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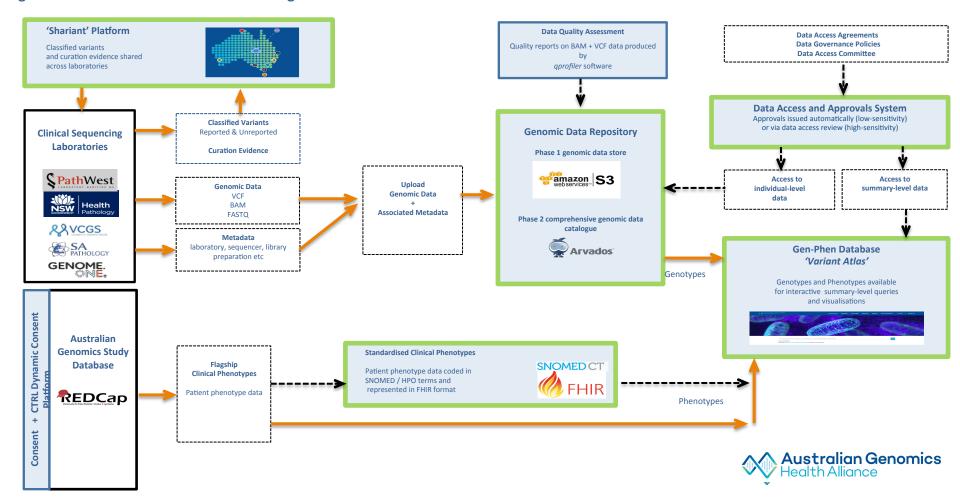
Supplemental Data

Australian Genomics: A Federated Model for

Integrating Genomics into Healthcare

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Figure S1: Australian Genomics data management work flow and infrastructure



Supplementary Table 1: Australian Genomics programs and sub-projects.

| PROGRAM 1: National diagnostic and research network | PROGRAM 2: National approach to data federation and analysis | PROGRAM 3: Evaluation, Policy and Ethics | PROGRAM 4: Workforce and Education |
|--|--|---|--|
| National clinical genomic consent Develop a single shared consent process and supporting documentation for clinical genomic testing for all indications. | Study database (REDCap) Manage participant clinical, demographic and survey data | Evaluate progress across the Flagship projects to ensure research outcomes are being effectively translated into practice. Create a research base for getting evidence into practice. | Workforce status Professional status survey of genetic counsellors and clinical geneticists, including current activity and education in genomics. |
| CTRL – Participant portal and dynamic consent platform Empower research participants with granular, dynamic consent options; updates about the research they've contributed to; and access to information, and patient support. Real-time feedback loop to researchers as consent preferences change. | Standardized, computer-readable clinical data Translate clinical data into standardized codes (phenotype ontologies: HPO/SNOMED) to allow computerized search, discovery and coding | Social network analysis Map and analyze interconnections between members to capture the collaborations amongst the genomic community, to document learning, assess Australian Genomics' influence, and identify key players in implementation. | Existing genomics education Map existing Australian education and training for health care professionals. |
| Virtual gene panels Develop a national approach to | Genomic data repository The cloud-based storage of | Genomics policy Mapping of major genomic | Needs assessment A portfolio of projects aiming to understand health professionals' |

| the design, maintenance and use of virtual gene panels in genomic analysis to minimise duplication of effort and ensure consistency of data analysis. | genomic sequencing data (BAM/VCF/FASTQ) together with a subset of the participant's clinical data to facilitate meaningful analysis and sharing. | medicine policy frameworks and demonstration projects internationally. Review of the international genomics policy landscape, including reimbursement policies, genetic discrimination and data sharing. | current practice and needs for continuing medical education in genomics, including genetic specialists, medical specialists, and community practitioners from the perspectives of these professionals, educators and patients. |
|---|--|--|--|
| Clinical variant re-classification | Genotype-Phenotype platform: | Ethics and legal | Evaluating Education |
| Develop guidelines for managing | Variant Atlas | Review and make | Develop and pilot tools to |
| re-classification of genetic | Interactive Genotype-Phenotype | recommendations on the legal | support development of |
| variants, and systems to ensure | data platform to visualise | and ethical aspects of various | effective genomic education |
| reporting to referring clinicians | aggregated variant data, and | models and applications of | strategies and enable evidence- |
| and patients | filter by key clinical features to | genomic data sharing in Australia | based education practice. |
| | describe cohorts | and internationally | |
| | | Contribute to Australian | |
| | | Genomics initiatives in clinical | |
| | | consent, research consent and | |
| | | insurance discrimination. | |
| Mainstreaming genomic | National variant sharing | Health economic evaluation | Genetic Counseling for rapid |
| pathology reports | platform: Shariant | Measuring and valuing the short- | turnaround. |
| Develop recommendations for | Online platform for laboratories | and long-term benefits of clinical | Examine genetic counselling |
| plain-language genomic test | to share curated variant | genomics using diverse | processes in the acute care |
| reporting, making reports | classifications and detailed | approaches including cost- | setting and the psychosocial |
| accessible to non-genetic health | curation evidence in real-time | effectiveness, cost-benefit, cost- | impact of rapid genomic |

| professionals, and patients | i.e. prior to report generation. Automatic notification about classification differences, and submission of summary information to international databases. | utility, and budget analyses. | sequencing on families to inform service delivery models. |
|---|---|-------------------------------|--|
| MSAC application pipeline Apply evidence built through clinical Flagships to submit applications to MSAC seeking Federal funding recommendations for genomic testing | Data access agreements and policies Participants provide consent to both national and international data sharing, for the benefit of healthcare. Clinicians and researchers can request access to Australian Genomics datasets, and this will be granted according to the level of data sensitivity, the specific consent of the participant, and the researcher's HREC, where applicable. Australian Genomics subscribes to GA4GH policies and standards. | | Public perceptions and expectations Establish an evidence base to understand the Australian public's diversity of experiences, values, attitudes and expectations of genomics in research and healthcare. |
| Functional genomics network Linking clinicians and researchers, this project aims to provide catalyst grants for pilot | | | |

| projects aiming to incorporate | | |
|--------------------------------|--|--|
| functional genomic assays into | | |
| the diagnostic pipeline | | |
| Genomics in the community | | |
| Develop a suite of information | | |
| resources about genomics, | | |
| genomic testing and related | | |
| issues (eg data sharing), in | | |
| collaboration with patient | | |
| advocacy groups | | |