





Genome
of interest

Sites of
interest

cuRRBS

Maximum no. of
fragments to sequence

Minimum proportion of
sites of interest

order	combination to use	size range	CRF	robustness
1st		40-200 bp	205	0.95
2nd		100-135 bp	163	0.4
...		200-350 bp	97	0.8
30th		50-250 bp	86	1