Description

The present invention relates to a DNA probe for identifying bacterial strains of the Lactobacillus helveticus species, a process for producing such a probe and a method for identifying bacterial strains of this species with this DNA probe.

Lactobacillus helveticus is a very important bacteria for the fermentation of food. It is predominantly used in the fermentation of milk products and is found in starter cultures for some cheese and, in some countries. also for yoghurt production. Fermentation and maturation of these food products usually result from growth association and interaction of different bacteria, in most cases different Lactobacilli, Lactococci and other bacterial species. As most of these bacterial species have very similar nutritional requirements and grow under similar environmental conditions a clear identification within the Lactobacillus species is sometimes very difficult. In particular, it may be difficult to differentiate between the two yoghurt bacteria L. bulgaricus, which belongs to the species of L. delbrueckii, and L. helveticus (formerly also called L. jugurti; O.Kandler and N.Weiss, Bergey's Manual of Systematic Bacteriology, vol. 2, 1208-1260, 1986). So far the classification of all these species is tedious and involves unreliable criteria like sugar fermentation patterns and acid production. Due to these tests an identification of these different species remains doubtful and often arbitrary.

DNA hybridisation techniques, using specific DNA probes, are a very valuable tool for the identification of bacterial and viral strains and have already found application in clinical diagnostics. Such DNA probes have already been used for the identification of Yersinia enterocolitica (J.Jagow et al., Appl.Environ.Microbiol. 51, 441-443, 1986), Plasmodium falciparum (R.H.Barker et al., Science 231, 1434-1436, 1986), Salmonella typhi (F.A.Ruben et al., J.Clin.Microbiol. 22, 600-605, 1985), Bacillus subtilis - (J.Krauss et al., FEMS Microbiol.Lett. 33, 89-93, 1986), Haemophilus influenzae (F.Malouin et al., J.Clin.Microbiol. 26, 2132 -2138, 1988) and other bacterial species, of DNA viruses (J.Brandsma et al., Proc.Natl.Acad.Sci.USA, 77, 6851 -6855; P.Stolhandske et al., J.Clin.Microbiol. 15, 744 -747, 1982; P.Stolhandske et al., J.Med.Virol. 12, 213 -218, 1985), as well as RNA viruses (J.Flores et al., Lanceti, 555-559, 1983; M.Lin et al. , J.Virol. 55, 509 -512, 1985).

Antonie van Leeuwenhoek, vol. 49, 1983, pages 143-152 (P.L.Manachini et al.) disclose a comparative study of physiological characteristics, DNA base composition, DNA similarity and DNA restriction endonuclease, e.g. BamH1, EcoR1 and HindIII cleavage patterns from two strains of L.jugurti and four strains of L.helveticus and show that all these strains are closely related. As disclosed in this article the DNA-DNA hybridisations trials as well as the DNA cleavage patterns involve the whole DNA content of the strains.

In the species of Lactobacillus, DNA probes for L.curvatus (H.A.R. Petrick et al., Appl.Environ.Microbiol. 54, 405 -408, 1988) which is specifically associated with spoilage of vacuum-packed meat and for the L.delbrueckii species comprising L.bulgaricus, L.lactis and L.delbrueckii (M.Delley et al.) have been isolated. It certainly would be of use in the dairy industry not only to be able to quickly identify L.delbrueckii , as can be done by the procedure described in our European Patent Application No. 89106016.2, but also to have a quick and reliable method to detect and classify strains belonging to the L.helveticus species.

Accordingly, a first object of the present invention is to provide a specific DNA probe, which can be used in hybridisation procedures to specifically identify strains belonging to the Lactobacillus helveticus species either in bacterial cultures, or as food constituents or during food processing, e.g. during industrial fermentation.

A second object of the present invention is to provide a process for producing such a specific DNA probe.

A third object of the present invention is to provide a method for specifically identifying strains belonging to the L.helveticus species with this DNA probe.

To this end, a DNA probe for identifying bacterial strains of the Lactobacillus helveticus species according to the present invention is obtained by a process which comprises preparing a HindIII clonebank from the strain L.helveticus ATCC 15009 by digestion of DNA of this strain with HindIII, ligation into the plasmid pACYC184, and transformation of the ligation mixture into the E.coli strain HB101, and isolating therefrom a clone of which a HindIII DNA fragment is capable of specifically hybridising to DNA of strains of the L.helveticus species, this DNA fragment being the fragment sLH1 or the fragment sLH2, as defined in figures 1 and 2.

The method for identifying a bacterial strain of the L.helveticus species according to the present invention comprises preparing DNA of a strain to be identified and checking whether this DNA hybridises to the present probe produced by the present process. In a preferred embodiment of this method, the DNA is prepared by growing cells of the strain to be identified on a culture medium supplemented with a fermentable carbon source, incubating them in the presence of proteinases, treating them with an N-acetyl-

muramidase, further incubating them in the presence of an emulsifying agent, a chelating agent and a proteinase, phenol extracting DNA therefrom, ethanol precipitating the extracted DNA, treating this DNA with an RNase and chloroform extracting the RNase treated DNA.

BRIEF DESCRIPTION OF THE DRAWING

Fig. 1 is a restriction map of a cloned HindIII DNA fragment, sLH1, originating from L.helveticus ATCC 15009 and being about 1845 base pair long.

Enzymes which do not cut within this fragment are:

ApaI, AvaI, BamHI, BglI, BglII, BstBI, CfoI, EcoRI, Hae II, HindIII, KpnI, MluI, NarI, NcoI, NdeI, NheI, NotI, NruI, NsiI, PstI, PvuI, SacI, SacII, SalI, SmaI, SnaBI, SphI, SspI and StuI.

Fig. 2 is a restriction map of another cloned HindIII DNA fragment, sLH2, originating from L.helveticus ATCC 15009 and being about 1875 base pair long.

Enzymes which do not cut this fragment are:

AccI, AflII, AvaI, BalI, BglI, BglII, DraI, EcoRV, KpnI, NarI, NcoI, NheI, NotI, NruI, NsiI, PstI, PvuI, SacI, SacII, SalI, ScaI, SmaI, SphI, SspI and StuI.

In these two Fig., the numbers in brackets indicate the relative position of the appropriate restriction sites in base pairs.

DETAILED DESCRIPTION OF THE INVENTION

In order to obtain DNA probes which specifically identify strains belonging to the species of L.helveticus, it was possible to randomly clone HindIII DNA fragments originating from L.helveticus strains and to screen them for species specific hybridisation.

The preferred vector for transforming a clone bank of HindIII DNA fragments of L.helveticus was the plasmid pACYC184 which is capable of being transformed into an E.coli strain, e.g. in the preferred E.coli strain HB101.

Especially specific probes were made of HindIII DNA fragments originating from strain L.helveticus ATCC 15009, e.g. from two DNA fragments sLH1 and sLH2.

Beside the above mentioned preferred way of preparing DNA of a strain to be identified, it was also possible to prepare this DNA by simply lysing cells of this strain, either from cultures or on solid supports such as e.g. filter paper or nitrocellulose paper, and to sucessfully carry out identification tests on this DNA with the probes of the present invention.

In screening experiments applying dotblot hybridisation, over 50 different Lactobacillus strains from different origins were tested. It could be shown that probes according to the present invention only hybridised specifically to the L.helveticus species and that they had a very high sensitivity.

MATERIALS AND METHODS

Bacteria and Plasmid

Lactic acid bacteria used in the Example hereafter are shown on table I. The E.coli strain is HB101 (leuB6 proA2 recA13 thi1 ara14 lacY1 galK2 xyl5 mtl1 rpsL20 supE44 hsdS20). The plasmid used as vector is pACYC184 (A.C.Y. Chang and S.N.Cohen, J.Bacteriol. 134: 1141 -1156, 1978).

Media

For growth of the different Lactobacilli, Lactococci and Propionibacteria MRS broth (Difco Laboratories) was used, supplemented with 1% lactose. E.coli strain was grown in LB medium.

Preparation of DNA

i) DNA from Lactobacillus, Lactococcus and Propionibacteria

Cells were diluted from overnight cultures into 10 ml MRS, supplemented with 1% lactose and grown to mid-log phase at 43 °C. The cells were then harvested by centrifugation, washed once in cold 1M NaCl, and incubated for 1 h at 37 °C in the presence of Proteinase K (250 µg/ml) and Pronase E (500 µg/ml). The cells were washed in TE (10 mM Tris hydrochloride pH 7.4; 1 mM EDTA) and treated with Mutanolysin (200 µg/ml) in the presence of TE for 1h at 37 °C SDS. EDTA and Proteinase K were added

to a final concentration of 0.1%, 75 mM and 200 µg/ml, resp., and incubated for 4 h at 65 °C. The DNA was then phenol extracted, ethanol precipitated and spooled onto a sterile toothpick. The DNA was resuspended in TE in the presence of RNase A (200 µg/ml), chloroform extracted, reprecipitated in ethanol and spooled again onto a toothpick. The DNA was then resuspended in 100 µl of TE and stored at 4 °C.

ii) Plasmid DNA from E.coli

Plasmid DNA from E.coli was isolated and as needed purified on CsC1 gradients according to Maniatis et al. (Molecular cloning: a laboratory manual. Cold Spring Harbor Laboratory, Cold Spring Harbor, New York, 1982). Chemicals were purchased from E.Merck Chemicals Inc. and the enzymes from Sigma Chemical Co..

Preparing a HindIII clone bank from a strain of L.helveticus

DNA of L.helveticus ATCC 15009 was digested to completion with HindIII and ligated into vector pACYC184 which was linearised at its unique HindIII site and phosphatase treated. The ligation mixture was transformed into HB101, plated onto LB plates supplemented with 30 µg/ml chloramphenicol and grown at 37 °C overnight. Single colonies were grown in LB supplemented with chloramphenicol, their plasmids extracted, digested with HindIII and analysed by agarose gel electrophoresis. 90% of the analysed clones contained L.helveticus HindIII DNA fragments.

Labelling of DNA fragments

To radioactively label the DNA we used a fill-in-replacement DNA synthesis or nick-translation reaction (Maniatis et al.). The Klenow fragment of DNA polymerase I and E.coli DNA polymerase I were purchased from Boehringer-Mannheim, [α-^{32}P] dATP from Amersham Co.

Dotblot Hybridisation Procedures

Aliquots of 200 ng of chromosomal DNA in TE were denatured by heating for 5 min. at 95 °C. SSC was added to the aliquots to give a final concentration of 16X SSC and then the mixture was spotted onto 10XSSC wetted GeneScreen® paper and rinsed once with 20XSSC. A Bio-Rad dotblot apparatus was used. The filter was then ready for DNA hybridisation, applying standard procedures with hybridisation with 6XSSC at 65 °C and a subsequent wash with 0.1XSSC at 65 °C (E.Southern, J.Mol. Biol. 98, 503 -517, 1975). As DNA probes we used either total plasmids or HindIII fragment DNA which was ^{32}P labelled. To detect the hybridisation signals, the filters were used to expose X-ray films.

Restriction Mapping

Restriction enzymes were purchased from Boehringer-Mannheim Co. and New England Biolabs Co. and were used as recommended by the suppliers.

EXAMPLE

Isolation of L.helveticus DNA probes

HindIII digested chromosomal DNA of L.helveticus ATCC 15009 has been used to prepare a clone bank as described above. Several clones were isolated and used as DNA probes in dotblot hybridisation experiments using filters containing DNA samples of a selected number of different lactic acid bacteria. The hybridisation results revealed that about 70% of the tested L.helveticus clones hybridised only to the different L.helveticus samples on the dotblot filters. The other 30% of the clones tested hybridised also to a variety of other unrelated species and genera of the lactic acid bacteria. For the following Examples, two DNA probes were selected which were very specific for the L.helveticus species. They are referred to as "sLH1" and "sLH2" in the present specification and claims.

Restriction mapping of sLH1 and sLH2

A restriction map of the DNA fragments sLH1 and sLH2 was determined using several restriction enzymes. The results are shown in Fig. 1 and 2 (see brief description of the drawing here above).

Specificity and sensitivity of the DNA probes sLH1 and sLH2

The two DNA probes sLH1 and sLH2 were tested against many different representatives of the Lactobacillus genus and other lactic acid bacteria with dotblot hybridisations. For these hybridisation experiments, only the HindIII DNA fragments originating from L.helveticus, isolated from the plasmids, were used as DNA probes. The results of the dotblots showed that both DNA probes only hybridise to DNA derived from the bacterial species of L.helveticus. All other strains tested of different Lactobacillus species, of Lactococcus and Propionibacteria did not show any hybridisation . The only exceptions we observed, were two strains of L.acidophilus, N21 and N22, which gave a weak hybridisation signal with sLH1 and sLH2. However, these weak signals can clearly be differentiated from the much stronger hybridisation signals of strains belonging to the species of L.helveticus . The complete results are summarised in table I.

In order to test the sensitivity of our dotblots we made filters of serial dilutions of DNA from different Lactobacillus strains and used them for hybridisation with the probes sLH1 and sLH2. The results can be summarised as follows:

i We easily could detect positive hybridisation signals with all the different strains of L.helveticus at DNA quantities as low as 2 ng per spot.

ii All other bacterial strains tested as listed in table I do not show hybridisation signals at DNA quantities of 200 ng per spot.

iii At DNA amount between 200 ng and 1.6 µg per spot many strains of the L.acidophilus species started to give rise to hybridisation signals.

iv The following strains have been tested at DNA quantities of 1.6 µg per spot and did not show any hybridisation signals: L.lactis N4, L.fermentum N7, L.plantarum N24, L.buchneri N25, L.brevis N26, L.casei N27, L.bulgaricus N123, L.maltaromicus N207, L.reuteri N211 and E.coli HB101.

Table I: Bacterial strains

Species	Source	Strain[a)]	sLH1[b)]	sLH2[b)]
L. helveticus	ATCC 15009	N2	+	+
L. lactis (type)	ATCC 12315	N4	-	-
L. helveticus	NCDO 87	N6	+	+
L. fermentum	NCDO 1750	N7	-	-
L. delbrueckii (type)	NCIB 8130	N8	-	-
L. lactis	Liebefeld 125	N9	-	-
L. acidophilus	ATCC 4356	N12	-	-
L. acidophilus	Piacenza D 179	N16	-	-
L. acidophilus	Piacenza 188 S	N17	-	-
L. acidophilus	Piacenza 247 S	N18	-	-
L. acidophilus	Piacenza T 145	N21	+/-	+/-
L. acidophilus	Piacenza T 145, 125	N22	+/-	+/-
L. plantarum	ATCC 8041	N24	-	-
L. buchneri	ATCC 4005	N25	-	-
L. brevis	ATCC 14869	N26	-	-
L. casei (type)	ATCC 393	N27	-	-
L. helveticus	ATCC 7994	N47	+	+
L. helveticus	ATCC 7995	N48	+	+
L. helveticus	ATCC 12278	N50	+	+
L. helveticus	ATCC 13866	N51	+	+
L. bulgaricus	NCDO 1006	N52	-	-
L. bulgaricus	NCDO 1373	N54	-	-
L. lactis	NCDO 270	N62	-	-
L. lactis	NCDO 297	N64	-	-
L. bulgaricus	NCDO B 15	N95	-	-
L. bulgaricus	NCDO B 19	N96	-	-
L. bulgaricus	NCDO B 24	N99	-	-
L. helveticus	NCDO 2395	N106	+	+
L. helveticus	ATCC 11977	N122	+	+
L. bulgaricus (type)	NCDO 1489	N123	-	-
L. bulgaricus	ATCC 21815	N124	-	-
L. helveticus	ATCC 27558	N125	+	+
L. helveticus	ATCC 33409	N126	+	+
L. bulgaricus	Piacenza CO 13	N139	-	-
L. bulgaricus	Piacenza CO 14	N141	-	-

Species	Source	Strain[a)]	sLH1[b)]	sLH2[b)]
L. plantarum	ATCC 14917	N186	-	-
L. delbrueckii	ATCC 9649	N187	-	-
L. maltaromicus	ATCC 27865	N207	-	-
L. reuteri (type)	DSM 20016	N211	-	-
L. helveticus	Piacenza b 50	N213	+	+
L. helveticus	Zürich S 400	N217	+	+
L. helveticus	our collection	N222	+	+
L. helveticus	NCDO H 1	N254	+	+
L. helveticus	NCDO H 2	N255	+	+
L. helveticus	NCDO H 3	N256	+	+
L. helveticus	NCDO H 6	N257	+	+
L. helveticus	NCDO H 9	N258	+	+
L. helveticus	NCDO H 10	N259	+	+
L. helveticus	NCDO H 11	N260	+	+
L. helveticus	NCDO H 13	N261	+	+
L. helveticus	NCDO H 16	N262	+	+
L. helveticus	NCDO H 17	N263	+	+
L. helveticus	NCDO H 18	N264	+	+
L. helveticus	NCDO H 19	N265	+	+
L. helveticus	NCDO J 3	N266	+	+
L. helveticus	NCDO J 4	N267	+	+
L. helveticus	NCDO J 5	N268	+	+
L. helveticus	NCDO J 6	N269	+	+
L. helveticus	NCDO J 11	N270	+	+
L. helveticus	NCDO J 14	N271	+	+
L. helveticus	NCDO J 15	N272	+	+
L. helveticus	NCDO J 17	N273	+	+
L. helveticus	NCDO J 18	N274	+	+
L. helveticus	NCDO J 19	N275	+	+
propionibacterium freudenreichii	our collection	PP21	-	-
propionibacterium shermanii	our collection	PP13	-	-
Lactococcus therm.	our collection	ST1	-	-
Lactococcus lactis	our collection	SL9	-	-
E. coli	our collection	HB101	-	-

a) Code of the Nestlé Research Centre

b) Probe used for dotblot hybridisation:
(+) indicates hybridisation; (-) no hybridisation

Claims

1. A DNA probe for identifying bacterial strains of the Lactobacillus helveticus species, obtained by a process which comprises preparing a HindIII clonebank from the strain L.helveticus ATCC 15009 by digestion of DNA of this strain with HindIII, ligation into the plasmid pACYC184, and transformation of the ligation mixture into the E.coli strain HB101, and isolating therefrom a clone of which a HindIII DNA fragment is capable of specifically hybridising to DNA of strains of the L.helveticus species, this DNA fragment being the fragment sLH1 or the fragment SLH2, as defined in figures 1 and 2.

2. A method for identifying a bacterial strain of the L.heveticus species, which comprises preparing DNA of a strain to be identified and checking whether this DNA hybridises to a probe according to claim 1.

3. A method according to claim 2, which comprises preparing said DNA by lysing cells of said strain to be identified.

4. A method according to claim 3, which comprises preparing said DNA by lysing cells of said strain to be identified on a solid support.

Species	Source	Strain[a)]	sLH1[b)]	sLH2[b)]
L. plantarum	ATCC 14917	N186	-	-
L. delbrueckii	ATCC 9649	N187	-	-
L. maltaromicus	ATCC 27865	N207	-	-
L. reuteri (type)	DSM 20016	N211	-	-
L. helveticus	Piacenza b 50	N213	+	+
L. helveticus	Zürich S 400	N217	+	+
L. helveticus	our collection	N222	+	+
L. helveticus	NCDO H 1	N254	+	+
L. helveticus	NCDO H 2	N255	+	+
L. helveticus	NCDO H 3	N256	+	+
L. helveticus	NCDO H 6	N257	+	+
L. helveticus	NCDO H 9	N258	+	+
L. helveticus	NCDO H 10	N259	+	+
L. helveticus	NCDO H 11	N260	+	+
L. helveticus	NCDO H 13	N261	+	+
L. helveticus	NCDO H 16	N262	+	+
L. helveticus	NCDO H 17	N263	+	+
L. helveticus	NCDO H 18	N264	+	+
L. helveticus	NCDO H 19	N265	+	+
L. helveticus	NCDO J 3	N266	+	+
L. helveticus	NCDO J 4	N267	+	+
L. helveticus	NCDO J 5	N268	+	+
L. helveticus	NCDO J 6	N269	+	+
L. helveticus	NCDO J 11	N270	+	+
L. helveticus	NCDO J 14	N271	+	+
L. helveticus	NCDO J 15	N272	+	+
L. helveticus	NCDO J 17	N273	+	+
L. helveticus	NCDO J 18	N274	+	+
L. helveticus	NCDO J 19	N275	+	+
propionibacterium freudenreichii	our collection	PP21	-	-
propionibacterium shermanii	our collection	PP13	-	-
Lactococcus therm.	our collection	ST1	-	-
Lactococcus lactis	our collection	SL9	-	-
E. coli	our collection	HB101	-	-

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b) Probe used for dotblot hybridisation:
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Claims

1. A DNA probe for identifying bacterial strains of the Lactobacillus helveticus species, obtained by a process which comprises preparing a HindIII clonebank from the strain L.helveticus ATCC 15009 by digestion of DNA of this strain with HindIII, ligation into the plasmid pACYC184, and transformation of the ligation mixture into the E.coli strain HB101, and isolating therefrom a clone of which a HindIII DNA fragment is capable of specifically hybridising to DNA of strains of the L.helveticus species, this DNA fragment being the fragment sLH1 or the fragment SLH2, as defined in figures 1 and 2.

2. A method for identifying a bacterial strain of the L.heveticus species, which comprises preparing DNA of a strain to be identified and checking whether this DNA hybridises to a probe according to claim 1.

3. A method according to claim 2, which comprises preparing said DNA by lysing cells of said strain to be identified.

4. A method according to claim 3, which comprises preparing said DNA by lysing cells of said strain to be identified on a solid support.

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5. A method according to claim 2, which comprises preparing said DNA by growing cells of said strains to be identified on a culture medium supplemented with a fermentable carbon source, incubating them in the presence of proteinases, treating them with an N-acetyl-muramidase, further incubating them in the presence of an emulsifying agent, a chelating agent and a proteinase, phenol extracting DNA therefrom, treating this DNA with an RNase and chloroform extracting the RNase treated DNA.

Patentansprüche

1. DNA-Sonde zum Identifizieren von Bakterienstämmen der Spezies Lactobacillus helveticus, welche nach einem Verfahren erhalten worden ist, welches das Herstellen einer HindIII-Genbank aus dem Stamm L.helveticus ATCC 15009 durch Verdauen der DNA dieses Stammes mit HindIII, das Ligieren in das Plasmid pACYC184 und das Einführen des Ligationsgemisches in den E.coli-Stamm HB101 sowie das Isolieren eines Klons aus demselben umfaßt,von welchem Klon ein HindIII-DNA-Fragment befähig ist, sich spezifisch an DNA aus Stämmen der Spezies L.helveticus zu hybridisieren, wobei dieses DNA-Fragment das in den Fig. 1 und 2 definierte Fragment sLH1 oder das Fragment sLH2 ist.

2. Verfahren zum Identifizieren eines Bakterienstammes der Spezies L.helveticus, welches das Präparieren von DNA aus dem zu identifizierenden Stamm und die Prüfung umfaßt, ob sich diese DNA an eine Sonde gemäß Anspruch 1 hybridisiert.

3. Verfahren nach Anspruch 2, welche das Präparieren dieser DNA durch Lysieren von Zellen des zu identifizierenden Stammes umfaßt.

4. Verfahren nach Anspruch 3, welches das Präparieren dieser DNA durch Lysieren von Zellen des zu identifizierenden Stammes auf einem festen Träger umfaßt.

5. Verfahren nach Anspruch 2, welches das Präparieren der DNA durch Vermehren von Zellen der zu identifizierenden Stämme auf einem Kulturmedium, welches mit einer fermentierbaren Kohlenstoffquelle ergänzt worden ist, das Inkubieren der Zellen in Gegenwart von Proteinasen, das Behandeln der Zellen mit einer N-Acetyl-muramidase, das weitere Inkubieren der Zellen in Gegenwart eines Emulgators, eines Chelatbildners und einer Proteinase, das Extrahieren der DNA aus den Zellen mit Phenol, das Behandeln dieser DNA mit einer RNAse und das Extrahieren der mit der RNAse behandelten DNA mit Chloroform umfaßt.

Revendications

1. Sonde d'ADN pour l'identification de souches bactériennes de l'espèce Lactobacillus helveticus obtenue par un procédé qui comprend la préparation d'une banque de clones HindIII à partir de la souche L. helveticus ATCC 15009 par une digestion de l'ADN de cette souche avec HindIII, une ligation dans le plasmide pACYC184, une transformation du mélange de ligation dans la souche E. coli HB101, et l'isolement à partir de cette souche d'un clone dont un fragment d'ADN HindIII est capable de s'hybrider spécifiquement à l'ADN de souches de l'espèce L. helveticus, ce fragment d'ADN étant le fragment sLH1 ou le fragment sLH2, ces fragments étant définis sur les figures 1 et 2.

2. Procédé d'identification d'une souche bactérienne de l'espèce L. helveticus, qui comprend la préparation d'ADN d'une souche à identifier et le contrôle de l'hybridation éventuelle de cet ADN à une sonde suivant la revendication 1.

3. Procédé suivant la revendication 2, qui comprend la préparation de l'ADN par lyse de cellules de la souche à identifier.

4. Procédé suivant la revendication 3, qui comprend la préparation de l'ADN par lyse de cellules de la souche à identifier sur un support solide.

5. Procédé suivant la revendication 2, qui comprend la préparation de l'ADN par culture de cellules des souches à identifier sur un milieu de culture supplémenté avec une source de carbone fermentable, la mise en incubation de ces cellules en présence de protéinases, le traitement de ces cellules avec une N-acétyl-muramidase, une incubation supplémentaire de ces cellules en présence d'un agent émulsion-

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nant, d'un agent chélatant et d'une protéinase, une extraction par le phénol de l'ADN de ces cellules, le traitement de cet ADN avec une RNAse et l'extraction par le chloroforme de l'ADN traité à la RNase.