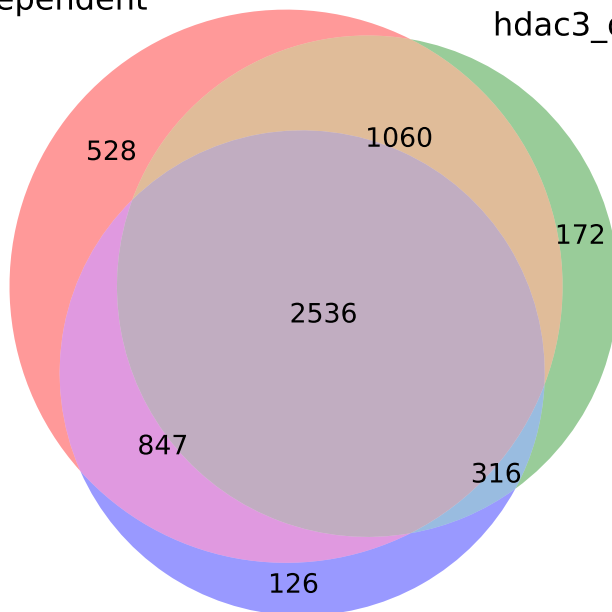


rAMP_fold_diff > 1.5 or phase_diff > 3

nr1d1_dependent

hdac3_dependent



nfil3_dependent