

Table S23.

Seed length	Average coverage		Number of fixed or polymorphic bases		Number of singleton bases	
	30	45	30	45	30	45
k1n3	25.4	25.1	15801	15336	74753	72716
k2n5	30	29.9	19737	19522	149263	148169
k3n7	32	32	22457	22306	157904	157193
k4n9	32.4	32.4	23891	23744	160718	160294

s = seed length k = number of errors allowed in the seed

n = number of errors allowed in the entire read