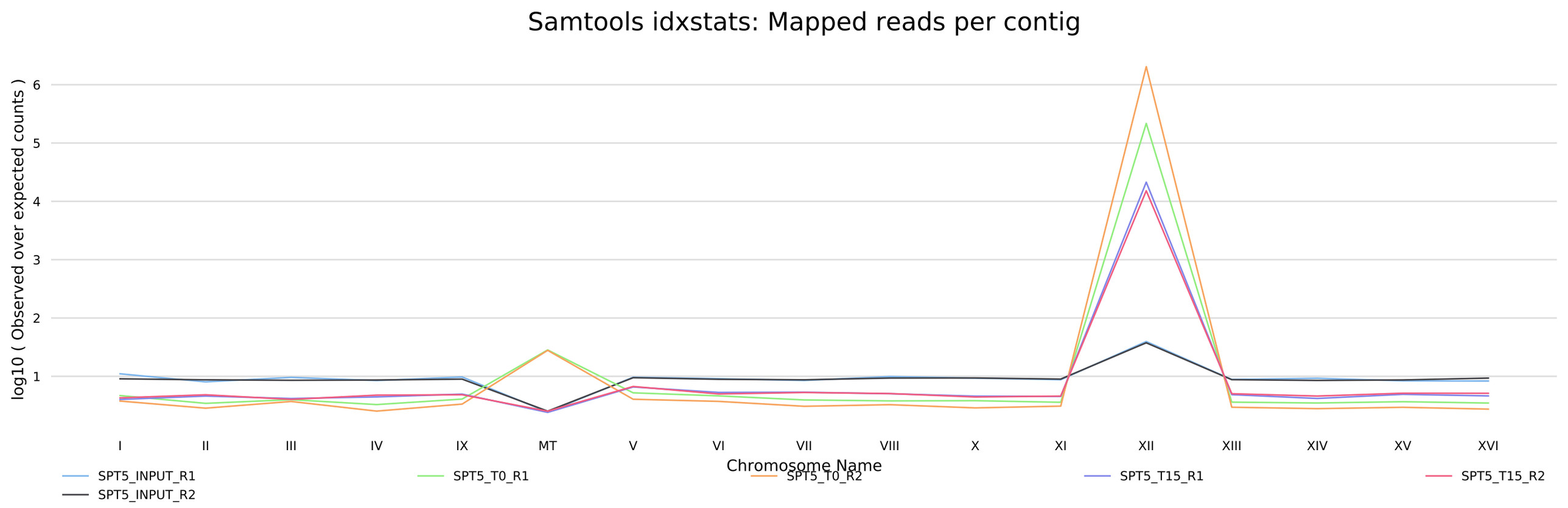
Samtools idxstats: Mapped reads per contig
log10 (Observed over expected counts)
6
5
4
3
2
1
I
II
III
IV
IX
MT
V
VI
VII
VIII
X
XI
XII
XIII
XIV
XV
XVI
Chromosome Name
SPT5_INPUT_R1
SPT5_INPUT_R2
SPT5_T0_R1
SPT5_T0_R2
SPT5_T15_R1
SPT5_T15_R2