

# DEEPIKA T N

Master of Science in Computer Science Student

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## SUMMARY

I am a Master of Science in Computer Science student with a passion for software development, database management, and web technologies. Proficient in Python, HTML, SQL, DBMS, and Excel, with a strong foundation in problem-solving, data structures, and algorithms. Seeking internship opportunities to apply technical skills in real-world software projects and enhance my expertise in backend and frontend development.

## EDUCATION

### Master of Science in Computer Science

#### MOUNT CARMEL COLLEGE

08/2024 - 06/2026 Bengaluru

### Bachelor of Science in Mathematics, Statistics and Computer Science

#### REVA University : 87%

12/2021 - 06/2024 Bengaluru

### Pre-University Education

#### Nagarjuna Pre-University College : 90%

05/2019 - 03/2021 Bengaluru

### School Education

#### Jnana Jyothi Educational Institution:91%

06/2011 - 03/2019 Bengaluru

## LANGUAGES

English  
Native



Kannada  
Native



Hindi  
Proficient



## INTERESTS

### Hobbies

Enjoys listening to music and exploring creativity through drawing and pencil shading

### Extracurricular Activities

Participated in cultural activities and volunteered in local cancer hospitals to support patients

## CERTIFICATIONS & ACHIEVEMENTS:

NPTEL Certificate

DBMS , Fundamentals of OOPS

 Cognitive Class.AI Powered by IBM Developer Skills Network.  
Python Data Science

## SKILLS

Python Data Structures SQL

Excel Tableau Pandas

Java

## PROJECTS

### Blood Bank Management System

01/2023 - 05/2023

A project to manage blood donations and inventory effectively

- Developed a database-driven web application to manage blood donations and inventory
- Implemented secure login authentication, donor registration, and hospital management features

Technologies used:

PHP MySQL ,HTML ,CSS ,JavaScript

### Computational Drug Discovery Using Machine Learning 2025

Predictive Modeling of Drug Potency (pIC50) for Alzheimer's Treatment

- Developed a QSAR-based Random Forest regression model to predict pIC50 values from molecular structure data, targeting Human Acetylcholinesterase (AChE).
  - Benchmarked model performance using R<sup>2</sup> and MAE metrics; deployed and tested in Google Colab using LazyPredict for comparative analysis.
- Languages:** Python
- Tools & Libraries:** PaDEL-Descriptor, LazyPredict, Scikit-learn, Pandas, NumPy, ChEMBL Database