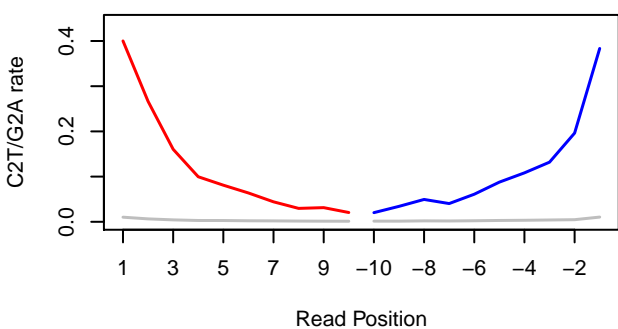


Damage plot for uncultured\_bacterium node



uncultured_bacterium;_HE681350.1 100%
Clostridiales_bacterium_oral_taxon_G74;_GU432185.1 100%
ostreptococcaceae_bacterium_canine_oral_taxon_030;_JN713194.1 100%
Eubacterium_sp._oral_clone_FX028;_AY134903.1 100%
Eubacterium_sp._oral_taxon_B60;_GU429169.1 100%
ostreptococcaceae_bacterium_feline_oral_taxon_028;_KM461989.1 100%
Clostridiales_bacterium_oral_taxon_G74;_GU432184.1 100%
Methanobrevibacter_millerae;_CP011266.1 100%
Saccharofermentans_sp._Marseille-P3694;_LT725666.1 100%
uncultured_Gemella_sp.;_EU071466.1 100%

Node	uncultured_bacterium
Top Reference	uncultured_bacterium
assigned reads	12675
all reads	394
nonDup	385
readDis	0.003
nonStacked	0
destacking?	Off
C>T_1	0.4001
G>A_-1	0.3835
mean length (sd)	54 (22.442)