Figure S11 - Transcriptional comparison of skin vs vaginal DC subsets. a. 1,007 transcripts differentially expressed between skin and vaginal APC. HLA-DR- vaginal cells are also displayed. The data are normalized to the median of all APC (HLA-DR+) cells. b. Bar chart representing GO biological process enrichment in skin and vaginal APC. c. IPA network analysis of the 156 transcripts over-expressed in vaginal HLA-DR- cells as compared to HLA-DR+ cells. d. IPA network analysis of the 123 transcripts over-expressed in vaginal HLA-DR+ cells as compared to HLA-DR- cells.