

## Phage display-based identification and potential diagnostic application of novel antigens from subsp. small colony type

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2	novel antigens from Mycoplasma mycoides subsp. mycoides small colony type.						
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#### 29 Abstract

30

31 Contagious Bovine Pleuropneumonia caused by Mycoplasma mycoides subsp. 32 *mycoides* small colony type is a respiratory disease of considerable economic importance in sub-Saharan Africa; control of the disease in Africa is hampered by 33 34 diagnostic tests which are suited for herd-level but not for individual animal 35 diagnostics. In the work presented we identified 22 potential immunogenic 36 antigens of the Kenyan outbreak strain B237 by using phage display technology. 37 We determined the relative strength of immunogenicity, the discriminatory 38 capacity between bovine positive and negative sera, and the cross reactivity with 39 rabbit hyperimmune sera directed against 15 different mycoplasmal species. The 40 three best-performing antigens, a conserved hypothetical protein (MSC\_0636), a 41 glycosyl transferase (MSC\_0108), and an acyl carrier protein phosphodiesterase 42 (MSC\_0029) were considered candidate diagnostic proteins. They were expressed 43 as GST-fusion proteins in E. coli, purified, and used in an ELISA as solid phase 44 antigens. The diagnostic potential of the recombinant antigens was tested using 45 the sera of ten experimentally infected animals and six control animals. This 46 prototype test resulted in 100 % diagnostic sensitivity and specificity. In 47 comparison, the complement fixation test and the competitive ELISA performed 48 with a diagnostic sensitivity of 70 % and 60 %, respectively. 49

50 *Keywords: Mycoplasma mycoides* subsp. *mycoides* small colony type; phage
51 display; diagnostic antigens

#### 52 **1. Introduction**

53

54 Contagious Bovine Pleuropneumonia (CBPP) is a highly infectious respiratory 55 disease in cattle caused by Mycoplasma mycoides subsp. mycoides small colony 56 type (*MmmSC*) and solely transmitted by animal to animal contact. The disease is 57 of major economic importance in subsaharan Africa (FAO, 2003) with a constant 58 risk of reintroduction into other parts of the world. Diagnosis is primarily based 59 on serology currently done by complement fixation test (CFT), competitive 60 ELISA (cELISA) (Le Goff and Thiaucourt, 1998) and a lipoprotein Q-based 61 ELISA (Bruderer et al., 2002). Both ELISA tests closely correlate with the CFT 62 reported to have a diagnostic sensitivity of 64 to 70 % and a diagnostic specificity 63 of 98 % (Amanfu et al., 2000; Marobela-Raborokgwe et al., 2003). For a reliable 64 diagnosis of disease on the individual animal level the diagnostic sensitivity needs to be improved without losing specificity. To achieve this goal, novel diagnostic 65 66 antigens need to be identified. Recently it has been shown that this goal can be achieved effectively using a non-67 68 lytic phage display approach based on phage M13 and designated as pHORF-69 system (Kügler et al., 2008); it improves the panning and screening process 70 compared to the lytic lambda-based phage display system (Beghetto et al., 2009; 71 March et al., 2006; Kügler et al., 2008). Furthermore, the included open reading 72 frame selection and the packaging into "Hyperphage", a helperphage not encoding 73 pIII protein on its genome, improve the subsequent panning process (Hust et al., 74 2006; Kügler et al., 2008; Soltes et al., 2007; Rondot et al., 2001). 75 In the work presented, the phage M13-based pHORF system was used to identify 76 novel immunogenic epitopes of *Mmm*SC which are potentially suitable not only 77 for herd-level but also for individual animal diagnostics.

3

#### 78 2. Materials and Methods

- 79
- 80 2.1 Bacterial strains, phages, growth conditions, plasmids and sera
- 81
- 82 The strains, plasmids, primers and sera used in this work are listed in Table 1.
- 83 Escherichia coli strains were cultured in Luria-Bertani medium supplemented
- 84 with the appropriate antibiotics (ampicillin,  $100 \mu g/l$ , kanamycin,  $30 \mu g/l$ ). The
- 85 *Mmm*SC field isolate B237 was grown as stationary culture for three days in
- 86 modified PH medium (Kirchhoff and Rosengarten, 1984), supplemented with
- 87 20 % equine serum (WDT, Garbsen, Germany) at 37 °C in a 5 % CO<sub>2</sub> atmosphere.
- 88 The 10 bovine anti-MmmSC sera used in this study were derived from
- 89 experimentally infected cattle (Jores et al., 2008). A positive control serum was
- 90 obtained by pooling equal volumes of these sera. The six negative sera used were
- 91 obtained from uninfected cattle from a CBPP free region within Kenya. A
- 92 negative control serum was generated by pooling equal volumes of these sera. The
- 93 rabbit sera directed against different mycoplasmal species had been raised

94 previously against the respective whole cell bacterins.

- 95
- 96
- 97 2.2 Construction of an MmmSC genomic phage display-library and enrichment of98 open reading frames
- 99
- 100 Site-specific mutation of the phagemid vector pHORF3 (Kügler et al., 2008) was
- 101 performed using primers oMHLacZPro\_f, oMHORF3\_pme\_rX1 and
- 102 oMHORF3\_pme\_rXX2 (Table 1) such that the unique PmeI site used for insertion
- 103 of foreign DNA is shifted by 1 and 2 bp resulting in plasmids pHORF3X and

104	pHORF3XX, respectively (Fig. 1). Genomic DNA of the MmmSC African field
105	strain B237 was isolated as described previously (Meens et al., 2006) and digested
106	using five different blunt end-cutting restriction endonucleases (AluI, PsiI, SspI,
107	HpyCH4V, RsaI). DNA fragments with a size of 100-1000 bp were isolated after
108	separation in an agarose gel using the Qiagen PCR DNA and gel band purification
109	kit (Qiagen, Hilden, Germany) according to the manufacturer's instructions, and
110	ligated into PmeI-restricted vectors pHORF3, pHORF3X and pHORF3XX (Fig.
111	1). The genomic MmmSC phage library was constructed as described previously
112	(Kügler et al., 2008).
113	
114	
115	2.3 Selection of phage encoding immunogenic oligopeptides of MmmSC and
116	production of individual phage clones for screening
117	
118	The panning of phage presenting immunogenic peptides on the surface was
119	performed using the bovine positive control serum pre-absorbed to Hyperphage in
120	order to prevent unspecific reactivity with antibodies directed against phage-
121	surface epitopes. The preabsorbed serum was coupled to MaxiSorb <sup>®</sup> 96 well
122	microtitre plates (Nunc, Wiesbaden, Germany) via rabbit anti-bovine IgG (Sigma,
123	Deisenhofen, Germany), and four successive panning rounds were performed as
124	previously described (Kügler et al., 2008). Individual clones were picked from the
125	titration plates of the last panning round.
126	For production of individual phage clones, polypropylene 96-well U-bottom
127	plates (Greiner bio-one, Frickenhausen, Germany) containing 175 $\mu$ l of 2 x TY-
128	GA medium (1 % Bacto-Yeast Extract, 1.6 % Bacto Tryptone, 0.05 % NaCl; 100
129	mM glucose, 100 µg/ml ampicillin) per well were inoculated with single clones

130	and incubated at 37 °C with constant shaking at 250 rpm overnight. Cultures were
131	expanded to 1.5 ml of 2 x TY-GA medium, infected with 5 x $10^9$ cfu Hyperphage
132	per tube and incubated at 37 °C without shaking for 30 min, followed by 30 min
133	incubation with shaking at 250 rpm. Cells were centrifuged at 3,220 x g for
134	10 min and the supernatants were discarded. The bacterial pellets were
135	resuspended in 1.5 ml of 2 x TY medium containing ampicillin (100 $\mu$ g/ml) and
136	kanamycin (30 $\mu$ g/ml) and incubated at 30 °C at 250 rpm overnight for phage
137	production.
138	In order to confirm the antigenicity of individual phage clones PolySorb <sup>®</sup> 96 well
139	microtitre plates (Nunc) were coated overnight with 10 $\mu$ l (1 x 10 <sup>10</sup> pfu/ml) of
140	individual phage in 100 $\mu$ l of carbonate buffer (100mM NaHCO <sub>3</sub> , 46 mM
141	Na <sub>2</sub> CO <sub>3</sub> , pH 9.6). The ELISA was developed using the positive control serum
142	(diluted 1:100), goat anti-bovine IgG conjugated with horseradish peroxidase
143	(HRP; Jackson ImmunoResearch Laboratories, Dianova, Hamburg, Germany) as a
144	conjugate and 2,2-azino-di-[3-ethylbenzithiazoline sulfonate] (ABTS; Roche
145	Diagnostics, Mannheim, Germany) as a substrate. The staining reaction was
146	stopped by adding 100 $\mu l$ of 50% (v/v) methanol. The absorbance at 405 nm and
147	scattered light at 492 nm were measured using a Sunrise <sup>®</sup> microtiter plate reader
148	(Tecan, Crailsheim, Germany).
149	

- 150

151 2.4 Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and
152 Western blot

153

154 SDS-PAGE was performed following standard procedures using a Protean II

155	Minigel system (BioRad Inc, Munich, Germany). Western blotting of the SDS-
156	PAGE minigels was done using the Mini Trans-Blot® system (BioRad). The
157	bovine positive and negative control serum and a rabbit anti-GST serum were
158	used as primary antibodies; blots were developed using alkaline phosphatase-
159	conjugated goat anti-bovine or goat anti-rabbit IgG antibody (Jackson
160	ImmunoResearch Laboratories) and BCIP (5-bromo-4-chloro-3-indolyl
161	phosphate) and NBT (nitroblue tetrazolium) as substrate.
162	

163

164 2.5 Preparation of recombinant fusion proteins and ELISA set-up

165

166 The MmmSC genes encoding immunogenic epitopes as assessed by ELISA of 167 individual phage clones were identified by database analysis. The respective parts 168 of the genes located between two TGA codons were amplified by PCR using 169 primer pairs with an EcoRI- (upstream primer) or NotI-restriction endonuclease 170 site (downstream primer) at their 5'ends (Table 1). The corresponding PCR 171 fragments were digested with EcoRI and NotI and ligated into pGEX-5x-2, 172 resulting in plasmids pMSC0029-500, pMSC0108-500 and pMSC0636-500. 173 Recombinant fusion proteins were prepared as inclusion bodies as described 174 previously (Gerlach et al., 1992) and dissolved in guanidine hydrochloride (4 M). 175 To remove small-sized contaminating material, the solution was washed twice in 176 guanidine hydrochloride (4M) using a 10 kDa cut-off (MicroCentricon 10 system, 177 Amicon, Beverly, Mass., U.S.A.). Using BSA as reference the concentration of the 178 purified protein (desalted by precipitation with trichloroacetic acid [TCA]) was 179 then determined by SDS PAGE and staining with Coomassie blue. The structure

180 and immunogenicity of the fusion proteins was confirmed by Western blot 181 analysis using positive and negative control sera as well as an anti-GST serum. 182 For use in an ELISA, fusion proteins in guanidine hydrochloride were pipetted in 183 a petri dish, diluted by adding the appropriate volume of carbonate coating buffer 184 with a 10 ml pipette, and used as solid phase antigens on a PolySorb 96 well 185 microtiter plate (Nunc) coating 100 µl volumes per well overnight at 4 °C. The 186 coating concentration was optimized by checkerboard titration of the antigen, 187 development of the ELISA using 2-fold serial dilutions of the positive and 188 negative control sera, goat-anti-bovine peroxidase, and ABTS, and analysis of the 189 log-log-transformed data. Prior to use in the ELISA sera were absorbed with 190 whole-cell lysate of *E. coli*-pGEX-5x-2 transformants for 1 h at room temperature 191 in order to remove antibodies directed against contaminating E. coli components 192 in the semi-purified antigen preparation. 193

194

- 195 3. Results
- 196

197 3.1 Identification of immunogenic peptides

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After performing four panning rounds with the positive control serum a total of 147 clones (49 from each of the three libraries based on vectors with the cloning site in three different reading frames with respect to the 3' pIII protein-encoding ORF) were sequenced. One hundred and thirty four clones contained analyzable sequence and showed an ORF consisting of a *Mmm*SC-derived genomic DNA fragment in frame with both the *pelB* signal sequence and the PIII proteinencoding ORF. Upon NCBI Blast analysis (http://www.ncbi.nlm.nih.gov/blast/

206 <u>Blast.cgi</u>) 22 different clones were identified with each clone present at least twice

207 in the random selection of sequenced clones (Table 2).

208

209

210 3.2 Selection of clones with diagnostic potential

211

212 The individual clones were amplified and the relative strength of serum reactivity 213 was determined by phage ELISA. Eight clones showing a reactivity of more than 214 30 OD% with respect to MmmSC whole cell lysate (Table 2) were considered to 215 be of potential diagnostic value and were subjected to further investigation. The 216 clones were individually amplified, and the discrimination between positive and 217 negative control serum was investigated by ELISA. For six of the eight clones 218 (MSC\_0029, 0108, 0636, 0656, 1009 and 1062) at least a 3-fold difference in titre was observed between the positive and the negative control serum (data not 219 220 shown). These six clones were tested with respect to their discriminatory potential 221 using the ten individual positive sera and the six individual negative sera 222 contained in the respective control sera. Four of the clones (MSC\_029, 108, 636 223 and 1062) reacted with all individual positive sera; no overlap in OD%-values was 224 seen between positive and negative sera (data not shown). 225 These four clones were investigated for cross reactivity to other mycoplasmal 226 species using rabbit hyperimmune sera previously raised against 15 different 227 mycoplasmal species. All four clones (MSC\_0029, MSC\_0108, MSC\_0636, and 228 MSC 1062) were highly positive with the hyperimmune serum directed against 229 MmmSC. Two clones showed clear cross-reactivity (i.e. more than 1.5-fold of the 230 average reactivity with the negative control serum) with the serum directed against M. capricolum subsp. capricolum; for clone MSC\_1062 an additional 231

232 cross-reactivity against A. modicum and M. bovis was observed (Table 3) and, 233 therefore, this clone was excluded from further analyses. The three remaining 234 clones (MSC\_0029, MSC\_0108 and MSC\_0636) were expressed as GST fusion 235 proteins. 236 237 3.3 Potential of recombinant GST-fusion proteins as diagnostic antigens 238 239 240 All fusion proteins had a molecular mass corresponding to the calculated masses 241 of 32.5, 32.0, and 45.0 kDa, respectively. Approximately 90% of the fusion 242 protein preparations consisted of recombinant antigen (Fig. 2A). The proteins 243 reacted strongly positive with both the anti-GST and the positive control serum 244 but did not show any reaction with the negative control serum (Fig. 2B-D). 245 The purified fusion proteins, upon checkerboard titration performed to determine 246 the suitable coating concentration of  $2.5 \,\mu$ g/ml, were used as solid-phase antigens. 247 In a first ELISA, using rabbit hyperimmune sera previously raised against 248 *Mmm*SC, a 32-times higher titre of the positive versus the negative serum was 249 observed (data not shown) thereby showing suitability of the antigens in principal. 250 In a second ELISA, serial two-fold dilutions of the bovine positive and negative 251 control sera were used after absorption against a lysate of E. coli pGEX-5x-2 252 transformants. The slope of the regression curve of the log-log transformed data 253 obtained with the positive control serum was determined to be 0.72 thereby 254 confirming the suitability of the coating concentration; in addition, an at least 255 three-fold difference in titre and a 2.5-fold difference of OD%-values was 256 observed between positive and negative control serum thus confirming the discriminatory power of the antigens. 257

258 Using these antigens and coating concentrations the ELISA was performed three

times in duplicate for 1:500 dilutions of the individual positive and negative sera.

260 The intra-assay variation for all three ELISAs was determined to be below

261 5 OD% and the inter-assay variation below 20 OD%. Applying a cut-off range of

262 50 OD% all three ELISAs resulted in an analytical sensitivity and specificity of

263 100 % as compared to a 70 % sensitivity for the complement fixation test and

264 60 % for the competitive ELISA commercially available (Fig. 3).

265

266

#### 267 **4. Discussion**

268

269 The control of the continuous spread of CBPP in sub-Saharan Africa is severely 270 limited by the lack of sufficiently sensitive and specific diagnostic tests. In an 271 attempt to improve the diagnostic possibilities we set out to identify novel 272 MmmSC-derived antigens using a non-lytic M13-based phage display system 273 (Kügler et al., 2008). The construction of phage display libraries in three different 274 reading frames using five different restriction endonucleases with a 4mer 275 recognition sequence was considered to result in a reasonable coverage of the 276 MmmSC genome although the three vectors facilitate the generation of a 277 functional pIII fusion protein only for one of the three possible reading frames 278 (Fig. 1). Thus, in silico analysis of the MmmSC PG1 genome revealed that the restriction enzyme digests resulted in a total of 9,178 restriction endonuclease 279 280 fragments of 100 to 1000 bp in length. In total these fragments cover 3,413,598 bp 281 which corresponds to 2.8 times the *Mmm*SC genome size. The finding that each 282 peptide was present at least twice in the pool of 147 clones analyzed implied that 283 the number of panning rounds was sufficient to eliminate non-specific binders

284	(Eshaghi et al., 2005; Kügler et al., 2008). When comparing the open reading
285	frames (ORFs) identified with the results of a previous study by March et al.
286	(March et al., 2006) only ORF MSC_0266 encoding a pyruvate dehydrogenase
287	was identified in both studies. This is likely due to the different methodologies
288	used; thus, March et al. (March et al., 2006) used a lytic bacteriophage Lambda
289	ZAP Express vector with insert sizes of 6 to 8 kb and screened using rabbit IgG
290	antibodies.
291	Some of the initially identified antigens are intracellular proteins. This is in
292	accordance with many other studies, where intracellur proteins were found to be
293	immunogenic (Kügler et al., 2008; March et al., 2006; Delvecchio et al., 2006;
294	Cho et al., 2008; Kodzius et al., 2003). A possible explanation for this
295	phenomenon could be the release of intracelluar proteins by cell lysis or by
296	programmed cell death (Lewis, 2000).
297	Four of 22 identified antigens were consistently recognized by all positive sera
298	and showed sufficient discrimination of all negative sera. Using rabbit
299	hyperimmune sera directed against 15 other bovine mycoplasmal species these
300	four antigens were tested for serological cross-reactivity which might limit
301	diagnostic specificity. The cross-reactivity of all four antigens with M. capricolum
302	subsp. capricolum appeared acceptable since sequence analysis of the 16S rRNA
303	(Pettersson et al., 1994, 1996) and of five conserved protein-coding sequences
304	(Manso-Silván et al., 2007) showed a high sequence similarity and, consequently,
305	a close phylogenetic relationship between MmmSC, M. mycoides subsp. capri
306	and M. capricolum subsp. capricolum, and the other members of the "mycoides-
307	cluster". In addition, it has been reported that M. capricolum subsp. capricolum
308	can cause CBPP-like disease in cattle (Ajuwape et al., 2003). Acholeplasma
309	modicum and M. bovis, however, are not members of the "mycoides-cluster";

310 since serological cross-reactivity particularly with the frequently occurring bovine 311 pathogen *M. bovis* is unacceptable for a diagnostic test, ORF MSC 1062 was 312 excluded from further analyses. Immunoblot analysis of the three remaining 313 antigen-GST fusion proteins showed a strong reaction particularly of antigens 314 MSC\_0108 and MSC\_0636 (Fig. 2) thereby confirming the initial panning results 315 where these two peptides had elicited the highest reactivity. 316 The testing of the cohort sera (16 infected and six non-infected) resulted in a 317 diagnostic sensitivity and specificity of 100 % each in comparison to a diagnostic 318 sensitivity of 70 % for the CFT and 60 % for the cELISA (Fig. 3). This result 319 implies that all three antigens are potentially suitable to improve the reliability of current serological testing which is primarily hampered by low diagnostic 320 321 sensitivity (Marobela-Raborokgwe et al., 2003). However, the cohort sera used 322 were by no means representative; in addition, MmmSC is known to show a high 323 antigenic variability (Gonçalves et al., 1998; Persson et al., 2002) and different 324 breeds of cattle may show a distinct serological response to MmmSC infection. 325 Therefore, in order to support the potential suitability of these antigens for field 326 application, additional sera from experimentally infected cattle as well as field 327 sera from different geographical regions need to be investigated in the future.

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## 428 Figure captions:

429

430	Fig. 1. Cloning vector pHORF3 and derivatives. (A) Schematic of the phage
431	display vector pHORF3. Abbreviations: Lac Pr.: promoter of the bacterial lac
432	operon; RBS: ribosome binding site; <i>pelB</i> : sequence encoding the signal peptide
433	of bacterial pectate lyase, mediating protein secretion into the periplasmic space;
434	ochre: ochre stop codon; amber: amber stop codon; tag: hexahistidine and c-myc
435	tags; terminator: sequence terminating transcription; bla: $\beta$ -lactamase gene for
436	ampicillin resistence; ColE1: bacterial origin of DNA replication; M13 ori:
437	intergenic region of phage f1. (B) Comparision of the PmeI cloning site of
438	pHORF3, pHORF3X and pHORF3XX.
439	
440	Fig. 2. Characterisation of GST fusion proteins. (A) SDS PAGE showing
441	aggregate preparations of recombinant fusion proteins of E. coli transformants
442	containing pMSC0029-500, pMSC0108-500, pMSC0636-500 and the cloning
443	vector pGEX-5x-2 (Lines 1 to 4). (B) Western blot using anti-GST antibody
444	(1:2000), (C) Western blot using the positive control serum obtained from CBPP-
445	infected animals (1:100) and (D) Western blot using the negative control serum
446	(1:100). All three clones pMSC0029-500, pMSC0108-500, pMSC0636-500 were
447	found to strongly react with the positive control serum and not with the negative
448	control serum. In addition, GST showed only minor reactivity with the positive
449	control serum.
450	

- 451 Fig. 3. ELISA results using GST fusion proteins. Sera of ten individual CBPP-
- 452 infected and six uninfected animals were used in the commercial cELISA and in
- 453 ELISAs with the GST-fusion proteins as solid phase antigens. The horizontal line
- 454 indicates the cut-off given for the cELISA.. Solid symbols indicate a deviating
- 455 outcome of CFT test and ELISA for the respective sera. The OD%-values give the
- 456 arithmetic mean of three independent experiments.

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Α

## ACCEPTED MANUSCRIP



## Β

### pHORF3:

3' pelBPmeIHis Tagamber5' gIIIatg gcg gttt|aaac cat cat cat cat cat cat cat cat taggga tcc aaa gat atc aga gctMAV\*HHHH\*GSKDIR

#### pHORF3X:

3' pelBPmeIHis Tagamber5' gIIIatg gcg gtttlaaaca cat cat cat cac cat cat cat taggga tcc aaa gat atc aga gctMAV\*HHHHKDIR

#### pHORF3XX:

3' g	belB		PmeI				Ηİ	is Ta	ag		ambe	er			5' g]	III			
atg	gcg	gtt	t aa	aca	а	cat	cat	cac	cat	cat	cat	tag	gga	tcc	aaa	gat	atc	aga	gct
М	А	V	*			Н	Η	Η	Η	Η	Η	*	G	S	K	D	D	I	R



Fig. 2: Characterization of the GST fusion proteins.



Fig. 3: ELISA results using GST fusion proteins.

### Table 1

### Bacterial strains, plasmids, primers and sera used in this work

Strains, plasmids, primers or sera	Characteristic(s)	Source or Reference			
Strains					
Mmm SC	<i>Mmm</i> SC B237, isolated from an outbreak in Thika/Kenya in 1997	Jores et al., 2008			
E. coli EC-TS40.6	K12 strain	AG Dübel			
<i>E. coli</i> DH5αF′	F' endA1 hsdR17 ( $r_{K}m_{K}$ ) supE44 thi-1 recA1 gyrA (Nal <sup>r</sup> ) relA1 $\Delta(lacZYA-argF)U169$ deoR [ $\Phi$ 80d $lac\Delta(lacZ)M15$ ]	New England Biolabs, Frankfurt, Germany			
M13K07	Stratagene, Amsterdam, The Netherland				
"Hyperphage"	Helperphage carrying a kanamycin resistance determinant and truncated gIII	Rondot et al., 2001			
Plasmids					
pHORF3	pHORF3 used for insertion of foreign DNA	Kügler et al., 2008			
pHORF3X	pHORF3 was modified such that the singular PmeI site used for insertion of foreign DNA is shifted by 1bp, allowing the in-frame fusion with gIII in an additional reading frame	This work			
pHORF3XX	pHORF3 was modified such that the singular PmeI site used for insertion of foreign DNA is shifted by 2bp, allowing the in-frame fusion with gIII in an additional reading frame	This work			
pGEX5x2	<i>E. coli</i> expression vector carrying an bla resistance determinant, for construction of GST fusion proteins	GE Healthcare, Munich, Germany			
pMSC0029-500	plasmid encoding the GST-MSC_0029 fusion protein	This work			
pMSC0108-500	plasmid encoding the GST-MSC_0108 fusion protein	This work			
pMSC0636-500	plasmid encoding the GST-MSC_0636 fusion protein	This work			

### Table 1 - continued

Primers				
MHLacZPro_f	GGCTCGTATGTTGTGTGG; control for phagemids; binds with LacZ promoter	Kügler et al., 2008		
MHgIII_r1	CTAAAGTTTTGTCGTCTTTCC; control for phagemids; binds with <i>pIII</i>	Kügler et al., 2008		
oMHORF3_Pme_rX1	CTTTGGATCCCTAATGATGATGGTGATGA TG; primer for modification of pHORF3 by adding 1bp (pHORF3X)	This work		
oMHORF3_Pme_ rXX2	CTTTGGATCCCTAATGATGATGGTGATGA TGG; primer for modification of pHORF3 by adding 2bp (pHORF3XX)	This work		
oMSC0029A	GATCGAATTCAATGGGGACAACACACTCA ATAT; forward primer to clone MSC_0029	This work		
oMSC0029B	CGATGCGGCCGCTTAAAAAGTTCTAGCAG CTTCTAT; reverse primer to clone MSC_0029	This work		
oMSC0108A	ATAATCGAATTCAATGGATTTCAAGTGCT TATTCAATTCA	This work		
oMSC0108B	C0108B TCCGATGCGGCCGCTCATCAATTAGTTTG; reverse primer to clone MSC_0108			
oMSC0636A	GACGAATTCAAATGGTTTATTATGCAAGA AACAATGAACCG; forward primer to clone MSC_0636	This work		
oMSC0636B	ATATGCGGCCGCTCATGATTTTAATTGTTC TTTTAATTG; reverse primer to clone MSC_0636	This work		
Sera				
Hyperimmune sera	hyperimmune sera raised in rabbits against different mycoplasmal species	FLI, Jena, Germany		
CBPP negative sera	preimmune sera obtained from uninfected cattle	ILRI Nairobi Kenya		
CBPP positive sera	sera from CBPP infection trials, ILRI Nairobi, Kenya	Jores et al., 2008		
PC C				

#### Table 2

Immunogenic peptides identified in *Mycoplasma mycoides* subsp. *mycoides* small colony type (*MmmSC*)

No	Fq <sup>1)</sup>	Genome position	Product from <i>Mmm</i> SC	Insert length (bp)	Locus tag	OD% <sup>2)</sup>
1	3	733268 - 733109	Conserved hypothetical	159	MSC_0636	56.7 %
2	2	127879 - 127937	Glycosyl transferase	59	MSC_0108	51.4 %
3	7	696995 - 697122	ATP dependent protease	128	MSC_0613	43.7 %
4	7	1153378 - 1153506	DNA directed RNA polymerase beta subunit	199	MSC_1009	39.5 %
5	8	32410 - 32351	Acyl carrier protein phosphodiesterase	60	MSC_0029	37.3 %
6	2	1205835 - 1205959	AAA family ATPase	124	MSC_1062	36.9 %
7	6	1032859 - 1032922	Amino acid permease	64	MSC_0908	34.3 %
8	5	750456 - 750516	Conserved hypothetical	60	MSC_0656	32.7 %
9	2	304438 - 304508	Pyruvate dehydrogenase	71	MSC_0266	30.0 %
10	5	218668 - 218814	Oligopeptide ABC trans- porter, component OppA	147	MSC_0184	30.0 %
11	8	159499 - 159643	Hypothetical transmembrane protein	144	MSC_0135	29.0 %
12	4	1160317 - 1160401	Glucose inhibited division protein A	85	MSC_1017	28.3 %
13	2	23668 - 23767	Mannitol-1-phosphate 5- dehydrogenase	99	MSC_0017	26.8 %
14	26	77529 - 77641	Transcriptional regulator; NifR3 family	184	MSC_0068	26.7 %
15	3	940581 - 940693	Conserved hypothetical	112	MSC_0826	25.2 %
16	19	1117031 - 1117172	UDP glucose 4-epimerase	141	MSC_0978	24.5 %
17	5	44289 - 44393	D-Lactate dehydrogenase	105	MSC_0034	24.5 %
18	4	701239 - 701317	Prolipoprotein	78	MSC_0617	23.8 %
19	6	1210760 - 1210841	Conserved hypothetical transmembrane protein	82	MSC_1066	22.3 %
20	4	698434 - 698492	Putative hydrolase of HAD family	69	MSC_0614	22.1 %
21	3	385965 - 386234	tRNA pseudoridine syntase B	269	MSC_0336	21.0 %
22	3	97781 - 98005	Prolipoprotein ; putative phosphate ABC transport	224	MSC_0079	21.0 %

<sup>1)</sup> Frequency of clones among the random sample of 134 clones sequenced.

<sup>2)</sup> The optical density obtained with *Mmm*SC whole cell lysate used as solid phase antigen served as positive control (100 %).

#### Table 3

ELISA cross-reactivity of four clones with rabbit sera directed against 16 different

### Mycoplasma species

Serum raised aganist	culture collection number	OD% +/- standard deviation			
		MSC_029	MSC_0108	MSC_636	MSC_1062
A. axanthum S 743	NCTC 10138	$28\pm8.6$	$26\pm~4.1$	35 ± 2.3	45 ± 1.8
A. laidlawii PG 8	NCTC 10116	$39\pm8.3$	$43\pm0.2$	42 ± 3.5	39 ± 1.7
A. modicum PG 49	NCTC 10134	$32\pm 6.8$	$26\pm0.6$	$32 \pm 0.4$	51 ± 1.2
M. alkalescens PG 51	ATCC 29103	$17\pm0.6$	21 ± 2.1	43 ± 1.3	$41\pm0.5$
M. arginini G 230	NCTC 10129	$21\pm1.6$	$21 \pm 0.3$	26 ± 1.6	$32 \pm 0.3$
M. bovigenitalium PG 11	ATCC 19852	$17 \pm 0.3$	$17 \pm 1.9$	$22 \pm 1.1$	$25\pm0.3$
M. bovirhinis PG 43	NCTC 10118	$36 \pm 4.8$	$40 \pm 2.1$	44 ± 14.9	$44 \pm 3.2$
M. bovis Donetta PG 45	ATCC 25523	44 ± 13.7	$39 \pm 2.5$	$41\pm5.6$	$49 \pm 2.3$
<i>M. bovoculi</i> M 165/69	NCTC 10141	$44 \pm 0.4$	$42 \pm 8.8$	44 ± 7.1	33 ± 2.9
M. californicum ST-6	ATCC 33461	21 ± 2.5	$22\pm0.9$	$32\pm0.6$	$32 \pm 2.2$
<i>M. capricolum</i> subsp. <i>capricolum</i> california kid	NCTC 10154	$45\pm21.7$	$48\pm0.5$	45 ± 1.4	$57 \pm 4.9$
M. gallinarum PG 16	ATCC 19708	$19\pm1.9$	$28\pm2.4$	45 ± 1.1	$44\pm0.3$
M. gatae C 8	ATCC 23392	$30 \pm 4.2$	$21\pm4.0$	$27 \pm 1.5$	$31 \pm 2.0$
M. leachii PG 50	NCTC 10133	$46\pm 6.0$	$27\pm2.0$	33 ± 4.3	$45\pm7.3$
M. mycoides subsp. capri y-goat	NCTC 10706	$30 \pm 1.2$	27 ± 2.3	$30\pm0.5$	34 ± 3.4
MmmSC PG 1	NCTC 10114	$100\pm4.6$	$100 \pm 12.1$	$100\pm2.5$	$100 \pm 8.9$
Negative serum	-	$27\pm3.0$	$30 \pm 2.6$	$29\pm1.0$	$36 \pm 0.7$