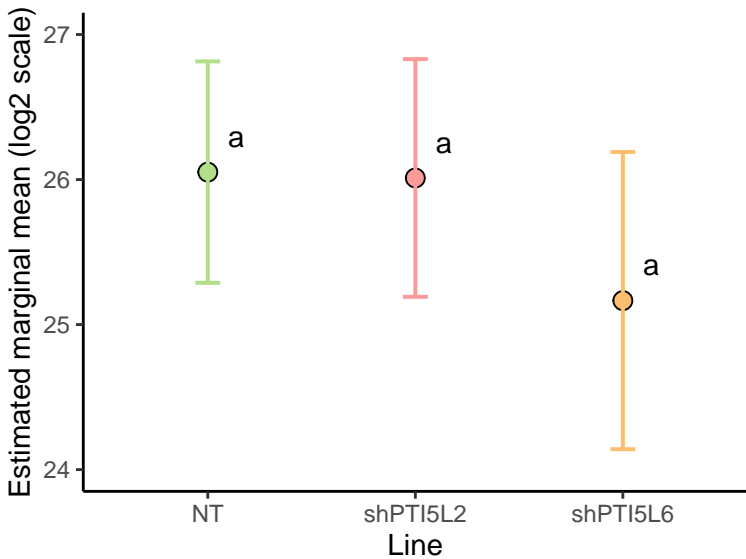


Estimated Marginal Means by Genotype with CLD



Genotype  NT  shPTI5L2  shPTI5L6