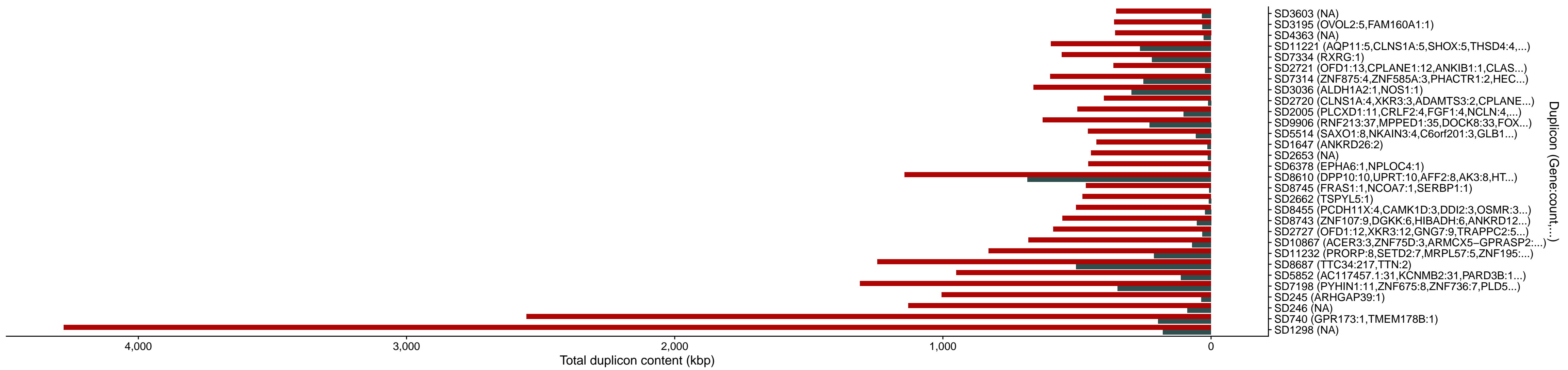


30 duplicons with largest change between  
T2T CHM13 and GRCh38



**b** SDs unannotated by DupMakser (Red)

