

Supplemental Figures

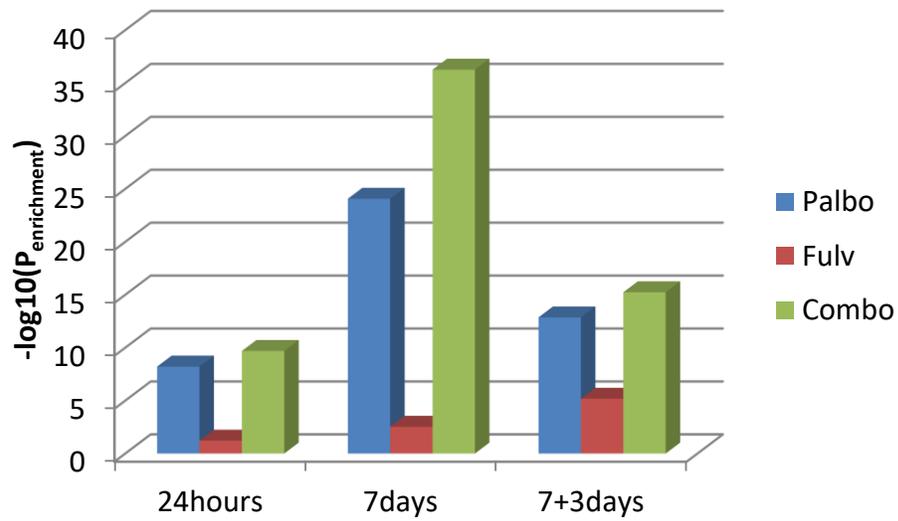


Figure S1. The enrichment of metabolic genes among those modulated by drug treatments. The statistical significance of over-representation was determined using cumulative hypergeometric distribution.

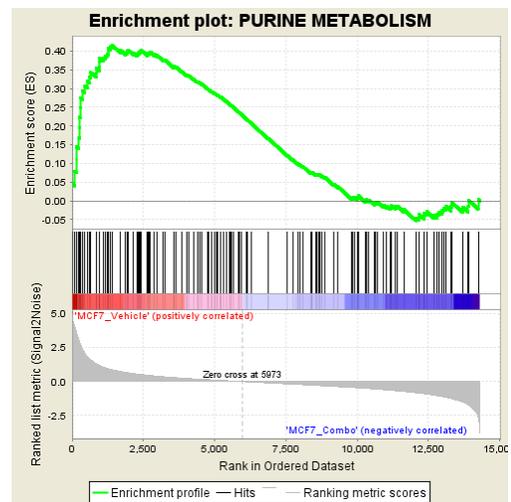
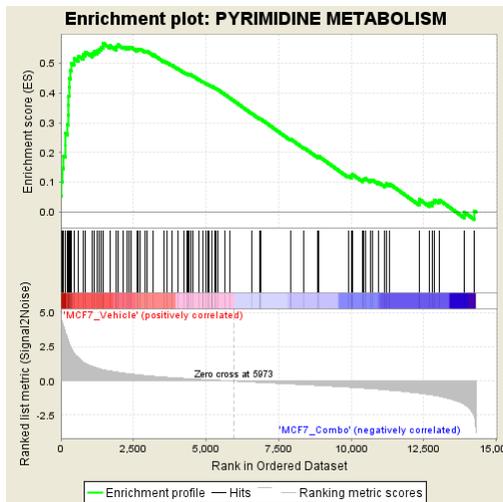


Figure S2. GSEA enrichment plot of pyrimidine metabolism and purine metabolism (FDR $q = 4.92E^{-5}$ and $3.83E^{-2}$), 24hr combo treatment.