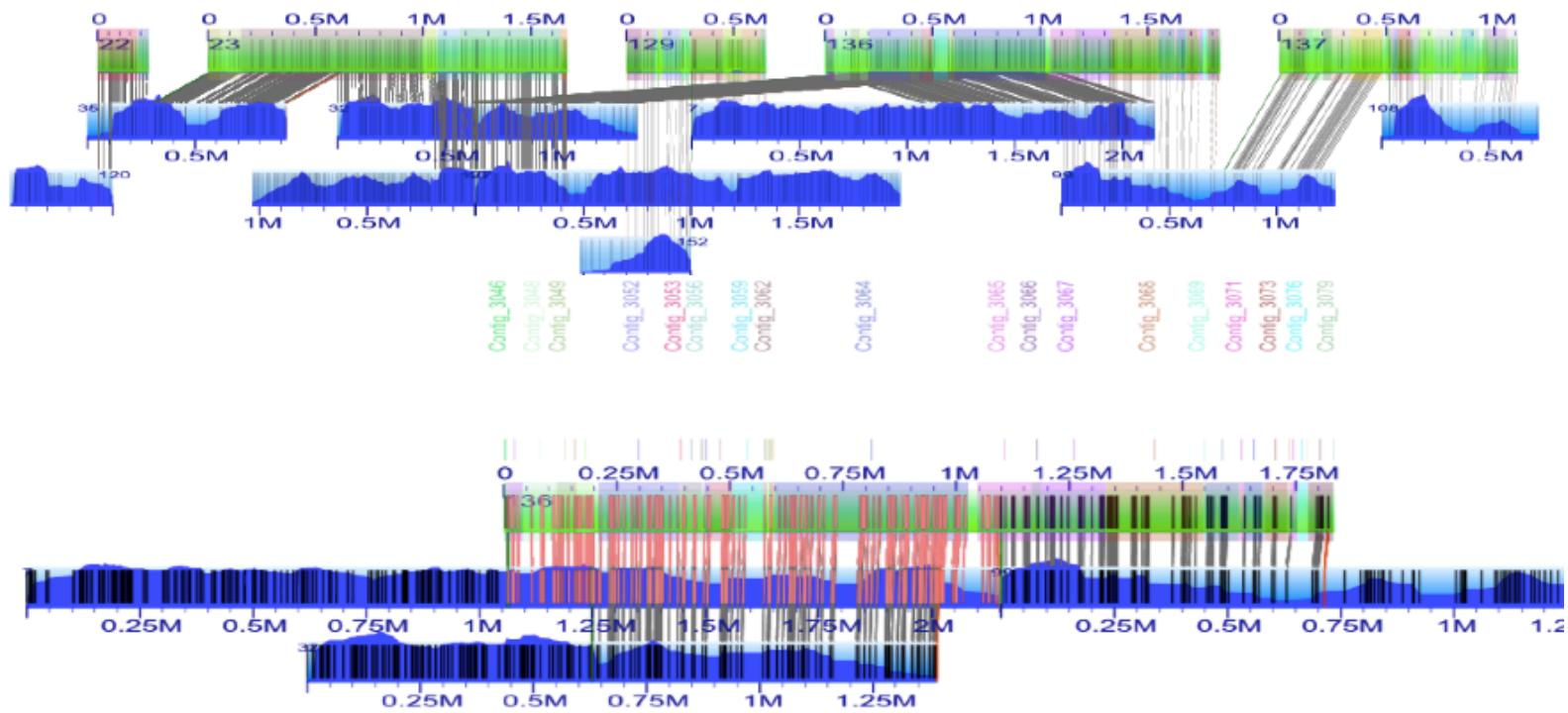


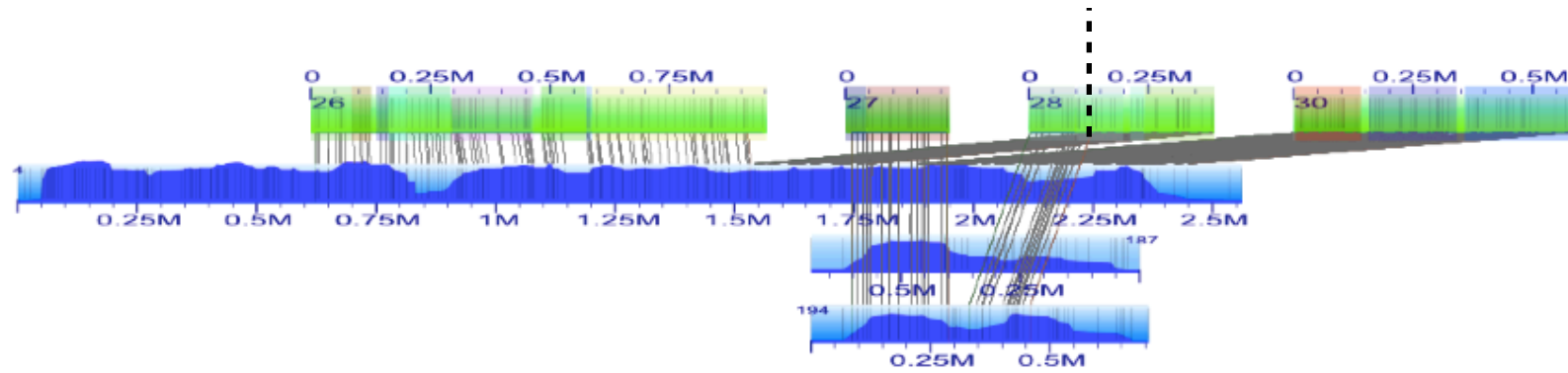
Is part of scaffold_23 connected to 136?

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.

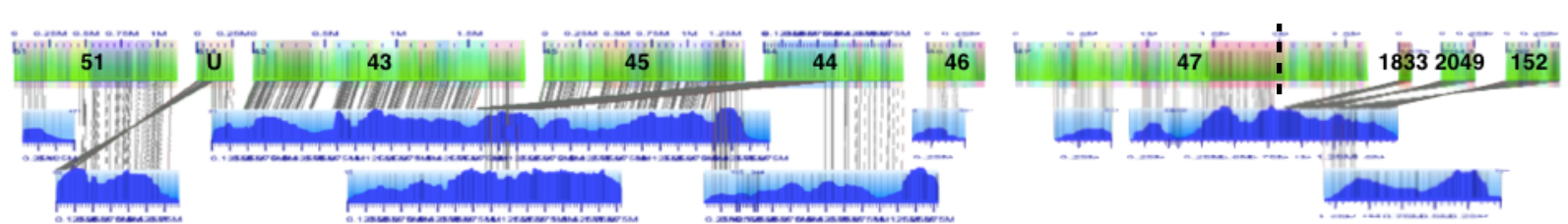
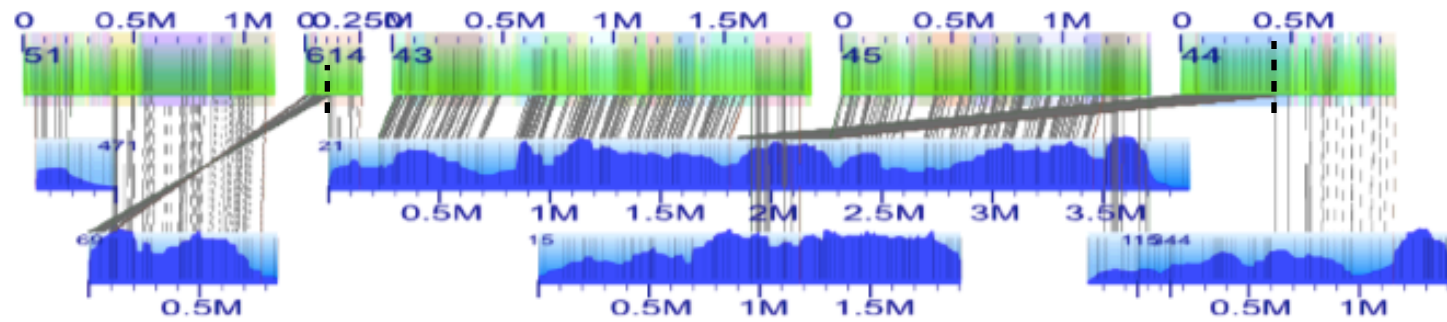
In bottom alignment of 136 you can see that a large section of the BNG map 32 (which joins 23 to 136) is a duplicate in the BNG assembly?



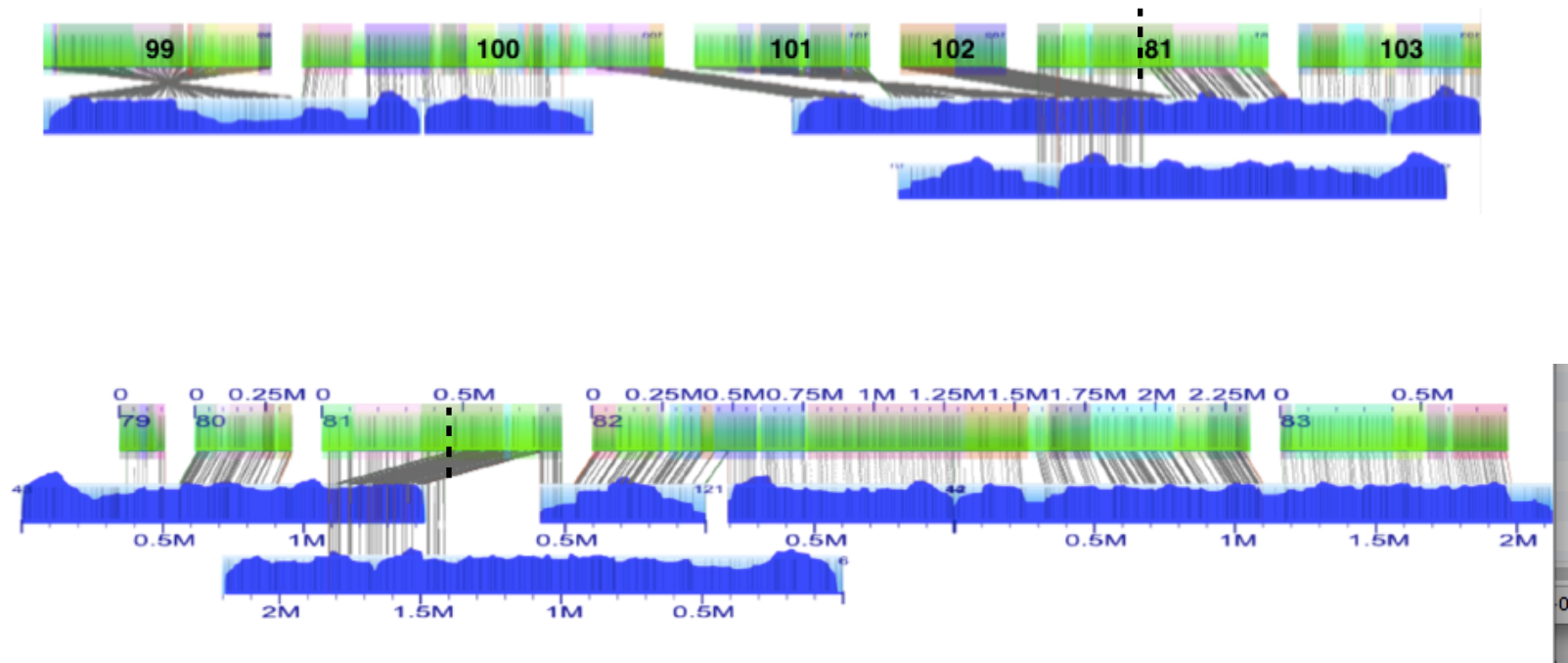
Does 28 need to be broken ?



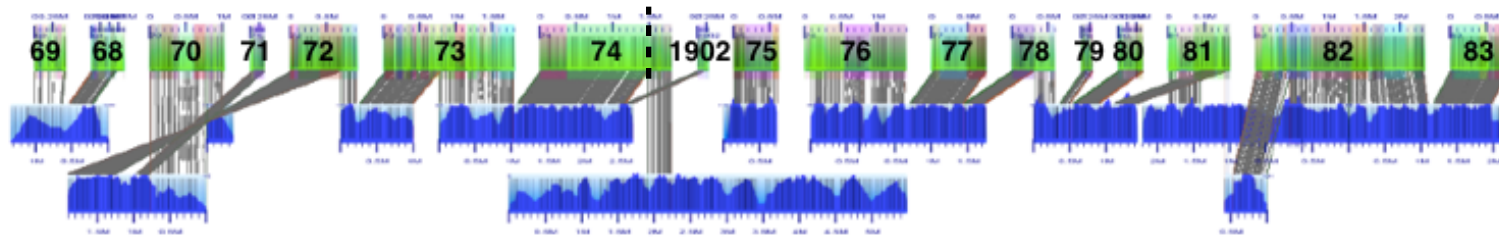
scaffold_614 looks like it may need to be split part appears to be within scaffold_51 and half could be linked with a gap to 43?
scaffold_44 looks like it should be split half going between scaffold_43 and scaffold_45 and half on its own?
scaffold_47 looks like it should be split and joined with scaffold_1833 scaffold_2049 scaffold_152



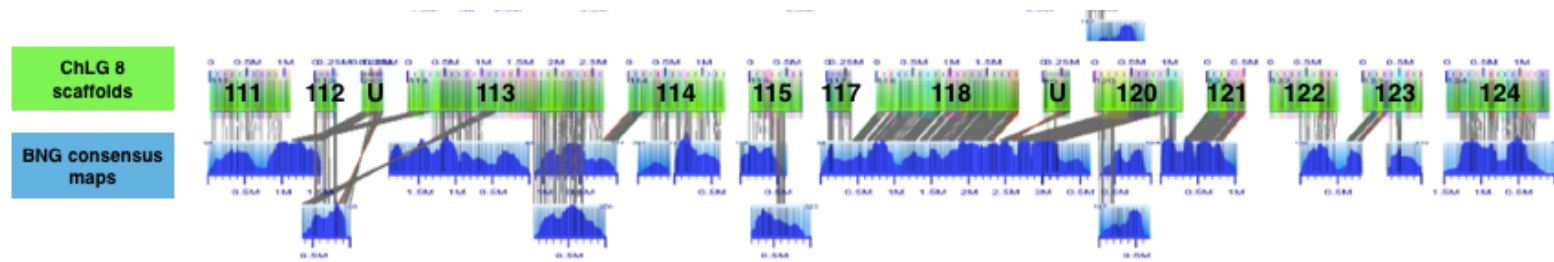
Half of scaffold_81 aligns with ChLG7



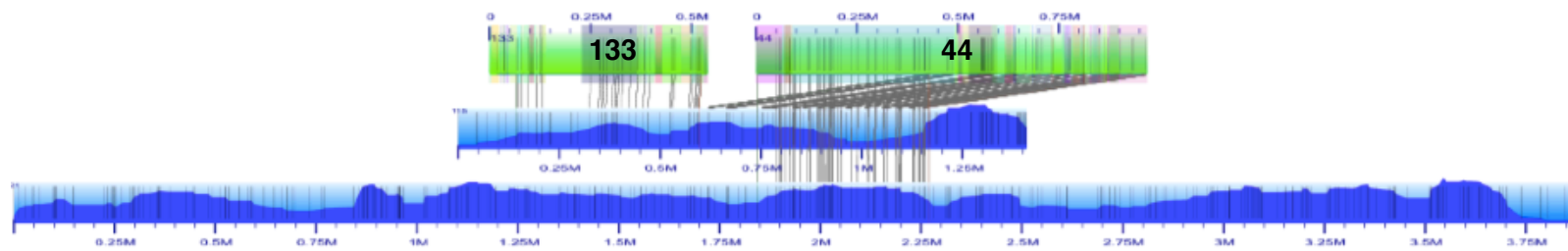
does scaffold_74 need to be broken and part of it joined with scaffold_1902?



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 ??

