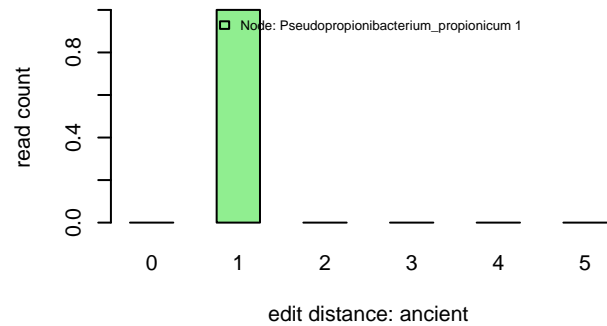
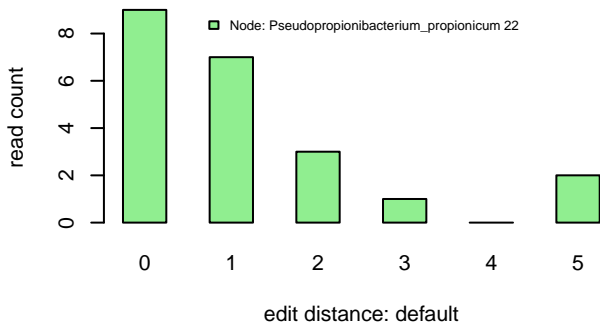
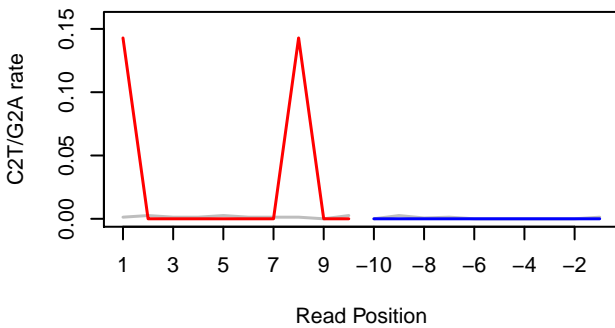


0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur0101\_S0\_L000\_R1\_000.fastq.merged.prefixed.hg19ur



### image plot for Pseudopropionibacterium\_propionicum



Node	Pseudopropionibacterium_propionicum
Top Reference	Pseudopropionibacterium_propionicum_F0230a
assigned reads	22
all reads	21
nonDup	21
readDis	0.681
nonStacked	11
destacking?	Off
C>T_1	0.1429
G>A_-1	0
mean length (sd)	50 (13.841)

Pseudopropionibacterium_propionicum_F0230a;_CP002734.1	100%
Pseudopropionibacterium_propionicum;_NR_102999.1	081%
Pseudopropionibacterium_propionicum;_MF574814.1	054%
Pseudopropionibacterium_propionicum;_MF574792.1	036%
Pseudopropionibacterium_propionicum;_LC102212.1	018%
uncultured_Propionibacterium_sp.;_AM420135.1	009%
uncultured_Propionibacterium_sp.;_AY807199.1	009%
uncultured_Propionibacterium_sp.;_FJ976260.1	009%
uncultured_Propionibacteriaceae_bacterium;_GU227181.1	009%
Pseudopropionibacterium_propionicum;_GU425642.1	009%