Figure S3. Correlation of number of potential binding neoepitopes with number of nonsynonymous mutations. The number of potential binding neoepitopes (IEDB score $\leq 500$) are highly correlated with the number of nonsynonymous mutations for all three subtypes of breast cancer. In all the plots, a linear regression model is used to fit the data; the fitted line is shown in red and 95% CIs are shown as grey shaded area around the line.