

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 (Multiplex)
Input Material	DNA/RNA
Minimum Input	depends on Material
Standard Input	depends on Material
Maximum Volume	5,0 μL
Minimal Concentration	depends on Material
0, 1, 1,0	depends on Material
Standard Concentration	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.

Erstellt: Alexander Krüger	Geändert: Alexander Krüger	Geprüft: Sandra Oster	Freigegeben:
Datum: 28.09.2021	Datum: 25.02.2022	Datum:	Datum:

Version 03 Seite 1 von 1