

Supported by: German Cancer Research Center University Hospital Carl Gustav Carus Dresden Carl Gustav Carus Faculty of Medicine, TU Dresden Helmholtz-Zentrum Dresden-Rossendorf

## Core Unit for Molecular Tumor Diagnostics

## **Input Recomendation for selected Protocol**

Protocol	Digital Droplet PCR	
Input Material	DNA/RNA	
Minimum Input	depends on Material	
Standard Input	depends on Material	
Maximum Volume	5,0 μL	
Minimal Concentration	depends on Material	
Minimal Concentration Standard Concentration	depends on Material depends on Material	

## For more Information please visit the manufacturers homepage

https://www.stillatechnologies.com/

https://www.gene-pi.com/

The CMTD is not responsible for the consequences/results, if an ordering work group wants to sequence RNA/DNA samples that failed the predefined quality parameters for sequencing!

Possible, risks which can occur are for example bad sequencing quality, no sequencing possible due to failure in library generation, introducing false signals of gene expression or allele frequency.

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