In the figure, we observe the normalized response (Inorm) of different glycine concentrations for various protein variants. The x-axis represents the glycine concentration in mM, ranging from 1E-3 to 1000. The y-axis shows the normalized response (Inorm) ranging from 0.0 to 1.0. The graph includes data points and error bars for four protein variants: WT, S267I, R271K, and R271Q. Each variant is represented by a different symbol, and the data is fitted using the Hill-Fit model. The Hill-Fit model curve is shown as a black line.