Report

# contigs (>= 1000 bp)		Einal access
# contigs (>= 5000 bp) 310 # contigs (>= 10000 bp) 164 # contigs (>= 25000 bp) 24 # contigs (>= 50000 bp) 0 Total length (>= 10000 bp) 3878820 Total length (>= 50000 bp) 2831064 Total length (>= 5000 bp) 3878820 Total length (>= 5000 bp) 727737 Total length (>= 50000 bp) 727737 Total length 725000 bp) 727737 Total length 72500 5000 bp) 727737 Total length 72500 5000 bp) 727737 Total length 72500 5000 5000 5000 5000 5000 5000 500	H continue (c. 1000)	final.contigs
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NGA75 6722 LA50 116	NGA50	12252
LA50 116	NA75	
	NGA75	6722
10150	LA50	
	LGA50	118
LA75 238		238
LGA75 243	LGA75	243

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Misassemblies report

	final.contigs
# misassemblies	1
# relocations	1
# translocations	0
# inversions	0
# possibly misassembled contigs	0
# misassembled contigs	1
Misassembled contigs length	7679
# local misassemblies	0
# mismatches	16
# indels	0
# short indels	0
# long indels	0
Indels length	0

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).

Unaligned report

	final.contigs
# fully unaligned contigs	0
Fully unaligned length	0
# partially unaligned contigs	0
# with misassembly	0
# both parts are significant	0
Partially unaligned length	0
# N's	0

All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).















