

Supplementary Materials: Influence of Intratumor Microbiome on Clinical Outcome and Immune Processes in Prostate Cancer

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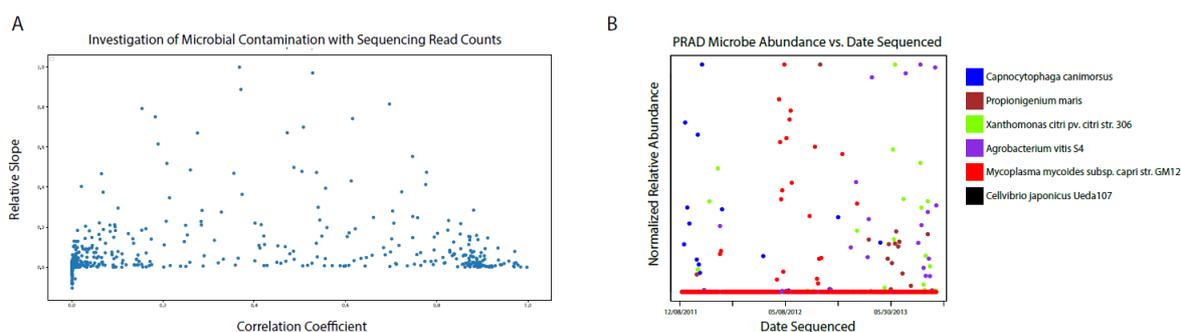


Figure S1. (A) Microbial contamination correction with sequencing read counts. R-squared and slope was calculated to detect contaminants, microbes with large slopes. (B) Contamination correction by date sequenced.

Table S1. Microbes in PC. Columns “Role in PC” and “Correlation” are written according to results found in this study. Column “Known Function” is written according to prior research.

Microbe	Role in PC	Known Function	Correlation
<i>Pediococcus pentosaceus</i>	Anti-tumor	Mitigate colon cancer	Gleason score
<i>Listeria monocytogenes</i> (Lm)	Anti-tumor	Stimulate proinflammatory cytokines and trigger immune response	Gleason score
<i>Lactobacillus crispatus</i>	Anti-tumor	Causes reduction in tumor size in breast cancer	Gleason score
<i>Bacillus halodurans</i>	Anti-tumor		Gleason score
<i>Nevskia ramosa</i>	Pro-tumor		Gleason score
<i>Methylobacterium radiotolerans</i> JCM 2831	Anti-tumor		TNM staging
<i>Meiothermus silvanus</i> DSM 9946	Anti-tumor		TNM staging
<i>Rhodococcus erythropolis</i> PR4	Anti-tumor	Associated with immunosuppression	TNM staging
<i>Methylobacterium radiotolerans</i>	Anti-tumor	Associated with immunosuppression	TNM staging
<i>Stenotrophomonas maltophilia</i>	Anti-tumor	Associated with infection	TNM staging
<i>Thermus thermophilus</i>	Pro-tumor	Anti-tumor according to past research; inhibits cancer growth	PSA
<i>Campylobacter concisus</i>	Pro-tumor	Induces inflammation	PSA
<i>Streptococcus pneumoniae</i>	Pro-tumor		PSA
<i>Xanthomonas albilineans</i> GPE PC73	Anti-tumor		PSA
<i>Herminiimonas arsenicoxydans</i>	Anti-tumor		PSA

<i>Pseudarthrobacter chlorophenicus</i>	Anti-tumor		PSA
<i>D. acidovorans SPH-1</i>	Pro-tumor	Associated with immunosuppression and infection	Genomic alterations and dysregulated IA genes
<i>N. hamburgensis X 14</i>	Pro-tumor		Genomic alterations and dysregulated IA genes
<i>S. aureus</i>	Pro-tumor	Causes prostate abscess/inflammation	Genomic alterations, dysregulated IA genes, PCSC gene expression
<i>G. vaginalis</i>	Pro-tumor	Strongly associated with infection	Genomic alterations and dysregulated IA genes
<i>Bradyrhizobium elkanii</i>	Anti-tumor		AR expression
<i>Ochrobactrum anthropi ATCC 49188</i>	Anti-tumor		AR expression
<i>Bradyrhizobium japonicum</i>	Anti-tumor		AR expression
<i>Escherichia coli ETEC H10407</i>	Pro-tumor		AR expression
<i>Escherichia coli str. K-12 substr. MG1655</i>	Pro-tumor		AR expression
<i>Paraburkholderia phymatum STM815</i>	Pro-tumor		PCSC gene expression
<i>Pseudomonas putida F1</i>	Pro-tumor		PCSC gene expression
<i>Haemophilus parainfluenzae T3T1</i>	Pro-tumor		PCSC gene expression
<i>Stackebrandtia nassauensis DSM 44728</i>	Pro-tumor	Breaks down uric acid	
<i>Mycoplasma hyorhinitis HUB-1</i>	Pro-tumor	Activates the NLRP3 Inflammasome and MMP2, phosphorylation of EGFR, promotes migration and invasion of gastric cancer cells	



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