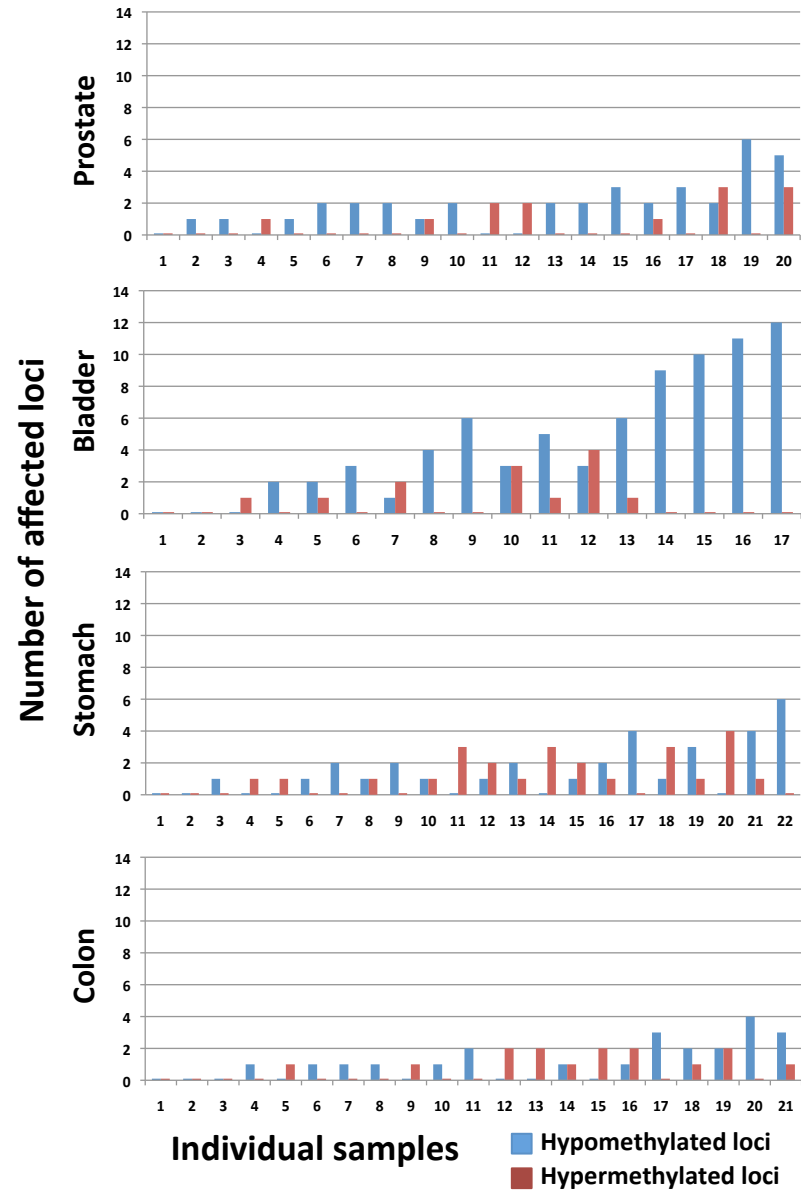


A) Number of hyper- (>10%) and hypomethylated loci (<-10%) in each tumor samples



B) Number of patients vs. number of affected loci

