

Differences in the Stool Microbiome Before and After Colorectal Cancer Treatment

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Supplemental

Table S1: P-values of Differences between Initial and Follow Up Samples for Richness, Shannon Diversity, and Evenness.

| Comparison | Alpha Metric | P-value | BH Adjusted P-value |
|------------|--------------|-----------|---------------------|
| all | sobs | 0.2542809 | 0.3536011 |
| all | shannon | 0.3536011 | 0.3536011 |
| all | evenness | 0.3251909 | 0.3536011 |
| adn | sobs | 0.7195598 | 0.7195598 |
| adn | shannon | 0.6168494 | 0.7195598 |
| adn | evenness | 0.4485818 | 0.7195598 |
| crc | sobs | 0.1573652 | 0.1573652 |
| crc | shannon | 0.0407472 | 0.0611208 |
| crc | evenness | 0.0236109 | 0.0611208 |

Table S2: Summary of Most Important Variables For the Model

| Variable | Lowest Taxonomic ID | Total Model Appearances |
|-------------|-----------------------|-------------------------|
| fit_result | NA | 100 |
| Age | NA | 100 |
| Otu000013 | Bacteria | 100 |
| Otu000018 | Escherichia/Shigella | 100 |
| Otu000020 | Bacteria | 100 |
| BMI | NA | 100 |
| Otu000017 | Ruminococcus | 99 |
| Otu000153 | Porphyromonas | 99 |
| Otu000012 | Blautia | 99 |
| Otu000010 | Blautia | 98 |
| Otu000007 | Bacteroides | 98 |
| Otu000003 | Roseburia | 97 |
| Gender.mYes | NA | 92 |
| Otu000006 | Bacteria | 92 |
| Otu000045 | Bacteria | 90 |
| Otu000114 | Clostridium_XIVa | 88 |
| Otu000015 | Parabacteroides | 88 |
| Otu000008 | Faecalibacterium | 86 |
| Otu000016 | Bacteroides | 86 |
| Otu000046 | Bacteroides | 85 |
| Otu000061 | Bacteria | 83 |
| Otu000032 | Bacteroides | 81 |
| Otu000089 | Ruminococcus | 80 |
| Otu000231 | Alistipes | 69 |
| Otu000002 | Anaerostipes | 67 |
| Otu000088 | Bacteria | 65 |
| Otu000005 | Bacteroides | 65 |
| Otu000124 | Bacteria | 61 |
| Otu000034 | Clostridium_XIVa | 57 |
| Otu000220 | Clostridium_IV | 55 |
| Otu000036 | Phascolarctobacterium | 54 |
| Otu000049 | Barnesiella | 52 |

Table S3: Summary of Probability Assignment of Follow Up Individuals

| Initial Positive Probability | Follow Up Positive Probability | Disease Free on Follow Up | Initial Diagnosis |
|---------------------------------|-----------------------------------|------------------------------|----------------------|
| 0.9855 | 0.9275 | No | cancer |
| 0.9570 | 0.7255 | Yes | cancer |
| 0.9300 | 0.7975 | Unknown | cancer |
| 0.9635 | 0.8785 | Yes | cancer |
| 0.9820 | 0.7580 | Unknown | cancer |
| 0.9380 | 0.3340 | Unknown | cancer |
| 0.9500 | 0.4930 | Yes | cancer |
| 0.9610 | 0.3465 | Unknown | cancer |
| 0.9845 | 0.7365 | Yes | cancer |
| 0.8090 | 0.6040 | Yes | cancer |
| 0.9515 | 0.8870 | Yes | cancer |
| 0.9705 | 0.6820 | Yes | cancer |
| 0.9785 | 0.7600 | Yes | cancer |
| 0.9655 | 0.5860 | Yes | cancer |
| 0.9715 | 0.3440 | Yes | cancer |
| 0.9190 | 0.9290 | Yes | cancer |
| 0.8060 | 0.3600 | Yes | cancer |
| 0.9730 | 0.9370 | Yes | cancer |
| 0.7785 | 0.2710 | Yes | cancer |
| 0.8855 | 0.4750 | Yes | cancer |
| 0.9350 | 0.9020 | Yes | cancer |
| 0.9325 | 0.3420 | Unknown | cancer |
| 0.9850 | 0.3250 | Yes | cancer |
| 0.9595 | 0.6970 | Yes | cancer |
| 0.9590 | 0.8770 | Yes | cancer |
| 0.9900 | 0.4170 | Yes | cancer |
| 0.9595 | 0.6190 | Yes | adv_adenoma |
| 0.8070 | 0.4295 | Yes | adenoma |
| 0.7265 | 0.4200 | Yes | adenoma |
| 0.7895 | 0.4710 | Yes | adv_adenoma |
| 0.7390 | 0.4180 | Yes | adenoma |
| 0.7845 | 0.5715 | Yes | adenoma |
| 0.9570 | 0.7940 | Yes | adenoma |
| 0.8310 | 0.4080 | Yes | adenoma |
| 0.8995 | 0.8525 | Yes | adenoma |
| 0.8840 | 0.7940 | Yes | adenoma |
| 0.8625 | 0.7740 | Yes | adenoma |
| 0.9460 | 0.8350 | Yes | adenoma |
| 0.9110 | 0.6160 | Yes | adv_adenoma |
| 0.9545 | 0.8710 | Yes | adenoma |

| Initial Positive Probability | Follow Up Positive Probability | Disease Free on Follow Up | Initial Diagnosis |
|---------------------------------|-----------------------------------|------------------------------|----------------------|
| 0.8350 | 0.6095 | Yes | adv_adenoma |
| 0.9590 | 0.9185 | Yes | adv_adenoma |
| 0.7410 | 0.4550 | Yes | adenoma |
| 0.8825 | 0.7670 | Yes | adv_adenoma |
| 0.9265 | 0.8790 | Yes | adenoma |
| 0.9735 | 0.6035 | Yes | adv_adenoma |
| 0.8425 | 0.2030 | Yes | adenoma |
| 0.8290 | 0.6525 | Yes | adv_adenoma |
| 0.8035 | 0.3405 | Yes | adv_adenoma |
| 0.9940 | 0.5190 | Yes | adenoma |
| 0.9775 | 0.9715 | Yes | adv_adenoma |
| 0.9435 | 0.5615 | Yes | adv_adenoma |
| 0.8220 | 0.7395 | Yes | adv_adenoma |
| 0.9225 | 0.9285 | Yes | adv_adenoma |
| 0.9370 | 0.3410 | Yes | adenoma |
| 0.7685 | 0.5425 | Yes | adv_adenoma |
| 0.9275 | 0.7890 | Yes | adv_adenoma |
| 0.7620 | 0.3775 | Yes | adenoma |
| 0.8775 | 0.7920 | Yes | adenoma |
| 0.7660 | 0.3530 | Yes | adenoma |
| 0.9425 | 0.3385 | Yes | adv_adenoma |
| 0.9550 | 0.4755 | Yes | adenoma |
| 0.7710 | 0.8150 | Yes | adv_adenoma |
| 0.9735 | 0.9295 | Yes | adenoma |
| 0.9870 | 0.9120 | Yes | adenoma |
| 0.7530 | 0.4330 | Yes | adv_adenoma |

Table S4: Summary of OTUs in both Lesion and Initial/Follow Up Models

| OTU | Lowest Taxonomic ID | P-value | BH Corrected P-value |
|-----------|----------------------|-----------|----------------------|
| Otu000012 | Blautia | 0.0037523 | 0.0300185 |
| Otu000005 | Bacteroides | 0.2330135 | 0.2663011 |
| Otu000008 | Faecalibacterium | 0.1647056 | 0.2196075 |
| Otu000018 | Escherichia/Shigella | 0.0431082 | 0.1149553 |
| Otu000002 | Anaerostipes | 0.2694179 | 0.2694179 |
| Otu000114 | Clostridium_XIVa | 0.0425792 | 0.1149553 |
| Otu000124 | Bacteria | 0.1006246 | 0.2012493 |
| Otu000020 | Bacteria | 0.1390492 | 0.2196075 |

Table S5: Summary of P-values for Specific CRC OTUs between Initial and Follow Ups

| Taxonomic ID | CRC P-value | CRC BH Corrected P-value | Adenoma P-value | Adenoma BH Corrected P-value |
|---------------------------------------|----------------|--------------------------------|--------------------|---------------------------------|
| Fusobacterium Nucleatum | 0.06848 | 0.2739 | 0.370 | 1.000 |
| Parvimonas Micra | 0.01385 | 0.0554 | 0.159 | 0.635 |
| Peptostreptococcus Assacharolytica | 0.00457 | 0.0183 | 1.000 | 1.000 |
| Porphyromonas Stomatis | 0.00384 | 0.0154 | 0.236 | 0.944 |