

# Voice Enabled Artificial Intelligence for Detection of Pathologic Gastroesophageal **Reflux Disease and Barrett's Esophagus**

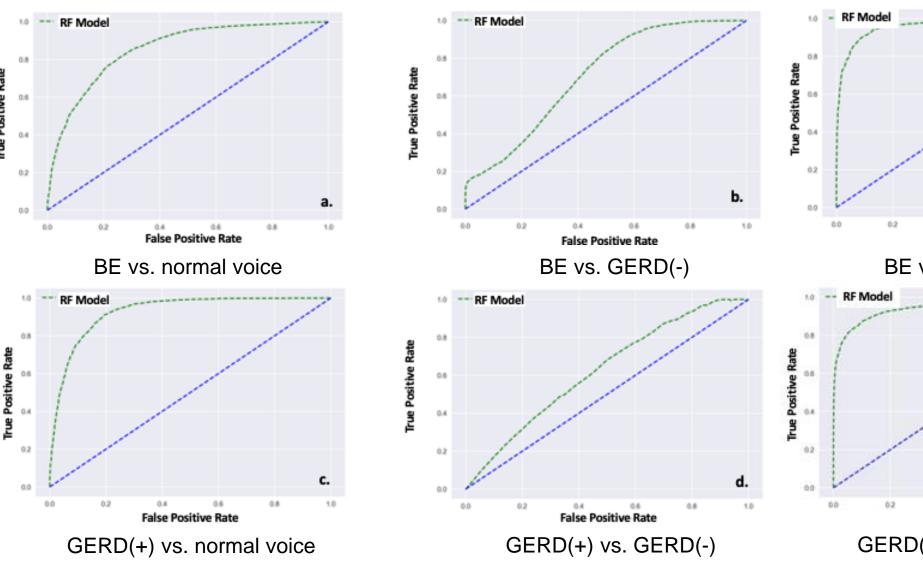
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### BACKGROUND

- reflux Gastroesophageal disease (GERD) can lead to voice alterations, including hoarseness.
- The study aim was to identify specific voice biomarkers associated with pathologic GERD using advanced machine learning tools.
- Voice biomarkers reflect the periodicity of the voice signal, which mirrors the quality of the voice. The lesser the periodicity, the more noisy the sound quality (e.g., hoarseness, breathiness, roughness, creakiness).
- Detection of pathologic GERD, including Barrett's esophagus (BE), with voice biomarkers can serve as a simple noninvasive screening tool.

## **STUDY SUBJECTS**

- Voice recordings were obtained from patients undergoing clinically indicated esophagogastroduodenoscopy (EGD) and/or ambulatory pH monitoring studies.
- Patients were excluded if they had another condition (pulmonary, cardiac, neurologic, etc) associated with voice disturbance.
- Voice recording consisted of a 6sentence standard script read over 25-45 seconds.



#### **BASELINE CHARACTERISTICS**

Subgroup	Sex	Number of patients	Mean age, years (SD)
Barrett's	Female	13	65 (15)
esophagus	Male	22	67 (9)
GERD (+)	Female	23	51 (14)
	Male	11	53 (19)
GERD (-)	Female	48	55 (15)
	Male	30	59 (15)
Vocally	Female	64	28 (11)
normal	Male	34	34 (15)

 
 Table 1. Baseline patient characteristics in study
subgroups.

Figure 1. Receiver Operating Characteristics for a Random Forest (RF) Model for males (a-d) and females (e-h).

#### **STUDY GROUPS**

- GERD(+) patients were defined as those with erosive esophagitis (LA grade B-D) or peptic stricture or acid exposure time >6%.
- BE was defined as columnar mucosa >1 cm with confirmed specialized intestinal metaplasia.
- · Patients without these findings were considered GERD(-).
- · A vocally normal group consisting of individuals with normal voice as judged by speech pathology evaluation was used as an independent control group.

### **STATISTICAL ANALYSIS**

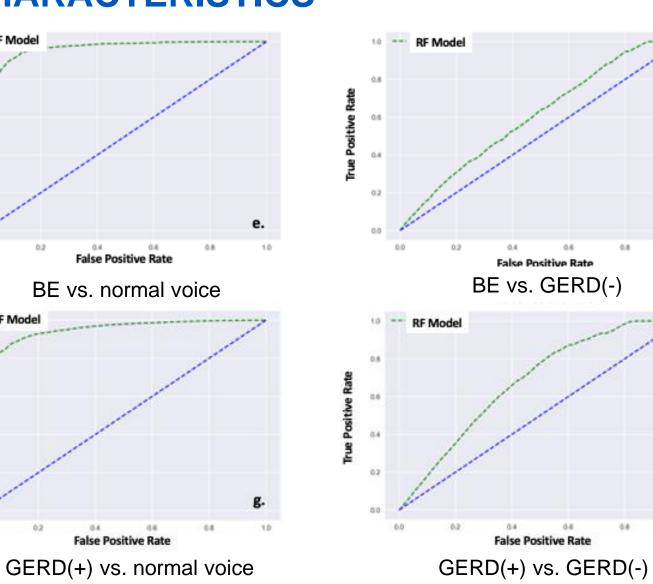
- Random forest models were trained using a balanced number of subjects per condition using random participant selection from the majority class.
- Using a 5-fold nested cross validation strategy, features were selected and ranked within fold, and a series of models were trained within each fold using recursive feature elimination.
- The average F1 score, a harmonic mean of precision and recall (range 0-100), across all folds was reported to assess performance.

### RESULTS

- from 60-70.

# **RECEIVER OPERATING CHARACTERISTICS**

**False Positive Rate** 







• The study sample consisted of 245 patients (vocally normal, n=98; GERD(-), n=78; GERD(+), n=34; BE, n=35).

Feature rankings suggested voice quality differences between groups relating to voice signal periodicity.

• The model demonstrated excellent ability to discern BE and GERD(+) from the vocally normal group with F1 scores 82 (males) and 89 (females) and 80 (males) and 80 (females) for BE and GERD(+) respectively.

• There was also a good voice signal distinguishing BE and GERD(+) groups from GERD(-) with F1 scores ranging

#### DISCUSSION

 These results suggest that voice biomarkers may be useful as a non-invasive tool in the detection of pathologic GERD/BE.

• A deep learning diagnostic model will be developed using the identified voice biomarkers.