

Diagnostic Expert Advisor: Progress

- Last experiments were conducted with Horowitz only. A new model was trained per patient.
- Now all available data is utilized and a global model is trained.

Previously	Now
Only Horowitz Index	All available timeseries data
No static data	Static data included (Age, Sex, Weight, ...)
No Surprise Loss	Surprise Loss included
New model per patient	Consistent model trained on all patients

Diagnostic Expert Advisor: Setup and new Results

- Cohort:
 - Roughly 1.000 ARDS patients
 - Small cohort to limit training times
- Prediction:
 - 70%/10%/20% split for training, testing and validation
 - Prediction Horizon up to 12 hours
- Temporal Fusion Transformer (TFT):
 - Multi-Horizon, Uncertainty Modeling, Flexible history length, ...
 - Easy to use data format



Diagnostic Expert Advisor: Setup and new Results

Model Name	RMSE	MAE
Linear Regression	78	71
ARIMA	55	45
Facebook Prophet	301	266
Neural Prophet	70	63
Autoregressive Deep Learning	76	66
TFT Hyperparameters tuned, transfer learning no static data, no SL	68	40
TFT (preliminary) transfer learning static data, SL	54	<u>25</u>

Old Results

New Results

Diagnostic Expert Advisor: Setup and new Results

- Observation:

- MAE improved with ML Model using all available data
- RMSE barely changed

Model	RMSE	MAE
ARIMA	55	45
TFT	54	<u>25</u>

- Interpretation:

- Models focuses on fitting to stationary parts of Horowitz
- We should focus on the phase transitions!
- The improvement is misleading (we mostly improve stationary), but we can trust that the method can learn effectively!
- *Now we need to teach it to learn the right thing!*

Diagnostic Expert Advisor: Next Steps

- Split prediction task into two phase:
 - Phase I: Predict volatile regions
 - Phase II: Predict the direction of change
- Leaving out the stationary parts for the training of Phase II and changing the prediction task for Phase I
- The two phases can both be handled by the existing, known to be working, model architecture, altering the target functions and input data

Diagnostic Expert Advisor: Hardware Requirements

- Approximations based on current experiences training/running the DEA
- Training/Development mostly done with ~1.000 patients, as using all available 18.000+ patients increases training times to days even on the Cluster (10+ 24-core machines, Tesla V100 GPUs)
- Without optimization (e.g. SL approximation) running the model for a hundred beds can quickly require around 80 CPU hours to complete

Diagnostic Expert Advisor: Hardware Requirements Summary

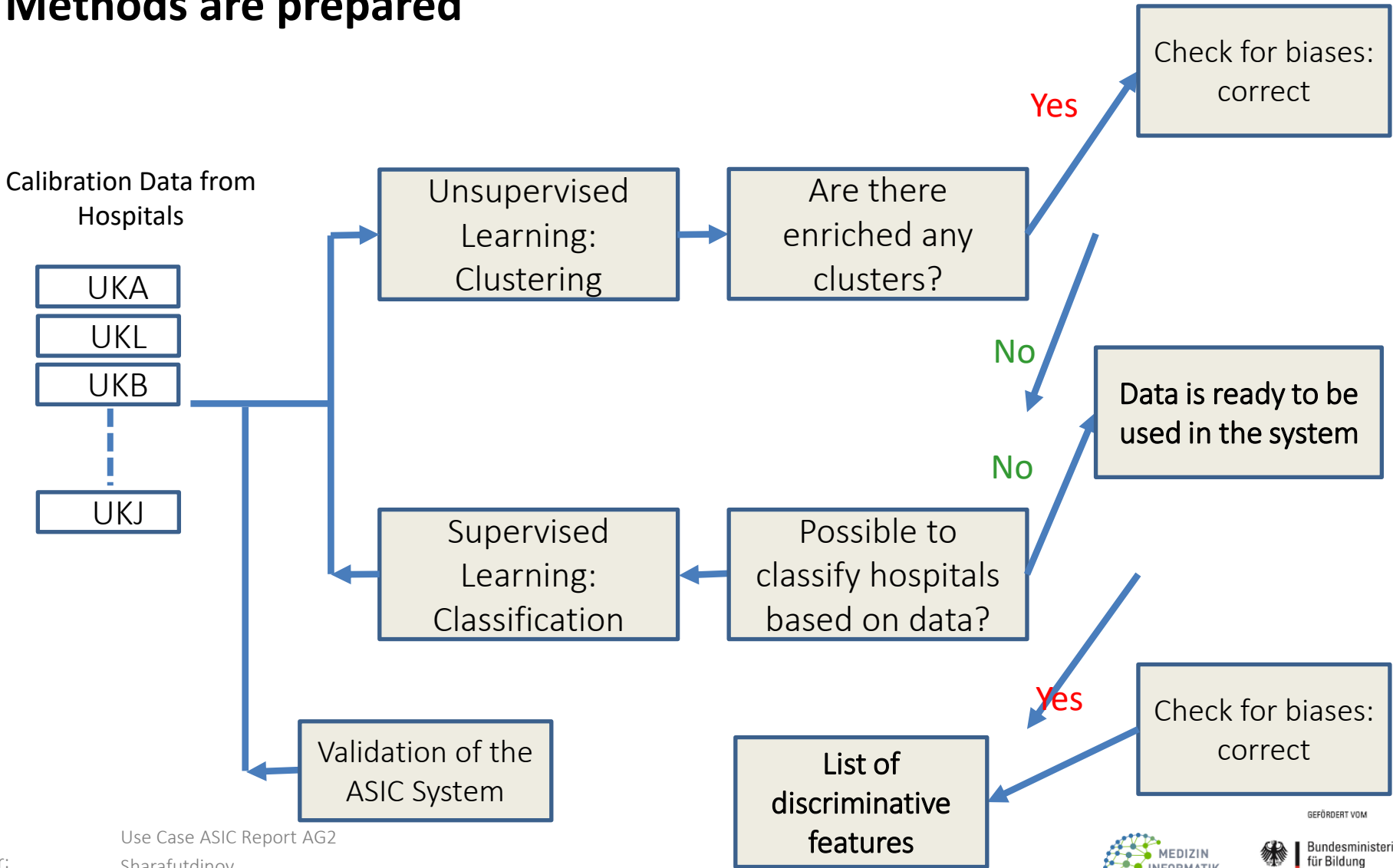
- Optimizations (e.g. ablation studies) can bring down required resources for inference (in-hospital prediction)
- Surprise Loss is computationally expensive and needs to regularly be updated
- Depending on how frequent the predictions should be updated this step alone would require multiple dedicated state-of-the-art workstations
- These estimation don't yet factor in any kind of local training, which would increase the resource requirements even more.

Calibration data

- Data available from 4 sites (5th site - only 150 patients)
- Sample sizes:
 - uk_00 - 13068 patients
 - uk_01 - 1024 patients
 - uk_02 - 2977 patients
 - uk_03 - 1367 patients
- Overall 18000 MV patients satisfying inclusion criteria for the ASIC use case
- Still a lot of opened questions for some of sites --> reminders have been sent on 26.04.2021

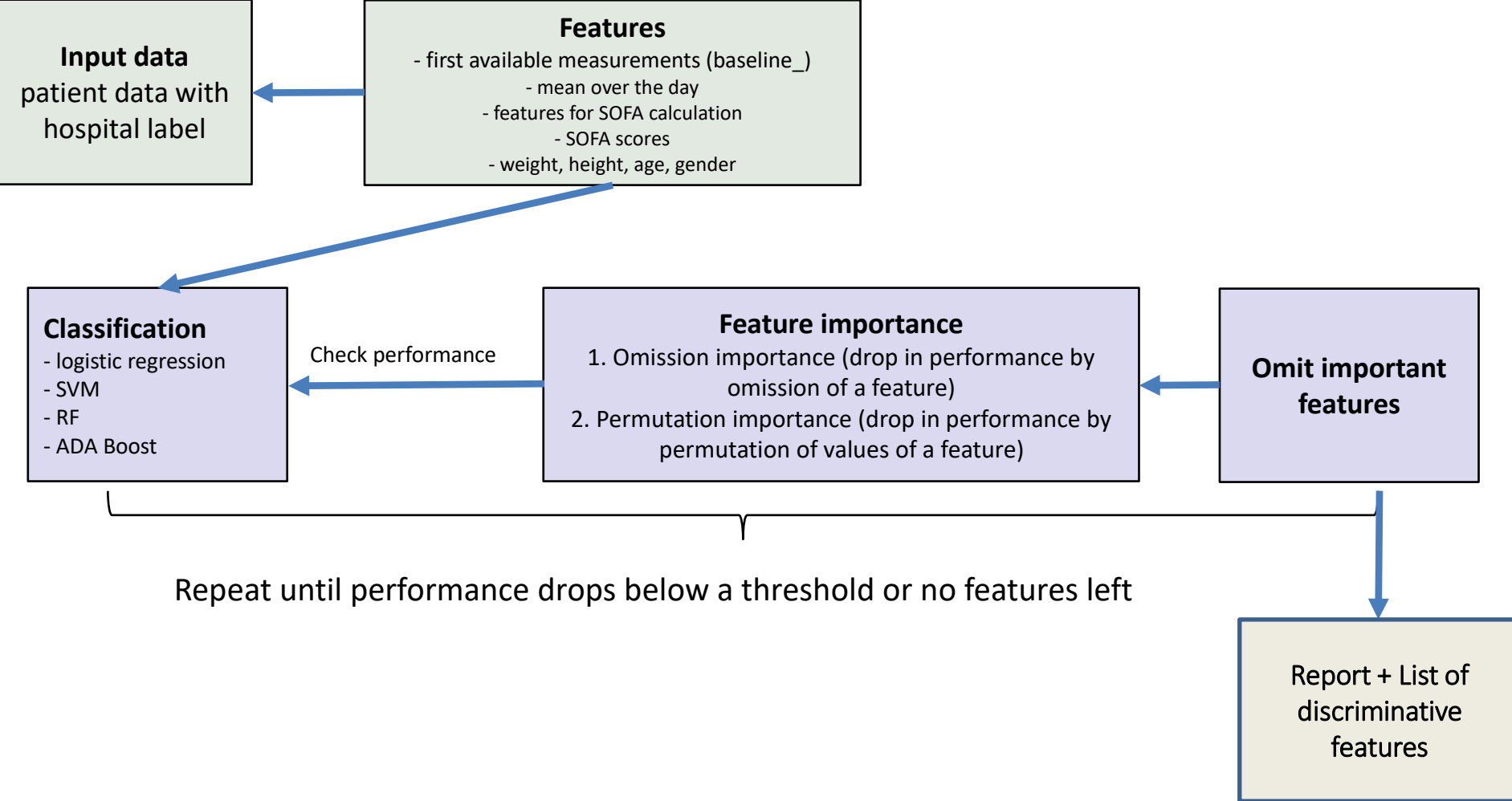
Calibration data: Strategy

- Methods are prepared



Calibration data: Supervised learning

- Methods are prepared



Calibration data: Convex hull analysis

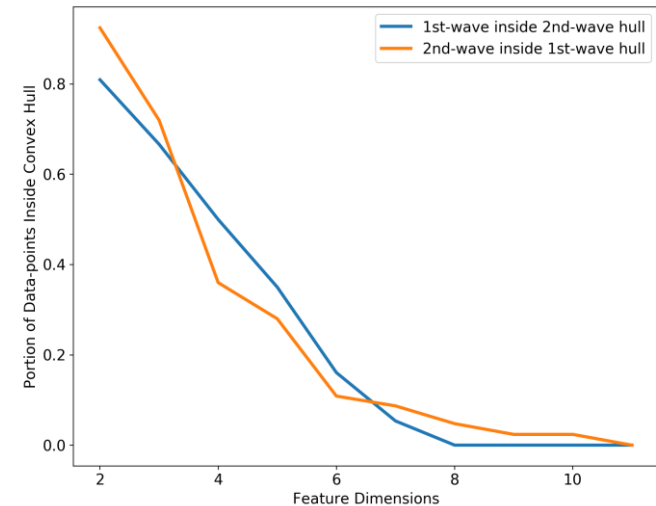
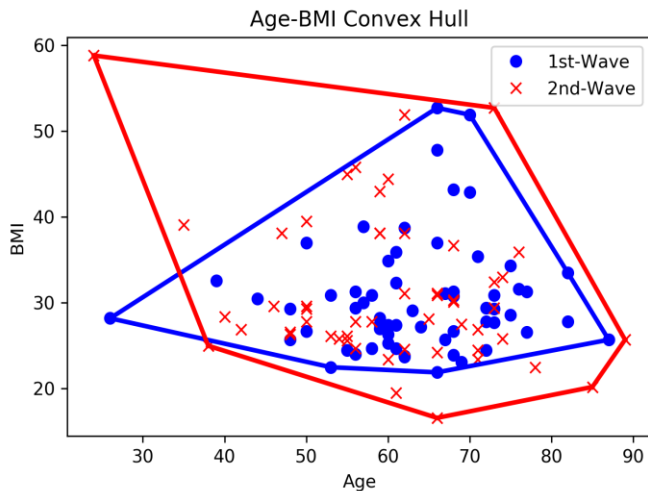
- the smallest convex set enclosing all data points
- data-driven method can make predictions inside the convex hull of the given training data
- transfer learning is possible if **data of new hospital is inside of the convex hull of research data repository**

Convex hull of COVID-19 Data

- 1st and 2nd wave MV COVID-19 patients consisting of 63 and 54 patients
- for low-dimensional feature spaces, e.g. 2D Age and BMI, convex hulls cover each other

Convex Hull in Higher Dimensions

- for higher dimensional feature spaces, the convex hulls of 1st and 2nd wave data get separated
- results are shown for the most important features for mortality classification
- larger data sets are needed for a classification task using data-driven models



Calibration data: next steps

- Final curation after feedback from the sites
- Parameters for calibration found
 - test with other features/on different cohorts (MIMIC/ICCA, COVID, etc..)
- Decide how to treat respective parameters
- Convex hull analysis of available calibration data - general tool for population comparison and transfer learning

Parameters to optimize for

1. Define an initial state of the VP with **measurable global parameters** of the real patient:

	CO	FatmO2	Hb	IE	PaCO2	PaO2	PEEP	Temperature	VentRate	Vtidal	HCO3a	Bea	scPressure	SaO2	pHa
1	10000.0	0.500000	143.755008	0.343615	3.946272	10.652268	8.582611	36.312500	17.041667	505.296	20.4	-2.45	24.675008	93.075	7.4325

2. Optimization parameters: (anatshunt, RQ, VO₂, m₁, m₂, m₃, alpha, c₁, c₂)

Non-measurable global parameters:

- Anatomical Shunt
- Respiratory quotient
- Metabolic rate of O₂ in the body

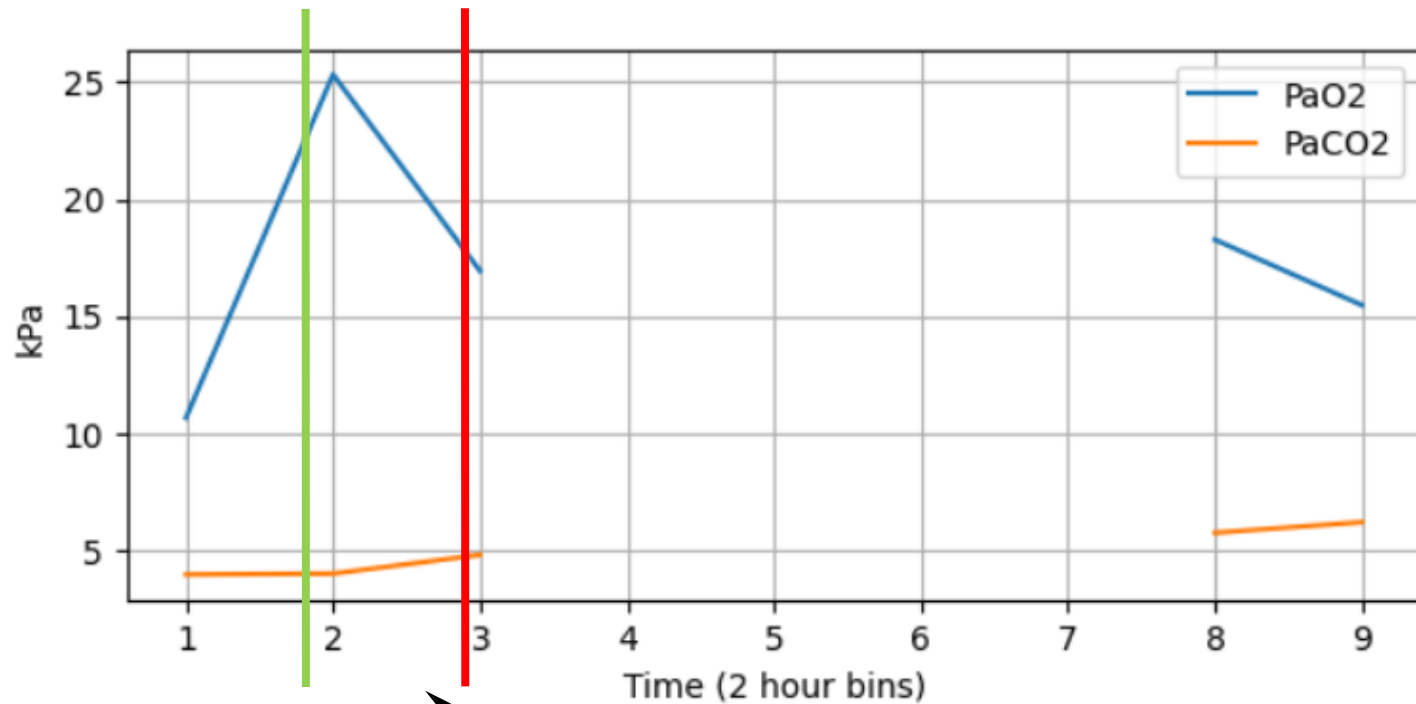
Non-measurable parameters reflecting distributions of compartmental parameters:

- # of closed compartments (controlled through Pext - alpha closed, 1-alpha opened compartments)
- m₁, m₂, m₃
- where $R_{ards} = m_1 * N + c_1$, $VR_{ards} = m_2 * N + c_2$, $STalv_{ards} = m_3 * STalv_{healthy}$

3. Objective function:

$$\min_{\text{params}} ((PaO2_{\text{real}} - PaO2_{\text{sim}})^2 + (PaCO2_{\text{real}} - PaCO2_{\text{sim}})^2)^{0.5}$$

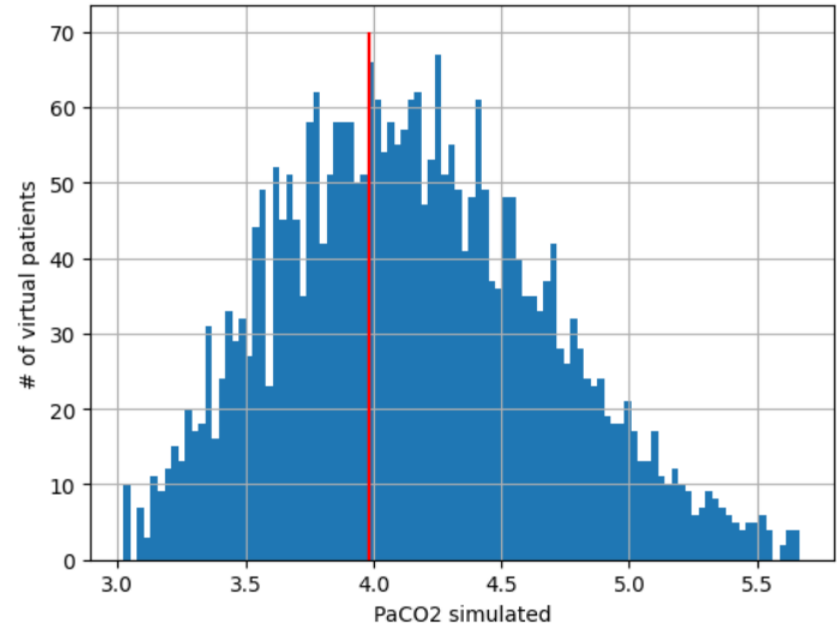
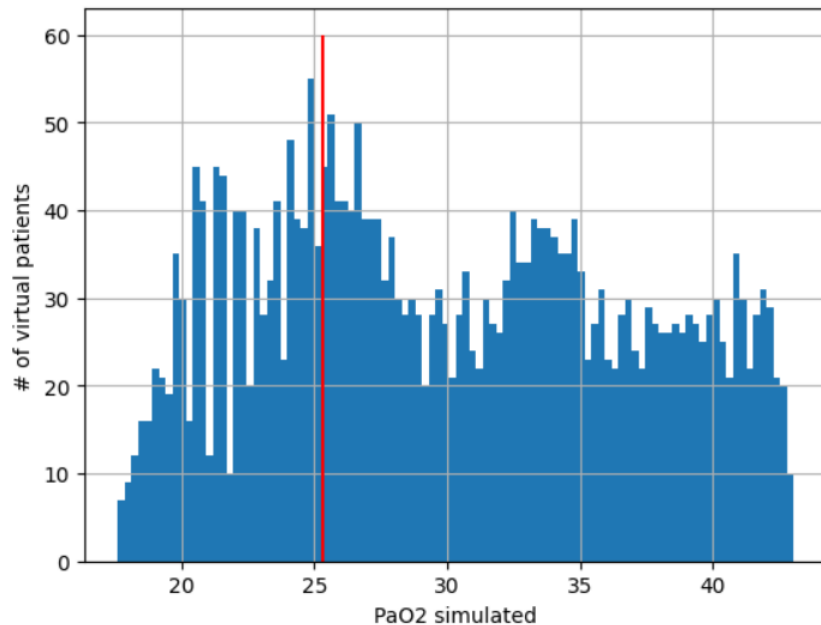
Objective function: $\min_{\text{params}} ((\text{PaO2}_{\text{real}} - \text{PaO2}_{\text{sim}})^2 + (\text{PaCO2}_{\text{real}} - \text{PaCO2}_{\text{sim}})^2)^{0.5}$



Simulation for the next 2 hours

- Potentially for every time step we can find a VP configuration, which suits
- Multiple time points increase confidence for a particular VP configuration
- Final configuration defines the VP

- Currently defined a cohort of 3000 VP based on one real patient and varying optimization parameters
- Investigated whether simulations cover values of a real patient
 - A lot of VPs, which showed similar behaviour --> not unique solution
- Next step: implement objective function and perform optimization algorithm



Issues:

- Ill-posed problem (solution is not unique)
- Multiple time points - multiple configurations possible
 - Dynamic configuration - parameters of VP do change with time
 - Sort optimization parameters based on the time scale for changes (i.e. Shunt changes slowly - should be fixed, N closed compartments can change)

Work in progress:

- Objective function, which parameters to choose
- Define final set of parameters for optimization
- Choose and implement an optimization algorithm

Steps:

- Prepare a dedicated environment (storage, processors) for source data and for outputs.
- Set up platform to run scripts.
- Implement multirun matlab script on Cluster.
- Prepare outputs for model training.

The screenshot displays the JupyterLab interface. On the left, a file explorer window shows the directory structure of a project. A red box highlights the following files and folders:

Name	Last Modified
mimic	3 months ago
outputs	3 months ago
Pulmonary_Simulator	25 days ago
length.csv	7 months ago
NaNmeans.csv	7 months ago
patients_list.csv	6 months ago
simulation.ipynb	a day ago

The main area of the interface is the 'Launcher' grid, which contains various application icons. A red box highlights the 'Octave-6.1.0' icon, which is used for running MATLAB scripts. The launcher grid includes icons for Python 3, Bash, C++17, gpu_kernel, Javascript (Node.js), Julia 1.5.2, Octave-6.1.0, PyDeepLearning-1.0, PyParaView-5.8.1, PyQuantum-1.1, R-4.0.2, Ruby 2.7.1, and Xpra Desktop [~].

Roadblocks:

- Since no Matlab implementation exists on the cluster we try to adapt the available Octave implementation.
- Had to import several packages that weren't originally available.
- Made changes to the original script to adapt it to these packages.

```
[3]: run('Pulmonary_Simulator/scripts/multi_run/multiRun_main.m')

warning: genvarname is obsolete; use matlab.lang.makeValidName or matlab.lang.makeUniqueStrings instead
T = dataframe with 24 rows and 28 columns
Src: ../input_table/patients_ID.csv
_1 MATLAB_NO          ID anatShunt chPressure   CO ColVol FatmO2   Hb   IE  Ncomp PaCO2  PaO2  PEEP  RQ run_time scPres
sure Temperature VDphys VentRate  VO2 Vtidal          comp_ID          P_ext          Rcomp          ST_alv          Stick_t
ime_rand             TOPcomp          VRcomp
double double double double char double double double double double double char char double double double double
char char char char char char char char char char char char char char char char char char char char char char
1 1 37.200 50 12.253 294.25 A 0.010000 1 11100 6 0.8000 105 0.2759 100 - - 5 0.6000 120
20 A 37.200 50 12.253 294.25 A 0.010000 1 11100 A 6 0.8000 105 0.2759 100 A - - 5 0.6000 120
3 2 37.200 60 15.800 300.00 B 0.080000 1 7200 6 0.9000 108 0.2500 100 - - 5 0.6500 120
20 B 37.200 60 15.800 300.00 B 0.080000 1 7200 B 6 0.9000 108 0.2500 100 B - - 5 0.6500 120
4 3 37.200 60 12.000 250.00 H 0.020000 1 5000 6 0.2100 160 0.3300 100 - - 5 0.8000 120
20 H 37.200 60 12.000 250.00 H 0.020000 1 5000 H 6 0.2100 160 0.3300 100 H - - 5 0.8000 120
5 4 37.200 50 12.253 294.25 A 0.010000 1 11100 6 0.8000 105 0.2759 100 4.1 13.95 5 0.6000 120
20 A 37.200 50 12.253 294.25 A 0.010000 1 11100 A 6 0.8000 105 0.2759 100 A 4.1 13.95 5 0.6000 120
6 5 37.200 60 15.800 300.00 B 0.080000 1 7200 6 0.9000 108 0.2500 100 4.1 13.95 5 0.6500 120
20 B 37.200 60 15.800 300.00 B 0.080000 1 7200 B 6 0.9000 108 0.2500 100 B 4.1 13.95 5 0.6500 120
7 6 37.200 60 12.000 250.00 H 0.020000 1 5000 6 0.2100 160 0.3300 100 4.1 13.95 5 0.8000 120
20 H 37.200 60 12.000 250.00 H 0.020000 1 5000 H 6 0.2100 160 0.3300 100 H 4.1 13.95 5 0.8000 120
8 7 37.200 50 12.253 294.25 A_permuted 0.010000 1 11100 6 0.8000 105 0.2759 100 4.1 13.95 5 0.6000 120
20 rmutd 37.200 50 12.253 294.25 A_permuted 0.010000 1 11100 A_permuted 6 0.8000 A_permuted A_permuted A_permuted A_permuted A_permuted A_permuted A_permuted A_permuted
8 8 37.200 60 15.800 300.00 B_permuted 0.080000 1 7200 6 0.9000 108 0.2500 100 4.1 13.95 5 0.6500 120
20 rmutd 37.200 60 15.800 300.00 B_permuted 0.080000 1 7200 B_permuted 6 0.9000 B_permuted B_permuted B_permuted B_permuted B_permuted B_permuted B_permuted B_permuted
```

Result:

Next Steps:

- Performance analysis on the cluster:
 - Single run duration in JupyterLab.
 - Multi-run duration.
 - Adapting scripts to the cluster.
- Tweaking and testing parallelisation methods.

Thank you for your attention!