



**Figure 5. Purging cell cycle genes from a signature does not rule out proliferation signals.** Distribution of the correlations with meta-PCNA of genes in the Embryonic Stem Cell Module (blue, ref. [15]), of the correlations of the same module with its cell cycle genes removed (red) and of all of the genes spotted on the microarray (black). doi:10.1371/journal.pcbi.1002240.g005