

Practical Aspects of Running a Bioinformatics Core Facility

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Together we will beat cancer



Structure

- Team structure/composition (9.4 FTE)



- Customers

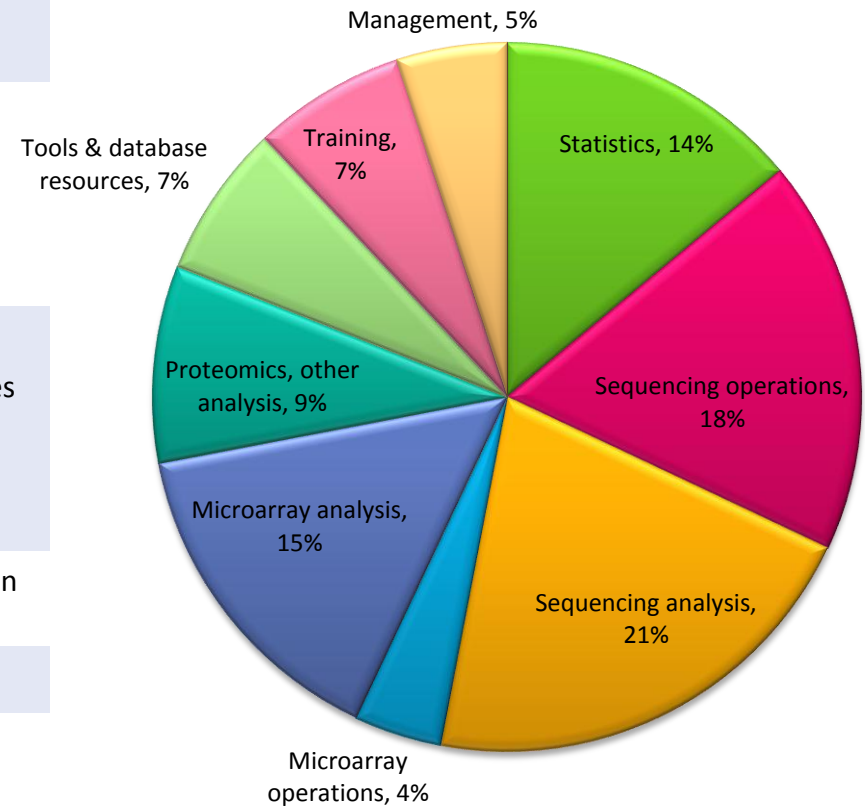
- 300 research scientists in 22 research groups
- Other core facilities – Genomics, Proteomics
- Sequencing operation for 3 collaborating institutes in Cambridge

- CompBio @ CRI

- Core (10) + 2 research groups (20) + 10 “embedded” in wet-lab groups

Services

Statistics	Drop-in clinics
Experimental Design	Experimental design clinics
Next gen sequencing	LIMS Data management, processing, QC Analysis pipeline development Analysis projects – ChIP-seq, variation
Microarrays	Operational support, data QC Analysis pipeline, Bioconductor packages <ul style="list-style-type: none"> ▪ beadarray package, probe annotation Analysis projects <ul style="list-style-type: none"> ▪ Illumina expression, Affymetrix SNP, exon, ...
Proteomics	Analysis projects – SILAC MS/MS, protein arrays
Other analysis	Motif enrichment
Analysis tools & data resources	Galaxy, Ensembl (local installations) Commercial databases, tools
Training courses	NGS, microarrays, motif analysis, functional/pathway analysis



Funding

- Cross charging since April 2011
 - Hourly rate for consultation and analysis support directly attributable to research group or grant
 - Does not include overheads (seminars, office space, computing resources)
 - No sign-off or approval process

- Exemptions
 - Drop-in clinics, experimental design meetings
 - Training courses
 - Operational support – LIMS, analysis pipeline development
 - Commercial software and database licenses

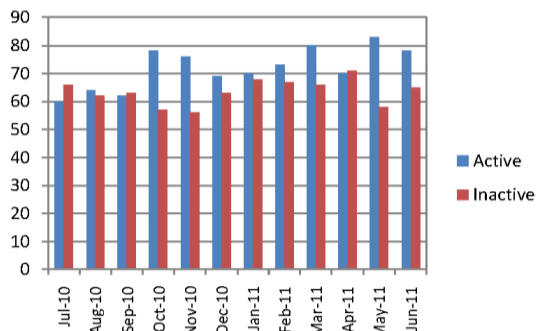
- Approx. 40% recorded time is cross-charged

Monitoring

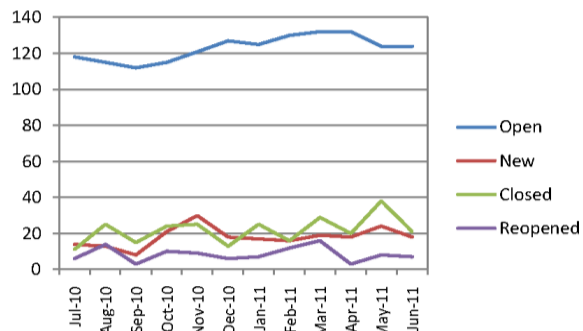
- Group objectives (yearly)
- Update to Steering committee (quarterly)
- Scorecard (monthly)
 - Metrics to monitor health/performance of Core
- External review (every 3 – 4 years)
 - User questionnaire/survey
 - Detailed report
 - Assessment by external review panel

Bioinformatics Core Scorecard – June 2011

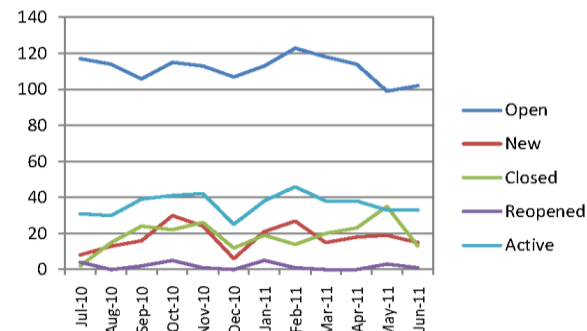
Research Group Projects/Tasks



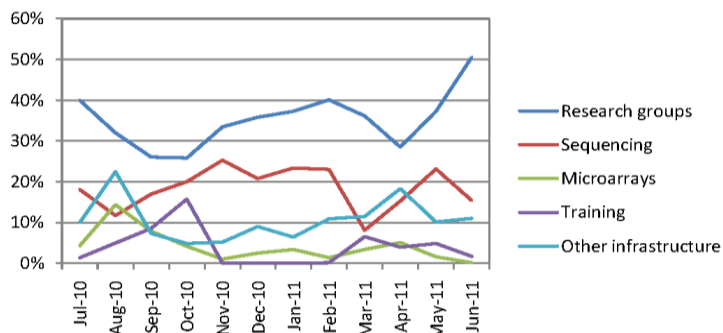
Research Group Projects/Tasks



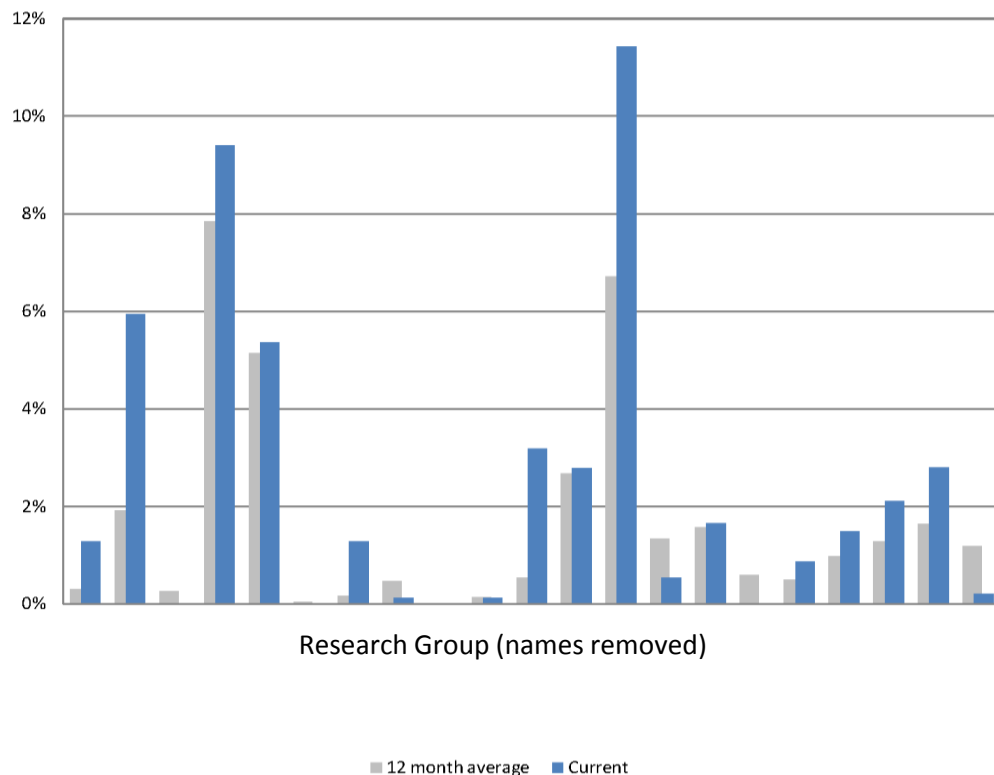
Infrastructure and Other Projects/Tasks



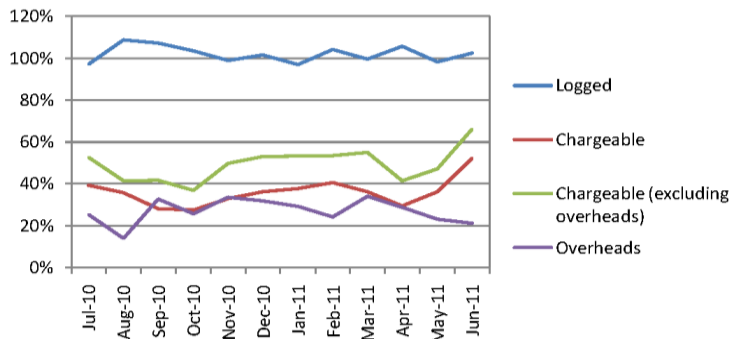
Time Breakdown



Time Recorded – Research Group Projects/Tasks



Time Recorded – Chargeable and Overheads



Challenges

Challenge	Strategy/Response
Large numbers of projects and support requests	<ul style="list-style-type: none">▪ Project tracking/management
High workload/demand	<ul style="list-style-type: none">▪ Analysis pipeline development▪ Balancing workload across team▪ Cross charging
Frustration with turnaround/responsiveness	<ul style="list-style-type: none">▪ Improved communication, expectation setting▪ Enabling biologists to be more self-sufficient (training courses, access to tools, databases)
Evolving bioinformatics service model	<ul style="list-style-type: none">▪ Changing interactions with users, remit and nature of service

Project tracking

Challenge: volume of projects/requests

- 70 – 80 chargeable projects/tasks active each month
 - largely excludes infrastructure, software development
- Project management tool (redmine)
 - Capture project details, status
 - Time logging for monitoring, charging
- Projects largely scheduled on FIFO basis but no formal queue
- Questions:
 - *Do other cores operate a more formal queuing/scheduling process?*
 - *How are other cores measuring response/turnaround times?*

Workload management

- **Communication with researchers**
 - Clearly define scope/deliverables, set realistic expectations
 - Communicate often about project status and progress
- **Balance workload within team**
 - Project monitoring and reassignment
 - Skills development across team
- **Analysis pipeline development and automation**
 - Significant investment but yields efficiency gains and consistent, high-quality deliverables
- **Access to analysis tools/resources and training**
 - Galaxy including roll-out of analysis tools developed in-house
 - Various commercial databases and associated tools

Challenge: high workload/demand

Challenge: turnaround, responsiveness

Future of the Bioinformatics Core

Challenge: evolving service model

- Increasing adoption of “embedded bioinformatician” model
- Bioinformatics Editorial, March 2011, Kallioniemi, Wessels & Valencia
Author’s institutes:
 - Core: 1 FTE per 100 scientists, institutionally-funded
 - Embedded: 5–10x working in research groups funded by outside grantsChanging nature of Core service:
 - More emphasis on more general, institute-wide support activities, less on specific research-oriented work
- Questions:
 - *What impact would this have on the Core’s ability*
 - *to deliver services, e.g. training, pipeline development?*
 - *to retain high-calibre staff?*