



Federated Learning in Computational Toxicology

Our Perspective on the Effiris Hackathon

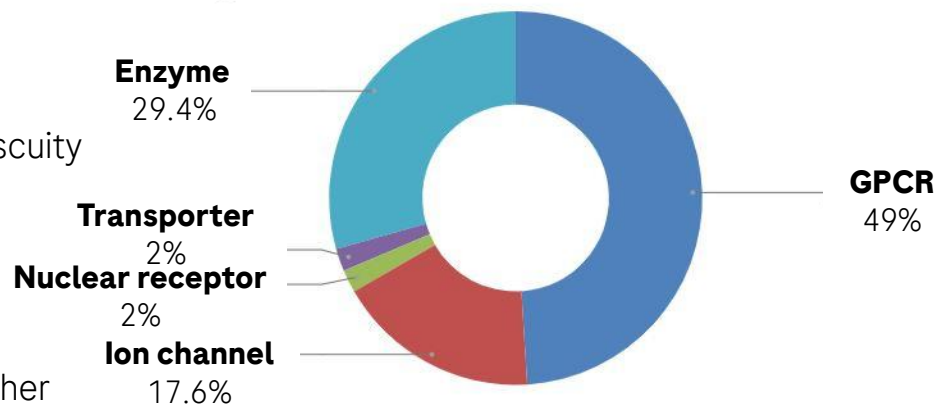
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Secondary pharmacology

Off-target interactions which may occur at related or structurally distinct targets

- Secondary Pharmacology screening is used from Candidate Selection to post Marketing
- Regulatory submissions require secondary pharmacology data to be included
- Panel-based screens are used to assess promiscuity and identify potential safety flags
- Roche recurrent panel for 50 targets
- Panel testing typically uses only a single test concentration at 10 μ M but may need to be higher

Distribution of targets in the Roche Recurrent Profiling panel



Application areas of secondary pharmacology prediction models

Prediction of toxicity to

- Guide optimization of a drug candidate
- Prioritise between different chemical classes, i.e. with very different scaffolds
- Save money - run in silico before submitting to panel:
 - Only submit to panel if in silico hits, or if in silico cannot make an informative prediction
 - Do not submit if compound is predicted to be highly promiscuous

Toxicological (in vitro) data

Can we overcome data challenges using federated learning (FL)?

Challenges of in vitro toxicology data

- Comparably small datasets
- Typically high data imbalances between numbers of active vs. inactive compounds
- Often challenging translatability from binding to mechanistic (e.g. agonist/antagonist) to in vivo data

	Model built on		
	Proprietary data	Public data	Prop. data from other company
Chemical space	limited	broad	limited, but different
Data reliability	high	low	high
Access	easy	easy (need curation)	difficult

Effiris

A secondary pharmacology model suite using federated learning (FL)



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Article

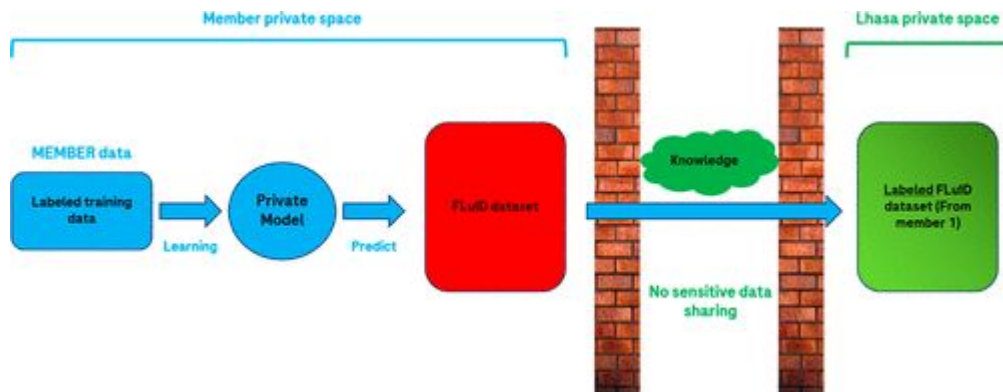
Federated Learning in Computational Toxicology: An Industrial Perspective on the Effiris Hackathon

Davide Bassani, Alessandro Brigo, and Andrea Andrews-Morger*

- Can we increase the chemical space of machine learning (ML) training data - without sharing sensitive data - using FL?
- Effiris federated learning software to predict secondary pharmacology developed by Lhasa Limited, UK
- 50 ML models for off-target prediction
- Organised Effiris Federated Learning hackathon: 9 off-targets, 7 pharmaceutical companies
- Compare models built with Effiris to internally built ML models

Effiris federated learning approach

Student-Teacher approach



No sharing of sensitive (proprietary) data

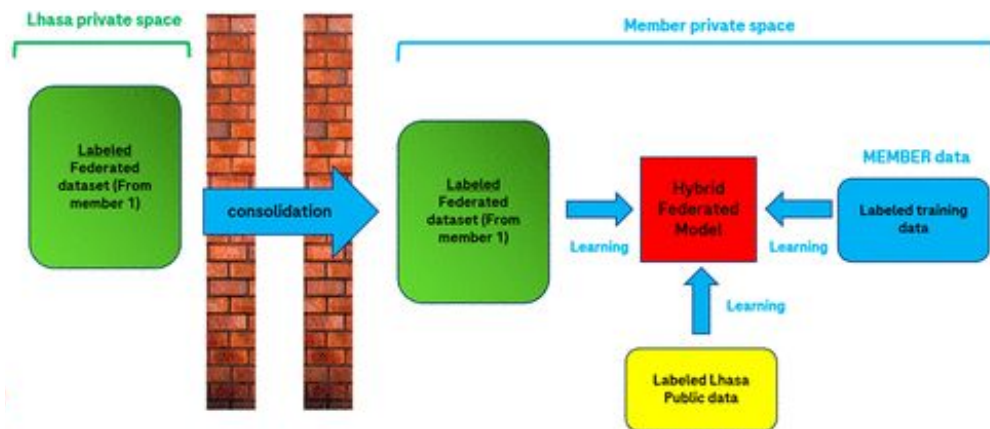
Project information from internal data to 300 K public and unlabelled dataset

Consolidate predicted labels from industry partners

Hypothesis: size and chemical space of dataset more important than accuracy of data

Effiris federated learning approach

Student-Teacher approach



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Differences Effiris - MELLODDY

	Effiris	MELLODDY
Federated learning approach	Student-teacher model	Machine learning ledger orchestration for drug discovery
Endpoints	1 endpoint per model, defined and known	“All available data” used, assay/target names not disclosed
# tasks per model	Single-task	Multi-task
Sharing of	Predicted labels on FLuID dataset	Gradients
Training set	Consolidated, predicted labels for FLuID dataset + internal data	Internal data

Collection of internal data

Effiris evaluation work performed on five off-targets

Effiris: Currently 50
classification models available,
regression models under
development

Test set: time-split 2020-2022

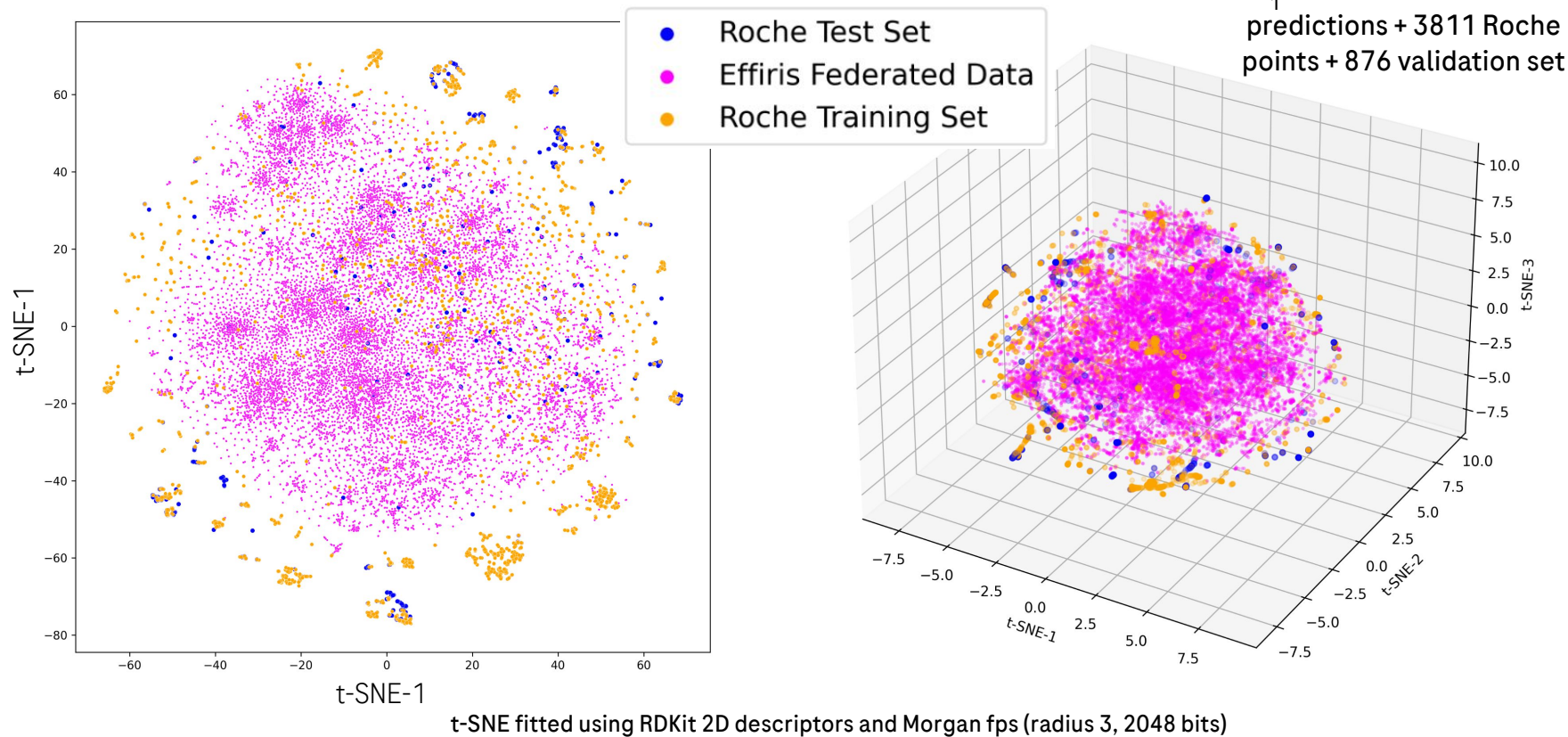
1. Collect internal activity data
2. Binary models: PCT_INHIB data, threshold: $\leq 50\%$ ($10\ \mu\text{M}$)
3. Validate Effiris model on Lhasa and Roche test set
4. Build model using own algorithms on internal and federated data

Target	Assay type	Training set (inactive/active)	Test set (inactive/active)
ACh M ₁	binding	3811 (2829/982)	876 (745/131)
GABA _A Benz	binding	3614 (3216/398)	877 (819/58)
5-HT _{2B}	binding	3374 (2447/927)	876 (709/167)
hERG	Electrophysiology (SynchroPatch)	2285 (1388/897)	556 (309/247)
COX-2	enzymatic	3042 (2815/227)	881 (809/72)

Chemical space analysis

Chemical space comparison between internal & federated data using t-SNE

Example of M_1 : Internal space largely but not completely covered by federated data

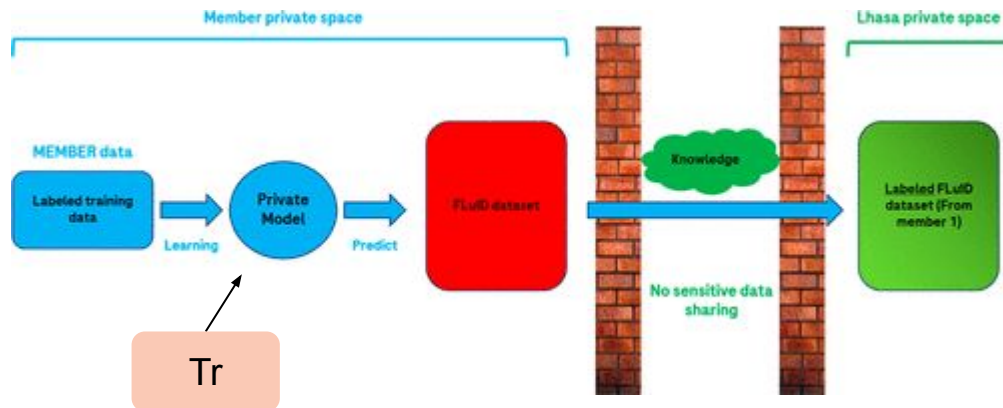


SOHN approach

SOHN

Self Organising Hypothesis Network Classifier

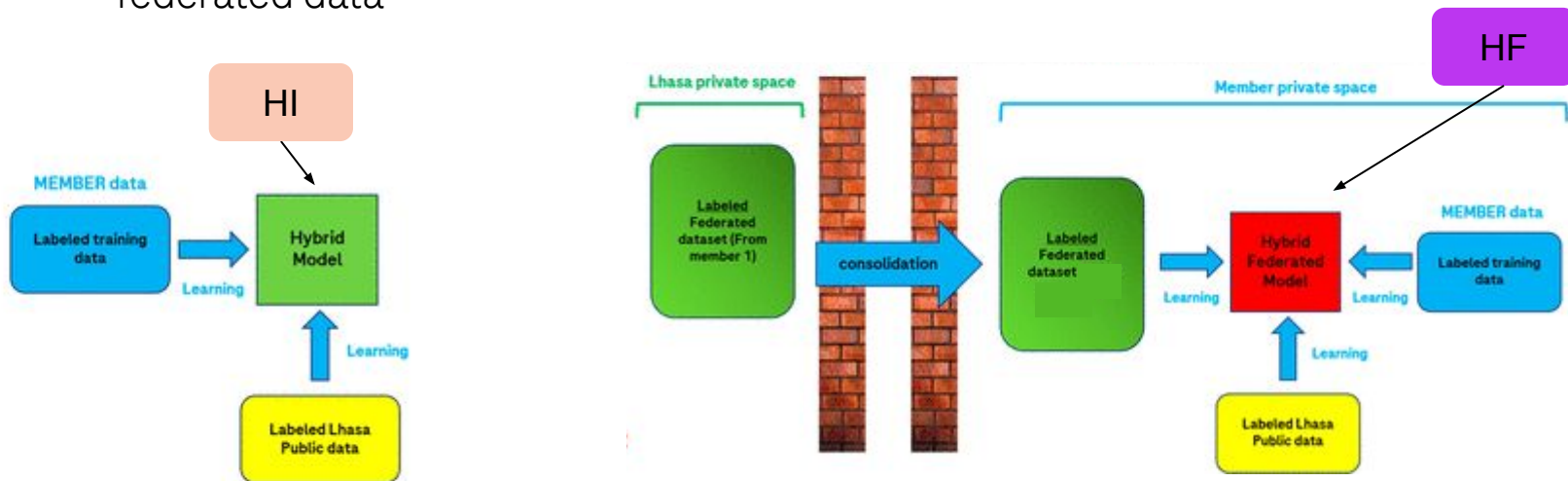
- Upload internal data (Member data) to build SOHN Teacher model (Tr)
- Make predictions on FLuID dataset (external, 350K) and share predicted labels with Lhasa
- Lhasa consolidated labels across teacher models and selected 20K compounds to be used as federated dataset



SOHN

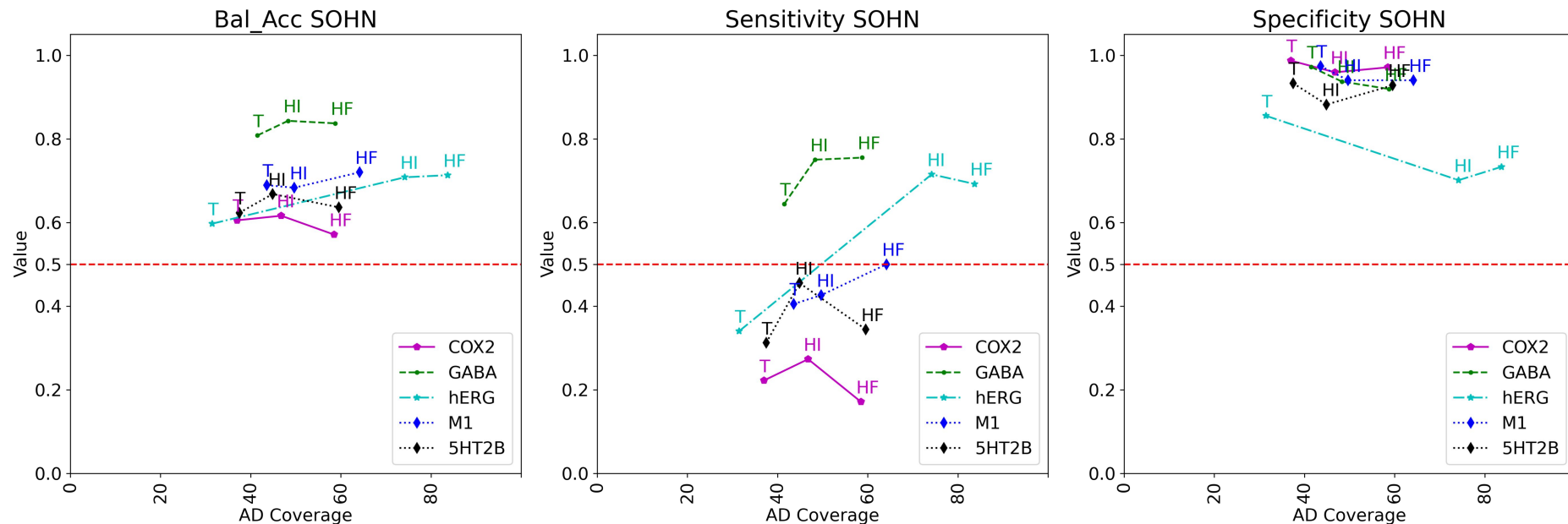
Self Organising Hypothesis Network Classifier

- Hybrid model I (HI): build hybrid Model based on internal and Lhasa public data
- Hybrid model II (HF): build hybrid model based on internal, Lhasa public, and federated data



Impact of Effiris on model performance and applicability domain - SOHN

Applicability domain increase, sensitivity > 0.5 for GABA_A and hERG



T = teacher with internal data

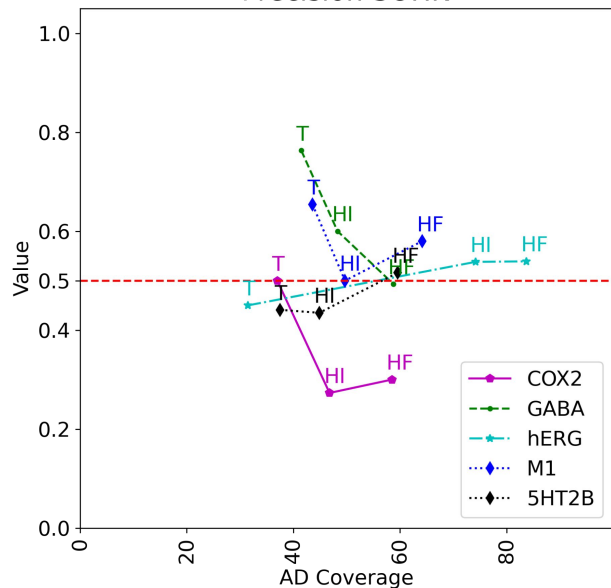
HI = hybrid with public data

HF = hybrid with Lhasa Federated data

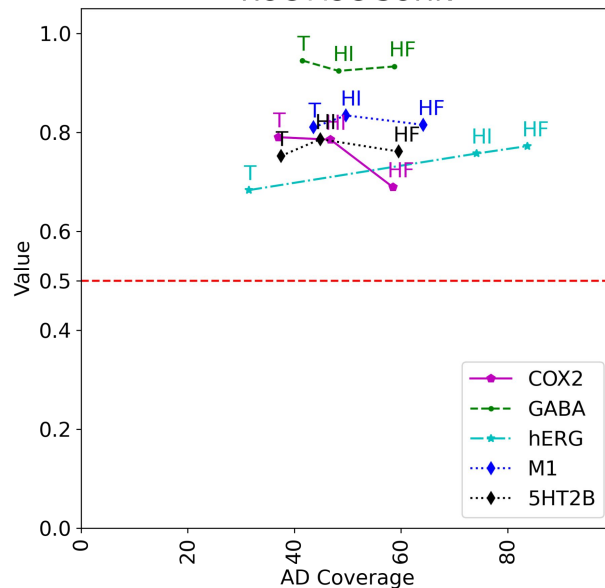
Impact of Effiris on model performance and applicability domain - SOHN

Change of performance measure is target-dependant

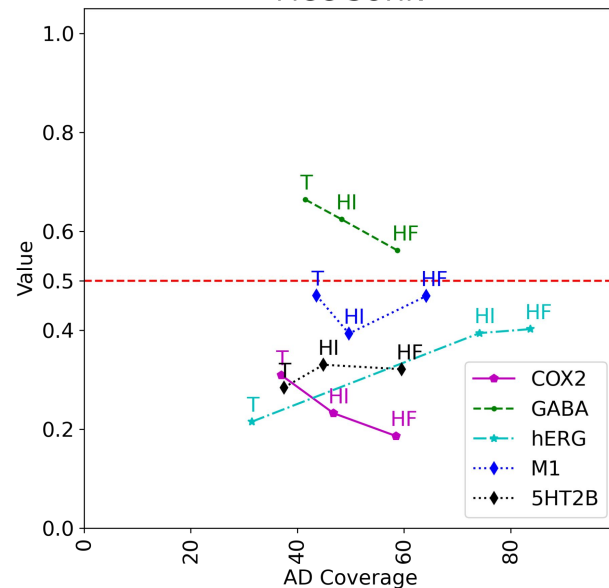
Precision SOHN



ROC AUC SOHN



MCC SOHN



T = teacher with internal data

HI = hybrid with public data

HF = hybrid with Lhasa Federated data

Building internal models

On internal data and federated data

Methods to build internal models

Internal model
building,
benchmark, and
analysis performed
by Davide Bassani



Datasets

Used different sets for
each target: **internal**, **Lhasa**
Federated (20K), **Lhasa**
FLUID (300K+)

ACh M ₁
GABA _A Benz
5-HT _{2B}
hERG
COX-2

RDKit Descriptors:
1D+2D Molecular Descriptors
+
Morgan Fingerprints
from RDKit (2048 bits, radius 3)

ML Models

Gaussian NB
K-Neighbors
Decision Tree
Random Forest
ExtraTrees
AdaBoost
Gradient Boosting
XGBoost
MLP
SGD
SVC

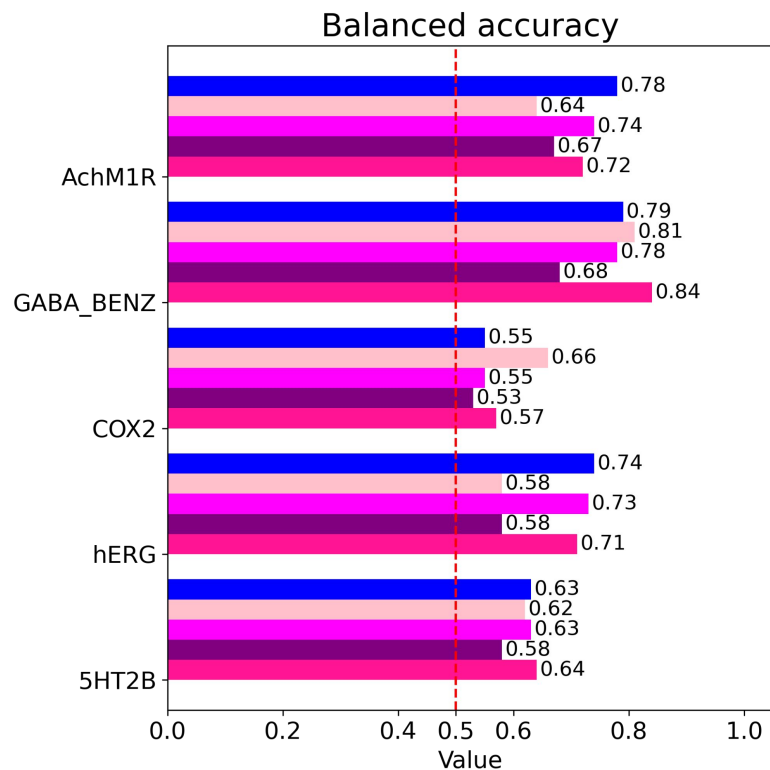
**Hyperparameter
Optimization**
(5-fold, 10 repeats
Stratified CV)

**Prediction on
internal test set
+
comparison with
Lhasa models**

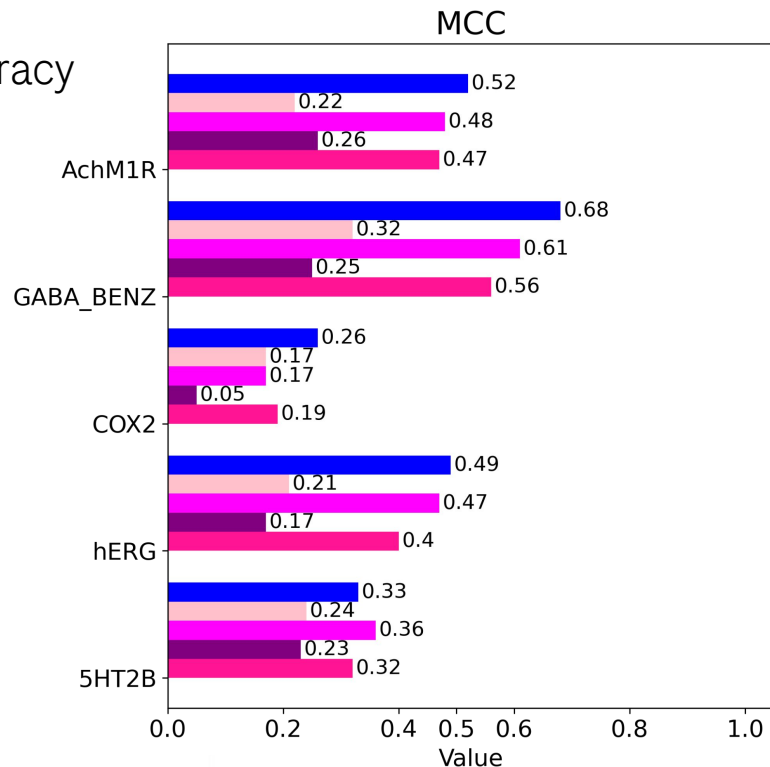
(usually SVC and XGBoost were best-performing)

Lhasa vs. internal ML models - validation on internal test set

Performance of Lhasa (SOHN) and Roche models is very similar

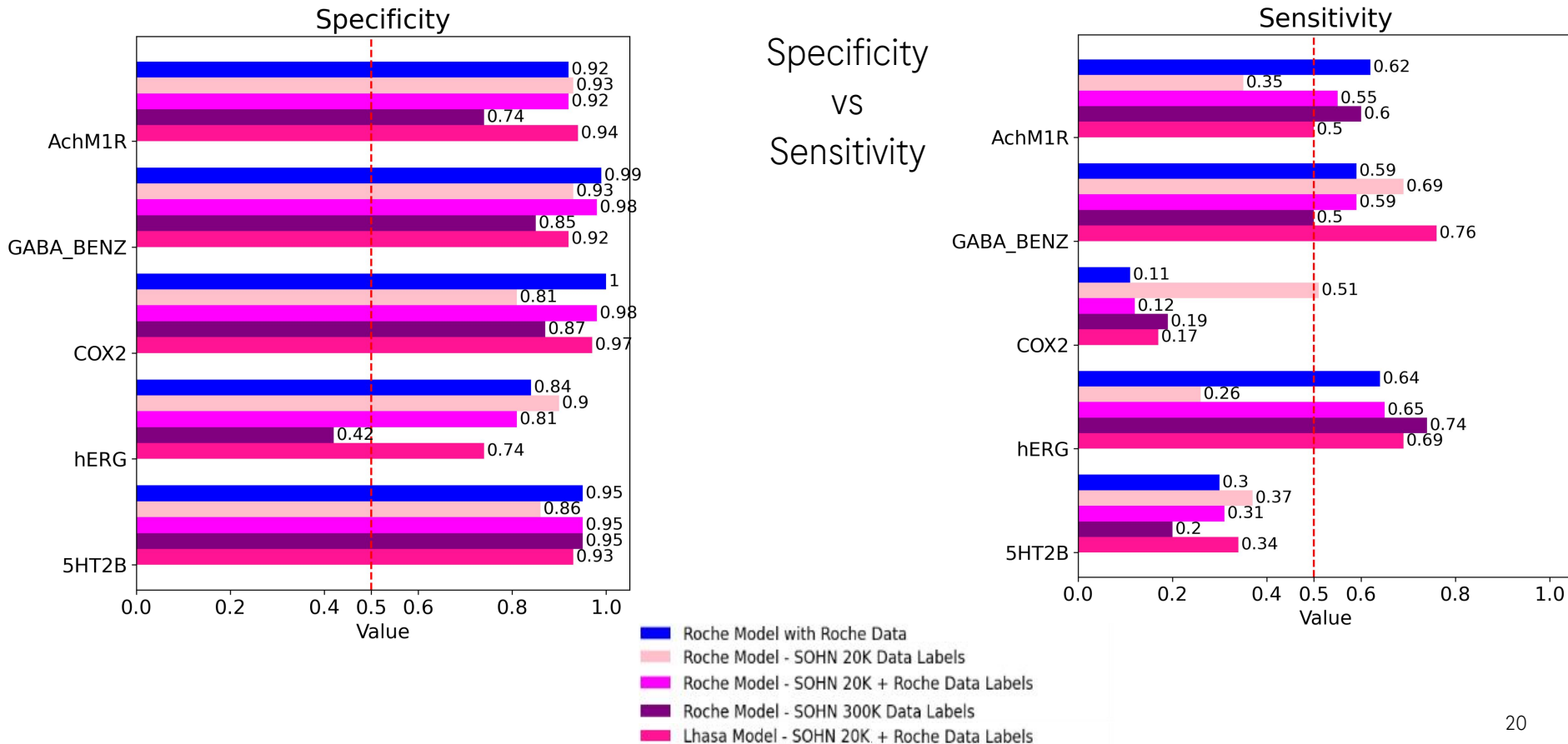


Balanced Accuracy
VS
MCC



Lhasa vs. internal ML models - validation on internal test set

Also in this case, the performance of Lhasa (SOHN) and Roche models is very similar...



Conclusions and discussion

Advantages of Effiris

- Uses knowledge from other industry partners (provided that they join Effiris)
- Larger training sets
- Increased Applicability Domain (AD)
- New models for ca. 50 targets
- Models being regularly updated with new member data (uncertain update schedule over time)
- Maintenance @ Lhasa (Exception: model updates, integration in to internal systems)

Limitations and Challenges

- Majority of training set consists of predictions → Approach is more technical than toxicologically-/biologically-oriented
- Target-dependency of performance: reduced confidence in models not validated with internal data
- As of now: only classification models are available (regression models planned)
- Data produced with different assay protocols (e.g. different temperature for hERG)
- Input from other companies conditional to their participation in Effiris
- FLulD dataset could be more tailored towards drug-like space

Summary

- Modelling secondary pharmacology data has several challenges
- Student-Teacher approach to leverage data from different sources without sharing sensitive information
- Model performance
 - Federated learning shows applicability domain increase
 - Lhasa model performance (SOHN & MLP) very similar to internal models
 - Target-dependant performance change, hard to transfer validation results to other models
- More tailored FLuiD dataset towards drug-like space could potentially increase performance of federated learning

Acknowledgements

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Doing now what patients need next