

### cDNA synthesis and PCR protocol for Metagenomics

# **Reagents and Supplies**

Q5® High-Fidelity DNA Polymerase SuperScript<sup>TM</sup> IV First-Strand Synthesis

dNTP

Micropipette set

**Primers** 

Tips

Lab coat

Powder-free gloves

PCR tubes 0.2ml

Tube racks 0.2ml

Thermocycler

Laminar flow chapel

## Procedures to cDNA synthesis

a) Add in a 0.2mL tube for each reaction:

RLB RT 9N (2 uM) - 1  $\mu$ L

 $dNTPs (10 \text{ mM}) - 1\mu L$ 

Treated RNA - 10 μL

- b) Place the tube to incubate in a heat block for 5 minutes at 65 °C.
- c) Then place the tube **immediately on the ice** to prevent the formation of secondary structures.
- d) On each tube of the previous reaction, add  $8 \mu L$  of the Master Mix

below:

SSIV buffer (5x) -4  $\mu$ L

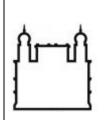
DTT (100 mM) -1  $\mu$ L

RNase OUT (inhibitor) -1μL

SS IV RTase (200U/ul) -1  $\mu$ L

RLB **TSO** (2  $\mu$ L) -1  $\mu$ L

 $TOTAL = 20\mu L$ 



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e) Then place the tubes in the thermocycler following the configuration represented in the table below:

Number of cycles	Temperature (°C)	Time
1	42	1h30 minutes
1	70	10 mins
1	4	$\infty$

### **Procedures to Metagenomics PCR**

a) Prepare the PCR Mix by following the volumes for 1 reaction shown in the table below:

Component	Amount for 1 reaction	
Ultra pure water	15.75 μL	
Q5 reaction buffer (5x)	5 μL	
dNTPs (10uM)	0.5 μL	
RLB PCR 20 uM primer	1 μL	

- b) After calculating the number of reactions to be prepared for the Master mix, **transfer** 22.5 μL of the mix to the 0.2mL PCR tubes and then, in another physically separated work area, add 2.5 μL of cDNA to each PCR tube containing the Master mix.
- c) Place the tubes in the thermocycler, following the configuration represented in the table below:

Number of cycles	Temperature (°C)	Time
1	98	45 seconds
26	98	15 seconds
	56	15 seconds
	65	5 mins
1	65	6 mins
	4	$\infty$

#### References

Chiu C, Greninger AL, Naccache SN, Federman S, Yu G, Mbala P, Bres V, Stryke D, Bouquet J, Somasekar S, Linnen JM. Rapid metagenomic identification of viral pathogens in clinical samples by real-time nanopore sequencing analysis.