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# Cloning and expression of cultural filtrate proteins from novel and native strains of *Mycobacterium avium* subspecies *paratuberculosis* and their application in ELISA based sero-diagnosis of Johne's disease

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Johne's disease (JD), caused by Mycobacterium avium subspecies paratuberculosis (MAP), is endemic in livestock leading to low per animal productivity. MAP as survives pasteurization, poses a public health problem because of high exposure to animals and humans. There is an urgent need for newer diagnostic tests with high specificity and sensitivity as the current ones suffer from lower sensitivity and specificity. In present study, six Mycobacterium avium subspecies paratuberculosis (MAP)-specific culture filtrate proteins (CFPs) were produced and evaluated for sero-diagnosis of MAP infection in goat and cattle herds in India. Genes encoding for six MAP-CFPs were amplified and cloned into easy cloning vector pJET1.2/pTZ57R followed by sub-cloning into expression vector pET28a (+)/pET22b (+) containing C-terminal Histidine. Recombinant CFPs (r-CFPs) expressions were optimized in Escherichia coli (Rosetta cells) and purified using Ni-NTA affinity chromatography. In SDS-PAGE, MAP CFPs viz., 1693c, 2168c, ModD, 85C, Pep AN and Pep AC showed 22, 24, 55, 38, 20 and 25 kDa molecular masses, respectively. Identity of these r-CFPs was further confirmed by immunoblotting. We developed six different ELISAs using the six individual r-CFPs and one additional ELISA i.e. cocktail ELISA (c-ELISA) was prepared using cocktail of all 6 r-CFPs. The performance of all seven ELISAs were further evaluated against whole cell protoplasmic based indigenous ELISA (i-ELISA). c-ELISA showed almost similar sensitivity as shown by i-ELISA. However, individual r-CFP based ELISA could not reach up to the sensitivity of cocktail of six r-CFPs. None of the r-CFP showed any false positive (as compare to i-ELISA) thereby specificity was 100%. Results of ELISA tests based on cocktail of r-CFPs, ModD and 85C were quite similar to i-ELISA from goat sera whereas in cattle serum c-ELISA was comparable with i-ELISA. Our study showed a comparable specificity of c-ELISA for the diagnosis of JD and it may have applicability in region where disease is endemic. Future validation of c-ELISA against gold standard or confirmatory tests would give a better insight on its diagnostic potential over i-ELISA.

Keywords: Cattle, Culture filtrate proteins (r-CFPs), c-ELISA & i-ELISA, Goats, Livestock

Johne's disease (JD) is chronic incurable granulomatous enteritis characterized by persistent or intermittent diarrhoea, progressive weight loss, debilitation and death<sup>1</sup>. It is caused by *Mycobacterium avium* subspecies *paratuberculosis* (MAP) that has wide host distributions, infecting livestock<sup>2</sup>, wild animals<sup>3</sup> (rabbits<sup>4</sup>, Blue-bull<sup>5</sup>) as well as primates<sup>6</sup> and human beings<sup>7</sup>. The disease is endemic in

livestock leading to low per animal productivity<sup>2</sup>. Since MAP gets secreted in milk of affected domestic livestock and possibly survives during pasteurization as well, with the high exposure to animals and humans, it has emerged as a 'public health issue'<sup>7</sup>. Diagnostic tests may be helpful in identification and elimination of exposed animals in the early stage of infection<sup>8</sup>. JD control programs are hampered largely due to lower sensitivity and specificity of current diagnostic tests. Tests are based on 'antigen detection' (culture, ZN-staining, PCR, *in situ* hybridization and Immunofluorescence), 'antibody detection' (Cell mediated as well as humoral Immune response) and

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pathology based detection (gross and histopathology). Specificity and sensitivity of these tests varies and is often not consistent<sup>9</sup>. Therefore, newer diagnostic tests with high specificity and sensitivity need to be developed.

Culture filtrate proteins (CFPs)/secretary proteins are predictable to be immune-reactive or, immunedominant due to their occurrence in the extracellular environment of cells where they are more possible to encounter the sensitized immune cells<sup>10</sup>. Cho et al.<sup>11</sup> checked infected serum by immune-blot analysis collected from MAP infected cattle, showed strong reactivity to CFPs in compare to antigens of intracellular origin. Using CFPs for sero-diagnostics of MAP, sensitivity of the tests has enhanced and similar results came using CFPs of other mycobacteria<sup>12</sup>. As compared to conventional ELISAs, the use of CFPs led to increase in sensitivity of the antibody detection by 20.0% 13. Studies support the CFPs as better antigens for solid-phase ELISA assay performing better sensitivity 10,14. Pradenas et al. 15 also reported that most of the CFPs having low molecular weight (>70 kDa) showed strong reactivity with serum from culture-positive animals. They have also observed a drastic alteration in immune reactivity of CFPs with MAP heavy shedders cattle and cattle having clinical cases of paratuberculosis. MAP infected small ruminants (goats and sheep) serum also displayed good reactivity with MAP CFPs examined by immune-blot assays<sup>16</sup>. However, selection of antigen remains a challenge, since no single CFP based antigen can recognize all the stages of infection especially sub-clinical infection. The results of earlier studies support the use of cocktail of CFPs and it could be a better candidate for sero-diagnosis of MAP infection.

In the present study, we cloned, expressed, and purified six MAP CFPs having potential sero-diagnostic value. Diagnostic potential of rCFPs were assessed by developing the different ELISAs (using individual CFP and cocktail of all 6 CFPs) and its comparison with whole cell protoplasmic antigen

(derived from of native MAP strain - 'S-5 Indian Bison Type') based indigenous ELISA (i-ELISA).

#### **Materials and Methods**

#### Reference strain and sample collection

MAP reference strain 'S 5', 'Indian Bison Type' was a novel biotype obtained from Microbiology laboratory, Central Institute for Research on Goats (CIRG), Makhdoom, Mathura, UP, India. Culture was propagated as per standardized protocols<sup>17</sup>. Serum samples from 90 goats and 120 cattle were included for evaluation of different diagnostic ELISAs. Animals were from native livestock breeding farms of three states (Uttar Pradesh, Haryana and Madhya Pradesh) of India.

#### Targeted immunogenic CFPs

In this study, we targeted 6 CFPs encoding genes viz. 1693c; 2168c, ModD; 85C; Pep AN and Pep AC for cloning, expression, purification and for developing ELISA tests. Immunoreactivity of these selected CFPs have been demonstrated in earlier studies<sup>18</sup> (Table 1).

### **Cloning of amplified gene segment of respective CFPs genes** *PCR amplification of CFPs genes*

Genes encoding MAP CFPs were amplified using MAP genes specific primers in PCR (Table 2). Cycling parameters were: Initial denaturation was done at 94°C, 5 min., followed by 37 cycles of denaturation at 94°C, 30 s, annealing at indicated temperatures (Table 2) for 30 s, extension at 72°C for 1.0 min, and final extension at 72°C for 5 min<sup>17</sup>. Amplified PCR products of MAP CFPs genes were purified from gel by agrose gel extraction DNA purification kit (GeneJet gel extraction kit, Thermo scientific, USA)<sup>19</sup>.

Easy cloning

Purified PCR products were subjected to restriction digestion (Table 2), ligation into appropriate cloning vector, followed by transformation (Table 3) and plating on ampicilin-containing Luria Bertani (LB) agar plates<sup>20</sup>. Positive clones were confirmed by restriction digestion and sequence analysis. Products

Table 1 — Profile of secretome identified as potential antigens in native strain of MAP							
Secretary protein	Putative function	Gene size	Approx. size of protein	Approx. size of proteins appeared with 6 histidine amino acids			
MAP 1693c	Peptidyl-prolylcis-trans isomerase	474 bp	19 kDA	22 kDa			
MAP 2168c	Hypothetical protein	453 bp	21 kDA	24 kDa			
MAP ModD	Fibronectin attachment	998 bp	52 kDA	55 kDa			
MAP 85C	Mycolyltransferase	999 bp	35 kDA	38 kDa			
MAP Pep AN	Serine proteinase	423 bp	17 kDA	20 kDa			
MAP Pep AC	Serine proteinase	540 bp	22 kDA	25 kDa			

Table 2 — Culture filtrate proteins (CFP)s Primers used in the study							
MAP	Direction	Sequence	Restriction	Product	Annealing		
Gene			Enzyme	Length	temperature		
MAP	Forward	5'-GAA TTC ATG GCC GAC TCC TGC CCG ACC GCC-3'	EcoRI	474 bp	64°C		
1693c	Reverse	5'-CTC GAG CTA GGT CGT GGC GCC GAG GAT C-3'	XhoI				
MAP	Forward	5'-GAA TTC ATG GCA GCG CCG ACC GGT CTG C-3'	EcoRI	453 bp	66°C		
2168c	Reverse	5'-CTC GAG TCA GTT CAT CGC GGT GGC CGC C-3'	XhoI				
$MAP^{\#}$	Forward	5'-CAT ATG GAT CCC GAG GTC CCG ACC-3'	NdeI	998 bp	54°C		
Mod D	Reverse	5'-CTC GAG GGC CGA GAG GGT CTG CTG C-3'	XhoI				
MAP 85C	Forward	5'-GGA TCC ATG GCG GGT GGC TCC CCC GTC-3'	BamHI	999 bp	68°C		
	Reverse	5'-AAG CTT TCA GGT GGC GGG CTG GGC C-3'	HindIII				
MAP	Forward	5'-GGA TCC ATG GCA CCG TCG GGC CTG GC-3'	BamHI	423 bp	58°C		
PepAN	Reverse	5'-CTC GAG TCA GAC GAC CTT GCC GGC CAC C-3'	XhoI	•			
MAP	Forward	5'-GGA TCC ATG GCG CTC AAC CAG AGC GTC-3'	BamHI	540 bp	58°C		
PepAC	Reverse	5'-CTC GAG TCA GGC CGG CGG CCC CTC C-3'	XhoI	•			
[#Cho et al., 2007 <sup>ref.18</sup> ]							

	Table 3 — Cloning and expression vector and kits, restriction enzyme used in this study							
MAP CFPs		Easy Cloning	Cloning in Expression Vector					
Gene	Vector / kit	Antibiotics and supplements	Cells used in	Expression	Cells used in			
			Transformation	vector	Transformation			
MAP 1693c	pJET1.2/blunt vector, CloneJET	Ampicillin	XL-1- Blue	pET28a(+)	XL-1- Blue			
	PCR Cloning kit (#K1231)	$(100  \mu g/mL)$	competent cells	expression vector	competent cells			
MAP 2168c	pJET1.2/blunt vector, CloneJET	Ampicillin	XL-1- Blue	pET28a(+)	XL-1- Blue			
	PCR Cloning kit (#K1231)	$(100  \mu g/mL)$	competent cells	expression vector	competent cells			
MAP Mod D	pTZ57R/T Cloning vector,	Ampicillin (100 μg/mL),	XL-10 Ultra-	pET22b(+)	XL-10 Ultra-			
	InsTAclone PCR Cloning Kit	X-gal (20 µg/mL) and IPTG	competent cells	expression vector	competent cells			
	(#1214)	$(20\mu g/mL)$						
MAP 85C	pTZ57R/T Cloning vector,	Ampicillin (100μg/mL),	XL-10 Ultra-	pET28a (+)	XL-10 Ultra-			
	InsTA clone PCR Cloning Kit	X-gal (20µg/mL) and IPTG	competent cells	expression vector	competent cells			
	(#1214)	$(20\mu g/mL)$						
AP Pep AN	pJET1.2/blunt vector, CloneJET	Ampicillin	XL-1- Blue	pET28a(+)	XL-10 Ultra-			
	PCR Cloning kit (#K1231)	$(100\mu g/mL)$	competent cells	expression vector	competent cells			
AP Pep AC	pJET1.2 / blunt vector, Clone	Ampicillin	XL-1- Blue	pET28a (+)	XL-10 Ultra-			
	JET PCR Cloning kit (#K1231)	$(100\mu g / mL)$	competent cells	expression vector	competent cells			

were named as pJET-MAP1693c, pJET-MAP2168c, pTZ57R/T-Mod D, pTZ57R/T-85C, pJET-MAP Pep AN and pJET-MAP Pep AC.

Cloning in the expression vector

CFP genes were transferred from cloning vector to expression vector<sup>20</sup>. Briefly, expression vector (pET28a(+)pET22b(+)was digested with appropriate restriction enzyme (Table 2), gel purified and dephosphorylated by alkaline phosphatase enzyme (Fermentas, USA). Gel purified and dephosphorvlated vector and respective fragment (digested from cloned product) were ligated using T4 DNA Ligase (Fermentas, USA) and transformed into XL-1/XL-10 competent cells followed by plating on kanamycin (pET28a vector)/ampicillin (pET22b vector) containing LB agar plates. Positive clones were confirmed by colony PCR and sequencing using T7 universal primer. Products were named as pET28a-MAP1693c, pET28a-MAP2168c, pET22b-Mod D, pET22a-85c,

pET28a-MAP Pep AN as well as pET28a-MAP Pep.

#### AC.Expression of recombinant proteins

To express recombinant CFP proteins, cloned constructs (pET28a-MAP1693c, pET28a-MAP2168c, pET22b-Mod D, pET22a-85c, pET28a-MAP Pep AN as well as pET28a-MAP Pep AC) vectors were further transformed into Rosetta competent cells, and a single colony for each CFP clone was picked up and propagated in 10 mL LB broth supplemented with antibiotics {kanamycin/ampicillin (50/100 µg per mL), respectively} at 37°C in shaking mode (200 rpm) for 16 h. For expression, 500 mL LB broth supplemented with antibiotic was inoculated with 1% culture (overnight grown) and propagated for 5 h till absorbance reached between 0.5-0.6 at 595 nm. Then, isopropyl-β-D thiogalactopyranoside {IPTG} was added to a optimized concentration of 0.3 mM for induction of protein expression. Bacterial cells were incubated at indicated time and temperature (Table 4)

before being harvesting. Both soluble form (medium) and inclusion bodies (cell pellet) were assessed for optimum production of recombinant protein.

Recombinant MAP proteins that formed inclusion bodies (MAP 1693c and MAP 85C) were purified as follows: Pellet of Rosetta cells harbouring recombinant MAP CFPs were resuspended by gentle stirring in 15 mL equilibration buffer (50 mM, Tris HCl, 200 mM NaCl, 5 mM DTT, 1.0 mM PMSF, pH 8.0). The 15 mL suspension subjected was to sonication (at optimized conditions). MAP CFPs purification was done using affinity chromatography method on a Nickel-nitrilotriacetic acid (Ni-NTA) gel matrix (Qiagen). A 50 mL conical tube containing 12 mL sonicated suspension equilibrated with 1.0 mL Ni-NTA resin in horizontal position for 8 h at 4°C. After incubation, the equilibrated suspension of MAP CFPs were centrifuged at 515 g, 10 min at 4°C. Supernatants (flow through) were removed and pellets were washed thrice with wash buffer (50 mM, Tris HCl, 200 mm NaCl, 5 mm DTT, 1.0 mm PMSF, pH 8.0 and 20 mM Imidazole) and supernatant was saved. All recombinant proteins was then eluted by increasing concentration of the imidazole to 250 mM (50 mM, Tris.HCl, 200 mM NaCl, 5 mM DTT, 1.0 mm PMSF, pH 8.0 and 250 mM). Fractions contained recombinant proteins was visualized by SDS-PAGE electrophoresis followed by Coomassie brilliant blue staining. For purification of soluble proteins (MAP 2168c; MAP Mod D; MAP Pep AN and MAP Pep AC), the medium was directly subjected to Ni-NTA affinity chromatography. The yield of proteins were in the range of 12 mg to 200 mg/L of bacterial culture (Table 4)

### $\label{lem:confirmation} \textbf{Confirmation, Immuno-reactivity and estimation of recombinant proteins}$

For confirmation, purified products were run in 12% SDS-PAGE and subjected to Coomassie blue staining to analyze the size of desired products. In addition to this, recombinant His-fusion proteins were analyzed by Western blot.

Confirmation of proteins and their size

Nitrocellulose membrane was incubated with 20 mL of peroxidase-conjugated anti-His<sub>6</sub> peroxidase (Cat. No. 11965085001, Roche Diagnostic GmBH, Germany) in 100 mU/mL for 1h at room temperature (25-30°C on shaker, followed by five times washing with Transfer buffer saline with Tween (TBST). Visualization of immune-reactive proteins bands were done using 2,3-diaminobenzidine (DAB) (Sigma-Aldrich).

#### Measure of the immunoreactivity

Nitrocellulose membrane was incubated with 20 mL of diluted sera (primary antibody) in 1:200 dilution and the membrane was kept at 4°C on rocker for overnight. Further, Nitrocellulose membrane was washed thrice using TBST buffer for 10 min. Reactivity was recorded by incubating nitro-cellulose membrane with per-oxidase-labelled rabbit based anti-species (Goat/Cattle) whole IgG antibody (Sigma Aldrich) in 1:2000 dilutions and incubated for 1.0 h at room temperature on rocker, followed by washing with TBST five times. Visualization of immunogenic proteins bands were completed using 2,3-diaminobenzidine (DAB) reactivity (Sigma-Aldrich).

#### Concentration of purified rCFPs

Concentration of purified rCFPs was estimated using Bradford protein estimation method at 595 nm using ELISA plate spectrophotometer<sup>21</sup> (Table 4).

#### Dot ELISA (d-ELISA)

Serum based d-ELISA was standardized for CFPs as per Singh et al., 2016<sup>22</sup>. Briefly, plastic d-ELISA strips (Cat No: CCN-12, MDI, India) were coated with either 0.5µg in 1.0 µL of standard antigen (MAP 'S 5' strain') used as 'positive control' or individual MAP recombinant protein used as 'test antigen'. Antigencoated strips were blocked with 3% skimmed milk in 1XPBS (1h, 37°C). Test and positive control antigens were reacted with positive and negative control serum (1 h at 37°C). Rabbit anti-goat HRP conjugate was added (30 min at 37°C) and finally antigen antibody reaction was observed with absence of brown dot by adding substrate (DAB). Positive and negative controls were used to read the results of two antigens.

Table 4 — Conditions optimized for expressed recombinant CFPs proteins and their concentrations								
Proteins	Expression form	Temperature and Time	Protein concentration	Protein quantit/litre of culture				
MAP 1693c	Inclusion bodies	30 h at 27°C	0.25 μg/μL in 2mL	50 mg				
MAP 2168c	Soluble	30 h at 24°C	0.12 μg/μL in 2mL	24 mg				
MAP Mod D	Soluble	16 h at 37°C	0.239 μg/μL in 4mL	92 mg				
MAP 85C	Inclusion bodies	30 h at 27°C	0.06 μg/μL in 2mL	12 mg				
MAP Pep AN	Soluble	16 h at 37°C	0.25 μg/μL in 4mL	100 mg				
MAP Pep AC	Soluble	16 h at 37°C	1 μg/μL in 2mL	200 mg				

#### **Recombinant CFP-based ELISA**

'Indigenous ELISA' (i-ELISA) using sPPA as developed in goats by Singh et al.23 and in cattle by Sharma et al. 24 was employed. Either cocktail of 6 r-CFPs; MAP1693c; MAP 2168c; MAP ModD; MAP 85C; MAP Pep AN and MAP Pep AC were used at 6 μg of six r-CFPs (1.0 μg of each r-CFPs) to develop cocktail ELISA (c-ELISA); and 6 µg of each individual r-CFP to develop recombinant-ELISA (r-ELISA). Serum collected from MAP culture positive and culture negative animals were used as serum positive and serum negative controls, respectively. OD values of the samples were expressed as sample to positive (S/P) ratios<sup>25</sup>. Animals came positive and strong positive categories of the S/P ratios were considered positive for MAP infection. Sensitivity and specificity of c-ELISA and r-ELISAs were compared with i-ELISA.

#### S/P ratio and corresponding status of JD

S/P ratios revealed corresponding status of JD as between 0.00-0.09- Negative (N), 0.1-0.24-Suspected/Borderline (S), 0.25-0.39- Low Positive (LP), 0.4-0.99- Positive (P) and 1.0-10.0- Strong Positive (SP).

#### Sensitivity and Specificity

Sensitivity = (True Positive / True Positive + False Negative)  $\times$  100

Specificity = (True Negative / True Negative + False Positive) × 100

#### Statistical analysis

McNemar's test and kappa agreement was used to measure the statistical significance between new tests (c-ELISA and r-ELISAs) and i-ELISA (GraphPad software, USA)

#### Results

#### Cloning and expression of CFPs

Genomic DNA was isolated from 'S5' strain of novel 'Indian Bison Type' of MAP. Genes encoding 6 CFPs (MAP1693c; MAP2168c; MAP ModD; MAP 85C; MAP PepAN and MAP PepAC) were successfully amplified (Fig. 1) and cloned into pJET1.2/blunt or, PTZ57R/T cloning vector (Fig. 2) and then sub-cloned into expression vector viz., pET28a (+)/ pET22b(+). Recombinant clones of CFPs genes were confirmed by colony PCR, restriction digestion and nucleotide sequencing. Purified clones (plasmids) were transformed into competent cells (*E. coli* Rosetta) to evaluate expression of r-CFPs. Optimum r-CFPs production was observed at 0.3 mM

IPTG concentration, 24-37°C temperature and 16-30 h of incubation (Table 4) and all expressed r-CFPs was purified using Ni-NTA resin and confirmed by SDS-PAGE (Fig. 3).

## Comparison of sero-reactivity of six individual r-CFPs and their cocktail of six r-CFPs in Immunoblot and dot-ELISA system

SDS-PAGE analysis of purified r-CFPs revealed single thick band without detection of any other protein band suggesting optimum purity of recombinant products. Identity of r-CFPs was further assessed in immuno-blotting using anti-His antibody, where all the six r-CFPs showed their expected molecular mass (Figs. 4 & 5 A-C) except ModD, which seemed to be higher (55 kDa) than the expected size of 32 kDa. Immunoreactivity of these six r-CFPs was measured with positive serum of goats and cattle by immunoblot analysis. MAP 1693c, MAP 2168c and MAP Mod D, r-CFPs showed similar reactivity

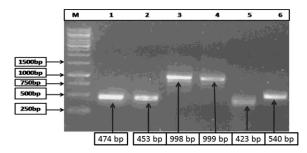


Fig. 1 — Amplification of Six MAP CFPs genes by PCR: M=1kb DNA ladder (#SM0313, Fermentas); lane 1= amplified product of MAP 1693c (474 bp); lane 2= amplified product of MAP 2168c (453 bp); lane 3= amplified product of MAP ModD (998 bp); lane 4= amplified product of MAP 85C (999 bp); lane 5= amplified product of MAP Pep AN (423 bp); lane 6= amplified product of MAP Pep AC (540 bp)

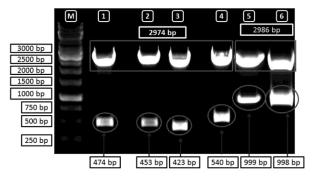


Fig. 2 — Easy cloning and confirmation in pJET 1.2 / pTZ57R/T vector: M=1kb DNA ladder (#SM0313, Fermentas); lane 1 = Xho I and Eco RI digested pJET-1693c; lane 2=Xho I and Eco RI digested pJET-2168c; lane 3 = Bam HI and Xho I digested pJET-Pep AN; lane-4 = Bam HI and Xho I digested pJET-Pep AC; lane 5= Bam HI and Xho I digested pTZ57R/T.85c; lane 6= Nde I and Xho I digested pTZ57R/T.Mod D

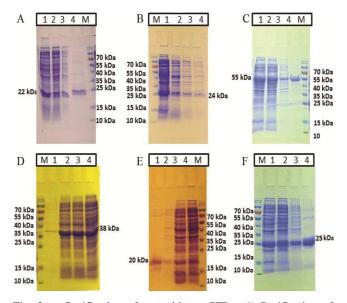


Fig. 3 — Purification of recombinant CFPs: (1) Purification of recombinant MAP 1693c protein expressed in Rosetta expression strain of E.coli (Left to right): lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product, lane M: Marker (#SM0671); (2) Purification of recombinant MAP 2168c protein expressed in Rosetta expression strain of E.coli (Left to right): lane M: Marker (#SM0671), lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product; (3) Purification of recombinant MAP Mod D protein expressed in Rosetta expression strain of E.coli (Left to right): lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product, lane M: Marker (#SM0671); (4) Purification of recombinant MAP 85C protein expressed in Rosetta expression strain of E.coli (Left to right): lane M: Marker (#SM0671), lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product; (5) Purification of recombinant MAP Pep AN protein expressed in Rosetta expression strain of E.coli (Left to right): lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product, lane M: Marker (#SM0671); (6) Purification of recombinant MAP 85C protein expressed in Rosetta expression strain of E.coli (Left to right): lane M: Marker (#SM0671), lane 1. Crude sonicated extract, lane 2. Supernatant, lane 3. Wash buffer, lane 4. Eluted product.

with goats and cattle serum. Whereas, Pep AN and Pep AC showed higher reactivity with goat serum and 85c showed more reactivity in serum of cattle as compared to goats (Fig. 5 B & C). All individual recombinant CFPs showing good antigenicity by individual antigen based dot-ELISA (Fig. 6).

#### Comparison of sero-reactivity of six individual r-CFPs and their cocktail of six r-CFPs with sPPA in ELISA system

OD values of six individual r-ELISA, cocktail ELISA and i-ELISA were expressed as S/P ratio and positivity was determined by comparing S/P ratio of positive and negative controls<sup>25</sup> (Table 5). c-ELISA showed almost similar sensitivity as shown by i-ELISA (Table 6). However, individual r-CFPs based ELISA could not reach up to the sensitivity of cocktail of six r-CFPs in c-ELISA (Table 6). Sero-reactivity of individual r-CFPs to positive goat serum was in the order of ModD > 85C > 2168c > PepAC > PepAN > 1693c (Fig. 7) while in

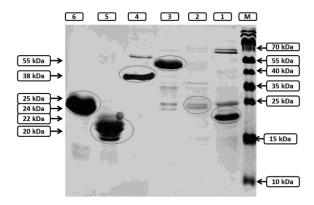


Fig. 4 — Purified expressed recombinant CFPs (Right to left): lane M= Pre-stained page ladder (#SM0671), lane 1= MAP 1693c purified protein, lane 2= MAP 2168c purified protein, lane 3= MAP Mod D purified protein, lane 4= MAP 85C purified protein, lane 5= MAP Pep AN purified protein, lane 6= MAP Pep AC purified protein.

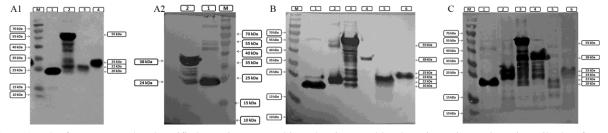


Fig. 5 — (A) The four expressed and purified proteins were subjected to immunoblot detection using anti-6 His antibody. (from left to right): lane M- Pre-stained ladder (#SM0671), lane 1- MAP 1693c, lane 2- MAP Mod D, lane 3- MAP Pep AN and lane 4- MAP Pep AC; (A1) The two expressed and purified proteins were subjected to immunoblot detection using anti-6 His antibody. (from right to left): lane M- Pre-stained ladder (#SM0671), lane 1- MAP 2168c, lane 2- MAP 85C; and (B & C) Imuunoreactivity of secretory proteins with goat & cattle positive serum by Western Blot analysis (from left to right): lane M- Pre-stained ladder (#SM0671), lane 1- MAP 1693c, lane 2- MAP 2168c, lane 3- MAP Mod D, lane 4- MAP 85c, lane 5-MAP Pep AN and lane 6- MAP Pep AC

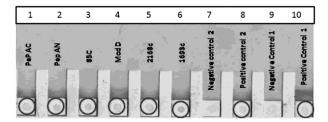


Fig. 6 — Dot-ELISA test for MAP recombinant CFPs showing brown dot for positive samples for MAP infection. (Left to Right): lane 1. MAP Pep AC, lane 2. MAP Pep AN, lane 3. MAP 85C, lane 4. MAP Mod D, lane 5. MAP 2168c, lane 6. MAP 1693c, lane 7. Negative control 2, lane 8. Positive control 2, lane 9. Negative control 1, and lane 10. For lane 1 to 6, 8 and 10 positive sera was used of MAP infected goats, and for Lane 7 and 9 negative sera was used from healthy goats. For lane 7 to 10 sPPA was used as an antigen.

Table 5 — Screening of goats and cattle serum using sPPA based i-ELISA and six recombinant CFPs based six individual r-ELISA and a c-ELISA

and a C-LEIST							
Comparison	Goats - 90		Cattle – 120				
	Positive	Negative	Positive	Negative			
sPPA- i-ELISA	58	32	86	34			
c-ELISA	57	33	83	37			
1693c-ELISA	24	66	77	43			
2168c-ELISA	40	50	78	42			
Mod D-ELISA	56	34	80	40			
85C-ELISA	54	36	79	41			
Pep AN-ELISA	34	56	75	45			
Pep AC-ELISA	37	53	76	46			

Table 6 — Sensitivity and specificity of recombinant CFPs-based ELISA in detecting anti-MAP antibodies in goat sera

Compari-	Animal	+ve	-ve	False	False	Sensitivity and
son	species			+ve	-ve	specificity (%)
c-ELISA	Goat	57	32	0	1	98.0 & 100.0
	Cattle	83	34	0	3	96.5 & 100.0
1693c	Goat	24	32	0	34	41.4 & 100.0
	Cattle	77	34	0	9	89.5 & 100.0
2168c	Goat	40	32	0	18	69.0 & 100.0
	Cattle	78	34	0	8	90.7 & 100.0
Mod D	Goat	56	32	0	2	96.6 & 100.0
	Cattle	80	34	0	6	93.0 & 100.0
85C	Goat	54	32	0	4	93.1 & 100.0
	Cattle	79	34	0	7	91.9 & 100.0
Pep AN	Goat	34	32	0	24	58.6 & 100.0
	Cattle	75	34	0	11	87.2 & 100.0
Pep AC	Goat	37	32	0	21	63.8 & 100.0
	Cattle	76	34	0	10	88.4 & 100.0

cattle it was ModD > 85C > 2168c > 1693c > PepAC > PepAN (Fig. 8).

### Sensitivity and specificity of recombinant CFPs based ELISA in detecting anti-MAP antibodies in comparison to i-ELISA

Sensitivity of recombinant CFPs based ELISA tests was compared with i-ELISA. None of the recombinant CFPs reacted to produce false positive results thereby considered as 100% specific as

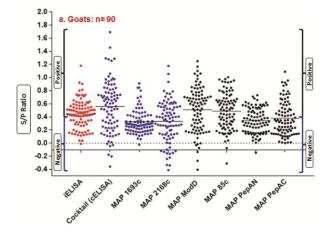


Fig. 7 — Preliminary results using the expressed proteins as solid phase antigens for screening of sera from goats. [Representative results for six individual recombinant proteins based ELISA with traditional i-ELISA and c-ELISA (Cocktail of all six recombinant secretory proteins)]

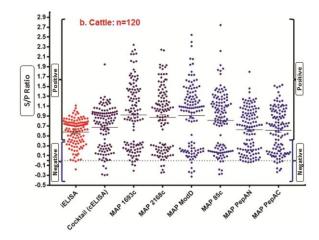


Fig. 8 — Preliminary results using the expressed proteins as solid phase antigens for screening of sera from cattle. [Representative results for six individual recombinant proteins based ELISA with traditional i-ELISA and c-ELISA (Cocktail of all six recombinant secretory proteins)]

compared to i-ELISA. However, sensitivity varied from 41.4 to 98% in serum of goats and 87.2 to 96.5% in bovine serum samples. None of them could provide up to 100% sensitivity even with the use of the cocktail of six CFPs (Table 6).

The correlation between two different ELISA tests was done to detect anti-MAP antibodies, sensitivity and specificity, Mc-Nemar calculation as well as kappa agreement. Results of ELISA tests based on cocktail of r-CFPs, Mod D and 85C were quite similar to i-ELISA in detecting anti-MAP antibodies in goat sera whereas in cattle serum c-ELISA was comparable with i-ELISA (Table 7).

Table 7 — Statistical evaluation between r-CFP/(s) based ELISA and i-ELISA in detection of MAP infection in goats and cattle Recombinant CFP/(s) used in ELISA Relative agreement with i-ELISA

		Goat			Cattle			
	Kappa	95% CI	P value	Kappa	95% CI	P value		
c-ELISA	$0.976\pm0.024$	0.929-1.000	1.000	$0.940\pm0.034$	0.873-1.000	0.2482		
1693c	$0.334 \pm 0.067$	0.202-0.466	< 0.0001	$0.829\pm0.054$	0.723-0.935	0.0077		
2168c	$0.612\pm0.075$	0.465- 0.760	< 0.0001	$0.847 \pm 0.052$	0.745-0.948	0.0133		
Mod D	$0.952\pm0.033$	0.887-1.000	0.4795	$0.883\pm0.046$	0.793-0.974	0.0412		
85C	$0.906\pm0.046$	0.816-0.996	0.1336	$0.865\pm0.049$	0.768-0.961	0.0233		
Pep AN	$0.502\pm0.076$	0.354-0.650	< 0.0001	$0.794 \pm 0.058$	0.681-0.908	0.0026		
Pen AC	0.556+0.076	0.407-0.705	< 0.0001	0.812+0.056	0.702-0.921	0.0044		

[Kappa value (0.0-0.20, poor; 0.21-0.40, fair; 0.41-0.60, moderate; 0.61-0.80, substantial and 0.81-100, perfect); P value showing 0.05 or less is considered as significant variation between the two ELISAs; CI: confidence interval]

#### **Discussion**

Specific and early diagnosis of MAP infection is a major challenge for the control of MAP infection in livestock. Increased production of interferon gamma in the view of MAP infection is the earliest and first immune response (cell mediated), and is slowly replaced by response by humoral immunity (antibody based)<sup>26</sup>. However, few studies established that some elevated level of antibodies is recognized considerably earlier and hence serological ELISA could also be used as early marker for MAP diagnosis in livestock<sup>27</sup>. Selection of the particular early secretary protein (antigen) is challenging, since no single MAP CFPs can detect all the MAP infected animals, especially those in sub-clinical stages of disease 28,29. However, in view of enhanced seroreactivity to MAP infected serum, secretary CFPs are the focus of research for the development of DIVA based diagnostics and vaccines<sup>30,31</sup>.

In the present study, six MAP CFPs known to seroreact with anti-MAP serum were successfully cloned and expressed in bacterial expression system (pET-28a/pET-22b). Yield (12 mg to 200 mg/L of culture) of the purified recombinant proteins was sufficient enough to establish ELISA test. No significant amount of undesired protein band was observed in the purified products suggesting their purity. However, solubility of purified proteins was variable, some of them were soluble proteins (MAP 2168c, MAP ModD, MAP Pep AN and MAP Pep AC) whereas other formed inclusion bodies (MAP 1693c and MAP 85C). Expressed products were detected in Western blot using both anti-His antibody and anti-MAP serum suggesting their in-frame fusion as well as intact immunoreactivity.

Predicted molecular mass of maximum of the recombinant purified CFPs, based on their amino acid composition, matched with observations accomplished by SDS-PAGE assay and immune-blot analysis. Only exception was observed with MAP ModD protein that moved slowly in SDS-PAGE than predicted speed. This unusual movement pattern was probably due to its high proline composition in its peptide sequence, and due to this reason we observed the higher molecular weight band on SDS-PAGE than expected size.

It is rationale to use mixture (cocktail) of MAPspecific immunogenic CFPs that might upsurge the sensitivity and specificity of JD sero-diagnosis. In the development of six individual r-ELISA and cocktail ELISA, we assessed these six recombinant secretory protein antigens (MAP 1693c; MAP 2168c; MAP ModD; MAP 85C; MAP Pep AN and MAP Pep AC) with i-ELISA. c-ELISA showed almost similar sensitivity as shown by i-ELISA (Table 6). Whereas, individual r-CFPs based ELISA could not reach up to the sensitivity of cocktail of six r-CFPs in c-ELISA (Table 6). Results showed that single antigen may not be sufficiently sensitive during entire course of infection and therefore future experiments with cocktails of MAP-specific recombinant protein antigens might improve the test sensitivity and enable detection of infected animals at different stages of JD. Among the six r-CFPs, ModD and 85C revealed high sensitivity of 96.6 and 93.1% in goats and 93.0 and 91.9% in bovine sera, respectively, while, 1693c, Pep AN and Pep AC showed lowest sensitivity of 41.4, 58.6, 63.8% in goats and 89.5, 87.2 and 88.4% in cattle, respectively.

Several other studies supports the sero-diagnostic value of CFPs. Dheenadhayalan *et al.*<sup>32</sup> has reported the immunogenicity of five recombinant antigens (rAgs), MAP 2411; ClpP (MAP 2281c); Ppa (MAP 0435c), MAP 0593c and GreA (MAP 1027c). These antigens were tested with serum from MAP infected (41) and non-infected (41) control sheep where these

antigens, MAP0593c and ClpP, reacted with 58.5% and 46.3% positive serum and 12.1 and 4.9% of the negative serum, respectively. Another study also tested rAgs of Ag85A, Ag85B, Ag85C as well as SOD in ELISA with serum from 60 MAP shedders and 22 non shedders cows<sup>33</sup>. Ag85 complex and SOD were showing high reactivity with serum of MAP infected cows and very low reactivity with non-infected. Conserved proteins of Ag85 complex were observed in all other mycobacterial species<sup>34</sup>. At protein level, there were three Ag85 components were presented in MAP and M. avium subspecies avium (MAA) sharing 99% sequence identity<sup>35</sup>. Bannantine et al.<sup>36</sup> have identified 21 potential coding sequence of MAP by comparative genomic approach. All the 21 genes were successfully cloned and five of the 21 genes were only expressed and getting purified in its denaturing conditions and examined by immune blotting with serum of MAP immunized rabbits and mice found immunogenic. Western blot analysis with serum of 9 healthy and 10 MAP infected cattle showed the same five antigens recognized in context of disease. Leroy et al.37 have identified 25 antigen candidates for diagnosis of MAP infection. They tested diagnostic potential of 5 antigens from 25 antigen candidates in an ELISA assay with small panel of serum collected from fields and their mixture (all three antigens) were contested in performance with other assays available commercially, getting a diagnostic specificity of 97.9% and sensitivity of 94.7%. Mon et al.<sup>28</sup> expressed 54 proteins by recombinant technology and spotted on nitrocellulose membrane and gave exposure with serum of 25 MAP infected, 10 healthy and 8 M. bovis infected animals. Out of 54, only 7 MAP antigens (MAP 2513, 1693, 2020, 0038, 1272, 0209c and Map 0210c) reacted with serum of MAP infected animals. Developed ELISA using Cocktail of these antigens reacted with 18 serums of the 25 MAP infected animals and did not showed any cross-reactivity with healthy animals and showed low reactivity with serum from diseased cattle from bovine tuberculosis. Gupta et al. 16 reported that most of the CFPs governs low molecular mass (>70 kDa) could be used as potential targets for development of new diagnostics for paratuberculosis carrying high specificity and improved sensitivity in comparison to crude protoplasmic antigens (PPA) prepared by sonication of whole cell. Hatamifar et al.<sup>38</sup> have developed a secretary proteins based ELISA and evaluated it's sensitivity and specificity as 70 and 100%, respectively, which can detect the infection in its early stage. In another study, a recombinant MAP 1637c

(r20.8) protein was used in development of an ELISA for the diagnosis of bovine paratuberculosis, reported high specificity and sensitivity when compared with the PPD skin test<sup>39</sup>. Facciuolo *et al.*<sup>40</sup> have identified four MAP specific proteins were uniquely reacted with serum from MAP infected cows. They demonstrated the specificity of these proteins by calf and uninfected cow sera containing no detectable antibodies to these antigens. So, above reports given by other authors also suggest the recombinant CFPs incorporated single multiplex (cocktail) ELISA to be valuable for detection of the disease at all stages of infection in animals<sup>16</sup>.

#### Conclusion

Six genes encoding MAP CFPs were cloned, expressed, purified and evaluated in ELISA to detect anti-MAP antibodies in infected animals. While sensitivity of individual CFP-based ELISA was low, the ELISA developed by cocktail of six recombinant CFPs antigen had high sensitivity and it was comparable to whole cell protoplasmic i-ELISA. Our study shows c-ELISA has comparable specificity than i-ELISA and may have better specificity to evaluate further against gold standard test and provide a better understanding on its diagnostic potential over i-ELISA for the diagnosis of JD and it's applicability in region where infection is widespread.

#### **Ethical approval**

Central Institute for Research on Goats. Makhdoom, Mathura ethical committee chaired by Member Secretary, Institutional Animal Ethics committee (IAEC) and The Committee for the Purpose of Control and Supervision of Experiments on Animals (CPCSEA), New Delhi has approved works were performed under Indian Council of **Project** Medical Research grant number 5/8/5/28/TF/2013/ECD-I], ICMR, New Delhi, India under reference number IAEC/CIRG/16-17 dated 12.05.2016 and confirmed that this project do not have any ethical issue. Serum samples were collected/ received only for laboratory analysis. We have avoided unnecessary pain and suffering of the animals. Samples were not collected from endangered or protected species.

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#### **Conflict of interest**

Authors declare no conflict of Interest.

#### References

- 1 Tiwari A, VanLeeuwen JA, McKenna SLB, Keefe GP & Barkema HW, Johne's disease in Canada: Part I: Clinical symptoms, pathophysiology, diagnosis, and prevalence in dairy herds. *Can Vet J*, 47 (2006) 874.
- 2 Singh SV, Singh PK, Singh AV, Sohal JS, Kumar N, Chaubey KK, Gupta S, Kumar A, Bhatia AK, Srivastav AK & Dhama K, Bio-load and bio-type profiles of *Mycobacterium avium* subspecies paratuberculosis infection in the farm and farmer's herds/flocks of domestic livestock: a 28 years study (1985-2013). *Transbound Emerg Dis*, 61 (2014) 43.
- 3 Chaubey KK, Gupta RD, Gupta S, Singh SV, Bhatia AK, Jayaraman S, Kumar N, Goel A, Rathore AS, Sahzad, Sohal JS, Stephen BJ, Singh M, Goyal M, Dhama K & Derakhshandeh A, Trends and advances in the diagnosis and control of paratuberculosis in domestic livestock. *Vet Q*, 36 (2016) 203.
- 4 Singh SV, Singh AV, Gupta S, Rajindran AS, Swain N, Singh PK, Singh H, Sohal JS & Kumar N, Interspecies sharing of 'Indian Bison Type', a novel predominant genotype of *Mycobacterium avium* subspecies *paratuberculosis* between naturally infected and endemic flocks of Bharat Merino sheep and a colony of rabbits (*Oryctolagus cuniculus*) raised on the same ecosystem in South India. *Res Rev J Life Sci*, 2 (2012) 1.
- 5 Kumar S, Singh SV, Singh AV, Singh PK, Sohal JS & Maitra A, Wildlife (Boselaphus tragocamelus)\_small ruminant (goat and sheep) interface in the transmission of 'Bison type' genotype of Mycobacterium avium subspecies paratuberculosis in India. Comp Immunol Microbiol Infect Dis, 33 (2010) 145.
- 6 Singh SV, Singh AV, Singh PK, Kumar A & Singh B, Molecular identification and characterization of Mycobacterium avium subspecies paratuberculosis in free living non-human primate (Rhesus macaques) from North India. Comp Immunol Microbiol Infect Dis, 34 (2011) 267.
- 7 Chaubey KK, Singh SV, Gupta S, Singh M, Sohal JS, Kumar N, Singh MK, Bhatia AK & Dhama K, Mycobacterium avium subspecies paratuberculosis an important food borne pathogen of high public health significance with special reference to India: an update. Vet Q, 37 (2017) 282.
- 8 Chaubey KK, Gangwar N, Pawaiya RS, Jatav GP, Sohal JS, Singh SV, Singh M, Gupta S, Kumaresan G, Kumar N & Jayaraman S, Evaluation of newly developed 'six recombinant secretary proteins based 'cocktail ELISA' and 'whole cell lysate' based 'indigenous ELISA' and tissue microscopy' with 'Gold standard' histo-pathology for the

- diagnosis of Johne's disease in slaughtered goats and buffaloes. *Comp Immunol Microbiol Infect Dis*, 66 (2019) 10138.
- 9 Chaubey KK, Singh SV & Bhatia AK, Comparison of 'Indigenous ELISA kit' with 'Ethanol Vortex ELISA kit (USA)' for the detection of *Mycobacterium avium* subsp. *paratuberculosis* infection in cattle population in 'Reverse Ice-burg' environment: In search of a 'globally relevant kit'. *Indian J Exp Biol*, 56 (2018) 279.
- 10 Cho D & Collins MT, Comparision of the proteosomes and antigenicities of secreted and cellular proteins produced by mycobacterium paratuberculosis. Clin Vaccine Immunol 13 (2006) 1155.
- 11 Cho D; Sung N & Collins MT, Identification of proteins of potential diagnostic value for bovine paratuberculosis. *Proteomics*, 6 (2006) 5785.
- 12 Wadhwa A, Hickling GJ & Eda S, Opportunities for improved serodiagnosis of human tuberculosis, bovine tuberculosis, and paratuberculosis. *Vet Med Intern*, 2012 (2012) 674238.
- 13 Shin SJ, Cho D & Collins MT, Diagnosis of bovine paratuberculosis by a novel enzyme-linked immunosorbent assay based on early secreted antigens of *Mycobacterium avium* subsp. *Paratuberculosis*. *Clin Vaccine Immunol*, 15 (2008) 1277.
- 14 Willemsen PT, Westerveen J, Dinkla A, Bakker D, van Zijderveld FG & Thole JE, Secreted antigens of Mycobacterium avium subspecies paratuberculosis as prominent immune targets. Vet Microbiol, 114 (2006) 337.
- 15 Pradenas M, Jara MC, Hernandez N, Zambrano A, Collins MT & Kruze J, Antibody recognition to secreted proteins of *Mycobacterium avium* subsp. *paratuberculosis* in sera from infected ruminants. *Vet Microbiol*, 138 (2009) 378.
- 16 Gupta S, Chaubey KK, Singh SV, Bhatia AK, Kumar N, Goel A, Sachan TK, Rawat KD, Sohal JS & Dhama K, Immunoreactivity to culture filtrate proteins of *Mycobacterium avium* subspecies *paratuberculosis* in naturally infected goat and sheep sera. *Adv Anim Vet Sci*, 3 (2015) 347.
- 17 Singh N, Singh SV, Gupta VK, Sharma VD, Sharma RK & Katoch VM, Isolation and identification of *Mycobacterium paratuberculosis* from naturally infected goatherds in India. *Indian J Vet Pathol*, 20 (1996)104.
- 18 Cho D, Shin SJ, Talaat AM & Collins MT, Cloning, expression, purification and serodiagnostic evaluation of fourteen *Mycobacterium paratuberculosis* proteins. *Protein Expr Purif*, 53 (2007) 411.
- 19 Chaubey KK, Singh SV & Bhatia AK. Recombinant secretory proteins based new 'Cocktail ELISA' as a marker assay to differentiate infected and vaccinated cows for Mycobacterium avium subspecies paratuberculosis infection. Indian J Exp Biol, 57 (2019) 967.
- 20 Chaubey KK, Kumaresan G, Gupta RD, Singh SV, Bhatia AK, Rathore AS, Gupta S & Singh M, Molecular Diversity and Homology in Six CFPs Genes in the Novel Bio-Type, 'Indian Bison Type' of *Mycobacterium avium* subspecies paratuberculosis of Goat Origin vs. other Biotypes. Mycobact Dis, 8 (2018) 267.
- 21 Bradford MM, A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the

- principle of protein-dye binding. Anal Biochem, 72 (1976) 248.
- 22 Singh SV, Audarya SD, Singh M, Stephen BJ, Chhabra D, Chaubey KK, Gupta S, Sahzad, Pachoori A, Jayaraman S, Aseri GK, Sohal JS, Bhatia AK & Dhama K, Development of new 'Indigenous Dot-ELISA Kit' as sensitive field based herd screening test for the diagnosis of john's disease in the domestic buffalo population. *Asian J Anim Vet Adv*, 11 (2016) 1.
- 23 Singh SV, Singh AV, Singh PK, Gupta VK, Kumar S & Vohra J, Sero-prevalence of paratuberculosis in young kids using Bison type, *Mycobacterium avium* subsp. paratuberculosis antigen in plate ELISA. Small Rumin Res, 70 (2007) 89.
- 24 Sharma G, Singh SV, Sevilla I, Singh AV, Whittington RJ, Juste RA, Kumar S, Gupta VK, Singh PK, Sohal JS & Vihan VS, Evaluation of indigenous milk ELISA with m-culture and m-PCR for the diagnosis of Bovine Johne's disease (BJD) in lactating Indian dairy cattle. *Res Vet Sci*, 84 (2008) 30
- 25 Collins MT, Interpretation of a commercial bovine paratuberculosis enzyme-linked immune-sorbent assay by using likelihood ratios. *Clin Vaccine Immunol*, 9 (2002) 1367.
- 26 Subharat S, Shu D, Neil Wedlock D, Price-Carter M, de Lisle GW, Luo D, Collins DM & Buddle BM, Immune responses associated with progression and control of infection in calves experimentally challenged with *Mycobacterium avium* subsp. paratuberculosis. Vet Immunol Immunopathol, 49 (2012) 225.
- 27 Begg DJ, De Silva K, Carter N, Plain KM, Purdie A & Whittington RJ, Does a Th1 over Th2 dominancy really exist in the early stages of *Mycobacterium avium* subspecies paratuberculosis infections? *Immunobiol*, 216 (2011) 840.
- 28 Mon M, Viale M, Baschetti G, Pinedo FA, Gioffre A, Traveria G, Willemsen P, Bakker D & Romano MI, Search for *Mycobacterium avium* subspecies *paratuberculosis* antigens for the diagnosis of paratuberculosis. *Vet Med Int*, 2012 (2012) 1.
- 29 Mikkelsen H, Aagaard C, Nielsen SS & Jungersen G, Review of *Mycobacterium avium* subsp. *paratuberculosis* antigen candidates with diagnostic potential. *Vet Microbial* 152 (2011) 1.
- 30 Deb R & Goswami PP, Coexpression of PPE 34.9 antigen of *Mycobacterium avium* subsp. *paratuberculosis* with murine interferon gamma in HeLa cell line and study of their immunogenicity in murine model. *Biotechnol Res Int*, 2011 (2011) 1.

- 31 Malamo M, Sakoda Y, Ozaki H & Kida H, Development of ELISA to detect antibodies specific to *Mycobacterium avium* subsp. *paratuberculosis* with truncated 34 kDa proteins. *Jpn J Vet Res*, 54 (2006) 99.
- 32 Dheenadhayalan V, Shin KS, Chang CF, Chang CD, Wang SJ, McDonough S, McDonough P, Stehman S, Shin S, Torres A & Chang YF, Cloning and characterization of the genes coding for antigen 85A, 85B and 85C of Mycobacterium avium subsp. Paratuberculosis. DNA Seq, 13 (2002) 287.
- 33 Bannantine JP, Stabel JR, Lamont EA, Briggs RE & Sreevatsan, S, Monoclonal antibodies bind a SNP-sensitive epitope that is present uniquely in *Mycobacterium avium* subspecies *paratuberculosis*. Front Microbiol, 2 (2011) 1.
- 34 Gumber S, Taylor DL & Whittington RJ, Evaluation of the immunogenicity of recombinant stress-associated proteins during Mycobacterium avium subsp. Paratuberculosis infection: implications for pathogenesis and diagnosis. Vet Microbiol 137(2009) 290.
- 35 Rosseels V, Marche S, Roupie V, Govaerts M, Godfroid J, Walravens K & Huygen K, Members of the 30- to 32-kilodalton mycolyl transferase family (Ag85) from culture filtrate of *Mycobactelium avium* subsp. *paratuberculosis* are immunodominant Th1-type antigens recognized early upon infection in mice and cattle. *Infect Immun*, 74 (2006) 202.
- 36 Bannantine JP, Hansen JK, Paustian ML, Amonsin A, Li LL, Stabel JR & Kapur V, Expression and immunogenicity of proteins encoded by sequences specific to *Mycobacterium avium* subsp. *Paratuberculosis*, *J Clin Microbiol*, 42 (2004) 106
- 37 Leroy B, Roupie V, Noel-Georis I, Rosseels V, Walravens K, Govaerts M, Huygen K & Wattiez R, Antigen discovery: a postgenomic approach to paratuberculosis diagnosis. *Proteomics*, 7 (2007) 1164.
- 38 Hatamifar M, Mosavari N & Kazemi J, Designing of Indirect ELISA system using secreted antigens of *Mycobacterium avium* subsp. *paratuberculosis* for Diagnosis of paratuberculosis. *Iran J Med Microbiol*, 11 (2017) 26.
- 39 Goswami PP, Chand G, Prasad NS, Deb R & Basagoudanavar SH, Recombinant 20.8-kDa protein of *Mycobacterium avium* subsp. *paratuberculosis*-based sero-diagnosis of paratuberculosis. *3 Biotech*, 7 (2017) 156.
- 40 Facciuolo A, Kelton DF & Mutharia LM, Novel Secreted Antigens of *Mycobacterium paratuberculosis* as Serodiagnostic Biomarkers for Johne's Disease in Cattle. *Clin Vaccine Immunol*, 20 (2013) 1783.