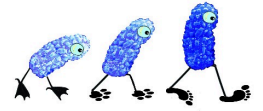


Jan 06, 2020

## Isolation of Klebsiella strains from food samples

DOI

[dx.doi.org/10.17504/protocols.io.baxtifnn](https://dx.doi.org/10.17504/protocols.io.baxtifnn)



MedVetKlebs consortium<sup>1</sup>

<sup>1</sup>Institut Pasteur

Klebsiella Research and ...



Carla Rodrigues

Institut Pasteur

OPEN  ACCESS



DOI: [dx.doi.org/10.17504/protocols.io.baxtifnn](https://dx.doi.org/10.17504/protocols.io.baxtifnn)

External link: <https://doi.org/10.1128/spectrum.02376-21>

**Protocol Citation:** MedVetKlebs consortium 2020. Isolation of Klebsiella strains from food samples. **protocols.io**  
<https://dx.doi.org/10.17504/protocols.io.baxtifnn>

### Manuscript citation:

Rodrigues C, Hauser K, Cahill N, Ligowska-Marzeta M, Centorotola G, Cornacchia A, Fierro RG, Haenni M, Nielsen EM, Piveteau P, Barbier E, Morris D, Pomilio F, Brisse S, High Prevalence of in European Food Products: a Multicentric Study Comparing Culture and Molecular Detection Methods. Microbiology Spectrum 10(1). doi: [10.1128/spectrum.02376-21](https://doi.org/10.1128/spectrum.02376-21)

**License:** This is an open access protocol distributed under the terms of the **[Creative Commons Attribution License](#)**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

**Protocol status:** Working

**We use this protocol and it's working**

**Created:** January 03, 2020

**Last Modified:** January 06, 2020

**Protocol Integer ID:** 31443



**Keywords:** isolation; *Klebsiella* spp.; *Klebsiella pneumoniae*; food samples

## Abstract

This protocol is intended for isolation of *Klebsiella* strains from different food sources. It is derived from the initial description of the SCAi medium (van Kregten E, Westerdaal, N. A. C., and Willers, J. M. N. New, simple medium for selective recovery of *Klebsiella pneumoniae* and *Klebsiella oxytoca* from human feces. Journal of Clinical Microbiology. 1984;20:936-41) and its validation across a diversity of *Klebsiella* strains (Passet V, Brisse S. 2015. Association of tellurite resistance with hypervirulent clonal groups of *Klebsiella pneumoniae*. J Clin Microbiol. 53(4):1380-2).

The protocol entails enrichment using Buffered Peptone Water (BPW), and plating on SCAi (Simmons Citrate with Inositol) agar.

This protocol was optimized by the MedVetKlebs consortium using chicken meat and salad samples.

## Guidelines

There is no commercial availability of SCAi agar plates (in 2019). Plates must be prepared locally and can be stored several weeks at 4°C.



# Materials

## MATERIALS

✕ myo-inositol Merck MilliporeSigma (Sigma-Aldrich) Catalog #I5125-50 g

✕ Simmons citrate Bio-Rad Laboratories Catalog #64834-500g

### 1. Stomacher or blender

### 2. Myo-inositol solution preparation

Catalog: Sigma-Aldrich I5125-50 g

Preparation of myo-inositol at 10 %

Steps:

Weigh 10 g of myo-inositol and dissolve in 100 ml of water. Sterilize by filtration.

### 3. Simmons Citrate Agar

Catalog: Bio-rad 64834-500g or Dutscher 777388-500g

For Simmons citrate agar from BioRad:

Steps:

Suspend 21 g of the powder in 1 liter of purified water. Mix thoroughly. Heat with frequent agitation and boil for 1 minute to completely dissolve the powder. Autoclave at 121°C for 20 minutes. Cool to 45-55 °C and take 900 ml Simmons Citrate Agar. Add 100 ml myo-inositol at 10 % (leading to a final concentration of 1%). Distribute into sterile Petri plates and store at 4 °C.

Note: Simmons citrate can also be ordered at Conda (ref. Simmons Citrate agar ISO 10273, catalog Number 1014). In that case 24.3 g of powder must be used for one liter, instead of 21 g.

## Before start

Prepare Buffered Peptone Water (BPW) solution and SCAi agar plates.

Pre-warm the BPW at room temperature before use.

## Pre-treatment of sample

- 1 Cut sample into small slivers to a final weight of 25g.
- 2 Dilute the 25g portion in 225ml of **Buffered Peptone Water (BPW) broth** (1:10 dilution).
- 3 Mix the sample using a stomacher for 30 seconds or pulse using a blender.
- 4 Incubate the suspension at **37°C ± 1°C** for **24 h ± 1 h**.

## Streak a SCAi medium agar plate

- 5 Following 24h incubation, using a **10µl loop**, streak for single colonies onto the surface of a small petri dish (90 mm) of SCAi medium and incubate at **44°C ± 1°C** for **48h ± 1 h**.

Sometimes, typical colonies (yellow, moist, dome-shaped) can be recognized after 24h culture on plates, but 48 h is much better to discriminate *Klebsiella*-looking colonies from other ones (*E. coli* colonies are typically white because they do not use inositol).

It can happen that the medium, which should initially be blue, turns completely yellow, when there are many inositol-fermenting colonies (typically *Klebsiella*). In these cases, discriminating yellow colonies is less easy. Diluting before streaking could help in these cases.

## Purification and identification of suspect *Klebsiella* colonies

- 6 Typical *Klebsiella* spp. colonies are yellow on SCAi medium.

Select suspect *Klebsiella* colonies for subculture and bacterial identification.

Streak the selected colonies onto the surface of a non-selective agar medium (e.g., LB or TSA) in a manner which will allow isolated colonies to develop. Incubate plates at 37 °C ± 1°C for 24 h ± 1 h.

Note: If colonies are numerous and close to each other, re-isolate the colony on another SCAi agar plate to control for purity. Incubate for up to 48h.



- 7 Determine species of purified suspect *K. pneumoniae* colonies using MALDI-TOF mass spectrometry and/or species-specific PCR.

### Mixed-colony storage for future studies (additional)

- 8 If desired, after picking selected colonies, sweep the remaining SCAi plate content and freeze it at -80°C (e.g., for mixed colonies sequencing) using CryoBank tubes or equivalent (e.g., in house BHI + 15% glycerol medium).

### Storage of bacterial strains

- 9 Freeze strains confirmed as *Klebsiella pneumoniae* (or its related species, which also grow on SCAi) at -80°C using CryoBank tubes or equivalent (e.g. BHI + 15% glycerol medium).

If several morphotypes are available, you may want to store one colony per morphotype.