Protocol: Illumina RNA Fusion

**Input Material**: RNA

**Minimum Input**: 10,0 ng

**Standard Input**: depends on Quality

**Maximum Volume**: 8,5 µL

**Minimal Concentration**: 1,18 ng/µL

**Standard Concentration**: depends on Quality

Extraction derived from: FFPE/Non-FFPE

**RNA Quality parameters**

- **RIN**: not considered
- **DV200**: >30%

For more information please visit the manufacturers homepage:


[https://support.illumina.com/sequencing/sequencing_kits/trusight-rna-fusion-panel.html](https://support.illumina.com/sequencing/sequencing_kits/trusight-rna-fusion-panel.html)

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.