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 |
|------------|--------------|-----------|---------------------|
| lesion | sobs | 0.4294106 | 0.6550031 |
| lesion | shannon | 0.5549842 | 0.6550031 |
| lesion | evenness | 0.1412681 | 0.4238043 |
| adn | sobs | 0.6909558 | 0.6909558 |
| adn | shannon | 0.4962226 | 0.6550031 |
| adn | evenness | 0.5378507 | 0.6550031 |
| crc | sobs | 0.5822250 | 0.6550031 |
| crc | shannon | 0.0890290 | 0.4006306 |
| crc | evenness | 0.0072909 | 0.0656181 |

Table S2: Summary of OTUs in both Lesion and Initial/Follow Up Models

| OTU | Lowest Taxonomic ID | Lesion Model MDA | Initial Model MDA |
|-----------|----------------------|------------------|-------------------|
| Otu000015 | Anaerostipes | 4.749411 | 0.7342620 |
| Otu000006 | Blautia | 4.004671 | 0.8124998 |
| Otu000061 | Bacteroides | 3.938754 | 0.9090068 |
| Otu000005 | Ruminococcus | 3.759063 | 0.7485388 |
| Otu000004 | Collinsella | 3.710266 | 0.8403637 |
| Otu000058 | Faecalibacterium | 3.693178 | 0.6764208 |
| Otu000018 | Ruminococcaceae | 3.651454 | 0.8432880 |
| Otu000001 | Roseburia | 3.627399 | 0.9493613 |
| Otu000037 | Roseburia | 3.612381 | 0.8905661 |
| Otu000014 | Ruminococcaceae | 3.463134 | 0.7600457 |
| Otu000010 | Escherichia/Shigella | 3.440272 | 0.8503705 |
| Otu000013 | Streptococcus | 3.430315 | 0.6954449 |
| Otu000002 | Faecalibacterium | 3.393844 | 0.6779158 |
| Otu000007 | Blautia | 3.349713 | 0.7652515 |
| Otu000009 | Bacteroides | 3.314681 | 0.7102074 |
| Otu000003 | Roseburia | 3.308582 | 0.7094481 |
| Otu000020 | Lachnospiraceae | 3.296764 | 0.7812310 |
| Otu000040 | Blautia | 3.213801 | 0.9582915 |
| Otu000084 | Anaerostipes | 3.146415 | 0.6561690 |
| Otu000026 | Bacteroides | 3.099604 | 0.9000035 |
| Otu000041 | Clostridiales | 3.098968 | 0.6728631 |
| Otu000059 | Ruminococcus | 3.078183 | 0.7190440 |
| Otu000109 | Clostridiales | 3.062303 | 0.7107204 |
| Otu000108 | Bacteroides | 3.031357 | 0.7637155 |

| OTU | Lowest Taxonomic ID | Lesion Model MDA | Initial Model MDA |
|-----------|---------------------|------------------|-------------------|
| Otu000016 | Blautia | 3.031011 | 0.6952266 |
| Otu000085 | Blautia | 3.027084 | 0.6717972 |
| Otu000148 | Bacteroides | 2.938932 | 0.9272185 |
| Otu000180 | Lachnospiraceae | 2.903063 | 0.7038127 |
| Otu000028 | Lachnospiraceae | 2.814039 | 0.9395611 |
| Otu000025 | Dorea | 2.787955 | 0.6563795 |
| Otu000605 | Clostridium_XI | 2.529424 | 0.6223713 |
| Otu000167 | Dorea | 2.372671 | 0.6050109 |

Table S3: Summary of Comparisons based on Chemotherapy or Radiation Received

| Variable Tested | Treatment | P-value | BH Corrected P-value |
|----------------------|--------------|-----------|----------------------|
| Positive Probability | Chemotherapy | 0.0091511 | 0.036604 |
| Positive Probability | Chemotherapy | 0.0333166 | 0.088844 |
| Positive Probability | Chemotherapy | 0.0006355 | 0.005084 |
| Positive Probability | Chemotherapy | 0.0001309 | 0.002095 |
| Positive Probability | Radiation | 0.0815484 | 0.163097 |
| Positive Probability | Radiation | 0.0754788 | 0.163097 |
| Positive Probability | Radiation | 0.0089848 | 0.036604 |
| Positive Probability | Radiation | 0.0213082 | 0.068186 |
| Thetayc | Chemotherapy | 0.9187938 | 0.933448 |
| Thetayc | Radiation | 0.6279206 | 0.717624 |
| Sobs | Chemotherapy | 0.5946906 | 0.717624 |
| Shannon | Chemotherapy | 0.1822094 | 0.291535 |
| Evenness | Chemotherapy | 0.1628138 | 0.289447 |
| Sobs | Radiation | 0.9334483 | 0.933448 |
| Shannon | Radiation | 0.3838308 | 0.558299 |
| Evenness | Radiation | 0.4965131 | 0.662017 |