| ± = Epigenome intensity percentage scale: 0% | 100% PC1: -35.9 42.9 TADs: GeneHancer   | •   |
|--|---|---|
| 222.91Mb                                     | TLR5_1: -6.87   | 223.71Mb<br>promoter/enhancer   |
|  | Promoters and enhancers are shown in the top track, which are coloured according their PCs. The rest of the track are | Bcell-H3K27ac Bcell-H3K27ac Bcell-H3K27ac CLL-H3K27ac CLL-H3K27ac CLL-H3K27ac CLL-H3K27ac CLL-H3K27ac                               |
|  | coloured in different topologically associating domains (TADs).   | CLL-H3K27ac             |
|  |   | Bcell-H3K4me1<br>Bcell-H3K4me1<br>Bcell-H3K4me1<br>Bcell-H3K4me1  |
|  |   | CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1 |
|  |   | CLL-H3K4me1<br>CLL-H3K4me1<br>CLL-H3K4me1   |