

Coordination of TB diagnostics research:
Enabling standards and sharing of data on the molecular basis of drug resistance

Senate House, London, UK, 3 - 4 February 2014

Action points

1. Compile short-term tools
 - Suggest fields for metadata
 - List of datasets, content, strain availability
 - List of unsequenced strains highly characterized available for sequencing, ideally including outcome data
 - List of strain collections and criteria for inclusion in a strain bank
 - Rough summary of gaps in current datasets (geographic representation, resistance coverage)
2. Establish rules for quality assessment of sequencing and phenotype
3. By email collect larger potential uses for such a database
4. Have IT consultant propose architecture that would meet all DST needs
5. Reach-out to the larger TB R&D community
6. Have a follow-up meeting