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Efficacy of Black Quarter Vaccine on Cattle: A Spectroscopic Approach

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In this work, normal healthy pre vaccinated blood samples (zero day) and post vaccinated (7th, 14th, 21st day after vaccination) blood samples are analyzed by employing FTIR and UV-Vis spectroscopic techniques. The internal standards among the application peaks are calculated in both the methods. Variation is expected due to the production of antibodies in animal. Among the various techniques to study the antibody production, ELISA (enzyme-linked immuno sorbent assay) is considered a better one for the present, but it can be done only in sophisticated laboratories. Hence there is a need to try for other techniques to know the immune status of vaccinated animals. The present work can be extended and compared to the ELISA technique.

Key Words: Black quarter vaccine, UV-visible and FTIR spectra, Internal standard.

INTRODUCTION

Vaccinology as a scientific field and vaccine production as a commercial activity are areas of rapid growth. Vaccines are generally considered the most-efficient tool of public health, research and development in vaccinology may be considered a major public concern. Vaccination is the best and cheapest method to protect the body against bacterial and viral diseases. Among the various bacterial diseases affect the animal, Blackleg is one of the fatal disease of young cattle and sheep and occasionally in other animal species. Clostridium chauvoei is the causative organism in most cases. Some affected animals have also been found to be affected with Clostridium feseri. These two microbes are gas-producing bacteria. They also form spores, which can live in soils for many years. The microbes cannot spread the disease Blackleg from one animal to another simply by contact. The common symptoms are cattle not being able to walk properly, loses interest in food, high temperature, swelling that develops in shoulder, back, neck and makes a cracking sound under pressure. Gas is formed inside the enlarged swellings. Once affected with Blackleg, the animal can die within 48 h. Without proper treatment, cattle can be found dead with no time to react. After the initial symptoms, the bacterium causes a gradual

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poisoning reaction in the body. In sheep, Blackleg is mostly found after instances of physical wounds. The spores of the bacterium might be present in the gut of the cattle for sometime. These spores do not immediately result in the disease. Medical research is ongoing to find what conditions cause these ingested spores to give rise to the disease. Naz et al.1 have taken thigh muscle samples from 6 cows and 2 buffalo died in outbreaks of Black Quarter (BQ) at different districts of Punjab. Clostridium chauvoei was isolated from all the samples, caused death in guinea pigs within 24 to 78 h with rapid development of symptoms identical to the symptoms of (BQ) in cattle. It is evident from their study that *Clostridium chauvoei* alone is isolated from the affected muscle samples of cattle and buffalo which causes Blackleg disease. Sudarsanam et al.² assessed the result of cell mediated immune response and humoral immune response combinedly in production against Black Quarter. They recorded the result of monitoring those responses in guinea pigs vaccinated with black quarter vaccine and correlating the responses with protection to virulent challenge afforded by these animals along with unvaccinated controls. Srinivasan et al.³ assessed the serological response of bovines to combined vaccine containing foot and mouth disease virus, rabies virus, Pasteurella multocida and Clostridium chauvoei antigens and individual component vaccines also. Serological response of the calves was assayed on days 21 and 90 post vaccination. There was no significant variation in the serological response elicited by individual component vaccines and combined vaccine containing all four antigens. In serological study, the sample used is very less and hence it is a best method. The fluid portion plasma contains a large number of organic and inorganic substances such as proteins, vitamins, minerals, lipids, etc. Although the main function of blood is to transport various minerals to all cells of the body, blood also provides the temperature regulating a defense mechanism^{4,5}. In this work, normal healthy pre vaccinated blood samples (zero day) and post vaccinated (7th, 14th, 21st day after vaccination) blood samples are analyzed by employing FTIR and UV-Vis spectroscopic techniques.

EXPERIMENTAL

The experiments were carried out in a village Kaveripakkam, Vellore District, Tamilnadu. Ten healthy cattle were used. Blood samples were collected from each of them from jugular vein. They were housed in a clean shed with good ventilation. No antibiotics were given during the experimental period. Pre-vaccinated blood samples were taken. Then they were immunized with 0.5 mL of black quarter vaccine. Subsequent blood samples were collected on 7th day, 14th day and 21st day from the same vaccinated animal. After collecting the blood, the serum was separated. Using the conventional method, the samples could be prepared by spreading a small volume of serum on an IR-transparent material, allowing drying and measuring the absorption spectrum of the film. The accuracy of the method may be compromised by any variation in the amount of serum successfully deposited on the KBr window, particularly with the manual sample preparation. In order to make up for this variation Vol. 22, No. 4 (2010)

and to assess its impact on the overall accuracy of the method, a standard solution is added to each serum sample. The solution is chosen in such a way that it respond to IR radiation at the point where serum sample contains no absorption peak. Shaw et al.⁶ reported that the IR absorption spectrum of thiocyanate ion includes absorption at 2060 cm⁻¹ in a spectral region where sera samples and subsequently normalizing all of the spectra to equal intensities therefore compensated for the imprecision in the film preparation. A volume of 1 mL of serum was diluted with an equal volume of 4mg/L aqueous potassium thiocyanate solution 20 µL of each diluted sample was spread evenly over the surface of a circular KBr window (9 mm diameter and 2 mm thickness). Infrared spectra in the region 4000-500 cm⁻¹ were recorded on an ABB BOMEM MB SERIES-one FTIR spectrometer equipped with an air-cooled DTGS (Deuterated triglycine sulphate) detector. It has already mentioned that the strong absorption band of water in the mid IR region is hindered and to eliminate in the same, the serum samples are air dried to form a thin uniform film on the KBr pellet^{6,7}. IR transparent KBr material without the samples was scanned as background for each spectrum and 23 scans were co added at a spectra resolution of 4 cm⁻¹. The collected signal was transferred to the PC. The data were processed by windows based data program-spectrum software. The spectra were base line corrected and they were normalized to acquire identical area under the curves and the maximum absorbance values of the corresponding characteristics bands were noted. For UV-vis spectroscopic measurements, the whole blood was used. The whole blood was diluted with normal saline at a concentration of 0.9 % and the spectra were recorded using Shimadzu UV 1601 spectrometer in the region 200 to 700 nm. Both the spectra were taken at Dr. Ceeal Analytical Lab., Chennai, India.

RESULTS AND DISCUSSION

FTIR Spectral analysis: The spectra of pre and post-vaccinated sera samples were all distinct from one another but were dominated mainly by the absorption of the protein constituents which provides the selectivity in infrared based serum analysis. Table-1 presents the vibration band assignment of serum. The vibration band at 3300 cm⁻¹ is due to the N-H stretching vibration of the secondary amides of protein. The asymmetric and symmetric stretching vibrations of the methyl group of proteins and lipids are found to be present at 2956 and 2896 cm⁻¹, respectively. The other two vibration bands in C-H stretching region are found to be present near 2922 and 2851 cm⁻¹, which are due to the asymmetric and symmetric stretching vibration of the methylene group. The strong absorption band present at 1655 cm⁻¹ is attributed to C=O stretching of amide-I of the proteins. In the same way the presence of the band at 1548 cm⁻¹ is due to the amide-II or N-H bending vibration that are strongly coupled to the C-N stretching vibrations of the protein amide groups. The peaks at 1456, 1400 and 1315 cm⁻¹ are considered to be asymmetric and symmetric deformations of the methyl group of proteins. The peak at 1400 cm⁻¹ may also considered due to COO⁻ stretch of ionized amino acid chains, suggesting an increased contribution

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TABLE-1
INFRARED VIBRATIONAL BAND FREQUENCY ASSIGNMENT OF SERUM

Vibration	Assignment				
band (cm^{-1})	Assignment				
3296	N-H stretching of secondary amides of protein: amide A				
2960	CH ₃ asymmetric stretching of proteins and lipids				
2936	CH ₂ /CH stretching				
2874	CH ₃ symmetric stretching of proteins and lipids				
2851	CH ₂ /CH stretching				
1660	C=O stretching (80 %) weakly coupled with C-N stretching (10 %) and NH deformation (10 %)-amide I				
1545	NH deformation (60 %) strongly coupled with C-N stretching (40 %) amide II				
1457	CH ₃ asymmetric deformation				
1398	CH ₃ asymmetric deformation COO ⁻ stretching of amino acids				
1315	CH ₃ symmetric deformation				
1240	Asymmetric PO ₂ stretching of lipid phosphates				
1169	C-O stretching				
1128	C-O stretching				
1081	C-O stretching				
955	PO ₂ symmetric stretching of lipid phosphates				
701	NH asymmetric deformation coupled with CH ₂ rocking amide V				
625	O=C-N deformation (40 %) coupled with other ring deformation (60 %) amide IV				

from carboxalate. The lipid phosphate band due to the asymmetric P-O stretching vibration is found to occur at 1240 cm⁻¹. The spectral region 1250-925 cm⁻¹ is predominantly occupied by the C-O stretching vibrations of glucose. The absorption peaks present at 1169, 1153, 1107, 1079 and 1035 cm⁻¹ are considered to be due to the different C-O stretching vibrations of C-O-H and C-O-C bonds. The weak absorption band at 955 cm⁻¹ is considered to be due to P-O symmetric stretching of the phosphate bond of proteins. The medium strong vibration bond present at 702 cm⁻¹ is assigned as N-H out-of-plane bending with the contribution of C-N torsional vibrations.

The FTIR spectra of all the sera sample both pre and post vaccinated show the corresponding absorption bands in their specific regions qualitatively. But quantitatively there is a considerable difference in the spectra between the pre and post-vaccinated sera. The absorbance was directly proportional to the concentration. Hence the different sera samples were analyzed quantitatively by calculating the intensity ratio among the absorption peaks. In order to quantify the spectral difference, three intensity ratio parameters have been introduced. They were $R_1 = I_{3296}/I_{2960}$ due to the N-H stretching of secondary amides of protein, amide A and CH₃ asymmetric stretching of proteins and lipids, $R_2 = I_{1398}/I_{1457}$ due to CH₃ asymmetric deformation COO⁻ stretching of amino acids and CH₃ asymmetric deformation and $R_3 = I_{1660}/I_{2874}$ due to the ratio of the intensities of amide-I and CH₃ symmetric stretching of proteins and lipids. Though the infrared spectra of all the sera samples were similar, considerable differences were found to be present in the internal standards among the absorption peaks in pre and post vaccinated samples.

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Table-2 summarizes the internal standard calculations of the pre and post-vaccinated samples. The values of R_1 were found to be around 1.32 to 1.47 for pre-vaccinated serum. But the values of R_1 , during the period of 7th and 14th day changed due to the production of antibodies. On 21st day of vaccination the values increases to the range of 1.46 to 1.8. Similarly R_2 values were from 1.026 to 1.078 in the pre-vaccinated state and it changes to 1.137 to 1.179 in the post vaccinated state. The values of R_3 were found to be around 1.516 to 2.375 for pre-vaccinated serum. But the values of R_3 during the period of 7th and 14th day changed due to the production of antibodies. On 21st day of vaccination the values increased to the range of 2.126 to 2.473. These variations occur due to the antibodies produced by the way of vaccination.

The UV-Vis spectra have been recorded for the pre and post vaccinated blood samples. The UV-Vis spectra of all the blood samples exhibit the presence of two strong absorption peaks at 417 and 576 nm. But there was a marked difference in the absorption levels of pre and post vaccinated blood samples. Table-3 summarizes the internal standard representation among the absorption peaks for the pre- and post-vaccinated blood samples. The internal standard among the absorption peaks was found to be around 6.555 to 8.324 in pre-vaccinated blood sample. But it decreased in the 7th day and increased in 14th and 21st day after vaccination. But its value was less than the pre vaccinated state. These variations occur due to the production of antibodies between zero day and 21st day. The antibodies were produced at the maximum level on the 21st day. The immunity following vaccination was established in about 10 days and is expected to confer protection against natural infection for a period of 6 to 9 months.

Conclusion

Animal diseases cause enormous economic loss through mortality, inefficient production and increase in the stock replacement rates, which all require additional resources. Control and treatment of the diseases also contribute to the losses⁸. Control measures in present-day programme include quarantine of imported animals; cooperation of agencies in the study and control of animal diseases; inspection of red meat and poultry to minimize the danger of spread of animal disease to human beings; inspection and evaluation of vaccines and other pharmaceutical and biological products as to purity, efficacy and safety; inspection of the mass slaughter of animals and the destruction of carcasses. Universities and other research institutions conduct studies on the many disease problems that arise.

Spectroscopy has been employed as a diagnostic tool in the study of blood. FTIR and UV-Visible spectroscopic methods have been employed to study the pre and post vaccinated blood samples. The internal standards among the absorption peaks were calculated. By studying this, the potency of the vaccine can be justified. Present work is a serological analysis in infrared spectroscopy. In future this study can be compared with the ELISA serological procedure.

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Category of sample	Days	$R_1 = I_{3296} / I_{2960}$	$R_2 = I_{1398} / I_{1457}$	$R_3 = I_{1660} / I_{2874}$
Cattle 1	Pre	1.493	1.026	2.224
	Post 1	1.328	1.012	1.638
	Post 2	1.297	1.018	1.368
	Post 3	1.657	1.137	2.392
Cottle 2	Pre	1.603	1.136	2.375
	Post 1	1.325	1.006	1.712
Cattle 2	Post 2	1.604	1.120	2.274
	Post 3	1.611	1.136	2.473
	Pre	1.320	1.038	1.677
Cottle 2	Post 1	1.847	1.096	3.360
Cattle 5	Post 2	1.646	1.138	2.447
	Post 3	1.823	1.087	3.344
	Pre	1.453	1.078	2.080
C-#1- 4	Post 1	1.578	1.136	2.360
Cattle 4	Post 2	1.661	1.130	2.292
	Post 3	1.442	1.179	2.179
	Pre	1.475	1.062	2.048
	Post 1	1.576	1.040	2.524
Cattle 5	Post 2	1.274	1.018	1.268
	Post 3	1.320	1.038	1.677
	Pre	1.322	1.039	1.582
	Post 1	1.577	1.042	2.451
Cattle 6	Post 2	1.921	1.143	2.661
	Post 3	1.466	1.256	2.023
	Pre	1.471	1.039	2.058
	Post 1	1.442	1.035	2.063
Cattle /	Post 2	1.223	1.028	1.476
	Post 3	1.502	1.028	2.282
	Pre	1.279	1.032	1.516
Cattle 8	Post 1	1.474	1.053	2.126
	Post 2	1.415	1.011	2.009
	Post 3	1.569	1.04	2.491
	Pre	1.748	1.024	2.671
Cottle 0	Post 1	1.491	1.053	2.116
Caule 9	Post 2	1.408	1.966	1.830
	Post 3	1.571	1.040	2.598
	Pre	1.446	1.106	1.841
Cattle 10	Post 1	1.583	1.040	2.530
Cattle 10	Post 2	1.334	1.012	1.601
	Post 3	*	*	*

TABLE-2 INTERNAL STANDARD CALCULATIONS OF THE PRE AND POST-VACCINATED SAMPLES OF CATTLE

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Category of sample	Days	Wavelength	Wavelength (nm) and absorbance value		
	Pre	417	576	A 417/ A 570	
Cattle 1	Post 1	0.7662	0.1018	7.526	
	Post 2	0.3642	0.0976	3.732	
	Post 3	0.3517	0.0931	3.778	
Cattle 2	Pre	0.3187	0.0902	3.533	
	Post 1	0.8382	0.1007	8.324	
	Post 2	0.9728	0.1810	5.374	
	Post 3	0.9646	0.1627	5.929	
	Pre	0.9562	0.1542	6.201	
	Post 1	0.9052	0.1381	6.555	
Cattle 3	Post 2	0.9637	0.1722	5.596	
	Post 3	0.9528	0.1658	5.747	
	Pre	0.9372	0.1582	5.924	
	Post 1	1.0675	0.1310	8.149	
Cattle 4	Post 2	0.8704	0.1310	5.212	
	Post 3	0.8610	0.1670	5.595	
	Pre	0.8541	0.1539	6.308	
	Post 1	0.9129	0.1354	7.717	
Cattle 5	Post 2	1.0942	0.1183	5.744	
	Post 3	1.0812	0.1905	5.788	
	Pre	1.0524	0.1868	6.261	
	Post 1	0.7362	0.1067	6.900	
Cattle 6	Post 2	1.1101	0.1987	5.587	
	Post 3	1.1091	0.1798	6.168	
	Pre	1.1016	0.1683	6.545	
	Post 1	0.8768	0.1082	8.104	
Cattle 7	Post 2	0.6717	0.1415	4,747	
	Post 3	0.6661	0.1355	4.916	
Cattle 8	Pre	0.6546	0.1215	5.388	
	Post 1	0.9092	0.1314	6.919	
	Post 2	1.2489	0.2113	5.911	
	Post 3	1.2391	0.2031	6.101	
	Pre	1.2185	0.1925	6.330	
	Post 1	1.0342	0.1409	7.340	
Cattle 9	Post 2	0.8619	0.1574	5.476	
	Post 3	0.8540	0.1498	5.701	
	Pre	0.8361	0,1245	6.716	
	Post 1	0.9371	0.1285	7 293	
Cattle 10	Post 2	1 0572	0 1672	6 3 2 3	
	Post 2	0.9192	0.1551	5 777	
	Post 2	0.0172	0.1331	6 720	

TABLE-3

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