

Supplementary Materials: Regulation of Small RNAs and Corresponding Targets in Nod Factor-Induced *Phaseolus vulgaris* Root Hair Cells

Damien Formey, José Ángel Martín-Rodríguez, Alfonso Leija, Olivia Santana, Carmen Quinto, Luis Cárdenas and Georgina Hernández

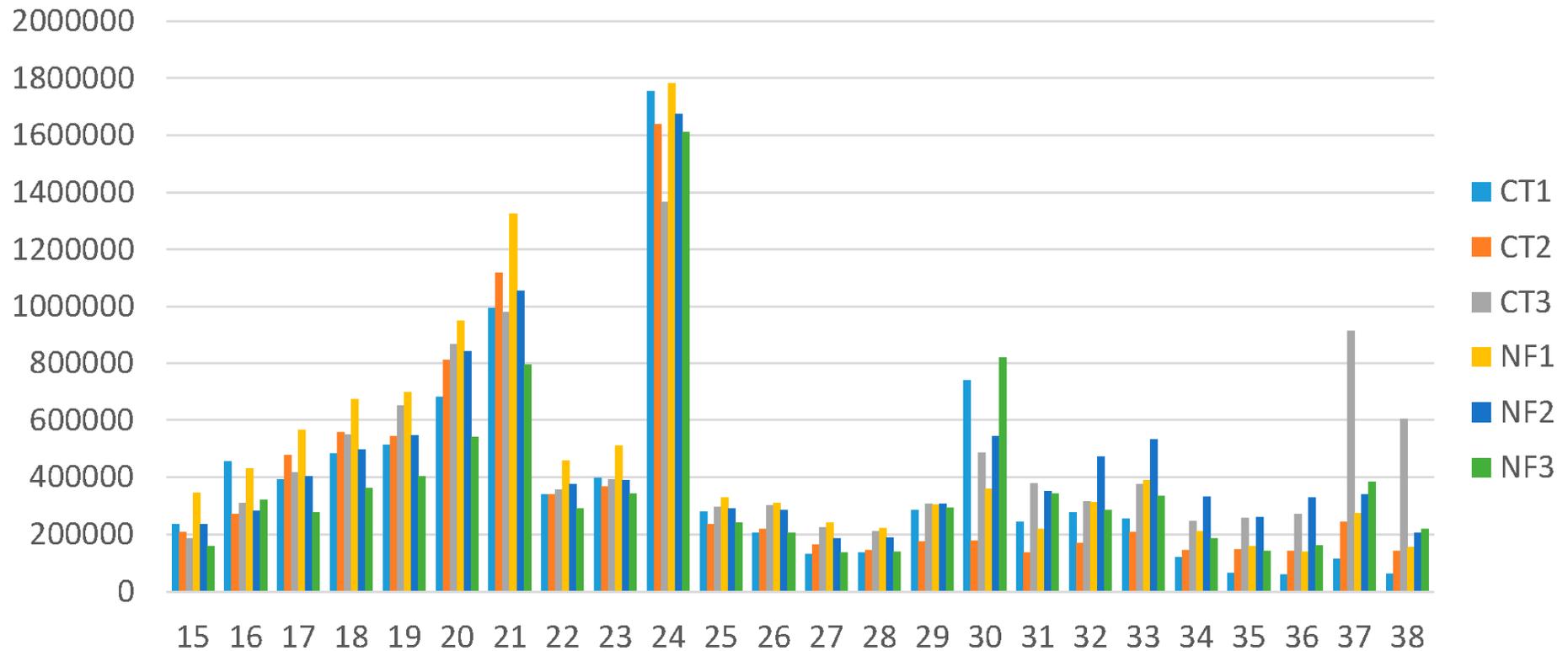


Figure S1. Size distribution of the sequencing reads of each library. X axis represents the read sizes (nt) and the Y axis represents the number of reads for each read size. The 6 histograms represent the 6 libraries sequences.