ORAL ECOLOGIX

invivo

REPORT ID:

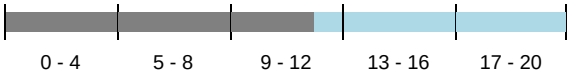
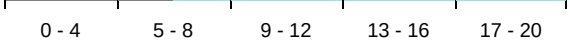
TEST REPORTED: 27/09/2021
TEST RECEIVED: 15/09/2021
PATIENT NAME: SAMPLE PATIENT
PATIENT DOB:
GENDER: FEMALE

REPORT STATUS: COMPLETED
CLINICIAN NAME: SAMPLE CLINICIAN
ACCESSION NO:
SAMPLE TYPE: SALIVA

Lab Director:
Eviatar Natan, PhD

CareQuality Commission Registered

Commensal Bacteria
(Orange Complex)

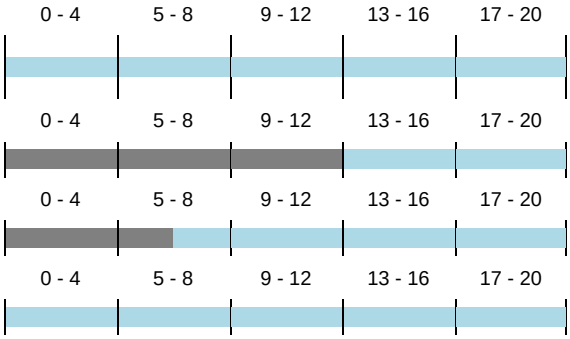
	RESULTS:		0 - 4 / 5 - 8 / 9 - 12 / 13 - 16 / 17 - 20	RANGE:
Campylobacter rectus	11.5	HIGH		3.5-10.5
Eubacterium nodatum	<DL			<1.0
Fusobacterium nucleatum	15.9			3-16.5
Lactobacillus spp.	6.1	HIGH		< 5.6
Parvimonas micra	11.1			2.2-11.3
Peptostreptococcus anaerobius	13.5			6.5-15.0
Prevotella intermedia	10.2	HIGH		<4.9
Prevotella nigrescens	6.2	HIGH		1.4-5.9
Streptococcus mutans	7.2	HIGH		<3.8

Scan for more information and resources on Oral EcologiX

***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.*

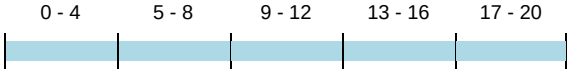
Pathogens (Red Complex)

	RESULTS:		**RANGE:**
Aggregatibacter actinomycetemcomitans	<DL		<5.1
Porphyromonas gingivalis	12.4	HIGH	<6.8
Tannerella forsythia	5.9	HIGH	< 5.6
Treponema denticola	<DL		<2.5



Fungi

	RESULTS:	**RANGE:**
Candida albicans	<DL	<2.6



The Oral EcologiX™ 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.

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