# **NoSQL** Database

Hands-On: R Flow Control Database Design Project

BMI701 Introduction of Biomedical Informatics Lab Session 4

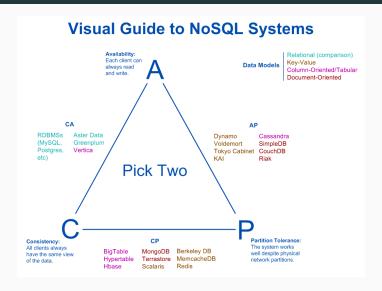
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HMS DBMI — MGH LCS





#### **Database Selection**



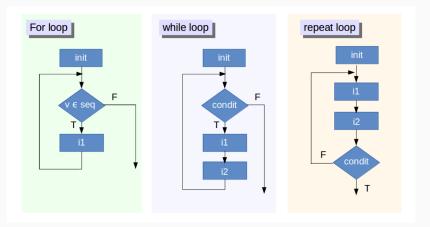
### **NoSQL**

- Key-value Stores
  - Hash table: key: value
  - Redis: data in RAM, not HD (so limited to RAM)
- Column Family Stores
  - Also key-value hash table, but one key may map to different columns (not the single value)
  - Cassandra: super fast write, slow read: e.g. facebook LIKE)
  - HBase: best to run MapReducr with Hadoop/HDFS stack, good for realtime queries (logs)
- Document Databases
  - JSON like document (BSON: binary JSON)
  - Nested: {Key:{Key:Value, Key:Value}}
  - MongoDB: very good for general use
  - CouchDB: can use RESTful HTTP API, use request
- Graph Databases
  - Node / Relation / Property
  - Neo4J

## MongoDB

- Download and install MongoDB
- Open your terminal
  - brew update
  - brew install mongodb
  - mkdir -p /mongo/db
  - mongod --dbpath /mongo/db
- Running MondoDB in R
  - install.packages("rmongodb")
  - library(rmongodb)
  - Then we can follow the cheat sheet

#### Flow Control



Courtesy by Dr. Mujeeb Basit

- for, while, if/else, next/break, stop
- Sample codes

### R: Apply Function

- apply
  - apply a given function to the rows (index 1) or columns (index 2) of a matrix
- lapply
  - apply a given function to every element of a list and obtain a list
- sapply
  - apply a given function to every element of a list and obtain a vector
- tapply
  - function to process vector subsets

### R: More Apply

- Map() and mapply()
  - iterate over multiple input data structures in parallel
- mclapply() and mcMap()
  - parallel versions of lapply() and Map()
  - Only for Linux or Mac
- clusterApply() and clusterApplyLB()
  - Linux, Mac, Windows

### R: Other Useful Packages/Functions

- foreach, parallel, doParallel
- sqldf
- system()
- Sample codes

### Take Home Message

- NoSQL summary
- Good luck on your presentation!
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