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**Simulated HIV mix (20 000x, 2x150 bp)**

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	# contigs $\geq 500$ bp	largest contig	N50	MAC length (%)	target genomes (%)	N-rate (%)	mismatches (%)	indels (%)
SAVAGE-de-novo	76	5914	1244	0	98.4	0	0.006	0
SAVAGE-h-ref	59	4093	1612	0	90.6	0	0	0
SAVAGE-b-ref	86	2622	1222	0	95.5	0.002	0.004	0

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Supplemental Table S3: Assembly results for SAVAGE on the 20.000x simulated 5-HIV-mix with 2x150 bp reads.