

# NGS

## VIASURE



### Efficient Amplification

**OneStep multiplex qPCR** for covering V1, V2, V3, V4 regions of the 16SrRNA gene.



**Quantitative Precision**  
qPCR ensures immediate and accurate library quantification.



### Streamlined Process

**Easy qPCR, quantification, pooling and purification.**



**Comprehensive Analysis**  
**V-Xplora** software for seamless data interpretation.



/01  
ENRICHMENT & INDEXING



/02  
POOLING



/03  
FINAL LIBRARY



/04  
SEQUENCING



/05  
SEQUENCE DATA ANALYSIS

**certest**

NGS/16SV14-0525EN

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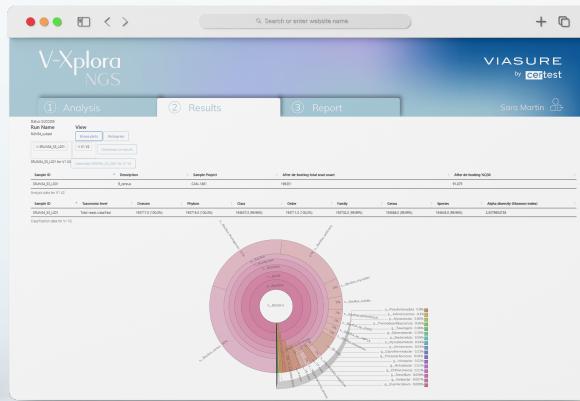


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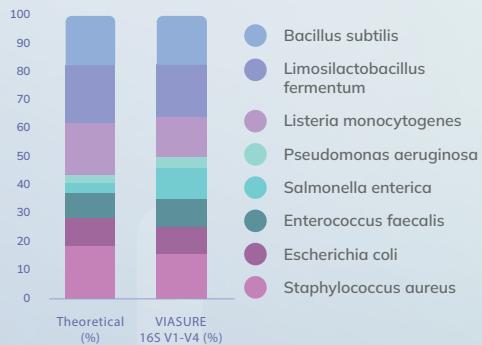
# VIASURE 16S V1-V4

## NGS Solution

(VS-16S-V1448PHRUO)\*



Achieve reliable and precise results effortlessly!



Enrichment Method	Amplicon-based enrichment
Sequencing platforms	Illumina (MiSeq) and Element Biosciences-AVITI
Library size	500-600 bp
Input DNA requirements	5 µL of your sample. No need to worry about input
Nº of pools per panel	1
Supported sample types	Biological samples or culture
Sequencing Analysis	Software V-Xplora
Plexity	up to 48
Hands on time	<30 min

AMPLIFY  
YOUR RESEARCH  
*to excellence*

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