cost (if any)			
Work Place*	PIEAS		

^{*}define %age of work to be done in PIEAS or at some other place

Supervisor Information

Supervisor information				
Name of supervisor and designation	Dr. Fayyaz ul Amir Afsar Minhas, Senior Scientist			
Department\Division and Organization	Department of Computer Science, PIEAS			
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Project Details

Synopsis	Motivation: Infections result from the
2,110,000	interaction between proteins from host and
	pathogen. For example, the interaction
	shown on the right between the Tat protein
	from HIV with the human P-TEFb protein is
	involved in AIDS. Identifying these protein-
	protein interactions is the first step in the
	development of medicinal drugs to counter infectious diseases
	which are responsible for 17% of all deaths in the world. For more
	details visit: http://faculty.pieas.edu.pk/fayyaz/hpi/index.html
	Objectives: The objective of this project is to develop machine
	learning models for predicting host-pathogen protein interactions.
	The model will be give protein sequences of two proteins, one from
	the host and the other from a pathogen, as input. The model's
	output is a score reflecting the tendency of the proteins to interact
	with each other. Such a computational model can pave the way for
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	future biology discovery through wet-lab experimentation.
	Skills Resulting from the project: Big data analysis, machine
	learning, Bioinformatics, Python Programming, Text analysis and
	Interactomics, Multidisciplinary Research methodologies and
	publishing.

Goals of the Project

4 th Semester (3 credit hour per week are available)	 I. Development of understanding of proteins and their interactions II. Learning to handle protein data on the computer III. Study of properties of proteins useful in prediction of interactions IV. Understanding the challenges developing a machine learning predictor for host-pathogen interactions V. Hands-on understanding of machine learning techniques VI. Developing a baseline predictor
5 th Semester (12 credit hour per week are available)	I. Analysis of features and classification schemes for Host-pathogen protein interaction prediction. II. Benchmarking the predictor III. Development of a webserver for the prediction

Instructions

- > All the columns of project proposal forms are mandatory to be filled.
- > The minimum qualification for a supervisor is eighteen years of education plus two years service.
- > The minimum requirement for a co-supervisor is eighteen years of education.
- In case of external supervisor (outside DCIS), a co-supervisor will be mandatory from faculty.
- Wherever the project work is carried out (including classified establishments), the defense (only examiners and supervisor) and final presentation (open to all) will be held at PIEAS. The presence of supervisor is necessary for all presentations of fourth and fifth semesters.
- > DCIS has right to reject or accept any project.

Signature of Supervisor

Signature Head of the Department

(With name and designation)