

## Eco-friendly skin preservation and depilation

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## Historical context

- Schlosser et al. (1986) using an undisclosed Lactobacillus culture to preserve hides, based on the fact many lactic acid bacteria were used in the preservation of food.
  - Hides were incubated in this culture,
  - The hides were preserved, but also depilated over time.
  - No further research was carried out to understand the mechanisms.

### Historical context

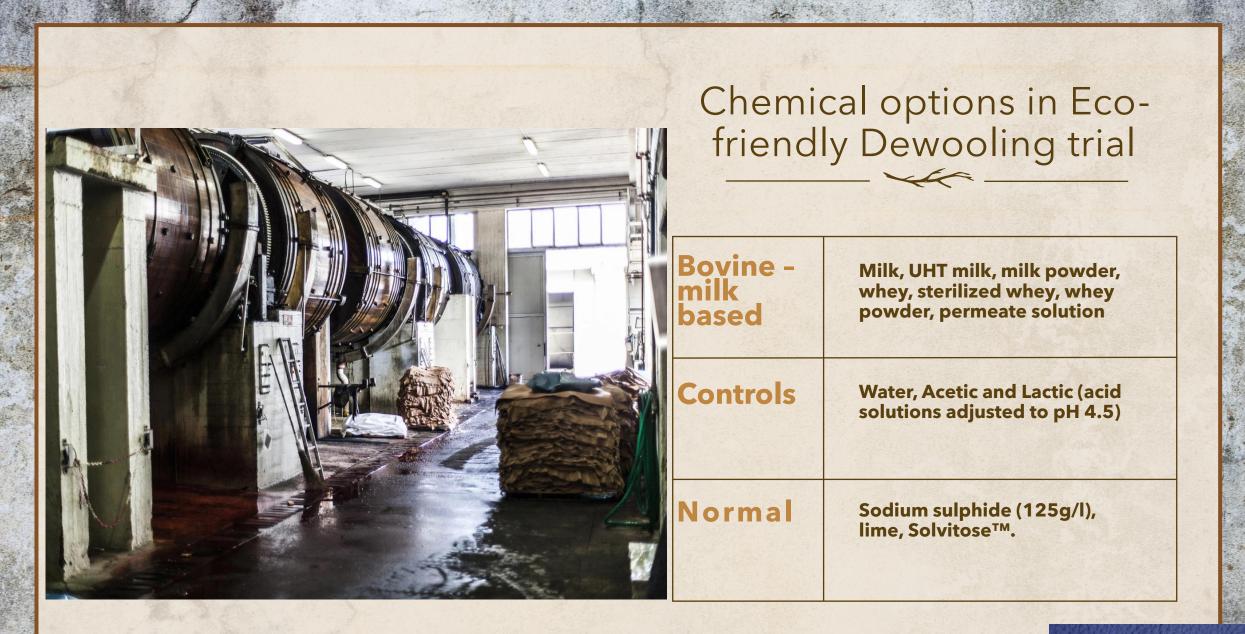
 Dettmer et al. (2013) used a Bacillus enzyme preparation to depilate hides and reduce COD, BOD and the total nitrogen and sulphides in the wastewater. However, the grain was damaged and sulphide was needed to increase the depilatory activity. Others found the same result (De Souza & Guterres, Wahuntari & Hendrawati, Lopéz, et al)

## Eco-friendly process

- Removes wool from sheepskins using whey or whey permeate, omitting the need for sulphide, lime, enzymes.
- Prevents microbial degradation of skins for up to a week at ambient temperatures,

• Potentially avoids the need for salt and biocides for temporary preservation.

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	Initial pH	Final	Deve to	Smell	Depiloted
	of solution	pH	Days to depilate	Smell	Depilated skin condition
Milk	6.8	4.5	3	Fermented/	Pink, shiny,
UHT milk	6.7	4.5	3	sour milk	plump, and
12.5% (w/v) milk powder solution	6.8	4.5	3		very smooth
Whey	6.2	4.5	3		
Sterilised whey	6.0	4.5	3		
Sterilised permeate	6.0	4.5	3		
6.25% (w/v) whey protein solution	6.5	6.0	5	Putrefied and rotten	Grey, easily broken into pieces, and rigid
H <sub>2</sub> O	7.0	7.5	5		-
H <sub>2</sub> O with acetic acid	4.5	7.5	3		
H <sub>2</sub> O with lactic acid	4.5	8.0	3		

Results

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-	CR primer sets for bacteria 16S and fungal 18S ribosomal ribonucleic acid RNA) encoding gene amplification.			
Primer name	Sequence $(5' \rightarrow 3')$	Target gene	Amplicon length (bp)	
27f	AGAGTTTGATCCTGGCTCAG	Full length	1400	
1492r	GGTTACCTTGTTACGACTT	16S rRNA		
Eub338	ACTCCTACGGGAGGCAGGAG	V3 region of	200	
Eub518	ATTACCGCGGCTGCTGCTGG	16S rRNA		
nu-SSU-0817-5'	TTAGCATGGAATAATRRAATAGGA	Nuclear small	420	
nu-SSU-1196-3'	TCTGGACCTGGTGAGTTTCC	subunit of		
		18S rRNA		

#### PCR primers for bacterial and fungal amplification



# Identification of viable microorganisms



TSB, LB, MRS, malt, and Wilson's media were used to culture microorganisms

Microbial genomic DNA (gDNA) was extracted from cells

Sequences were analysed using the BLAST algorithm against the NCBI nucleotide collection



NCBI BLAST identification	Accession	Primer used	% Identity	Culturing media	Depilation media
Bacteroides xylanolyticus	MT192666.1	27f/1492r	98.6%	Malt	Sterilised permeate
	MT459291.1	Eub338/Eub518	100.0%		
Citrobacter europaeus	NR_156052.1	Eub338/Eub518	98.8%	Malt	Whey
Enterobacter sp.	MH477686.1	Eub338/Eub518	99.4%	Wilson's media	Sterilised permeate
Enterococcus faecalis	MT158867.1	Eub338/Eub518	100.0%	Wilson's media	Whey
Empedobacter falsenii	MN198120.1	Eub338/Eub518	98.7%	MRS	Sterilised permeate
Escherichia fergusonii	NR_074902.1	27f/1492r	99.1%	LB	Whey
	MT645516.1		100.0%	MRS	Sterilised permeate
Escherichia coli	CP066366.1	27f/1492r	100.0%	LB	Whey
	MW846276.1	Eub338/Eub518	100.0%	MRS	Sterilised permeate
Hafnia alvei	LR699008.1	27f/1492r	98.9%	TSB	Whey
	KC210872.1		99.1%	Wilson's media	Sterilised permeate
			98.3%		Raw milk
	KX674363.1		99.4%		Milk
Hafnia paralvei	MT470952.1	27f/1492r	99.2%	TSB	Whey
	NR 116898.1		99.7%	MRS	Sterilised permeate
			98.8%	Wilson's media	Milk
	NR 025334.1		99.2%		Milk powder
	MT470952.1	Eub338/Eub518	99.4%		Milk
	MN868256.1		99.3%	Malt	Sterilised permeate
Klebsiella aerogenes	MW784626.1	Eub338/Eub518	99.3%	Wilson's media	Sterilised permeate
Kurthia gibsonii	MN966854.1	27f/1492r	99.5%	Malt	Sterilised whey
	MK898830.1	Eub338/Eub518	100.0%		
Lactobacillus brevis	NR_116238.1	27f/1492r	99.5%	MRS	Whey
	MG722900.1	Eub338/Eub518	99.4%		
		200000/200010	100.0%	Malt	Sterilised permeate
Lactobacillus curvatus	MT645312.1	Eub338/Eub518	99.4%	MRS	Sterilised permeate
Lactobacillus graminis	MN640858.1	Eub338/Eub518	99.3%	MRS	Milk
Lactobacillus plantarum	MF623219.1	27f/1492r	97.9%	MRS	Raw milk
	MK652787.1	Eub338/Eub518	100.0%		
	EU931245.1	200000, 200010	99.4%		Milk
	KT626385.1		100.0%		Sterilised permeate
	A1020000.1		99.4%	Malt	Sterilised whey
Lactococcus lactis	NR_113960.1	27f/1492r	100.0%	TSB	Whey
Laciococcas iacias	MR_113900.1	2/1/14921	99.2%	Malt	Milk
	GQ337875.1		98.5%	Walt	Sterilised whey
	0.000000000		98.2%		Sterilised permeate
	MT545096.1		99.5%	MRS	sternised permeate
	MT597705.1		99.9%	MRB	Raw milk
	M1397703.1		99.9%	Wilson's media	Sterilised whey
	MH666046.1	Eub338/Eub518	100.0%	MRS	Raw milk
	MH000040.1	E00336/E00316	98.6%	MRS	Sterilised permeate
			98.0%	Wilson's media	oternised permeate
Tanta and India and an annual	NID O (COTA A	076/1400-		Wilson's media Wilson's media	
Lactococcus lactis subsp. cremoris	NR_040954.1	27f/1492r	99.3%	Wilson's media Malt	Milk powder
Lactococcus lactis subsp. lactis	MF108188.1	27f/1492r	99.7%		Sterilised permeate
Leuconostoc holzapfelii	NR_042620.1	Eub338/Eub518	98.8%	Malt	Milk
Lysinibacillus macroides	NR_114920.1	27f/1492r	97.9%	MRS	UHT milk
Pseudomonas sp.	KJ496054.1	27f/1492r	100.0%	Malt	Sterilised permeate
	MW844014.1	Eub338/Eub518	100.0%		
Proteus vulgaris	KX867797.1	27f/1492r	98.1%	MRS	Sterilised whey

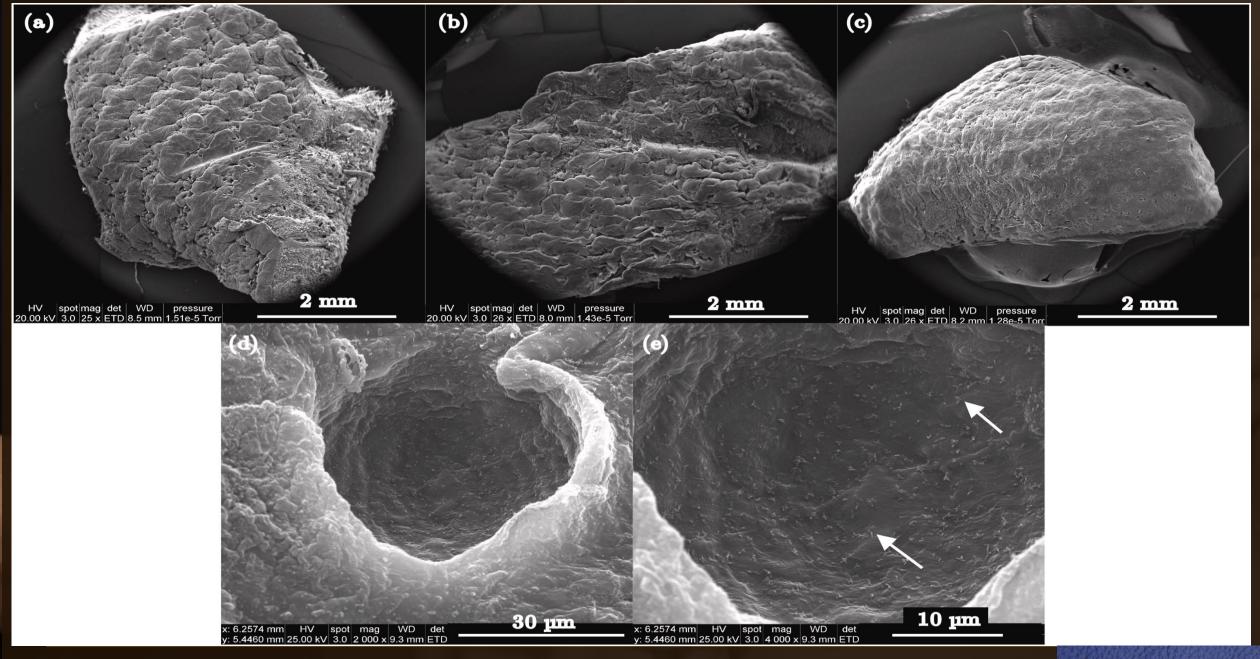


NCBI BLAST identification	NCBI Accession	% Identity	Culturing media	Depilation media
Galactomyces	KY457577.1	99.7	Wilson's media	Milk
candidum		99.5		UHT milk
		99.7	TSB/LB/MRS/Malt/ Wilson's media	Fresh whey
Geotrichum candidum	KY977411.1	99.7	MRS/Malt	Sterilised permeate
		99.7	Malt	Sterilised whey
Pichia insulana	NG_063091.1	99.5	Malt	Sterilised whey
Rhodotorula spp.	MT569975.1	99.5	Malt	UHT milk
Trichosporon lactis	NG_070852.1	99.7	Wilson's media	Milk powder
Yarrowia lipolytica	MH545931.1	99.2	Wilson's media	Sterilised permeate
		99.1	Malt	Sterilised whey
	NG_013120.1	99.7		Milk

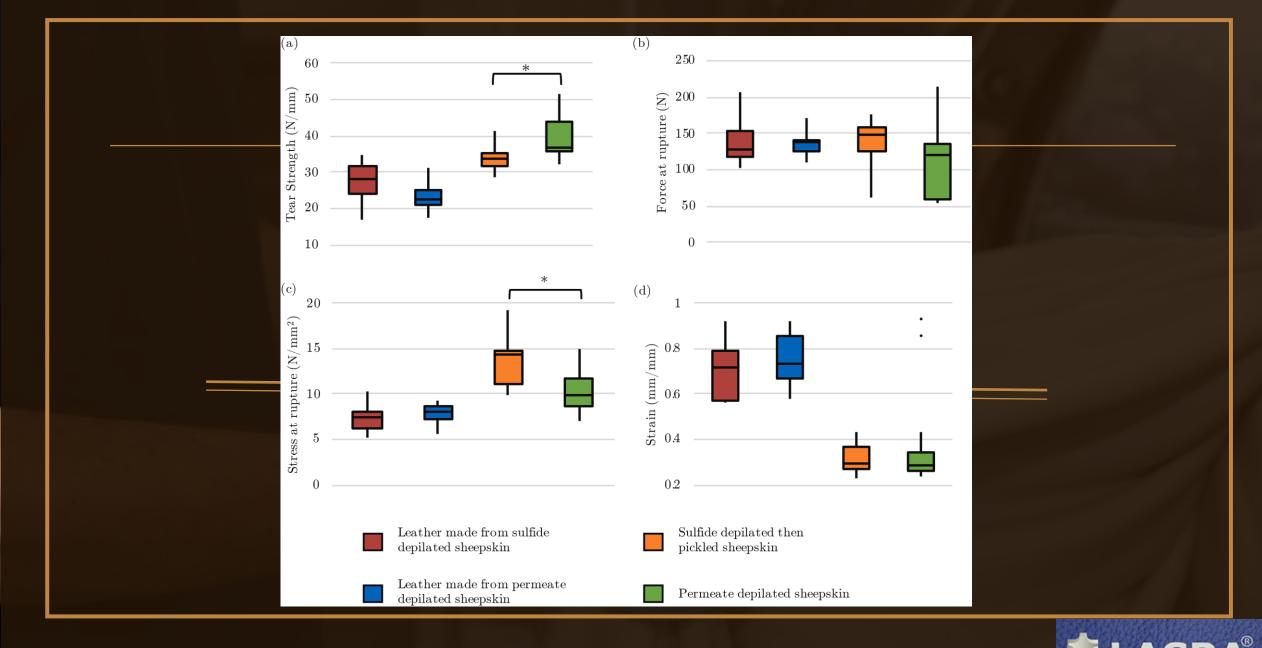
Fungal species identified in the sheepskin depilation trials using various milk products and by-products.











## Eco-friendly Process

Prevention of putrefaction likely due to the high concentrations of lactose at the initial stages of the process.

- Bacterial growth on the unwashed skin and wool controlled by restricting the carbon source at the early stages of depilation in combination with a gradual reduction in pH of the depilation liquid, due to the production of lactic acid.
- Lactobacillus and Lactococcus species are known to produce many antimicrobial substances



## Eco-friendly Process

 Bacteria and fungi are known to secret proteases and glycosidases.

• Those found in this environment are known to secrete enzymes that could preferentially attack the structure around the hair follicle, loosening it to allow easy removal, but without disrupting the structure of the skin.

• Watch this space!

