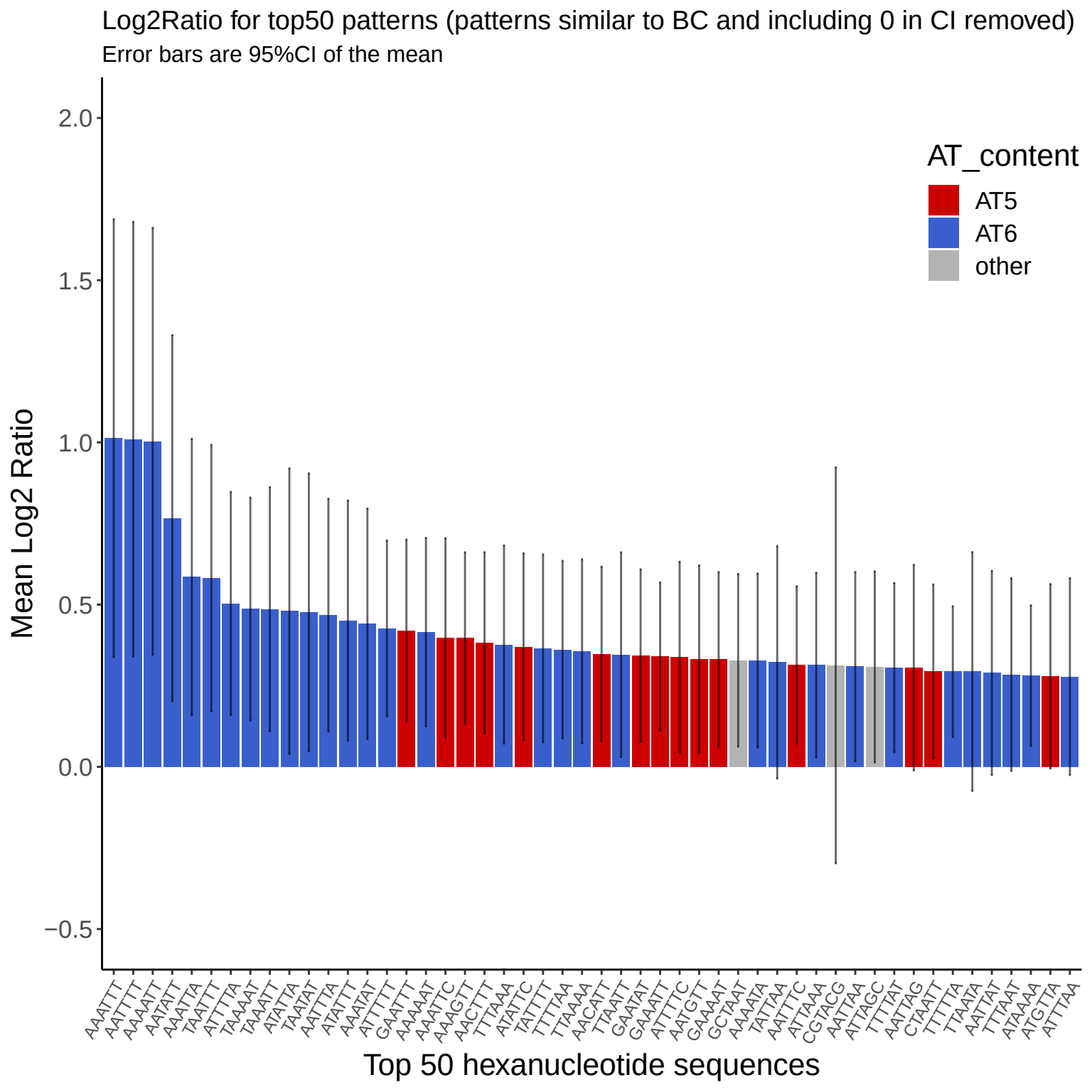
Log2Ratio for top50 patterns (patterns similar to BC and including 0 in CI removed)
Error bars are 95%CI of the mean
Mean Log2 Ratio
2.0
1.5
1.0
0.5
0.0
−0.5
AT_content
AT5
AT6
other
AAATTT
AATTTT
AAAATT
AATATT
AAATTA
TAATTT
ATTTTA
TAAAAT
TAAATT
ATATTA
TAATAT
AATTTA
ATATTT
AAATAT
ATTTTT
GAATTT
AAAAAT
AAATTC
AAAGTT
AACTTT
TTTAAA
ATATTC
TATTTT
TTTTAA
TTAAAA
AACATT
TTAATT
GAATAT
GAAATT
ATTTTC
AATGTT
GAAAAT
GCTAAT
AAAATA
TATTAA
AATTTC
ATTAAA
CGTACG
AATTAA
ATTAGC
TTTTAT
AATTAG
CTAATT
TTTTTA
TTAATA
AATTAT
TTTAAT
ATAAAA
ATGTTA
ATTTAA
Top 50 hexanucleotide sequences