S4 Fig. Association between IgVH mutation status and BAGS subtype. For each subtype the proportion of samples with mutated and unmutated IgVH are shown. (a) All subtypes, (b) All subtypes divided into early, naïve and late subtypes. The Munich, Duke, Padova, and IDFCI cohorts were used (n = 337). Unclassified samples were not included. Tests for significantly different distributions between groups were calculated using Fisher’s exact test ((a) P = < .001 and (b): P = < .001, early vs. late; P = < .001, naïve vs. late; P = .01).