

# FinOMOP Swarm Learning: Deep learning for patient-specific modelling of Acute Myeloid Leukemia based on longitudinal clinical laboratory data and OMOP

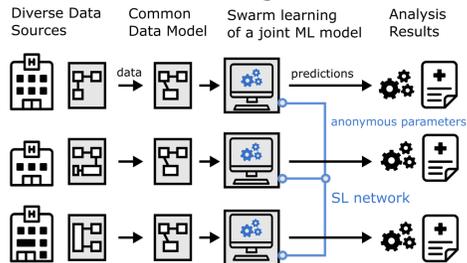
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## INTRO:

- Deep learning **is powerful** but data hungry!
- Data sharing hindered by privacy concerns
- Federated learning hindered by heterogeneous data formats
- **Solution:** Use common data model and learn models *at the edge* without sharing source data using Swarm learning (SL)

## METHODS

### 1. OMOP + Swarm Learning => Joint model



### 2. Cohort: AML use case

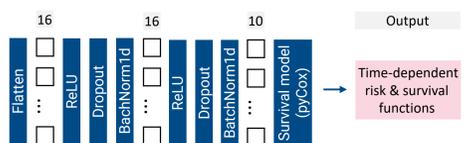
All patients with AML diagnoses and at least 3 blast measurements within 21 days of diagnosis

### 3. Data

Endpoint: Overall patient survival  
Features: Blood count measurements, timepoints up to 21 days after diagnosis

### 4. Model architecture v1:

pyTorch, pyCox, DeepHit



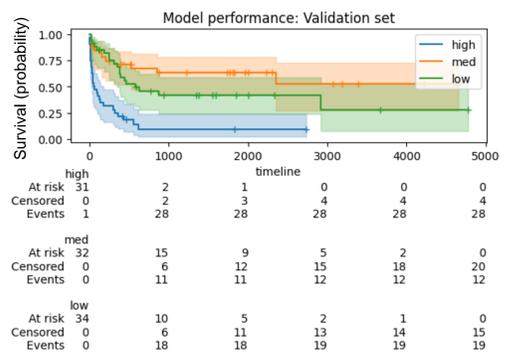
## RESULTS

### Developed FinOMOP-SL framework:

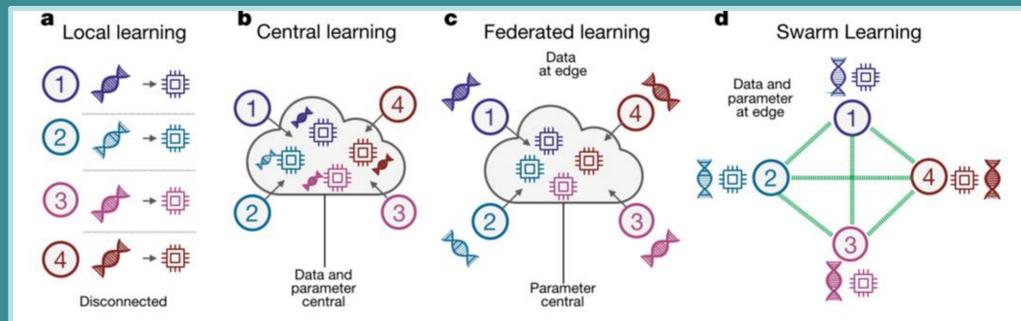
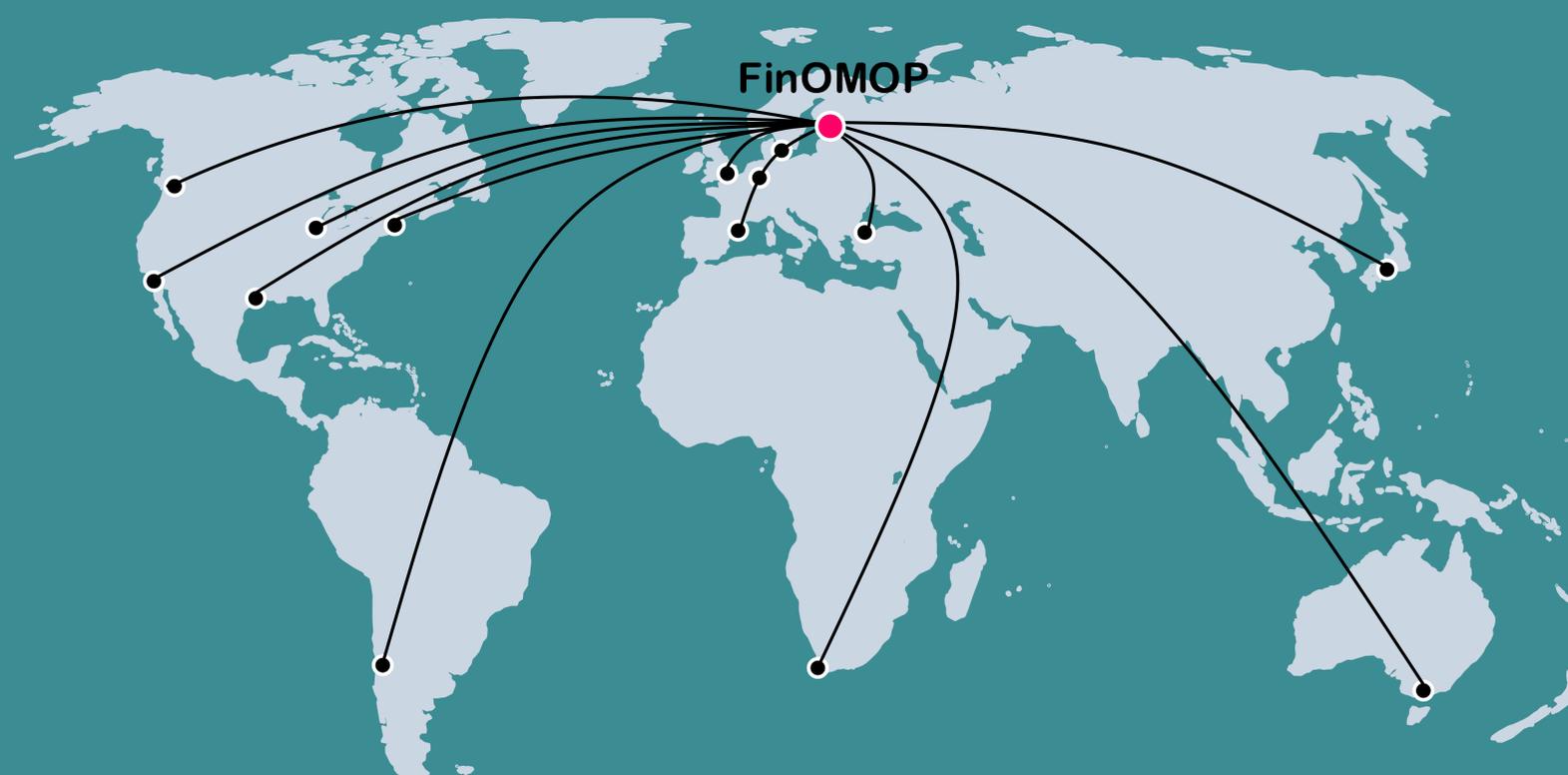
- Findata compliant implementation at three sites: **Helsinki, Turku, Tampere**
- Integrating OMOP & swarm learning

### Proof of concept: use case AML

- Predictive power model v1, so far only trained on HUS data::



# Build deep, predictive models for precision medicine together in global networks.



- Limited dataset  
- Bias  
- Low accuracy  
- Disconnected

+ Enlarged dataset  
+ Better accuracy  
- Aggregation  
- Monopolisation

+ Enlarged dataset  
+ Better accuracy  
+ No data movement  
+ No central custodian  
+ Shared insights

## AMMO BAR

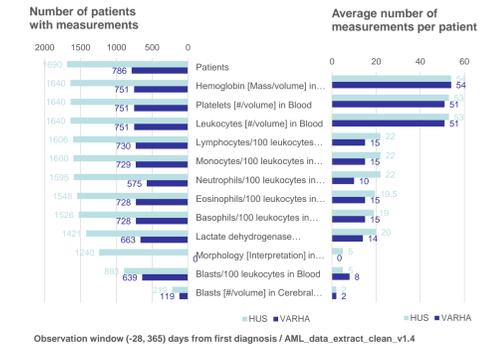
### CONCLUSIONS

- Routine short-term follow up data – blood count measurements (21 days) – can predict long-term prognosis in AML
- Federated/Swarm learning of a joint model feasible
- Routine data available at all three sites

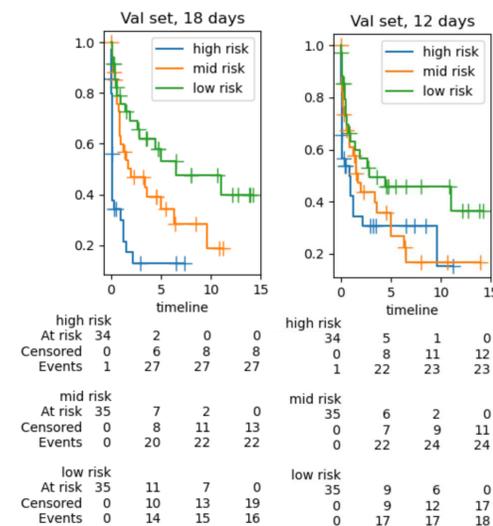
### Next:

- Estimate and include pCR & RFS
- Integrate genomic data
- Advanced model architectures, e.g. autoencoders + survival models

### Data availability HUS & VARHA (Turku)



### Performance based availability of timepoints



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