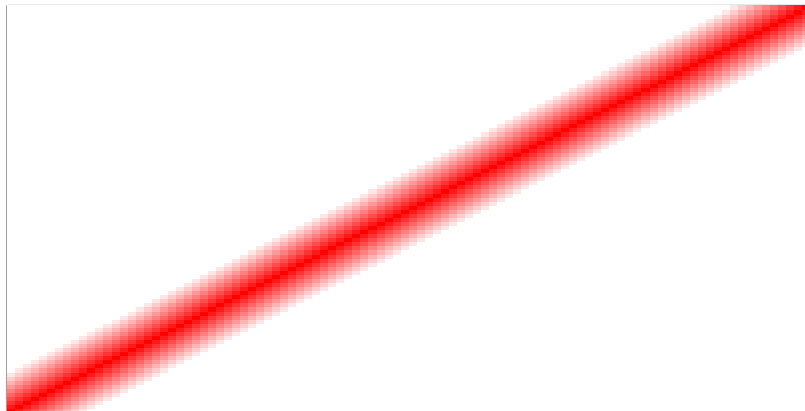
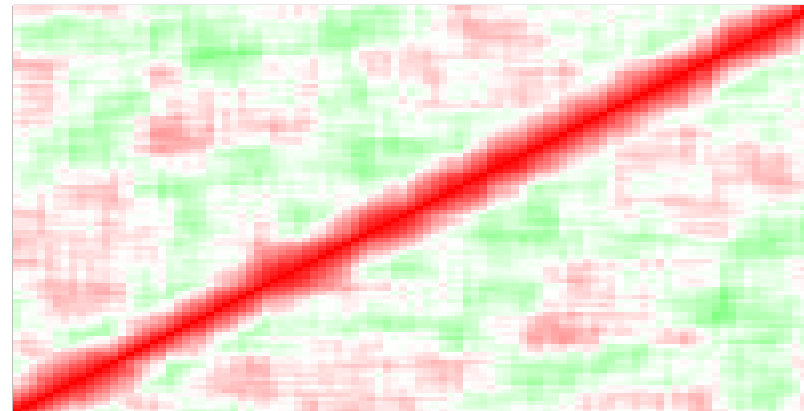


Toeplitz matrix (n = 50, p=100)

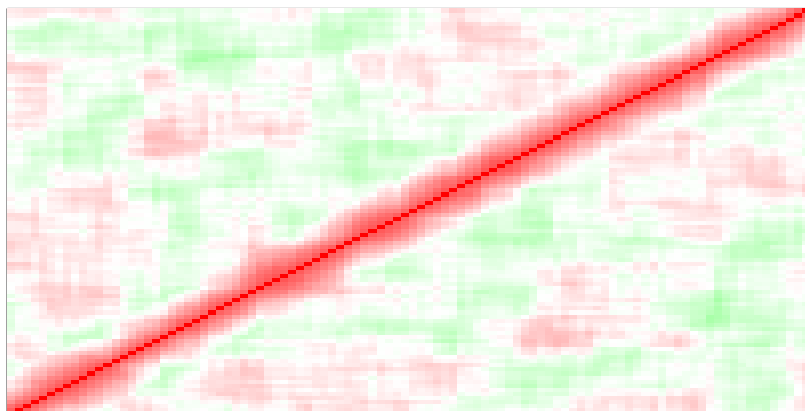
pop corr:



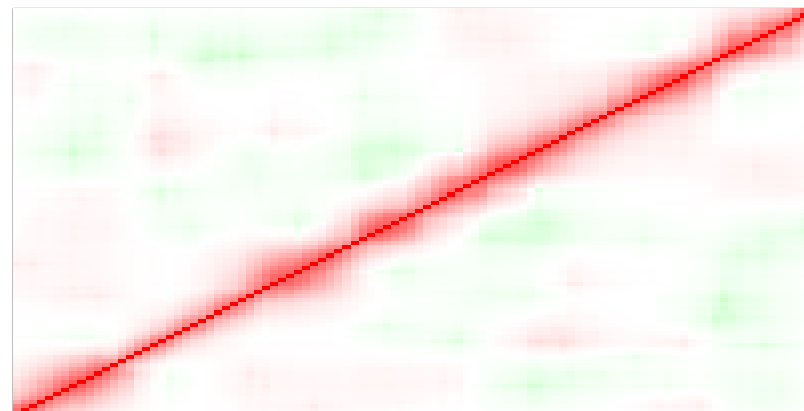
sample corr:



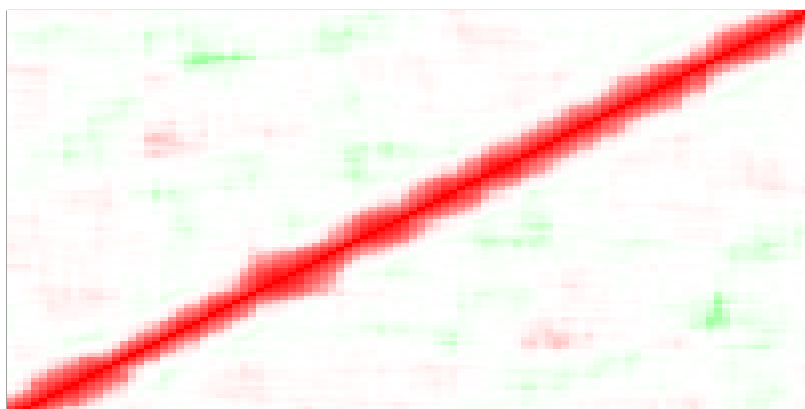
shafer strimmer:



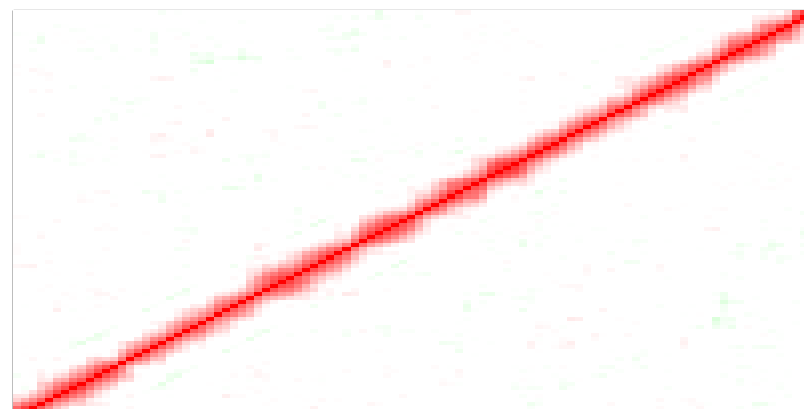
glasso :



corshrink:

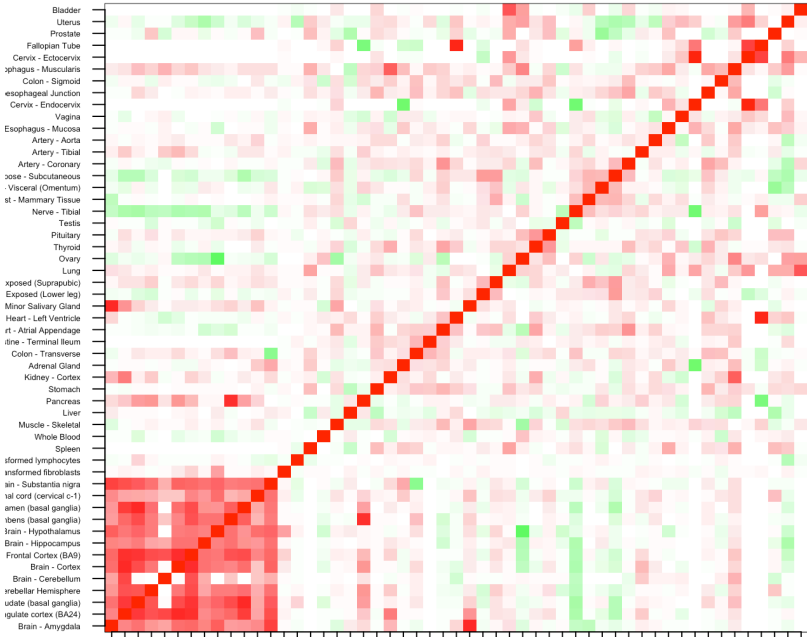


spcov:

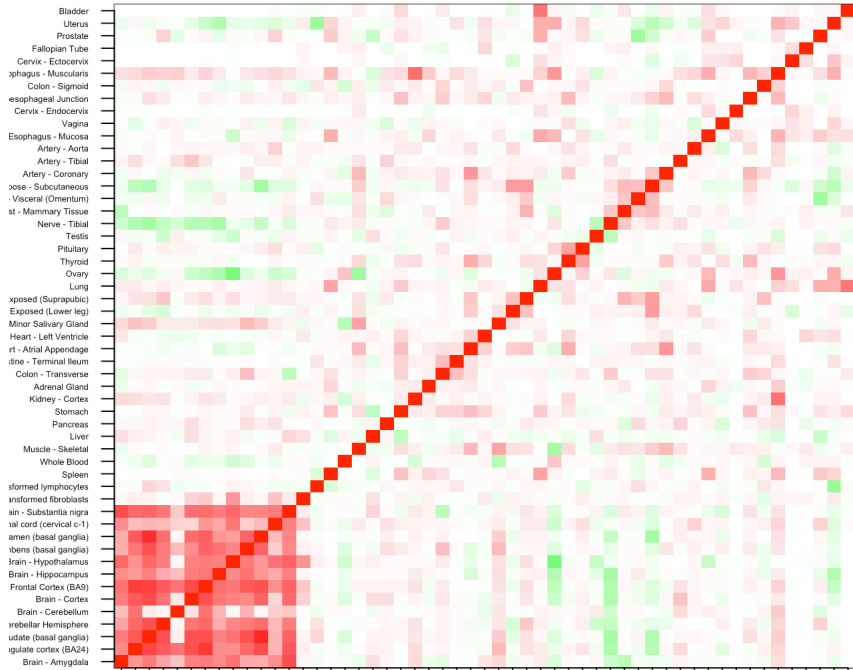


sample tissue-pair correlations

tissue-wide
CorShrink (mode = 0)



gene-wide
CorShrink (mode = 0)



tissue-wide
CorShrink (mode estimated)



gene-wide
CorShrink (mode estimated)

