

Personalized medicine

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SAMEN
GRENZEN
VERLEGGEN

Conflict of interest

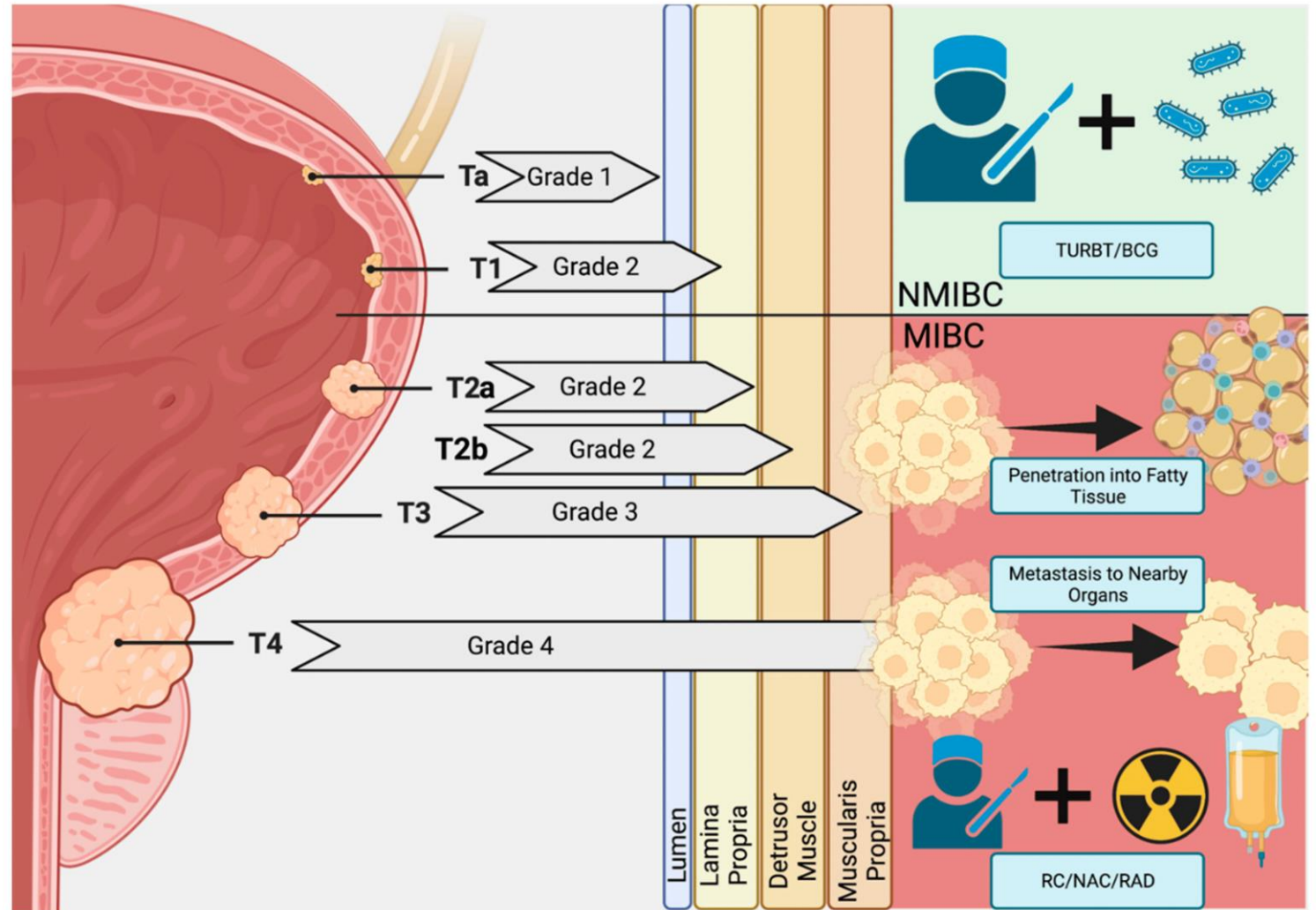


None

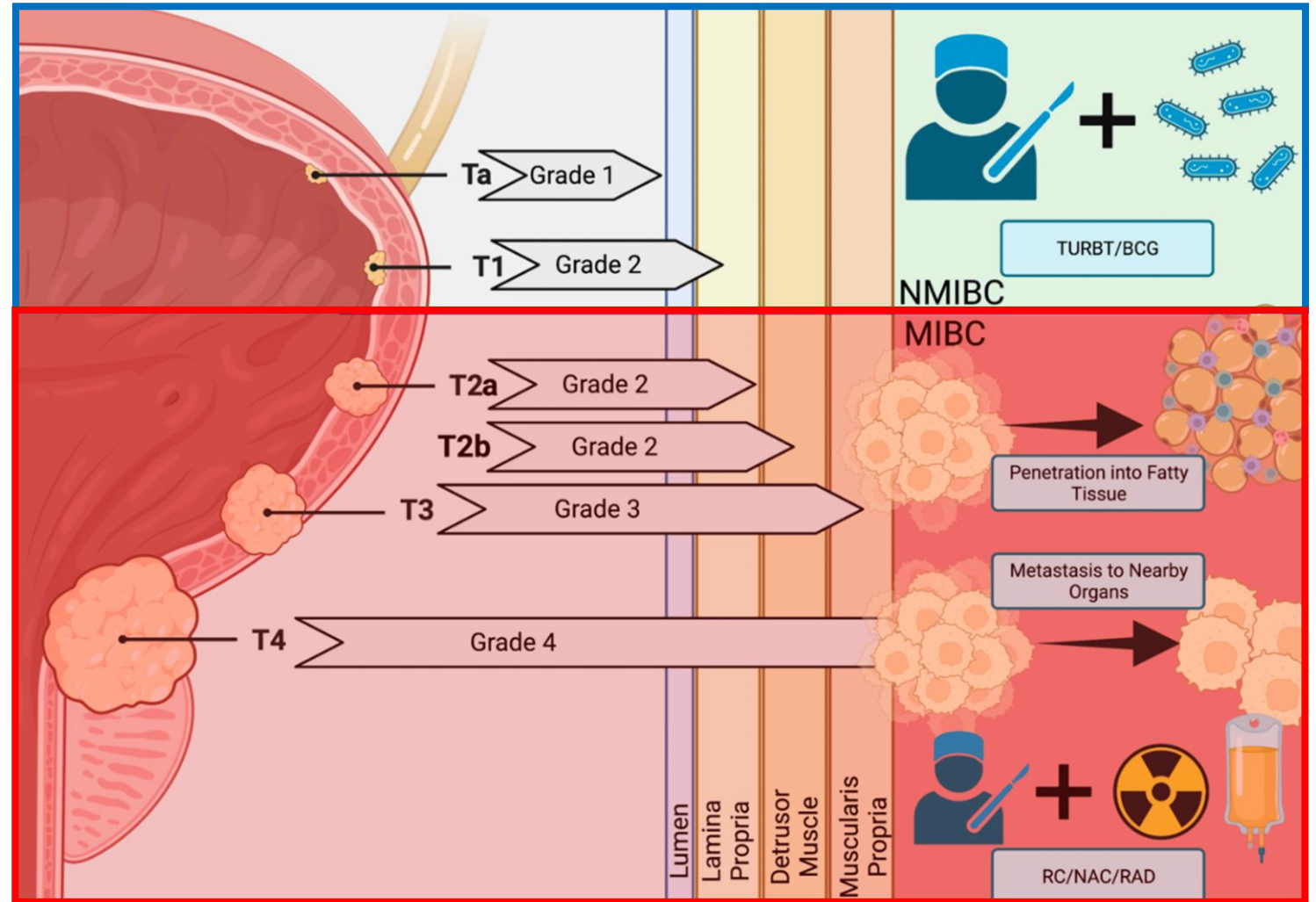


~~DON'T~~
GO RED.
 GO TO A
DOCTOR.

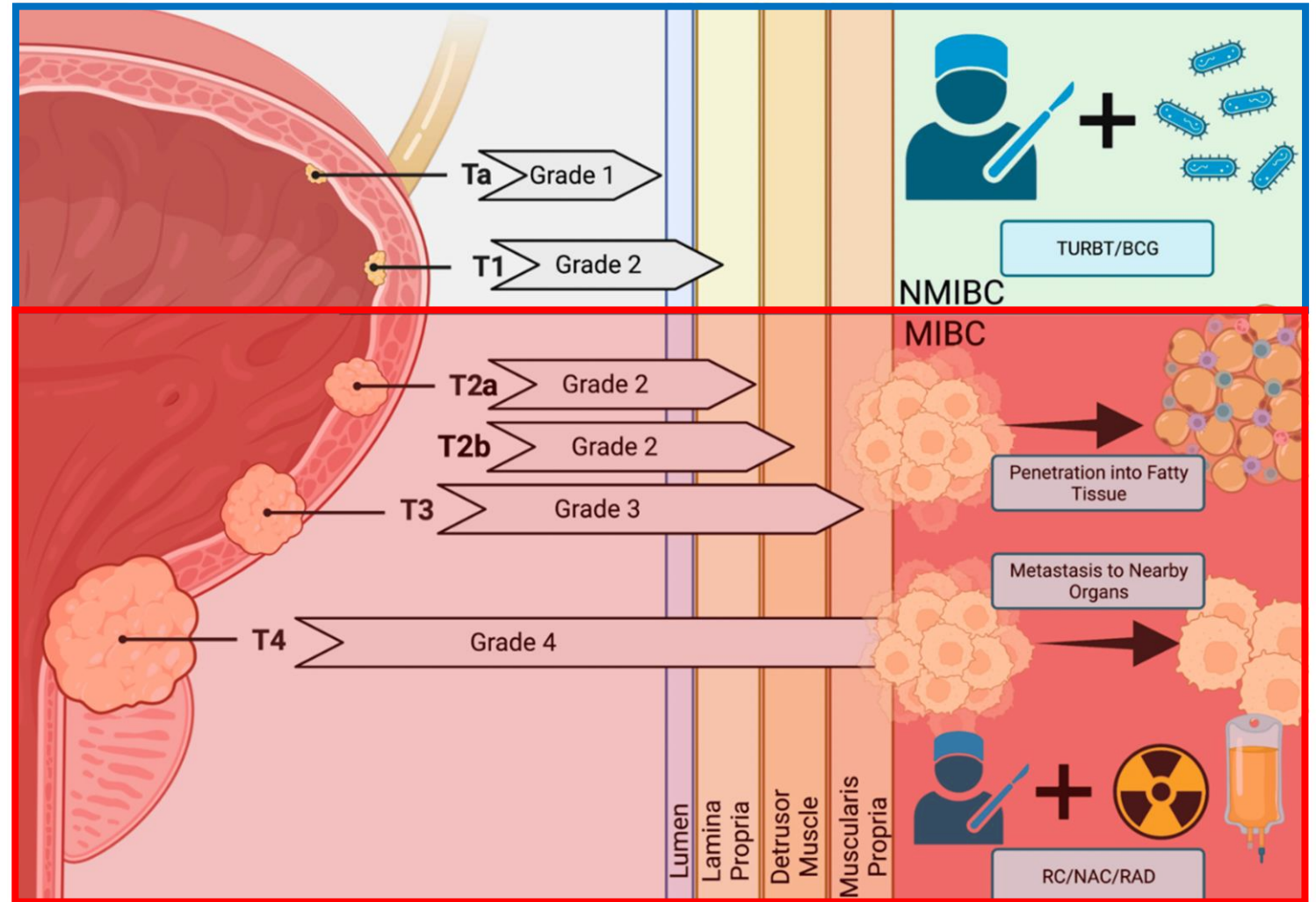
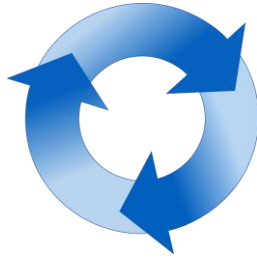
Brought to you by the World Bladder Cancer Patient Coalition



70% NMIBC



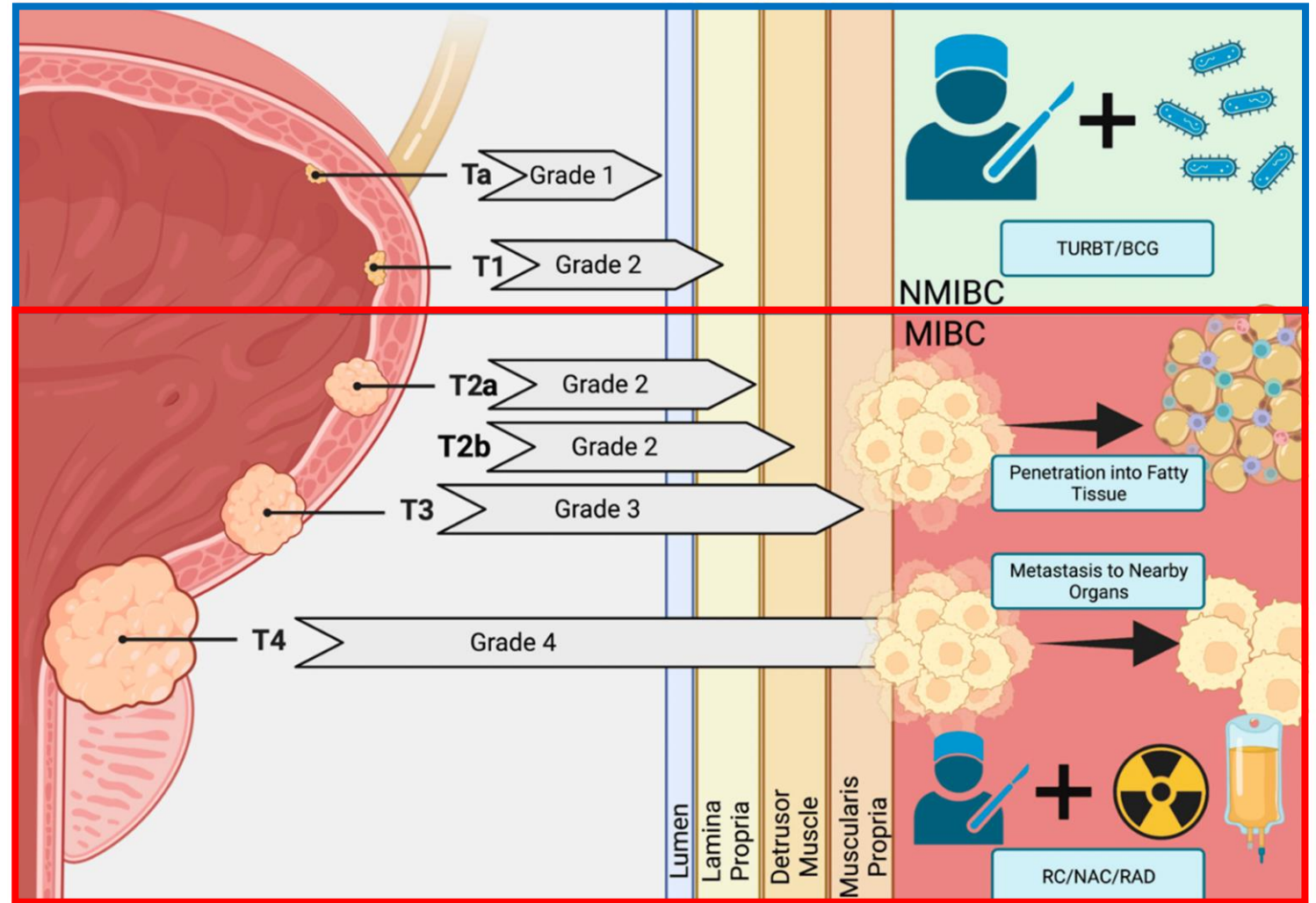
70% NMIBC
Recurrence



70% NMIBC

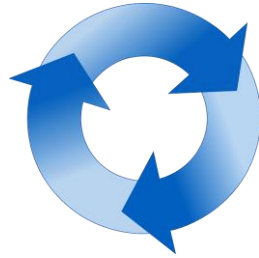
Recurrence

Progression



70% NMIBC

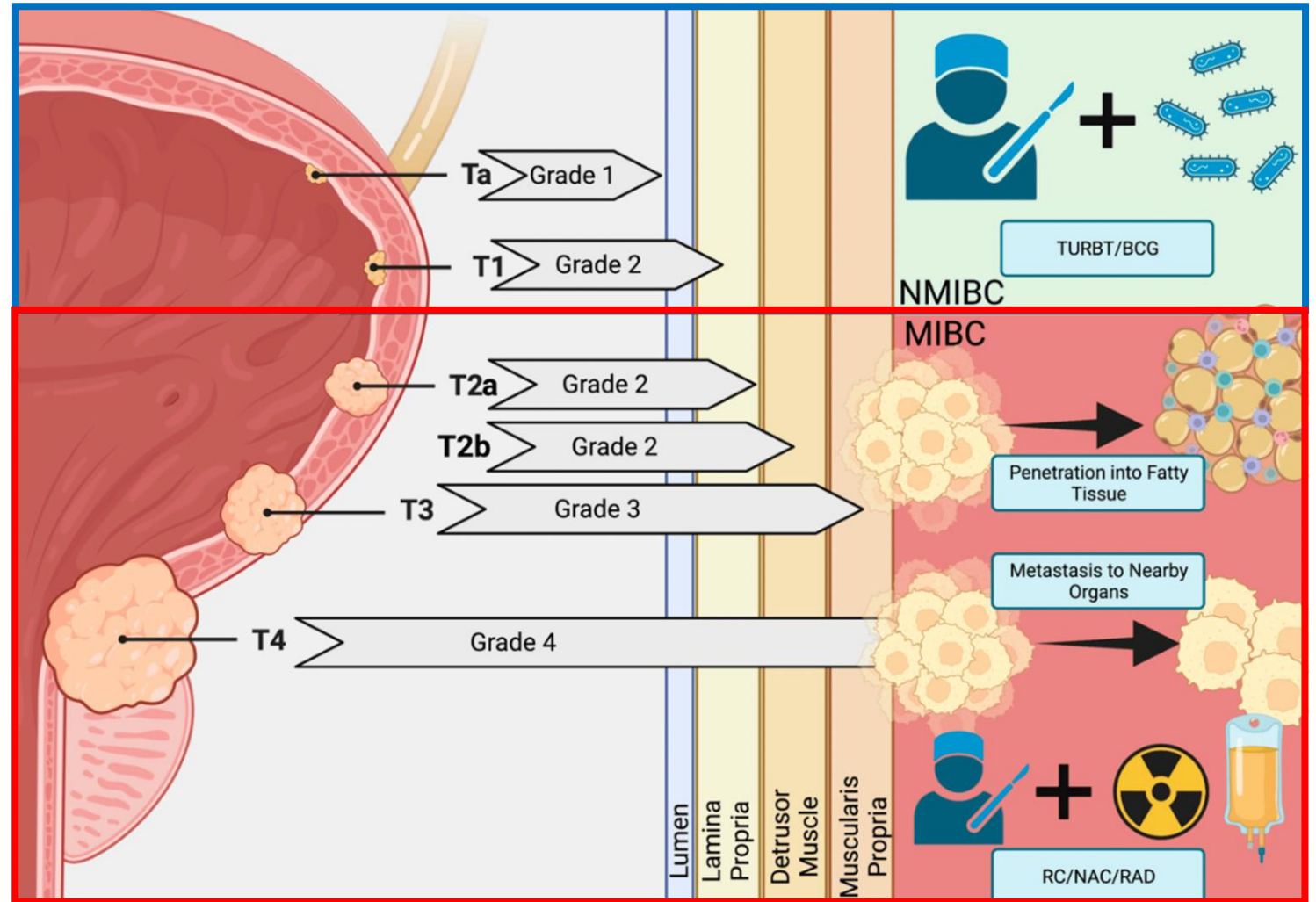
Recurrence



Progression

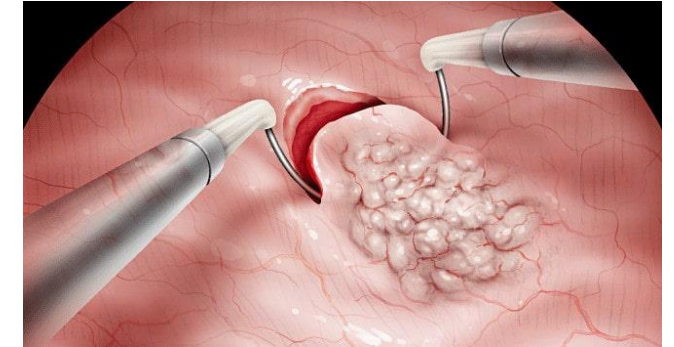
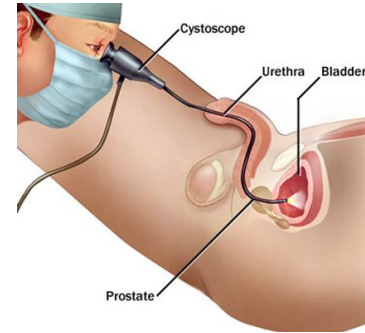


Lethal disease

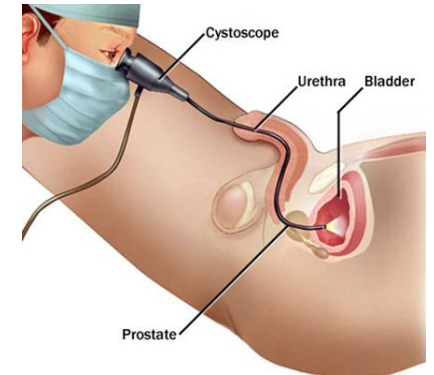
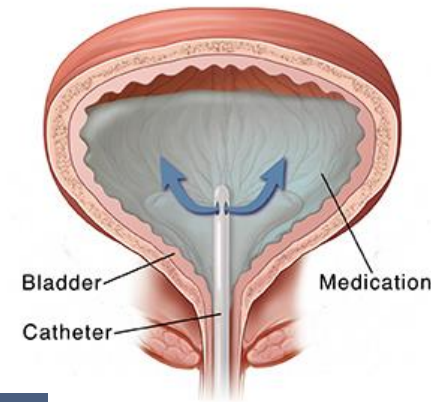




Recurrent cystoscopies
Recurrent TURBT procedures
Bladder instillations



Progression
Cystectomy
Chemotherapy



Prevalent – omnipresent in urological practices

Bothersome due to many interventions (recurrence)

Expensive due to many interventions (recurrence and progression)

Potentially *dangerous* (progression)

Bothersome due to potentially dangerous (progression)

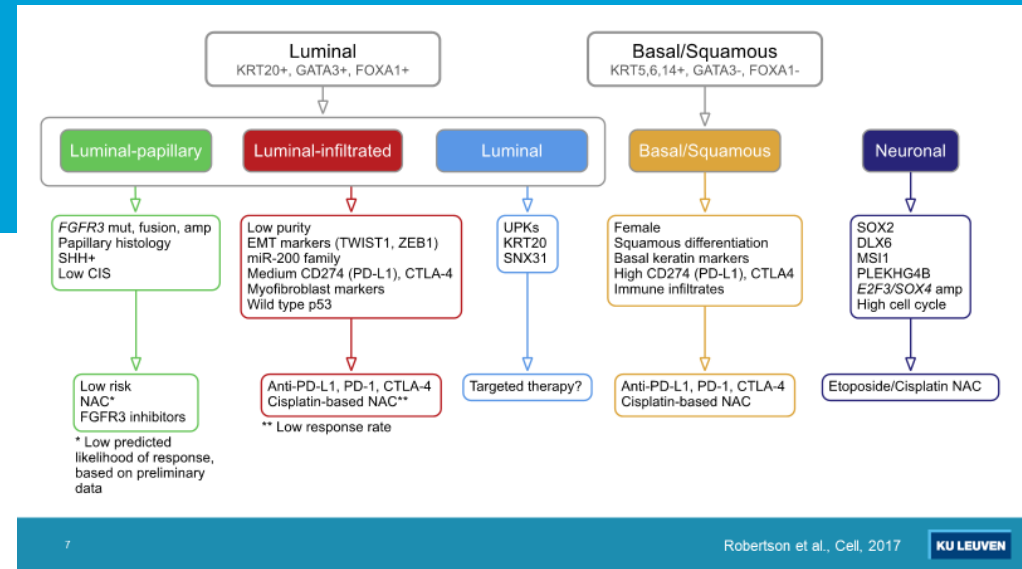
Typically a disease with a *longitudinal* trajectory

Notorious for *practice variation* and non-adherence to guidelines

→ Can we consistently extract data from the EHR?

→ Can we visualize these data?





OMICS have entered the field

OMIC classification of disease has been proposed (not widely accepted or used in clinical practice)

New medical *disease targets* are currently present

- Can we incorporate this knowledge in disease management?
- Can we link this knowledge to current disease risk groups?
- Can we refine current disease risk groups?



Standardized data collection in different hospitals

Generation of a data catalogue (EHR → OMOP) in different hospitals

(automated) data extraction → generic ready to analyse worksheet for multifactorial analysis and machine learning

Visualisation of longitudinal patient journey

Generate OMIC data on a retrospective dataset

Generate OMIC data on a prospective dataset

Scale project for use in multihospital setting



Q1: Is there a **difference in transcriptomic profiles** (either by single-cell RNA sequencing or bulk sequencing) between patients

Patients with low-/intermediate-/high-risk NMIBC/m(-) MIBC/m(+) MIBC at first diagnosis

Patients with intermediate-/high-risk NMIBC who progress to MIBC

Patients with BCG-naïve intermediate-/high-risk NMIBC recurrence

Q2: Is there a **difference in transcriptomic profiles** (either by single-cell RNA sequencing or bulk sequencing) between patients

Patients with intermediate-/high-risk NMIBC recurrence during or after BCG

Patients with intermediate-/high-risk NMIBC recurrence after anti-PD-1/PD-L1 treatment or nadofaragene firadenovec (Instiladrin®), FGFR inhibitor



Q3: What is the **natural history & epidemiology of DNA alterations** (as FGFR or other biomarkers) in bladder cancer (logitudinal from NMIBC – MIBC – mUC)?

How do they change through the evolution of the disease?

How will certain treatments effect these changes?

Q4: Can we **predict recurrence/progression timelines & outcomes** in correlation with epigenetic/transcriptomic biomarkers & clinical/pathological data for intermediate-/high-risk NMIBC and MIBC?

Is there genetic predisposition for development/progression or treatment response

Panels available UZ Leuven, MSK-IMPACT, Illumina

Regarding existing risk models/calculators (EORTC/CUETO)





Legal framework



ATHENA as a **pilot** project for (secondary) use of RWD
In the setting of a VLAIO project

Needed a **specific trajectory** through ethical – data protection –
legal departments/boards

*What are the
implications?*

Took a lot of time





Legal framework



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Legal framework



ATHENA as a **pilot** project for (secondary) use of RWD
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Needed a specific trajectory through the ethical and data protection
BUT no approval for image analysis

*What are the
implications?*

Took a lot of time





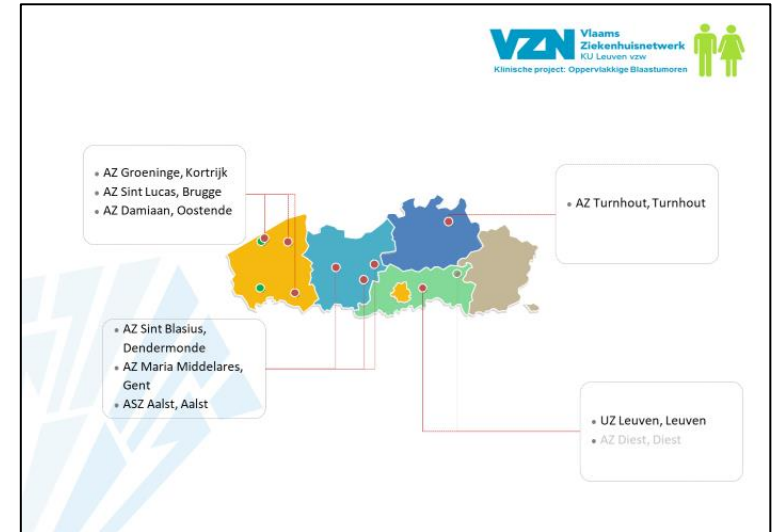
Data collection

Flemish bladder cancer registry

70-90% data completeness in **best** centers

other centers...

no data on MIBC or M+ BC



Data catalogue

link between EHR parameters and OMOP (also omic data)

implies **choices!**

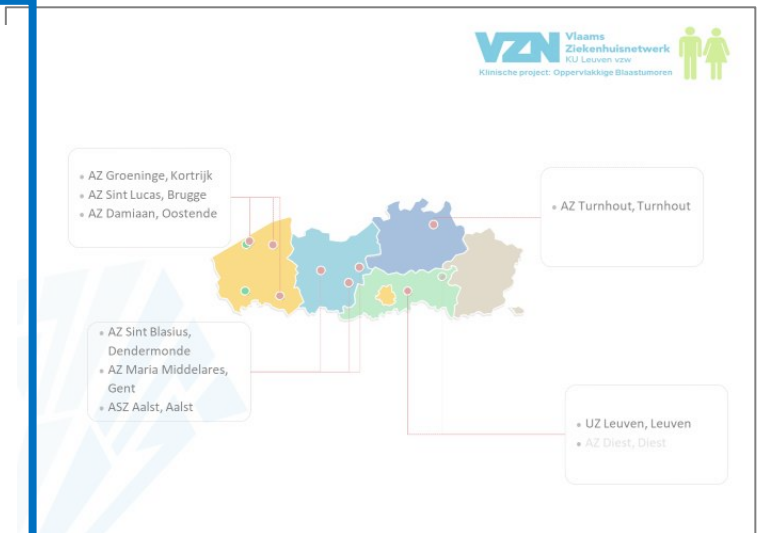
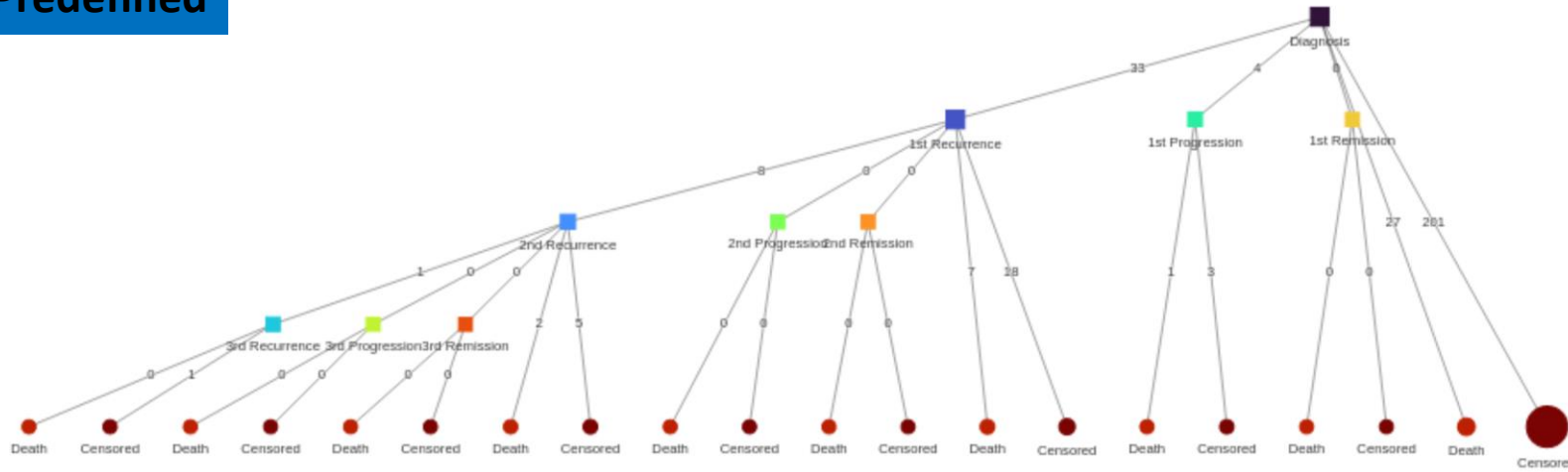
Data extraction

large datafiles, missing data,... → needs **manual** data iteration





Predefined



Data

Work in Progress

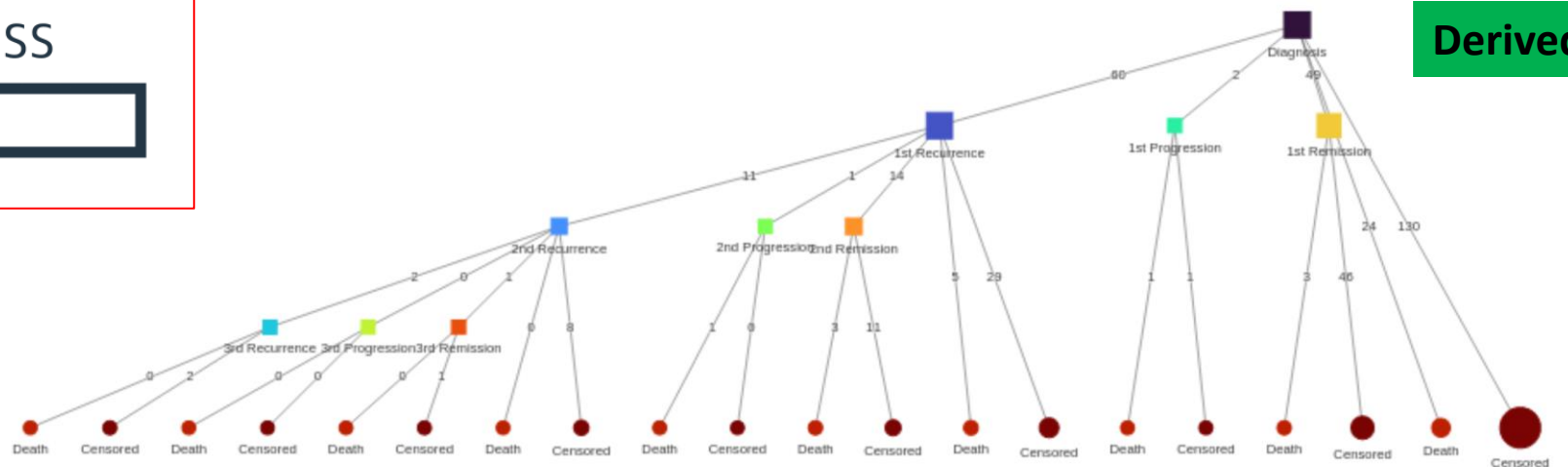


implies choices

Data extraction

large datafiles,

Derived





Biobanking – omics data generation

Tissue collection in NMIBC can be **challenging**

small lesions (enough tissue?)

immediate processing

not part of routine clinical practice

*What are the
implications?*

- Presence of research **personnel** required
- Active cooperation of **surgeon** required





Biobanking – omics data generation

Omics data generated

TSO 500

Bulk RNA seq

} 102 patients (retro)

analysis through available pipelines ongoing

WP3

methylation profiling ongoing (50%)

single nucleus sequencing planned (Q1 2024)

→ Active cooperation of **surgeon** required

What
impli



(NMI)BC is a challenging case that fits well in the scope of the ATHENA project

Many hurdles taken, but several hurdles ahead

We are not running a sprint, but a marathon

ATHENA can serve as a baseline project



UZ Leuven team

Murat Akand

Jan Cornelissen

Andries Clinkaert

Loic Baekelandt

all CABLE participants

all clinicians that use(d) format

Agnes Van den Berghe, Ann Lisa

Hilde Steeno

...

**THANK
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Joris Vermeesch

...

JNJ

Flavio Cammarone

Michel Van Speybroeck & team

Bart Vannieuwenhuyse

Tinne Lewi

Patricia Van Rompuy

Valerie Vandeweerd

Dario Masullo ...

And many more...

questions?

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