



Additional file 3, Supplementary Figure 1. RNAs from the 13 indicated tumors were assayed for relative mRNA content for 9 key genes: *c-Myc*, *Socs1*, *Socs2*, *Abl1*, *Jak1*, *Xbp1*, *Sdc1*, *Irf4*, and *Pax5* (*BSAP*) as well as for the “housekeeping gene,” GAPDH, using primers designed in consultation with the manufacturer (Applied Biosystems, Foster City, CA). All data were normalized to GAPDH, and these values were then calibrated against the sample with the lowest expression and represented as $-\Delta\Delta Ct$ values with standard errors.