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Federated Learning - Introduction

ISMB 2022, Federated Learning in Biomedicine (Tutorial)

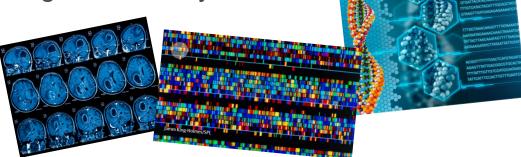


Large-scale AI in Biomedicine

 Machine learning (ML) allows for classification and predictions in different domains

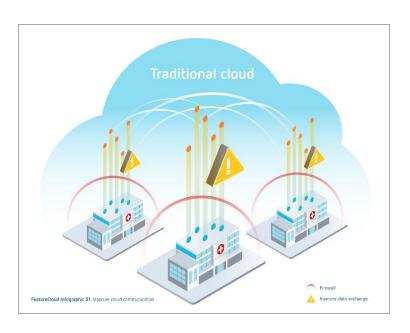
Benefits hugely from larger quantities of data

Amount of data is growing continuously





Large-scale Al in Biomedicine - Classical Approach



- Merge data into big datasets
- Train ML models on the whole set
- Share models with data contributors



Problem: Privacy regulation



- General Data Protection Regulation (GDPR) and others
- Patients must have control over their data
- Data cannot leave the hospitals without big legal efforts



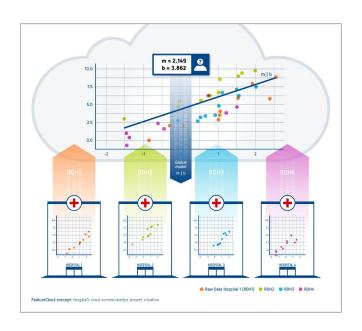
Idea: Federated Learning

- Data remains where is has been collected
- Split training between local training and global aggregation

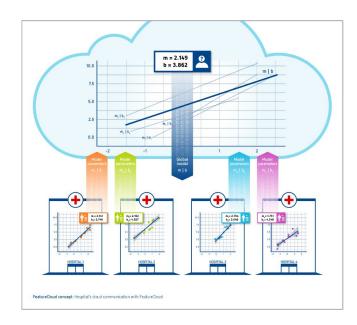
"Bring the model training to the data, not the other way round"



Large-scale AI in Biomedicine - Classical Approach









Federated Learning Platforms



- Isolate data
- Impose restrictions on communication
- Provide assistance for development of federated applications



What's the price?

- Loss of performance due to communication overhead
- Loss of accuracy (in most cases) due to higher difficulty of generalization
- Less control over data quality



Is Federated Learning privacy-preserving?

- Depends on the method and the data
- Privacy leaks can still occur (e.g., due to overfitting)
- But: huge improvement over sharing raw data
- Can be enhanced with additional techniques
 - Differential privacy
 - Secure multiparty computation
 - Homomorphic encryption

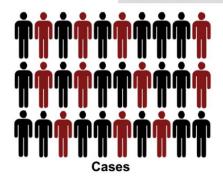


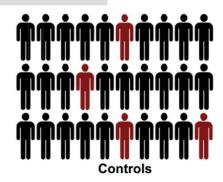
Application example: GWAS

- Case/control studies on SNP data
- Each SNP examined individually (LR)
- Find SNPs related to diseases

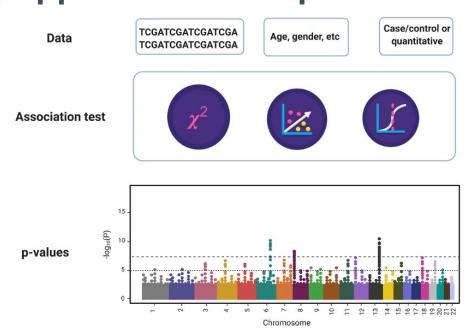
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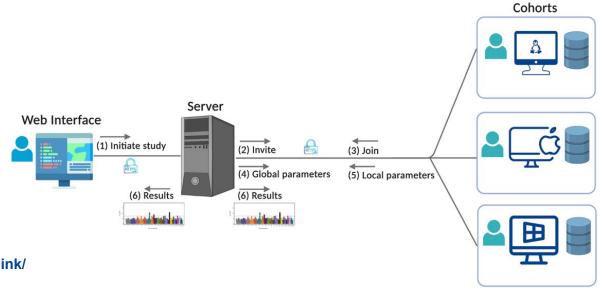






- Chi-square test
- Linear regression
- Logistic regression

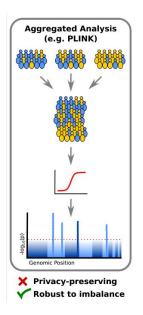


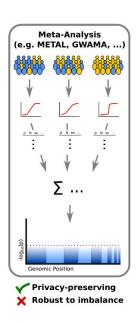


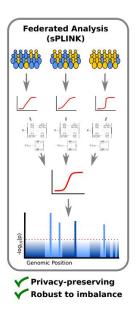
URL: exbio.wzw.tum.de/splink/

Reza Nasirigerdeh et al. (published in Genome Biology)

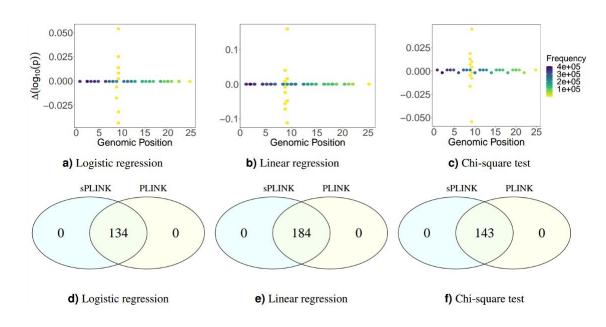














Other examples

Flimma

Differential expression analysis based on the limma voom pipeline

URL: exbio.wzw.tum.de/flimma/ | Olga Zolotareva et al. (published in Genome Biology)

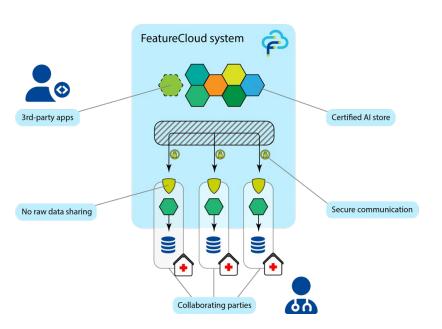
Partea

Time-to-event studies

URL: exbio.wzw.tum.de/partea/ | Julian Späth et al.



Platform example: FeatureCloud



- Increase convenience for developers (including testing and deployment)
- Increase trust in apps (certification, consent management)
- Create a community (developers, researchers, patients)

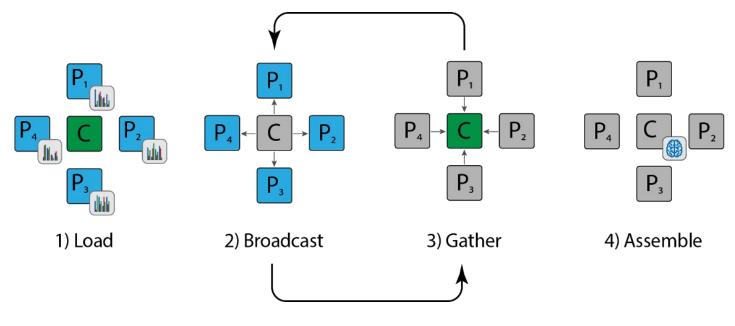


App communication

- Star-based federated learning
 - Gather-broadcast
- Peer-to-peer
- Secure multi-party computation

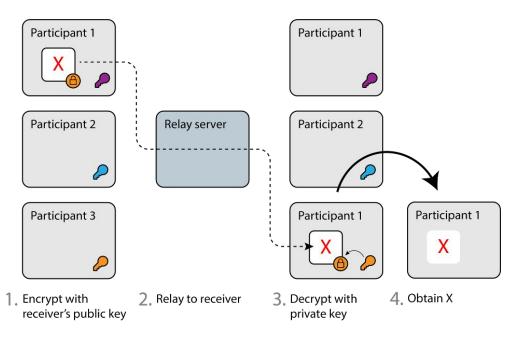


Star-based communication





Peer-to-peer communication





Continuous vs. discrete models

- Continuous models can usually be federated 'straightforward'
 - Gradient descent
 - Federated average
 - One or multiple interactions
- Discrete models are more difficult
 - Example: Decision tree
 - Possible solutions: Ensemble techniques



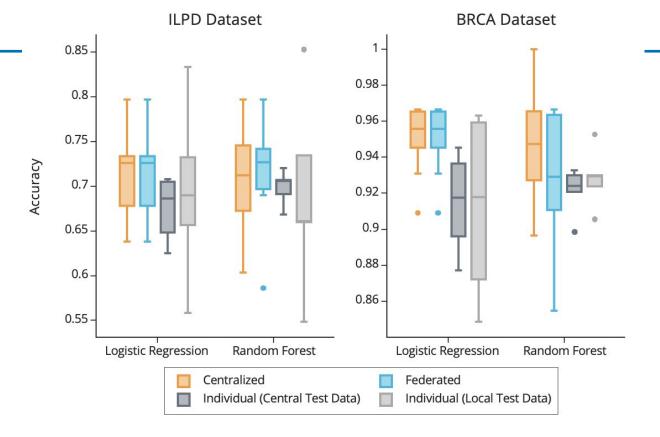
Example: Random forest

- Train decision trees using bootstrapping on each site
- Transmit models and merge into big ensemble → Random Forest

Federated Random Forests can improve local performance of predictive models for various healthcare applications, Anne-Christin Hauschild, Marta Lemanczyk, Julian Matschinske, et al., Bioinformatics, Volume 38, Issue 8, 15 April 2022, Pages 2278–2286

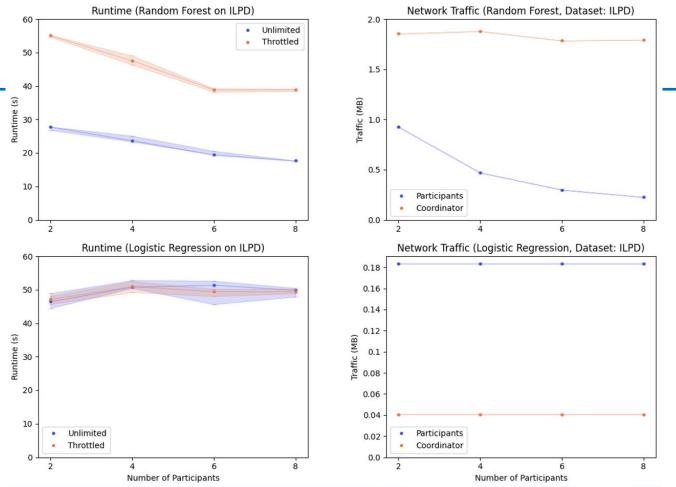


Evaluation





Evaluation





Thank you!

→ Questions, comments, thoughts?