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

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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