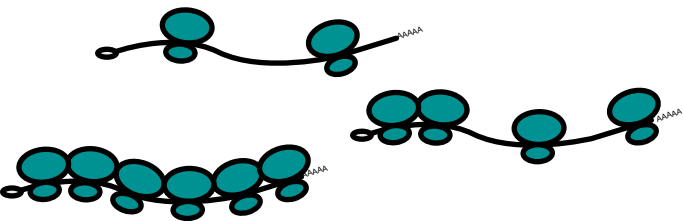
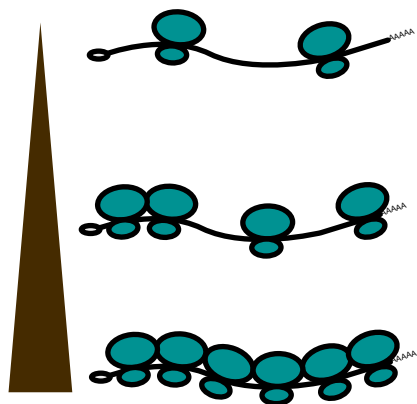


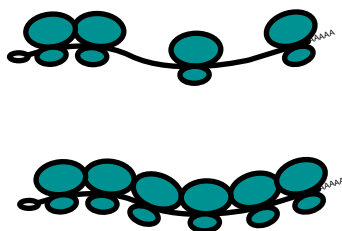
Polysome profiling



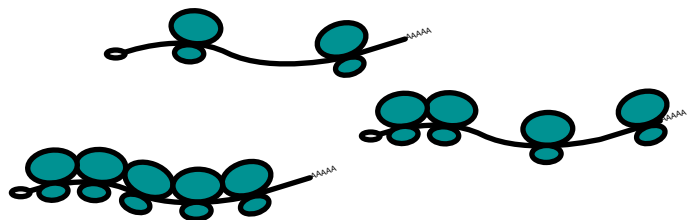
Separation of mRNAs based on the number of bound ribosomes



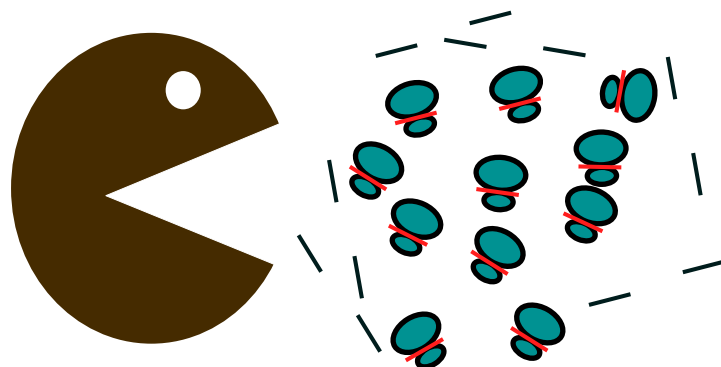
Extraction of polysome-associated mRNAs



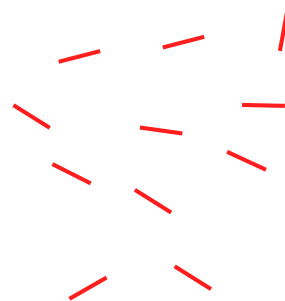
Ribosome profiling



RNA digestion to create ribosome protected fragments



Isolation of ribosome protected fragments



Quantification, e.g. via RNA sequencing