

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	Agilent Mouse Exom
Input Material	DNA
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 different versions of Exom available

## For more Information please visit the manufacturers homepage

https://www.agilent.com/en/product/next-generation-sequencing/hybridization-based-next-generation-sequencing-ngs/exome-probes/sureselect-non-human-exomes-232868

https://www.agilent.com/en/product/next-generation-sequencing/hybridization-based-next-generation-sequencing-ngs/dna-seq-reagents-kits-library-preparation-kits/sureselect-xt-hs2-dna-reagent-kit-783974

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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