### Product Description:

sparQ RNA-Seq HMR Kit simultaneously depletes rRNA and globin mRNA while generating stranded RNA-seq libraries in 4.5 hours. In a single step and tube, RNA fragmentation and depletion of abundant ribosomal and globin transcripts (human, mouse, and rat) are integrated. The proprietary, highly optimized enzymes and streamlined workflow generate high quality, directional, transcriptome NGS libraries from either intact or degraded RNA samples, with key improvements for low input and FFPE samples.

### Component Part Numbers:

1126261 Frag Prime RG Depletion Mix  
1126264 1st Strand Enzyme Mix  
1126267 2nd Strand Buffer  
1126270 2nd Strand Enzyme Mix  
1126273 Rapid Ligation Buffer (5X)  
1126276 T4 DNA Ligase  
1126279 HiFi Plus Master Mix (2X)  
1126282 Primer Mix  
1129849 Bead Booster  
1126285 UDI Dilution Buffer

### Quality Control Analysis and Specifications:

**Library Prep Functional Assay:** Quality of the sparQ RNA-Seq HMR Kit is tested functionally by preparation of a cDNA library from high quality reference total RNA. The differences in library yield and profile among different lots must be within 15%. Sequencing of the amplified cDNA library must yield mapped reads >90%.

Enzyme components were tested prior to assembly and free of contaminating endonucleases and exonucleases. Enzyme purity was >95% as determined by SDS-PAGE and negligible *E. coli* genomic DNA contamination was confirmed by qPCR.