

Robust Airway Generation Labelling with Airway Segmentation for Reliable Airway Assessment

Pooja K N ¹, Teju S V ²

¹ Assistant Professor, Department of MCA, BIET, Davanagere

² Student, 4th Semester MCA, Department of MCA, BIET, Davanagere

Abstract

This study focuses on the accurate classification of airways within the human respiratory system, which is characterized by a complex, tree-like structure with diverse patterns. While recent advancements in deep learning show potential for automatic airway segmentation, further research is necessary to make these methods practical for assessing a patient's airway status and enabling personalized treatment. A thorough investigation into airway generation is essential for improving diagnostics and therapies, as it aids in understanding specific structures and identifying issues such as airway narrowing and lung compliance. Current techniques, including template matching and machine learning approaches that treat airway classification as a problem, have limitations in capturing the complexities of higher-generation airways. To address these challenges, our study introduces a novel methodology. The Prim algorithm is employed to establish a minimum spanning tree from the centerline, creating an accurate tree structure that reflects airway connections. A new secondary branch is generated when a main branch with an outdegree of 1 is identified, ensuring precise labeling of airway generation.

Keywords: Airway classification, human respiratory system, tree-like structure, deep learning, automatic airway segmentation, airway generation, diagnostics, personalized treatment, template matching, machine learning, Prim algorithm, minimum spanning tree, airway connections, false branch generation, trunk lines, pruning method, subtree length, segmentation results, patient health monitoring, disease prediction, and prevention.

I. INTRODUCTION

The accurate classification of airways in the human respiratory system is essential for effective diagnostics and treatment planning. The intricate, tree-like structure of the airways presents significant challenges for segmentation and assessment, particularly in higher-generation regions where

small airways are located. Recent advancements in deep learning have shown promise for automatic airway segmentation; however, existing methods still face limitations in capturing the complexity of airway structures and ensuring reliable assessments.

Current approaches to airway segmentation typically involve skeletonization and tree structure

detection, template matching, or a combination of both. While these methods can provide valuable insights, they often suffer from issues such as false branch generation and inaccuracies in detecting variable branch forms, particularly in the presence of bumpy airway surfaces. Errors in lower-generation airways can propagate through the segmentation process, leading to further inaccuracies in higher-generation assessments. Additionally, template matching methods are constrained by their

reliance on predefined templates, which may not accommodate the diverse topological structures of higher-generation airways across different individuals.

To address these challenges, our proposed framework introduces a novel approach to airway generation that leverages the Prim algorithm to establish a minimum spanning tree from the centerline. This method allows for the accurate representation of airway connections while minimizing the generation of false branches. By implementing a tree-pruning technique based on subtree length, we can effectively manage segmentation results from deep learning, ensuring that only valid airway branches are retained.

The proposed system not only enhances the reliability of airway assessments in higher-generation regions but also contributes to improved automatic diagnostic capabilities and treatment planning. By providing precise identification of airway generations, our approach holds significant potential for advancing patient health monitoring, disease prediction, and prevention. Through

experimental validation on public datasets and clinical outcomes from our private dataset, we demonstrate the effectiveness of our framework in overcoming the limitations of existing methods and achieving robust airway generation labeling.

2. LITERATURE REVIEW

S. Gupta, K. R. Sharma, and D. Jain, "Airway assessment: Predictors of difficult airway," *Indian J. Anaesthesia*, vol. 49, no. 4, pp. 257–262, 2005.

Expertise in airway management is crucial across all medical specialties. Ensuring a patent airway is vital for adequate oxygenation and ventilation, as even a brief failure to maintain it can be life-threatening. Respiratory events are the most frequent injuries associated with anesthesia, following dental damage. The primary causes of respiratory-related injuries include inadequate ventilation, esophageal intubation, and challenging tracheal intubation. Difficult tracheal intubation accounts for 17% of these respiratory-related injuries and can lead to significant morbidity and mortality. In fact, up to 28% of all anesthesia-related fatalities are attributed to the inability to effectively mask ventilate or intubate patients.[1]

D. Couper, L. M. LaVange, M. Han, R. G. Barr, E. Bleeker, E. A. Hoffman, R. Kanner, E. Kleerup, F. J. Martinez, P. G. Woodruff, and S. Rennard, "Design of the subpopulations and intermediate outcomes in COPD study (SPIROMICS): Table 1," *Thorax*, vol. 69, no. 5, pp. 492–495, May 2014.

The Subpopulations and Intermediate Outcomes in COPD Study (SPIROMICS) is a multicenter observational study focused on chronic obstructive pulmonary disease (COPD). Its aim is to inform the

future development of COPD therapies by establishing robust criteria for subclassifying participants into groups that are most likely to benefit from specific treatments in clinical trials. Additionally, the study seeks to identify biomarkers and phenotypes that can serve as intermediate outcomes to reliably predict clinical benefits during therapeutic trials. The objective is to enroll 3,200 participants across four strata. Each participant will undergo a baseline visit followed by three annual follow-up examinations, along with quarterly telephone check-ins. The study will also include the adjudication of exacerbations and mortality.[2]

R. D. Miller and R. E. Hyatt, "Evaluation of obstructing lesions of the trachea and larynx by flow-volume loops," *Amer. Rev. Respiratory Disease*, vol. 108, no. 3, pp. 475–481, Aug. 1973.

This study investigates the diagnostic value of flow-volume loops in identifying obstructive lesions within the trachea and larynx. By analyzing airflow patterns during forced inspiration and expiration, the research demonstrates how specific changes in loop morphology can help differentiate between upper airway obstructions. The findings emphasize that flow-volume loops serve as a non-invasive and effective method to evaluate central airway narrowing, thereby aiding clinicians in the timely diagnosis and management of tracheal and laryngeal obstructions.[3]

A. P. Kiraly, W. E. Higgins, G. McLennan, E. A. Hoffman, and J. M. Reinhardt, "Three-dimensional human airway segmentation methods for clinical virtual bronchoscopy," *Academic Radiol.*, vol. 9, no. 10, pp. 1153–1168, Oct. 2002.

This paper presents advanced segmentation techniques for constructing three-dimensional models of human airways, facilitating the use of clinical virtual bronchoscopy. The authors detail image processing algorithms that extract airway structures from high-resolution CT scans with improved accuracy. The study highlights how these 3D segmentation methods enhance the visualization of bronchial pathways, allowing clinicians to perform non-invasive diagnostics, plan bronchoscopic procedures more effectively, and detect abnormalities that may not be visible through traditional imaging techniques.[4]

V. R. de Water, J. K. Saridin, F. Bouw, M. M. Murawska, and M. J. Koudstaal, "Measuring upper airway volume: Accuracy and reliability of dolphin 3D software compared to manual segmentation in cranosynostosis patients," *J. Oral Maxillofacial Surgery*, vol. 72, no. 1, pp. 139–144, Jan. 2014.

This study evaluates the accuracy and reliability of Dolphin 3D imaging software in quantifying upper airway volume in cranosynostosis patients, comparing it with traditional manual segmentation methods. The findings demonstrate that Dolphin 3D offers high reproducibility and efficiency, suggesting it is a reliable alternative for clinical assessment and surgical planning. The study underscores the growing role of automated tools in craniofacial and airway evaluation, particularly in complex anatomical conditions.[5]

N. A. Alsufyani, A. Hess, M. Noga, N. Ray, M. A. Q. Al-Saleh, M. O. Lagravère, and P. W. Major, "New algorithm for semiautomatic segmentation of nasal cavity and pharyngeal airway in comparison

with manual segmentation using cone-beam computed tomography,” *Amer. J. Orthodontics Dentofacial Orthopedics*, vol. 150, no. 4, pp. 703–712, Oct. 2016.

The authors propose a novel semiautomatic algorithm for segmenting the nasal and pharyngeal airways using CBCT imaging, aimed at improving efficiency and reducing operator dependency. The algorithm’s performance is benchmarked against manual segmentation, showing high correlation and reduced processing time. This advancement contributes to the development of reliable airway analysis tools in orthodontics and respiratory diagnostics.[6]

A. Weissheimer, L. M. D. Menezes, G. T. Sameshima, R. Enciso, J. Pham, and D. Grauer, “Imaging software accuracy for 3-dimensional analysis of the upper airway,” *Amer. J. Orthodontics Dentofacial Orthopedics*, vol. 142, no. 6, pp. 801–813, Dec. 2012.

This research investigates the precision of various imaging software programs used for three-dimensional upper airway analysis. It compares the software outputs with manually derived measurements, assessing discrepancies and reliability across different platforms. The study highlights the variability among tools and emphasizes the need for validated, standardized software for consistent and accurate upper airway assessment in orthodontic and medical practice.[7]

J.-P. Charbonnier, E. M. V. Rikxoort, A. A. A. Setio, C. M. Schaefer-Prokop, B. V. Ginneken, and F. Ciompi, “Improving airway segmentation in computed tomography using leak detection with

convolutional networks,” *Med. Image Anal.*, vol. 36, pp. 52–60, Feb. 2017.

This study introduces a deep learning-based approach for enhancing airway segmentation in CT images, employing convolutional neural networks (CNNs) to detect and correct segmentation leaks. The methodology leverages a dual-stage pipeline combining initial segmentation with a CNN-based leak detector, significantly improving accuracy and robustness. The findings mark a significant step toward automating complex medical image analysis tasks using artificial intelligence.[8]

3. EXISTING SYSTEM

These kinds of methods generally include (1) airway segmentation skeletonization and detecting tree structure from skeleton, (2) template matching, and (3) a combination of both approaches. The methods based on the first approach often suffer from errors related to false branches and variable forms of branches (e.g., trifurcation). This is primarily because the skeleton of the airway used in tree detection is generated inaccurately when the airway has bumpy surfaces. When these methods are employed to evaluate the generation of airways, errors that occur in lower generations propagate through, leading to additional errors in higher generations. Therefore, it is essential to design a process that filters such errors for these methods.

The methods based on the second approach are more robust to such errors, but they have considerable limitations, as they can only detect branches included in the template. Since higher-generation airways exhibit varied topological structures among individuals, they cannot be templated and are

undetectable with these methods. This makes the methods using the second approach inappropriate for small airway assessment.

As various machine learning techniques (e.g., deep neural networks (DNNs)) have been developed, numerous studies on airway labeling that exploit deep neural networks have been published. Some of these methods use skeletonization, similar to some methods in the previous section, to find all candidate branch points (including false branches). These candidate branch points are then provided as input to the DNN along with some information about each candidate branch point. These methods define the problem as a classification problem that classifies which type the given candidate branch point belongs to. The 'types' of branch points are predefined in this context. Candidate branch points that do not correspond to the predefined types or are false branch points are considered not to be branch points by the DNN. Only low-generation branch points that commonly appear in every individual's airway are considered, while others remain undefined.

Other research defines the problem as a regression problem that finds the coordinates of predefined branch points. This method directly obtains the position of branch points from 3D CT images before performing airway segmentation or skeletonized airway segmentation. After this, airway segmentation is conducted using a region-growing algorithm, guided by the calculated branch points. These methods have the advantage of being less likely to misclassify false branch points as real branch points and can produce consistent results across topologically diverse airway segmentations. However, a significant limitation of these methods

is that they are not capable of detecting high-level branches that are not predefined. Since we aim to assess small airways in higher generations with varied topological structures, these methods are not very useful for our purpose.

3.1 DISADVANTAGE OF EXISTING SYSTEM

The intricacy of data presents a significant challenge, as most current machine learning models must effectively analyze extensive and complex datasets to identify Robust Airway Generation Labeling. Additionally, data accessibility plays a crucial role; many machine learning models rely on substantial volumes of data to generate precise predictions, and insufficient availability can lead to diminished accuracy. Furthermore, mislabeling issues can severely impact model performance, as the effectiveness of these models is directly tied to the quality of the training data. If the dataset contains incorrect labels, the model's predictive capabilities will be compromised, ultimately affecting its reliability and accuracy.

4. PROPOSED SYSTEM

In this paper, we introduce a new framework for airway generation that precisely identifies airways in higher generation regions, ensuring dependable airway mapping by flexibly managing the outcomes of the centerline analysis. Firstly, we incorporate the Prim algorithm to establish a minimum spanning tree by connecting each point of the centerline. In the minimum spanning tree, the emergence of a primary branch connecting to the centerline with an outdegree of 1 results in the formation of an unnecessary secondary branch. In addition, to

mitigate the generation of false branches, we propose a method that calculates the cost of trunk lines connected to a centerline with an outdegree of 1.

Next, we introduce a tree-pruning method based on subtree length to effectively manage segmentation results predicted by deep learning. This method prevents the occurrence of false branches by removing vertices from the tree whose subtree airway lengths are shorter than their sibling vertices. Therefore, our approach rectifies the main issue of false branch generation, which is prevalent in the existing centerline-based airway generation labeling method. As a result, assessment in the higher-generation sections is possible, enabling effective airway evaluation, particularly for small airways. This enhancement contributes to the improvement of automatic diagnostic capabilities and treatment planning.

4.1 Advantages of Proposed System

We propose a novel framework for airway generation that addresses the challenges in higher generation areas (i.e., small airways) by mitigating the generation of false branches. The proposed framework facilitates robust airway assessment by accurately identifying airways in higher generation areas, enhancing automatic diagnostics and treatment planning. The experimental results on public datasets and the clinical outcomes of airway assessment on our private dataset validate the effectiveness of our approach in airway generation.

4.1 System Architecture

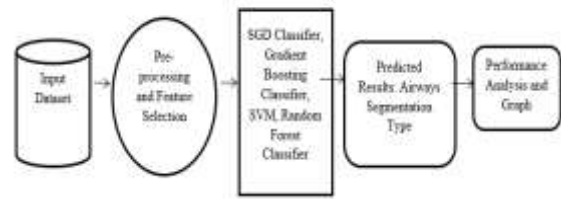


Fig:4.1.1 System Architecture

5. MODULE DESCRIPTION

5.1 Remote User:

The remote user component of this project allows users to securely log in and interact with the system by uploading data from the dataset to predict airway segmentation types. This interface is designed to be user-friendly, enabling users to efficiently submit their input data and receive accurate predictions of airway segmentation based on the trained machine learning models. The remote user can focus on data input and obtaining results without needing to manage the underlying algorithms or system details, making it accessible for users with varying levels of technical expertise.

5.2 Service Provider:

The service provider component serves as the administrative and analytical hub of the project. It displays comprehensive information about all the algorithms implemented within the system, including detailed metrics on the accuracy and performance of each algorithm. Additionally, service providers can view the results of airway segmentation predictions made by remote users, facilitating validation and quality control. This interface also provides a complete overview of all

remote users, offering access to their details for management and monitoring purposes. Together, these features empower service providers to oversee the system's operation, optimize algorithm performance, and ensure overall reliability and transparency.

6. RESULTS

The algorithm accuracy results for the SGD Classifier, Gradient Boosting Classifier, SVM, and Random Forest Classifier can be effectively visualized in a pie chart, which highlights the performance of each model. The SGD Classifier achieved an accuracy of 51.5%, indicating a moderate level of effectiveness in predicting airway segmentation types. In contrast, the Gradient Boosting Classifier recorded an accuracy of 45.5%, suggesting that while it is a popular choice for many tasks, it may not be the best fit for this specific dataset. The SVM (Support Vector Machine) attained an accuracy of 49.5%, demonstrating its potential but also revealing some limitations in this context. Similarly, the Random Forest Classifier also achieved an accuracy of 49.5%, showing comparable performance to SVM but falling short of the SGD Classifier. This visual representation allows stakeholders to quickly assess which algorithms are performing well and which may require further tuning or alternative approaches to enhance their predictive capabilities.

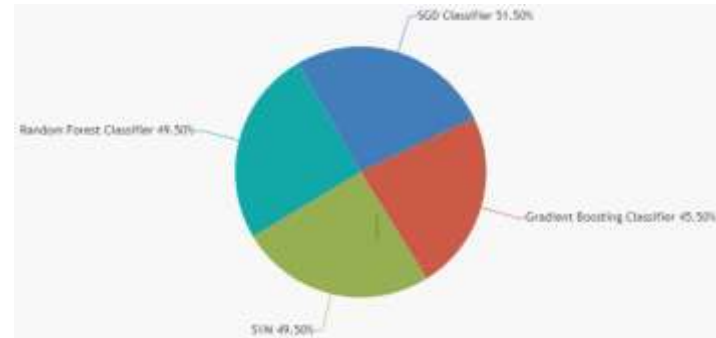


Fig: 6.1 Resultant graph

7. CONCLUSION

In conclusion, this study presents a novel framework for airway generation that significantly enhances the accuracy and reliability of airway classification within the complex structure of the human respiratory system. By leveraging the Prim algorithm to establish a minimum spanning tree and implementing a tree-pruning method based on subtree length, our approach effectively addresses the prevalent issues of false branch generation and inaccuracies in higher-generation airway assessments. The proposed methodology not only improves the precision of airway segmentation but also facilitates better diagnostics and treatment planning, particularly for small airways that are often challenging to evaluate. Through experimental validation on both public datasets and clinical outcomes from our private dataset, we demonstrate the robustness and effectiveness of our framework. Ultimately, this advancement holds substantial promise for improving patient health monitoring, disease prediction, and prevention, paving the way for more personalized and effective treatment strategies in respiratory care.

8. REFERENCES

- [1] S. Gupta, K. R. Sharma, and D. Jain, "Airway assessment: Predictors of difficult airway," *Indian J. Anaesthesia*, vol. 49, no. 4, pp. 257–262, 2005.
- [2] D. Couper, L. M. LaVange, M. Han, R. G. Barr, E. Bleecker, E. A. Hoffman, R. Kanner, E. Kleerup, F. J. Martinez, P. G. Woodruff, and S. Rennard, "Design of the subpopulations and intermediate outcomes in COPD study (SPIROMICS): Table 1," *Thorax*, vol. 69, no. 5, pp. 492–495, May 2014.
- [3] R. D. Miller and R. E. Hyatt, "Evaluation of obstructing lesions of the trachea and larynx by flow-volume loops," *Amer. Rev. Respiratory Disease*, vol. 108, no. 3, pp. 475–481, Aug. 1973.
- [4] A. P. Kiraly, W. E. Higgins, G. McLennan, E. A. Hoffman, and J. M. Reinhardt, "Three-dimensional human airway segmentation methods for clinical virtual bronchoscopy," *Academic Radiol.*, vol. 9, no. 10, pp. 1153–1168, Oct. 2002.
- [5] V. R. de Water, J. K. Saridin, F. Bouw, M. M. Murawska, and M. J. Koudstaal, "Measuring upper airway volume: Accuracy and reliability of dolphin 3D software compared to manual segmentation in craniosynostosis patients," *J. Oral Maxillofacial Surgery*, vol. 72, no. 1, pp. 139–144, Jan. 2014.
- [6] N. A. Alsufyani, A. Hess, M. Noga, N. Ray, M. A. Q. Al-Saleh, M. O. Lagravère, and P. W. Major, "New algorithm for semiautomatic segmentation of nasal cavity and pharyngeal airway in comparison with manual segmentation using cone-beam computed tomography," *Amer. J. Orthodontics Dentofacial Orthopedics*, vol. 150, no. 4, pp. 703–712, Oct. 2016.
- [7] A. Weissheimer, L. M. D. Menezes, G. T. Sameshima, R. Enciso, J. Pham, and D. Grauer, "Imaging software accuracy for 3-dimensional analysis of the upper airway," *Amer. J. Orthodontics Dentofacial Orthopedics*, vol. 142, no. 6, pp. 801–813, Dec. 2012.
- [8] J.-P. Charbonnier, E. M. V. Rikxoort, A. A. A. Setio, C. M. Schaefer-Prokop, B. V. Ginneken, and F. Ciompi, "Improving airway segmentation in computed tomography using leak detection with convolutional networks," *Med. Image Anal.*, vol. 36, pp. 52–60, Feb. 2017.