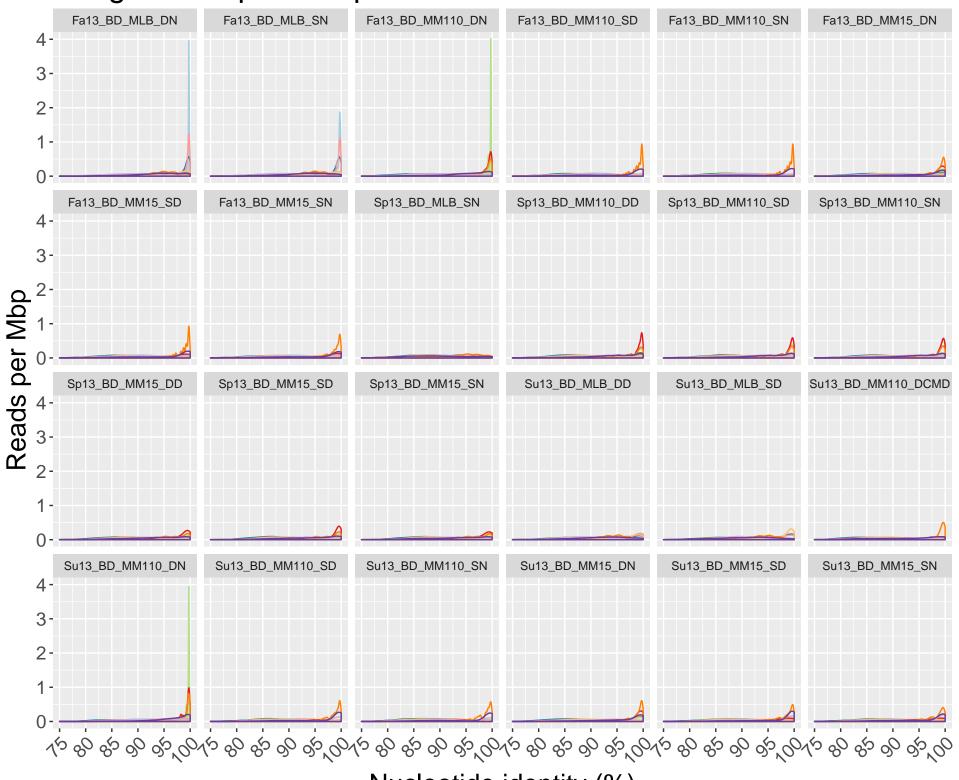
Merged bins per sample



Nucleotide identity (%)

