

Mild hypothermia provides Treg stability

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INVENTORY OF SUPPLEMENTARY DATASET

Figure S5

Figure S5. Top regulator effect network based on differentially expressed genes in Tregs expanded at 33°C. The network was generated in Ingenuity Pathway Analysis and yielded the highest consistency score of 51.5. The middle row consists of molecules that are connected with upstream regulators (transcription regulators: IRF8, KLF2, CDKN2A, GFI1, NFKB1, BCL6 and RELA, presented in the top row) and downstream diseases and functions (bottom row).

