

k__Bacteria; p__Proteobacteria; c__Gammaproteobacteria; o__Pseudomonadales; f__Pseudomonadaceae; g__Pseudomonas; s__veronii featureID: ddf1f3e5f93ba45f39c7a066375ba07f **Correlation with all samples** relative abundance (%) ••• 27 33 Cq value SampleType Blank-extraction Correlation within sample type SampleType: REF-DIC SampleType: REF-DIM $log_e(S) = 6.516$, p = 0.510, $\rho_{Spearman} = 0.172$, $Cl_{95\%}$ [-0.301, 0.675], n = 17 $log_e(S) = 5.939$, p = 0.007, $\rho_{Spearman} = 0.608$, $Cl_{95\%}$ [0.285, 1.004], n = 18relative abundance (%) relative abundance (%) 25 26 31 32 22 23 24 27 Cq value Cq value SampleType: IM100-DIC SampleType: IM100-DIM $log_e(S) = 5.930, p = 0.083, \rho_{Spearman} = 0.447, Cl_{95\%} [0.058, 0.920], n = 16$ $log_e(S) = 6.797$, p = 0.763, $\rho_{Spearman} = 0.076$, $Cl_{95\%}$ [-0.532, 0.635], n = 18relative abundance (%) 0.0 2.0 0.0 0.0 0.0 relative abundance (%) 25.0 27.5 28 30.0 30 32 Cq value Cq value

k_Bacteria; p_Actinobacteria; c_Actinobacteria; o_Actinomycetales; f_Propionibacteriaceae; g_Propionibacterium; s_acnes featureID: f17d3da8eb98435912cf7cfb2d1bfbdb **Correlation with all samples** 1.25 0.00 24 27 30 33 Cq value SampleType Blank-extraction IM100-DIC Correlation within sample type SampleType: REF-DIC SampleType: REF-DIM $log_e(S) = 5.308$, p = 0.533, $\rho_{Spearman} = -0.224$, $Cl_{95\%}$ [-0.883, 0.457], n = 10 $log_e(S) = 4.682$, p = 0.328, $\rho_{Spearman} = 0.345$, $Cl_{95\%}$ [-0.155, 0.878], n = 10uelative abundance (%) 0.75 relative abundance (%) 0.5 0.0 -0.5 23 25 26 22 24 27 31.5 31.8 32.1 32.4 Cq value Cq value SampleType: IM100-DIC SampleType: IM100-DIM $log_e(S) = 4.407$, p = 0.138, $\rho_{Spearman} = 0.503$, $Cl_{95\%}$ [0.009, 1.037], n = 10 $log_e(S) = 3.989$, p = 0.939, $\rho_{Spearman} = 0.036$, $Cl_{95\%}$ [-0.976, 0.954], n = 7relative abundance (%) 1.0 0.2 -0.5 0.0 0.0 -0.5 -0.2 25 27 30 23 29 31 28 32 Cq value Cq value

relative abundance (%)