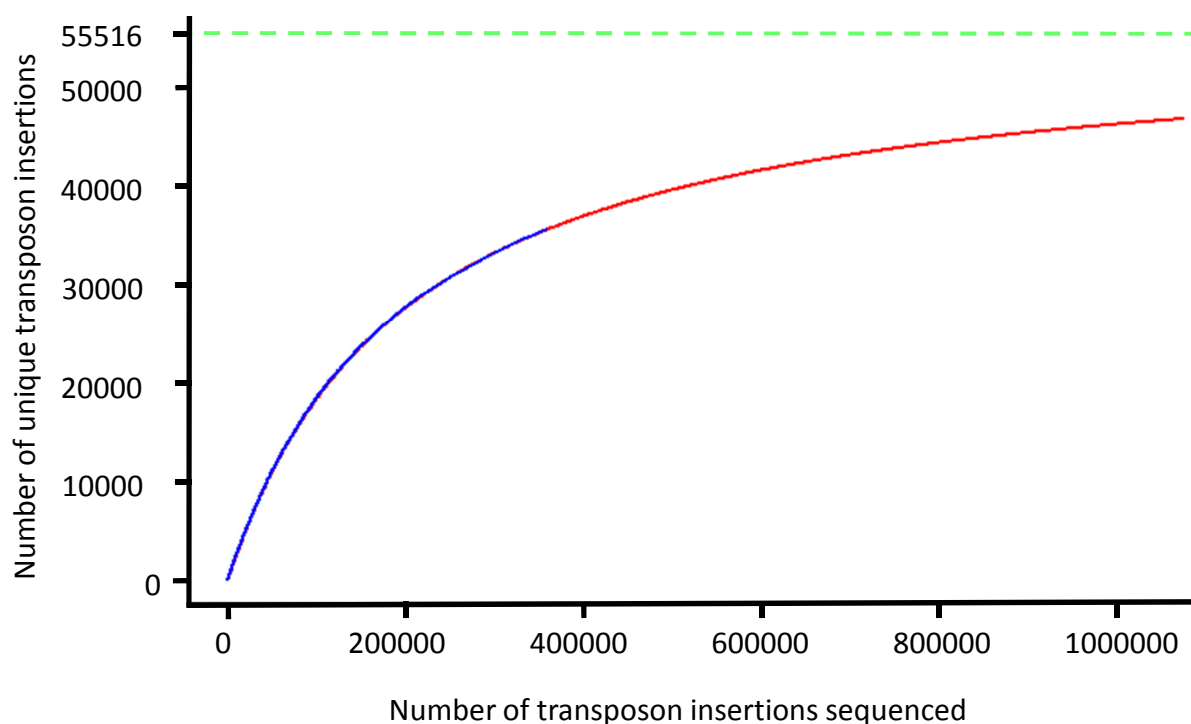


Additional Figure 1



Additional Fig. 1. An accumulation curve of the transposon insertion sites in the *M. tuberculosis* library. To determine the library coverage the number of genes or TA sites mutated was plotted against the number of insertions sampled. A Michaelis-Menten curve was fitted to the data (blue line) and extrapolating (red line) to give an asymptotes (green dashed line) that predicts that the library contains 55,176 unique transposons mutants (out of a possible 62,418 predicted TAs in the GC1237 genome).