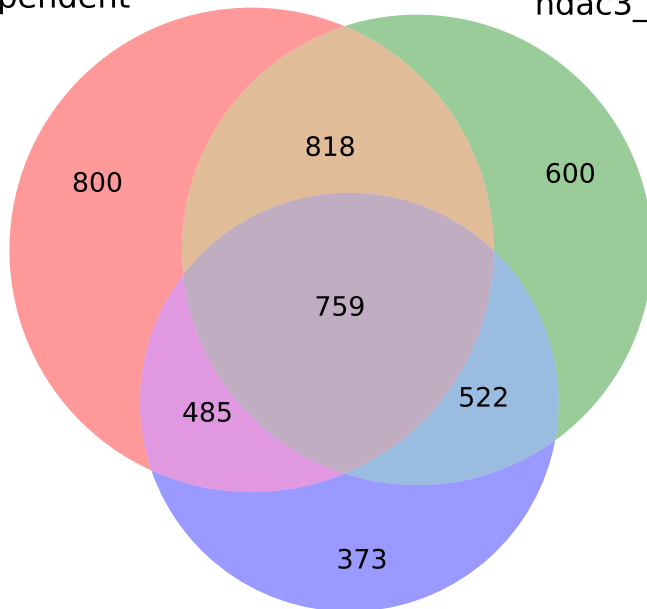


## 2\_6\_0.2B dependent genes overlap

nr1d1\_dependent

hdac3\_dependent



nfil3\_dependent