Imperial College London centre for antimicrobial optimisation Health (

Co-morbidity Representation in Artificial Intelligence: **Tapping into Unused** Clinical Knowledge

William Bolton^{1, 2, 3, 4}, Pantelis Georgiou^{3, 4, 5}, Alison Holmes^{3, 4, 6, 7}, Timothy Rawson^{3, 4}

¹Department of Computing, Imperial College London, UK.²UKRI Centre for Doctoral Training in AI for Healthcare, Imperial College London, UK. ³Centre for Antimicrobial Optimisation, Imperial College London, UK ⁴National Institute for Health Research, Health Protection Research Unit in Healthcare Associated Infections and Antimicrobial Resistance, Imperial College London, UK. ⁵Centre for Bio-inspired Technology, Department of Electrical and Electronic Engineering, Imperial College London, UK. ⁶Faculty of Health Life Sciences, University of Liverpool, UK. ⁷Department of Infectious Diseases, Imperial College London, UK.

Why?

- **Co-morbidities** defined here as chronic long-term medical conditions are a major challenge in healthcare¹
- **Challenges exist** with representing and using co-morbidity data in AI systems¹:
 - **Combinatorial complexity** due to the large number of unique diseases
 - Sparsity, missingness and a lack of data particularly for those with rare diseases or complex co-morbidity combinations
 - Heterogeneity in how chronic conditions are recorded
- Existing AI research on co-morbid patients does not tackle these problems and therefore lacks appropriate representation

How?

- Processed **SNOMED CT**² a comprehensive clinical healthcare terminology into a connected undirected graph
- Generated **disease embeddings** through Node2vec³ with optimization to reduce the mean SNOMED distance (shortest path length) between each node and their nearest neighbor
- Tested disease embeddings as sole features to **supervised learning models** for clinically relevant predictions
- Defined **co-morbid patient embeddings** as the mean of all the

Aim

Creating meaningful embeddings from external medically grounded knowledge, to help overcome such challenges and support downstream AI applications

Results

SNOMED disease embeddings for a particular individual

- Evaluated co-morbid patient embeddings through a **task to** retrieve the most similar patient for any given unique comorbid patient, where no identical match was possible
 - Retrieved similar patients in our method through **nearest neighbor lookup** based on euclidian distance
 - Utilized two **novel metrics** and **human experts** as evaluators



nearest neighbor SNOMED distance (hyperparameter optimisation resulted in a mean of 1.23) and [B] high-level disease groups displayed.

Table 2: Mean evaluation results for the similar patient retrieval task.

	SNOMED	Charlson
Method	similarity	Jaccard
	score	index
One hot encodings	4.40 (SD 2.32)	0.88 (SD 0.30)
Rocheteau's method ⁴	3.52 (SD 3.26)	0.69 (SD 0.20)
Co-morbid patient embeddings	1.78 (SD 1.90)	0.84 (SD 0.34)

Discussion

- We developed a **novel pipeline** to extract and utilize untapped medical knowledge and demonstrated its utility in classification and similar patient retrieval tasks with automatic and human evaluation
- Our approach is generalizable and can overcome some problems with using disease data in AI systems as the embeddings are not influenced by dataset size, the number of diseases or their rareness and are adaptable to variation in clinical documentation
- Future work includes, considering **temporal** aspects and embedding additional clinical data such as demographics and medications

Figure 2: Equations used to determine SNOMED similarity score and Charlson Jaccard index.

Co-morbidities



in question and the similar patients retrieved by each method. The co-morbidities similarity of IS through a traffic light indicated coloring scheme.



Figure 3: Proportion of human expert votes for patients identified by each method, for each question, in a survey. Co-morbid patient embeddings obtained the most votes for 6 questions.

