

PCR for metagenomics

Using Platinum II polymerase for 16S, ITS and mcrA gene analysis

Prep:

- 01 Thaw an aliquot of **PCR-grade water**.
- 02 Thaw an aliquot of **Platinum II polymerase**, keep it on ice.
- 03 Thaw **10 μM primer** aliquots, keep them on ice.
- 04 Dilute reasonable volumes of primers you will need to **2.5 μM**
i.e. 2 μL of primer per sample plus excess
- 05 **Calculate** the master mix composition you will need for all the samples together.
- 06 Compose the **master mix** for the whole PCR reaction. Don't forget to include negative controls, where instead of DNA you add pure water.

Master mix	1 \times	\times
Polymerase 2 \times	10,0 μL	
FW primer 2.5 μM	2,0 μL	–
REV primer 2.5 μM	2,0 μL	–
DEPC-H ₂ O	9,0 μL	
DNA sample	2,0 μL	–

- 07 Pipette **19 μL** of the master mix into the wells.
- 08 Pipette **2 μL** of the **2.5 μM forward** primer. One row, one primer.
- 09 Pipette **2 μL** of the **2.5 μM reverse** primer. One column, one primer.
- 10 Pipette **2 μL** of your **DNA** sample into the well
- 10 Cover the wells with strong adhesive foil. Make sure it **adhered properly**.
- 11 Do a quick vortex to mix the reactions and **quick spin**.
- 12 Put the plate into the cycler and run the following programme. You need to change the **ramp rate of the annealing to 60 %**.

Amplification programme	Temperature	Time	
Initial denaturation	95 °C	3 min	
Denaturation	95 °C	45 s	35\times
Annealing	52 °C	60 s	
Extension	72 °C	90 s	
Final extension	72 °C	5 min	
	12 °C	Hold	