



Fig. S11. Sensitivity and specificity of Resphera Insight on the HMP reference isolate collection. *In silico* sensitivity and mis-assignment rates of Resphera Insight compared to QIIME RDP and QIIME UCLUST for **a** a collection of representative species from the HMP reference genome set (N = 190), **b** species listed by Human Oral Microbiome Database (N = 123), and **c** species within the genus *Bacteroides* (N = 17). See Methods for validation description.