

Figure S1: Flow diagram underlying the study selection process.

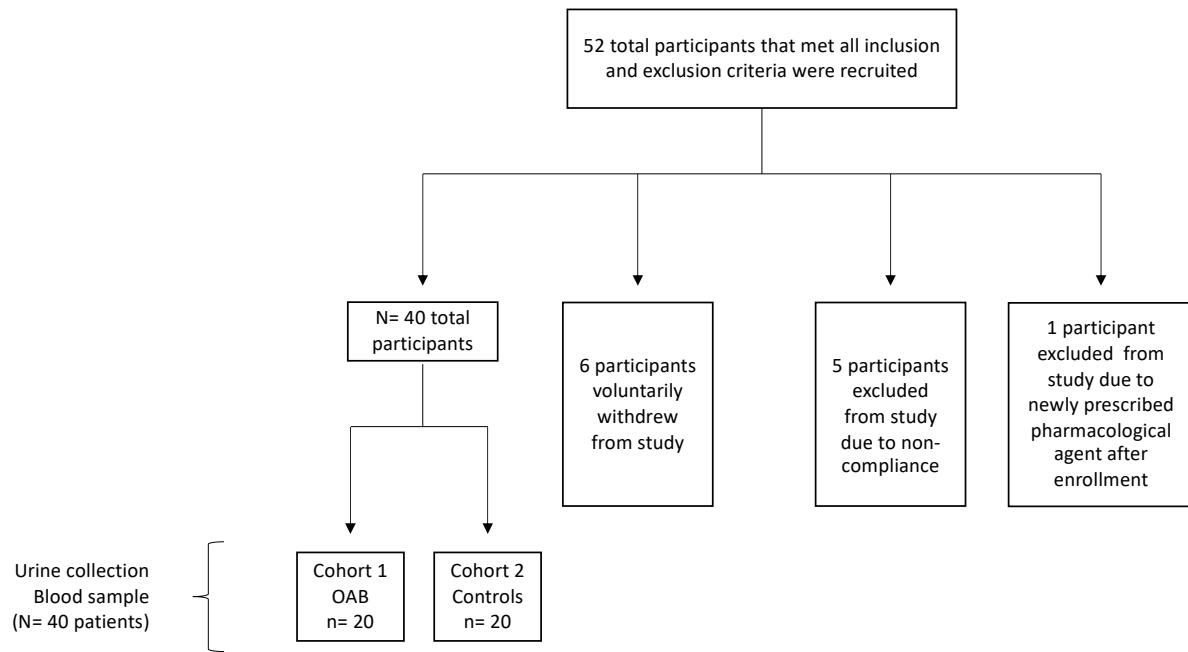


Table S1: qPCR miRNA information.

Sequence accession number	
<b>miR-26b-5p</b>	MIMAT0000083
<b>miR-26a-5p</b>	MIMAT0000082
<b>miR-10a-5p</b>	MIMAT0000253
<b>miR-491-5p</b>	MIMAT0002807
<b>miR-103a-3p</b>	MIMAT0009196
<b>miR-15b-5p</b>	MIMAT0000417
<b>miR-142-3p</b>	MIMAT0000433
<b>miR-202-3p</b>	MIMAT0002810
<b>miR-124-5p</b>	MIMAT0004591
<b>miR-152-5p</b>	MIMAT0026479
<b>snU6</b>	NR 004394.1

Table S2: Primer sequences used for qPCR

primers	
<b>miR-26b-5p</b>	ACACTCCAGCTGGGTTCAAGTAATTCAAGG
<b>miR-26a-5p</b>	ACACTCCAGCTGGGTTCAAGTAATCCAGG
<b>miR-10a-5p</b>	ACACTCCAGCTGGGTACCCTGTAGATCCG
<b>miR-491-5p</b>	ACACTCCAGCTGGGAGTGGGAACCCCT
<b>miR-103a-3p</b>	ACACTCCAGCTGGGAGCAGCATTGTACAG
<b>miR-15b-5p</b>	ACACTCCAGCTGGTAGCAGCACATCATG
<b>miR-142-3p</b>	ACACTCCAGCTGGGCATAAAGTAGAAAGC
<b>miR-202-3p</b>	ACACTCCAGCTGGGAGAGGTATAGGGCAT
<b>miR-124-5p</b>	ACACTCCAGCTGGCGTGTTCACAGCGGA
<b>miR-152-5p</b>	ACACTCCAGCTGGGCATAAAGTAGAAAGC
<b>universal primer</b>	CTCACAGTACGTTGGTATCCTTGTG
<b>snU6 forward</b>	CTCGCTTCGGCAGCACATATACT
<b>snU6 reverse</b>	ACGCTTCACGAATTGCGTGTC

Table S3: Correlation between miR-491-5p and p75<sup>ECD</sup> with symptom questionnaires' scores and voiding diary parameters:

miRNA	Variables	Correlation coefficient (r)	p value
miR-491-5p <sup>\$</sup>	OABSS	-0.405	<b>0.011</b>
	ICIQ-SF	-0.379	<b>0.019</b>
	IIQ-7	-0.494	<b>0.001</b>
p75 <sup>ECD</sup>	OABSS	-0.372	<b>0.018</b>
	ICIQ-SF	-0.539	<b>0.0001</b>
	IIQ-7	-0.372	<b>0.018</b>

The p value for Spearman correlation, only p <0.05 are reported as considered statistically significant.

<sup>\$</sup> Data on miR-491-5p were published in Metabolites (same special issue), Cammisotto et al (2022).