



Bioinformatics Training

(in the context of a core)

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Harvard Chan Bioinformatics Core

<http://bioinformatics.sph.harvard.edu/>



Harvard Chan Bioinformatics Core

- Focus on Next-Generation Sequencing (NGS) data



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- Consulting/Analysis services use a fee-for-service model
 - Different types of subsidies available for specific groups based on internal funding
- Training Program is fully subsidized by internal funding - assigned FTE for “dedicated” training personnel



Interplay between Training and Consulting

- Outreach,
- Community building,
- Resources for client education



Training

Consulting



- Current best practices,
- New technologies,
- Workshop topic selection

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 - ✦ Additional time spent on material development, organization, administration, outreach, community building

Training goals and method highlights

Training focused on enabling participants to-

- Utilize best practices for experimental design and data analysis
- Analyze their own data more independently
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Methods highlights-

- Small class sizes - 25 participants on average with 3 instructors
- Incoming students at similar level of computational skills (not enforced)
- Use of local research computing resources for continuity
- Easily updatable and accessible training/learning materials on GitHub

Acknowledgements and Contact Information

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Meeta Mistry & Mary Piper!



Contact information-

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