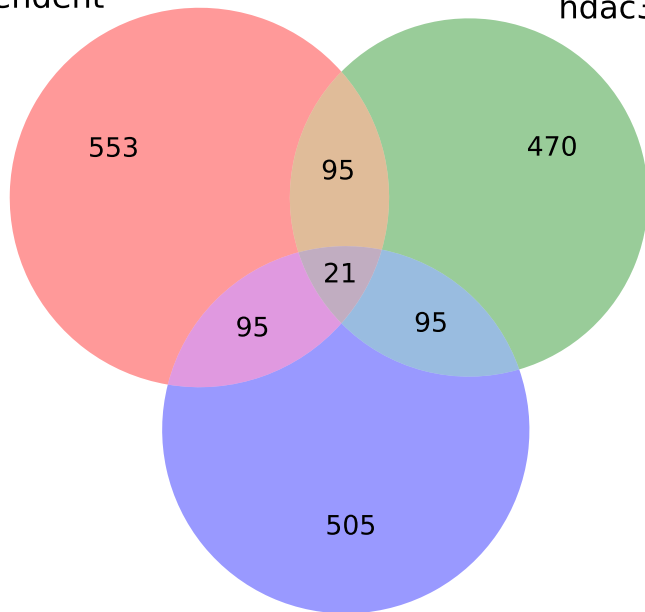


flip_2_0.2 dependent genes overlap

nr1d1 dependent

hdac3 dependent



nfil3 dependent