

Supplementary Table S11: Number of sites nominated by the three genome-wide methods outside of up to 5 MM 0 gap, or up to 3 MM 1 gap homology space but within 7 MM, 5 gap space

gRNA	GUIDE-seq	CIRCLE-seq	SITE-seq
CCR5	5	3083	154
FANCF	5	2309	617
HEK1	6	1771	782
HEK2	6	7907	829
HEK3	8	5721	570
IL2RG	13	5425	3153
RNF2	14	1216	970
VEGFA1	37	7504	5908