

The graph displays two data series: the ratio of reads with a 5' poly(A) tail (red line) and the ratio of reads with a 3' poly(A) tail (blue line). The x-axis represents the genomic position in base pairs (bp), ranging from 176964100 to 176964200. The y-axis represents the ratio, ranging from 0 to 1.0. The red line is consistently higher than the blue line, indicating a higher ratio of reads with a 5' poly(A) tail. Both lines show significant fluctuations, with peaks around 176964125 bp and 176964175 bp.

Position (bp)	Ratio of reads with a 5' poly(A) tail (red)	Ratio of reads with a 3' poly(A) tail (blue)
176964100	0.45	0.15
176964105	0.55	0.18
176964110	0.55	0.15
176964115	0.50	0.15
176964120	0.75	0.35
176964125	0.85	0.45
176964130	0.65	0.25
176964135	0.75	0.45
176964140	0.90	0.50
176964145	0.75	0.35
176964150	0.70	0.38
176964155	0.70	0.40
176964160	0.85	0.45
176964165	0.95	0.55
176964170	0.75	0.35
176964175	0.85	0.45
176964180	0.65	0.25
176964185	0.45	0.15
176964190	0.35	0.10

