

Supp Fig S7: Significance of similarity of CMINT modules inferred from 1 million 2000 bp genomic regions of the hematopoietic lineage. Shown are the significance of similarity of pairs of modules for pairs of cell types in the hematopoiesis lineage. Similar to **Supp Fig S6**, but using the negative log of p-value of the overlap significance (computed similarly to **Fig 4A**). Two different scales are used, Red: similarity for diagonal modules; Blue: similarity for off-diagonal modules. The more red or blue an entry, the more significant is the similarity of the corresponding modules.

Supp Fig S7

