



Inequalities as Barriers for Global Sustainability: DNA and the Human Variome Project Sir John Burn

MD FRCP FRCPE FRCPCH FRCOG FMedSci Professor of Clinical Genetics, Newcastle University Genetics Lead, UK National Institute of Health Research Co-chair Scientific Advisory Gp., Human Variome Project Medical Director, QuantumDx group

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Jou wêreld, Jou koerant∗∗

Woensdag 15 Mei 2013 R7,50

moet weer wag 2 Eugene de Kock Ekstra bylae oor jagbedryf Binne

Aussie-l Cheeta



2 aspirin heredi

Long-term effect hereditary colore randomised cont

John Burn, Anne-Marie Gerdes, Finlay M Eamonn R Maher, Lucio Bertario, Marie-Patrick J Morrison, Victoria Murday, Raj I Faye Elliott, Mohammad Movahedi, Kirs of the CAPP2 Investigators

1993

rs at 5 yrs

k in carriers of from the CAPP2

ne Olschwang, Diane Eccles, D Gareth Evans, V Hodgson, Annika Lindblom, Jan Lubinski, Hans F Vasen, Gail Barker, Gillian Crawford, Jynch, John C Mathers, D Timothy Bishop, on behalf

Lancet 378 December 11th 2011

43 CAPP2 recruiting centres in 16 countries recruited 1009 mismatch repair gene defect carriers

UK

Aberdeen, Edinburgh, Glasgow, Newcastle, Leeds Sheffield, Manchester, Liverpool, Birmingham, Cardiff, Belfast,Oxford, Bristol, St Marks, St Georges, Guys-London Southampton, Exeter, Guildford, Worthing,

Rest of Europe

Finland, Sweden, Denmark, Germany, Belgium, Poland, Netherlands, France, Hungary, Switzerland, Portugal, Spain, Italy



Human Genomics Strategy Group report 2012



- Government response
 - 100,000 whole genomes
 - £100 million investment
 - Genomics StrategyBoard (NHS nowHEE)



Functional classification of JDW

SNPs

SNP T	Type
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Missense

Nonsense

Synonymous

UTH

Intror

2421

5

2261

7102

381924

August 2010 China National Highway 110 traffic jam, China, world's worst traffic jam ever....

more than 100 kilometres from August 14 - 26, inc. at least 11 days of total gridlock.



We must collect, store and interpret 3 million variants per person....

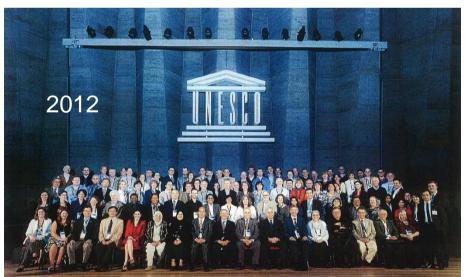




LOVD

THE HUMAN VARIOME PROJECT

An online reference database which will provide free access to all variants which have been assessed and their functional significance validated



www.thehumanvariomeproject.org

Nat Genet, 2007. 39(4): p. 433-6

Family at risk of breast cancer due to p53 mutation

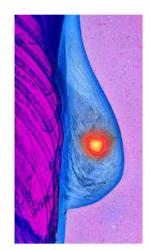


Hereditary cancer syndromes: opportunities and challenges

Patricia Ashton-Prolla

From São Paulo Advanced School of Comparative Oncology Águas de São Pedro, Brazil. 30 September - 6 October 2012

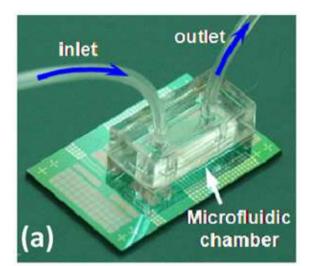


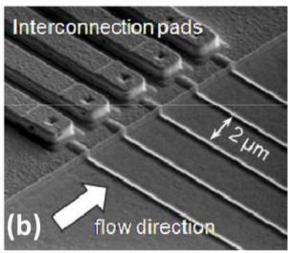


1 in 300 women in South Brazil carry p.R337H founder mutation in tp53 – a single DNA letter change causes 1 in 12 cases of all breast cancers

The much higher rate of childhood adrenal cancers has led to understanding how assembly of the protein is pH dependent



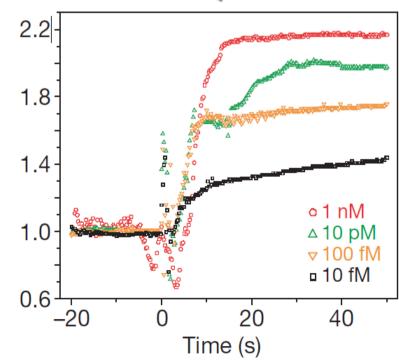




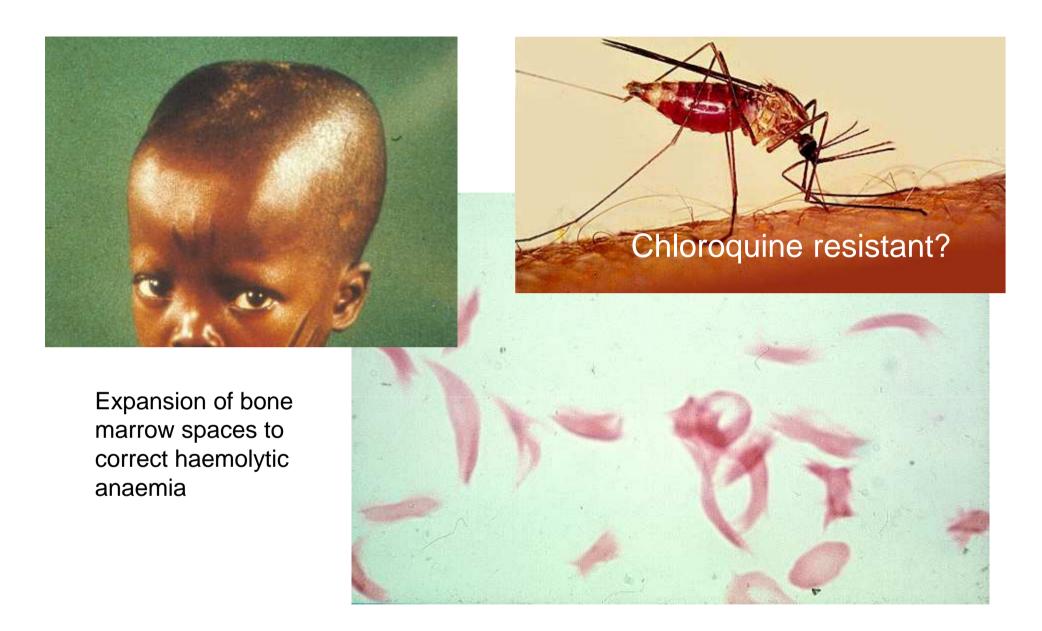
We will produce a cheap handheld device capable of analysing DNA at the point of care in 10 minutes



QuantuMDs



conductance change



Sickle Cell Disease Homozygous for a missense mutation



In memory of Fred Sanger

UNESCO DECLARATION 1997

Article 12 a) Benefits from biology, genetics and medicine, concerning the human genome, shall be made available to all, with due regard for the dignity and human rights of each individual.



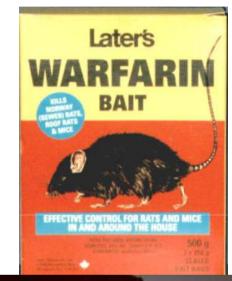
Double Nobel laureate 1918-2013

The man who showed us how to sequence DNA

pharmacogenetics

3 SNPs predict loading dose and likely dose needed to achieve INR

Population varies 20 fold yet we Treat everyone the same





Sequencing 2005

Using AB 3730 XL*



Washington University
Genome Sequencing Center



Baylor College of Medicine Human Genome Sequencing Center

*fluorescent Sanger chemistry on capillary sequencers