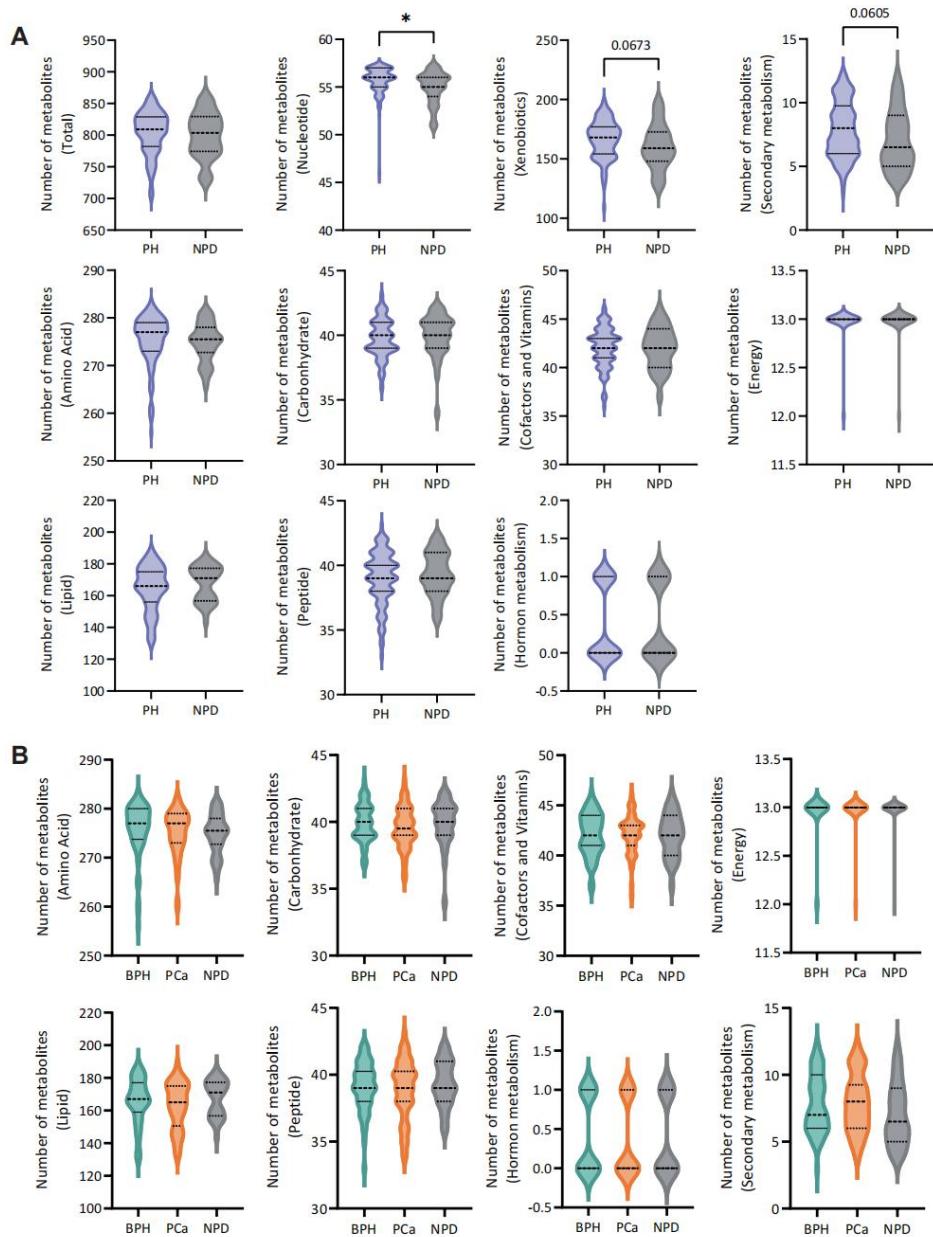


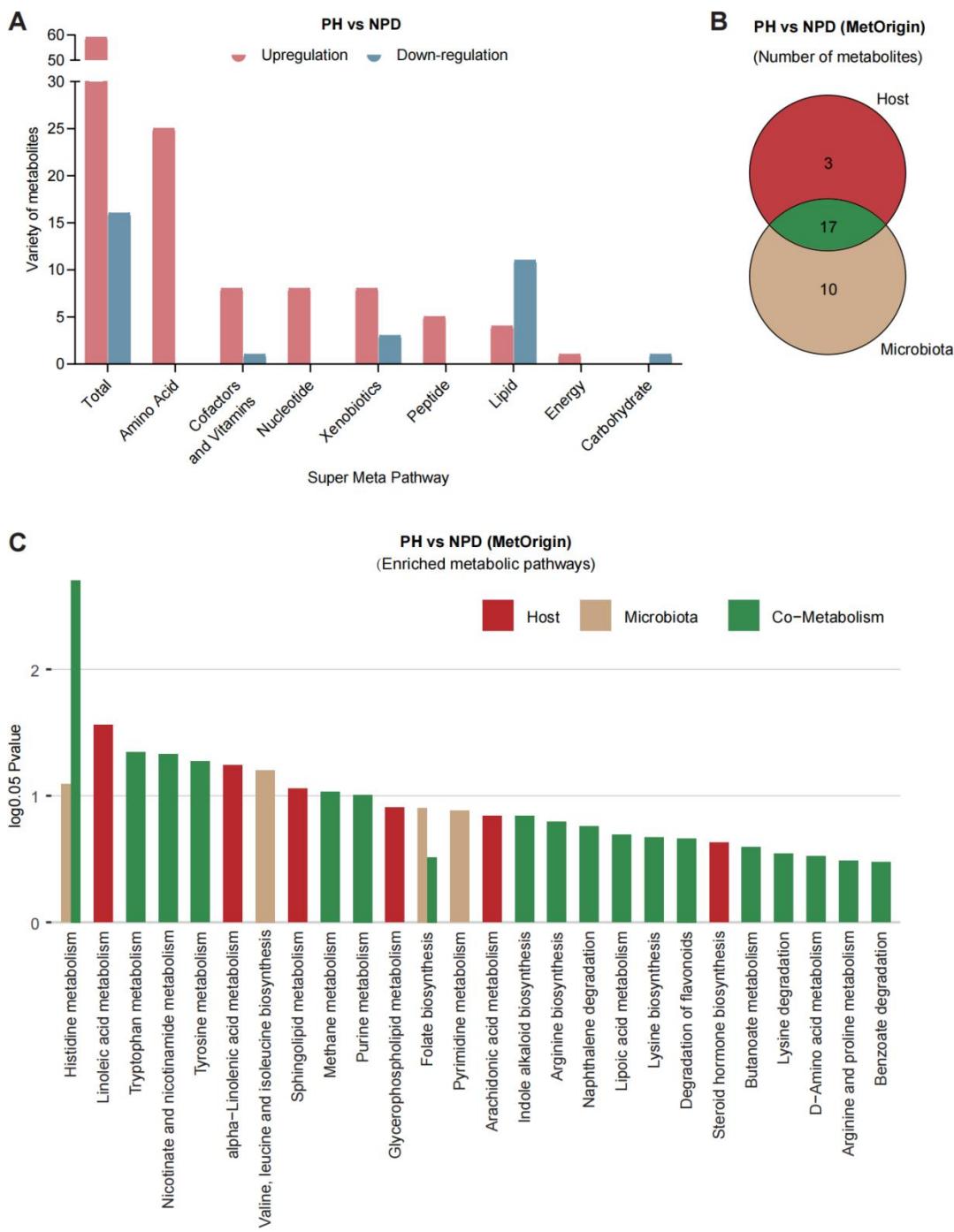
## Supplementary Material

### 1 Supplementary Figures and Tables

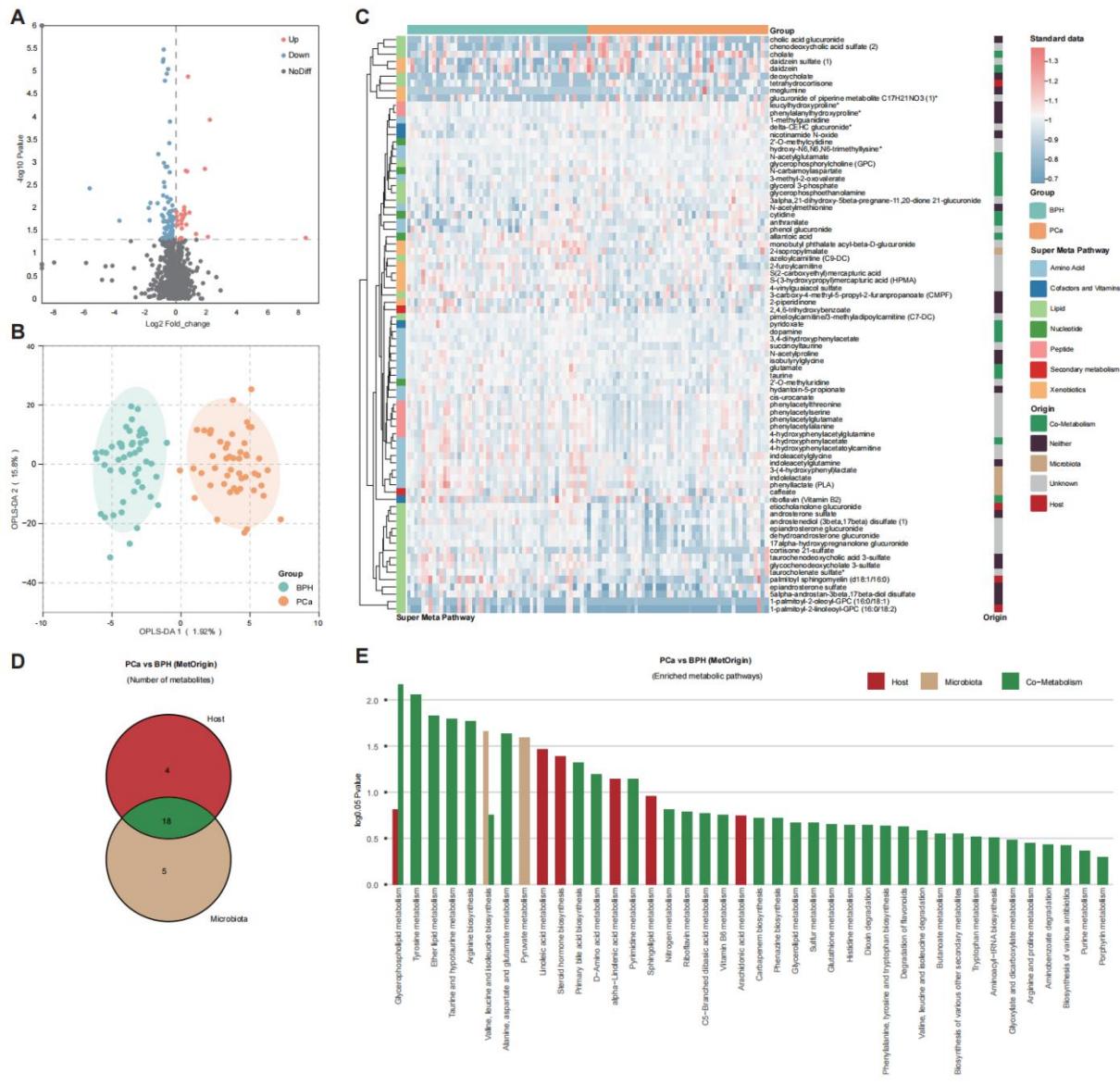
#### 1.1 Supplementary Figures



**Supplementary Figure 1.** Number of detected metabolites in each category from each group. \* $p < 0.05$ .



**Supplementary Figure 2. Category and origin of differential metabolites between PH and NPD.** (A). Bar plot of detected varieties of key metabolites in each category. (B) Venn plot of MetOrigin between PH and NPD. (C). Enriched pathways of MetOrigin between PH and NPD.



**Supplementary Figure 3. Discriminate analysis between PCa and BPH.** (A). Volcano plot of changing pattern of metabolites displayed with fold change and p-value. (B). Plot of structural separation after OPLS-DA. (C). Heatmap of 79 differential metabolites (OPLS-DA VIP >1.7). (D). Venn plot of MetOrigin between PCa and BPH. (E). Enriched pathways of MetOrigin between PCa and BPH.

## 1.2 Supplementary Tables

**Supplementary Table 1 List of key metabolites to discriminate PCa from BPH (except those of xenobiotics)**

Index	Super Meta Pathway	HMDB ID	VIP	AUC
etiocholanolone glucuronide	Lipid	HMDB0004484	2.59	0.775
2'-O-methyluridine	Nucleotide	HMDB0240328	3.31	0.760
palmitoyl sphingomyelin (d18:1/16:0)	Lipid	HMDB0010169	2.06	0.746
17alpha-hydroxypregnanolone glucuronide	Lipid	/	3.43	0.738
1-palmitoyl-2-oleoyl-GPC (16:0/18:1)	Lipid	HMDB0007972	3.57	0.734
epiandrosterone glucuronide	Lipid	/	2.20	0.715
pimeloylcarnitine/3-methyladipoylcarnitine (C7-DC)	Lipid	/	2.92	0.705
androsterone sulfate	Lipid	HMDB0002759	2.26	0.704
1-palmitoyl-2-linoleoyl-GPC (16:0/18:2)	Lipid	HMDB0007973	2.88	0.704
succinyltaurine	Amino Acid	/	2.54	0.696
dehydroandrosterone glucuronide	Lipid	/	2.01	0.673
1-methylguanidine	Amino Acid	HMDB0001522	2.50	0.671
N-acetylproline	Amino Acid	HMDB0094701	2.22	0.671
taurochenodeoxycholic acid 3-sulfate	Lipid	HMDB0002486	2.02	0.663
cholic acid glucuronide	Lipid	HMDB0002577	2.20	0.661
allantoic acid	Nucleotide	HMDB0001209	2.15	0.660
taurocholenate sulfate	Lipid	/	2.14	0.652
phenylacetylglutamate	Peptide	HMDB0059772	2.11	0.651
phenylalanylhydroxyproline	Peptide	HMDB0011176	2.10	0.651
pyridoxate	Cofactors and	HMDB0000017	2.88	0.651

Index	Super Meta Pathway	HMDB ID	VIP	AUC
Vitamins				
2'-O-methylcytidine	Nucleotide	/	1.83	0.650
glycerol 3-phosphate	Lipid	HMDB0000126	2.41	0.648
cortisone 21-sulfate	Lipid	/	2.01	0.645
androstenediol (3beta,17beta) disulfate	Lipid	/	1.88	0.644
leucylhydroxyproline	Peptide	HMDB0028930	2.08	0.643
chenodeoxycholic acid sulfate	Lipid	/	2.20	0.639
3-carboxy-4-methyl-5-propyl-2-furanpropanoate (CMPF)	Lipid	HMDB0061112	1.72	0.638
phenyllactate (PLA)	Amino Acid	HMDB0000779	2.02	0.637
N-acetylglutamate	Amino Acid	HMDB0001138	1.79	0.634
azeloylcarnitine (C9-DC)	Lipid	/	2.15	0.632
4-hydroxyphenylacetate	Amino Acid	HMDB0000020	2.22	0.629
5alpha-androstan-3beta,17beta-diol disulfate	Lipid	HMDB0094682	2.17	0.628
delta-CEHC glucuronide	Cofactors and Vitamins	/	2.12	0.627
phenylacetylalanine	Peptide	/	1.87	0.623
2,4,6-trihydroxybenzoate	Secondary metabolism	HMDB0029649	2.15	0.623
isobutyrylglycine	Amino Acid	HMDB0000730	2.15	0.619
indolelactate	Amino Acid	HMDB0000671	1.81	0.619
hydroxy-N6,N6,N6-trimethyllysine	Amino Acid	/	1.78	0.617
cis-urocinate	Amino Acid	/	1.72	0.616
4-hydroxyphenylacetylglutamine	Peptide	/	1.93	0.614
phenylacetylthreonine	Peptide	/	1.80	0.612

Index	Super Meta Pathway	HMDB ID	VIP	AUC
deoxycholate	Lipid	HMDB0000626	1.72	0.612
taurine	Amino Acid	HMDB0000251	1.74	0.610
glycochenodeoxycholate 3-sulfate	Lipid	HMDB0002497	1.82	0.610
nicotinamide N-oxide	Cofactors and Vitamins	HMDB0002730	2.46	0.608
phenylacetylserine	Peptide	/	1.92	0.606
epiandrosterone sulfate	Lipid	HMDB0062657	2.44	0.602
3alpha,21-dihydroxy-5beta-pregnane-11,20-dione 21-glucuronide	Lipid	/	2.36	0.602
anthranilate	Amino Acid	HMDB0001123	1.85	0.602
cholate	Lipid	HMDB0000619	1.75	0.601
N-carbamoylaspartate	Nucleotide	HMDB0000828	1.96	0.600
4-hydroxyphenylacetatoylcarnitine	Amino Acid	/	2.06	0.600
glutamate	Amino Acid	HMDB0000148	1.75	0.598
indoleacetylglutamine	Amino Acid	HMDB0013240	2.09	0.598
3-(4-hydroxyphenyl)lactate	Amino Acid	HMDB0000755	1.73	0.595
tetrahydrocortisone	Lipid	HMDB0000903	1.73	0.591
glycerophosphoethanolamine	Lipid	HMDB0000114	1.74	0.588
hydantoin-5-propionate	Amino Acid	HMDB0001212	1.70	0.583
indoleacetylglycine	Amino Acid	/	2.05	0.576
caffeate	Secondary metabolism	HMDB0001964	2.15	0.555
phenol glucuronide	Amino Acid	HMDB0060014	1.80	0.555
dopamine	Amino Acid	HMDB0000073	1.71	0.534
riboflavin (Vitamin B2)	Cofactors and	HMDB0000244	1.90	0.531

<b>Index</b>	<b>Super Meta Pathway</b>	<b>HMDB ID</b>	<b>VIP</b>	<b>AUC</b>
Vitamins				
3-methyl-2-oxovalerate	Amino Acid	HMDB0000491	2.41	0.515
cytidine	Nucleotide	HMDB0000089	1.90	0.512
N-acetylmethionine	Amino Acid	HMDB0011745	1.91	0.466
glycerophosphorylcholine (GPC)	Lipid	HMDB0000086	1.80	0.465
3,4-dihydroxyphenylacetate	Amino Acid	HMDB0001336	2.02	0.446