Text mining to identify potential mechanisms linking sedentary behaviour to disease outcomes



## **Understanding mechanisms**



- Improve causal inference
- Easier/more effective to intervene on mediator
- Alter exposure in way to maximise mediation effect



# How to identify plausible mechanisms?

Search for common terms (potential mechanisms) in literature using PubMed®



A overlapping terms identified by Boolean operator 'and'



# How to identify plausible mechanisms?

Search for common terms (potential mechanisms) in literature using text mining



- A overlapping articles (with common terms)
- B non-overlapping articles (with common terms) associated with both risk factor and disease





# **Example - BDNF**

- Protein expressed by hypothalamus
- Also excreted by skeletal muscle during exercise
- Exercise and cognitive function research (humans)
- Mouse models have shown upregulation of BDNF reduces tumour burden (various cancers)



## **Text-mining tools**

â

University of BRISTOL

**MRC Integrative Epidemiology Unit** 

Data and analysis tools

## Current students Current staff Alumni

### Integrative Cancer Epidemiology Programme

#### Research themes

**Research outputs** 

→ Publications

⊢ Grants

### → Data and analysis tools

People

Collaborations

Impact and engagement

Contacts





### **MR Base**

MR Base enables online Mendelian randomization analysis using a comprehensive manually curated database of GWAS studies.



### MELODI

MELODI is a literature mining platform to identify potential mechanistic pathways between exposures and disease outcomes.

BRIAD BARRIE



"MR Base has transformed the ability of researchers world-wide to run rapid two-sample Mendelian randomization analyses, which can form the basis for in-depth analyses of many important questions regarding possible prevention or treatment of disease"

- PROF GEORGE DAVEY SMITH



### <u>TeMMPo</u>

**TeMMPo** is a literature search tool to quantify the literature for specific disease mechanism.

Equesre A Planadas A Cutore

### LD Hub

LD Hub supports online LD score regression analysis using a comprehensive manually curated database of GWAS studies.

# **TeMMPo tool**

- Text Mining for Mechanism Priorisation
- Uses Medical Subject Headings (MeSH) system
- Identifies co-ocurrence of MeSH headings in publications
   → Link 'mechanism' to exposure and/or outcome
- Targeted approach: specify a priori potential mechanism terms







International Journal of Epidemiology, 2018, 1–11 doi: 10.1093/ije/dyx251 Software Application Profile



Software Application Profile

### MELODI: Mining Enriched Literature Objects to Derive Intermediates

### Benjamin Elsworth,<sup>1</sup>\* Karen Dawe,<sup>1</sup> Emma E Vincent,<sup>1</sup> Ryan Langdon,<sup>1</sup> Brigid M Lynch,<sup>2-4</sup> Richard M Martin,<sup>1</sup> Caroline Relton,<sup>1</sup> Julian P T Higgins<sup>1</sup> and Tom R Gaunt<sup>1</sup>

<sup>1</sup>MRC Integrative Epidemiology Unit, University of Bristol, Bristol, UK, <sup>2</sup>Cancer Epidemiology and Intelligence Division, Cancer Council Victoria, Melbourne, VIC, Australia, <sup>3</sup>Centre for Epidemiology and Biostatistics, University of Melbourne, Melbourne, VIC, Australia and <sup>4</sup>Physical Activity Laboratory, Baker Heart and Diabetes Institute, Melbourne, VIC, Australia

\*Corresponding author. MRC Integrative Epidemiology Unit (IEU), Bristol Medical School: Population Health Science, Oakfield House, Oakfield Grove, University of Bristol, Bristol BS8 2BN, UK. E-mail: ben.elsworth@bristol.ac.uk

Editorial decision 2 November 2017; Accepted 3 January 2018



- A text mining platform designed to identify and prioritise intermediates between two datasets.
- Uses PubMed to create datasets (exposure and outcome).
- MELODI looks for overlapping
  - MeSH terms
  - Single SemMedDB concepts
  - SemMedDB 'triples' (subject predicate object) datasets.
- Enrichment step compares frequency of terms within datasets to background frequency in PubMed.



- Enrichment using OR and two-tailed FET.
- P-values corrected for multiple testing using the Benjamini/Hochberg (non-negative) correction with a cutoff of p<1e-5.</li>
- The results (corrected p-value and OR) used to filter results.
- Also uses frequency of predicate term (SemMedDB Triple) and minimum position in the MeSH hierarchy (MesH method).



- Visualisation of results via Sankey plot.
- SemMedDB Triples method also shows directed network diagram.
- These graphs are 'live' and change as filters applied.
- Once spurious intermediates removed and filters applied need to manually curate results.



Search results are retained if they represent a biological marker that could plausibly be associated with both the exposure and outcome. Two independent extractions!

Exclusion criteria for intermediates:

•synonyms, antonyms or similar terms for the exposure or outcome

•scientific methods, diagnostic tests, therapies (including drugs), anatomical or physiological nomenclature, comorbid conditions.



## **Example application**

Identifying known and novel mechanisms underpinning association between sedentary behaviour and breast cancer risk





## **Strengths and limitations**

- Strengths:
  - Sophisticated text-mining techniques
  - Mechanism discovery
  - Combining literature from different fields



- Limitations:
  - Co-occurrence in article  $\rightarrow$  not necessarily association
  - Multiple intermediates in pathway: Not really possible to combine



## **Acknowledgements**

### **University of Bristol, UK**

Sarah Lewis Richard Martin Tom Gaunt Benjamin Elsworth Luke Robles Karen Dawe Julian Higgins







**NBCF** Career Development Fellowship 2015 – 2018



**UICC** Yamagiwa-Yoshida Memorial International Cancer Study Grant Jan – Mar 2017









## brigid.lynch@cancervic.org.au



http://www.bristol.ac.uk/integrative-epidemiology/faciliitiesresources/software/



## **Further enquiry**

