

# Exploring DNA methylation biomarkers associated with bladder cancer in patients with neurogenic bladder

<b>Submission date</b> 11/06/2023	<b>Recruitment status</b> No longer recruiting	<input type="checkbox"/> Prospectively registered <input type="checkbox"/> Protocol
<b>Registration date</b> 14/06/2023	<b>Overall study status</b> Completed	<input type="checkbox"/> Statistical analysis plan <input checked="" type="checkbox"/> Results
<b>Last Edited</b> 30/07/2025	<b>Condition category</b> Cancer	<input type="checkbox"/> Individual participant data

## Plain English summary of protocol

### Background and study aims

The term neurogenic lower urinary tract dysfunction (NLUTD, also known as neurogenic bladder) refers to abnormal function of the bladder and urinary system. Previous studies have shown that NLUTD patients might have an increased risk of bladder cancer (BCa), which is diagnosed at a younger age and at a more advanced stage in comparison to the general population. Also, there is a higher risk of dying from BCa among NLUTD patients compared to the general population. For these reasons, early diagnosis and definitive treatment of BCa are more crucial in this patient group. However, there is currently scarce research on the most appropriate screening method and follow-up for these patients. Risk factors and possible biomarkers for BCa in patients with NLUTD have to be identified, with future research carried out to provide guidance for screening.

BCa shows frequent alterations in DNA methylation. Nowadays, with the application of modern methods DNA methylation can be recognized in body fluids with high sensitivity and specificity. Several studies support the use of DNA methylation in urine for the early detection of BCa. This study will evaluate the presence of DNA methylation of a panel of five BCa biomarkers in the urine of patients with NLUTD in comparison with healthy controls, and their association with the patients' clinical characteristics/demographics as well as with histopathological findings and DNA methylation of the panel of genes from bladder tissue samples from the same patients.

### Who can participate?

1. Adults aged 18 years or older with NLUTD (or neurogenic bladder) for at least 5 years who have been scheduled for cystoscopy with cold cup biopsy and wash cytology for bladder cancer screening
2. Adults aged 18 years or older who do not have urinary symptoms or disease

### What does the study involve?

Patients with NLUTD provide urine and tissue samples for DNA methylation analysis. Volunteers without lower urinary tract symptoms provide a urine sample for DNA methylation analysis.

What are the possible benefits and risks of participating?

Participants are contributing to an effort to investigate new early non-invasive methods of diagnosing bladder cancer in NLUTD patients. There is no risk associated with participating in this study.

Where is the study run from?

Papageorgiou Hospital of Thessaloniki (Greece)

When is the study starting and how long is it expected to run for?

August 2019 to December 2023

Who is funding the study?

Investigator initiated and funded

Who is the main contact?

Dr Periklis Koukourikis, perikliskoukourikis@gmail.com

## Contact information

**Type(s)**

Scientific

**Contact name**

Dr Periklis Koukourikis

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## Additional identifiers

**EudraCT/CTIS number**

Nil known

**IRAS number**

**ClinicalTrials.gov number**

Nil known

**Secondary identifying numbers**

PhD/28/08/2019

## Study information

**Scientific Title**

A study of DNA methylation of bladder cancer markers in urine and bladder tissue of patients with neurogenic lower urinary tract dysfunction

**Study objectives**

The aim of the study is to evaluate the presence of DNA methylation of a panel of five gene promoters, previously associated with bladder cancer, in the urine and bladder tissue of patients with neurogenic lower urinary tract dysfunction and in comparison with controls, their association with the patients' clinical characteristics/demographics and their potential as urine biomarkers for bladder cancer screening and diagnosis.

**Ethics approval required**

Ethics approval required

**Ethics approval(s)**

Approved 29/07/2020, Aristotle University of Thessaloniki, Greece, Faculty of Health Sciences, School of Medicine, Ethics Committee (Administration Building (opposite AHEPA Hospital), University Campus, Thessaloniki, 54124, Greece; +30 (0)2310 999338; sakkageor@auth.gr), ref: 6665

**Study design**

Single-center case-control study

**Primary study design**

Observational

**Secondary study design**

Case-control study

**Study setting(s)**

Hospital, Laboratory

**Study type(s)**

Diagnostic, Screening

**Participant information sheet**

Not available in web format, please use contact details to request a participant information sheet

**Health condition(s) or problem(s) studied**

Neurogenic lower urinary tract dysfunction (neurogenic bladder), screening for bladder cancer

**Interventions**

This study will recruit two groups of participants.

1. NLUTD Group: patients suffering from neurogenic lower urinary tract dysfunction (NLUTD) for at least 5 years, who have been scheduled for cystoscopy with cold cup biopsy and wash cytology for bladder cancer screening. Patients will provide a urine sample via spontaneous voiding, clean intermittent catheterization or indwelling catheterization, depending on their voiding habits, and part of their bladder biopsy for DNA methylation analysis.

2. Control Group: volunteers without lower urinary tract symptoms. Patients' attendants and hospital staff members will be asked to participate in the study as controls. Volunteers will provide a free-voiding midstream urine sample for DNA methylation analysis and will be assessed with a transabdominal ultrasonography of the urinary system.

Demographics from all participants will be recorded.

The following additional clinical data will be recorded from the NLUTD group:

1. Neurological disease and duration
2. Duration of NLUTD
3. Method of bladder voiding (spontaneously, clean intermittent catheterization [CIC], indwelling catheter [IDC])
4. Years of IDC or CIC usage
5. History of bladder stone(s)
6. History of hematuria
7. History of recurrent UTIs
8. Histopathology results
9. Urine wash cytology results
10. Cystoscopy results

In all urine samples the content of the Urine Preservative Single Dose (Cat#18124, NORGEN BIOTEK CORP., Canada) will be added, which stabilizes DNA, RNA and proteins at room temperature for over 2 years without the need for immediate freezing of the samples. The tissue samples from the NLUTD group that will be subjected to molecular processing will be placed in microvials and immediately cooled in liquid nitrogen, which will then be stored in a deep freeze (-80 °C) until their processing. The samples will be processed, in a blinded fashion, at the Laboratory of Biological Chemistry of the School of Medicine of the Aristotle University of Thessaloniki (Au.Th.). DNA will be extracted from urine and bladder tissue samples using the Cells and Tissue DNA Isolation Kit (Norgen Biotek Corp) and will be modified with sodium bisulfite employing the EZ DNA Methylation-GoldKit (Zymo Research). Quantitative Methylation Specific PCR (qMSP) will be performed to detect bisulfite-induced changes at the promoters of the investigated genes. qMSP will be performed with Luna Universal Probe qPCR Master Mix (New England Biolabs, USA), using the Applied Biosystems StepOnePlus Real Time PCR System (Thermo Fisher Scientific, Inc.).

## **Intervention Type**

Other

## **Primary outcome measure**

DNA methylation of a panel of five genes' promoters (Ras-Association domain Family member 1 [RASSF1], Retinoid Acid Receptor beta [RARβ], Death-Associated Protein Kinase [DAPK], Telomerase Reverse Transcriptase [TERT] and Adenomatous Polyposis Coli [APC]) in urine measured using Quantitative Methylation Specific PCR at a single timepoint

## **Secondary outcome measures**

DNA hypermethylation of the panel of five genes' promoters (RASSF1, RARβ, DAPK, TERT and APC) in the bladder tissue of patients suffering from neurogenic lower urinary tract dysfunction (NLUTD) measured using Quantitative Methylation Specific PCR at a single timepoint

## **Overall study start date**

28/08/2019

**Completion date**

31/12/2023

## Eligibility

**Key inclusion criteria**

Neurogenic lower urinary tract dysfunction group:

1. Neurogenic lower urinary tract dysfunction duration for at least 5 years scheduled for cystoscopy, wash cytology and bladder biopsy
2. Age 18 years or older
3. Consent to provide urine sample and bladder tissue

Control group (healthy volunteers):

1. Age 18 years or older
2. Consent to provide urine sample

**Participant type(s)**

Healthy volunteer, Patient

**Age group**

Adult

**Lower age limit**

18 Years

**Upper age limit**

90 Years

**Sex**

Both

**Target number of participants**

35 participants in each group

**Total final enrolment**

76

**Key exclusion criteria**

Neurogenic lower urinary tract dysfunction group:

1. Prior diagnosis and treatment of bladder cancer
2. History of other urinary tract malignancy

Control group (healthy volunteers):

1. Lower urinary tract symptoms
2. History of urinary tract malignancy

**Date of first enrolment**

01/08/2020

**Date of final enrolment**

31/12/2023

## **Locations**

### **Countries of recruitment**

Greece

### **Study participating centre**

2nd Department of Urology Aristotle University of Thessaloniki, Papageorgiou University

Hospital of Thessaloniki

Ring Road

Nea Efkarpia

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Greece

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## **Sponsor information**

### **Organisation**

Aristotle University of Thessaloniki

### **Sponsor details**

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### **Sponsor type**

University/education

### **Website**

<https://www.med.auth.gr/>

### **ROR**

<https://ror.org/02j61yw88>

## **Funder(s)**

### **Funder type**

Other

**Funder Name**  
Investigator initiated and funded

## Results and Publications

**Publication and dissemination plan**  
Planned publication in a high-impact peer reviewed journal.

**Intention to publish date**  
01/07/2023

**Individual participant data (IPD) sharing plan**  
The datasets generated during and/or analysed during the current study are/will be available upon request from Dr Periklis Koukourikis MD, FEBU (perikliskoukourikis@gmail.com). Raw data in Excel sheets of demographics, clinical characteristics and results of methylation analysis would be available upon request after the completion of the study. All the participants of the study receive a number during recruitment and participation for anonymization reasons. Signed consent for data availability in an anonymous way was received from the participants.

**IPD sharing plan summary**  
Available on request

Study outputs					
Output type	Details	Date created	Date added	Peer reviewed?	Patient-facing?
<a href="#">Results article</a>	DNA hypermethylation results	23/05/2024	30/07/2025	Yes	No
<a href="#">Results article</a>	Results	12/08/2023	30/07/2025	Yes	No