

WashU Epigenome Browser - Bug # 87

Status:	Closed	Priority:	Normal
Author:	Xin Zhou	Category:	
Created:	10/31/2013	Assignee:	
Updated:	11/18/2013	Due date:	
Subject:	SNP track showing all SNPs to be on reverse strand		
Description	happens to all SNPs in hg19 SNP track		

History

#1 - 10/31/2013 10:46 am - Xin Zhou

- Status changed from New to Resolved

Fixed on public server, still visible on test server

Problem is that SNP track use +/- to indicate SNPs while the browser only accepts >/<

Now the browser accepts both marks, but for annotation track only

#2 - 11/18/2013 10:33 pm - Daofeng Li

- Status changed from Resolved to Closed

- could observe SNPs in reverse strand!