

Machine learning to identify people who inject drugs for hepatitis C surveillance

Carol El-Hayek, Thi Nguyen, Margaret Hellard, Michael Curtis, Anna Wilkinson, Rachel Sacks-Davis, Nick Scott, Jason Asselin, Paul Dietze, Annie Madden, Htein Linn Aung, Rebecca Guy, Mark Stoové, Douglas Boyle, Jane Hocking, Adam Dunn

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AT BURNET INSTITUTE, WE PROUDLY ACKNOWLEDGE
THE BOON WURRUNG PEOPLE OF THE KULIN NATIONS
AS THE TRADITIONAL CUSTODIANS OF THE LAND ON
WHICH OUR OFFICE IS LOCATED. WE PAY OUR
RESPECT TO ELDERS PAST AND PRESENT, AND EXTEND
THAT RESPECT TO ALL FIRST NATIONS PEOPLE.





Hepatitis C elimination

PEOPLE WHO INJECT DRUGS

- Primary risk group
- Priority population

HEPATITIS C SURVEILLANCE

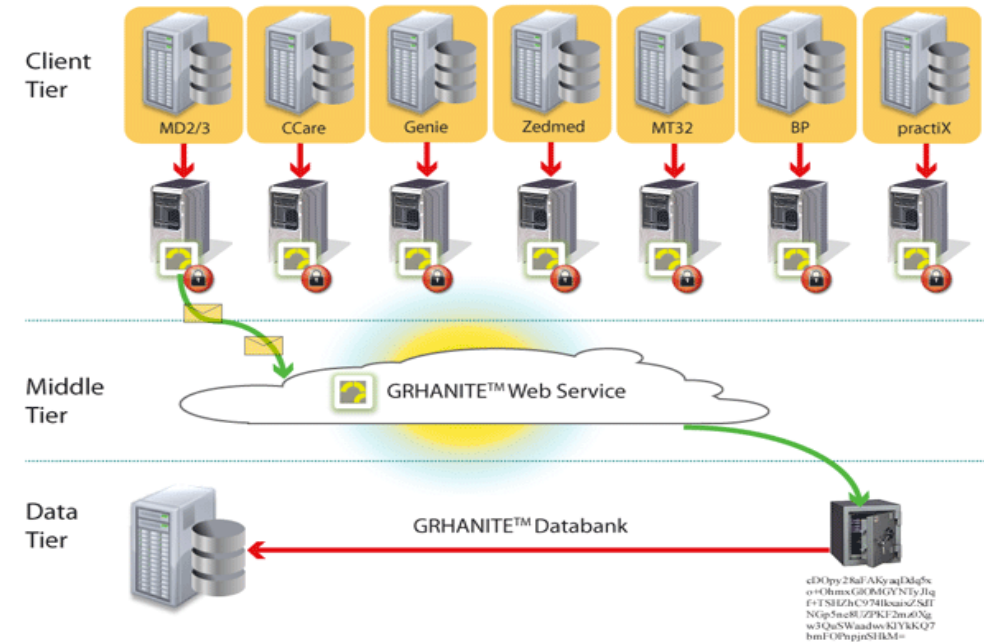
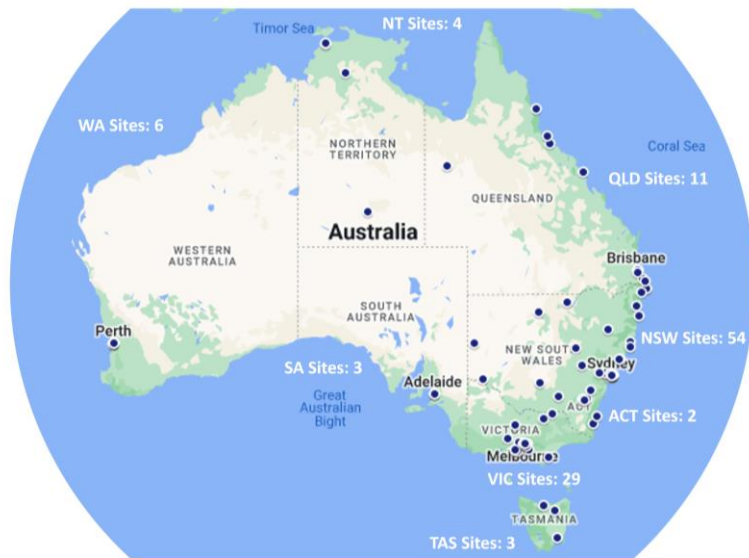
- Inform tailored intervention
- Ongoing monitoring and evaluation
- Patient level information





Australian Sentinel Surveillance of Blood Borne Viruses and Sexually Transmissible Infections

Monitor HIV, viral hepatitis and STIs in Priority populations



Strengths and limitations of ACCESS



Patient exposures (risk factors) linked to outcomes



Hundreds of EMR variables extracted



Longitudinally linked data from >120 sites representing millions of patient visits



Skill and capacity to process, manage and analyse the data



Limited and inconsistently recorded behavioural risk factor information



No access to important risk information from patient progress notes



Large volumes of data cannot be manually audited for risk factors



Relies on human effort and expertise to develop algorithms to identify risk groups



Can machine learning help?

EXPERT-DRIVEN ALGORITHMS

Define risk using proxy indicators

Relies on experts to produce all the possible solutions

Requires human effort to program all the rules

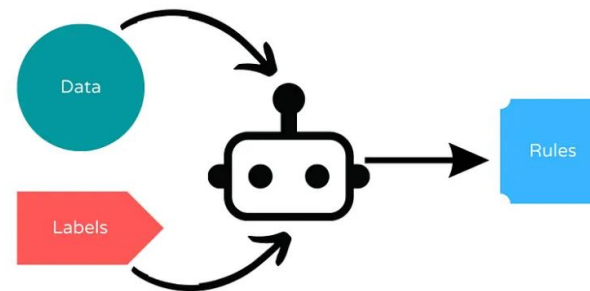


DATA-DRIVEN ALGORITHMS

Computer learns from the data

Recognises patterns and relationships between variables

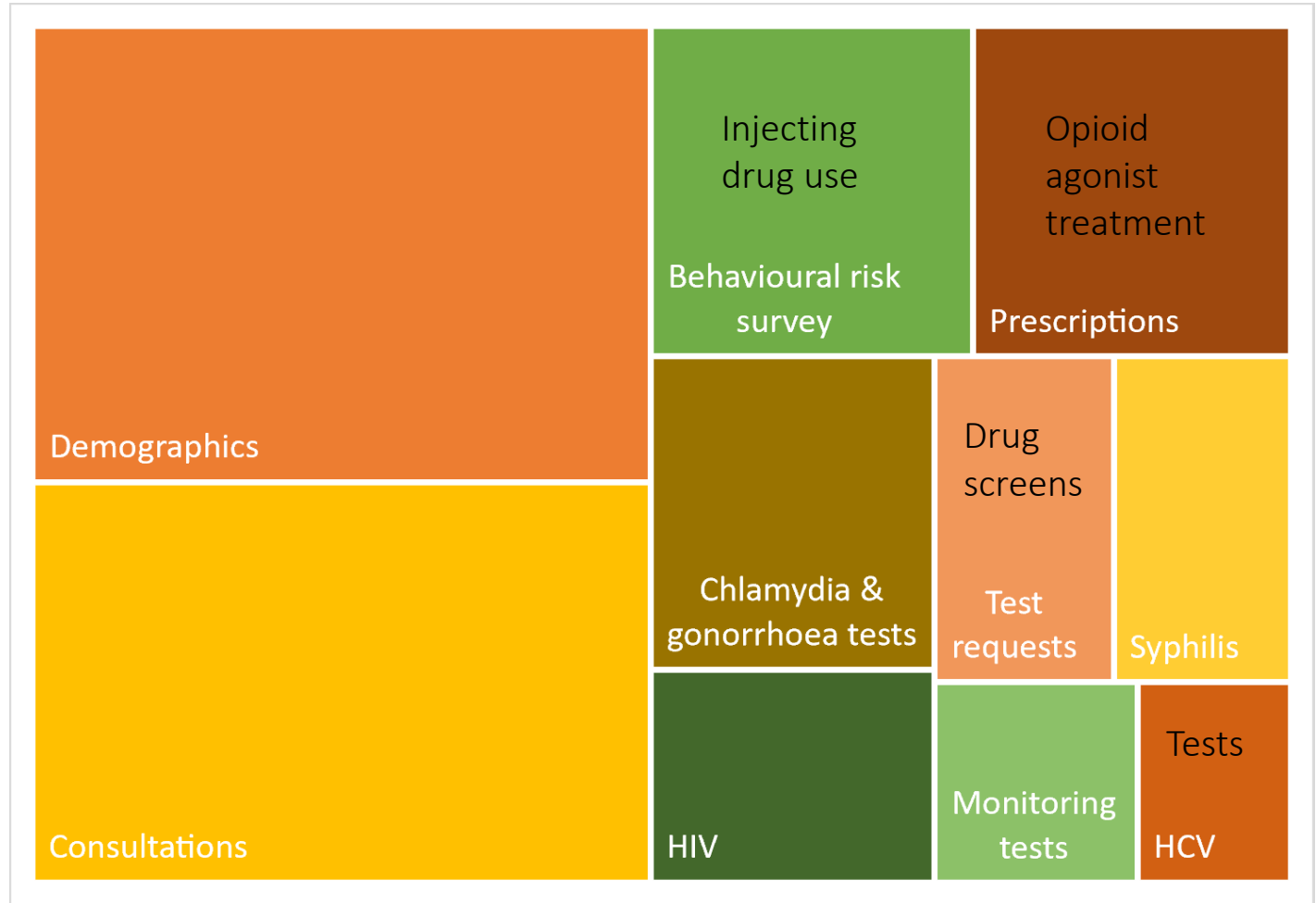
No need to program rules



Available ACCESS data and variables

Expert-driven method uses limited variables based on known predictors

Data-driven method uses all available variables plus expert knowledge

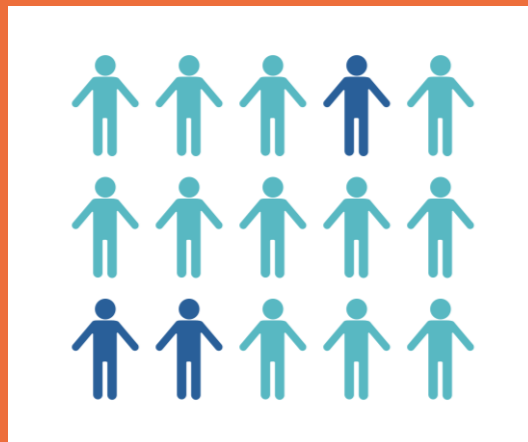




Objectives and method

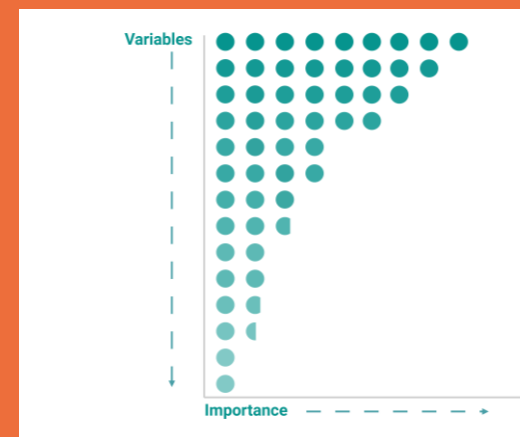
Develop a model to
classify people who
inject drugs

OBJECTIVE 1



Find out which variables
were important to the
classification

OBJECTIVE 2





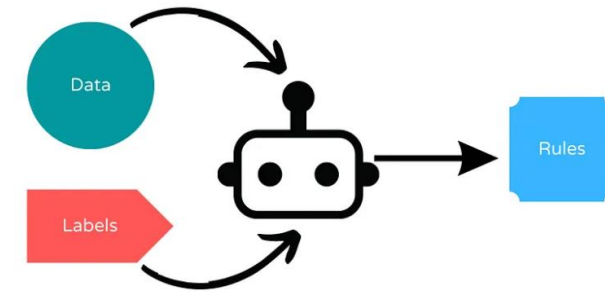
Training the machine learning model

SAMPLE OF LABELLED DATA FOR TRAINING AND TESTING

Labels

1 = People who inject drugs

0 = Random sample of patients



Data

88 features derived from variables in patient clinical records

	label	total_clinics	total_types	gp_clinic	ch_clinic	gbm_gp	sh_clinic	hosp_clinic	total_visits	person_time
0	0	3.0	2.0	0.0	0.0	1.0	1.0	0.0	2.0	0.038330
1	0	1.0	1.0	0.0	0.0	0.0	1.0	0.0	2.0	0.156057
2	1	1.0	1.0	1.0	0.0	0.0	0.0	0.0	145.0	13.382615
3	0	1.0	1.0	0.0	1.0	0.0	0.0	0.0	3.0	0.232717
4	1	7.0	4.0	0.0	1.0	1.0	1.0	1.0	74.0	12.123203
5	1	1.0	1.0	1.0	0.0	0.0	0.0	0.0	43.0	2.937714
6	1	4.0	3.0	1.0	1.0	0.0	1.0	0.0	259.0	13.259412
7	1	3.0	2.0	1.0	1.0	0.0	0.0	0.0	5.0	0.665298



Results – predictions and important features



Classification of people who inject drugs

n=1454	POSITIVE LABEL	NEGATIVE LABEL
POSITIVE PREDICTION	700	55
NEGATIVE PREDICTION	49	650

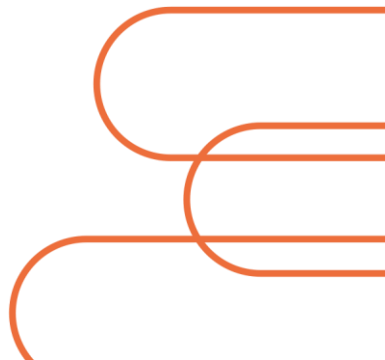
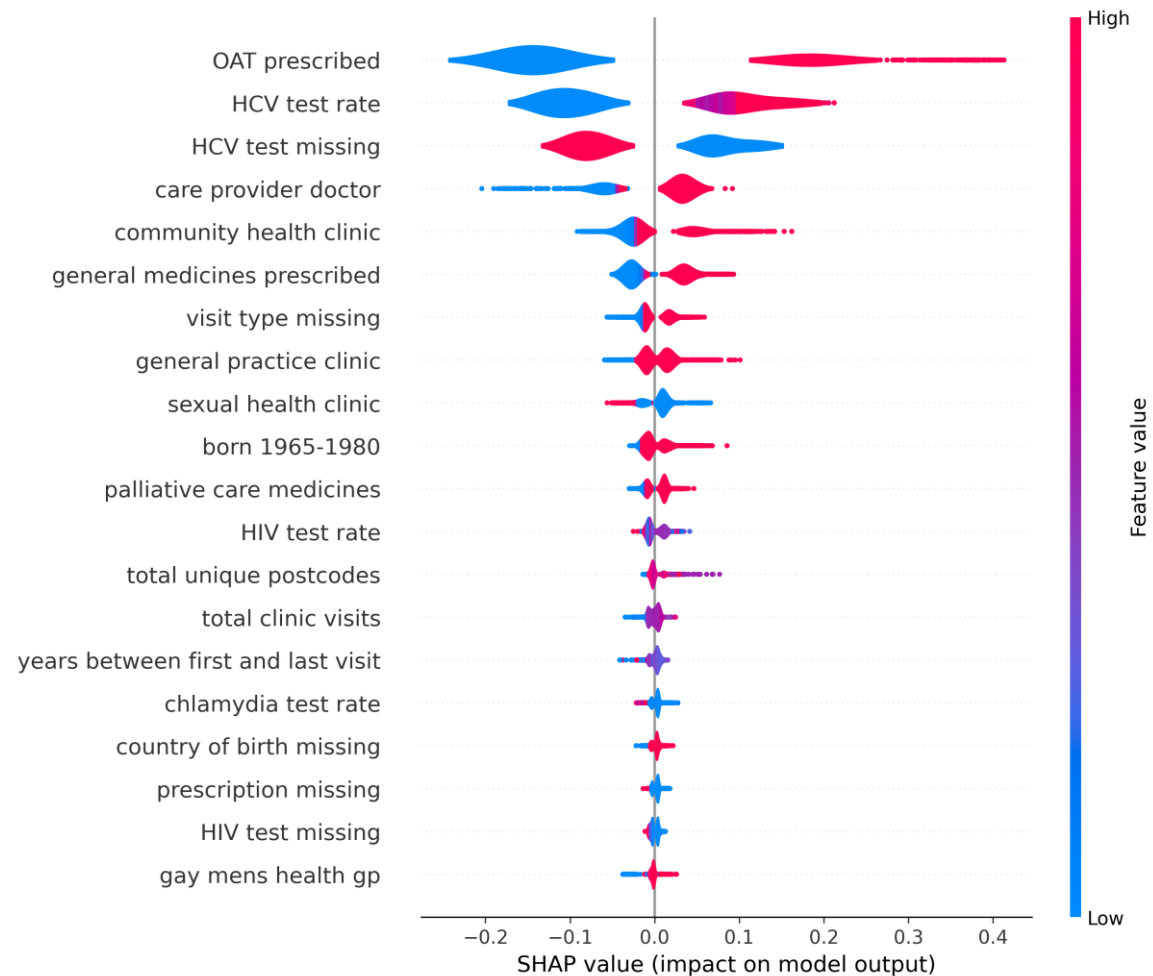
METRIC	DEFINITION	PERFORMANCE
Accuracy	Correct predictions	93%
Precision (PPV)	Positive predictions are truly positive	93%
Recall (Sensitivity)	Positive labels are predicted positive	93%



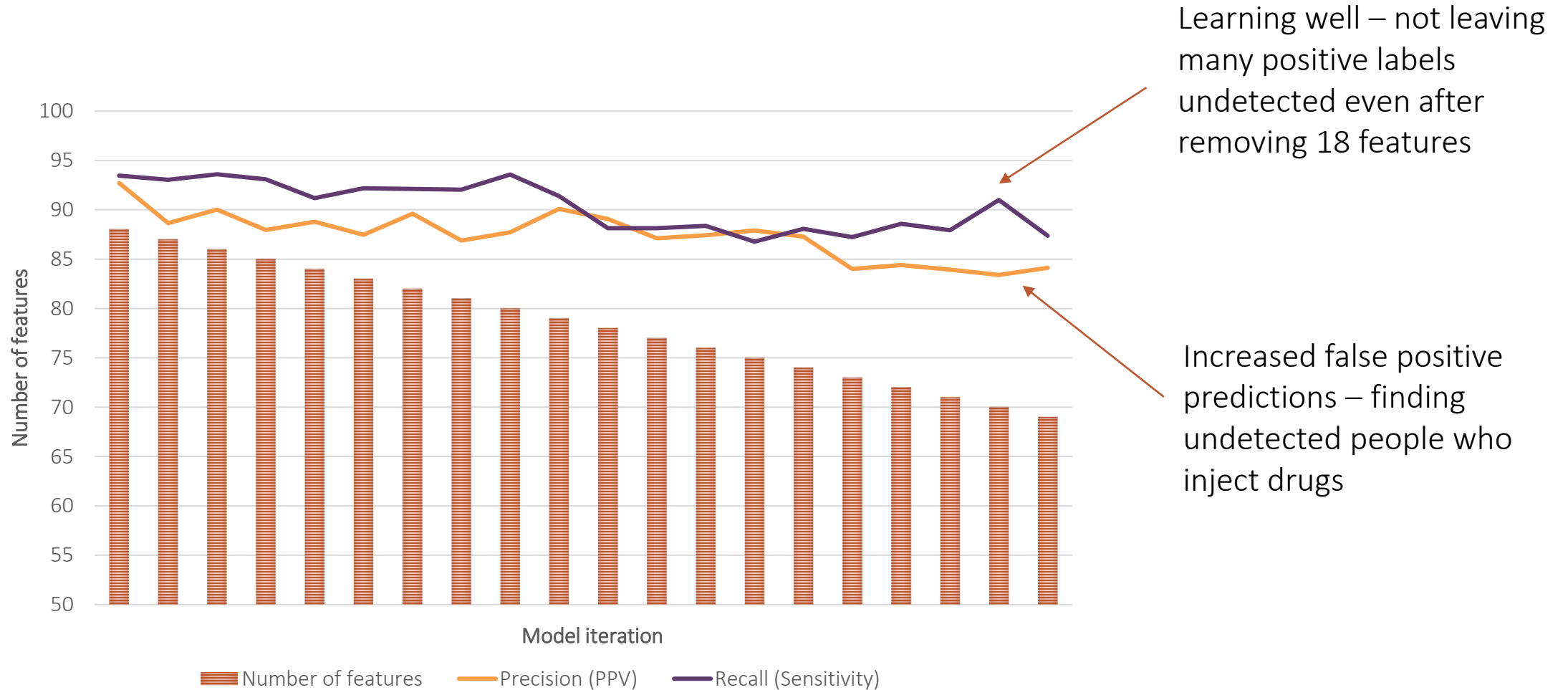
Contributions to model predictions

Top 20 contributing features

- First 5-10 have either a large positive or negative influence on the prediction
- Others work in combination to influence the model prediction



Model predictions by number of features





Summary



Implications for surveillance and future direction

- We built a highly predictive model
- Works when known predictors are unavailable
- Increased pool of candidate people who inject drugs
- We have a new way to classify risk groups
- Shows the suitability of machine learning for these tasks
- Machine learning has its limitations
- Needs to be evaluated on unseen data and real-world scenarios
- Algorithmic bias should also be assessed



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burnet.edu.au

carol.el-hayek@burnet.edu.au

