THE USE OF HIGH PERFORMANCE LIQUID CHROMATOGRAPHY IN SEPARATION AND PURIFICATION OF LP 64 KDA BAND OF MYCOPLASMA GALLISEPTICUM IN LARGE QUANTITIES

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SUMMARY

High performance liquid chromatography (HPLC) size exclusion technique (SEC) was used for separation and purification of the LP 64 kDa band of Mycoplasma gallisepticum (MG) in large quantities. We found that the use of 210 nm was the best wave length for isolation and gave highest peak and the best resolution. The use of 200 mM of NaH2PO4 as a mobile phase improved chromatographic profile. Injection volume of 200 µl resulted in the highest peak and the best resolution. The flow rate of 5ml /min increased the peak height and profile resolution. The use of pH 5.7 for the Mycoplasma gallisepticum culture also improved the height of the chromatographic peak and the profile of resolution and was the optimum pH for production and purification. The use of HPLC-SEC could be adapted to extract and purify the intended LP64 kDa band of MG.

INTRODUCTION

Mycoplasma gallisepticum is one of the most important disease agents affecting poultry causing high losses to poultry farms. The LP64 kDa is a lipoprotein playing major role in cytadherence of Mycoplasma gallisepticum (Forsyth et al., 1992). Many methods for isolation and purification of this lipoprotein were used such as pH of the medium, buffer and salts, detergents and chelators or metal ions but the most accurate is high performance liquid chromatography (HPLC) (Jansen and Rydén, 1998). Size-exclusion chromatography (SEC) has widely been used for separation and purification of proteins. SEC is an entropically controlled separation technique that depends on the relative size of macromolecules with respect to the size and shape of the pores of the packing. High performance size-exclusion chromatography (HP-SEC) columns are favored because of their speed, high resolution and high sensitivity (Laing et al., 2001). Optimization of HP-SEC conditions such as wave length of the detection, ionic strength of the mobile phase, sample injection volume, flow rate and pH of the MG culture (Engelhardt and Ahr, 1983). Our work was to optimize the HPLC-size exclusion technique for production and purification of the LP 64 kDa band of Mycoplasma gallisepticum.

MATERIALS AND METHODS

Materials:

A Model P 580 pump and Model UVD 170 S detector were used (Dionex, USA). The column employed for separation by size exclusion (SEC) was GF-250Xl column 21.2 mm ID X 250 mm (Zorbax). The fraction collector was Foxy Jr (Isco,Inc., USA).

Methods:

1- Preparation of media:

The basal broth dehydrated culture media was added to 100 ml distilled water, then sterilized by autoclaving at 121°C for 15 min. After cooling to 50°C other enrichment ingredients; horse serum, yeast extract, DNA and inhibitors (Thallium and penicillin G sodium), were added aseptically. Sterility was checked by incubation of the media at 37°C over night then stored at 4°C for use within 2-3 weeks (Frey et al., 1968)

2- Culture passage

M. gallisepticum, strains F (vaccinal strain), R, PG31 (ATCC type strain) and A5969 (high -

passage chicken isolate), were grown in Frey's medium with 12% swine serum. All cultures were grown at 37°C and harvested by centrifugation at 12000x g for 20 min.

3-Integral membrane proteins extraction

Extraction of integral membrane protein was accomplished with triton X-114 by the method of (Bricker et al, 1988), withsome modifications briefly pellets of M. gallisepticum (1.25 mg protein/pellet) were resuspended in 1 ml of ice cold. 1.0% (v/v) Triton X-114 was added in (10mM Tris, 150 mM NaCl buffer, pH 7.49 (TS buffer) 2mM phenylmethylsulphonyl floride with (PMSF, Sigma chemical co.) these were incubated in at 4°C for 30 min. Insoluble materials was pelleted by two cycles of centrifugation at 4°C for 15 min. at 13000 g. Supernatants were incubated at 37°C for 8 min. and centrifuged/at 10000 g for 5 min at room temperaturers The aqueous phase was readjusted with 10% Triton X-114 to 1.0% (v/v). The detergent phase was brought to the original volume with TS buffer. The previous method was repeated five times and the detergent phase was used for HPLC after dialyzing against three changes of 150 mM phosphate buffer saline (PBS) and stored at - 20°C until used.

4- Optimization of HPLC-SEC conditions:

4-1-Standard protein

64 kDa protein was isolated by electroelution after SDS-PAGE electrophoresis and then was used as slandered protein for the separation by HPLC- sEC procedure. The retention time of the protein was determined (from 6-11 min.) and 3 fractions could be separated having the same retention time.

4-2-Optimization of the wavelength.

Different wavelength 210 nm, 220 nm, 240 nm and 280 nm were used to optimize the wavelength, the column employed for separation by size exclusion was GF-250Xl column 21.2 mm ID X 250 mm (Zorbax). The mobile phase is 0.2 M di-sodium phosphate, pH 7.0.

4-3-Optimization of the strength of mobile phase.

Two mobile phase 100 mM and 200 mM NaH₂PO₄ were used to improve the chromatographic profile of separated protein, size exclusion GF-250Xl column 21.2 mm ID X 250 mm (Zorbax) was used. The injected sample was 200µl, the flow rate 5 ml/min. and detection was by UV absorption at 210 nm.

4-4-Optimization of the sample injection volume.

Different volumes of the protein samples (20, 40, 200 and 300 μ l) were injected, the column employed for separation by size exclusion was GF-250Xl column 21.2 mm ID X 250 mm (Zorbax). The mobile phase is 0.2 M disodium phosphate, pH 7.0, The flow rate 5 ml/min. and de-

tection was by UV absorbtion at 210 nm. The fraction collector was Foxy Jr (Isco,Inc., USA).

4-5-Optimization of the flow rate.

Different flow rates (3, 5 and 15ml/min) were used to improve the chromatographic profile of separated protein, the column employed for separation by size exclusion was GF-250Xl column 21.2 mm ID X 250 mm (Zorbax). The mobile phase is 0.2 M di-sodium phosphate, pH 7. 0. injected sample 200 µl. and detection was by UV absorption at 210 nm. The fraction collector was Foxy Jr (Isco, Inc., USA)

4-6-Optimization of the pH of M. gallisepticum culture

Different pH of M. gallisepticum culture (7.4, 6.4, 5.7, 5.45, 5.25 and 5 pH) were used to obtain the highest peaks of separated protein, the column employed for separation by size exclusion was GF-250Xl column 21.2 mm ID X 250 mm (Zorbax).The mobile phase is 0.2 M sodium phosphate, pH 7.0 . injected sample 200 μ l. and The flow rate 5 ml / min detection was by UV absorbtion at 210 nm. The fraction collector was Foxy Jr (Isco, Inc., USA)

5-Analysis of Eluted Fractions:

A-SDS-PAGE. for the eluted fractions.

Electrophoresis was performed as described by Laemmli (1970). The gels were stained with Coo-

massie brilliant blue R-250 (Sigma), and destained with mixture of 45% methanol, 10% acetic acid and 45% distilled water.

B-Immunological activity of the eluted frac-

Electrophoretic transfer to nitrocellulose (NTC) paper was accomplished as described by Towbin et al., (1979). After transfer to NTC, the membranes were stained with Ponceau's (Sigma), and the molecular weight standards were marked. The NTC blots were blocked with 5 % bovine serum albumin (BSA), 20 % fetal bovine serum, in Trisbuffer saline (10 mM Tris, pH 7.3, 0.9% NaCl), for 1 hr at 37°C with gentle rocking, The Mg antisera was diluted (1:500) in TBST (10 mM Tris pH 7.2, 150 mM NaCl, 0.05% Tween 20). Secondary antibody (horse radish peroxidase) conjugate was diluted 1:1000. The blots were developed for 5-10 minutes using H₂O₂ and 4-chloro-1-naphthol substrate.

6-Calculations: resolution for two adjacent peaks (m&n) can be calculated as follows:

Resolution = $2(t_n-t_m)$

 $W_n + W_m$

(t_n,t_m retention times wn,wm-band width)

RESULTS AND DISCUSSION

The LP 64 kDa is a lipoprotein playing major role in cytadherence of *Mycoplasma gallisepticum* (Forsyth et. al., 1992), so it is the part of cell

responsible for its virulence. For obtaining this lipoprotein by HPLC the most suitable method was the size exclusion technique in order to obtain this lipoprotein according to its molecular weight (64 kDa).

Optimization of the HPLC in separation and purification of LP 64 kDa band was studied including: wavelength, the mobile phase, sample injection, flow rate and pH of *M. gallisepticum culture*.

Optimization of the HPLC of wavelength on peak height using different wavelengths 210 nm, 220nm, 240 nm and 280 nm., results revealed that 210 nm gave the best peak height 58.91 mAU (Table 1) and the best resolution (Fig. 3). These results were in agreement with Calam and Davidson (1984), Laing et al., (2001) and Hayakawa et al., (2001).

To overcome the electrostatic effects between mobile phase and size-exclusion column matrix to improve the chromatographic profiles two aqueous buffers containing 100 mM NaH₂PO₄ and 200 mM NaH₂PO₄ were used; the aqueous buffer containing 200 mM NaH₂PO₄ improved the chromatographic profiles (Table, 2). The use of buffer ionic strength in range of (0.05-0.5M) was found to be suitable to avoid interaction between solute and matrex, this was in agreament with Jansen and Ryden (1998), also Laing et al, (2001) discovered that albumin or myoglobin had proper

size exclusion behavior at cocentrations from 25 to 300 mM.

Different injection volumes were tried 20, 40, 200, 300 μl to optimize the injection volume of protein sample on peak height. The injection of 200 μl resulted in peak height of 657.21 mAU and increased the resolution of the band (Fig. 4), while in case of injection of 20 μl,40 μl,300 μl the peak heights were 28.47,58.91 and 510 mAU respectively (Table 3).

The effect of varying flow rates on peak height, protein profile resolution revealed that the decrease of flow rate from 15 ml/min to 5 ml/min, increase peak height from 55.41 mAU to 114.63 mAU and also increase in the resolution of the curve (Table 4) and (Fig. 5).

The effect of pH of Mycoplasma gallisepticum culture used for separation revealed that the optimum pH was 5.7 which gave best result of peak height 173.55 mAU and best curve resolution (Table 5) and (Fig. 6).

Electrophoretic pattern and immunoblot of M. gallisepticum protein fractions separated by HPLC revealed the presence of LP 64 kDa in fraction 3 with high concentration and pure form (Figures 1 and 2).

In conclusion LP 64 kDa lipoprotein of Mycoplasma gallisepticum have been separated by size-exclusion (SEC) HPLC (GF-250X1 column 21.2 mm ID X 250 mm (Zorbax)). Mobile phase was 0.2 M sodium phosphate, pH was 5.7. The flow rate was 5 ml / min. and detection was by UV absorption at 210 nm. and recovered with its immunological activity as seen by immunoblot. The results obtained by SDS-PAGE suggests that this may be a valuable method for isolation and purification as we could separate the target lipoprotein in a pure form, so we can use HPLC (SEC) as a simple, convenient, rapid, reproducable, and reliable method for seperation of *M. gallisepticum* LP 64 kDa in highly pure form.

Chromatogram	Integration			Lan	
		l) metal	957) (1	a Maria	ma.I
2.7276 g.11.163	No.	Rct. time	Peak Name	Height mAU	Ret. Area %
> W \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	1	6.800	P1	12.957	19.61
: \	2	9.721	P2	10.914	24.69
$= 11 \text{ Y} \setminus 11$	3	10.981	P3	9.379	55.70
60 700 100 7th	Tot al			033.25	100.00
	AU WVL239 AN 2 1276 9 11,109	No. 1 2 3 Tot al	No. Ret. time min 1 6.800 2 9.721 3 10.981 Tot al	No. Ret. time Peak Name 1 6.800 P1 2 9.721 P2 3 10.981 P3 Tot al	No. Ret. time Name MAU 1 6.800 P1 12.957 2 9.721 P2 10.914 3 10.981 P3 9.379 Tot 033.25

Fig. (1): Electrophoretic pattern of M. gallisepticum protein fractions separated by HPLC.

Lane 1: low molecular weight standard (Pharmacia)

Lane 2: M. gallisepticum whole cell protein

Lane 3: M. gallisepticum (fraction 1, batch 1)

Lane 4: M. gallisepticum (fraction 2, batch 1)

Lane 5: M. gallisepticum (fraction 3, batch 1)

Lane 6: M. gallisepticum (fraction 1, batch 2)

Lane 7: M. gallisepticum (fraction 2, batch 2)

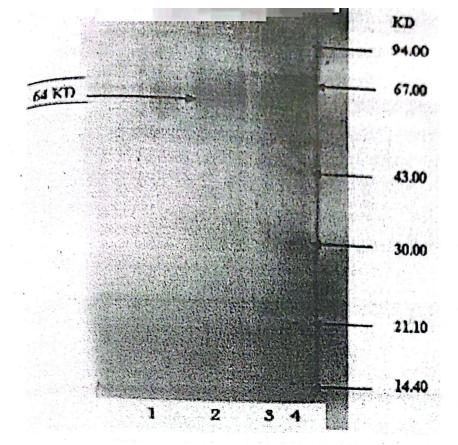
Lane 8: M. gallisepticum (fraction 3, batch 2)

Lane 9: M. gallisepticum whole cell protein.

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ractions

HPLC against M. gallisepticum positive serum.

ntl: M. gallisepticum (Fraction 3).

m2: M. gallisepticum (Fraction 2).

m3: M. gallisepticum (Fraction 1).

#4: Low Molecular weight standard (Pharmacia).



Table (1): Optimization of the wavelength.

	Wave	Chromatogram	Integration					
L	length	1 mm 1 m	_					
	*	Marie Company of the		No.	Ret.	Peak Name	Height mAU	Ret. Arca %
		A TOP TO THE PARTY OF THE PARTY	r	1	Time min 5.99	P1	15.93	10.74
				2	7.31	P2	13.25	23.91
1	210 nm			3	9.78	P3	13.69	20.05
				4	11.11	P4	16.05	45.30
	8.0			Total			58.91	100.00
		一				e *		
-							-	-
1		64 CELAN		No.	Ret. Time min	Peak Name	Height mAU	Ret.
		1		1	5.99	P1	9.38	Area % 11.79
	•	2.7.200 2.0073		2	7.30	P2	7.38	24.81
	220 nm			3	9.77	P3	7.53	20.89
	220 1411		1	4	11.09	P4	8.17	42.51
			Ì	Total		1	32.46	100.00
					1			
H		17.00 do 150 300	-			4 - 14 1 - 2	and the	
		AND THE PERSON NAMED IN COLUMN TO TH		No.	Ret. Time min	Peak Name	Height mAU	Ret. Area %
l	•			1	5.99	P1	6.28	51.35
		, are		2	7.03	P2	1.25	6.77
1	240 nm	\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \		3	9.77	P3	3.68	40.38
				4	11.09	P4	. 0.61	1.50
				Total	The shape		12.26	100.00
	0 6	als de de de					y-	
_	,	\$110,850,080,030,000		9			1 / 5	1.151
		WAL 200 mg	Ì	No.	Ret.	Peak	Height	Ret.
		1300			Time min	Name	mAU	Area %
			}	1	5.99	Pl	13.31	53.82
	280 nm			2	7.39	P2	1.9	5.44
	200 mii .			3	9.75 11.12	P3	6.08	19.36
					11.12		4.10	21.38
		Ab.	l	Total		L	25.39	100.00
		FEBRUARY WAS 10.0 DO.D						'
		te na ilo 200						

Table (2): Optimization of mobile phase concentration.

Mobile phase Conc.	Chromatogram		In	tegratio	on	
	10 mAU WA.218 se			*		,
	1.002	No.	Ret. Time min	Peak Name	Height mAU	Ret. Area %
A1 -No U DO		1	6.87	Pl	25.96	85.14
0.1 mNa H ₂ PO ₄	Jun 1	2	11.04	P2	4.56	4.44
		3.	19.72	P3	0.00	0.42
	201987	Total			30.53	00.00
						17
			Ret	Peak	Height	Ret.
	2003 MAIL NV.223 MI	No.	Ret. Time min	Peak Name	Height mAU	Ret. Area %
	11129	No.	Time	Name P1	mAU 30.062	Area % 24.12
	1 (4,22)	No. 1 2	Time min 6.522 10.490	Name P1 P2	30.062 17.573	24.12 22.25
0.2 mNa H ₂ PO ₄	11129	1	Time min 6.522 10.490 11.838	P1 P2 P3	30.062 17.573 44.219	Area % 24.12 22.25 37.64
0.2 mNa H ₂ PO ₄	1 (4,22)	1 2 3 4	Time min 6.522 10.490 11.838 13.402	Name P1 P2 P3 P4	30.062 17.573 44.219 4.128	24.12 22.25 37.64 1.12
0.2 mNa H ₂ PO ₄	10.00 1.1539 10.00 1.1539 10.00 1.1539	1 2 3 4 5	Time min 6.522 10.490 11.838 13.402 15.761	P1 P2 P3 P4 P5	30.062 17.573 44.219 4.128 4.386	24.12 22.25 37.64 1.12 5.33
0.2 mNa H ₂ PO ₄	1 (4,22)	1 2 3 4	Time min 6.522 10.490 11.838 13.402	Name P1 P2 P3 P4	30.062 17.573 44.219 4.128	24.12 22.25 37.64 1.12 5.33 9.54

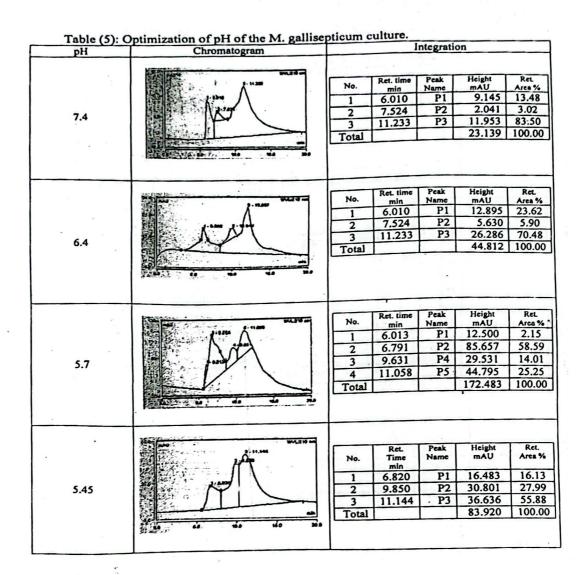
Table (1): Optimization of the wavelength.

Wave length	Chromatogram	Integration .				
			• 1			
	3007 WC210 an	No.	Ret. Time min	Peak Name	Height mAU	Ret. Area %
	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	1	5.99	P1-	15.93	10.74
	Well	2	7.31	P2	13.25	23.91
210 nm		3	9.78	P3	13.69	20.05
		4	11.11	P4	16:05	45.30
		Total	of regard		58.91	100.00
	to the state of th		- H			
	BARRIE		Ret.	Peak	Height	Ret.
	12.5 mAj (WV. 230 nm	No.	Time min	Name	mAU	Area %
	J. 8 DOZ	1	5.99	P1	9.38	11.79
	2.7.300 2.0703	2	7.30	P2	7.38	24.81
220 nm		3	9.77	P3.	7.53	20.89
		4	11.09	P4	8.17	42.51
		Total			32.46	100.00
	10 min		Za Terrestina			1 30
	EDD WILESO AND		Ret.	Peak	Height	Ret.
	4-6.907 WVL 240 am	No.	Time min	Name	mAU	Area %
	1	1	5.99	P1	6.28	51.35
	2-7.020	2	7.03	P2	1.25	6.77
240 nm	2.00	3	9.77	P3	3.68	40.38
		4	11.09	P4	0.61	1.50
	1.00 man	Total			12.26	100.00
	2.9 4.0 10.0 14.0 20.0					
	10.5] [AAU WV. 200 nm]	No.	Ret.	Peak	Height	Ret.
	J-8 900	1	Time min	Name	mAU	Area %
	100-	1	5.99	P1	13.31	53.82
280 nm	3.0790	2	7.39	P2	1.9	5.44
200 IIII	100	3	9.75	P3	6.08	19.36
		4 Total	11.12	P4	4.10 25.39	21.38
		l Oto			175 30	1 100.00

Table (2): Optimization of the mobile phase concentration.

Table (4): Optimization of flow rate.

	ptimization of flow rate.					
flow Rate	Chromatogram			Integrat	ion	
Flow 15 ml/ min		No. 1 2 3 4 5 Total	Ret. time min 1.26 2.18 2.44 3.40 3.86	Peak Name P1 P2 P3 P4 P5	Height mAU 4.11 14.05 13.58 11.75 11.24	Ret. Area % 5.37 10.05 27.96 22.82 33.51
	The second secon	10121		-	55.41	100.00
Flow 5 ml/ min	24 WASS WASS WAS CO.	No. 1 2 3 4 5 Total	Ret time min 6.310 6.342 6.370 16.061 19.449	Peak Name P1 P2 P3 P4 P5	Height mAU 36.81 36.65 0.843 6.77 33.18 114.63	Ret. Area % 13.60 16.73 0.14 13.61 55.91 100.00
Flow 3 ml/ min	20.0 4.10.204 10.224 10.	No. 1 2 3 4 5 Total	Ret. time min 10.795 11.592 17.414 19.684 22.480	Peak Name P1 P2 P3 P4 P5	Height mAU 21.476 1.320 7.830 '29.135 2.539 62.300	Ret. Area % 46.07 1.52 12.70 37.76 1.95 100.00



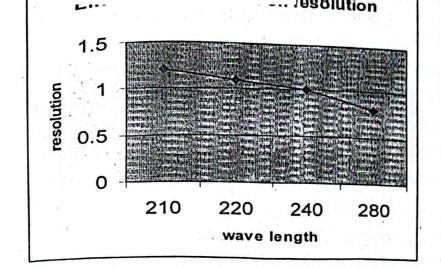


Fig. (3): Effect of wave length on the resolution.

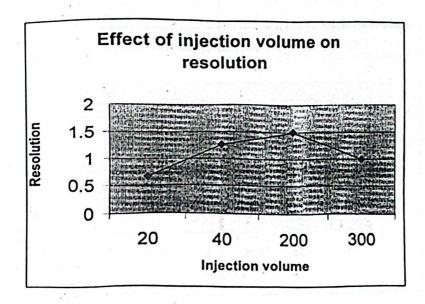


Fig. (4): Effect of injection volume on the resolution.

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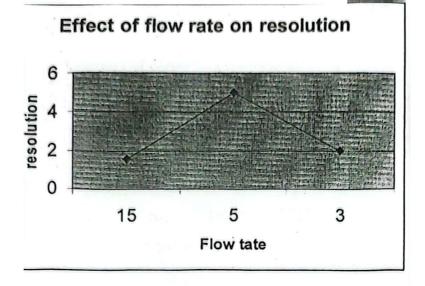
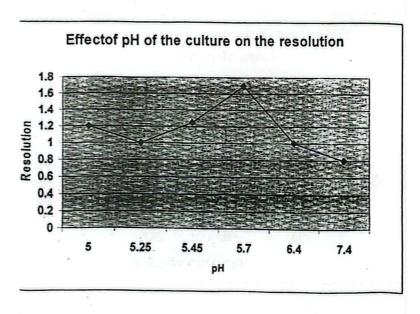


Fig. (5): Effect of flow rate on the resolution.



ig. (6): Effect of pH of the culture on the resolution.

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