

Figure 7: Rearrangements



Figure 8: Inversion

## 5.4.7 Inversions

#CHROM

Similarly an inversion such as in Figure 8:

POS

can be described equivalently in two ways. Either one uses the short hand notation described previously (recommended for simple cases):

| $\overset{\prime\prime}{2}$     | 321682 | 321681                             | INV0         | <b>T</b> - <u>G</u> <in< th=""><th>IV&gt; 6</th><th>PA</th><th>SS SVTYPE=INV;END=421681</th></in<> | IV> 6 | PA     | SS SVTYPE=INV;END=421681           |  |  |  |
|---------------------------------|--------|------------------------------------|--------------|--|-------|--------|------------------------------------|--|--|--|
| or one describes the breakends: |        |                                    |              |  |       |        |                                    |  |  |  |
| #CHROM                          | POS    | ID                                 | REF          | ALT  | QUAL  | FILTER | INFO                               |  |  |  |
| 2                               | 321681 | $\mathrm{bnd}_{	ext{-}}\mathrm{W}$ | G            | G[2:421681]  | 6     | PASS   | SVTYPE=BND;MATEID=bnd_U;EVENT=INV0 |  |  |  |
| 2                               | 321682 | $bnd_{-}V$                         | ${ m T}$     | [2:421682[T  | 6     | PASS   | SVTYPE=BND;MATEID=bnd_X;EVENT=INV0 |  |  |  |
| 2                               | 421681 | $bnd_U$                            | A            | A]2:321681]  | 6     | PASS   | SVTYPE=BND;MATEID=bnd_W;EVENT=INV0 |  |  |  |
| 2                               | 421682 | bnd X                              | $\mathbf{C}$ | [2:321682[C  | 6     | PASS   | SVTYPE=BND:MATEID=bnd V:EVENT=INV0 |  |  |  |

QUAL

FILTER

## 5.4.8 Uncertainty around breakend location

ID

REF

ALT

It sometimes is difficult to determine the exact position of a break, generally because of homologies between the sequences being modified, such as in Figure 9. The breakend is then placed arbitrarily at the left most position, and the uncertainty is represented with the CIPOS tag. The ALT string is then constructed assuming this arbitrary breakend choice.

The figure above represents a nonreciprocal translocation with microhomology. Even if we know that breakend U is rearranged with breakend V, actually placing these breaks can be extremely difficult. The red and green dashed lines represent the most extreme possible recombination events which are allowed by the sequence evidence available. We therefore place both U and V arbitrarily within the interval of possibility:

| #CHROM | POS    | $^{\mathrm{ID}}$                     | $\operatorname{REF}$ | ALT          | $_{ m QUAL}$ | FILTER | INFO                              |
|--------|--------|--------------------------------------|----------------------|--------------|--------------|--------|-----------------------------------|
| 2      | 321681 | $\mathrm{bnd}_{	extsf{-}}\mathrm{V}$ | ${ m T}$             | T]13:123462] | 6            | PASS   | SVTYPE=BND;MATEID=bnd_U;CIPOS=0,6 |
| 13     | 123456 | $bnd_{-}U$                           | A                    | A]2:321687]  | 6            | PASS   | SVTYPE=BND;MATEID=bnd_V;CIPOS=0,6 |

Note that the coordinate in breakend U's ALT string does not correspond to the designated position of breakend V, but to the position that V would take if U's position were fixed (and vice-versa). The CIPOS tags describe the uncertainty around the positions of U and V.

The fact that breakends U and V are mates is preserved thanks to the MATEID tags. If this were a reciprocal translocation, then there would be additional breakends X and Y, say with X the partner of V on Chr 2 and Y