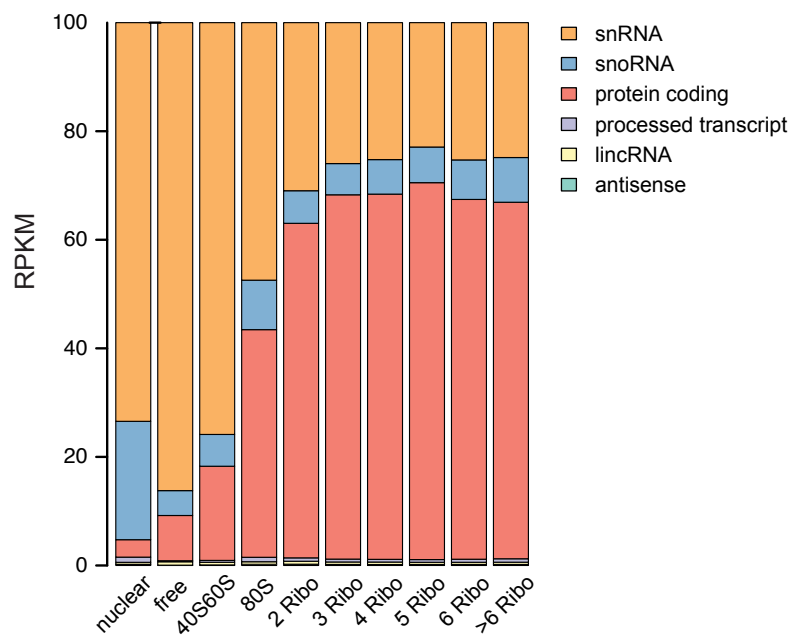




Additional file 4) Reads per kilobase per million (RPKM) normalized content of each sequenced fraction. Types of transcripts (both coding and noncoding) are indicated with different colors.