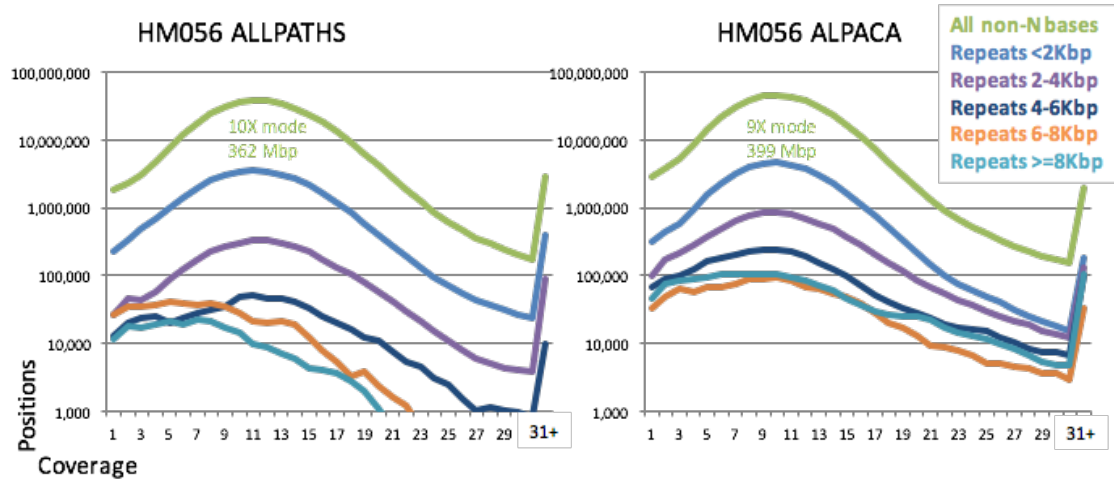




Supplemental Figure S1. Coverage distributions for same-scaffold repeats. Uncorrected long reads were mapped to ALLPATHS and Alpaca assemblies of *M. truncatula* HM056. Horizontal axis: bins representing scaffold positions covered by 1 or more reads. Vertical axis: Number of assembly positions having the given coverage. Colors: coverage over the genome as a whole (green) or over repeats binned by size.