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# **Project Report**

## **HUMAN SKELETAL SYSTEM**

### **1. Introduction**

#### **1.1 Purpose**

The purpose of this project is to study and organize the bones in the human body by applying various data structures and algorithms. The project will serve as a learning tool for understanding the number and specific sequence of bones in an adult human being and the application of computational techniques to biological data.

#### **1.2 Document Conventions**

- All headings and subheadings follow a hierarchical numbering system.
- Code snippets for algorithms are included in monospace font.
- Diagrams are provided where necessary.

#### **1.3 Project Scope**

##### **1.3.1 Scope Definition**

This project covers:

- Identifying and organizing the bones in the adult human body.
- Applying algorithms (insertion, merge, heap, quick sort) and data structures (array, stack, queue, tree, hash) to the data.

##### **1.3.2 Core Features**

- Implementation of sorting algorithms for ordering bones.
- Data structure operations to manage bone information.

##### **1.3.3 Subsequent Releases**

Future versions may include:

- Enhanced visualization of the skeletal system.
- Integration with 3D modeling tools.

### **1.3.4 Alignment with User and Business Goals**

The project aligns with educational and research objectives, making data analysis for biological systems more accessible and structured.

## **1.4 References**

1. Gray's Anatomy: The Anatomical Basis of Clinical Practice.
  2. Introduction to Algorithms (Fourth Edition).
  3. Human Anatomy and Physiology (Marieb).
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## **2. Overall Description**

### **2.1 Product Perspective**

#### **2.1.1 Product Context**

The project integrates anatomy and computational techniques to enhance understanding of skeletal organization.

#### **2.1.2 Product Origin**

This is an educational project aimed at bridging biology and computer science.

#### **2.1.3 Product Relationship to Existing Systems**

Relates to anatomy educational software and algorithm simulation tools.

#### **2.1.4 Product Ecosystem**

It fits into learning management systems and standalone computational tools for anatomy.

### **2.2 User Classes and Characteristics**

#### **2.2.1 Tech Enthusiasts**

Interested in data structures and algorithms.

#### **2.2.2 Casual Shoppers**

Biology students aiming for a structured study guide.

#### **2.2.3 Favored User Class**

Educators and researchers in interdisciplinary fields.

#### **2.2.4 Alignment with User Needs**

Provides practical examples of algorithms and their application to real-world data.

### **2.3 Operating Environment**

#### **2.3.1 Hardware Platform**

Minimum: Laptop/Desktop with modern processor and 4 GB RAM.

#### **2.3.2 Operating Systems and Versions**

Windows 10+, macOS 10.14+, Linux (Ubuntu 18.04+).

### **2.4 Design and Implementation Constraints**

#### **2.4.2 Database Technology**

SQLite for managing bone-related data.

#### **2.4.4 Third-Party Integrations**

No external libraries beyond Python's standard library.

#### **2.4.5 User Interface Design**

Simple command-line interface for interaction.

### **2.5 Assumptions and Dependencies**

#### **2.5.1 Assumptions**

- User has basic knowledge of biology and programming.
- Python 3.8+ is pre-installed.

#### **2.5.2 Dependencies**

- Python standard libraries for data processing.

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## **3. System Features**

- Detailed list of all 206 bones in the adult human body.
  - Sequence management using sorting algorithms.
  - Operations on bones using arrays, stacks, queues, and trees.
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## 4. Number and Sequence of Bones

The adult human skeleton consists of 206 bones, organized as follows:

### 4.1 Axial Skeleton

- **Skull:** 22 bones
  - Cranial bones: 8 (e.g., frontal, parietal, occipital)
  - Facial bones: 14 (e.g., maxilla, mandible, zygomatic)
- **Hyoid Bone:** 1 bone
- **Vertebral Column:** 26 bones
  - Cervical: 7
  - Thoracic: 12
  - Lumbar: 5
  - Sacrum: 1 (fused)
  - Coccyx: 1 (fused)
- **Thoracic Cage:** 25 bones
  - Ribs: 24
  - Sternum: 1

### 4.2 Appendicular Skeleton

- **Pectoral Girdle:** 4 bones
  - Clavicles: 2
  - Scapulae: 2
- **Upper Limbs:** 60 bones
  - Humerus: 2
  - Radius: 2
  - Ulna: 2
  - Carpals: 16
  - Metacarpals: 10
  - Phalanges: 28
- **Pelvic Girdle:** 2 bones
  - Hip bones (coxal bones): 2
- **Lower Limbs:** 60 bones
  - Femur: 2
  - Tibia: 2
  - Fibula: 2
  - Patella: 2
  - Tarsals: 14

- Metatarsals: 10
  - Phalanges: 28
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## **5. External Interface Requirements**

### **5.1 User Interfaces**

#### **5.1.1 Design Standards and Guidelines**

Follows a modular programming approach.

#### **5.1.2 Screen Layout and Resolution**

Text-based outputs for easy readability.

#### **5.1.3 Standard Interface Elements**

Command-line input and output.

### **5.2 Software Interfaces**

#### **5.2.4 Non-Functional Requirements**

- Code readability and maintainability.
- Efficient algorithms with optimal time complexity.

### **5.3 Hardware Interfaces**

#### **5.3.1 Supported Device Types**

PCs and laptops.

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## **6. Quality Attributes**

### **6.1 Performance**

Efficient sorting and data structure manipulation within  $O(n \log n)$  complexity.

### **6.2 Reliability**

Correctly manages and organizes all 206 bones.

### 6.3 Usability

Intuitive interface for non-expert users.

### 6.4 Security

No sensitive data is processed.

### 6.5 Maintainability

Code is modular and well-documented.

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## Appendix B: Analysis Model

- **Data Representation:** Each bone is stored as an object containing its name, location, and properties.
  - **Sorting Algorithms:** Applied to organize bones alphabetically or by position.
  - **Data Structures:**
    - **Array:** Stores bone names.
    - **Stack:** Allows reverse sequencing of bones.
    - **Queue:** Simulates a lineup of bones for processing.
    - **Tree:** Represents hierarchical relationships (e.g., skull -> cranial bones).
    - **Hash Table:** Fast retrieval of bone information.
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