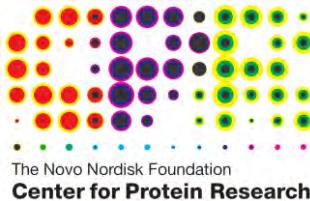


Linking two temporal big data domains: The clinical and the omics worlds



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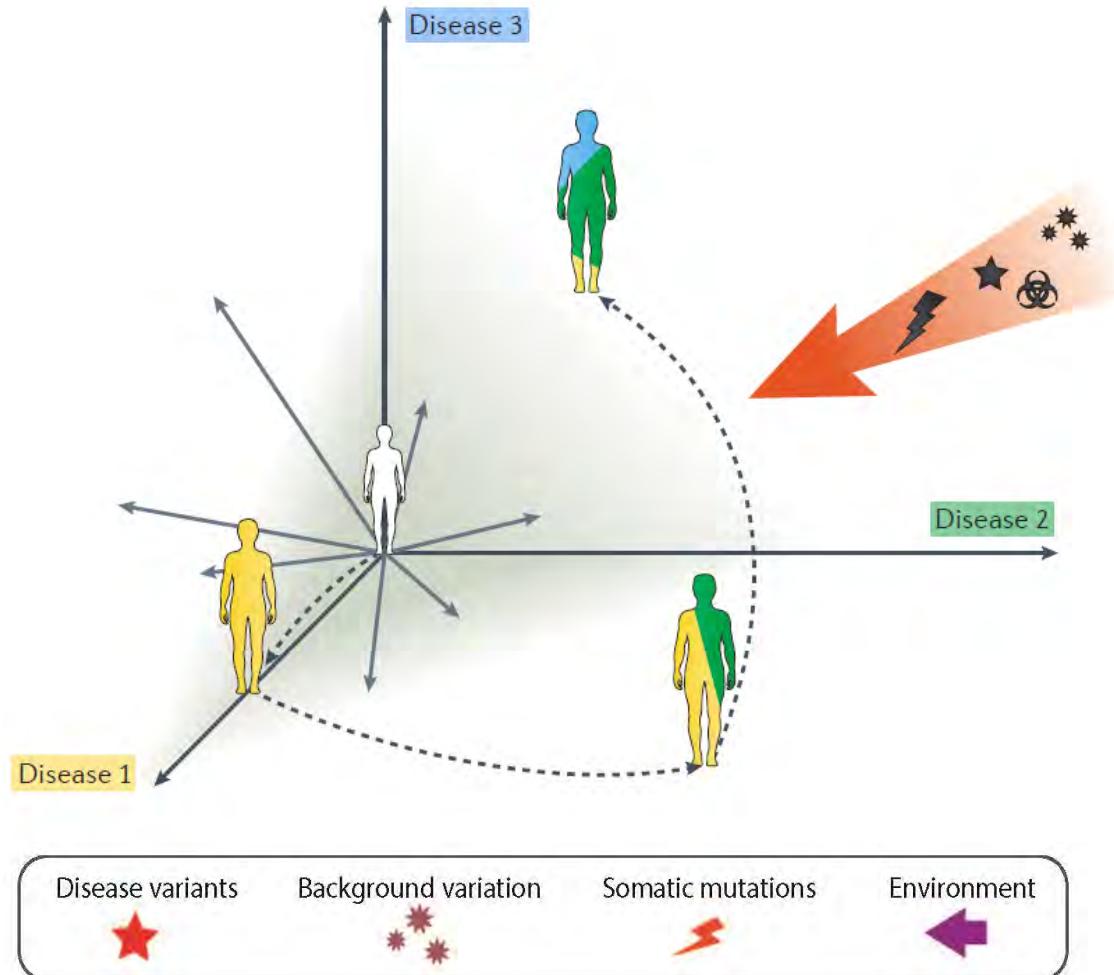
Søren Brunak

Novo Nordisk Foundation Center for Protein Research
University of Copenhagen
soren.brunak@cpr.ku.dk

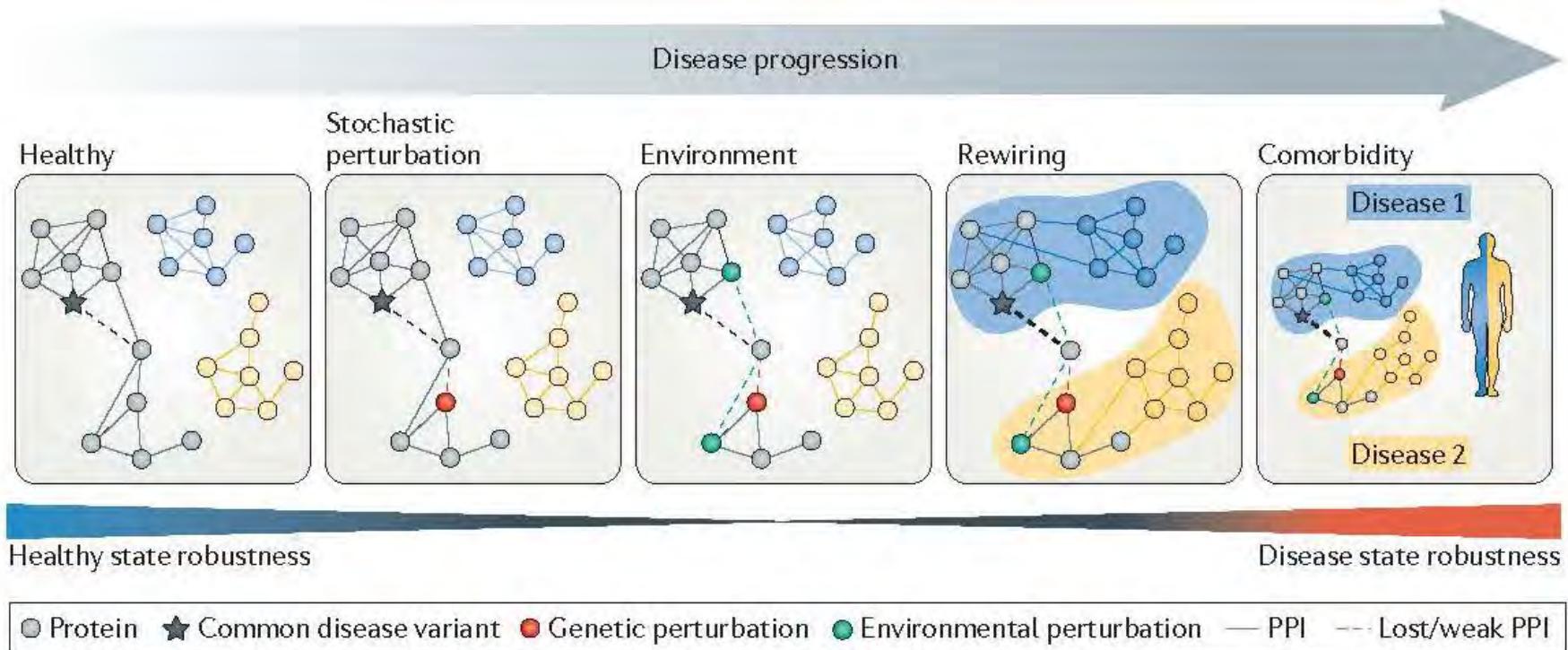
Rigshospitalet
soeren.brunak@regionh.dk

Center for Biological Sequence Analysis
Technical University of Denmark
brunak@cbs.dtu.dk

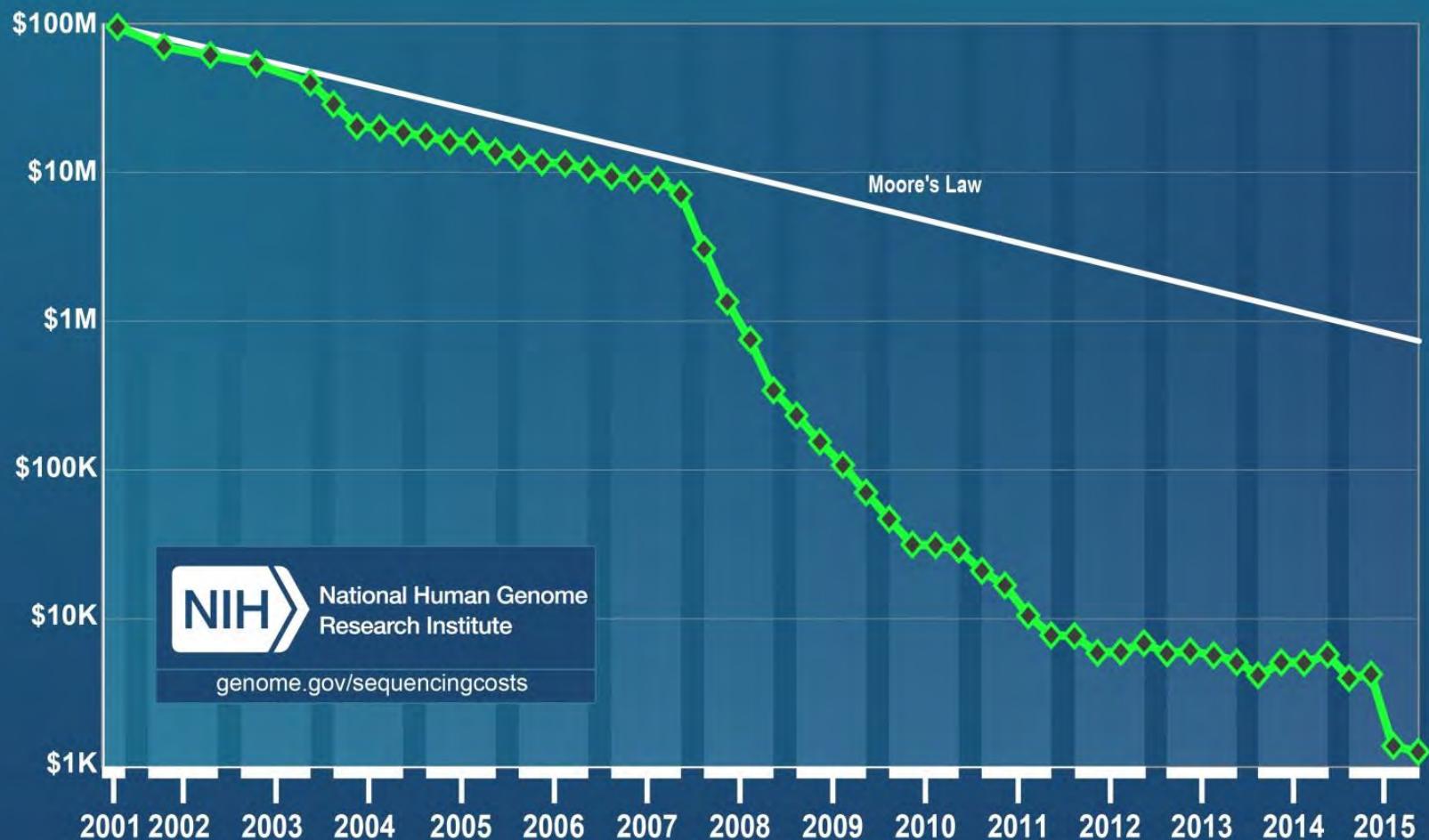
Lifelong multimorbidity journeys in disease space



Rewiring



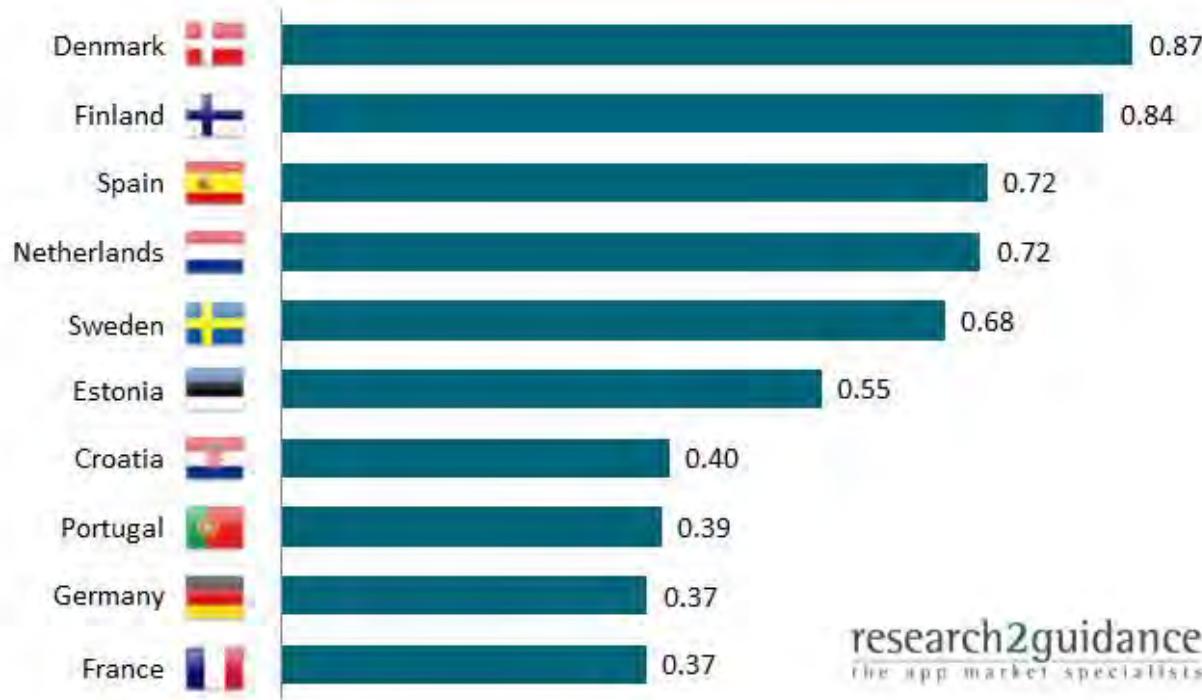
Cost per Genome





DENMARK IS THE LEADING COUNTRY IN EHEALTH ADOPTION

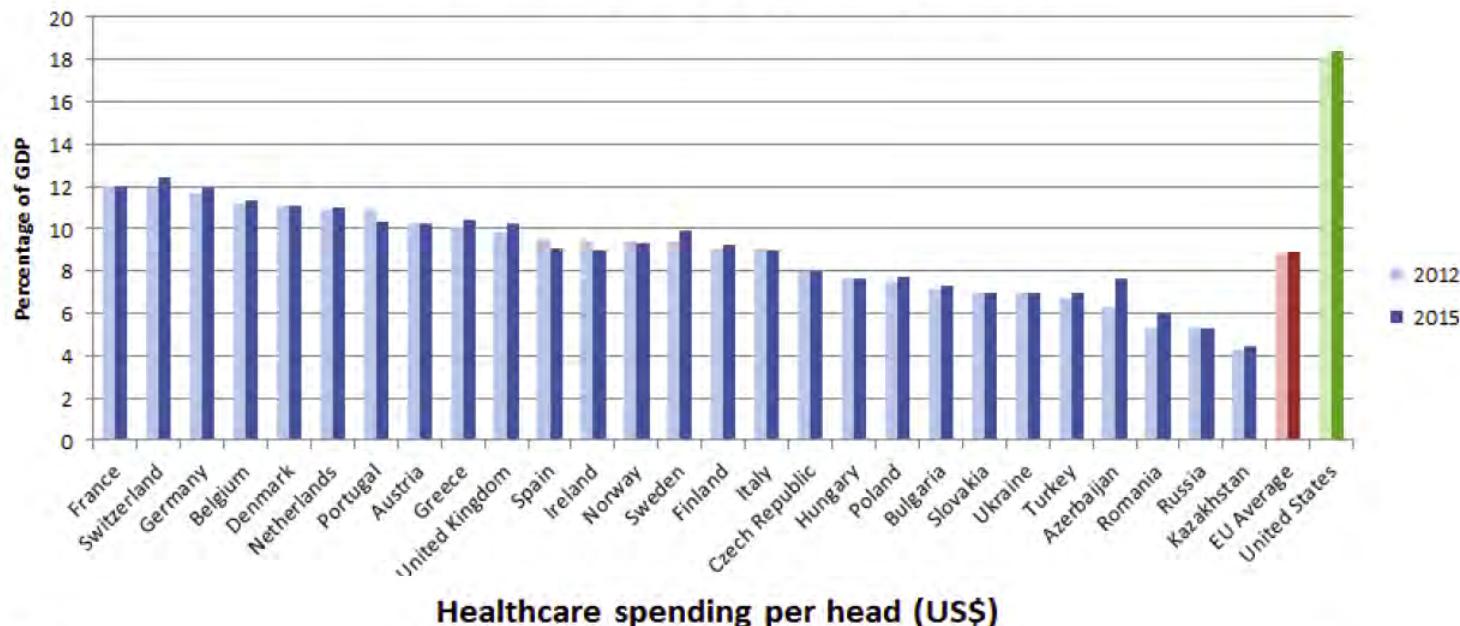
Top 10 EU countries by eHealth adoptions of patients and doctors



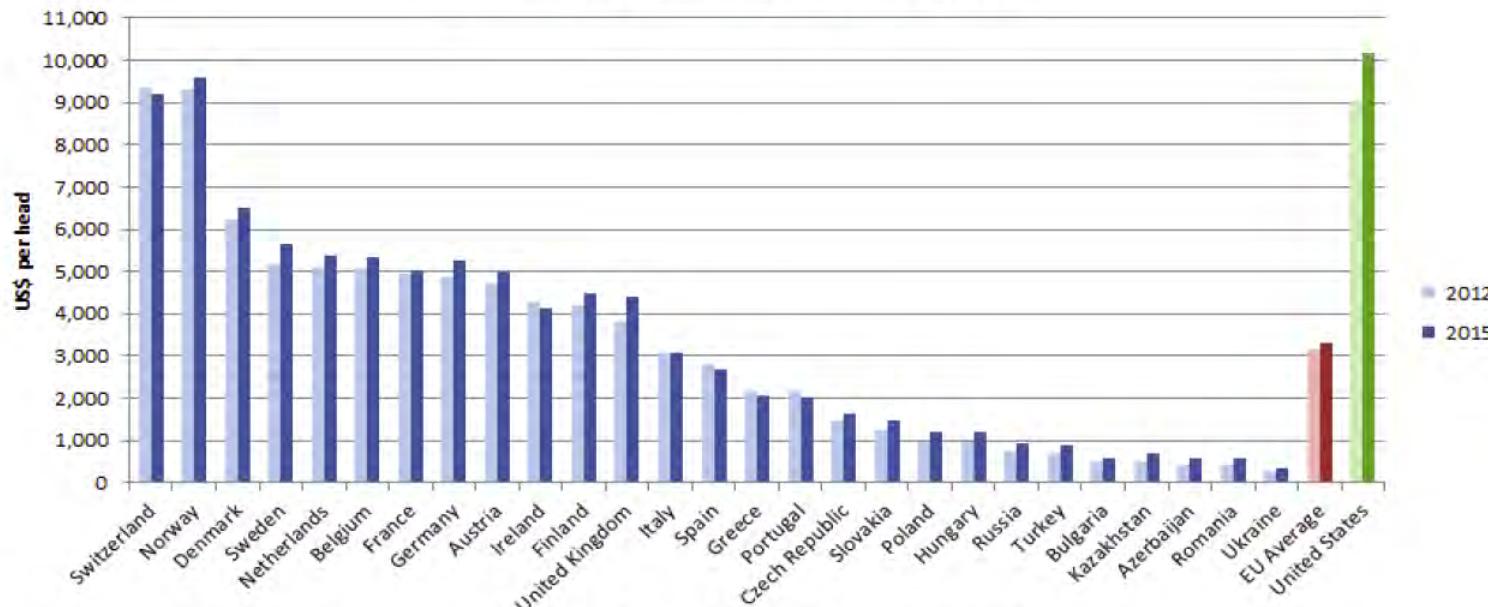
research2guidance
the app market specialists

eHealth adoption – doctors transferring prescription electronically, doctors electronically exchanging medical patient data with other healthcare professionals , patients making appointment via website, patients seeking online information about health

Healthcare spending as percentage of GDP



Healthcare spending per head (US\$)



Source: The Economist Intelligence Unit (2012)

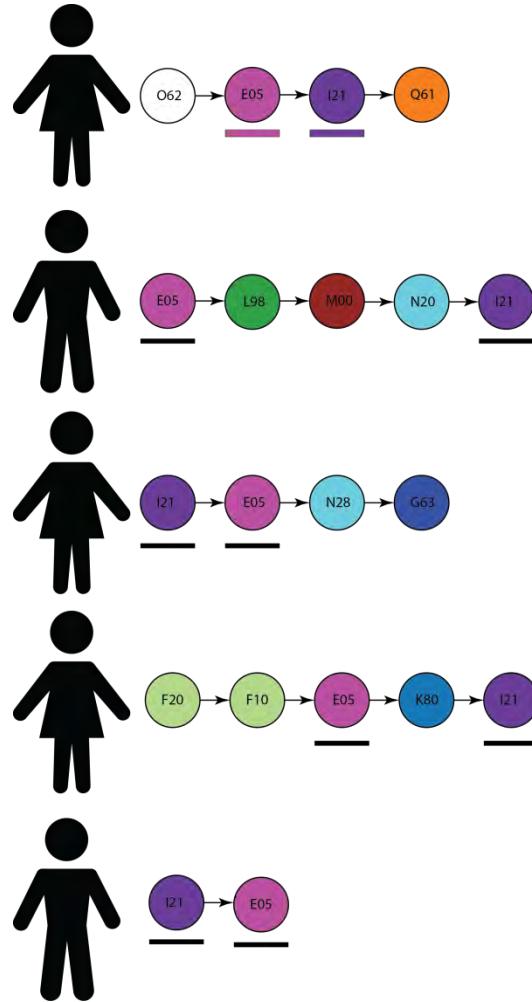
Disease risk

- journey
versus
destination

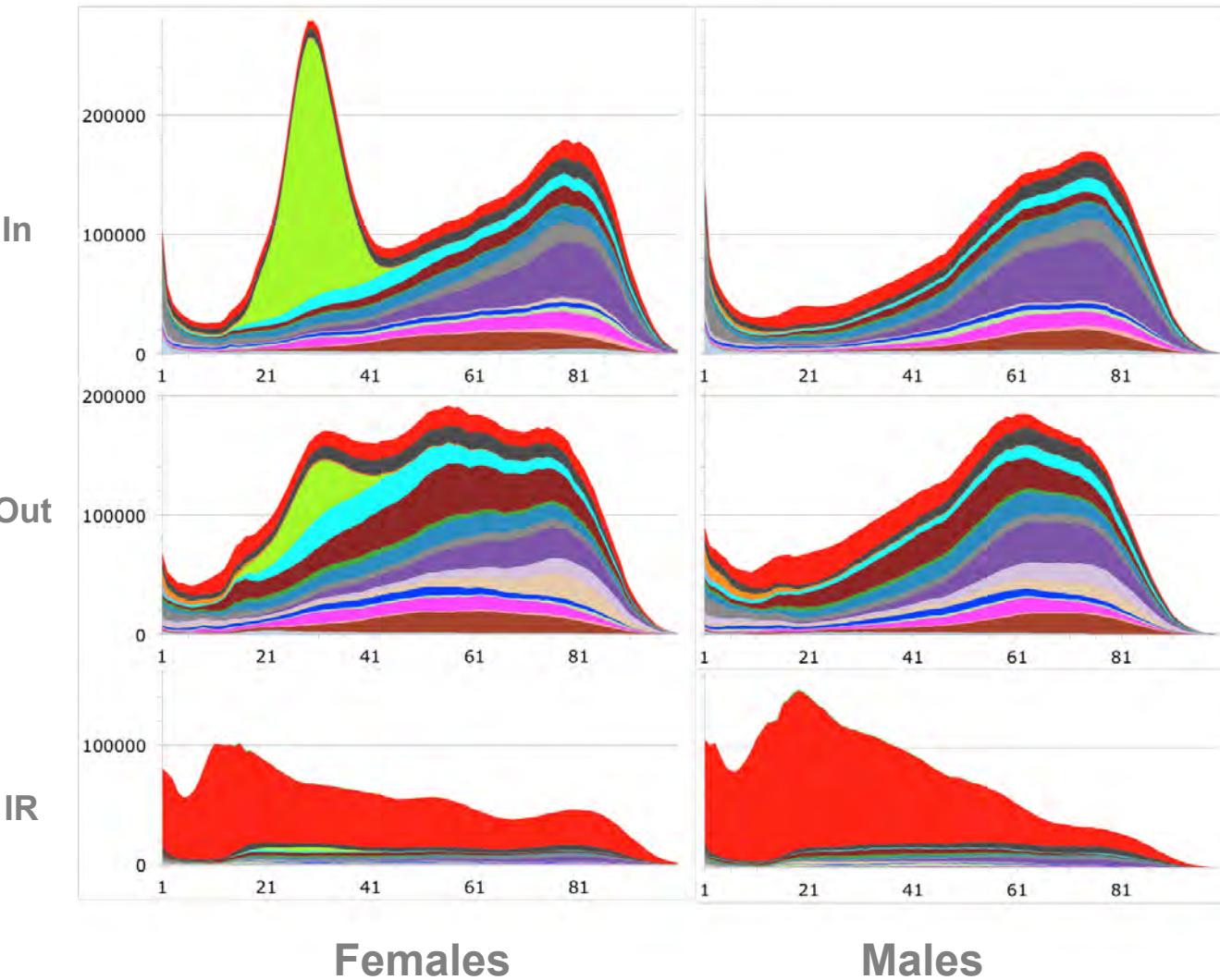
Cancer
Diabetes
Obesity
Mental disorders

Diagnosis trajectories across 6.2 million Danish individuals

(ICD-10 era, 1994-2015)



National Patient Registry (6.2M Danes) ICD10 diagnoses as a function of age



ICD 10 chapter coloring

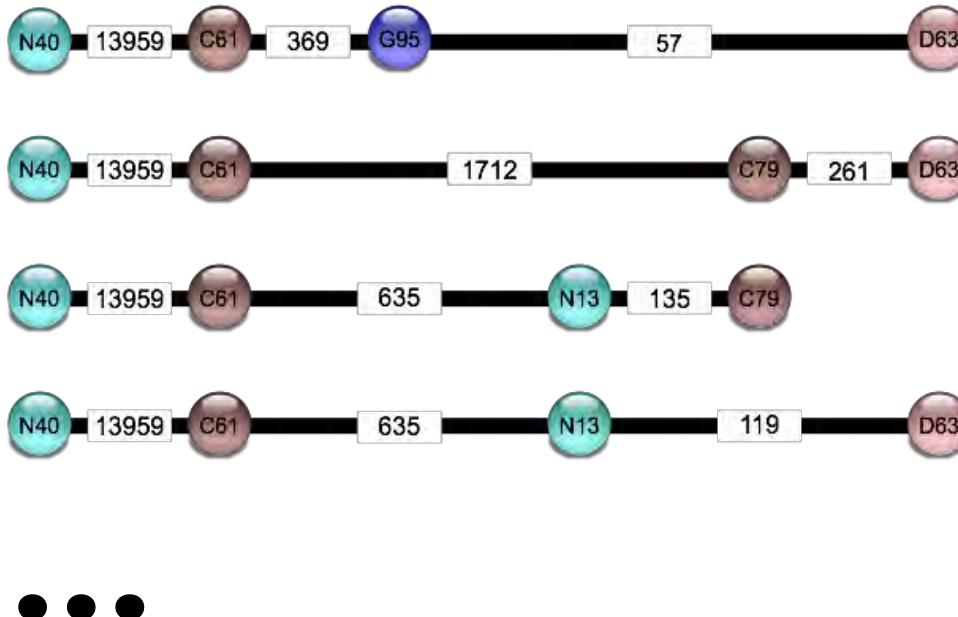
- 1: Certain infectious and parasitic diseases
- 2: Neoplasms
- 3: Diseases of the blood and blood-forming organs and certain disorders involving the immune mechanism
- 4: Endocrine, nutritional and metabolic diseases
- 5: Mental and behavioural disorders
- 6: Diseases of the nervous system
- 7: Diseases of the eye and adnexa
- 8: Diseases of the ear and mastoid process
- 9: Diseases of the circulatory system
- 10: Diseases of the respiratory system
- 11: Diseases of the digestive system
- 12: Diseases of the skin and subcutaneous tissue
- 13: Diseases of the musculoskeletal system and connective tissue
- 14: Diseases of the genitourinary system
- 15: Pregnancy, childbirth and the puerperium
- 16: Certain conditions originating in the perinatal period
- 17: Congenital malformations, deformations and chromosomal abnormalities
- 18: Symptoms, signs and abnormal clinical and laboratory findings, not elsewhere classified
- 19: Injury, poisoning and certain other consequences of external causes
- 20: External causes of morbidity and mortality

6-7 million trajectories

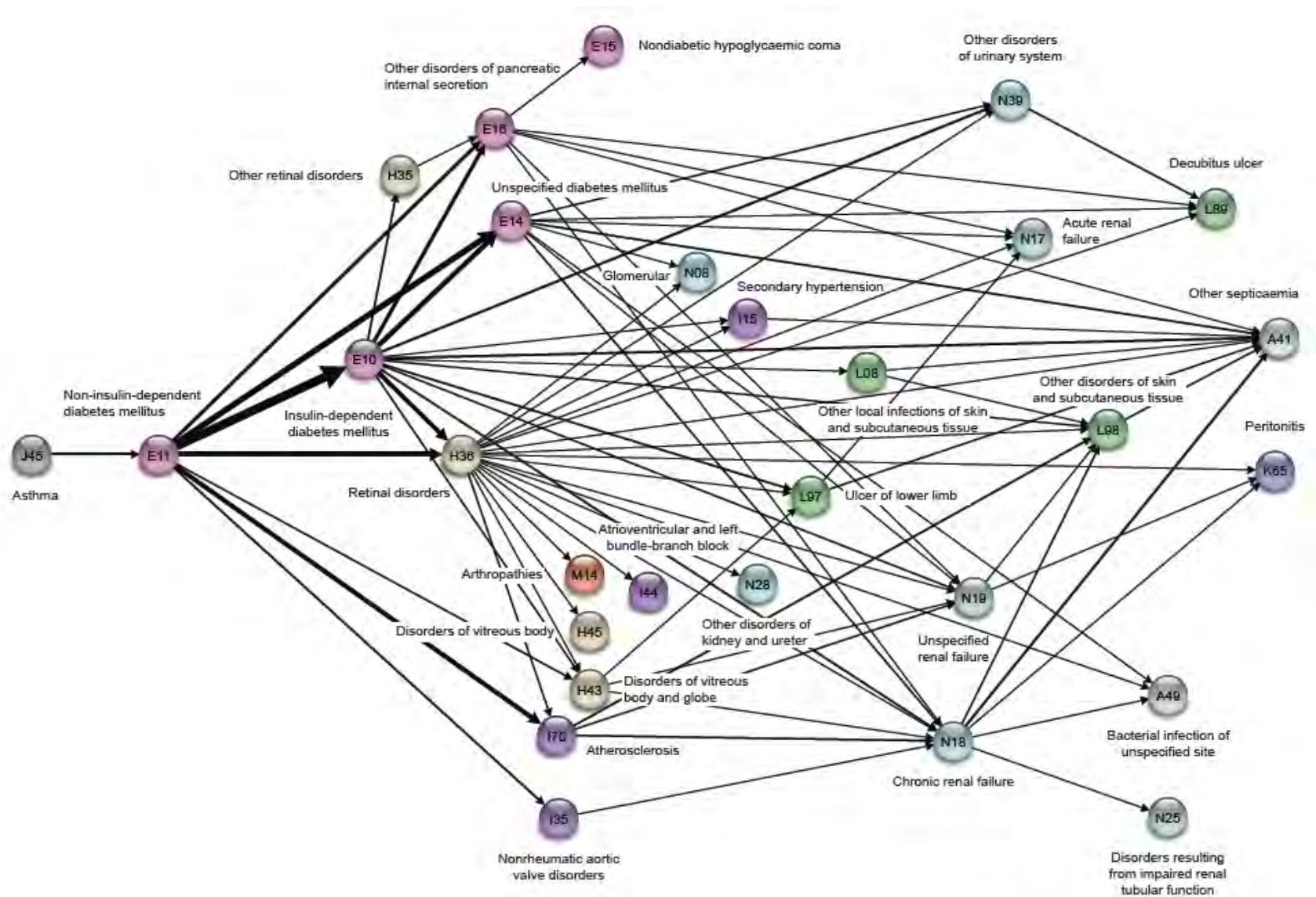


?

6.2 million individual trajectories condensed into 1,171 “recurrent” ones

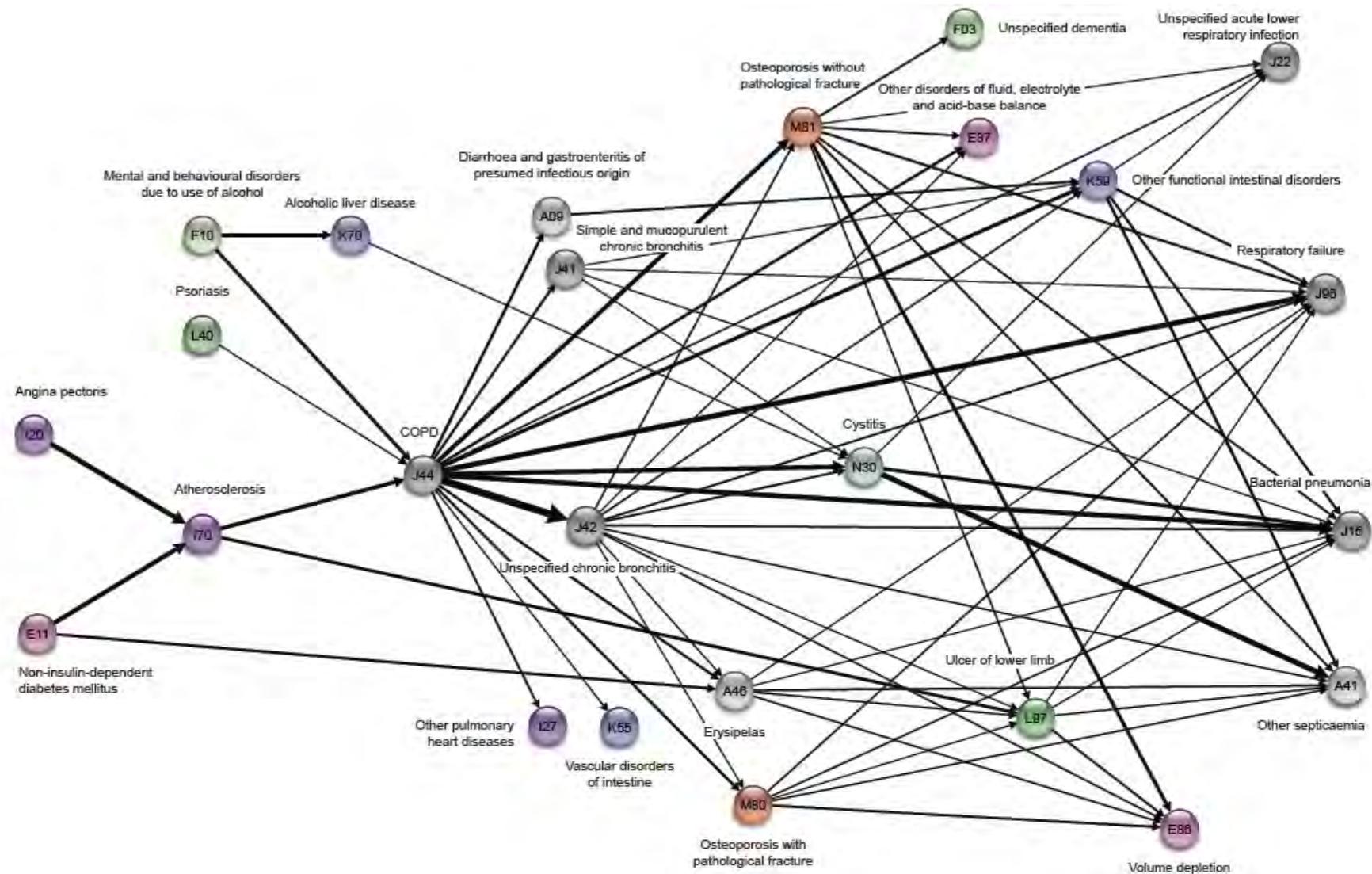


Diabetes trajectory network



COPD trajectory cluster

with five preceding diagnoses leading
to COPD and some of the possible outcomes



DISEASE TRAJECTORY SEARCH:

ALL DIAGNOSES (UNION)

SEARCH:

FILTERS

EDGE ANNOTATION:

PATIENTS RELATIVE RISK OFF

NODE ANNOTATION:

ICD CODE TEXT DESC. NONE

 INSTANT SEARCH PERFORMANCE ISSUES?

SEARCH

Information

Data from: Danish National Patient Register (Landspatientregisteret)

Population: ~6,900,000 people

Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci

David Ellinghaus^{1,49}, Luke Jostins², Sarah L Spain², Adrian Cortes^{3,4}, Jörn Bethune¹, Buhm Han⁵, Yu Rang Park⁶, Soumya Raychaudhuri^{7–10}, Jennie G Pouget^{11,12}, Matthias Hüenthal¹, Trine Folseraa^{13–16}, Yunpeng Wang¹⁷, Tonu Esko^{18–20}, Andres Metspalu¹⁸, Harm-Jan Westra^{7–10}, Lude Franke²¹, Tune H Pers^{7,20,22,23}, Rinse K Weersma²⁴, Valerie Collij²⁴, Mauro D'Amato^{25,26}, Jonas Halfvarson²⁷, Anders Boeck Jensen²⁸, Wolfgang Lieb^{29,30}, Franziska Degenhardt^{31,32}, Andreas J Forstner^{31,32}, Andrea Hofmann^{31,32}, The International IBD Genetics Consortium (IIBDGC)³³, International Genetics of Ankylosing Spondylitis Consortium (IGAS)³³, International PSC Study Group (IPSCSG)³³, Genetic Analysis of Psoriasis Consortium (GAPC)³³, Psoriasis Association Genetics Extension (PAGE)³³, Stefan Schreiber^{1,34}, Ulrich Mrowietz³⁵, Brian D Juran³⁶, Konstantinos N Lazaridis³⁶, Søren Brunak²⁸, Anders M Dale^{17,37}, Richard C Trembath³⁸, Stephan Weidinger³⁵, Michael Weichenthal³⁵, Eva Ellinghaus¹, James T Elder^{39,40}, Jonathan N W N Barker⁴¹, Ole A Andreassen^{42,43}, Dermot P McGovern^{44,45}, Tom H Karlsen^{13–16}, Jeffrey C Barrett², Miles Parkes⁴⁶, Matthew A Brown^{47,48,50} & Andre Franke^{1,50}

We simultaneously investigated the genetic landscape of ankylosing spondylitis, Crohn's disease, psoriasis, primary sclerosing cholangitis and ulcerative colitis to investigate pleiotropy and the relationship between these clinically related diseases. Using high-density genotype data from more than 86,000 individuals of European ancestry, we identified 244 independent multidisease signals, including 27 new genome-wide significant susceptibility loci and 3 unreported shared risk loci. Complex pleiotropy was supported when contrasting multidisease signals with expression data sets from human, rat and mouse together with epigenetic and expressed enhancer profiles. The comorbidities among the five immune diseases were best explained by biological pleiotropy rather than heterogeneity (a subgroup of cases genetically identical to those with another disease, possibly owing to diagnostic misclassification, molecular subtypes or excessive comorbidity). In particular, the strong comorbidity between primary sclerosing cholangitis and inflammatory bowel disease is likely the result of a unique disease, which is genetically distinct from classical inflammatory bowel disease phenotypes.

Genome-wide association studies (GWAS) have shown overlap in the genetic susceptibility to human diseases that affect a range of

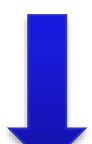
In this study, we combined Immunochip genotype data for 52,262 cases and 34,213 controls of European ancestry, currently the largest available

**Immunochip genotype data:
52,262 cases**

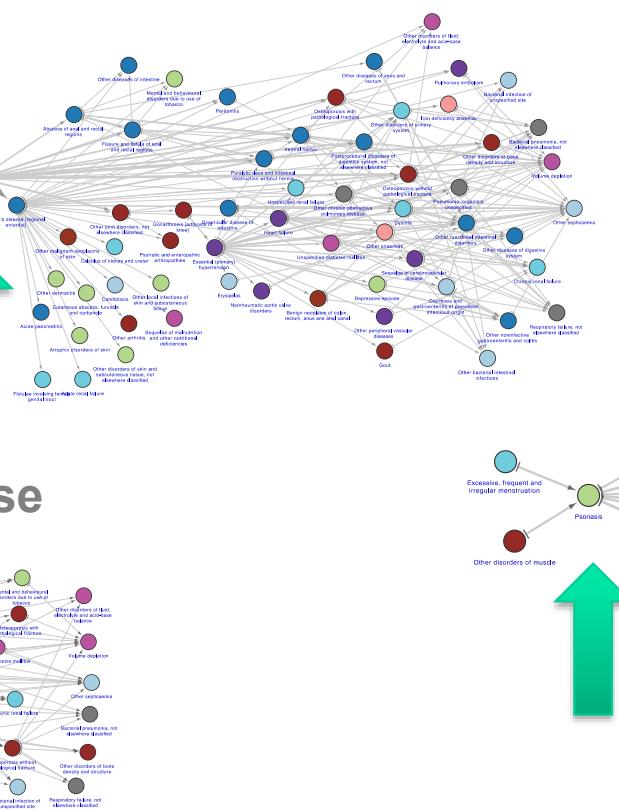
Ankylosing spondylitis (8,726)
Crohn's disease (19,085)
Psoriasis (6,530)
Primary sclerosing cholangitis (3,408)
Ulcerative colitis (14,413)

34,213 healthy controls

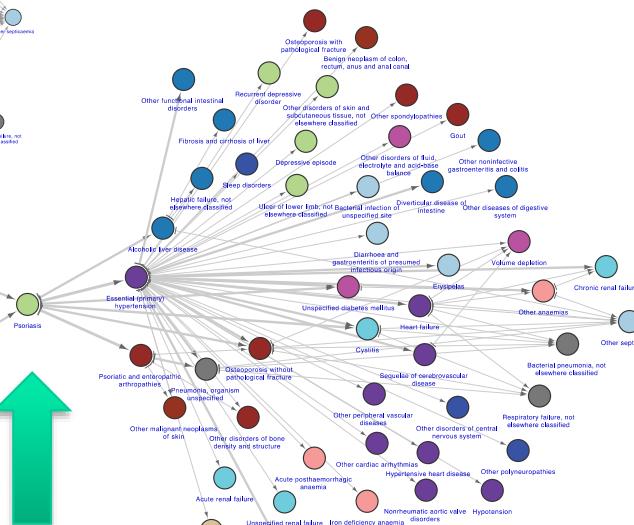
Ulcerative colitis



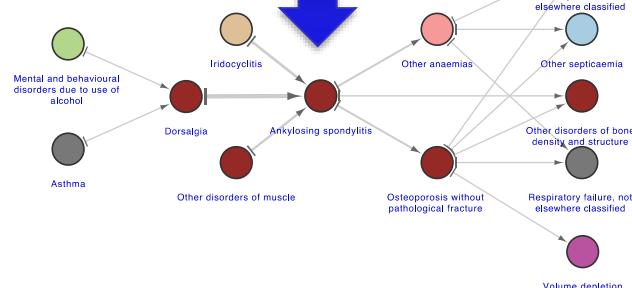
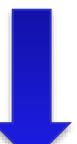
Crohn's disease



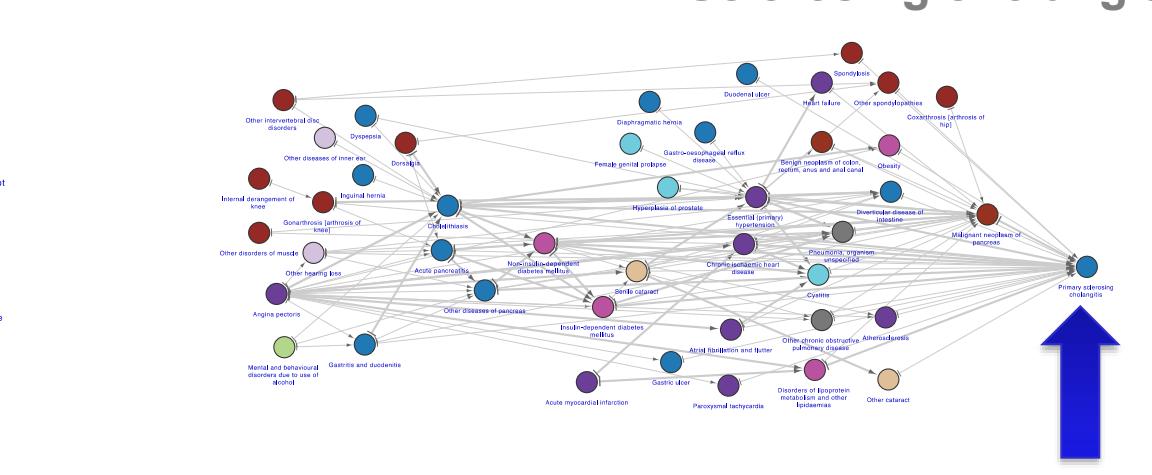
Psoriasis



Ankylosing spondylitis

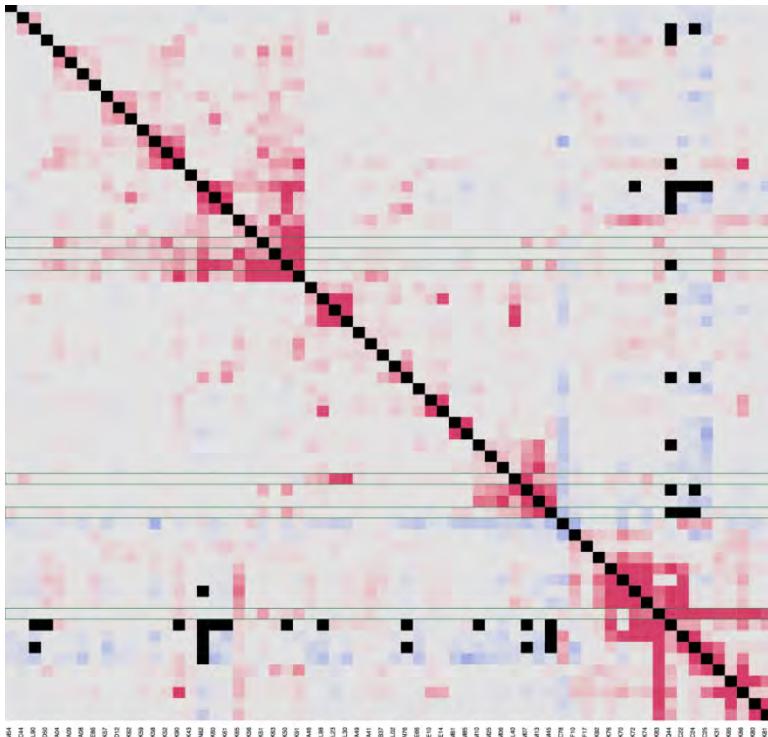


Sclerosing cholangitis

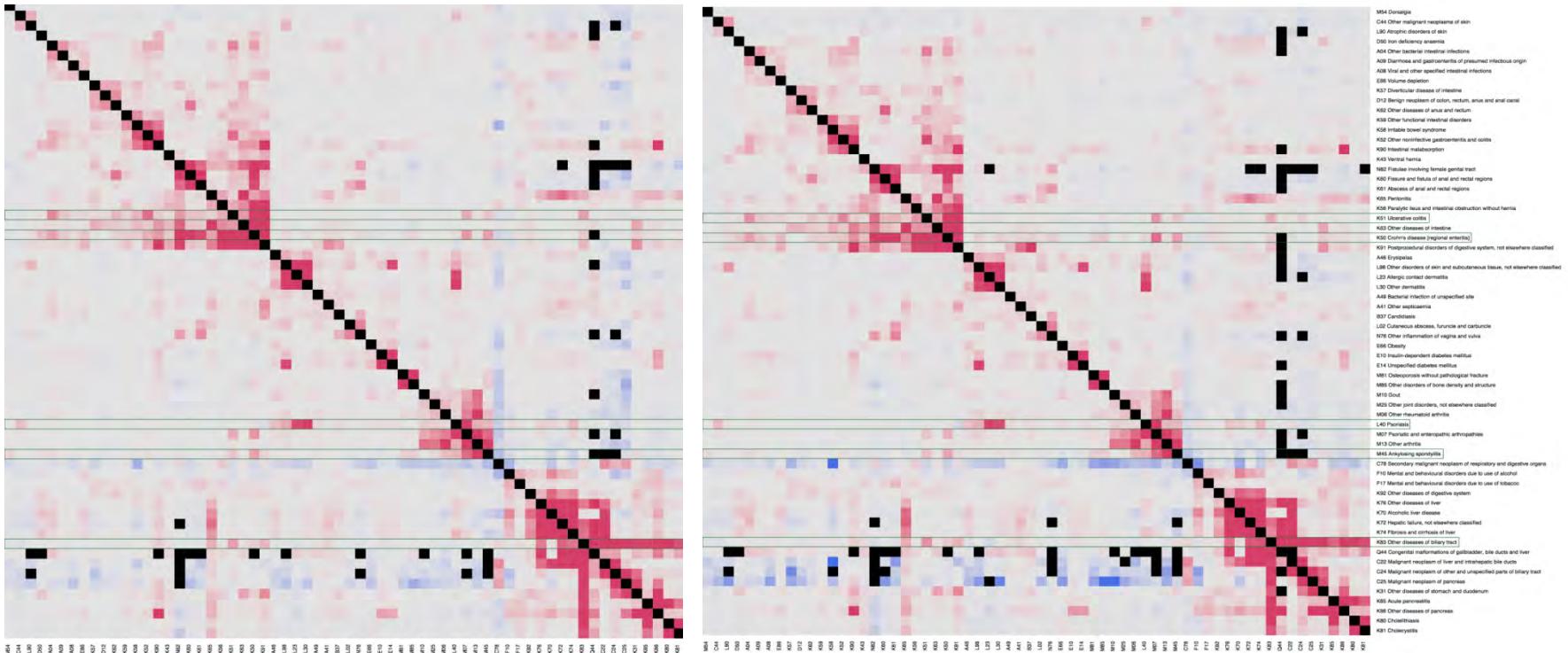


Temporal disease associations

A and B



A before B



Shared or distinct genetic etiology?

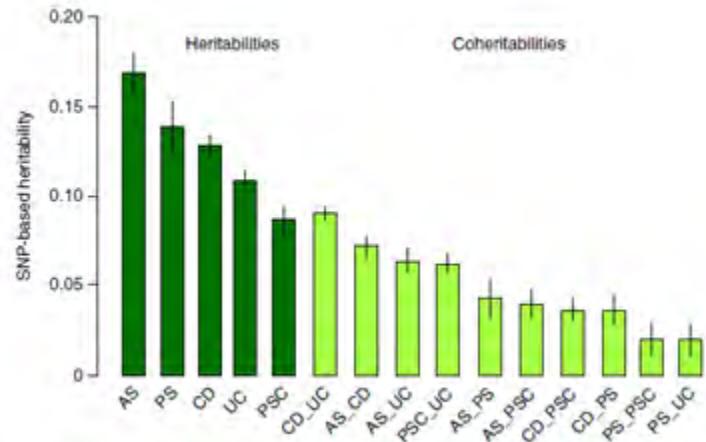
Pleiotropy (sharing of risk alleles by disease A and disease B) or **Heterogeneity** (a subgroup of disease A cases has a higher load of risk alleles for disease B)?

Genetic and healthdata overlap analyses shows

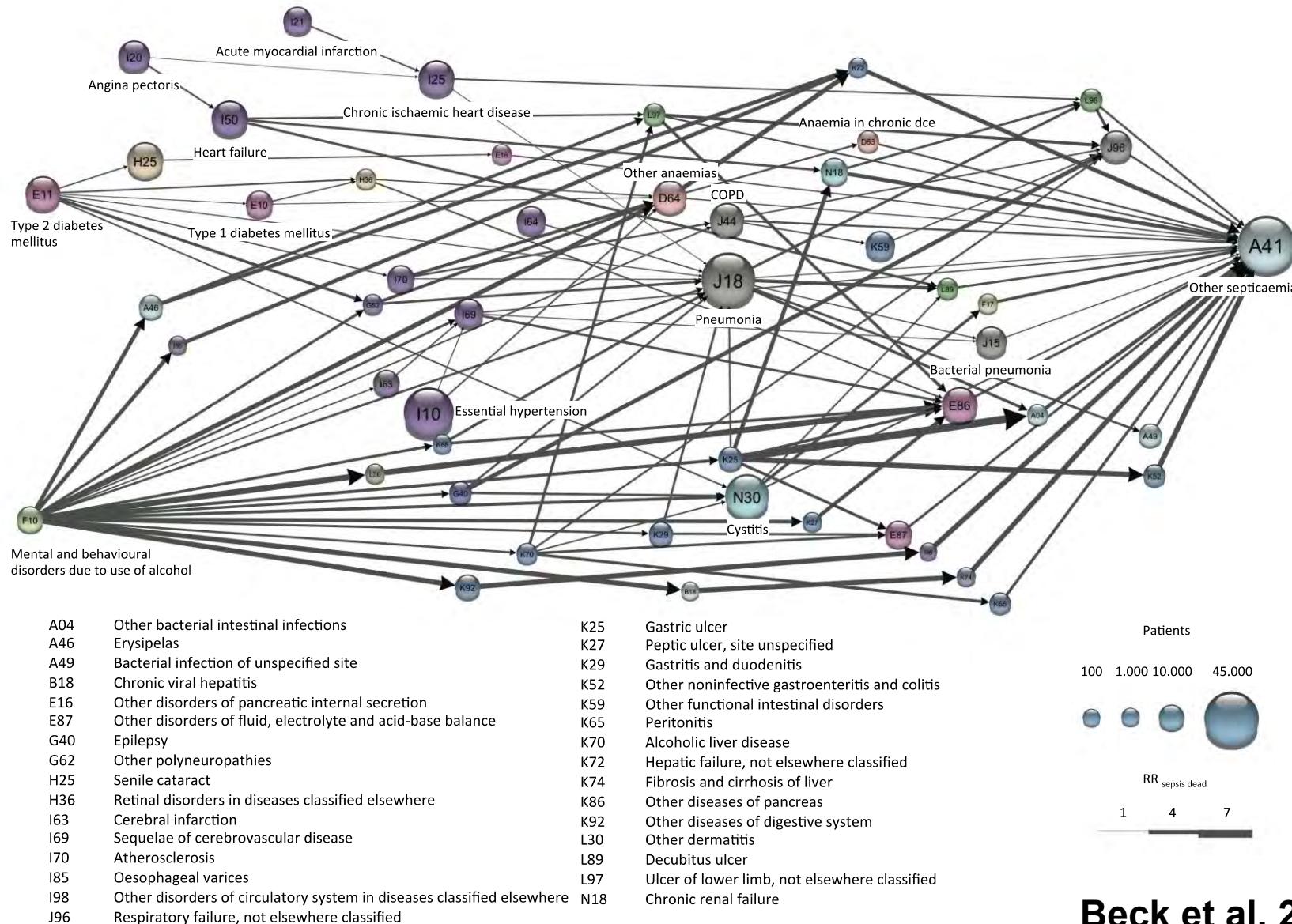
- **that shared pathophysiological pathways are the basis for clinical co-occurrence**
- that patients with concomitant syndromes are genetically distinct from patients without concomitant syndromes.

Immunochip-wide pleiotropy estimates

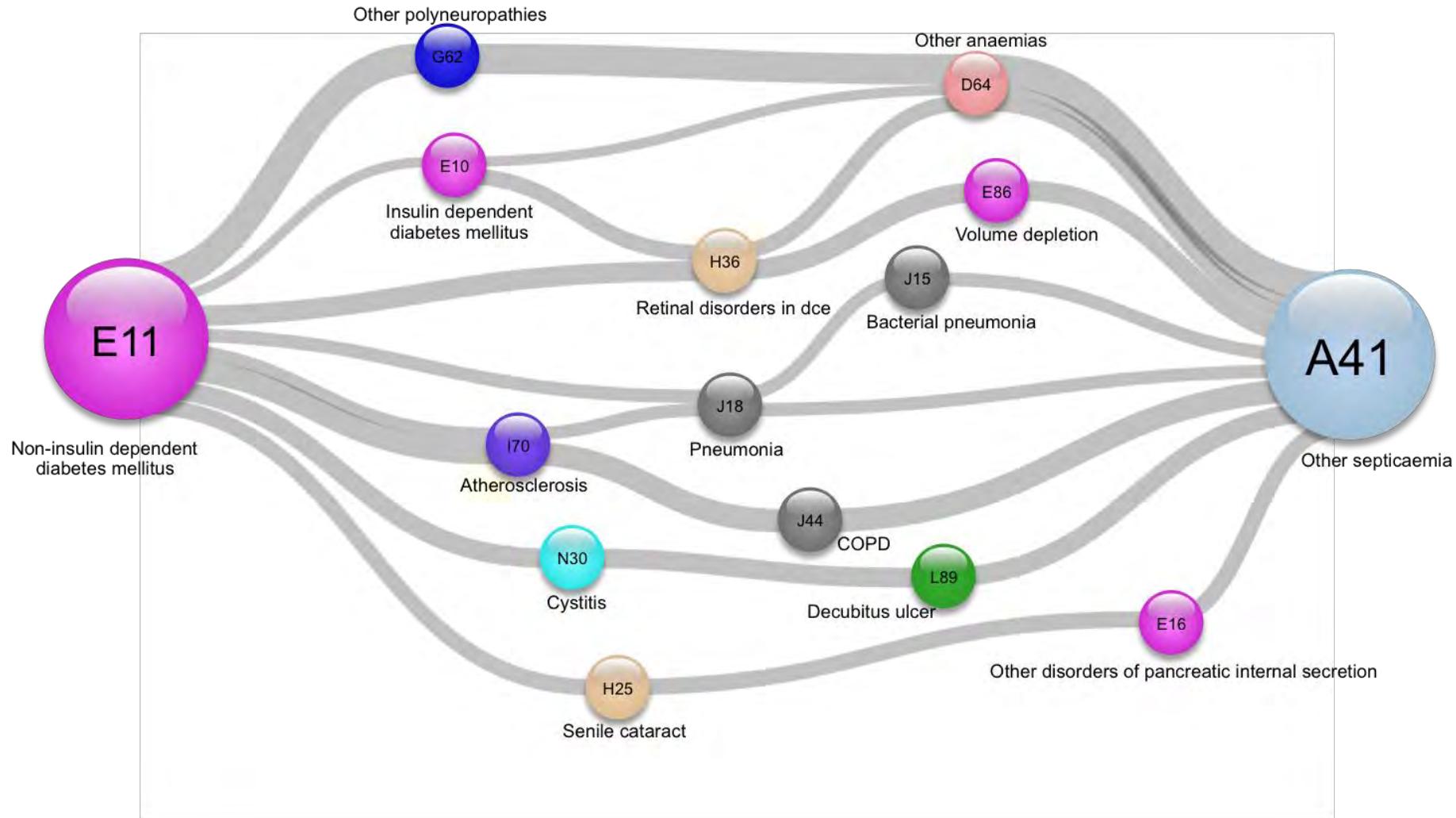
Ellinghaus et al. Nature Genetics, 48:510-518, 2016



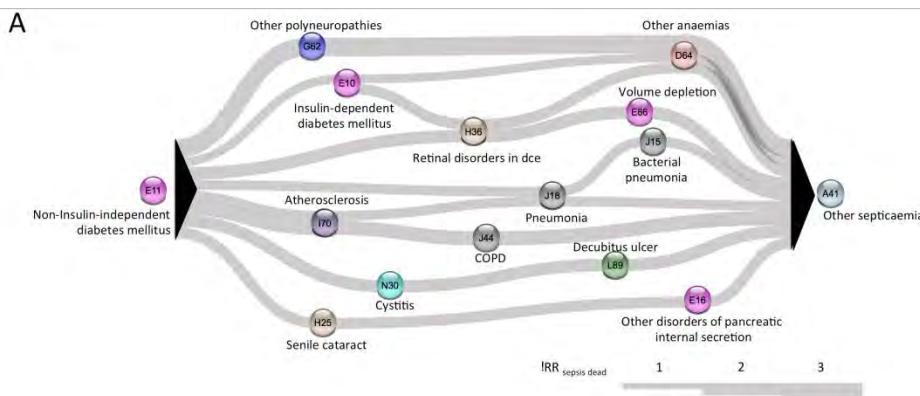
Sepsis survival across pre-history (120,000 patients, 56 significant trajectories)



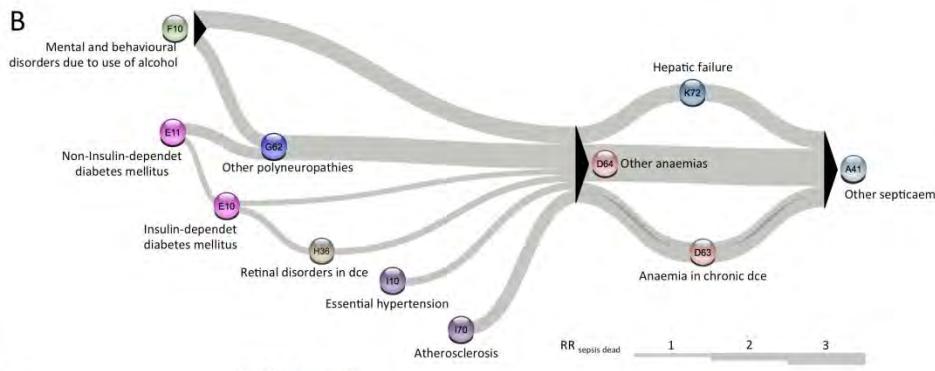
From diabetes to sepsis



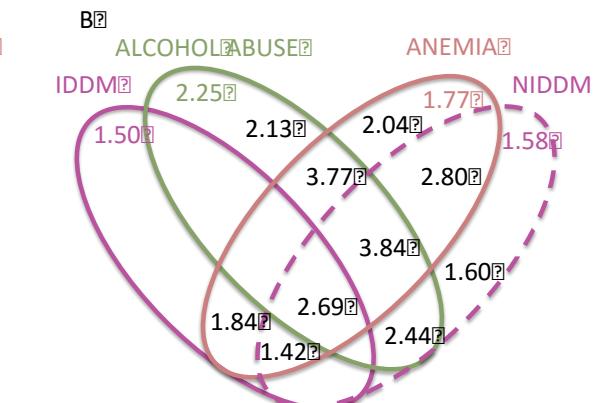
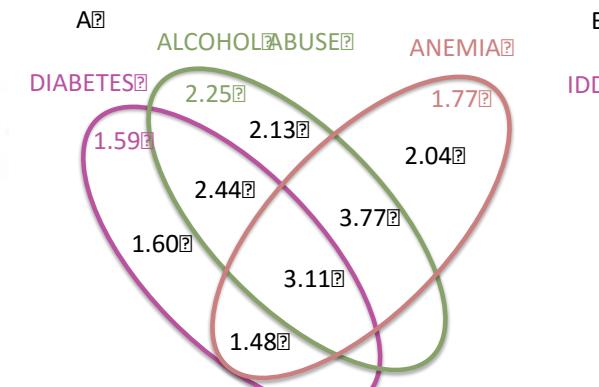
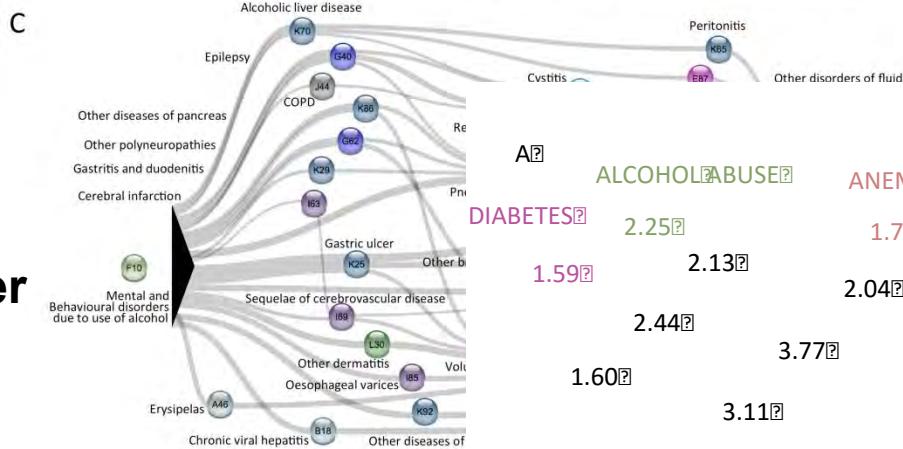
Diabetes



Anaemia



Mental disorder





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CLINICAL IMPLICATIONS OF BASIC RESEARCH

A Wake-up Call for Type 2 Diabetes?

Shanta J. Persaud, Ph.D., and Peter M. Jones, Ph.D.

N Engl J Med 2016; 375:1090-1092 | September 15, 2016 | DOI: 10.1056/NEJMcibr1607950

Share:

This study examines an established association between a variant in a melatonin-receptor gene and type 2 diabetes, yielding insights into how the variant confers susceptibility to the disease.

Disclosure forms provided by the authors are available at NEJM.org.

SOURCE INFORMATION

From the Diabetes Research Group, Division of Diabetes and Nutritional Sciences, King's College London, London.

MEDIA IN THIS ARTICLE

FIGURE 1



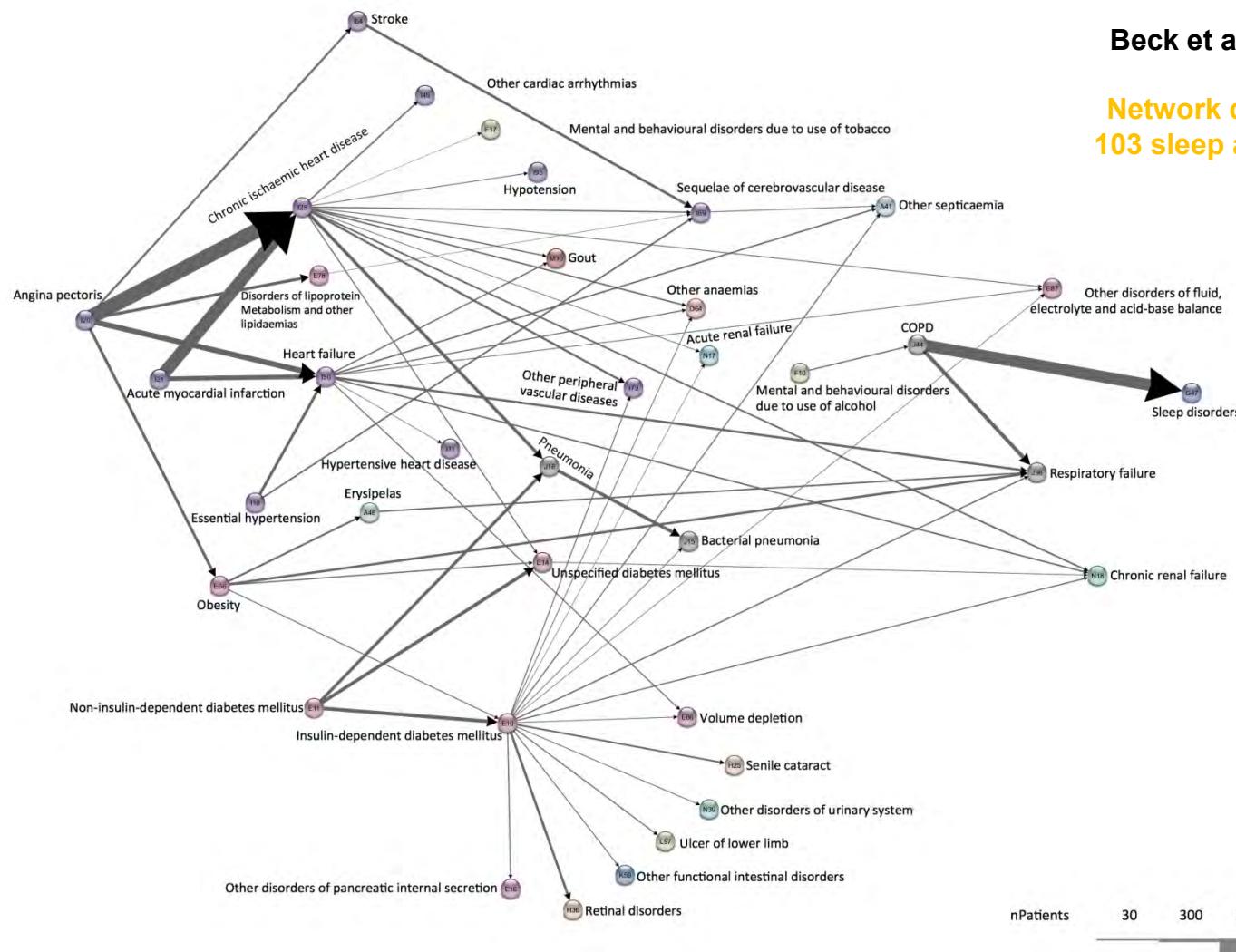
Effect of Variant *MTNR1B* on Melatonin Signaling in Islet Beta Cells.

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Sleep Apnea and Diabetes

Beck et al., to appear 2017

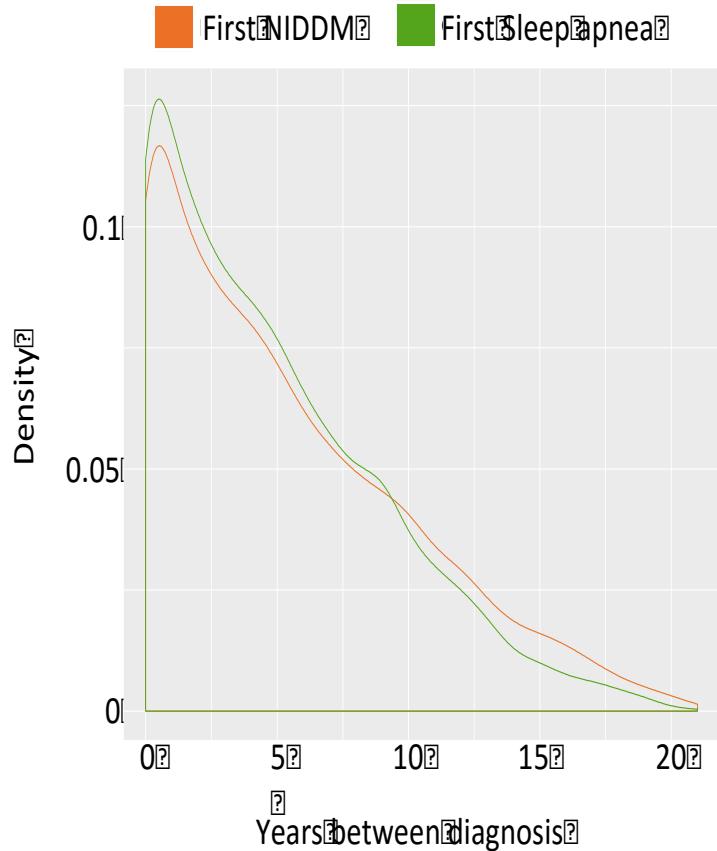
Network constructed from
103 sleep apnea trajectories



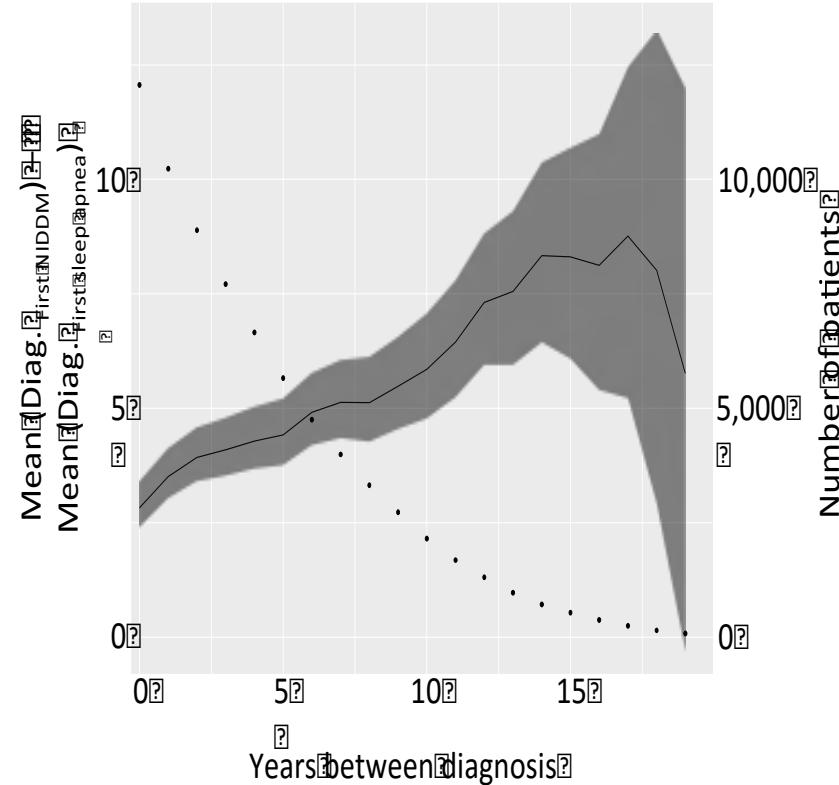
No significant diagnosis direction, but patients diagnosed with diabetes before sleep apnea have significantly more comorbidities than patients diagnosed with sleep apnea before diabetes.

Excess number of comorbidities if diabetes is first

A



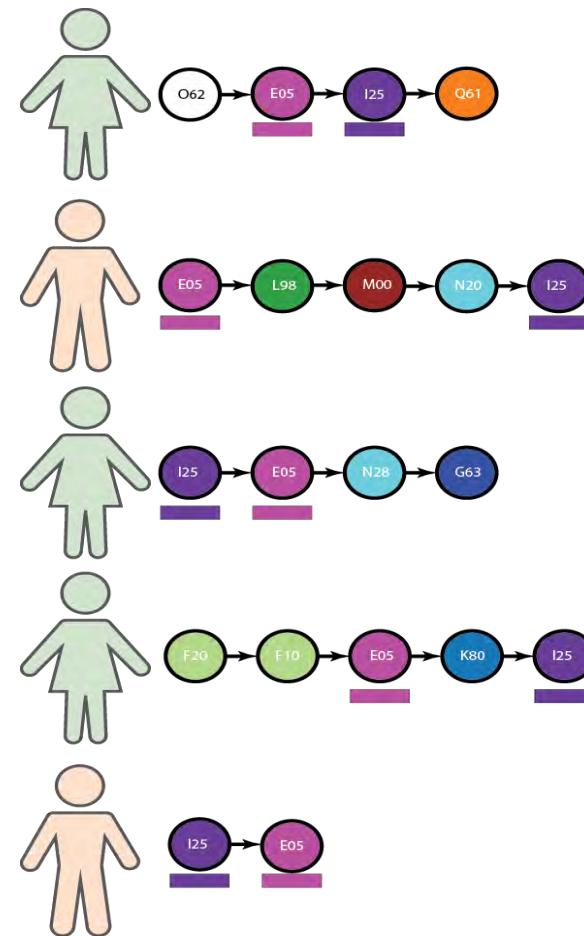
B



(A) Distribution of years between NIDDM and sleep apnea for patients diagnosed with NIDDM first (orange) and for patients diagnosed with sleep apnea first (green). (B) The plot shows the excess number of comorbidities for patients diagnosed with NIDDM first compared to those diagnosed with sleep apnea first (black line) with the 95% confidence interval (grey area). The dots indicate the number of patients having the minimum years between the two diagnoses.

Gender medicine: Gender specific disease trajectories

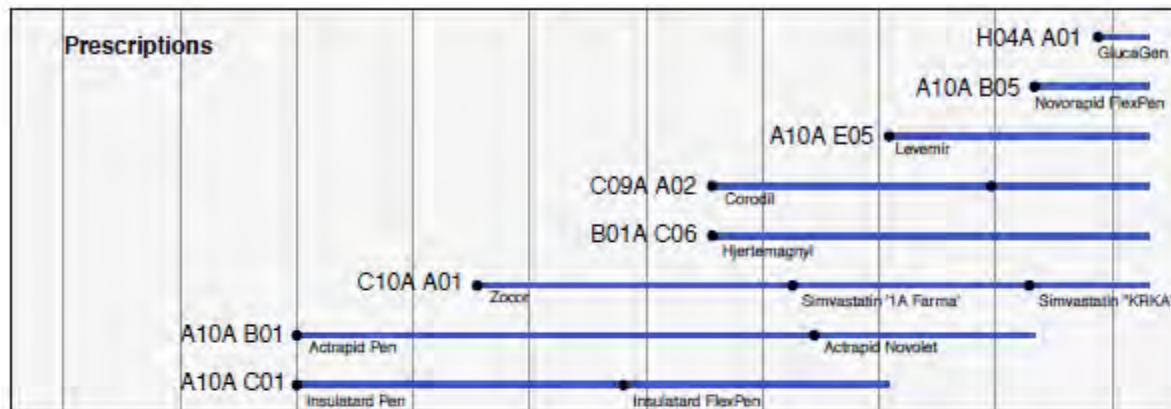
- Males and females have some differences in physiology which may predispose or protect against certain diseases and their comorbidities



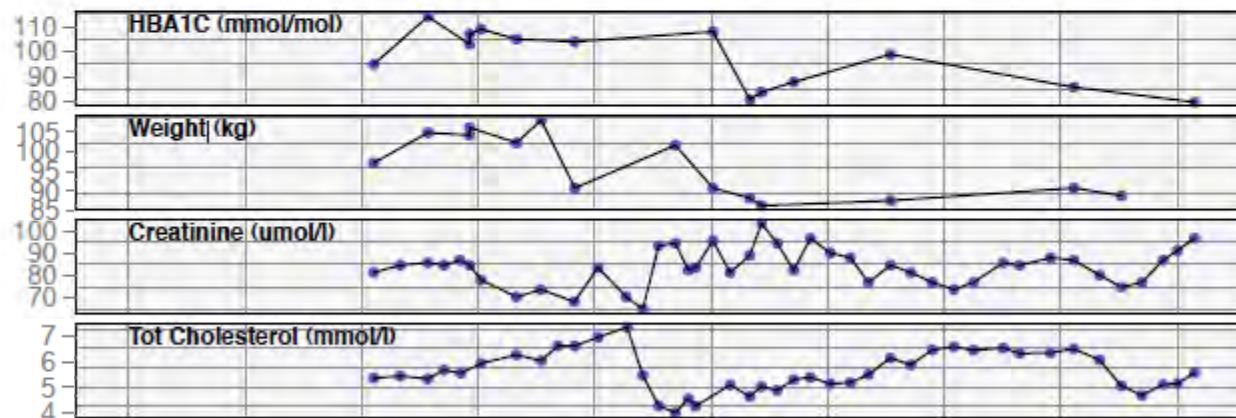
Diagnoses



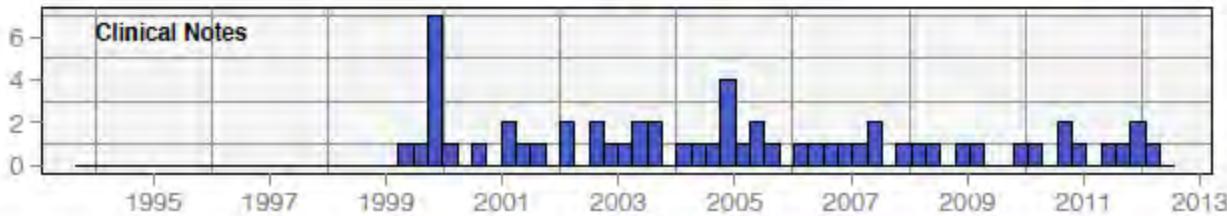
Drugs



Lab tests



Text

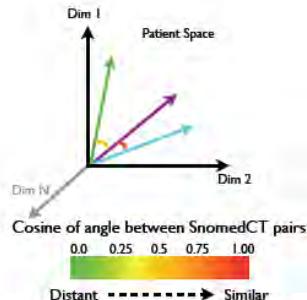


Deep-phenotyping by text mining of ICD10 terms in patient records

F20

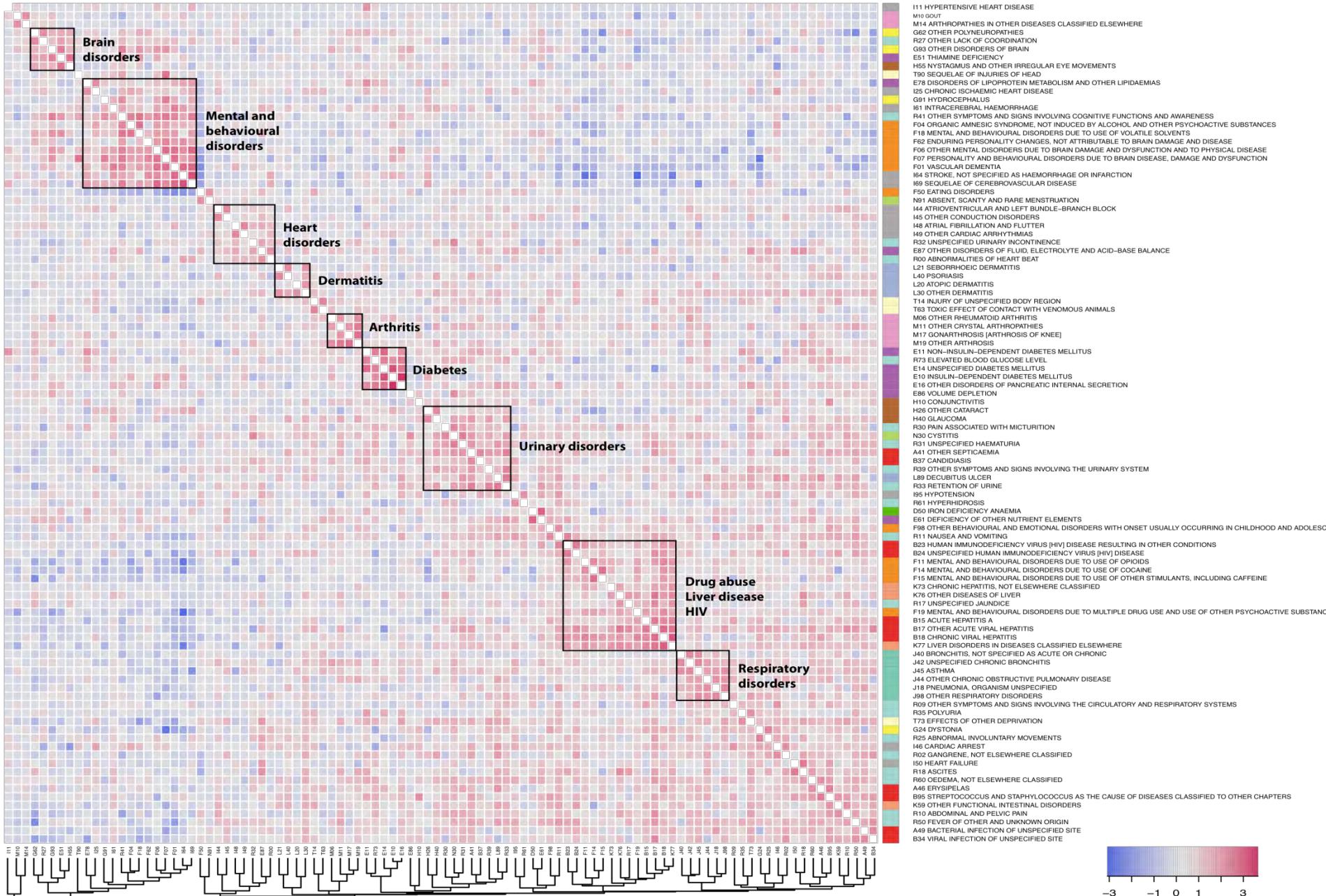
F200

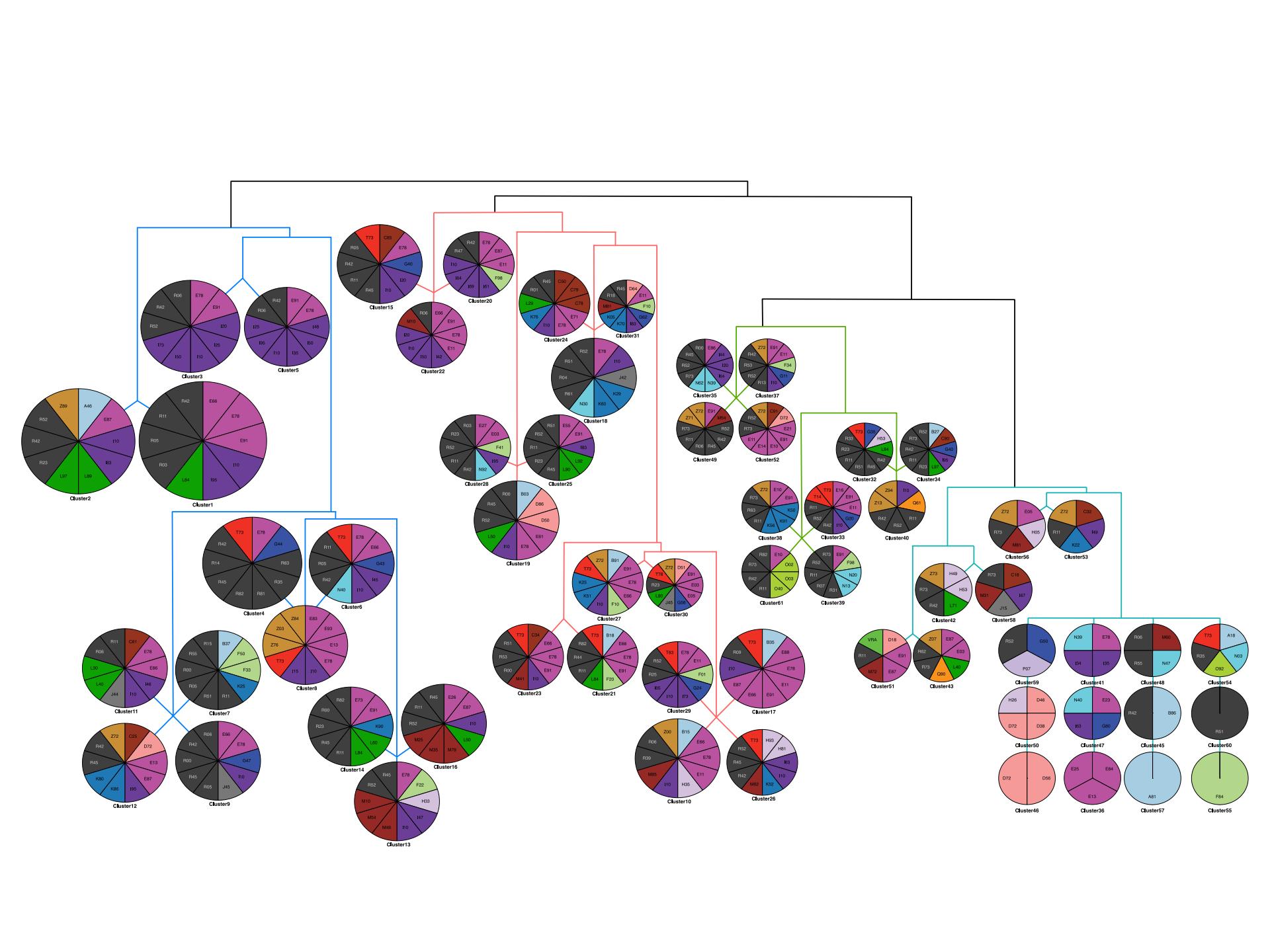
det drejer sig om en 36-årig sygemeldt mand der overflyttes fra frederiksberg hospital, afdeling m.h.p. længerevarende rehabiliteringsophold. , er allergisk overfor kat og parfume, men tåler penicillin. er i besiddelse af en vis indsigt og virker svært forpint. ang. det at vi tilråder, at hun har brug for at være mere i afd. , siger hun til det, at det for hende er som at vælge mellem pest eller kolera. Har stadig mange spørgsmål omkring **skizofreni** og er meget bekymret for hvordan hendes fremtid ser ud. er meget plaget af tanketræghed og er bange for at det er et led i sygdommen. der siges til hende at det godt kan være bivirkning af risperdal men at der ikke laves om på medicinen, før vi har lært hende bedre at kende. Har **aldrig** haft **hallucinationer** på nogen af sanserne har været til lægesamtale idag. der snakkes en del om diagnose og at pernille har svært ved at forholde sig til at have **diagnosen skizofreni**, det virker som om pernille er blevet lidt mere afslappet, selvom hun stadig har gang i mange ting. pt. møder til samtale i dag, hvor vi gennemgår mit udkast til erklæringen til pensionskassen. endvidere udspørges der til pt.s diverse symptomer på **paranoid skizofreni**. i denne beskriver hun at "hendes største problem nok er den manglende sociale evne, som er en følge af sygdommen (**paranoid skizofreni**) og henviser til contras beskrivelse" Pt. Nævner sin **mor**, som han mener har en nervøs lidelse, muligvis **social fobi** pt. har her til aften angivet tiltagende **bivirkninger** i form af trækninger i nakken, indre uro og stivhed af fingre. pt. har fået svar på sit ekg, som viser sinus rytme med enkelte **ventrikulære ekstrasystoler** uforandret fra tidl. med baggrund i oplysninger om tidligere maniske episoder præget af irritabilitet, hyperaktivitet og øget seksuel interesse revurderes diagnosen til **bipolar affektiv sindslidelse**. følges i distrikt vest med psykologsamtaler. har i dag tydeligvis brug for en faglig forklaring på hendes symptomer. det drejer sig om **paranoia**, uvirkelighedsfølelses, influenssympt. og koncentrationsbesvær. det største problem er dog samværet med andre. det er specielt om natten det påvirker hendes **astma**, klg. desuden over uro i benene. ,xxx nævner på et tidspunkt, hun er bange for, tidlige tiders **spiseforstyrrelser** er ved at dukke op igen. xxx har haft **søvnbesvær** og har af vagtlægen i aftes fået tabl. imovane 7,5 mg med god effekt. kl 19, pinex, tabletter 500 mg indtaget dosis: 1 gram for hovednigne. pt. er henviset til at



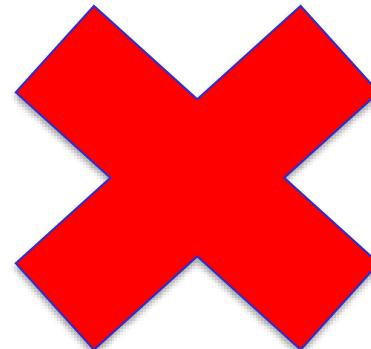
Negation

Family





Which disease-disease and symptom correlations are treatment related?



Text mining of drug names, ADE/ADRs, diagnoses, ...

Removed ADR - no corresponding structured data

Behandlet med Zyprexa 5 mg fra 3. til 24.6.99 og 10 mg fra 24. til 29.6.99 med nogen effekt på vægtstygmelser, men seponeret pga appetitoøgning. Herefter Risperdal 2 mg stigende til 4 mg i perioden 29.6. til 12.7.99, men seponeret på grund af uro i kroppen og "osteklokkefornemmelse". Herefter Oraj 2 mg fra 2.8. stigende til 3 mg fra 30.8.99 med god effekt på tankeekko og tankemylder. Behandlet med Zoloft 50 mg fra maj 98 til maj 99 med noget virkning på depressive symptomer; men seponeret på grund af hatlig svættendens. Siden 14.7.99 Elxior 75 mg med nogen effekt på antallet og sværhedsgraden af kortvarige depressive episoder.

se venligst under allergier. Desuden forsøgt beh med Zyprexa, sep grundet vægtøgning, træthed og manglende effekt. Risperdal ord med nogen effekt tillagt dogmat(1999). Efterfølgende aurix beh seponeret i 1999. Startede istedet reme. Aktuel medicindosis, jvf udskrivningsnotat fra UI2 samt EPJmedicinliste.

tbl. Ieponex 100+0+0+200 mg.tbl. rivotril 0,5 +0+1+0 mg.tbl. arintaprin 0+0+0+30 mg.tbl. clomipramin 0+0+0+25 mg.tbl. imocdone 7,5 mg nocte.tbl. rivotril 2 mg p.n. max x 1 dgl. tbl. marenvan a 2,5 mg efter skema.tbl. magnesia 1 g p.n. laxoberaldr 7,5 mg /ml 15 dr.p.n. mix. link 150 mg /ml 15 ml p.n. max x 3 dgl. Figensaft 20+0+20+0 ml.Pt. er aktuelt,CAVE,tricykliske antidepressiva. Dette kan dog ikke bekræftes og pt. har tidl. fået imipramin, som han har fået godt, hvorfor der er ansøgt om opnævelser af denne cave på højere niveau. Har tidl. fået zyprexa som blev sep. grundet vægtøgning, træthed og manglende effekt

Removed ADR - negation and subject identification

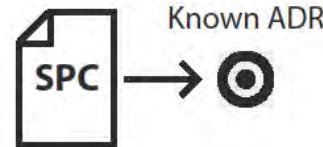
–Jeg mener fortsat, at han har brug for medicin, da han i går fx var meget vred og følte sig utryg og angst og har haft svært ved at sove. Dette synes pt. at accepterer. Jeg tilbyder herefter Zyprexa i stedet for Risperdal, pt. avisør dette, da han ved patient har fået denne medicin og har fået angst, dette vil han ikke. Har ikke tidligere fået antipsykotisk medicin. Accepterer herefter Cisordin, startende på en lille dosis. Accepterer også angstdæmpende medicin i dagtiden. Angiver, at når han bliver vred kan han godt styre det. Synes det hjalp i går noget at få Nozinar. Virker fortsat garderet. Siger intet upfordret. Sparsomt sygdomsindsigt. Er ..., predilader. Er i dag ..., heller latent aggressive...

Text mining Adverse Drug Reactions (using 7,500 drug names and 21,000 ADRs)

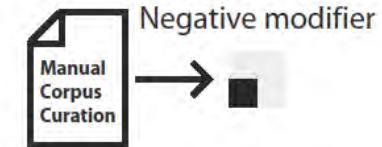
Identification of indications



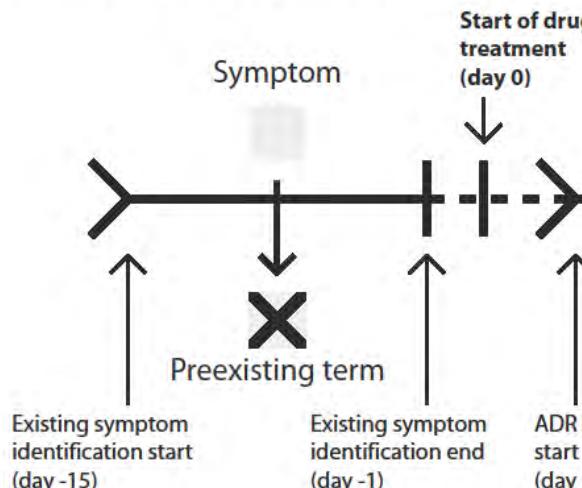
Identification of known ADRs



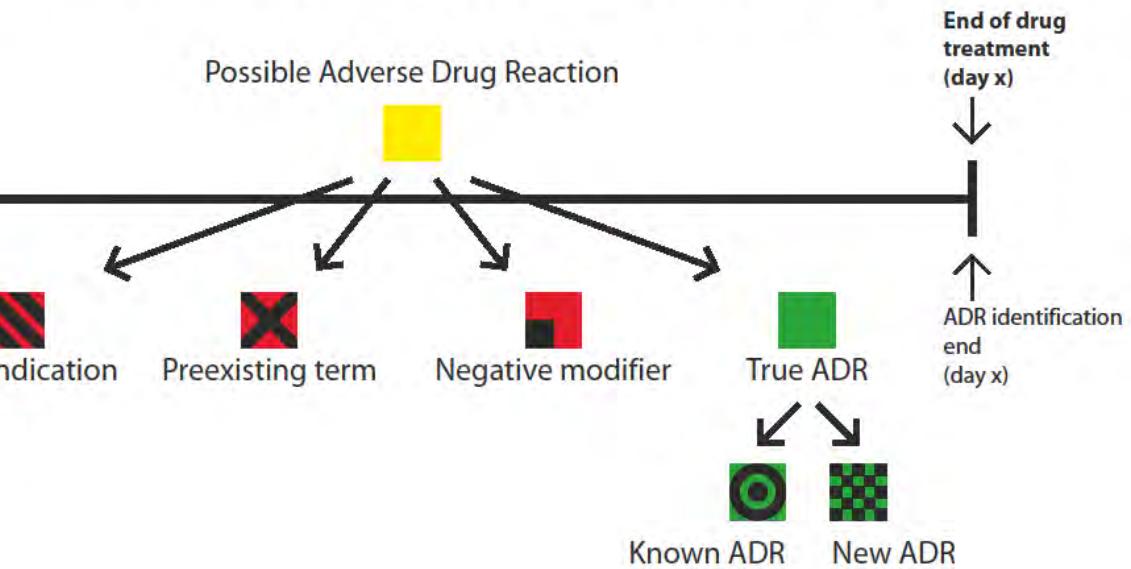
Creation of negative modifiers



Identification of existing symptoms



Identification of possible Adverse Drug Reactions

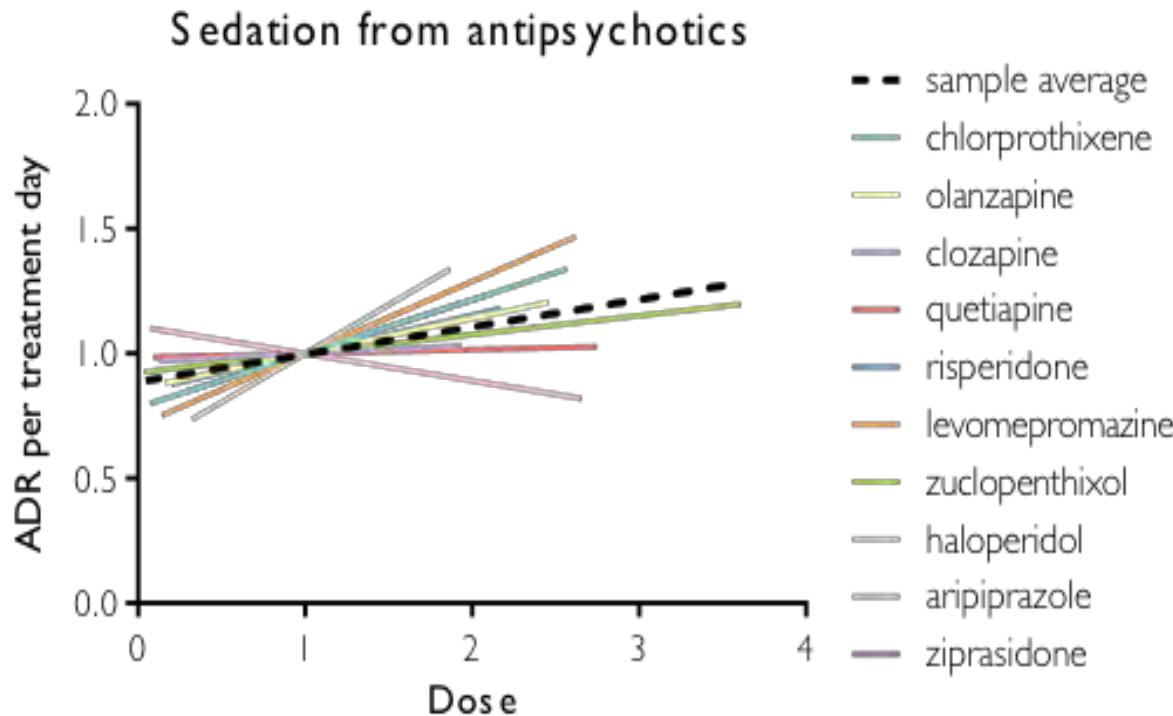


Data from two Danish regions



ADR-dose dependencies

Dosages from structured medication data



ADRs and doses are normalized on multiples of the minimum dose prescribed of each drug.

Plot for 21 days steady dosage data is visualized, sample average slope 0.1105 (95% CI, 0.03085-0.1901), non-zero slope p-value was 0,0074, all individual drug slopes are positive except for haloperidol.

△ population health data

- **Health data driven:**
 - Redefine phenotypes as trajectories
 - Enable prediction using predictable trajectories?

- **Include what is not in patient records in new ways:**
 - Diet,
 - Income, ...
 - Education, grades in exams, ...
 - Behavior, risk-taking, ...





Personlig Medicin og Individualiseret Behandling

Oplæg til en samlet dansk indsats

22. juni 2015



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EPR and registry analysis

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TRANSLATIONAL GENETICS

Mining electronic health records: towards better research applications and clinical care

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Abstract | Clinical data describing the phenotypes and treatment of patients represents an underused data source that has much greater research potential than is currently realized. Mining of electronic health records (EHRs) has the potential for establishing new patient-stratification principles and for revealing unknown disease correlations. Integrating EHR data with genetic data will also give a finer understanding of genotype–phenotype relationships. However, a broad range of ethical, legal and technical reasons currently hinder the systematic deposition of these data in EHRs and their mining. Here, we consider the potential for furthering medical research and clinical care using EHR data and the challenges that must be overcome before this is a reality.

Clinical decision support (CDS). Software systems providing support for decision making to physicians through the application of health knowledge and logical rules to patient data.

Biobanks
Central repositories of biological material that are mainly used for research. They facilitate the re-use of collected samples in different research projects.

Information technology has transformed the way health care is carried out and documented. Presently, the practice of health care generates, exchanges and stores huge amounts of patient-specific information. In addition to the traditional clinical narrative, databases in modern health centres automatically capture structured data relating to all aspects of care, including diagnosis, medication, laboratory test results and radiological imaging data.

This transformation holds great promise for the individual patient as richer information, coupled with clinical decision support (CDS) systems, becomes readily available at the bedside to support informed decision making and to improve patient safety^{1,2}.

especially interesting when traditional health-care-sector data is linked with biobanks and genetic data⁴.

Despite the great potential, researchers who wish to analyse large amounts of patient data are still faced with technical challenges of integrating scattered, heterogeneous data, in addition to ethical and legal obstacles that limit access to the data^{5,6}. It is hoped that large-scale adoption of health information technology (HIT) infrastructure in the form of electronic health records (EHRs) and agreed standards for interoperability and schemes for privacy and consent, will improve this situation (TABLE 1). With incentives for improved public health and the expected health budget savings^{7,8}, these matters