



✓ We were given a booth of our own at the conference to spread the word about our initiatives and activities.

 RSG-India also featured in the participant handbook and several other publicity material of HiPC.

 RSG-India also shared the exhibition space with major companies such as Wipro, CRL, Platform computing which are actively involved in providing infrastructure support to Computational Biologists for carrying out high performance computing.



MESSAGE BOAR

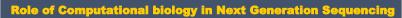
 ARTICLE – <u>Mastering Your</u> <u>Ph.D. Defending Your Thesis</u> <u>With Flair</u>
 Research Fellowship -<u>CCMB Summer Training</u> <u>Program for the year 2011</u>
 Scholarship - <u>Sir Doradji</u> <u>Tata Trust – Young Researcher</u> <u>Award – 2011</u>
 EVENT - <u>Bioinformatics</u> <u>National Certification 2011</u> (<u>BINC 2011) Examination</u>
 Research Position - <u>Career</u> <u>Opportunities @ JNU</u>

6. OPENING - <u>POST-</u> <u>Doctoral fellowship in</u> <u>systems biology and</u> <u>bioinformatics</u>
7. QUERY - <u>perl script for</u> <u>cartesian to internal</u> <u>coordinates</u>
8. ADMISSION - <u>Admission</u> to Ph. D. Program @ IIT

<u>Hyderabad</u> 9. CONFERENCE - <u>OMICS</u> Hyderabad-International

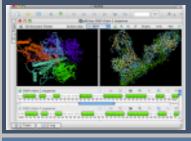
Conference and Exhibition on Proteomics & Bioinformatics 2011

10. MISC - <u>1000 Genomes</u> <u>Project Tutorial Videos</u>



NGS are the set of new techniques, algorithms, and software leading to better and high-end implementation of genomic applications. The flowchart depicts the flow of the discussion. Please click on the elements to view the corresponding discussion.

Path to NOS Advantages of Niss



'BIO – Innovation' - <u>UGENE</u>

UGENE is free, open-source bioinformatics software. It has a friendly user interface and runs on Linux, Windows, MacOS and Solaris. It also contains a library of highly optimized algorithms, and has very moderate hardware requirements to run. The primary features include MSA using KAlign, PCR Primer design, ORF Finder, Protein back translation, etc.



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