



# Genetics of endometriosis

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## Other disclosures

Lecture fees Merck, Exeltis, Besins Health Care, Stragen



# *Endometriosis*

## **CHRONIC MULTIFACTORIAL SYSTEMIC DISEASE**

Inflammation / Progesterone resistance  
Infertility / CVD / Low BMI / Fatigue

### **HORMONAL TREATMENT**

Amenorrhea

Fertility treatments

***New!***

## **CHRONIC PAIN**

Dysmenorrhea / Pelvic, abdominal and back pain /  
Migraine

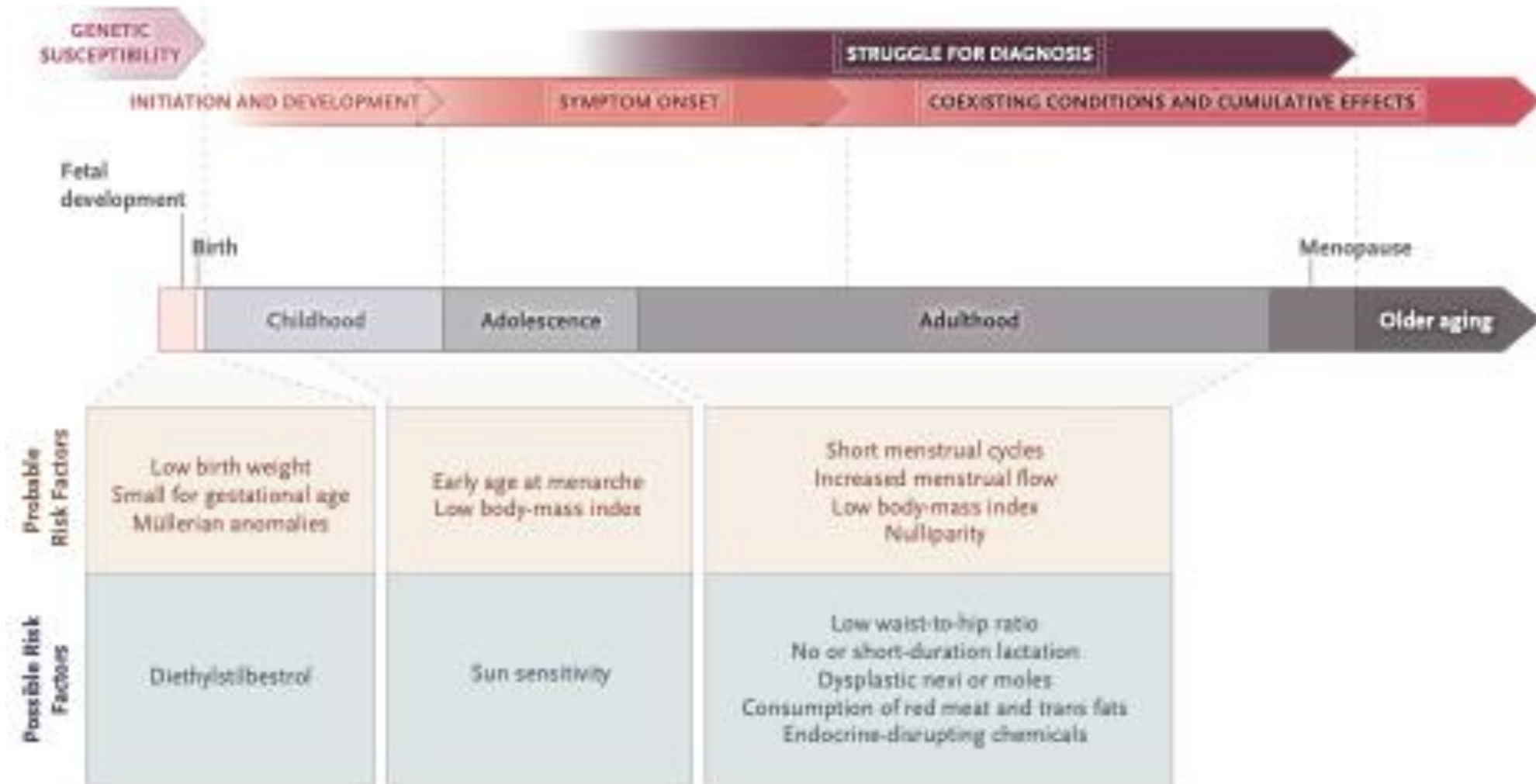
### **PAIN MANAGEMENT PLAN**

Chronic pain

Acute pain



# Endometriosis across the Life Course



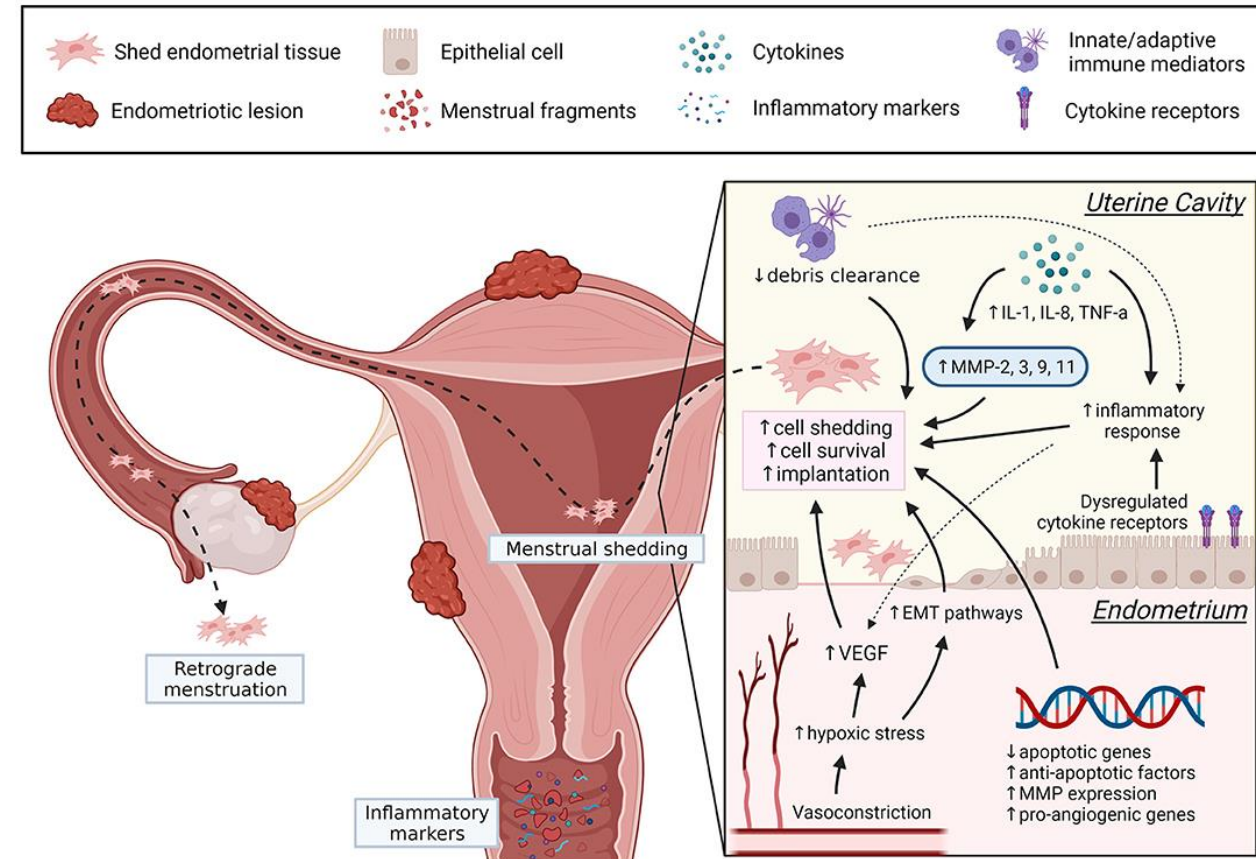


# Pathogenesis of endometriosis

- 1) Endometrial tissue (endometriosis lesions) located outside the uterus, consisting of
  - a. endometrial glands
  - b. stromal cells
  - c. fibrosis

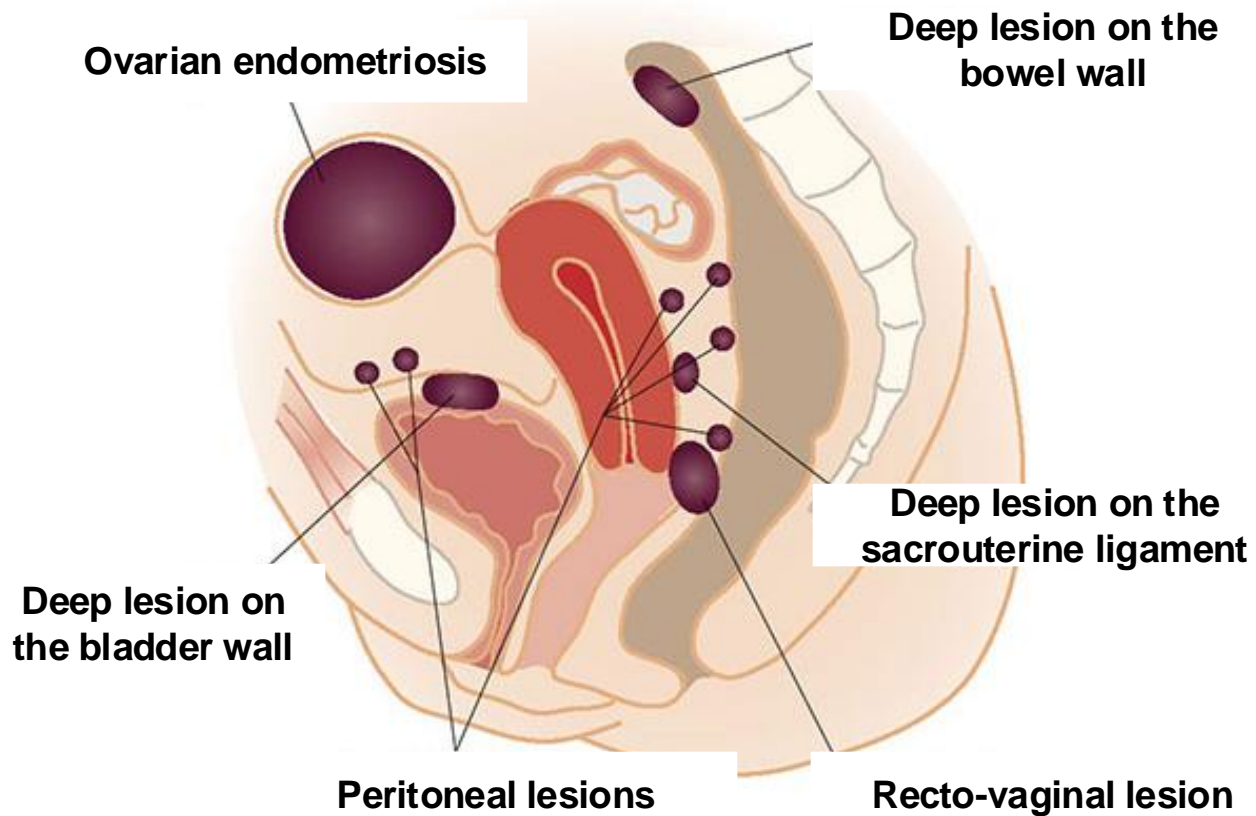
→ Endometrial cells implant and form a cyst cavity bordered by glandular cells
- 2) Several dysregulated menstrual processes: angiogenesis, EMT, apoptotic regulators, inflammation, debris clearance, MMP  

→ May promote endometrial cell detachment, prolong endometrial cell survival and increase the likelihood for implantation following retrograde menstruation.
- 3) Causes adhesions between abdominal organs
- 4) Cyclic estrogen and progesterone secretion regulates glandular function. Endometriosis cells also produce their own estrogen via aromatase.





# ***Endometriosis lesion sites***



Ovaries

Recto-vaginal space (Fossa Douglas)

Sacrouterine ligaments

Rectum and sigma

Conjunctive of the bladder

Ligament latum

Fallobian tubes

→ **SAMPSON'S THEORY (RETROGRADE MENSTRUATION)**

Extraperitoneal sites:

bowel, urinary tract, pulmonary system, CNS, skin

→ **VIA STEM CELLS/METAPLASIA**





Minimal  
peritoneal  
**LESIONS**  
(white arrows)

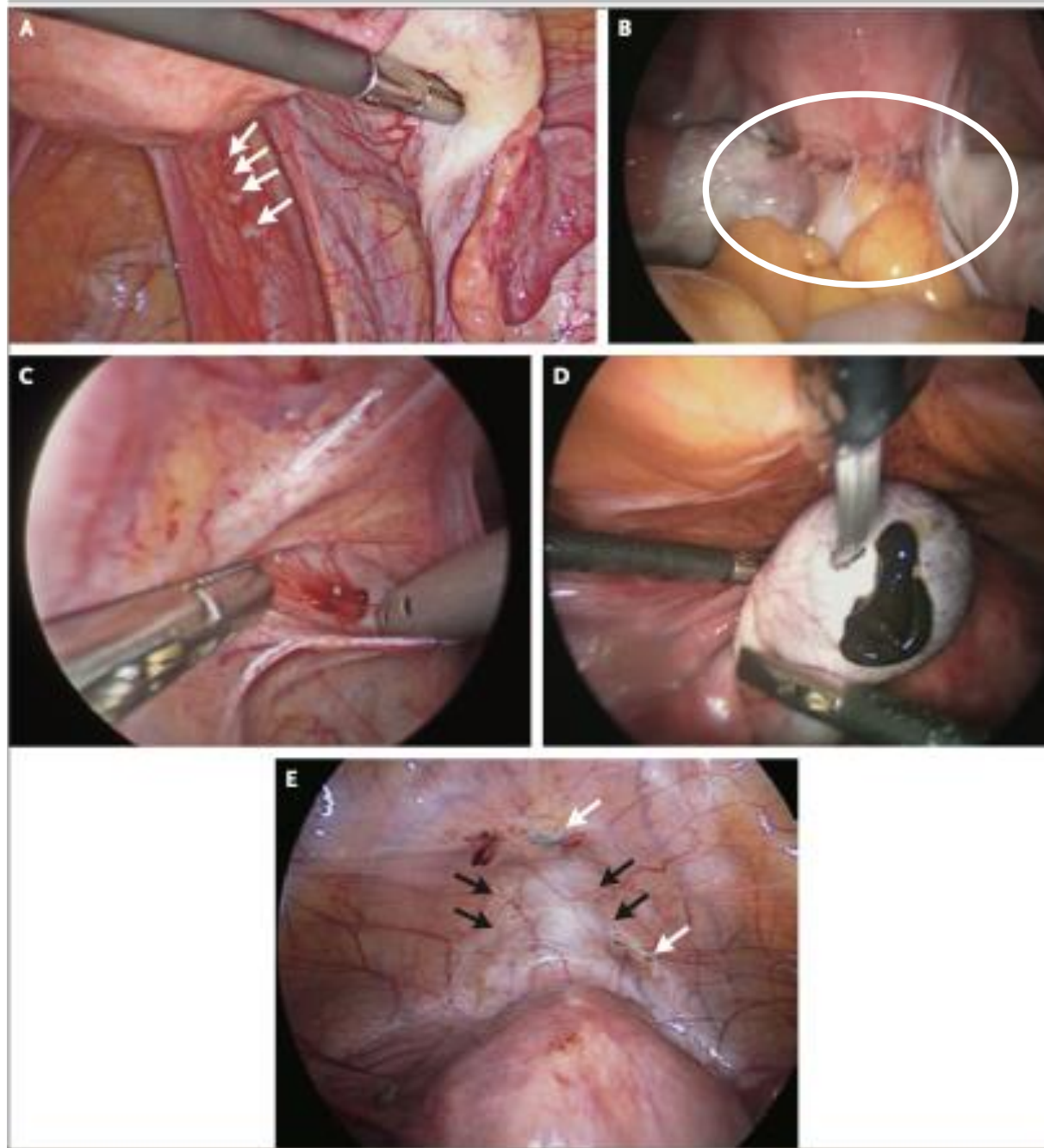
Superficial red  
peritoneal  
**LESION** and  
hyperemia

### CHARACTERISTICS OF ENDOMETRIOSIS LESIONS

Vascularization of lesions  
Innervation of lesions  
Scarring or adhesions within  
the pelvic cavity

### LESION SIZE

Varies from millimeters to  
centimeters



Bowel  
**ADHESIONS**  
to uterus,  
obliteration of  
recto-vaginal  
space

**ENDOMETRIOMA**  
(ovarian  
endometriosis),  
"chocolate cyst"

**DEEP** bladder lesions  
(black arrows) and red,  
brown and black  
peritoneal **LESIONS**  
(white arrows)



# CLASSIFICATION OF ENDOMETRIOSIS

According to lesion site

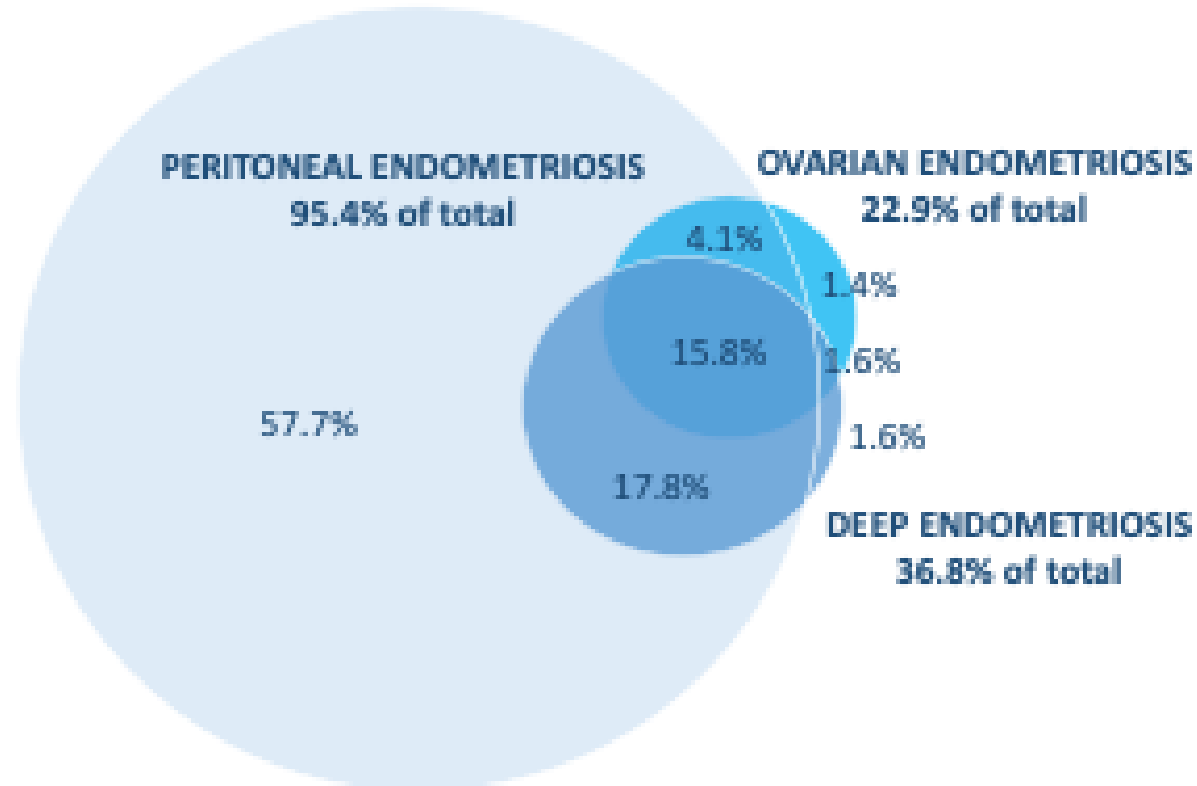


Figure 1. Presentation of the prevalence and overlap of endometriosis lesion sites using venn diagrams. Most patients with endometriosis have only peritoneal lesions, but those with ovarian endometriosis often have additionally peritoneal lesions and deep lesions. Patients with deep lesions often have peritoneal and ovarian lesions.





# CLASSIFICATION OF ENDOMETRIOSIS

Grading according to lesion site, size, tissue depth and adhesions

## Revised ASRM

Stage I – minimal	1-5
Stage II – mild	6-15
Stage III – moderate	16-40
Stage IV – severe	>40



AMERICAN SOCIETY FOR REPRODUCTIVE MEDICINE  
REVISED CLASSIFICATION OF ENDOMETRIOSIS

Patient's Name \_\_\_\_\_ Date \_\_\_\_\_  
 Stage I (Minimal) : 1-5  
 Stage II (Mild) : 6-15  
 Stage III (Moderate) : 16-40  
 Stage IV (Severe) : >40  
 Total \_\_\_\_\_  
 Laparoscopy \_\_\_\_\_ Laparotomy \_\_\_\_\_ Photography \_\_\_\_\_  
 Recommended Treatment \_\_\_\_\_  
 Prognosis \_\_\_\_\_

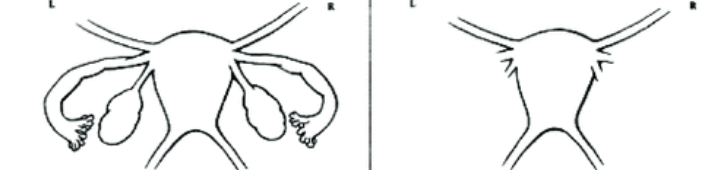
	ENDOMETRIOSIS	< 1cm	1-3cm	> 3cm
		1	2	4
PERITONEUM	Superficial	1	2	4
	Deep	2	4	6
	R Superficial	1	2	4
	Deep	4	16	20
OVARY	L Superficial	1	2	4
	Deep	4	16	20
	POSTERIOR CULDESAC OBSTRUCTION	Partial 4		Complete 40
	ADHESIONS	< 1/3 Enclosure	1/3-2/3 Enclosure	> 2/3 Enclosure
OVARY	R Filmy	1	2	4
	Dense	4	8	16
	L Filmy	1	2	4
	Dense	4	8	16
TUBE	R Filmy	1	2	4
	Dense	4*	8*	16
	L Filmy	1	2	4
	Dense	4*	8*	16

\*If the fimbriated end of the fallopian tube is completely enclosed, change the point assignment to 16.  
 Denote appearance of superficial implant types as red (R), red, red-pink, flame-like, vesicular blots, clear vesicles, white (W), opacifications, peritoneal defects, yellow-brown, or black (B) black, hemosiderin deposits, blue]. Denote percent of total described as R\_\_\_\_%, W\_\_\_\_%, and B\_\_\_\_%. Total should equal 100%.

Additional Endometriosis: \_\_\_\_\_ Associated Pathology: \_\_\_\_\_

\_\_\_\_\_

To Be Used with Normal Tubes and Ovaries To Be Used with Abnormal Tubes and/or Ovaries

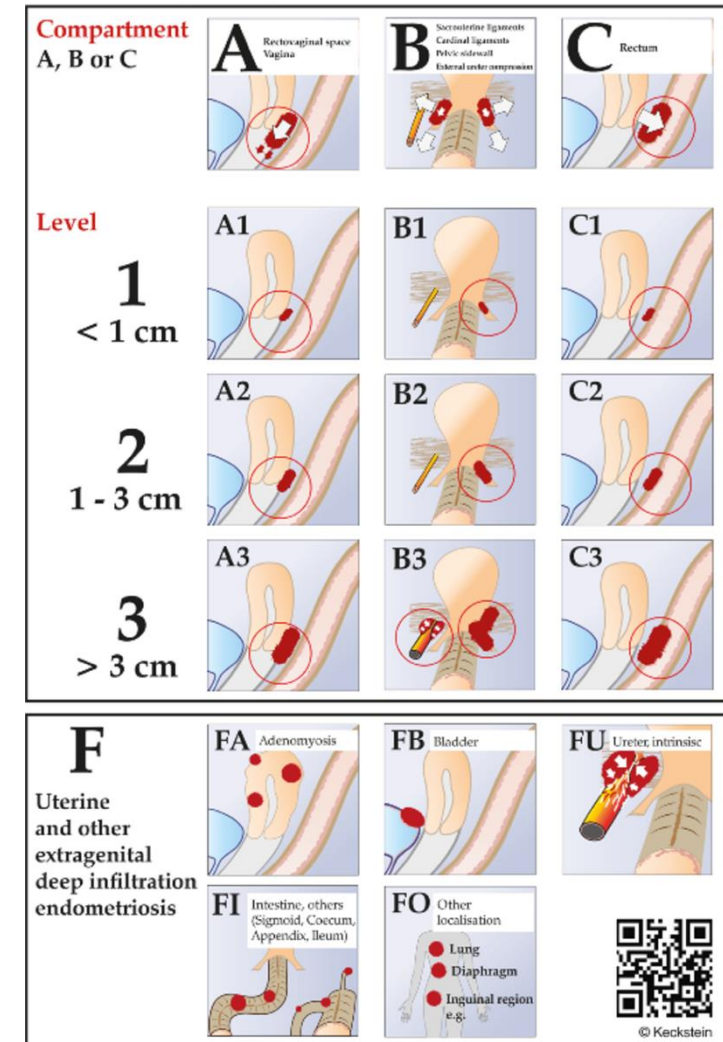




# CLASSIFICATION OF ENDOMETRIOSIS

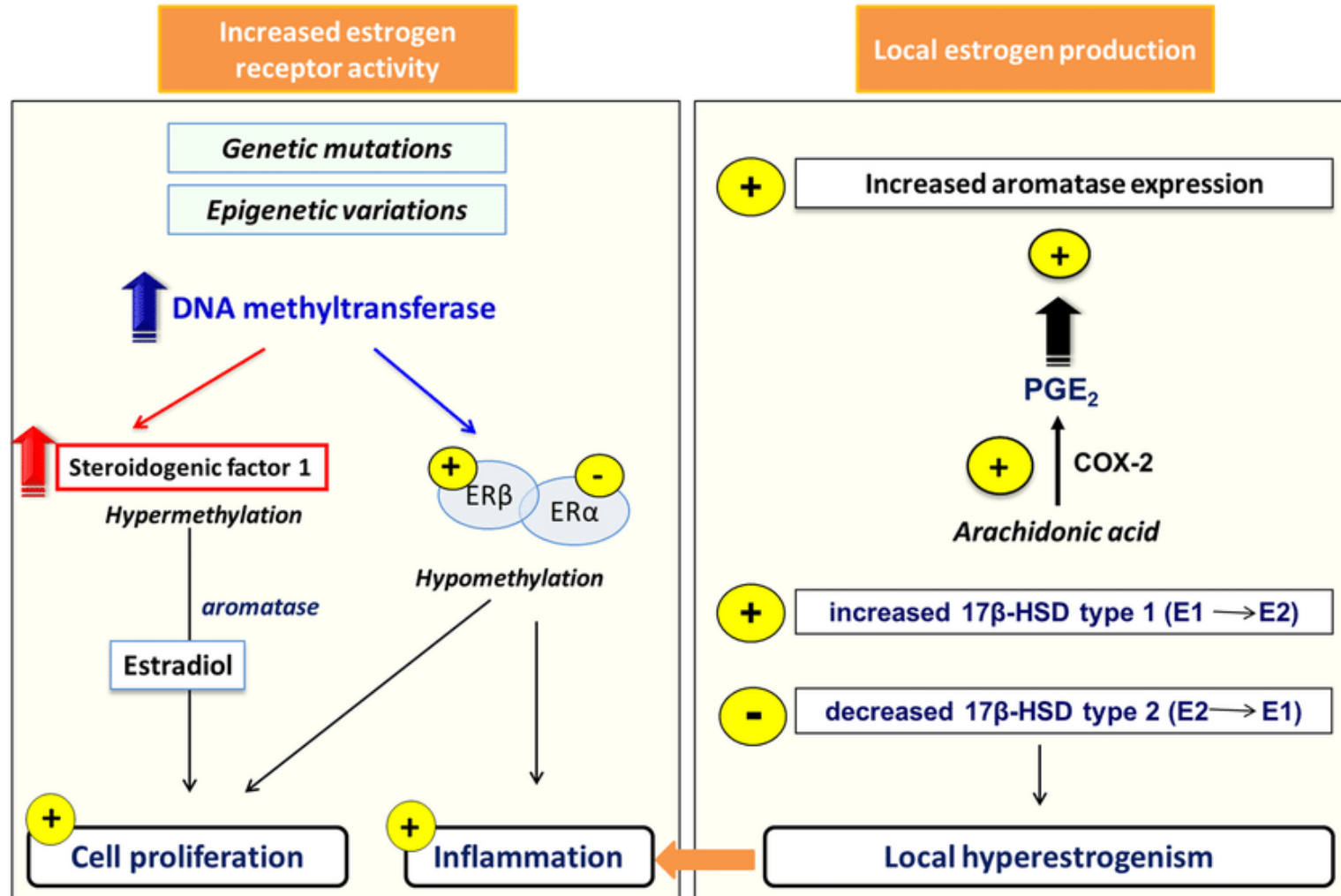
According to lesion site and tissue depth, specifically recto-vaginal disease

## The ENZIAN-Score





# ESTROGEN: a dominant role in endometriosis pathogenesis

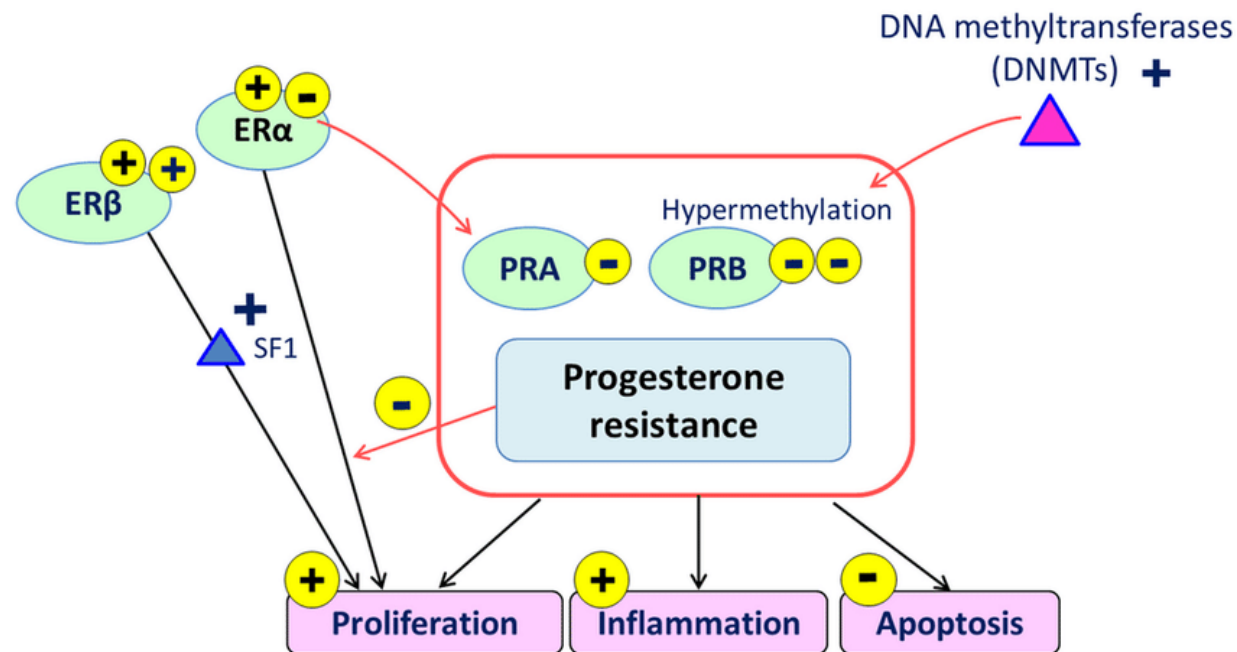




# Mechanisms of progesterone-resistance in endometriosis

- 1) Dysregulation of progesterone signaling
- 2) Endometrial tissue inability to appropriately respond to progesterone

*Well-established in both the endometriotic lesions and eutopic endometrium of women with endometriosis!*









# Quality of life



## COMMON SYMPTOMS (prevalence)

Dysmenorrhea (pain during menses) (70-80%)

Dyspareunia (pain during intercourse) (70-75%)

Chronic abdominal or pelvic pain (60%)

Dyschezia (pain during bowel movement) (40-70%)

Dysuria (pain while urinating) (6-68%)

Infertility (30-50%)

## Rare symptoms

Shoulder stab or upper abdominal pain during menses

Acute abdominal pain

Painful lump in abdominal wall

Bleeding from belly button

Sciatic pain

Loss of function of one kidney

Pneumothorax



Artist Ellie Kammer. An Australian artist hopes to raise global awareness about endometriosis by painting powerful images detailing her struggle with the illness.

**Artist With Endometriosis Channels 'Desperation' Into Striking Portraits To Raise Awareness** *'Painting made me feel powerful again.'* **By Rachel Moss**  
Huffington Post UK 24/10/2017 12:23pm BST | **Updated** October 24, 2017

Suivitie. Milloin epäillä endometrioosia? [When to suspect endometriosis]

<sup>14</sup> Lääkärilehti 14-15/2020





# COMMON DISEASE

**10% of reproductive-age women → ~190 million women worldwide**

2 to 11%	asymptomatic women
5 to 50%	infertile women
5 to 21%	women hospitalized for pelvic pain
49 to 75%	symptomatic adolescents, chronic pelvic pain and unresponsive pain to medical treatment

Zondervan, Becker, Missmer. Endometriosis. NEJM 2020

## **COMPARISON**

**Prevalence of type 2 diabetes 6.28%**

Khan et al. Epidemiology of Type 2 Diabetes – Global Burden of Disease and Forecasted Trends. J Epidemiol Glob Health. 2020

**Prevalence of asthma 5.4%**

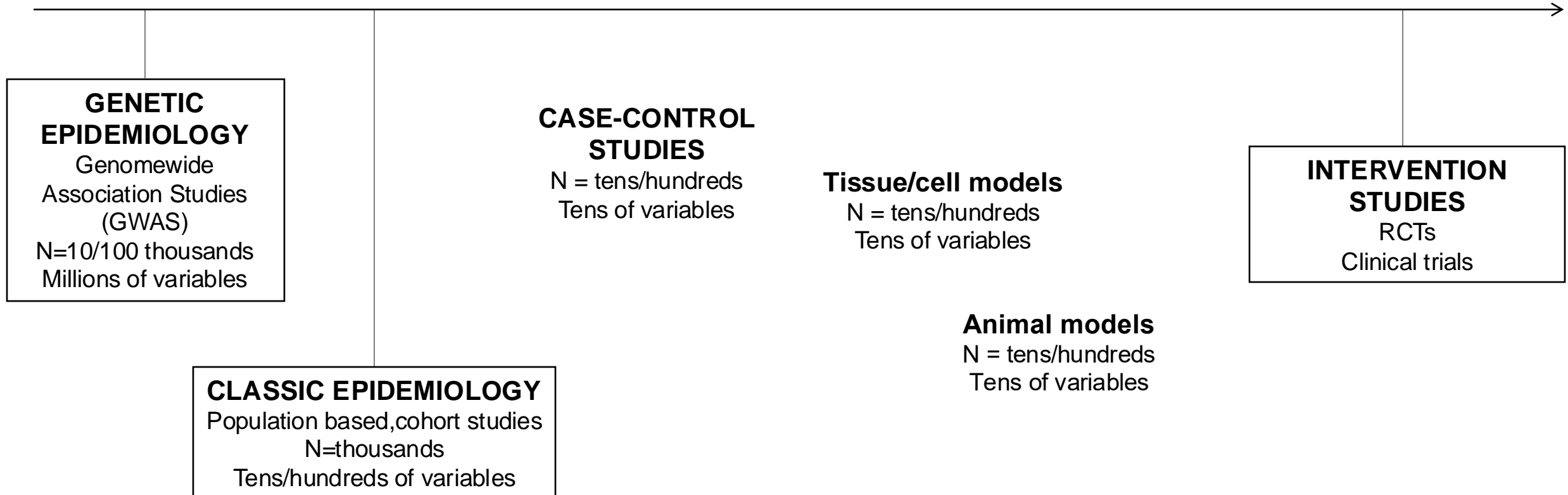
Song et al. Global, regional, and national prevalence of asthma in 2019: a systematic analysis and modelling study. J Glob Health. 2022



# Research study designs by hypothesis level

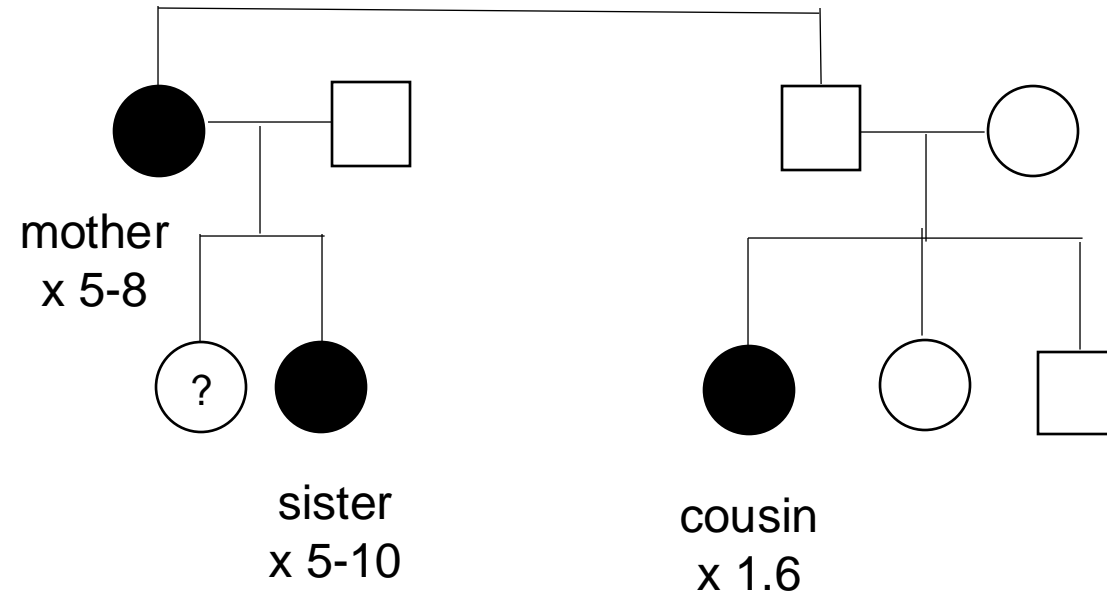
*Hypothesis free approach*

*Hypothesis driven approach*





# Increased disease risk by family history



Moen et al. The familial risk of endometriosis. ACTA Obstet Gynecol Scand 1993  
Stefansson et al. Genetic factors contribute to the risk of developing endometriosis. Hum Reprod 2002  
Kashima et al. Familial risk among Japanese patients with endometriosis. Int J Gynaecol Obstet 2004  
Matalliotakis et al. Familial aggregation of endometriosis in the Yale series. Arch Gynecol Obstet 2008



# Potential high-risk genetic variants for endometriosis

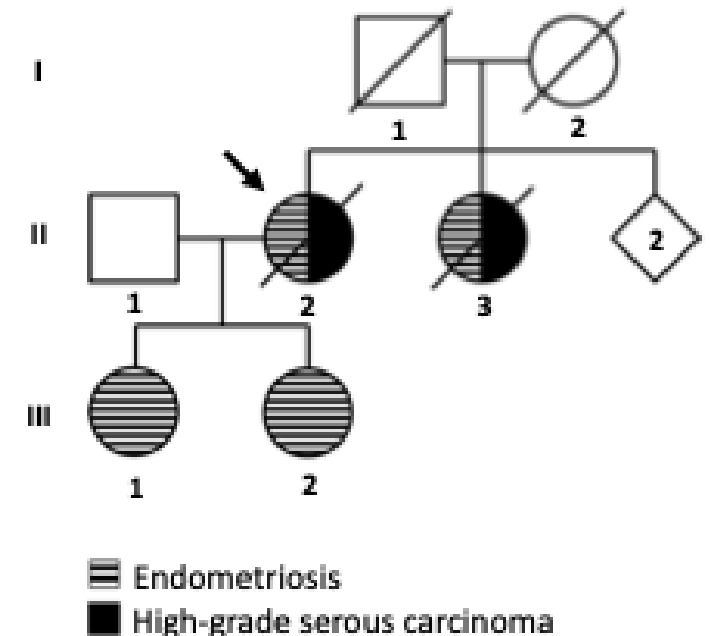
Only a few potential high-risk genetic variants have been identified:

- 1) Variants causing amino acid changes in the NPSR1 gene with association to reduction of inflammatory cell infiltrate and abdominal pain, have been identified in nine different families with endometriosis.

(Tapmeier et al. Neuropeptide S receptor 1 is a nonhormonal treatment target in endometriosis. Sci Transl Med 2021)

- 2) Identified variants in three genes (FGFR4, NALCN, NAV2) by whole-exome sequencing that segregate with endometriosis using Finnish family (Fig. 1) data.

(Nousiainen et al. Whole-exome sequencing reveals candidate high-risk susceptibility genes for endometriosis. Hum Genomics 2023)



**Fig. 1** Pedigree of the Finnish endometriosis family. Four women in two generations have been diagnosed with surgically verified endometriosis and two also with high-grade serous carcinoma (HGSC). The index patient is marked with an arrow. The pedigree has been slightly modified for anonymity



# Endometriosis heritability

*The relative contribution of genetic and environmental factors to the observed variation in a trait within a population*

According to **twin studies**, the **heritability** of endometriosis (the proportion of disease risk due to genetic variation) **~ 50%**

Saha et al. Heritability of endometriosis. Fertil Steril 2015;104:947-52.

**Common genetic variation** accounts for **~ 26% of the risk of endometriosis**

Lee et al. Estimation and partitioning of polygenic variation captured by common SNPs for Alzheimer's disease, multiple sclerosis and endometriosis. Hum Mol Genet 2013; 22:832-41.

**Common risk variants** for multifactorial disease are detectable by sufficiently powered, population-based genome-wide association studies (GWAS)

Visscher et al. 10 Years of GWAS discovery: biology, function, and translation. Am J Hum Genet 2017;101:5-22.



## Genome-wide association studies (GWAS)

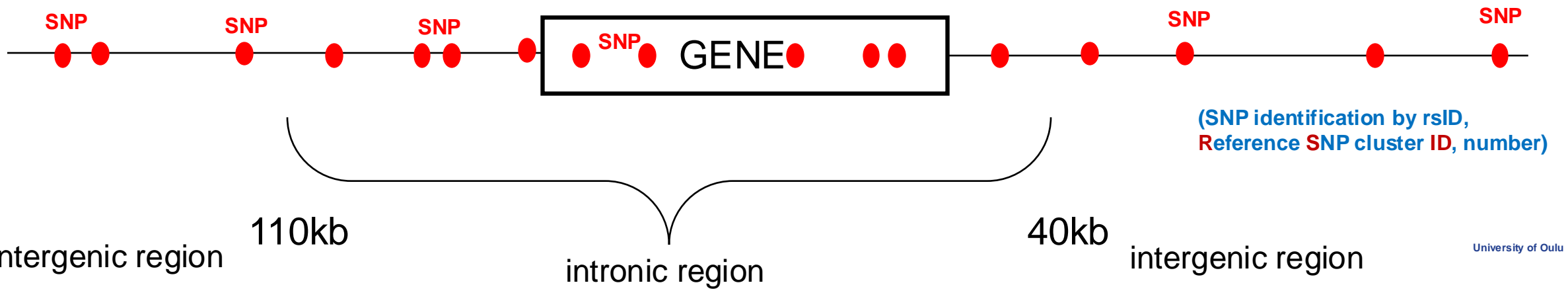
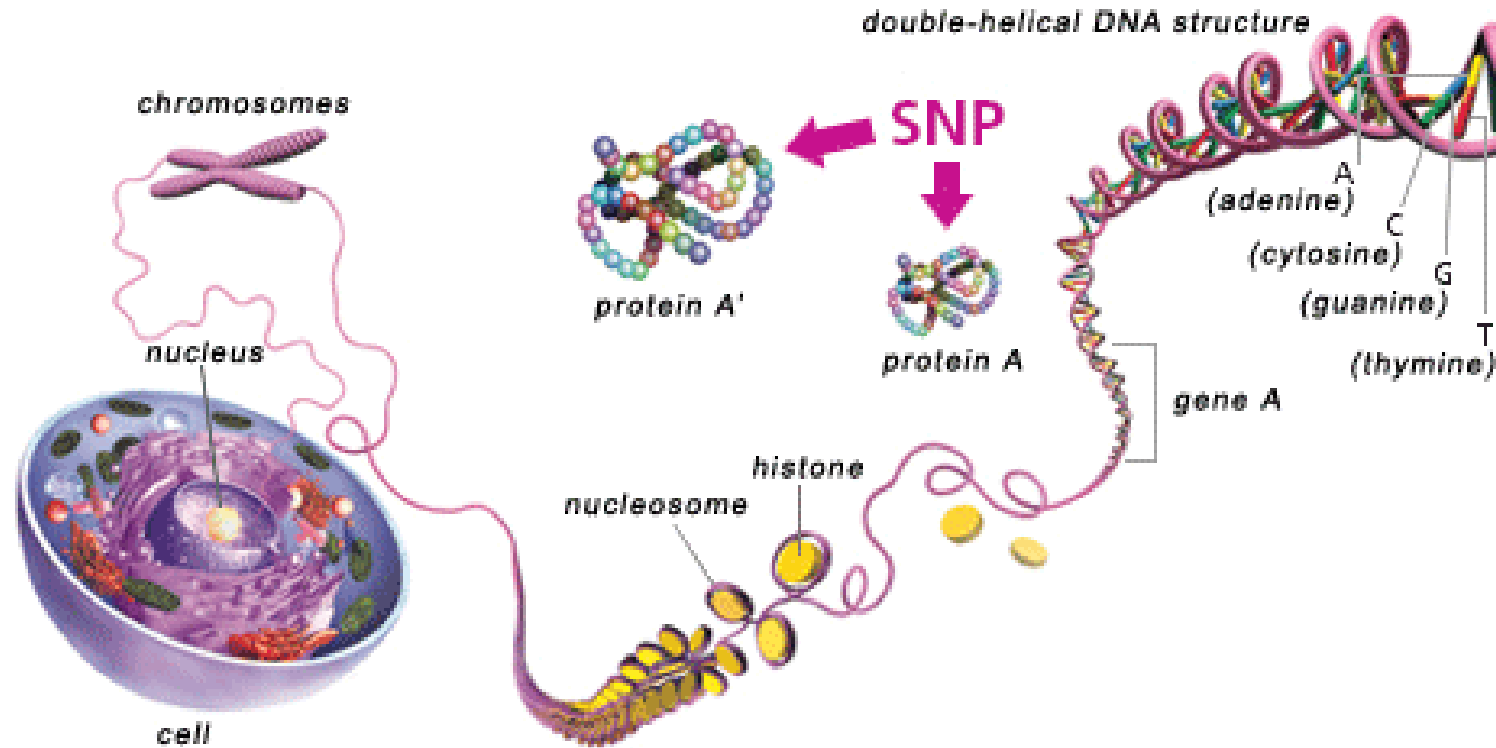
test millions of genetic variants (SNPs)  
across many genomes (sample sets of hundreds of thousands)  
to find those variants that are statistically ( $P < 5 \times 10^{-8}$ )  
associated with a specific trait or disease

**SNP** single nucleotide variations (polymorphism)

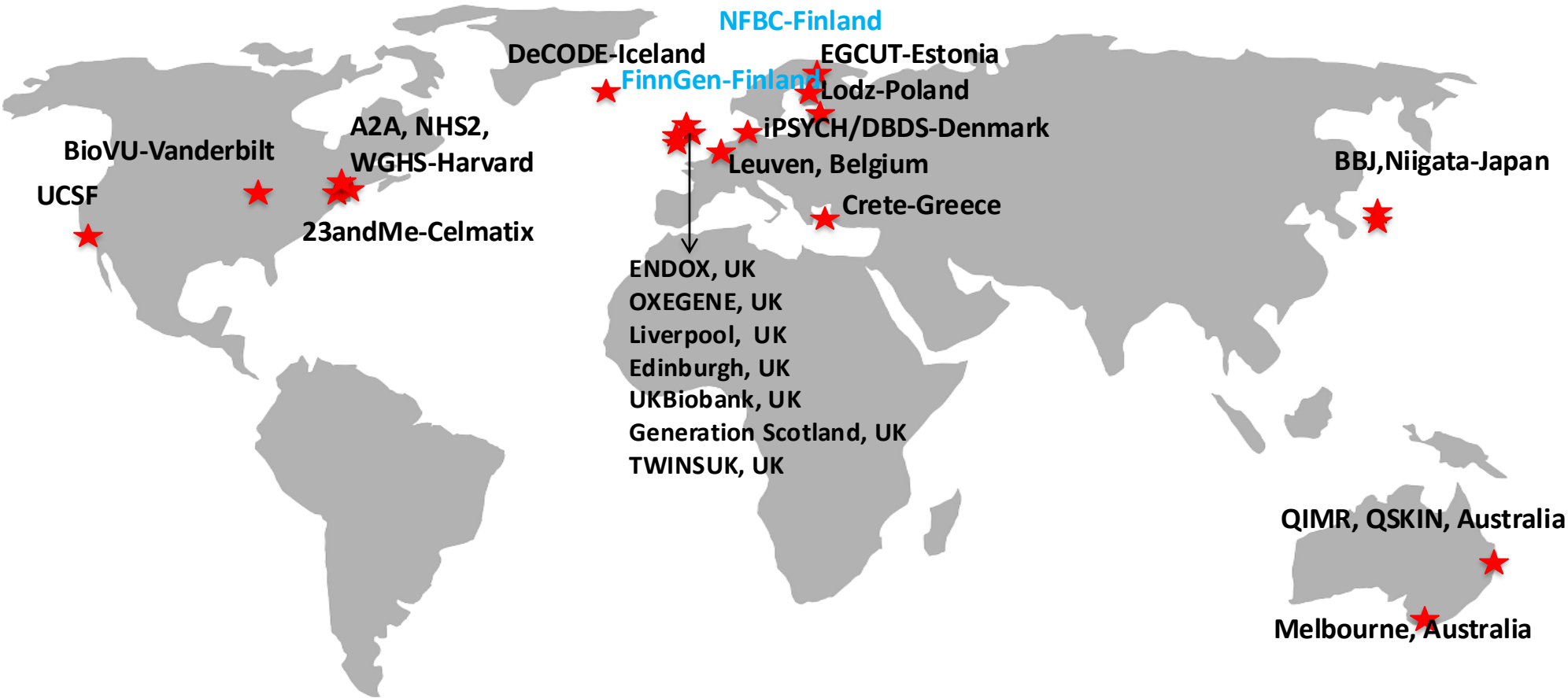
- one SNP for every 500-1000 bases
- 3-5 million SNPs in the entire human genome, of which 500,000 SNPs are in gene regions
- are different between individuals

SNP in a gene can cause changes in the function, timing, or amount of the protein produced by that gene.





# International Endometriosis Genomics Consortium (IEGC)



Prof Stacey Missmer, Michigan State Uni, USA



Prof Grant Montgomery, Uni of Queensland, Australia



Prof Andrew P. Morris, Uni of Liverpool, UK



Prof Krina Zondervan, Uni of Oxford, UK

GWAS	All Cases:Controls	Stage III/IV: Co	Stage I/II: Co	Infertile: Co
Total N (24 Studies)	60,674 : 701,926	4,045 : 379,898	3,916 : 184,014	3,060 : 221,674



	<b>IEGC</b>	<b>International Endometriosis Genomics Consortium</b>			
	<b>Study acronym</b>	<b>Study name</b>	<b>Ancestry</b>	<b>Total Case N</b>	<b>Total Control N</b>
1	23andMe	23andMe dataset	European	37 182	251 255
2	FinnGen	FinnGen	European	6 502	57 407
3	UKBiobank	UK Biobank dataset	European	6 611	196 188
4	QIMRHCS	Queensland Institute of Medical Research and Hunter Community Study	European	2 262	2 923
5	NHS2	Nurses' Health Study II	European	2 104	5 854
6	DECODE	DeCODE Genetics	European	1 857	132 978
7	EGCUT	The Estonian Biobank Cohort	European	1 716	35 176
8	WGHS	The Women's Genome Health Study	European	1 494	15 033
9	BBJ	BioBank Japan	Japanese	1 423	1 318
10	QSKIN	QSkin Sun and Health Study	European	1 038	7 604
...					
				~65,000	~760,000



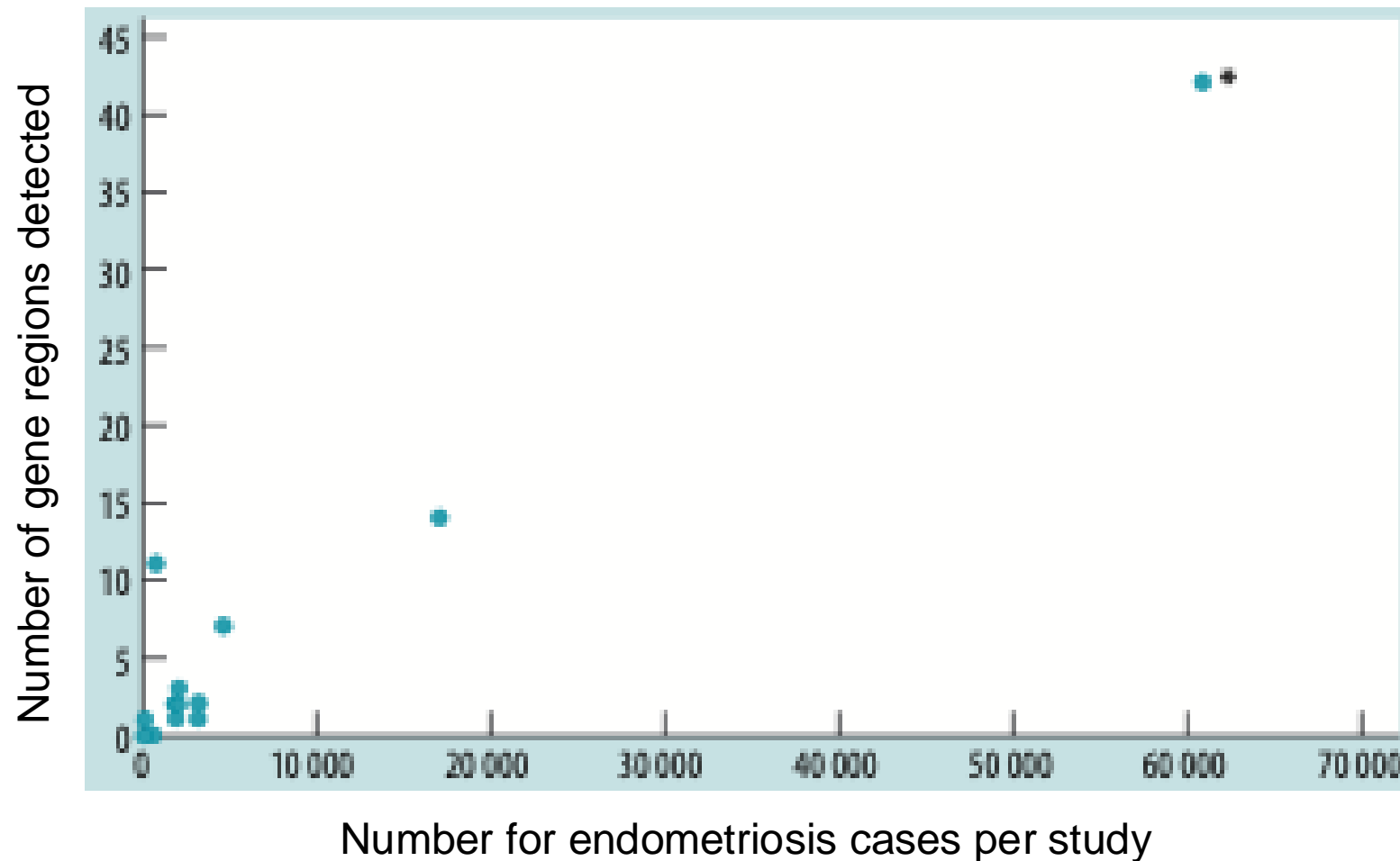
# FINNGEN: AN EXPEDITION INTO GENOMICS AND MEDICINE

FinnGen is a research project in genomics and personalized medicine. It is a large public-private partnership that has collected and analysed genome and health data from 500,000 Finnish biobank donors to understand the genetic basis of diseases. FinnGen is now expanding into understanding the progression and biological mechanisms of diseases. FinnGen provides a world-class resource for further breakthroughs in disease prevention, diagnosis, and treatment and an outlook into our genetic make-up.

## The final FinnGen cohort consists of over 500 000 individuals

The combined amount of the legacy samples and newly collected samples is 520 000. The median age of the participants when donating was 53 years and 43% are men, and 57% women.





**FIGURE 1.** Number of endometriosis-susceptible genetic regions found in GWAS studies relative to the sample size of each study. Genome-wide studies require large sample sizes to detect significant associations. To date (as of January 15, 2024), the sample sizes of endometriosis GWAS studies published have been small, with the exception of the most recent study (Rahmioglu et al. Nat Genet 2023).



# The genetic basis of endometriosis and comorbidity with other pain and inflammatory conditions

Received: 1 December 2021

Accepted: 27 January 2023

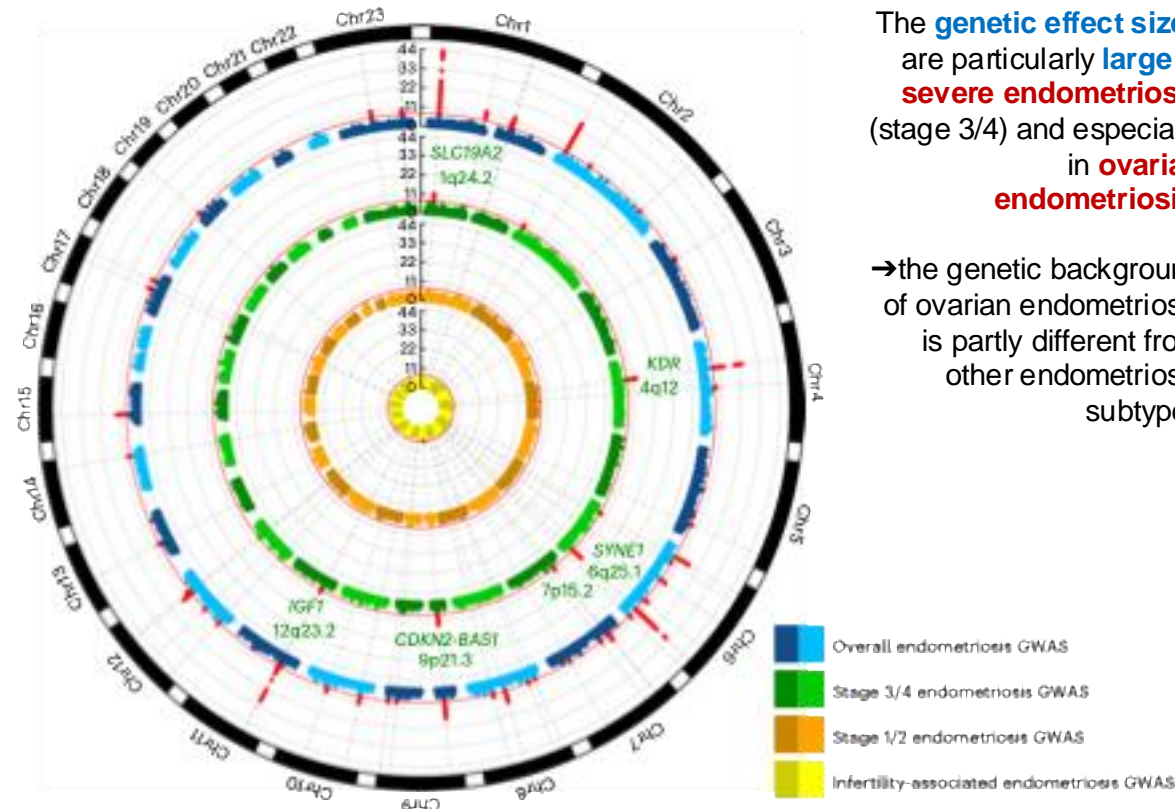
Published online: 13 March 2023

 Check for updates

A list of authors and their affiliations appears at the end of the paper

Endometriosis is a common condition associated with debilitating pelvic pain and infertility. A genome-wide association study meta-analysis, including 60,674 cases and 701,926 controls of European and East Asian descent, identified 42 genome-wide significant loci comprising 49 distinct association signals. Effect sizes were largest for stage 3/4 disease, driven by ovarian endometriosis. Identified signals explained up to 5.01% of disease variance and regulated expression or methylation of genes in endometrium and blood, many of which were associated with pain perception/maintenance (*SRP14/BMF*, *GDAP1*, *MLLT10*, *BSN* and *NGF*). We observed significant genetic correlations between endometriosis and 11 pain conditions, including migraine, back and multisite chronic pain (MCP), as well as inflammatory conditions, including asthma and osteoarthritis. Multitrait genetic analyses identified substantial sharing of variants associated with endometriosis and MCP/migraine. Targeted investigations of genetically regulated mechanisms shared between endometriosis and other pain conditions are needed to aid the development of new treatments and facilitate early symptomatic intervention.

42 genome-wide significant loci ( $P < 5 \times 10^{-8}$ ) identified for overall endometriosis, of which 31 novel



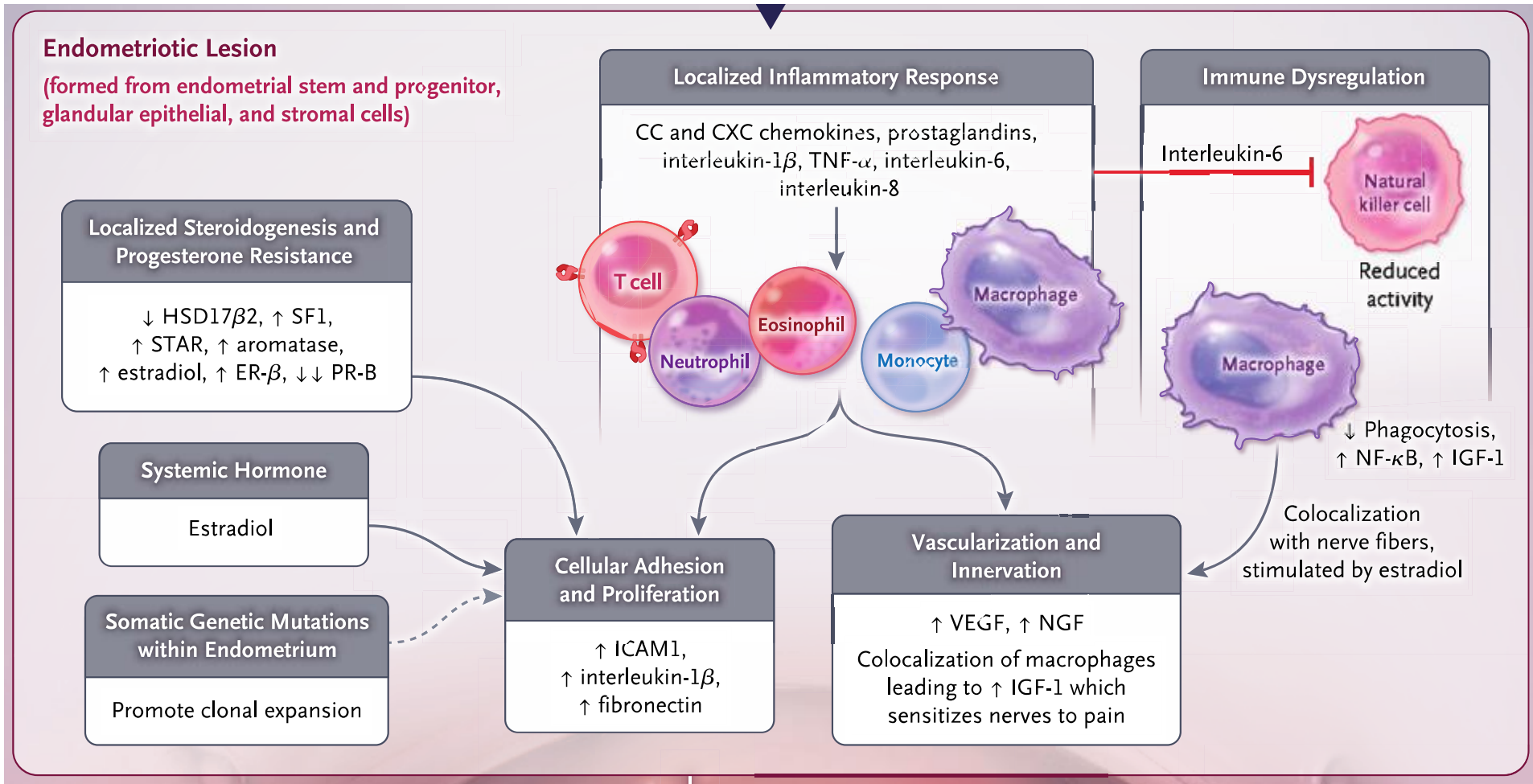






Analyses of genes closest to the GWAS loci point to biological pathways that have previously been thought to be associated with the development of endometriosis:

**Wnt signaling, cell adhesion and proliferation, angiogenesis, inflammation, and hormonal pathways.**



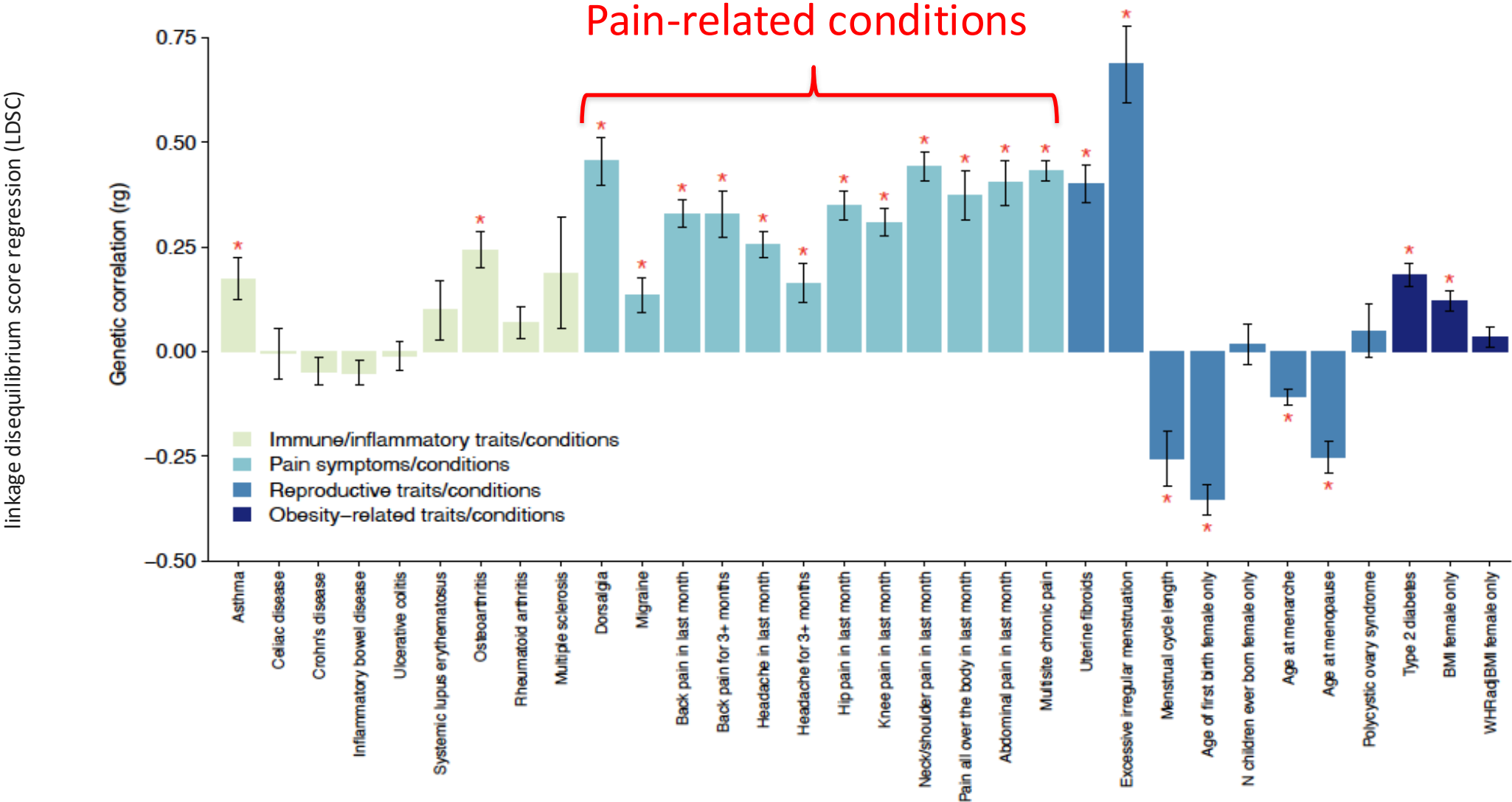


**TABLE. Genetic regions predisposing to endometriosis and pain regulation phenotypes (migraine and multifocal pain).** Data are based on multitrait analysis of GWAS (MTAG) (Rahmioglu et al. Nat Genet 2023). Gene and protein function descriptions are taken from <https://www.ncbi.nlm.nih.gov/gene/> and <https://www.uniprot.org>

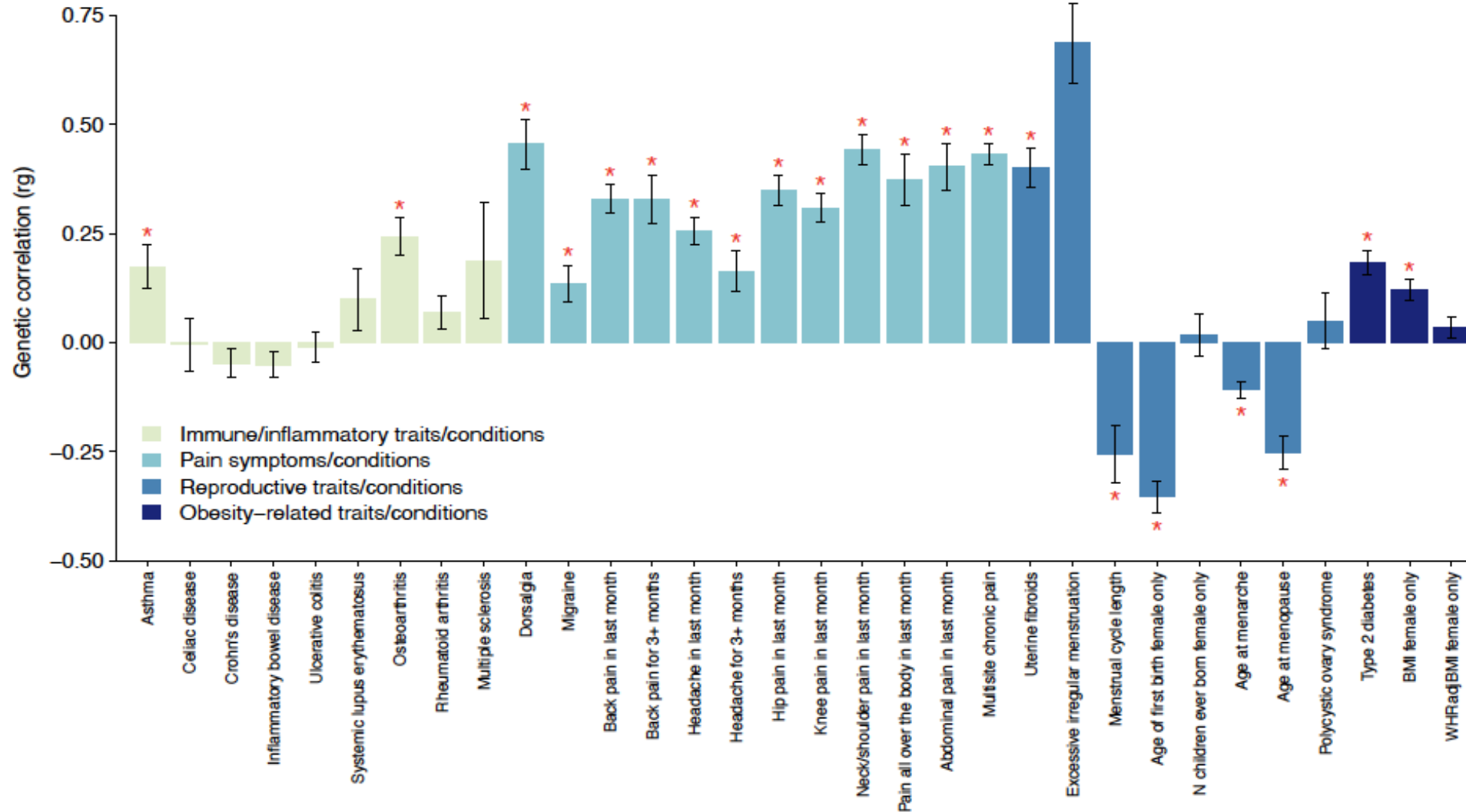
Gene region/gene	Complete gene name and function
1p32.3/ <i>FAF1</i>	Fas Associated Factor 1: initiation and enhancement of apoptosis, expressed in skeletal muscle and heart
1p13.2/ <i>NGF</i>	Nerve growth factor: development and maintenance of the sympathetic and sensory nervous system, participates in the regulation of neuronal proliferation, differentiation and survival. Affects pregranulosa cells in the ovaries and participates in the differentiation stages of stem cells in the spinal cord, neurons and brain
1q21	<b>Support for the HYPOTHESIS: endometriosis causes pain through both <u>hormonal mechanisms</u> and <u>activation of immune and neuronal cells</u>, as has been observed in other chronic pain conditions.</b>
2q37	
3p21	
5q21	
5q31	
10p15	
11p15	
12p12	
13q14.3/ <i>OLFM4</i>	Olfactomedin 4: antiapoptotic factor, promotes tumor growth and affects cell adhesion
20q11.21/ <i>NOL4L</i>	Nucleolar Protein 4 Like: encoded by protein located in the cytosol and nucleoplasm

# Genetic correlation between endometriosis and various traits

GENETIC CORRELATION – The proportion of variance that two traits share due to genetic causes

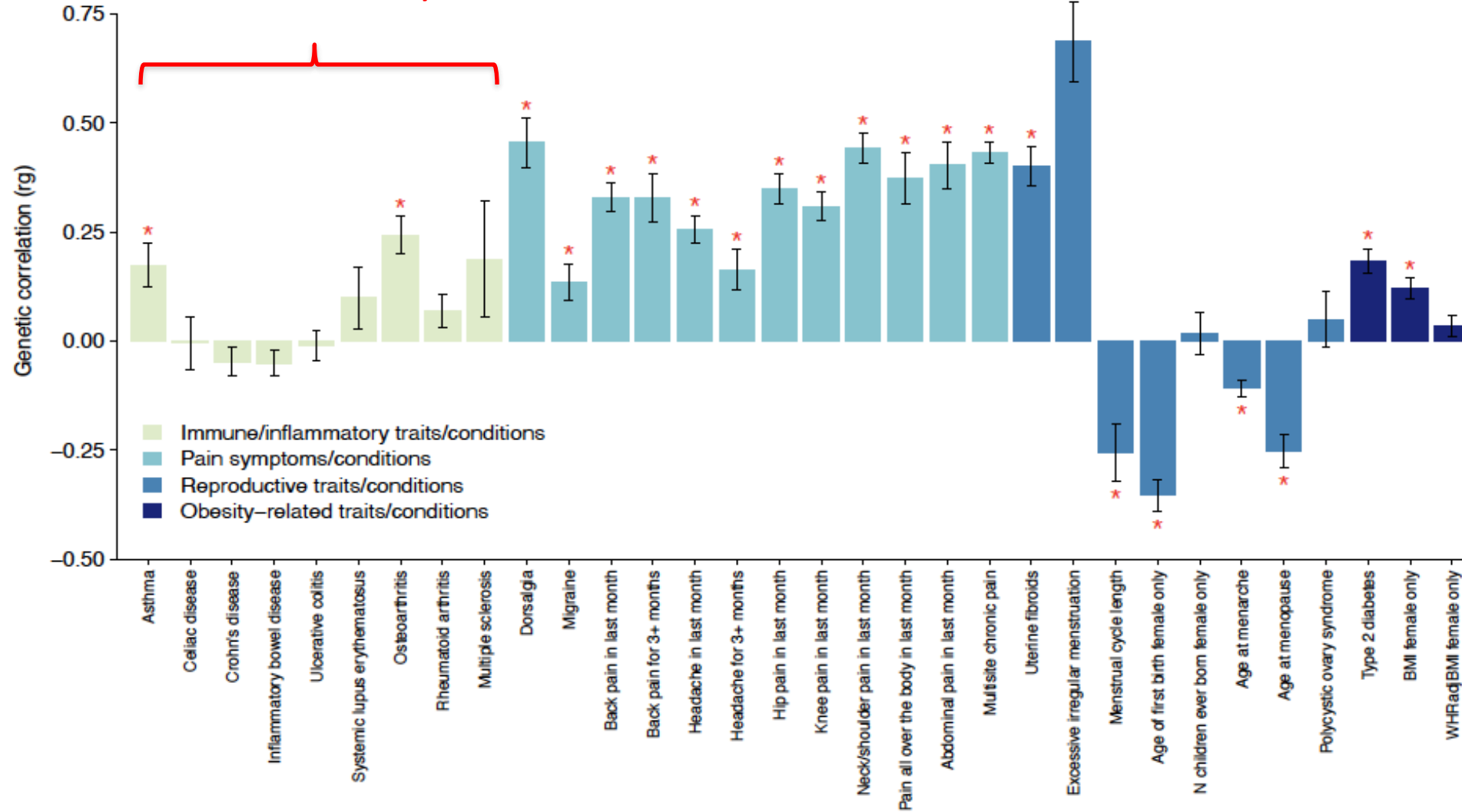


## Reproductive traits/conditions



**NEGATIVE CORRELATION**  
when the correlated trait decreases with the original trait increasing

## Inflammatory/immune conditions





# What is the significance of GWAS variants in risk mapping?

- Disease variation of common GWAS variants: only 2.2% (5.6% stage III/IV)
- Combined tools based on family history, genetic variants and other risk factors may in the future provide screening tools for girls/women with a family history of endometriosis
- The validity of such tools in independent cohorts and clinical utility need to be carefully evaluated
- Primary benefit of genetic results: discovery of a new therapeutic target

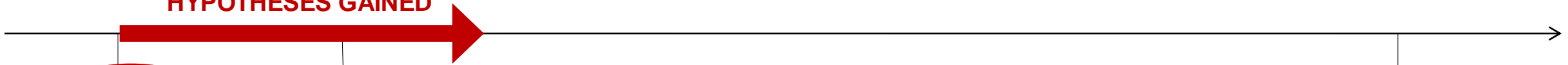


# Research study designs by hypothesis level

*Hypothesis free approach*

*Hypothesis driven approach*

**HYPOTHESES GAINED**



## **GENETIC EPIDEMIOLOGY**

Genomewide  
Association Studies  
(GWAS)  
N=10/100 thousands  
Millions of variables

## **CASE-CONTROL STUDIES**

N = tens/hundreds  
Tens of variables

## **Tissue/cell models**

N = tens/hundreds  
Tens of variables

## **INTERVENTION STUDIES**

RCTs  
Clinical trials

## **CLASSIC EPIDEMIOLOGY**

Population based, cohort  
studies  
N=thousands  
Tens/hundreds of variables

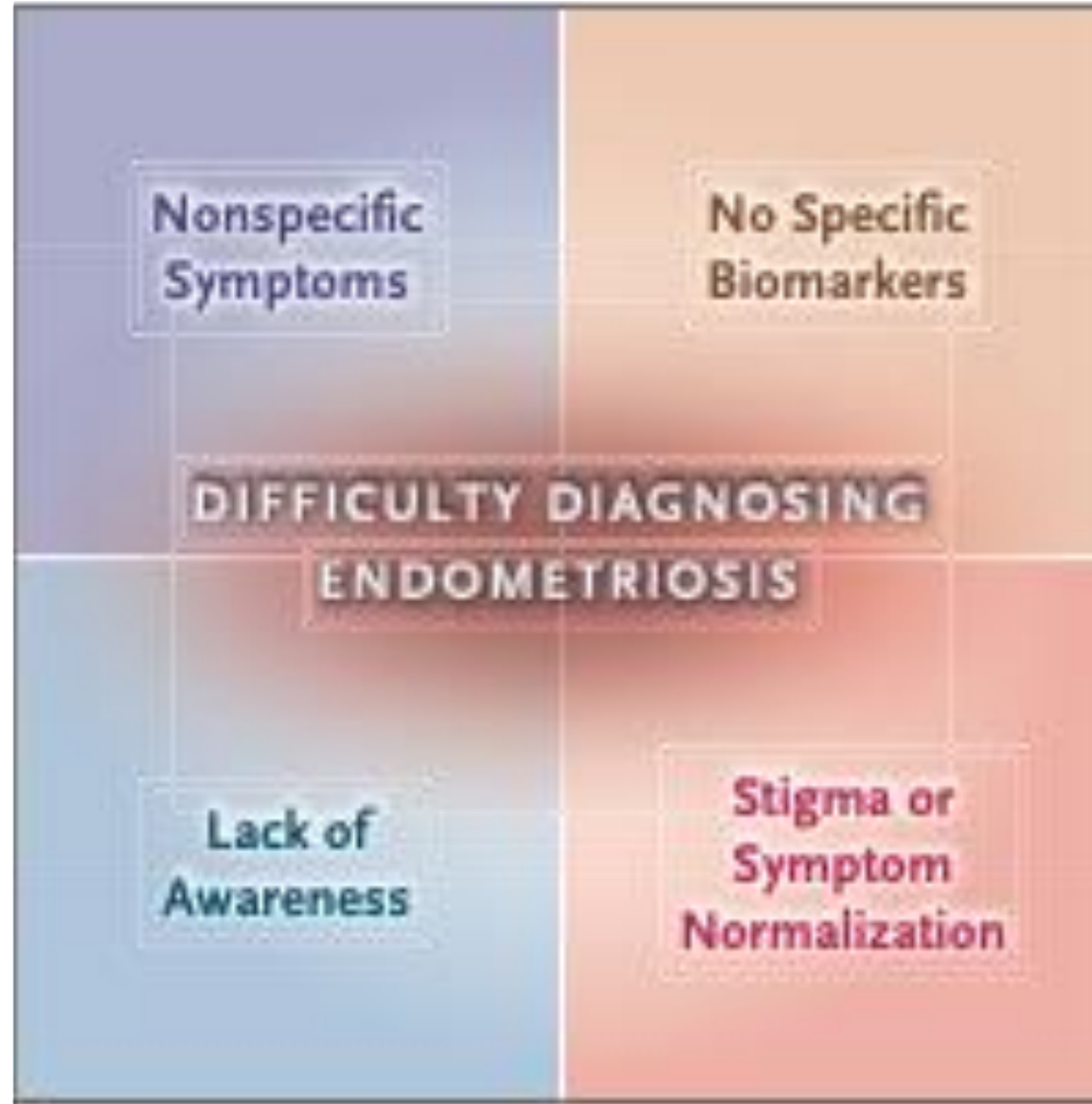
## **Animal models**

N = tens/hundreds  
Tens of variables



## KEY MESSAGES

1. Endometriosis is a common chronic pain disease in women, for which nearly 50 low-risk genetic variants have been identified in GWAS studies.
2. Genetic factors predisposing to endometriosis are related to hormonal, immunological and nervous functions, such as pain regulation.
3. Endometriosis partly shares a genetic basis with several pain diseases.
4. The genetic background of ovarian endometriosis appears to be somewhat different from other endometriosis.
5. Family history should be taken into account when investigating chronic pelvic and abdominal pain.





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FINNGEN



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Celmatrix-USA (Caterina Clementi, David-Emlyn Parfitt, Genevieve Galameau, Tina Hu-Seliger, [Piraye Yurttas Beim](#))  
Crete U-Greece (Charoula Matalliotaki, Michail Matalliotakis, [George Goulielmos](#))  
DeCODE-Iceland (Lilja Stefansdottir, Thorleifsson Gudmar, Reynir T Geirsson, [Valgerdur Steinthorsdottir](#), [Stefansson Kari](#))  
Edinburgh U-UK (Andrew Horne, Philippa Saunders)  
FinnGen (Eeva Sliz, Venla Kurra, Liisu Saavalainen, Päivi Härkki, Oskari Heikinheimo, Johannes Kettunen, Hannele Laivuori, [Outi Uimari](#))  
Generation Scotland-UK (Archie Campbell, Alison Murray, [Caroline Hayward](#))  
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WGHS-Harvard U-USA (Rebecca Danning, [Daniel Chasman](#))