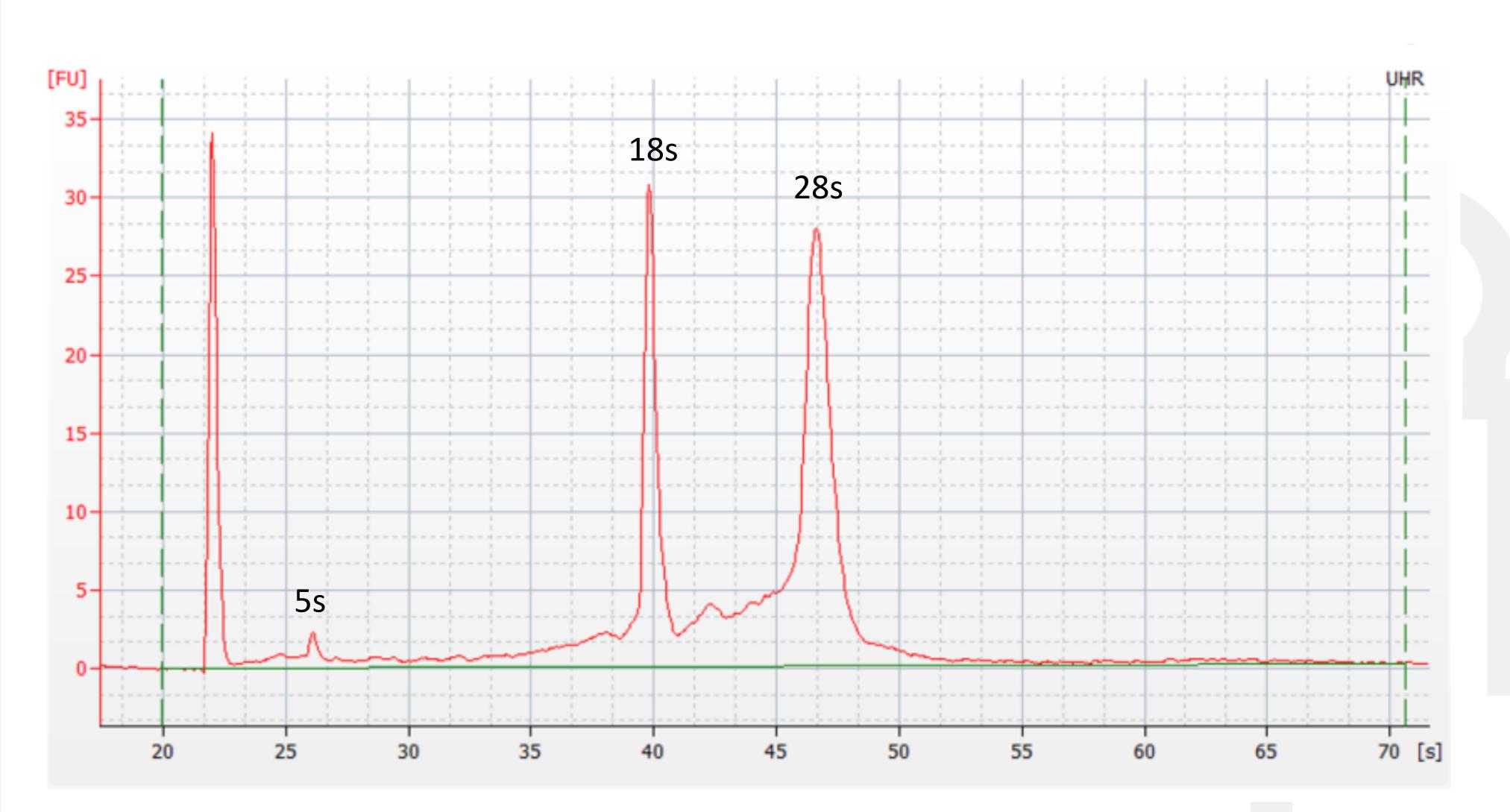
## Integridade do RNA



Avaliada por uma métrica arbitrária: razão 18s/5s



## Um protocolo para cada objetivo



Application	Product	Benefits
Total RNA Sequencing	Illumina Stranded Total RNA Prep	<ul> <li>Integrated enzymatic RNA depletion removes both rRNA and globin mRNA in a single, rapid step</li> <li>One tube to deplete abundant transcripts from multiple species provides flexibility for mixed samples</li> <li>Use with human, mouse, rat, bacteria and epidemiology samples</li> <li>Broad RNA input range, rapid protocol, cost-effective sequencing with up to 384 UDIs</li> <li>Ligation method results in high coverage uniformity and reliability even from degraded samples</li> <li>Measure fold changes in FFPE and low-quality samples</li> </ul>
mRNA Sequencing	Illumina Stranded mRNA Prep	<ul> <li>Cost-effective, scalable RNA sequencing of coding transcriptome with precise measurement of strand orientation</li> <li>Broad RNA input range, rapid protocol, cost-effective sequencing with up to 384 UDIs</li> <li>Wide dynamic range for more accurate measurement of gene expression</li> <li>Poly(A) capture, ligation-based addition of adapters and indexes</li> </ul>
Targeted RNA Sequencing	Illumina RNA Prep with Enrichment	<ul> <li>Exceptionally fast tagmentation-based library prep enables deep insights into many transcripts of interest, including the RNA exome</li> <li>Single hybridization step</li> <li>No mechanical shearing needed</li> <li>Benefit of panel modularity</li> <li>Compatible with low quality / degraded / FFPE tissue</li> <li>Respiratory virus detection with oligo panel</li> </ul>

https://www.illumina.com/techniques/sequencing/ngs-library-prep/rna.html