

# Leveraging Machine Learning to Identify Risk Factors for HPV-Related Oropharyngeal Cancer and Sex-Based Differences

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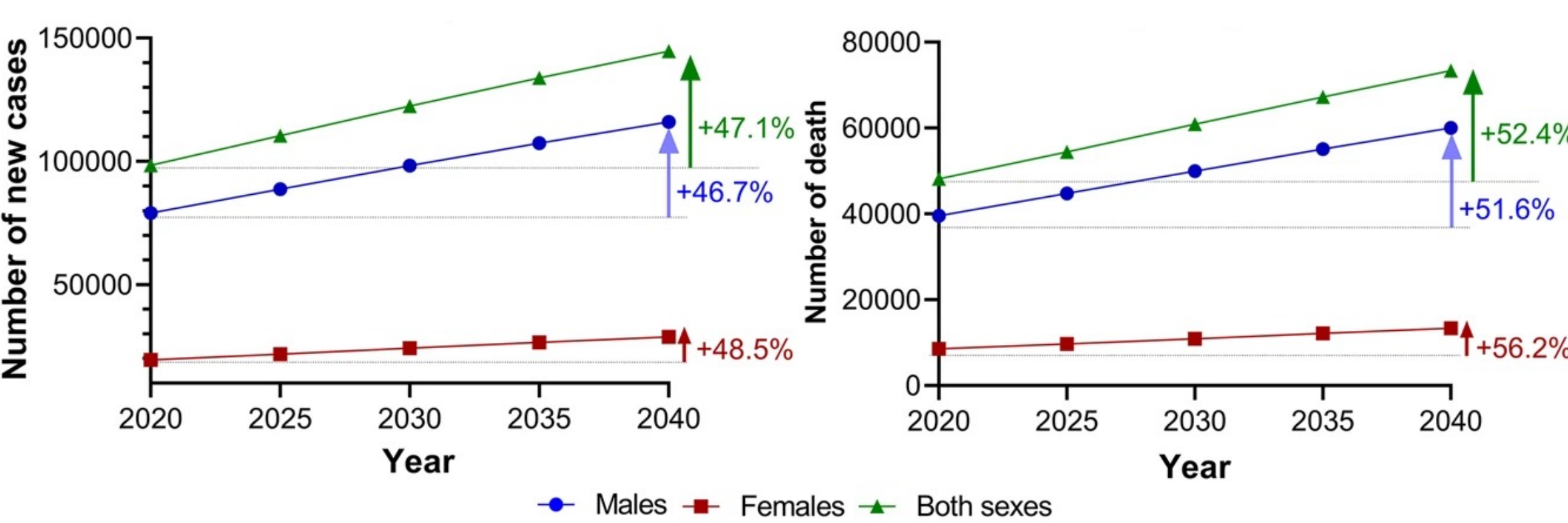
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## Background

Globally, the incidence and mortality rates of **oropharyngeal cancers** are expected to **rise by ~50%** in the next 20 years. In the US, the incidence of HPV-associated oropharyngeal cancer has surpassed that of cervical cancer, with a marked and unexplained sex disparity favoring **higher rates in men**.

20-Year Projection of Global Oropharyngeal Cases and Deaths<sup>1</sup>



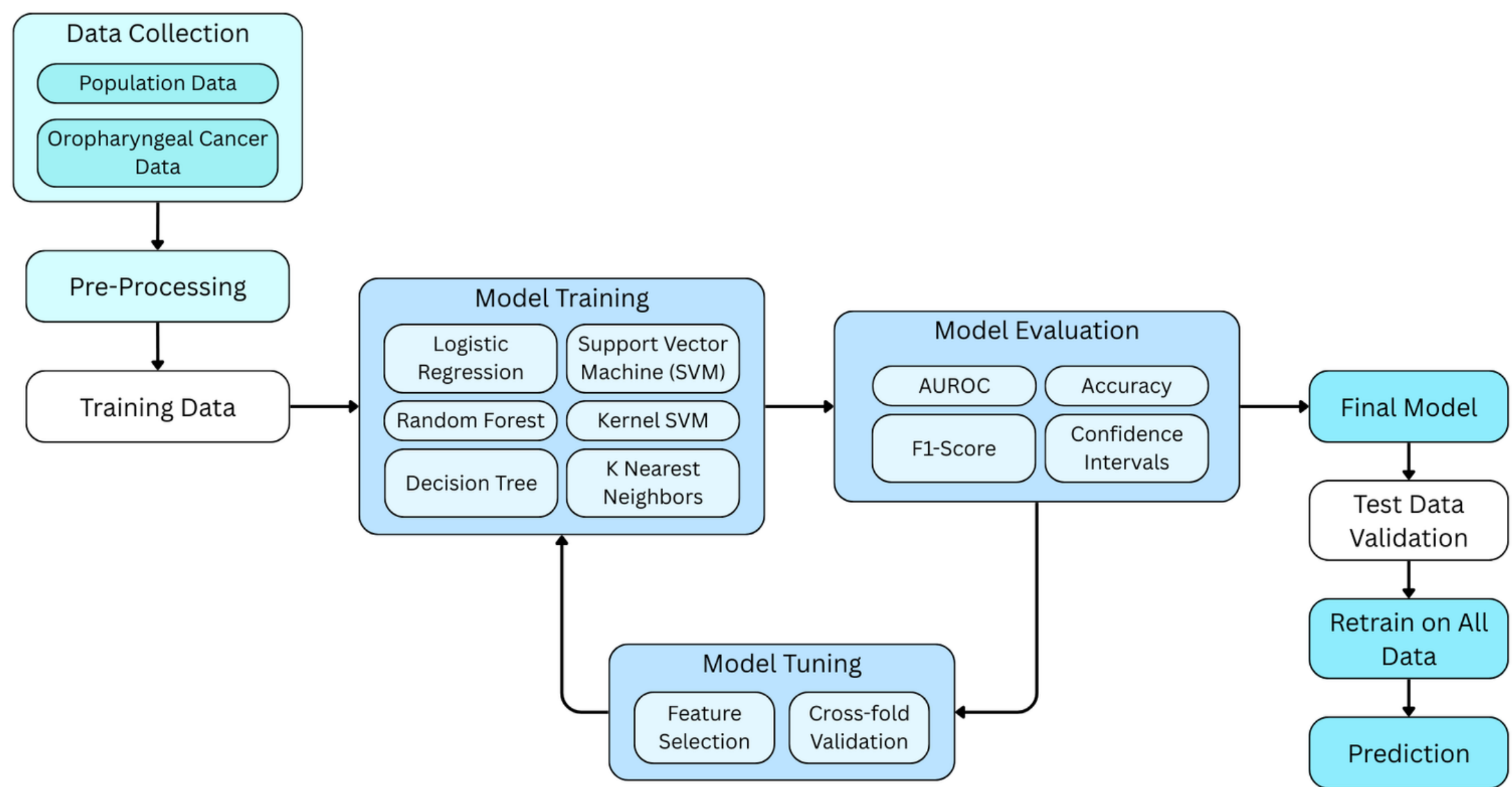
The underlying mechanisms likely involve complex sociodemographic, behavioral, and environmental interactions.

To better understand these contributors, this study employed **machine learning (ML)** techniques to identify **sex-specific community-level risk factors associated with oropharyngeal cancer incidence** in Massachusetts between 2001 and 2022.

## Methods

We performed a **retrospective analysis** of Massachusetts public health data from 2001 to 2022.

- Population data across 352 towns and oropharyngeal cancer incidences were sourced from the Massachusetts Health Data and Environmental Public Health Tracking Tools.
- Six ML models were trained, evaluated, and tuned (dimensionality reduction, GridSearchCV) on training data only, then validated on test data.
- The most predictive models were used to identify key risk factors stratified by sex.



## Results

Across all Massachusetts data, the K Nearest Neighbors model performed best for males:

- AUROC: 0.666 (CI: 0.589–0.738)
- Accuracy: 0.663 (CI: 0.596–0.725)
- F1-Score: 0.504 (CI: 0.403–0.600)

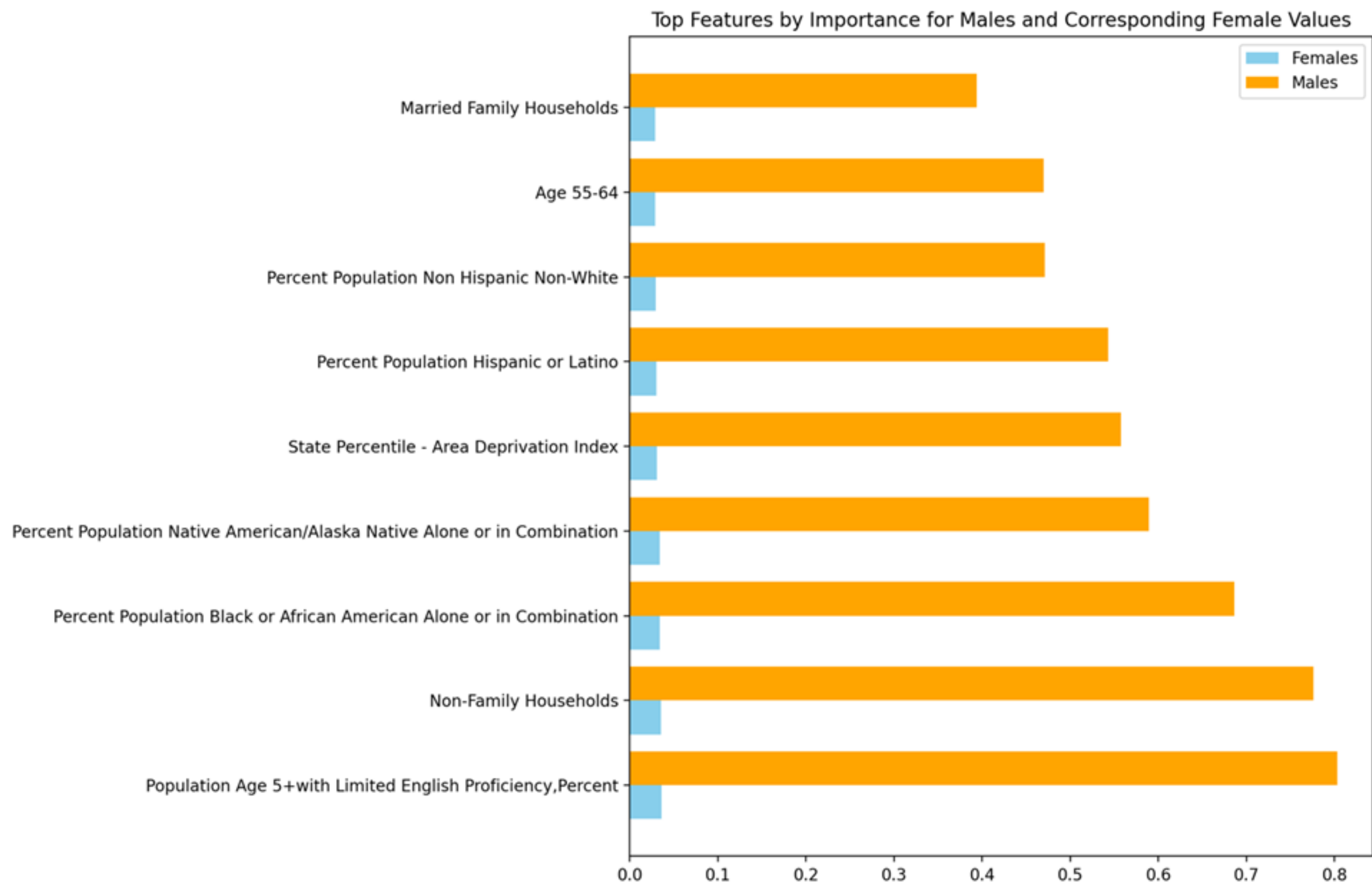
For males, key risk factors included:

- Limited English proficiency
- Low educational attainment
- Poverty
- Physical Inactivity
- Being of non-white race
- Residing in non-family households

For females, the Random Forest model showed the best performance:

- AUROC: 0.636 (CI: 0.559–0.709)
- Accuracy of 0.636 (CI: 0.554–0.683)
- F1-score of 0.503 (CI: 0.411–0.584)

However, no significant predictors were identified for females, highlighting distinct epidemiological pathways.



Coefficient magnitudes of the top 9 features driving increased oropharyngeal cancer incidence in males, alongside their corresponding coefficient magnitudes for females using the same variable set, demonstrating distinct epidemiological pathways of malignancy.

**Key takeaway:** Strong community-level predictors of oropharyngeal cancer incidence were identified in males, but associations remain unclear for females, pointing towards a potential sex disparity in how environmental and social factors influence cancer risk.

## Discussion

The risk factors identified in this study reflect **broader social determinants** that influence behaviors like tobacco and alcohol use, which are known contributors to HPV-related oropharyngeal cancer.<sup>2</sup>

While it is well-known that HPV-associated oropharyngeal cancers predominantly affect males, this **sex disparity remains unclear** and cannot be solely explained by behavioral patterns.<sup>3</sup> In Massachusetts, this disparity in HPV-associated cancers is especially pronounced, with rising male oropharyngeal cancer rates surpassing national trends, while cervical cancer rates among females continue to decline.<sup>4</sup>

## Future Directions

- Further Investigation: Underlying Drivers of Sex Disparities
  - Explore the complex, sex-specific interactions among HPV exposure, behavioral risk factors, and access to preventive care.
- Public Health Efforts: HPV Vaccination Coverage and Preventative Services
  - Emphasize the vaccine's role in preventing both cervical and oropharyngeal cancers and promote vaccine completion, particularly among racial and ethnic minorities.

## Acknowledgements

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For any further questions, please feel free to contact Austin Coppinger ([austin\\_coppinger@brown.edu](mailto:austin_coppinger@brown.edu)).

### References:

- [Figure adapted from] Lim, Y.X., D'Silva, N.J. HPV-associated oropharyngeal cancer: in search of surrogate biomarkers for early lesions. *Oncogene* 43, 543–554 (2024). <https://doi.org/10.1038/s41388-023-02927-9>
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