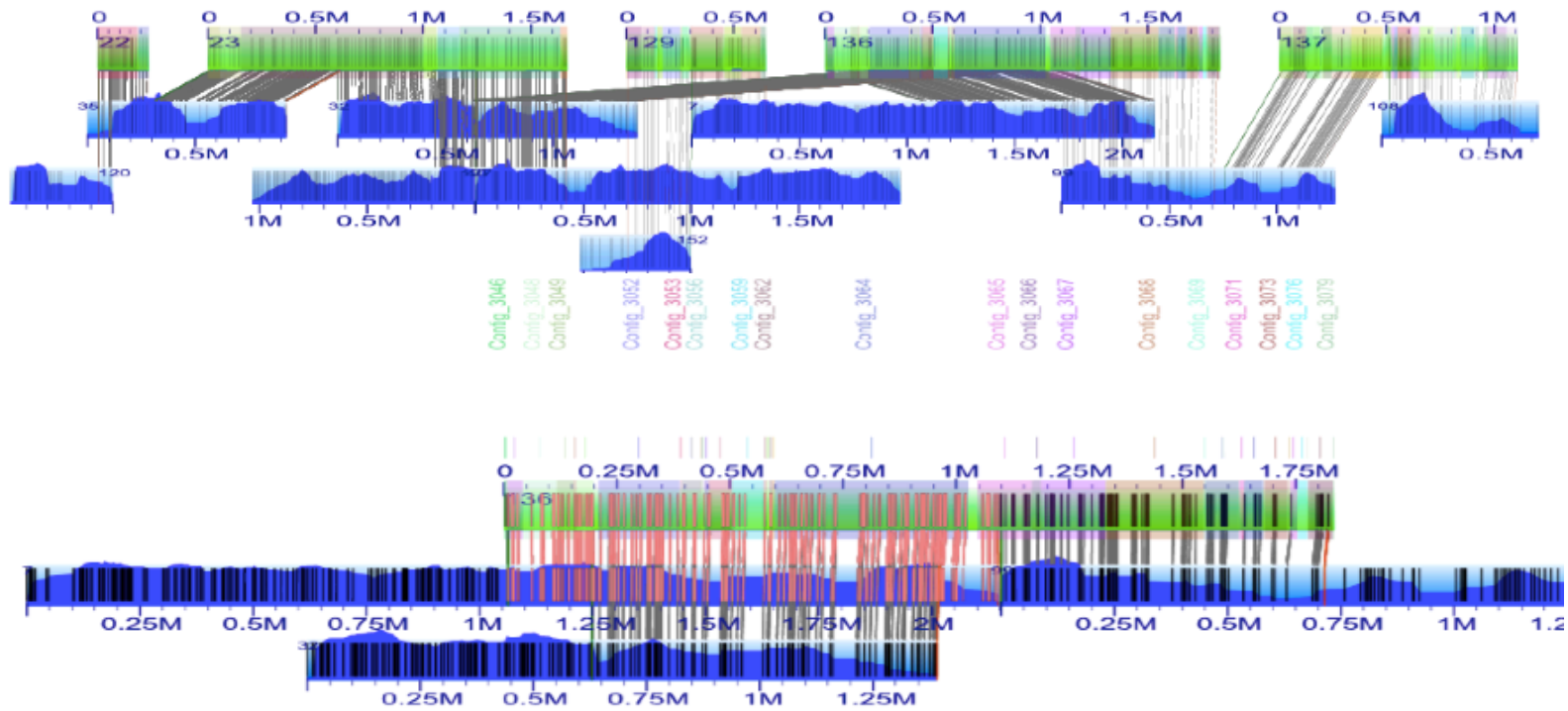


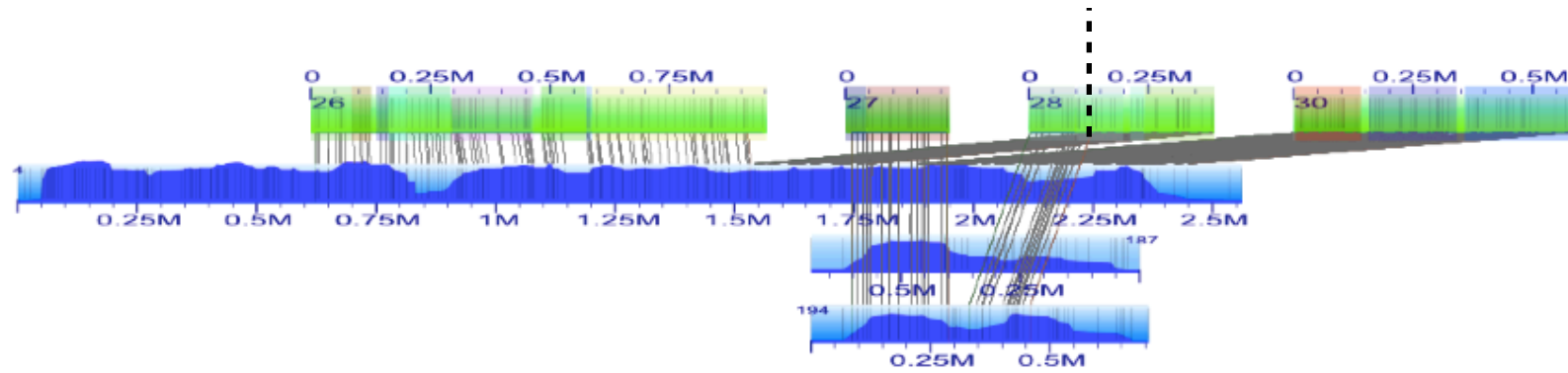
Is part of scaffold\_23 connected to 136?

I went with the second alignment (21-26 together and 136-137 together because it is supported by genetic maps) but we should check these assemblies.

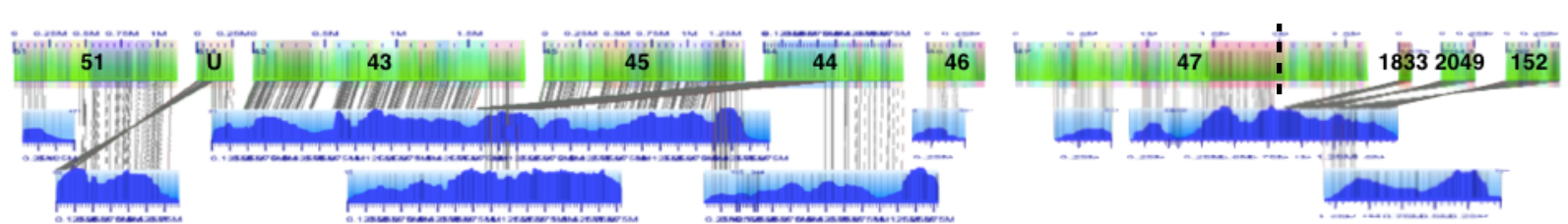
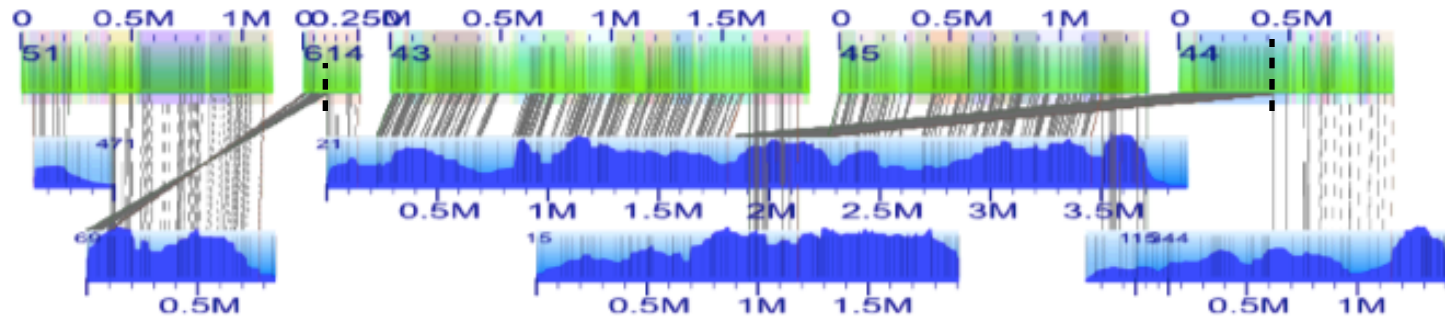
In bottom alignment of 136 you can see that a large section of the BNG map 32 (which joins 23 to 136) is a duplicate in the BNG assembly?



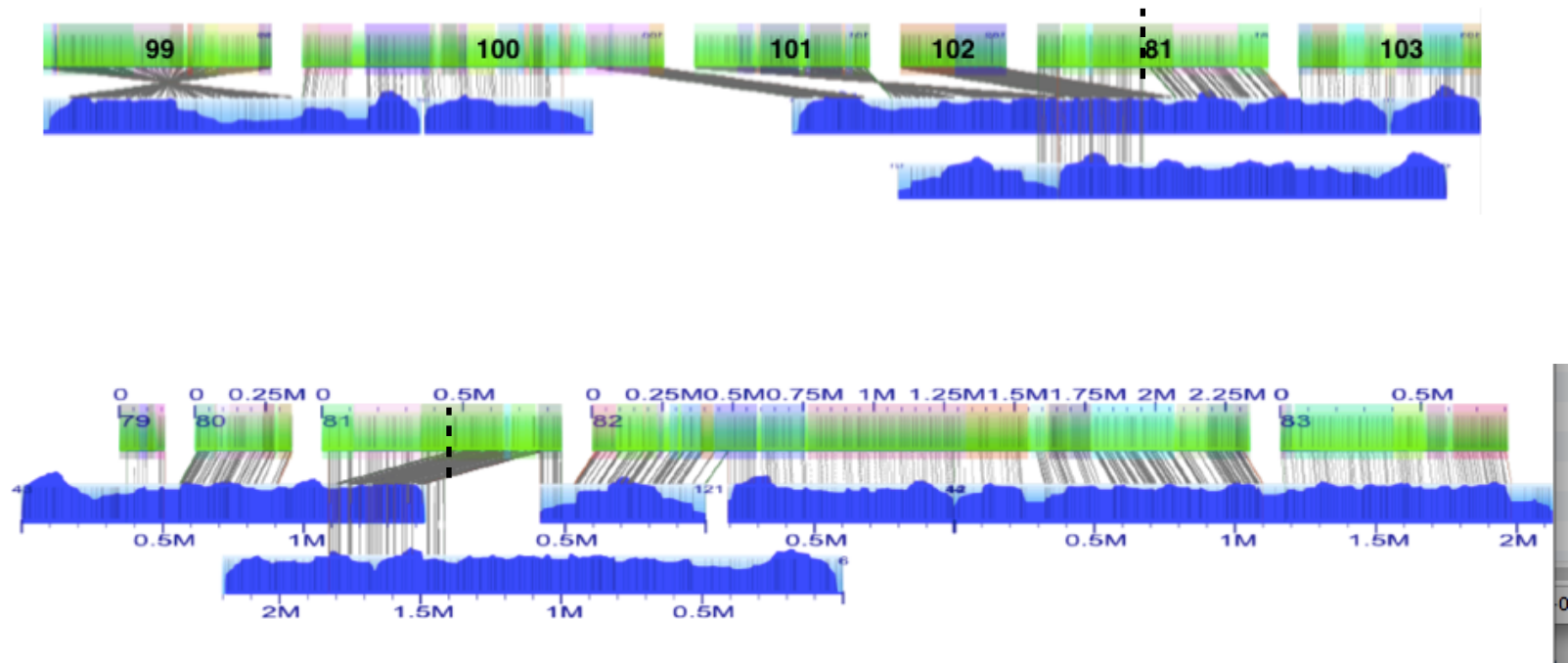
Does 28 need to be broken ?



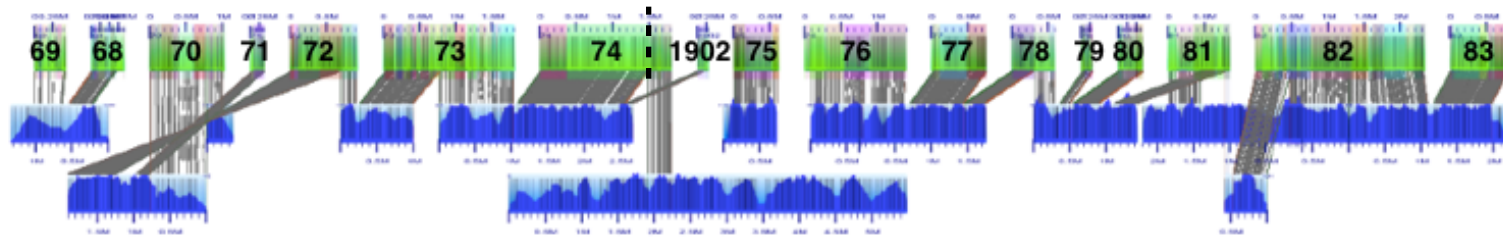
scaffold\_614 looks like it may need to be split part appears to be within scaffold\_51 and half could be linked with a gap to 43?  
scaffold\_44 looks like it should be split half going between scaffold\_43 and scaffold\_45 and half on its own?  
scaffold\_47 looks like it should be split and joined with scaffold\_1833 scaffold\_2049 scaffold\_152



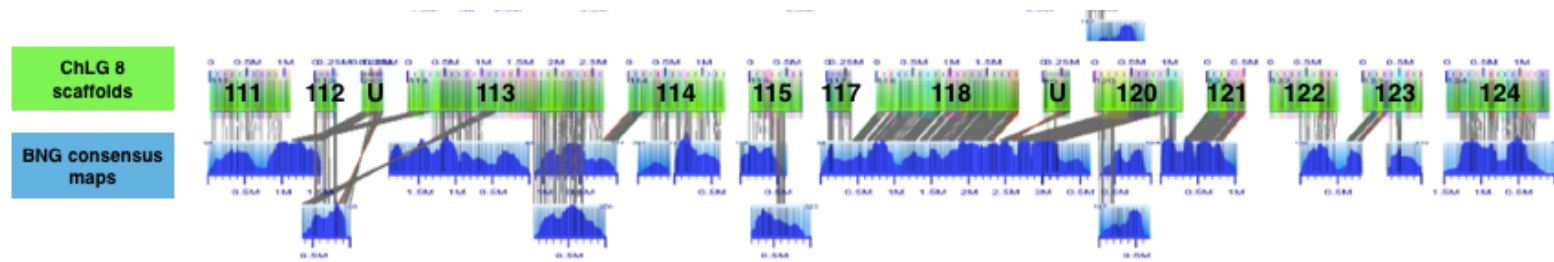
Half of scaffold\_81 aligns with ChLG7



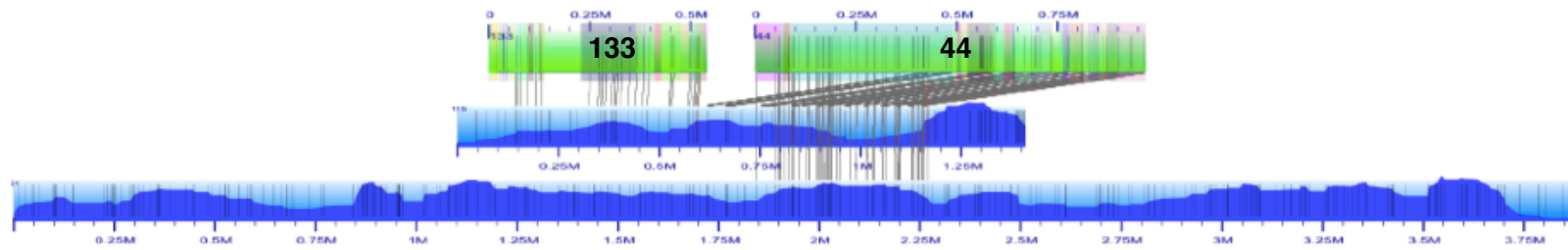
does scaffold\_74 need to be broken and part of it joined with scaffold\_1902?



scaffold\_118, scaffold\_86 and scaffold\_120 seem scrambled ?



Does part of scaffold\_44 belong with scaffold 133?



Does scaffold\_155 overlap unplaced scaffold\_424 ??

