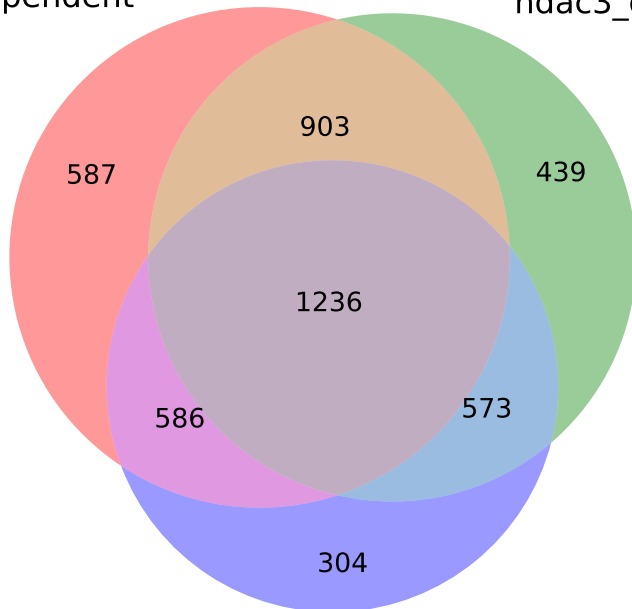


# 1.6\_5 dependent genes overlap

nr1d1\_dependent

hdac3\_dependent



nfil3\_dependent