positive control screen RNase MRP • 30 -5S rRNA Significant covariations **RNA** class 20 -U2 (S. pombe) H/ACA (29) other (7) snRNA (6) tRNA (39) U3 10 -SRP RNA U5L RNase P 5.8S rRNA U2 U6 0 -U1 Telomerase RNA

50

Percentage of base pairs expected to covary (%)

75

100

25