

# Digital Transformation of Healthcare

## Data Cleaning

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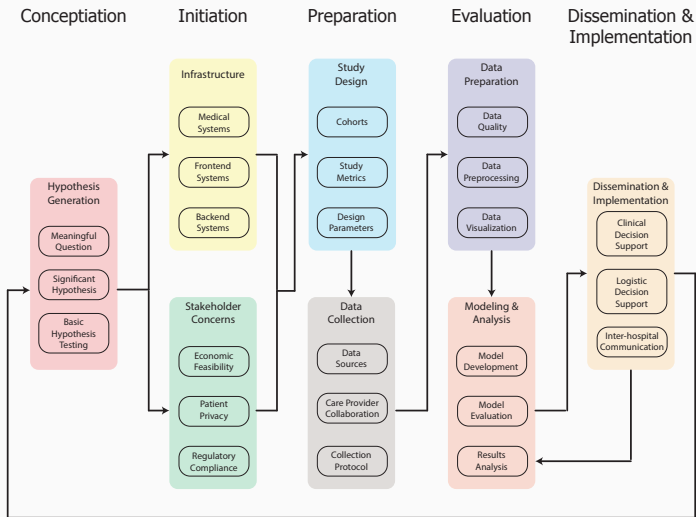
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Center for Health Data Innovations

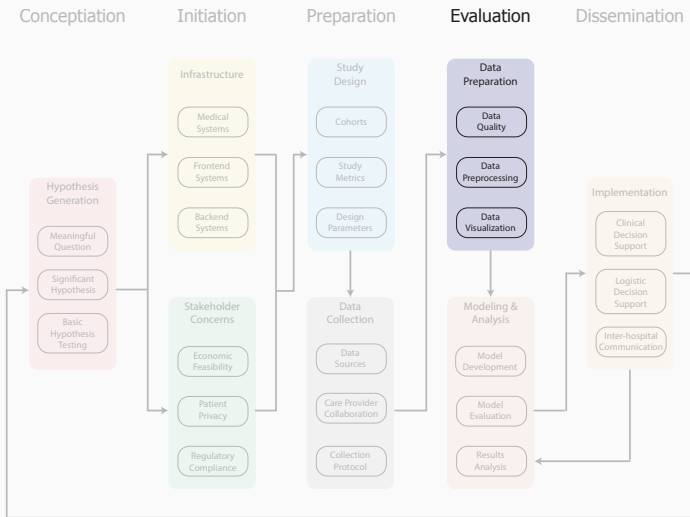
After this lecture students will be able to

- Discuss and apply the steps involved in cleaning data for modeling
- Design a process for the imputation of missing data
- Build a bioinformatics pipeline starting from given data

# Bioinformatics Pipeline

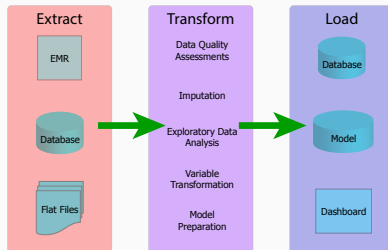


# Data Cleaning



# Scenario

- You are trying to optimize the use of pain medication regimens for pediatric sickle cell patients
- You have just collected all the data on all peds hem-onc patients on the floor for the past month.
- How can you systematically assess, organize and prepare the data for modeling and analysis?
  - What are the steps to transform raw data into a usable format



# Imputation and Extrapolation

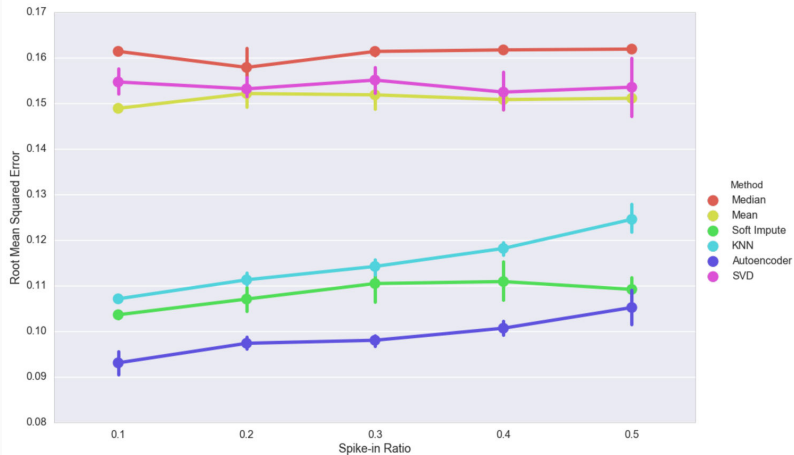
- what are the different reasons why data might be missing
- What are the different ways that data could be missing
- Can we develop a systematic way to deal with missing data
  - pain score
  - pain medication usages
  - retic count
  - infection status
  - imaging results
- How do you evaluate imputation

## Missing Data Imputation in the Electronic Health Record Using Deeply Learned Autoencoders

- Researchers started from an ALS clinical trials database of 10,723 patients
- The dataset includes patient demographic data, family history, concomitant medications, vital sign measurements, laboratory results, and patient clinical history
- They removed data using either an MCAR or MNAR approach
- For end metrics they considered both accuracy in imputation and ALS functional rating scale

# Imputation Example

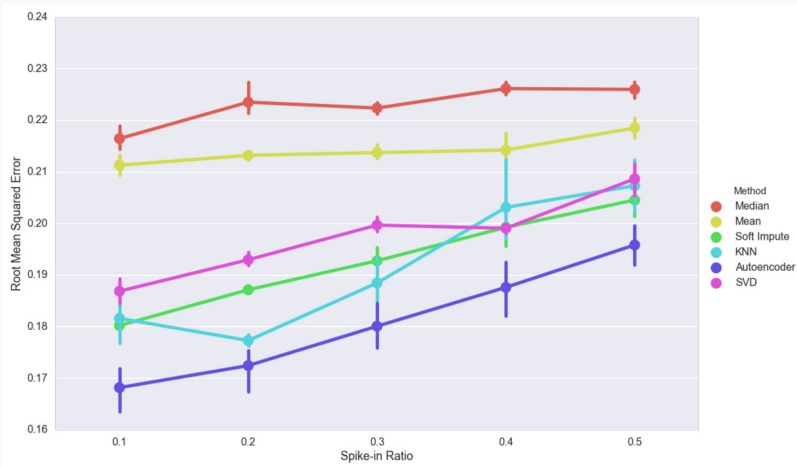
## Missing Completely at Random



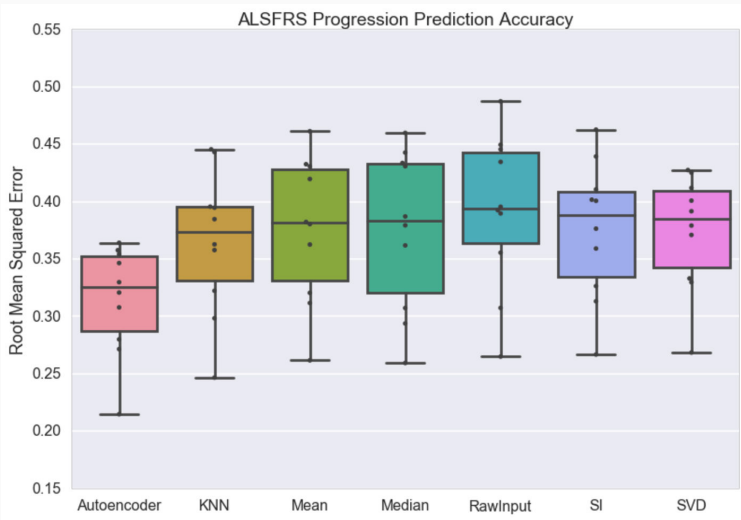


# Imputation Example

## Missing Not at Random



# Imputation Example



# Pipeline

- Using the pediatric sickle cell example let's walk through building the second half of the bioinformatics pipeline

