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Production and Purification of Antimicrobial Bioactive Peptides from Camel and Goat milk by a potential proteolytic probiotic Lactobacillus Strain



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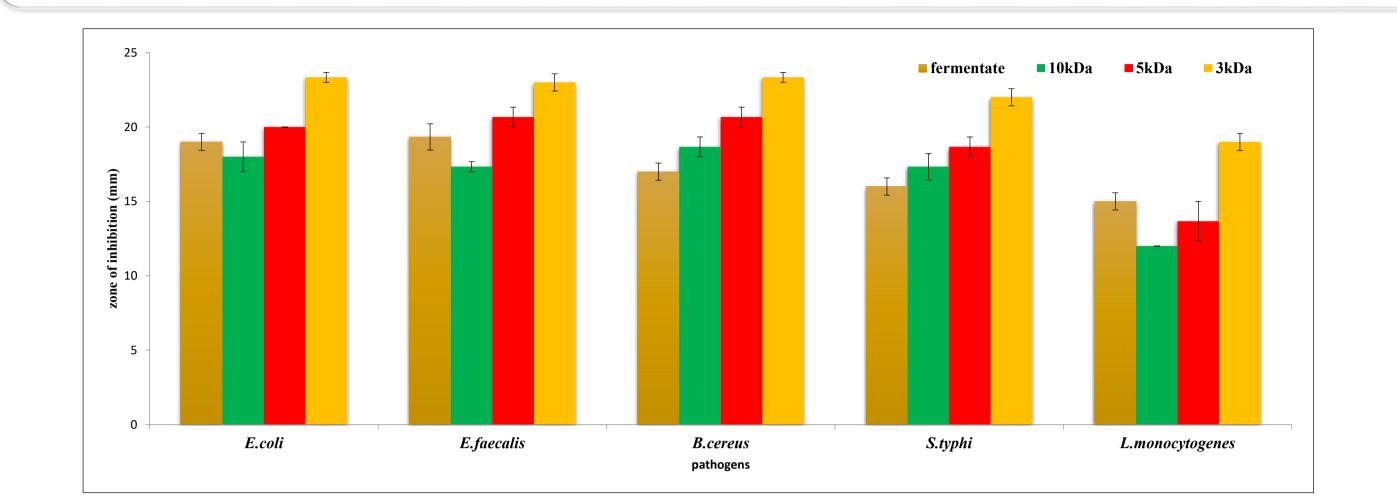
Introduction

Bioactive peptides have been defined as specific protein fragments that are nascent or encrypted in the primary sequences of proteins and have a positive impact on body functions or conditions and may ultimately influence health. They are considered to promote diverse activities, including, antioxidant, immunomodulatory, antihypertensive, antimicrobial, opiate-like, mineral binding and antithrombotic actions.

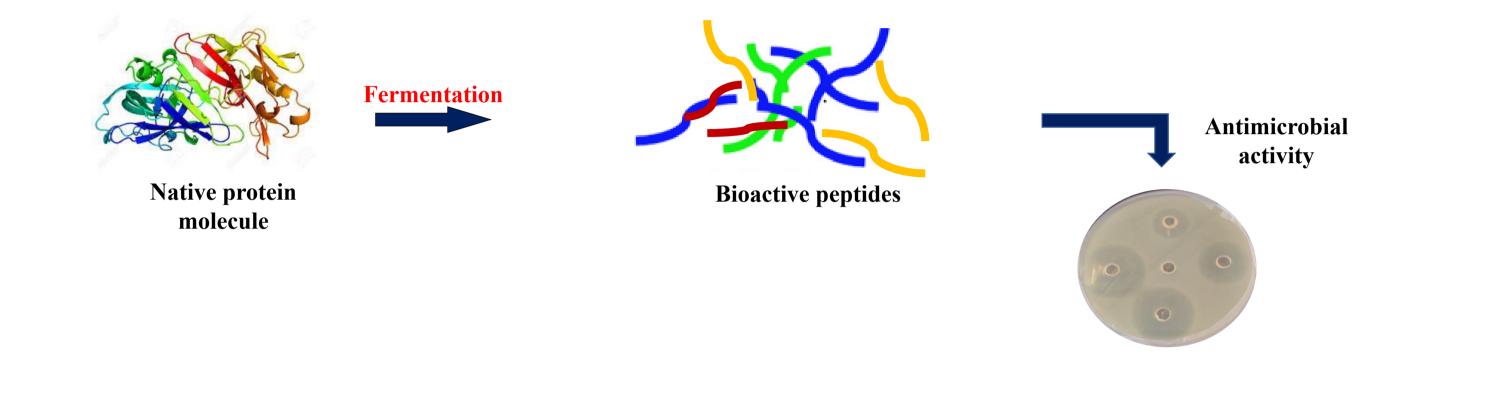
They can be released through gastrointestinal digestion or food processing or fermentation from plant and animal proteins such milk, soy or fish proteins and usually include 2-20 amino acid residues per molecule



Antimicrobial activity of camel milk fermentates 50 kDa, 10 kDa, 5kDa & 3kDa fractions



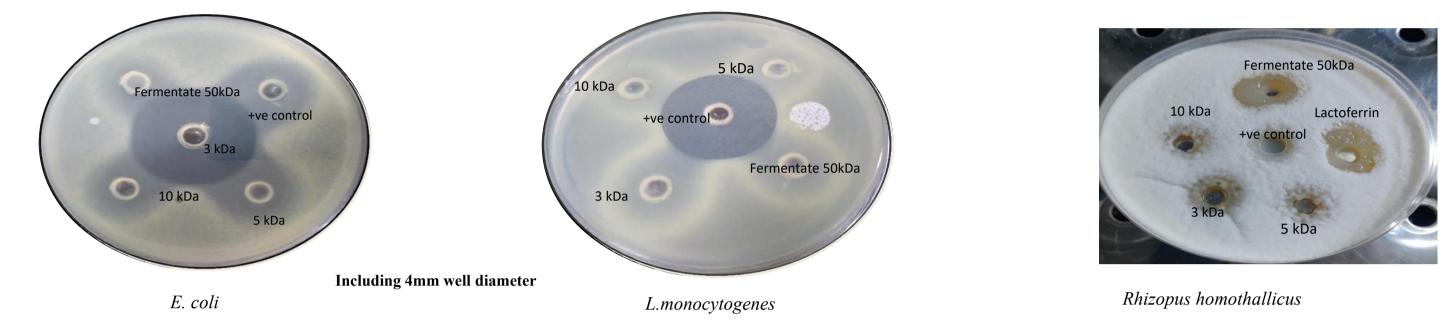
Antimicrobial bioactive peptides (AMPs) interact specifically with bacterial membrane and kill the cell by causing leakage of its content.



Objective: Purification and identification of antimicrobial bioactive peptide derived from Camel and Goat milk by microbial fermentation

Methods

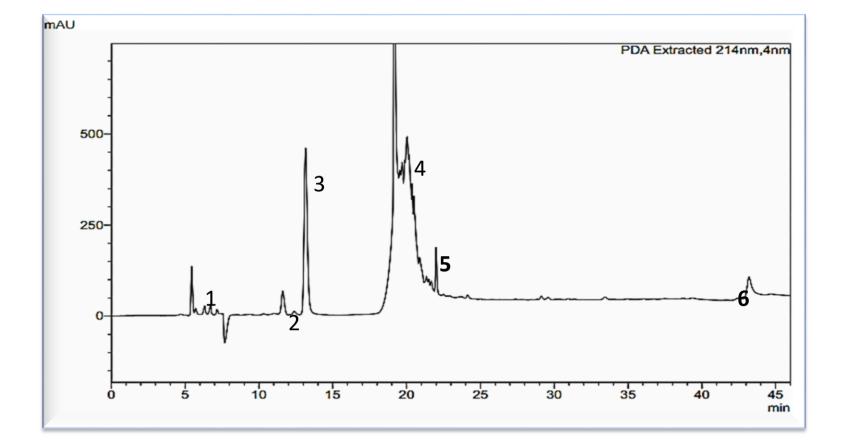
Preparation of 50 kDa, 10kDa, 5kDa and 3kDa fractions from fermented camel mik

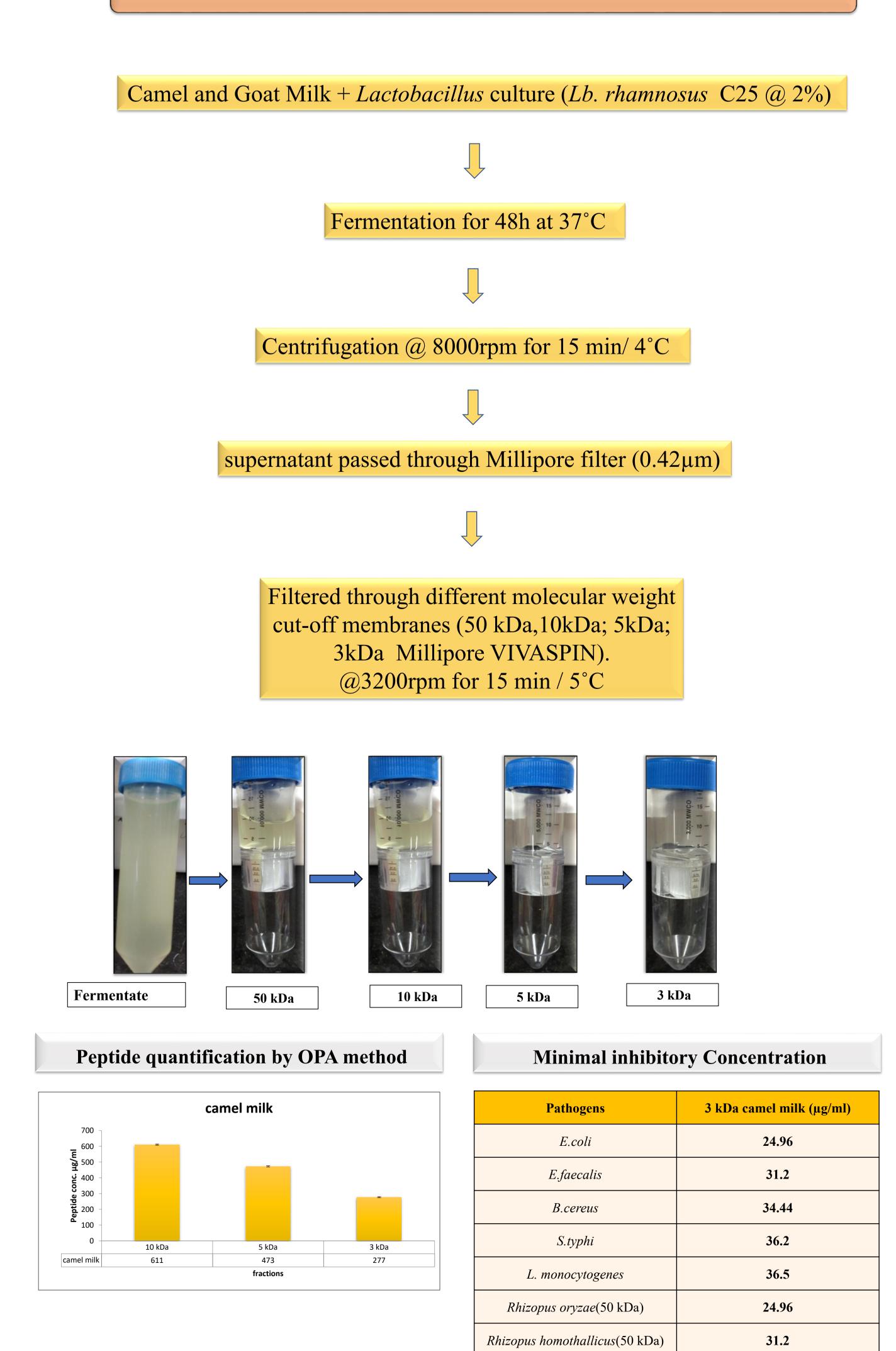


RP-HPLC analysis of camel milk bioactive peptides

HPLC peak of 3kDa fraction from Camel Milk-

- Bioactive peptides of 3kDa for camel milk fractionated and collected by Reverse phase-HPLC.
- HPLC peaks from 3kDa camel milk showed highest antimicrobial activity in peak number 1 & 3 against E.coli.





LC-MS/MS analysis of camel and goat milk bioactive peptides

- One mL of sample of 3 kDa fractions of fermented camel has been outsourced •
- **Obtained Raw Files were searched in Mascot engine using** *Bos taurus* database

LC-MS/MS peaks of 3kDa Camel milk peptide

Identified Proteins using peptides generated through camel milk

um 1120 juliers 1000 (Accession	Protein	MW [kDa]	pI	#Peptid es	Scores
	IPI00721027	Tax_Id=9913 Gene_Symbol=ZCCHC6 zinc finger, CCHC domain containing 6	171.2	6.8	3	55.4 (M:55.4
	IPI00838436	Tax_Id=9913 Gene_Symbol=CASC5 hypothetical protein	268.3	5.5	3	55.0 (M:55.0
	IPI00713748	Tax_Id=9913 Gene_Symbol=SERPINA7 Thyroxine-binding globulin	46.0	5.5	2	62.0 (M:62.0
milk protein analysis results	IPI00825160	Tax_Id=9913 Gene_Symbol=LOC789339 myosin light chain 2a-like, partial	12.3	4.4	2	49.7 (M:49.7
Search Engine Database Ident. Compounds whey_shveta_23.5.16/CAM/ Mascot, 2.5.0 IPI_bovine, 500/8600 roteinAnalysisResults.mgf IPI_bovine_3.73.fasta 500/8600	IPI00692457	Tax_Id=9913 Gene_Symbol=GPD2 Uncharacterized protein	81.0	6.6	2	47.9 (M:47.9
Score: 117.49 MW [kDa]: 18.30 pl: 5.14 No. of Peptides: 5 50 60 70 80 90 100 110 120	IPI01003719	Tax_Id=9913 Gene_Symbol=FREM3 FRAS1 related extracellular matrix 3	239.1	5.0	2	47.1 (M:47.1
Site P Range Sequence Modification Type [%] 0 106-113 E.NLHLPLPL.L CID	IPI00728345	Tax_Id=9913 Gene_Symbol=BAZ2A bromodomain adjacent to zinc finger domain protein 2A	210.0	6.3	2	46.9 (M:46.9
22 0 106-113 E.NLHLPLPLL CID 02 0 106-115 E.NLHLPLPLLQ.S CID 27 0 106-116 E.NLHLPLPLLQS.L CID 95 0 107-113 N.LHLPLPLL CID 03 0 146-161 Y.QEPVLGPVRGPFPIIV CID	IPI00843435	Tax_Id=9913 Gene_Symbol=LRRC29 leucine rich repeat containing 29-like	66.2	6.6	2	46.7 (M:46.7
protein analysis spectrum report	IPI01002995	Tax_Id=9913 Gene_Symbol=LOC100337431 solute carrier family 12, member 7-like	117.8	8.7	2	46.3 (M:46.3
3147.d/ProteinAnalysisResults.mgf A7 Thyroxine-binding globulin Parent m/z: 803.41876, 2+ Score: 46.73	IPI00963807	Tax_Id=9913 Gene_Symbol=SLC12A4 solute carrier family 12 (potassium/chloride transporters), member 4	120.5	6.2	2	44.3 (M:44.3)
b (1)++ 965.32 903.48 903.48 1125.58	IPI01028244	Tax_Id=9913 Gene_Symbol=SETDB1 histone- lysine N-methyltransferase SETDB1	142.9	5.8	2	43.6 (M:43.6)
Sin.27 0.00 2- 737.39 0.02- 136.70 b.0.2 b.0.1 500 600 700 800 900 1000 1.000 m/z 900 743. +MS2/803.4186), 31.1eV, 32.9 min #4434 (cmpd 3743, +MS2/803.4186), 31.1eV, 32.9 min #4434) 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 m/z Printed: June 6, 2016 2 </td <td>IPI00907863</td> <td>Tax_Id=9913 Gene_Symbol=- Uncharacterized protein (Fragment)</td> <td>35.1</td> <td>6.3</td> <td>2</td> <td>43.1 (M:43.1)</td>	IPI00907863	Tax_Id=9913 Gene_Symbol=- Uncharacterized protein (Fragment)	35.1	6.3	2	43.1 (M:43.1)

Conclusion

- * This work has evaluated the presence of antimicrobial bioactive peptide in camel milk fermented by Lactobacillus strain of dairy origin.
- ★ 34 peptides were identified by LC-MS/MS in 3kDa peptide fractions of camel milk fermented with *L.rhamnosus* C25.
- Consequently, it may be concluded that camel and goat milk proteins underwent an extensive degradation process during fermentation and generated a large number of bioactive peptides.
- ✤ The identified peptide sequences from camel milk were as follows: QIITYRDYLPLVLGREAMR, HGVTFGYEGQKPLFK, PAVKAHLFAAEPGGR, QKAVPYPQRD and goat milk were GQPQVVPVEGSRR, DPFLPKQPILVPQR, AKYIPIQYVLSR, VLLFMVFDEK, ESKESVPSLPQSPVK, REEPRVPPLK.
- Moreover, use of fermented camel and goat milk and their bioactive peptides for nutraceuticals and functional food or inclusion in therapeutic diet for patients with diseases linked to immunity system could be functional.