# Quantabio

# Product Specifications 95191-096 Rev 01

Product Information		
sparQ DNA Library Prep Kits		
Part Number	95191-096	
Number of Reactions	96 Reactions	
Storage Temperature	-25ºC to -15ºC	
Lot Number	66215580	
Reference Number	041222,081722,128100422, 120821B,020322,275754832	
Expiration Date	02/29/24	

# Components:

84515 DNA Polishing Enzyme Mix, 0.960mL 84517 DNA Polishing Buffer, 0.480mL 84519 DNA Ligase, 0.480mL 84521 DNA Ligation Buffer, 0.960mL 84523 HiFi PCR Master Mix (2X), 1.250mL 84525 Primer Mix, 0.288mL

# **Quality Control Analysis and Specifications:**

# Product Description:

The sparQ DNA Library Prep Kit provides components for the rapid construction of DNA libraries from fragmented doublestranded DNA for sequencing on Illumina® NGS platforms. The streamlined workflow can be completed in under 3 hours with minimal hands-on time. The DNA polishing reactions are combined in a single step to convert fragmented DNA into 5'-phosphorylated and 3'-dA-tailed DNA fragments suitable for direct ligation of sequencing adapters without the need for an intervening cleanup. The 2X HiFi PCR Master Mix and Primer Mix allow the optional, unbiased amplification of fragments with appropriate adapters ligated to both ends. The kit is compatible with multiple sample types and facilitates efficient and consistent library construction from a wide range of input amounts from 0.25 ng to 1000 ng DNA.

Product Specifications	
95191	
Assay	Library Functional Assay
Specification	Functional

**Library Prep Functional Assay**: Quality of the sparQ DNA Library Prep Kit is tested functionally by preparation of a DNA library from bacterial genomic DNA with GC-content of 10-80%. The differences in library yield and profile among different lots must be within 15%. Sequencing of the amplified library must yield mapped reads >90% and normalized coverage between 0.7 and 1.3 across the full GC spectrum.

Enzyme components were tested prior to assembly and free of contaminating endonucleases and exonucleases. Enzyme purity was >95% as determined by SDS-PAGE and negligible *E. coli* genomic DNA contamination was confirmed by qPCR.

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