

Learning Classifier Systems in an Automated Machine Learning Framework

Ryan Urbanowicz, PhD

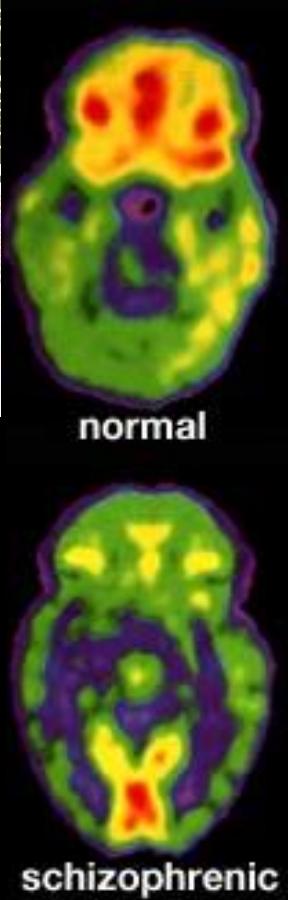
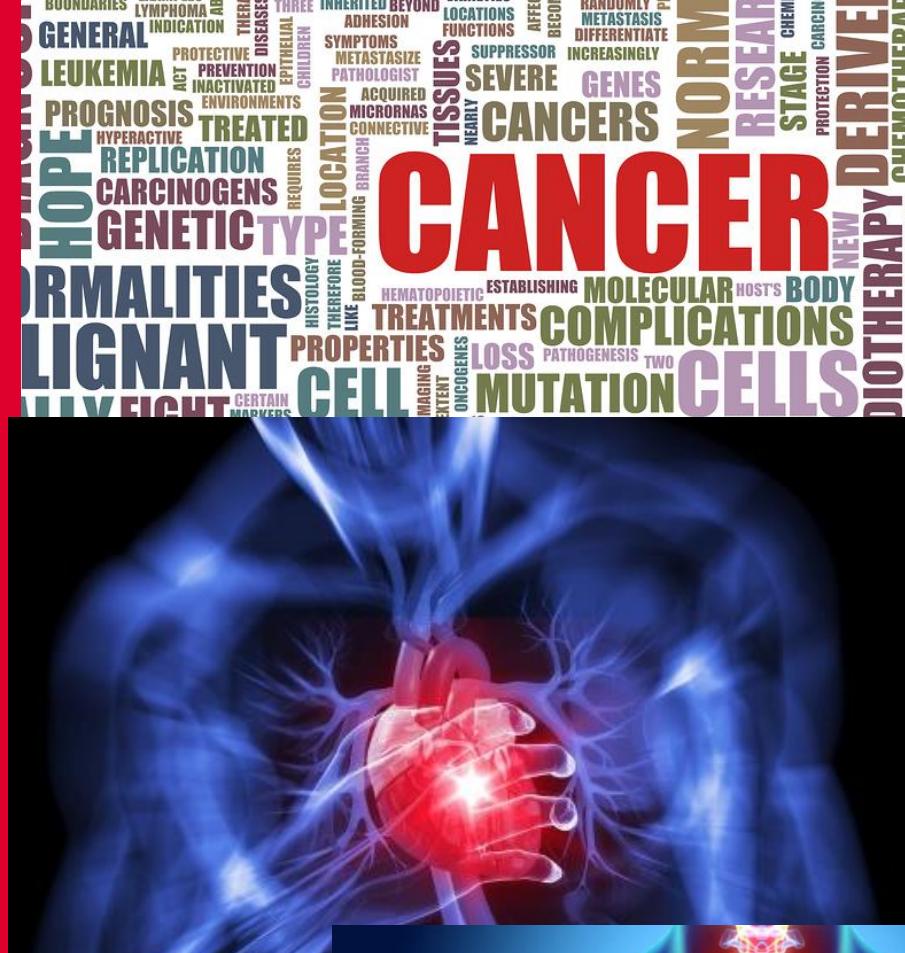
Department of Biostatistics, Epidemiology, and Informatics

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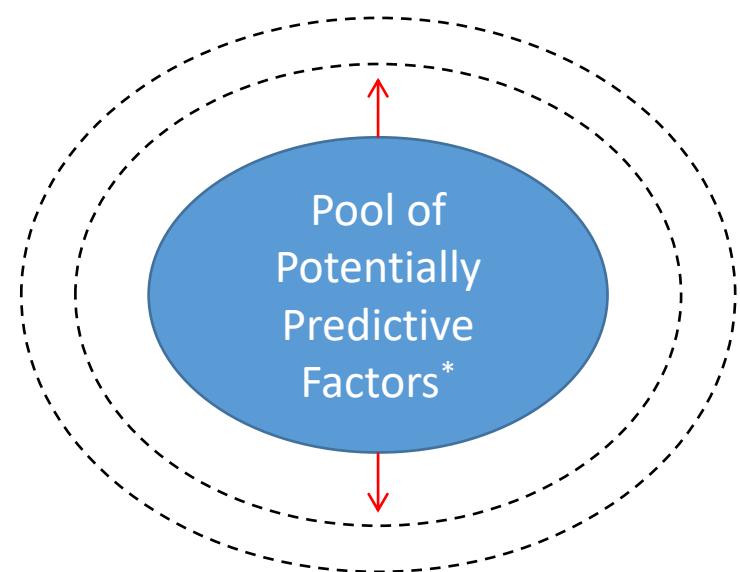
Twitter: @DocUrbs

Lab Website: www.med.upenn.edu/urbslab/

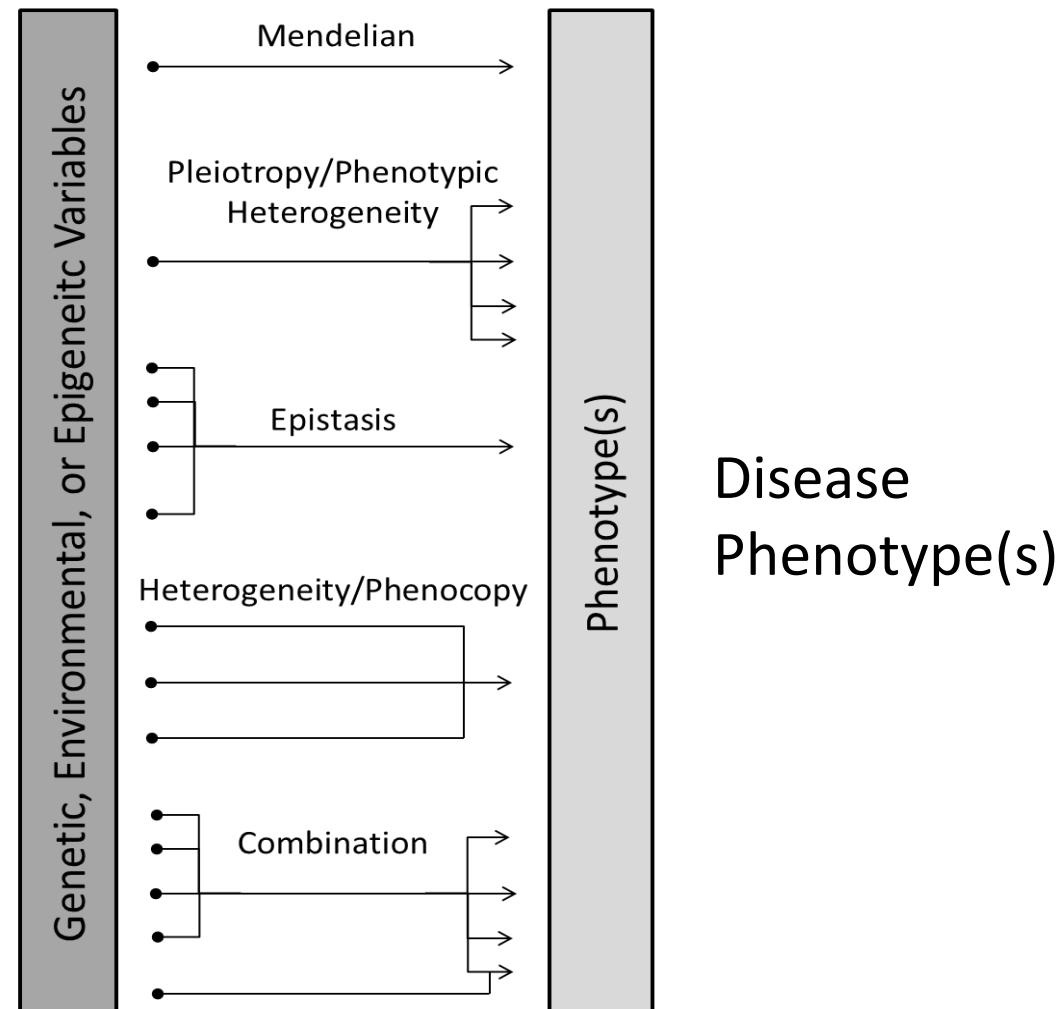
MAY 6, 2002



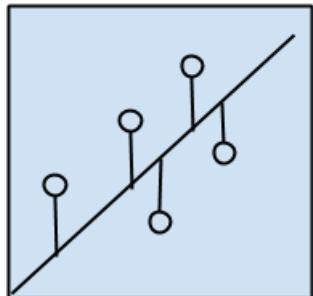
Challenge: Biomedical Data Analysis



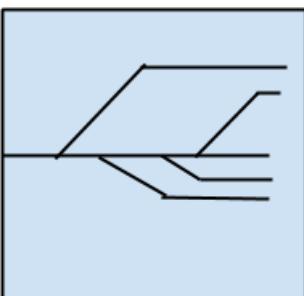
Do we have [all] the relevant features collected/available?



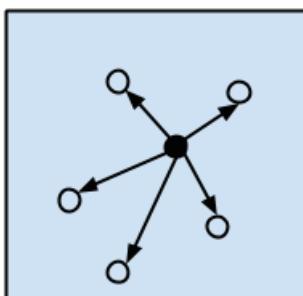
Machine Learning Algorithm Families



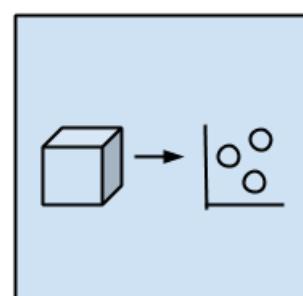
Regression Algorithms



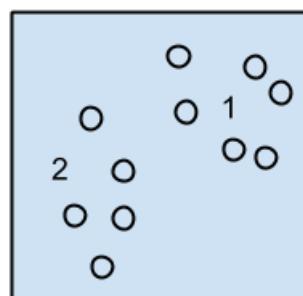
Regularization Algorithms



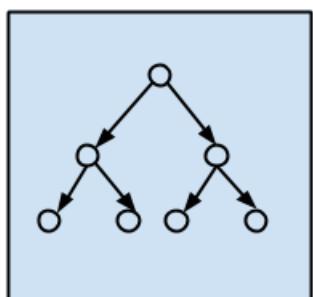
Instance-based Algorithms



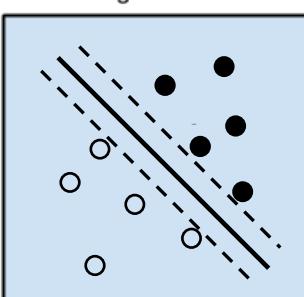
Dimensional Reduction Algorithms



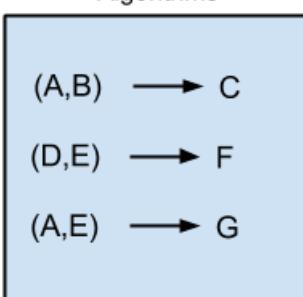
Clustering Algorithms



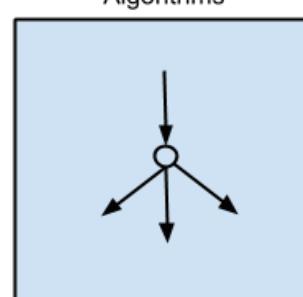
Decision Tree Algorithms



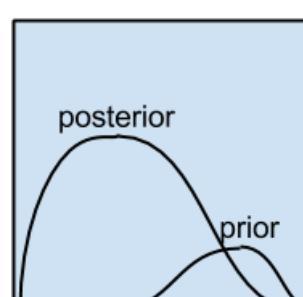
Support Vector Machines



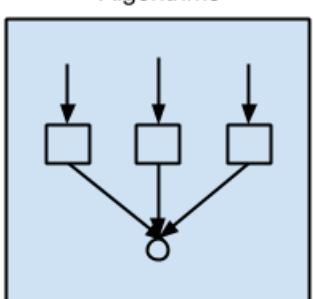
Association Rule Learning Algorithms



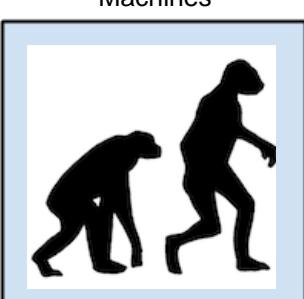
Artificial Neural Network Algorithms



Bayesian Algorithms

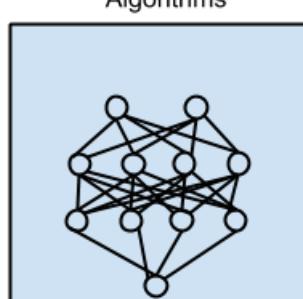


Ensemble Algorithms

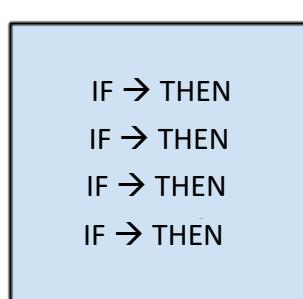


Evolutionary Algorithms

Non-exhaustive
list of ML families



Deep Learning Algorithms



Rule-Based Machine Learning

Learning Classifier Systems

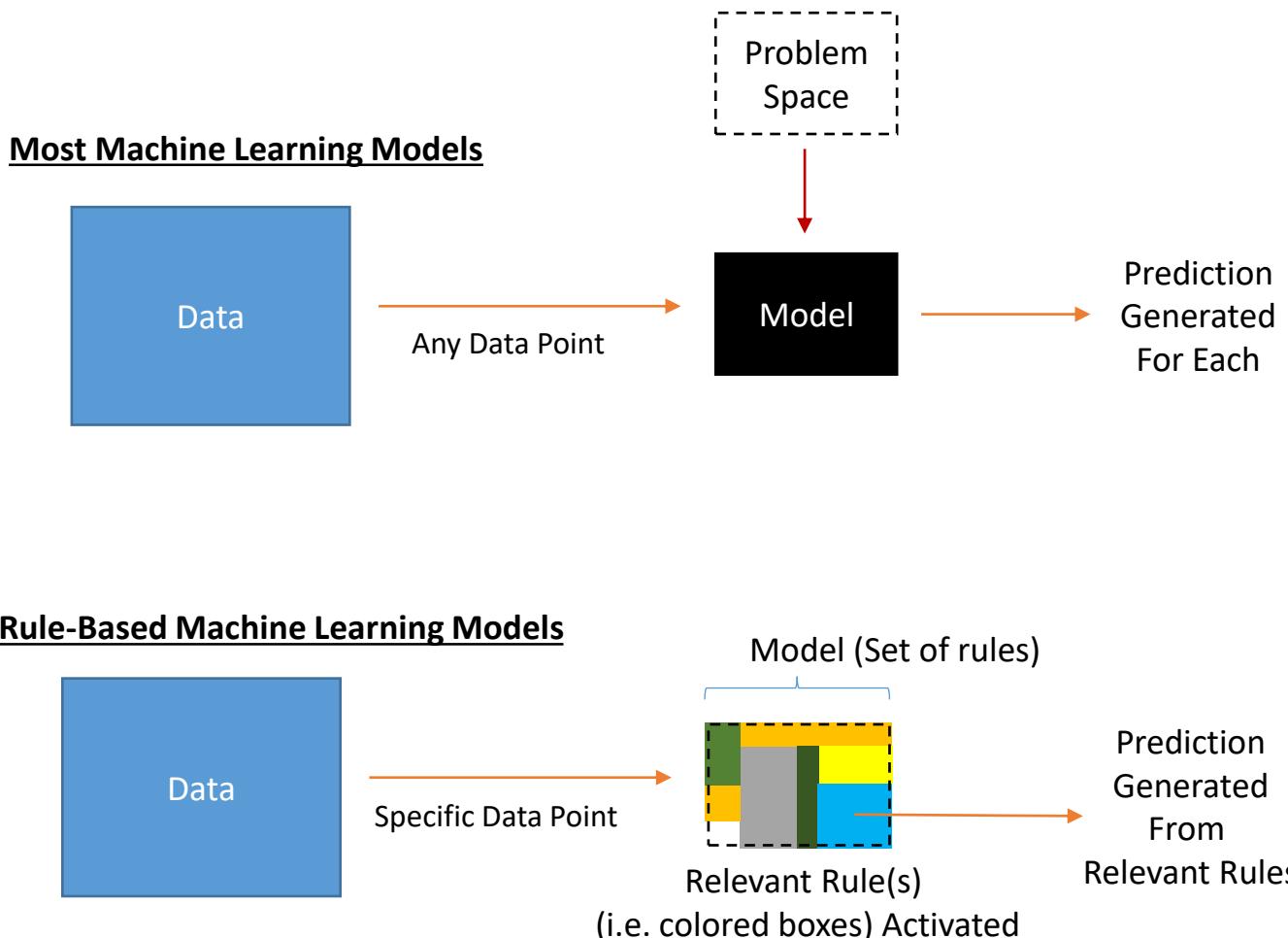
- LCS as an RBML
 - The intuition and **interpretability** of a rule-based system
 - Data-driven **improvement of performance** of ML
 - **Open ended discovery** of evolutionary algorithms
 - Focused primarily on classification tasks
- ‘Piece-wise’ Learning
 - Ideal for directly **modeling heterogeneous associations**
- A different modeling perspective than most ML algorithms
 - Part of a valuable ML toolkit



John H. Holland

- Set of IF:THEN rules
 - IF SNP_A = 0 AND BMI < 20 THEN Control
 - IF SNP_X = 1 THEN Case
 - IF 30 < BMI > 40 THEN Case
 - IF 20 < BMI > 30 THEN Control
 - IF 20 < BMI > 30 AND SNP_Y = 2 THEN Case
 - ...

Challenging the Paradigm of Modeling



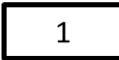
Rule Representation

Quaternary Rule Representation

Rule Condition



Classification



Mixed Discrete-Continuous Attribute-List Knowledge Representation

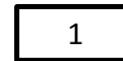
Attribute Reference



Rule Condition



Classification



KEY: Continuous Discrete

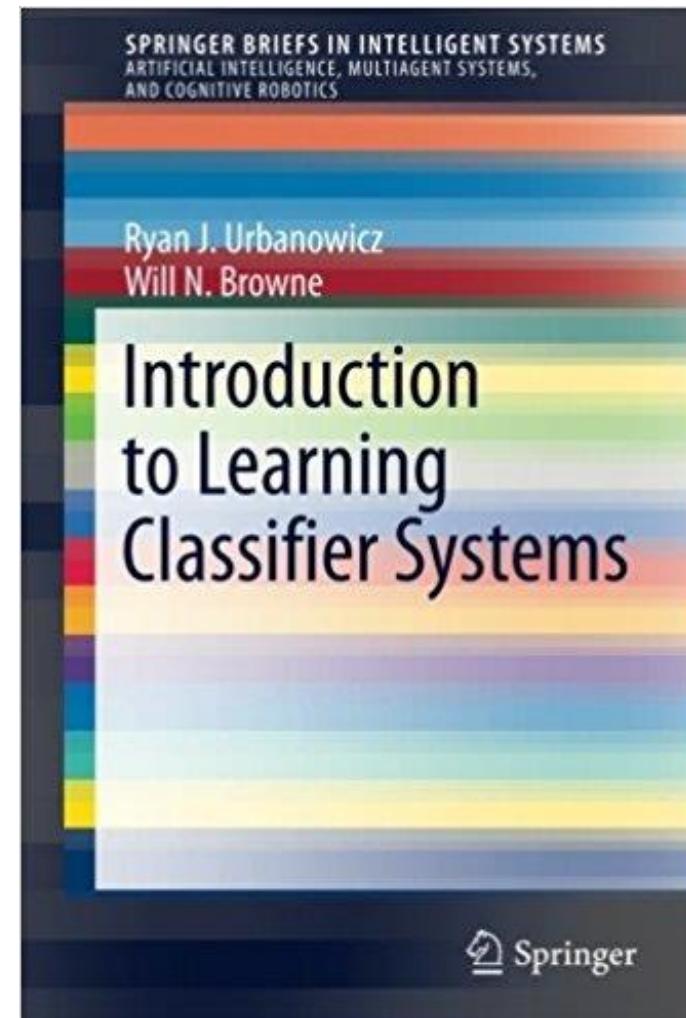
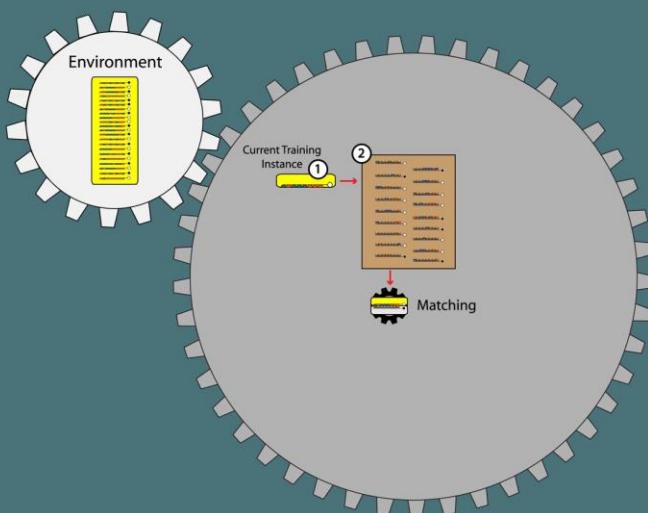
Very similar to ALKR(Attribute List Knowledge Representation): [Bacardit et al. 09]

Urbanowicz et. al 2015 – ExSTraCS 2.0: Description and Evaluation of a Scalable LCS

Learning Classifier Systems - Resources

Review Paper: Urbanowicz & Moore 2009 –
*Learning Classifier Systems: A complete
introduction, review, and roadmap*

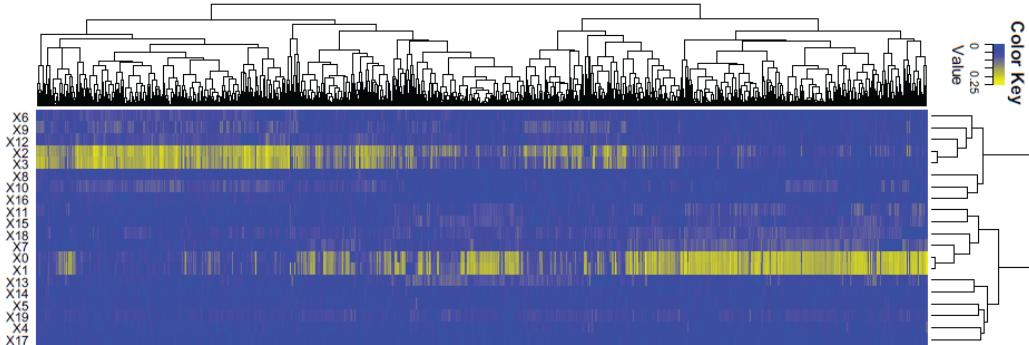
Learning Classifier Systems in a Nutshell



eLCS

Improving Performance and Interpretability

- Attribute Tracking/Feedback
 - Characterize heterogeneity
 - Speed up learning with useful building blocks.
- Expert Knowledge Guided Rule Discovery
 - EK probabilistically weighting mutation, crossover, and covering.
- Rule Compaction

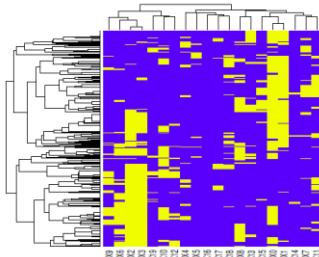


Using Expert Knowledge to Guide Covering and Mutation in a Michigan Style Learning Classifier System to Detect Epistasis and Heterogeneity

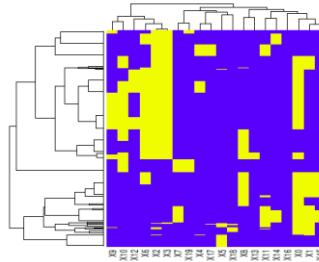
Ryan J. Urbanowicz, Delaney Granizo-Mackenzie, and Jason H. Moore

Computational Genetics Laboratory, Department of Genetics
Dartmouth Medical School, Lebanon, NH, USA
{ryan.j.urbanowicz,jason.h.moore}@dartmouth.edu
<http://www.epistasis.org/>

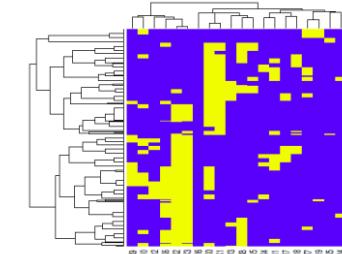
NONE



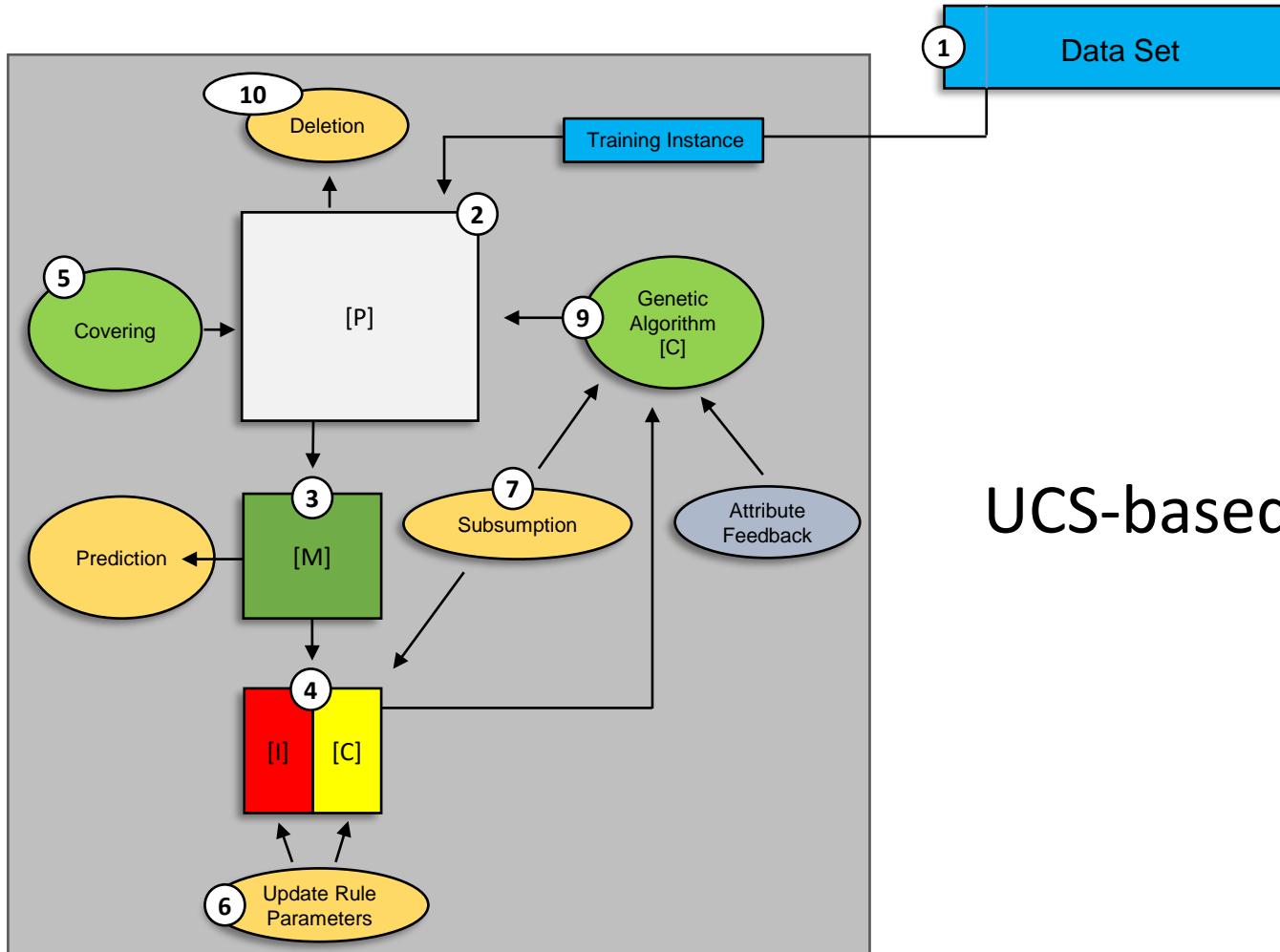
Fu2



PDRC

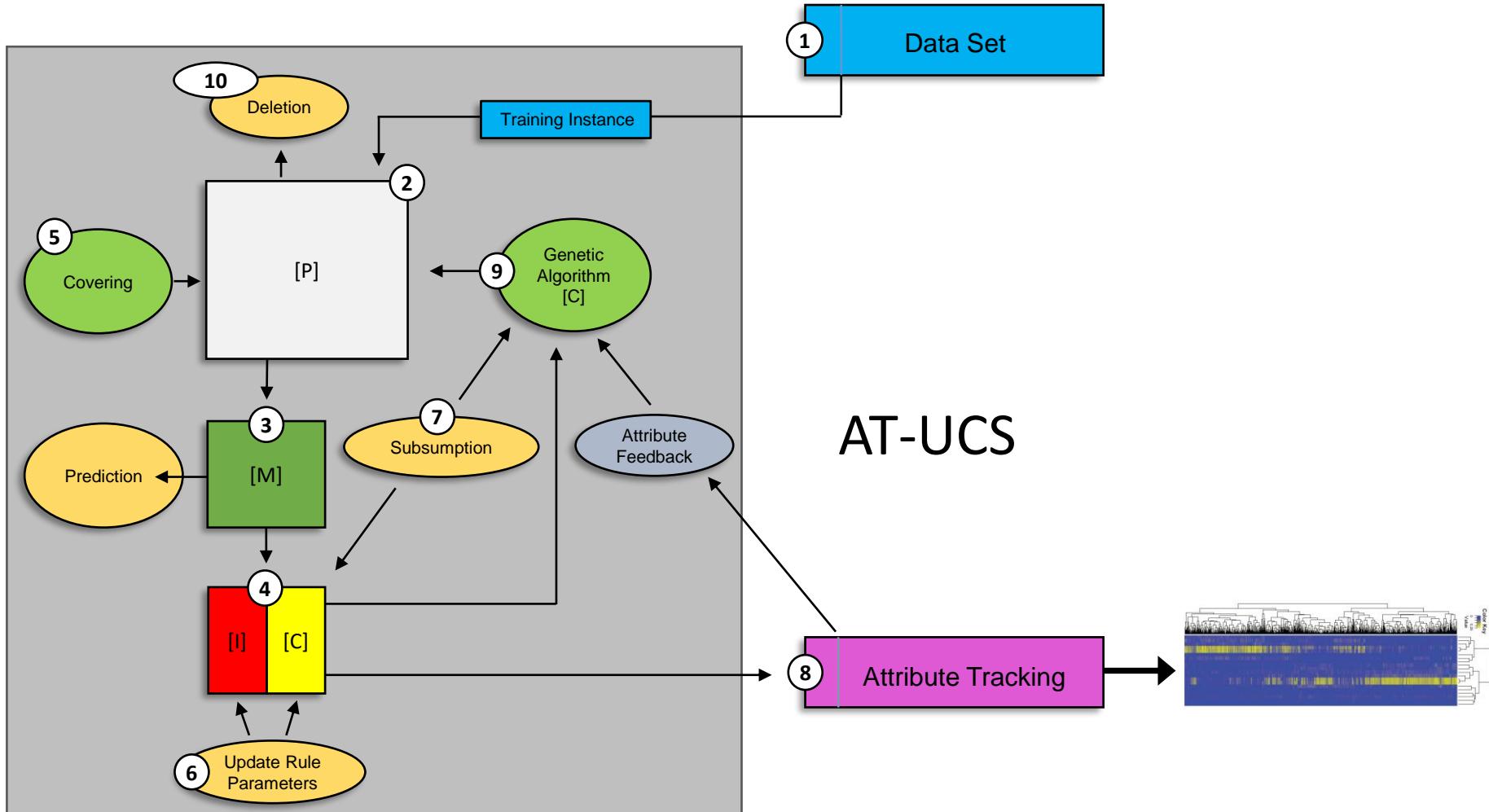


Stepping back: combining the heuristics

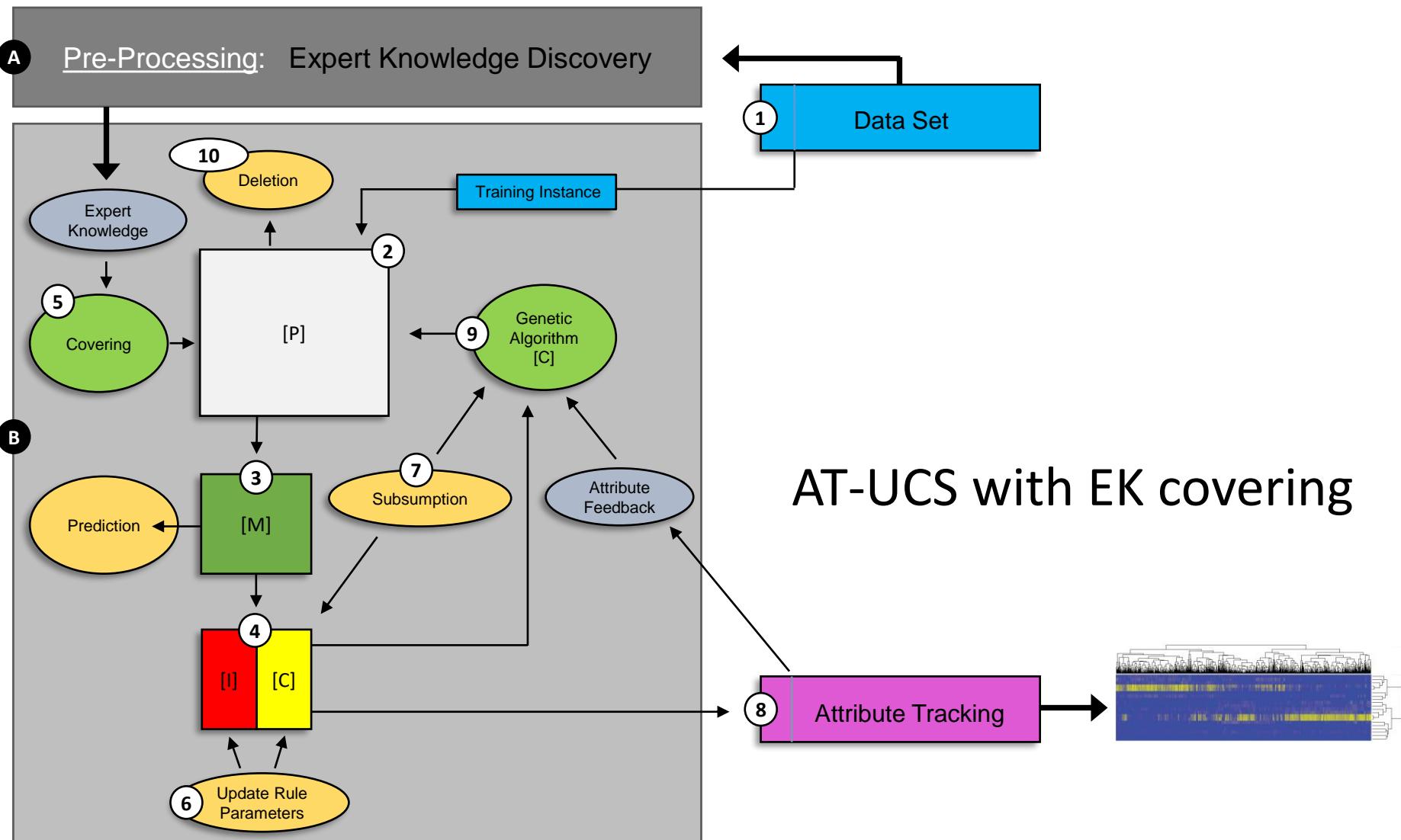


UCS-based

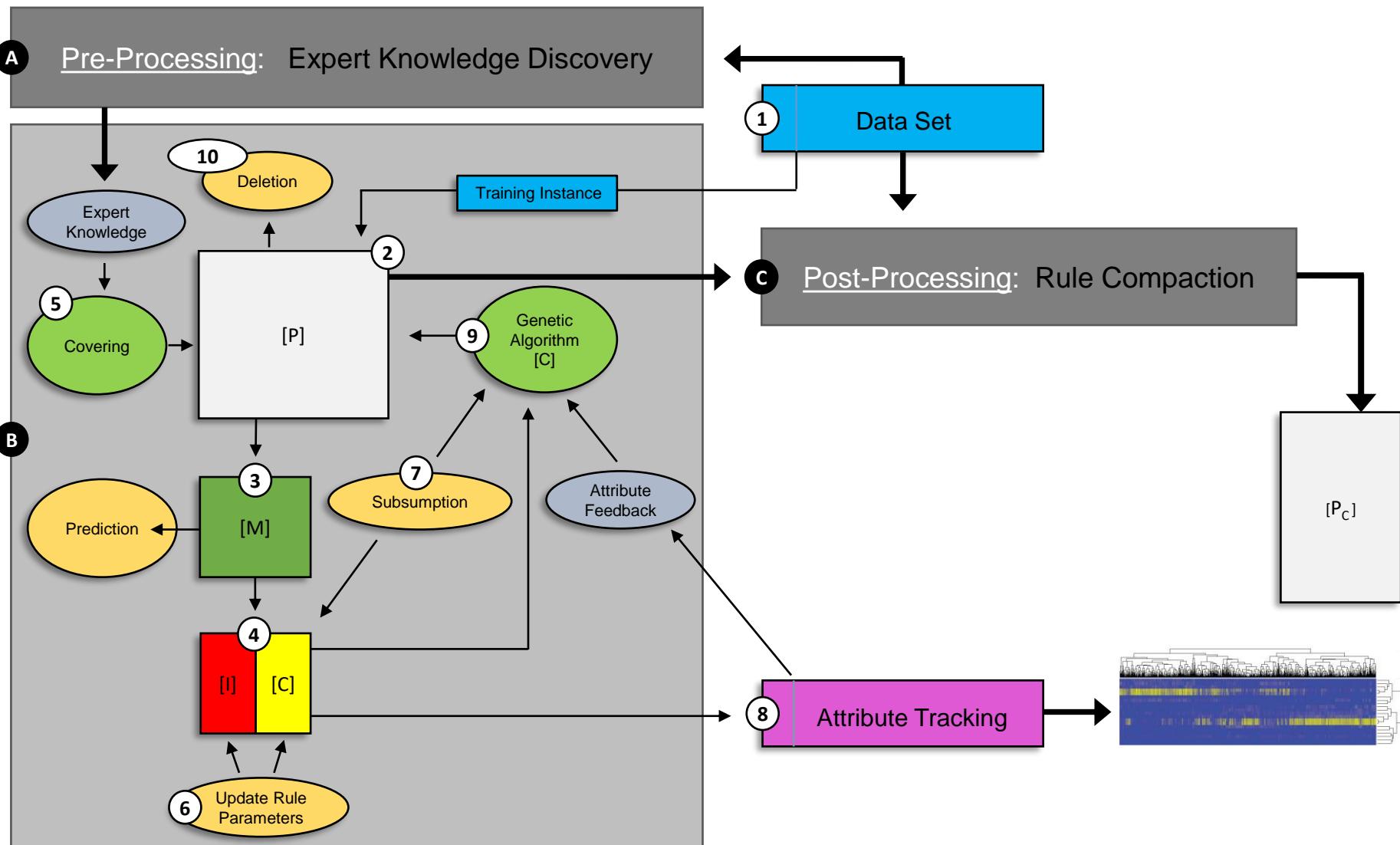
Stepping back: combining the heuristics



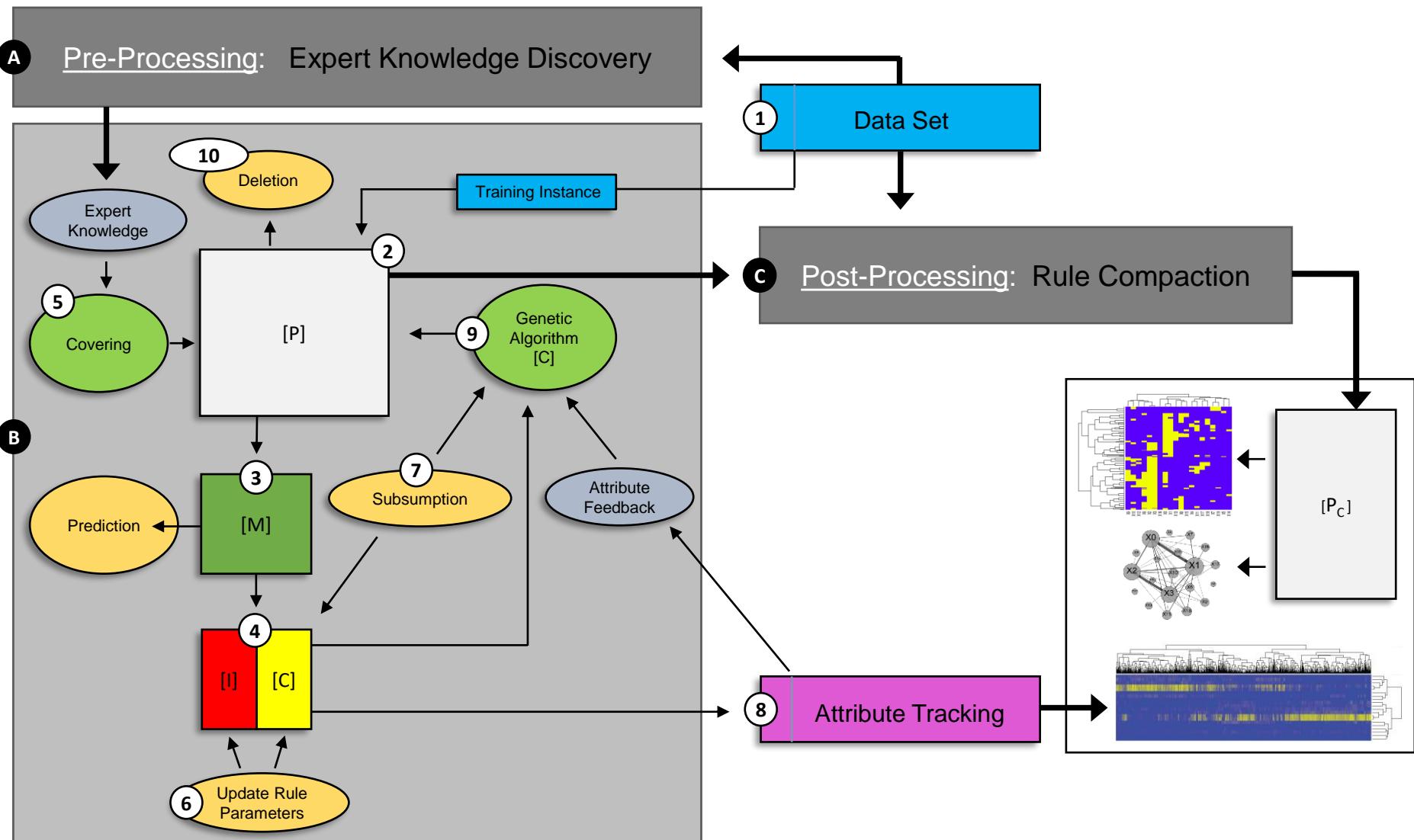
Stepping back: combining the heuristics

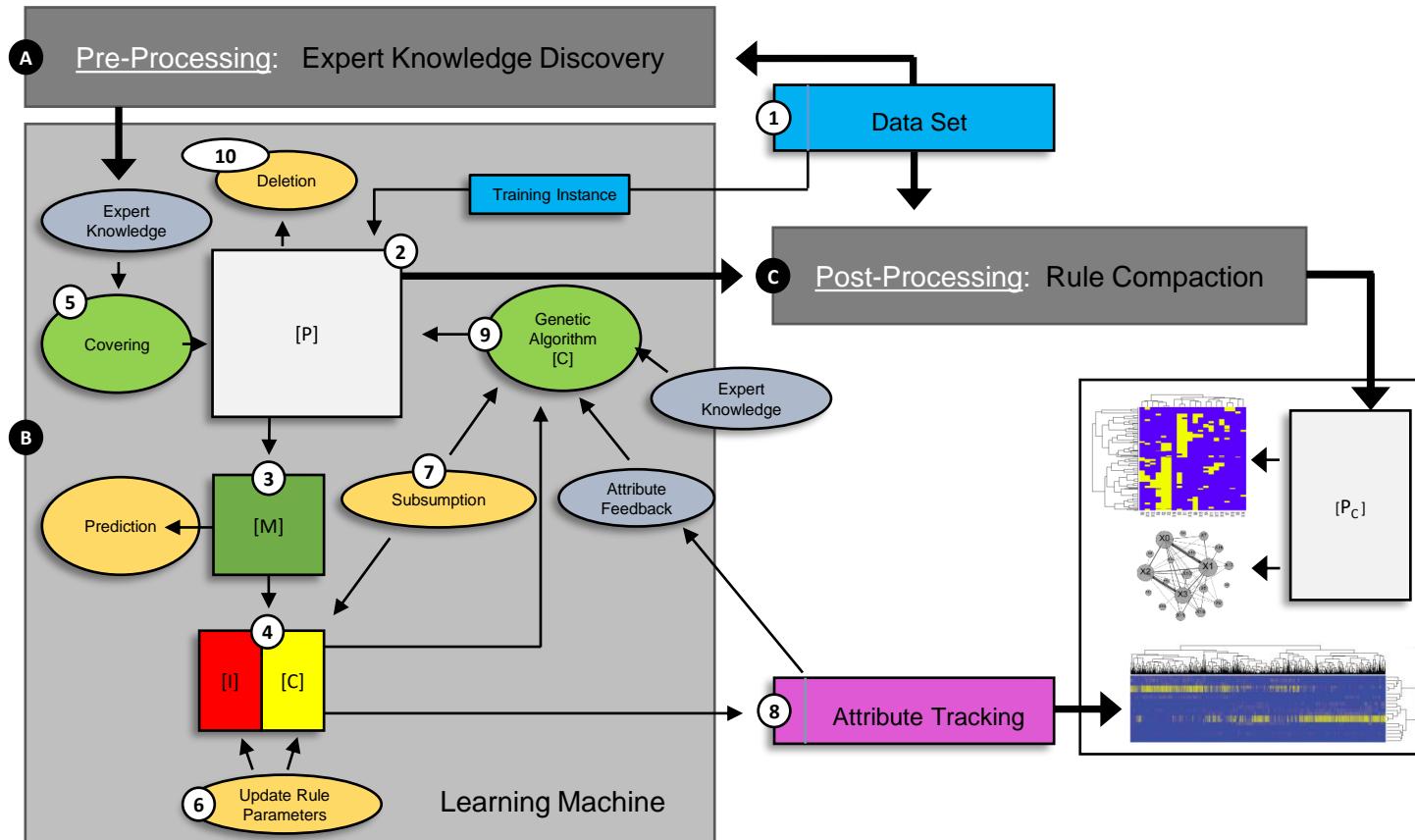


Stepping back: combining the heuristics



Stepping back: combining the heuristics

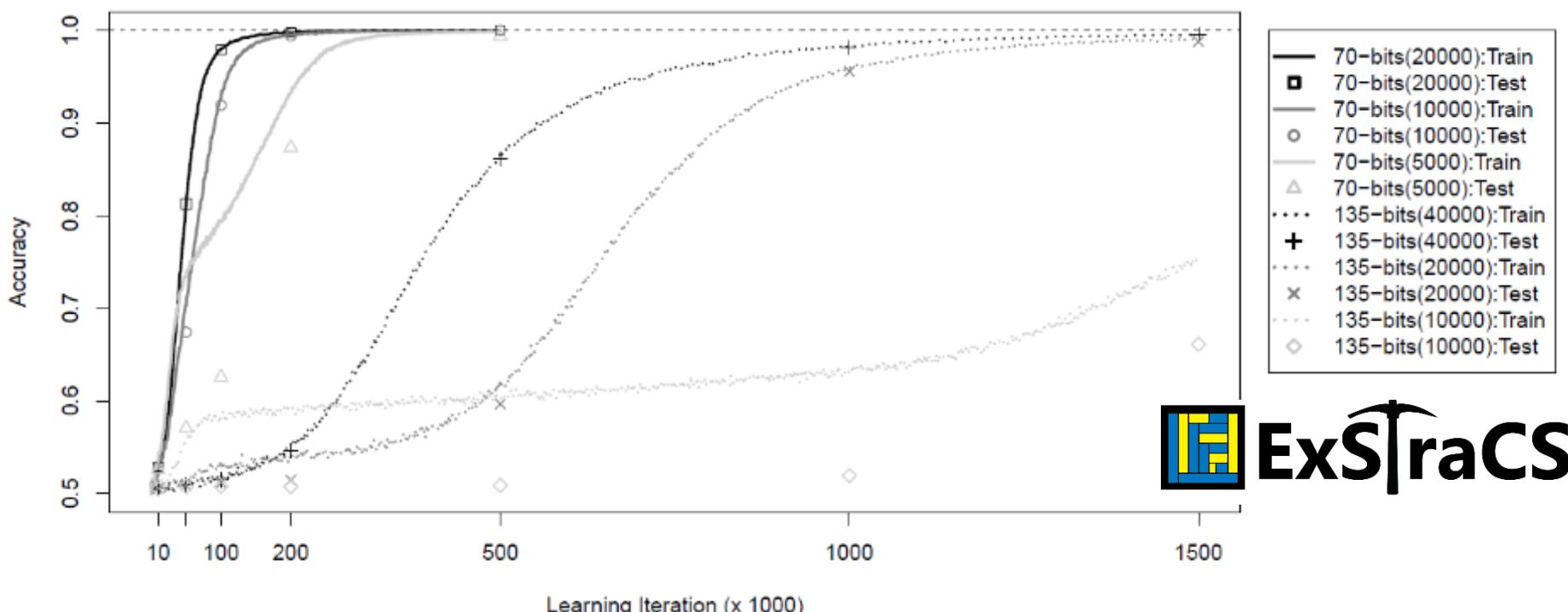




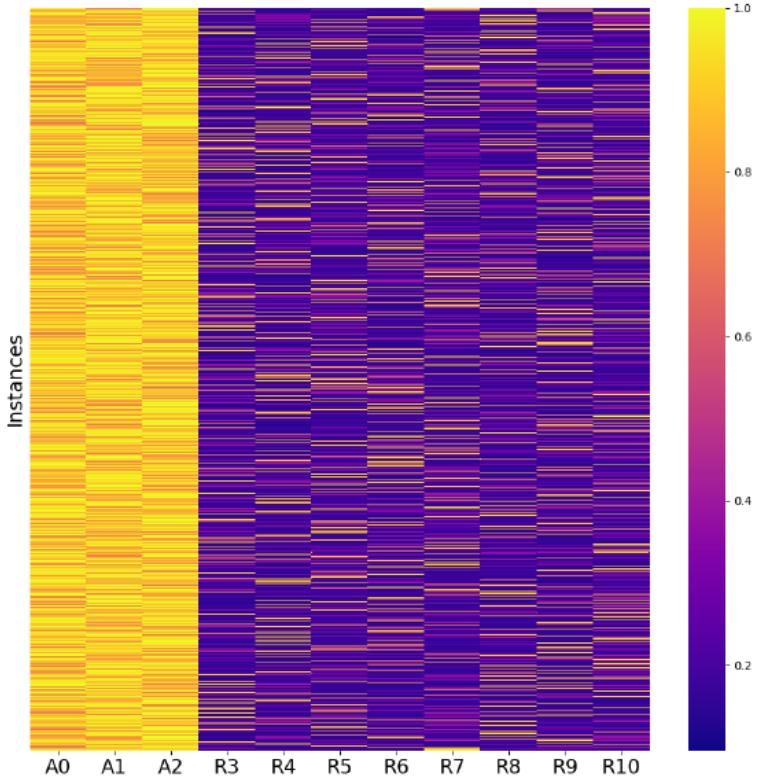
Urbanowicz, Ryan J., and Jason H. Moore. "ExSTraCS 2.0: description and evaluation of a scalable learning classifier system." *Evolutionary intelligence* 8.2-3 (2015): 89-116.

Directly solving the 135-bit Multiplexer

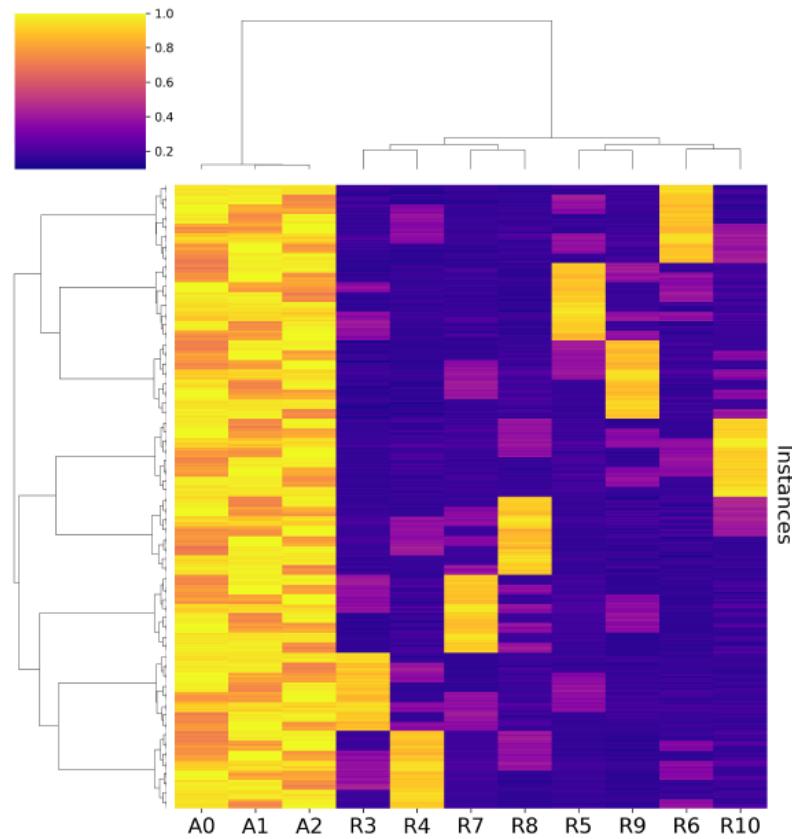
- TO SOLVE: 135-bit Multiplexer
 - Find 128 heterogeneous 8-way interactions
 - All 135 features are predictive in at least some subset of the dataset.
 - 256 optimal rules (in a traditional XCS-based algorithm)



LCS-DIVE: AT Clustering (11-bit MUX)

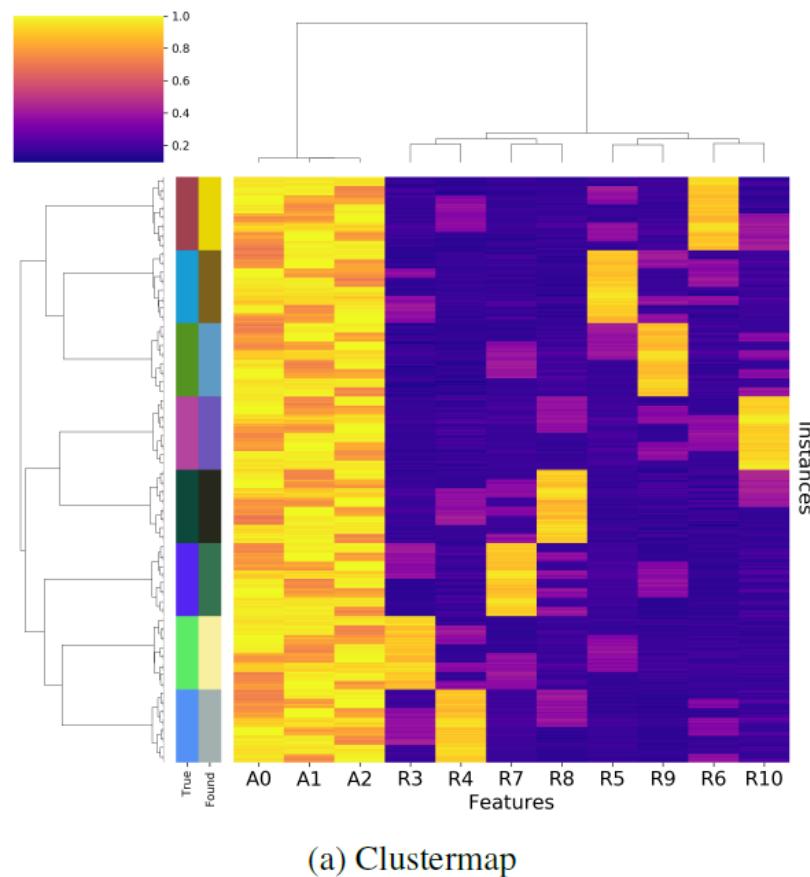
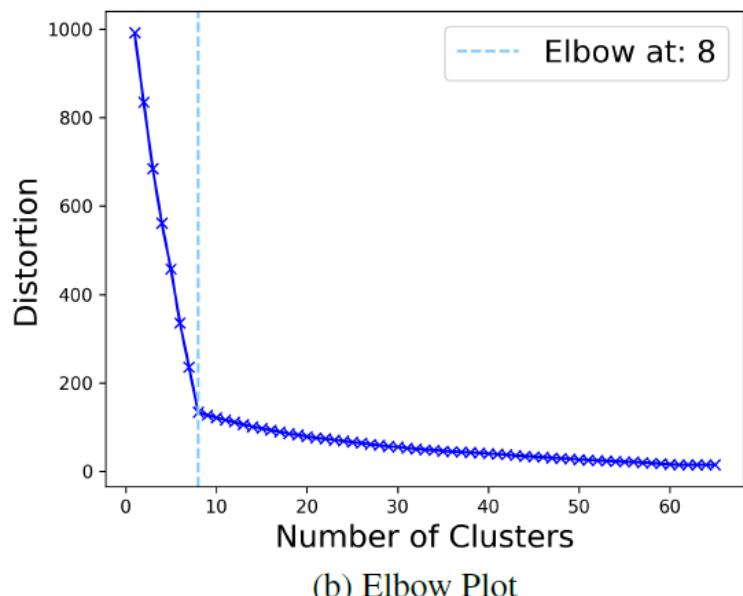


(a) Unclustered Heatmap

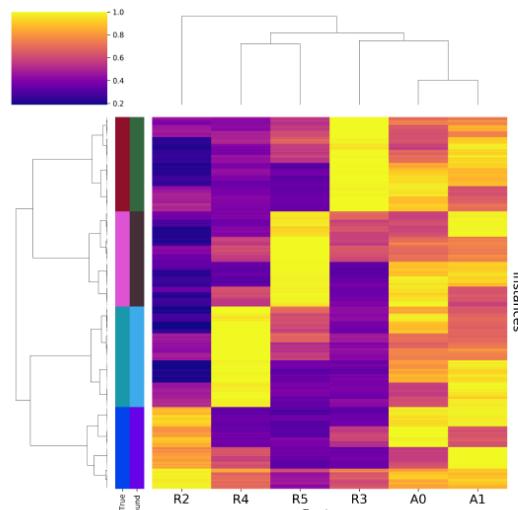


(b) Clustered Heatmap

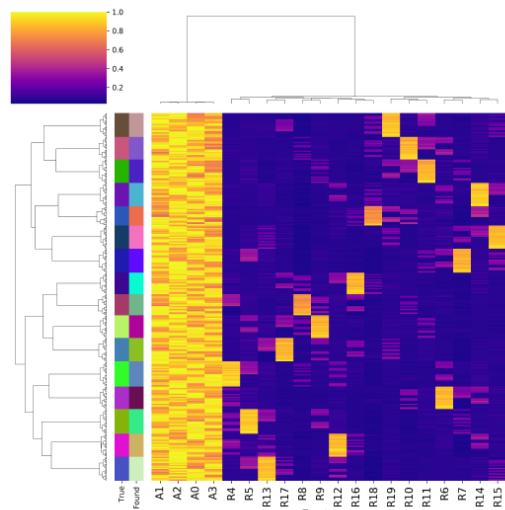
LCS-DIVE: AT Clustering (11-bit MUX)



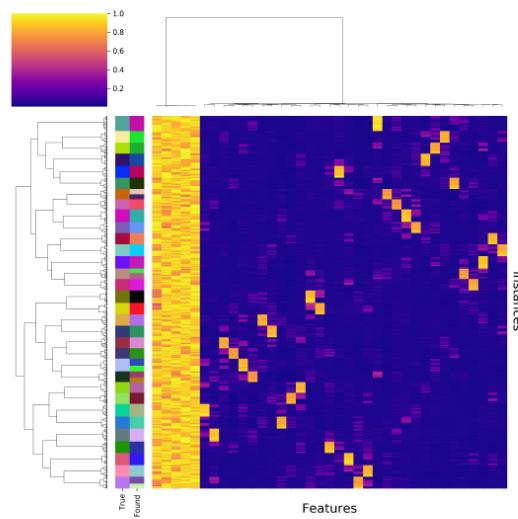
LCS-DIVE: MUX Problems



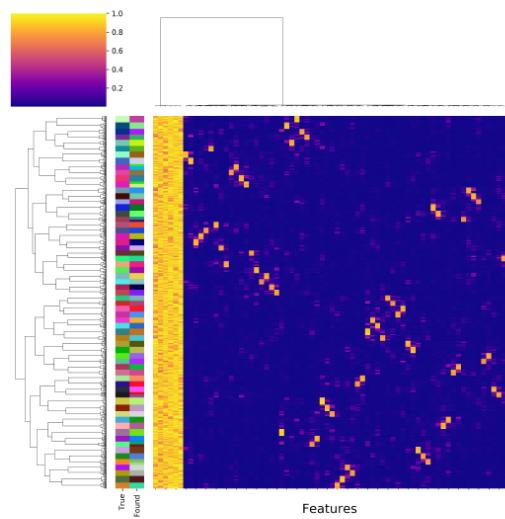
(a) 6-bit MUX



(b) 20-bit MUX



(c) 37-bit MUX

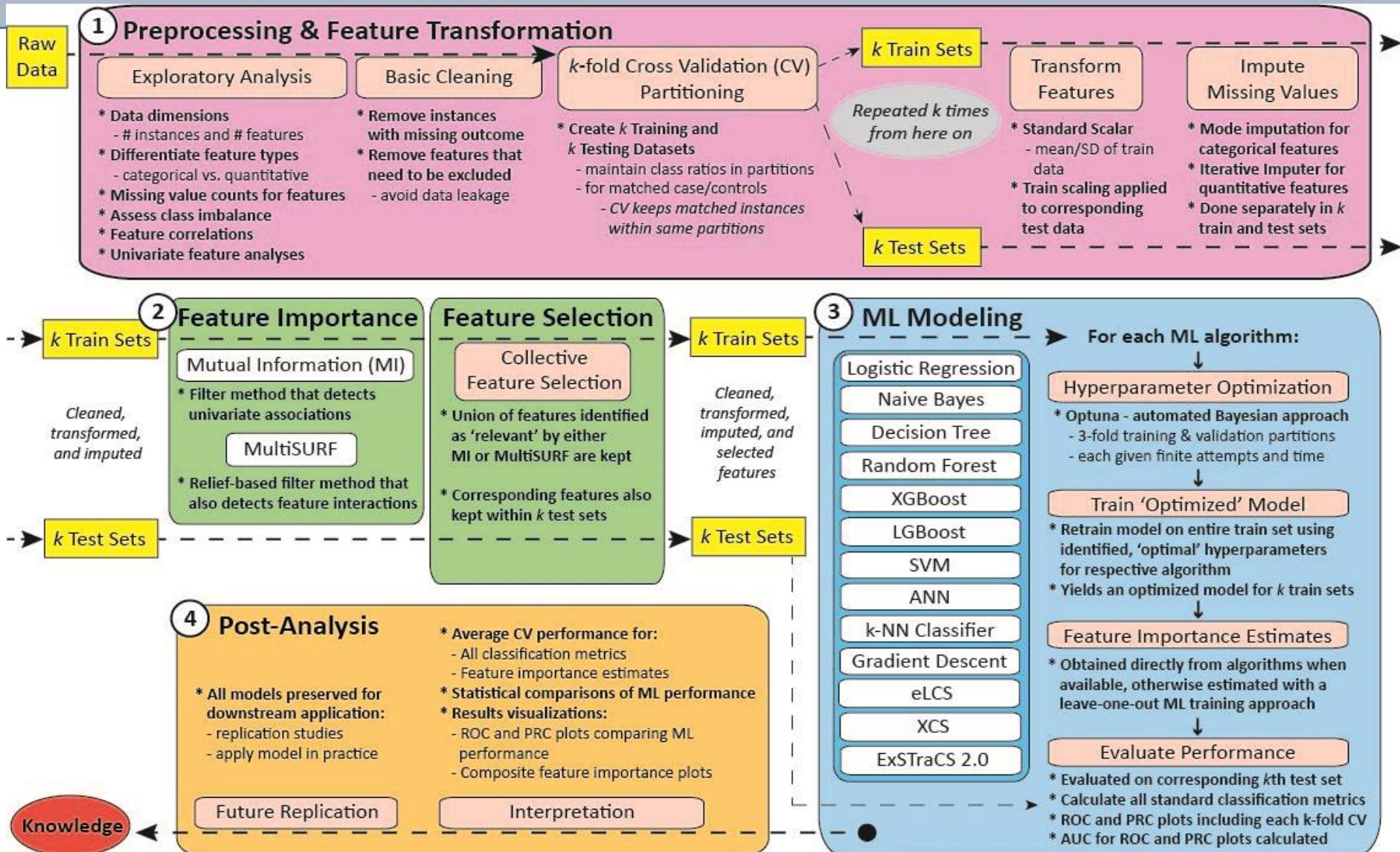


(d) 70-bit MUX

Scikit-Learn LCS Implementations

- Scikit-learn – is a very popular Python ML algorithm library
- Scikit-eLCS
 - <https://github.com/UrbsLab/scikit-eLCS>
- Scikit-XCS
 - <https://github.com/UrbsLab/scikit-XCS>
- Scikit-ExSTraCS
 - <https://github.com/UrbsLab/scikit-ExSTraCS>

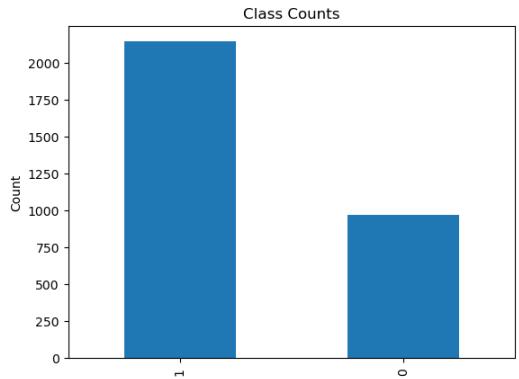
Automated ML Analysis Pipeline



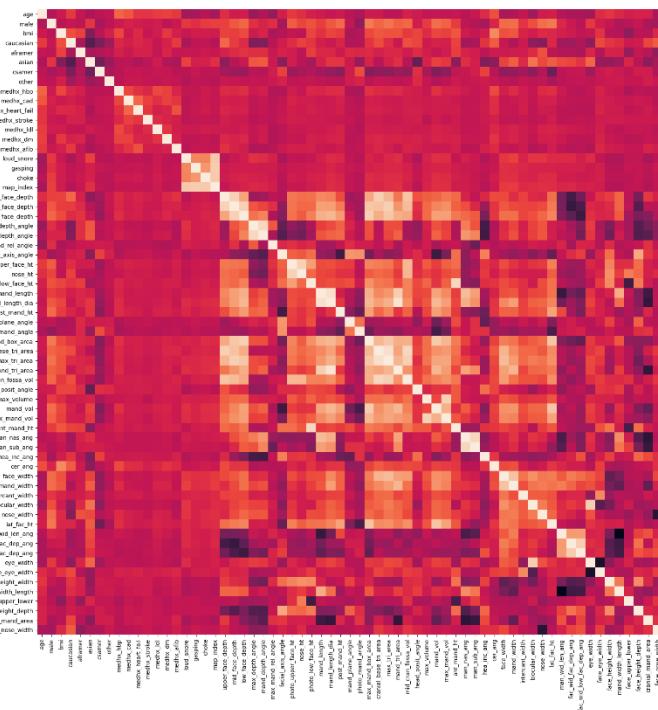
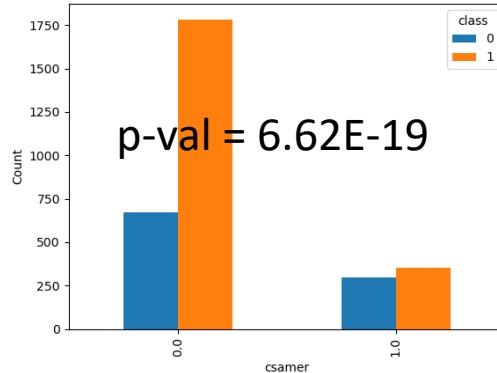
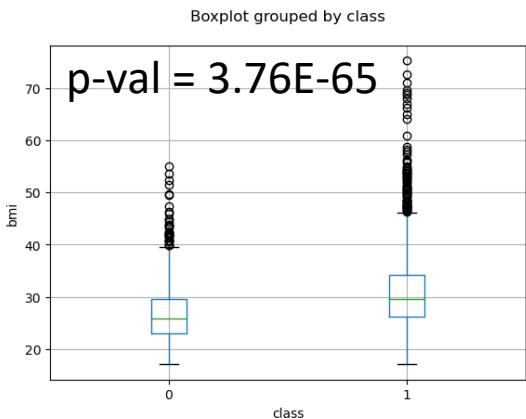
Preprint: Urbanowicz et al. A Rigorous Machine Learning Analysis Pipeline for Biomedical Binary Classification: Application in Pancreatic Cancer Nested Case-control Studies with Implications for Bias Assessments

AutoMLPipe-BC: Exploratory Analysis

AHI>5: DEM+DX+SYM+CF



Variable	Count
instances	3111
features	65
categorical_features	16
quantitative_features	49



Categorical → Chi Square
Quantitative → MannWhitney U-Test

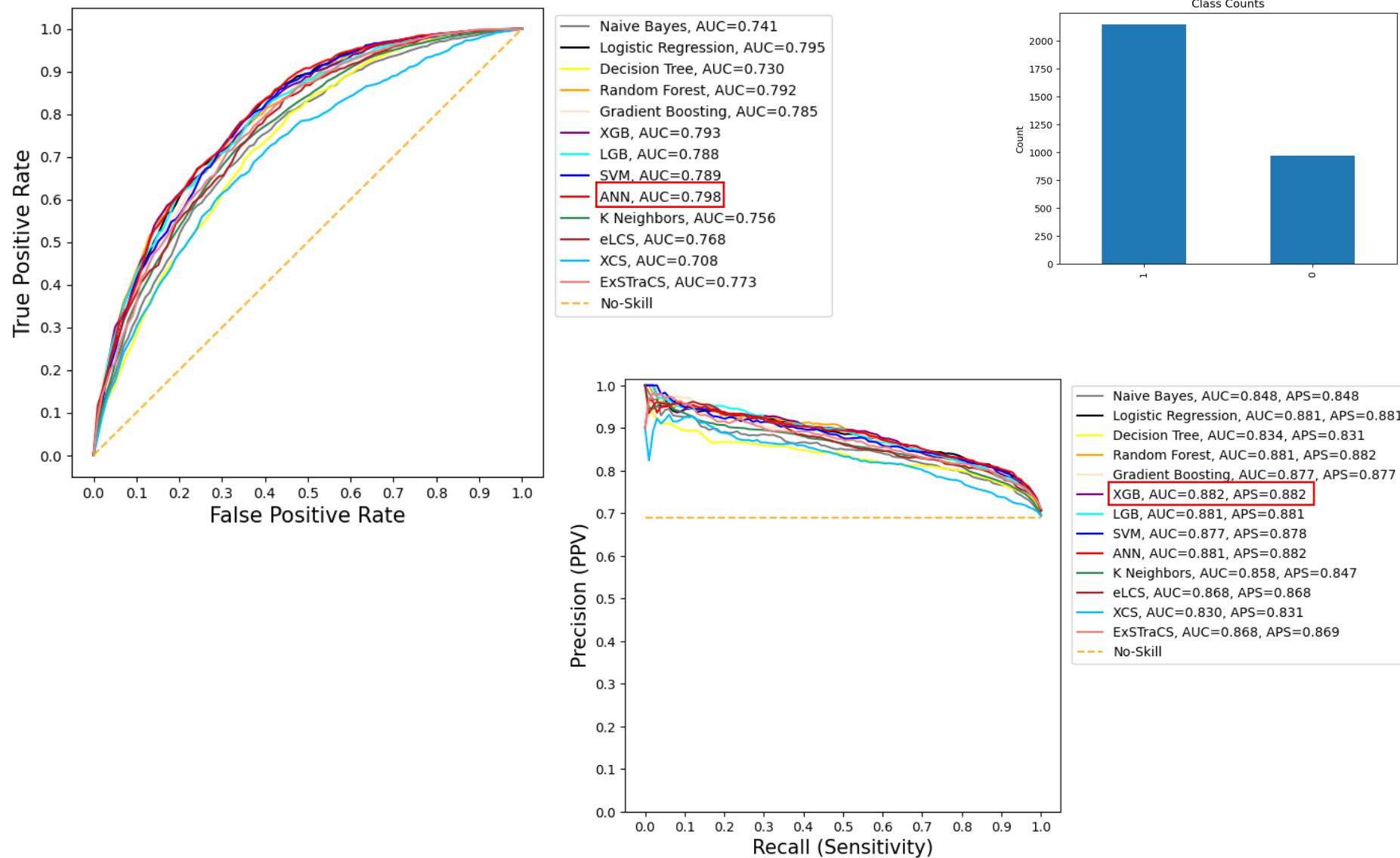
Modeling: Classification Evaluation Metrics

Average Model Prediction Statistics: D7 = SAGIC_GE5_DEMDXSYMCF_2021-04-28

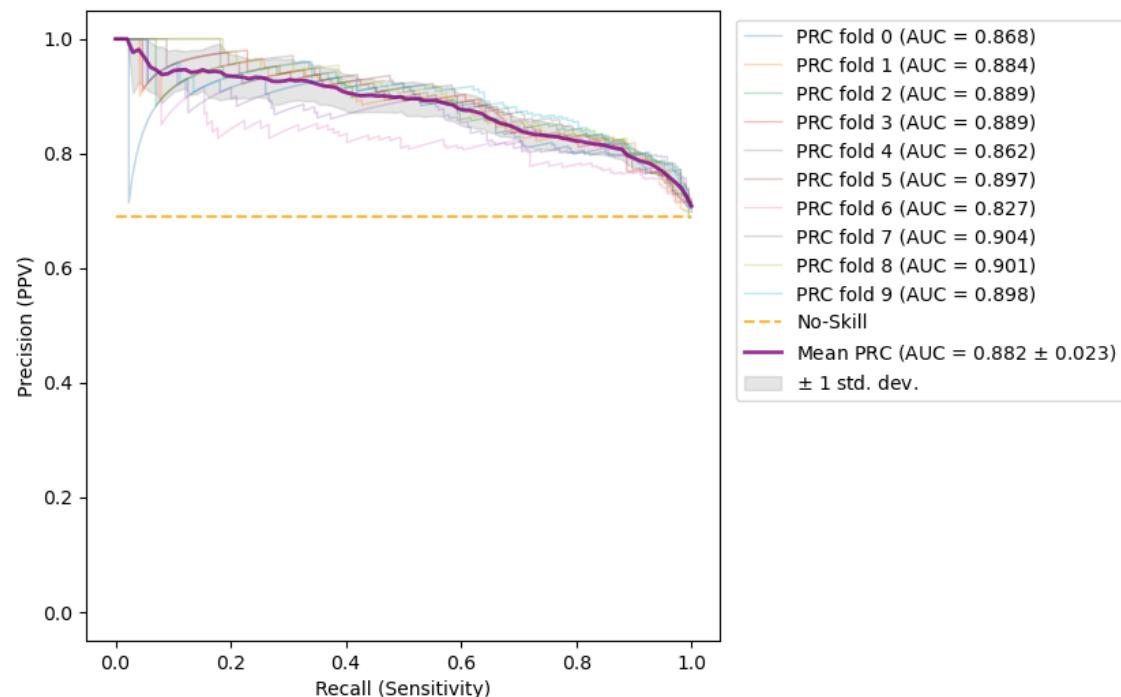
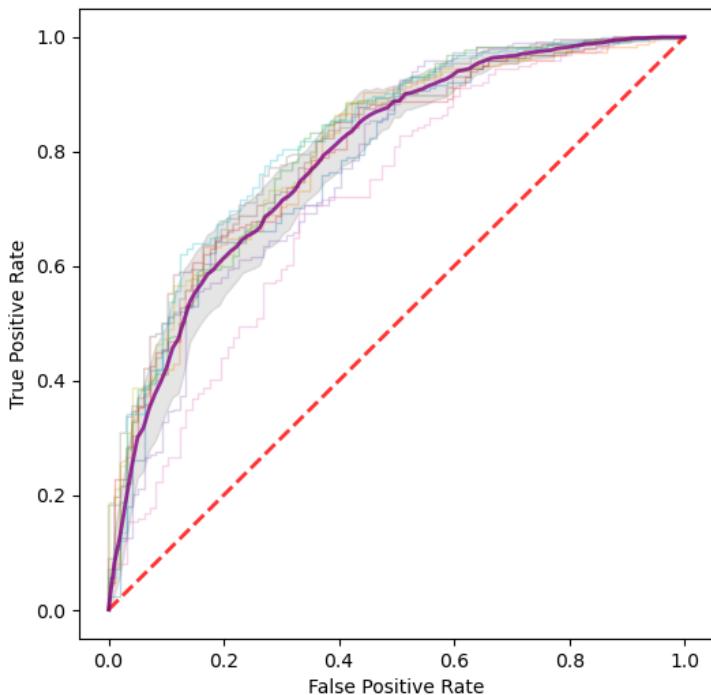
index	Balanced Accuracy	Accuracy	F1 Score	Sensitivity (Recall)	Specificity	Precision (PPV)	TP	TN	FP
Naive Bayes	0.679	0.6805	0.7464	0.683	0.6749	0.8234	146.3	65.4	31.5
Logistic Regression	0.7186	0.721	0.7812	0.725	0.7122	0.8478	155.3	69.0	27.9
Decision Tree	0.6684	0.6815	0.7519	0.7031	0.6337	0.8092	150.6	61.4	35.5
Random Forest	0.7026	0.7445	0.8143	0.8137	0.5914	0.8152	174.3	57.3	39.6
Gradient Boosting	0.685	0.7631	0.8383	0.8922	0.4778	0.7909	191.1	46.3	50.6
XGB	0.7059	0.7448	0.8134	0.8091	0.6027	0.8185	173.3	58.4	38.5
LGB	0.7063	0.7274	0.7935	0.7624	0.6502	0.8286	163.3	63.0	33.9
SVM	0.7122	0.7258	0.7894	0.7484	0.676	0.8361	160.3	65.5	31.4
ANN	0.703	0.774	0.8442	0.8912	0.5148	0.8031	190.9	49.9	47.0
K Neighbors	0.6737	0.7351	0.8123	0.8366	0.5107	0.7926	179.2	49.5	47.4
eLCS	0.692	0.7563	0.8298	0.8627	0.5212	0.7997	184.8	50.5	46.4
XCS	0.5729	0.7078	0.8136	0.9308	0.215	0.7265	199.4	20.8	76.1
ExSTraCS	0.6939	0.7268	0.7972	0.781	0.6068	0.815	167.3	58.8	38.1

index	FN	NPV	LR+	LR-	ROC AUC	PRC AUC	PRC APS
Naive Bayes	67.9	0.4905	2.1464	0.4717	0.741	0.8477	0.8479
Logistic Regression	58.9	0.541	2.5892	0.3878	0.7953	0.8808	0.8814
Decision Tree	63.6	0.4931	1.9659	0.4739	0.7302	0.834	0.8313
Random Forest	39.9	0.5893	2.0209	0.3177	0.7915	0.8815	0.882
Gradient Boosting	23.1	0.6692	1.7202	0.2265	0.785	0.8766	0.8766
XGB	40.9	0.5905	2.0615	0.3175	0.7929	0.8819	0.8824
LGB	50.9	0.5549	2.2172	0.3657	0.7883	0.8806	0.8811
SVM	53.9	0.551	2.3251	0.3727	0.7888	0.8775	0.878
ANN	23.3	0.6921	1.8611	0.2075	0.7981	0.8814	0.8822
K Neighbors	35.0	0.5963	1.7544	0.3116	0.7563	0.8578	0.8466
eLCS	29.4	0.6326	1.8282	0.2669	0.7677	0.8676	0.8683
XCS	14.8	0.5754	1.221	0.2482	0.7081	0.8299	0.8312
ExSTraCS	46.9	0.558	2.0161	0.3614	0.7728	0.868	0.8688

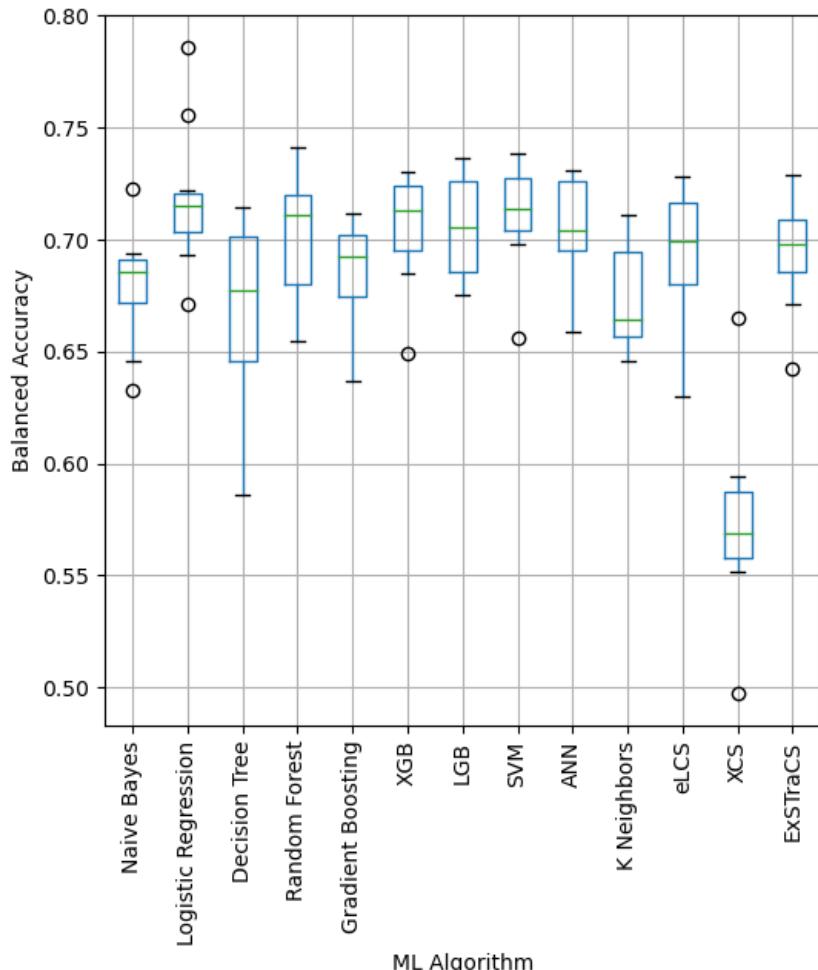
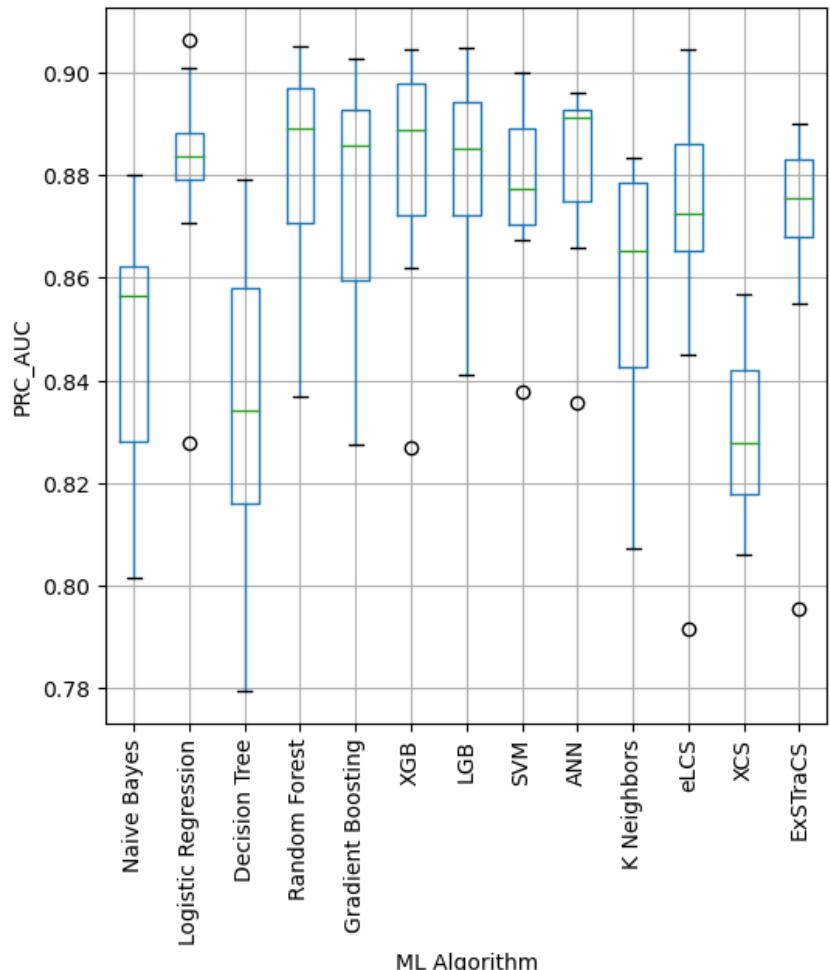
Modeling: Average - ROC/PRC



Modeling: XGB – ROC/PRC

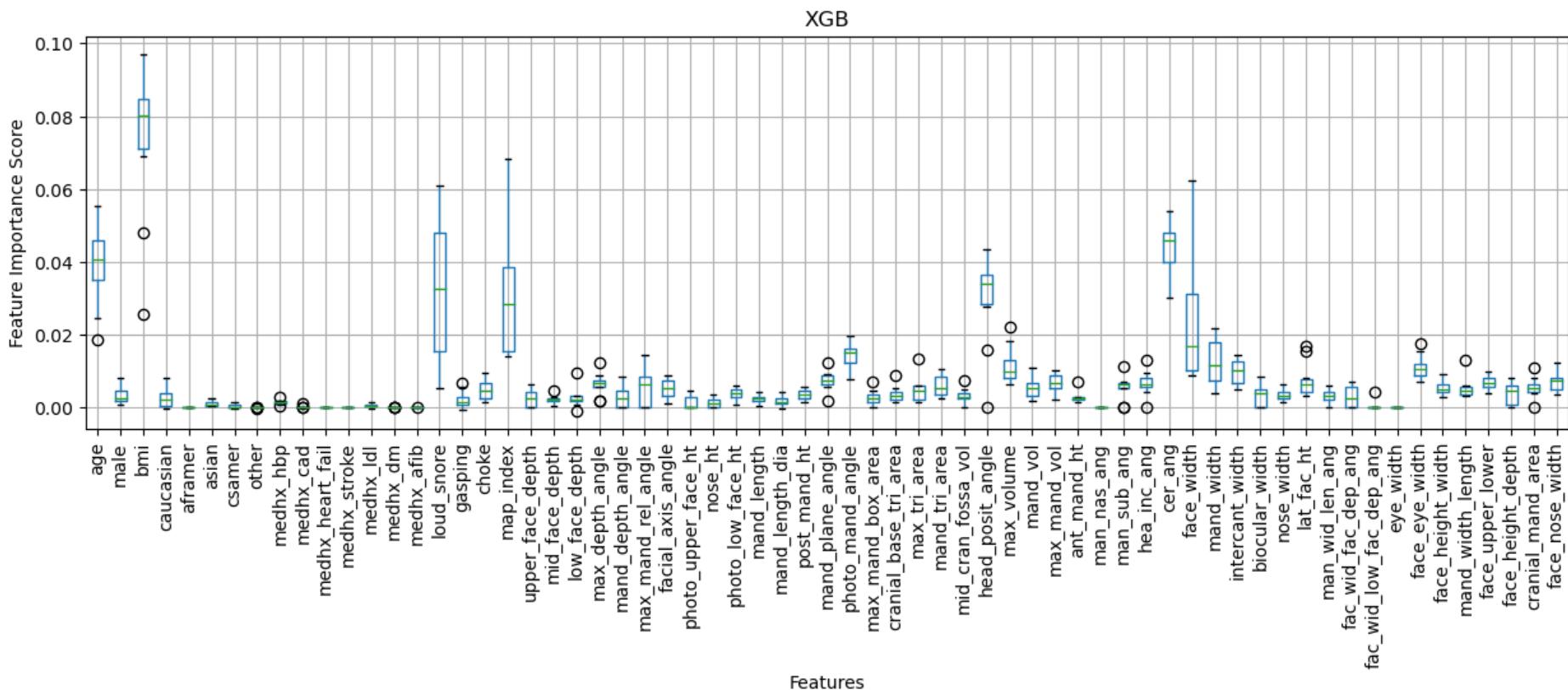


Modeling: Comparing ML Statistics

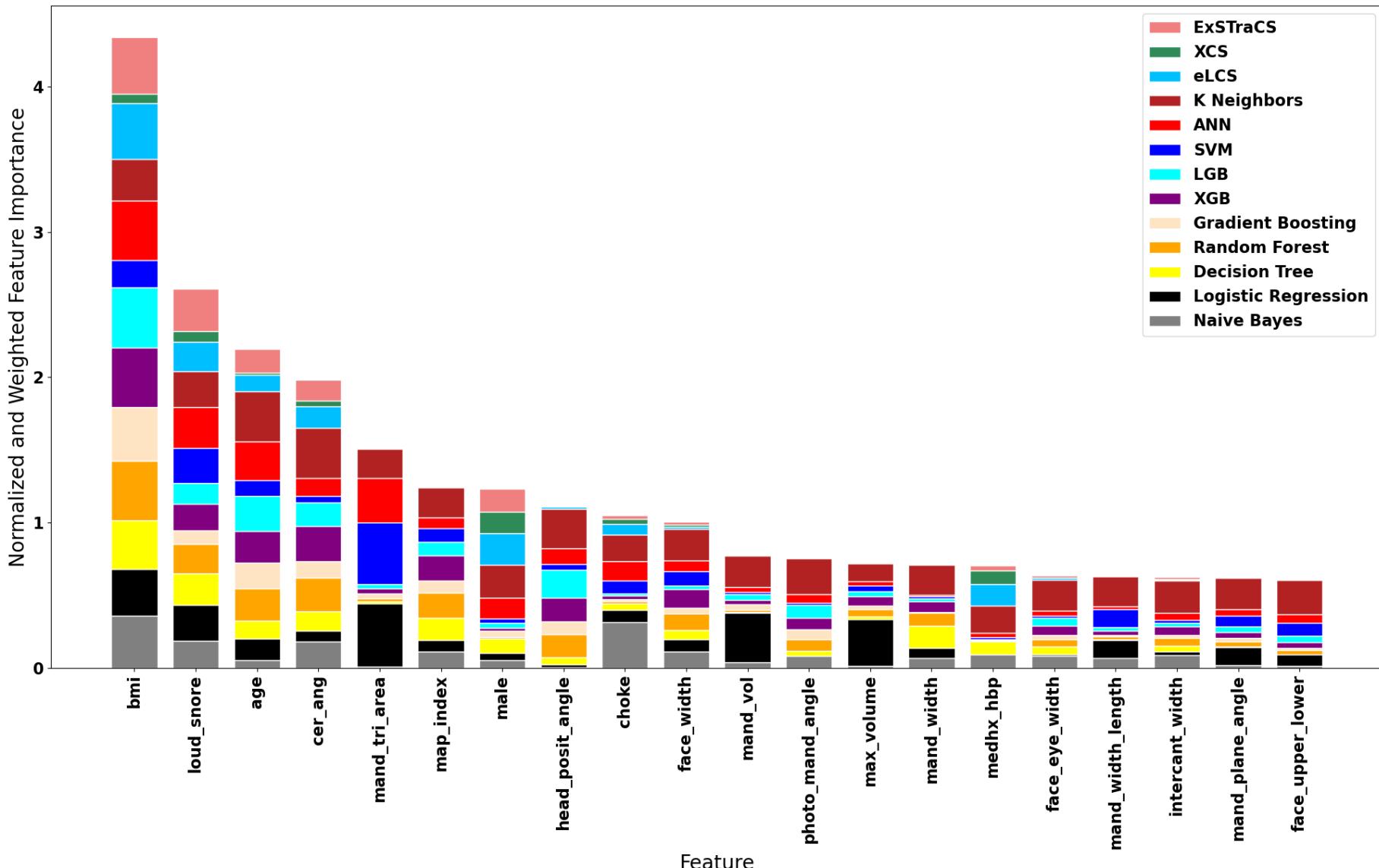


Modeling: Feature Importance

- Internal Feature Importance Estimators (if available)
- Permutation Feature Importance Estimation
 - Applied for all ML algorithms or whenever internal estimators unavailable



Modeling: Composite FI Plot



Statistical Comparisons

- Nonparametric statistical comparisons
 - Kruskall Wallis: one-way ANOVA on ranks is a non-parametric method for testing whether samples originate from the same distribution
 - Wilcoxon Rank Sum: test whether two samples (pairwise) are likely to derive from the same population
- Performed for
 - For each metric (across all algorithms)
 - For each metric (between datasets)
 - For each metric (comparing best performing algorithms)

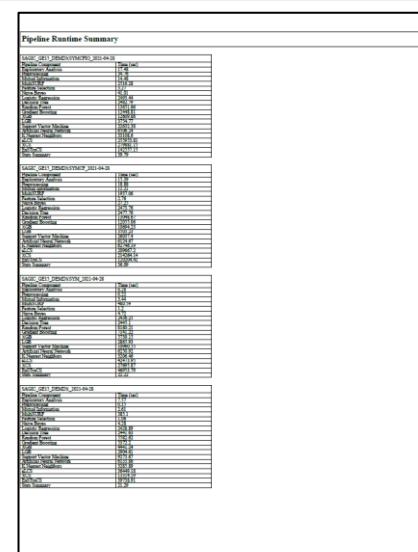
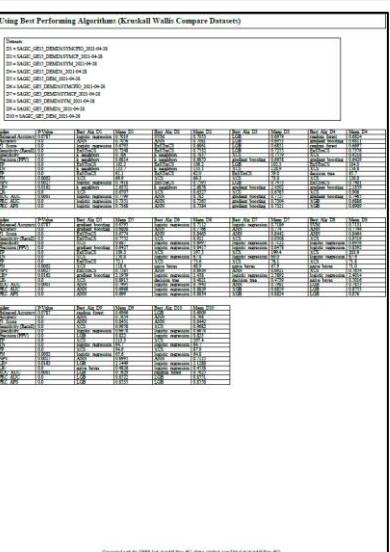
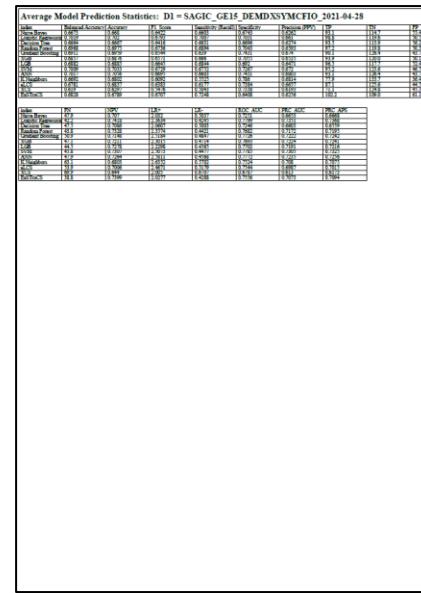
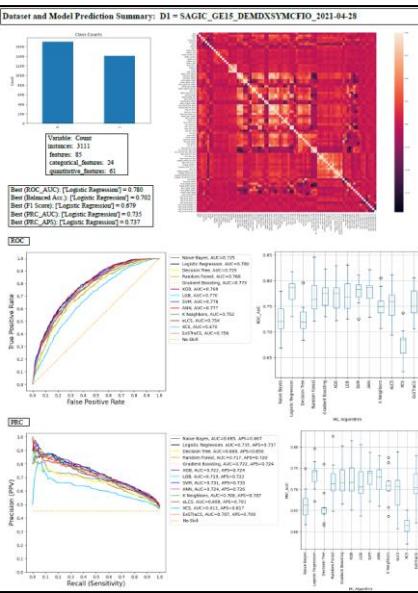
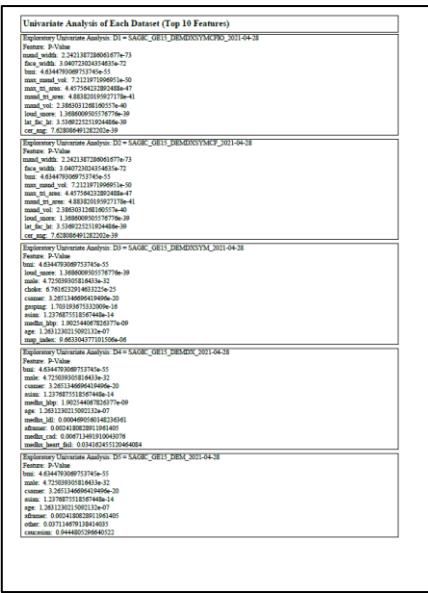
PDF Pipeline Summary Report

AutoMLPipes-BC Training Summary Report: 2021-08-18 14:20:45.409734	
Pipeline Settings:	ML Modeling Algorithms:
class_label: chd	SGD: True
max_iter: 1000	LGB: True
random_state: 42	RF: True
regression: False	GB: True
statistical significance cutoff: 0.05	XGB: True
categorical_imputation: True	LR: True
partition method: 5	ADA: True
gradient boosting: True	AB: True
specified categorical variables: None	ETC: True
data scaling: True	SGD: True
internal information: True	RFECV: True
TUFS: False	ETC: True
TUFS_RF: 0.5	SGD: True
MultiRF: 1000	RFECV: True
max_features_to_be_kept: 1000	ETC: True
max_depth: 1000	SGD: True
primary_metric: balanced_accuracy	RFECV: True
number_of_folds: 5	ETC: True
number_of_repetitions: 1000	SGD: True
number_of_trees_per_stage: 1000	RFECV: True
hyperparam_tuner: SGD	ETC: True

Datasets:

- D1 = SAGIC_GEI1_DEMDNSYMCIO_2021-04-28
- D2 = SAGIC_GEI1_DEMDNSYMCIO_2021-04-28
- D3 = SAGIC_GEI1_DEMDNSYMCIO_2021-04-28
- D4 = SAGIC_GEI1_DEMDNSYMCIO_2021-04-28
- D5 = SAGIC_GEI1_DEMS_2021-04-28
- D6 = SAGIC_GEI1_DEMS_2021-04-28
- D7 = SAGIC_GEI1_DEMDNSYMCIO_F_2021-04-28
- D8 = SAGIC_GEI1_DEMDNSYMCIO_F_2021-04-28
- D9 = SAGIC_GEI1_DEMS_2021-04-28
- D10 = SAGIC_GEI1_DEMS_2021-04-28

Generated with the F800-Lab-AutoML-Pipe-BC: <https://github.com/Urbslab/F800Lab>



Implementation

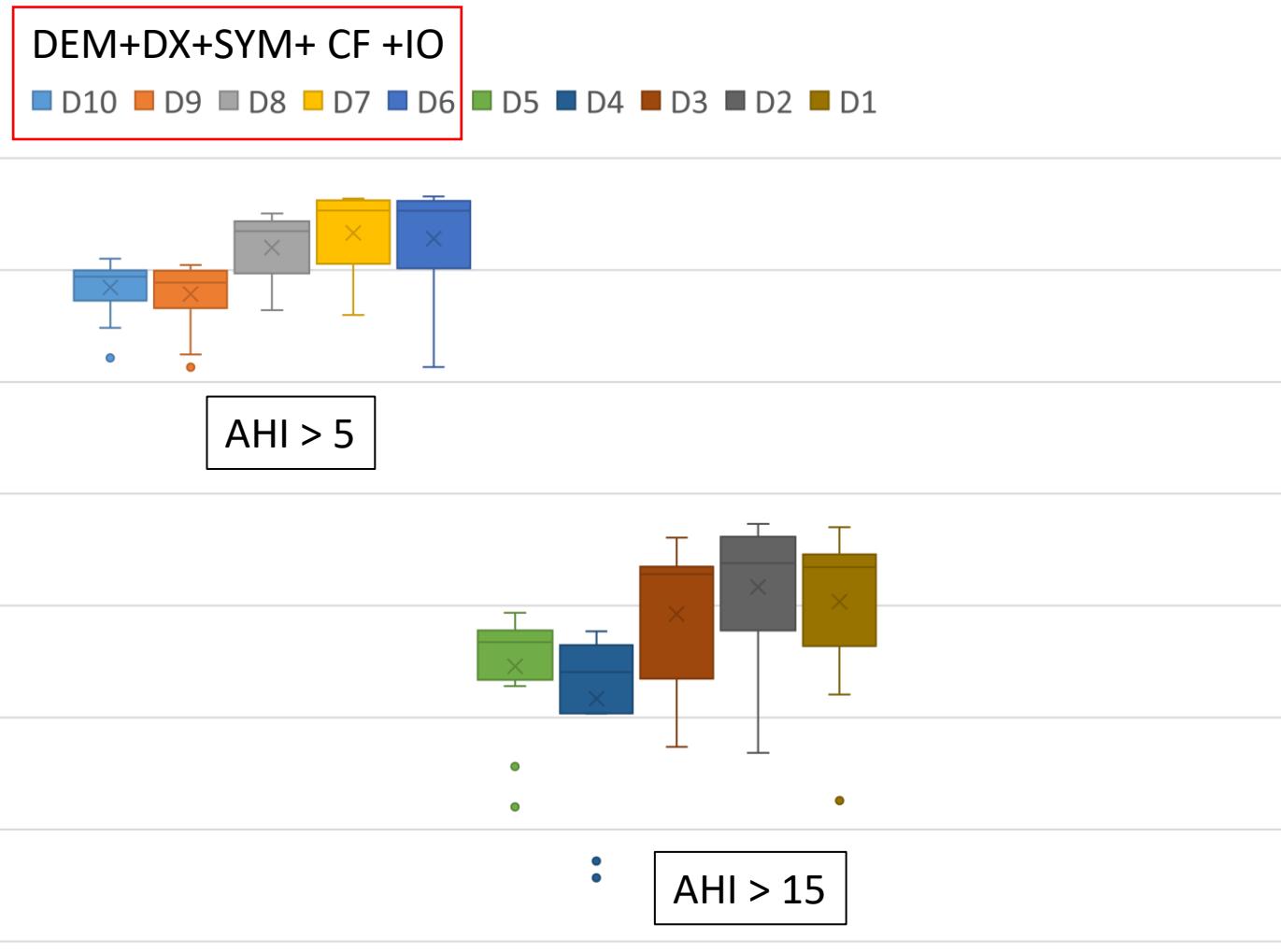
<https://github.com/UrbsLab/AutoMLPipe-BC>

- Detailed README
 - Accessible to use for those with very basic Python experience
- Local (serially)
 - Jupyter notebook
 - Command line
- Computing Cluster (In Parallel)
- PostHoc Jupyter Notebooks:
 - Regenerate figures
 - Evaluate models using alternative decision thresholds
 - Show decision tree models

Phase 1: Exploratory Analysis
Phase 2: Data Preprocessing
Phase 3: Feature Importance Evaluation
Phase 4: Feature Selection
Phase 5: Machine Learning Modeling
Phase 6: Statistics Summary
Phase 7: [Optional] Compare Datasets
Phase 8: [Optional] Copy Key Files
Phase 9: [Optional] Generate PDF Training Summary Report
Phase 10: [Optional] Apply Models to Replication Data
Phase 11: [Optional] Generate PDF 'Apply Replication' Summary Report

Average PRC-AUC Across All Datasets

Each box includes average PRC-AUC for all 13 ML Algorithms



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DEPARTMENT OF
**BIOSTATISTICS
EPIDEMIOLOGY &
INFORMATICS**



Institute for
Biomedical Informatics

Lab GitHub: <https://github.com/UrbsLab>

YouTube: <https://www.youtube.com/channel/UCHIKWNLhgIKHJpkxw993sKQ>

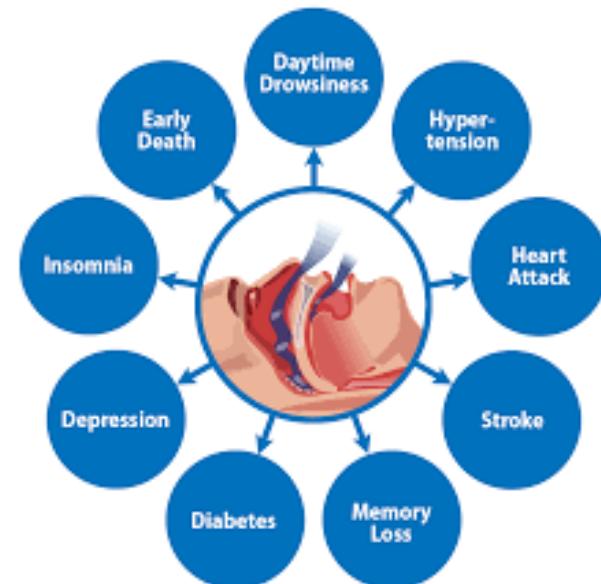
Funding: (NIH) R01 HL134015

(DOD) W81XWH-17-1-0276

UPenn award for Faculty Mentoring of Undergraduate Research

Application to Obstructive Sleep Apnea (OSA)

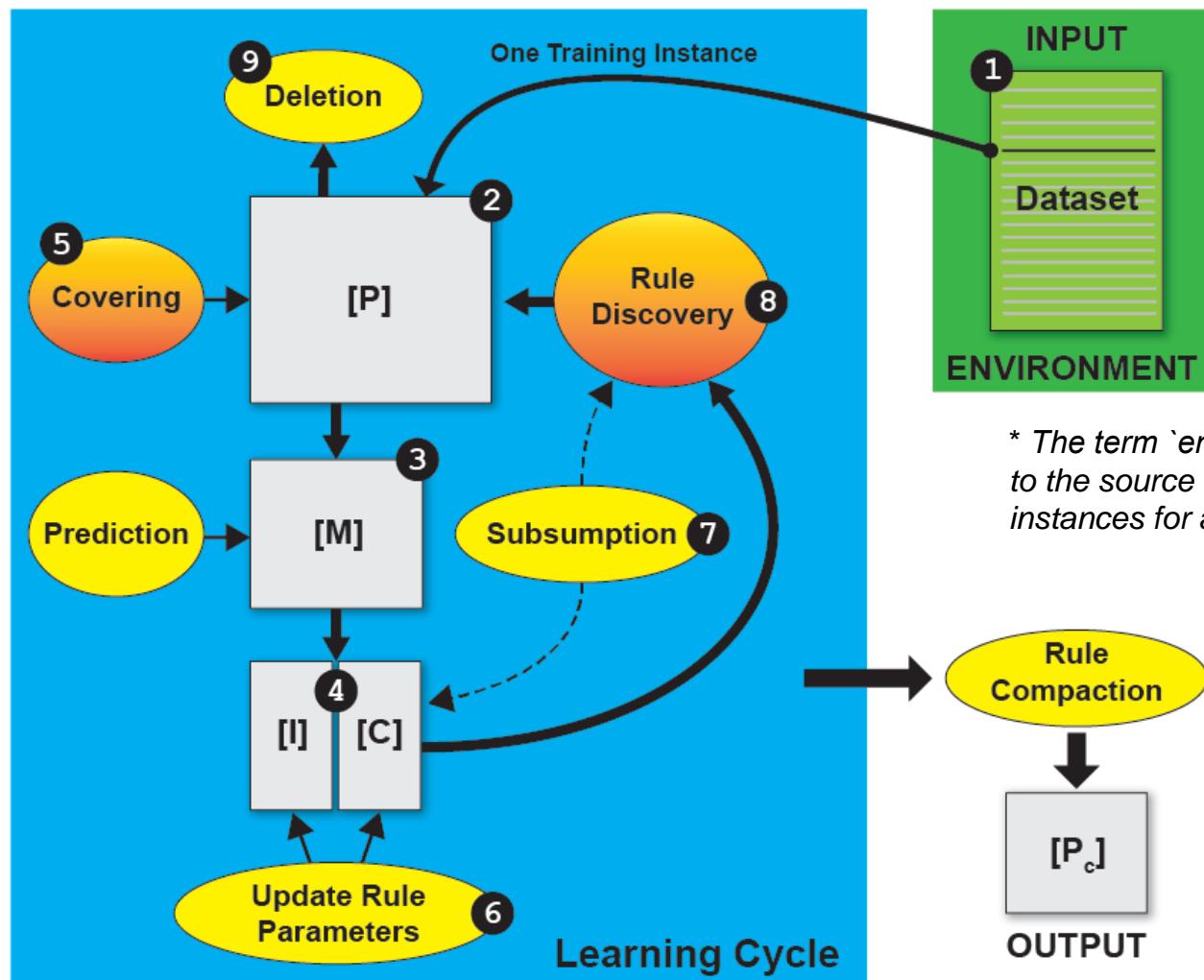
- Sleep Apnea Global Interdisciplinary Consortium (SAGIC)
 - 3111 de-identified subjects who underwent polysomnography (PSG)
- Target Outcome: Apnea-hypopnea index (AHI)
 - Events/hour
 - OSA diagnosis:
 - Liberal Cutoff: AHI > 5 (Controls: 969, Cases: 2142)
 - Conservative Cutoff: AHI > 15 (Controls: 1410 , Cases: 1701)
- Features (5 categories, 85 total):
 - Demographics (DEM) - 8
 - Comorbidities (DX) -7
 - Symptoms (SYM) - 4
 - Craniofacial Photos (CF) - 46
 - Intraoral Photos (IO) - 39
- Comparing sequential addition of feature categories
 - DEM + DX + SYM + CF + IO



AutoMLPipe-BC Goals

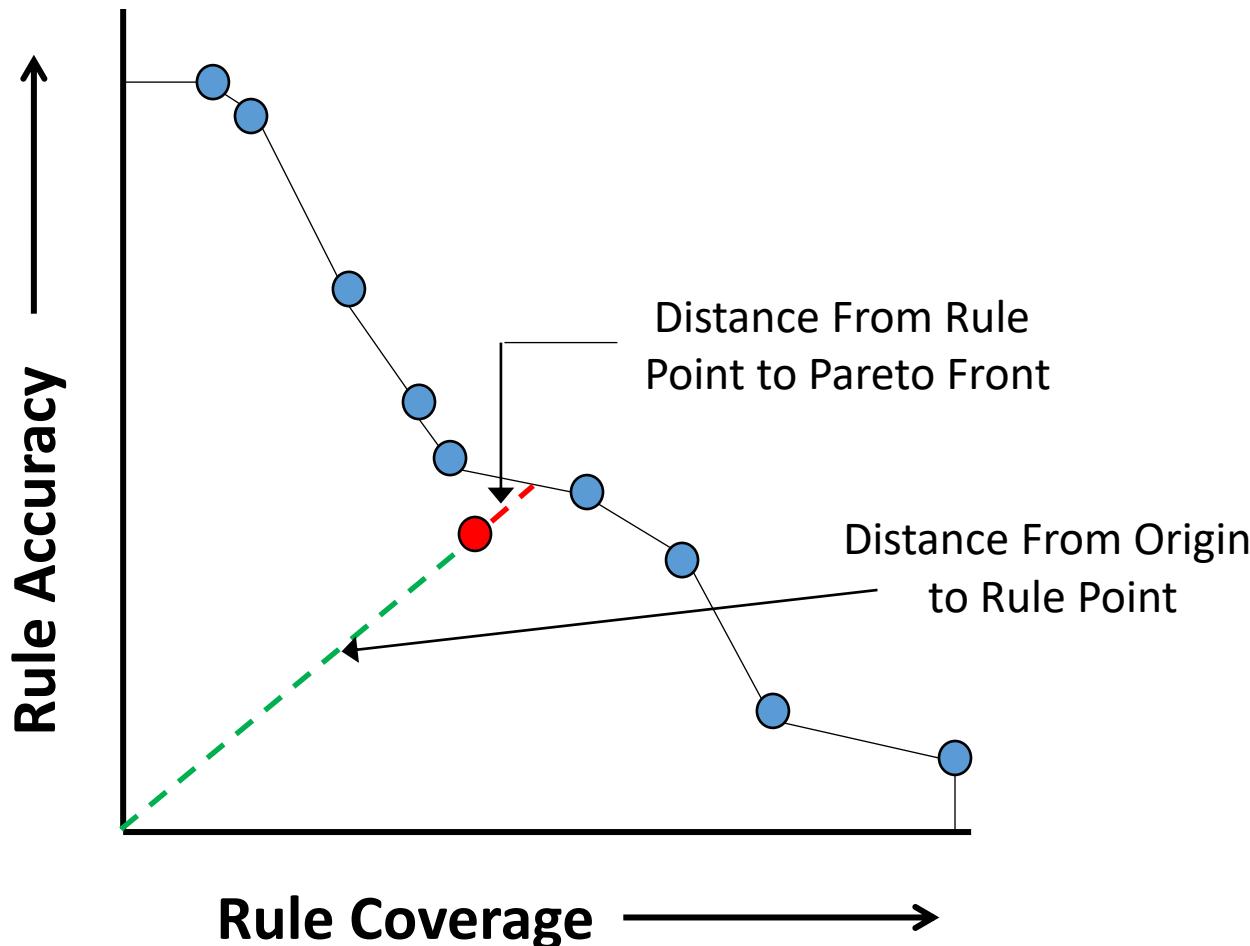
- Automate and parallelize phases of ML analysis pipeline for binary classification (BC)
 - Obtain a rigorous and thorough look at performance
- Capability to handle big data
 - Feature selection is key
- Capability to deal with typical data challenges
 - Missing data
 - Imbalanced data
 - Data normalization for ML efficacy and interpretability
- Capability to take complex associations into account
- Provide standard of comparison for other AutoML tools
- Compare different ML modeling algorithm performance
- Compare performance between target datasets
 - Addition of different feature types
 - Adjusting (or not) for covariates
- Interpret models
 - Feature importance (within and across models/algorithms)
 - Characterizing associations
- Apply models to future data

LCS – A General Schematic (Supervised)

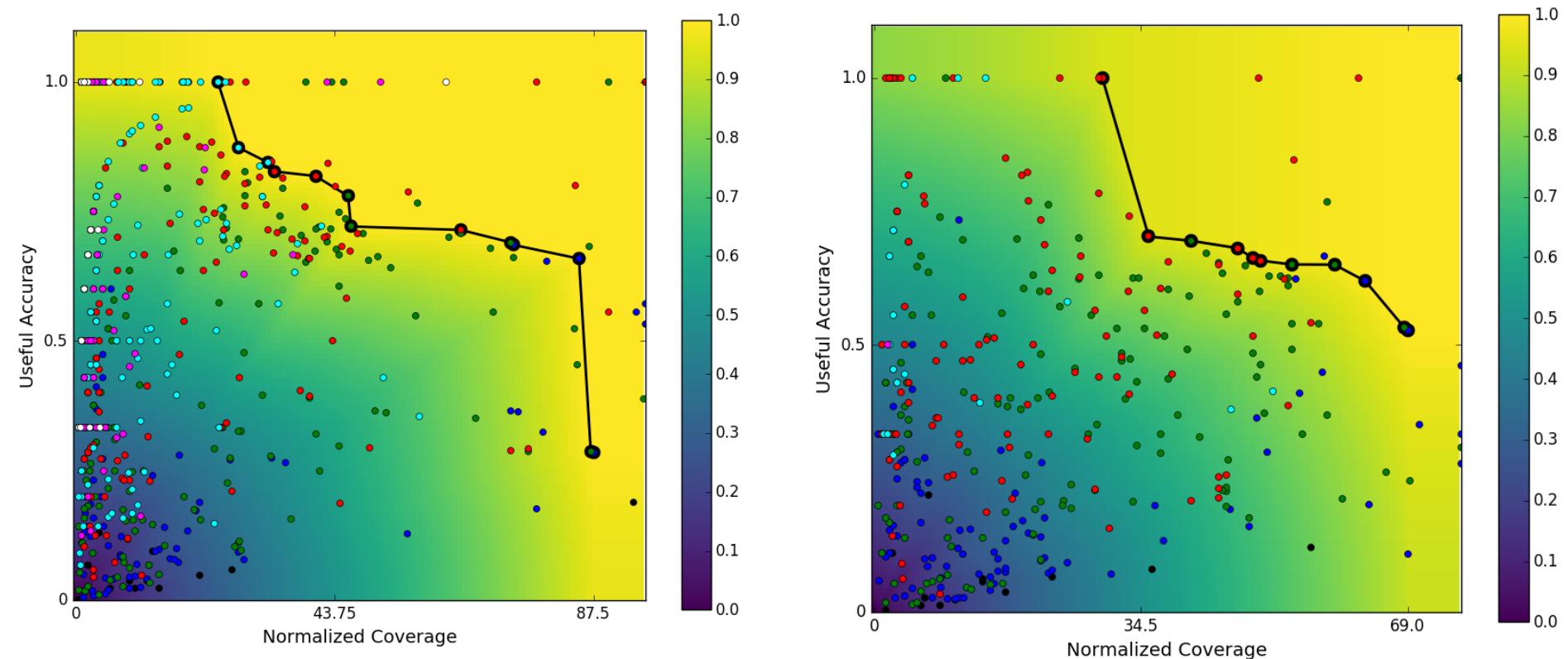


Urbanowicz, Ryan J., and Will N. Browne. *Introduction to learning classifier systems*. Springer, 2017.

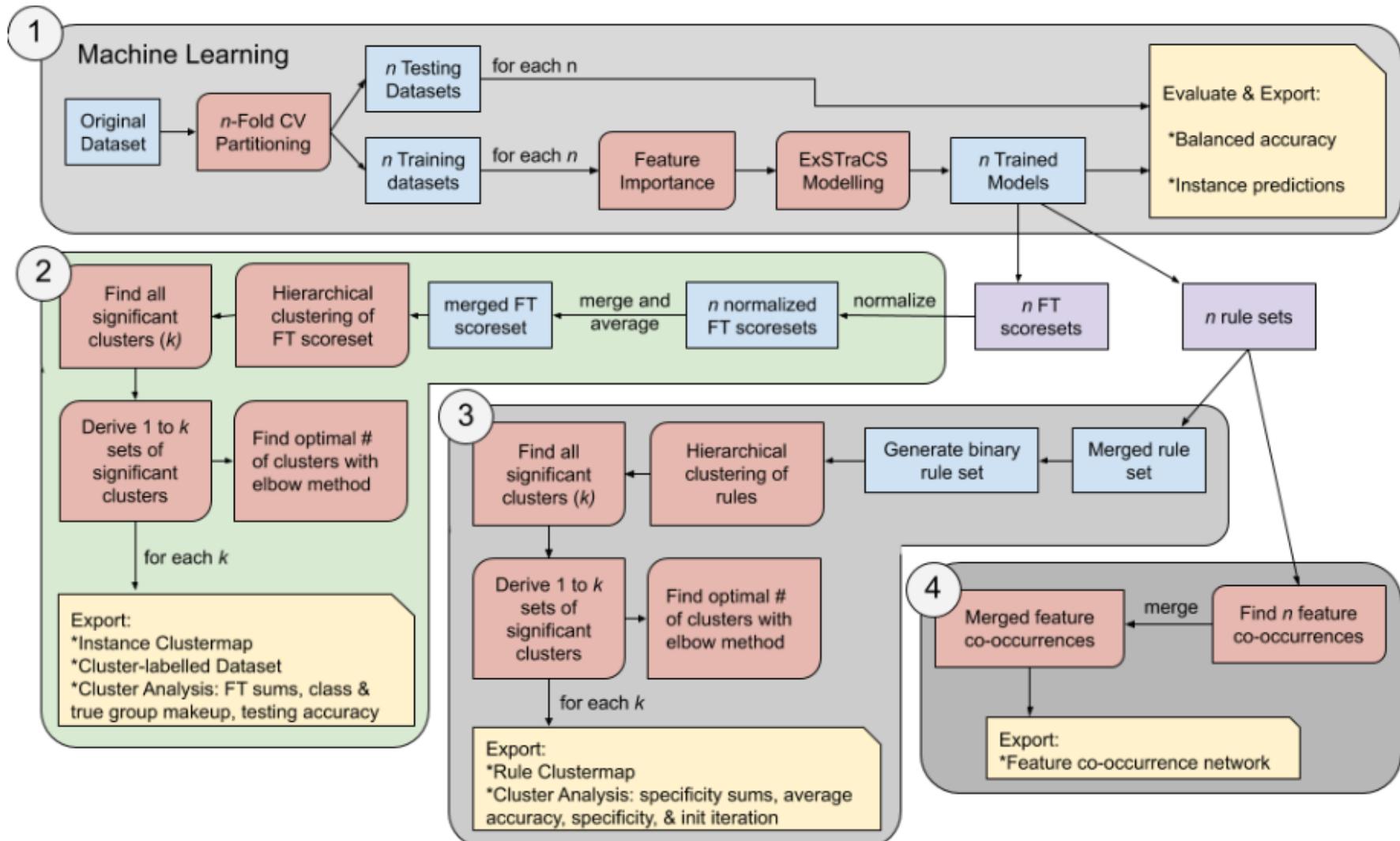
LCS Rule-fitness ‘front’ – Calculate fitness?



Data-set specific rule-fitness gradient



LCS-DIVE: Automated Rule-based Machine Learning Visualization Pipeline

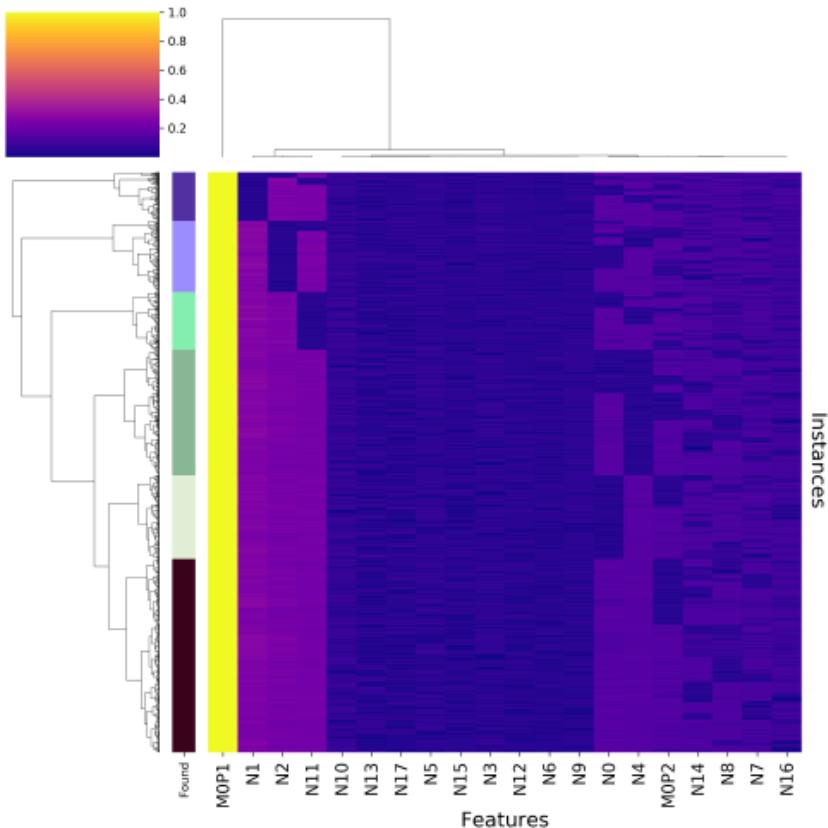


LCS-DIVE: Association Signatures

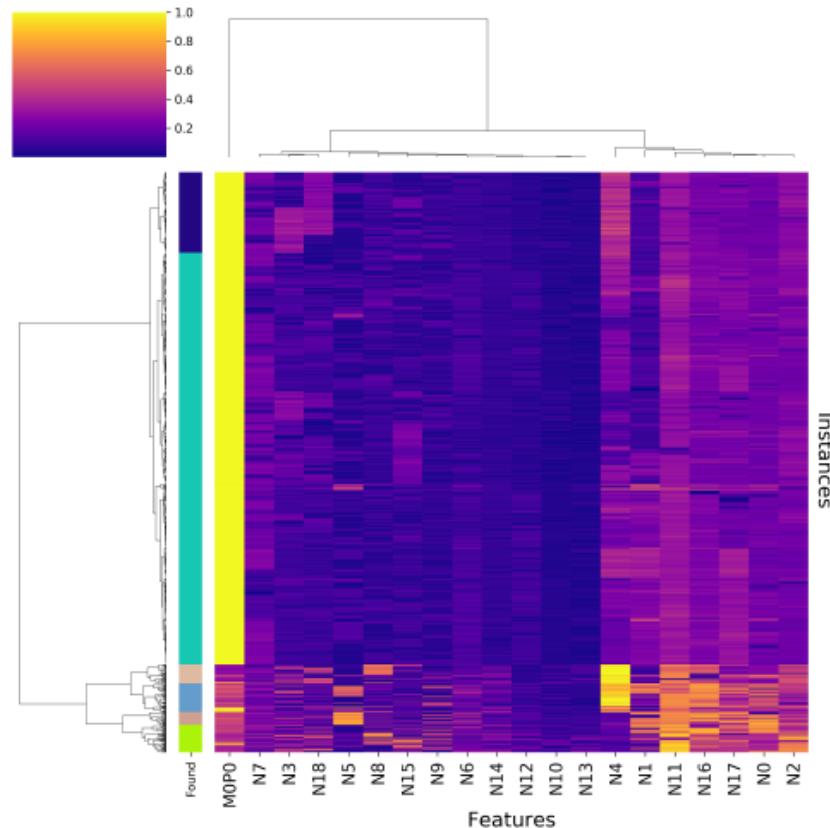
Table 2: Characteristics of the 21 GAMETES Simulated Datasets

Dataset ID	Underlying Association	Predictive Features	# of Models	Model Ratio	Model Heritability	Instances
1 & 2	univariate	1	1	NA	1 & 0.4	1600
3 & 4	additive	2	2	equal	1 & 0.4	1600
5 & 6	additive	4	4	equal	1 & 0.4	1600
7 & 8	2-way pure epistasis	2	1	NA	1 & 0.4	1600
9 & 10	3-way pure epistasis	3	1	NA	1 & 0.2	3200
11 & 12	additive (2-way epistasis + univariate)	4	3	equal	1 & 0.4	1600
13 & 14	additive (2, 2-way epistasis)	4	2	equal	1 & 0.4	1600
15 & 16	heterogeneity (univariate)	2	2	equal	1 & 0.4	1600
17 & 18	heterogeneity (univariate)	4	4	equal	1 & 0.4	1600
19 & 20	heterogeneity (2, 2-way epistasis)	4	2	equal	1 & 0.4	1600
21	heterogeneity (2, 2-way epistasis)	4	2	75 : 25	0.4	1600

LCS-DIVE: Univariate (1 Feature)

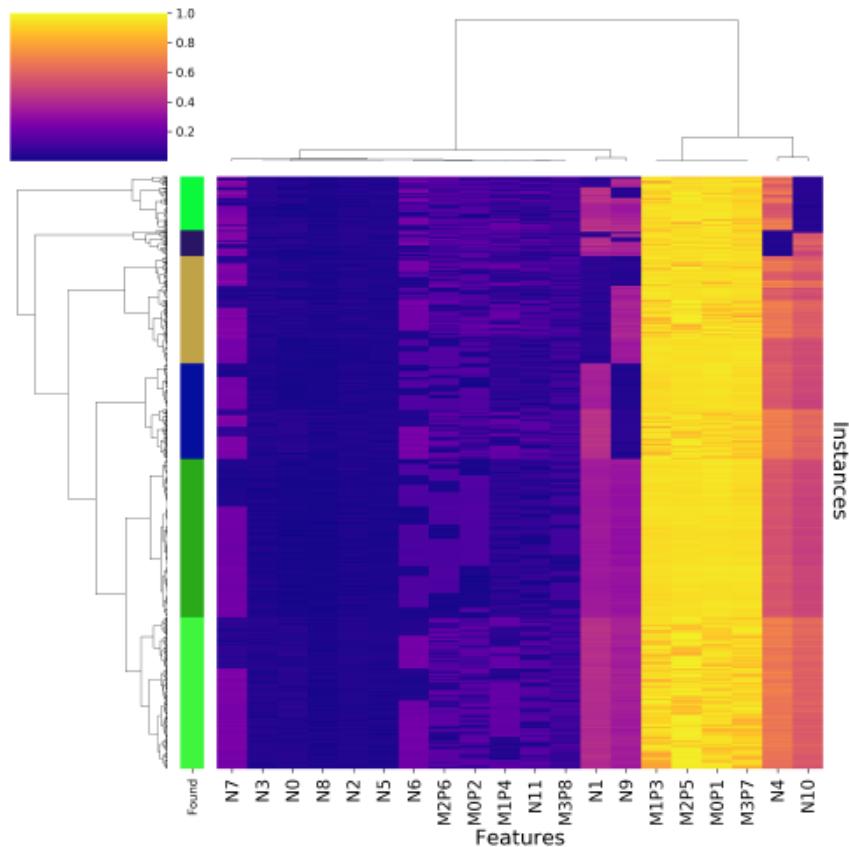


(a) D1: Clean Univariate

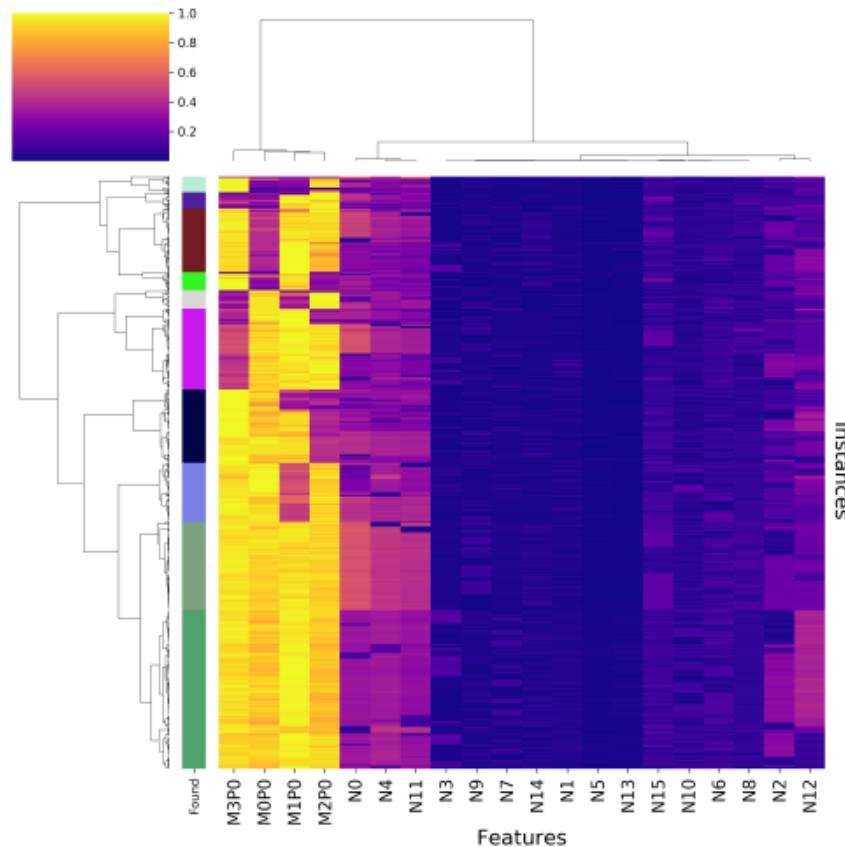


(b) D2: Noisy Univariate

LCS-DIVE: Univariate-Additive (4 Features)

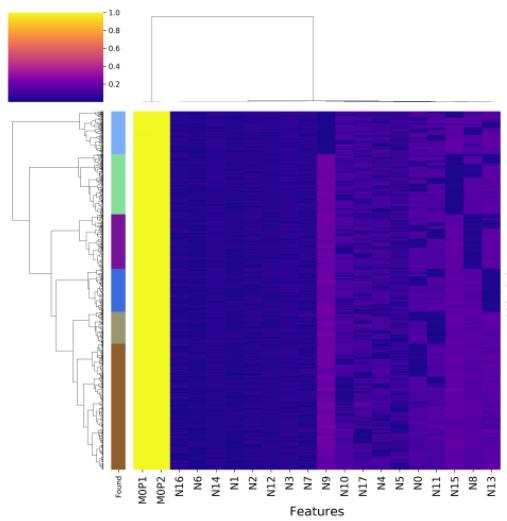


(e) D5: Clean 4-feature additive

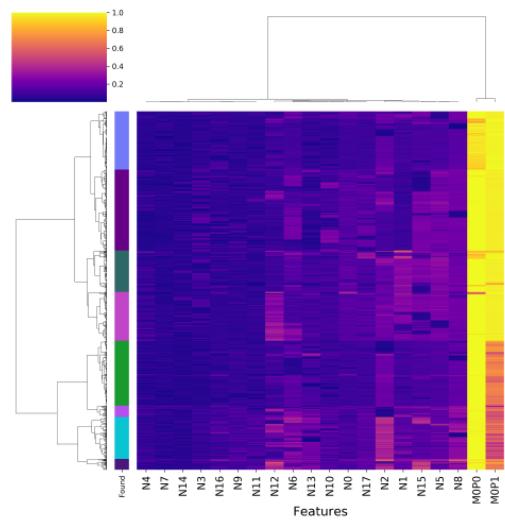


(f) D6: Noisy 4-feature additive

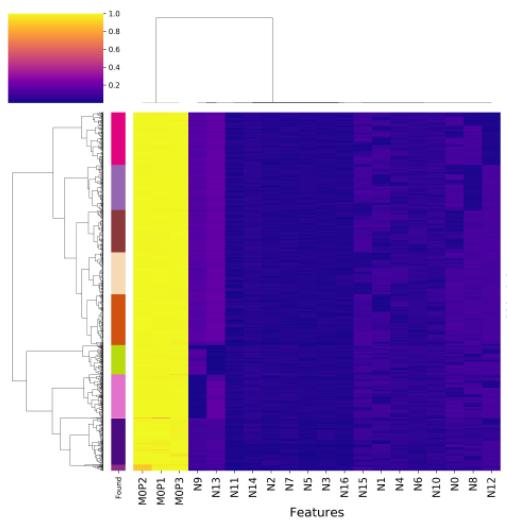
LCS-DIVE: Epistasis



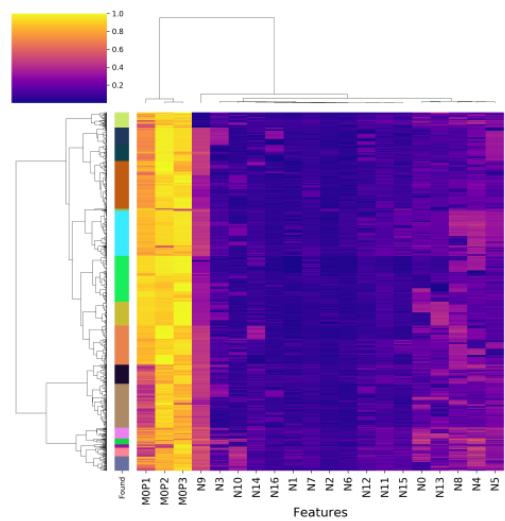
(a) D7: Clean 2-way pure epistasis



(b) D8: Noisy 2-way pure epistasis

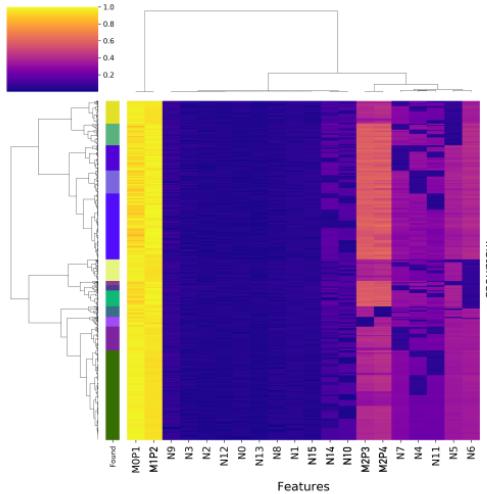


(c) D9: Clean 3-way pure pistasis

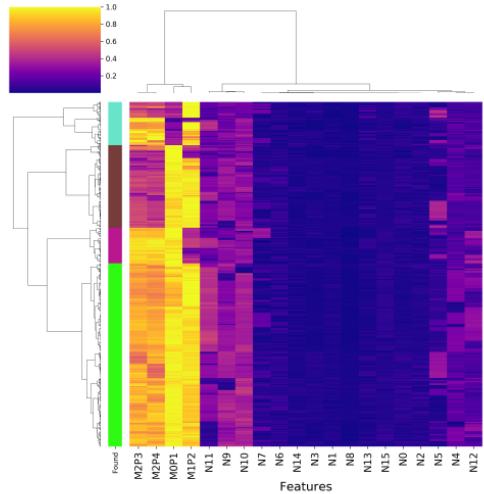


(d) D10: Noisy 3-way pure epistasis

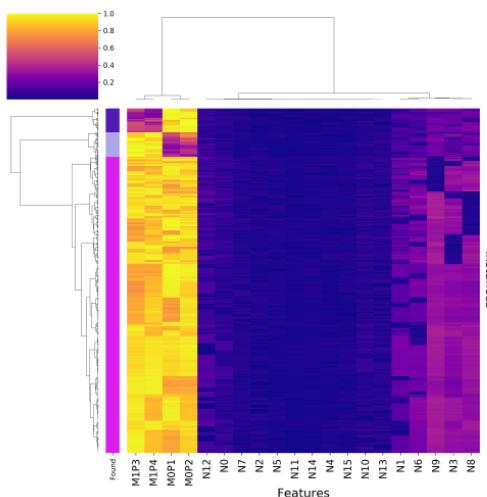
LCS-DIVE: Epistasis+Additive



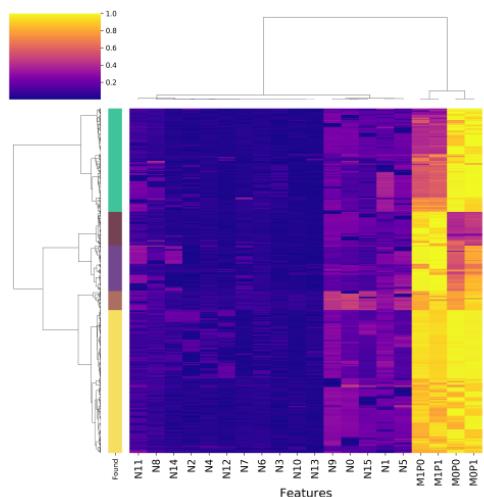
(a) D11:Clean additive (2-way epistasis+univariate)



(b) D12: Noisy additive (2-way epistasis+univariate)

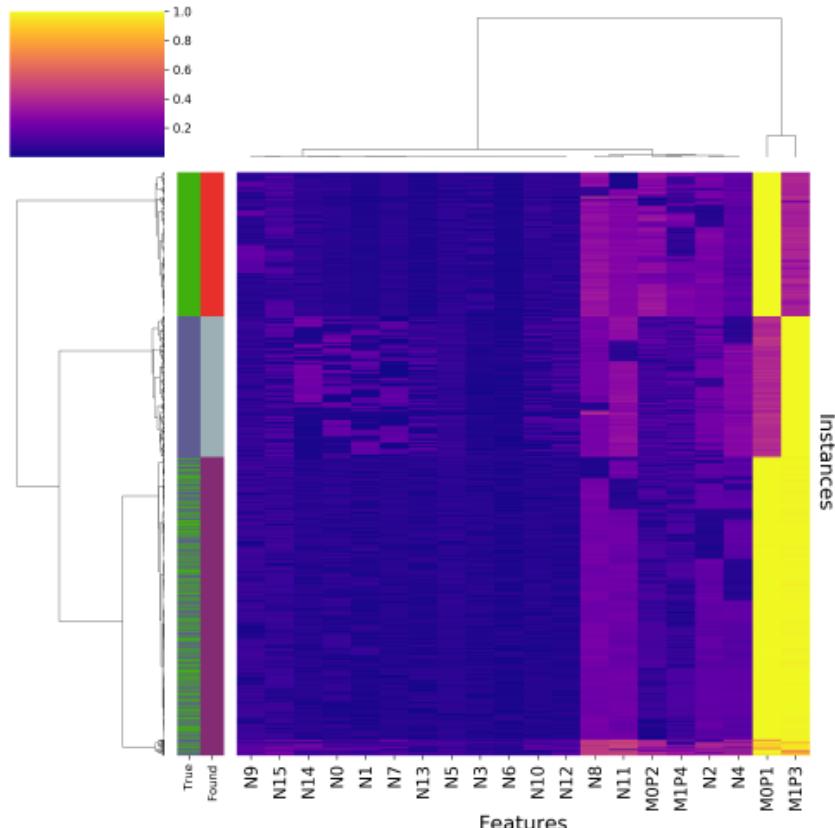


(c) D13: Clean additive (2, 2-way epistasis)

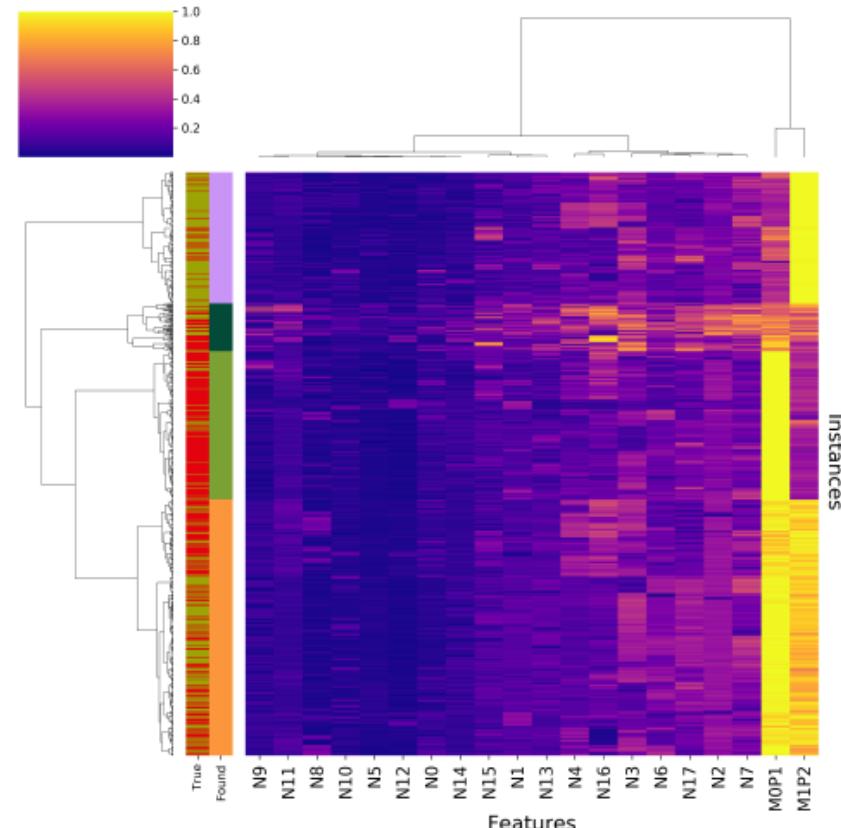


(d) D14: Noisy additive (2, 2-way epistasis)

LCS-DIVE: Heterogeneous-Univariate

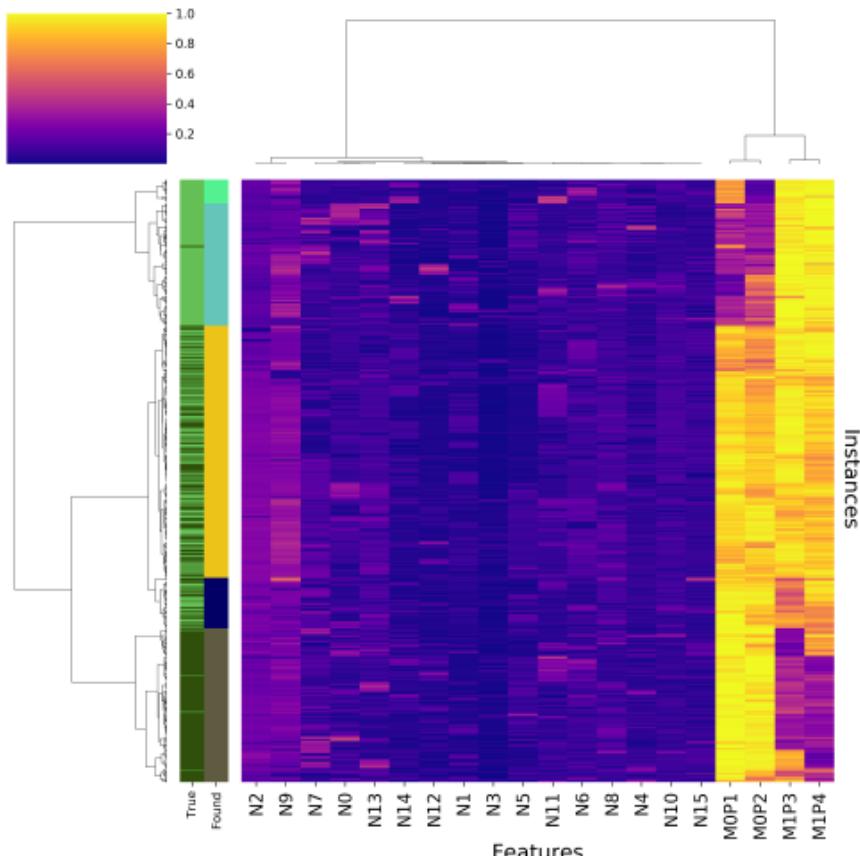


(a) D15: Clean 2-feature het. univariate

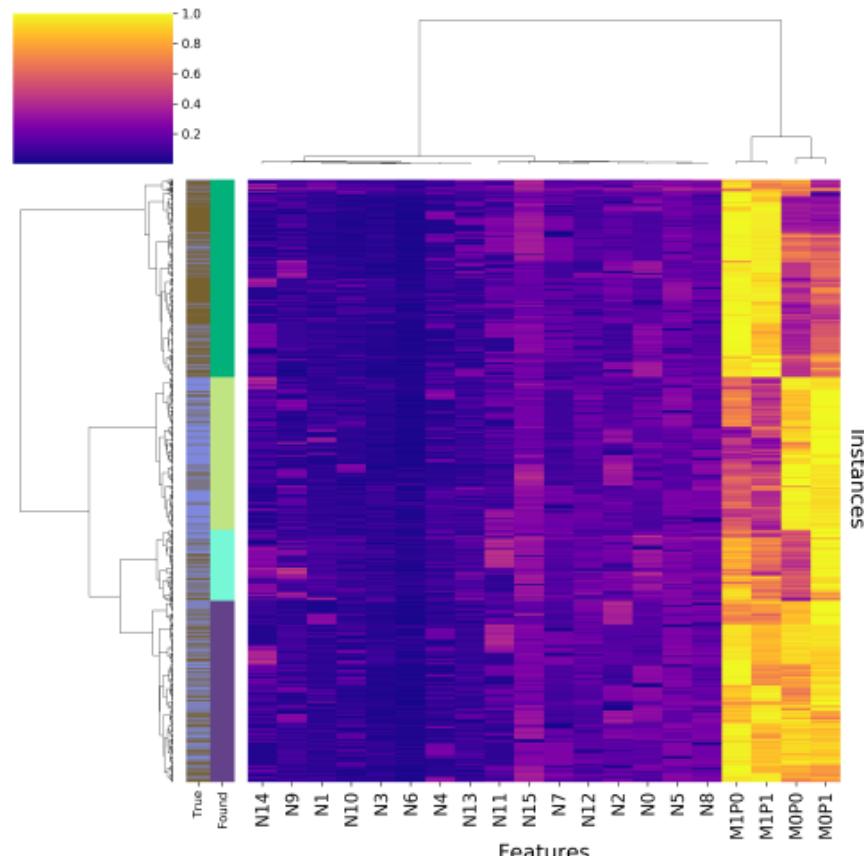


(b) D16: Noisy 2-feature het. univariate

LCS-DIVE: Heterogeneous-Epistasis



(e) D19: Clean 2, 2-way Epistasis, het.



(f) D20: Noisy 2, 2-way Epistasis, het.