

Supplementary Table 4. Random forest classifier analysis of urine samples by groups

Act. group	Pred. group (train set)				Correctly	Pred. group (test set)				Correctly
	Control	LG	HG	GC	classified	Control	LG	HG	GC	classified
Control	72	0	0	0	100.0%	28	0	0	0	100.0%
LG	6	43	11	9	62.3%	2	20	2	7	64.5%
HG	3	5	56	4	82.4%	4	1	25	2	78.1%
GC	9	5	5	52	73.2%	3	1	3	22	75.9%
Overall correct class. Rate					79.6%					79.2%

The random forest classifier was performed using *Cutibacterium acnes*, *Ralstonia insidiosa*, and *Pseudomonas antarctica*, identified as group-discriminative by MaAsLin2 analysis of urine samples. To address sample imbalance, SMOTE was applied, generating a balanced dataset of 100 samples.

Pred., predicted group; Act., actual group; LG, low-grade dysplasia; HG, high-grade dysplasia; GC, gastric cancer.