

a Data Resources



For Development

All of Us
RESEARCH PROGRAM



For Validation

biobank^{uk}



Electronic Health Record

e.g.,
• conditions
• medications

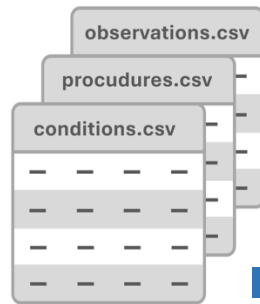


Whole Genome Sequencing

1257 identified independent genome-wide significant SNPs

b Data Preprocessing

EHR Data under OMOP CDM

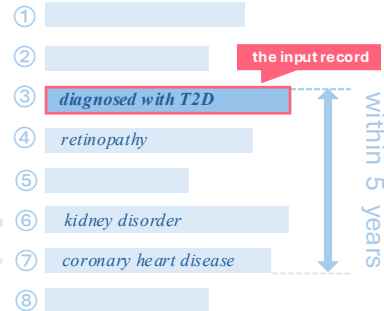


Flattening

multi-hot labeling



Per-patient Visit Timeline

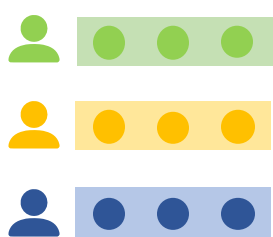


c Hypergraph-based Prediction & Subtyping

Model Input

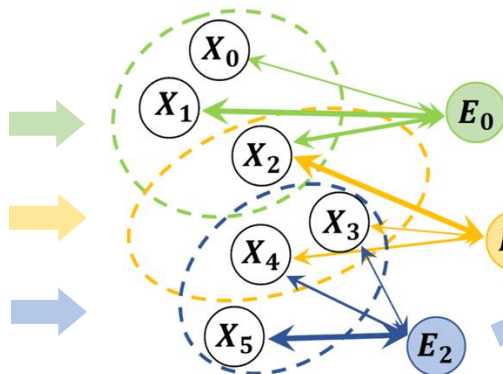
initial T2D diagnosis visit
for each patient
(diabetic codes excluded)

Records



Hypergraph Learning

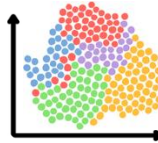
EHR codes as nodes,
patient visits as hyperedges



Self-supervised Clustering



clusters
assignment



Predicting Complications



predicted labels

0	0	1	1	0	1
1	1	0	0	0	0
1	0	0	1	0	1