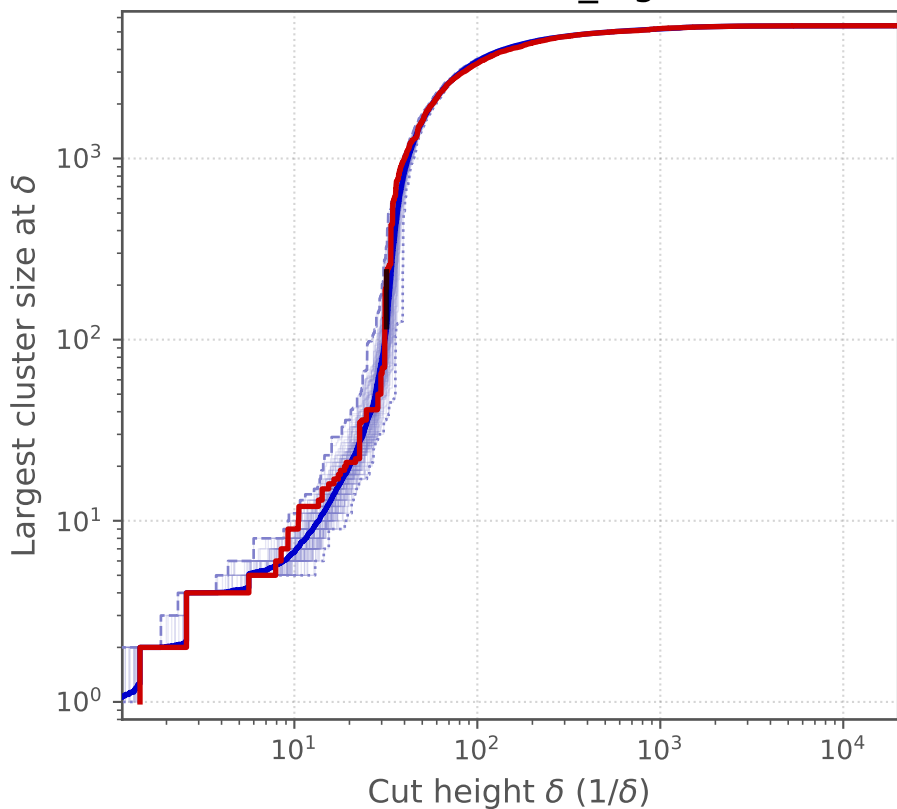


# Cluster sizes across hierarchy cuts for STRING BRCA\_logFC



— Observed sizes

— Expected sizes

— Chosen cut

..... Permuted sizes (minimum)

— Permuted sizes (all)

- - - Permuted sizes (maximum)