

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	[Custom] Agilent cfDNA-GBM-Panel
Input Material	cfDNA
Minimum Input	10,0 ng
Standard Input	200,0 ng
Maximum Volume	7,0 μL
Minimal Concentration	1,43 ng/μL
Standard Concentration	28,57 ng/μL
Extraction derived from	Non-FFPE

Special Considerations

please contact the CMTD for further consulting. Custom Panel requiers at least 96 Samples.

For more Information please visit the manufacturers homepage

https://www.agilent.com/en/product/next-generation-sequencing/hybridization-based-next-generation-sequencing-ngs/ngs-custom-target-enrichment-probes/ngs-custom-target-enrichment-probes-232874

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