TraitHunter

Mapping and extraction of biomedical traits via text embeddings

IEU Monthly Meeting

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https://gwas.mrcieu.ac.uk/datasets/ieu-b-40



body mass index Dataset: ieu-b-40

Download VCF	ownload index View report		
PMID	30124842	Top 30 related datasets 3	
Year	2018	ieu-b-40: body mass index	
Category	Continuous	ieu-a-1089: Body mass index	
Sub category	Anthropometric	ieu-a-974: Body mass index	
Population	European	ieu-a-95: Body mass index	
Sex	Males and Females	ebi-a-GCST004904: Body mass	
Sample size	681,275	index	
Number of SNP	2,336,260	ebi-a-GCST006368: Body mass index	
Unit	NA	bbj-a-2: Body mass index	
Author	Yengo, L	ieu-a-835: Body mass index	
Consortium	GIANT	ieu-a-2: Body mass index	

- Originally powered by an internal trait recommender service, *Vectology*
 - Elsworth, Liu, Gaunt, 2019, 1st International "Alan Turing" Conference on Decision Support and Recommender Systems
- Applied methods / models in what we call today as "Large Language Models" (LLMs) or Foundation Models
 - BioSentVec
 - BERT, BioBERT, BlueBERT
- The service is now powered from EpiGraphDB
- That was 2019. We are now almost 2025. Time for an upgrade.

Today's talk

The main aim for today's talk is to showcase our next-gen web service

TraitHunter

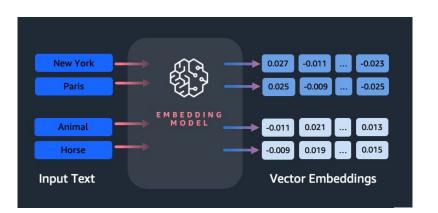
https://traithunter.epigraphdb.org for mapping and identifying biomedical trait.

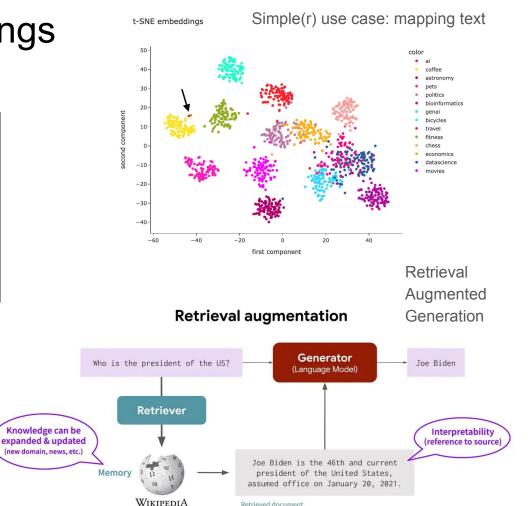
This is very preliminary, however feedbacks on various aspects are deeply appreciated.

Outline

- Background
- Key methods
- TraitHunter live demo
- Next steps

Concept: Text embeddings





Convert text into their (semantic) vector representations via an encoder model

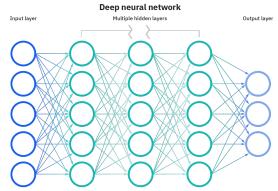
Retrieved document

The Free Encyclopedia

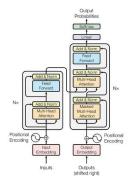
Knowledge can be

(new domain, news, etc.)

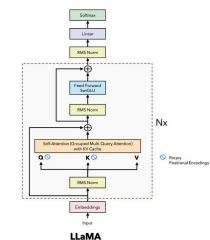
Concept: Large Language Models (LLM)

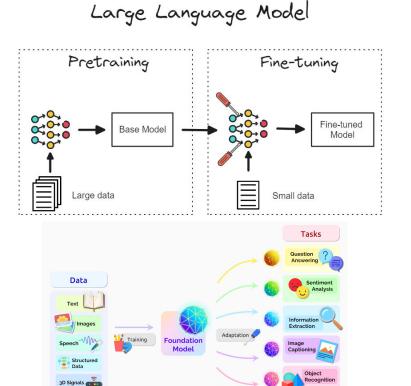


Transformer vs LLaMA



Transformer ("Attention is all you need")





Instruction

Following

Background: previous works

In Liu, et al. (2023) we approached the trait mapping problem with a two stage approach:

- Identify a sub set of related traits with naive text embeddings, from the broad set
- Map the traits using an LLM task model finetuned on ontologies

This approach is put into use in the ASQ platform (Liu & Gaunt, 2024) for querying EpiGraphDB evidence.

JOURNAL ARTICLE

Using language models and ontology topology to perform semantic mapping of traits between biomedical datasets 3

Yi Liu, Benjamin L Elsworth, Tom R Gaunt 🕿 🛛 Author Notes

Bioinformatics, Volume 39, Issue 4, April 2023, btad169, https://doi.org/10.1093/bioinformatics/btad169 Published: 03 April 2023 Article history ▼

JOURNAL ARTICLE ACCEPTED MANUSCRIPT Triangulating evidence in health sciences with Annotated Semantic Queries Yi Liu S. Tom R Gaunt S

Bioinformatics, btae519, https://doi.org/10.1093/bioinformatics/btae519
Published: 22 August 2024 Article history
→

Limitations:

- Locked to a specific ontology (EFO)
- Finetuned task model not widely adaptable
- Just using naive embeddings might be preferable with the latest-and-greats of LLMs (in ASQ next gen)

Why yet another LLM based recommender service

Models



- LLaMA3

and other good foundation models

Tooling

- Huggingface and other tools (ollama) are now much more mature
- Elasticsearch only recently supports
 4095 dim dense vector embeddings

Research

- We collaborated with a few research teams on the problem of trait mapping for pre-select traits of interest
- So we probably should have a formal service and short paper up so collaborators can cite it
- In preparation for a few research projects

TraitHunter https://traithunter.epigraphdb.org

- Map **traits** across curated biomedical **dictionaries**
- Typical example: the trait recommender on OpenGWAS --OpenGWAS to OpenGWAS
- UKBiobank to ALSPAC
- OpenGWAS to EFO

A search engine for biomedical traits

- Identifying a trait from biomedical sources
- · "Trait"
 - a label from a source
 - MONDO_0007254 breast cancer
 - ieu-b-40 body mass index
 - a *named entity* based on the NER model
 - "age", "body", "BMI"
- Examples:
 - Find / map a UKBiobank variable with OMIM text
 - If a grant project is associated a trait

TraitHunter: concepts

Dictionary

A vocabulary set of trait entities

Regular examples:

- UK Biobank variables
- Trait names of OpenGWAS studies
- EpiGraphDB terms

Ontologies

- Hierarchical from general to specific concepts

Trait entity

- ID
- Label
 - Concept singleton
 - Complex trait
- (optional) Description
- Named entities
- optional for ontologies
 - Synonyms
 - Ontological parents, children, and siblings

Glycogen storage disease due to aldolase A deficiency MONDO:0012747

Glycogen storage disease due to aldolase A deficiency is an extremely rare glycogen storage disease characterized by hemolytic anemia with or without myopathy or intellectual deficit. Myopathy can be severe enough to result in fatal rhabdomyolysis in some patients. A family with episodic rhabdomyolysis (triggered by fever) without hemolytic anemia has recently been reported.

Report Entry Issue

breast cancer IMPORTED

Export Associations

🗹 http://purl.obolibrary.org/obo/MONDO_0007254 🍺 Copy

A primary or metastatic malignant neoplasm involving the breast. The vast majority of cases are carcinomas arising from the breast parenchyma or the nipple. Malignant breast neoplasms occur more frequently in females than in males.

Defined by		IDO			
Also appe	ars in	CPONT GENEPIO	ОВА		
Synonym	BC	breast cancer 🛈	breast tumor	breast tumou	r ① cancer of breast ①
malignar	nt brea	ast neoplasm 🕕 🛛 🖿	nalignant breast tur	nor O malig	nant breast tumour

114480

BREAST CANCER

Alternative titles; symbols

BREAST CANCER, FAMILIAL

Other entities represented in this entry:

BREAST CANCER, FAMILIAL MALE, INCLUDED

Description

Breast cancer (referring to mammary carcinoma, not mammary sarcoma) is histopathologically and almost certainly etiologically and genetically heterogeneous. Important genetic factors have been indicated by familial occurrence and bilateral involvement.

Clinical Features

Cady (1970) described a family in which 3 sisters had bilateral breast cancer. Together with reports in the literature, this suggested to him the existence of families with a particular tendency to early-onset, bilateral breast cancer. The genetic basis might, of course, be multifactorial. ◆

Anderson (1974) concluded that the sisters of women with breast cancer whose mothers also had breast cancer have a risk 47 to 51 times that in control women, a revised estimate was 39 times (Anderson, 1976). The disease in these women usually developed before menopause, was often bilateral, and seemed to be associated with ovarian function. About 30% of daughters with early-onset, bilateral breast cancer inherited the susceptibility. The risk of breast cancer to women with affected relatives is higher when the diagnosis is made at an early age and when the disease is bilateral. Ottman et al. (1983) provided tables that give the cumulative risk of breast cancer to mothers and sisters at various ages. The highest risk group is sisters of premenstrual probands with bilateral disease. Among the sisters of women with bilateral disease, an affected mother (25 +/-7.2%), or an affected sister (28 +/-11%). The risks were reduced to 18 +/-3.3% and 14 +/-2.6%, respectively, with unilateral disease. An early example of familial breast cancer was provided by

TraitHunter: data curation

Curated

- OpenGWAS
- ICD10
- HPO
- UKBiobank

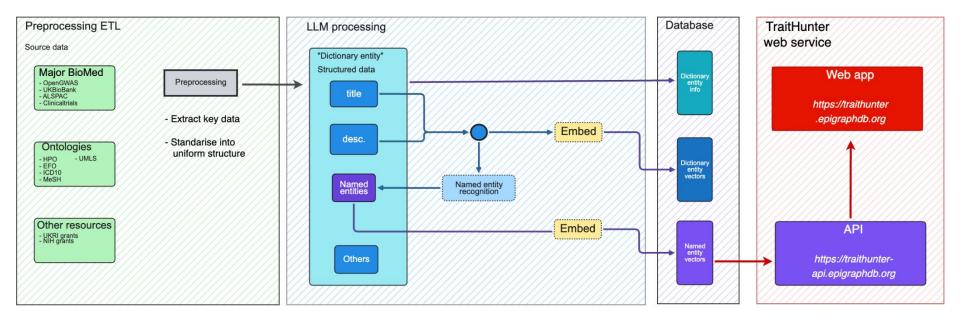
In preparation

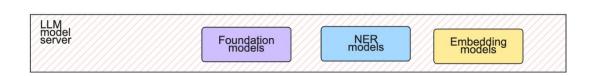
- EFO
- MeSH
- UKRI grants
- NIH grants
- Clinicaltrials

Next step

- UMLS
- OMIM
- All of Us

TraitHunter architecture





- Web app: <u>https://traithunter.epigraphdb.org</u>
- API: https://traithunter-api.epigraphdb.org
- code: https://github.com/MRCIEU/traithunter

TraitHunter: Demo

About Misc. information for this platform	Trait Hunter TraitHunter is a platform to search and map biomedical traits across
Trait mapping Map a trait to other traits by semantic similarity	various major dictionaries. The dictionaries we curate and offer search functionalities include: • OpenGWAS traits (2024-08)
Pairwise similarity Compute the pairwise semantic similarity of	 ICD10 codes (2024-08) HPO ontology terms (2024-08) UKBiobank variables (2024-08)

Trait map

Target entity

Select dictionary of the target entities

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Explore curated data (work in progress)	
--	--

API connected: true

Data explorer

		С В	RL Didemiology Unit	BRISTOL		Select dictionary
					Pairwise similarity	Select dictionary
Trait Hunter					Compute the pairwise semantic similarity of	Step 2: Configure par
TraitHunter is a platform to search and map biomedical traits across various major dictionaries.					Data explorer	Other parameters
The dictionaries we curate and offer search functionalities include:					Explore curated data (work in progress)	Select embedding model bge
OpenGWAS traits (2024-08) (CD10 codes (2024-08) HPO ontology terms (2024-08) UKBiobank variables (2024-08)					API connected: true	
Trait mapping						
Configure search parameters Search and map a trait to trait in other dictionaries via the text embeddings of its label (k-Nearest Neighbour search).					Data Evala	ror (pilot)
How to use 1. Select the dictionary of the source entity 2. Search for the source entity of interest by its label 2. Select the dictionary of the target entities 3. Configure other variables 4. Click of the source bumb tottom					Data Explor Data explorer for entities NOTE: this is just a prev	
Source entity		Other parameters Select embedding model			Select dictionary hpo	
Select dictionary	-	bge		*		
Choose a dictionary first	Ŧ	Top K neighbors			Entity id	

		C T T DB	Unit Unit	BRISTOL
About Misc. information for this platform Trait mapping Map a trait to other traits by semantic similarity	Pairwise cosine similarities Compute the pairwise cosine similarity scores for the indued list of trait entities via the text embeddings of their labels. Step 1: Search for entities to include in the comparison			
Pairwise similarity Compute the pairwise semantic similarity of	Select dictionary Choose a dictionary first Step 2: Configure parameters		-	ADD TO LIST
Data explorer Explore curated data (work in progress)	Other parameters Select embedding model bge ~			
PI connected: true	SUBMIT			

Graph University of Endemnider

change.

Select dictionary hpo	•	UPDATE TABLE
Entity id	Entity label	Description
http://purl.obolibrary.org/obo/CHEBI_131604	Mycoplasma genitalium metabolite	Any bacterial metabolite produced during a metabolic reaction in Mycoplasma genitalium.
http://purl.obolibrary.org/obo/CHEBI_131604- -1	Mycoplasma genitalium metabolites	Any bacterial metabolite produced during a metabolic reaction in Mycoplasma genitalium.
http://purl.obolibrary.org/obo/CHEBI_131619	C27-steroid	A steroid compound with a structure based on a 27-carbon (cholestane) skeleton.
http://purl.obolibrary.org/obo/CHEBI_131619- -1	C27-steroids	A steroid compound with a structure based on a 27-carbon (cholestane) skeleton.
http://purl.obolibrary.org/obo/CHEBI_131621	C19-steroid	A steroid compound with a structure based on a 19-carbon (androstane) skeleton.
http://purl.obolibrary.org/obo/CHEBI_131621- -1	C19-steroids	A steroid compound with a structure based on a 19-carbon (androstane) skeleton.
http://purl.obolibrary.org/obo/CHEBI_131702	stigmastane derivative	Any steroid (or derivative) based on a stigmastane skeleton.

Next steps, in infra development

Assessment of embedding quality with benchmarks

 How well does an embedding model objectively perform for the task of mapping biomedical traits Named entity recognition

- Identification and extraction of a trait entity from full text docs.

(maybe) Molecular symbols

- Special treatment for gene names, etc.

Next steps, in research

The TraitHunter paper

- Describing the service and methods

Your input is needed on what this platform can help with your research.

Clustering analysis

Investigation on the different clusters of biomedical relationships for novel insights

- Semantics
- Genetic associations

RAG

RAG on knowledge graphs to investigate about biomedical assertions (i.e. ASQ-next-gen).

Acknowledgements

MRC IEU

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- Zhaozhen Xu
- Gibran Hemani
- (alumni Benjamin Elsworth)

Thanks for listening. A preprint will be up this month.

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Naomi Cornish

Andrew Mumford

Questions / feedback welcome

web app: https://traithunter.epigraphdb.org

Million Veteran Program

- Brian Ferolito
- Daniel Golden
- Alexandre Pereira



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