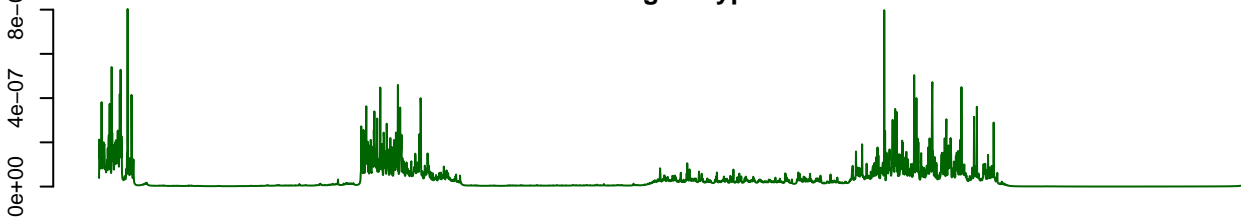
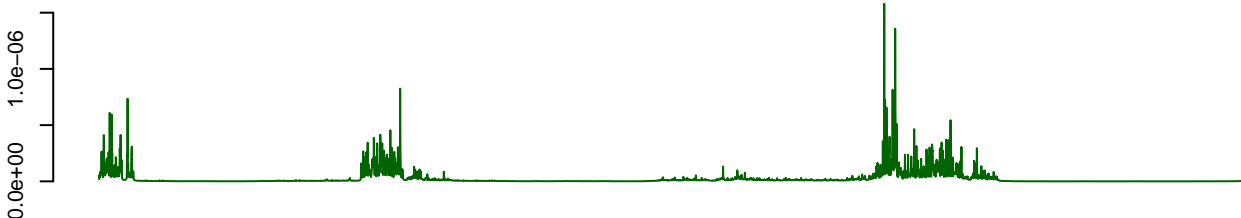


**baselinegenotype AA**



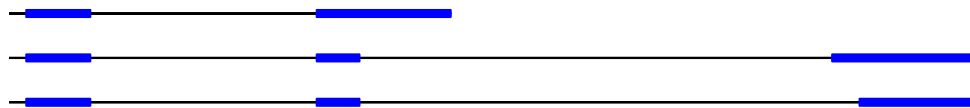
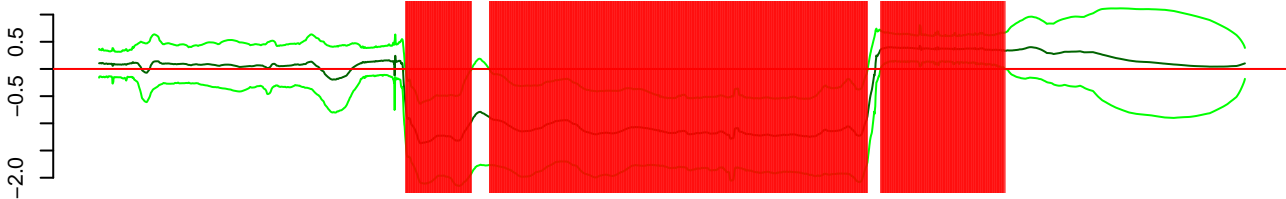
**baselinegenotype AG**



**baselinegenotype GG**



**effect +/- 2 s.d.**



Position (Mb) on chr12  
113.355 113.356 113.357 113.358