

14:05-14:25 Foodomics 2.0 Wim Van Criekinge Ghent University



























Lab for Bioinformatics and computational genomics

30 "genome hackers" mostly engineers (statistics)





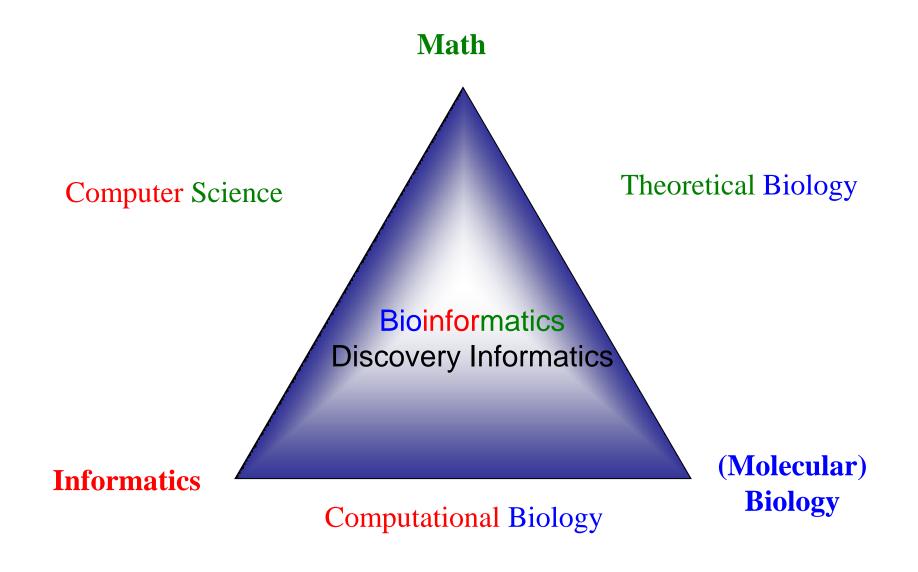
scientists, technicians, geneticists, clinicians

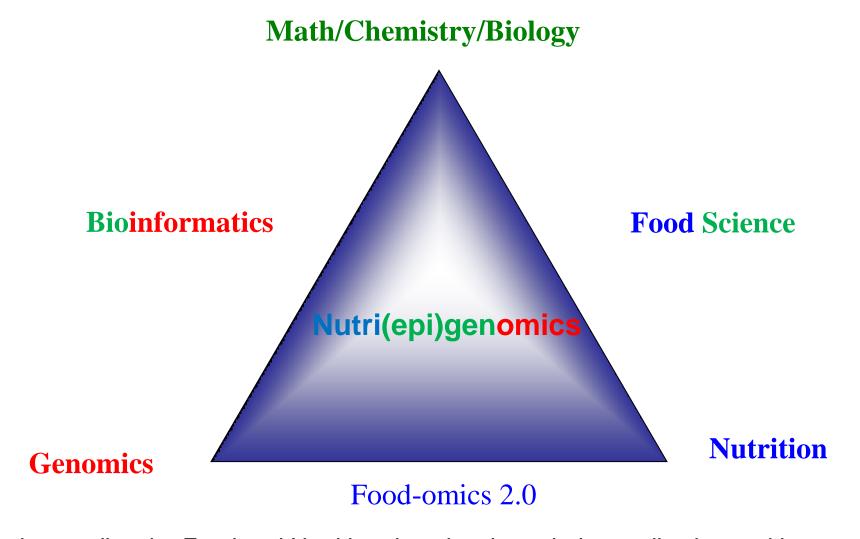


dewpal/aerolis



>100 people Hardware/software engineers, mathematicians, molecular biologists





"a discipline that studies the Food and Nutrition domains through the application and integration of advanced -omics technologies to improve consumer's well-being, health, and knowledge".



"The future is already here,... it's just not evenly distributed."

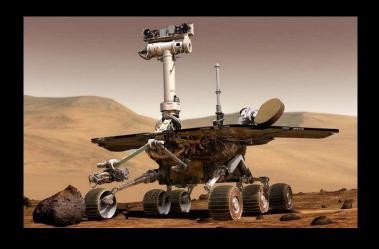
William Gibson, Futurist | Cyber-Visionary





"The future is already here,... it's just not evenly distributed."

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Foodomics 2.0

- Genetics
 - Nutrigenomics
 - Examples
- Epigenetics
 - Nutri-epigenetics
 - Examples
- Metagenomics
 - Nutri-metagenomics
 - Examples
- Foodomics 3.0
 - Technology
 - Big Data / Al

5 years ago ...





Home / Makers C4M / Personalized Proteins

Personalized Proteins



Wobblebase develops apps that allow the user to manage and explore his or her own genome. The apps aid in the interpretation of those variations /mutations with the highest probability of having a phenotype. In order to make this more tangible we allow the user to 3D print his own protein sets.

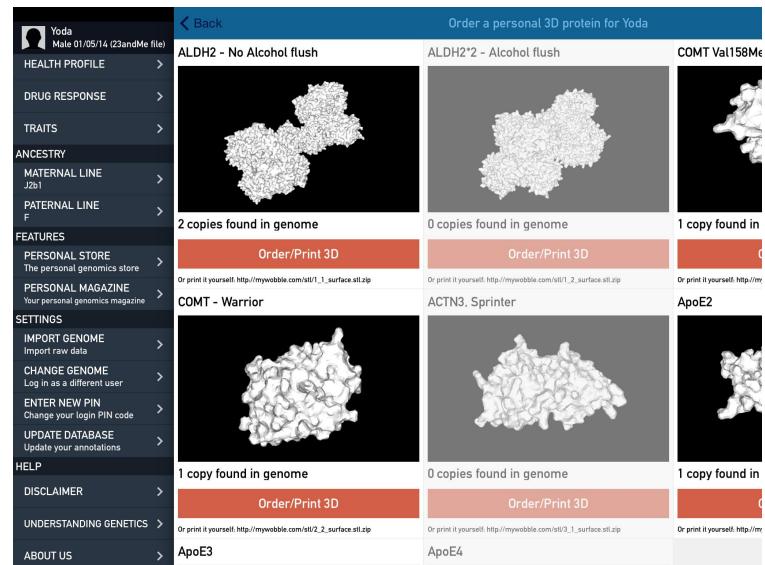
Team

Personalized Proteins

Geert Trooskens, ?Bioinformatician and app developer

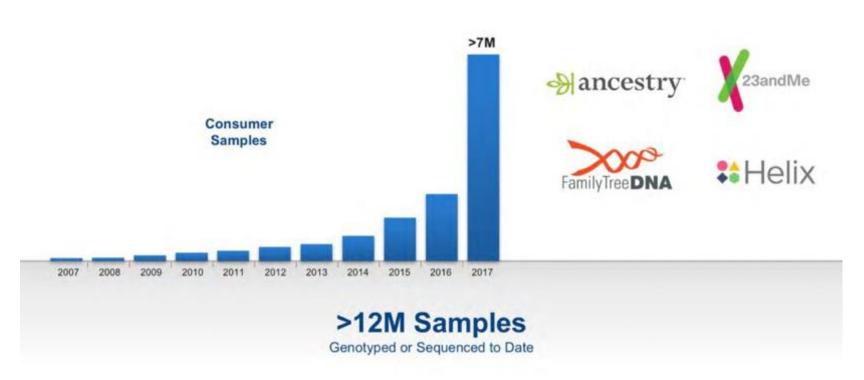
Peter Schotte, Senior scientist with biotech experience Wim Van Criekinge, Bioinformatician and entrepreneur

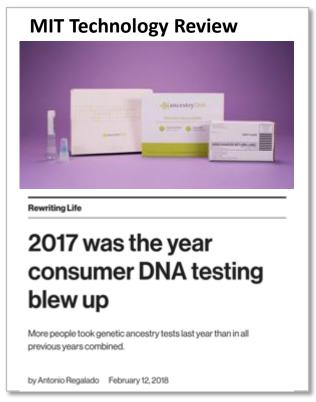
Website



Consumer Genomics Market Expanding Rapidly

Consumer Genomics Inflecting





What can Consumer Genetic Testing Offer?

Applied & Translational Genomics 8 (2016) 16-22



Contents lists available at ScienceDirect

Applied & Translational Genomics

journal homepage: www.elsevier.com/locate/atg



'Only a click away — DTC genetics for ancestry, health, love...and more: A view of the business and regulatory landscape'



IIIC

Andelka M. Phillips

DPhil Candidate, Faculty of Law, University of Oxford, United Kingdom



Categories of tests offered by DTC companies.

Category	Number of companies	Percentage
Ancestry	74	30%
Athletic	38	15%
Child talent	4	2%
Matchmaking	3	1%
Surreptitious	34	14%
Nutrigenetic	74	30%
Non-legal paternity	88	36%
Legal paternity	83	34%
Genetic relatedness	92	37%
Carrier	27	11%
Only health testing	31	13%
Total companies analyzed	246	

38.6%

Sub-Saharan African

20.5%

East Asian

Source: https://www.sciencedirect.com/science/article/pii/S2212066116300011

Wobblebase's mission is to bridge the gap between consumers and medical professionals to leverage genetic information to its fullest potential.



All-in-One Interface



Genetic Traits



Health Risk

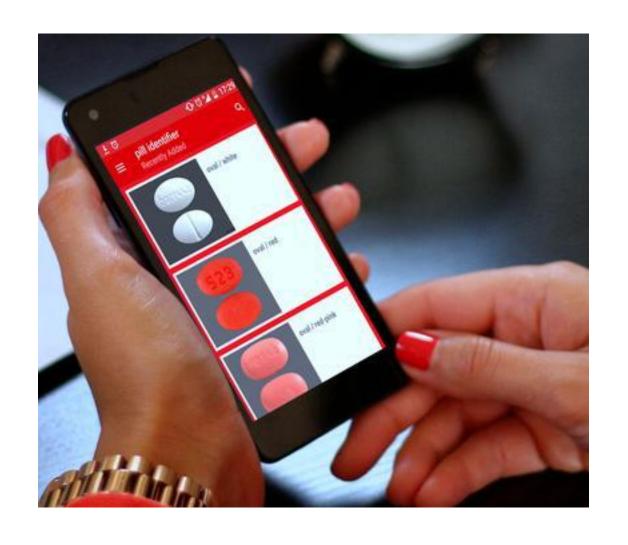


Pharmacogenetics



Genealogy





Use camera to identify Pill

Get "Insert" from database

Get pharmacogenomics SNPs

Check your SNPs

AR / Report



Use barcode to identify product

Get ingredients

Check your SNPs against ingredients

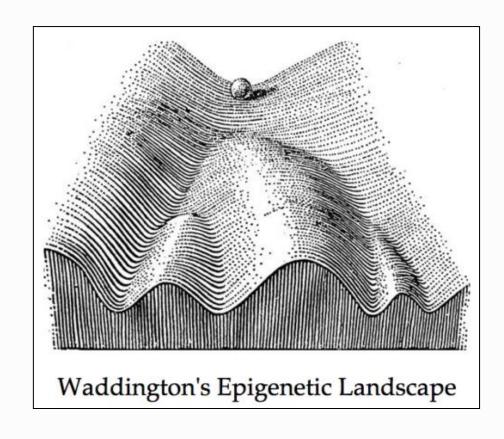
- Sensitivities
 - eg peanut
 - allergens
- Metabolic compatibility

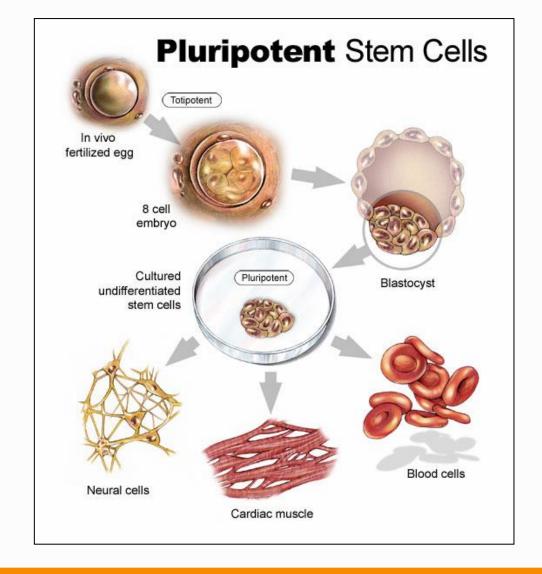
AR / Report

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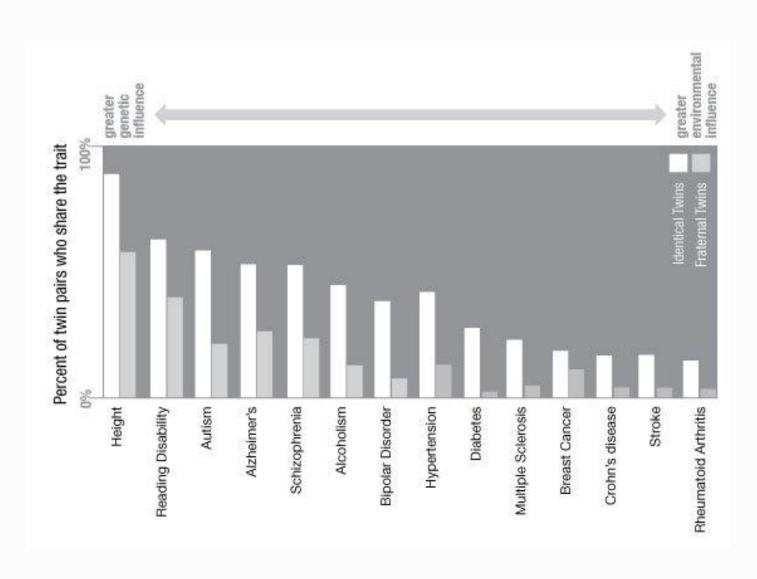
Relative importance of (Epi)genetics (i)





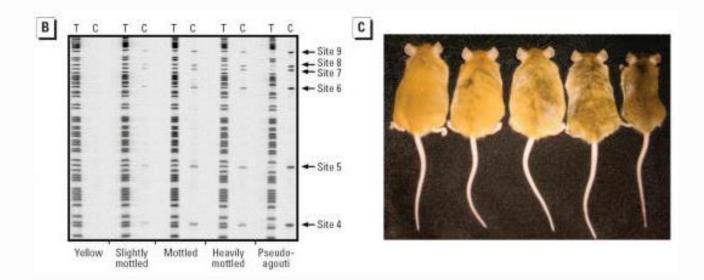
Relative importance of (Epi)genetics (ii)

- Epigenetics enables to reuse one genome for many different purposes
- Epigenetics driving etiology of many human diseases

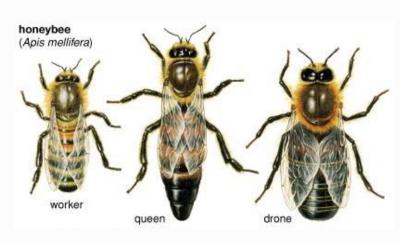


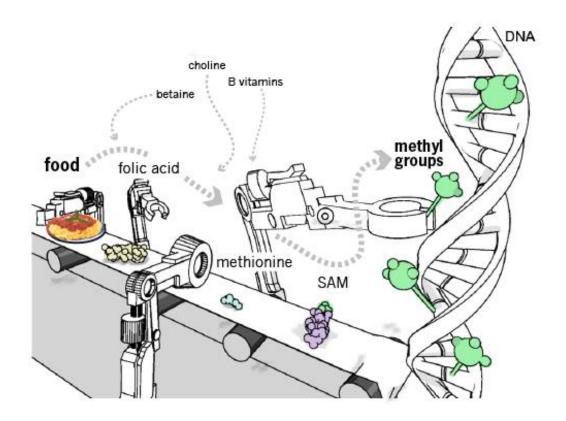
Relative importance of (Epi)genetics (iii)

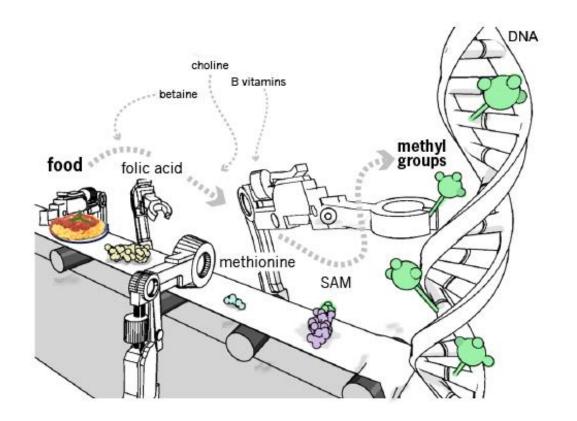
 Actionability* and integration of intrinsic with environmental signals



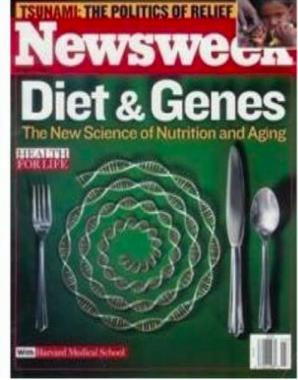


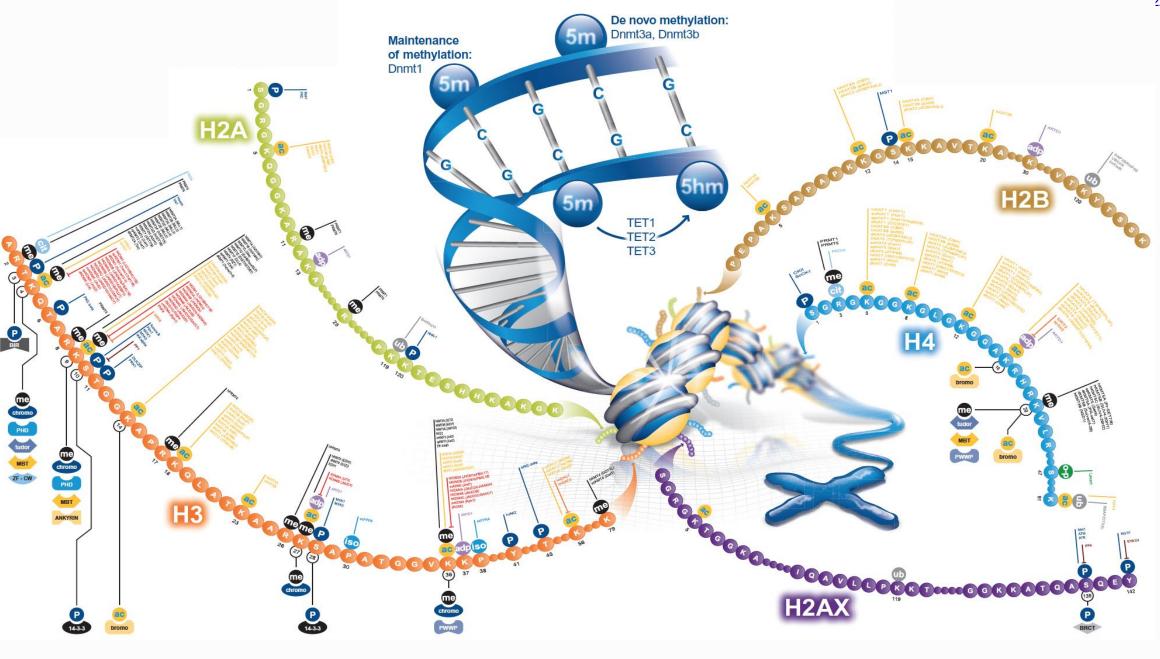












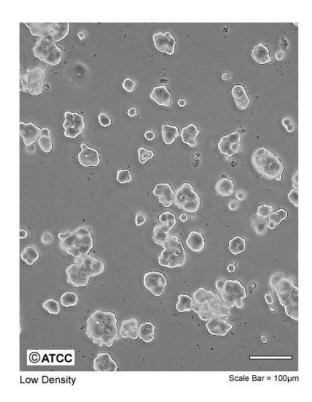
Phyto-oestrogens



Structure – natural products are

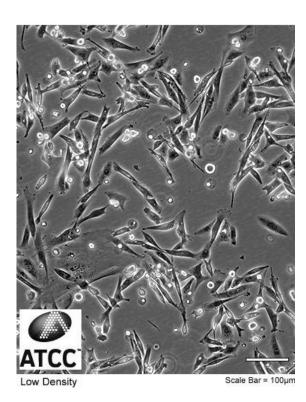
- α- en β-receptors
- Epigenetic modifier ?

Method



MCF-7 cellen

ER+

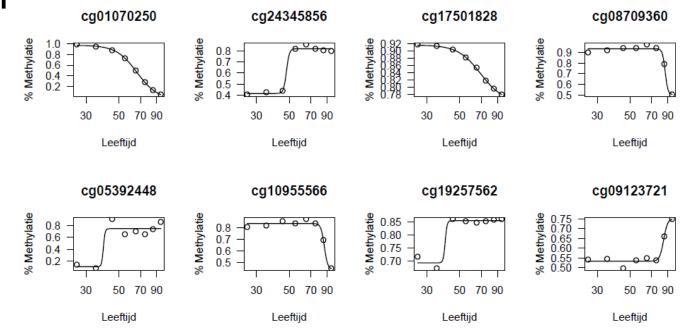


MDA-MB-231 cellen

ER-

Data analysis

- 450K Beadchip array (Illumina)
- Dose-response curves with active compound (R-PNI)





Received: 23 April 2018

Accepted: 24 August 2018

Published online: 11 September 2018

OPEN Exploratory analysis of the human breast DNA methylation profile upon soymilk exposure

Louis Coussement¹, Selin Bolca¹, Wim Van Criekinge^{1,2,3}, Geert Trooskens¹, Klaas Mensaert¹, Katrien Poels⁴, Nathalie Roche⁵, Phillip Blondeel⁵, Lode Godderis⁴, Herman Depypere⁶ & Tim De Meyer^{1,2,3}

Upon soy consumption, isoflavone metabolites attain bioactive concentrations in breast tissue possibly affecting health. Though in vitro epigenetic activity of soy metabolites has been described, the in vivo impact on the epigenome is largely unknown. Therefore, in this case-control study, the breast glandular tissue DNA methylome was explored in women undergoing an aesthetic breast reduction. After a run-in phase, 10 generally healthy Belgian or Dutch women received soymilk for 5 days. MethylCap-seq methylation profiles were compared with those of 10 matched controls. Isoflavones and their microbial metabolites were quantified in urine, serum, and glandular breast tissue (liquid chromatography-mass spectrometry) and 17β -estradiol in glandular breast tissue (immunoassay). Global DNA methylation levels were obtained for 6 cases and 5 controls using liquid chromatographymass spectrometry. Although lower MethylCap-seq coverages were observed, mass spectrometry results and computational LINE-1 methylation analysis did not provide evidence supporting global methylation alterations upon treatment. At a false discovery rate of 0.05, no differentially methylated loci were identified. Moreover, a set of previously identified loci was specifically tested, but earlier reported results could not be validated. In conclusion, after a 5-day soymilk treatment, no major general epigenetic reprogramming in breast tissue could be found in this exploratory study.

Evidence based phytotherapy



Fermented Soy

Active metabolite **Equol**

Equal: natural agonist **ER-beta** receptor

anti-proliferation and anti-inflammatoir
effect



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We Are More Bacteria Than Human

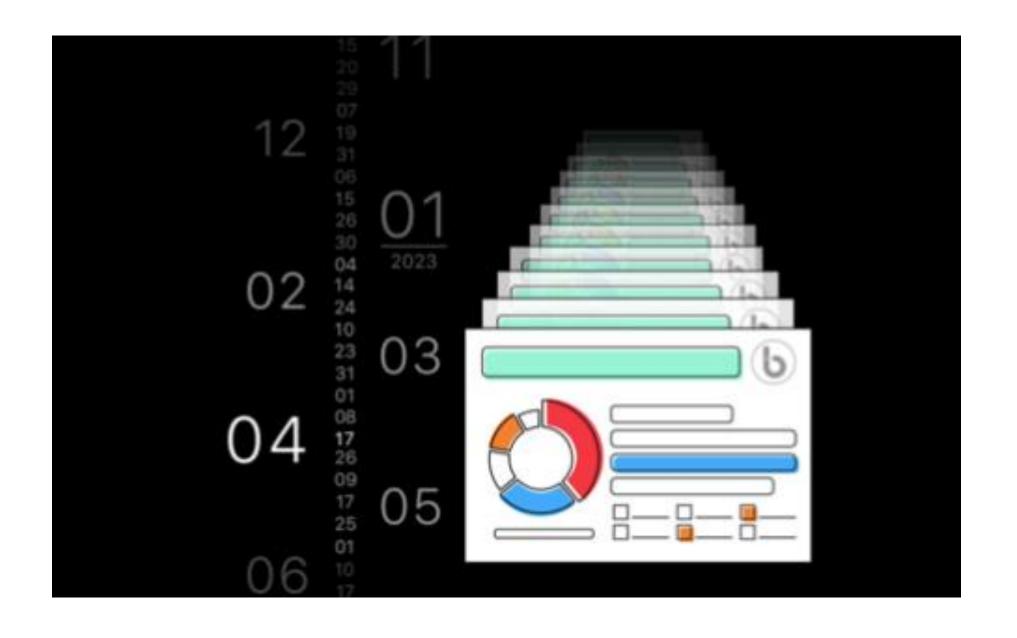
- Healthy adult harbors ~100 trillion bacteria in gut alone (10x the number of human cells we possess)
- Communal gut microbial genome (microbiome) is
 ~150 times larger than human genome
- Reasonable to view microbiome as an organ
- Is dominated by 4 large groups of bacteria: Actinobacteria, Bacteroidetes, Firmicutes, Proteobacteria

uBiome











bactoyou bac-up pills

takes you bac to FEBRUARY 24TH, 2023

100mg x 7 of sample # 000118632

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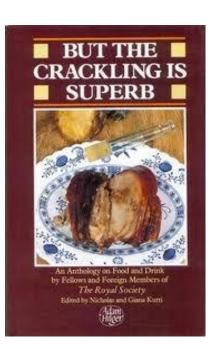


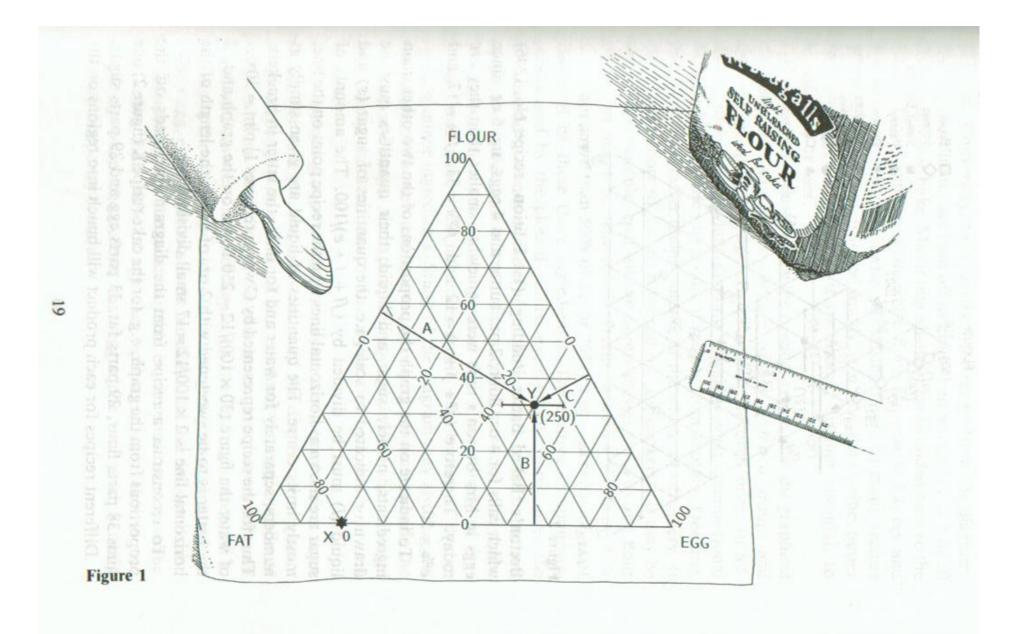


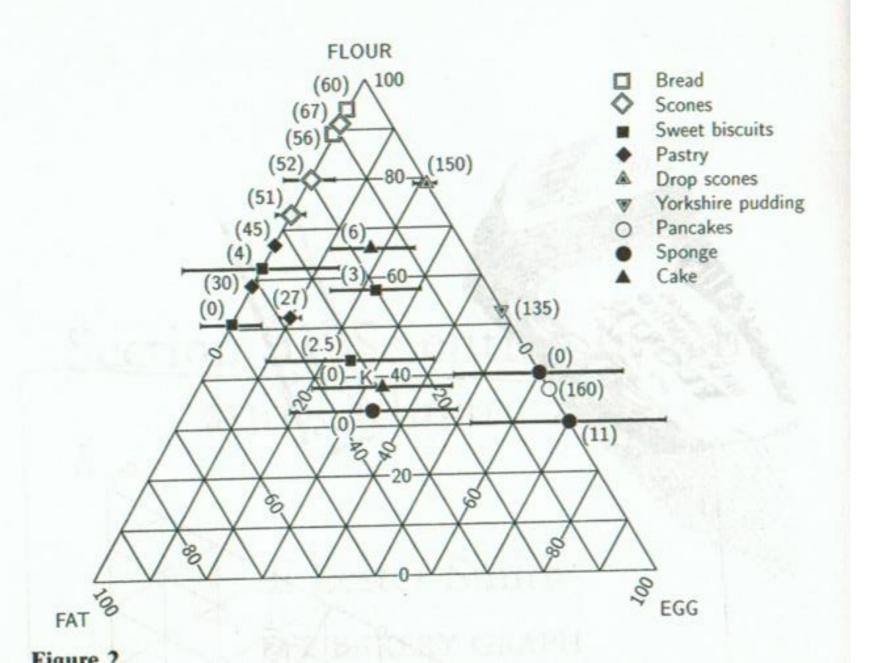












A SURVEY OF FOOD RECOMMENDERS

A PREPRINT

Carl Anderson

Weight Watchers International New York, USA carl.anderson@weightwatchers.com

September 18, 2018

ABSTRACT

Everyone eats. However, people don't always know what to eat. They need a little help and inspiration. Consequently, a number of apps, services, and programs have developed recommenders around food. These cover food, meal, recipe, and restaurant recommendations, which are the most common use cases, but also other areas such as substitute ingredients, menus, and diets. The latter is especially important in the area of health and wellness where users have more specific dietary needs and goals.

In this survey, we review the food recommender literature. We cover the types of systems in terms of their goals and what they are recommending, the datasets and signals that they use to train models, the technical approaches and model types used, as well as some of the system constraints.

Keywords: Personalization · Food recommendation · Recommendation systems · Collaborative filtering · Content-based recommenders · Expert systems

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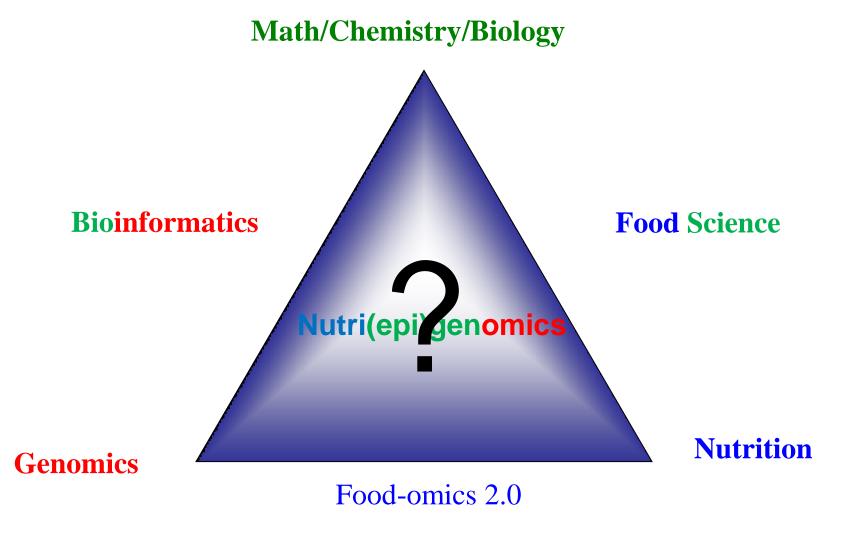
 $\textit{Keywords:}\ \text{Personalization} \cdot \text{Food recommendation} \cdot \text{Recommendation systems} \cdot \text{Collaborative filtering} \cdot \text{Content-based recommenders} \cdot \text{Expert systems}$

Add textmining (patents), phylogenetic data

The Science behind Foodpairing®

Foodpairing is a scientific method to identify which foods & drinks go well together. To understand why ingredients match it's important to know how humans perceive flavour.





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