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<u>Protocol for DNA extraction from saliva</u> <u>samples, used for shotgun microbiome analysis</u>

Other SOPs for shotgun microbiome analysis can be accessed at: <u>http://www.microbiome-standards.org/</u> & <u>http://mgps.eu/standard-operating-procedure/</u>





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1. OBJECTIVE

This protocol is an adaptation of MGP_SOP 001 V1 for saliva samples collected with DNA Genotek's OMNIGene·ORAL. MGP_SOP 001 V1 is a standardized stool DNA extraction protocol, recommended as a replacement for International Human Microbiome Standards (IHMS) SOP 006 as QIAGEN no longer makes QIAamp DNA stool kit commercially available. Similarly to MGP_SOP 001, this protocol requires the QIASymphony DSP Virus/Pathogen Midi Kit and the QIASymphony SP instrument from the same supplier.

This protocol is of first interest for automated and quick saliva samples DNA extraction in order to characterize the oral microbiota by metagenomic profiling.

2. PRINCIPLE

The step-by-step description of this protocol will allow to generate lysates from saliva samples using a triple lysis: thermal, chemical and mechanical. The DNA from these lysates will then be purified using magnetic beads and a custom QIASymphony SP protocol developed by QIAGEN and MGP.

3. PERSONS ENTITLED TO USE THE PROCEDURE

This protocol applies to any person involved in saliva samples DNA extraction. This person can be a trainee, fellow, technician or the engineer in charge of saliva samples DNA extraction.

4. PRELIMINARY STEPS, SPECIFICITIES

Protocol should be approved by an ethics committee according to national regulations. Protocol should be declared on international database (e.g. <u>https://clinicaltrials.gov</u>).

Volunteers and patients should have signed an informed consent according to the approved protocol.

For the preparation of the nucleic acids, aliquots are prepared from appropriately identified and collected samples using DNA Genotek's OMNIGene·ORAL. Following collection and delivery to the laboratory, saliva samples must be heated for 2 hours at 50 °C in an incubator, then aliquoted in 500 μ L subsamples into 2 mL tubes and stored at -80 °C.

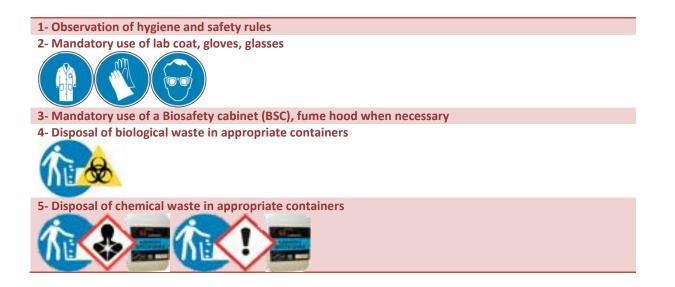
Moreover, it must be kept in mind that the specific area of nucleic acids preparation does see constant evolutions and improvements, such that it is hardly conceivable to definitely "freeze" a protocol that will be considered as optimal in the long term.





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5. CONDITIONS AND USAGE CONSTRAINTS TO FOLLOW



6. MATERIALS, REAGENTS, PRODUCTS, KITS, SOLUTIONS

Materials:

Materials	Risks	Prevention	Providers Names and references
0.1 mm glass beads			152016 DUTSCHER
Screw cap microtubes, 2 mL, steriles			73.693.005 SARSTEDT
Safelock Eppendorf tubes, 2 mL, steriles			033297 DUTSCHER
Toothpicks, steriles			046003 DUTSCHER
1000, 200, 100 and 10 μL micropipettes			Eppendorf Research
1000 μL filter tips			S1122-1830 STARLAB
200-100 μL filter tips			S1120-8810 STARLAB
100 μL filter tips			S1121-3810 STARLAB
10 μL filter tips			S1121-3810 STARLAB
Sample Prep Cartridges 8-well			997002 QIAGEN





INRAE METAGENOPOLIS	SALIVA SAMPLES D EXTRACTION	NA	Version: 1 Date: 2023-03-08 Page n°4 of 9	Last Contributor: Christian Morabito
8-Rod C	Covers			997004
Filter Time 1500 vil	Onum (D (1024)			QIAGEN
Filter-Tips, 1500 μl	l, Qsym SP (1024)			997024 QIAGEN
Filter-Tips, 200 µl,	. Osym SP (1024)			990332
· · · · · · · · · · · · · · · · · · ·	, ()			QIAGEN
Mixer mill (Ret	tsch MM400)	A A	Beware of	20750001
	L	nd crush hazard	balancing	RETSCH
Benchtop mic		nd crush hazard		DUTSCHER
Benchtop refrige	rated centrifuge	^	Beware of	5409000217
		4	balancing	Eppendorf
70 °C dr	y bath	Burn hazard		060917 DUTSCHER
Vortex	mixer	A		250158 DUTSCHER
Biosafety	cabinet	A		1046-2614 FISHER SCIENTIFIC

<u>Kits:</u>

Materials	Risks	Prevention	Providers Names and references
QIAsymphony DSP Virus/Pathogen Midi Kit, version 1			937055 QIAGEN

Products/Reactives:

Materials	Risks	Providers Names and references





INRAE METAGENOPOLIS	SALIVA SAMPLES DNA EXTRACTION	Version: 1 Date: 2023-03-08 Page n°5 of 9	Last Contributor: Christian Morabito
Guanidine T	hiocyanate	G663 SIGN	
Tris-HCl 1	M pH 8	15893 Thermo so	
N-Lauroyl Sarcosine 20 %		L7414-5 SIGN	
N-Lauroyl S	Sarcosine	L9150- SIGM	•
Na	Cl	S304 SIGN	
EDTA 0.5	М рН 8	E7889-1 SIGN	
PBS 1X	pH 7.4	15374 Thermo so	
PVF	P	7762 SIGN	27

7. STEP-BY-STEP PROCEDURE

- 1. Add 250 µL de Guanidine Thiocyanate 4 M Tris-HCl 0,1 M to each frozen sample.
- 2. Add 40 μL N-lauroyl sarcosine 10 % and thaw.
- 3. Add 500 μL N-lauroyl sarcosine 5 % PBS 1X and vortex to mix well.
- 4. Incubate at 70 °C in a dry bath for 1 hour.
- 5. Add 750 mg glass beads (0.1 mm) in each tube and vortex vigorously.
- 6. Shake for mechanical disruption:
 - a. with Bead Beater™:
 - i. Turned on (medium speed) for 5 min
 - ii. Stopped for 10 min
 - iii. Turned on again (medium speed) for 5 min
 - b. with MixerMill MM400 (Retsch): run Program 1, 10 min at 25 s⁻¹ (Hz)
- 7. Add 15 mg PVPP (powder) per sample and vortex vigorously.

<u>Possible variation</u>: steps 4, 6 and 7 can be replaced by 1 hour, 70 °C incubation at 1400 rpm in a thermoshaker. In this case, it should be performed after step 5.





INRAE SALIVA SAMPLES DNA METAGENOPOLIS EXTRACTION	Version: 1 Date: 2023-03-08 Page n°6 of 9	Last Contributor: Christian Morabito	
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- 8. Centrifuge at 18,200 ×g for 5 min, 4 °C.
- 9. Recover the supernatant in a sterile 2 mL tube and set aside.
- 10. Add 500 μ L TENP (resuspend before use) to the pellet and vortex vigorously. Use a toothpick if necessary.
- 11. Centrifuge at 18,200 ×g for 5 min, 4 °C.
- 12. Recover the supernatant and pool with the first.
- 13. Pipet 800 μ L lysate in a 2 mL screw cap Sarstedt tube. The remaining lysate can be stored at 80 °C for up to at least 3 months.
- 14. Turn on, prepare, and run the QIASymphony according to the manufacturer's instructions and use the protocol **COBL1200_CR23506_ID4502**. Elution will be performed in 110 μL buffer, in 0.5 mL Matrix tubes contained in a 96-well SBS rack (*« Deep well »* then *« TS#3744 2D storage tubes »*).

The QIAGEN client can install the protocol used on demand. One cartridge is enough to extract 110 samples. Users will need to plan a training with QIAGEN to learn how to use the QIASymphony.

15. Measure DNA concentration with a fluorometer (e.g. Qubit) and check for DNA quality with a 1 % Agarose gel or Fragment Analyzer (Agilent).





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Annex. Preparation of solutions

TEN (50 mM Tris-HCl, 20 mM EDTA, 10 mM NaCl) / TENP (TEN + 1 % PVPP)

- 150 mL Tris-HCl 1 M pH 8
- 100 mL EDTA 0.5 M pH 8
- 14.6 g NaCl
- H₂O q.s. 2.5 L

Mix until totally dissolved. Filter through 0.2 microns Millipore filter and store at 4 °C protected from light. For TENP, add 0.1 g PVPP powder to 10 mL TEN and mix (it will not dissolve).

N-Lauroyl Sarcosine 10 %

- 50 mL N-Lauroyl Sarcosine 20 %
- H₂O q.s. 100 mL

N-Lauroyl Sarcosine 5 % PBS 1X

- 50 g N-Lauroyl Sarcosine
- 1 L PBS 1X ph 7.4

Guanidine Thiocyanate 4 M Tris-HCl 0,1 M

- 236.32 g Guanidine Thiocyanate
- 50 mL Tris-HCl 1 M pH 8
- H₂O q.s. 500 mL

Mix until totally dissolved, protected from light by aluminum foil. Filter through 0.2 microns Millipore filter and store at 4 °C protected from light.