

Fig XYZ. Assessment Framework. (A) Count tables evaluated by the assessment framework are generated from the assessment dataset using marker-gene survey bioinformatic pipelines. Count table rows are features identified by the bioinformatic pipeline and column are samples, four PCR replicates (A-D) for PRE and POST and titrations, to simplify the diagram only three titrations are shown. The seven features represent the seven feature types observed. (B) Feature type definitions along with pictoral depiction of abundance values and which features are used for the qualitative and quantitative assessments, indicated by grey filled cells in table. (C) Assessment metrics calculated for example count table and example features.