









# 2<sup>nd</sup> National Workshop on Bioinformatics and Computational Biology

Dates: 18-20 December, 2016 Venue: CVASU and USTC Campus, Chittagong, Bangladesh

### **Organized Jointly By:**

Bangladesh Bioinformatics and Computational Biology Association (BBCBA)
Bioinformatics Research Group of Rajshahi University (BioRGRU)
Chittagong Veterinary and Animal Sciences University (CVASU)
University of Science and Technology Chittagong (USTC)

#### **Supported By:**

Higher Education Quality Enhancement Project (HEQEP:CP-3603, CP-3215) of UGC

#### **Program**

Day-1. Sunday 18 December 2016

**Venue: CVASU New Conference Room (Academic Building-2, 2<sup>nd</sup> Floor)** 

| 09.00-09.30 am   | Registration  |
|------------------|---|
| 09.30-10.30 am S | Professor, Dept. of Statistics, Rajshahi University  Introducing the Keynote speaker: Prof. Dr. AMAM Zonaed Siddiki, General Secretary, BBCBA and Professor, Dept. of Pathology and Parasitology, CVASU  Key presentation: "Application of Bioinformatics in Epidemiology and Control of Infectious Diseases" Prof. Dr. Probodh Borah, Head, Biotechnology Dept, Assam Agricultural University, India  Speech from Special Guest: Mrs. Mahmuda Khatun, Director, BCSIR, Chittagong.  Speech from Special Guest: Prof. Dr. Provat Chandra Barua, Hon'ble Vice Chancellor, University of Science and Technology Chittagong (USTC)  Speech from Chief Guest: Prof. Dr. Goutam Buddha Das, Hon'ble Vice Chancellor, Chittagong Veterinary and Animal Sciences University  Speech from the Chairperson: Prof. Dr. Nurul Absar, Pro-Vice Chancellor, University of Science and Technology Chittagong (USTC) |

| 10:30-10:45am  | Tea Break   |
|----------------|---|
| 10.45-01.00 pm | Session 2: Presentation session Venue: Video conference Room, Academic Building (3 <sup>rd</sup> Floor), CVASU Facilitator: Prof. Dr. Probodh Borah, AAU, India Theoretical sessions  1. Fundamentals of Bioinformatics and its Applications 2. Sequence Alignment and Molecular Phylogeny  |
| 01.00-02:00 pm | Lunch break   |
| 02:00-05:00 pm | Session 3: Hands on Practical session Facilitator: Prof. Dr. Probodh Borah Theoretical Sessions  1. Protein Structure Prediction and Molecular Modeling 2. Bioinformatics in Drug Discovery Hands-on Sessions:  1. Protein Sequence Retrieval from UniProtKB (with keywords tumor necrosis factor, human)  2. Protein Structure Visualization using RASMOL/UCSF Chimera 3. Protein Primary Structure Analysis - ProtParam 4. Protein secondary structure prediction by SOPMA/ExPAsy Tools 5. Protein Tertiary structure prediction by homology modeling using Swiss Model workspace/Modeller 6. Protein Tertiary structure analysis by RAMPAGE – PDBsum/ PROCHECK |
| 05:00 pm       | Closing   |

Day-2. Monday 19 December 2016 Venue: USTC New Campus (Dept of Pharmacy, 6<sup>th</sup> Floor Lecture Room)

| 09.00-10.00 am | Session 4: Presentation and Hands on training session   |
|----------------|---|
|                | Lecture 1   |
|                | Topic: Bioinformatics as a tool for regeneration biology: Retinal regeneration study                                  |
|                | perspective   |
|                | Resource Person: Dr. Md. Rafiqul Islam  |
|                | Chairman, Dept. of Genetic Engineering and Biotechnology, Chittagong University.                                      |
| 10:00-10:15 am | Tea Break   |
| 10.15-01.00 pm | Hands on training session   |
|                | Lecture 2   |
|                | <b>Topic:</b> Introduction to computational genomics and hands-on training.   |
|                | Resource Persons: Md. Jahangir Alam <sup>1</sup> and Dr. Hafizur Rahman <sup>2</sup>                                  |
|                | <sup>1</sup> Bioinformatics Lab, Dept. of Statistics, <sup>2</sup> Institute of Biological Sciences (IBSc.), Rajshahi |
|                | University  |
| 01.00-02.00 pm | Lunch Break   |
| 02:00-05:00 pm | Session 5: Presentation and Hands on training session   |
|                | Lecture 3   |
|                | <b>Topic</b> : Introduction to computational transcriptomics and hands-on training.                                   |
|                | Resource Persons: Mr. Md. Shahjaman and Prof. Dr. Md. Nurul Haque Mollah  |
|                | Bioinformatics Lab, Dept. of Statistics, Rajshahi University  |
|                | Lecture 4   |
|                | Topic: Introduction to computational metabolomics and hands-on training.  |
|                | Resource Persons: Mr. Nishith Kumar and Prof. Dr. Md. Aminul Hoque  |
|                | Bioinformatics Lab, Dept. of Statistics, Rajshahi University  |

## Day-3.Tuesday 20 December 2016 Venue: USTC New campus (Dept of Pharmacy, $6^{\rm th}$ Floor Lecture Room)

| 09.00-10.00 am  | Consign (. Durgentation and Hands on two ining against  |
|-----------------|---|
| 09.00-10.00 am  | Session 6: Presentation and Hands on training session   |
|                 | Lecture 5   |
|                 | Bioinformatics tools for whole genome sequence (WGS) data analyses  |
|                 | Resource Person: Dr. Asutosh Das  |
|                 | Associate Professor, Dept. of Animal Breeding and Genetics, CVASU and Fellow, Aarhus                                |
|                 | University, Denmark   |
|                 |   |
| 10:00-10:15 am  | Tea Break   |
| 10.17.01.00     |   |
| 10.15-01.00 pm  | Lecture 6   |
|                 | Topic: Next generation drug discovery: Designing Drugs by Computer.   |
|                 | Resource Person: Mr. S. M. Zahid Hosen  |
|                 | Scientific Officer, BCSIR, Chittagong   |
|                 |   |
| 01.00-02.00 pm  | Lunch Break   |
|                 |   |
| 02:00-04:00 pm  | Session 7: Presentation and Hands on training session   |
|                 | Lecture 7   |
|                 | <i>Topic: Introduction to computational toxicogenomics and hands-on training.</i>                                   |
|                 | Resource Persons: Md.Nazmol Hasan <sup>1</sup> , Md. Masud Rana <sup>1</sup> and Dr. Moizur Rahman <sup>2</sup>     |
|                 | <sup>1</sup> Bioinformatics Lab, Dept. of Statistics, <sup>2</sup> Dept. of Veterinary and Animal Science, Rajshahi |
|                 | University  |
|                 | University  |
| 04.00-04.30 pm  | Lecture 8   |
| 04.00-04.30 pm  |   |
|                 | Topic: Next generation sequencing (NGS) data analyses pipeline: An introduction                                     |
|                 | Resource Person: Prof. Dr. AMAM Zonaed Siddiki  |
|                 | Dept. of Pathology and Parasitology, CVASU and Fulbright Scholar, The George Washington                             |
|                 | University, USA   |
|                 |   |
| 04.30- 05.00 pm | Session 7: Closing session  |
|                 | Feedback session  |
|                 | BBCBA Membership drive  |
|                 | Certificate awarding ceremony   |
|                 |   |

<sup>\*</sup>All trainee participants are requested to bring their own laptop PC, one multi-plug for AC power and Hi-speed wifi modem for Hands-on training sessions. The copy of all presentations will be available online in Dropbox and relevant information will be forwarded by email to all registered participants.

<sup>\*\*</sup>For any queries, please contact Prof. AMAM Zonaed Siddiki @ 01717718884 or Mr. Abul Hashem @ 01819179169.