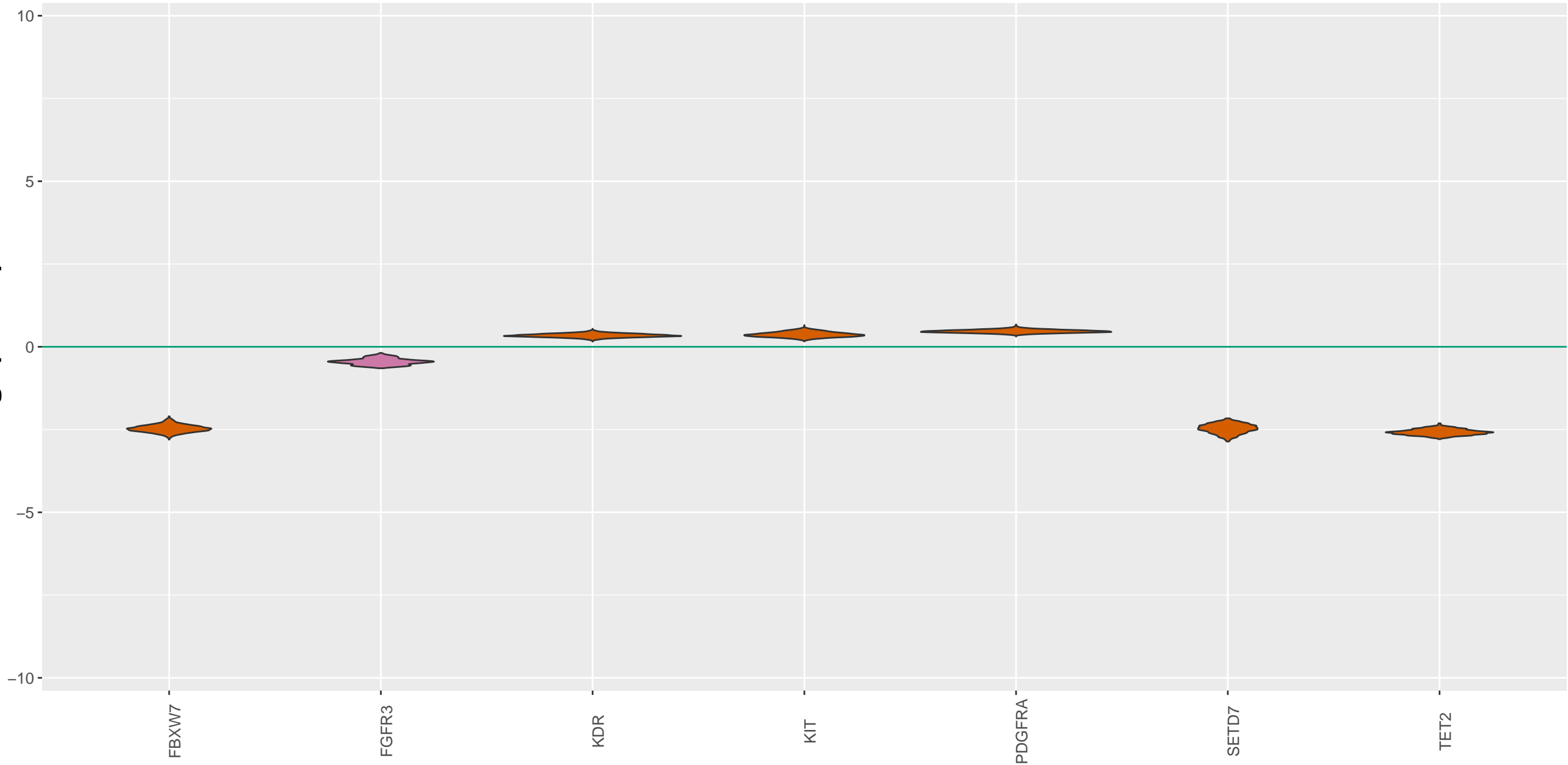


T394Biopsy_aligned.bam

log2(ratios)



CNV Reliability

- NonReliableChange
- ReliableChange

Gene Names