**Figure S2** – Global density distribution of single-site CpG methylation levels in splenic naïve B cells with and without *Dnmt3a*. Density distribution (bin size, 0.05) of single-site mCG/CG levels for all validated CpG sites with read depth of at least 10 for (A) *Dnmt3a*+/+ (N = 17,295,282) and (B) *Dnmt3a*−/− (N = 16,696,664) samples.