



Supplementary Figure 1. Reproducibility of Ad5 induced cellular gene expression changes. Significance analysis of microarrays (SAM) was used to identify 505 down-regulated (A) and 1113 significantly up-regulated (B) genes at 30 hours after infection (hpi) with Ad5 (false discovery rate = 0.9%). Expression responses for these probes, shown in blue (down) and yellow (up), were isolated from a joint dataset containing the response profiles from the previous time-course experiment labeled in red and the current experiment (black). In the former, the black triangles from 18 to 56 indicate increasing periods of infection.