```
browser position chrY:104002-867875
track name="chromosomeY" description="chromosome Y" visibility=2 color=0,180,0 useScore=1
chrY    293529 867875 250388 450
chrY    104002 239292 280673 450
chrY    732030 744392 287905(3/3)    950
```

EXPLANATION:
browser position --> it's just the position initially displayed (if you don't write anything, the most recently setting position will be displayed; then you can move from this position to the interesting position)
track name --> name
description --> not necessary
color --> indicates that the personal track data (track name, region IDs, etc...) are highlighted in green color
useScore=1 --> indicates that the user will set a color score to display the different features in different shades of grey (white-grey-black).
If you wrote just "useScore=1" the browser will give automatically a shade of grey to each features: if you want display all your features in just two colors (for example for dividing all the features in two groups), you can write a specific number (es. 450 = grey) near to the regions which belong to one group, and similarly you can write another number (950 = black) near to the regions which belong to the other group.

Regions: CHR   START   END   ID(region-name)   ColorScore