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

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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.

Protocol status: Working
We use this protocol and it's working

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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 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	TACGTGGAC CGACAAGAA CAGT	58.1°C	0,0075uM	2,5µl	1
DENV1SA_1_RIGHT	ACTATCATRT GTGGCTCTC CCC	57.2°C	0,0075uM	2,5µl	1
DENV1SA_3_LEFT	CACACGTGG GACTTGGTC TAGA	58.4°C	0,01125uM	7,5µl	1
DENV1SA_3_RIGHT	ACACACAAA GTTTCGCGTC TTGT	57.6°C	0,01125uM	7,5µl	1
DENV1SA_5_LEFT	CCTCACATT GGACTGCTC ACCT	59.1°C	0,01125uM	7,5µl	1
DENV1SA_5_RIGHT	TGCACTARR ACAGTTCCA TGCT	56.6°C	0,01125uM	7,5µl	1
DENV1SA_7_LEFT	CGAGGAGCA CGAAGGATG GC	60.6°C	0,0075uM	2,5µl	1
DENV1SA_7_RIGHT	ATGATGTTCT CAAGACGCG TGG	57.5°C	0,0075uM	2,5µl	1
DENV1SA_9_LEFT	TGGGAAGTT GAGGACTAY GGGT	58.7°C	0,015uM	5µl	1
DENV1SA_9_RIGHT	TGTRGTTCTG AGRGATGGA CCTC	57.8°C	0,015uM	5µl	1
DENV1SA_11_LEFT	GATGACTGG AACACTGGC TGTT	57.4°C	0,030uM	10µl	1

A	B	C	D	E	F
DENV1SA_11_RIGHT	CACCGGAAG CCATGTTGTT TTT	56.7°C	0,030uM	10µl	1
DENV1SA_13_LEFT	AASAAGAAG CAGAACACT CCGG	57.3°C	0,015uM	5µl	1
DENV1SA_13_RIGHT	ACTGGCCCA GCTTGGTTC CAG	62.4°C	0,015uM	5µl	1
DENV1SA_15_LEFT	ATGGAGTGG TGACAACAA GTGG	57.4°C	0,015uM	5µl	1
DENV1SA_15_RIGHT	GCTGGATCG GTAAARTGT GCTTC	57.4°C	0,015uM	5µl	1
DENV1SA_17_LEFT	ACGGGTRAT YCAAYTGAG CAGRA	58.5°C	0,01125uM	7,5µl	1
DENV1SA_17_RIGHT	CCTCTTCTCA TGAGCTCCA CA	56.3°C	0,01125uM	7,5µl	1
DENV1SA_19_LEFT	AGTGTCTCA GGTGACCTA ATATTGGA	57°C	0,0075uM	2,5µl	1
DENV1SA_19_RIGHT	RGCTGCCAC TGTCAGTAT CATG	57.5°C	0,0075uM	2,5µl	1
DENV1SA_21_LEFT	YGCAAAYCA GGCWGCYAT ATTGAT	57.6°C	0,015uM	5µl	1
DENV1SA_21_RIGHT	GATGTTTGC CATGGACAC TGCT	58.2°C	0,015uM	5µl	1
DENV1SA_23_LEFT	ACAACCAAA CATGCAGTG TCGA	57.5°C	0,015uM	5µl	1
DENV1SA_23_RIGHT	TTTCGCACT AGCATCCCT CCAT	58.5°C	0,015uM	5µl	1
DENV1SA_25_LEFT	ACCTAGATA TYATTGGCC AGAGGA	55.9°C	0,030uM	10µl	1
DENV1SA_25_RIGHT	ACCTTTCGT CTTCCACTG CTTC	57.3°C	0,030uM	10µl	1
DENV1SA_27_LEFT	TGGAAGGAG AAGGACTGC ACAA	58.4°C	0,030uM	10µl	1
DENV1SA_27_RIGHT	CACRCAATC ATCTCCGCT RATT	55.5°C	0,030uM	10µl	1
DENV1SA_29_LEFT	ATGGAGCCT GAGAGAAAC TGCT	58.2°C	0,030uM	10µl	1

A	B	C	D	E	F
DENV1SA_29_RIGHT	GCYCCTTCG GGATCACTC TCAT	59.7°C	0,030uM	10µl	1
DENV1SA_2_LEFT	TGTTGAACA TAATRAACA GGAGGAA AAGA	55.9°C	0,01125uM	7,5µl	2
DENV1SA_2_RIGHT	GAATCCTGG GTGTCKCAA AGCC	59.5°C	0,01125uM	7,5µl	2
DENV1SA_4_LEFT	ACTGTGCAT TGAAGCTAA AATATCAA ACA	56°C	0,015uM	5µl	2
DENV1SA_4_RIGHT	ACCATTGTTT GTGGACAAG CCA	57.7°C	0,015uM	5µl	2
DENV1SA_6_LEFT	AAACTGACY TTARAGGGG ATGTCAT	56.1°C	0,015uM	5µl	2
DENV1SA_6_RIGHT	ATATGCRGT CCCAAAAAC CTGG	56.7°C	0,015uM	5µl	2
DENV1SA_8_LEFT	AGGCTGACT CCCCAAAAA GACT	58.5°C	0,015uM	5µl	2
DENV1SA_8_RIGHT	TTGATGGCA GCTGACATT AGCC	57.8°C	0,015uM	5µl	2
DENV1SA_10_LEFT	GCAGGGCCA TGGCACCTA GG	63.5°C	0,0075uM	2,5µl	2
DENV1SA_10_RIGHT	TCCCCATCC TGTCTGAAG CATT	58.4°C	0,0075uM	2,5µl	2
DENV1SA_12_LEFT	GGATTATGC ATGGAARAC AAYGGC	56.9°C	0,030uM	10µl	2
DENV1SA_12_RIGHT	GTGAGTGTR TCATCCCTYT CTTCA	56.2°C	0,030uM	10µl	2
DENV1SA_14_LEFT	AGGTCCCAA GTAGGAGTG GGAGT	61.2°C	0,015uM	5µl	2
DENV1SA_14_RIGHT	CACCTCRTC CTCAATCTCT GGT	57.2°C	0,015uM	5µl	2
DENV1SA_16_LEFT	GGGAGATAG TTGACCTCA TGTGCCA	60.3°C	0,015uM	5µl	2
DENV1SA_16_RIGHT	CCTGTCGGC CCGGAAATT TGC	61.7°C	0,015uM	5µl	2

A	B	C	D	E	F
DENV1SA_18 _LEFT	CAGAAGGGA TCATCCCAG CCCT	60.9°C	0,0075uM	2,5µl	2
DENV1SA_18 _RIGHT	CCTCCTTGTT CGGAATTGT GCA	57.9°C	0,0075uM	2,5µl	2
DENV1SA_20 _LEFT	GCTGCTCAT TCCAGARCC AGAC	59.2°C	0,015uM	5µl	2
DENV1SA_20 _RIGHT	ATGGGTTCA CCTGGGAAT AGCA	58.4°C	0,015uM	5µl	2
DENV1SA_22 _LEFT	TCCATCACA CTGGCTACT GGAC	58.6°C	0,015uM	5µl	2
DENV1SA_22 _RIGHT	CCCACAACC GAGGTCTAT GACT	58.4°C	0,015uM	5µl	2
DENV1SA_24 _LEFT	GCTYAGAGG AAACCAATT CTGCA	56.5°C	0,030uM	10µl	2
DENV1SA_24 _RIGHT	TGATCCTGA TGGYTTGAC CTCA	54.7°C	0,030uM	10µl	2
DENV1SA_26 _LEFT	CTGCACAAG AGAGGAGTT CACA	56.8°C	0,015uM	5µl	2
DENV1SA_26 _RIGHT	TATTCTTGTTG TCCCATCCG GCT	58.3°C	0,015uM	5µl	2
DENV1SA_28 _LEFT	GAAACCCCC AAYCTAGCT RAGA	56.4°C	0,030uM	10µl	2
DENV1SA_28 _RIGHT	TAGCCGCTA GTCTCAGGT CTCT	58.8°C	0,030uM	10µl	2
DENV1SA_30 _LEFT	GGGCCACYA ATATACAAG TAGCCA	57.6°C	0,030uM	10µl	2
DENV1SA_30 _RIGHT	CCCGCTGCT GCGTTATGT CT	60.4°C	0,015uM	10µl	2
DENV1SA_31 _RIGHT	CCTGTTGATT CAACAGCAC CATTCCA	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 concentration /	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