Position (kbp)

3526.4   3526.6   3526.8   3527.0   3527.2

"TRA at chr10:3526827−91865625"

Sample: NA12877_S1

Delly

TRA : 0/1 : 80.9 Mbp
TRA : 0/1 : 88.34 Mbp
TRA : 0/1 : 184.55 Mbp
TRA : 0/1 : 86.68 Mbp
TRA : 0/1 : 26.5 Mbp

Depth (reads/ bp)

forward mapping distance (bp)

reverse mapping distance (bp)

clipped
diffmol
orphaned
inverted
samestrand

Sample: NA12878_S1

Delly

TRA : 0/1 : 80.9 Mbp
TRA : 0/1 : 88.34 Mbp
TRA : 0/1 : 184.55 Mbp
TRA : 0/1 : 86.68 Mbp
TRA : 0/1 : 26.5 Mbp

Depth (reads/ bp)

forward mapping distance (bp)

reverse mapping distance (bp)

clipped
diffmol
orphaned
inverted
samestrand

Delly

TRA : AF = 0 : 80.9 Mbp
TRA : AF = 0 : 88.34 Mbp
TRA : AF = 0 : 35.64 Mbp
TRA : AF = 0 : 184.55 Mbp
TRA : AF = 0 : 86.68 Mbp
TRA : AF = 0 : 26.5 Mbp
TRA : AF = 0 : 248.94 Mbp

None

3526.4   3526.6   3526.8   3527.0   3527.2

Position (kbp)

Read MapQ

>= 30   < 30   = 0

Inferred Insert Size
Proportion in position x
with mapping distance y

0   1

Mapping Stats
proportion of reads in position x

0   1

SV Allele Frequency

INV
CNV
DUP
DEL

0   1

Bin size: 10   Num bins: 100   Date: 2016−11−28