The figure displays the distribution of 5' boundary of poly(A) regions around TTS for two different cell lines: HeLa_B2_R1 (A) and HEK293T_B2_R1 (B). The distributions are shown for different categories: shortest 20%, next shorter 20%, middle 20%, next longer 20%, and longest 20%. The y-axis represents the number of poly(A)-containing reads, and the x-axis represents the distribution of the 5' boundary of poly(A) regions around TTS.