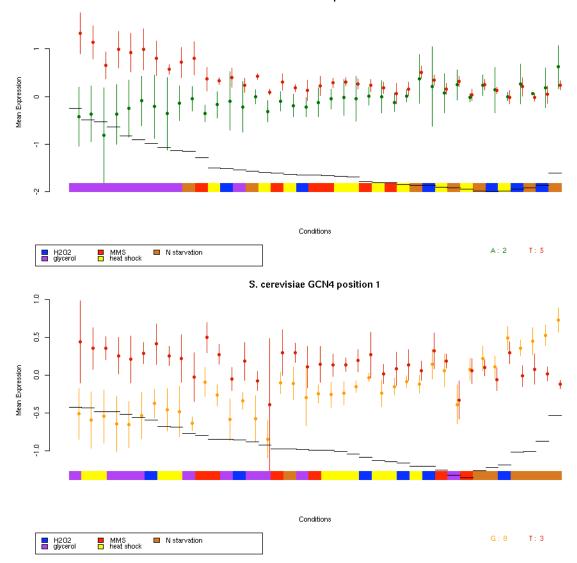
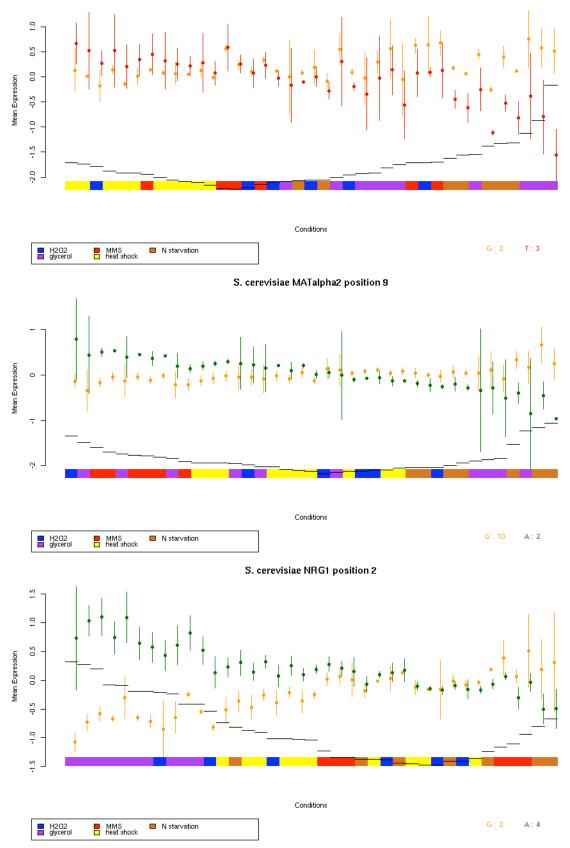
Figure S5. Complete set of figures showing comparison of average gene expression levels between genes with different functional variants of transcription factor binding site motifs in *S. cerevisiae, S. kudriavzevii, S. mikatae* and *S. paradoxus* based on expression data from Y6.4kv6 cDNA arrays. Mean expression levels for target genes of functional variants found at positions in TF binding sites using expression data from Y6.4kv6 cDNA arrays and stress conditions. Even if more than two variants exist at a position, only two are shown in each individual graph, and additional graphs show the pairwise comparison between each variant present at each position. The means are ordered across conditions according to the difference between mean expression of the two variants. Vertical lines extending from each point indicate the standard deviation of the mean. Horizontal black bars indicate the difference between the mean ranks. The significance of the functional heterogeneity was determined without reference to the segregation of experimental conditions, which are shown according to color along the x-axis. The number of targets for each variant graphed are shown at the bottom right hand of the graph.

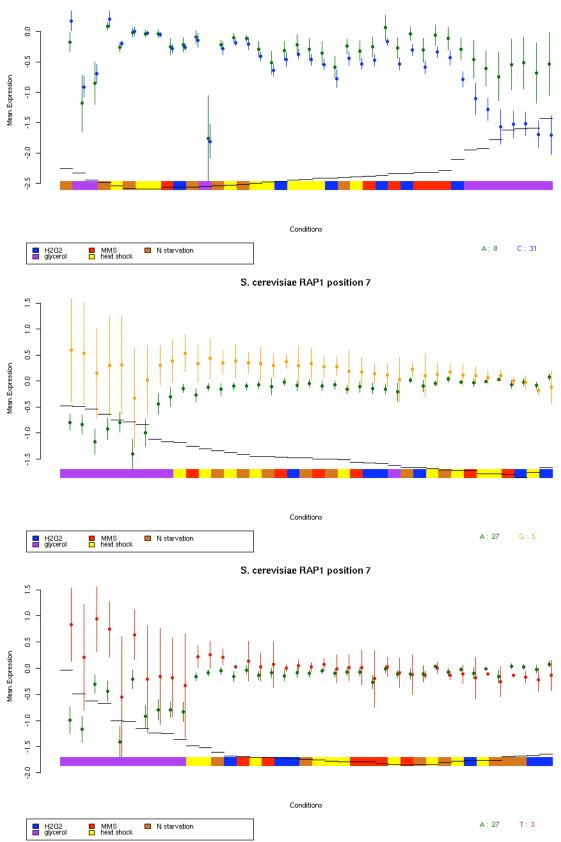


S. cerevisiae THI2 position 9

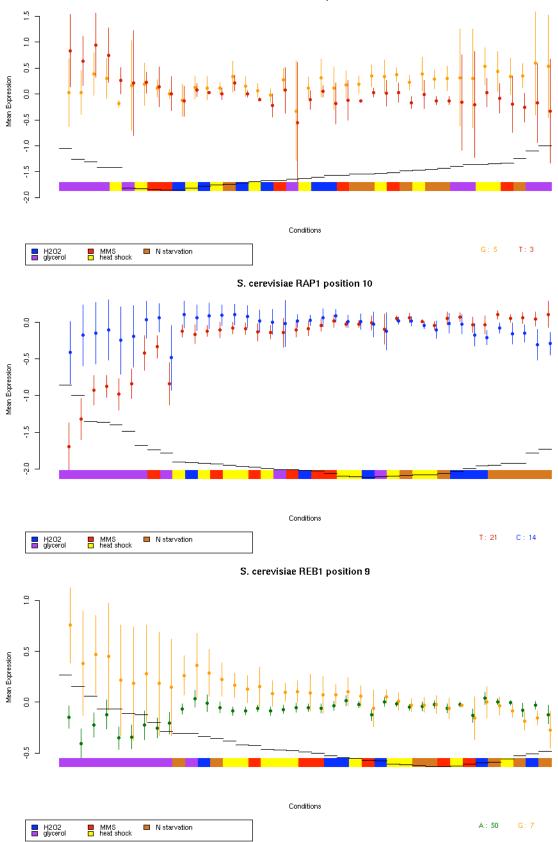
S. cerevisiae HSF1 position 7



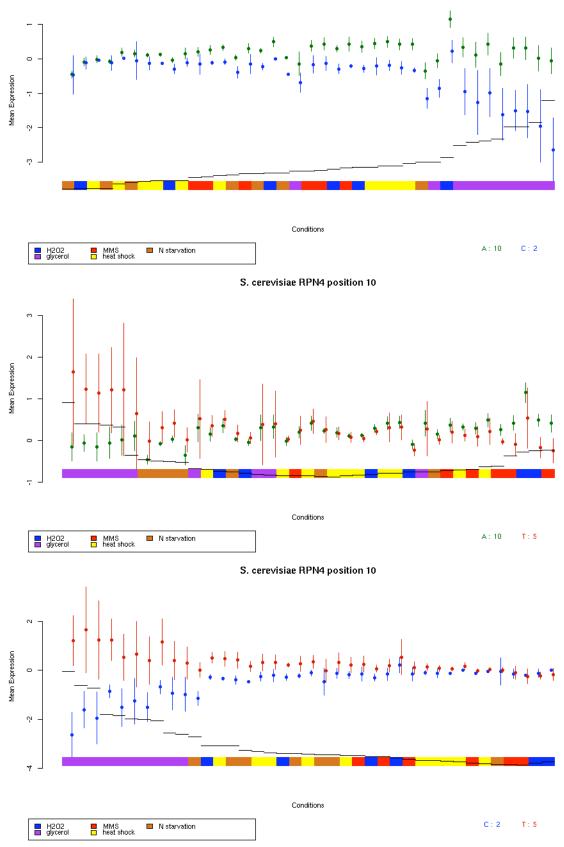




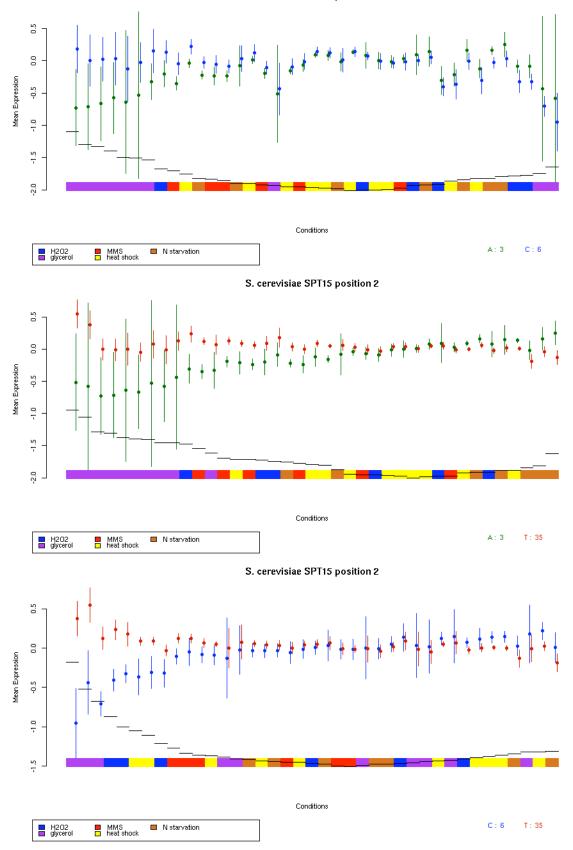
S. cerevisiae RAP1 position 7



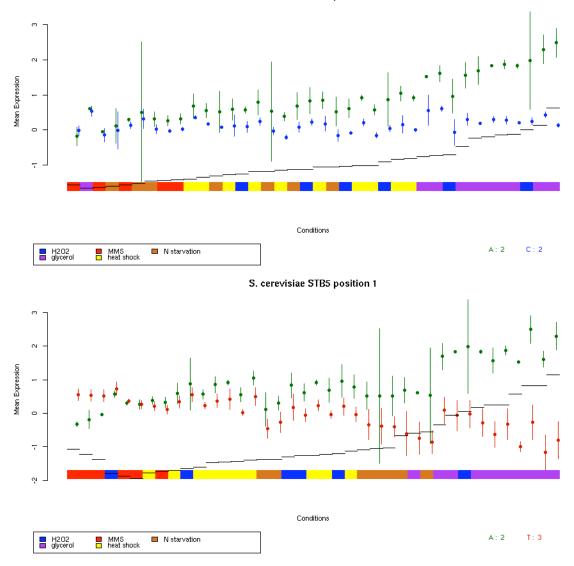
S. cerevisiae RPN4 position 10

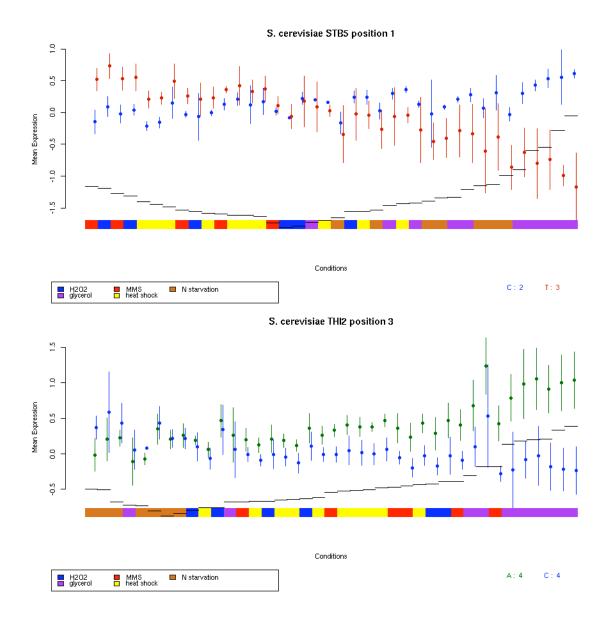


S. cerevisiae SPT15 position 2

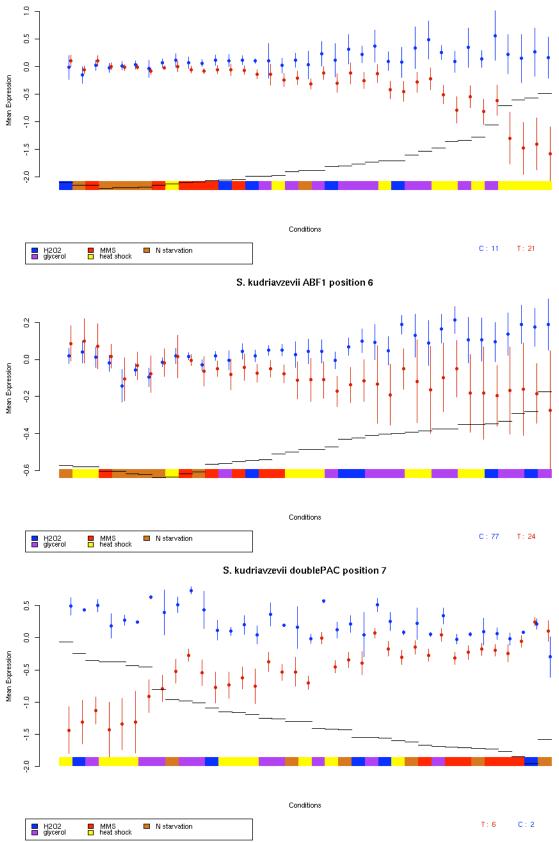


S. cerevisiae STB5 position 1

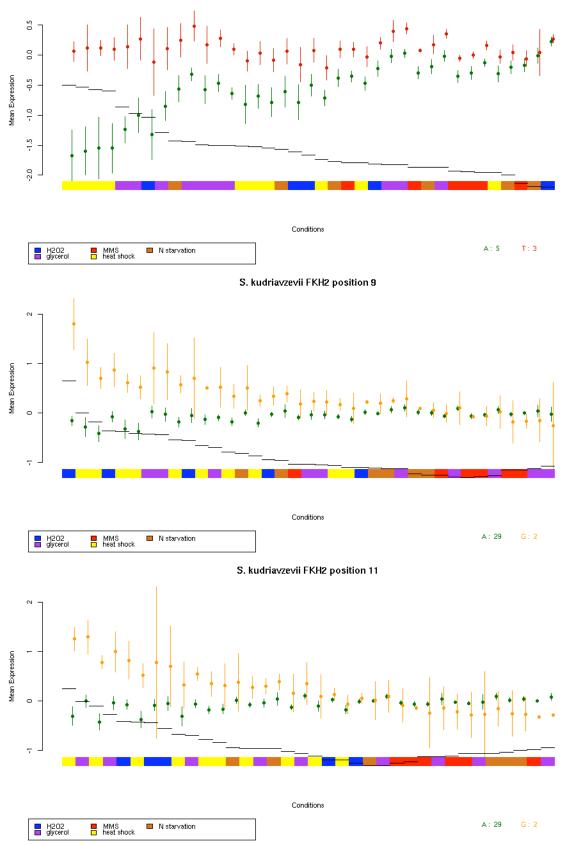


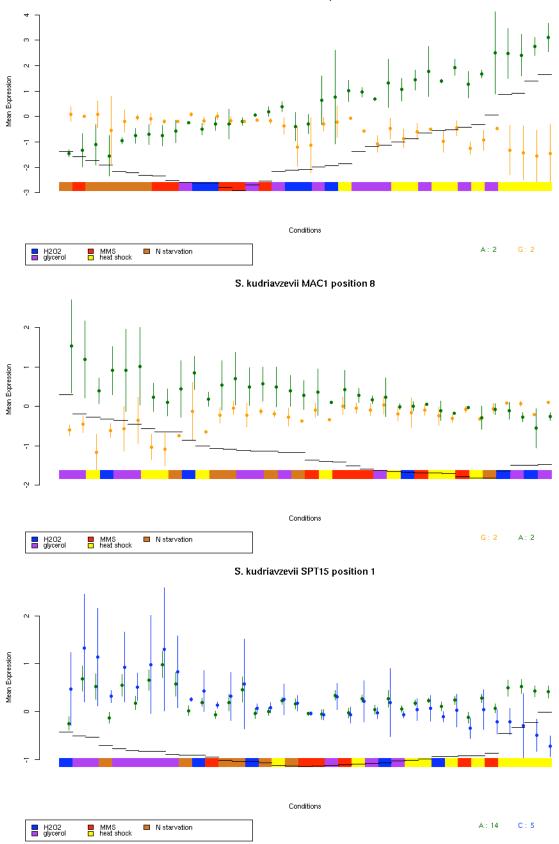


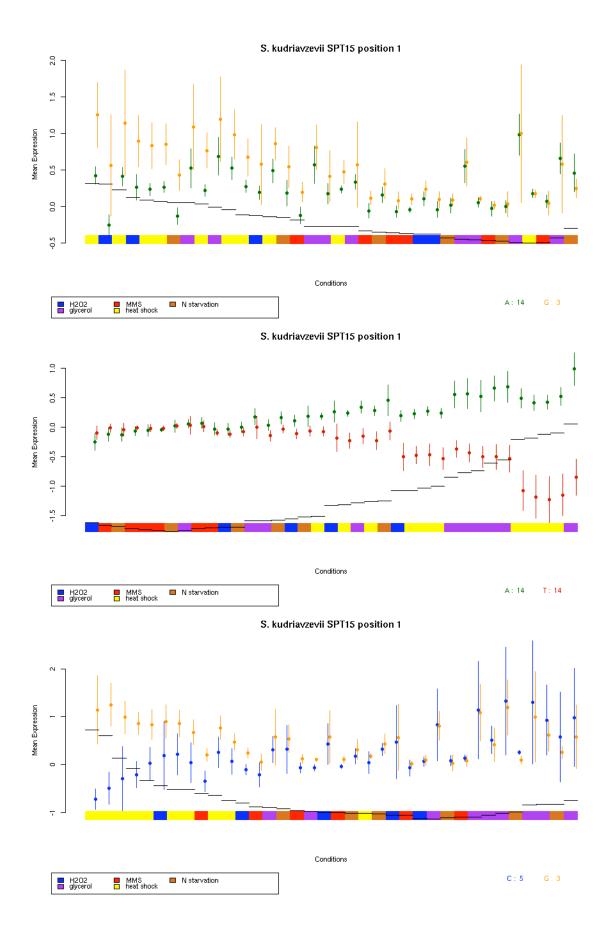


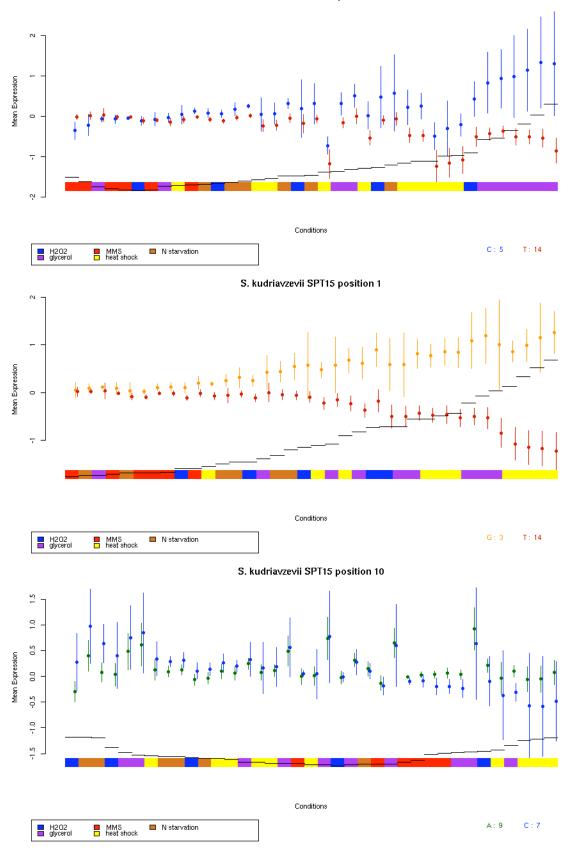


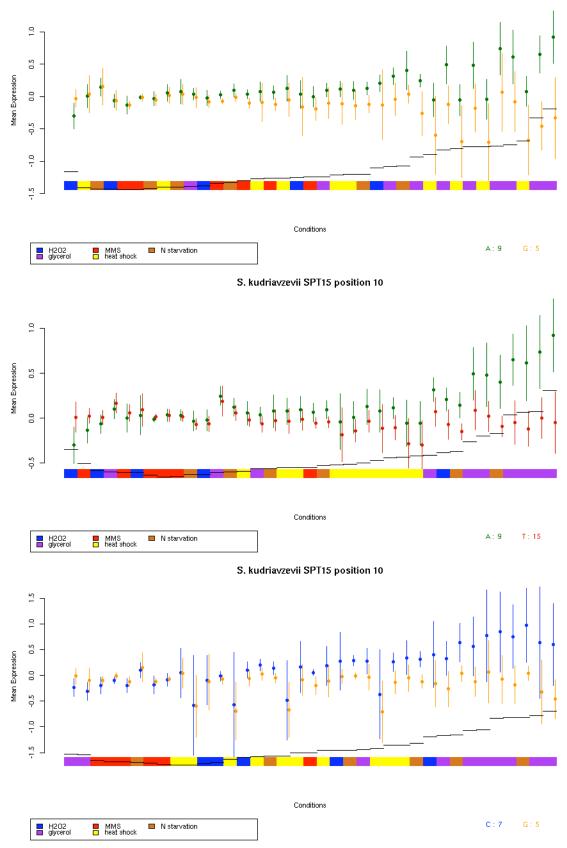


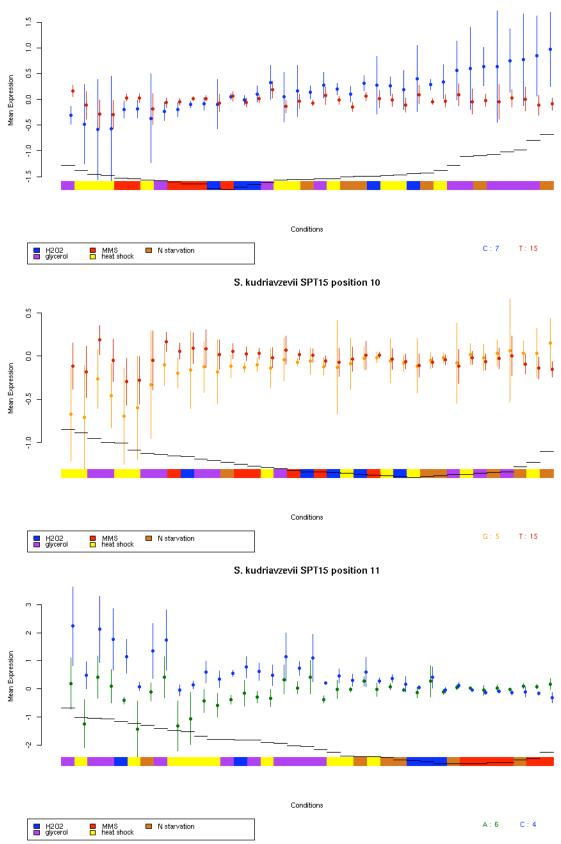


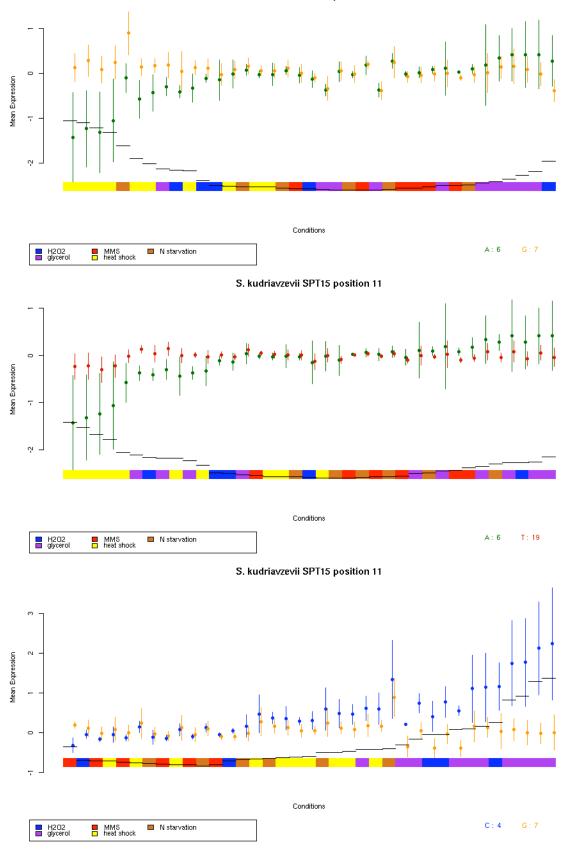


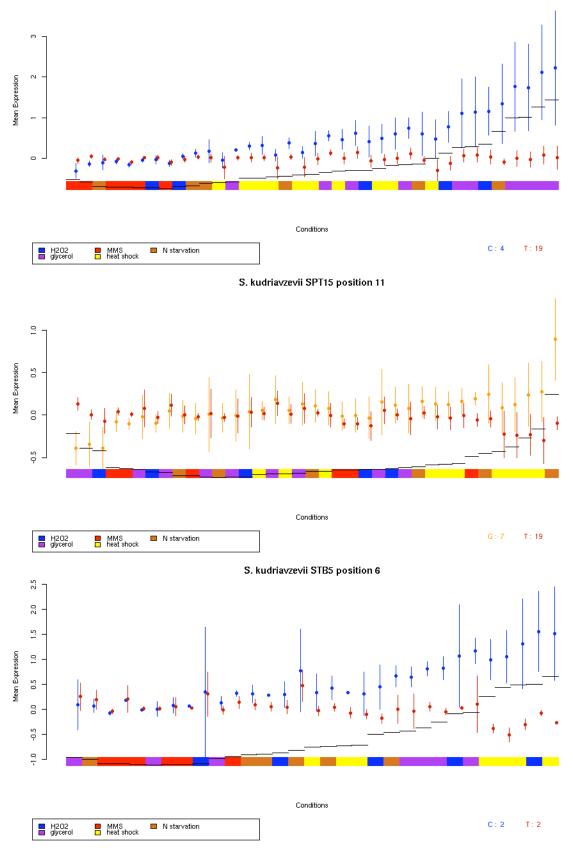




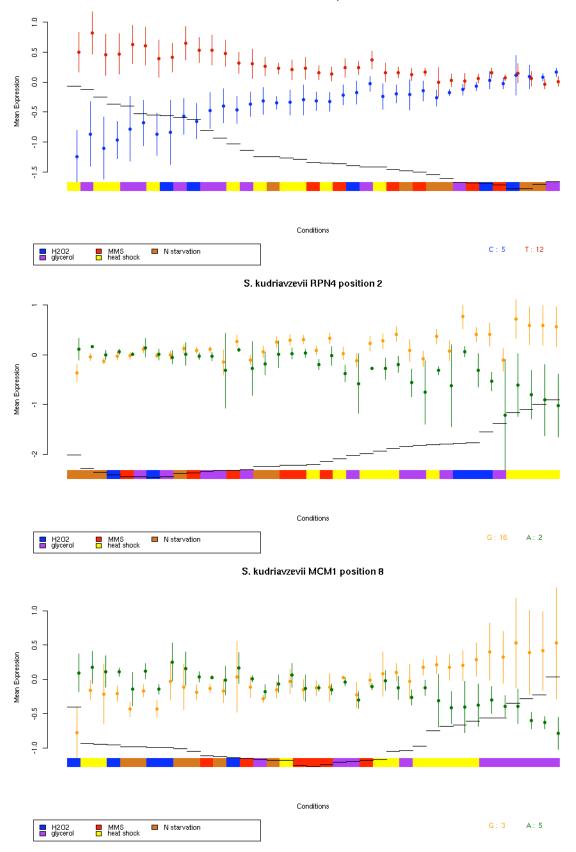




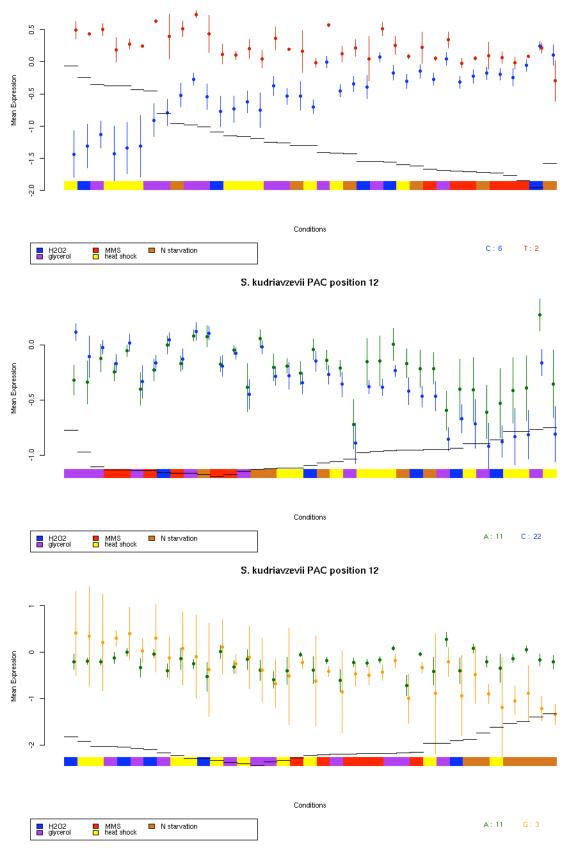




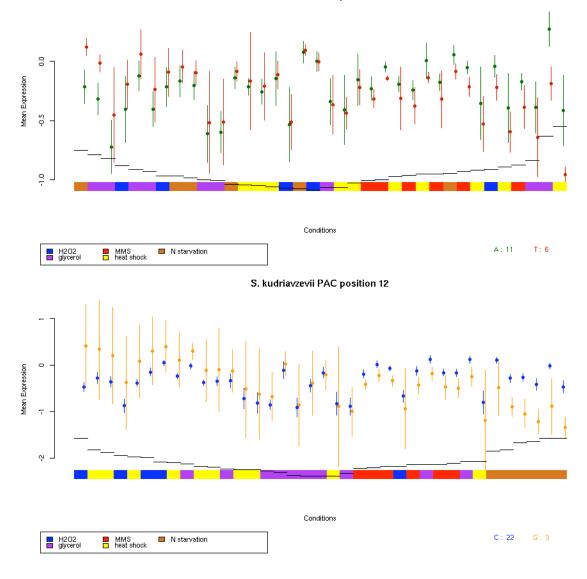
S. kudriavzevii CIN5 position 9



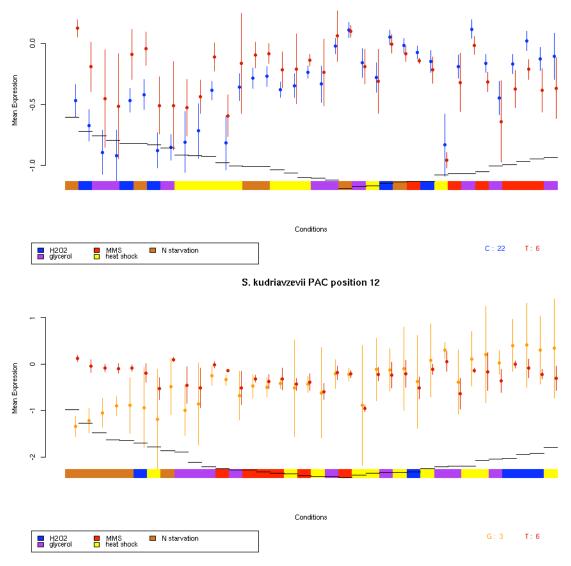
S. kudriavzevii doublePAC position 3

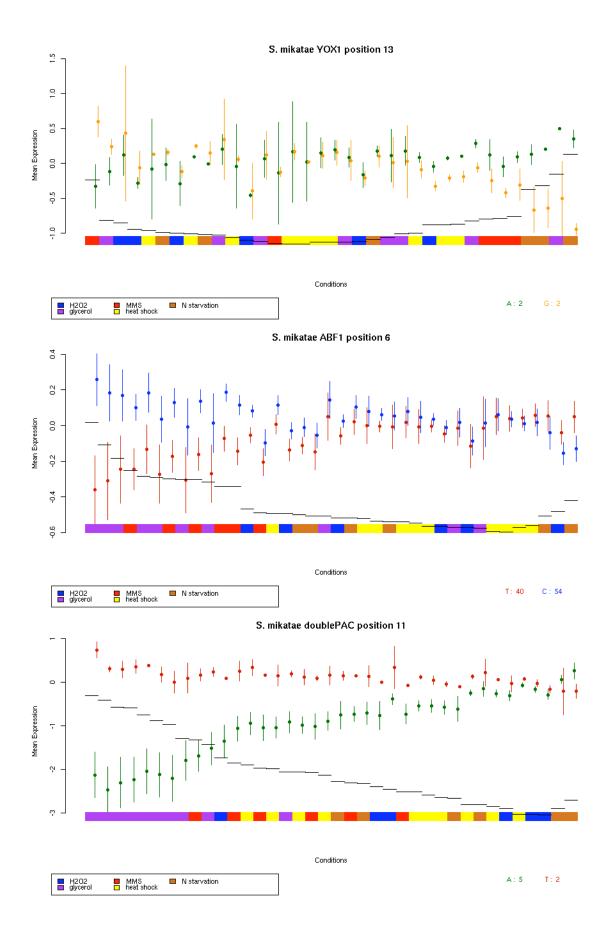


S. kudriavzevii PAC position 12

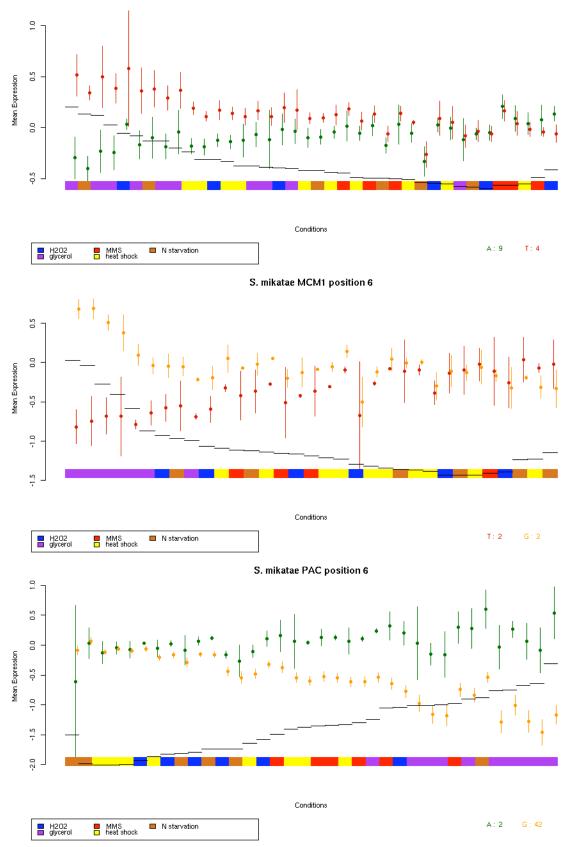


S. kudriavzevii PAC position 12

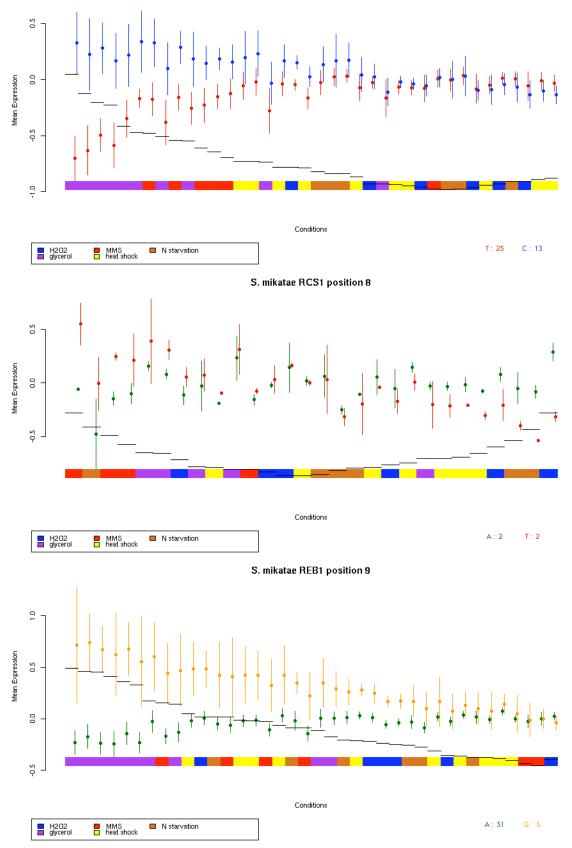


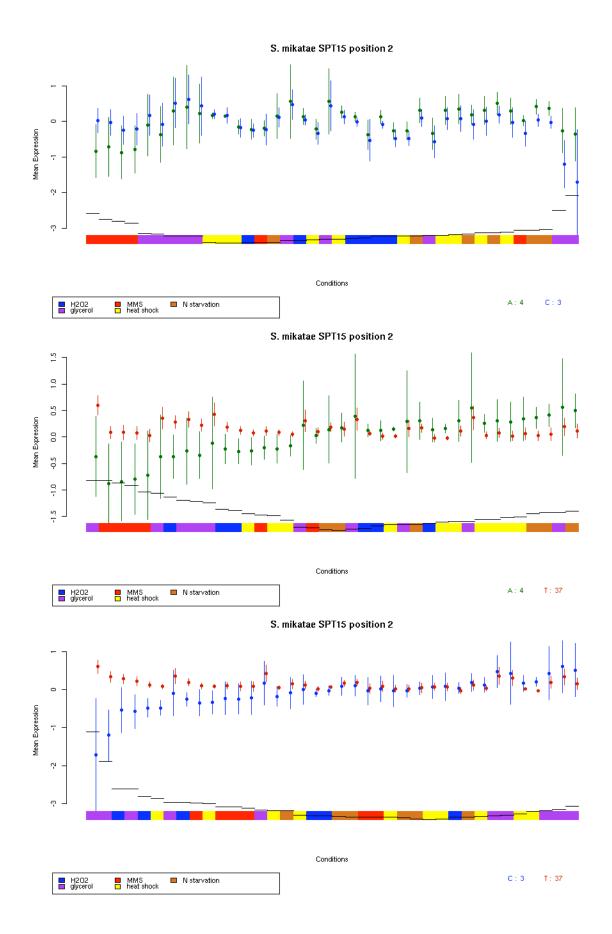




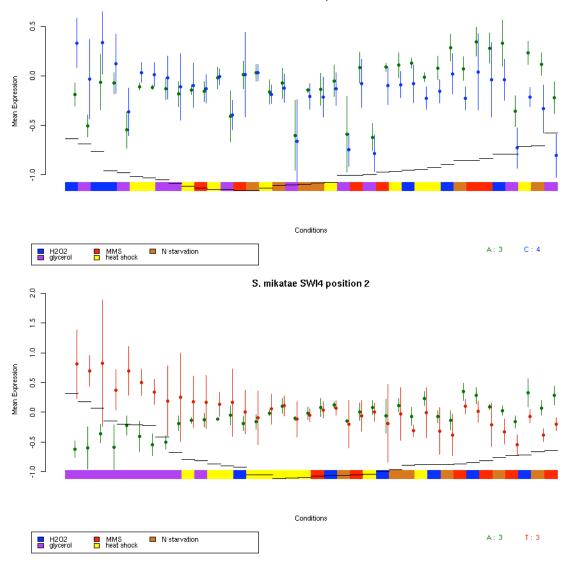


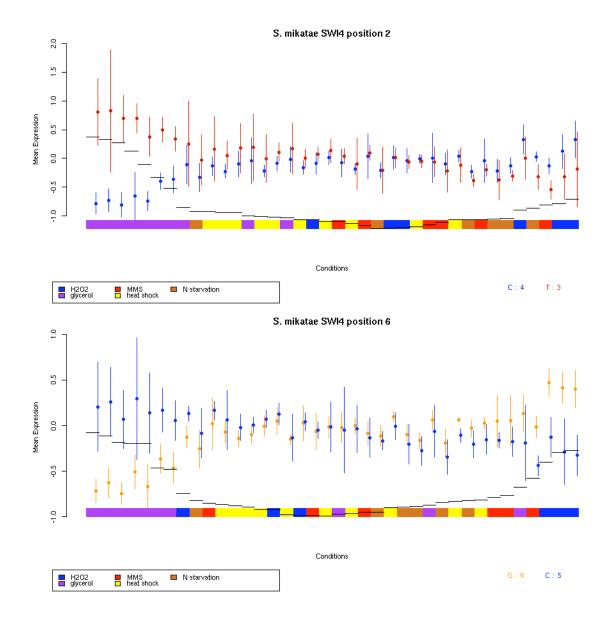


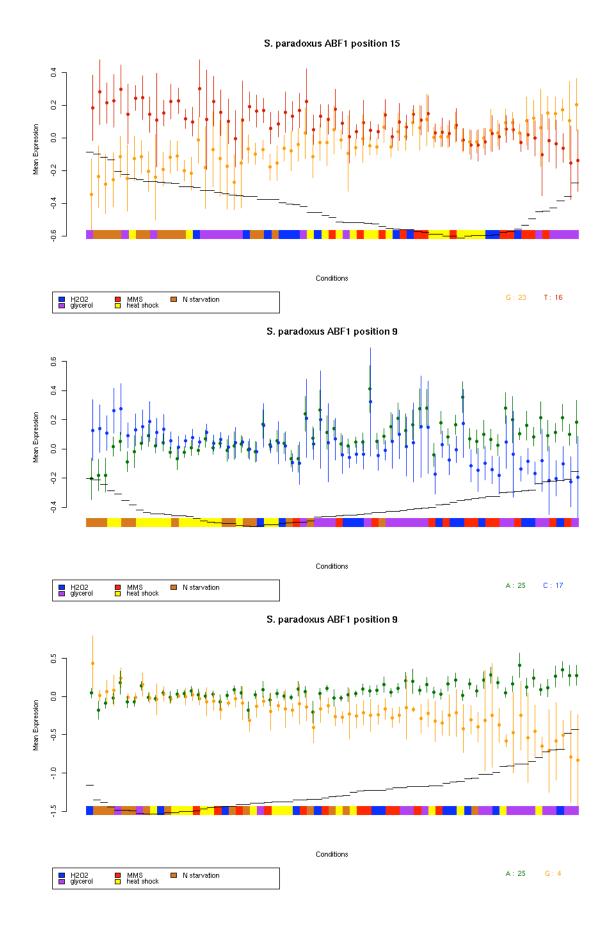


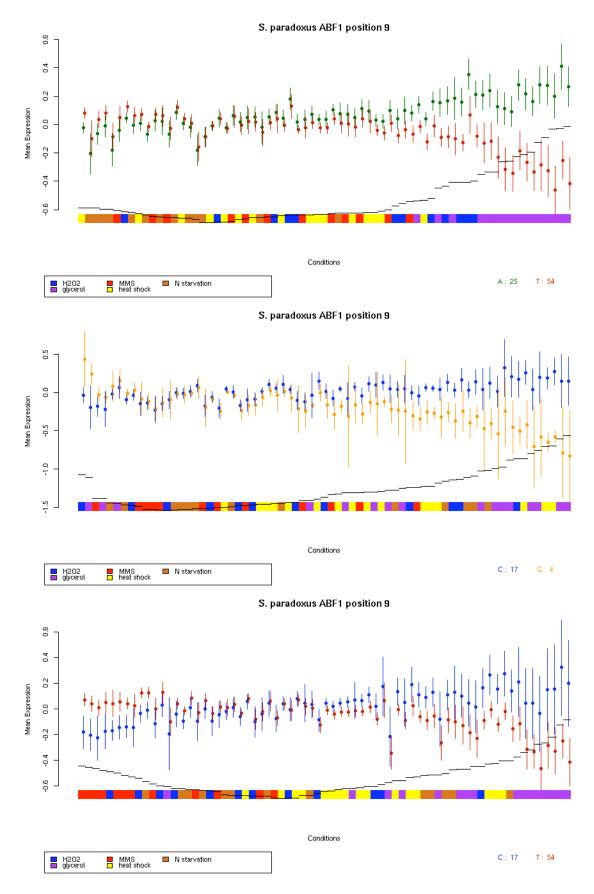


S. mikatae SWI4 position 2

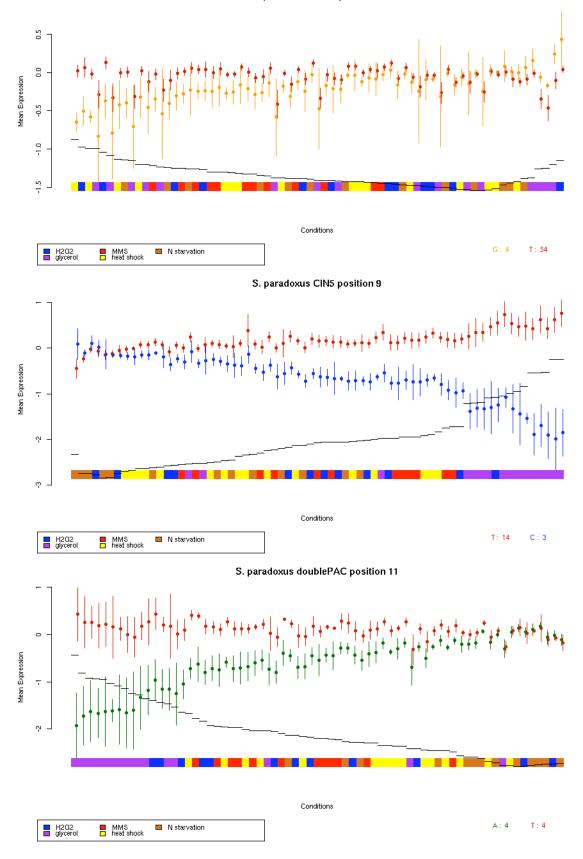




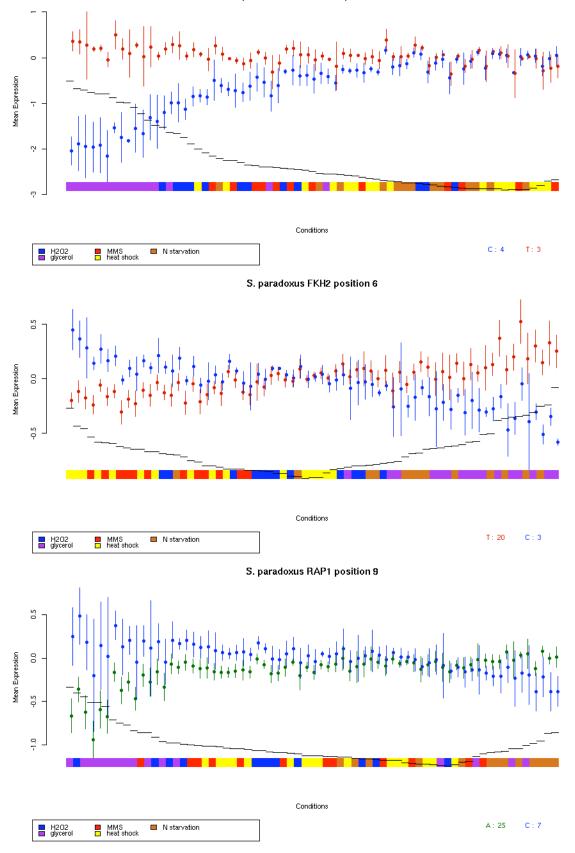


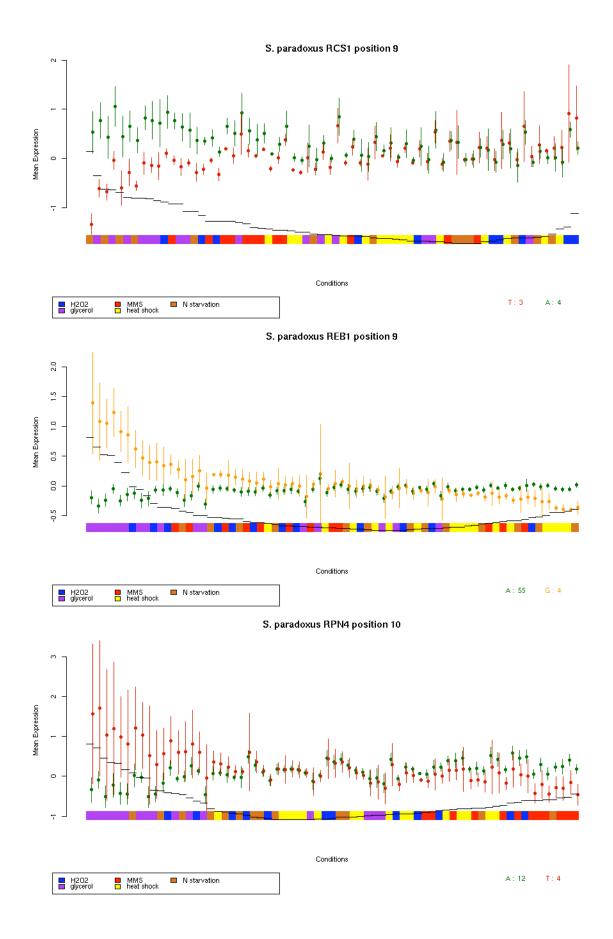


S. paradoxus ABF1 position 9



S. paradoxus doublePAC position 13





S. paradoxus STB5 position 1

