

Epigenetic Alterations in Barretts Esophagus: Novel screening test for Barretts esophagus and cancer

William Grady, MD, AGAF

Ming Yu, PhD

Fred Hutchinson Cancer Research Center
University of Washington School of Medicine



FRED HUTCH[™]
CURES START HERE

W
UNIVERSITY of
WASHINGTON

Early
Detection
Research
Network



 **BETRNet**

Disclosure Information

William M. Grady

I have the following financial relationships to disclose:

| | |
|----------|-----------------------|
| Freenome | Advisory Board Member |
|----------|-----------------------|

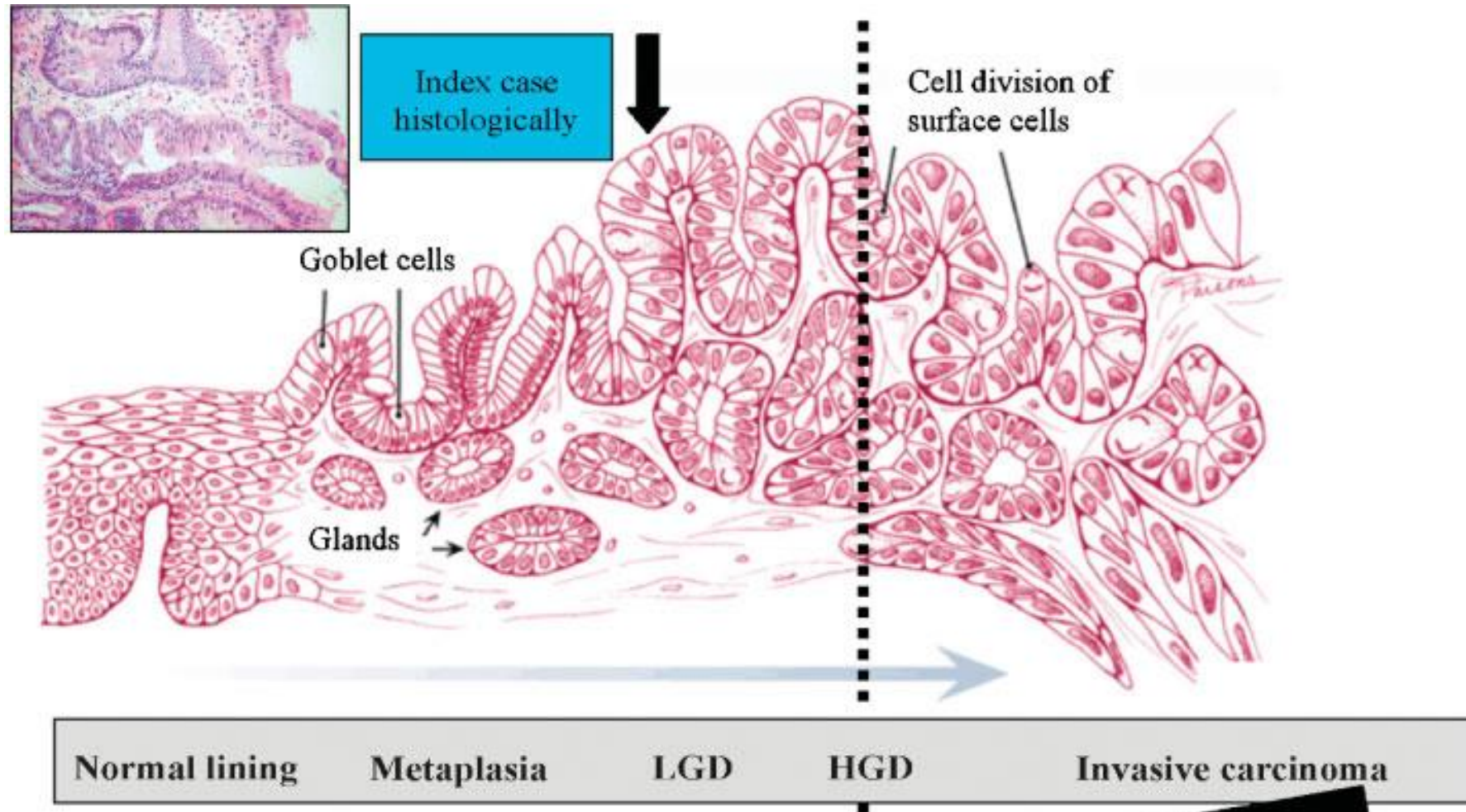
| | |
|---------|-----------------------|
| SEngine | Advisory Board Member |
|---------|-----------------------|

| | |
|-----------------|-----------------------|
| Guardant Health | Advisory Board Member |
|-----------------|-----------------------|

| | |
|----------|-----------------------|
| Diacarta | Advisory Board Member |
|----------|-----------------------|

| | |
|-------------------------|-------------------------------------|
| Clinical trial support: | Janssen, Tempus, Lucid Technologies |
|-------------------------|-------------------------------------|

BE→EAC progression sequence



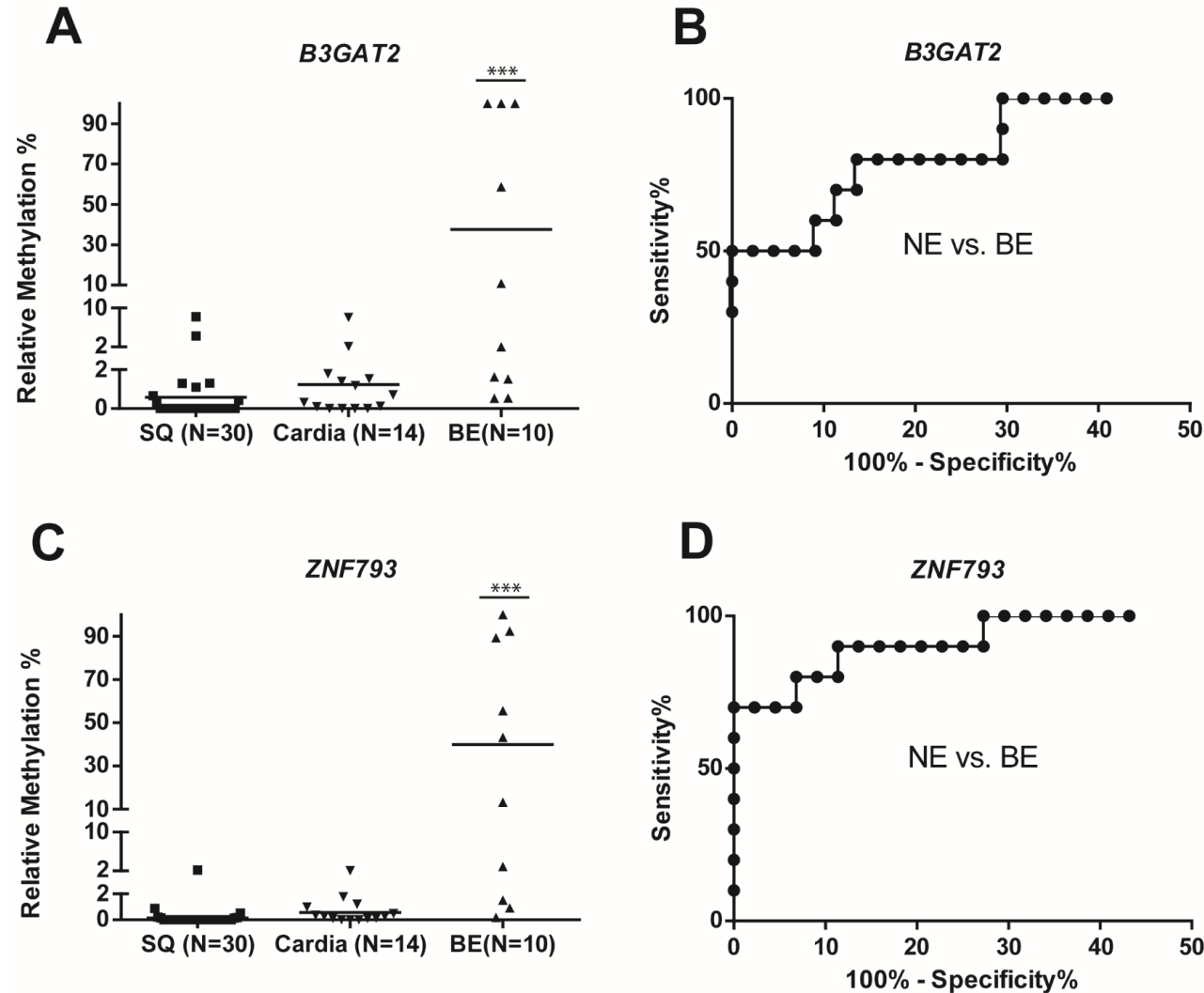
Driven by genetic and epigenetic alterations=**candidate biomarkers**

Methylated genes:

High potential for screening and surveillance biomarkers

- Aberrant methylation is common in both BE and EAC and more common in EAC
- Assay technology optimal for clinical assays
- Feasibility demonstrated for colon cancer screening
 - Cologuard assay
 - Epi proColon assay

Methylated *ZNF793* and *B3GAT2*: BE Detection Markers



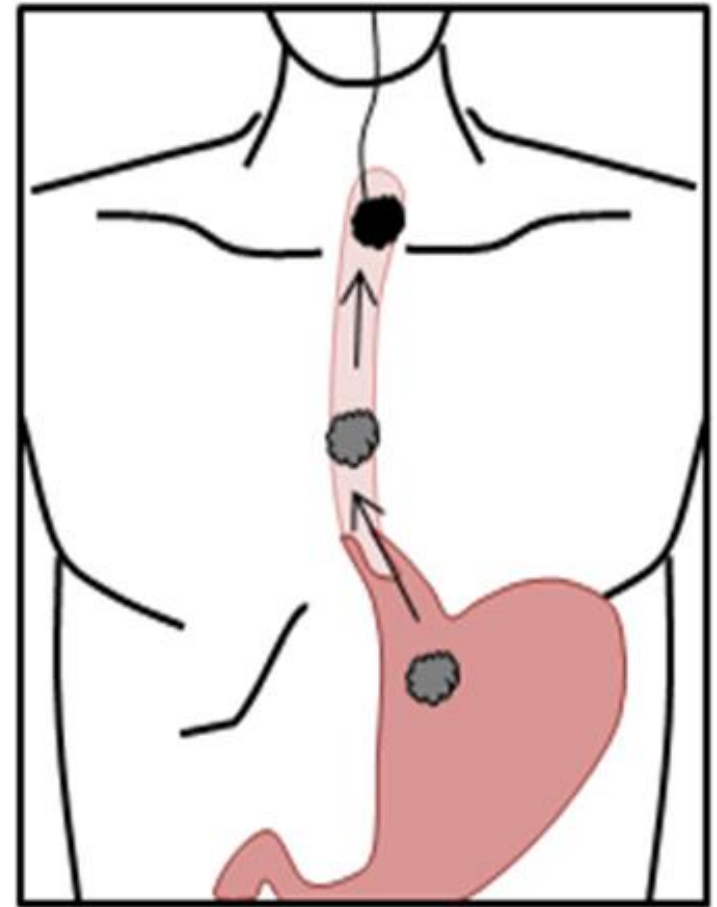
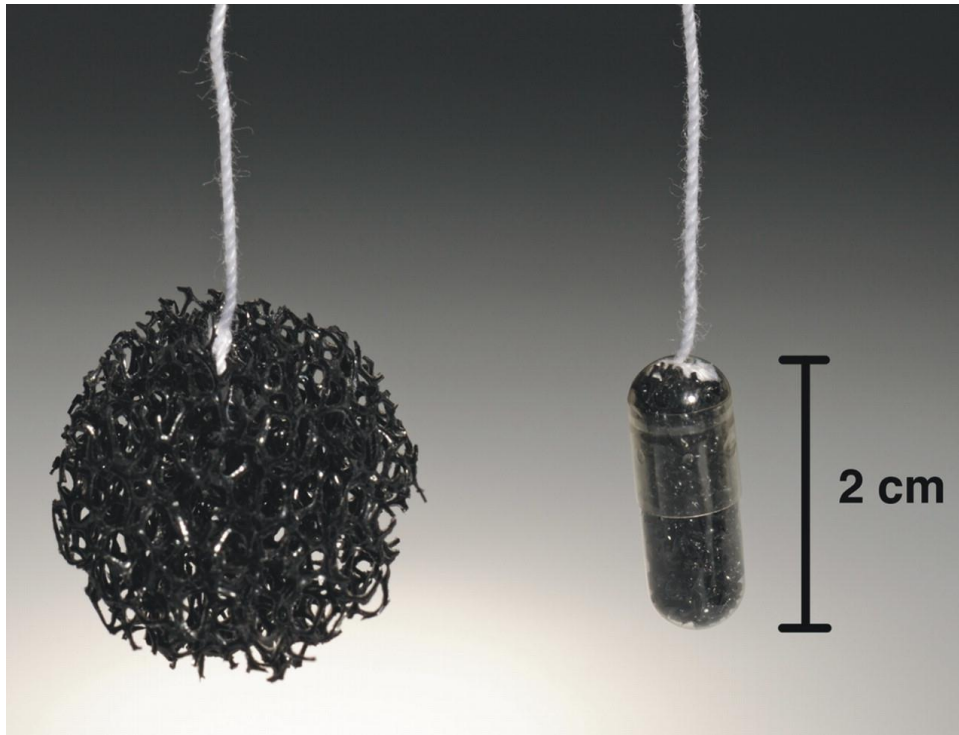
Promising candidate BE screening markers

EDRN Phase 2/3 Biomarkers

- TFF3
- Methylated *VIM* and *CCNA1*
- Methylated *B3GAT2* and *ZNF793*
- Clinical trials-EDRN and BETRNET sponsored

1st generation non-endoscopic tests for BE screening

Cytosponge



2nd generation device: Esocheck balloon

- **Esochek** combines a swallowed balloon to sample the esophagus with a DNA based biomarker test for diagnosing BE: **Esoguard** (FDA approved)
- Additional biomarkers to catch progression of BE to EAC in development
→ potential to create a panel for BE detection and continued monitoring

“ Esophageal Pap Smear to detect BE / EAC ”



Tethered, capsule sized balloon is swallowed



Balloon with structured surface is inflated in the stomach and withdrawn to resistance at distal esophagus to retrieve sample



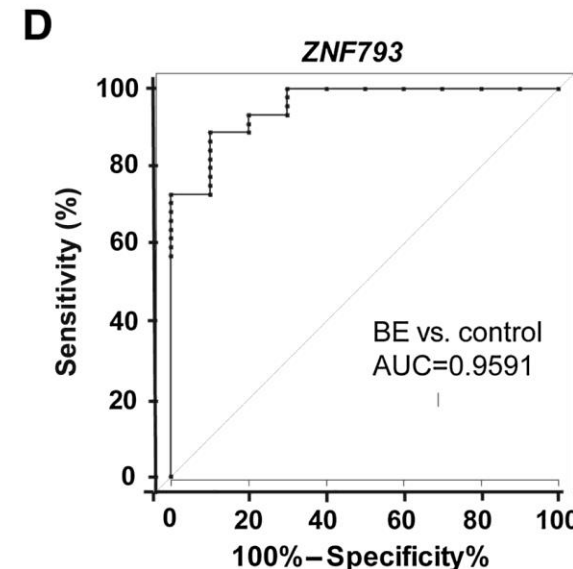
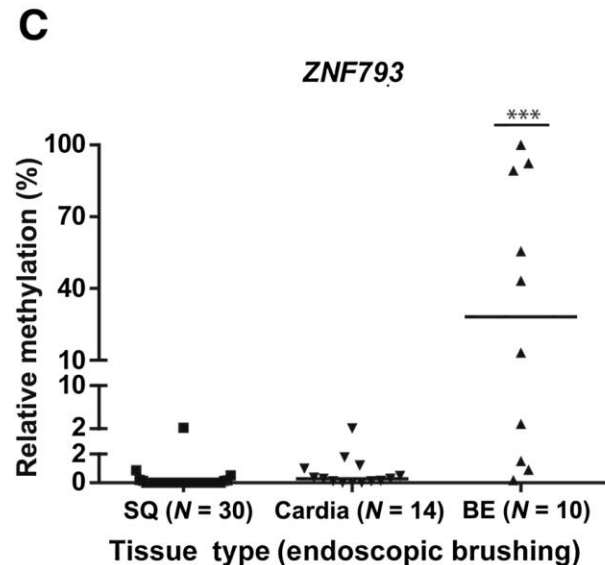
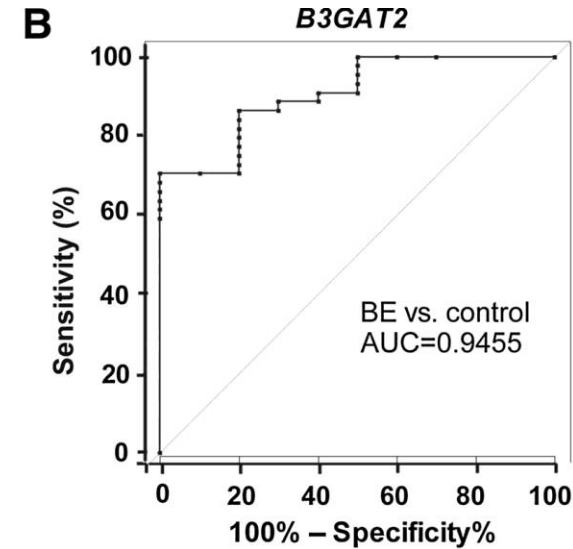
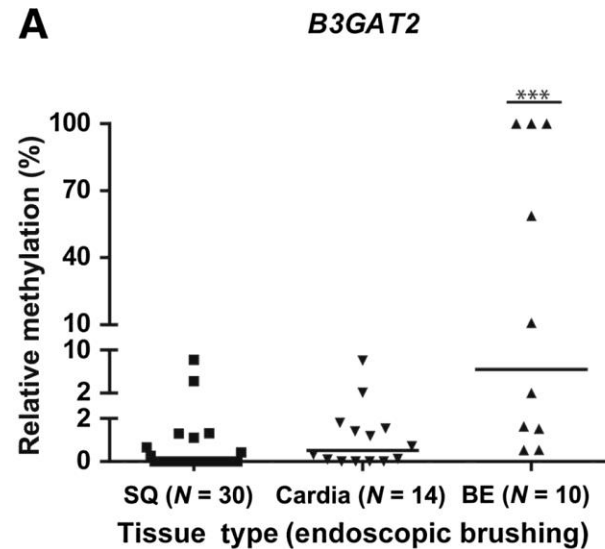
Inverted to protect sample on retrieval



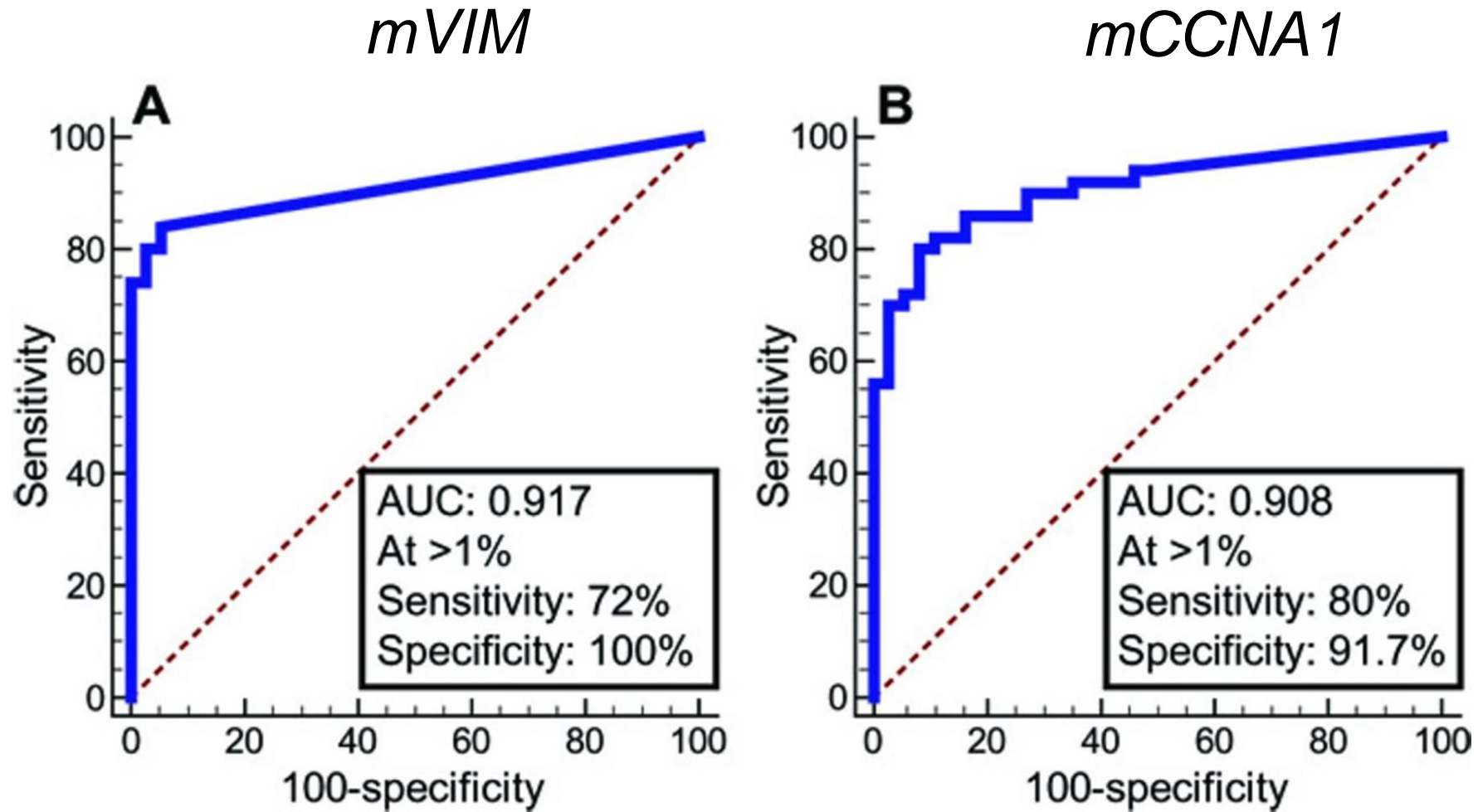
Sample sent to lab for DNA assay

Methylated *ZNF793* and *B3GAT2*: Accurate BE markers in esophageal brushings

Sensitivity=100%
Specificity =70-73%



mVIM and *mCCNA1*: accurate for BE screening



Sensitivity=90% and Specificity =91%

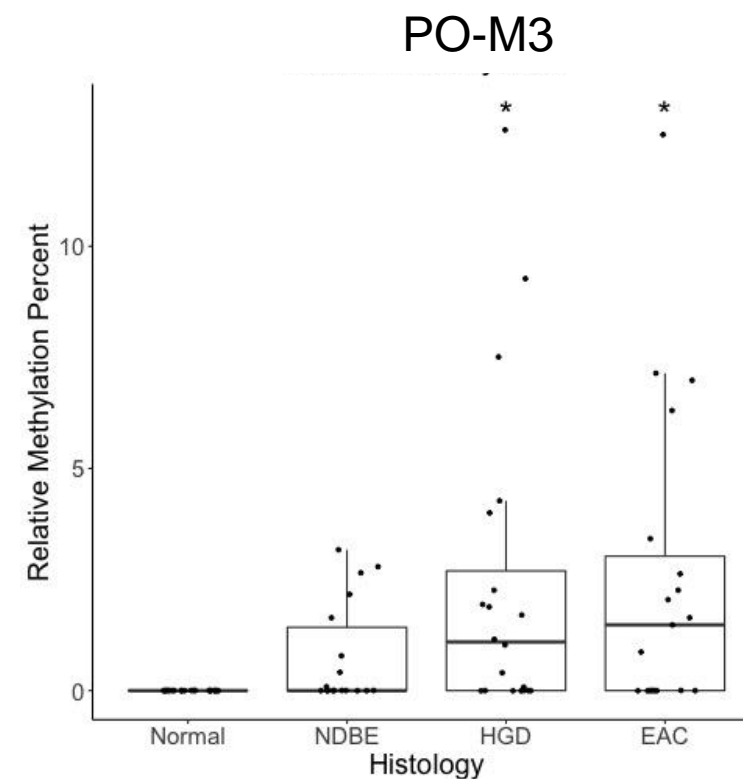
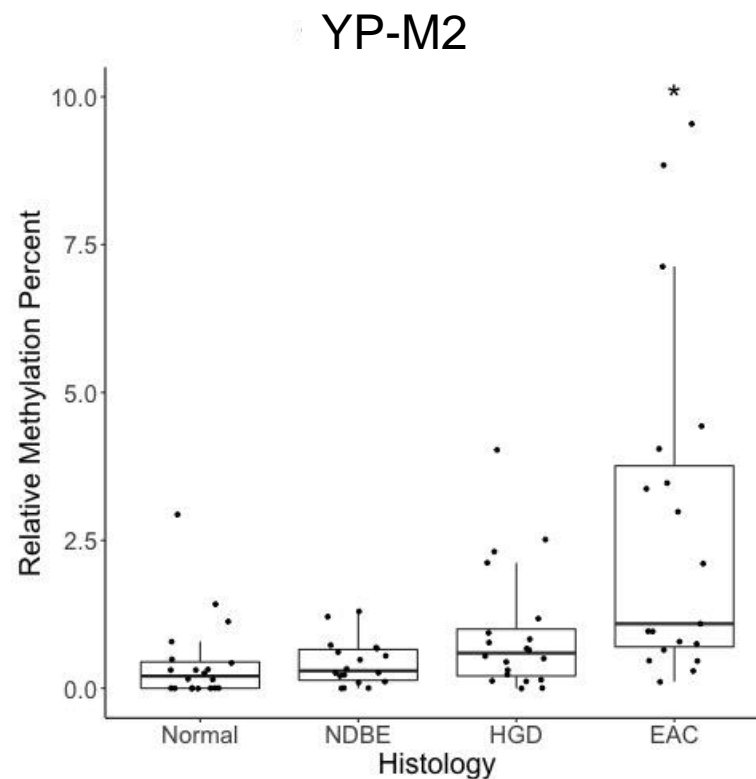
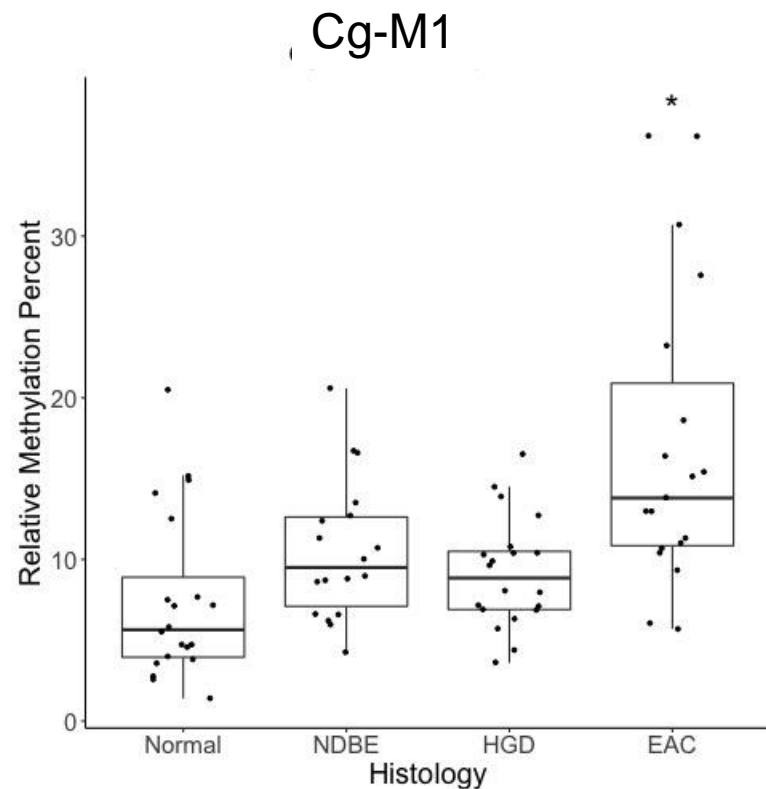
Candidate BE surveillance markers

Detection of high grade dysplasia or cancer

- P53 protein expression-IHC
- *TP53* mutations
- Aneuploidy/tetraploidy
- 17pLOH
- Methylated genes
 - *Cg65-M1*
 - *YP-M2*
 - *PO-M3*

Methylation status BE surveillance markers

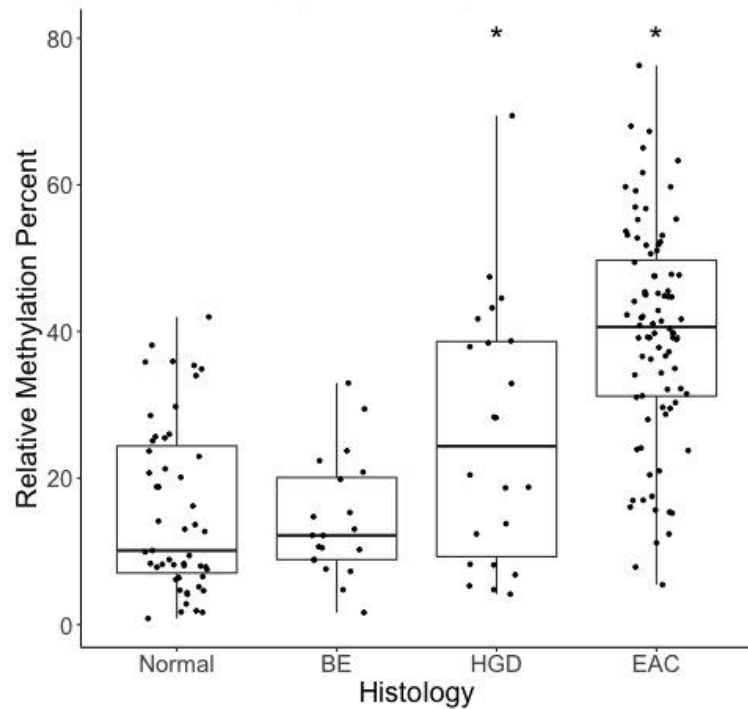
Training Set Cohort



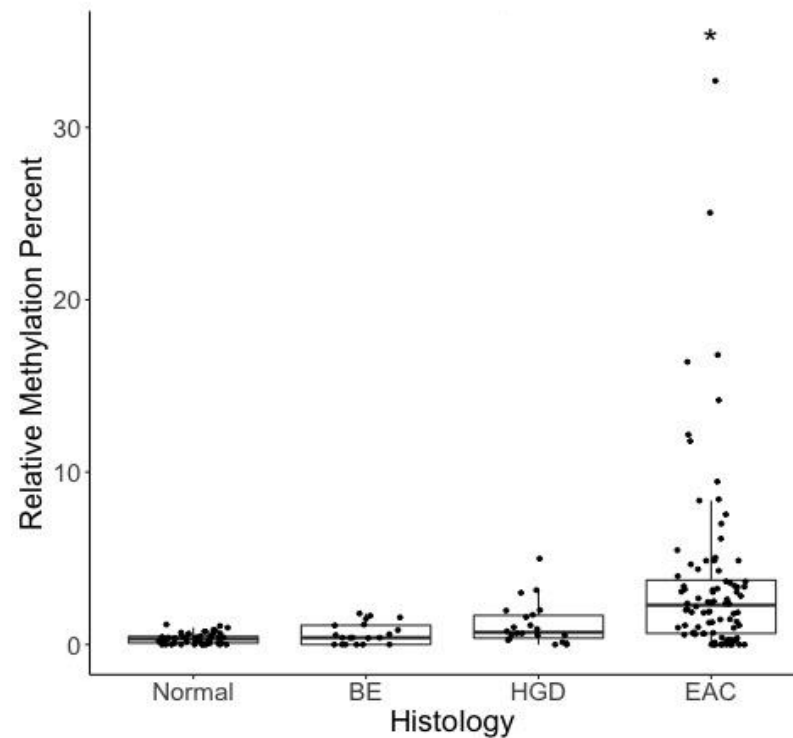
Methylation status BE surveillance markers

Validation Set Cohort

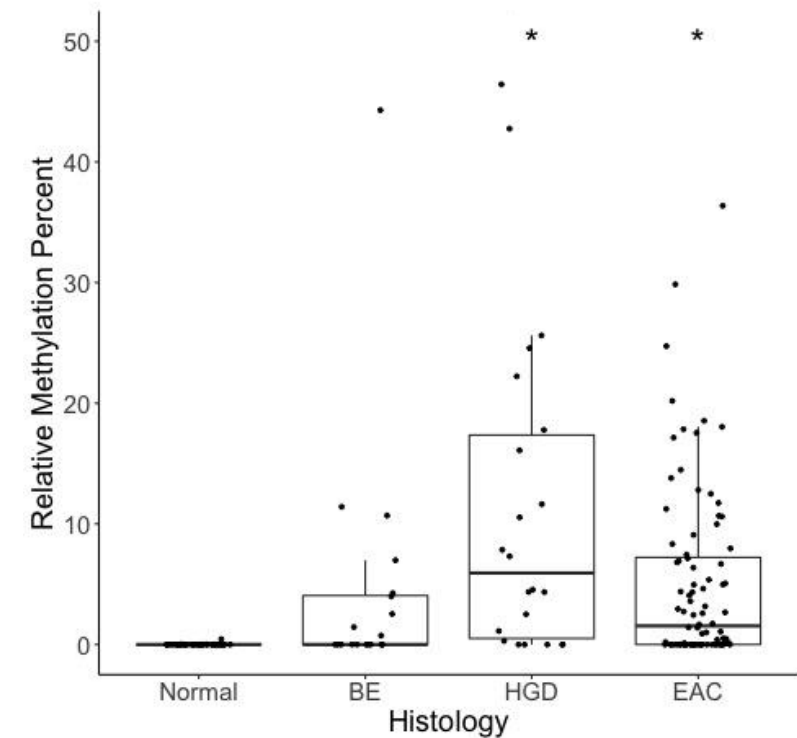
Cg-M1



YP-M2

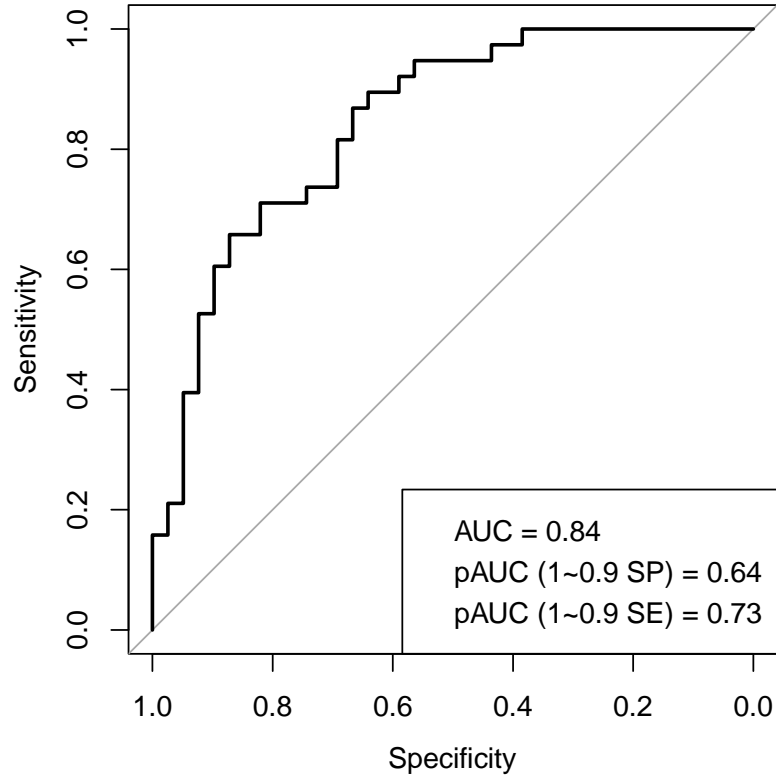


PO-M3

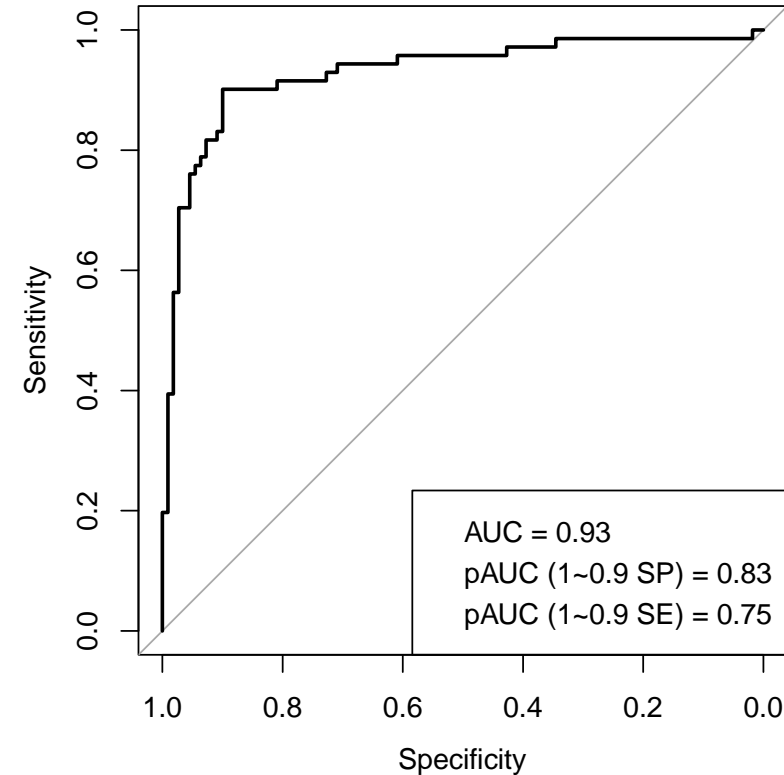


Candidate BE surveillance 3 marker panel

Training Set



Validation Set

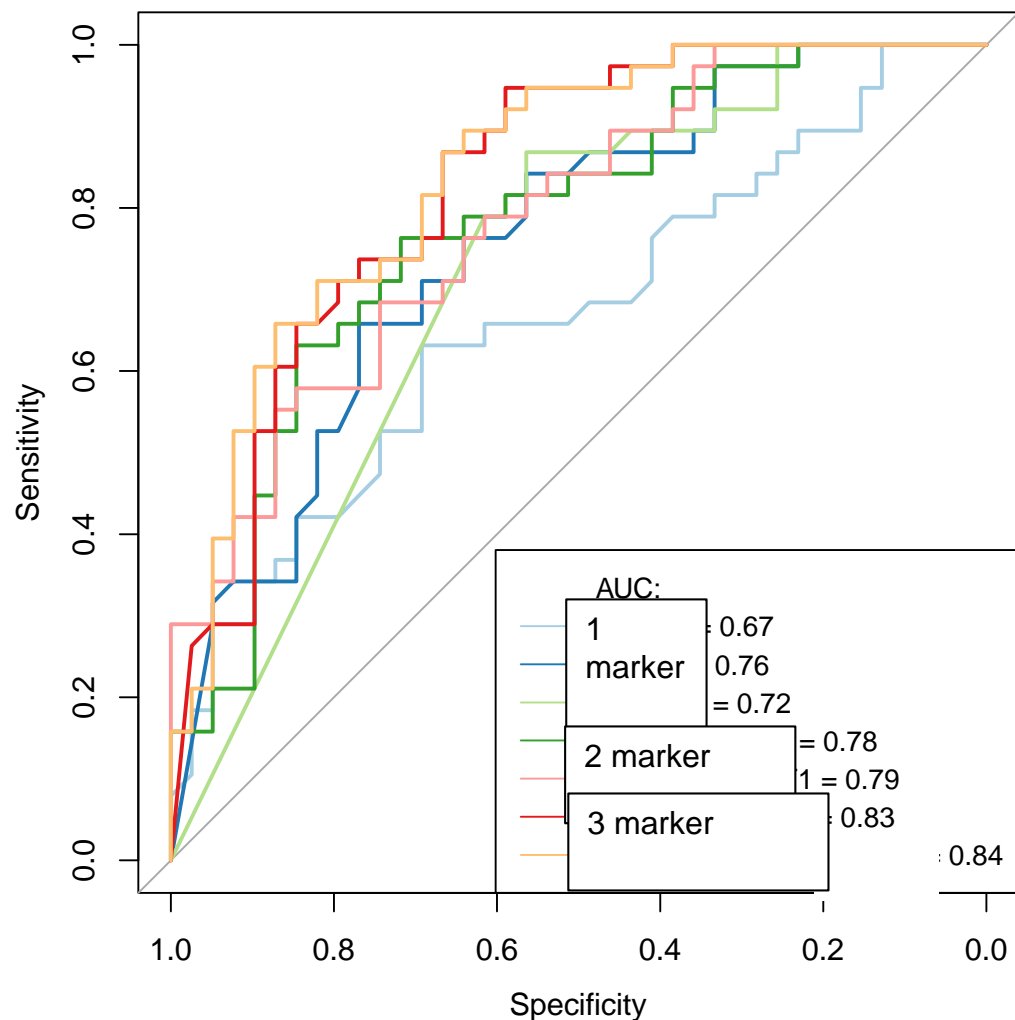


| Accuracy = 0.74 Sensitivity = 0.67 Specificity = 0.82 | | True | |
|---|---------|--------|------|
| | | HGDEAC | SQBE |
| Predicted | HGD+EAC | 26 | 7 |
| | SQ+BE | 13 | 31 |

| Accuracy = 0.88 Sensitivity = 0.93 Specificity = 0.80 | | True | |
|---|---------|---------|-------|
| | | HGD+EAC | SQ+BE |
| Predicted | HGD+EAC | 102 | 14 |
| | SQ+BE | 8 | 57 |

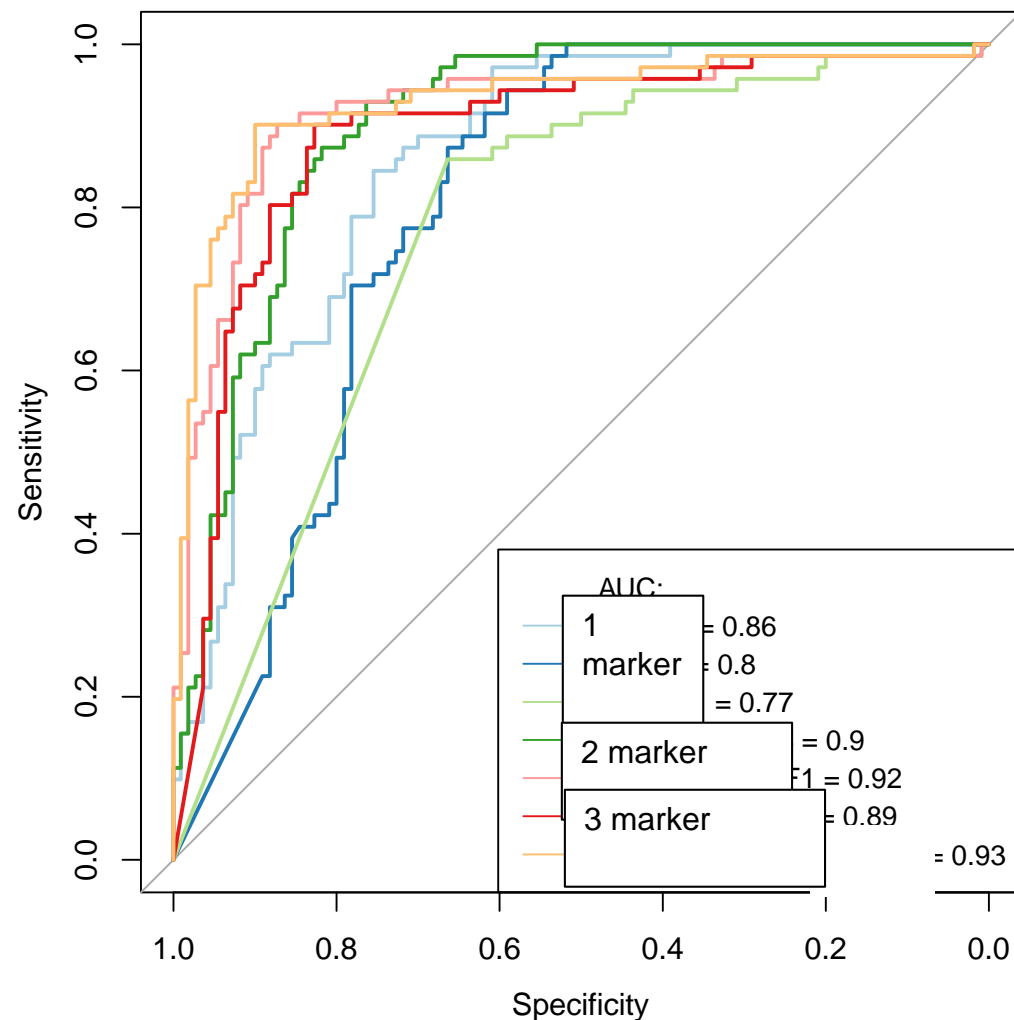
Candidate BE surveillance 3 marker panel ROC

Training Set ROC



Validation Set ROC

1 marker



Epigenetic alterations in Barretts esophagus: Insights into molecular mechanisms and their potential use in the clinic.

- **BE screening biomarkers**

- Identify at risk population using safe, cheap test
- Most promising marker: TFF3 cytology
- Emerging markers: methylated genes-*VIM*, *CCNA1*, *B3GAT2*, *ZNF793*

- **BE surveillance biomarkers**

- Identify high-risk BE patients
- Prevent overtreatment of low-risk BE patients
- Most promising markers: *TP53* mutations, aneuploidy, LOH events, methylated genes

Acknowledgements

FHCRC

- Ming Yu
- Sean Maden
- Jess Ayers
- Kelly Carter
- Ting Wang
- Amber Willbanks
- Tai Heizerling

Collaborators

Harvard/DFCI

- Adam Bass
- Matt Stachler

CWRU

- Amitabh Chak
- Joseph E. Willis
- Apoorva K. Chandar
- Robin Elliot
- Sandy Markowitz
- Helen Moinova

UW

- John Inadomi
- Mike Saunders
- Adam Templeton
- Deepti Reddi
- Maria Westerhoff
- Staff at UW, HMC, ESC GI clinics and units
- GICaRes Study Team: Wynn Burke, Brian Foerster, Colton Johnson, Lucilla Bella, Lito Morada



Table 1: Patient Demographic Information

| Subject Information | Discovery set | Training set | Validation set |
|-------------------------|---------------|--------------|----------------|
| Total | 167 | 76 | 181 |
| Gender | | | |
| Female | 32 | 10 | 55 |
| Male | 135 | 66 | 126 |
| Age | | | |
| Range (Mean) | 21-93 (64) | 46-82 (64) | 23-89 (64) |
| BMI | | | |
| Range (Mean) | 21-40 (29) | 19-57 (31) | NA |
| NA | 131 | 3 | |
| Smoking | | | |
| Yes (current or former) | 36 | 47 | 105 |
| Never | 15 | 25 | 68 |
| NA | 116 | 4 | 8 |
| NSAID Use | | | |
| Yes | NA | 39 | NA |
| No | | 35 | |
| NA | | 2 | |
| Diabetes (I or II) | | | |
| Yes | NA | 17 | NA |
| No | | 59 | |
| NA | | 0 | |

- There is no significant difference between HGD/EAC vs SQ/BE for any variable in the three data sets ($p > 0.05$). Chi-Square test for category variables, ANOVA F-test for numerical variables.
- BMI=Body Mass Index; NSAID=nonsteroidal anti-inflammatory drug.



5 minute Q&A

Chair/Co-Chair/NCI

feed Zoom Chat questions to presenter
and Track Time

NCI and Production Team

answer Chat questions not related to presentations
and use Slack



SWOT

Colon & Esophageal
40 Minutes