



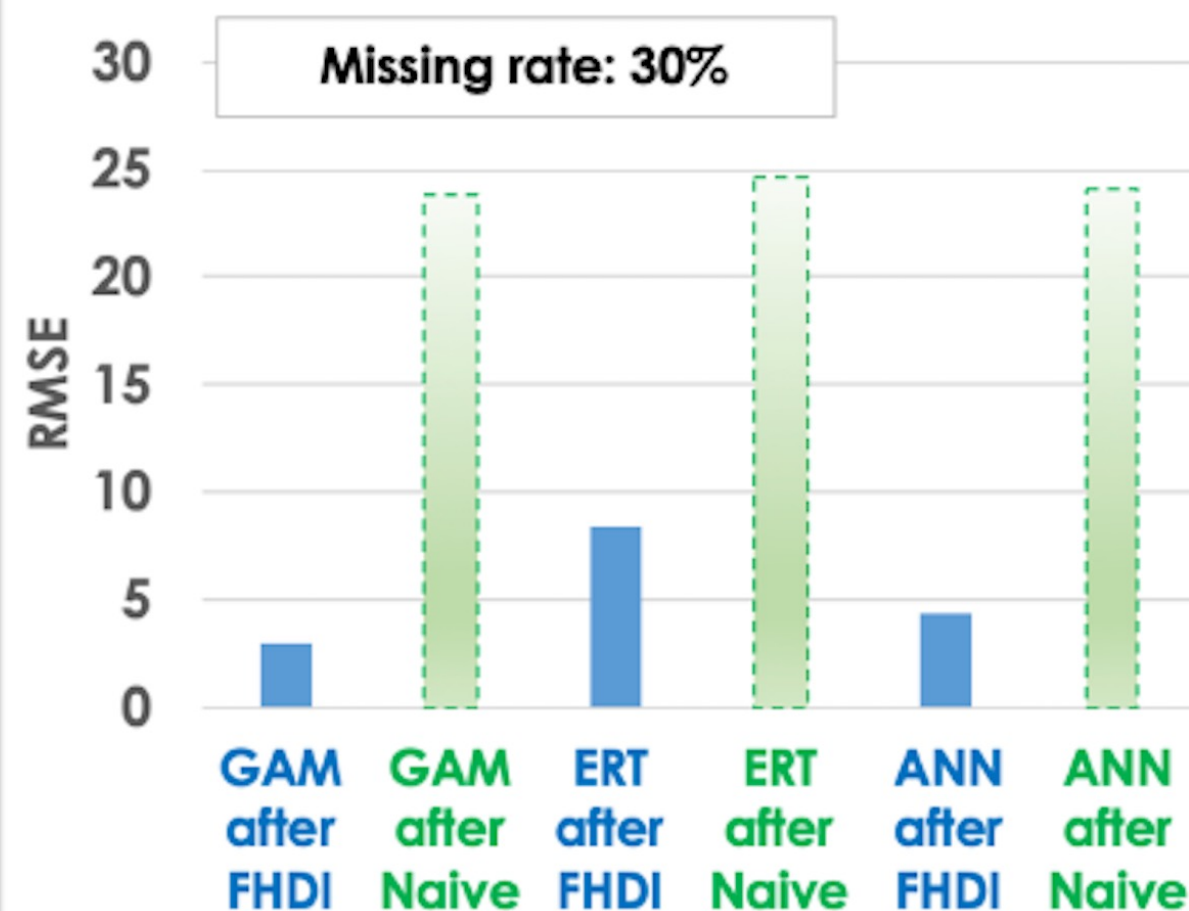
Award #: 1931380

# CSSI Elements: Development of Assumption-Free Parallel Data Curing Service for Robust Machine Learning and Statistical Predictions

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## Grand Challenges

- Incomplete data issue is everywhere in broad science and engineering
- Theories and methods of missing data curing (called “imputation”) is **limited to small data**
- **Naïve imputation** may substantially hamper the accurate machine learning (ML) and statistical learning (SL)-based predictions (see Fig below)
- **Lack of theories and software** for large/big incomplete data curing



**Fig. Positive impact of the proposed data curing method (FHDImputation) on statistical learning (SL) and ML predictions:** Generalized additive model (GAM); Extremely randomized trees (ERT); Artificial neural network (ANN). Root mean square error (RMSE) is shown.

## Research Objective

- Develop a new community-level **large data curing service** running on NSF Cyberinfrastructure (XSEDE) and local HPC
- **No restriction** of data sizes, types, high-dimensionality; No distributional assumptions or expert knowledge on data science required
- Pursue a purely data-driven imputation by developing the **ultra data-oriented parallel fractional hot deck imputation (UP-FHDImputation)**
  - Assumption-Free, General Data Curing; **Only Observed Data** Are Needed for Imputation (thus, “Hot-deck”)
  - Provide **“Cured” large/big data set** for convenient subsequent ML and SL

## Proposed Methods

Ultra Data-Oriented Parallel Fractional Hot Deck Imputation (**UP-FHDImputation**) [Ultra Data: Big-n & Big-p]

**[Step 0] Sure Independence Screening (SIS)**  
Selectively Done for **big-p** (high-dimensional) Data

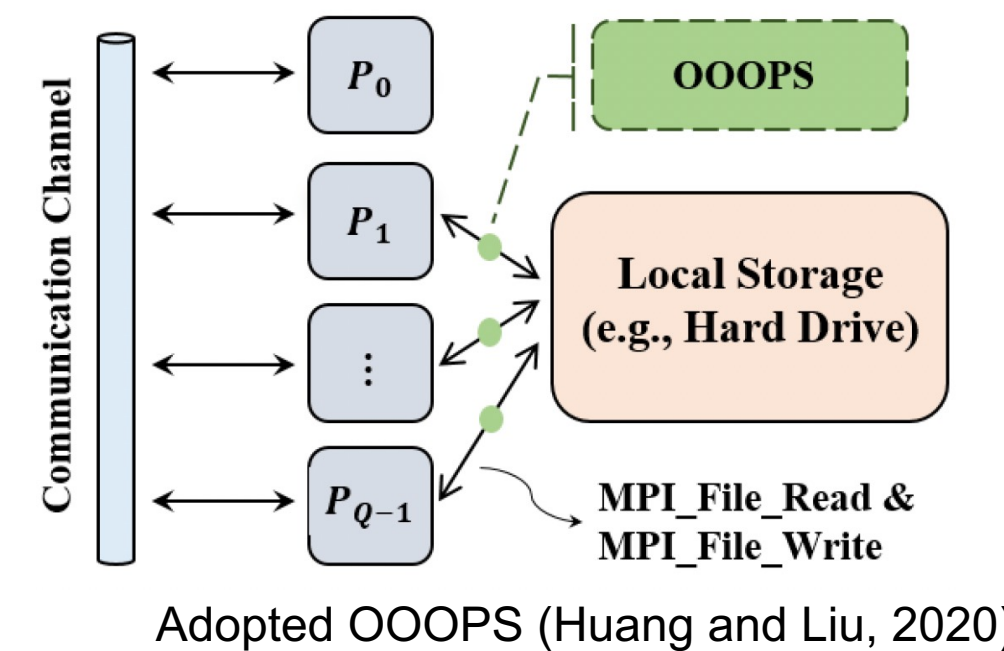
**[Step 1] Parallel Imputation Cell Construction**  
Hybrid Data (Continuous & Categorical)

**[Step 2] Imputation Cell’s Joint Probability**  
Parallelized Modified EM Algorithm

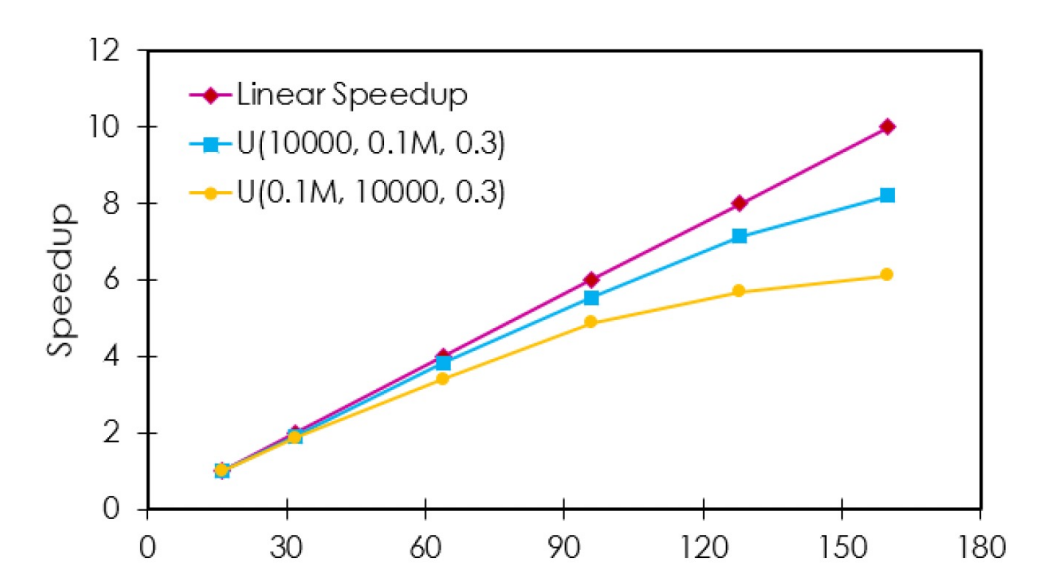
**[Step 3] Fractional Hot Deck Imputation**  
Parallelized Donor Selection with KNN

**[Step 4] Parallel Variance Estimation**  
Parallelized Jackknife & Parallel Linearization

## Results



Data: U(size, variables, missing rate)



**Fig. Scalability of UP-FHDImputation**

SD	UP-FHDImputation	Naive
3	0.052	0.062
5	0.088	0.100
8	0.134	0.160

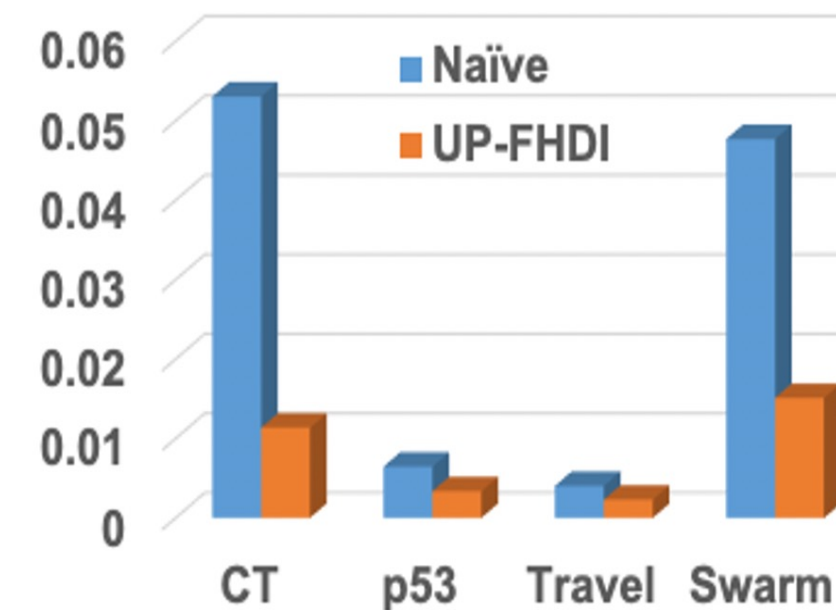
**Fig. Average Standard Error**

SD	UP-FHDImputation	Naive	Synthetic Big-p Data (n=10,000, p=0.1 million, Missing=0.3) with increasing randomness (SD) in data
3	0.077	0.081	
5	0.079	0.088	
8	0.080	0.092	

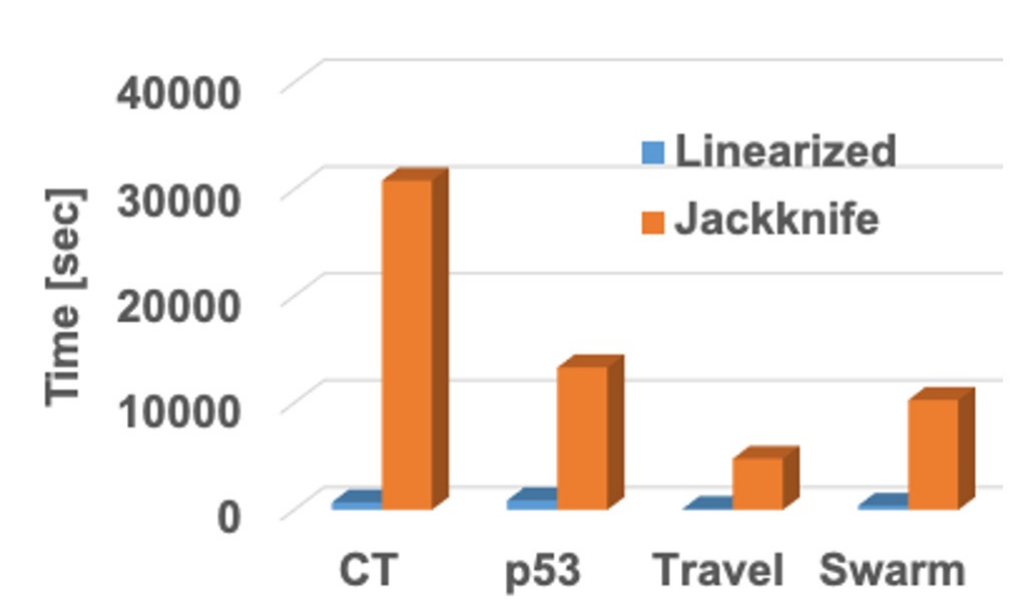
**Fig. RMSE**

Dataset name	Number of Instances	Number of Variables	Discipline	Data Source	Test Data Sets
CT [Graf et al. 2011]	53500	380	Medicine	UCI	
p53 [Lathrop 2010]	31159	5408	Genetics	UCI	
Travel [Gao et al. 2021]	23772	50	Transportation	IEEE Dataport	
Swarm [Abpeikar et al. 2020]	24016	2400	Biology	UCI	

Mean Abs Error



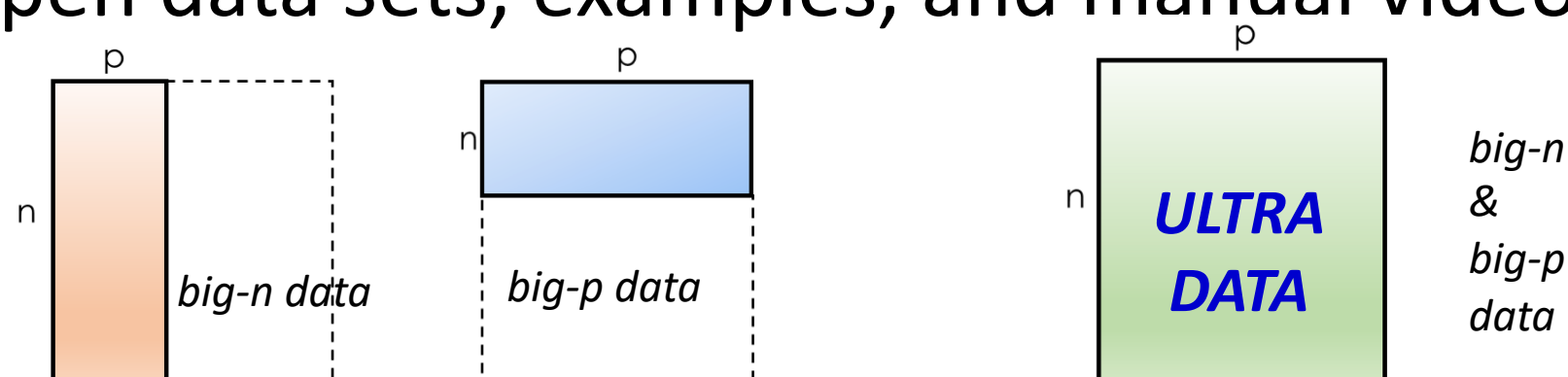
**Fig. Accuracy of UP-FHDImputation**



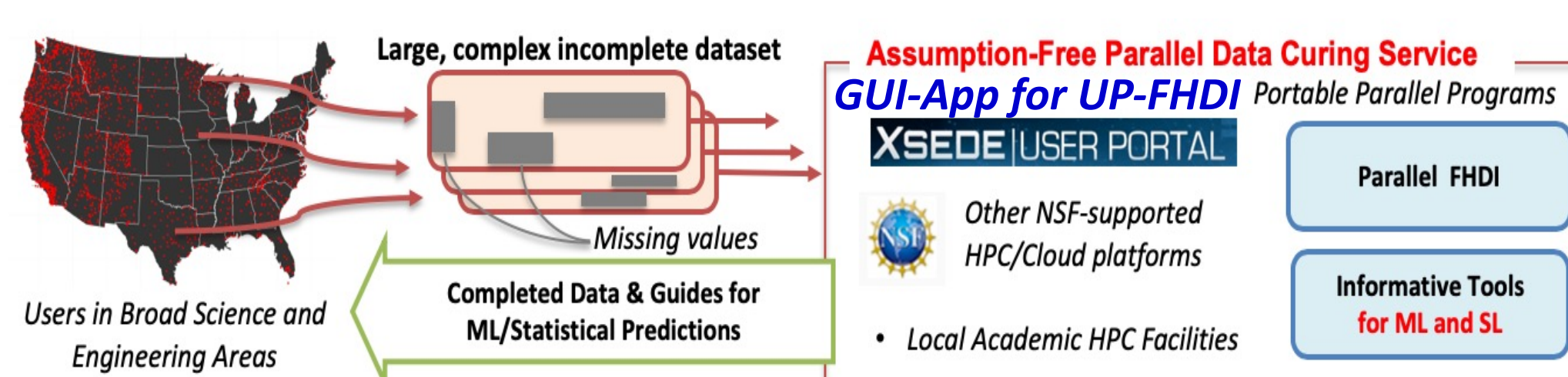
**Fig. New Variance Est. Method’s Efficiency**

## User-Friendly Service

- Deployment of the UP-FHDImputation on *NSF XSEDE*
- Graphical User Interface (GUI) for UP-FHDImputation
- Open data sets, examples, and manual videos



**Fig. UP-FHDImputation Can Tackle Three Data Types**



## Conclusions

- **UP-FHDImputation** has been developed for improving prediction accuracy of ML and SL with Ultra incomplete data (**up to millions of instances and 10,000 variables**)
- The program is deployable on NSF XSEDE and local HPC
- Serial version **R Package FHDImputation** available on CRAN

## References of the PIs

- Yang et al., 2022, *IEEE TKDE* (under 2<sup>nd</sup> review)
- Yang et al., 2020, *IEEE TKDE*
- Song et al., 2019, *IEEE TKDE*
- Im et al., 2018, *The R Journal*

## Acknowledgement

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