Supplementary figure 6



Supplementary figure S6. Cold shifted cells display the opposite flow cytometry profile shift compared to heat shocked cells. (A) Heat map of mRNA expression changes during a heat shock and cold shock time series including all transcripts changing more than 1.7-fold and p<0.05 in any single time point compared to t=0. (B) Time series of flow cytometry profiles after heat shock. (C) Time series of flow cytometry profiles after cold shock. (D) Average magnitude of gene expression changes as shown in (A) above for genes with increased (yellow) or decreased (blue) expression during heat shock. (E) Average magnitude of gene expression changes during cold shock, as shown in (A) above, colours as in (D). (F) Fraction of 1N cells during heat shock. (G) Fraction of 1N cells during cold shock.