Ashley D'Souza

■ akd6@rice.edu ■ 512-228-4140 ■ <u>Home</u> ■ <u>Blog</u> ■ <u>Github</u> ■ <u>LinkedIn</u>

Education ————————————————————————————————————	
·	17 - present
 BS Computer Science & BA Cognitive Science, graduating May 2021, GPA: 3.40 	Houston, TX
 Westwood High School GPA: 4.0/4.0 (5.58/5.0 weighted), National Merit Scholar, Graduated May 2017 	2013 - 2017 Austin, TX
Work Experience	
	019-present Houston, TX
 UT Austin, College of Natural Sciences, Vertebrate Interactome Lab Summer Research Intern DNA recombineering of DHX35 gene in E. coli, and computational Flux Balance Analysis on cyanobacteria 	2016 Austin, TX
Conference Presentations ————————————————————————————————————	
PyoFuel: Using Python and Pathway Tools to engineer synthetic biofuel Sole Author more ■ Int'l. Soc. for Computational Biology / 2016 Rocky Mountain Bioinformatics Conference (poster session)	Dec 2016 Colorado
Software Development Skills ———————————————————————————————————	
Programming Languages & Frameworks 20	13 - present
Proficient: Python, Java, ReasonML/OCaml, React	
Basic: C, Racket, Pyret, Elm, Html, Hasura's Postgres + Graph Software Design 20	12 procent
 Software Design Systematic Program Design - Designing Data & Functions, EdX course based on HtDP2 	13 - present
Typed functional programming, Object-Oriented Design	
Software Testing • Test-driven development, unit & property-based tests, Python: Pytest, Java: JUnit 5 & QuickTheories	17 - present
Projects	
Lentil - A Web-Based Musician's Feedback Service Independent Project repo demo 20 ■ Musicians submit performances and receive pointed feedback from others (ReasonML, React, Postgres)	18 - present
Physics Sunset Independent Project repo demo	2017
• Design and implement a browser-based interactive graphical simulation of a physics problem (ReasonML)	
Disease Transmission Analysis from Outbreak Data	2018
 Infer disease transmission tree from disease outbreak genetic + epidemiological data using RDMST (Python DNA Sequence Alignment)
	2010
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene 	2018
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) 	2018
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene 	
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene Phylogenetic (Evolutionary) Trees Infer evolutionary tree, given DNA sequences for leaf taxa and plausible mutations (Python) Hidden Markov Models and Part-of-Speech Tagging (NLP) 	
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene Phylogenetic (Evolutionary) Trees Infer evolutionary tree, given DNA sequences for leaf taxa and plausible mutations (Python) Hidden Markov Models and Part-of-Speech Tagging (NLP) Implement statistical learning of HMM using training corpus of pre-tagged sentences (Python) 	2018
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene Phylogenetic (Evolutionary) Trees Infer evolutionary tree, given DNA sequences for leaf taxa and plausible mutations (Python) Hidden Markov Models and Part-of-Speech Tagging (NLP) Implement statistical learning of HMM using training corpus of pre-tagged sentences (Python) Implement Viterbi algorithm to assign part-of-speech tags to new sentences using trained HMM 	2018 2018
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene Phylogenetic (Evolutionary) Trees Infer evolutionary tree, given DNA sequences for leaf taxa and plausible mutations (Python) Hidden Markov Models and Part-of-Speech Tagging (NLP) Implement statistical learning of HMM using training corpus of pre-tagged sentences (Python) 	2018
 Design and implement dynamic programming solutions to DNA sequence alignment problems (Python) Align human and fruit-fly protein sequences to identify the PAX domain within the "eyeless" gene Phylogenetic (Evolutionary) Trees Infer evolutionary tree, given DNA sequences for leaf taxa and plausible mutations (Python) Hidden Markov Models and Part-of-Speech Tagging (NLP) Implement statistical learning of HMM using training corpus of pre-tagged sentences (Python) Implement Viterbi algorithm to assign part-of-speech tags to new sentences using trained HMM Impact of Language on Perception 	2018 2018

Organizations and Activities CSters: Women in Computer Science, Rice University CS Club, Rice University Society of Women Engineers (SWE), Rice University 2017 - present 2017 - present

SASE: Society of Asian Scientists and Engineers, Rice University
 Club Tennis, Rice University
 2017 - present

• Music: sing, record, perform, take lessons soundcloud youtube 2007 - present

2015 - 2017

• PyLadies: Austin community of women Python programmers, team programming, presentations