



# Family Health History Breakout Report

G2MC Virtual Session  
May 8, 2020



Global Genomic  
Medicine Collaborative



## Grant Opportunities and other Funding Ideas for a Shared Family Health History Project

- Dr. Theodora Katsila (Greece) outlined EU grants (digital biomarkers for public health, young investigators) postponed until later this year due to COVID-19. She will provide an outline of the funding opportunity and her aims.
- Geoff Ginsburg (USA) would like to see funding that would support G2MC hosting a FHH resource center for members to use: knowledge experts, examples of established implementations and solutions, papers, access to shared FHH tools, grant opportunities, etc.



## Grant Opportunities and other Funding Ideas for a Shared Family Health History Project

- Other sources of funding could include –
  - Genealogy companies,
  - Cloud providers and Software tech - could provide in-kind support (personnel, etc) or funding,
  - Genetic testing labs,
  - Direct-to-consumer companies (e.g. 23andme, etc.)
  - Rare disease foundations (need to suggest specific projects closer to their interests)
- Need to create marketing stories for each potential funding partner

# GA4GH and HL7 Related Activities

- Global Alliance for Genomics & Health Pedigree subgroup working on a minimum core dataset with common data elements and definitions for collection and sharing of FHH information.
- Reviewed current HL7 FHIR Pedigree standards.



---

# Sharing of Use Cases

FHH Breakout



# Common Family Health History Use Cases

- Familial risk assessment for a disease of interest
- Using FHH as an indication for genetic testing, or for genetic analysis post-genotyping/sequencing
- Using FHH, genotype, and phenotype (clinical) data of a patient or relative, to determine if the patient needs further testing or sequence analysis, and/or if a relative needs the same, which is called cascade testing
- Tracking in pedigrees inherited genetic disorders, such as both autosomal recessive and dominant, codominant, x-linked, and mitochondrial

## Some shared stories

- [Tomoharu Tokutomi \(Japan\)](#) new tools one being in English, French, and Japanese
- [Theodora Katsila \(Greece\)](#) discussed her work in using FHH coupled with PGx for drug toxicity evaluation
- [Sonia Margarit \(Chile\)](#) described how the medical school is teaching student to administer pedigree analysis
- [Franciso Domingues \(Italy\)](#) described population study looked at chronic, age-related diseases. They are still interested in integrating FHH tools into this population. They are conducting a sub-study on COVID-19
- [Geoff Ginsburg \(USA\)](#) suggested that using FHH tools in the realm of infectious diseases
- [Anne Ochoa \(Philippines\)](#) is a Genetic Counseling student from Manila says they take family histories by hand, manually drawing family pedigrees up to third generation.
- [Yusuf Danasabe Jobbi \(Nigeria\)](#) shared that they take a short family history while doing case files and appreciates the chance to learn and have the opportunity to introduce more tools in their system.





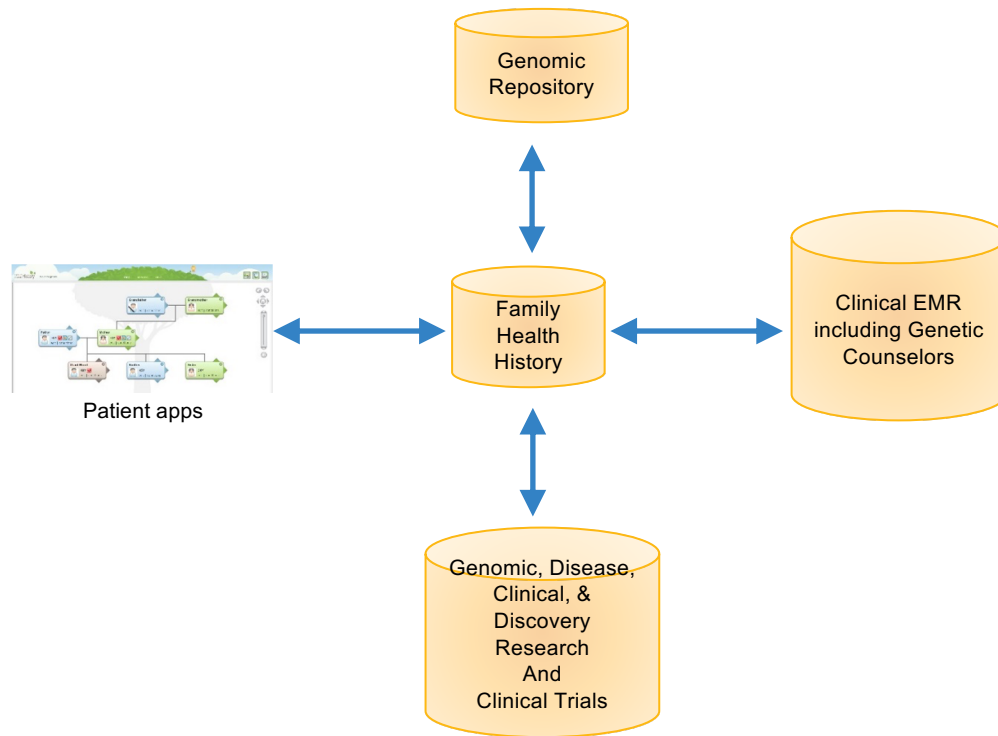
## Ongoing Collection of FHH Use Cases Using a Webform

- ▶ Describe what FHH collection tools are currently in use and need to be created
- ▶ What is the targeted patient population and main data focus (disease, genetic variants, drug, etc.)
- ▶ Explain the main goal and high-level clinical outcome you want to achieve





# Universal FHH Use Case for Data Sharing





# Goals review and discussion

## Short term – infrastructure

- Discuss what tools can be shared, or must be local
- Local data storage requirements
- Discuss how data could be accessed/shared across countries
- List other local requirements (language, devices, etc.)



# Goals review and discussion

## Long term – Desired outcomes

- Utilize common or standardized method/process to capture data
- Generate consistent FHH data across multiple different research and care sites
- Analyze FHH data to develop clinical decision support algorithms