

Blueprint Genetics

Rare3k

December 4th, 2019

Introduction to Blueprint Genetics

Dedicated Laboratory for Human Rare Diseases

- Founded in 2012 as a spin-off from Stanford University
- Specialty genetic testing business focused on broad hereditary genetic diseases
- End-to-end service from sample preparation to report delivery
- Single gene analysis, gene panels, Whole Exome Sequencing
- Unrivalled clinical interpretation practice supported by a highly qualified team
- Proprietary software platform and machine learning / AI capabilities

Key Facts



170 Employees
Founded in 2012
>4,000 Clinicians in 70 countries

Core Operations In



Helsinki
Seattle

Quality Management



CLIA

Clinical laboratory certification by US regulatory authority

CAP

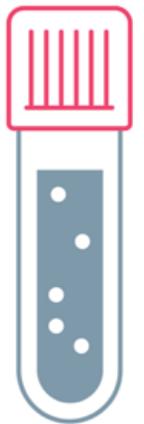
All laboratory functions accredited by College of American Pathologists

ISO 15189

Processes and documentation comply with the most widely used standards worldwide

CE-IVD

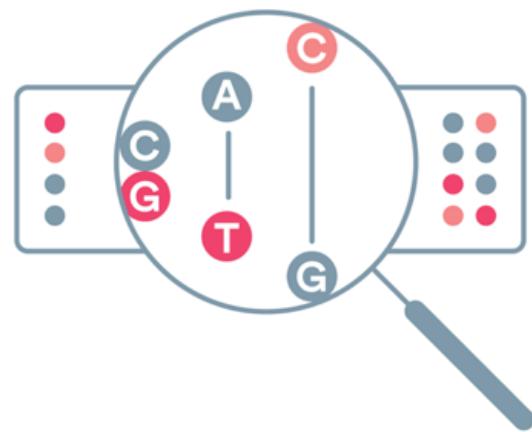
All Blueprint Genetics' panel tests are CE marked, general IVD devices



Patient
sample



DNA
sequencing



Analysis and
interpretation



Clinical
report

Superior analytic and clinical performance

PERFORMANCE	WHOLE EXOME	CLINICAL EXOME
Single nucleotide variants		
Sensitivity	99.65%	99.83%
Specificity	>99.99%	>99.99%
Insertions, deletions and indels by sequence analysis		
1-10 bps sensitivity	96.94%	99.33%
1-10 bps specificity	>99.99%	>99.99%
11-50 bps sensitivity	99.07%	99.16%*
11-50 bps specificity	>99.99%	>99.99%
Copy number variants (exon level dels/dups)		
1 exon level deletion (heterozygous) sensitivity	NA	100% (20/20)
1 exon level deletion (homozygous) sensitivity	NA	100% (5/5)
1 exon level deletion (het or homo) sensitivity	92.3% (24/26)	100% (25/25)
2-7 exon level deletion (het or homo) sensitivity	96.0% (24/25)	100% (44/44)
1-9 exon level duplication (het or homo) sensitivity	NA	75% (6/8)
Microdeletion-duplication sdrs (large CNVs)		
Size range (0.1-47 Mb)	100% (37/37)	100% (25/25)
Simulated CNV detection		
5 exons level deletion/duplication sensitivity	98.63%	98.7%
5 exons level deletion/duplication specificity	99.98%	100.00%

*11-30 bps

**Clinical sample indel sensitivity 100.0% (1-80bps)



The rare disease challenge

Over **350** million people affected

It takes on average **5.6** years for the illness to be properly identified

Visiting **8** doctors during diagnostic odysseia

Only **10%** have an approved therapy

Our solution

Digitalizing rare disease diagnostics:

AI programs can analyze huge quantities of information from medical records to identify high-risk individuals

Genome analysis and clinical interpretation can efficiently identify the underlying pathogenic mutations to establish a molecular diagnosis

Rare3K - Introduction

- Helsinki Biobank, HUS and Blueprint Genetics are launching **a research** initiative to sequence a cohort of 3 000 donors representing different rare disease groups.
- The study establishes pilot programs for diagnosing patients with Wilson disease, aTTR polyneuropathy, achondroplasia, inherited retinal disorders (IRD) and age-related macular degeneration (AMD).
- **This collaborative** project aims to **test, whether combination of** HUS data lake, Helsinki Biobank specimen collections **and AI tools** and Blueprint Genetics' clinical genetic testing resources **improves identification of rare disease patients**.
- This pilot aims to assess the impact of the biobank approach for diagnostics of rare diseases.
- **The Rare3k approach creates an accredited clinical path from sample to diagnosis by combining resources in the biobank infrastructure, clinical data in data lake and clinical whole exome sequencing for efficient diagnostics of rare diseases.**

Research objectives

- **The project is defined as biobank research, in which:**
 - patients with biobank consent and permission to recontacting (94%) are analyzed,
 - biobank donors want to receive “terveydelle merkityksellistä tietoa” (99%),
 - **with help of the participating clinical experts**, we aim to be able to recontact the patients to improve their diagnostics.
- **The Rare3k project aims to identify 3 000 donors in the Helsinki Biobank that have a clinical suspicion of a rare inherited disease by:**
 - developing algorithms to identify potential rare disease patients based on clinical information in the HUS data lake and
 - applying computational methods and machine learning to identify disease subclasses and establish disease classifiers.
- **The project' objective is to perform whole exome sequencing and clinical interpretation for the 3000 donors with suspected rare disease to identify the underlying pathogenic mutations**

Pilot programs in Rare3k

Wilson disease

- Underdiagnosed disease that leads to copper build-up affecting liver and brain

TTR amyloidosis

- Neurodegenerative disorder with several phenocopies

Achondroplasia

- 80% of mutations are *de novo* and lead to short stature

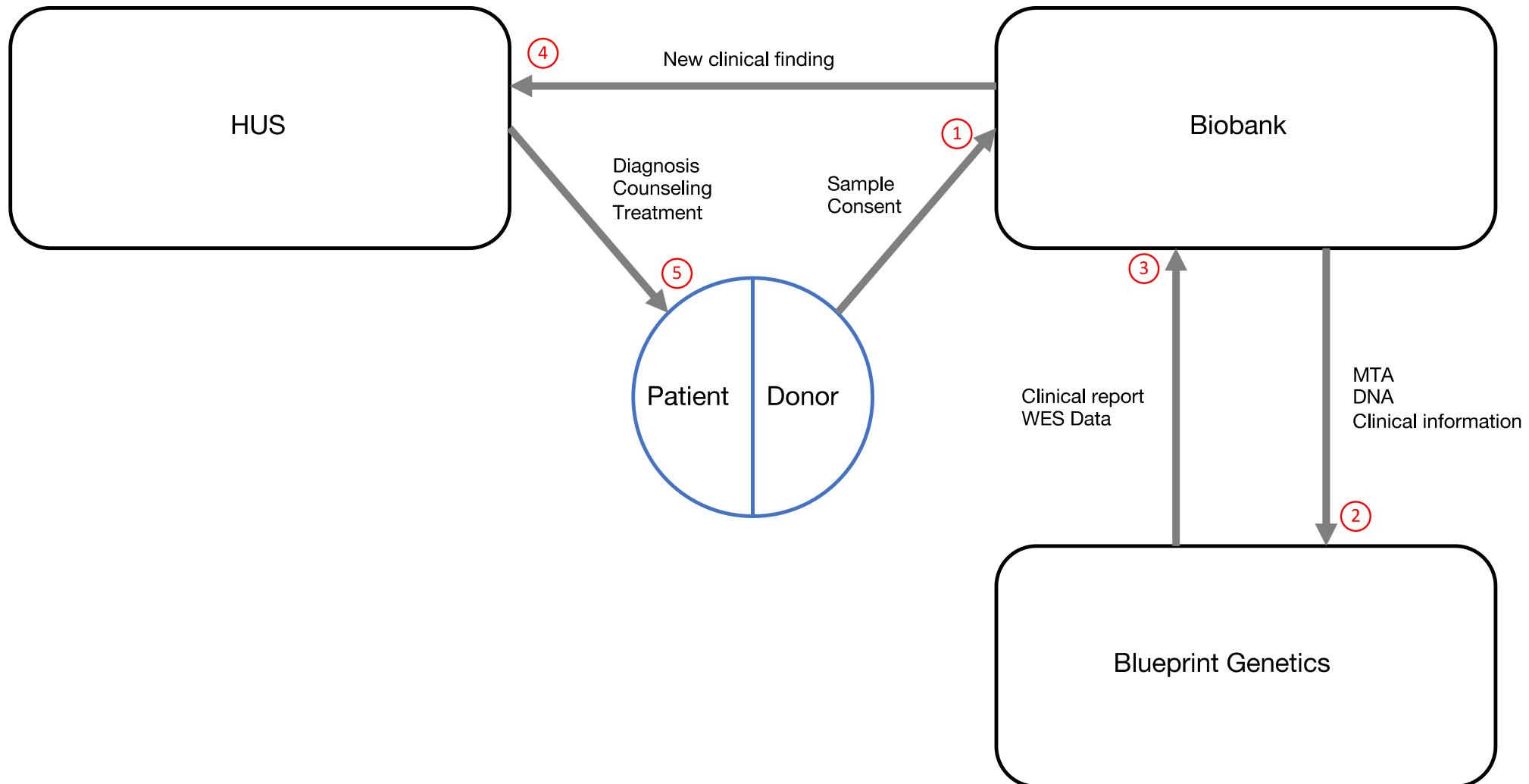
Inherited retinal disorders

- Heterogeneous disorders affecting retina where accurate molecular diagnosis is key in selecting treatments

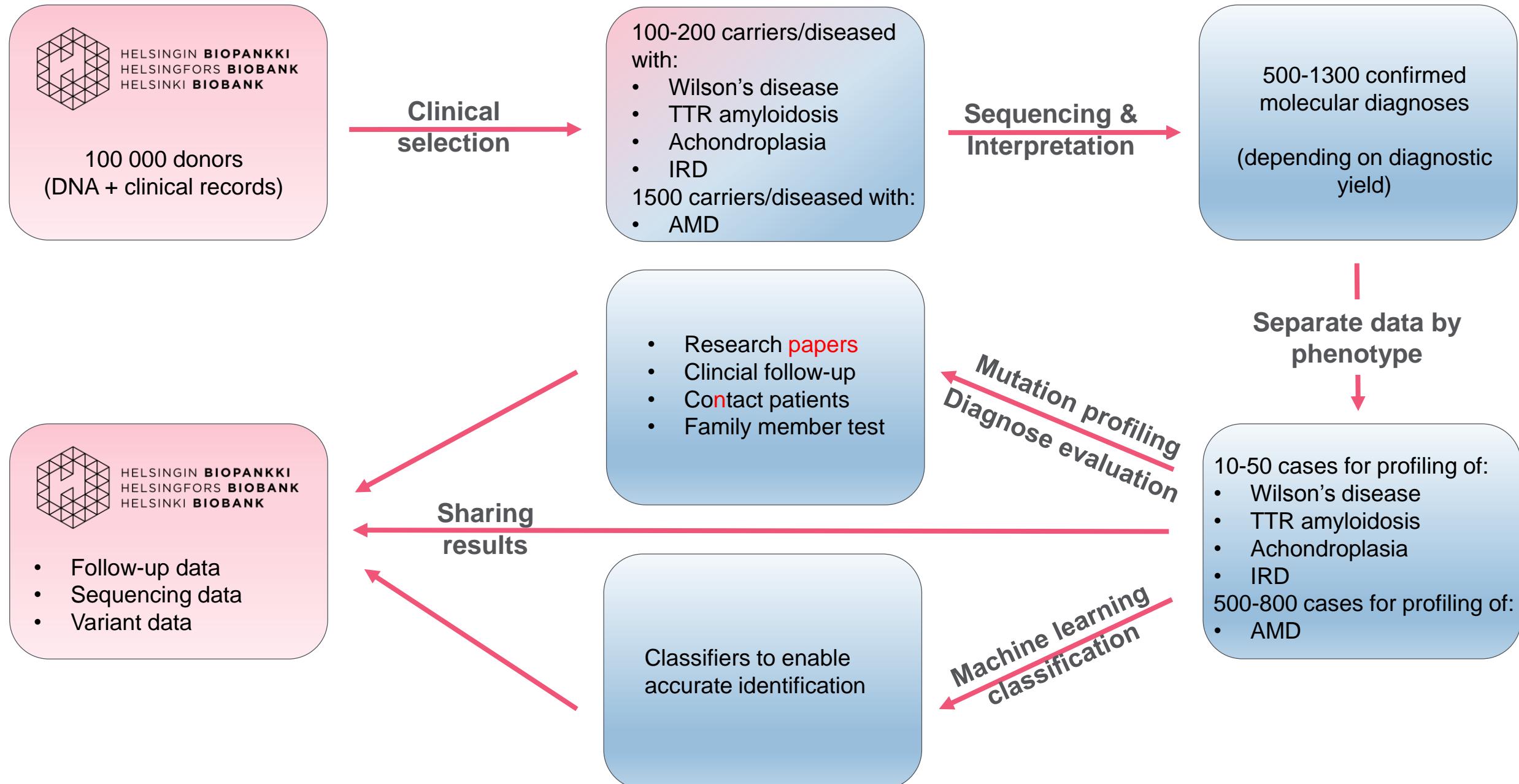
Age-related macular degeneration

- Progressive disease affecting macula where genetic factors explain up to 71% of disease variation

Collaboration model



Project work-flow





Team

Helsinki Biobank

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