



max. family size =
absolute frequency=
relative frequency=

AB
85
1
0.06250

BA
332
1
0.06250

total nr. of tags=

total numbers *
16

total nr. of reads=

1,312

1892_2398= 3
1141_1564= 3
656_1143= 5
87_636= 5

* In the plot, both family sizes of the ab and ba strands were used.

Whereas the total numbers indicate only the single count of the tags per region.