
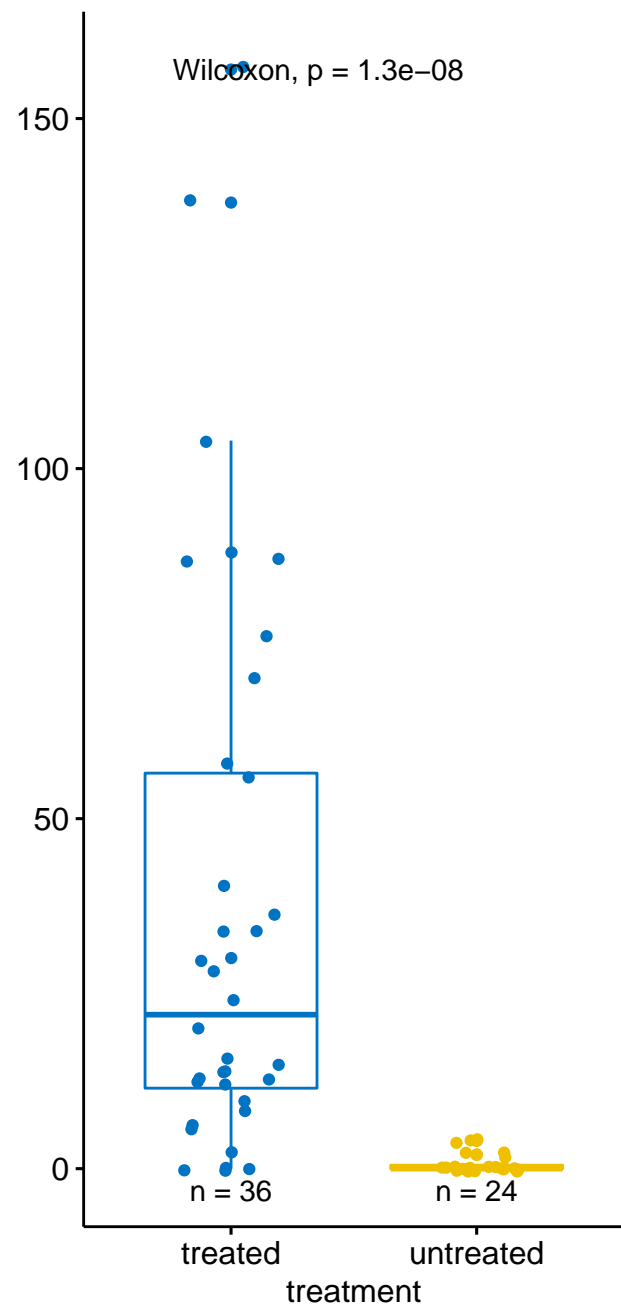



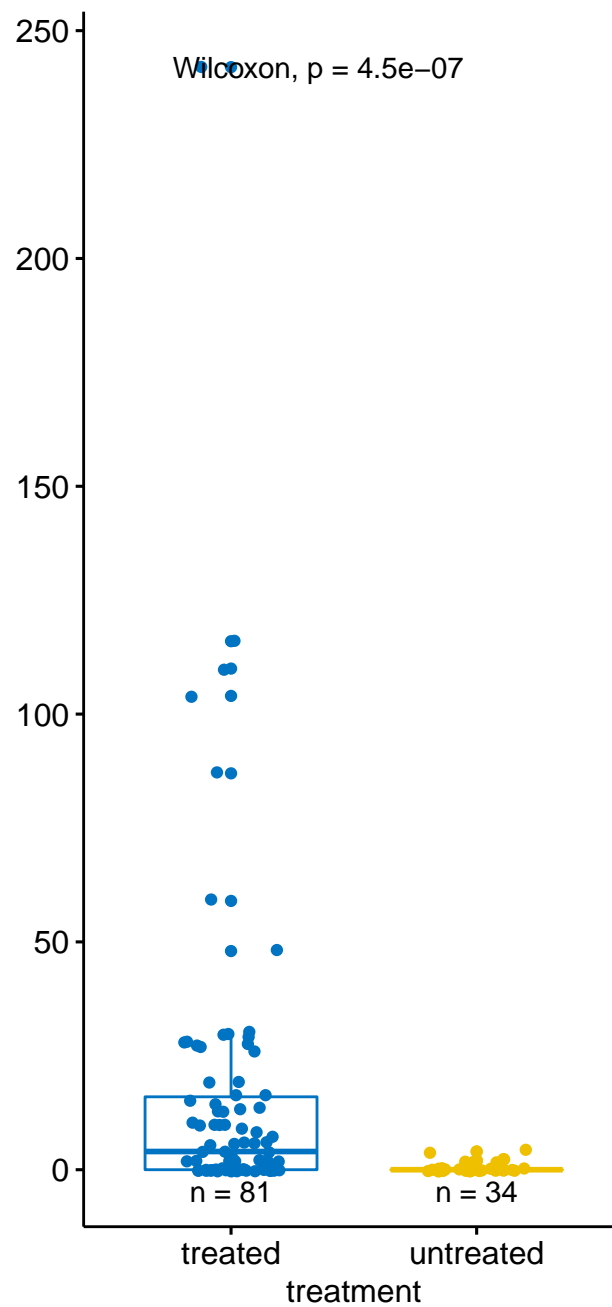
A # unique insertions per sample
Min Freq = 0.01 %

treatment  treated  untreated





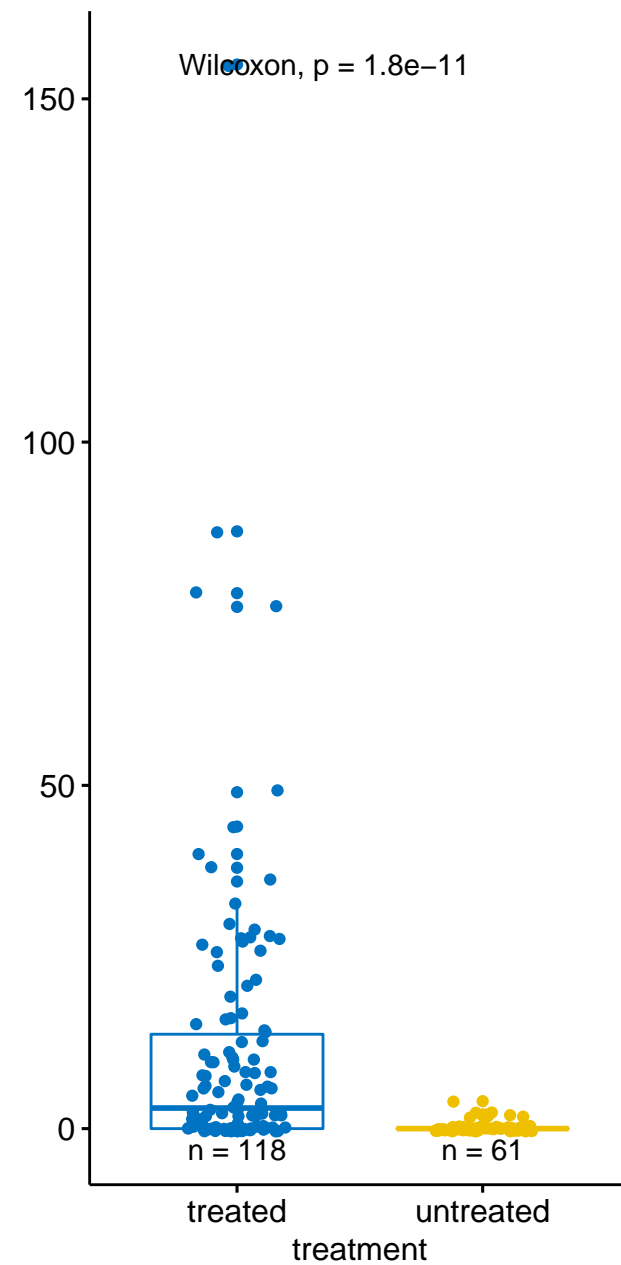
B # unique insertions per sgRNA
Min Freq = 0.01 %

treatment  treated  untreated





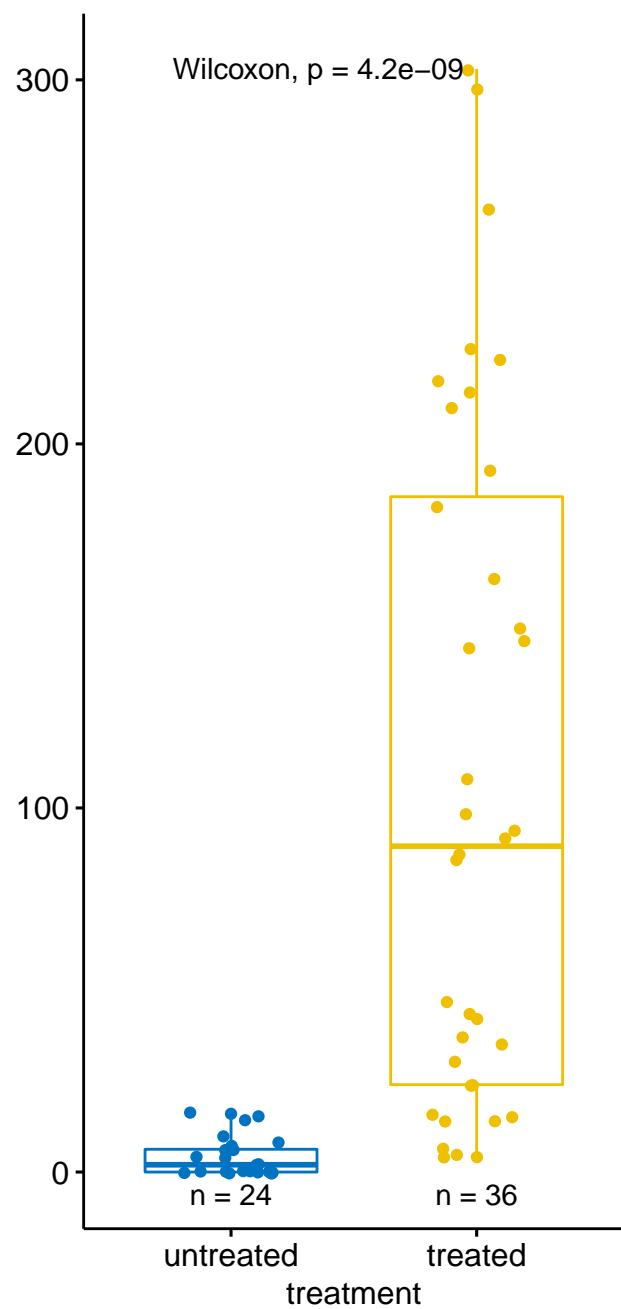
C # deletions by sgRNA & sample
Min Freq = 0.01 %

treatment  treated  untreated





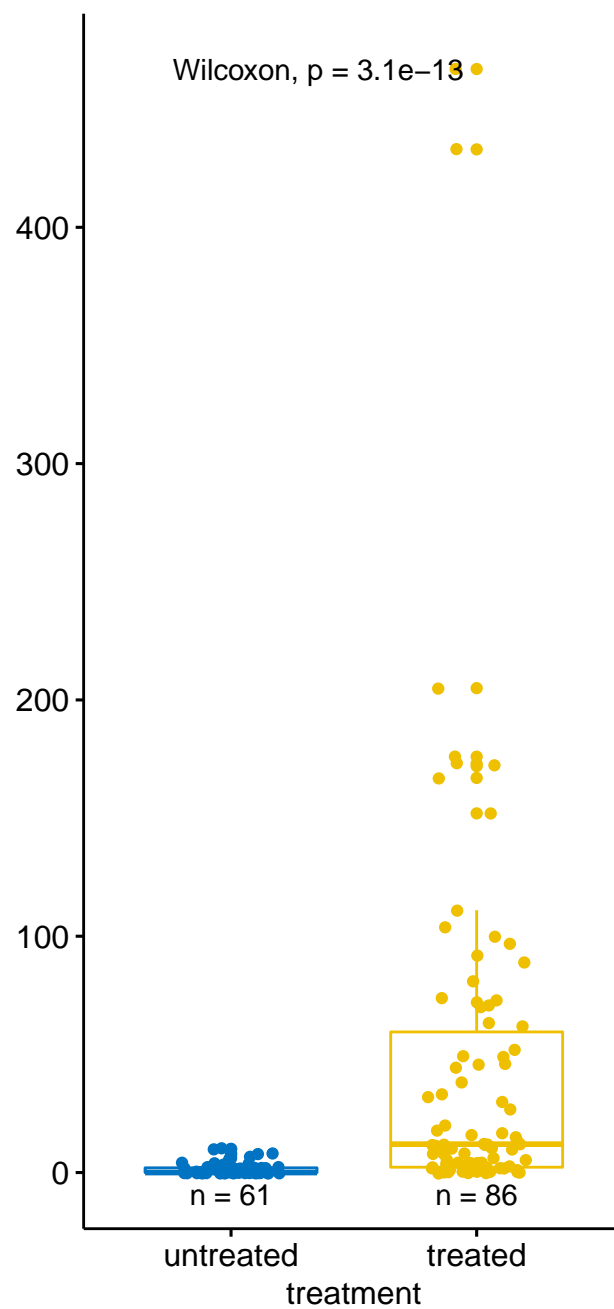
A # unique insertions per sample
Min Freq = 0.003 %

treatment  untreated  treated





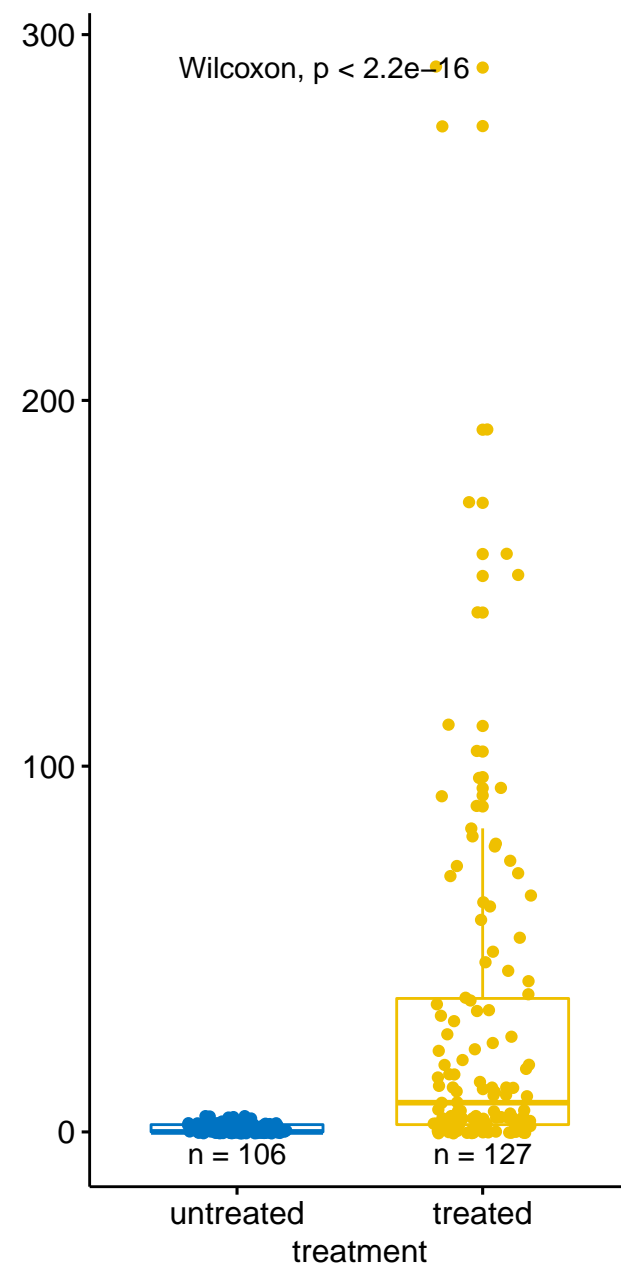
B # unique insertions per sgRNA
Min Freq = 0.003 %

treatment  untreated  treated





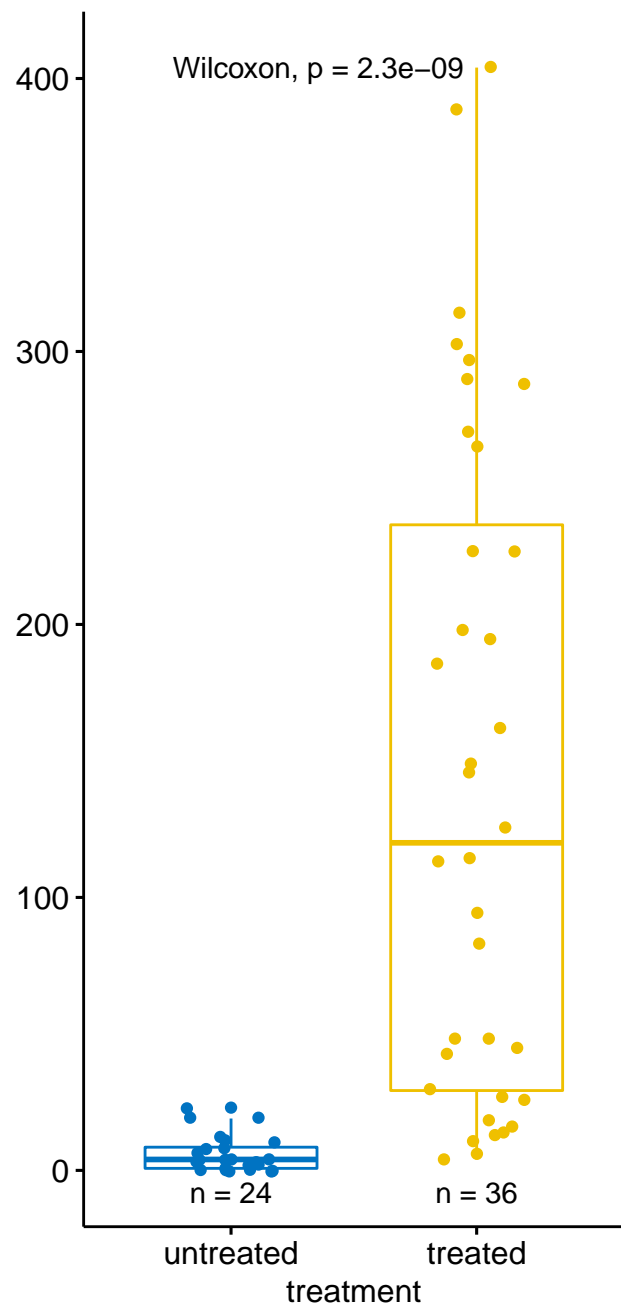
C # deletions
by sgRNA & sample
Min Freq = 0.003 %

treatment  untreated  treated





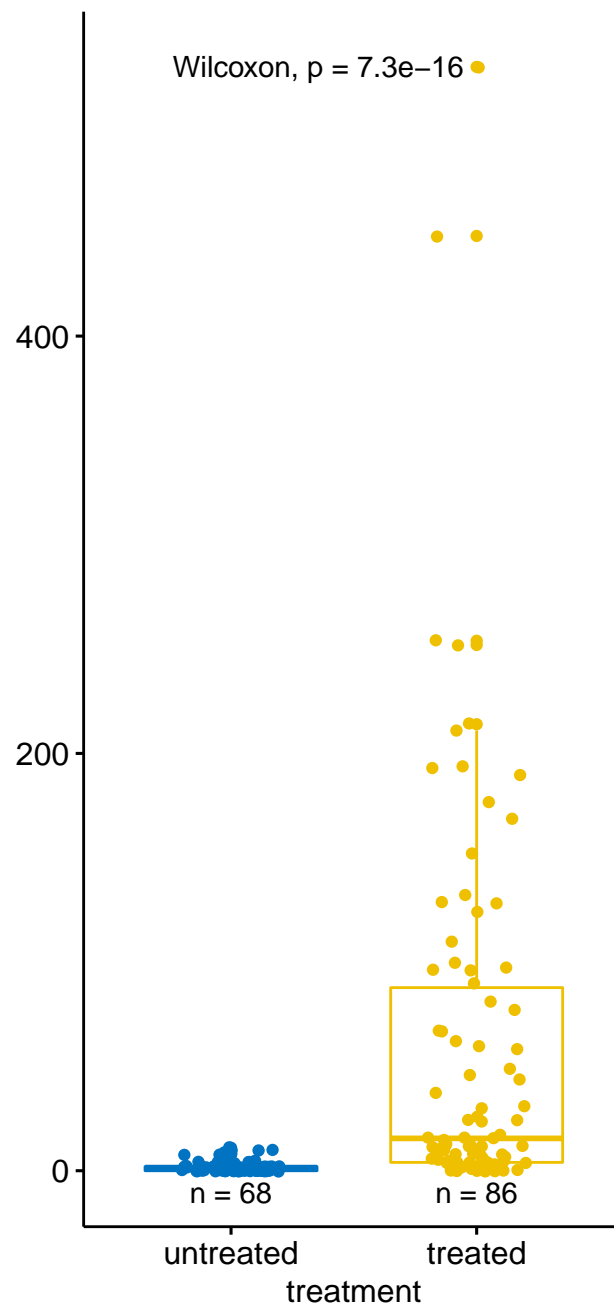
A # unique insertions per sample
Min Freq = 0.002 %

treatment  untreated  treated





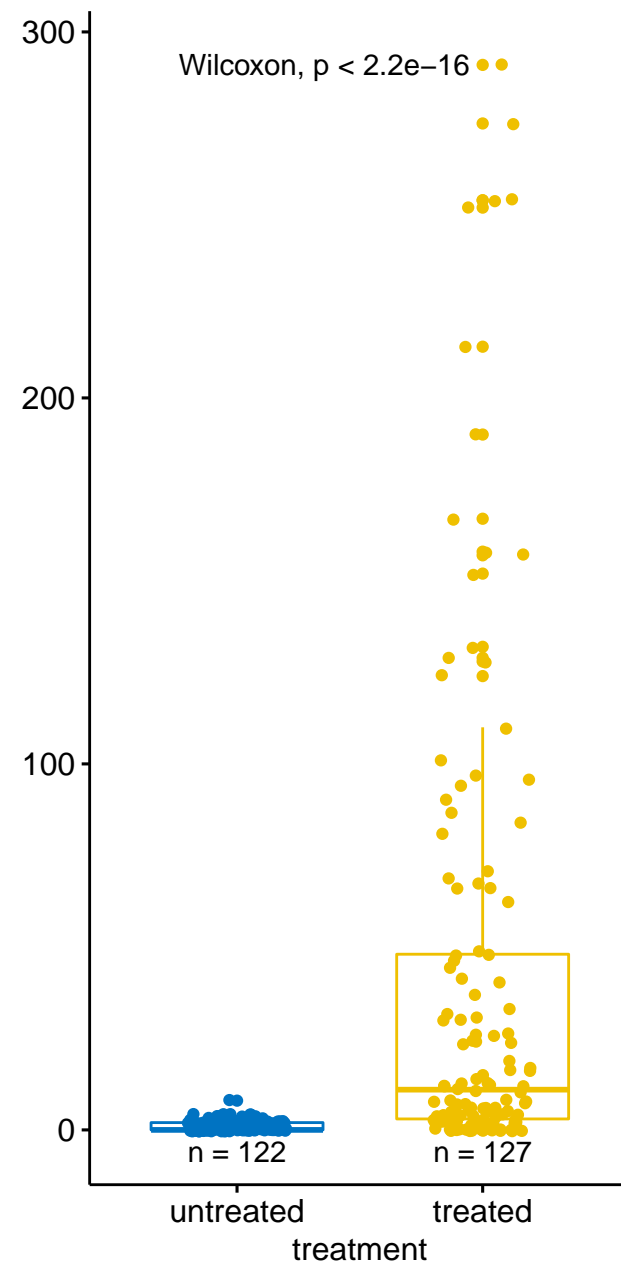
B # unique insertions per sgRNA
Min Freq = 0.002 %

treatment  untreated  treated





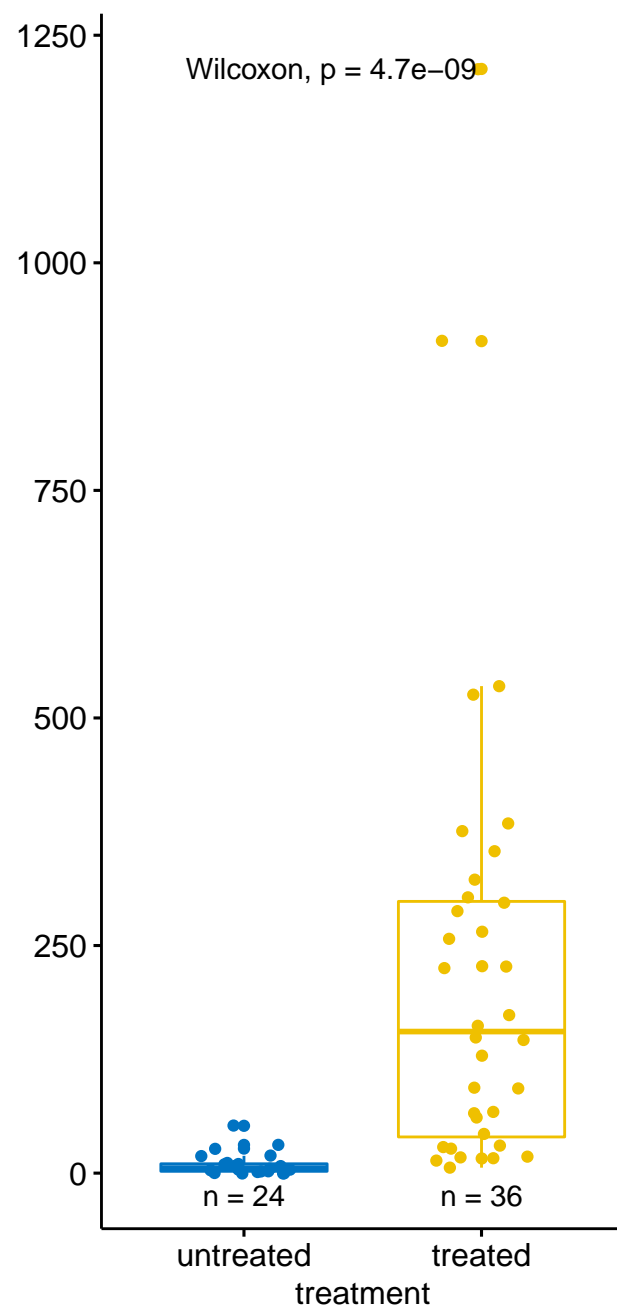
C # deletions
by sgRNA & sample
Min Freq = 0.002 %

treatment  untreated  treated



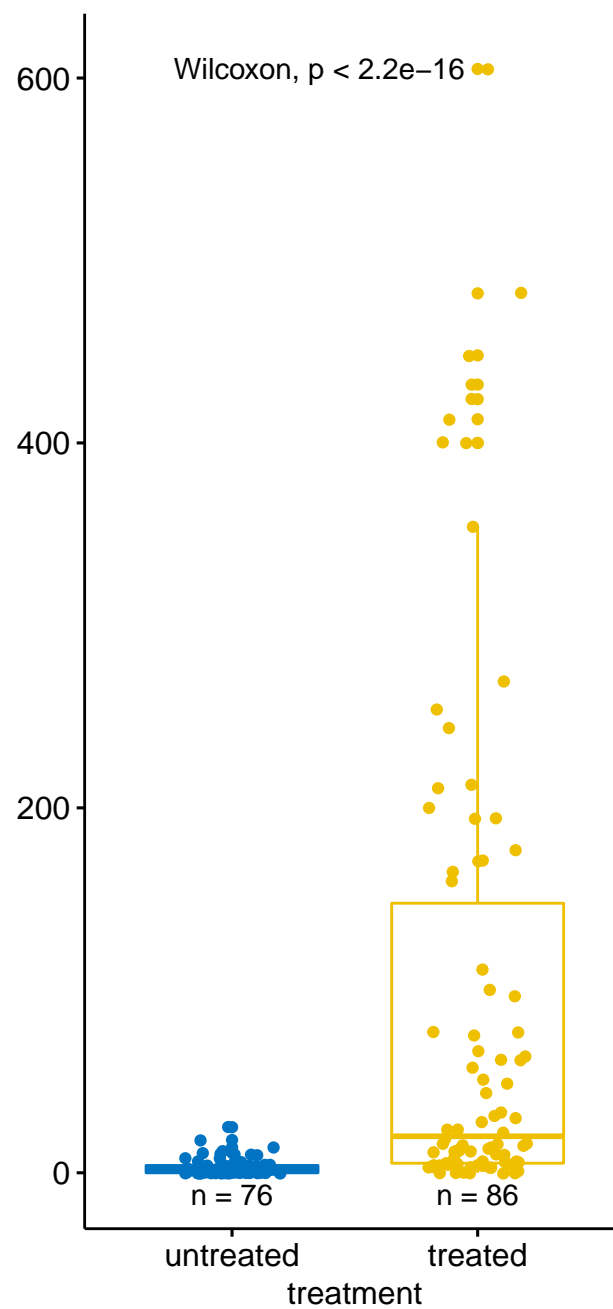
A # unique insertions per sample
Min Freq = 0.001 %

treatment  untreated  treated





B # unique insertions per sgRNA
Min Freq = 0.001 %

treatment  untreated  treated



C # deletions
by sgRNA & sample
Min Freq = 0.001 %

treatment  untreated  treated

