Why did all biomarker identification studies fail in endometriosis?

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Diagnosis of endometriosis

• Current gold standard:

 Laparoscopic visualization of endometriotic lesions, followed by histological confirmation

Advantage

- Can remove the lesion after seeing it

Downside

- Invasive
- Carries risks of morbidity and, very rarely, mortality
- Expensive
- The accuracy depends on the experience and skill levels of the surgeon

Dx delay

- Not suitable for screening purpose
- Incidence estimate difficult
- Makes the monitoring of treatment difficult
- Problem in determining recurrence

The need for non-invasive biomarkers

- Non-invasive
- Typically much less expensive than laparoscopy
- Shorten or eliminate the diagnostic delay

Applications of biomarkers

- Diagnosis
- Prognosis
 - Fertility outcome
 - Recurrence risk
- Screening
- Patient stratification
 - Precision medicine
- Monitoring the response to treatment
- Course of the disease

Types of biomarkers

• Sources

- Peripheral blood
 - Serum, plasma
- Urine
- Saliva
- Endometrial fluid
- Menstrual debris
- Peritoneal fluid

• Purpose

- Diagnosis
- Prognosis
- Variety
 - miRNA
 - DNA
 - RNA
 - Cytokines
 - Chemokines
 - Hormones
 - ...

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Current status

- A hot topic for extensive reviews
 - May et al. HRU 2010
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 - Fassbender et al. Biomed Res Int 2015
 - Liu et al. Cochrane Database Syst Rev 2015
 - Gupta et al. Cochrane Database Syst Rev 2016
 - Nisenblat et al. Cochrane Database Syst Rev 2016a
 - Nisenblat et al. Cochrane Database Syst Rev 2016b
 - Agrawal et al. Mol Sci 2018
- Consensus: Not a single biomarker has been clinically validated

What went wrong?

Failure is the mother of success.

-- A Chinese Proverb

Failure is infinitely more educational than success.

Challenges

- Heterogeneity
 - Different subtypes of endometriosis
 - Age
 - Premenarcheal
 - Cyclic
 - Postmenopausal
 - Variation in symptomology
 - Pain
 - Different kind of pain
 - Infertility
 - Both
- Co-morbidities
 - Adenomyosis
 - Uterine fibroids
 - Autoimmune diseases
 - Other chronic diseases

Ahn et al. F&S 2017

What have been done wrong?

A revisit of 70 studies on biomarkers

- These studies were evaluated by Nisenblat et al. 2016
- Published between 1986-2014
- Evaluated diagnostic performance of 47 blood biomarkers
- Total number of cases: 5,356
- Total number of controls: 3,470
- Highest sensitivity reported: 0.81
- Highest specificity reported: 0.87

Number of published biomarker studies 1986-2014





Cumulative # of published biomarker studies 1986-2014

Number of publications by continent



Number of publications by country



Distribution of #cases per study



of cases

Distribution of #controls per study



of cases

Single-gate design

Two-gate design



N=36 (51.4%)

N=34 (48.6%)

Population sampled



Indications for surgery



Combidities



Exclusions



Target condition severity



Index test vs. biomarkers



Biomarker novelty



Reference histology



Other features

- 23 of 70 (32.9%) studies did not give any information on age for cases
- 22 of 70 (31.4%) did not give any info on age for controls.
- 67/70=95.7% used rASRM classification system, only 4.3% unreported
- 62/70 (88.6%) did not report whether the operator(s) was/were blinded or not
- 43/70=61.4% specified the threshold. 38.6% did not.
- 64/70=91.4% did not validate their results in an independent sample
- 65 of 70 (92.9%) studies did not retest
- 62/70=88.6% of studies had missing data that could affect the interpretation of the results.
- 64/70=91.4% of studies did not provide diagnostic criteria

An analysis of 41 studies published in 2015-2018

- These studies were published after the Ephect guidelines in 2014
 - 2015: 9
 - 2016: 15
 - 2017: 10
 - 2018 (up to now): 7
- Studies on biomarker for recurrence were excluded
- Total number of cases: 2,572
- Total number of controls: 1,839

Other features

- 4 of 41 (9.8%) studies did not give any information on age for cases or controls
- 16 of 41 (39.0%) used single-gate (cohort) design, and the rest used two-gate (case-control) design
- 6/41=14.6% did not do histological confirmation, and 2 (5%) unreported
- 36/41=(12.2%) did not report whether the operator(s) was/were blinded or not
- Only 1/41=2.4% specified the threshold. 97.6% did not.
- 64/70=91.4% did not validate their results in an independent sample
- 39 of 41 (95.1%) studies did not retest
- 11/41=26.8% of studies did not report if there are missing data

Geographic distribution



Number of studies by country



Co-occurrance with adenomyosis



Co-occurrance with uterine fibroids



Co-occurrance with other comorbidities



of cases per study



of cases

of controls per study



of controls

Summary

- Typically small or moderate sample sizes
 - Not able to cover all stages
 - Impossible to cover all combinations of important factors: age, subtype, menstrual phase, comorbidity, symptomology
- Lacks detail
- Susceptible to biases
- No validation/retest
- No proof that the marker is lesion-specific

What should we consider?

- Where does the biomarker come from?
 - Any in vivo data?
 - Is the change
 - Subtype-dependent
 - progression-dependent?
 - Symptomology dependent?
 - Specific to endometriosis?
- How can we eliminate the "Humburg effect"?
- Can we have preclinical evidence first?

Basic requirements

- Large sample sizes
 - to account for variations in age, menstrual phase, rASRM stage, subtypes, symptomology and severity, and common co-morbidity
- Endometriosis-specific
 - Once all visible lesions are removed, there is a change; Retest
 - Correlation with lesional histology
 - Stage of progression
 - Support from in vivo data

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 - Correlation with lesional histology
 - Stage of progression
 - Support from in vivo data
- Validation
- Harmonization all protocols for phenotyping, biobanking, and experimentation

Conclusions

- The biomarker studies published during the 1986-2018 period overall have low quality
- Most of them can easily succumb to various biases
- Most of them are difficult to be generalized to more general population/subtypes
- The statistical power is typically low
- To maximize our chance in identifying biomarkers in the future, there are certain requirements need to be fulfilled

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