

# Parth S. Patel

Data Scientist  
parthspatel.me | parthspatel.nj@gmail.com

## EDUCATION

### DREXEL UNIVERSITY

#### BS IN BIOMEDICAL ENGINEERING

Sept 2014 - June 2018

Concentrated in Bioinformatics

College of Biomedical Engineering

## LINKS

Website: [parthspatel.me](http://parthspatel.me)

Github: [parthspatel](https://github.com/parthspatel)

LinkedIn: [parthspatel96](https://www.linkedin.com/in/parthspatel96)

## PROJECT LINKS

TMRemote: [beta.tmremote.io](https://beta.tmremote.io)

## COURSEWORK

### UNDERGRADUATE

Data Mining & Machine Learning

Data Structures & Algorithms

Genome Information Engineering

Computational Bioengineering

Database Management Systems

Linear & Dynamic Systems

Statistics

Biosimulation

## ONLINE

Stanford: Machine Learning

Stanford: Convolutional Neural Networks for

Visual Recognition

## SKILLS

### PROGRAMMING

Python (scikit-learn, TensorFlow, PyCUDA)

SQL (Transact & Postgres)

R (shiny & ggplot2)

C++ • C • MATLAB • Node.js • Java • Batch

TEX • Visual Basic • Android

### SOFTWARE

Tableau • Excel(VBA, formulas, pivot tables)

Teradata Studio

Android Studio • Access • SharePoint

## HOBBIES

Origami • Digital Painting • Cooking

## REFERENCES

Jason Pan | Data Scientist | Pfizer

Ashima Varshney | Associate Director CMC |

Pfizer

Bing Yao, Ph.D. | Physicist | Kessler

Foundation

## EXPERIENCE

### Wylei | DATA ANALYST

August 2018 – Present | Newport, NJ

- Detect & filter duplication patterns for marketing campaigns using ml techniques.
- Automated report generation and designed dashboards, to present campaign performance to clients using python & Node.js, reducing delays.
- Simplified and optimized the report configuration process by developing a tool using Node.js, reducing turnaround time for new report by days.

### Pfizer Global Business Intelligence | DATA SCIENTIST

Sept 2016 – March 2017 | Manhattan, NY

- Developed a prescription shopping outlier detection system based on unpredictability. Employed Shannon's entropy and clustering for 70M patients on opiates in the OptumRX dataset
- Trained the larger team in anomaly detection algorithms and techniques
- Employed Kaplan-Meier Estimator and Greenwood's Formula to predict probability of persistence on therapy for oncology drugs, presented via a persistency chart
- Automated the weekly vendor data refresh process using shell scripting and Python

### Kessler Foundation | SUMMER INTERN

June 2015 – Sept 2015 | West Orange, NJ

- Successfully designed and implemented an EEG-fMRI study on visual networks in the brain of multiple sclerosis patients

### Rutgers University WINLAB | SUMMER INTERN

June 2014 – Aug 2014 | New Brunswick, NJ

- Developed an Android music streaming application using the Mobility First API and Android Studio in a team of five

## PROJECTS

### Deep learning for CRISPR-Cas9 Activity Prediction | May 2018

Developed a convolutional neural network architecture using TensorFlow trained on Cas9 modification frequencies aggregated from literature. Demonstrated that our models perform better at predicting Cas9 activity than the CNN Seq-Deep model (Kim et al. 2018) and that deep learning methods are better than deterministic predictors.

### TMRemote - Real Time Data Collection & Manager | March 2018 – Present

Developed a dedicated desktop and web application to remotely monitor and update bot applications, utilizing QT, Python, PostgreSQL, and PHP. Currently under beta testing. ([link](#))

### CRISPR Excision Therapy Screening | Sept 2017 – May 2018

Designed a time-efficient, high throughput, and accurate screening strategy for a CRISPR-spCas9 excision therapy.

- Designed a custom microarray, describing a patient's HIV genome
- Maintained and utilized the crisptree Python module to determine Cas9 binding and cleavage potential
- Developed a Python algorithm to derive patient suitability from microarray results and crisptree

The developed methods and tools allow researchers to redesign the current treatment to target and treat more patients.

### Prostate Cancer Predictor | August 2017

Created a prostate cancer predictor in Python, using biomarker concentrations data. Utilized random forests and kNN for classification and 5-fold cross-validation for optimization.