

Heatmap showing the correlation of transcription factors (rows) with cell cycle genes (columns). The color scale ranges from dark blue (low correlation) to dark red (high correlation). The transcription factors are CREB1, DDIT3, 62(mod), EP300, ZNF326, SNAIL, SPI1, and BHLHE40. The cell cycle genes are pDS269, pDS263, pBA581, pDS268, pDS262, pDS266, pDS255, and pDS258. The diagonal elements are all dark red, indicating high self-correlation. The off-diagonal elements show varying degrees of correlation, with some transcription factors showing high correlation with multiple cell cycle genes.

- 0.15

- 0.10

- 0.05

- 0.00