



Figure 3 - Supplemental Figure 10: Chromatin remodeling not a mechanism driving novel Aim2 isoform. (A) mm9 genome browser shot between chr1:175,348,283-175,351,422 indicating ATAC-seq of cells from hematopoietic tree. (B) mm9 genome browser shot between chr1:175,348,283-175,351,422 indicating ATAC-seq of BMDMs +/- Lipid A for 2 hrs. (C) Schematic summarizing promoter accessibility of novel Aim2 isoform, red=accessible and black=not accessible.