A comprehensive, genetic, laboratory developed test for identifying ALL stages of bladder cancer

GALEAS / BLADDER DETECT | REPORT | MONITOR

Bladder cancer causes >200K deaths globally each year

	UK	Europe	APAC	USA
Hematuria cases*	1,000,000	10,000,000	20,000,000	9,000,000
Cystoscopies*	110,000	2,030,000	2,080,000	900,000
Bladder Cancer cases	11,000	203,000	208,000	90,000

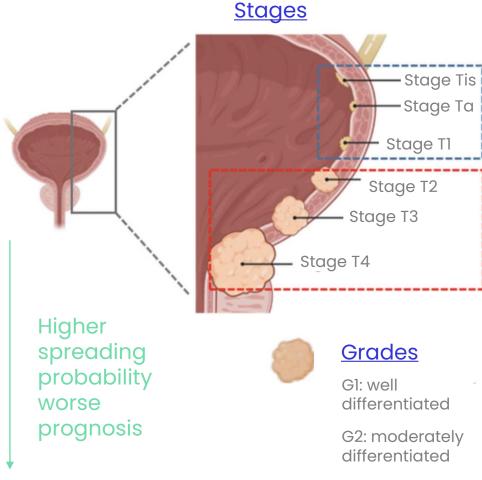
- **535,000** People living with bladder cancer in the USA (compared to 371,000 lung cancer)
 - 70% Of bladder cancer cases are lower grade **non-muscle invasive disease** (NMIBC) and require active surveillance for 5 year to life. 70% will recur within two years.
 - **30%** Of cases are **muscle invasive disease** (MIBC) of which 50% progress/ die within 5 years.

BUT early detection can significantly impact survival

Early diagnosis results in >80% survival at 5 years compared to <10% <u>late</u> diagnosis.

Stages of Bladder cancer

Two "types" of bladder cancer (BC)



G3: poorly differentiated

NMIBC

Tis: non-invasive

Ta: non-invasive

T1: tumor invades inner lining and connective tissue

<u>MIBC</u>

T2: tumor invades muscle

T3: tumor invades perivesical fat and lymph nodes

T4: tumor spreads to lymph nodes and other organs

Non Muscle Invasive (NMIBC)

- Accounts for 75% of BC
- High recurrence rate (60%-70%)
- Low progression rate (20%-30%)
- High 5-year survival (96%)

Treatment : BCG/Mytomicin C

Muscle Invasive (MIBC)

- Accounts for 25% of BC
- Low 5-year survival (45%)
- 50% progression post curative surgery

Treatment : Surgery, Radiotherapy, Chemotherapy (cisplatin), Immunotherapy

Blood in the urine is best clinical indicator of bladder cancer

For a very long time, hematuria (or blood in the urine) has been known to be the most common symptom of bladder cancer:





But hematuria can be due to many things, not just bladder cancer

For example:

- Bladder infection
- A Stones in the kidneys or bladder
- Inflammation of the kidneys (nephritis)
- Urinary tract injuries
- Blood disorders (e.g. sickle cell disease, clotting disorders, anticoagulant and anti-platelet drugs)
- Other causes, including less common infections (e,g. TB, schistosomiasis)

In fact **only 10%** of all patients presenting with hematuria will actually have bladder cancer .



How do healthcare providers currently identify bladder cancer?

Firstly, a simple dipstick test is most commonly used to test for two types of hematuria:

- Visible (macro haematuria)
- Non-visible (micro haematuria)

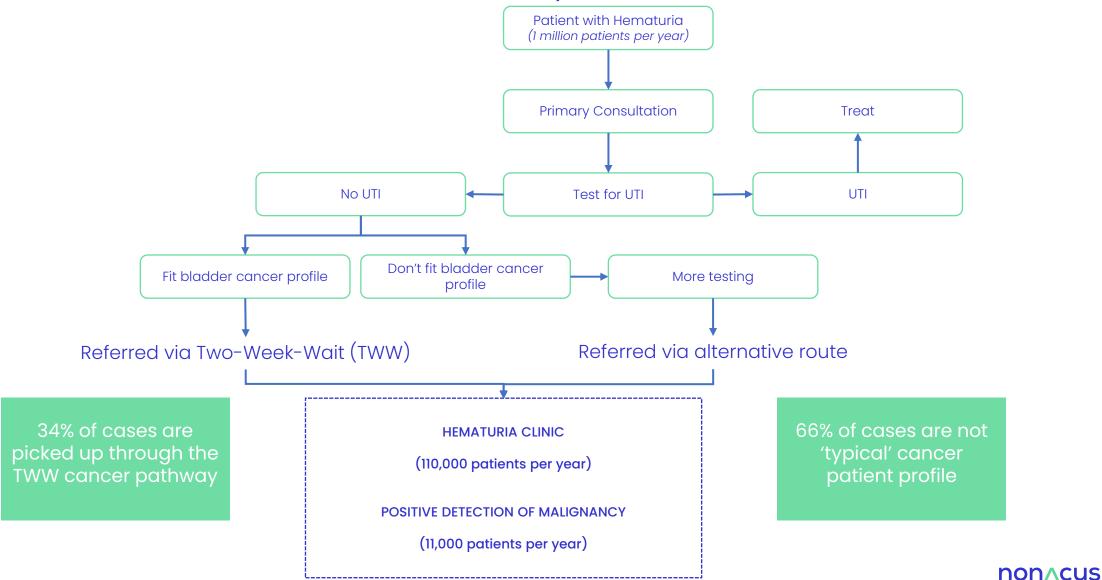
Secondly, patients are tested for a urinary tract infection (UTI).

Thirdly, patients who test negative for UTI are referred by doctors to a hematuria clinic to check to see if they have cancer.

The current standard of care to identify cancerous cells is a visual inspection in the bladder – an often painful, invasive, hospital-based procedure called a cystoscopy.

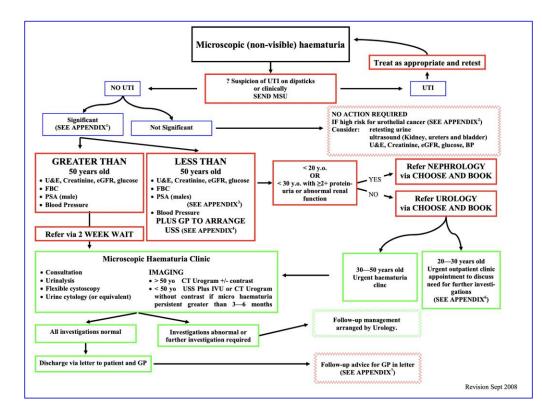


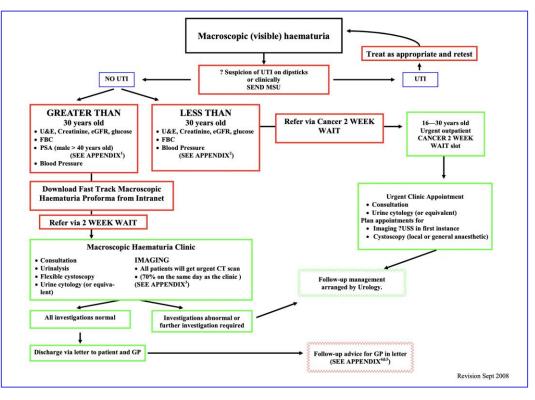




*Simplified representation. For actual pathway see next slide.

Current UK NHS Hematuria Care Pathway





Why is the current standard of care failing patients?

I have had bladder cancer 4 times. The cystoscopy is so invasive, uncomfortable and embarrassing I actually dread it for weeks before.

> NHS patient, commenting on a CR-UK blog post regarding the Nonacus Bladder cancer test

- It is at best uncomfortable, at worst painful and invasive
- It is inconvenient as it requires a hospital visit
- It costs the healthcare provider time and money (£88M* per year in the UK)
- Is not without complications
- Sensitivity and specificity is operator dependant.

AND

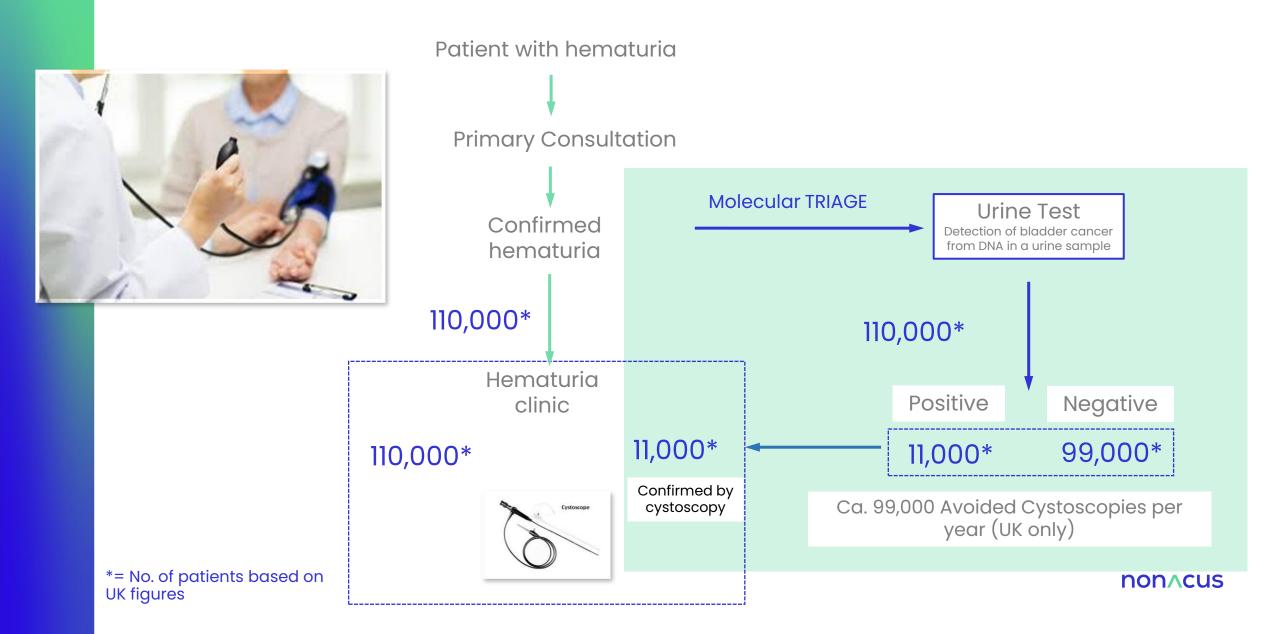
• 90% of patients referred don't actually need it

* Cost of cystoscopy only (NHS Tariff £330), plus staff time and ancillary costs for £470. Does not represent the full pathway.

How can Doctors assess which patients referred with blood in their urine have bladder cancer and need surgery ...

...without sending thousands of patients for painful invasive cystoscopy tests?

Molecular Triage - Reducing the Burden



Introducing GALEAS[™] Bladder

A comprehensive, laboratory developed, genetic test for identifying all stages of bladder cancer from a urine sample

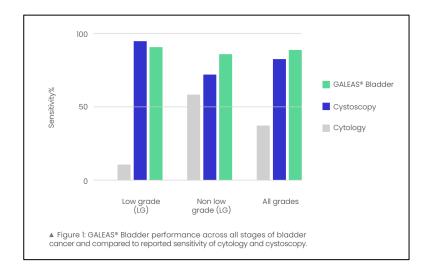
- Allows patients to provide a sample in the comfort of their own home. No hospital visits, no painful procedure.
- Offers equivalent sensitivity and specificity to cystoscopy.
- A Reduces the number of unnecessary cystoscopies reducing costs and resource burden on clinics.
- Leverages targeted next generation sequencing chemistry to accurately detect somatic mutations from 23 genes in over 96% of bladder cancer cases.

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GALEAS™ Bladder: Performance Data

- Validated in ~770 urine samples from 3 UK clinical cohorts
 - 382 positive bladder cancer cases
 - A 388 negative i.e. no cancer at the time of cystoscopy

	Sensitivity	Specificity	PPV	NPV
рТа	86%	86%	75%	93%
τ1	95%	86%	66%	99%
T2+	89%	86%	60%	97%
Gl	76%	86%	47%	96%
G2	92%	86%	67%	97%
G3	92%	86%	78%	95%
NMIBC	89%	86%	83%	92%
MIBC	89%	86%	60%	97%



• Equivalent sensitivity and specificity to cystoscopy across all grades and stages.

• Reducing cystoscopies by ~90%

References

Svatek RS, Hollenbeck BK, Holmäng S, Lee R, Kim SP, Stenzl A, Lotan Y. The economics of bladder cancer: costs and considerations of caring for this disease. Eur Urol. 2014 Aug;66(2):253-62. doi: 10.1016/j.eururo.2014.01.006. Epub 2014 Jan 21. PMID: 24472711. Zheng C, Lv Y, Zhong Q, Wang R, Jiang Q. Narrow band imaging diagnosis of bladder cancer: systematic review and meta-analysis. BJU Int. 2012 Dec;110(11 Pt B):E680-7. doi: 10.1111/j.1464-410X.2012.11500.x. Epub 2012 Sep 18. PMID: 22985502.

GALEAS[™] Bladder: from urine sample to test results

The test consists of four components

		ğ	
Urine Collection Device	gDNA Extraction kit	DNA Library Prep and Capture kit	Patient Report
Product Name: Urine GALEAS™ Urine Collection Device Product code: PRE_GAL_UCD	Product Name: GALEAS™ Bead Xtract: Urine gDNA 96 sample Product code: PRE_GAL_BXG_96	Product Name: GALEAS [™] Bladder Kit Product Code: NGS_GAL_BCP_FR_96_(A,B,C,D format)	Product Name: GALEAS™ Bladder Analysis Product code: NGS_GAL_GBA
Description: Urine collection in preserver tubes for safe delivery to lab	Description: Magnetic, bead- based automatable extraction of gDNA from urine cell pellet	Description: Library preparation and hybridisation and capture kits for targeting SNVs associated with bladder cancer	Description: Analysis software for generating report based on variant profile
 Steps: Doctor/Clinician complete a test requisition form. Doctor/Service lab provide patient with collection device. Patient uses device and posts preserver tubes to service lab Service lab scans tube into LIMS system and sends sample for gDNA extraction 	 Steps: Service lab spins urine sample and collects cell pellet Service lab follows protocol to extract gDNA NB. DNA concentration and yield must be of suitable quality to progress to next step. 	 Steps: Service lab prepares libraries using extracted gDNA Service lab pools libraries and capture the targeted regions for sequencing DNA library is sequenced using Illumina sequencers 	 Steps: Service lab analyses sequencing data through GALEAS bioinformatic pipeline Positive or Negative patient report is generated (<i>either PDF</i> or JSON format) Report submitted to Urologist Urologist organises patient consultation.

GALEAS[™] Bladder Urine Collection Device

- ^ An intuitive collection device
 - Flat packed ensures collection device is fit for purpose and not damaged when received by user
 - ^ Less storage and shipping space required reducing postage costs
 - Easy assembly and intuitive use ensures good sample taking process and return rates
- 50ml Falcon tube
- Order with test
 - Dispatch to patient is responsibility of service/LDT laboratory
- A LIMS tracking Unique ID barcode on tube provides options:
 - 1. Use unique tube ID barcode and associate with patient or
 - 2. Add specific patient ID barcode for sample tracking



GALEAS[™] Bladder: Urine Bead Xtract gDNA

gDNA is extracted from urinary cell-pellets

- Simple, magnetic bead-based protocol
- Optimised for >20ng of cell-pellet genomic DNA
- Quick and easy workflow
- Supports manual or automated preparation of 1–96 samples in a single batch



GALEAS™ Bladder Kit

Library Prep and Hybridization and Capture kits

- A GALEAS™ branded kits
 -specific GALEAS™ Bladder only control
- 96 sample format only
- Optimised for 20ng of cell-pellet genomic DNA
- Quick and easy workflow automation protocol available
- Preparation of 1-96 samples in a single batch
- Tracking SNPs and control regions included



Comprehensive NGS panel

Biomarkers were identified by Dr Rik Bryan and Dr Doug Ward at Birmingham University UK

- NGS sequencing panel that targets promoter and exonic regions of 23 of the most relevant genes associated with bladder cancer.
- Identified by a combination of publicly available data and deep exome sequencing

-Exome studies were performed on Caucasian populations

- The panel have been shown to accurately detect somatic mutations in over 96% of bladder cancers in over 770 clinical samples.
- Compatible with all Illumina sequencers
- 384 patient/sample indexes ensure that customer can use the GALEAS® Bladder Cancer panel on the smallest to the largest output sequencers.

AKT1	ERBB2	NRAS
BRAF	ERBB3	PIK3CA
C3orf70	ERRC2	RHOB
CDKNIA	FBXW7	RXRA
CDKN2A	FGFR3	SF3B1
CREBBP	HRAS	TERT (promoter)
CTNNB1	KDM6A	TP53
ELF3	KRAS	

GALEAS™ Software – Cloud based, end to end solution for bioinformatics

- Bulk upload of sequencing data (FASTQ files) via simple application; no minimum sample number
- Automated 'sample to report' analysis pipeline, requiring no specialist training to interpret data
- Simple report 'Yes' or 'No' to likely presence of Bladder cancer
- Detailed report of somatic variants identified
- Report available as .pdf or .JSON and sent directly to service lab
- Download batches of results
- Reports on failed samples

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GALEAS Bladder

Patient ID: Patient281123-2-POS Patient Name: Anon Anon Date of Birth:	Sample ID: N0131 Received: 20/12/2022 Processd: 30/11/2023 Report Date: 04/12/2023 14:58 Sample Type: Unine Pellet	Clinician: Dr Bob Loblaw Address: Coombs Ford Slowmarket Ipswich IP145QR UK	Customer: Nonacus Address: Not provided
Result Summary: P	ositive		
Apositive test result indicates that, at the present.	e time of GALEAS Bladder testing, cancer as so	ciated variants were detected and there	is a high likelihood that cancer is
Appropriate clinical follow up is require	d to confirm a clinical diagnosis.		
Variant Details			
Gene	HGVS	ic .	VAF 2.5%
Test Description	I NA	I	2.379
GALEAS Bladder data, generated on over and sensitivity of 89% for the detection of	770 patient urine samples, has determined a to all stages of bladder cancer (ref 1.2.3).	est positive predictive value (PPV) of 87%	i, negative predictive value (NPV) of 96%

The variants in this test have been validated as part of the GALEAS Bladder Triage Haematuria test only. They have not been validated as predictive markers for disease stratification or for the informing treatment decisions.

Positive Explanation

GALEAS Bladder tests for somatic variants in selected regions from across 23 genes. The presence of somatic variants in these regions in urinary DNAhas been shown to associate with the presence of bladder cancer. The detection of one of more of somatic variants indicates a high likelihood that cancer is present.

Negative Explanation

AGALEAS Bladder negative test result, at the time of testing, is determined by the lack of detection of cancer associated genomic variants in the urine sample, suggesting the presence of bladder cancer in unlikely. However, this does not completely exclude the presence of cancer now or in the future.

QC Status

PASS

QC Status Explanation

There was sufficient read depth across the regions to confidently determine a result

Test Limitations

The has not been validated as predictive biomarker for disease stratification or for informing treatment decisions.

Methodology

DNAwas extracted from urine derived cell pellets and collected using the GALEAS Bladder Home Collection Kit. Extracted genomic DNAsubsequently underwent target enrichment using the GALEAS Bladder Target Enrichment protocol with sequencing performed using Illumina sequencing by synthesis chemistry.

Data analysis was performed using the GALEAS Bladder analysis pipeline GALEAS Bladder version 23.12.1

References

Nonacus Limited. Quinton Business Park, Unit 5, Ridgeway, Quinton, Birmingham B32 1AP, United Kingdom, Registered No: 9590278

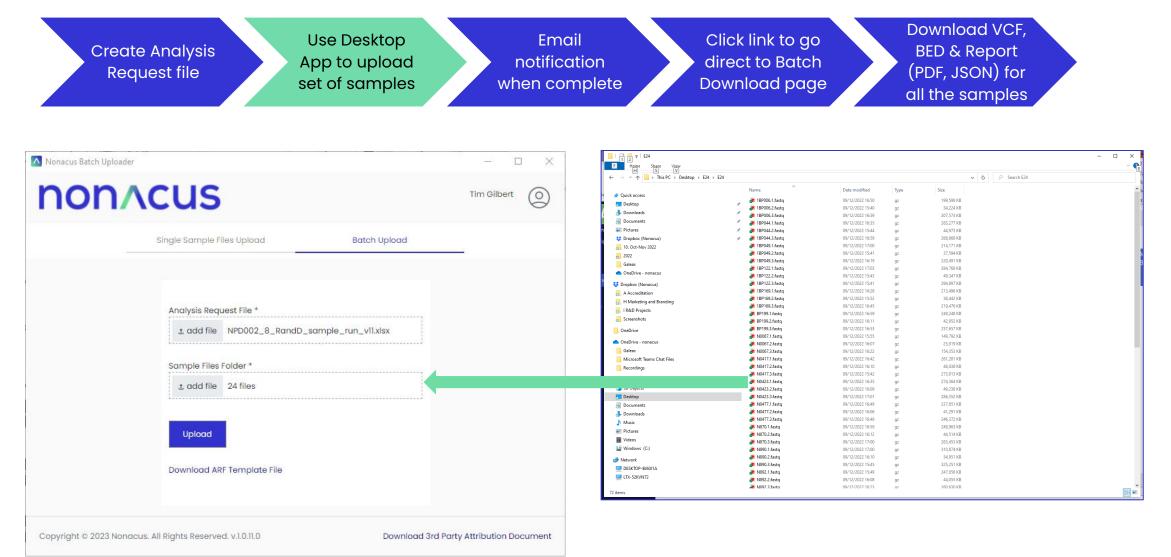


Create Analysis Request file Use Desktop App to upload set of samples

Email notification when complete

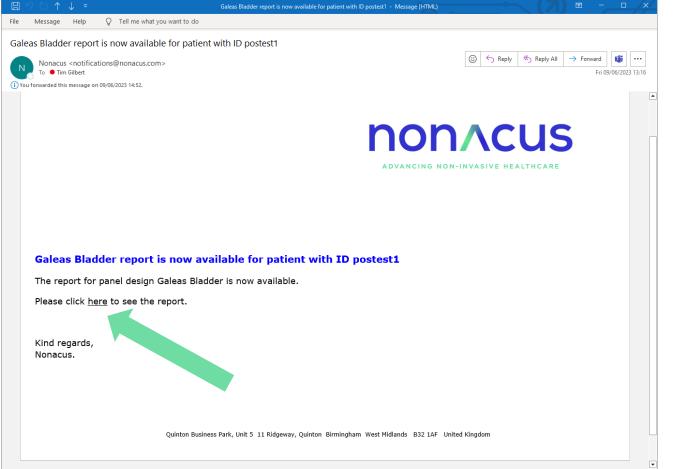
Click link to go direct to Batch Download page Download VCF, BED & Report (PDF, JSON) for all the samples

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3	BP044	Urine Pellet	969	Test BP044	01/12/2022	Test BP044				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
4	BP049	Urine Pellet	969	Test BP049	01/12/2022	Test BP049				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
5	BP122	Urine Pellet	969	Test BP122	01/12/2022	Test BP122				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
6	BP169 BP199	Urine Pellet Urine Pellet	969 969	Test BP169	01/12/2022 01/12/2022	Test BP169 Test BP199				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR IP145QR	07/12/2022 07/12/2022			
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2	s0581	Urine Pellet	969	Test s0581	01/12/2022	Test s0581				NPD002	Anon			Bladder Cancer Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
	s0601	Urine Pellet	969	Test s0601	01/12/2022	Test s0601				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
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;	s0873	Urine Pellet	969	Test S0873	01/12/2022	Test S0873				NPD002	Anon			Bladder Cancer		Dr Bob	Loblaw	Coombs Ford	Stowmarket	lpswich	UK	IP145QR	07/12/2022			
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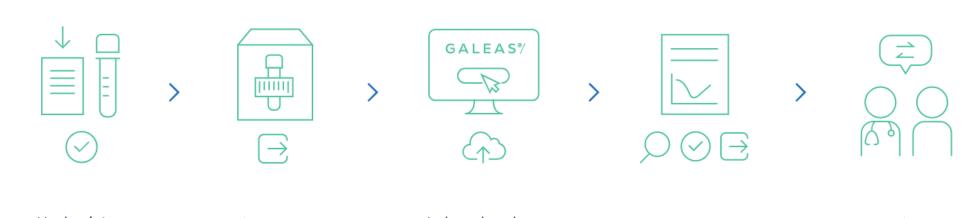
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Workflow



Urologist requests urine collection device for patient Patient sends barcoded urine sample to service laboratory Lab uploads sequencing results to secure cloudbased GALEAS® platform Results analysed, report generated and shared directly with urologist Urologist shares results with patient and discusses next steps

Support/Controls

LDT validation kit

- A Range of positive and negative 'non-cancer' DNA controls
- Full workflow validation coming later

Probe for control regions

- Targeting variants with high VAF
- Included in panel design
- 24 tracking SNPs included in panel design



Advantages of GALEAS[™] Bladder

NOW

- NGS assay with high analytical sensitivity equivalent to cystoscopy across all stages of bladder cancer (Other genomics tests are able to detect low grade Bladder cancer but not high grade).
- Inlike other genomics tests, NGS panels with a wider range of markers overcome tumour heterogeneity so you are less likely to miss a cancer.
- Automated and scalable workflow which can be **run in any NGS capable laboratory**
- **Sample to answer solution** including bioinformatics and reporting
- Part of a suite of products for oncology and liquid biopsy

And in the FUTURE, the broad range of mutations detected by GALEAS™ Bladder will allow:

Future advantages of GALEAS™ Bladder

Disease stratification

Ne are building the clinical evidence to demonstrate that GALEAS™ Bladder can stratify low- and high-grade bladder cancer which will be key in improving patient outcomes

Companion diagnostics

Many Pharma companies are using genetic signatures to target therapies E.g. FGFR3 inhibitors

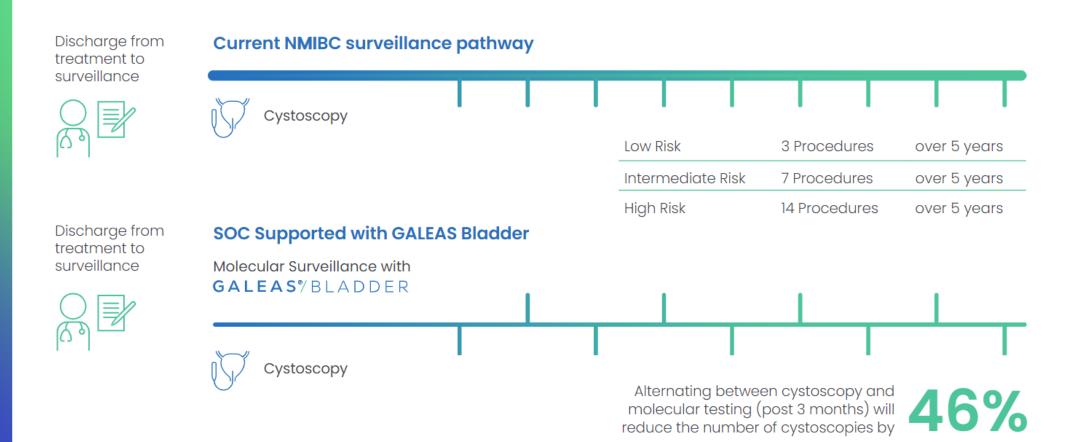
Whole care pathway

▲ Bladder cancer has one of the highest recurrence rates of any cancer. GALEAS[™] Bladder has the potential to be used as an alternative to cystoscopies in monitoring patients for recurrence without changing workflows (see next slide)



NMIBC Surveillance

GALEAS™ Bladder could be used to reduce the number of cystoscopies in the surveillance pathway by 46%

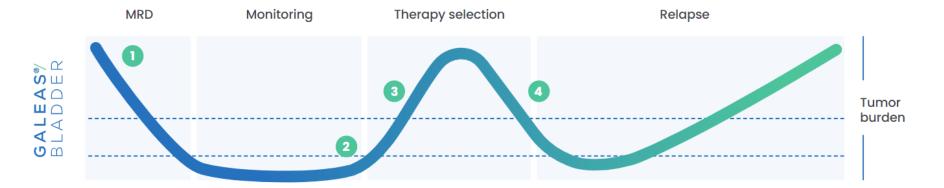


Circulating tumor DNA (ctDNA) guided monitoring for **MIBC**

The unique design of GALEAS[®] Bladder allows the test to be used as a blood-based test, post surgical resection to detect circulating tumor DNA and monitor treatment response and resistance in MIBC.

- De-escalation of adjuvant therapy post surgical resection in ctDNA negative patients^(3.4.).
- 2 Earlier identification of disease recurrence or progression.

- 3 Selection of patients for immunotherapy^(5.).
- 4 Monitoring of treatment response or resistance.



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Abstract



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Keywords: Bladder cancer Urine DNA Sequencing Mutation Biomarker Diagnosis

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Background: There is an unmet need for an accurate, validated, noninvasive test for diagnosing and monitoring bladder cancer (BC). Detection of BC-associated mutations in urinary DNA via targeted deep sequencing could meet this need. Objective: To test the ability of mutational analysis of urinary DNA to noninvasively

Translational Science

Targeted deep sequencing of urothelial bladder cancers and associated urinary DNA: a 23-gene panel with utility for non-invasive diagnosis and risk stratification

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Objectives

To develop a focused panel of somatic mutations (SMs) present in the majority of prothelial bladder cancers (UBCs).

EXERC. Second VIRGE, BACK, C. 30770, CARDEN, CDKN24, and NRA5, 93–98.3% of UECs of all grades and stages harboured ≥ 1 SM (mean 2.5 SM/stumour), RAS mutations were associated with better overall anrival (P = 0.04). Mutations in RXRA, RHOB and TERT present in the majority of unotheal bladder cancers (URAs) to investigate the diagnostic and prognostic utility of this panel, and to compare the identification of SMs in uniary cell-pellet (cpiDNA and cell-free (cfDNA as part of the development of a non-invasive clinical assay. (promoter) were associated with shorter time to recurrence (P < 0.05) MAFs in urinary cfDNA and cpDNA Patients and Methods were highly correlated; using a capture-based approact

A panel of SMs was validated by targeted deep-sequencing of >94% of tumour SMs were detected in both cpDNA and cfDNA. A pinet of SMS was visuated by targeted deep-sequencing of tumour DNA from 956 patients with UBC. In addition, amplicon and capture-based targeted sequencing measured mutant allele frequencies (MAFs) of SMS in 314 urine opDNAs and 155 urine dDNAs. The association of SMs with gade, stage, and clinical outcomes was investigated by Conclusions SMs are reliably detected in urinary cpDNA and cfDNA. The technical capability to identify very low MAFs is essential to minibly detect UBC, regardless of the use of cpDNA or cfDNA. This 23-gene panel shows promise for the non-invasive diagnosis and risk stratification of UBC.

univariate and multivariate Cox models. Concordance between SMs detected in tumour tissue and cpDNA and cfDNA was assessed. Results

The panel comprised SMs in 23 genes: TERT (promoter), PGFR3, PIK3CA, TP53, ERCC2, RHOB, EREB2, HRAS,

Introduction

BJU Ini 2019; 124: 532-544

performance and/or poor evidence [1-3]. Many tests are based on levels of proteins or RNA and, as these are not unique to UBC or causally linked to the disease, they tend to Introduction Despite intensive research into biomarkers for the non-invarve daposis of urothein's bladdr current (URC). Che ministry of districts means fields beyoncopy. Commercial urine tests exist however, nose have been widely or low-grade tumours [4]. The ideal nee-invarive test should

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Keywords

nonAcus

RXRA, ELF3, CDKN1A, KRAS, KDM6A, AKT1, FEXW7

EREE3. SF3E1. CTNNE1. BRAF. C3orf70. CREEEP.

mutations, diagnosis, prognosis, detection, urine, DNA, #Bladder cancer, #blcsm

accepted into routine clinical practice due to poor



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