

Nouvelle approche diagnostic d'une infection urinaire

Julien Favresse

- **Tigettes et sédiments urinaires**
- **Définition d'une infection urinaire**
- **Approche par statistiques « simples »**
- **Approches par statistiques « avancées » / machine learning**



- **Tigette**

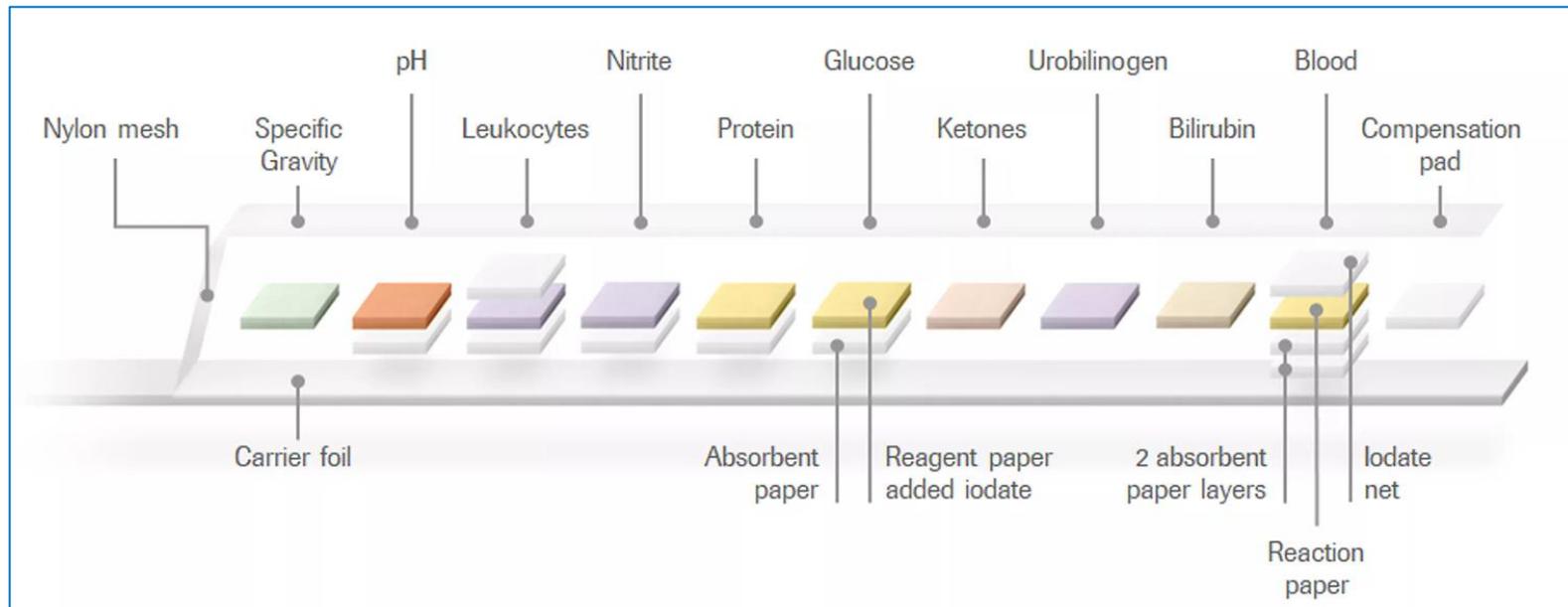
- Nitrites
- Leucocyte estérase
- Corps cétoniques
- Erythrocyte et hémoglobine
- Glucose
- Protéines
- Bilirubine
- Urobilinogène
- pH
- Densité
- Couleur
- Turbidité

- **Sédiment**

- Globules blancs
- Globules rouges
- Cellules épithéliales
 - Squameuses
 - Non-squameuses
- Cylindres
 - Pathologiques
 - Hyalins
- Bactéries
- Levures
- Mucus
- Cristaux

- **Tigette**

- Urine absorbée sur des petits PAD
- Réaction spécifique pour chaque analyte



- Tigette

- Me
- 5,5
- Lun



(5)
détecteur

- A Photomètre a
- B Zones réactive
- C Anneau de DI

- **Tigette**

- Réflectance # concentration de l'analyte dans l'urine
- Très utile pour définir un test REFLEX quantitatif

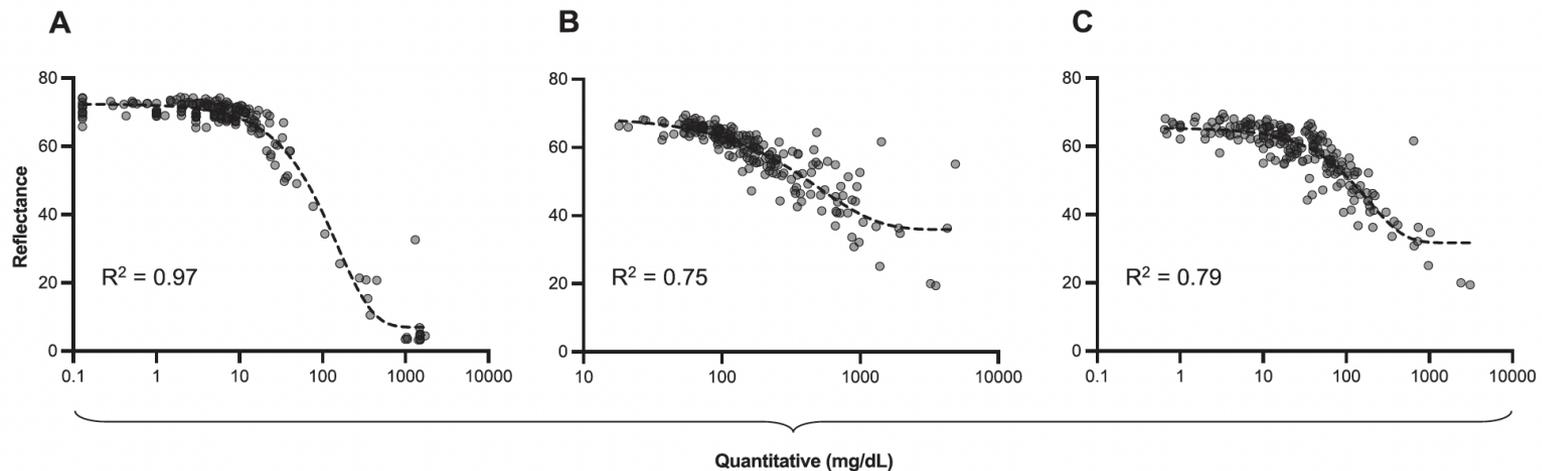
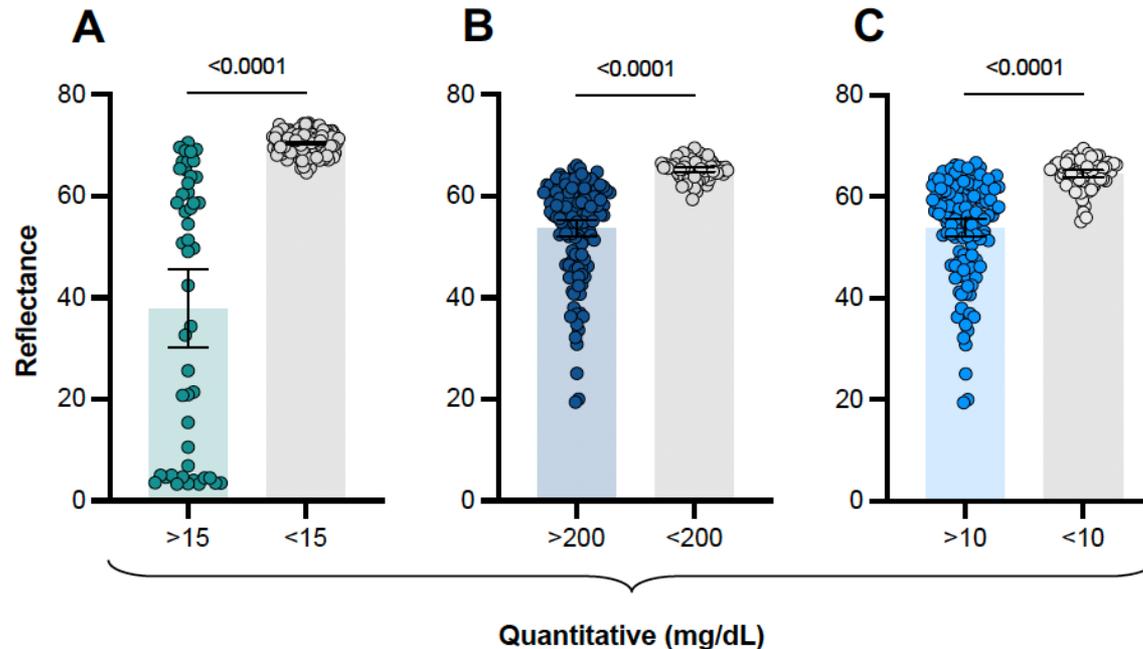


Fig. 2. Comparison of reflectance data to quantitative measurements. Correlation curves between reflectance data versus quantitative measurements. A = glucose; B = proteins; C = albumin.

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- Réflectance # concentration de l'analyte dans l'urine
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- **Tigette**

- Réflectance # concentration de l'analyte dans l'urine
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		cobas 601 ordinal scale arbitrary concentrations	Menarini ordinal scale arbitrary concentrations	cobas 601 reflectance (optimized ROC curve)	cobas 601 reflectance (optimized for sensitivity)
Glucose > 15 mg/dL	FP _D	0.00	0.00	3.90	51.30
	FN _G	48.28	79.31	17.24	0.00
	FN _C	0.00	0.00	0.00	0.00
Glucose ≥ 54 mg/dL	FP _D	6.63	1.66	0.00	0.00
	FN _G	0.00	0.00	0.00	0.00
	FN _C	0.00	0.00	0.00	0.00
Proteins ≥ 200 mg/L	FP _D	22.66	1.56	13.28	58.59
	FN _G	4.55	31.82	4.55	0.00
	FN _C	9.09	0.00	10.00	0.00
Albumin ≥ 10 mg/L	FP _D	6.00	0.00	4.00	86.00
	FN _G	62.07	90.80	49.41	0.00
	FN _C	2.94	26.47	1.49	0.00

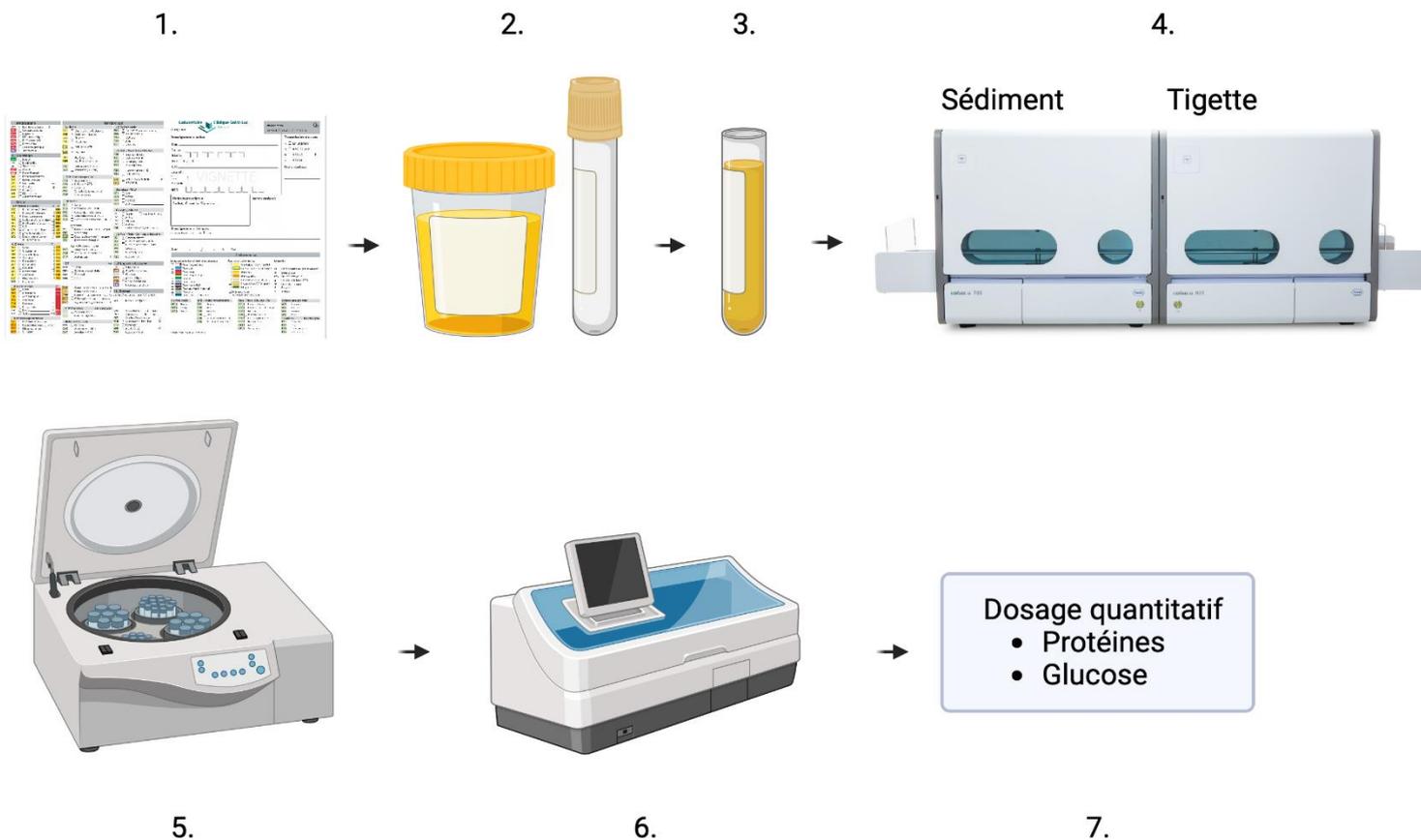
• Tigette

Table 1

Comparison of sensitivity and specificity for glucose, proteins, and albumin using reflectance values versus ordinal scale arbitrary concentrations. Confidence intervals (95%) are represented in brackets. ROC: Receiver Operating Characteristic; AUC: Area Under the Curve; Se: Sensitivity; Sp: Specificity; PPV: Positive Predictive Value; NPV: Negative Predictive Value; NA: Not Applicable.

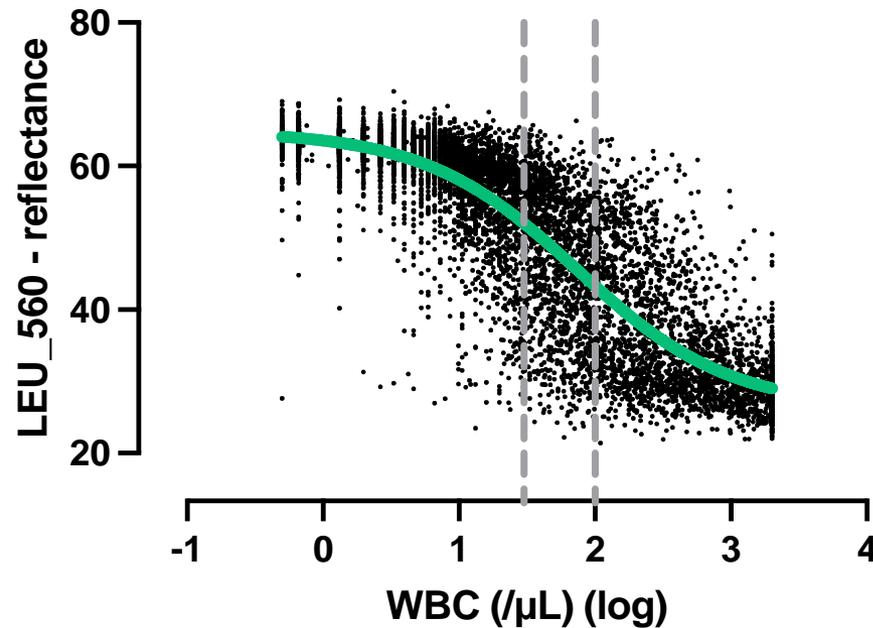
		cobas u 601 ordinal scale arbitrary concentrations	UC-MAX ordinal scale arbitrary concentrations	cobas u 601 reflectance (optimized ROC curve)	cobas u 601 reflectance (optimized for sensitivity)
Glucose > 15 mg/dL	Reflectance cut-off	NA	NA	≤ 66.96	≤ 70.57
	AUC	NA	NA	0.965 (0.929—0.985)	NA
	Se	71.4 (56.7—83.4)	53.1 (38.3—67.5)	89.8 (77.8—96.6)	100 (92.7—100)
	Sp	100 (97.7—100)	100 (97.7—100)	96.1 (91.7—98.6)	48.7 (40.6—56.9)
	PPV	100	100	87.9 (76.8—94.1)	38.13 (34.6—41.8)
	NPV	91.7 (87.7—94.5)	87.1 (83.3—90.1)	96.8 (92.8—98.6)	100
	Accuracy	93.1 (88.8—96.2)	88.7 (83.6—92.7)	94.6 (90.5—97.3)	61.0 (54.0—67.8)
Glucose > 54 mg/dL	Reflectance cut-off	NA	NA	≤ 42.42	≤ 42.42
	AUC	NA	NA	1.000 (0.982—1.000)	1.000 (0.982—1.000)
	Se	65.7 (47.8—80.9)	100 (85.2—100)	100 (85.2—100)	100.00 (85.2—100.0)
	Sp	100 (97.84—100)	98.3 (95.2—99.7)	100 (98.0—100)	100 (98.0—100)
	PPV	100	88.5 (71.4—95.9)	100	100
	NPV	95.8 (93.6—97.3)	100	100	100
	Accuracy	96.1 (92.5—98.3)	98.5 (95.8—99.7)	100 (98.2—100)	100 (98.2—100)
Proteins ≥ 200 mg/L	Reflectance cut-off	NA	NA	≤ 59.33	≤ 64.38
	AUC	NA	NA	0.959 (0.923—0.982)	NA
	Se	94.8 (87.2—98.6)	72.7 (61.4—82.3)	94.7 (87.1—98.5)	100 (95.3—100)
	Sp	77.3 (69.1—84.3)	98.4 (94.5—99.8)	86.7 (79.6—92.1)	41.41 (32.8—50.4)
	PPV	71.6 (64.5—77.7)	96.6 (87.6—99.1)	81.1 (73.3—87.0)	50.7 (47.0—54.3)
	NPV	96.1 (90.5—98.5)	85.7 (80.6—89.6)	96.5 (91.3—98.6)	100
	Accuracy	83.9 (78.1—88.7)	88.8 (83.6—92.8)	89.7 (84.7—93.5)	63.4 (56.4—70.0)
Albumin ≥ 10 mg/L	Reflectance cut-off	NA	NA	≤ 61.89	≤ 67.91
	AUC	NA	NA	0.900 (0.850—0.937)	NA
	Se	63.9 (55.8—71.4)	36.9 (29.4—45.0)	72.1 (64.3—79.0)	100 (97.6—100)
	Sp	94.0 (83.5—98.8)	100 (92.6—100)	96.0 (86.3—99.5)	14.0 (5.8—26.7)
	PPV	97.0 (91.6—99.0)	100	98.2 (93.3—99.5)	78.2 (76.2—80.0)
	NPV	45.8 (40.4—51.3)	34.0 (31.4—36.7)	52.3 (45.9—58.6)	100
	Accuracy	71.3 (64.5—77.4)	52.4 (45.3—59.4)	77.6 (71.2—83.1)	78.9 (72.7—84.3)

• Tigette



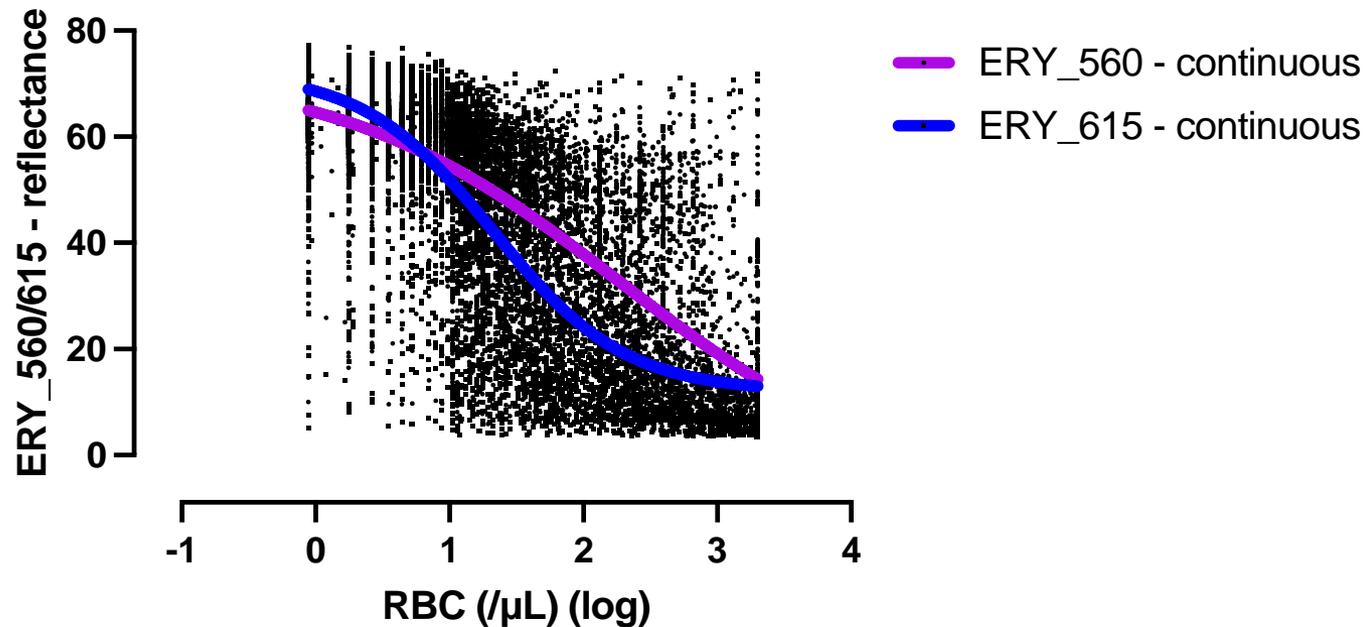
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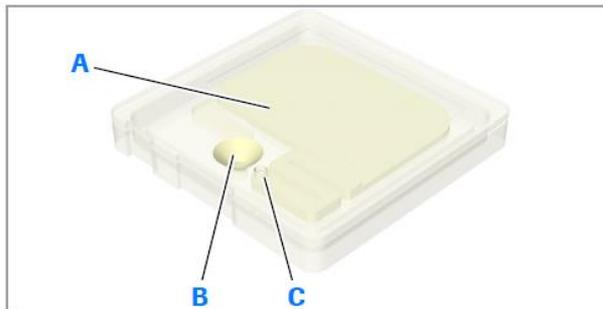
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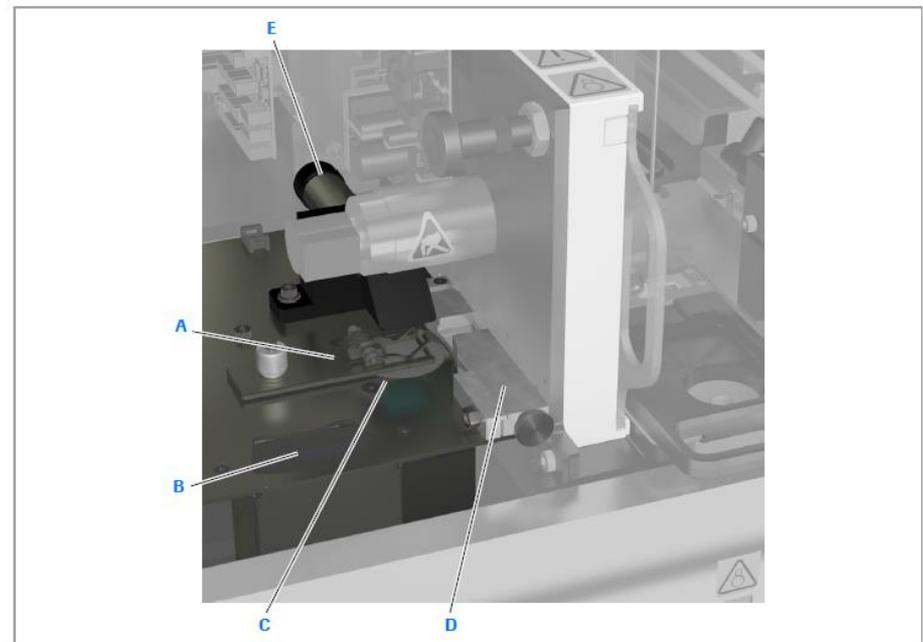


- **Sédiment**

- Cuvette avec lecture au microscope digital
- Éléments rassemblés en une seule couche



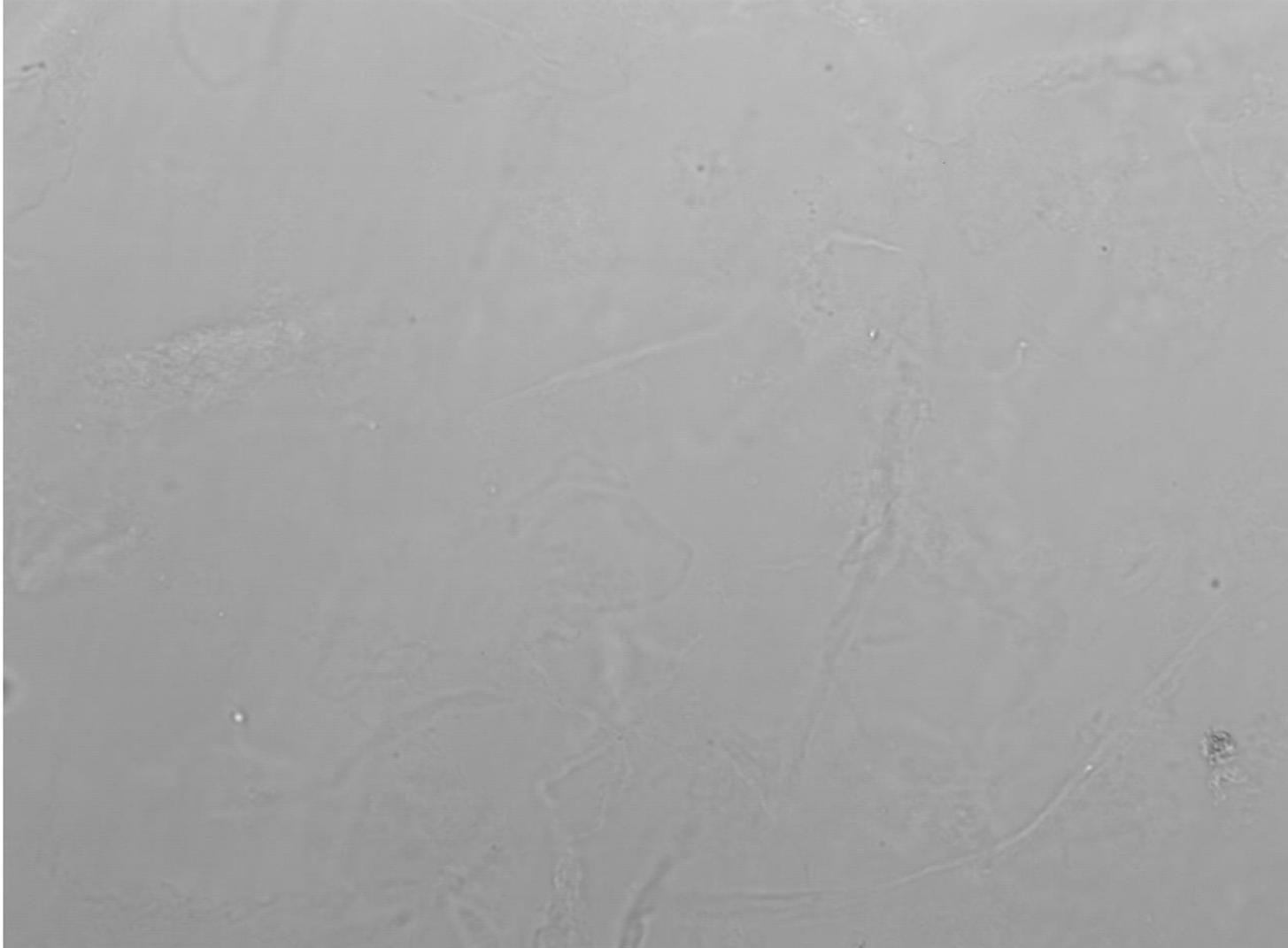
- A** Zone photographique, **C** Zone de débordement contient l'urine
B Orifice d'injection



- A** Bras du microscope
B Dévidoir à déchets
C Support microscope

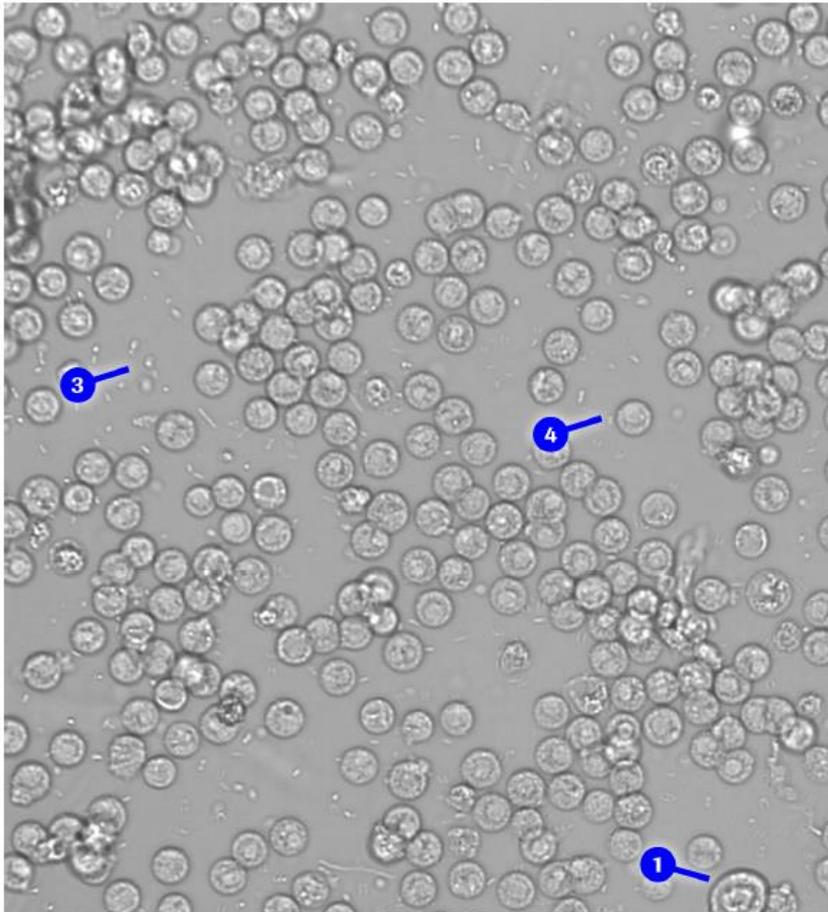
- D** Rail cuvettes
E Lampe du microscope

- **Sédiment** (négatif)

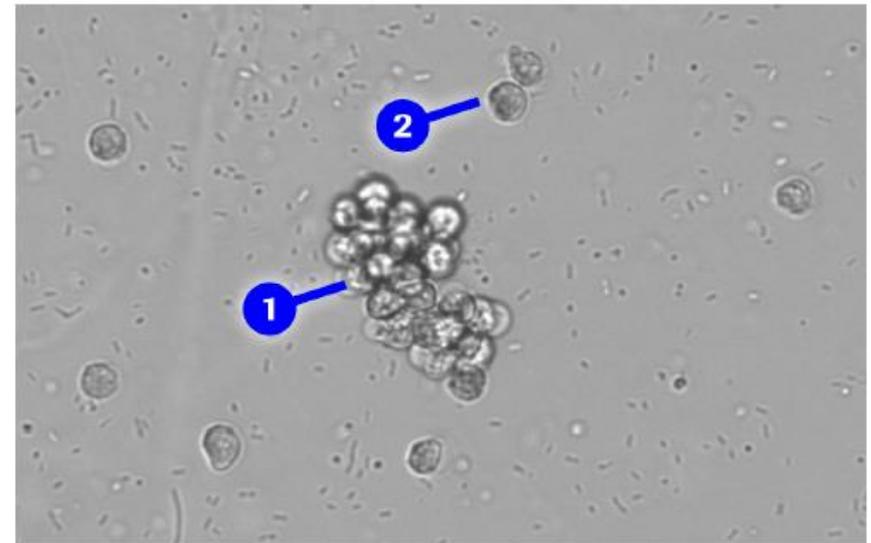


• Sédiment (GB)

WBC, RBC, NEC et BAC



Amas de WBC, WBC, BAC et MUC

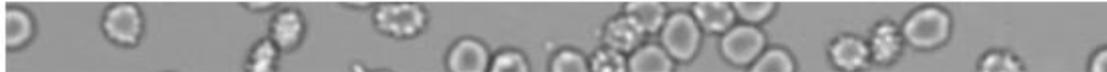


Amas de globules blancs (WBC) (1), globules blancs (WBC) (2) avec bactéries (BAC) et mucus (MUC). Les amas de WBC ne doivent pas être assimilés à des cylindres de WBC. Ces amas de WBC, qui apparaissent pendant la phagocytose des BAC et des particules WBC en décomposition, surviennent typiquement lors d'une infection des voies urinaires (UTI).

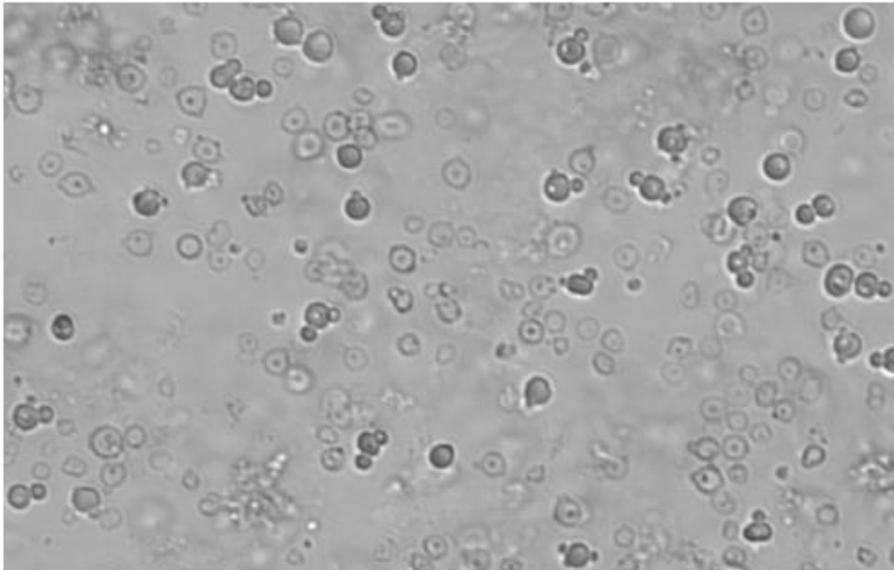
Cellule épithéliale non squameuse (NEC) (1), quelques globules rouges (RBC) (2) et quelques bactéries (BAC) (3), en plus de nombreux globules blancs (WBC) (4). Les neutrophiles polymorphonucléaires (12 – 15 µm) sont aisément reconnaissables par leur noyau segmenté.

- **Sédiment** (GR iso/dysmorphiques)

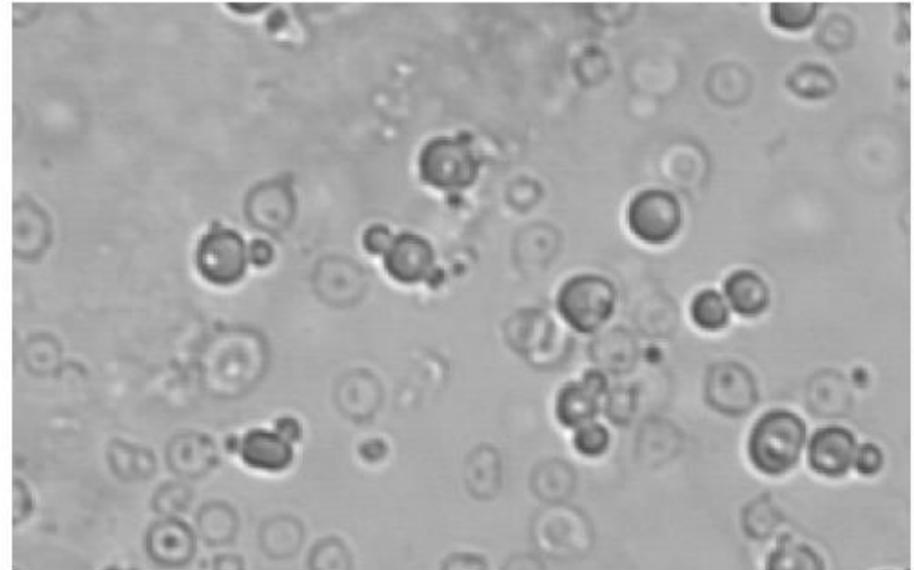
RBC



RBC



RBC, grossissement de 200 %

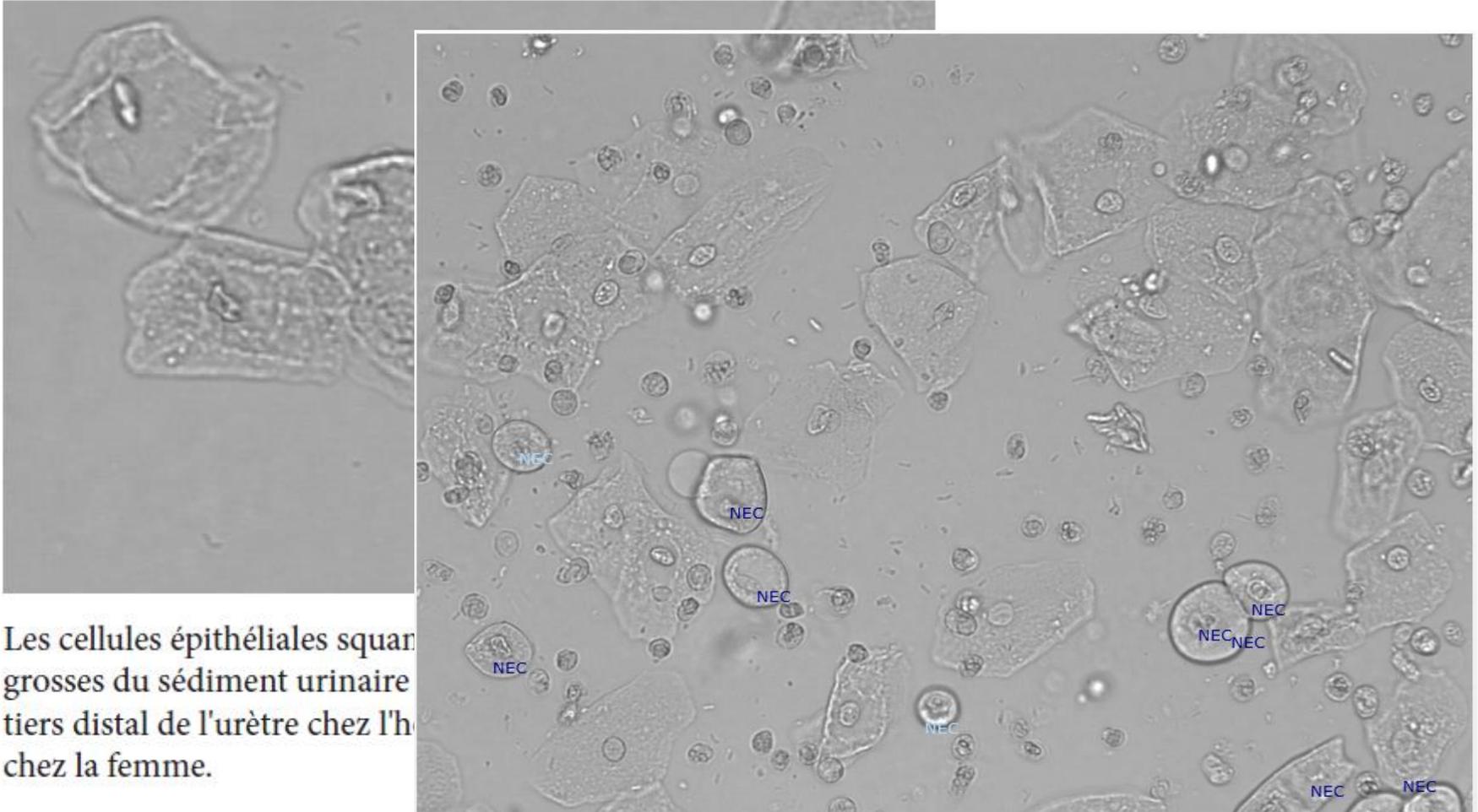


Globules rouges dysmorphiques (désignés par RBC). Les RBC dits « dysmorphiques » ont subi des changements morphologiques dans les reins (cellules glomérulaires). Les acanthocytes (décrits comme des RBC) sont considérés comme d'origine presque exclusivement glomérulaire.

Globules rouges dysmorphiques (désignés par RBC). Les RBC présentant diverses lésions membranaires (perte de cytoplasme, extrusion cytoplasmique de la membrane cellulaire, éléments granulaires sur la membrane cellulaire, etc.).

- **Sédiment** (SEC et NEC)

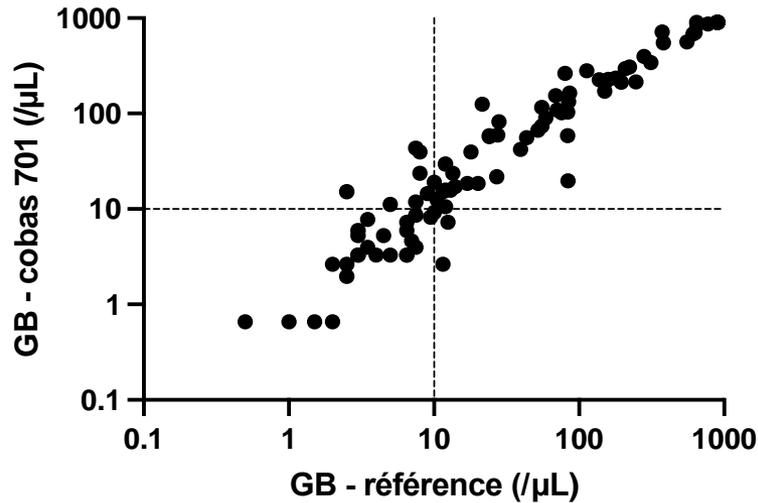
SEC



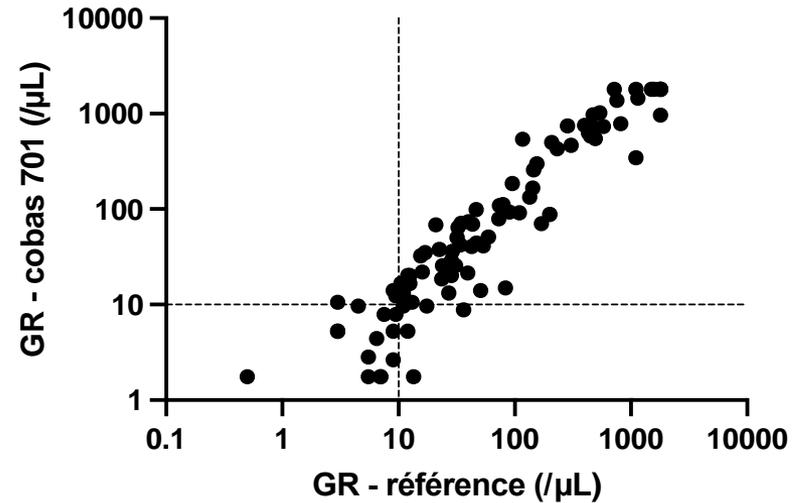
Les cellules épithéliales squar
grosses du sédiment urinaire
tiers distal de l'urètre chez l'h
chez la femme.

• Sédiment (manuel versus automatique)

Pearson $r = 0.98$
Spearman $r = 0.96$

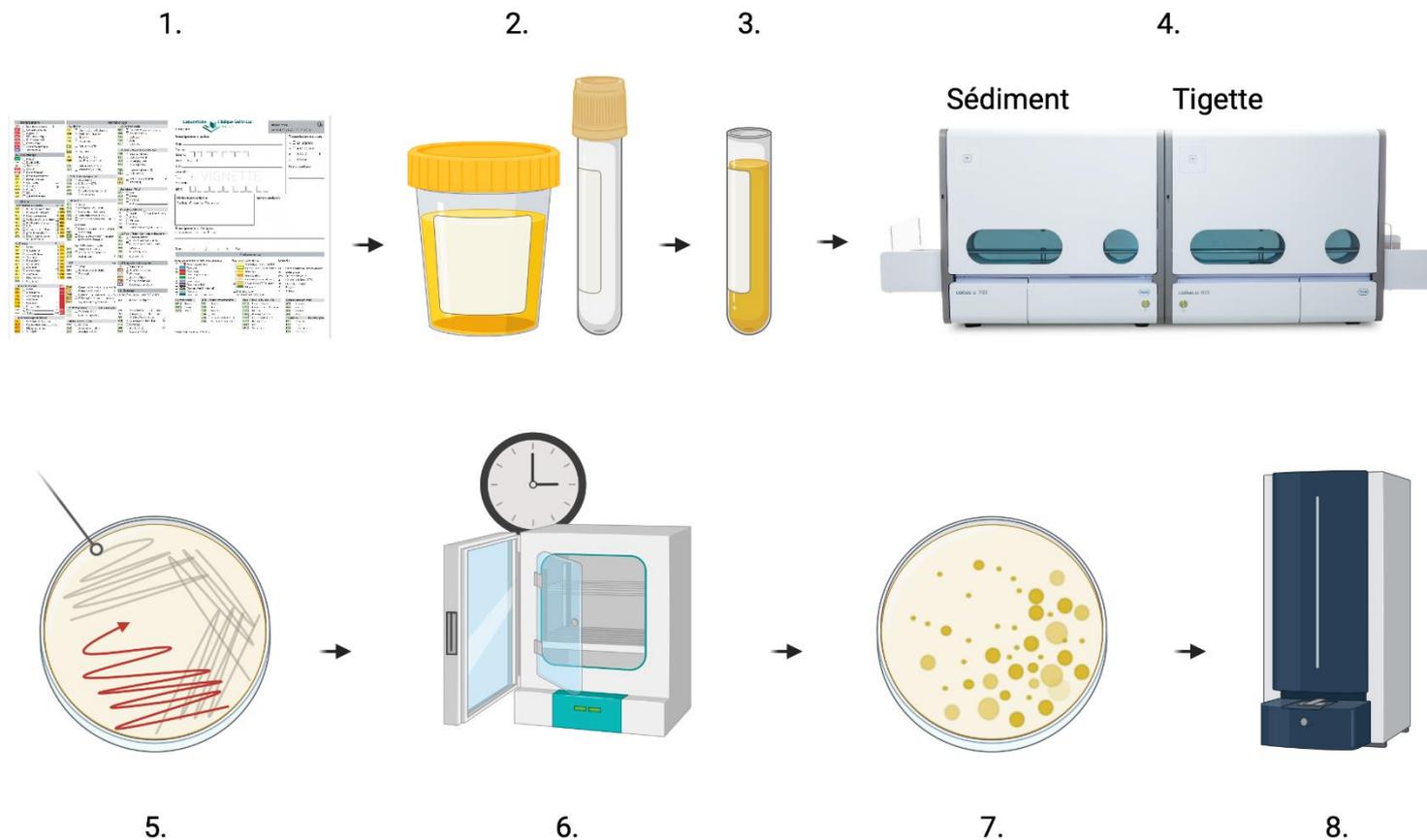


Pearson $r = 0.93$
Spearman $r = 0.94$



- 2 lecteurs indépendants au microscope (cellule de comptage uriglass)
- Comptage cobas 701
- 100 urines fraîches
- Critère EFLM = Spearman $r > 0.9$

• Sédiment



Délai de 24-48h pour identifier le germe impliqué!

• Protocole



Cas #1

- Femme
- 37 ans
- NIT +
- LEU +++
- GB 172/ μ L



Cas #2

- Femme
- 41 ans
- NIT +
- LEU +++
- GB 160/ μ L

Tigette/sédiment

Microbiologie

Culture négative

***E. coli* CFU >10⁵/mL**

- Définition d'une infection urinaire?
 - Culture
 - Germe(s) identifié(s)?
 - Quantité? CFU
 - Pure? Mixte? Prédominance?
 - Présence de globules blancs?



- Définition d'une infection urinaire?

Etude	Critère
Heckerling <i>et al.</i> 2007 IJMI	(1) $\geq 10^5$ CFU/mL et si uropathogène ou (1') $\geq 10^2$ CFU/mL et si uropathogène
Sterry-Blunt <i>et al.</i> 2015 JMM	(1) $\geq 10^4$ CFU/mL. <u>NB: flore mixte considérée comme négative</u>
Shimoni <i>et al.</i> 2017 Plos One	(1) $\geq 10^5$ CFU/mL, any bacteria
Taylor <i>et al.</i> 2018 PloS One	(1) $> 10^4$ CFU/HPF (2) flore mixte si <i>E. coli</i> présent
De la Tabla <i>et al.</i> 2019 Ann Clin Chem	(1) $\geq 10^4$ CFU/mL. <u>NB: ≥ 3 isolats = contamination</u>
Burton <i>et al.</i> 2019 BMC MIDM	(1) Tous les isolats avec ATBG associé. <u>NB: flore mixte considérée comme négative</u>
Dedeene <i>et al.</i> 2023 RBSLM poster	(1) $\geq 10^4$ CFU/mL et si uropathogène. <u>NB: flore mixte considérée comme négative</u>

- Définition d'une infection urinaire?

Open Forum Infectious Diseases

REVIEW ARTICLE



Definitions of Urinary Tract Infection in Current Research: A Systematic Review

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Defining urinary tract infection (UTI) is complex, as numerous clinical and diagnostic parameters are involved. In this systematic review, we aimed to gain insight into how UTI is defined across current studies. We included 47 studies, published between January 2019 and May 2022, investigating therapeutic or prophylactic interventions in adult patients with UTI. Signs and symptoms, pyuria, and a positive urine culture were required in 85%, 28%, and 55% of study definitions, respectively. Five studies (11%) required all 3 categories for the diagnosis of UTI. Thresholds for significant bacteriuria varied from 10^3 to 10^5 colony-forming units/mL. None of the 12 studies including acute cystitis and 2 of 12 (17%) defining acute pyelonephritis used identical definitions. Complicated UTI was defined by both host factors and systemic involvement in 9 of 14 (64%) studies. In conclusion, UTI definitions are heterogeneous across recent studies, highlighting the need for a consensus-based, research reference standard for UTI.

Keywords. complicated urinary tract infection; cystitis; definition; pyelonephritis; urinary tract infection.

- Définition d'une infection urinaire?

Table 7-3. The pathogenicity and frequency of example microorganisms in urine

Frequency (percent of isolates)		uUTI	cUTI	HA-UTI	CA-UTI
Pathogenicity in the urinary tract					
I. Primary pathogens	<i>E. coli</i>	70-75	55-65	45	30
	<i>S. saprophyticus</i>	3-6	-	-	-
II. Secondary pathogens	<i>Enterobacter</i> spp.	2	4	6	3
	<i>Enterococcus</i> spp.	4-5	6-11	10	10
	<i>Klebsiella</i> spp.	5-6	8-9	12	5
	<i>Proteus</i> spp.	2-4	2-5	6	11
	<i>P. aeruginosa</i>	1-3	2-7	9	11
	<i>S. aureus</i>	1-2	2-3	3	4
	<i>Citrobacter</i> spp.	2	3	1.5	5
	<i>M. morgani</i>	<1	5	<1	4
	<i>Serratia</i> spp.	<1	7	<1	<1
	<i>Aerococcus</i> spp.	1	1	-	-
	<i>Actinotignum schallii</i>	<0.1	<0.1	-	-
III. Doubtful pathogens	<i>C. urealyticum</i>	-	-	-	-
	<i>Streptococcus agalactiae</i>	3-4	2-3	<1	<1
	Yeast	1	3-7	2	7
	<i>Acinetobacter</i> spp.	<1	2	2	2
IV. Contaminants	<i>Staphylococcus</i> , coagulase negative (except <i>S. saprophyticus</i>)				
	<i>Corynebacterium</i> spp (except <i>C. urealyticum</i>)				
	<i>Gardnerella vaginalis</i>				
	<i>Lactobacillus</i> spp. (except <i>L. delbrueckii</i>)				

Abbreviations used: uUTI, uncomplicated urinary tract infection; cUTI, complicated urinary tract infection; HA-UTI, healthcare-associated urinary tract infection; CA-UTI, catheter-associated urinary tract infection; -, no data available.

• Définition d'une infection urinaire?

Définition 1	Définition 2	Définition 3	Définition 4	Définition 5
European association of urology 2015	Public Health England (PHE)/Health Protection Agency guidelines	POETIC protocol	EFLM (1)	EFLM (2)
$\geq 10^3$ CFU/mL uropathogène	$\geq 10^5$ CFU/mL uropathogène (prédominance dans flore mixte)	$\geq 10^5$ CFU/mL uropathogène (pure)	$\geq 10^3$ CFU/mL pathogène classe 1 + GB ≥ 30	$\geq 10^3$ CFU/mL pathogène classe 1 + GB ≥ 30
	$\geq 10^4$ CFU/mL uropathogène (pure)	$>10^4$ et $<10^5$ CFU/mL uropathogène (pure) + GB ≥ 30	$\geq 10^4$ CFU/mL pathogène classe 2 + GB ≥ 30	$\geq 10^4$ CFU/mL pathogène classe 2 + GB ≥ 30
	$\geq 10^3$ CFU/mL, <i>E. coli</i> ou <i>S. saprophyticus</i>	$\geq 10^5$ CFU/mL uropathogène (prédominance dans flore mixte)	$\geq 10^5$ CFU/mL pathogène classe 3 + GB ≥ 30	$\geq 10^5$ CFU/mL pathogène classe 3 + GB ≥ 30
		$\geq 10^5$ CFU/mL 2 uropathogènes (prédominance 1 vs 2) + GB ≥ 30	$\geq 10^5$ CFU/mL flore mixte + GB ≥ 30	
		$>10^4$ et $<10^5$ CFU/mL 2 uropathogènes (ensemble) + GB ≥ 30		
		10^4 à $<10^5$ CFU/mL uropathogène (prédominance dans flore mixte) + GB ≥ 30		

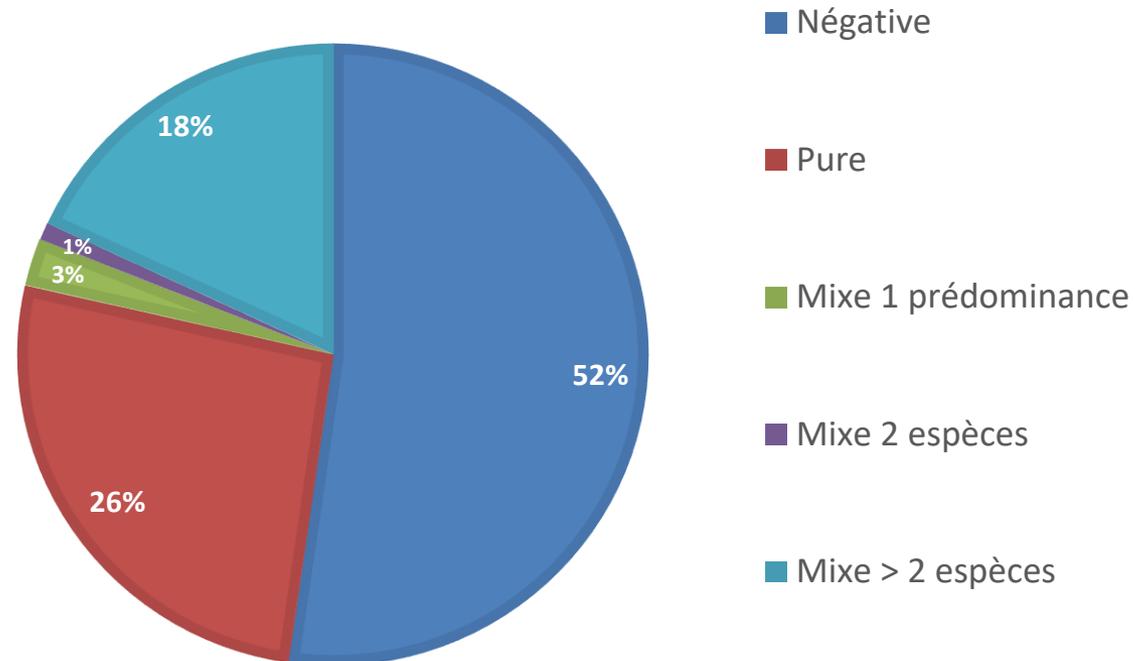
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- **Sédiment**

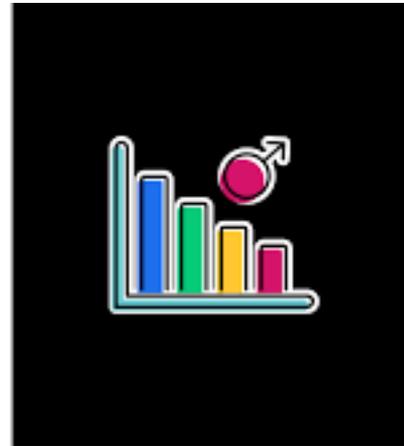
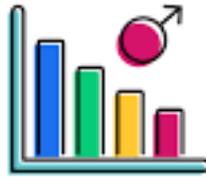
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- Base de données: 9,753 urines
 - Novembre 2023 – Janvier 2024
 - Ratio femme/homme: 5,871/3,884 (60.2%/39.8%)
 - Age: moyenne = 59 ans (min-max: 0-103 ans)
 - Femmes: moyenne = 57.2 ans
 - Hommes: moyenne = 61.8 ans



NB: 1,484 *E. coli* pure (~60%)

- Approche mono ou bi-paramétrique



Mono ou bi-paramétrique

	Test	Se/Sp (%)	VPN/VPP (%)	Accuracy (%)
EAU 2015	LEU_560	90.3/50.0	93.7/39.4	60.7
	NIT_560	41.5/95.3	82.0/76.0	81.1
	LEU_560 + NIT_560	91.8/49.7	94.4/39.6	60.8
	GB	89.3/54.4	93.4/41.3	63.6
	Test	Se/Sp (%)	VPN/VPP (%)	Accuracy (%)
PHE	LEU_560	91.6/49.4	94.8/36.9	59.7
	NIT_560	43.9/95.1	84.0/74.5	82.6
	LEU_560 + NIT_560	93.2/49.1	95.7/37.2	59.9
	GB	90.4/53.6	94.5/38.7	62.6
	Test	Se/Sp (%)	VPN/VPP (%)	Accuracy (%)
POETIC	LEU_560	95.7/49.7	97.5/35.9	60.1
	NIT_560	47.3/95.3	86.0/74.6	84.3
	LEU_560 + NIT_560	97.3/49.3	98.4/36.1	60.2
	GB	94.9/54.0	97.3/37.8	63.3

	Test	Se/Sp (%)	VPN/VPP (%)	Accuracy (%)
EFLM (1)	LEU_560	99.7/50.6	99.8/36.8	61.6
	NIT_560	44.9/94.7	85.9/71.2	83.8
	LEU_560 + NIT_560	99.8/49.8	99.9/36.4	61.0
	GB	100/55.2	100/39.1	65.2
	Test	Se/Sp (%)	VPN/VPP (%)	Accuracy (%)
EFLM (2)	LEU_560	99.8/49.6	99.9/34.1	60.0
	NIT_560	46.2/93.9	87.0/66.5	84.0
	LEU_560 + NIT_560	99.8/48.8	99.9/33.8	59.4
	GB	100/54.1	100/36.3	63.6

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Research article

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The urine dipstick test useful to rule out infections. A meta-analysis of the accuracy

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Conclusions: Overall, this review demonstrates that the urine dipstick test alone seems to be useful in all populations to exclude the presence of infection if the results of both nitrites and leukocyte-esterase are negative. Sensitivities of the combination of both tests vary between 68 and 88% in different patient groups, but positive test results have to be confirmed. Although the combination of positive test results is very sensitive in family practice, the usefulness of the dipstick test alone to rule in infection remains doubtful, even with high pre-test probabilities.

TABLE 2: Diagnostic performance of nitrite and leucocyte results relative to quantitative urine culture.

Culture	Sensitivity (%) (95% CI)	Specificity (%) (95% CI)	PPV (%) (95% CI)	NPV (%) (95% CI)	+LR (95% CI)	-LR (95% CI)
NIT+	27.7 (17.3–40.2)	98.4 (96.4–99.4)	75.0 (53.3–90.2)	88.4 (84.9–91.3)	16.8 (6.93–40.72)	0.74 (0.63–0.86)
LE+	60.0 (47.1–72.0)	73.9 (69.1–78.3)	29.1 (21.6–37.7)	91.2 (87.4–94.2)	2.30 (1.77–2.99)	0.54 (0.40–0.73)
NIT+ or LE+	72.3 (59.8–89.7)	72.5 (67.6–77.1)	32.0 (24.5–40.2)	93.6 (90.1–96.2)	2.63 (2.10–3.30)	0.38 (0.26–0.57)
NIT+ and LE+	16.9 (8.8–28.3)	99.7 (98.5–100)	91.7 (61.5–99.8)	87.1 (83.4–90.1)	61.6 (8.1–69.04)	0.83 (0.75–0.93)

NIT = nitrite, LE = leucocyte esterase, PPV = positive predictive value, NPV = negative predictive value, +LR = positive likelihood ratio, -LR = negative likelihood ratio.

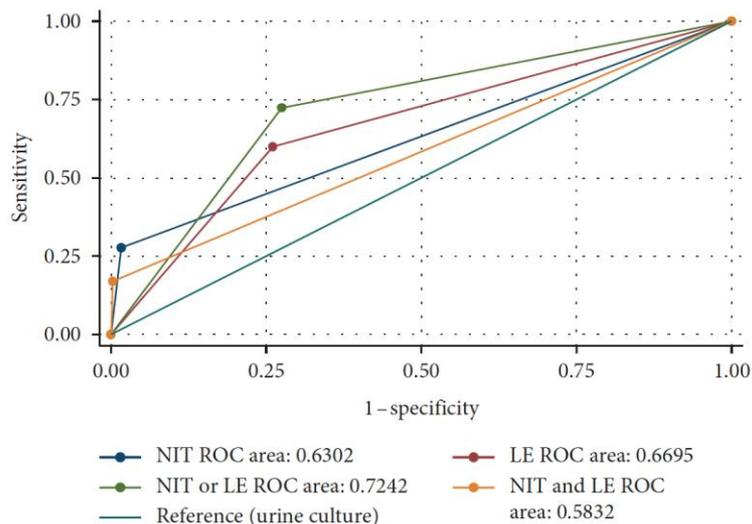


FIGURE 1: ROC curve for dipstick diagnosis with urine culture as a gold standard. NIT = nitrite, LE = leucocyte esterase.

Approche multi-paramètres

- 27 variables
 - Sédiment
 - Tigette
 - Age et sexe



1 Artificial Intelligence

Development of smart systems and machines that can carry out tasks that typically require human intelligence

2 Machine Learning

Creates algorithms that can learn from data and make decisions based on patterns observed
Require human intervention when decision is incorrect

3 Deep Learning

Uses an artificial neural network to reach accurate conclusions without human intervention

- Notions de « Machine learning »

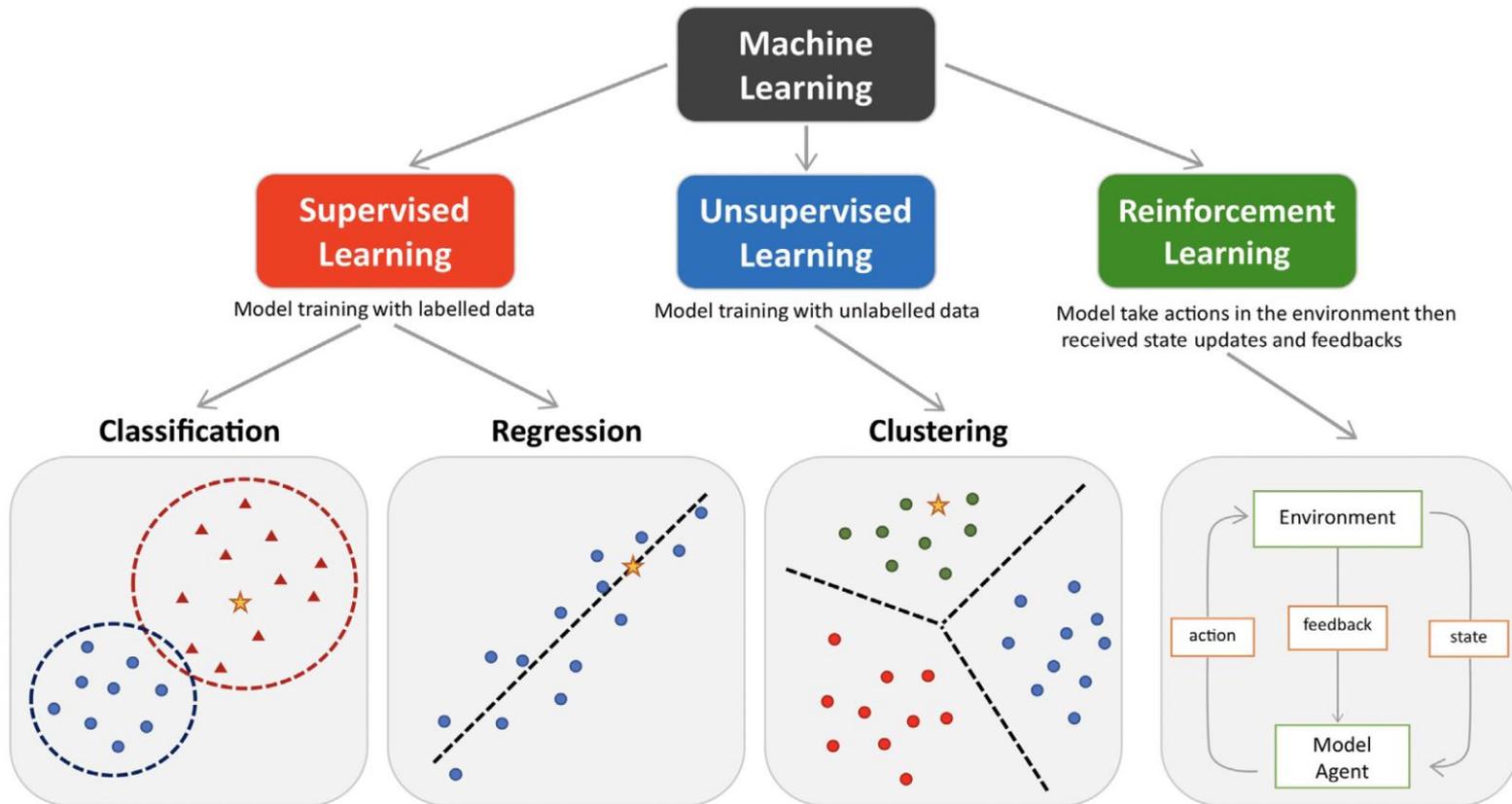


FIGURE 1 | The main types of machine learning. Main approaches include classification and regression under the supervised learning and clustering under the unsupervised learning. Reinforcement learning enhance the model performance by interacting with environment. Coloured dots and triangles represent the training data. Yellow stars represent the new data which can be predicted by the trained model.

- Notions de « Machine learning »

1. Obtention d'un modèle

A. Base de données pour l'apprentissage
(6,827 urines)



B. Obtention d'un modèle

C. Evaluation des performances

Features	Target
5.1 3.5 1.4 0.2	0
6.4 3.1 4.5 1.2	1
5.9 3.0 5.0 1.8	2
...	...
4.9 3.0 1.4 0.2	0

2. Validation du modèle

A. Base de données pour la validation
(2,926 urines)



B. Vérification des performances à partir du modèle

5.3 3.7 1.4 0.3	?
-----------------	---

- Etude SLBO: modèles utilisés:
 - **Multiple Logistic Regression**
 - **Support Vector Machine (SVM)**
 - **Random Forest**
 - **XGBoost**
 - **CatBoost**
 - **Multilayer Perceptron (MLP)**



Logistic regression

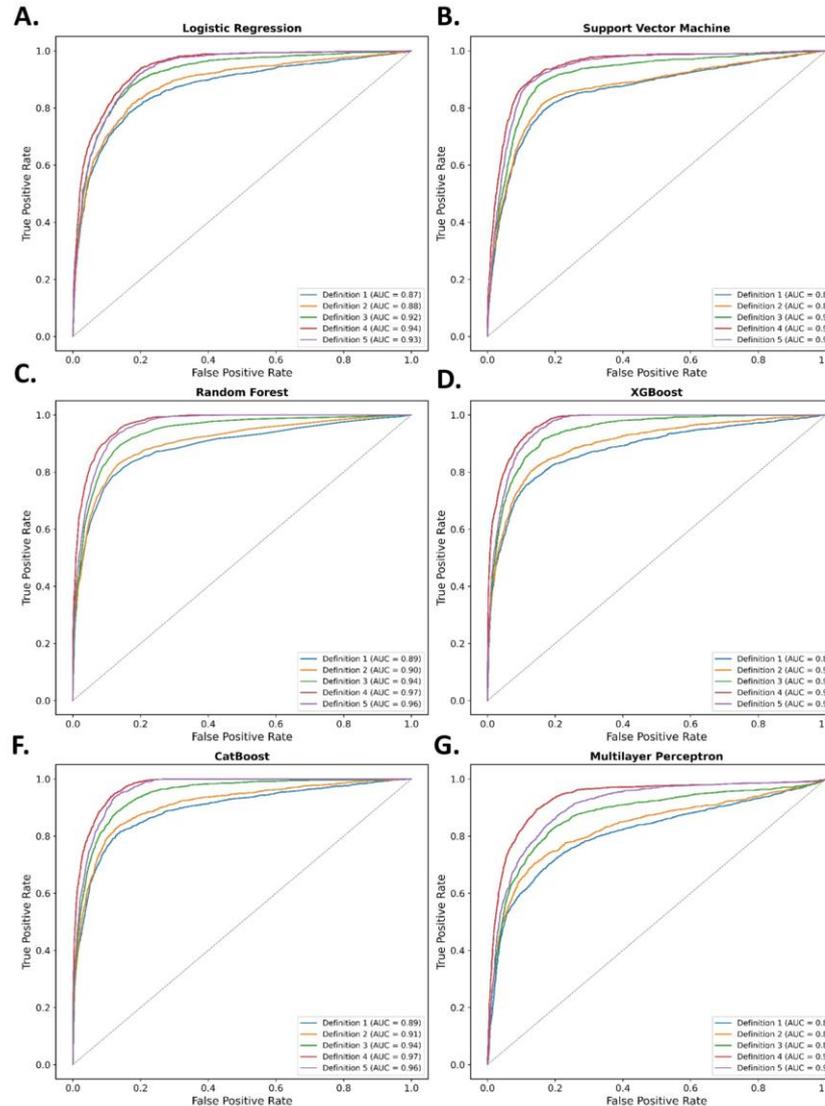
Définition 1: 0.87
Définition 2: 0.88
Définition 3: 0.92
Définition 4: 0.94
Définition 5: 0.93

Random forrest

Définition 1: 0.89
Définition 2: 0.90
Définition 3: 0.94
Définition 4: 0.97
Définition 5: 0.96

CatBoost

Définition 1: 0.89
Définition 2: 0.91
Définition 3: 0.94
Définition 4: 0.97
Définition 5: 0.96



Support Vector Machine

Définition 1: 0.86
Définition 2: 0.87
Définition 3: 0.91
Définition 4: 0.94
Définition 5: 0.93

XGBoost

Définition 1: 0.88
Définition 2: 0.90
Définition 3: 0.94
Définition 4: 0.97
Définition 5: 0.96

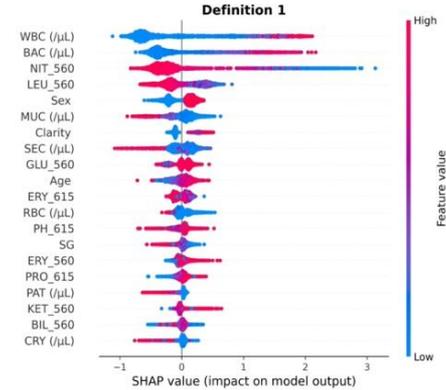
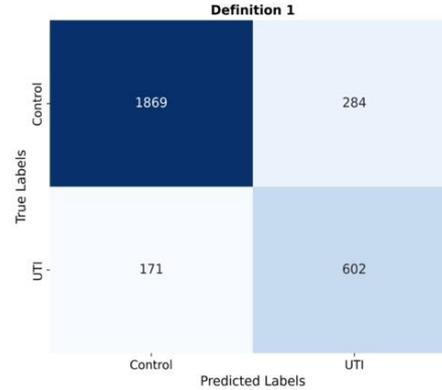
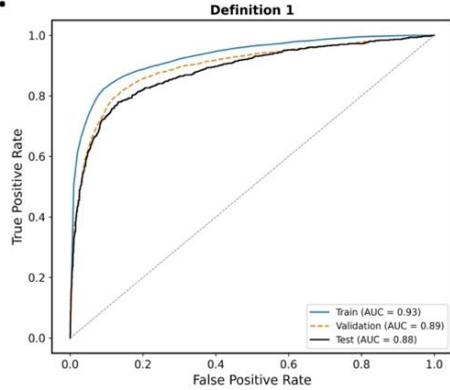
Multilayer Perceptron

Définition 1: 0.81
Définition 2: 0.83
Définition 3: 0.87
Définition 4: 0.93
Définition 5: 0.90

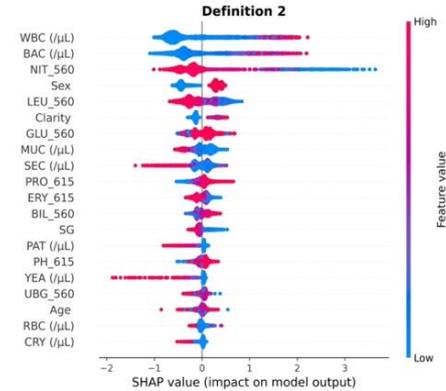
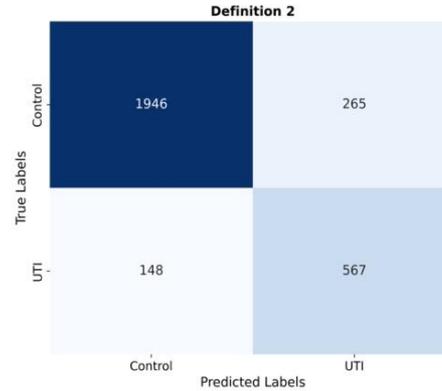
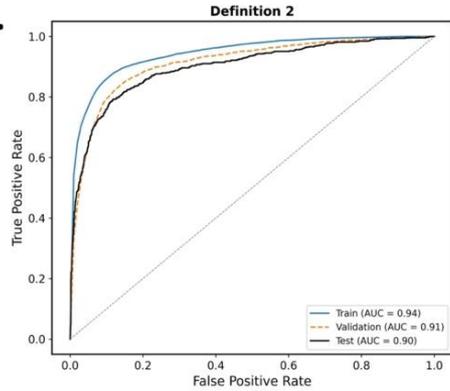
Table 1. Performance metrics of the CatBoost classifiers for the 7-fold stratified cross-validation procedure and predictions on the test set.

Model	Optimal threshold	Dataset	Sensitivity (% CI)	Specificity (% CI)	PPV (% CI)	NPV (% CI)	Accuracy (% CI)	AUC (% CI)
Definition 1	0.2575	Validation	82.2 (77.4-86.4)	86.5 (81.5-89.5)	68.8 (61.4-73.0)	93.1 (91.7-94.6)	85.3 (81.7-87.4)	89.5 (87.5-91.4)
		Test	77.9 (74.9-80.7)	86.8 (85.3-88.2)	68.0 (65.0-70.9)	91.6 (90.4-92.7)	84.4 (83.2-85.7)	88.2 (86.7-89.7)
Definition 2	0.2422	Validation	84.7 (81.5-90.0)	86.8 (84.5-90.4)	67.6 (64.2-73.8)	94.6 (93.5-96.4)	86.3 (84.4-88.8)	91.3 (89.9-92.7)
		Test	79.3 (76.3-82.1)	88.0 (86.7-89.3)	68.1 (65.2-71.2)	93.0 (91.8-94.0)	85.9 (84.7-87.1)	90.0 (88.5-91.3)
Definition 3	0.2318	Validation	90.2 (87.4-93.7)	87.3 (84.5-91.1)	67.9 (63.2-74.3)	96.8 (96.0-97.9)	88.0 (85.8-90.3)	94.5 (93.5-95.2)
		Test	87.9 (85.3-90.3)	87.8 (86.4-89.1)	67.9 (64.9-71.0)	96.1 (95.2-96.9)	87.8 (86.6-89.0)	94.7 (93.8-95.5)
Definition 4	0.2459	Validation	94.9 (92.7-95.9)	88.9 (87.7-90.1)	71.2 (69.2-73.0)	98.4 (97.7-98.7)	90.3 (89.5-91.1)	97.3 (96.7-97.7)
		Test	90.5 (88.4-92.6)	90.7 (89.5-91.8)	73.7 (70.7-76.7)	97.1 (96.4-97.7)	90.7 (89.6-91.7)	97.2 (96.7-97.6)
Definition 5	0.2333	Validation	94.5 (90.1-98.0)	87.4 (83.6-90.8)	66.5 (60.6-72.6)	98.4 (97.1-99.4)	88.9 (86.3-91.3)	96.3 (95.2-97.0)
		Test	91.7 (89.4-93.9)	90.0 (88.8-91.2)	70.6 (67.5-73.6)	97.7 (97.0-98.2)	90.4 (89.3-91.4)	96.8 (96.2-97.3)

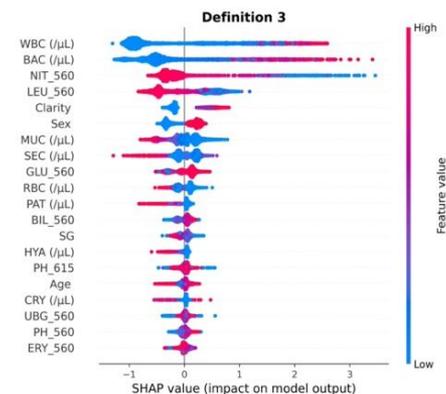
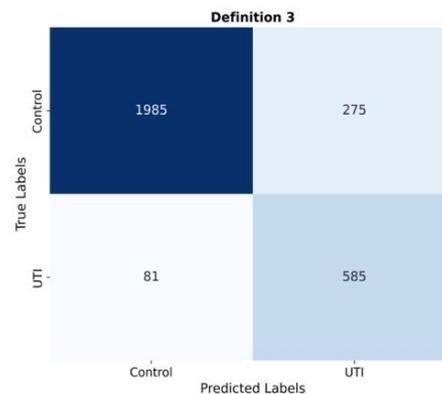
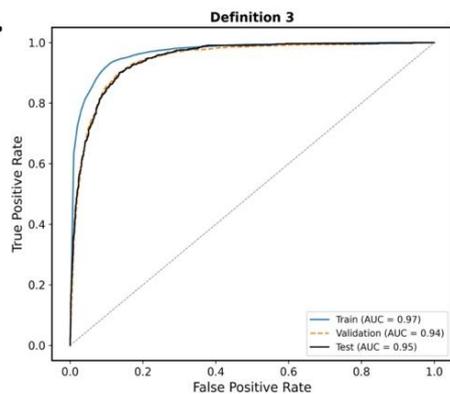
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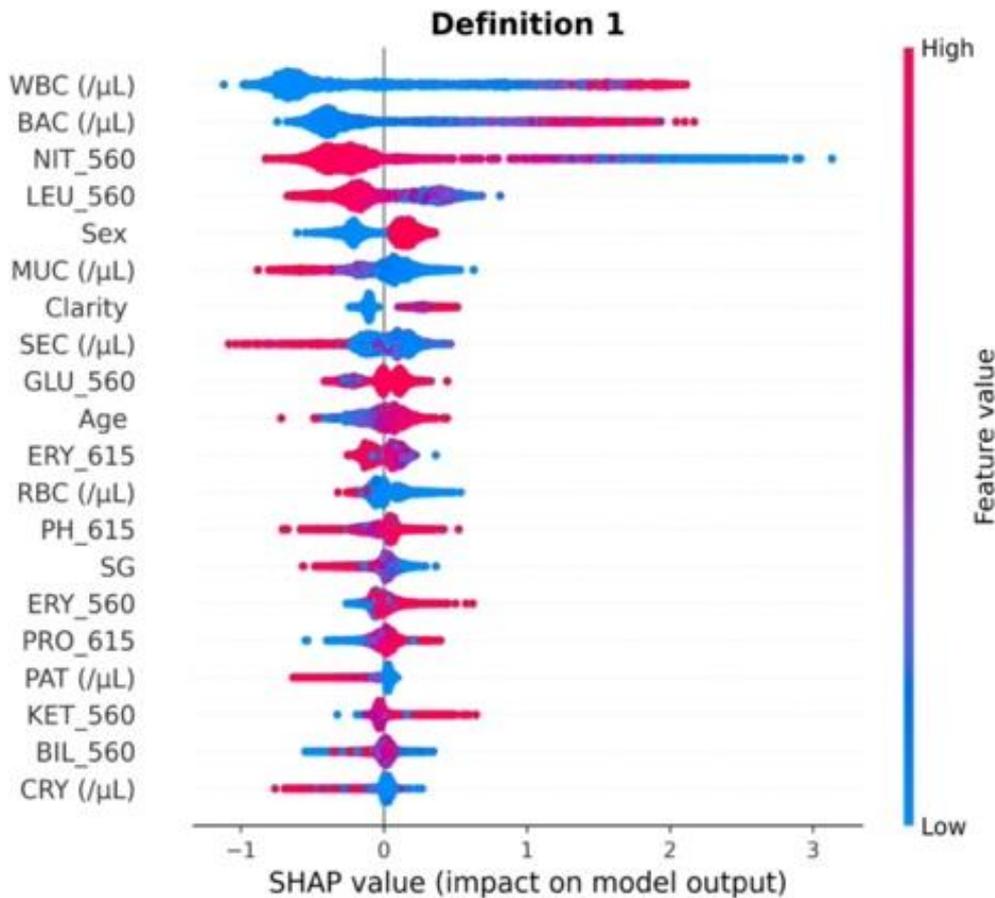


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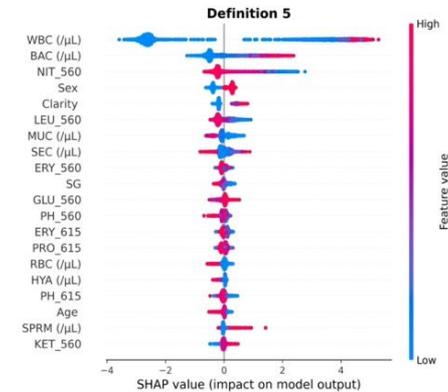
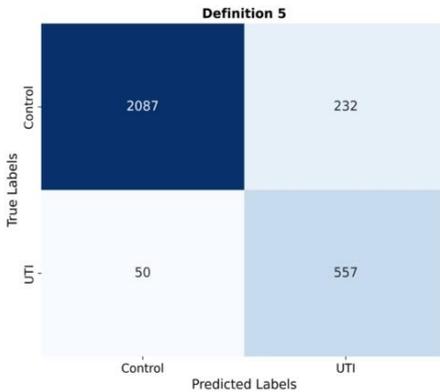
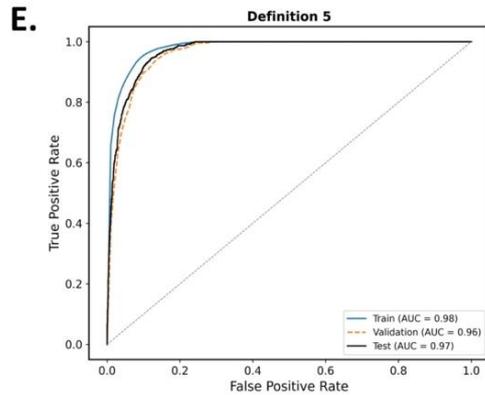
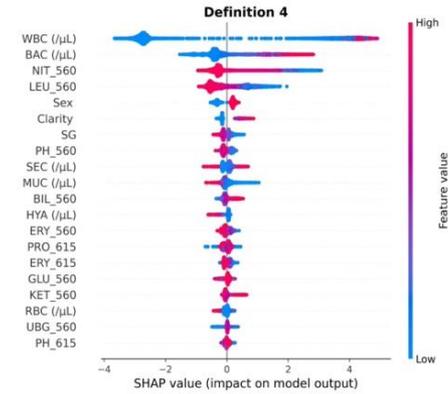
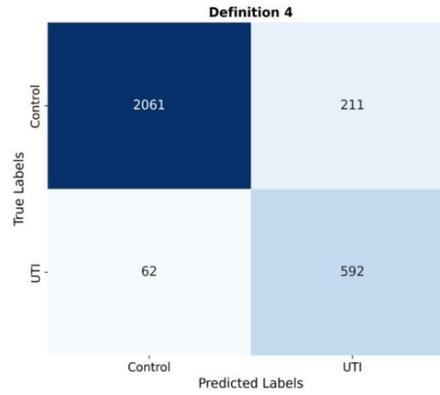
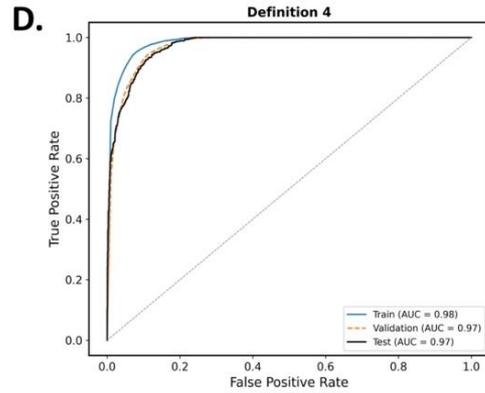
C.





- **Beeswarm plot:**

- SHAP = SHapley Additive exPlanations
- Contribution de chaque variable dans le modèle
- Impact élevé = en haut
- Bleu = valeur élevée
- Rouge = valeur faible

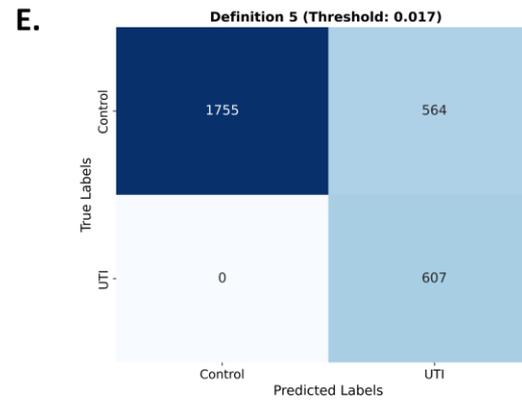
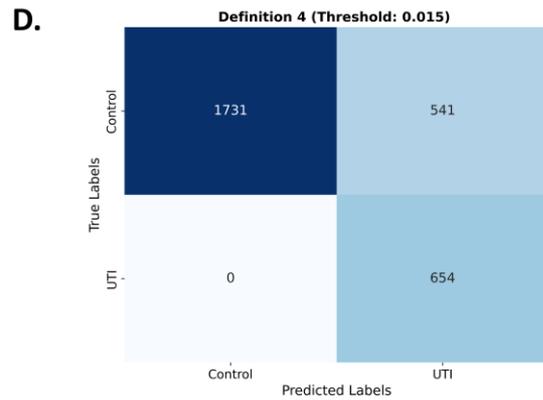
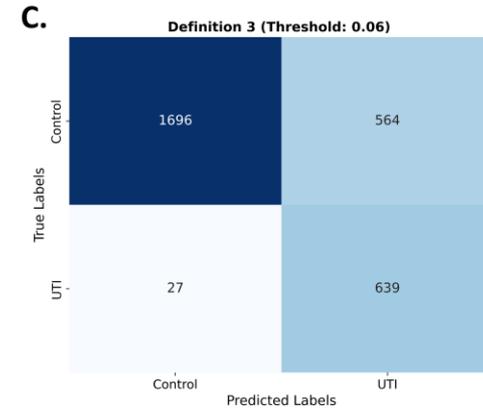
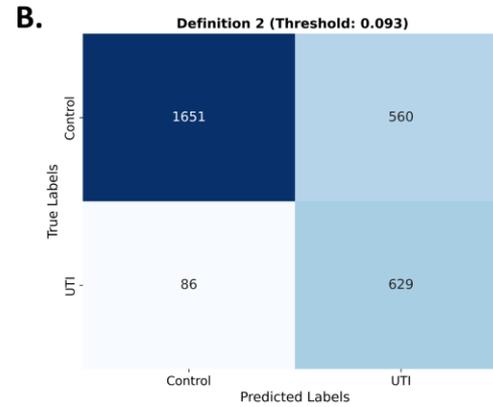
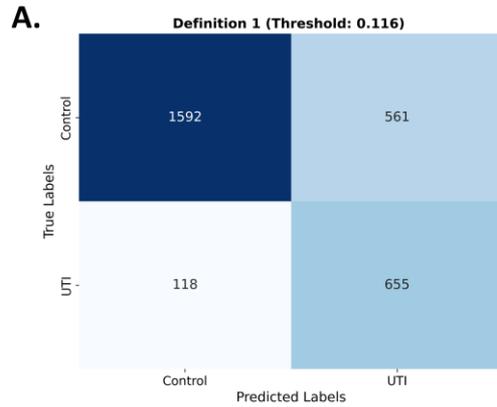


Statistiques simples versus avancées

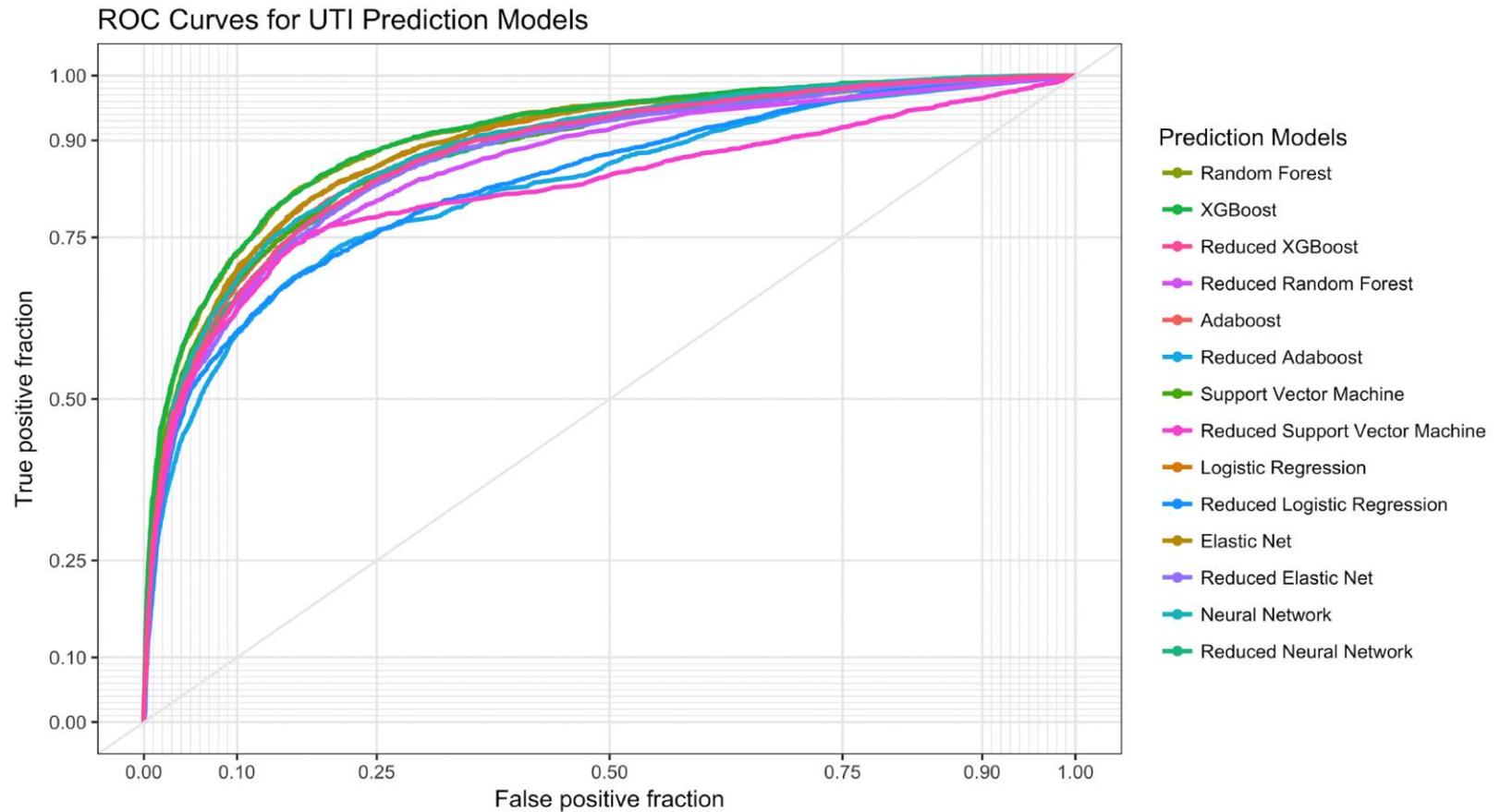
Accuracy (%)	Définition 1	Définition 2	Définition 3	Définition 4	Définition 5
LEU_560	60.7	59.7	60.1	61.6	60.0
NIT_560	81.1	82.6	84.3	83.8	84.0
LEU_560 + NIT_560	60.8	59.9	60.2	61.0	59.4
GB	63.6	62.6	63.3	65.2	63.6
ML (validation)*	85.3	86.3	88.0	90.3	88.9
ML (test)*	84.4	85.9	87.8	90.7	90.4

*CatBoost model

Accuracy (%)	Couleur
50-60	Red
60-70	Orange
70-80	Yellow
80-90	Light Green
90-100	Dark Green



Model	Sensitivity-tuned threshold	Sensitivity (% CI)	Specificity (% CI)	PPV (% CI)	NPV (% CI)	Accuracy (% CI)
Definition 1	0.116	84.6 (82.1-87.0)	74.0 (72.1-75.6)	53.9 (51.0-56.5)	93.1 (91.9-94.2)	76.8 (75.3-78.2)
Definition 2	0.093	88.0 (85.5-90.3)	74.6 (72.8-76.5)	53.0 (50.0-55.9)	95.0 (94.0-95.6)	77.9 (76.4-79.3)
Definition 3	0.06	95.6 (94.4-97.4)	75.1 (73.2-76.8)	53.2 (50.5-55.8)	98.4 (97.8-99.0)	79.8 (78.4-81.2)
Definition 4	0.015	100 (100-100)	76.2 (74.4-78.0)	54.7 (52.0-57.5)	100 (100-100)	81.5 (80.1-82.9)
Definition 5	0.017	100 (100-100)	75.7 (73.8-77.5)	51.8 (48.8-54.7)	100 (100-100)	80.7 (79.2-82.1)



- 212 versus 10 variables
- Outcome: $>10^4$ CFU/HPF (2) flore mixte si *E. coli* présent
- Modèle comparé au jugement clinique (AB pour IU ou catégorisation IU)

Table 4. Test characteristics of UTI prediction models on validation data*.

Models	AUC (95%CI)	Sensitivity (95% CI)	Specificity (95% CI)	+LR (95% CI)	-LR (95% CI)	Accuracy (95% CI)	P-value
XGBoost	.904(.898-.910)	61.7(60.0–63.3)	94.9 (94.5–95.3)	12.0(11.1–13.0)	.404(.387-.421)	87.5 (87.0–88.0)	NA
Random Forest	.902(.896-.908)	57.3(55.6–58.9)	96.0 (95.6–96.3)	14.3(13.0–15.6)	.445(.428-.462)	87.4 (86.9–87.9)	0.58
Adaboost	.880(.874-.887)	62.2(60.6–63.8)	92.3(91.8–92.7)	8.06(7.54–8.61)	.409(.392-.427)	85.6(85.1–86.2)	< .001
Support Vector Machine	.878(.871-.884)	49.6(47.9–51.2)	96.8(96.4–97.1)	15.3(13.8–16.9)	.521(.504-.538)	86.3(85.7–86.8)	< .001
ElasticNet	.892(.885-.898)	56.8(55.2–58.4)	94.9(94.5–95.2)	11.1(10.2–12.0)	.455(.438-.473)	86.4(85.9–87.0)	< .001
Logistic Regression	.891 (.884-.897)	57.5(55.8–59.1)	94.7(94.3–95.1)	10.9(10.0–11.8)	.449(.432-.466)	86.4(85.9–87.0)	< .001
Neural Network	.884 (.878-.890)	54.6(52.9–56.2)	95.3(95.0–95.7)	11.7(10.8–12.8)	.476(.460-.494)	86.3(85.8–86.8)	<.001
Reduced XGBoost	.877(.871-.884)	54.7(53.0–56.3)	94.7(94.3–95.1)	10.4(9.6–11.3)	.479(.462-.496)	85.9(85.3–86.4)	< .001
Reduced Random Forest	.861(.853-.868)	54.8(53.1–56.4)	94.3(93.9–94.7)	9.66(8.94–10.4)	.479(.462-.497)	85.5(85.0–86.1)	< .001
Reduced Adaboost	.826(.817-.834)	61.9(60.3–63.5)	88.8(88.2–89.3)	5.50(5.21–5.81)	.429(.412-.448)	82.8(82.2–83.3)	< .001
Reduced Support Vector Machine	.822(.813-.832)	49.4(47.8–51.1)	95.8(95.4–96.1)	11.7(10.7–12.9)	.528(.511-.546)	85.5(84.9–86.0)	< .001
Reduced Elastic Net	.870(.863-.877)	52.4(50.7–54.1)	95.2(94.8–95.5)	10.9(9.99–11.8)	.500(.482-.571)	85.7(85.1–86.2)	< .001
Reduced Logistic Regression	.870(.863-.877)	53.3(51.6–54.9)	94.8(94.4–95.2)	10.3(9.52–11.2)	.492(.476-.510)	85.6(85.0–86.2)	< .001
Reduced Neural Network	.873(.867-.881)	54.0(52.3–55.6)	95.0(94.6–95.4)	10.9(10.0–11.8)	.485(.468-.502)	85.9(85.4–86.5)	< .001

- XGBoost = meilleur modèle (full and reduced, $p > 0.05$)
- Modèle ML versus jugement clinique
 - FP ↓ 29.6-33.3%
 - FN ↓ 33.2-38.7%

Table 2. Selected variables for reduced models.

<u>Variable</u>	<u>References</u>
Age	[3, 21]
Gender	[3, 6, 21]
UA Leukocytes	[3, 6, 10, 12, 21]
UA Nitrites	[3, 6, 10, 12, 21]
UA WBC	[3, 6, 10, 12, 21]
UA Bacteria	[3, 6, 10, 12, 21]
UA Blood	[3, 6, 10, 12, 21]
UA Epithelial Cells	[3, 6, 10, 12, 21]
History of UTI	[3, 6, 21]
Dysuria	[3, 6, 21]

Table 2 Feature selection by recursive feature elimination using a Random Forest Classifier. Feature importance is shown as well as the individual AUC score

	RFE Ranking	RF Feature Importance	Individual AUC ^a
WBC count	1	0.30	0.82
Bacterial count	1	0.30	0.71
Age	1	0.12	0.63
Epithelial cell count	1	0.07	0.49
RBC count	1	0.06	0.56
# of positive cultures to date	1	0.03	0.60
Pyuria, no RBCs	1	0.02	0.57
Pregnant	1	0.02	0.57
Inpatient	1	0.01	0.53
Gender	1	0.01	0.53
Persistent/recurrent infection	1	0.01	0.55
# of positive cultures month prior	1	0.009	0.53
Positive for nitrates	1	0.008	0.52
Renal inpatient/outpatient	1	0.005	0.50
Pre-operative patient	1	0.004	0.51
Acute kidney disease	1	0.003	0.50
Immunocompromised	2	0.002	0.50
# of positive cultures week prior	3	0.002	0.51
Multiple Sclerosis	4	0.001	0.50
Offensive smell	5	0.0007	0.50
Haematuria, no WBCs	6	0.0001	0.50

^aIndividual AUC score is calculated from a Logistic Regression classifier, where the feature in question is the sole independent variable

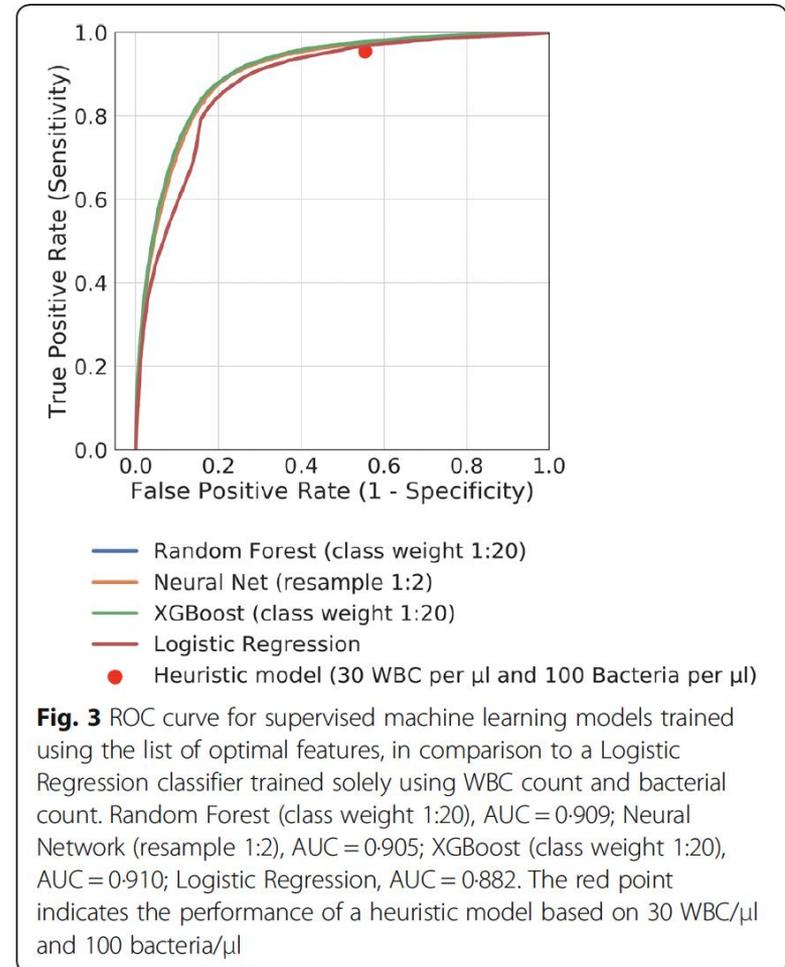


Fig. 3 ROC curve for supervised machine learning models trained using the list of optimal features, in comparison to a Logistic Regression classifier trained solely using WBC count and bacterial count. Random Forest (class weight 1:20), AUC = 0.909; Neural Network (resample 1:2), AUC = 0.905; XGBoost (class weight 1:20), AUC = 0.910; Logistic Regression, AUC = 0.882. The red point indicates the performance of a heuristic model based on 30 WBC/ μ l and 100 bacteria/ μ l

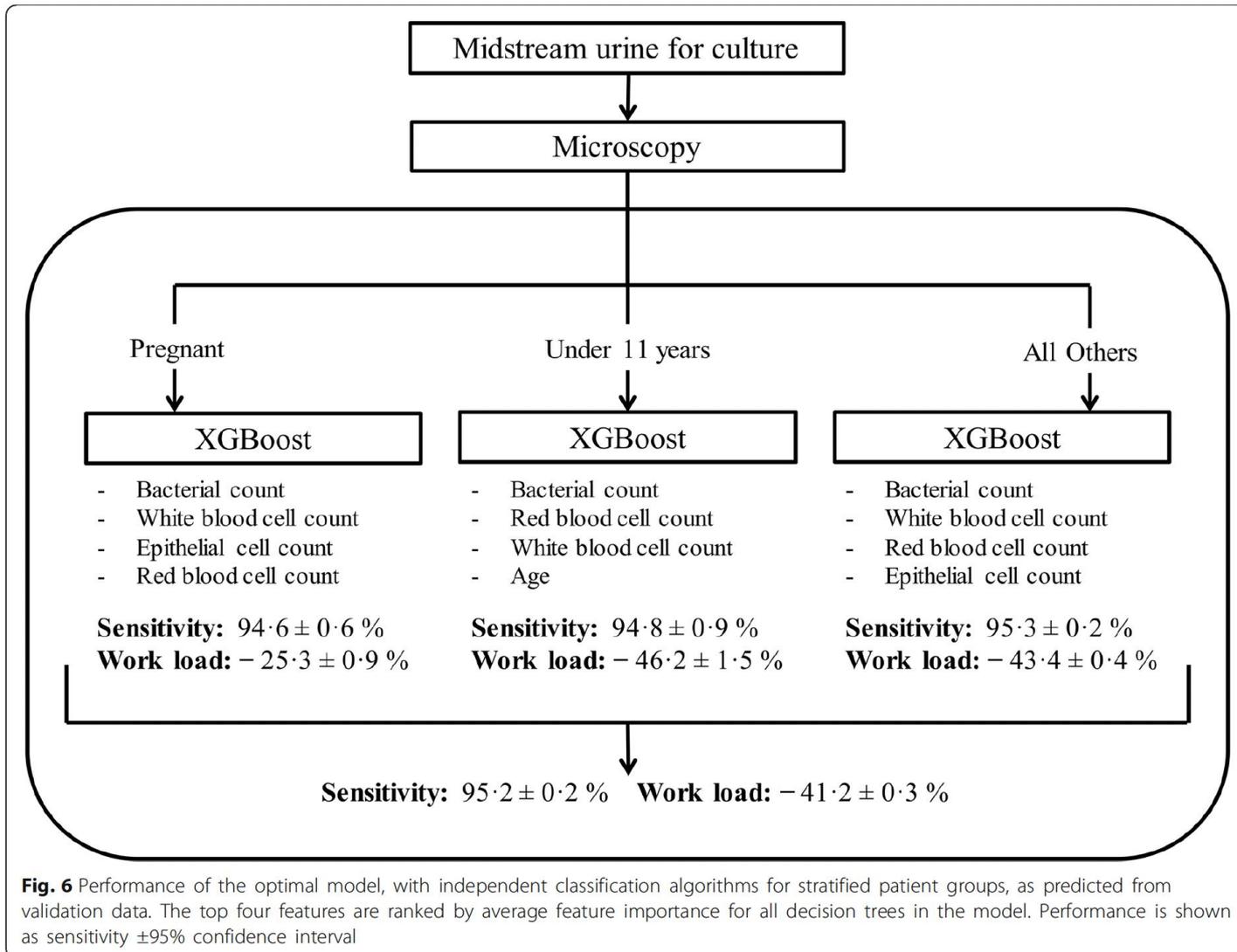
Table 3 Comparison of performance for heuristic and machine learning models tested on holdout data

Model Name	AUC Score	Accuracy (%)	<i>p</i> -value**	PPV	NPV	Sensitivity (%)			Specificity (%)	Relative Workload Reduction (%)
						All Patients	Pregnant	Children < 11 Yrs		
Heuristic model (30 WBC/ μ l or 100 bacteria/ μ l)		63.92	NA	42.73 [± 0.51]	97.01 [±0.28]	95.70 [± 0.15]	85.9 [± 0.72]	91.5 [± 0.92]	52.10 [± 0.36]	39.06 [± 0.38]
Random Forest (Class weight - 1:20)	0.908	71.96	< 0.001	40.47 [± 0.54]	97.67 [± 0.25]	95.95 [± 0.23]	70.5 [± 2.14]	89.8 [± 1.49]	63.40 [± 0.54]	47.58 [± 0.39]
Neural Network	0.906	85.00	< 0.001	71.70 [± 0.46]	90.18 [± 0.50]	74.03 [± 0.64]	27.6 [± 5.74]	69.3 [± 3.38]	89.09 [± 0.29]	71.98 [± 0.35]
Neural Network (with resampling*)	0.904	79.35	< 0.001	57.66 [± 0.74]	95.54 [± 0.19]	90.60 [± 0.35]	56.6 [± 3.43]	84.8 [± 2.04]	75.16 [± 0.44]	57.33 [± 0.38]
XGBoost (Class weight - 1:20)	0.910	65.68	< 0.001	44.05 [± 0.74]	97.77 [± 0.13]	96.70 [± 0.18]	77.1 [± 1.65]	93.1 [± 1.13]	54.14 [± 0.61]	40.36 [± 0.38]

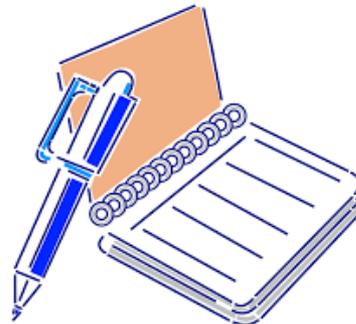
[95% Confidence Interval]

*Resampling (without replacement) at a ratio of 2:1 for positive samples to offset class imbalance

** *p*-values obtained by comparison to heuristic model by McNemar test



- Remarques:
 - Pas d'inclusion des symptômes cliniques
 - « *The diagnosis of UTI by clinical criteria alone has an error rate of approximately **33%**** »
 - Variabilité inter-individuelle/subjectivité
 - Prise d'antibiotiques? Problèmes pré-analytique?
 - Etude « *real-life* » sans exclusion de 0 à 104 ans
 - Trop dur à implémenter en routine?
 - Modèles compliqués. Logiciel/Middleware/application
 - Utiliser un modèle plus simple dans un LIS?



- GraphPad Prism Mac 10.2.0 (335)

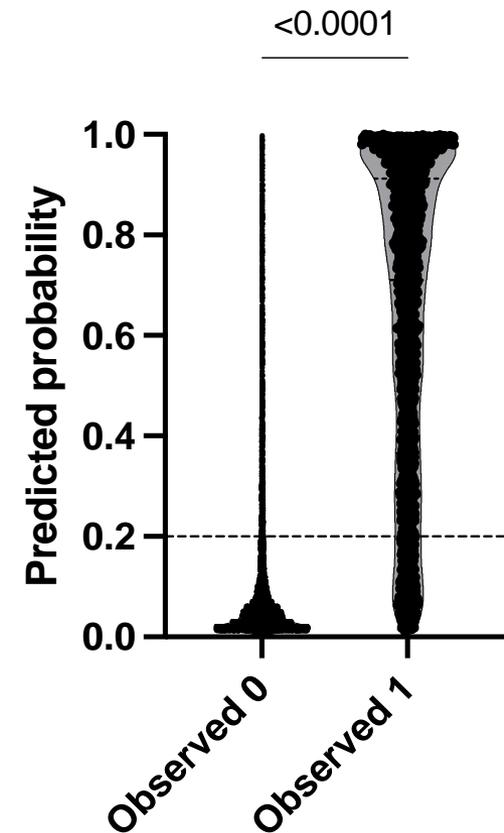


- Equation

$$\hat{p} = \frac{\exp(b_0 + b_1 X_1 + b_2 X_2 + \dots + b_p X_p)}{1 + \exp(b_0 + b_1 X_1 + b_2 X_2 + \dots + b_p X_p)}$$

- Calcul d'une probabilité de **0 à 1** pour chaque échantillon
- Probabilité comparée à l'outcome **0** ou **1**

Predicted vs Observed



- GraphPad Prism Mac 10.2.0 (335)  Prism

Area under the ROC curve				
Area	0,9284			
Std. Error	0,003044			
95% confidence interval	0,9225 to 0,9344			
P value	<0,0001			
Classification table	Predicted 0	Predicted 1	Total	% Correctly classified
Observed 0	6474	1060	7534	85,93
Observed 1	311	1908	2219	85,98
Total	6785	2968	9753	85,94
Negative predictive power (%)	95,42			
Positive predictive power (%)	64,29			
Classification cutoff	0,2			



Tigette/sédiment

Cas #1

- Femme
- 37 ans
- NIT +
- LEU +++
- GB 172/μL

Cas #2

- Femme
- 41 ans
- NIT +
- LEU +++
- GB 160/μL

Microbiologie

Culture négative

***E. coli* CFU >10⁵/mL**

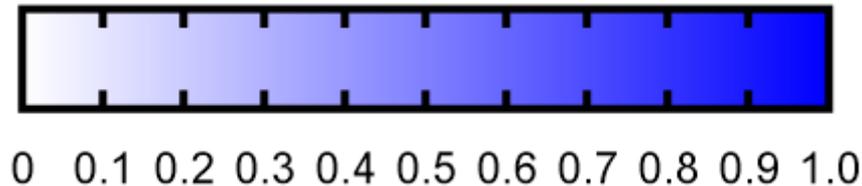
Prédiction machine learning (0 à 1)

**Score 0.003
→ négatif**

**Score 0.961
→ positif**

- **Intelligence artificielle comme aide au diagnostic**

- Possibilité d'exclure rapidement une infection urinaire sur base d'un score de prédiction



0-0.2: Très peu probable

0.2-0.5: Peu probable

0.5-0.8: Probable

0.8-1.0: Très probable

- Besoin d'une définition harmonisée
- Besoin d'une validation externe
- Evaluation prospective du réel apport (AB)