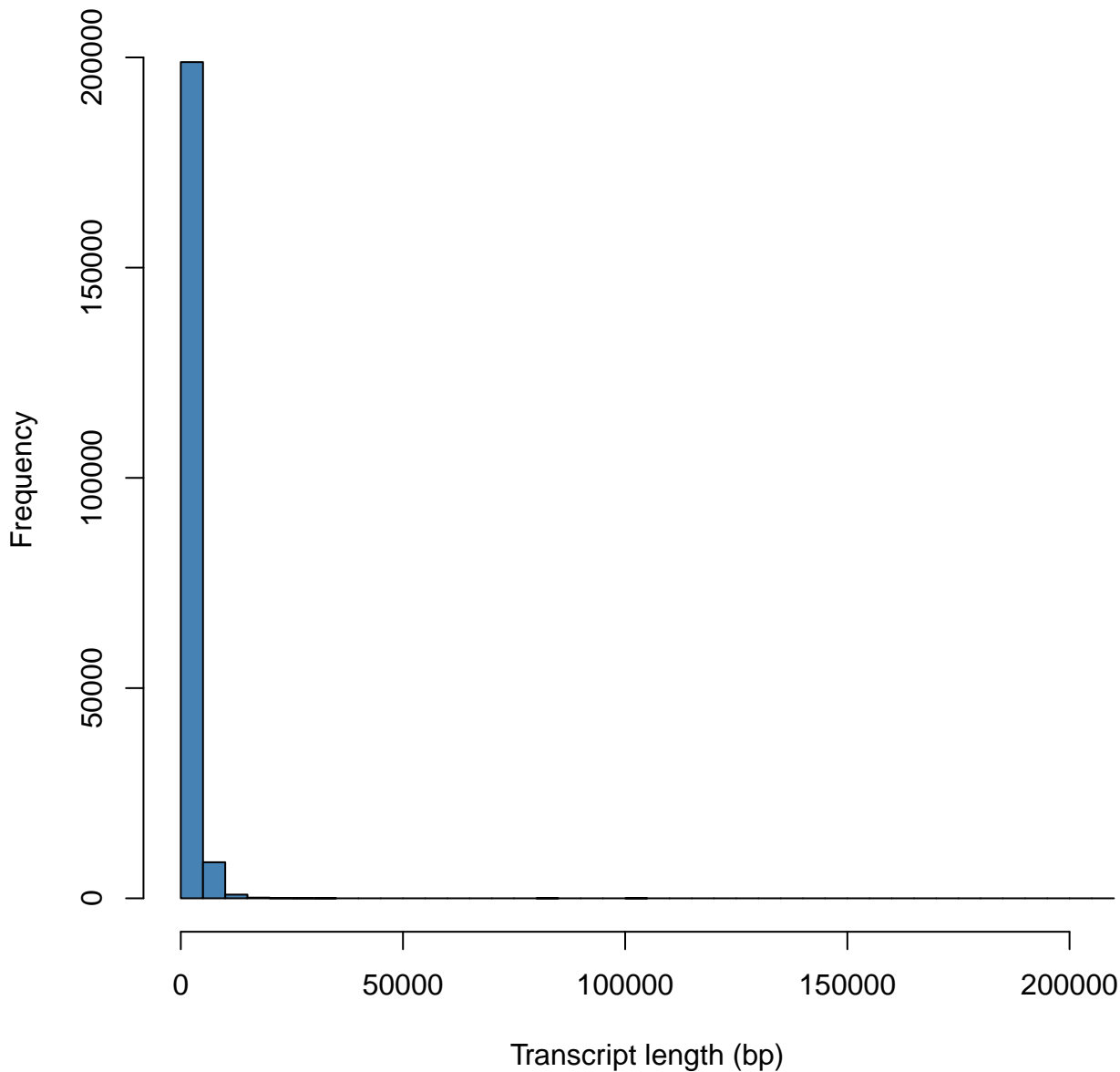


**Distribution of transcript lengths**



**Distribution of transcript count per gene**

