

A novel serum glycoprotein biomarker panel for screening of esophageal adenocarcinoma and surveillance of Barrett's esophagus

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Disclosures



- ▶ The biomarker concentrations were measured by Proteomics International, and the risk scores were calculated using a proprietary algorithm
- ▶ This analysis used archived samples received via a collaboration with the QIMR Berghofer Medical Research Institute
- ▶ The study was funded by Proteomics International
- ▶ Presenter disclosures: Lipscombe is an employee and shareholder of Proteomics International Laboratories Ltd, which is the license owner of a patent covering the use of the biomarkers, consequently, Lipscombe may receive financial benefit from the commercial use of any test

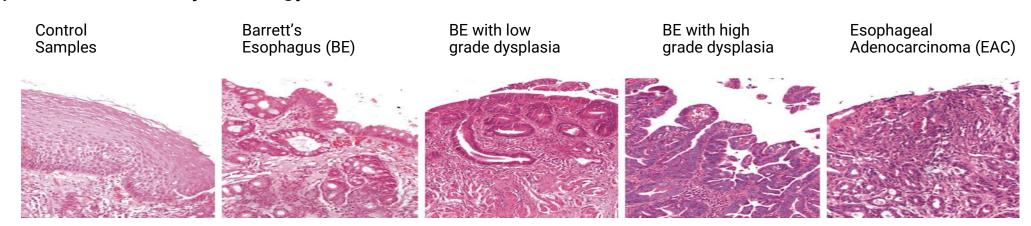




Introduction



- The overall five-year survival rate for esophageal cancer is approximately 20%
- An estimated 10-15% of patients with chronic acid reflux develop Barrett's Esophagus (BE), which has estimated prevalence of 1-2%.
- Currently patients with BE usually undergo endoscopy-biopsy surveillance with the degree of dysplasia assessed by histology



Aim

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- To develop a simple blood test for the diagnosis of esophageal adenocarcinoma (EAC) and Barrett's Esophagus (BE)
- To validate diagnostics models to screen EAC and BE based on a panel of glycoprotein biomarkers

Study Design



- **Development cohort (N=249):** Participants from The Progression of Barrett's Esophagus to Cancer Network (PROBE-NET) study, Australia
- **Validation cohort (N=49):** Participants collected at Ochsner Healthy System, New Orleans, United States

Group 1 - Controls Samples with normal endoscopy

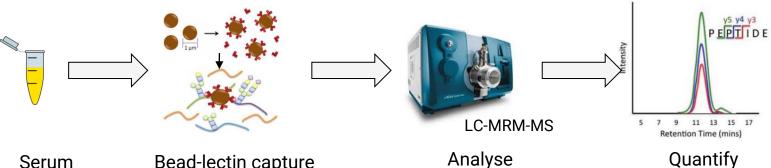
Group 2 BE without/with low grade dysplasia

Group 3 BE with high grade dysplasia

Group 4 Esophageal Adenocarcinoma

Samples tested by endoscopy and biopsy

- Potential biomarkers analysed using a proteomics workflow, adapted for glycoprotein selection using a lectin magnetic bead array followed by targeted mass spectrometry (LC-MRM-MS)
- Multivariate regression performed



Robust

- Intraday CV is 9.3%
- Interday CV is 11.5%

Results - I



Development cohort (N=249)

- Five glycoproteins were found to show statistical correlation with disease state progression
- Three of these proteins remained significant after adjusting for patient age and sex

Validation cohort (N=49)

Controls versus EAC (Group 1 vs Group 4):

Model performance (N=24)

Validation cohort:

AUC (95% CI)	0.94 (0.85 – 1.00)			
Sensitivity (%)	80%			
Specificity (%)	93%			
PPV(%)	89%			
NPV(%)	87%			

Controls versus BE-HGD & EAC (Group 1 vs Group 3+4):

Model performance (N=31)

Validation cohort:	
AUC (95% CI)	0.79 (0.70 - 0.88)
Sensitivity (%)	59%
Specificity (%)	93%
PPV(%)	91%
NPV(%)	65%

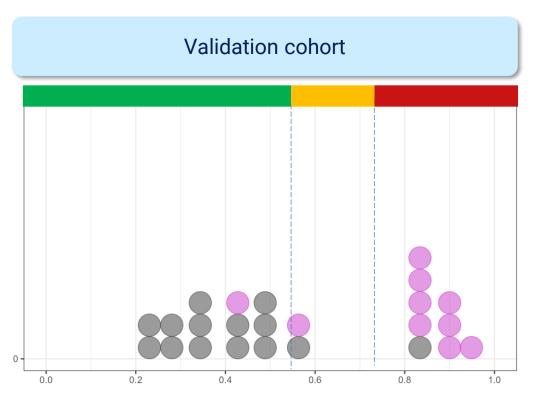


Results - II



Controls versus EAC (Group 1 vs Group 4):

Frequency dot plot of Control and EAC



n = 14 controls

n = 10 EAC

Classification	EAC	Controls	Accuracy
Red High probability	8	1	89%
Amber Moderate	1	1	
Green Low probability	1	12	92%

Traffic light system indicates if a patient is at risk of having EAC



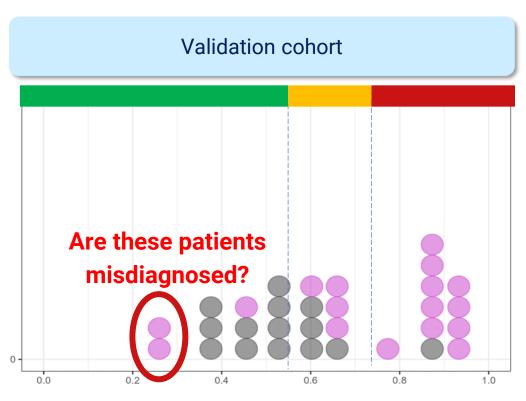
Results - III

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Controls versus BE-HGD & EAC (Group 1 vs Group 3+4):

Frequency dot plot of Control and BE-HGD & EAC



n = 14 controls

n = 17 BE-HGD & EAC

Classification	BE-HGD & EAC	Controls	Accuracy
Red High probability	10	1	91%
Amber Moderate	4	4	
Green Low probability	3	9	75%

Traffic light system indicates if a patient is at risk of having BE-HGD & EAC

Conclusions



- Panel of novel serum biomarkers (glycoproteins) were validated as showing significant correlation with esophageal adenocarcinoma and Barrett's esophagus with high grade dysplasia
- Diagnostic model developed as a potential simple screening tool for differentiating patients:
 - > with EAC from negative controls
 - > who required endoscopic surveillance (BE-HGD & EAC) from negative controls
- Further work required on larger cohorts to:
 - confirm diagnostic accuracy of test
 - > establish a potential diagnostic tool to differentiate patients who require endoscopic surveillance (BE-HGD & EAC) from patients having BE with *low* grade/no dysplasia









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