#### "Next-Generation Sequencing (NGS):future perspectives "

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#### Biologists modus operandi

- Observing a phenomenon that is in some way interesting or puzzling.
- Making a guess as to the explanation of the phenomenon.
- Devising a test to show how likely this explanation is to be true or false.
- Carrying out the test, and, on the basis of the results, deciding whether the explanation is a good one or not. In the latter case, a new explanation will (with luck) 'spring to mind' as a result of the first test.

http://www.biology.ed.ac.uk/archive/jdeacon/statistics/tress2.html

#### The Observed phenomenon



#### Selection of test times



#### But was is the real event?



#### Sometimes you could be lucky



"Positive" results are used "negative" rejected Why? Only positive results are publishable

## Things have changed: NGS

- 454 (Pyrosequencing)
- Illumina
- Solid
- Pacific Biosciences
- Oxford Nanopores
- Many more

#### Next Generation techniques



## Next Generation Sequencing



#### Ion Torrent: "Personal Genome Machine".



#### ■ LIFE TECHNOLOGIES CORPORATION

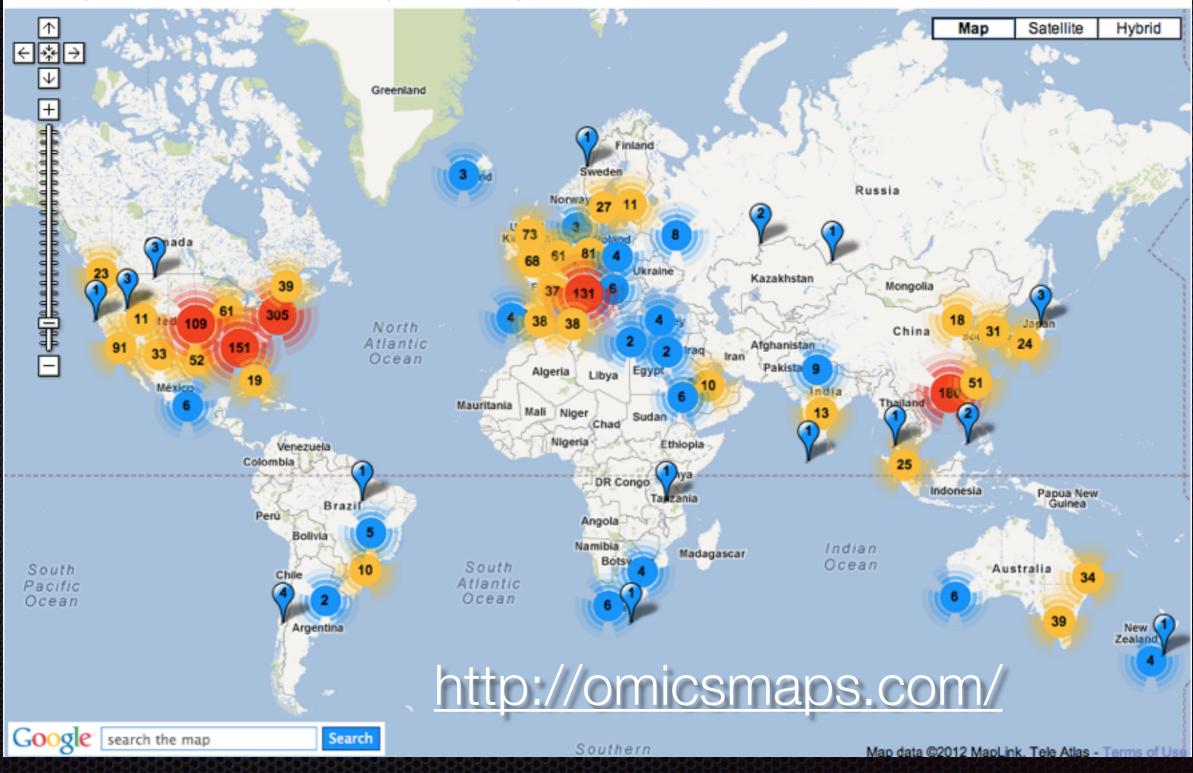
#### Oxford Nanopore



http://www.nanoporetech.com/

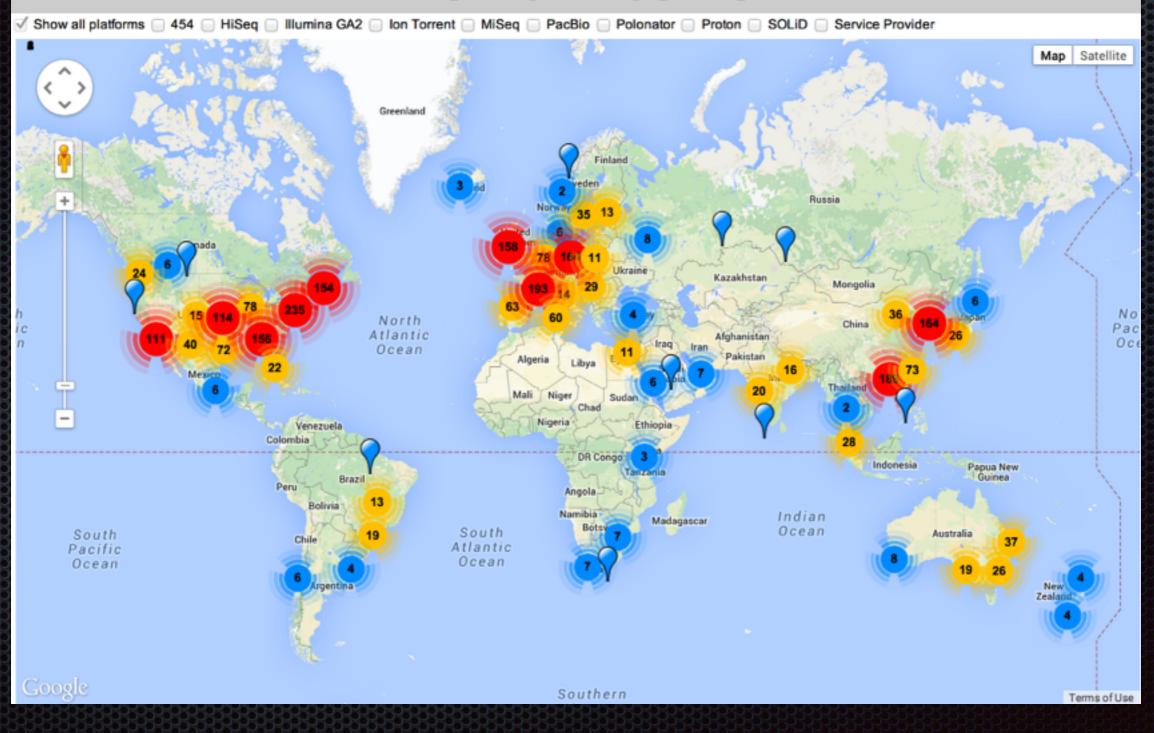
#### World NGS Map

Show all platforms 📃 Illumina GA2 📃 Illumina HiSeq 📃 Illumina MiSeq 📃 Ion Torrent 📃 PacBio 📃 Polonator 📃 Roche/454 📃 SOLiD 📃 Service Provider



#### Today

#### Next Generation Genomics: World Map of High-throughput Sequencers



#### NGS SUCCESS STORIES





Tomato Genome

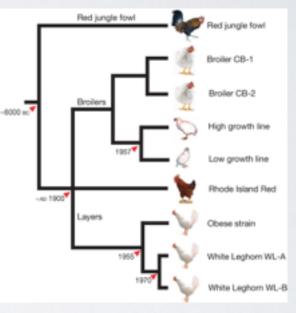


Mammoth Genome



nature

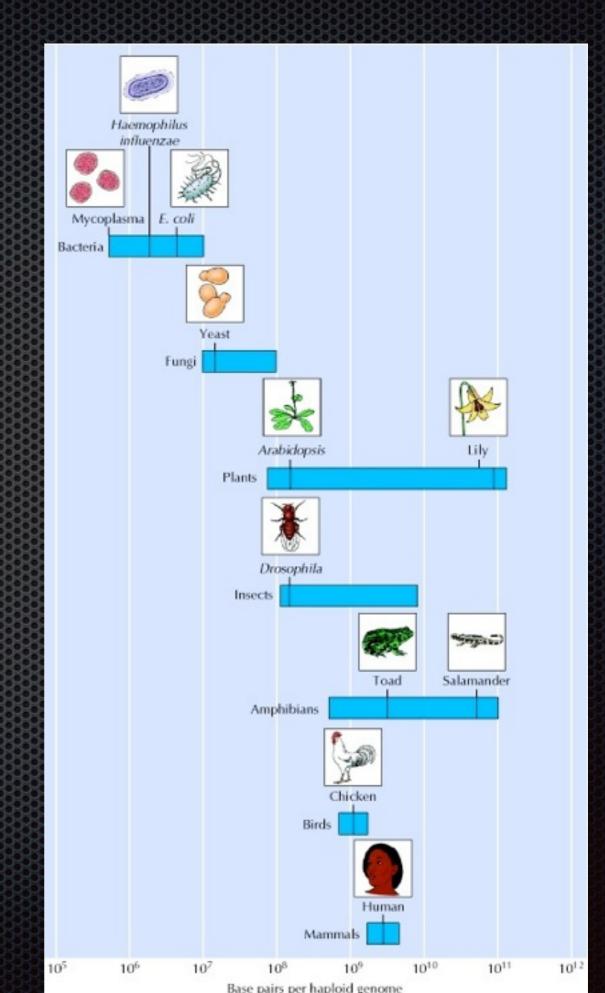
Mammoth Genome Chicken domestication



## One of the Centres: BGI

- The incredible productive efficiency of BGI has moved huge projects on human and other life forms variability from imagination to actual reality.
- The Center is taking a leading role in sequencing 10,000 vertebrates through the Genome 10K project; 5,000 insects and other arthropods through the i5k initiative; and more than 1,000 birds, including some extinct ones in a separate project.
- BGI has launched the Million Project, aimed to expand the sequenced samples to 1M for human, animals, plants and bacteria

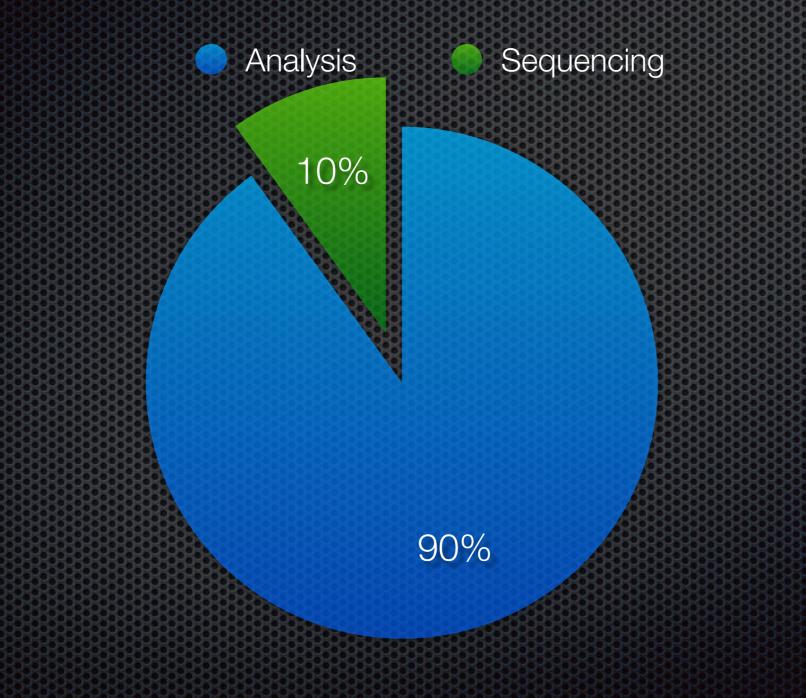
#### Genome sizes



## Is wonderful! Or?

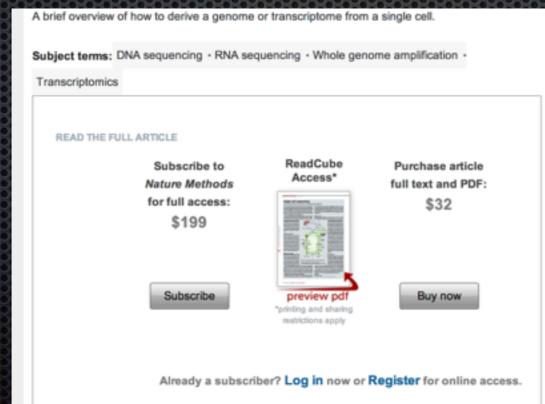
- Sequence without knowledge connected to it is worth:
- The deluge of data produced by these hordes of machines worldwide demand automatic workflows
- Complete new systems to shuffle data around
- Storage of never used amounts
- Machines with gigantic amounts of RAM
- BIOINFORMATICIANS!!

## COSTS



## OTHER PROBLEMS

- NOmenclature
- Publishing culture
- Moving target development
- Old ways of work and resistance to changes in culture



#### New challenges



#### today TB of data soon PB

#### UPPMAX/UPPNEX



#### 5 PB of Storage

#### So how much is a PB?

A petabyte contains enough information to fill a 45 mile high stack of CDs, the distance between SLAC and Berkeley Lab.

# We are in the ERA of EXABYTE

which is...

**1 EXABYTE** 

1024 PETABYTES 1,048,576 TERABYTES 1,073,741,824 GIGABYTES 1,099,511,627,776 MEGABYTES

## BIG DATA

#### WHAT IS BIG DATA?



Data Complexity: Variety and Velocity

#### EXOME

- Whole-exome sequencing has already been used for identifying the molecular defects of single gene disorders, for elucidating some genetically heterogeneous disorders and for improving the accuracy of diagnosis of patients.
- data for whole-exome sequencing is orders of magnitude smaller than for whole-genome sequencing
- On average, whole-exome sequencing identifies 12 000 variants in coding regions, of which ~90% are found in publicly available databases. In comparison, ~5 million variants, including 144 000 new variants, are reported on average by whole-genome sequencing.

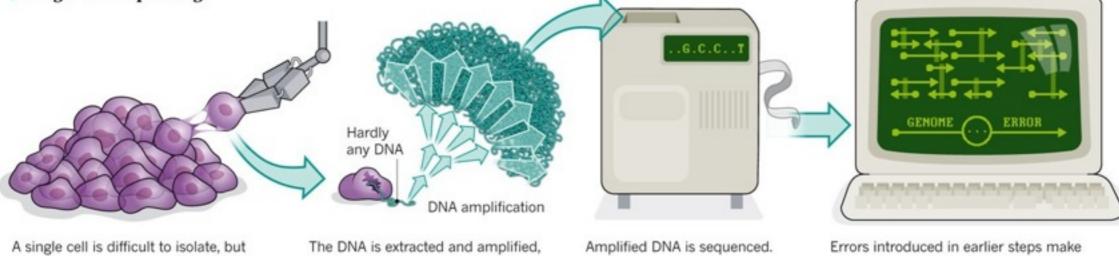
## EXOME

- OMIM the catalog of Mendelian disorders, lists >3000 disorders where the molecular basis has been reported. Still, >3500 disorders are listed where the genetic cause remains unknown and has yet to be identified
- Other main fields of interest for human geneticists are complex diseases and cancer. For example, the usage of NGS has led to the identification of driver mutations for specific types of cancer. Single cell sequencing.

#### Singel cell sequencing

#### **ONE GENOME FROM MANY**

Sequencing the genomes of single cells is similar to sequencing those from multiple cells - but errors are more likely. Standard genome sequencing A.T.C. Loads of DNA GENOME COMPLETE A sample containing thousands to DNA is extracted from all the nuclei. DNA is broken into fragments The sequences are assembled to give a millions of cells is isolated. and then sequenced. common, 'consensus' sequence. Single-cell sequencing

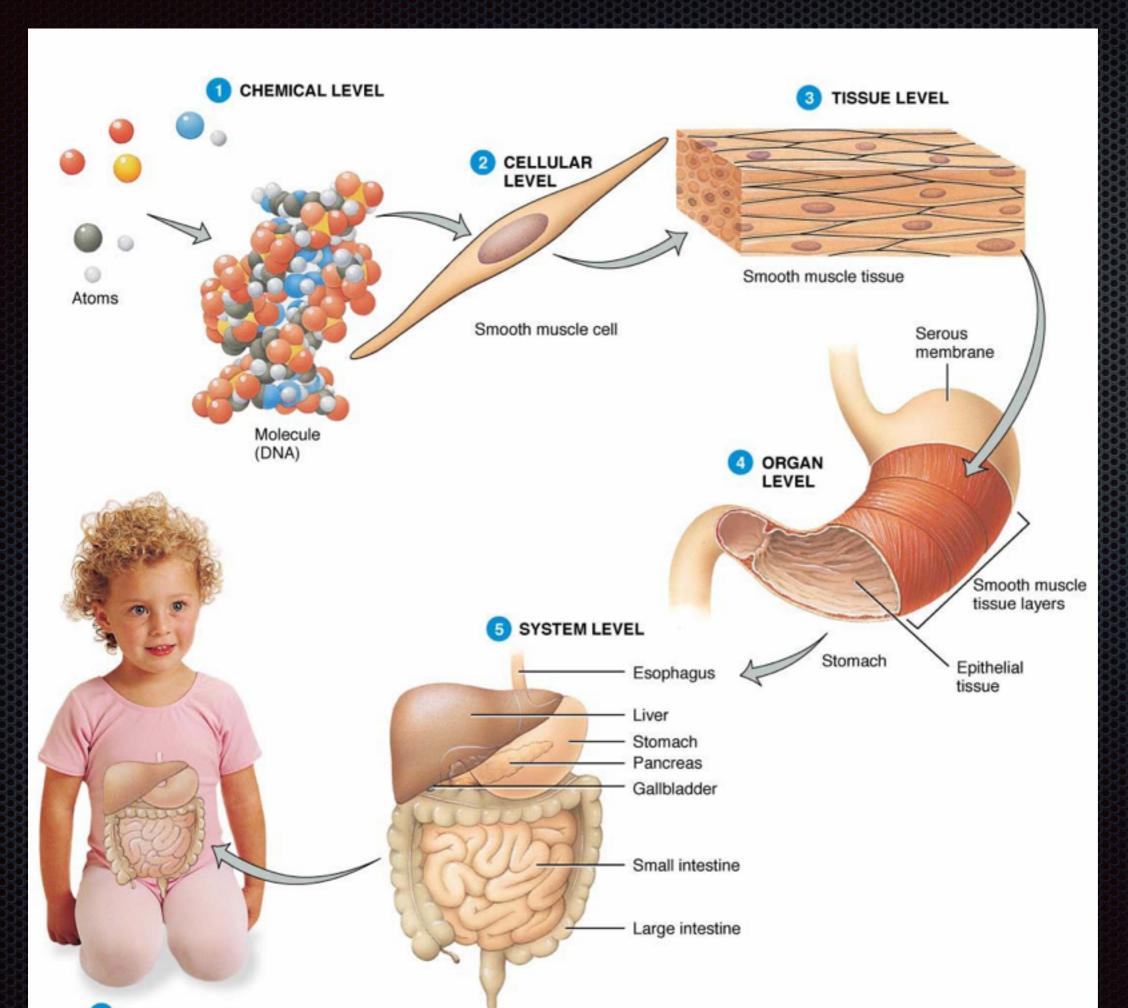


it can be done mechanically or with an automated cell sorter.

from www.nature.com

during which errors can creep in.

sequence assembly difficult; the final sequence can have gaps.



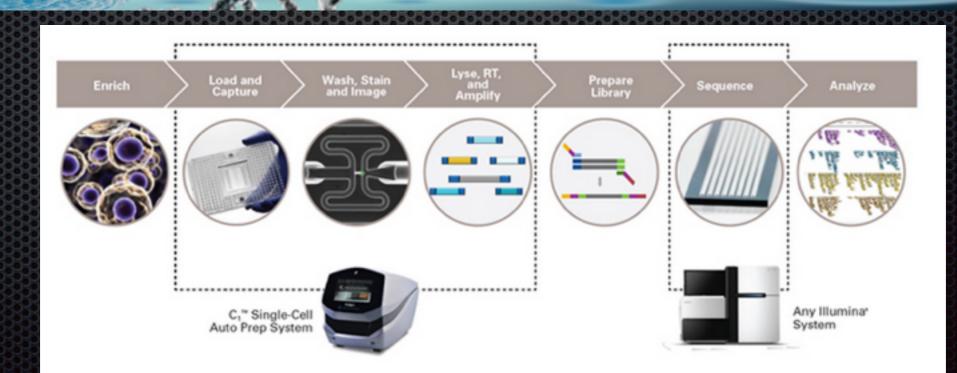


#### Singel Cell mRNA Sequencing

#### REVEAL HIDDEN VARIATION C1<sup>™</sup> Single-Cell DNA Sequencing

One workflow to discover genetic variants in individual cells





# Epigenomics and Epigenetics

- Epigenomics deals with the chemical modifications (e.g., 5' methylation) of DNA and RNA and the impacts of such changes on levels of gene expression.
- Tumor methylation profiles could also be detected in body fluid specimens showing a promising role as noninvasive markers for cancer diagnosis towards an early detection and potentially for the surveillance of cancer patients in a near future. However, the epigenomic exploration of cancer has only just begun.

#### Microbiome

A microbiome is the totality of microbes, their genetic elements (genomes), and environmental interactions in a particular environment.



http://www.secondgenome.com

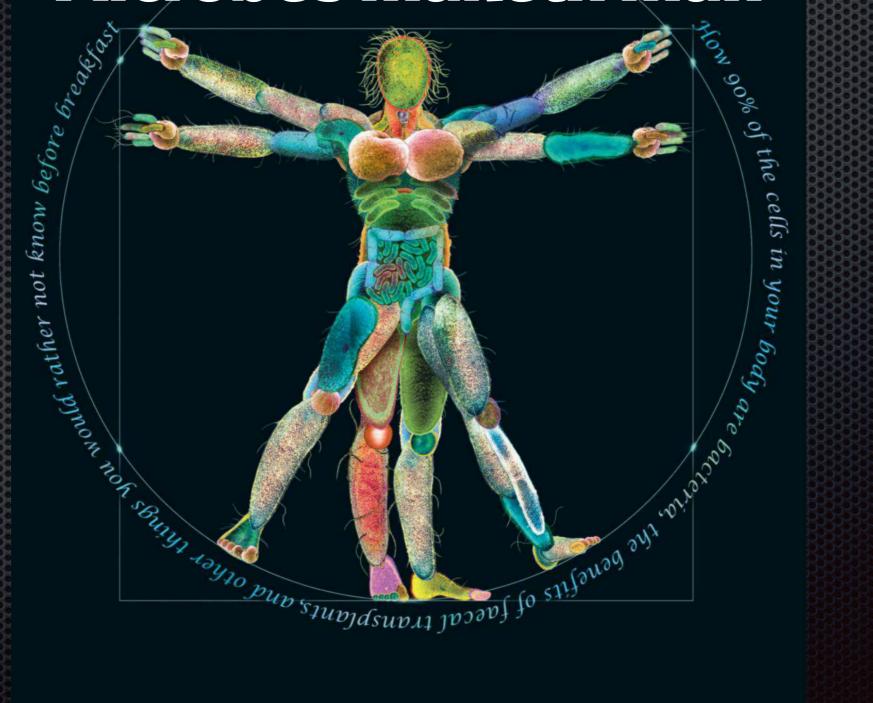
#### The Economist

The Catholic church's unholy mess Paul Ryan: the man with the plan Generation Xhausted China, victim of the Olympics? On the origin of specie

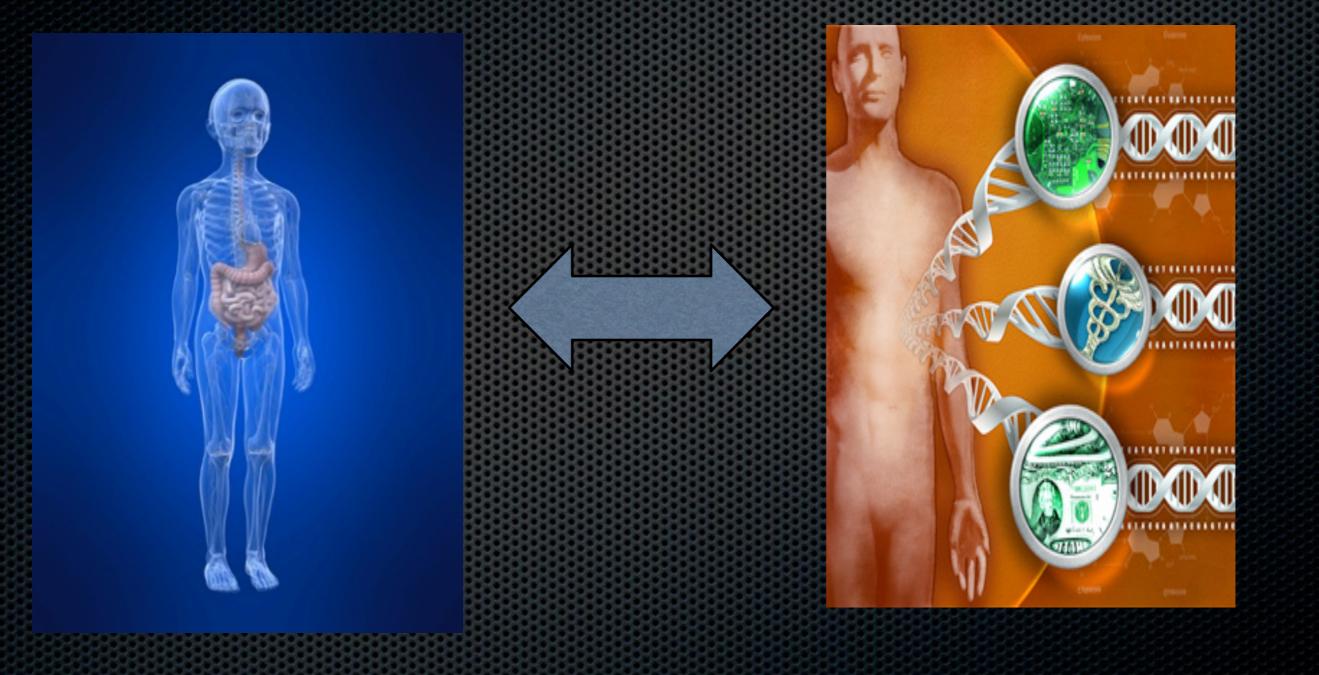
AUGUST 18TH-24TH 2012

#### Microbes maketh man

Economist.com



## Epigenetics



#### Fat and lean

- Metabolic effects of transplanting gut microbiota from lean donors to subjects with metabolic syndrome.
  A. Vrieze et al, EASD abstracts, 24 September 2012.
- The result was: Lean donor faecal infusion improves hepatic and peripheral insulin resistance as well as fasting lipid levels in obese individuals with the metabolic syndrome

#### New applications

Only imagination will put the limits of what its possible to be done using Next Generation Technologies!

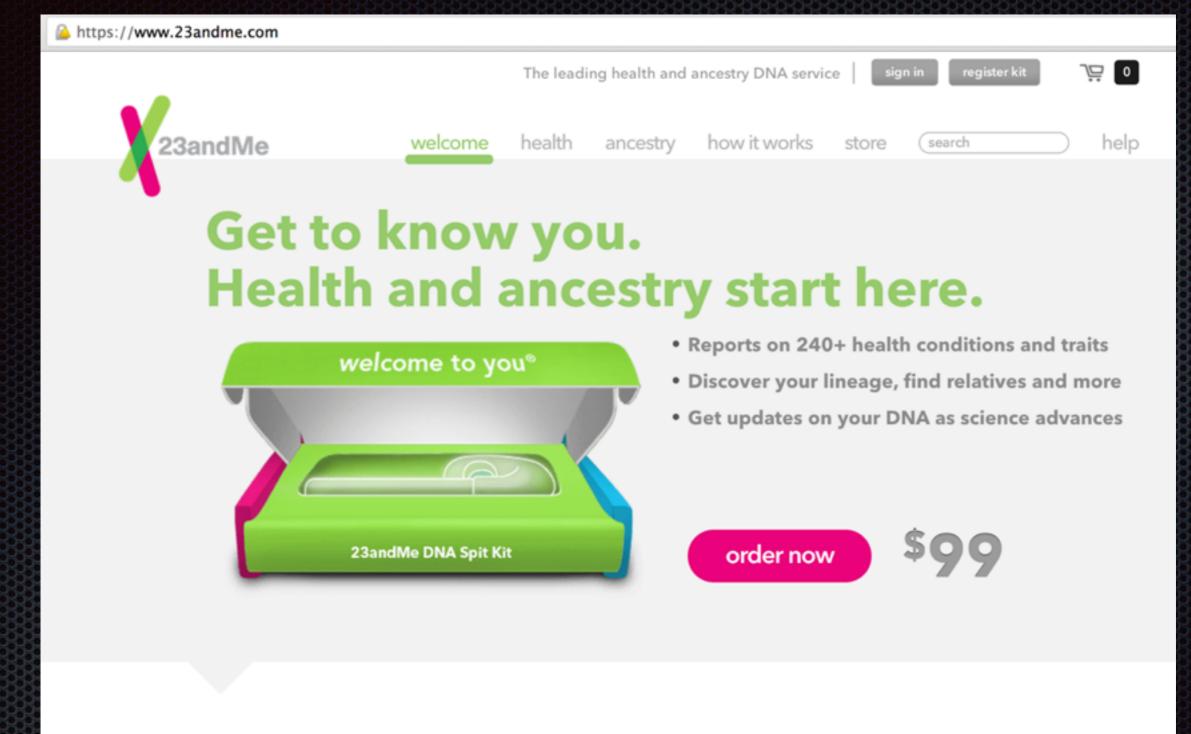
# Human Medicine Success Stories

- Whole-genome sequencing in autism identifies hot spots for de novo germline mutation. Cell. 2012 Dec 21;151(7):1431-42. doi: 10.1016/j.cell.2012.11.019.
- Genes contributing to pain sensitivity in the normal population: an exome sequencing study. PLoS Genet. 2012 Dec;8(12)

#### 23andMe: Linda Avey and Anne Wojcicki

andMe genetics jus	st got personal.				Search 23andMe		
home	our service	genetics 101	store	ab	about us		
23	BandMe Dem	nocratizes Pers	onal Gei	netics.			
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Oct. 8, 2008: 23andMe new, more informative Traits view, 1 new surv	Health and Cano	2, 2008 - 23andMe Announces Breast ser Initiative		How the Proce Frequently Ask			

#### Complete List of Health and Trail



#### Watch a life-changing story.

# Take a more active role in managing your health

Knowing how your genes may impact your health can help you plan for the future and personalize your healthcare with your doctor.

#### **Order Now** »





Plan for the future

Find out if your child will be at risk for 44 inherited conditions and learn about steps you can take. about carrier status



#### Be on the lookout now

Knowing your health risks will help you and your doctor figure out health areas to keep an eye on. about disease risks



.....

#### Plan with your doctor

Personalize your healthcare by knowing in advance how you will respond to certain medications like Warfarin. about drug response

### Future



- Cell-free fetus DNA sequencing in the era of NGS: New opportunities for pre-natal diagnosis
- The main advantages in detecting free fetal DNA from maternal blood are the quick and early response and the total absence of risk for the fetus itself. In fact fetal DNA could be detected as early as the 6th week of of pregnancy and the test simply requires a sample of mother's blood.

# Share your genome sequence using your smartphone

 A smartphone app that can store your DNA – and perhaps one day allow your partner, your relatives or your doctor to scan it



# Top 6 Deadly Diseases

- Trachea, Bronchus, Lung Cancers
- Malaria
- TBC
- Diarrheal
- Perinatal Conditions
- Chronic Obstructive Pulmonary Disease

### High technology everywhere!



#### **Field data collection**



### Connectomics

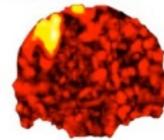


### Connectomics

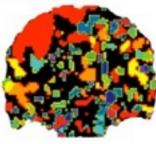


Student's t-test overlay

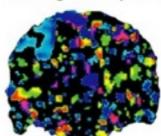
Fisher's g max statistic



Frequency of g max

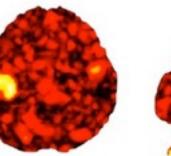


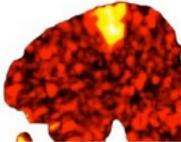
Phase of g max components



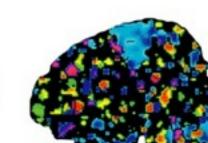


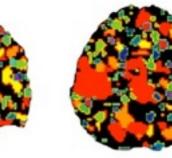




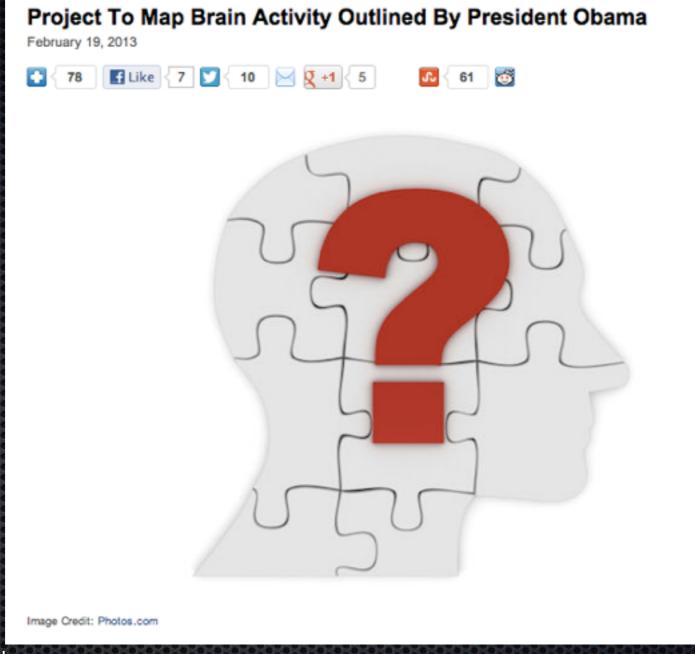








# USA:



#### HGP: \$3.8 billion over 13 years.

Map Brain project: \$3 billion

# EU:€100M/year for ten years



#### The Human Brain Project has been selected as a European Flagship!



#### EUROPEAN PARTNERS/



Press kit
The HBP Report

# THANKS!!