

Why didn't genetics change the world (yet)?

• P1: Complex genome structure and organization

- P2: Difficulties in definition of our phenotypes
- P3: Epigenetics
- P4: Cost
- P5: People...

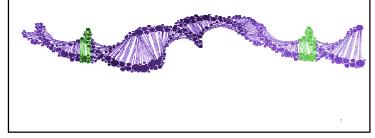


Problem 1: Genome structure

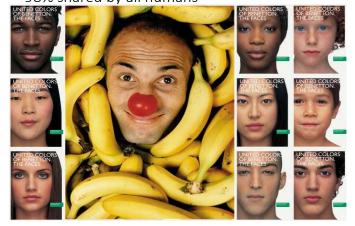
- A lot of data: 3×10⁹ base pairs (ACGT)
- About 2m long molecule (if assembled together)
- 23 chromosome pairs (22+XX/XY)
- 17-25k genes expected (based on homology with other species and gene structure)
- Gene: piece of DNA that has a (almost) unique start and end signals
- Furthermore ...

2% of DNA (3^*10^7 base pairs)

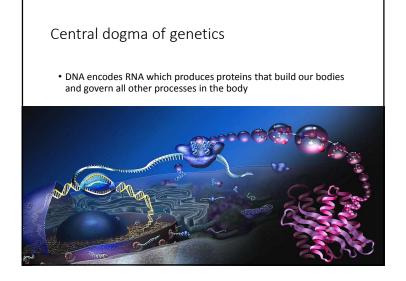
- All variation in humans is stored in those 2% called the exome
- The rest is "dark matter", of unknown function (so far) the introns
- Only recently we started producing some data related to the biological role of introns



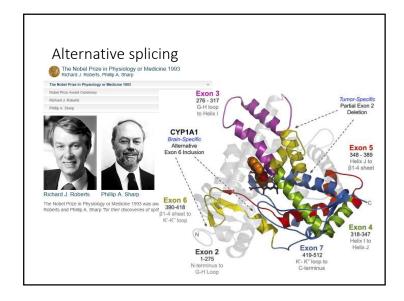
98% shared by all humans





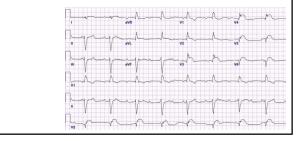


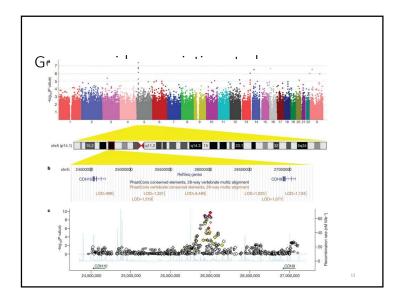
Problem 2: Difficult phenotypes Many disease occur as the result of numerous reasons Mycoardial infarction – occlusion of the blood vessel in heart that leads to death of a heart muscle Embolus (blood cloth, air bubble) Atherosclerosis (blood vessel hardening) Hyperlipidaemia (elevated lipids) Salt overuse* Chemical injury (homocysteine)

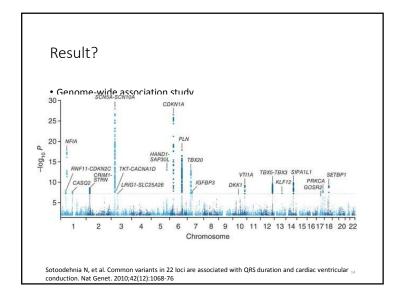


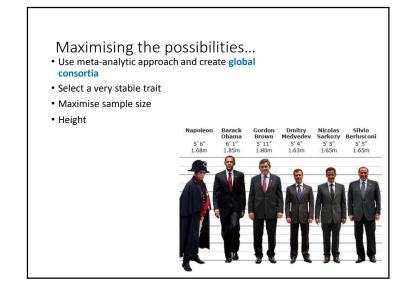
Solutions?

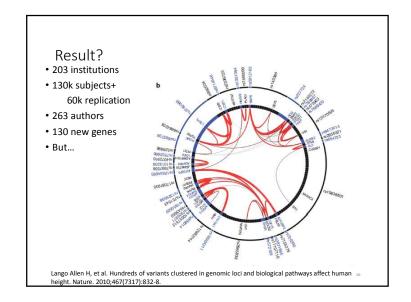
- Quantitative trait analysis
- Instead of going after the disease, why not investigate the trait that is related to the disease?
- ECG a recording of electrical impulse travel through the heart muscle

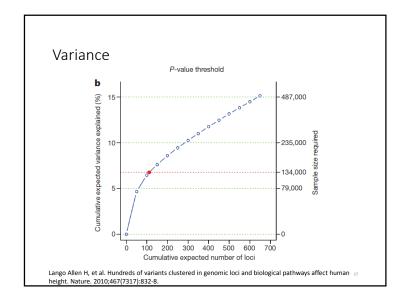


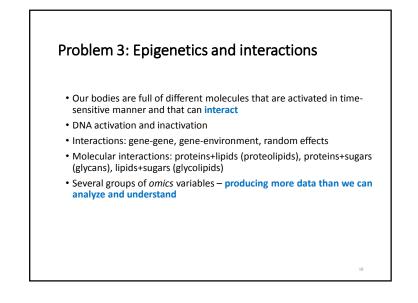


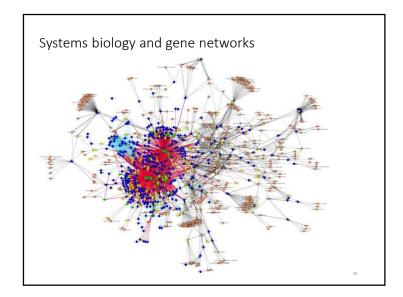


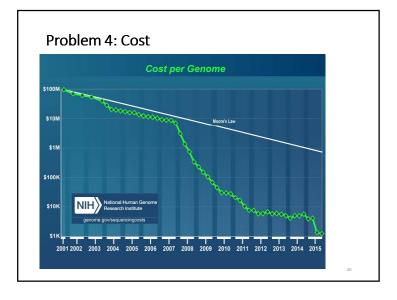






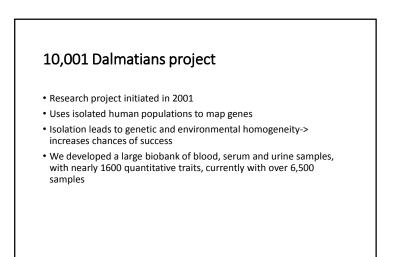










































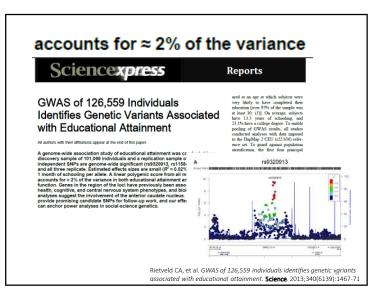


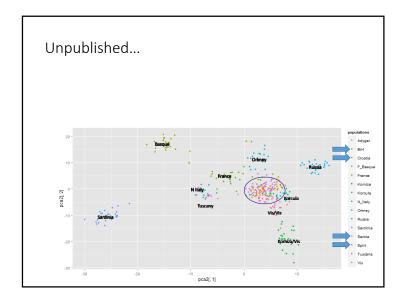












Elevated Risk @				
NAME	CONFIDENCE	YOUR RISK	AVG. RISK	COMPARED TO AVERAGE
Celiac Disease	****	2.7%	0.1%	23.02x †
Psoriasis	****	22.4%	11.4%	1.98x 🖛
Prostate Cancer O	****	28.2%	17.8%	1.58x 🖛
Atrial Fibrillation	****	33.9%	27.2%	1.25x 💳
Ulcerative Colitis	****	1.00%	0.77%	1.30x I
Parkinson's Disease	****	2.0%	1.6%	1.23x

