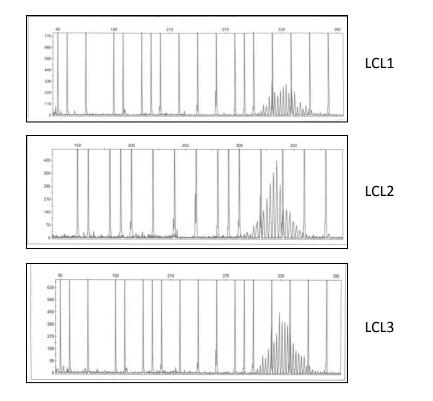


Supplementary figure 1 : Analysis of EBV gene expression following infection of GC B cells with EBV. Q-RT PCR showing Wp, Cp, Y3-U-K-spliced EBNA1, EBNA2, LMP1, LMP2A and BZLF1 expression in three GC B cell derived LCLs at different time-points post infection. Assays were performed in triplicate and results are presented as raw Ct values.



Supplementary figure 2: EBV infected GC B cells are polyclonal six weeks post infection. PCR was used for the detection of immunoglobulin heavy (IGH) chain gene rearrangements. Results are negative for clonal IGH chain gene rearrangement consistent with the presence of a polyclonal B-cell population. The method used is described in Linke *et al.,* 1997.