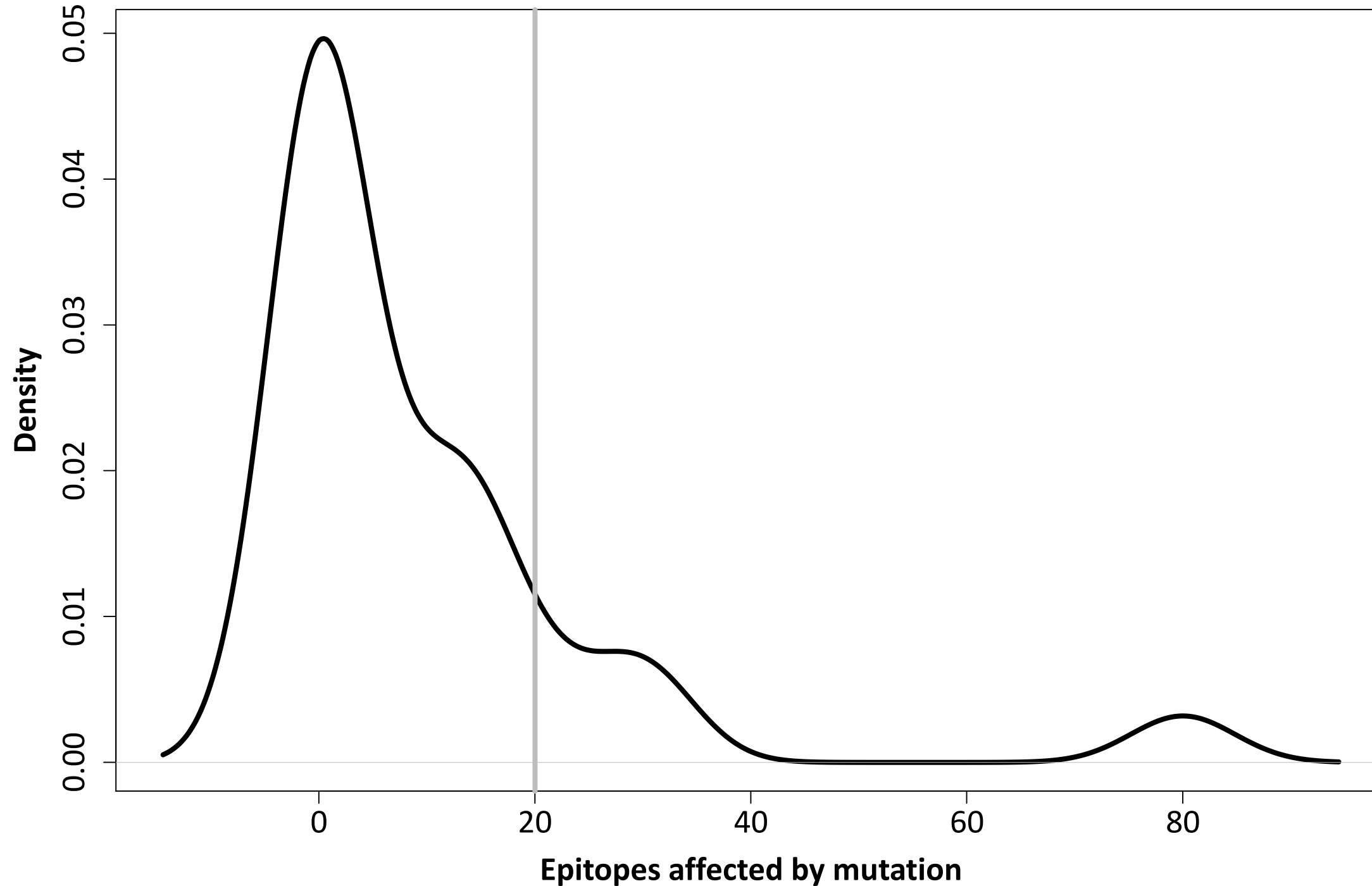**Figure S1: Density distribution of the affected epitopes.**
This plot visualizes the density or amount of data points (y-axis) around discrete epitope frequencies (x-axis).
The shape of the density curve highlights a three distinct clusters (Around 0, 20 and 80 epitopes). The second cluster, that is distinct from the 0 forms around an epitope frequency of 20. Therefore, we chose 20 as our threshold to distinguish low (n = 22) and high amount of epitopes (n=4).