

Salivary micro-RNA test for the diagnosis of endometriosis (Endotest)



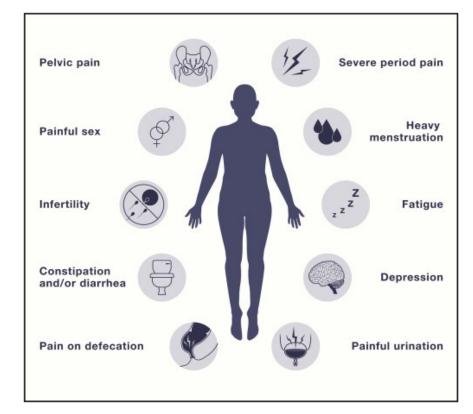
S. Bendifallah, A.Spiers, L.Delbos, M.Poilblanc, F.Golfier, S.Suisse, <u>Ph. Descamps</u> (*Paris-Lyon-Angers*)

Conflicts of interest

None

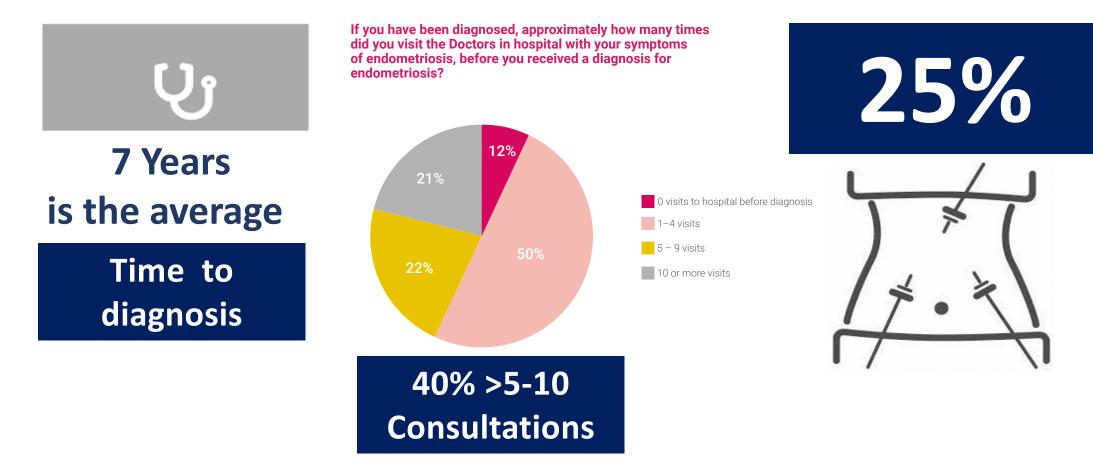
Endometriosis a Complex Disease

- Endometriosis is a common disease affecting 5-10% of women of reproductive age globally
- Endometrial-like tissue outside the uterine cavity
- Characterized mainly by symptoms of pain and infertility.
- Three subtypes ; peritoneal endometriosis, deep endometriosis and endometrioma.

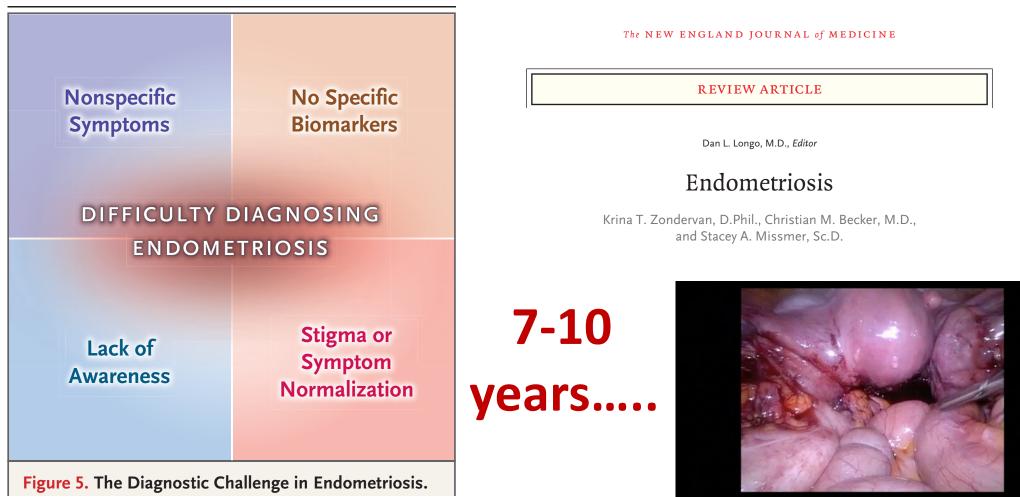


Saunders, Philippa T K, and Andrew W Horne. "Endometriosis: Etiology, pathobiology, and therapeutic prospects." *Cell* vol. 184,11 (2021): 2807-2824. doi:10.1016/j.cell.2021.04.041

A Delayed Diagnosis....



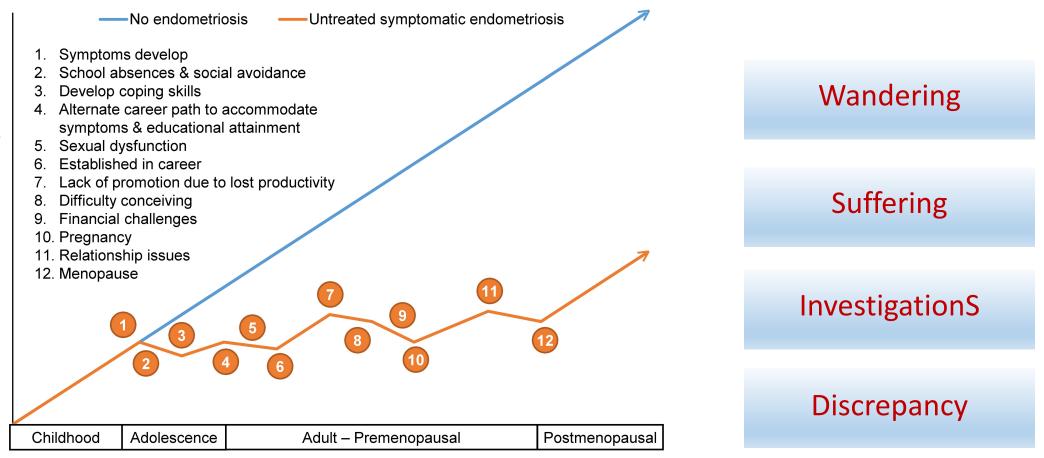
A Delayed diagnosis



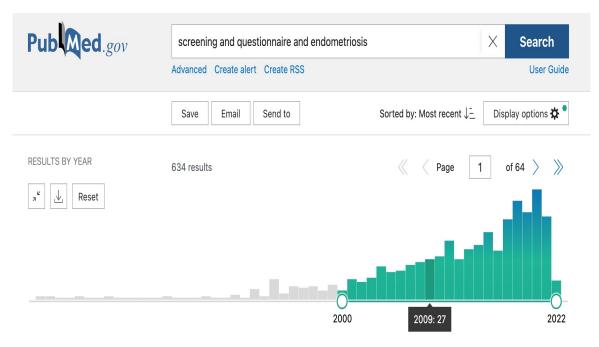


Consequences of Delayed Diagnosis....

Life Course Impact of Untreated Symptomatic Endometriosis



Questionnaires for Endometriosis screening



Patient-completed or symptom-based screening tools for endometriosis: a scoping review

Eric Surrey $^1\cdot$ Cathryn M. Carter $^2\cdot$ Ahmed M. Soliman $^3\cdot$ Shahnaz Khan $^4\cdot$ Dana B. DiBenedetti $^4\cdot$ Michael C. Snabes 3

Feb 2017

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Abstract

Purpose The objective of this review was to evaluate existing patient-completed screening questionnaires and/or symptom-based predictive models with respect to their potential for use as screening tools for endometriosis in adult women. Validated instruments were of particular interest.

Methods We conducted structured searches of PubMed and targeted searches of the gray literature to identify studies reporting on screening instruments used in all studies, as most studies focused on diagnosis versus screening.

Conclusions This literature review did not identify any fully validated, symptom-based, patient-reported questionnaires for endometriosis screening in adult women.

Keywords Endometriosis · Patient-reported · Screener · Self-administered · Symptoms

Endometriosis Diagnostics

Patient-completed or symptom-based screening tools for endometriosis: a scoping review

Eric Surrey¹ · Cathryn M. Carter² · Ahmed M. Soliman³ · Shahnaz Khan⁴ · Dana B. DiBenedetti⁴ · Michael C. Snabes³

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Cathryn M. Carter

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Methods We conducted structured searches of PubMed Self-administered - Symptoms and targeted searches of the gray literature to identify studies reporting on screening instruments used in endometriosis. Studies were screened according to inclu-

sion and exclusion criteria that followed the PICOS (population, intervention, comparison, outcomes, study design) Endometriosis is a painful, inflammatory condition charframework. Results A total of 16 studies were identified, of which 10 outside the uterus [1]. Endometriotic lesions may occur at

described measures for endometriosis in general, 2 various anatomic sites, including the pelvic peritoneum and described measures for endometriosis at specific sites, and the ovary [2]. Deep-infiltrating endometriosis occurs in the described measures for deep-infiltraing endometriosis.
 Only 1 study evaluated a questionnaire that was solely rarely, endometriosis lesions of the bladder, ureter, or patient-completed. Most measures required physician, imaging, or laboratory assessments in addition to patient-An estimated 10% of women of reproductive age are

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Surrey et al.

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completed questionnaires, and several measures relied on complex scoring. Validation for use as a screening tool in siderable clinical, economic, and humanistic burden. adult women with potential endometriosis was lacking in Clinical symptoms include chronic pelvic pain, dysmenorrhea, and infertility [3], and endometriosis may increase a woman's risk of cancer or autoimmune disorders [4, 5]. Numerous studies have demonstrated the considerable economic burden associated with endometriosis [6-8]. Hospitalizations, especially those related to surgical intervention, are a primary direct cost driver for endometriosis 7 9 101 Moreover, endometriosis has a significant social and psychological impact on the lives of women

acterized by the development of endometrial-like tissue

all studies, as most studies focused on diagnosis versu

(L) k

across several domains, including quality of life, intimate relationships, fertility, education and work, and emotional well-being [11, 12].

A new validated screening method for endometriosis diagnosis based on patient questionnaires

Charles Chapron,^{*a,b,c,1*} Marie-Christine Lafay-Pillet,^{*b,1*} Pietro Santulli,^{*a,b,c*} Mathilde Bourdon,^{*a,b*} Chloé Maignien,^{*b*} Antoine Gaudet-Chardonnet,^b Lorraine Maitrot-Mantelet,^b Bruno Borghese,^{a,b,c} and Louis Marcellin,^{a,b,c}

Early identification of women with endometriosis by means of a simple patient-completed questionnaire screening tool: a diagnostic study

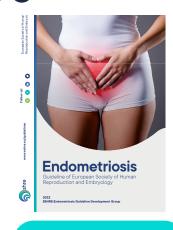
Arnaud Fauconnier, M.D., Ph.D., ^a Hocine Drioueche, M.Sc., ^b Cyrille Huchon, M.D., Ph.D., ^a Joseph Du Cheyron, B.Sc., ^b Emilie Indersie, Ph.D., ^c Yasmine Candau, M.B.A., ^c Pierre Panel, M.D., ^d and Xavier Fritel M.D. Ph.D.

Not reliable enough



No evidence that questionnaires reduce the time to diagnosis

2022



ESHRE guidelines

2000-2017

No fully validated questionnaire for

endometriosis screening

Diagnostic Algorythms in 2023





eshre.eu/guidelines

eshre

2022

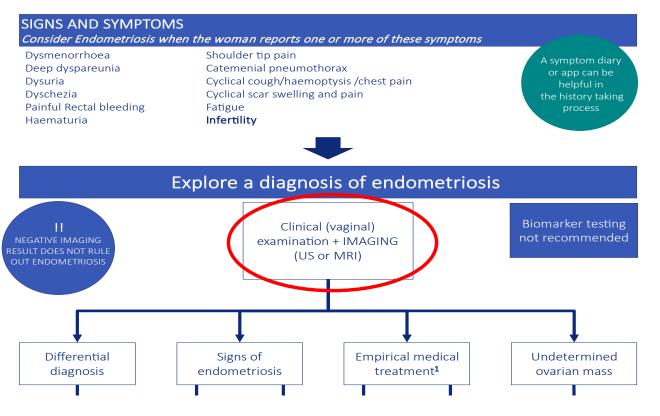


Endometriosis Guideline of European Society of Human

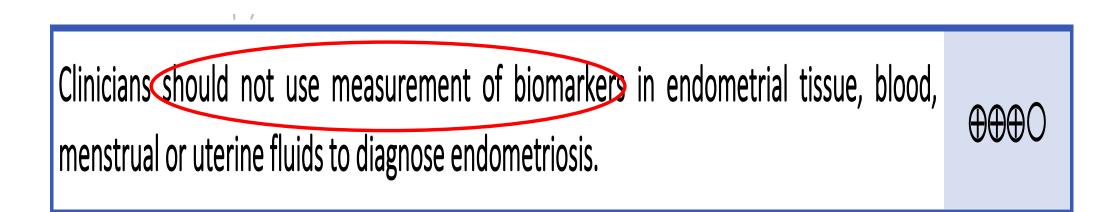
Reproduction and Embryology

ESHRE Endometriosis Guideline Development Group

DIAGNOSIS OF ENDOMETRIOSIS



2022 ESHRE recommendations: Diagnosis of endometriosis



ESHRE 2022 Guidelines

Recommendations (5-7)

Clinicians are recommended to use imaging (US or MRI) in the diagnostic work-up for endometriosis, but they need to be aware that a negative finding does not exclude endometriosis, particularly superficial peritoneal disease.

 $\oplus \oplus \bigcirc \bigcirc$

GPP

In patients with negative imaging results or where empirical treatment was unsuccessful or inappropriate, the GDG recommends that clinicians consider offering laparoscopy for the diagnosis and treatment of suspected endometriosis.

The GDG recommends that laparoscopic identification of endometriotic lesions is confirmed by histology although negative histology does not entirely rule out the disease.

Shre www.stre.eu/guidelines

Endometriosis

Guideline of European Society of

Reproduction and Embryology

ESHRE 2022 Guidelines



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Endometriosis

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Recommendations (5-7)

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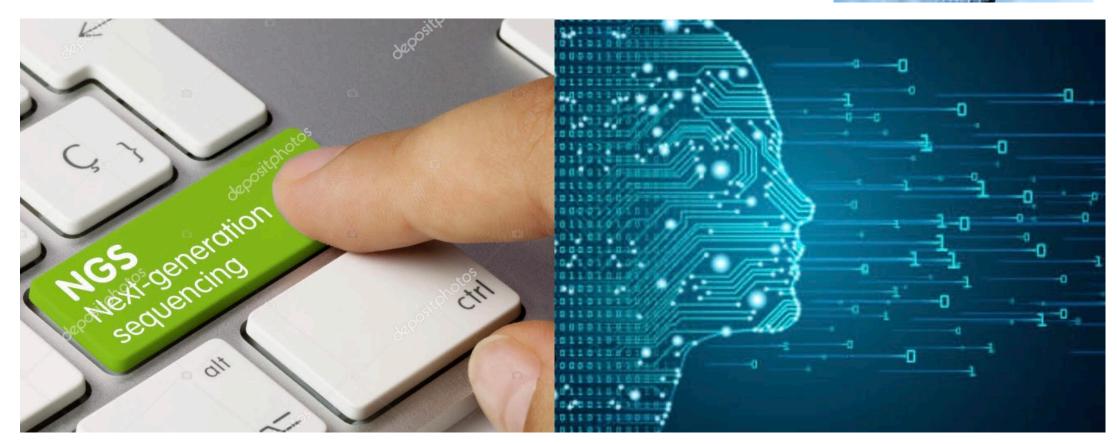
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GPP

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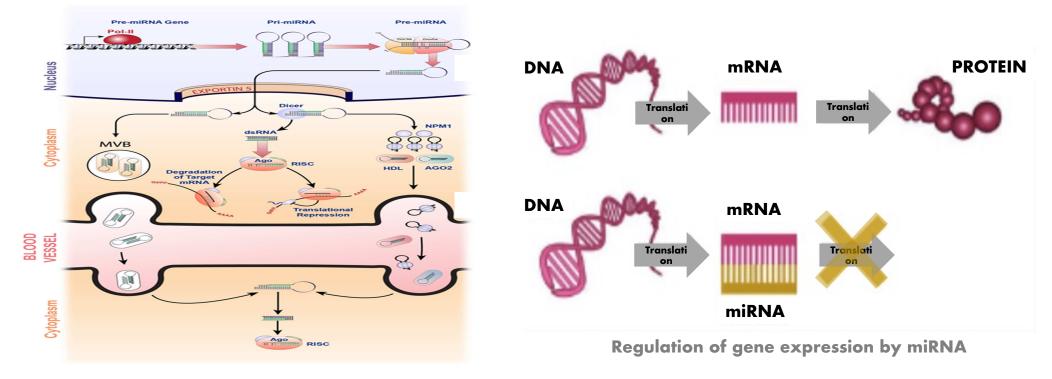
	Endometriosis Diagnostic Tests				
Endometriosis test					X al
	Medical questionnaire	Ultrasound imaging	MRI imaging	Blood test	Surgery and pathology
General value	++	++	+++	-	++++
Performance value - Sensitivity - Specificity	76-98% 20-58%	65-79% 91-95%	79% 72%	63% 69%	90-94% 40-79%
Reliability	Very Low specificity	Low accuracy for early stage lesion	Low accuracy for early stage lesion	-	Yes
	+	++	++		++++
Reproducibility	+	+	+++	-	+++
Be simple, safe	++++	++	+++	+++	+
Acceptability	++++	+++	++	++	+
Detect disease early in its natural history	+	++	+	-	++++

miRNA – NGS and Al

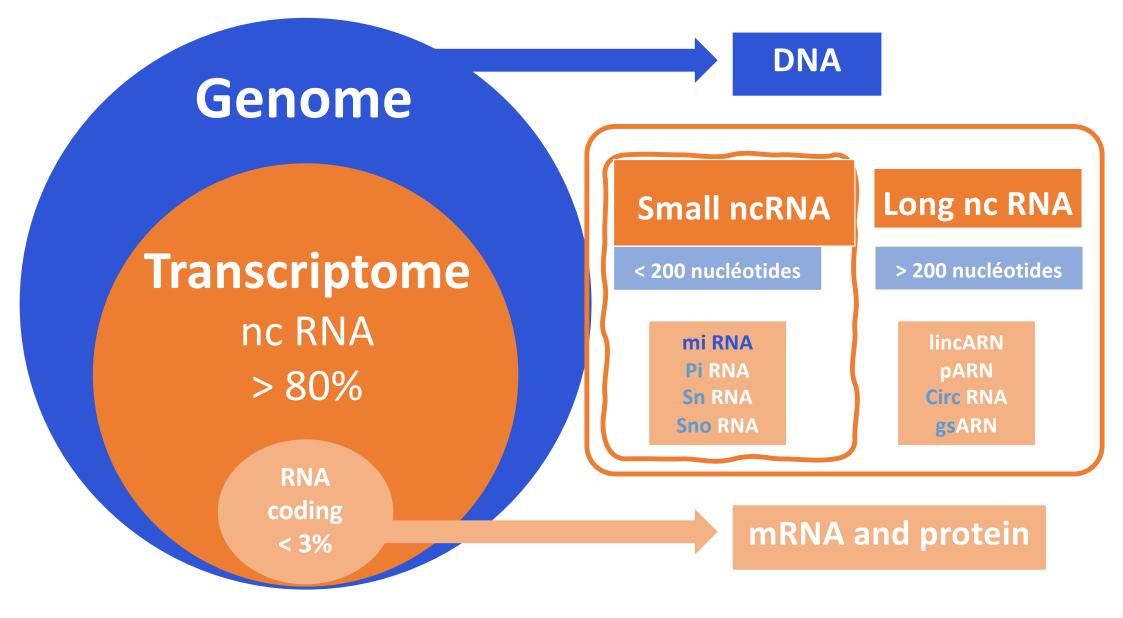




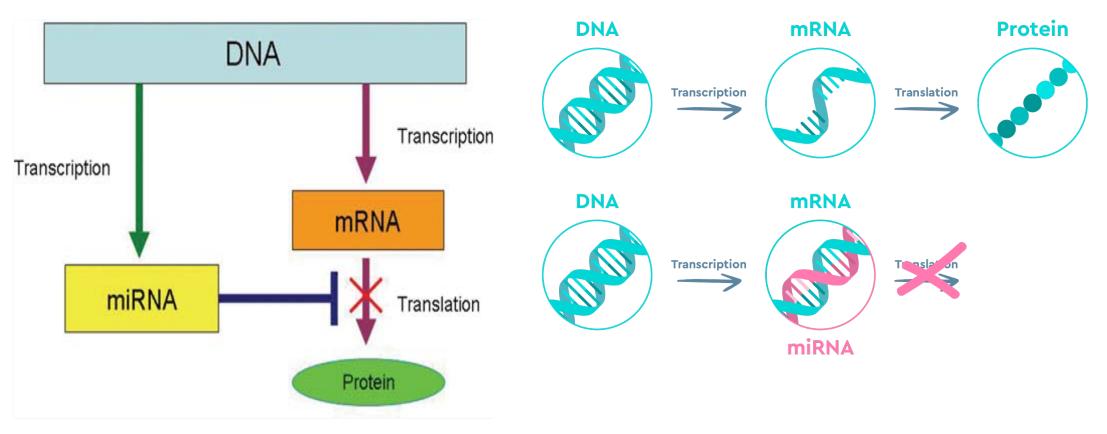
Biosynthesis of miRNAs Detection in biofluids



Nothnick et al. J Min Gynecol 2016

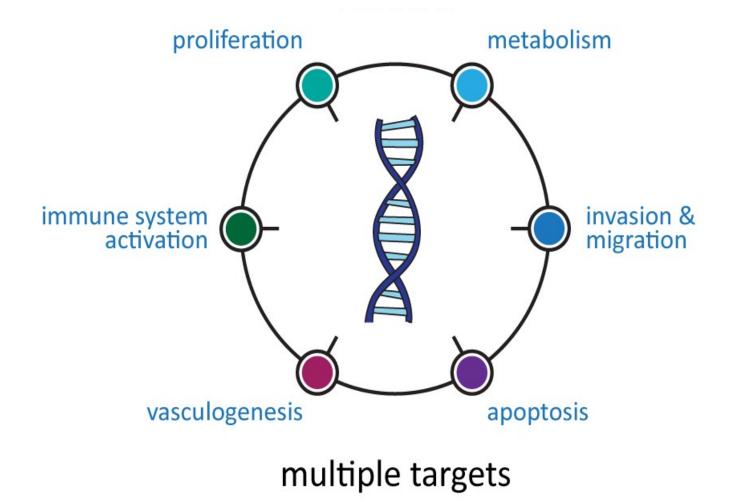


Regulation of Gene Expression by miRNA



Guo, Sun-Wei. "Epigenetics of endometriosis." *Molecular human reproduction* vol. 15,10 (2009): 587-607. doi:10.1093/molehr/gap064

miARNs function



Cancer Neurology Auto immune

ORIGINAL ARTICLE

Endocrine Research

The pioneers....

Individual analysis of 6 mi-RNA

No exhaustive analysis of mi-RNome

No NGS, no Al

Circulating MicroRNAs Identified in a Genome-Wide Serum MicroRNA Expression Analysis as Noninvasive Biomarkers for Endometriosis

Wen-Tao Wang,* Ya-Nan Zhao,* Bo-Wei Han, Shun-Jia Hong, and Yue-Qin Chen

Department of Obstetrics and Gynecology, Sun Yat-sen Memorial Hospital (Y.-N.Z., S.-J.H.), Key Laboratory of Gene Engineering of the Ministry of Education, and State Key Laboratory for Biocontrol (W.-T.W., B.-W.H., Y.-Q.C.), Sun Yat-sen University, Guangzhou 510120, China

Context: There is currently no reliable noninvasive biomarker for the clinical diagnosis of endometriosis. Previous analyses have reported that circulating microRNAs (miRNAs) can serve as biomarkers for a number of diseases.

Objective: The study aims to detect the serum miRNAs that are differentially expressed between endometriosis patients and negative controls to evaluate the potential of these miRNAs as diagnostic markers for endometriosis.

Design: A total of 765 serum miRNAs were profiled using a TaqMan microRNA array in a pool of 10 endometriosis patients and a pool of 10 negative controls, and a set of selected miRNAs were further analyzed in a validation cohort consisting of sera from 60 patients and 25 controls including 10 samples used in array profiling.

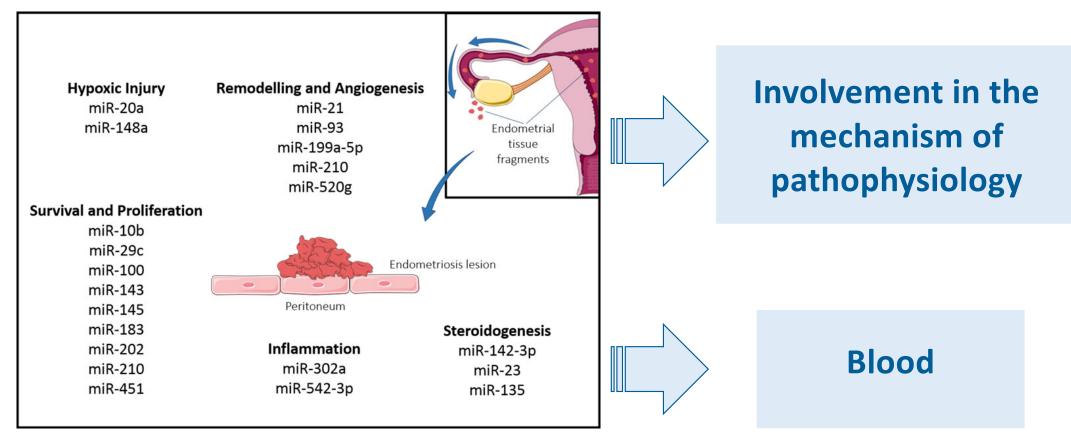
Results: The relative expression levels of miR-199a and miR-122 were found to be up-regulated in endometriosis patient samples compared with control samples, whereas miR-145*, miR-141*, miR-542-3p, and miR-9* were down-regulated in endometriosis patients. Importantly, the relative expression of miR-199a (P < 0.05) and miR-122 can be used to discriminate between severe and mild endometriosis. We also found that miR-199a is well correlated with pelvic adhesion and lesion distribution (P < 0.05) and associated with hormone-mediated signaling pathways. Furthermore, we investigated the diagnostic value of these molecules and confirmed the optimal combination of miR-199a, miR-122, miR-145*, and miR-542-3p with area under the curve of 0.994 (95% confidence interval = 0.984–1.000, P < 0.001) and a cutoff point (0.4950) of 93.22% sensitivity and 96.00% specificity.

Conclusions: Our study demonstrated that the circulating miRNAs miR-199a, miR-122, miR-145*, and miR-542-3p could potentially serve as noninvasive biomarkers for endometriosis. miR-199a may also play an important role in the progression of the disease. This is the first report that circulating miRNAs serve as biomarkers of endometriosis. *(J Clin Endocrinol Metab* 98: 281–289, 2013)

Non-coding RNAs in endometriosis: a narrative review

2018 Human Reprod. Update

Kavita Panir ¹, ^{*}, John E. Schjenken¹, Sarah A. Robertson¹, and M. Louise Hull^{1,2,3}



GYNECOLOGY Accurate diagnosis of endometriosis using serum microRNAs

2020 AJOG

Sarah Moustafa, MD; Martina Burn, MD¹; Ramanaiah Mamillapalli, PhD¹; Sepide Nematian, MD; Valerie Flores, MD; Hugh S. Taylor, MD

TABLE 2	
ROC analysis of individual miRNAs	

ROC model	Area	SE	95% Wald confidence limits	Optimal cutoff	Correct, %	Sensitivity, %	Specificity, %
miR_125b	0.73	0.05	0.63-0.83	0.084	68.0	56.1	78.0
miR_150	0.68	0.06	0.57—0.78	0.44	63.9	20.0	94.7
miR_342	0.92	0.04	0.86-0.99	0.085	90.8	90.0	91.2
miR_451a	0.84	0.04	0.76-0.92	0.35	79.8	90.0	72.9
miR_3613	0.76	0.05	0.66—0.85	0.014 ^a	74.0	92.7	61.0
let_7b	0.78	0.05	0.69—0.87	0.012 ^a	73.7	82.5	67.8



Article



Clues for Improving the Pathophysiology Knowledge for Endometriosis Using Serum Micro-RNA Expression

Yohann Dabi ^{1,2,3}, Stéphane Suisse ⁴, Ludmila Jornea ⁵, Delphine Bouteiller ⁶, Cyril Touboul ^{1,2,3}, Anne Puchar ¹, Emile Daraï ¹ and Sofiane Bendifallah ^{1,2,*}

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MDPI

Article

Clues for Improving the Pathophysiology Knowledge for Endometriosis Using Serum Micro-RNA Expression

Yohann Dabi 12.3, Stéphane Suisse 4, Ludmila Jornea 5, Delphine Bouteiller 6, Cyril Touboul 12.3, Anne Puchar 1, Emile Daraï 1 and Sofiane Bendifallah 12.*

Diagnostic Cartography



Jornea, L.; Bouteiller, D.; Touboul, C.; Puchar, A.; Dar, Bendifallah, S. Clues for Imp the Pathophysiology Knowle Endometriosis Using Serum Micro 175. https://doi.org/10.3390/ diagnostics12010175

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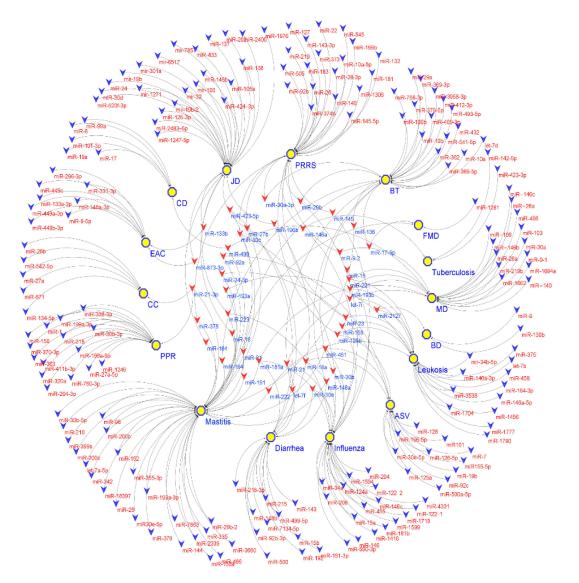


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Phenotypic Cartography

Evolutive Cartography





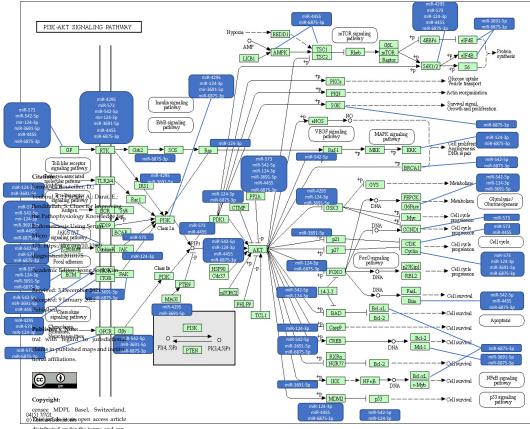
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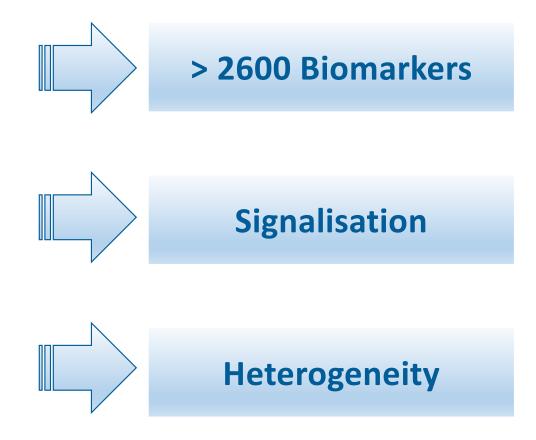
2022 Diagnostics

Article

Clues for Improving the Pathophysiology Knowledge for Endometriosis Using Serum Micro-RNA Expression

Yohann Dabi ^{1,2,3}, Stéphane Suisse ⁴, Ludmila Jornea ⁵, Delphine Bouteiller ⁶, Cyril Touboul ^{1,2,3}, Anne Puchar ¹, Emile Daraï ¹ and Sofiane Bendifallah ^{1,2,*}





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OPEN MicroRNome analysis generates a blood-based signature for endometriosis

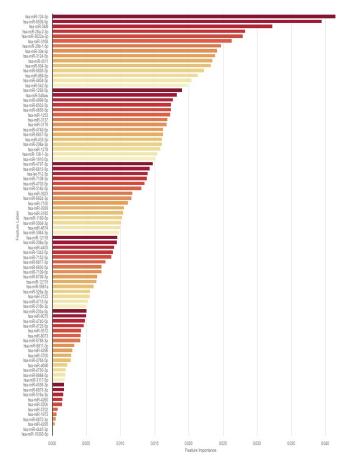
Sofiane Bendifallah^{1,22}, Yohann Dabi^{1,2,3}, Stéphane Suisse², Ludmila Jornea⁴, Delphine Bouteiller⁵, Cyril Touboul^{1,2}, Anne Puchar^{1,2} & Emile Daraï^{1,2}

Endometriosis, characterized by endometrial-like tissue outside the uterus, is thought to affect 2–10% of women of reproductive age: representing about 190 million women worldwide. Numerous studies have evaluated the diagnostic value of blood biomarkers but with disappointing results. Thus, the gold standard for diagnosing endometriosis remains laparoscopy. We performed a prospective trial, the ENDO-miRNA study, using both Artificial Intelligence (AI) and Machine Learning (ML), to analyze the current human miRNome to differentiate between patients with and without endometriosis, and to develop a blood-based microRNA (miRNA) diagnostic signature for endometriosis. Here, we present the first blood-based diagnostic signature obtained from a combination of two robust and disruptive technologies merging the intrinsic quality of miRNAs to condense the endometriosis phenotype (and its heterogeneity) with the modeling power of AI. The most accurate signature provides a sensitivity, specificity, and Area Under the Curve (AUC) of 96.8%, 100%, and 98.4%, respectively, and is sufficiently robust and reproducible to replace the gold standard of diagnostic surgery. Such a diagnostic approach for this debilitating disorder could impact recommendations from national and international learned societies.

nature scientific reports

MicroRNome analysis generates a blood-based signature for

endometriosis



2022 Nature Scientific Reports



2600 Biomarkers

86 Biomarkers

Sensitivity 96.7%, specificity 100%

Accuracy > 98%

Overview of miRNAs for the non-invasive diagnosis of endometriosis: evidence, challenges and strategies. A systematic review

	,				•	•		
Total	EU <i>versus</i> EN	EC versus EN	EC versus EU	Plasma	Serum	Blood	PF	
6	miR-145	miR-145	miR-145	miR-145	miR-145			Wang et al.,(et al., ⁽⁴⁵⁾ Cosa
5		miR-200b	miR-200b	miR-200b				Saare et al. Filighedduet
5	miR-424		miR-424		miR-424	miR-424		Braza-Boils e Wang et al. ⁽⁵¹
4	miR-199a	miR-199a			miR-199a		miR-199a	Wang et al.,(
4		miR-141	miR-141	mi R -141	miR-141			Wang et al., Rekkeret al. ^{(*}
4		miR-20a		mi <mark>l</mark> -20a	miR-20a	miR-20a		Zhao et al., ⁽³⁷
4		miR-200a	miR-200a	miF-200a				Saare et al., ⁽³ et al. ⁽⁵⁷⁾
3	miR-29c	miR-29c	miR-29c					Braza-Boils e
3	miR-34c	miR-34c	miR-34c					Braza-Boïls e
3		miR-200c	miR-200c					Liang et al., ⁽³
3		miR-21	miR-21			miR-21		Haikalis et al
3	miR-126		miR-126			miR-126		Liu et al., ⁽²⁰⁾ (
3			miR-16	miR-16		miR-16		Yang et al., ⁽⁴³⁾
3			miR-451a		miR-451a		miR-451a	Nothnick et a
3	miR-9		NiB.0		miR-9			Wang et al.,(

2021 Einstein Journal

Likewise, none of the papers examined investigated miRNAs in saliva. To date, there are no scientifically proven salivary biomarkers for endometriosis. Saliva is a suitable and desirable medium for biomarker detection^(96,97) and its applicability to the diagnosis of endometriosis has been explored previously.^(98,99) Saliva is widely available and can be easily collected in a non-invasive manner, at low cost and with minimal discomfort. Therefore, it is an ideal fluid for biomarker investigation and is attracting great interest in the public health sector. The use of saliva for miRNA identification could be a potential non-invasive solution to overcome current barriers to the diagnosis of endometriosis.

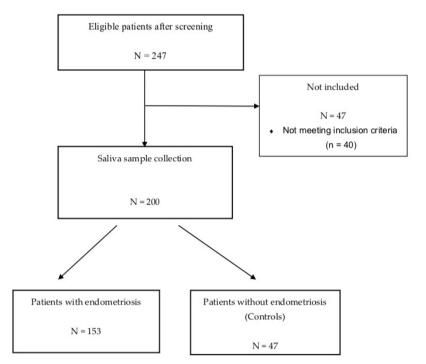


Clinical Medicine



Article Salivary MicroRNA Signature for Diagnosis of Endometriosis

Sofiane Bendifallah ^{1,2,*}, Stéphane Suisse ³, Anne Puchar ^{1,2}, Léa Delbos ^{4,5}, Mathieu Poilblanc ^{6,7}, Philippe Descamps ^{4,5}, Francois Golfier ^{6,7}, Ludmila Jornea ⁸, Delphine Bouteiller ⁹, Cyril Touboul ^{1,2}, Yohann Dabi ^{1,2} and Emile Daraï ^{1,2}



2022 Journal of **Clinical Medicine**





Figure 1. Flow chart of ENDO-miRNA study.





Artide Salivary MicroRNA Signature for Diagnosis of Endometriosis

Sofiane Bendifallah ^{1,2,*}, Stéphane Suisse ³, Anne Puchar ^{1,2}, Léa Delbos ^{4,5}, Mathieu Poilblanc ^{6,7}, Philippe Descamps ^{4,5}, Francois Golfier ^{6,7}, Ludmila Jornea ⁸, Delphine Bouteiller ⁹, Cyril Touboul ^{1,2}, Yohann Dabi ^{1,2} and Emile Daraï ^{1,2}

2022 Journal of Clinical Medicine

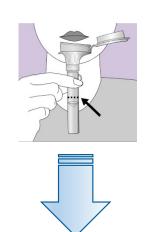
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- ⁹ Genotyping and Sequencing Core Facility, iGenSeq, Institut du Cerveau et de la Moelle Epinière, Institut du Cerveau, Hôpital Pitié-Salpêtrière, 47-83 Boulevard de l'Hôpital, 75013 Paris, France; delphine.bouteiller@icm-institute.org
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	Control Patients N (%) N = 47	Patients with Endometriosis N (%) N = 153		-
Age (years) (mean \pm SD)	30,92 (13.79)	31.17 (12.78)	0.1912	-
Age range - Less than 30 years - Over 30 years	72% (34) 28% (13)	63% (96) 37% (57)	0.294	-
BMI (body mass index) (mean \pm SD)	24.84 (11.10)	24.36 (8.38)	0.525	-
Infertility - Yes - No	17% (8) 83% (39)	24% (36) 76% (117)	0.556	-
rASRM classification - I–II - III–IV	<u>-</u>	52% (80) 48% (73)	-	-
Control diagnoses - No abnormality - Leiomyoma - Cystadenoma - Teratoma - Other gynecologic disorders	51% (24) 2% (1) 11% (5) 23% (11) 13% (6)	_	_	Population Characteristics
Dysmenorrhea	100%	100%		
Abdominal pain outside menstruation - Yes	66% (21)	71% (89)	0.6905	-
Pain suggesting sciatica - Yes	31% (10)	56% (70)	0.0214	
Lower back pain outside menstruation - Yes	62% (20)	81% (101)	0.0498	-
Right shoulder pain during menstruation - Yes	9% (3)	21% (26)	0.2184	_
Blood in the stools during menstruation - Yes	12% (4)	24% (30)	0.2425	
Blood in urine during menstruation - Yes	25% (8)	17% (21)	0.4172	-
Diagnostic method - Surgery - Magnetic Resonance Imaging	47 (100)	83 (54.2) 70 (45.8)	_	-

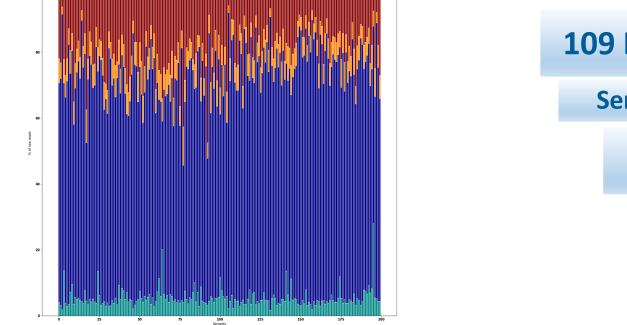


Article Salivary MicroRNA Signature for Diagnosis of Endometriosis

Sofiane Bendifallah ^{1,2,*}, Stéphane Suisse ³, Anne Puchar ^{1,2}, Léa Delbos ^{4,5}, Mathieu Poilblanc ^{6,7}, Philippe Descamps ^{4,5}, Francois Golfier ^{6,7}, Ludmila Jornea ⁸, Delphine Bouteiller ⁹, Cyril Touboul ^{1,2}, Yohann Dabi ^{1,2} and Emile Daraï ^{1,2}



MDPI





Sensitivity 96.7%, specificity 100%

All stages

Simplicity

Reliability





Article

A Bioinformatics Approach to MicroRNA-Sequencing Analysis Based on Human Saliva Samples of Patients with Endometriosis

Sofiane Bendifallah ^{1,2,3,*}, Yohann Dabi ^{1,2,3}, Stéphane Suisse ⁴, Ludmila Jornea ⁵, Delphine Bouteiller ⁶, Cyril Touboul ^{1,2,3}, Anne Puchar ¹ and Emile Daraï ^{1,2}

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- ² Clinical Research Group (GRC) Paris & Endometriosis Expert Center (C3E), Sorbonne University (GRC6 C3E SU), 75020 Paris, France
- ³ Cancer Biology and Therapeutics, Centre de Recherche Saint-Antoine (CRSA), Sorbonne University, INSERM UMR_S_938, 75020 Paris, France
- ⁴ Ziwig Health, 19 rue Reboud, 69003 Lyon, France; stephane@ziwig.com
- ⁵ Paris Brain Institute Institut du Cerveau-ICM, Sorbonne University, Inserm U1127, CNRS UMR 7225, AP-HP-Hôpital Pitié Salpêtrière, 75013 Paris, France; ludmila jorne a@icm-institute.org
- ⁶ Gentoyping and Sequencing Core Facility, iGenSeq. Institut du Cerveau et de la Moelle Épinière, ICM, Hôpital Pitié-Salpêtrière, 47-83 Boulevard de l'Hôpital, 75013 Paris, France; delphine.bouteiller@icm-institute.org
- * Correspondence: sofiane.bendifallah@aphp.fr; Tel.: +33-1-56-01-73-18

C check for updates

Citation: Bendifallah, S.; Dabi, Y.; Suisse, S.; Jornea, L.; Bouteiller, D.; Touboul, C.; Puchar, A.; Daraï, E. A Bioinformatics Approach to MicroRNA-Sequencing Analysis Based on Human Saliva Samples of Patients with Endometriosis. Int. J. Mol. Sci. 2022, 23, 8045. https:// doi.org/10.3390/ijms23148045

Academic Editor: David B. Alexander

Received: 3 June 2022 Accepted: 16 July 2022 Published: 21 July 2022

Publisher's Note: MDPI stays neutral with regard to jurisdictional claims in published maps and institutional affiliations. Abstract: Endometriosis, defined by the presence of endometrium-like tissue outside the uterus, affects 2-10% of the female population, i.e., around 190 million women, worldwide. The aim of the prospective ENDO-miRNA study was to develop a bioinformatics approach for microRNAsequencing analysis of 200 saliva samples for miRNAome expression and to test its diagnostic accuracy for endometriosis. Among the 200 patients, 76.5% (n = 153) had confirmed endometriosis and 23.5% (n = 47) had no endometriosis (controls). Small RNA-seq of 200 saliva samples yielded ~4642 M raw sequencing reads (from ~13.7 M to ~39.3 M reads/sample). The number of expressed miRNAs ranged from 1250 (outlier) to 2561 per sample. Some 2561 miRNAs were found to be differentially expressed in the saliva samples of patients with endometriosis compared with the control patients. Among these, 1.17% (n = 30) were up- or downregulated. Among these, the F1-score, sensitivity, specificity, and AUC ranged from 11-86.8%, 5.8-97.4%, 10.6-100%, and 39.3-69.2%, respectively. Here, we report a bioinformatic approach to saliva miRNA sequencing and analysis. We underline the advantages of using saliva over blood in terms of ease of collection, reproducibility, stability, safety, non-invasiveness. This report describes the whole saliva transcriptome to make miRNA quantification a validated, standardized, and reliable technique for routine use. The methodology could be applied to build a saliva signature of endometriosis.

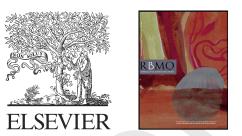
Keywords: endometriosis; miRNA; NGS; bioinformatics; saliva

2022 International Journal of Molecular Science

Saliva > Blood

- Ease of collection
- Non invasive
- Safety
- Stability
- Reproductibility





ARTICLE

Endometriosis-associated infertility diagnosis based on saliva microRNA signatures

Yohann Dabi 🛚 Stéphane Suisse 🔍 Anne Puchar 🔍 ... Cyril Touboul 🔍 Emile Daraï 🔍 Sofiane Bendifallah 🔗 🖂 🔍

October 2022



 $Log in Q \equiv$

FULL LENGTH ARTICLE | ARTICLES IN PRESS

Endometriosis–associated infertility diagnosis based on saliva microRNA signatures

Yohann Dabi • Stéphane Suisse • Anne Puchar
Léa Delbos • Mathieu Poilblanc • Philippe Descamps
Julie Haury • Francois Golfier • Ludmila Jornea •
Delphine Bouteiller • Cyril Touboul • Emile Daraï
Sofiane Bendifallah 옷 🖂 - Show less
Published: September 26, 2022 •
DOI: https://doi.org/10.1016/j.rbmo.2022.09.019

153 patients diagnosed with endometriosis 24 % were infertile, 76 % were fertile Of the 2561 known mi-RNA, the feature selection method generated a signature of 34 miRNAs linked to endometriosis associated infertility.

Those results still require external validation before using the signature in routine practise.

miRNA Salivary Signature for the Diagnosis of Endometriosis

NCT 05244668

« 1000 Study »

NIH National Library of Medicine

ClinicalTrials.gov

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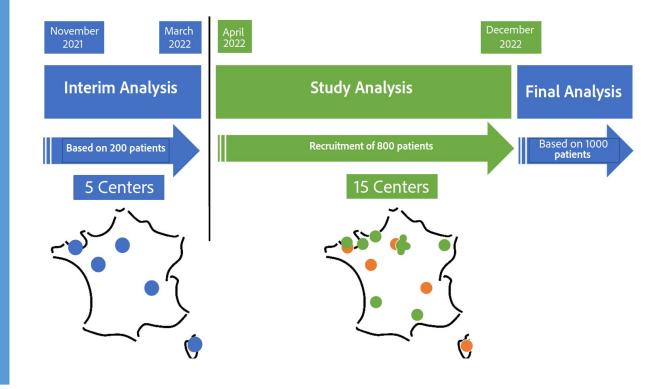
RECRUITING

ClinicalTrials.gov Identifier: NCT05244668

PRS Login

Multicenter Validation of the Salivary miRNA Signature of Endometriosis

Information provided by ZIWIG (Responsible Party) Last Update Posted: 2022-12-29





miRNA Salivary Signature for the Diagnosis of Endometriosis External Validation Study

NCT 05244668

Interim analysis

Study Design

Time persective : **Prospective** Locations : **Multicenter** (15 centres across France) **Cohort : 1150 patients**

Eligibility criteria :

- 18 to 43 years old (Adult)
- Suspected/ Diagnosed Endometriosis

Primary outcome :

Assess diagnostic accuracy of miRNA Salivary Signature

Epidemiologic, clinical, and saliva sequencing data were collected between **Nov. 2021 and March 2022.**

« » »



Publication in NEJM Evidence

June 9, 2023

DOI: 10.1056/EVIDoa2200282



Published June 9, 2023

DOI: 10.1056/EVIDoa2200282

ORIGINAL ARTICLE

Validation of a Salivary miRNA Signature of Endometriosis – Interim Data

Sofiane Bendifallah, M.D., Ph.D.,^{1,2,3} Yohann Dabi, M.D.,^{1,2,3} Stéphane Suisse,⁴ Léa Delbos, M.D.,^{5,6} Andrew Spiers, M.D.,⁴ Mathieu Poilblanc, M.D.,^{7,8} Francois Golfier, M.D., Ph.D.,^{7,8} Ludmila Jornea, Msc.,⁹ Delphine Bouteiller, M.D.,¹⁰ Hervé Fernandez, M.D., Ph.D.,¹¹ Alexandra Madar, M.D.,¹ Erick Petit, M.D.,¹² Frédérique Perotte,¹² Raffaèle Fauvet, M.D., Ph.D.,¹³ Michael Benjoar, M.D.,¹⁴ Cherif Akladios, M.D., Ph.D.,¹⁵ Vincent Lavoué, M.D., Ph.D.,¹⁶ Thomas Darnaud, M.D.,¹⁷ Benjamin Merlot, M.D.,¹⁸ Horace Roman, M.D., Ph.D.,¹⁸ Cyril Touboul, M.D., Ph.D.,^{1,2,3} and Philippe Descamps, M.D., Ph.D.,^{5,6}



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miRNA Salivary Signature for the Diagnosis of Endometriosis External Validation Study

NEJM Evidence

Published June 9, 2023

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ORIGINAL ARTICLE

Validation of a Salivary miRNA Signature of Endometriosis — Interim Data

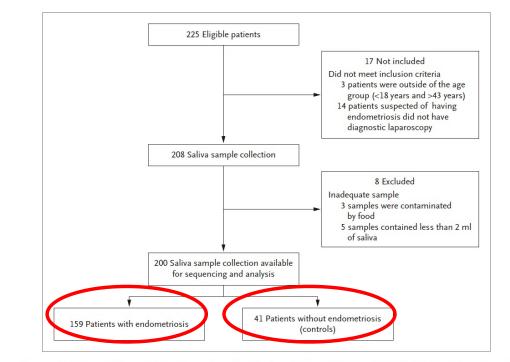


Figure 2. Flow Chart of the Interim Analysis of the ENDOmiRNA Saliva Test Study. This flow chart shows the recruitment of patients and provides the details of noninclusion and exclusion.

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miRNA Salivary Signature for the Diagnosis of Endometriosis External Validation Study

<mark>(Nејм</mark> Evidence

ORIGINAL ARTICLE

Validation of a Salivary miRNA Signature of Endometriosis — Interim Data

Accuracy of the Saliva-Based Diagnostic Signature for Endometriosis

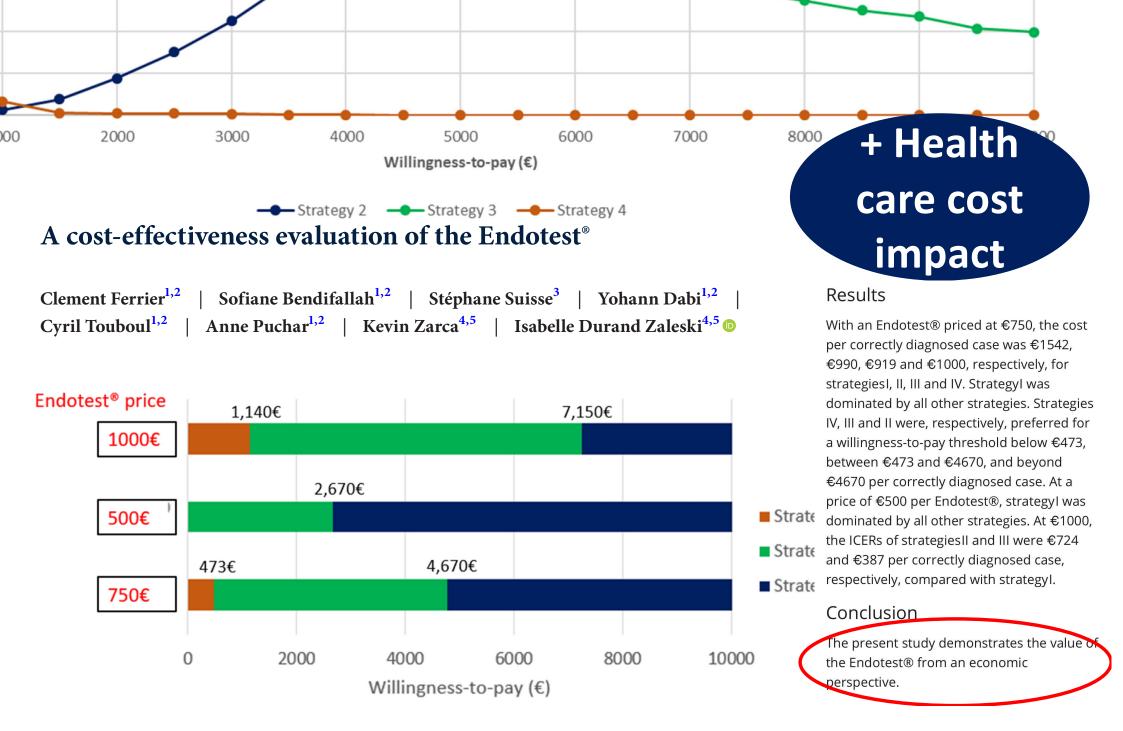
With an overall population prevalence of 79.5%, the diagnostic signature composed of 109 miRNAs (random forest model) against the validation cohort obtained a sensitivity of 96.2% (95% CI, 93.7 to 97.3%), specificity of 95.1% (95% CI, 85.2 to 99.1%), PPV of 95.1% (95% CI, 85.2 to 99.1%), NPV of 86.7% (95% CI, 77.6 to 90.3%), positive likelihood ratio of 19.7 (95% CI, 6.3 to 108.8), negative likelihood ratio of 0.04 (95% CI, 0.03 to 0.07), and AUC of 0.96 (95% CI, 0.92 to 0.98).

Interim Data Analysis - Results :

- Sensitivity of 96.2% (95% CI: 93.7%-97.3%)
- Specificity of 95.1% (95% CI: 85.2%-99.1%)



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miRNA Salivary Signature for the Diagnosis of Endometriosis Cost-Effectiveness Study

DOI: 10.1111/1471-0528.17348



Study Design

Comparison of 4 diagnostic pathways :

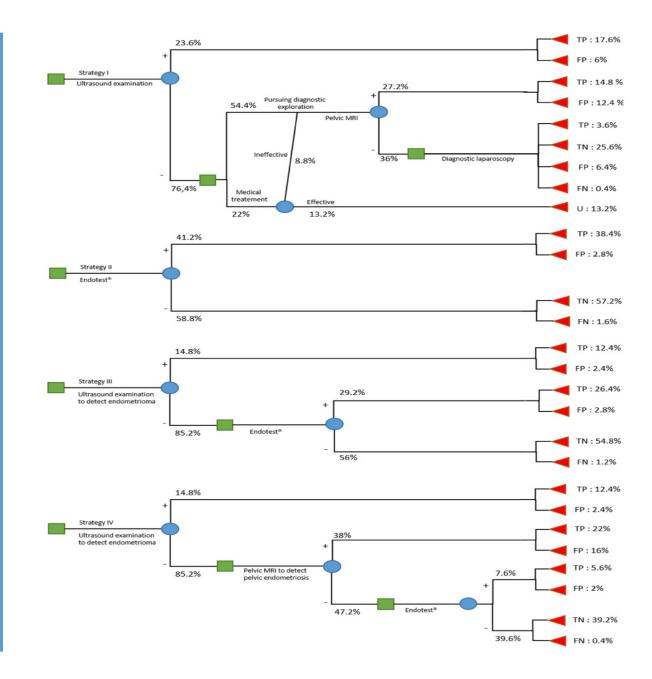
- **1** Current French Guidelines
- 2 Endotest
- 3 Ultrasound -> Endotest
- 4 Ultrasound -> MRI -> Endotest



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miRNA Salivary Signature for the Diagnosis of Endometriosis Cost-Effectiveness Study

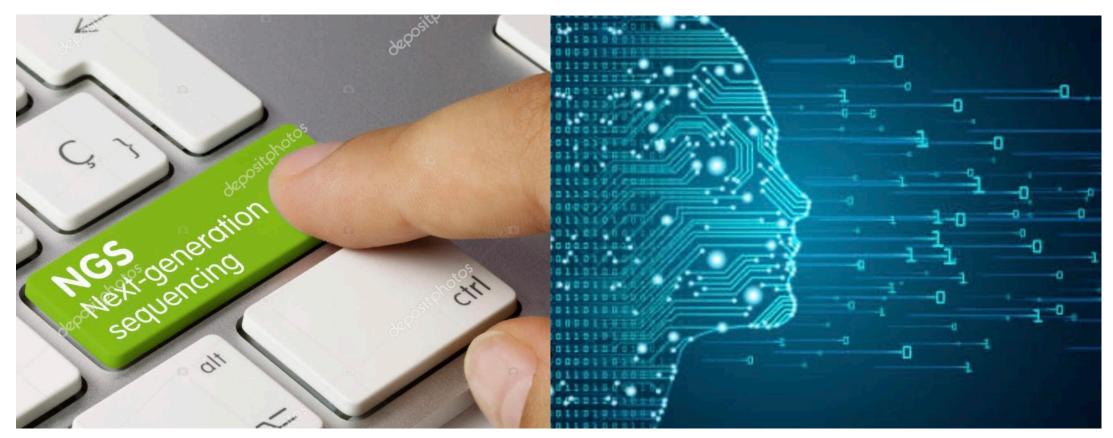
DOI: 10.1111/1471-0528.17348





Endotest: Indications

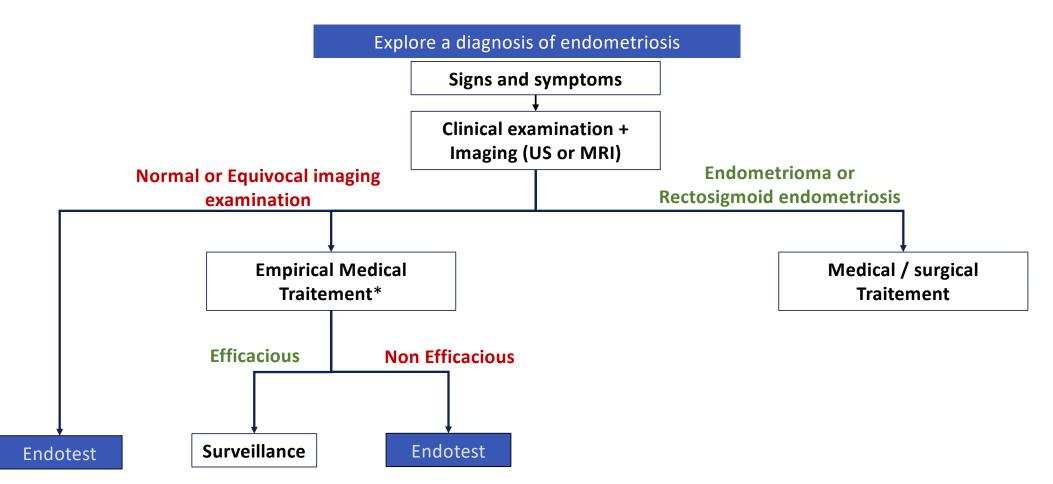




Endotest[®] Kit







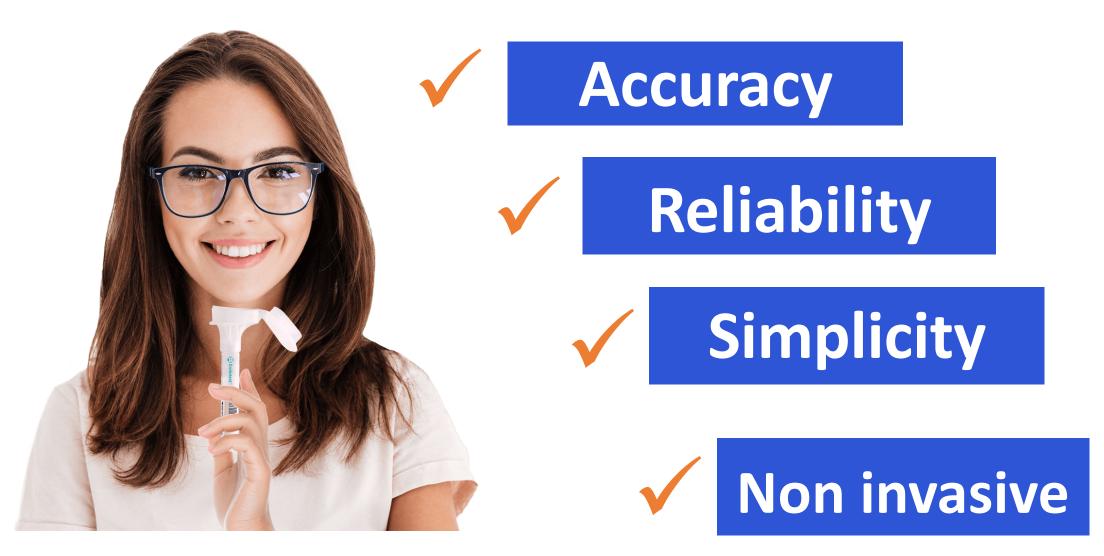
* The Endotest can be offered to patients who do not wish to take empirical medical treatment

Conclusions (1)

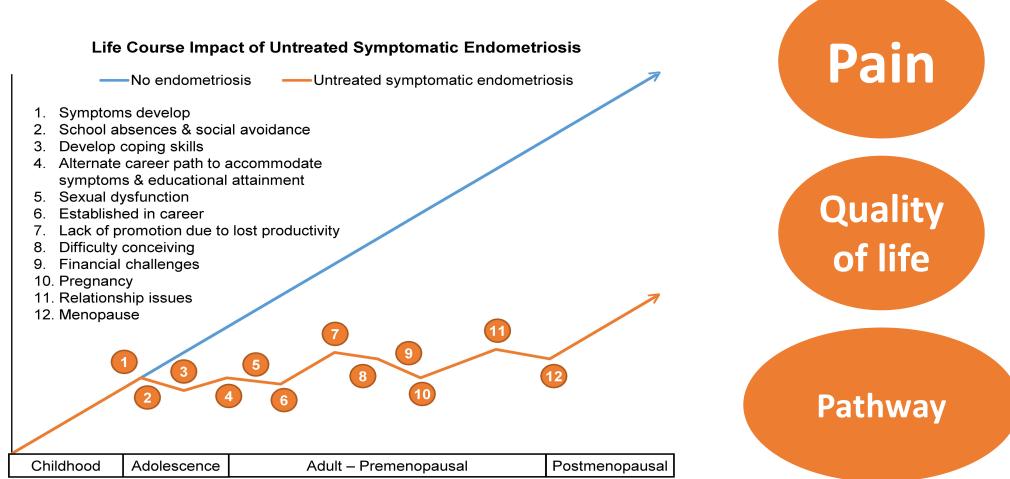
- Endotest is a salivary mi-RNA non invasive diagnostic test, should not be used for screening
- Interim results of the external validation confirm the relevance of the signature Spe : 95,1, Se :96,2 (June 9th 2023-NEJM-Evidence)
- Endotest is available in Switzerland since october 2022, in Germany since December 2022, and in total 10 European countries in 1st semester 2023

Endometriosis Diagnostic Tests

Endometriosis test	(Intersection of the second se	Ultrasound imaging	MRI imaging	Blood test	کری کر	Kerter Saliva Test
General value	++	++	+++	-	++++	+++++
Performance value - Sensitivity - Specificity	76-98% 20-58%	65-79% 91-95%	79% 72%	63% 69%	90-94% 40-79%	> 95% > 95%
Reliability	Very Low specificity	Low accuracy for early stage lesion	Low accuracy for early stage lesion	-	Yes	Yes
	+	++	++		++++	++++
Reproducibility	+	+	+++	-	+++	+++++
Be simple, safe	++++	++	+++	+++	+	+++++
Acceptability	++++	+++	++	++	+	+++++
Detect disease early in its natural history	+	++	+	-	++++	+++++







Conclusions (2)

- AI : Promizing path to revolutionize women's health
- Gyn RNA study : salivary test able to give several diagnosis
- Next steps :
 - . Informations from the 1000 study (Phenotypes)
 - . Teenagers study
 - . Fertility
 - . Adenomyosis
 - . Ovarian cancer

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2022-23



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International Journal of Molecular Sciences

<u>(Nejm</u> Evidence Published June 9, 2023

DOI: 10.1056/EVIDoa2200282



Validation of a Salivary miRNA Signature of Endometriosis — Interim Data

Sofiane Bendifallah, M.D., Ph.D.,^{1,2,3} Yohann Dabi, M.D.,^{1,2,3} Stéphane Suisse,⁴ Léa Delbos, M.D.,^{5,6} Andrew Spiers, M.D.,⁴ Mathieu Poilblanc, M.D.,^{7,8} Francois Golfier, M.D., Ph.D.,^{7,8} Ludmila Jornea, Msc.,⁹ Delphine Bouteiller, M.D.,¹⁰ Hervé Fernandez, M.D., Ph.D.,¹¹ Alexandra Madar, M.D.,¹ Erick Petit, M.D.,¹² Frédérique Perotte,¹² Raffaèle Fauvet, M.D., Ph.D.,¹³ Michael Benjoar, M.D.,¹⁴ Cherif Akladios, M.D., Ph.D.,¹⁵ Vincent Lavoué, M.D., Ph.D.,¹⁶ Thomas Darnaud, M.D.,¹⁷ Benjamin Merlot, M.D.,¹⁸ Horace Roman, M.D., Ph.D.,¹⁸ Cyril Touboul, M.D., Ph.D.,^{1,2,3} and Philippe Descamps, M.D., Ph.D.,^{5,6}

Awards 2022







BFM Business prize for the therapeutic advance of the year 2022



Innovation Award 2022





Académie Nationale de **Chirurgie**

et des pratiques opératoires innovantes

ANC Innovation Award 2022

Journal of Gynecology Obstetrics and Human Reproduction 50 (2021) 102157

M.H.R. Raja, N. Farooqui, N. Zuberi et al.

CONCLUSION (3)

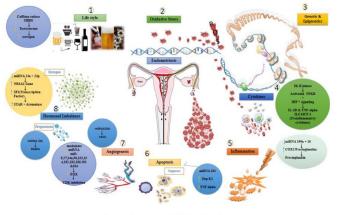


Fig. 1. Mechanism responsible for endometriosis

The mi-RNA revolution !....







Research paper

A diagnostic miRNA signature for pulmonary arterial hypertension using a consensus machine learning approach

Journal of Ambient Intelligence and Humanized Computing https://doi.org/10.1007/s12652-021-03091-2

ORIGINAL RESEARCH

MicroRNA expression classification for pediatric multiple sclerosis identification

Gabriella Casalino¹ · Giovanna Castellano¹ · Arianna Consiglio² · Nicoletta Nuzziello² · Gennaro Vessio¹



Salivary miRNAs as non-invasive biomarkers of hepatocellular carcinoma: a pilot study

Arshiya Mariam¹, Galen Miller-Atkins¹, Amika Moro², Alejandro I. Rodarte², Shirin Siddiqi², Lou-Anne Acevedo-Moreno², J. Mark Brown^{3,4}, Daniela S. Allende⁵, Federico Aucejo² and Daniel M. Rotroff^{1,6}

¹ Department of Quantitative Health Sciences, Cleveland Clinic, Cleveland, Ohio, United States

- ² Department of General Surgery, Cleveland Clinic, Cleveland, Ohio, United States
 ³ Department of Cardiovascular and Metabolic Sciences, Cleveland Clinic, Cleveland, Ohio,
- United States ⁴ Center for Microbiome and Human Health, Cleveland Clinic, Cleveland, Ohio, United States
- ⁵ Department of Pathology, Cleveland Clinic, Cleveland, Ohio, United States
- ⁶ Endocrinology and Metabolism Institute, Cleveland Clinic, Cleveland, Ohio, United States

Identifying Potential miRNA Biomarkers for Gastric Cancer Diagnosis Using Machine Learning Variable Selection Approach

Neda Gilani¹*, Reza Arabi Belaghi^{2,3}, Younes Aftabi⁴, Elnaz Faramarzi⁵, Tuba Edgünlü⁶ and Mohammad Hossein Somi⁵

The mi-RNA revolution !....

Check for updates



MicroRNA as a Potential Therapeutic Molecule in Cancer

Joanna Szczepanek ^{1,*}, Monika Skorupa ^{1,2} and Andrzej Tretyn ²

Table 4. Clinical trials of miRNA therapy in oncology (based on https://clinicaltrials.gov, accessed on 10 January 2022).

MDPI

Therapeutic Agent	Drug Name (Sponsor)	Clinical Trial Number	Phase Status	Cancer
miR-34 mimic	MRX34 (Mirna Therapeutics, Inc.)	NCT01829971	Terminated (Five immune-related serious adverse events) Withdrawn	Primary liver cancer, SCLC, lymphoma, melanoma, multiple myeloma, renal cell carcinoma, NSCLC
miR-34 mimic	MRX34 (Mirna Therapeutics, Inc.)	NCT02862145	Withdrawn (five immune-related serious adverse events in Phase 1 study)	Melanoma
miR-16 mimic	TargomiRs/MesomiR- 1 (Asbestos Diseases Research Foundation)	NCT02369198	Completed	Malignant pleural mesothelioma, non-small-cell lung cancer
anti-miR-155	Cobomarsen/MRG- 106/Vorinostat (miRagen Therapeutics, Inc.)	NCT03713320 NCT03837457	Terminated (terminated early for business reasons, not due to concerns regarding safety or lack of efficacy.) Terminated (study no longer needed because eligible subjects may receive treatment with cobomarsen in a crossover arm of the SOLAR clinical trial (NCT03713320))	Cutaneous T-cell lymphoma



What is the Next step ?







Review RNA-Targeting CRISPR–Cas Systems and Their Applications

Michal Burmistrz *⁰⁰, Kamil Krakowski and Agata Krawczyk-Balska⁰⁰

Department of Molecular Microbiology, Biological and Chemical Research Centre, Faculty of Biology, University of Warsaw, 02-089 Warsaw, Poland; k.krakowski@student.uw.edu.pl (K.K.); akra@biol.uw.edu.pl (A.K.-B.) * Correspondence: m.burmistrz@biol.uw.edu.pl

Received: 14 January 2020; Accepted: 4 February 2020; Published: 7 February 2020



Abstract: Clustered Regularly Interspaced Short Palindromic Repeats (CRISPR)–CRISPR-associated (Cas) systems have revolutionized modern molecular biology. Numerous types of these systems have been discovered to date. Many CRISPR–Cas systems have been used as a backbone for the development of potent research tools, with Cas9 being the most widespread. While most of the utilized systems are DNA-targeting, recently more and more attention is being gained by those that target RNA. Their ability to specifically recognize a given RNA sequence in an easily programmable way makes them ideal candidates for developing new research tools. In this review we summarize current knowledge on CRISPR–Cas systems which have been shown to target RNA molecules, that is type III (Csm/Cmr), type VI (Cas13), and type II (Cas9). We also present a list of available technologies based on these systems.

Keywords: CRISPR-Cas; RNA; Cas9; Cas13; Cmr; Csm

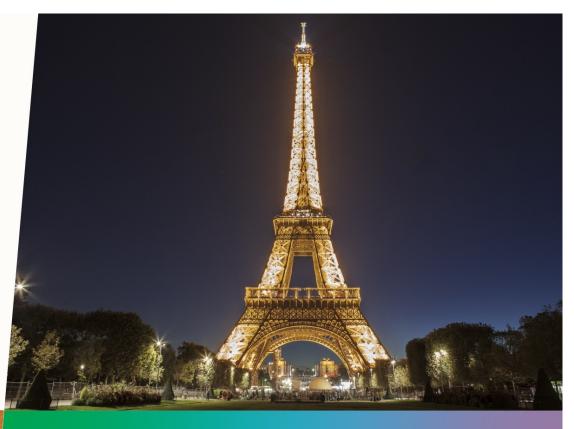
Thank you for your attention !





9-12 OCTOBER 2023 PARIS CONVENTION CENTRE







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