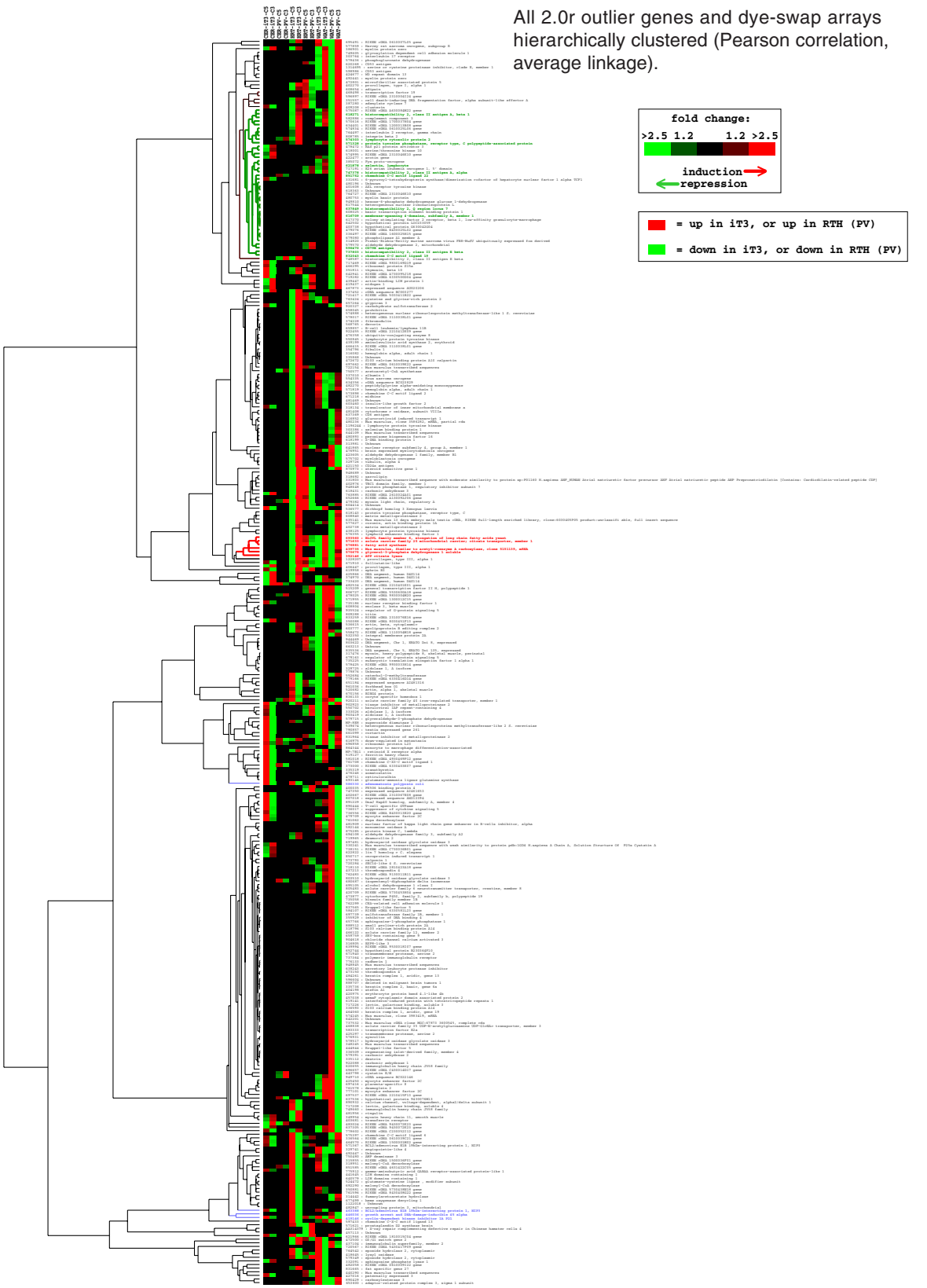

All 2.0r outlier genes and dye-swap arrays hierarchically clustered (Pearson correlation, average linkage).