10a Jornada de Recerca de l'ICS

11a Jornada de Recerca de l'IDIAP



La recerca amb grans bases de dades clíniques millora la salut

7 de juny de 2018

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A holistic approach to common diseases

Rafael de Cid Genomes for life-GCAT lab PMPPC -IGTP

> institut català de la salut ics.gencat.cat

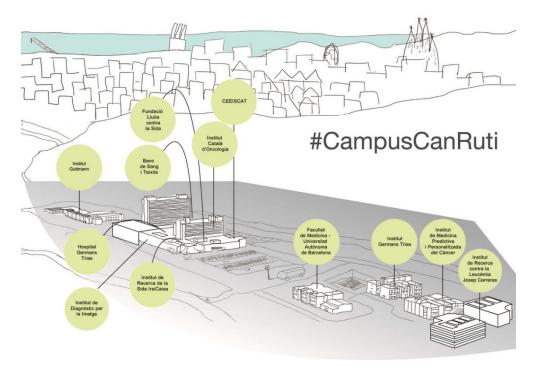






Personalized Medicine













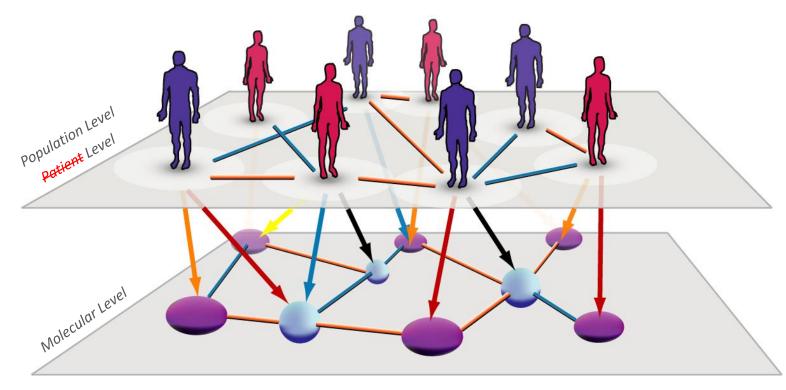






Common diseases





Personalized medicine

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Precision Medicine





Genome medicine

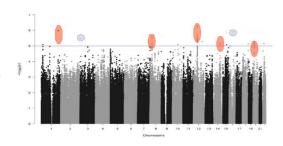


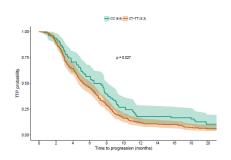
common diseases and all human biomedical traits

"Identifying genetic correlations between complex traits and diseases will are provideing useful etiological insights and help prioritize likely causal relationships"

Genomic profiling in advanced stage non-small-cell lung cancer patients with platinum-based chemotherapy identifies germline variants with prognostic value in *SMYD2*

Nán Galván-Femenía [®]편, Marta Guindo ^{b,}편, Xavier Duran [®]편, Slivia Calabuig-Fariñas ^{e, d, e,}편, Josep Maria Mercader ^{b, 1, 2}편, Jose Luis Ramirez ^f편, Rafael Rosell ^{f,}편, David Torrents ^{b, 0}편, Anna Carreras ^{a,}편, Takashi Kohno ^{h,}편, Eloisa Jantus-Lewintre ^{d, e, 1}편, Carlos Camps ^{e, d, j, k}편, Manuel Perucho ^{f,}편, Lauro Sumoy ^{f,}편, Jun Yokota ^{f,}편, Rafael de Cid ^a ⁸/₂ 편







Phenotyping



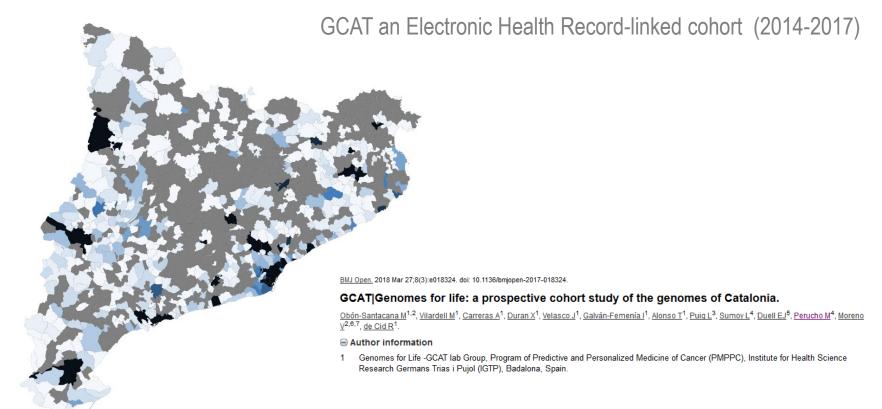
"The limiting factor in progress towards a more complete understanding of the relationship between genome and phenome will be was the availability of high-dimensional phenotype data"

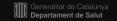
Gratten et Vischer. Genome Medicine, 2016.



GCAT | Genomes for Life. Cohort study



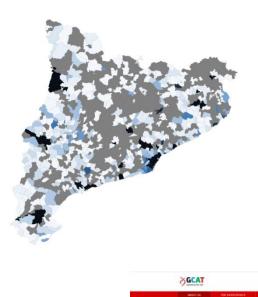






Open Active resource







DAMES VINCAUTED

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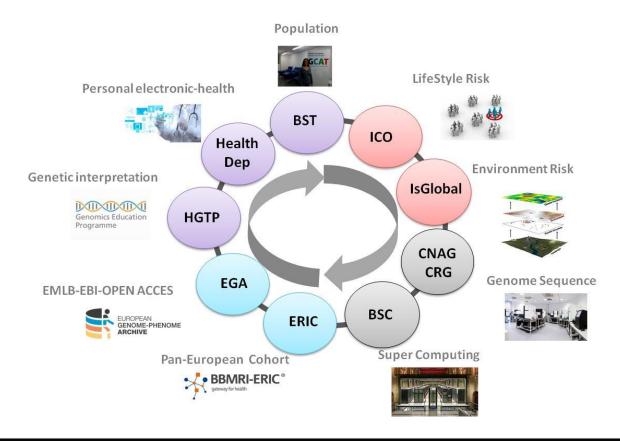
www.genomesforlife.com





Big Data / Complex Data



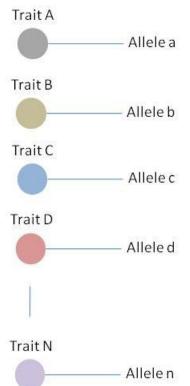


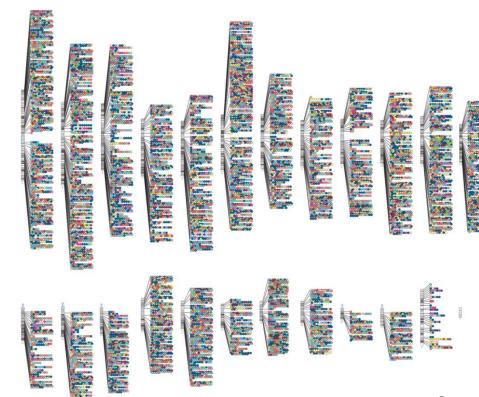


Genomic correlation



Scenario 1





GWAS Hits

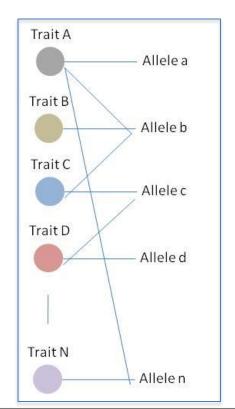


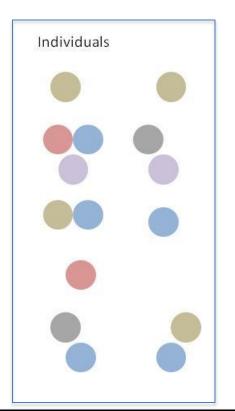


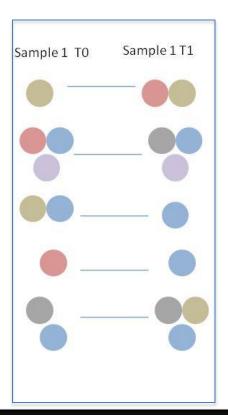
Pleiotropy Multimorbidity Trajectories



Scenario 2 Scenario 3 Scenario 4







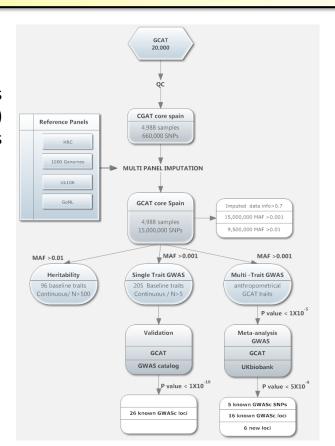


Shared Genomic Variants_population wide disease



GCAT core*

15.000.000 Variants 205 traits (baseline) 5.000 individuals



^{*}Multi-trait-genome association analysis identifies new susceptibility genes for human anthropometric variation in the GCAT cohort.

Galván-Femenía et al, J. Med Gen. 2018 under review

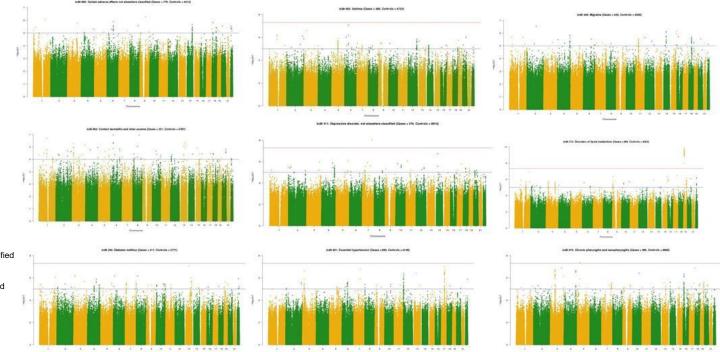




Self-reported_baseline



Identification of Shared Common Variants (ICD-9 coded diseases)



icd9 272: Disorders of lipoid metabolism

icd9 401: Essential hypertension

icd9 995: Certain adverse effects not elsewhere classified

icd9 346: Migraine

icd9 472: Chronic pharyngitis and nasopharyngitis icd9 311: Depressive disorder, not elsewhere classified

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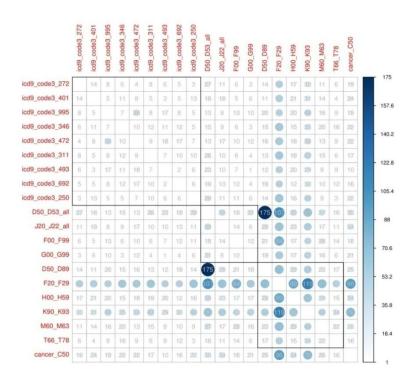
icd9 692: Contact dermatitis and other eczema

icd9 250: Diabetes mellitus



Pair-wise GWAs_disease linkages







icd9 272: Disorders of lipoid metabolism

icd9 401: Essential hypertension

icd9 995: Certain adverse effects not elsewhere classified

icd9 346: Migraine

icd9 472: Chronic pharyngitis and nasopharyngitis

icd9 311: Depressive disorder, not elsewhere classified

icd9 493: Asthma

icd9 692: Contact dermatitis and other eczema

icd9 250: Diabetes mellitus

D50-D53: Nutritional anaemias

F00-F99: Mental and behavioural disorders

G00-G99: Diseases of the nervous system

Guu-G99: Diseases of the nervous system

D50-D89: Diseases of the blood and blood-forming organs and certain disorders involving the immune mechanis

F20-F29: Schizophrenia, schizotypal and delusional disorders

H00-H59: Diseases of the eye and adnexa

K90-K93: Other diseases of the digestive system

M60-M63: Disorders of muscles

T66-T78: Other and unspecified effects of external causes

cancer C50: Breast cancer





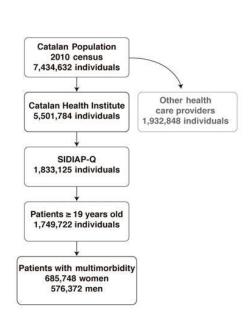


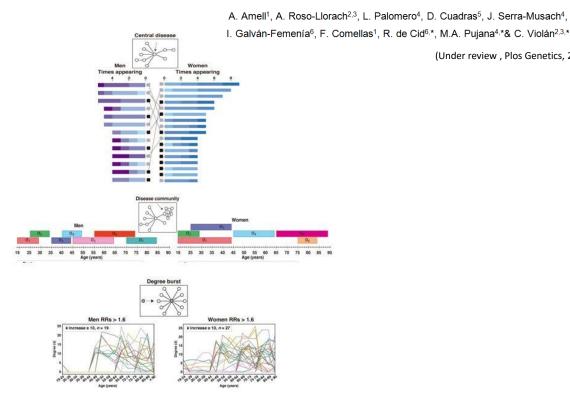
profiling of multimorbidity Key diseases



(Under review, Plos Genetics, 2018)

Topological analysis of disease networks in SIDIAP database

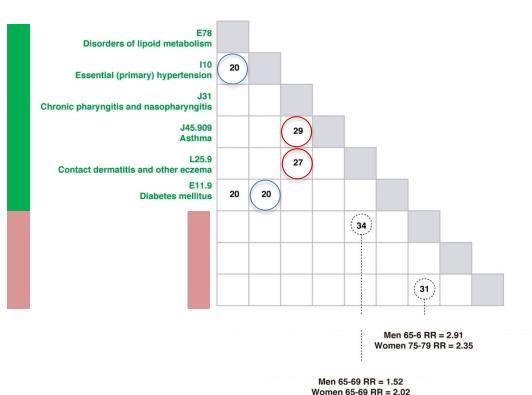






Genetic profiling_key diseases prediction





Common conditions

Known relationships

- E78-I10-E11.9 HTA-DM-TGD
- J45-J31
 Asthma- Chronic Pharyngitis

Key DiseasesUnexpected correlations
Nutrition-Nervous system

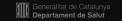




THM



- ■GCAT | Genomes for Life is a big-data project
 - Population based
 - Personal Omic Data
 - Personal Clinical data
 - Sensible data
- ■GCAT is a Resource Recovery and Safe Reuse (RRR)
 - Reuse information
 - Reduce inversions
- ■GCAT will represent a powerful tool for holistic approaches to disease
 - Stratification by disease trajectories and key diseases
 - Knowledge of hidden disease linkage association









Colaborators















Funding















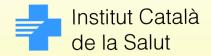




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