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| SAFIYA IMROZ  E- mail:  [safiya22imroz@gmail.com](mailto:safiya22imroz@gmail.com) **Contact Number:** Mobile: +91 89811 50605  Permanent Address:  39/1, Ismail Street, Kolkata -700014 Father's Name: Md. Hashim  **Date of Birth:**  08-09-1995  **Sex**:  Female  **Nationality:**  Indian  **Marital Status:**  Single  **Hobbies:**  Cooking  Reading Books  Watching Movies | **Aim / Career Objective**  To achieve a responsible position where I can apply my knowledge and skills with an opportunity for professional challenge and growth to support and enhance the objectives of the company.  **Education**   |  |  |  |  |  | | --- | --- | --- | --- | --- | | Examination/ Degree | Institution | Performance | | Year of completion | | B.Tech in Computer Science Engineering | Aliah University | 1st Sem | 6.67 | 2018 | | 2nd Sem | 5.59 | | 3rd Sem | 7.17 | | 4th Sem | 7.97 | | 5th Sem | 8.71 | | 6th Sem | 8.20 | | 7th Sem | 8.65 | | 8th Sem | 8.85 | | DGPA | 7.7 | | Higher Secondary Education (12th) (ISC) | Anglo Arabic Secondary School | 80% | | 2014 | | Secondary Education (10th) (ICSE) | Ling Liang High School | 70% | | 2012 |     **Computer / IT Skills**   |  |  | | --- | --- | | Programming Language | Java, C, MySQL | | Software | Microsoft Office (Word, Power point, Excel) |     **Participation/Achievements**   * Participated in Horlicks Wiz Kids (School Level). * Awarded Merit cum means Scholarship from WBMDFC for four consecutive years. * Participated in college cultural events.   **Languages**   * English * Hindi * Bengali * Urdu   **Subjects Of Interest**   * OS * Data Structure * Java |

**Training/ Industrial Visits**

* Training on Core java from Micro Pro(duration 3 months).
* Internship on Advance java Using JSP, Servlet organized by Prudential Infotech Pvt Ltd.

### Project Undertaken

* Project name: Disease Associated Gene Classification Using Supervised Learning.
* Software used: Orange, Gephi, RStudio and Matlab.
* Language involved: R & Python .
* Description: An approach based on supervised method to find if a particular gene is linked with

the disease or not. Using this approach, we can extract known disease genes and to predict unknown genes, which can be good candidates for experimental study.

Declaration

I hereby declare that the information given above is true to the best of my knowledge and belief and can be supported with reliable documents when needed.