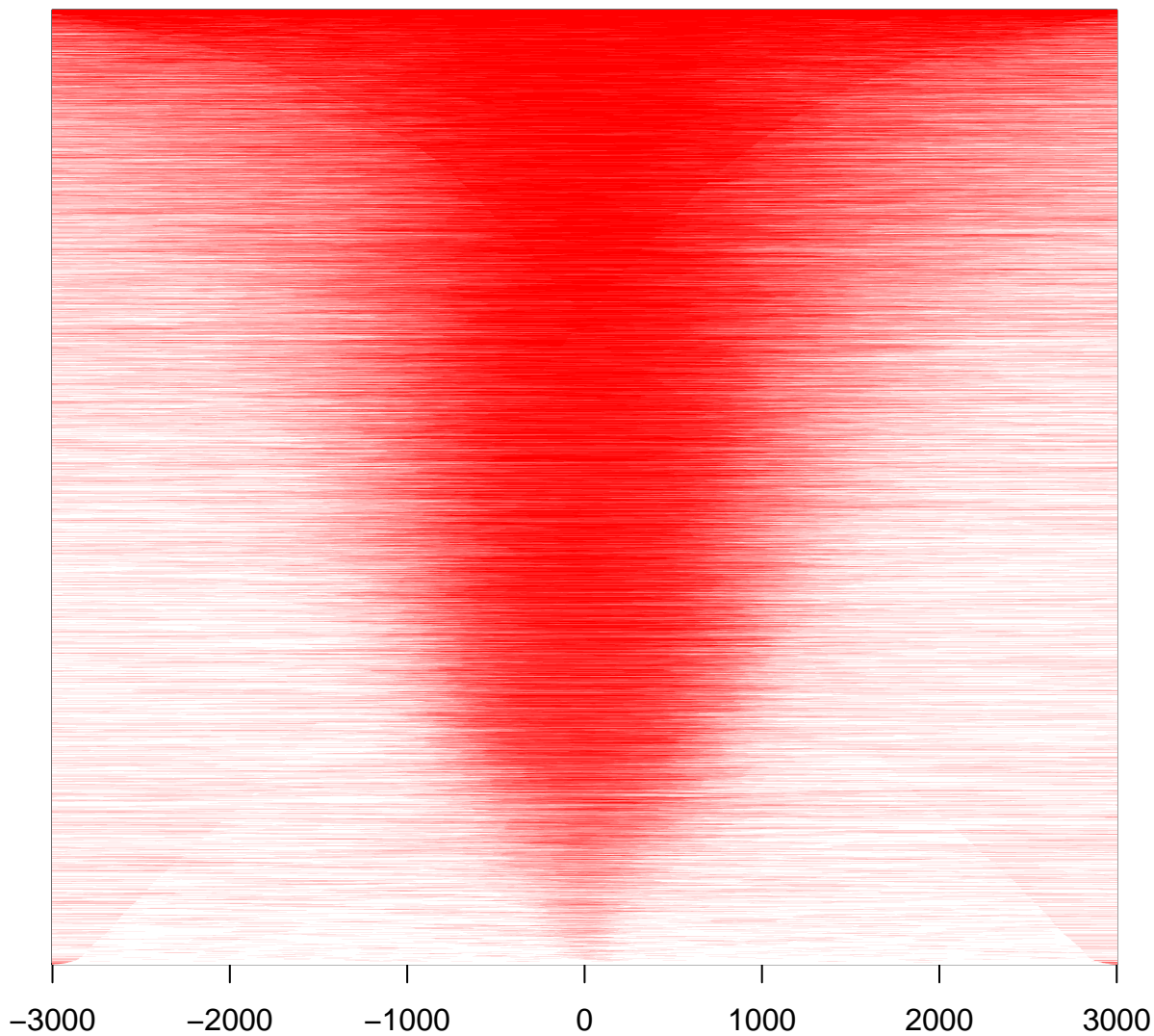
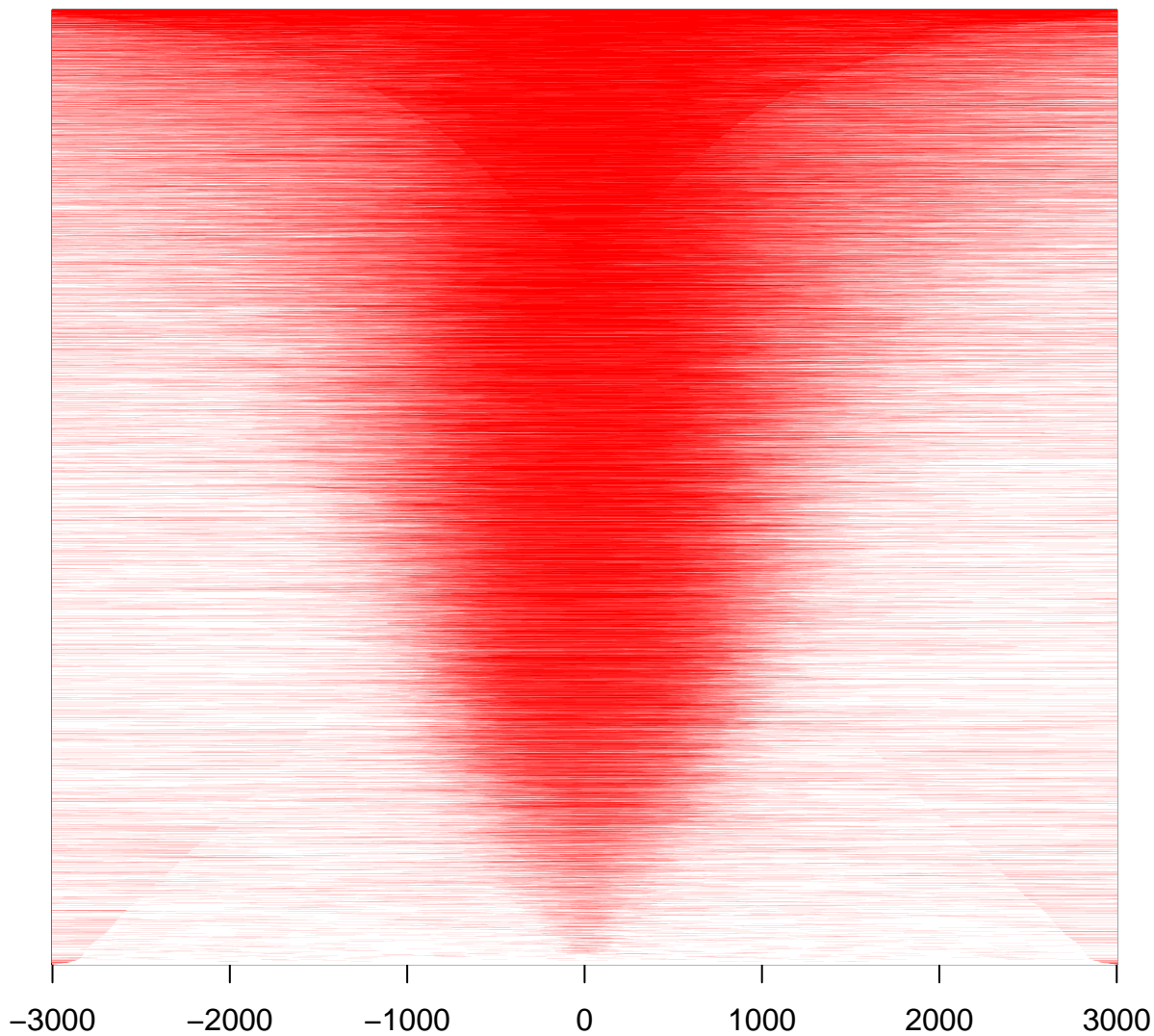


High_CD8



Low_CD8



CD8

Read Count Frequency

HighLow

High

Low

-2000

0

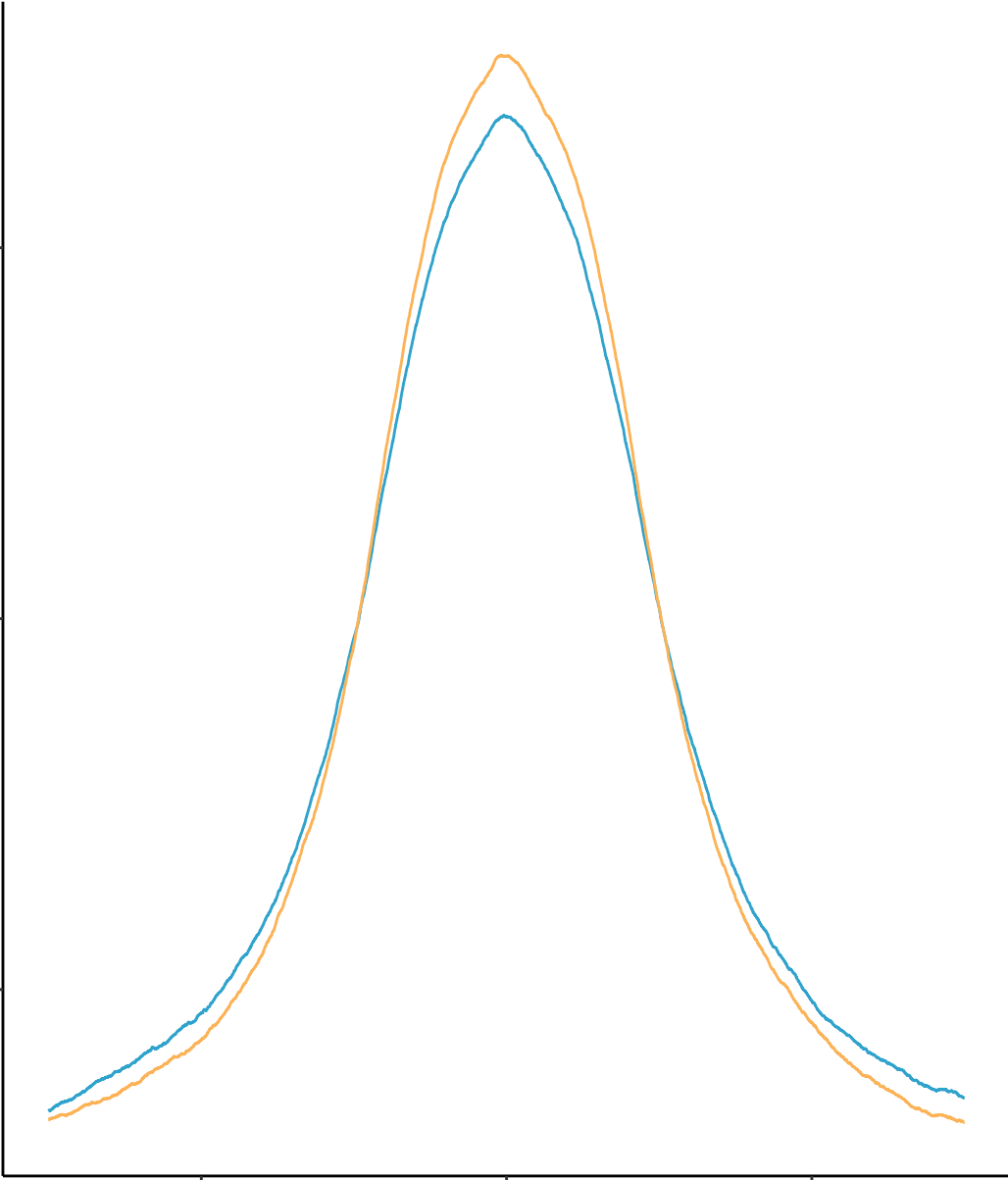
2000

Genomic Region (5'→3')

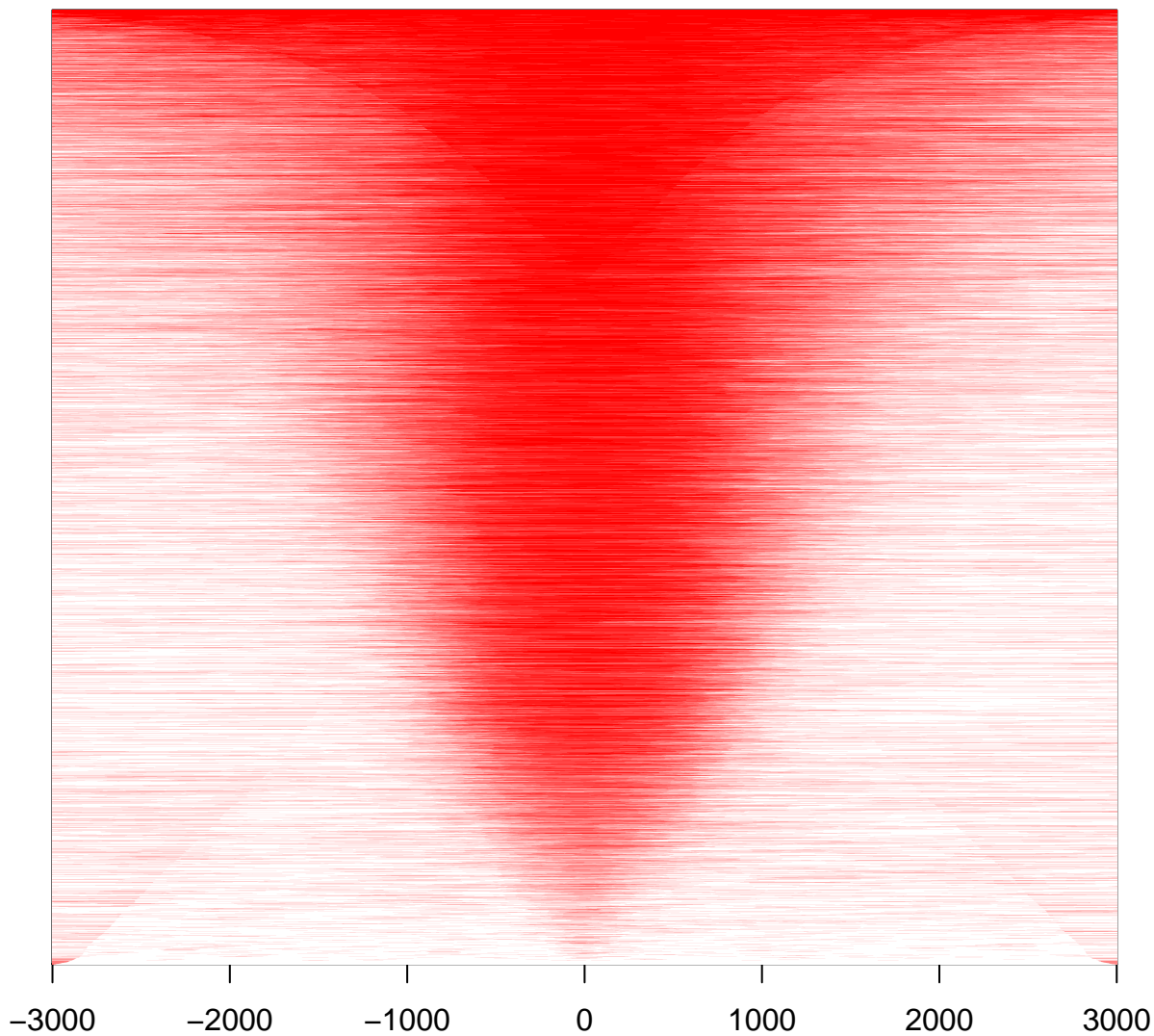
$3e-04$

$2e-04$

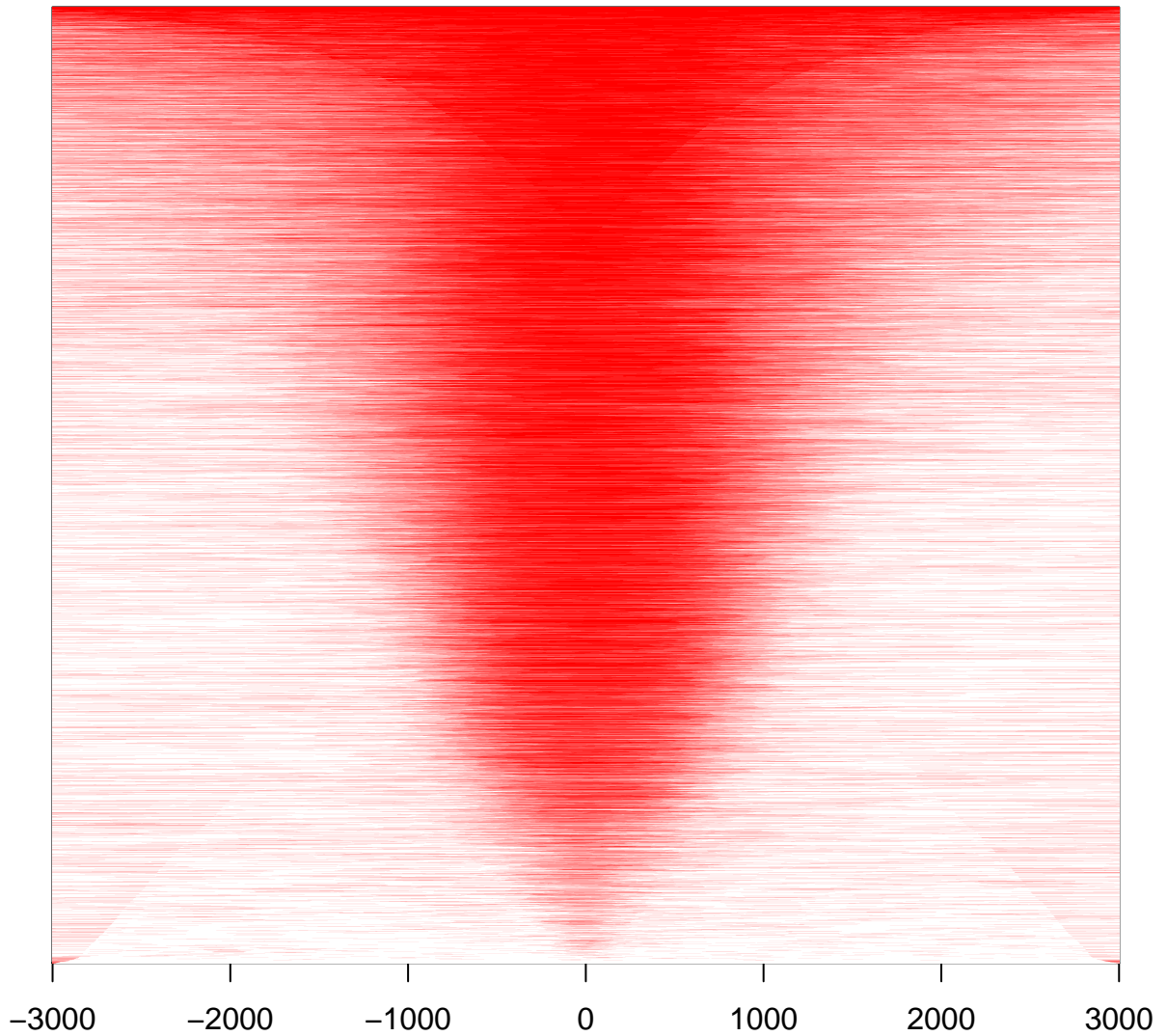
$1e-04$



High_CD4



Low_CD4



CD8

Read Count Frequency

HighLow

High

Low

-2000

0

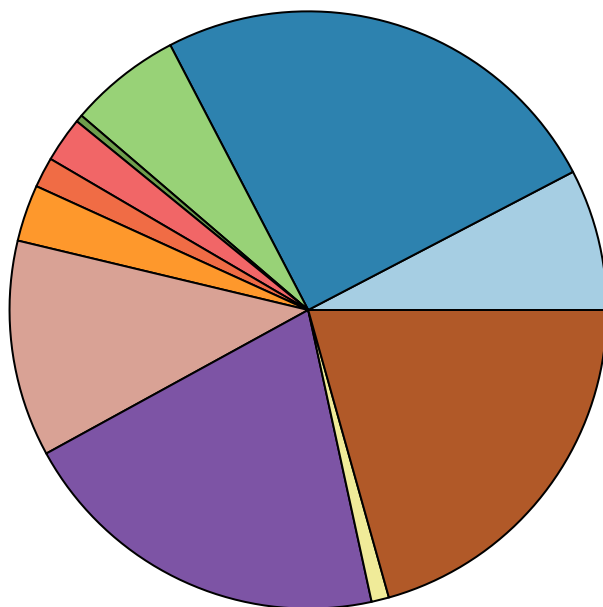
2000

Genomic Region (5'→3')

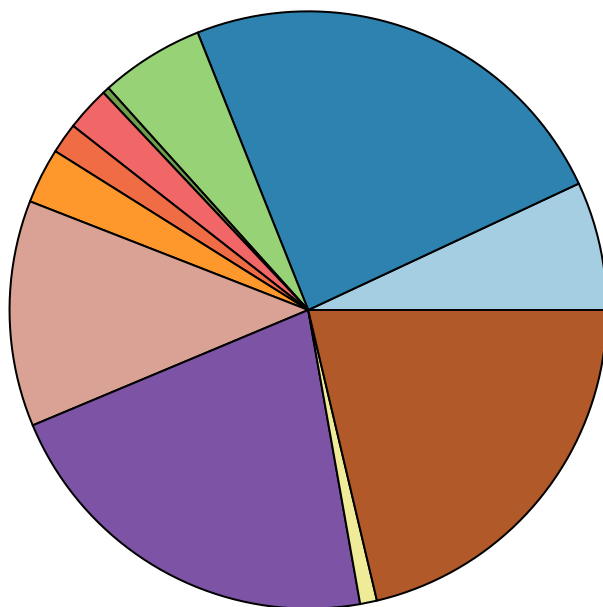
$3e-04$

$2e-04$

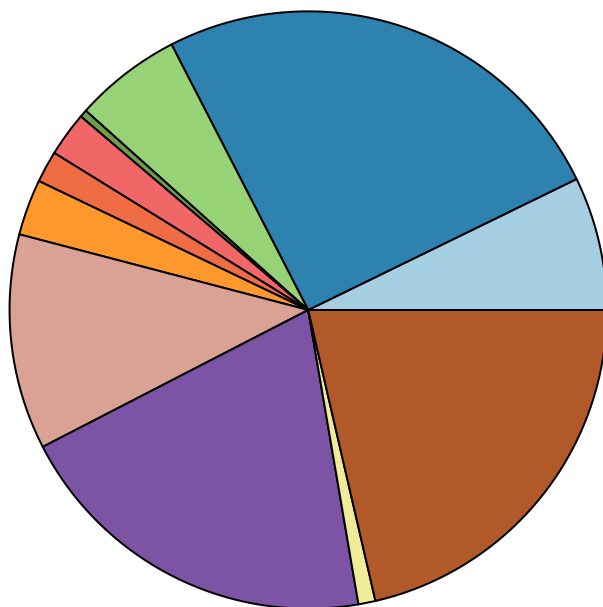
$1e-04$



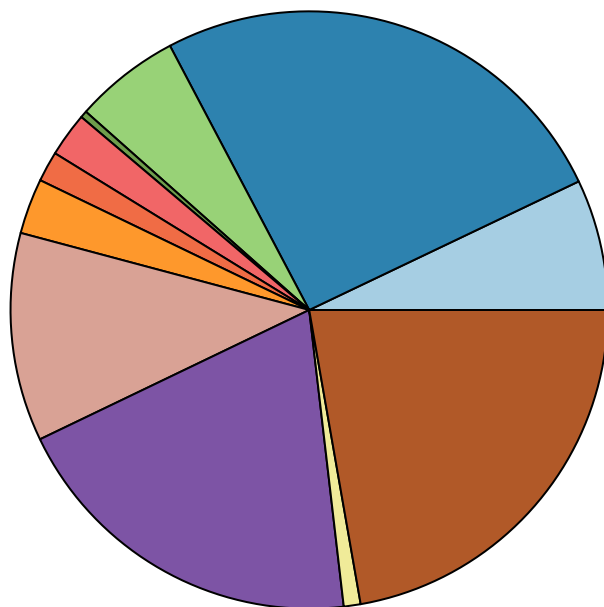
- Promoter (1-2kb) (7.65%)
- Promoter (<=1kb) (24.99%)
- Promoter (2-3kb) (6.08%)
- 5' UTR (0.39%)
- 3' UTR (2.45%)
- 1st Exon (1.64%)
- Other Exon (3.07%)
- 1st Intron (11.71%)
- Other Intron (20.43%)
- Downstream (<=300) (0.95%)
- Distal Intergenic (20.64%)



- Promoter (1–2kb) (6.96%)
- Promoter (≤ 1 kb) (24.08%)
- Promoter (2–3kb) (5.63%)
- 5' UTR (0.37%)
- 3' UTR (2.4%)
- 1st Exon (1.65%)
- Other Exon (3%)
- 1st Intron (12.24%)
- Other Intron (21.48%)
- Downstream (≤ 300) (0.91%)
- Distal Intergenic (21.3%)

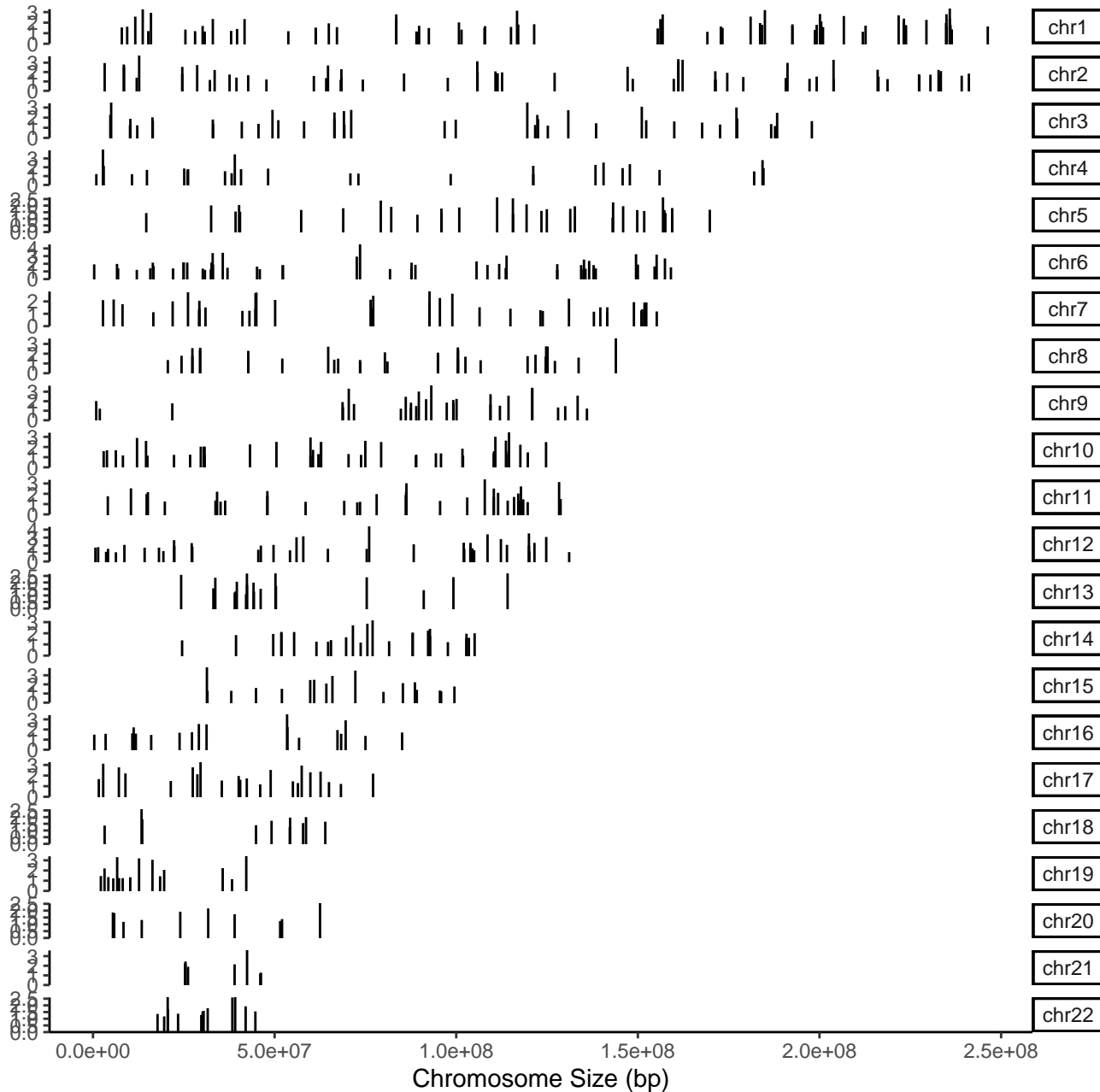


- Promoter (1–2kb) (7.23%)
- Promoter (<=1kb) (25.34%)
- Promoter (2–3kb) (5.81%)
- 5' UTR (0.41%)
- 3' UTR (2.41%)
- 1st Exon (1.71%)
- Other Exon (3.03%)
- 1st Intron (11.65%)
- Other Intron (20.12%)
- Downstream (<=300) (0.92%)
- Distal Intergenic (21.38%)



- Promoter (1–2kb) (7.09%)
- Promoter (<=1kb) (25.63%)
- Promoter (2–3kb) (5.72%)
- 5' UTR (0.37%)
- 3' UTR (2.39%)
- 1st Exon (1.64%)
- Other Exon (3%)
- 1st Intron (11.28%)
- Other Intron (19.72%)
- Downstream (<=300) (0.91%)
- Distal Intergenic (22.24%)

CD8



CD4

