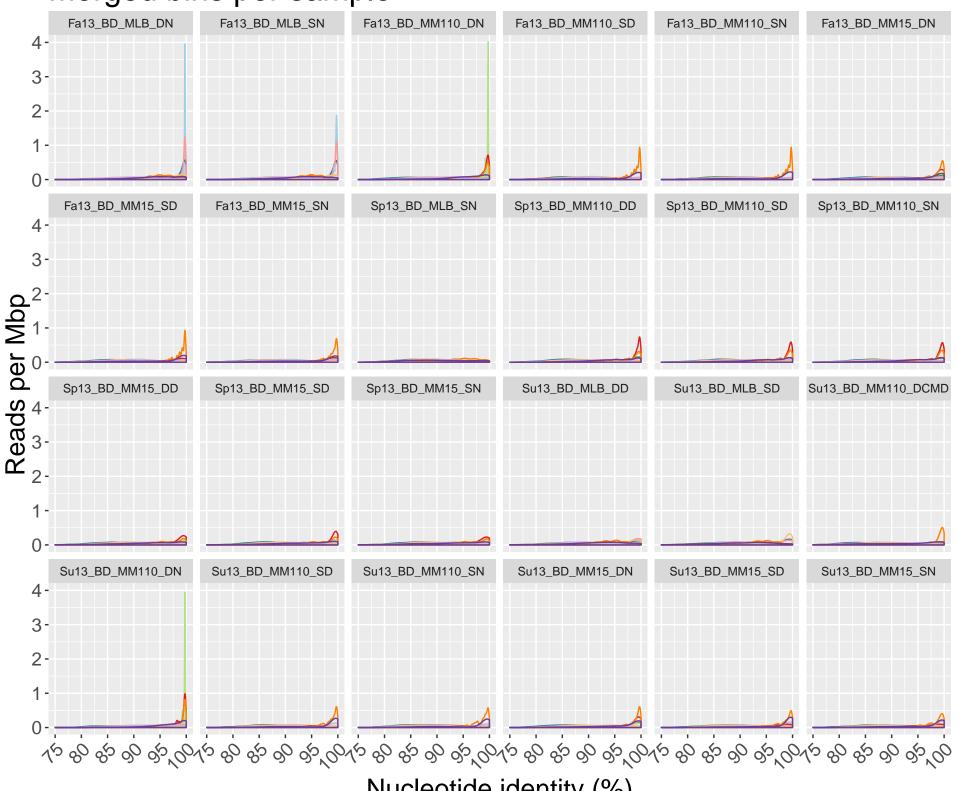
## Merged bins per sample



## Nucleotide identity (%)

