Information:

**Bed file:**

**chrom** - The name of the chromosome (e.g. chr3, chrY, chr2\_random) or scaffold (e.g. scaffold10671).

**chromStart** - The starting position of the feature in the chromosome or scaffold. The first base in a chromosome is numbered 0.

**chromEnd** - The ending position of the feature in the chromosome or scaffold. The chromEnd base is not included in the display

of the feature. For example, the first 100 bases of a chromosome are defined as chromStart=0, chromEnd=100, and span the bases

numbered 0-99.

Source: https://genome.ucsc.edu/FAQ/FAQformat#format1

chr22 1000 5000

chr22 2000 6000