

A

Sequence Display		
Junction sequence	> AAAAAAGGAAAGTGTGCCTATTCTTTTATGCATTGTATTTTCAACATATCAAACCTATTGGGGTCAGGTAATGTATTCTAAATACCCCTGTATTGTAATC	Strand: +/-
Virus sequence	CAAACCTATTGGGGTCAGGTAATGTATTCTAAATACCCCTGTATTGTAATC	Strand: -
Human sequence	AAAAAAGGAAAGTGTGCCTATTCTTTTATGCATTGTATTTTCAACATATCA	Strand: +
MH patterns	<pre> TTCACATAT.CA.AACTTATTGG TTCACATAT.CA.AACTTATTGG TTCACATAT.CA.AACTTATTGG </pre>	junctional microhomology/Snap-back junctional microhomology/Loop-out junctional microhomology/Loop-out
Microhomologies		

B

Sequence Display		
Junction sequence	> CTCCTCTTAGACAGATAGACACACACACACACACGCACACATCCAGTTCAGTTTCGCGAGGTGTGGAGGCTGCAATACAGATGGGTCAGTGAAG	Strand: +/-
Virus sequence	TCCAGTTTCGCGAGGTGTGGAGGCTGCAATACAGATGGGTCAGTGAAG	Strand: -
Human sequence	CTCCTCTTAGACAGATAGACACACACACACACACGCACACATCCAGT	Strand: +
MH patterns	CACATCCAGT-TCAGTTCT	apparent blunt join/Loop-out
<p>The end-join between human and virus sequence was displayed in three ways (junctional microhomology within dots; dash for apparent blunt join; lowercase for short insertion). The repeat, including primers (red) and microhomologies (blue), was underlined.</p>		