



Aug 11, 2023

Reverse transcription, primer pools preparation and multiplex PCR steps for DENV1 serotype for genomic sequencing

DOI

<https://dx.doi.org/10.17504/protocols.io.5jyl8pjyrg2w/v1>

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Protocol Citation: Laís Ceschini, Carla Julia da Silva Pessoa Vieira, Luisa Maria Inácio da Silva, Gustavo Lima, Raul Emídio, Tiago Graf, Gabriel Luz Wallau 2023. Reverse transcription, primer pools preparation and multiplex PCR steps for DENV1 serotype for genomic sequencing. **protocols.io** <https://dx.doi.org/10.17504/protocols.io.5jyl8pjyrg2w/v1>

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Protocol status: Working

We use this protocol and it's working

Created: August 10, 2023

Last Modified: August 11, 2023

Protocol Integer ID: 86328

Keywords: multiplex pcr steps for dengue serotype, complete genome of dengue serotype strain, dengue serotype strain, dengue serotype, multiplex pcr step, multiplex pcr condition, sequencing, multiplex pcr conditions with the main goal, cDNA synthesis, primer pools preparation, genomic, complete genome, transcription, strain

Funders Acknowledgements:

FIOCRUZ

Abstract

This step-by-step protocol describes the cDNA synthesis, primer pools preparation and multiplex PCR conditions with the main goal to sequence the complete genome of DENV1 serotype strains.

Materials

Reagents:

Reverse transcription: SuperScript™ IV First-Strand Synthesis System. (200 reactions) Cat: 18091200 Invitrogen

Multiplex PCR: Q5® High-Fidelity 2X Master Mix. Cat: M0492L NEB, H2O Ultrapure, primers described in table 1.



Reverse transcription

- 1 Using a 2mL tube prepare the **Mix 1** described below for 96 samples:

	A	B	C
	Mix 1 Reverse transcription	Vol. (1x)	96 samples (+2 = 98 to keep some extra due to pipetting issues)
	Random Hexamers (50μM)	1μL	98μL
	dNTPs mix (10mM each)	1μL	98μL
	Total	2μL	194μL

- 2 Using 0,2mL PCR tubes or 96 wells plates add 11-16μL of extracted RNA from RT-PCR positive samples. Add **2μL** of Mix 1 to the tube/well and take it to the thermocycler with the following set up

65°C ----- 5 minutes

- 3 Take the tubes/wells to ice for 1 minute (you can prepare a water bath with ice cubes to have a uniform temperature distribution).

- 4 Using a 2mL tube prepare **Mix 2**:

	A	B	C
	Mix 2 Reverse Transcription	Vol. (1x)	96 samples (+2 = 98 to keep some extra due to



	A	B	C
			pipetting issues)
	5x SSIV Buffer	4μL	392μL
	100mM DTT	1μL	98μL
	RNaseOUT or RNase Inhibitor	1μL	98μL
	SSIV Reverse Transcriptase	1μL	98μL
	Total	7μL	686μL

- 5 Add **7μL** of **Mix 2** to the tubes containing the **Mix 1** plus RNA and take it to the thermocycler following the set up below:

Step1:

42°C ---- 50 minutes

70°C ---- 10 minutes

4°C ---- Hold

- 6 Store the cDNA at -20°C.

Observation:.. As a suggestion, to improve the final results only samples RT-PCR positive showing a Ct value of < 30 should be used for cDNA conversion and genomic amplification.

Pools of primers

- 7 Select two 0,6mL tubes for each pool.
- 8 Using the original 100uM primer solution eluted individually, put them together following the table below containing each primer volume.
- 9 Pool 1 will have a final volume of 130μl and pool 2 of 170μl.

- 10 In order to prepare the solution to use in the Multiplex PCR, dilute each pool 1:10. That is, 10µl of pool 1 and 90µl of ultrapure water.

TABLE 1: Primers and pool order

	A	B	C	D	E	F
	Primer	Sequence	Tm	Concentration inside of the pool *	Volume of primer within the pool	POOL
	DENV1SA_1_LEFT	TACGTGGA CCGACAA GAACAGT	58.1°C	0,0075uM	2,5µl	1
	DENV1SA_1_RIGHT	ACTATCAT RTGTGGCT CTCCCC	57.2°C	0,0075uM	2,5µl	1
	DENV1SA_3_LEFT	CACACGTG GGACTTG GTCTAGA	58.4°C	0,01125uM	7,5µl	1
	DENV1SA_3_RIGHT	ACACACAA AGTTCGCG TCTTGT	57.6°C	0,01125uM	7,5µl	1
	DENV1SA_5_LEFT	CCTCACAT TGGACTGC TCACCT	59.1°C	0,01125uM	7,5µl	1
	DENV1SA_5_RIGHT	TGCACTAR RACAGTTC CATGCT	56.6°C	0,01125uM	7,5µl	1
	DENV1SA_7_LEFT	CGAGGAG CACGAAG GATGGC	60.6°C	0,0075uM	2,5µl	1
	DENV1SA_7_RIGHT	ATGATGTT CTCAAGAC GCGTGG	57.5°C	0,0075uM	2,5µl	1
	DENV1SA_9_LEFT	TGGGAAGT TGAGGACT AYGGGT	58.7°C	0,015uM	5µl	1
	DENV1SA_9_RIGHT	TGTRGTTC TGAGRGAT GGACCTC	57.8°C	0,015uM	5µl	1
	DENV1SA_11_LEFT	GATGACTG GAACACTG GCTGTT	57.4°C	0,030uM	10µl	1



	A	B	C	D	E	F
	DENV1SA_1 1_RIGHTDEN V1SA_11_RI GHT	CACCGGA AGCCATGT TGTTTTT	56.7°C	0,030uM	10µl	1
	DENV1SA_1 3_LEFT	AASAAGAA GCAGAACA CTCCGG	57.3°C	0,015uM	5µl	1
	DENV1SA_1 3_RIGHT	ACTGGCC CAGCTTGG TTCCAG	62.4°C	0,015uM	5µl	1
	DENV1SA_1 5_LEFT	ATGGAGTG GTGACAAC AAGTGG	57.4°C	0,015uM	5µl	1
	DENV1SA_1 5_RIGHT	GCTGGATC GGTAAART GTGCTTC	57.4°C	0,015uM	5µl	1
	DENV1SA_1 7_LEFT	ACGGGTRA TYCAAYTG AGCAGRA	58.5°C	0,01125uM	7,5µl	1
	DENV1SA_1 7_RIGHT	CCTCTTCT CATGAGCT CCACA	56.3°C	0,01125uM	7,5µl	1
	DENV1SA_1 9_LEFT	AGTGTCTC AGGTGACC TAATATTG GA	57°C	0,0075uM	2,5µl	1
	DENV1SA_1 9_RIGHT	RGCTGCC ACTGTCAG TATCATG	57.5°C	0,0075uM	2,5µl	1
	DENV1SA_2 1_LEFT	YGCAAAYC AGGCWGC YATATTGA T	57.6°C	0,015uM	5µl	1
	DENV1SA_2 1_RIGHT	GATGTTTG CCATGGAC ACTGCT	58.2°C	0,015uM	5µl	1
	DENV1SA_2 3_LEFT	ACAACCAA ACATGCAG TGTCGA	57.5°C	0,015uM	5µl	1
	DENV1SA_2 3_RIGHT	TTTCGCAC TAGCATCC CTCCAT	58.5°C	0,015uM	5µl	1
	DENV1SA_2 5_LEFT	ACCTAGAT ATYATTGG	55.9°C	0,030uM	10µl	1



	A	B	C	D	E	F
		CCAGAGG A				
	DENV1SA_2 5_RIGHT	ACCTTTTCG TCTTCCAC TGCTTC	57.3°C	0,030uM	10µl	1
	DENV1SA_2 7_LEFT	TGGAAGG AGAAGGAC TGCACAA	58.4°C	0,030uM	10µl	1
	DENV1SA_2 7_RIGHT	CACRCAAT CATCTCCG CTRATT	55.5°C	0,030uM	10µl	1
	DENV1SA_2 9_LEFT	ATGGAGCC TGAGAGAA ACTGCT	58.2°C	0,030uM	10µl	1
	DENV1SA_2 9_RIGHT	GCYCCTTC GGGATCAC TCTCAT	59.7°C	0,030uM	10µl	1
	DENV1SA_2 _LEFT	TGTTGAAC ATAATRAA CAGGAGG AA AAGA	55.9°C	0,01125uM	7,5µl	2
	DENV1SA_2 _RIGHT	GAATCCTG GGTGTCKC AAAGCC	59.5°C	0,01125uM	7,5µl	2
	DENV1SA_4 _LEFT	ACTGTGCA TTGAAGCT AAAATATC AA ACA	56°C	0,015uM	5µl	2
	DENV1SA_4 _RIGHT	ACCATTGT TTGTGGAC AAGCCA	57.7°C	0,015uM	5µl	2
	DENV1SA_6 _LEFT	AAACTGAC YTTARAGG GGATGTCA T	56.1°C	0,015uM	5µl	2
	DENV1SA_6 _RIGHT	ATATGCRG TCCCAAA AACCTGG	56.7°C	0,015uM	5µl	2
	DENV1SA_8 _LEFT	AGGCTGA CTCCCCA AAAAGACT	58.5°C	0,015uM	5µl	2



	A	B	C	D	E	F
	DENV1SA_8_RIGHT	TTGATGGC AGCTGACA TTAGCC	57.8°C	0,015uM	5µl	2
	DENV1SA_10_LEFT	GCAGGGC CATGGCAC CTAGG	63.5°C	0,0075uM	2,5µl	2
	DENV1SA_10_RIGHT	TCCCCATC CTGTCTGA AGCATT	58.4°C	0,0075uM	2,5µl	2
	DENV1SA_12_LEFT	GGATTATG CATGGAAR ACAAYGGC	56.9°C	0,030uM	10µl	2
	DENV1SA_12_RIGHT	GTGAGTGT RTCATCCC TYTCTTCA	56.2°C	0,030uM	10µl	2
	DENV1SA_14_LEFT	AGGTCCC AAGTAGGA GTGGGAGT	61.2°C	0,015uM	5µl	2
	DENV1SA_14_RIGHT	CACCTCRT CCTCAATC TCTGGT	57.2°C	0,015uM	5µl	2
	DENV1SA_16_LEFT	GGGAGATA GTTGACCT CATGTGCC A	60.3°C	0,015uM	5µl	2
	DENV1SA_16_RIGHT	CCTGTCTG GCCCGGA AATTTGC	61.7°C	0,015uM	5µl	2
	DENV1SA_18_LEFT	CAGAAGG GATCATCC CAGCCCT	60.9°C	0,0075uM	2,5µl	2
	DENV1SA_18_RIGHT	CCTCCTTG TTCGGAAT TGTGCA	57.9°C	0,0075uM	2,5µl	2
	DENV1SA_20_LEFT	GCTGCTCA TTCCAGAR CCAGAC	59.2°C	0,015uM	5µl	2
	DENV1SA_20_RIGHT	ATGGGTTC ACCTGGG AATAGCA	58.4°C	0,015uM	5µl	2
	DENV1SA_22_LEFT	TCCATCAC ACTGGCTA CTGGAC	58.6°C	0,015uM	5µl	2

	A	B	C	D	E	F
	DENV1SA_2 2_RIGHT	CCCACAA CCGAGGT CTATGACT	58.4°C	0,015uM	5µl	2
	DENV1SA_2 4_LEFT	GCTYAGAG GAAACCA ATTCTGCA	56.5°C	0,030uM	10µl	2
	DENV1SA_2 4_RIGHT	TGATCCTG ATGGYTTG ACCTCA	54.7°C	0,030uM	10µl	2
	DENV1SA_2 6_LEFT	CTGCACAA GAGAGGA GTTTACA	56.8°C	0,015uM	5µl	2
	DENV1SA_2 6_RIGHT	TATTCTTG TGTCCCAT CCGGCT	58.3°C	0,015uM	5µl	2
	DENV1SA_2 8_LEFT	GAAACCC CCAACTA GCTRAGA	56.4°C	0,030uM	10µl	2
	DENV1SA_2 8_RIGHT	TAGCCGCT AGTCTCAG GTCTCT	58.8°C	0,030uM	10µl	2
	DENV1SA_3 0_LEFT	GGGCCAC YAATATAC AAGTAGCC A	57.6°C	0,030uM	10µl	2
	DENV1SA_3 0_RIGHT	CCCGCTG CTGCGTTA TGTCT	60.4°C	0,015uM	10µl	2
	DENV1SA_3 1_RIGHT	CCTGTTGA TTCAACAG CACCATT CA	59.7°C	0,015uM	10µl	2

*approximate concentration of each primer in the 25µl PCR reaction.

Note: The primers were designed using the <https://primalscheme.com> (Brito, 2021) based on the JX669463.1 and KP188568 reference genomes.

Multiplex PCR

- 11 Prepare the **Mix 1** for a Multiplex PCR for each **Pool 1 and Pool 2** using a Falcon tube of 15mL (~96 amostras) or a 2mL tube.



	A	B	C	D
	Mix 1 Multiplex PCR	Vol. Pool 1 (1x)	Vol. Pool 2 (1x)	96 samples (+2) (pool1 or pool2)
	Q5 Master Mix High fidelity 2X	12,5 µl	12,5 µl	1.225 µl
	Pool primers (Pool1 ou Pool2) /Use concentrati on/	1,5 µl	1,5 µl	147 µl
	Ultra Pure Water	8,5 µl	8,5 µl	833 µl
	Total	22,5µl	22,5µl	2205µl

- 12 Add **2,5µl of cDNA** (totalling 5µl) in 22,5µl of the pool1 and pool2 reaction and take it to the thermocycler following the conditions bellow:

Step1:

98°C ---- 30 seconds

Step2: (45 cycles)

98°C ---- 15 seconds

58°C ---- 30 seconds

72°C ---- 5 minutes

Step3:

72°C ---- 2 minutes

Hold 4°C