

Type	Microhomology (gene)	Microhomology (non-gene)	p-value (gene vs non-gene)
Deletions	1/8 [12.5%]	0/4 [0%]	1
Deletion disagreements > 3 bp	8/32 [25%]	7/15 [46.7%]	> 0.55
Deletion disagreements < 3 bp	5/34 [14.7%]	3/16 [18.8%]	1
Insertion disagreements < 3 bp	10/20 [50%]	7/10 [70%]	> 0.44
Random location	2/47 [4.2%]	4/13 [30.8%]	N/A

**Supplemental Table S10. Comparison of microhomology frequency around deletions and duplex indel disagreements, for gene regions and non-gene regions.**

Data analysis as in Supplemental Table S9.