Figure S12: **Statistics for the clustered PC for *S.cerevisiae*.** Top left: histogram with the number of genes forming the PC; top right: histogram with the percentage of the PC-genes that are represented in the 1301 genes passing the correlation threshold in at least a case; bottom left: histogram of the number of clusters intersecting each PC; bottom right: histogram of the fraction of genes belonging to the "main" expression cluster for each PC.