Examples to adjust the output map

The following examples shows some options to tune the final map.

Change the ID position into "alternating".
> msb (M = atxn2, start = 111950277, end = 112036294, IDPos='alternating')

[Diagram showing the chromosome with SNP markers]

Rotate the direction of SNP ID label into vertical.
> msb (M = atxn2, start = 111950277, end = 112036294, rotation.item = 90, IDWd = 0.1)

[Diagram showing the chromosome with SNP markers rotated]
Change background color and background color of track title.

> msb (M = atxn2, start = 111950277, end = 112036294, rotation.item = 90, IDWd = 0.1,
   fontcolor.title = 'yellow', background.title = "lightblue")

Change background color of track title into "brown" and background color of track title into "white".

> msb (M = atxn2, start = 111950277, end = 112036294, rotation.item = 90, IDWd = 0.1,
   fontcolor.title = 'brown', background.title = "white")

Change the gene track name into "ATXN2".
Highlight the sixth SNP using red color.

```r
> color.snp = rep('black', 7); color.snp[6] = 'red'
```

```r
> msb (M = atxn2, start = 111950277, end = 112036294, rotation.item = 90, IDWd = 0.1,

geneName = 'ATXN2',

color.snp = color.snp)
```
Add more fine-grained tick marks, and change the positioning of the axis labels into "alternating".

> msb (M = atxn2, start = 111950277, end = 112036294, rotation.item = 90, IDWd = 0.1, littleTicks = TRUE, labelPos.axis = "alternating")