**Table 1:** Summary of Subsampling Comparisons in Figure 6

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Dataset | Bacteria detected | Samples per group (BH) | Samples per group (DS) | Proportion fewer samples | Reference |
| DME | 74 | 16 | 8 | 50 % | Dominguez-Bello et al, 2010 |
| UKT | 65 | 100 | 85 | 15 % | Beaumont et al, 2016 |
| DIBD | 87 | 65 | 42 | 35 % | Vazquez-Baeza et al, 2016 |
| CD | 713 | 320 | 260 | 19 % | Gevers et al, 2014 |
| AGP | 244 | 350 | 170 | 51 % | American Gut Project |
| AGA | 230 | 160 | 80 | 50 % | American Gut Project |