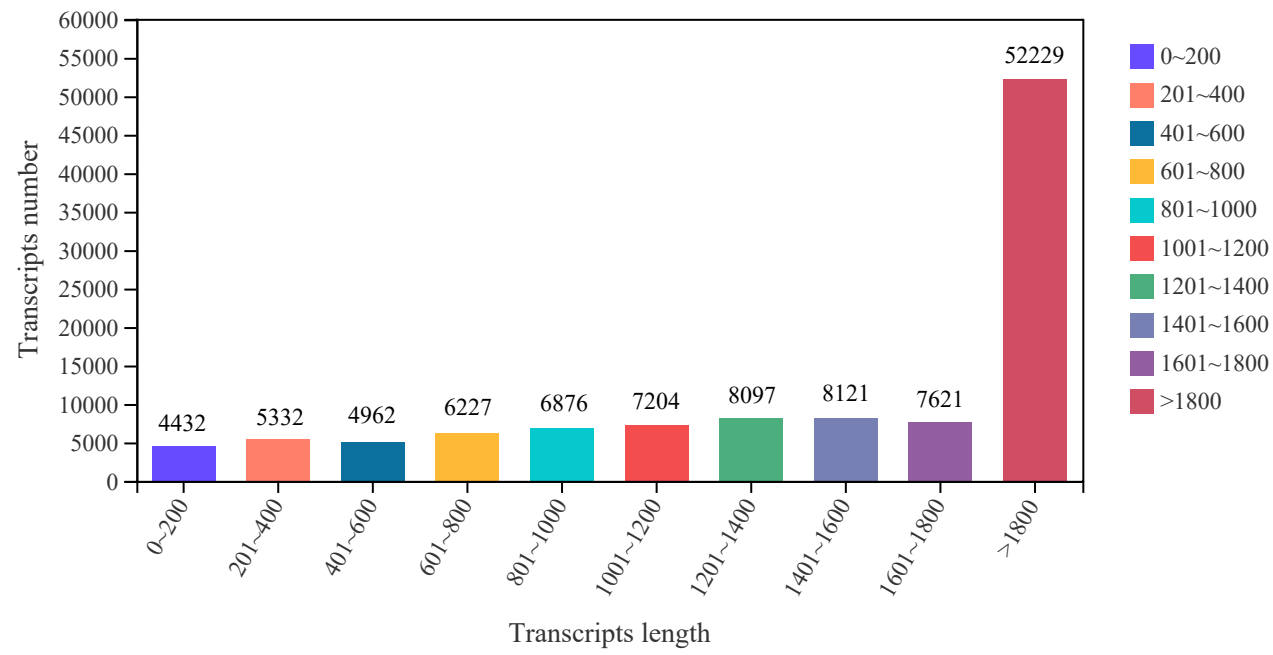


A

Length distribution of transcripts



B

New transcripts classification

