

The Human Microbiome Company

## **ORAL ECOLOGIX**

REPORT ID: S017058

invivo

TEST REPORTED: 28/10/2022 TEST RECEIVED: 28/10/2022 PATIENT NAME: FIRST LAST PATIENT DOB: 20/10/2022 CLINICIAN NAME: CLINICIAN

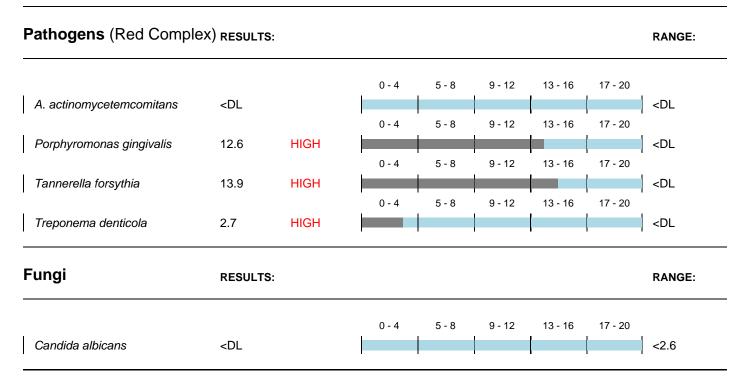
SAMPLE TYPE: SALIVA SEX: FEMALE Lab Director: Emma Beamish, PhD



## **Opportunistic Bacteria** (Orange Complex) **RESULTS:** RANGE: 0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 Campylobacter rectus 9.0 3.5-10.5 0 - 4 9 - 12 13 - 16 17 - 20 5 - 8 Capnocytophaga gingivalis 11.7 8.9-14.0 13 - 16 17 - 20 0 - 4 5 - 8 9 - 12 Enterococcus faecalis <DL <DL 0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 Fusobacterium nucleatum 14.1 3-16.5 0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 Lactobacillus spp. 6.9 HIGH < 5.6 0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 Parvimonas micra 11.8 HIGH 2.2-11.3 0 - 4 5 - 8 9 - 12 13 - 16 17 - 20 6.5-15.0 Peptostreptococcus anaerobius 11.7 13 - 16 17 - 20 5 - 8 9 - 12 0 - 4 Prevotella intermedia HIGH <4.9 13.3 5 - 8 9 - 12 0 - 4 13 - 16 17 - 20 Streptococcus mutans 2.7 <3.8



**Disclaimer**: This test was developed, and its performance characteristics determined by Invivo Diagnostics. This test is not intended for use by consumers or physicians as a means to cure, treat, prevent, diagnose or mitigate any disease or other medical condition. The information contained in this document is in no way to be taken as prescriptive nor to replace the physicians duty of care and personalised care practices.



The Oral EcologiX<sup>™</sup> profile utilises the highly sensitive quantitative PCR (qPCR) TaqMan technology for analysis of the oral microbiota. Microbial genes of interest are quantified within a sample and their abundances are normalised to an endogenous and highly conserved gene. The qPCR results are therefore reported as the relative abundance of a microorganism as proportional to the whole microbial community.

