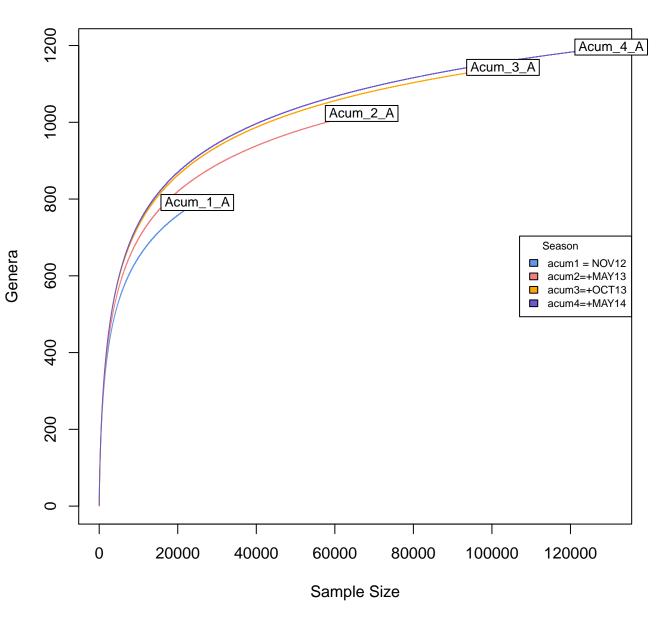
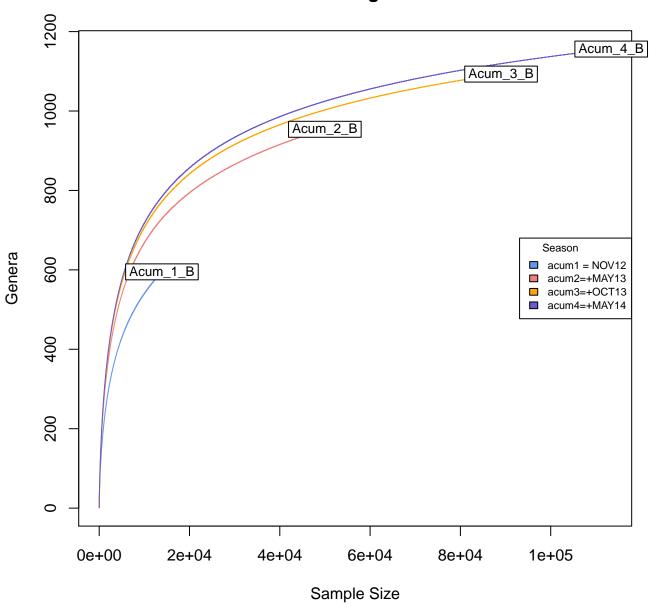
## Richness at genus level Site A



## Rarefaction curve at genus level Site B



## Rarefaction curve at genus level Site C

