

# Rajan Kapoor



rkapr.github.io



r.kapoor@tamu.edu



linkedin.com/in/rkapr



979-xxx-xxxx



github.com/rkapr

**OBJECTIVE:** Looking for internship/co-op in data science/bioinformatics.

## EDUCATION:

Texas A&M University, College Station, TX

PhD in Electrical and Computer Engineering (GPA: 3.82)

May 2017 – Dec 2021

Texas A&M University, College Station, TX

Master of Science in Electrical and Computer Engineering (GPA: 3.75)

Aug 2015 – May 2017

Indian Institute of Technology, Patna, India

Bachelor of Technology in Electrical Engineering (GPA: 8.57/10)

Aug 2010 – May 2014

## DATA SCIENCE PROJECTS / EXPERIENCE:

***Zeroinfl: Zero Inflated Poisson regression (ZIP) in Python*** (with Eric Chuu)

[Spring 2018]

- Modeled a process switching between perfect state with no errors and imperfect one with Poisson distributed errors using ZIP regression based on pscl package in R
- Implemented logit, probit, complementary log log, Cauchy and log link functions for GLM fit using OOP
- Wrote functions for likelihood, log-likelihood, gradient and maximum likelihood estimation using BFGS optimization, with expectation maximization-based initialization
- Wrote functions summary for pretty-printing results including p-value, standard error, z-statistics and predict for using fitted model for prediction
- Extended functionality by including zero inflated negative binomial and zero inflated geometric regressions
- **Key achievement:** Provided fully functional zero inflated regression functionality in Python

***Adaboost based face detection using Viola Jones framework***

[Fall 2019]

- Implemented five stage Adaboost classifier in Python with decision stump (one step decision tree) as weak learner for detecting face images using vertical, horizontal and diagonal Haar features
- Vectorized code to speed-up repeated processing, observed effects of FP/FN penalty on empirical error
- **Key achievement:** Achieved **3.5x improvement** in training time at cost of 2.25% reduction in empirical accuracy by filtering robust features

***Minimum description length (MDL) based Boolean network learning***

[Spring 2020]

- Implemented MDL based gene network learning algorithm from short Boolean time series by minimizing sum of error and model-description entropies
- Tested and verified the code on Boolean time series data generated using *BoolNet* package in R
- **Key achievement:** Released MATLAB code based on Dougherty et al. paper was for open source use

***Mixture of Poisson's for modeling number of daily deaths***

[Spring 2018]

- Derived, implemented and compared convergence of gradient descent and Newton Raphson optimization for maximum likelihood estimates
- Calculated *sympy* expressions for gradient and hessian in Python, then converted to *numpy* functions to avoid errors due to hard coding expressions

***Gaussian Mixture Modeling for Cancer Heterogeneity***

[Fall 2016]

- Modeled cancer heterogeneity as mixture of Boolean networks using Gaussian mixture models

- Estimated proportion of each subpopulation using expectation maximization with k-means initialization
- **Key achievement:** Results were published in IEEE/ACM TCBB journal

## RESEARCH PROJECTS:

### *Understanding nitrogen transport pathways related to grain protein content in wheat*

[Spring 2019]

- Worked with professor from Soil and Crop Sciences Dept. to identify problem of interest
- Proposed and performed WGCNA based gene clustering using R scripts on computing cluster
- Identified UMAMI transporters in wheat, surveyed, collected, cleaned publicly available gene expression datasets for different development series, stress and tissue types
- Created heatmaps and homeolog expression plots for wheat and Arabidopsis to identify phylogenetically conserved patterns
- Developed shell based optimized pipeline to download large CRAM alignment files for relevant SRA projects using globus-cli, and organize runs for a given project
- Extracted reads mapping to UMAMI genes using SAMtools and BAMtools for gene model verification
- **Key achievement:** The analysis discovered two modules involved in activating storage molecules including starch, lipids and protein, discovered multiple transcription factors as possible key candidate genes

### *Gaussian Graphical model with fused lasso penalty for learning causal transcriptional regulations of Sorgoleone biosynthesis genes in Sorghum*

[Fall 2020]

- Collaborated with professor from Dept. of Biochemistry and Biophysics to identify research problem
- Proposed Gaussian graphical modeling with fused penalty to uncover gene interactions
- Built end-to-end pipeline for cleaning, mapping, counting, DEseq analysis of RNAseq data
- Inferred causal gene interactions for sorghum circadian genes using unsupervised learning
- Researched literature to verify gene interactions of Arabidopsis, rice, maize orthologs
- **Key achievement:** Identified 21 potential regulators of sorgoleone biosynthesis genes, verified TF-gene interactions using motifs from plant transcription factor database and comparative genomics

## SKILLS:

- R, Python (numpy, pandas, scikit-learn, scipy, cvtools), shell scripting (SLURM, LSF), MATLAB, SQL, Jupyter/R Notebooks, C/C++
- ML models: SVM with kernels, k-means, decision trees, Adaboost, linear, logistic, generalized linear regression, parsimonious model selection (AIC/BIC), lasso, ridge regression, mixture models, mixed models, stochastic gradient, coordinate descent, random forests, time series analysis (ARMA/ARIMA/ARCH/GARCH), ANOVA

## RELEVANT COURSES/ CERTIFICATIONS:

- *Engineering/Statistics:* Regression Analysis (A), Statistical computing in R & Python (A), Applied Statistics & Data Analysis (B), Pattern Recognition (A), Distribution Theory (A), Information Theory (A)
- *Biology/Bioinformatics:* Bioinformatics (S), Bioinformatics Command Line (A)
- *Online Certifications:* Introduction to SQL, Learn the Command Line, Algorithms & Data Structures

## PUBLICATIONS:

- A Gaussian Mixture-Model Exploiting Pathway Knowledge for Dissecting Cancer Heterogeneity, IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018

## HONORS/ AWARDS:

- Texas Engineering Experiment Station (TEES) Research Assistantship, Sept 2015 - present
- Texas International Student Scholarship, Fall 2015, Spring 2016