



Hierarchical clustering of 930 probe sets that differentially expressed by >2.0 in the cells at 39°C for 0, 6, 12 and 24 h. Clustering was performed by GeneSpring software. Each probe set is represented by a single row of colored bars. Intensity in the green and red color range indicates down-regulated and up-regulated probe sets, respectively. Intensity in the black color indicates no change (relative expression: 1). The number in parentheses indicates the number of probe sets.