# **ANTO SUJIN J**

88/B,CHITHAMBARA NAGAR KOTTAR,NAGERCOIL 629002 9080763832 antosujinjoseph@gmail.com

## **SKILLS**

HTML: Semantic HTML5, Form Handling, Accessibility (ARIA), SEO Best Practices, Media Embedding.

**CSS**: Responsive Design, CSS Grid & Flexbox, Animations & Transitions, Frameworks (Bootstrap).

JavaScript: ES6+ Features (Arrow Functions, Promises, Async/Await), DOM Manipulation, Event Handling, Fetch API & Axios, JSON, Modular Programming (Import/Export), Error Handling, JavaScript Frameworks (React.js), State Management (Redux).

React.js: Component-Based Architecture, Functional Components & Hooks, State Management (Redux), Routing (React Router), Crud.

**Python**: Strong understanding of Python syntax, data types, control flow, and object-oriented programming concepts.

#### **LANGUAGES**

Tamil English Malayalam

### **PROFESSIONAL SUMMARY**

To be associated in a reputed organization in which I can contribute my knowledge and skills that dynamically works Towards the growth of organization.

#### **EDUCATION**

**SATHYABAMA UNIVERSITY,** CHENNAI—*Msc BIOINFORMATICS AND DATASCIENCE -73%*2022 - 2024

**SCOTT CHRISTIAN COLLEGE,** NAGERCOIL—*Bsc COMPUTER SCIENCE-61%*2018 -2021

D.V.D HIGHER SECONDARY SCHOOL, NAGERCOIL-69% 2016-2018

ST.XAVIER'S HIGH SCHOOL, NAGERCOIL-95% 2006-2016

### **PROJECTS**

## Mining TCGA database for genes associated in colon cancer

Employed Python and R programming languages to analyze gene expression data from the TCGA database. Utilized bioinformatics libraries (e.g., pandas, numpy, scipy, statsmodels) for data preprocessing, statistical analysis, and visualization. Conducted differential gene expression analysis using methods like t-tests and DESeq2. Annotated identified genes with functional information from GO and KEGG databases. Performed pathway enrichment analysis using tools like GSEA. Evaluated the prognostic significance of genes using survival analysis techniques.