



# **Inequalities as Barriers for Global Sustainability: DNA and the Human Variome Project**

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# Beeld

Jou wêreld, Jou koerant • •

Woensdag 15 Mei 2013 R7,50

Ekstra bylae oor  
jagbedryf **Binne**

Eugene de Kock  
moet weer wag **2**

Aussie-  
Cheeta



Die lewe het  
uitdagings.  
ons nie ban  
te maak nie  
wat ons kan  
waaroor on  
beheer kan

— ANGELINA JOLIE  
BESLUIT OM 'N  
MASTEKTOMIE T



2 aspirin tablets/day for 2 yrs reduced  
hereditary colorectal cancer risk at 5 yrs

Long-term effect  
hereditary colorectal cancer  
randomised controlled trial

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

risk 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,  
ynch, John C Mathers, D Timothy Bishop, on behalf*

1993

**Lancet 378**  
**December 11<sup>th</sup> 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

**Toronto**

**Sao Paulo**

**Cape Town**

**Hong Kong**

**Australia**

Melbourne  
Brisbane  
Newcastle  
Perth




# Human Genomics Strategy Group report 2012



- Government response
  - 100,000 whole genomes
  - £100 million investment
  - Genomics Strategy Board (NHS now HEE)

# Functional classification of JDW SNPs

SNP Type		Novel SNPs
Missense		2421
Nonsense		5
Synonymous		2261
UTR		7102
Intron		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](http://www.thehumanvariomeproject.org)***

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



# HVP country nodes and support

6

- Country Nodes
- Support



## Family at risk of breast cancer due to p53 mutation



**Diane Carol**

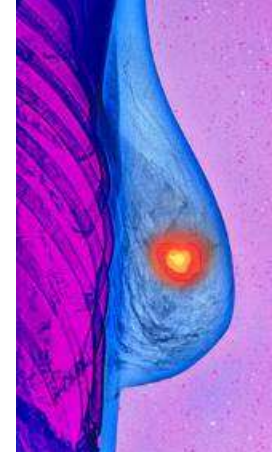
Varley et al Oncogene 1998



# 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

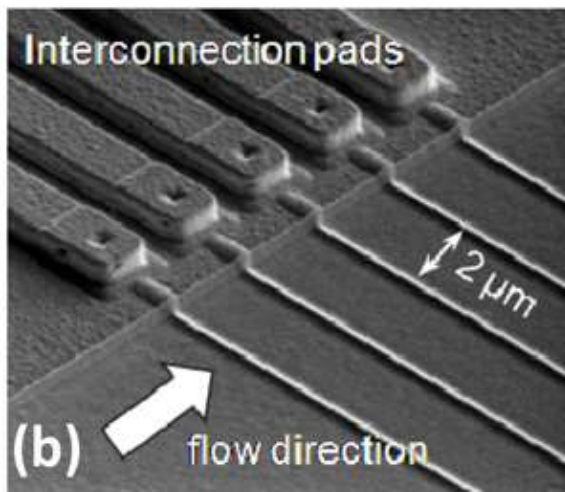
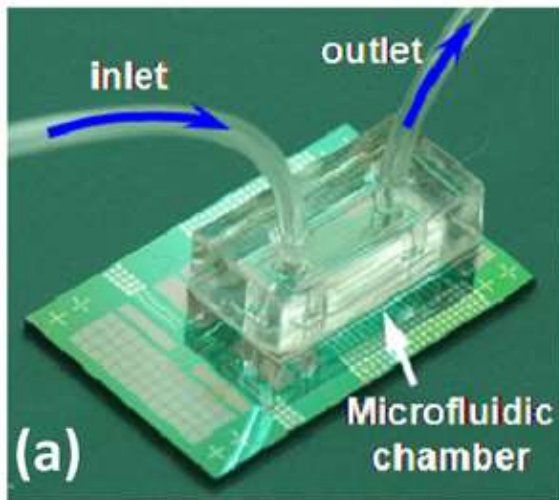


1 in 300 women in South Brazil carry  
p.R337H founder mutation in tp53 – a  
single DNA letter change causes  
1 in 12 cases of all breast cancers

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





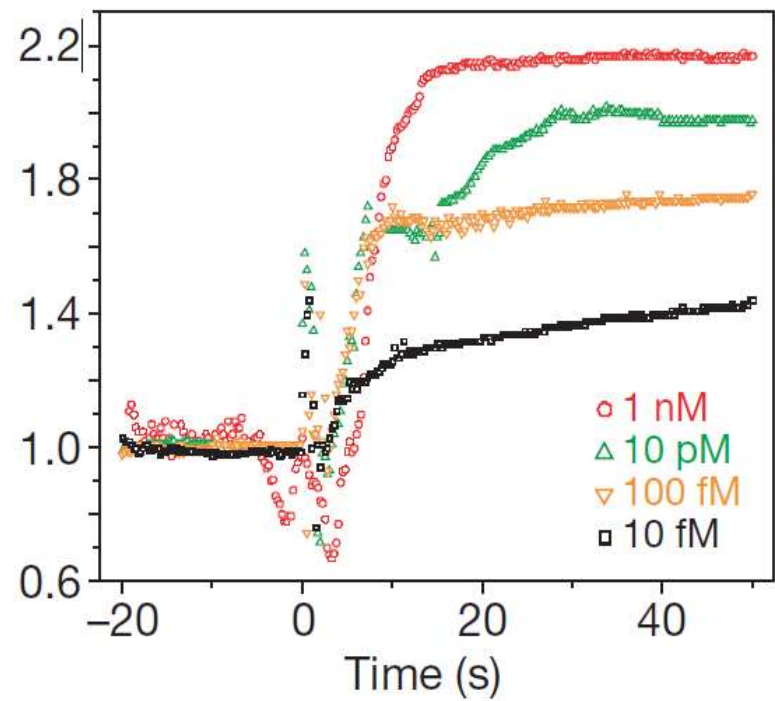


We will produce a cheap handheld device capable of analysing DNA at the point of care in 10 minutes



QuantuMD<sub>s</sub>

Normalised  
conductance change







Expansion of bone marrow spaces to correct haemolytic anaemia



# Sickle Cell Disease

## Homozygous for a missense mutation

T



Pe  
So



ld



# 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