

## Supplementary Materials

# FTIR Spectroscopic Imaging Supports Urine Cytology for Classification of Low- and High-Grade Bladder Carcinoma

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**Table S1.** Normal, LG and HG BC cell features according to the Paris system.

Cell features	Normal	LG BC	HG BC
<b>Nuclear to cytoplasmic ratio (N/C)</b>	<0,5	>0.5	>0,7
<b>Nuclear hyperchromasia</b>	-	++	+++
<b>Irregular nuclear membrane (chromatinic rim or nuclear contour)</b>	-	++	+++
<b>Irregular, coarse, clumped chromatin</b>	-	++	+++
<b>Pleomorphism</b>	-	++	+++
<b>Variation in size and shapes</b>	-	++	+++
<b>Scant, pale or dense cytoplasm</b>	-	+	+++
<b>Prominent nucleoli</b>	-	++	+++
<b>Mitoses</b>	-	++	+++
<b>Necrotic debris</b>	-	++	+++
<b>Inflammation</b>	-	+	+++
<b>Cytoplasmic homogeneity</b>	+++	+	-

BC – Bladder urothelial carcinoma; LG – low grade; HG – high grade, presence of the feature: + - low, ++ - moderate, +++ - severe.

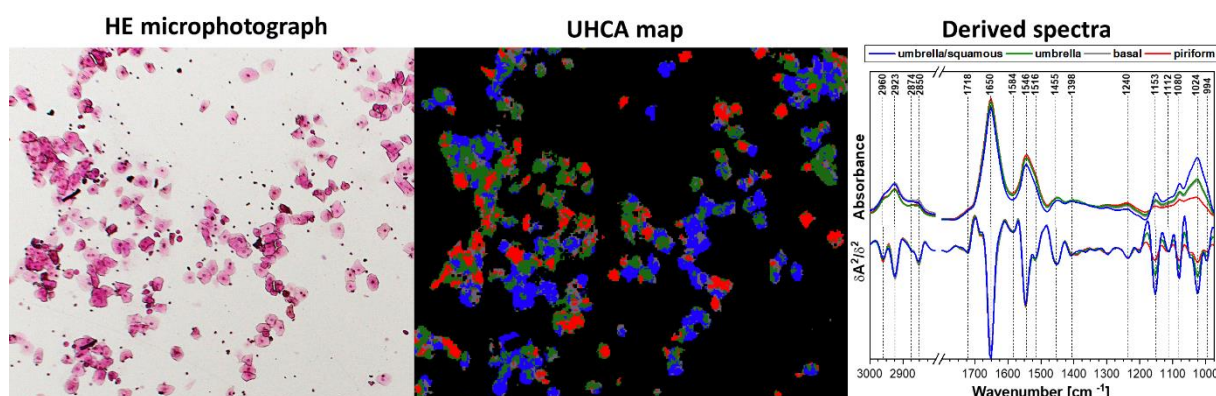
**Table S2.** Positions of IR bands observed in FTIR spectra of urine sediment with their assignment to vibrational modes and biomolecules [11,20-25].

Band [cm <sup>-1</sup> ]	Assignment
994	RNA and glycogen
1024	Carbohydrates and glycolipids (i.e. glycogen and sphingomyelin)
1052	DNA, polysaccharides, cholesterol, glycoproteins, glycolipids; $\nu(\text{C}-\text{O}-\text{O}-\text{C})$
1080	DNA, RNA and other phosphate containing molecules, glycogen; $\nu(\text{PO}_2^-)$ , $\nu(\text{CC})$
1112	Polysaccharides, ribose, lactate; $\nu(\text{C}-\text{O})$ , $\nu(\text{CC}-\text{OC})$
1153	Carbohydrates (predominantly glycogen); $\nu_{\text{as}}(\text{CO}-\text{O}-\text{C})$
1240	Nucleic acids, phospholipids, phosphoproteins; $\nu_{\text{as}}(\text{PO}_2^-)$
1398	Lipids, free amino acids; $\nu_{\text{as}}(\text{COO}^-)$
1455	Proteins, lipids, carbohydrates, lipopolysaccharides, phospholipids; $\delta(\text{CH}_2, \text{CH}_3)$
1516	Proteins; tyrosine ring $\nu(\text{CC})$
1546	Amid II and lipids
1584	Lipids, proteins, nucleic acids, lipopolysaccharides; $\nu_{\text{as}}(\text{COO}^-)$ , $\nu(\text{C}=\text{N}, \text{C}=\text{C})$
1623	Proteins (parallel $\beta$ -turns) ; $\text{C}=\text{O}$ , $\text{N}-\text{H}$
1652	Amide I of $\alpha$ -helix; $\text{C}=\text{O}$ , $\text{N}-\text{H}$
1718	Fatty acids and nucleic acids: T-A, G-C (Hoogsteen third strand binding) ; $\nu(\text{C}=\text{O})$
1740	Ester group in triglycerides, cholesterol and phospholipids, lipopolysaccharides; $\nu_s(\text{C}=\text{O})$
2850	Lipids and proteins; $\nu_{\text{as}}(-\text{CH}_2)$
2874	Phospholipids, esters
2923	Lipids and proteins; $\nu_{\text{as}}(-\text{CH}_2)$
2960	Lipids and proteins; $\nu_{\text{as}}(-\text{CH}_3)$

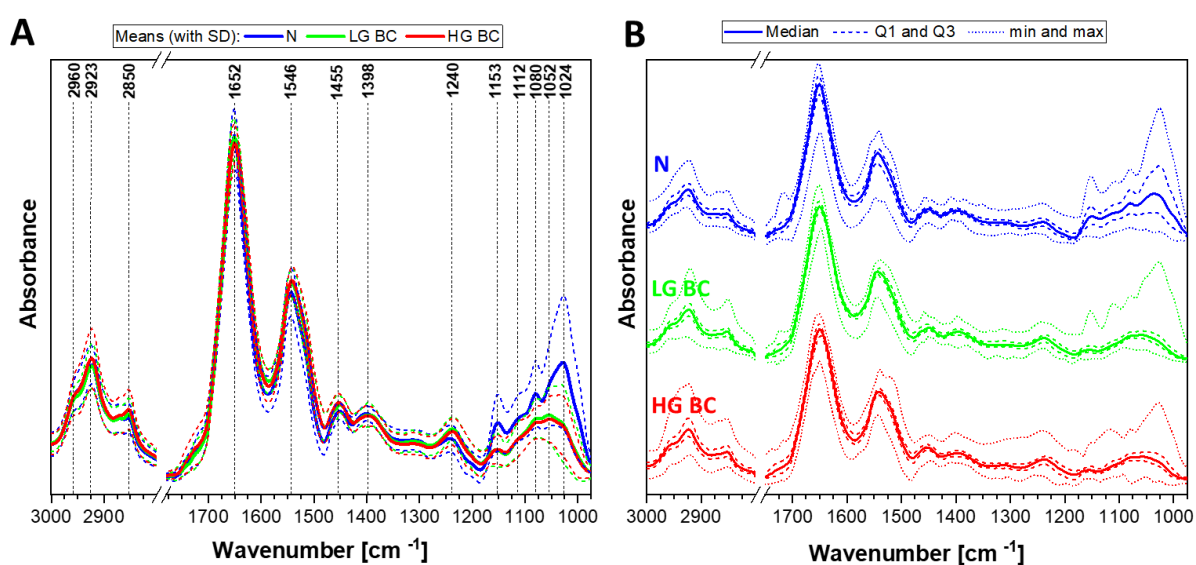
$\nu$  – stretching mode,  $\nu_{\text{as}}$  – antisymmetric,  $\nu_s$  – symmetric;  $\delta$  – in-plane deformations (bending);  $\gamma$  – out-of-plane deformations,  $\tau$  – twisting, (+) – intense signal from band (of whole cells), (+/-) – average intense signal of band, (-/+) – weak visible band, (-) – not observed.

**Table S3.** PLS-DA parameters for discrimination of LG and HG BC cytology from normal urothelial cells.

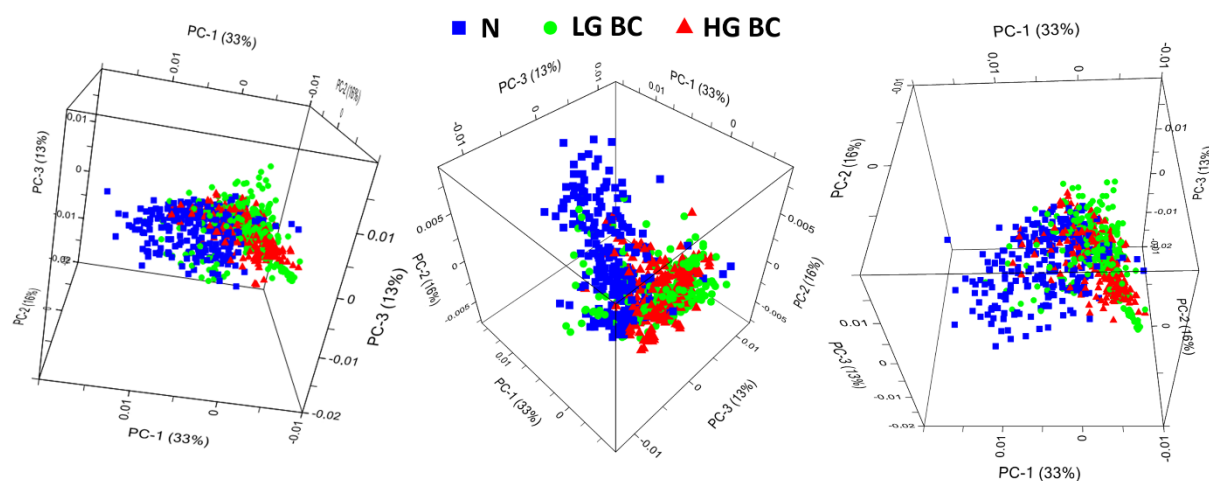
Classification	N vs LG	N vs HG
PLS parameters	RMSE / R <sup>2</sup>	RMSE / R <sup>2</sup>
Calibration	0.55 / 0.70	0.43 / 0.82
Validation	0.59 / 0.65	0.47 / 0.78
Prediction	0.55 / 0.70	0.47 / 0.78



**Figure S1.** The comparison of HE staining and UHCA-discriminated cells of normal urothelial cells showing their assignment; on right, mean FTIR spectra of UHCA classes (the colors of the spectra correspond to the colors of classes in the UHCA map).



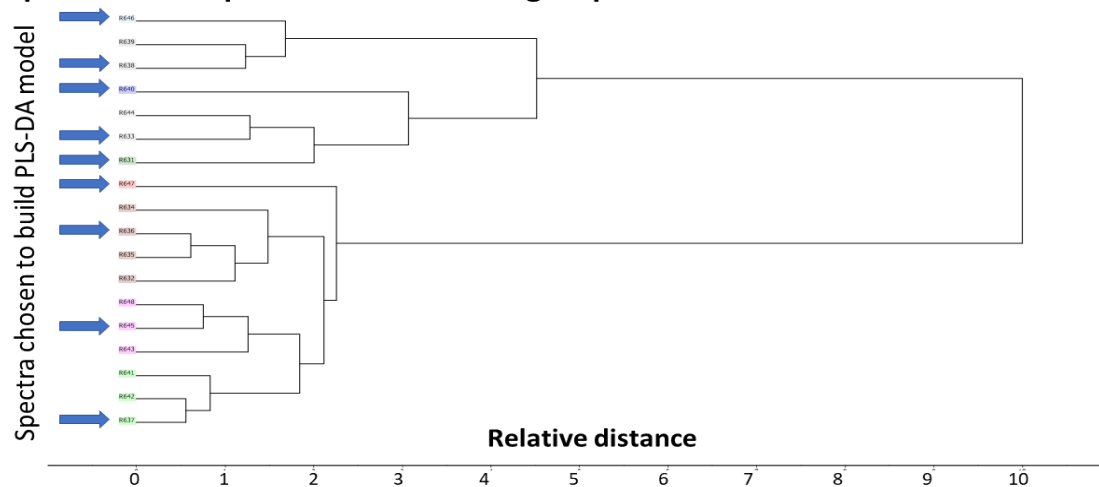
**Figure S2.** A. Averaged absorbance FTIR spectra (with SD marked as dashed lines). B. Median, 1<sup>st</sup> and 3<sup>rd</sup> quantile and min and max plots. All spectra were preprocessed after baseline correction and vector normalization.



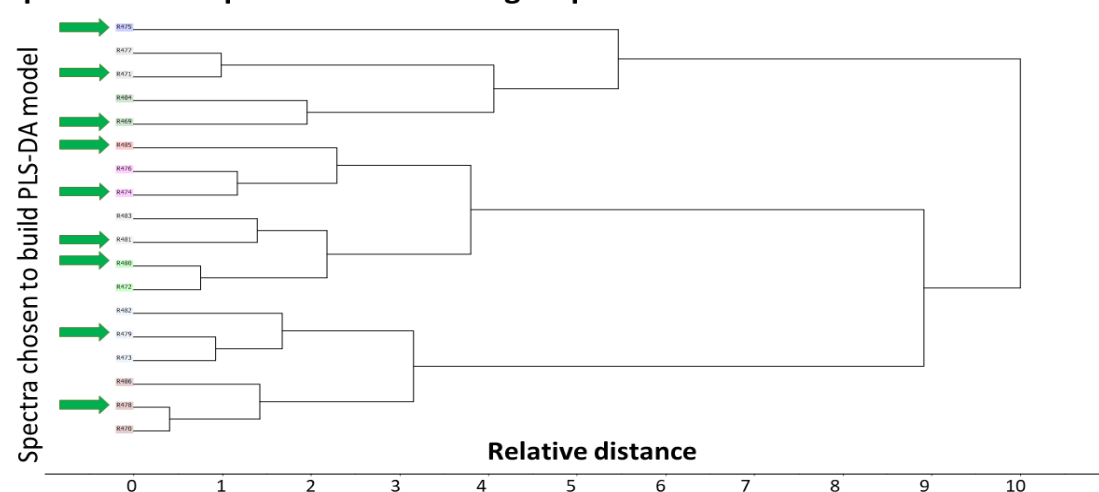
**Figure S3.** 3D PCA scores plots from Fig. 2B showed in different projections.

## Cluster Analysis (Ward's method using Squared Euclidean distance)

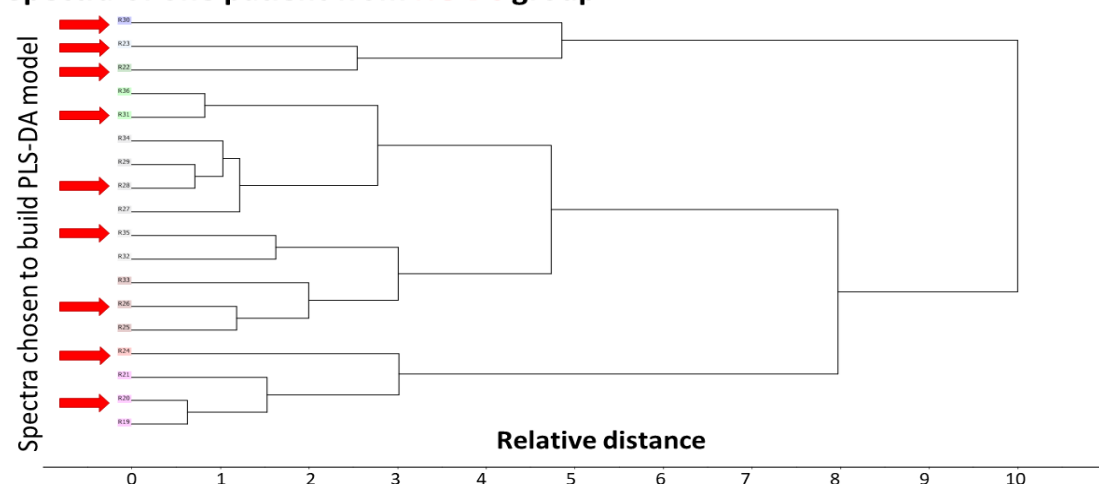
### Spectra of one patient from **normal** group



### Spectra of one patient from **LG BC** group



### Spectra of one patient from **HG BC** group



**Figure S4.** Example of cluster analysis from one of N, LG and HG BC patients. Spectra from the most different branches of Cluster Analysis were chosen to build the model, were marked with arrows and they represent 9 out of 18 spectra from the patient.