Additional Figure 4: Detailed normalized comparison between 16S samples obtained using 454 and SOLiD sequencing. Detailed normalized comparison tree obtained using MEGAN for '16S-454' dataset (cyan) and '16S-SOLiD' dataset (magenta) without considering the 'No hits' node. The tree is collapsed at 'genus' level of NCBI taxonomy. Circles are scaled logarithmically to indicate the number of summarized reads.