

IN VITRO EVALUATION OF SAFETY AND PROBIOTIC ATTRIBUTES OF LACTOBACILLI ISOLATED FROM HUMAN **AND GOAT MILK**

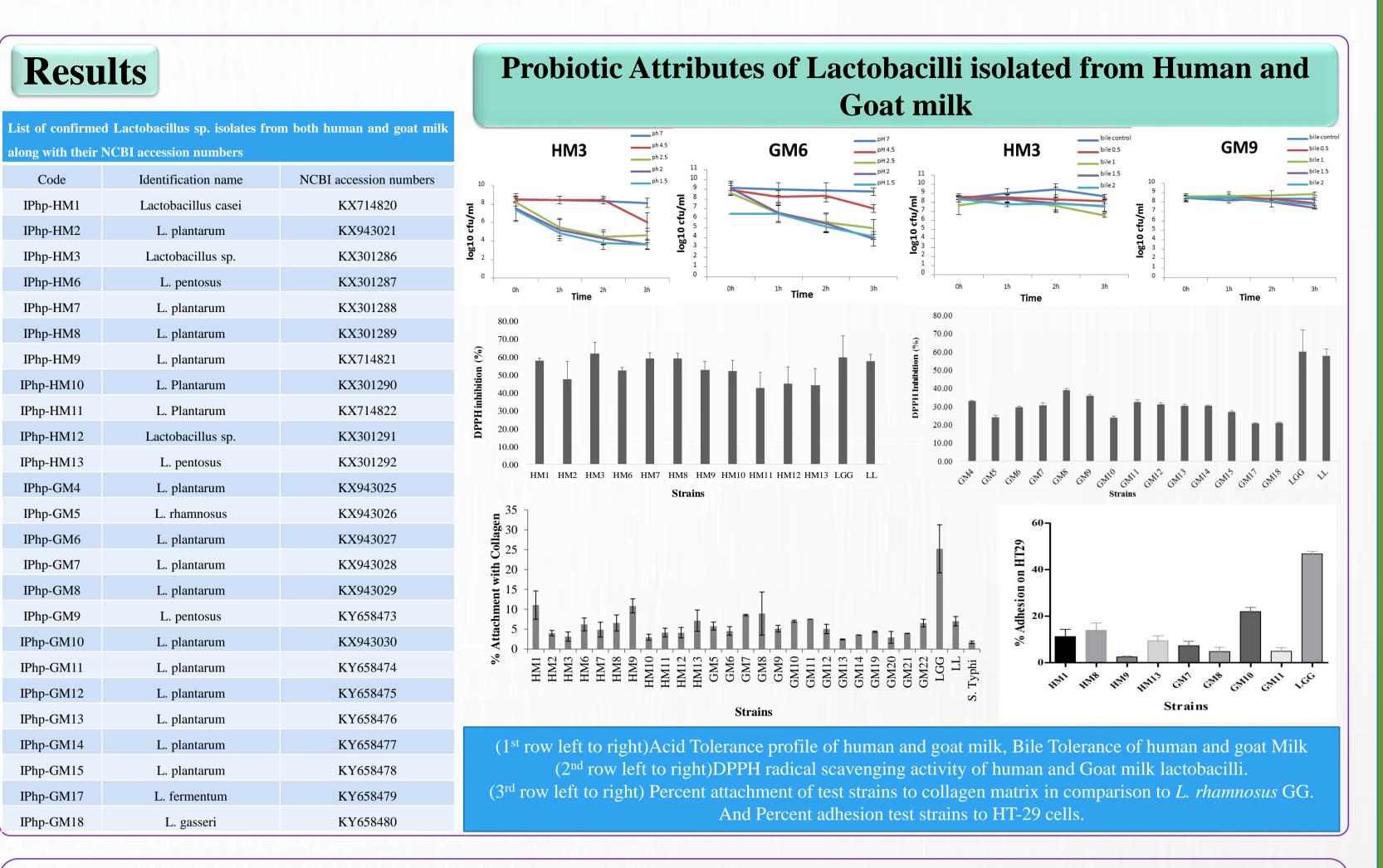


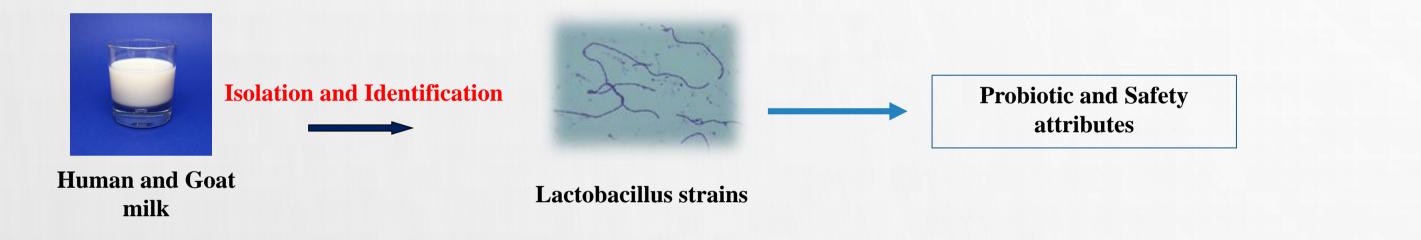
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Introduction: Probiotics, globally recognized as "Live microorganisms which when administered in adequate amount confer some health benefit to the host" (FAO/WHO2001), are the beneficial gut microbes that plays a crucial role in maintaining gut and human health (Marchesi et al 2016). Probiotic administration in adequate amount could improve human health through restoration of host normal microflora, re-establishing the intestinal barrier function, immune homeostasis and support of normal digestive functioning by providing several trace elements to the host (Fong et al 2015). The 'probiotic effect' is mainly mediated by suppression of pro-inflammatory cytokines, expression of anti-inflammatory and associated anti-oxidative and anti-microbial activity. The interference of probiotics with adhesion and colonization of pathogens also seems to be involved in probiotic action.

Microbial strains serving as candidate probiotic are mostly isolated from traditional fermented milk products (curd, lassi, cheese(s) etc.) and fermented fruits and vegetables. However, the isolation source varies between studies, regions and do have impact over functionality of isolates (Swain et al 2014). Microbial composition varies between regions, environmental conditions and fermentation type. Although milk, food and vegetables are explored a lot for isolation of probiotic strains, researchers believe that strains with human origin may survive better during human gastric transit compared to those of non-human origin (Ranadheera et al 2014). Keeping this in mind, healthy human milk samples are explored for selection of strains with rich probiotic potential. Safe transit through stomach and survival and colonization in the intestinal tract are the foremost parameters to qualify as a potential candidate for further screening for probiotic attributes (Kotzamanidis et al 2010). Microbes to be ingested as probiotic strain have to encounter various stress factors during the establishment in the gastro-intestinal tract (GIT). Probiotic bacteria must retain viability during its interaction with stomach acid, bile and high osmolarity in the small intestine (Franz and Holzapfel 2011). Keeping this in view, a bacterial strain to serve as potential probiotic should survive the pH stress of gastric acid. Tolerance towards bile acids is attributed to presence of bile salt hydrolases (BSH), product of *bsh* gene of bacteria (Begley et al 2006). Presence and expression of bsh gene is targeted as one of the criteria for probiotic strain selection (Patel et al 2010). After safe transit to the small intestine, probiotic strains need to adhere to the intestinal lining, determined by multiple factors viz. cell surface hydrophobicity (CSH); cell adhesion potential; aggregation. Adhesion to the intestinal epithelial cell lining along with bacterial CSH are among the most important characteristics of lactobacilli for selecting probiotic strains (Ouwehand et al 1999; Younes et al 2012). Higher bacterial hydrophobicity can be directly co-related to their stronger adherence capability (Pan et al 2006). Auto-aggregation potential of bacteria plays an important role in adhesion to intestinal cells (Dunne et al 2001). It tells about the activity of bacterial cells to interact with them in a non-specific way, which is pre-requisite for GIT colonization (del Re et al 2000). Anti-microbial activity is considered to be a significant functional criterion for competitively inhibiting the pathogenic intestinal microflora through production of organic acids, H2O2, bacteriocins etc. LABs possess strong anti-oxidative activity and decrease the risk of reactive oxygen species (ROS) accumulation (Achuthan et al 2012). Oxidative stress is caused by an imbalance between ROS or free radical production and body antioxidant defense, which alters the normal cellular functions, further leading to several clinical situations.





Objective: The aim of the present study was to establish the indigenous probiotic strains from human and goat milk along with their safety validation.

Antil	bact	eria	l pro	file	of La	ctob	acilli i	solat	ted 1	froi	n hı	ım	an a	and	Goa	t mi	lk sa	ampl	es			
		Antibacter	rial profile	of lacto	bacilli isolato	ed from hu	nan milk sam	ples					An	tibacteria	d profile o	of CFSs of	lactobac	illi isolated	from goa	t milk sam	ple	
	S. aureus	B. cereus	S. Mutams	L. monocytogenes	S. enterica Typhi	P. mirabilis	S. flexneri	P. aeruginosa	E. coli	K. pneumonia			. aureus	. Cereus	. mutams	. monocytogenes	. enterica Typhi	. mirabilis	S. flexneri	. aeruginosa	. coli	. pneumoniae
HM1	-	++	-	-	+	-	++	++	-	-			S.	В	S.	Г	S.	Ъ.	S	Ч.	н	K.
HM2	-	+++	-	-	++	-	++	+	-	-	GM	4	-	-	-	-	-	-	+	++	-	-
HM3	-	++	-	-	+	-	+	+	-	-	GM	5	-	-	-	-	-	-	++	++	-	-
HM6	-	++	-	-	+	-	++	+	-	-	GM	6	-	-	-	-	-	-	+	++	-	-
HM7	-	++	-	-	+	-	++	+	-	-	GM	7	-	-	-	-	-	+	-	++	-	-
HM8	-	+	-	-	+	-	-	+	-	-	GM	8	-	-	-	-	-	-	+	++	-	-
HM9	-	+	-	-	+	-	-	+	-	-	GM	9	+	-	-	-	-	+	-	++	-	-
HM10	-	+	-	-	+	-	-	+	-	-	GM		+	-	-	-	-	+	-	++	-	-
HM11	-	+	-	-	+	-	-	+	-	-	GM		+	-	-	-	-	+	-	++	-	-
HM12	-	+	-	-	+	-	+	+	-	-	GM	12	-	-	-	-	-	+	-	++	-	-
HM13	-	++	-	-	+	-	++	+	-	-	GM	13	+	-	-	-	-	-	+	+	-	-
LGG	+	+	-	-	+	-	++	+	-	-	GM	14	+	-	-	-	-	-	+	++	-	-
LL	-	+	-	-	-	-	-	-	-	-	GM	15	+	-	-	-	-	-	+	++	-	-
>11mm - No	activity (-)	; 12-15m <u>m</u> -	- Weak inhib	ition (+);	16-19 mm – M	oderate/Aver	age inhibition (+	-+); 20-24 - S	trong inhi	bition	GM	17	+	-	-	-	-	-	-	++	-	-
(+++); <25mr	n - Very st	trong inhibit	ion (++++)								GM	18	++	-	-	-	-	+	+	+++	-	-

Validation of safety parameters of candidate probiotic LAB Isolates

Antibiotic susceptibility assay (Human milk Isolates)

Antibacterial susceptibility profile of human milk lactobacilli to commercial antibiotics													
S.N	Antibiotics	Lactobacillus isolates											
0.		HM1	HM2	HM3	HM6	HM7	HM8	HM9	HM10	HM11	HM12	HM13	
1	Ampicillin	R	S	S	S	S	S	S	S	S	S	S	
2	Imipenem	S	S	S	S	S	S	S	S	S	S	S	
3	Meropenem	S	S	S	S	S	S	S	S	S	S	S	
4	Methicillin	R	R	R	R	R	R	R	R	R	R	R	
_	0 '11'	D	D	D	D	D	D	D	D	D	D	D	

Methods

Isolation and Identification of lactobacilli from Human and Goat Milk

Isolated from milk by standard serial plate standard dilution on MRS Agar

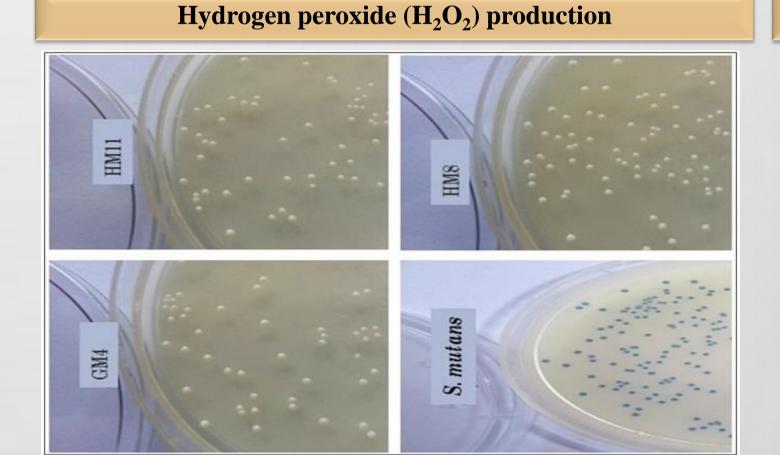
Morphological and Chemical Identification using Catalase test and Gram's staining and scanning electron micrographs

Molecular level identification by polymerase chain reaction (PCR) using Lactobacillus genus specific primers and 16S rDNA sequencing and MALDI-TOF

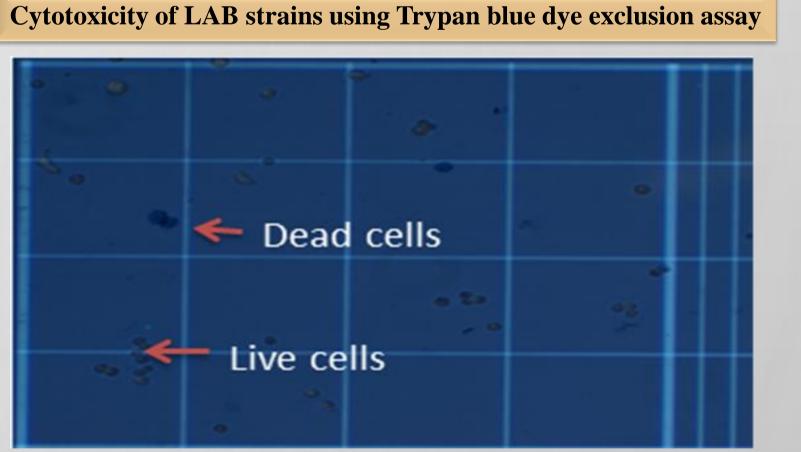
> Identified lactobacilli strains were checked for their Probiotic and Safety attributes.

	Oxaciiiii	К	K	K	K	K	ĸ	K	K	ĸ	K	К
6	Penicillin	R	S	15.3 ± 0.57	19 ± 0	S	17.6 ± 0.57	19 ± 1	S	16.3 ± 0.57	16.6 ± 0.57	S
7	Cefuroxime	17.6 ± 0.57	R	R	R	S	16.3 ± 0.57	R	16 ± 0	R	R	R
8	Cefoxitin	R	R	R	R	R	R	R	R	R	R	R
9	Ceftazidime	15.3 ± 0.57	R	R	R	18.3 ± 0.57	17.3 ± 0.57	16.3 ± 0.57	R	16.3 ± 0.57	16 ± 1	R
10	Cefotaxime	15.6 ± 0.57	S	S	S	S	S	S	S	S	S	S
11	Teicoplanin	R	R	R	R	R	R	R	R	R	R	R
12	Vancomycin	R	R	R	R	R	R	R	R	R	R	R
13	Ciprofloxacin	R	R	R	R	R	R	R	R	R	R	R
14	Ofloxacin	R	R	R	R	R	R	R	R	R	R	R
15	Gentamicin	17.3 ± 0.57	16.3 ± 0.57	16 ± 0	15.3 ± 0.57	16.6 ± 0.57	17.3 ± 0.57	17 ± 1	16.3 ± 0.57	15.6 ± 0.57	16.3 ± 0.57	R
16	Streptomycin	R	R	R	R	R	R	R	R	R	R	R
17	Tobramycin	R	R	R	R	R	R	R	R	R	R	R
18	Chloramphenic	17.3 ± 0.57	S	S	S	S	S	S	S	S	S	S
	ol											
19	Clindamycin	S	R	R	R	R	R	15.6 ± 0.57	R	R	16.3 ± 0.57	R
20	Erythromycin	17.6 ± 0.57	S	S	S	S	S	S	S	S	S	S
21	Fusidic acid	R	16.3 ± 0.57	17.3 ± 0.57	16.6 ± 0.57	17 ± 1	16.3 ± 0.57	R	17 ± 0	R	17.6 ± 0.57	15.6 ± 0.57
22	Nitrofurantoin	R	S	S	S	S	S	S	S	S	S	S
23	Tetracycline	19.3 ± 0.57	S	17.6 ± 0.57	18.3 ± 0.57	S	19 ± 0	17.3 ± 0.57	18.6 ± 0.57	16.3 ± 0.57	S	19 ± 0
24	Tigecycline	16.6 ± 0.57	S	S	S	S	S	S	S	S	S	S
25	Co-	R	R	R	R	R	R	R	R	R	R	R
	Trimoxazole											
26	Trimethoprim	R	S	R	R	R	R	R	R	R	19 ± 0	18.3 ± 0.57

signifies resistance; 'S' signifies sensitive; and numerical values \pm SD signifies intermediate resistance



Brucella agar plates showing negative H₂O₂ production by lactobacilli and positive (blue colonies) for S. mutans



Naebaur's hemocytometer indicating live (colorless) and dead (purple) HT29 cells after co-incubation with LAB strains for 2h



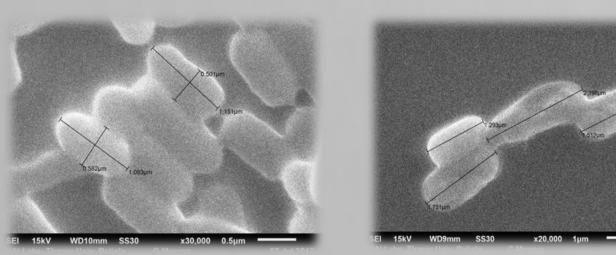
Morphological Identification of Isolated Lactobacilli





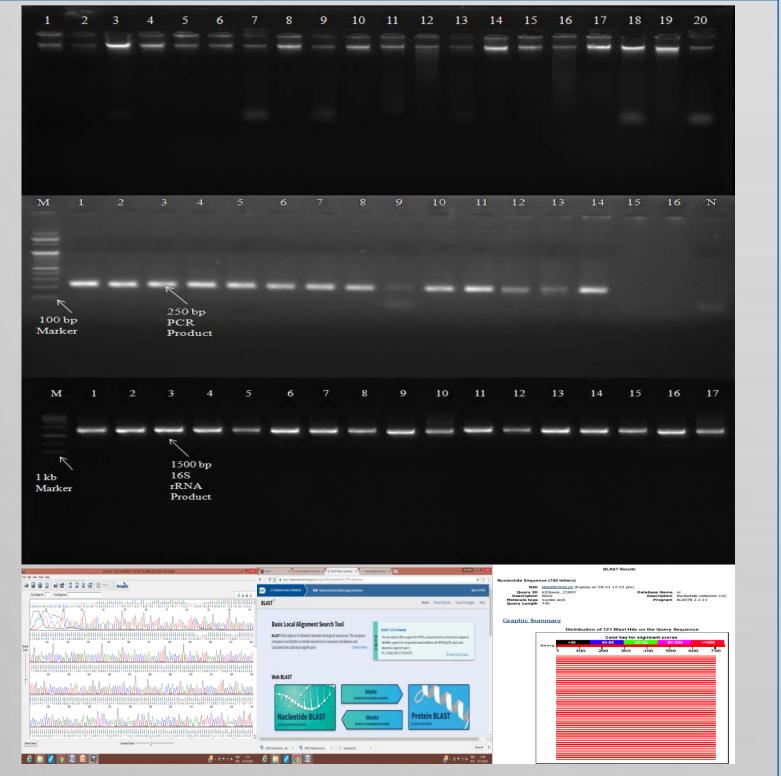






Gram's staining (First Row), negative staining (Second Row), and scanning electron micrographs (Third Row) for LAB isolates





Agarose gel representative for genomic DNA(1st row), Genus specific PCR product for Lactobacillus sp.(2nd row), 16S rDNA based PCR product(3rd row); and representative 16S sequence profile, BLAST search tool and BLAST results (last row). Lanes 'M' denotes marker i.e. ladder of 100bp and 1kb, numerical values denotes independent wells and samples

Hydrogen peroxide (H₂O₂) production

Blood agar plates showing negative hemolysis for lactobacilli and positive for S. mutans



Conclusion

We wanter and goat milk served as a good source of LAB strains falling under different species.

Wight HM3, HM6, HM8, HM9 and HM10 strains displayed higher acid tolerance for all pH ranges i.e. 7, 4.5, 2.5, 2 and 1.5 whereas, among goat milk isolates, GM6 showed highest acid tolerance at lowest studied pH (1.5) after 3h of incubation.

All human as well goat milk lactobacilli were negative for hemolytic activity and hydrogen peroxide production.

Section 4.1 Section 2.1 Secti complete resistance has been observed towards methicillin, oxacillin (β-lactams); teicoplanin, vancomycin (glycopeptides) and ciprofloxacin, ofloxacin (quinolones).

Will High percent viability (95% or more) of HT-29 cells was recorded with all the test strains after 2h of incubation, indicating zero toxicity and ensuring safety of LAB strains under in vivo conditions