

The standing of self-collected urine samples for DNA methylation in cervical screening

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Why first-void urine?

- Collects cervicovaginal secretions non-invasively¹
- Well-accepted^{2,3}
- Good HPV DNA agreement between first-void urine and cervical samples⁴
- Similar clinical accuracy (CIN2+; <CIN2) of hrHPV-DNA based PCR testing in (first-void) urine compared to clinician-collected cervical samples^{5,6}







¹Vorsters A et al. (2012) Eur J Clin Microbiol Infect Dis; ²Leeman A et al. (2017) BJOG; ³Rohner E et al. (2020) J. Women's Health; ⁴Pathak N et al. (2014) BMJ; ⁵Ørnskov D et al. (2021), BMJ Open; ⁶Van Keer S et al. (2021) Gynecol Oncol



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Review of the literature

(June 1st, 2022)

- PubMed ('All Fields Contain', English, Human)
- Web of Science ('Title/Keywords/Abstracts')
- Search terms: marker* OR biomarker*; AND HPV*, human papillomavirus*, human papilloma virus*, cervical cancer(s), cervical carcinoma*, OR cancer(s) of the cervix; AND urine



Updated from Van Keer S, et al. (2017) Eur J Obstet Gynecol Reprod Bio

Results according to phase of biomarker development



Growing evidence on the clinical performance of host and viral DNA methylation in urine for cervical cancer screening

Ressarch Articlo Molecular Triage of Prema Liquid-Based Cervical Cyt Cell-Free DNA from Urine, Methylated Human Papillo Genes 12 Nafael Guerrero-Preston ¹² , Blanca L. Valle ¹ , Oluwasina Folawiyo, Francesca Pirni ¹ , Fah Maartie Noordhuis ¹⁴ , Amanda Dziedzie ² , G Carolina Guerrero-Diaz ¹ , Edgar Do Jesus R José Rodribuis ²⁰ , Amanda Dziedzie ² , S Carolina Guerrero-Diaz ¹ , Edgar Do Jesus R José Rodribuis ²⁰ , Amanda Dziedzie ² , S	Cancer Prévention Research Diggnant Lesions in ology and Circulating , Using a Panel of oma Virus and Host Anne Jedlicka ³ , Nitesh Turaga ¹ , crina Lawson ¹ , Angelo Vergura ¹ , abriela Pérez, Marisa Renehan ¹ , odriguez ² , Teresa Diaz-Montes ² , , Josefina Romaguera ³ , Bruce J. Trock ⁷ ,	CLINICAL CANCER RE HPV and DNA Me Intraepithelial Ne Rianne van den Helder ^{1,2} , Ret Ming Y. Tjiong ³ , Ivonne Marti Maaike C.G. Bleeker ² , and Nie	SEARCH PR thylation To oplasia and nske D.M. Steenber n ⁴ , Fleur M.F. Rosi enke E. van Tromn	esting in Urine for Cervical Cervical Cancer Detection rgen ² , Annina P. van Splunter ² , Constantijne H. Mom ³ , er-van Dunné ⁵ , Irene A.M. van der Avoort ⁶ , nel ¹	Check for updates
I ELSEVIER journal	PapIllomavirus Research 9 (2020) 100193 Contents lists available at ScienceDirect Papillomavirus Research I homepage: www.elsevier.com/locate/pvr		scientifi	www.nature.com/scientificrepor	ts
Methylation analysis in urine fra detection Rianne van den Helder ^{a,b} , Nienke E. var Birgit I. Lissenberg-Witte ⁶ , Maaike C.G. *Amerikan UKC, Vije Untersite Anwerden, Degemene of *Anwerden Konscheholscheholmende *Amerikan UKC, Vije Untersite Anwerden, Degemene of Fehl SCIE	actions for optimal CIN3 and cervical cancer Partonmel ^b , Annina P. van Splunter ^a , Bleeker ^a , Renske D.M. Steenbergen ^{4,4} dogs, Cancer Caner Amerikan, Amerikan, he Noherlands forocologi concludge, Caned Ogneodig Concludy Amerikan, Amerikan, die Noherlands Manage and Busantins, Amerikan, he Noherlands CONTREFICTOR REE	www.nature.com/scientificreports	OPEN	Constraints Constrain	ates Des for claim
OPEN Received: 4 October 2018	Cervical cancer detec methylation analysis Barbara C. Snoek ¹ , Annina P. van Splunter ¹ , Maaike C. Daniëlle A. M. Heideman ¹ , W. Frederik Rurup ² , Wina V Mignon van Gent ⁴ , Nienke E. van Trommel ⁴ & Renske I	Stion by DNA S in urine G. Bleeker ¹ , Maartje C. van Ruiten ¹ , erlaat ¹ , Hans Schotman ³ , D. M. Steenbergen ¹	Evalu ZAR1 Rene H Hernán	aation of DNA methylation in promoter regions of SF in urine and plasma of women with cervical lesions offstetter ^{1,2,3*} , Ismael Riquelme ^{1,2*} , Alejandra Andana ^{1,2} , Carmen G. Ili ^{1,2} , Kurt Vargas ⁴ , Priscilla Brebi ^{1,2} , Juan Carlos Roa ⁵	RP4 and

Clinical performance host and viral gene methylation from literature

Methylation markers	Study population	Sample	AUC	Endpoint	Ref
ZNF516; FKBP6; INTS1; and HPV16-L1 (4-gene biomarker panel)	n=40 hrHPV+	ccfDNA 10ml urine (flash frozen)	0.86	CIN2+ vs. ≤CIN1	Guerrero-Preston R et al. (2016)
SFRP4 and ZAR1 (2-gene biomarker panel)	nSCC=31; nNILM=60 hrHPV	pellet ≥10ml first-void urine (crystal violet)	0.63	SCC vs. NILM	Hoffstetter R et al. (2017)
FAM19A4; GHSR; PHACTR3; PRDM14; SST; and ZIC1 (individual genes)	ncancer=38; ncontrol=47 82% hrHPV+	pellet 15ml urine	0.74-0.89	Cancers vs. healthy controls	Snoek BC et al. (2019)
ASCL1; GHSR; LHX8; SST; and ZIC1 (individual genes)	nCIN3+=47; ncontrol=27 hrHPV	pellet 15ml urine (EDTA)	0.87-0.97 0.62-0.79	Cancers vs. healthy controls CIN3 vs. healthy controls	van den Helder R et al. (2020)
ASCL1; ST6GALNAC5; GHSR; LHX8; SST; and ZIC1 (individual genes)	nCIN2+=19; n≤CIN1=14 hrHPV+	4ml first-void urine (UCM)	0.62-0.80	CIN2+ vs. ≤CIN1	Van Keer S et al. (2021)
ASCL1; GHSR; LHX8; SST; and ZIC1 (individual genes)* ASCL1/LHX8 (bi-marker panel)**	nCIN3+=181; ncontrol=64 87% hrHPV+	pellet 15ml urine (EDTA)	0.72-0.83* 0.84**	CIN3+ vs. healthy controls	van den Helder R et al. (2022)

All studies applied methylation specific qPCR on bisulphite converted DNA.

CIN: cervical intraepithelial neoplasia; SCC: squamous cell carcinoma; NILM: negative for intraepithelial lesion or malignancy.



Developing a complete cervical cancer screening solution based on first-void urine self-sampling – CASUS trial



Figure: Van Keer S et al. (2021) HPV World











Developing a complete cervical cancer screening solution based on first-void urine self-sampling – CASUS trial



WP 3 & 4 – Validate the first-void urine-based complete cervical cancer solution for detection of cervical (pre)cancer



Study design CASUS WP 3 & 4





Results CASUS WP 3 & 4

Unpublished data – cannot be distributed or published yet





A complete cervical cancer screening solution based on firstvoid urine self-sampling ?

- Similar clinical accuracy of hrHPV-DNA based PCR testing in urine compared to clinician-collected cervical samples
- Host cell methylation marker levels in first-void urine increase with increasing severity of disease
- Methylation of host cell genes discerns <CIN2 from CIN3+ and Cancer in first-void urine
- Collected at home



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