قسم: الفسيولوجيا. كلية: الطب جامعة أسيوط. رئيس القسم: أ.د ./ مصطفى جابر.

## تأثير العليقه الخضراء والعليقه الجافه المركزة على بكتريا الكرش في الماعسسز

سناء نصار ، محمد عوض ،عبد العزيزمحمود ، ابتسام حسن

كانت تغلب على بكتريا الكرش الأنواع موجبة الجرام وكروية الشكل في حالة العليقه الخضراء. بينما معظم البكتريا السالبه الجرام ذات الأشكال المختلفة تزد هـر على العليقـه المركزه الجافـه.

بعد ٢ ساعات من تناول العليقة الخضراء تناقصت كل أنواع البكتريـــا ماعدى العصويـه سالبة الجرام والكويـنز التي زادت.

بينما على العليقة الجافة بعد ٦ ساعات من تناول العليقة زادت البكتريا سالبة الجرام ماعدى الوادية الشكل والأسلوبيرا تناقصت.

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Dept. of Physiology., Fac. Of Med.; Assiut University. Head of Dept. Prof. Dr. M.G.Mohamed.

# EFFECT OF GREEN FODDER AND DRY CONCENTRATE RATION ON RUMEN MICROORGANISMS IN GOAT

(With 3 Tables and 3 Figures)

SANAA M. NASSAR, M.A. ALI, , A.A. HAHMOUD, IBTISAM M. HASSAN (Received at 7/7/1985)

#### SUMMARY

On green fodder, the Gram positive and coccoid form microorganisms were predominated. While most Gram negative and variable forms were flourished on dry ration. At 6 hours on green fodder, all forms of microorganisms decreased except Gram negative rods and Quins which were increased. On dry ration, at 6 hours Gram positive forms decreased and Gram negative increased except vibrio and oscillospira were decreased.

#### INTRODUCTION

It has been increasingly realized that more fundamental knowledge of ruminal microorganisms is needed to obtain more efficient rations and better control of metabolic disorders in ruminants (BRYANT, 1959). Little data is to be found on the significance of different types of rumen organisms in relation to the composition of the diet effects on the population in rumen.

BRYANT and BURKEY (1953); MAKI and FOSTER (1957); HUNGATE (1957); WARNER (1962) had shown that bacterial population in rumen of animals was conditioned to different diets.

The present paper has to study the pattern of changes in the various groups of micro organisms in goat rumen in relation to diet.

#### **MATERIALS** and METHODS

Four rumen fistulated Baladi goats were used. They were fed barseem as green fodder for period of 30 days and concentrates; Bran, Wheat straw and Sorghum for another 30 days as period of adaptation before taking samples.

Animals fed once per day at 9 oclock in the morning. Samples from rumen content were collected twice daily at 6 and 22 hours after feeding.

Samples were collected in air tight screw caped bottles. One sample was fresh and the other was fixed with 10% formalin in ratio of one part of the sample to one part of 10% formalin.

1) Fresh samples were incubated at 37°C with 1% starch to demonstrate the iodophilic micro-organisms.

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- 2) Fixed films were stained as follow:
  - Gram's stain as general bacterial stain.
  - Spores were stained by Ziehl Neelsen's carbolfucsin, decolourised by 5-10% sulphuric acid and counterstained with methylene blue.
  - Capsules were demonestrated by method of MUIR (1962).
- 3) For counting: On a fixed slide, we make differential count for 300 microorganisms from different field, twice daily for each case according to MOIR and MASSON scheme (1952).

#### RESULTS and DISCUSSION

The nutritional economy of ruminant animals depends on the absolute numbers of microorganisms present in the rumen. The pattern of change in concentration of different groups of micro organisms was studed in relation to diet.

On green fodder, the Gram positive and coccoid form microorganisms were predominated. While most Gram negative and Gram variable forms were flourished on dry ration (Table 3 & Fig. 2). This finding was supported by CHENG and COSTERTON (1975) and COSTERTON (1979).

On green ration, coccoid form and Quins oval predominates due to the high percent of crude fibers in the solid fraction of barseem. SUPERTEIGM, 1951; BRYANT and BURTREY, 1953 and BRYANT and DOETSCH, 1954, stated that Gram positive monococci represented by rumino coccus albus and rumino coccus flavifociens were usually cellulose digester forms. Quins oval increased after feeding as it grow on fresh clover and grasses (CLARKE, 1979).

On dry concentrate ration, the percent of Gram negative microorganisms were increased specially Gram negative rods, diplo-cocci, spiral, Vibrio and Selenomonas and Gram variable streptococcus (Tables 2, 3 & Fig. 2, 3). This finding was supported by OSBOURN, TERRY, CORELL and OUTEN, 1970; JOURNET, 1971 and STEWART, 1977. They stated that in ruminant maintained on green forage diet and changed to concentrate diet with large amount of starch, the cellulose producing population were decreased and amylase producing species proliferated.

The Gram negative rods were usually represented by Bacteroids amylophilus and Bacteroidsruminocola, as they were the main Gram negative starch digester rods (HALMIN and HUNGATE, 1956 and LEWIS, 1957). Gram variable streptococcus and Gram negative diplococci were increased. Strepto coccus increased in rumen where there is increased grains (HUNGATE, 1957). It is one of starch digester microorganisms which posses amylases (BRYANT, SMALL, BOURMA and ROBINSON, 1958).

Streptococcus bovis ferment large number of different carbohydrates with the formation of lactic acid (HOWARD, 1959). The Gram negative diplococci was usually represented by peptostrepto-coccus Elsdenii. It was in direct relationship with lactic acid content of rumen on concentrate diet (GUTIERREZ; DAVIS, TINDABLE and WARWICK, 1959). The spirillum was glucose and starch digester (KARULOV and KROTHOVA, 1971). Vibrio was represented by Butrivibrio fibrisolveno. It mainly utilizes starch in addition to cellulose (MAKIE and FOSTER, 1957; GILL and KING, 1958; MERGERITA and HANGATE, 1963). Selenomonas ruminatum was a starch digester microorganisms (HUNGATE, 1957).

On green fodder, at 6 hours after feeding, all forms of microorganisms decreased except Gram negative rods and QUINS OVAL (Tables 1, 3 & Fig. 1, 3).

The decrease in number of all forms of micro organism on feeding green fodder was due to diluting effect of barseem, as it contains high moisture content.

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#### DIET AND RMEN MICROORGANISMS

The Gram negative rods were usually bacteroid forms which act as glucose digester, specially bacterordes succinogen which digest cellulose beside glucose(KORULOV and KROTHOVA, 1971). Where most of the dry matter of Barseem is rich in soluble carbohydrates and fibers. Quins oval grow on fresh clover and grasses (CLARKE, 1979).

On green fodder, at 22 hour both Gram positive and Gram negative mono-cocci, Diplo-cocci and Vibrio were increased (Tables 1, 3 & Fig. 1, 3) as the main rumen content at that time was mainly cellulose.

On concentrate rations, at 6 hours Gram negative monococci, rods and spiral forms were increased which act as starch digesters. The number of streptococcus, Gram positive rods and vibrio discreased as they were utilized by special species of protozoa like Entodinium to check the progressive increase of lactic acid producing bacterial; strepto-coccus bovis; on concentrates (HUNGATE, DOUGERTY, BRYANT and CELLO, 1952; MANN, MASSON & OXFORD, 1954 and MANN, 1970).

From above, it is clear that rumen microorganisms are a well balanced system. On green fodder, where Gram positive and coccoid form increased, the Gram negative and rod forms decreased. Vice versa occures on concentrates.

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Table (1) The percentage of different types of bacteria in the rumen of goats fed on green (Barseem) ration.

Type of bacteria  Monococci  Small Gr (-) Small Gr (-) Large Gr (-) Large Gr (+)  Diplo  Small Gr (-) Small Gr (-) Small Gr (-) Small Gr (+) Small Gr (-) Large Gr (+)  Sarcina  Small Gr (-) Small Gr (+)  Chain  Shor Gr (+)	Animal No lat 2ndd3  14 16 18 20 19 6 21 1 1 2 - 6 12 10 6 11 2 - 6 1 2 - 7 1 1 1 - 1 1	At morn:  At 12 13  If 18 23 13 23  If 18 23 13 23  If 6 4 12 13  If 6 8 4 14 12  If 19 6 9  If 1 - 6 3  If 1  If 1	11 1 1 1 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	1 1 1 6 5 1 4 B 6 4 2 B	At morn 1 Animal No 2 1 Animal No 2 2 13 23 13 23 6 2 - 2 1 17 13 5 6 3 - 2 1 17 13 5 6 7 1 17 13 5 6 7 1 17 13 5 7 1 17 13 5 7 1 17 13 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	At morning all No 2 And 3rd 1s 23 19 13 23 19 12 13 2 10 2 17 117 13 2 2 10 2 1 2 1 2 1 2 1 2 1 2 1 2 1 2 1	19 20 2: 2 10 11 1 1 2 2 2 2 10 11 1 1 2 2 2 2 1 1 1 3	20 20 20 20 20 10 10 10 10 10 10 10 10 10 10 10 10 10	108 Animal No 3 1st 2nd 3rd 2 20 22 2 10 11 4 3 5 10 5 4 1 1 3 2 1 1 3 2 1		200 B B B B B B B B B B B B B B B B B B	Gree  Animal No 4  let 2nd Jrd  24 30 27  6 8 16  2 - 1  2 1 -  6 1 2  - 7 10  10 3 7°  6 1 2	Green ration  No 4  No 4  Jrd Mean  Jrd 10.6+1.7  1 1.9±0.4  - 3.3±1  7. 8.1±1.4  - 1.4±0.5  2 2.1±0.6  2 0.5±0.2  - 0.8±0.4  - 0.2±0.1	1 1 1 1 6 2 1 2 1 1 1 1 1 1 1 1 1 1 1 1	Animal No lat 2nd 31 1 22 26 11 22 26 11 8 2 2 5 10 6 4 6 1 - 1 - 6 1 - 2 - 3 1 - 2 - 3 1 - 3	No No No	1 1 2 1 1 7 1 1 20 materials	Animal Mo 2 lat 2nd 3rd 20 22 18 20 22 18 3 18 3 1 18 5 7 12 5 7 12 5	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	After 0 2 An 11 11 11 12 12 12 12 12 12 12 12 12 12	. 10 11	16 16 16 16 17 16 17 17 17 17 17 17 17 17 17 17 17 17 17	1 1 1 1 3 3 1 1 1 1 1 1 1 1 1 1 1 1 1 1	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	11 11 13 8 4 11 2 8 2 and .	Animal No 4  lat 2nd 3rd  16 28 16  2 2 3  2 2 3  2 1 - 2  1 1 - 2  1 - 2	P -	NO WE FARE LE	Mean 19.3±1.4 8.1±1.8 0.8±0.3 3.1+1.1 5.7+0.9 7.4+0.9 0.9+0.4 0.6+0.5 0.3+0.2 0.6+0.5
Small Gr (+) Chain	1	1	•	•	1	~	N	-		1	1	1	0.3±0.2		1		,				6	- 1	1 1	, -	1 1	1 1	0 0		0.6+0.5
Shor Gr (+) Long Gr (+)		1	1		1	1	-		- 1			1	0.2±0.1	м	1	. 1	,					1		,	1	1		1	1.0+
Rods	700		1 18		24	21	18	15	14	9	27	16	19.3+1.5	4	36	26	20	23	28	_	8 15		22 1		26	24	27.		27.9+2.5
Small Gr (-)	6 6	5 6					1 (	us i	6	2	12	w	3.6+1		6		9	10			7		, ,		1 0		1 4		4+1.1
Gr Gr		1 .	w 4	12	4	4	9	6	5	10	1	4	5±1.1	5	9	12	12	4	14		-	4 1		20	20	14	11.		11.6+1.6
Large Gr (-)	10	N 1		1 0	1 4	N 4	1 1	w	N	6	1	N	1.4+0.5		1	,	N					-	N	21		1	0.5+0.2	19	0.2
Large Gr (+)	The same												1 1	3												1	0 2		0.1
Filamentous form	1	1	•	1	1	4	м	ŧ	1	w	1	-	1.1+0.5	1	1	-	.4	20-		i		'	) 1	3 6		0 8	1.042.0		
Gr (-)	1	.h	6	3	4	6	4	6	5	5	4	u	3.9+0.6	6	N		w	_				-	N	7		8	3.4		3.4+0.8
Spiral Gr (-)	8	7 1	-	8	7	1	6	5	4	13	6	S	7.6+0.8	7	4	6	9	S	w			400	-	6	*	10	5.4		5.4+0.7
Vibrio Gr (-)	1 6	8 -				1	N	1	-	1-	1	,	0.3+0.2	N	1	2	-	1	-			5	5	-	-	1	1.4		0.5
Quin's oval Gr (-) Solenomonas Gr (-)	1	1	6	-	,	6	w	-	٣	N	-	-	1.9+0.6	1	-	1	S	N	N	N		N	S	S	-	•	2	10	2.3±0.6
Oecillospira Gr (-)		•		•		1	_	1		1	1	1	0.1+0.1	N	1	1	w		,	1			1	1	100	1	0.6	[14	0.6+0.3

Table (2) The percentage of different types of bacteria in the rumen of goats fod on dry concentrate (Wheatbrn and grain sorghums) ration

	1		-	-	1			-				610	ration	CO											
Type of			1	At	At morning	Bu		9	3					100	1	Afte	After noon	non				18	-		
bacteria	Ania	Animal No 1 Animal No 2 let 2nd 3rd 1st 2nd 3rd	No No	1 A	and No 1 Animal No 2	No d 31		Animal No 3 lst 2nd 3rd	al No 3	) Mean	9	Andi		No 1	al No 1 Anig	Animal No.2	No.	2 Ani	and No.2 Animal No 3	No Jrd		and 200	Animal No 4	Mean	Total mean
Monococci Small Gr (-)	12	15	21	16	14	23	62	3 16	17	17.4+1.3		=	33	1 5	=	3.	100	3	1	1:	1 :	1 8	1		+
Small Gr (+)	-	8	4	2	9						7	, ,	, ,	3	; ,	1 1	67	,	1	1 4	10	28	16	18.8+1.5	_
	_	-		, ,	1	1				4.010.0	0.0	-	- 1	0	-	-	6	٥	10	8	1	~	0	5.5+0.9	5.19+0.58
	-	1 4	0	0	1	, ,	<b>.</b>				4.0	1 (	~	1	~	1	~	t	7	1	2		1	0.8+0.3	
		•			1	1			•	2.140.2	> 1	~	-	2	1	8	-	5		1	4	1	,	1.8+0.7	2.23+0.64
Small Gr (-)	1	5	9	4	12	7	4	8	7	6.7+0.8	B	1	4	9	4	:			,						_
Small Gr (+)	6	~	2	9	9	7	2	7			8		0 0	2 -			9	^	2	0	*	•	5	7.4+0.9	7.09+0.6
Large Gr (-)	8	~	~		1	N	1	. 1	. 1		v		v 4	4 0	٠,	1	*	ν,	2	6	15	00	9	4.8+1.1	5.05+0.7
Large Gr (+)	7	8		ŧ	1	~	1	1		0.7+0 4		1		٧ .	٠,		1	- 0	8	N 1	1		1	1.1+0.4	1.1+0.29
Sarcina						1				1		8			v		8	0		~			,	1.3+0.7	1.0+0.4
Small Gr (-)	1		t	~	1	t	•	1	-	0.4+0.2	0	0	4								,				
Small Gr (+)	1		1	1	1	2	•	1	1	0.2+0.2	2	, ,	) 1	,	,		,	1	1 1	۱ ۱	-	1	~	1 +0.5	
Chain											-												_	,	0.110
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Long Gr (+)	,	*	1	-	1	1	1	1	1	0.1+0.1	7	,		-	-	,		, ,		4 1	- 1		,	2.0+0.0	1.38+0.48
Rods											9												-	1.7.0.0	0.24+0.09
_	21	8	38	12	2	53	34	32	35	28.6+2.7		36 3	33 4	40 3	1 90	18	9	13 4	0	1 21	2	6 96		4.0	
	12	ı	S	٠,	7	6	1	1	4	3.7+1.5						-		1					•	81.146.8	28.05+1.92
Large Gr (-)	11	10	7	16	10	1	10	1	4	7.9+1.6	_	18		, ,	, ,									2.2+0.7	2.81+0.75
large Gr (+)	1		,	8	-	1	1			1.4+0.6				•	Tr.			n (	0	202		20 1	14 110	10.8+1.8	9.57+1.25
Filamentous form										1			-3					v					,	0.3-0.2	0.81+0.3
Gr (-)	1	-	,	-	1	1				0.440.2	-							0					-		
Spiral Gr (-)	2	~		8	5	3	9	5	1	4 240 9	-								. '	. '				0.3+0.5	36
Vibrio Gr (-)	22	9	9	3	5	2	11	10	. 9	A 241 0		-		, ,	n (	1						8		6.0+9.5	\$ +0.64
Quin's oval Gr (-)	1	-		~					, ,	0 7.0 3				n .	· ·		15	~	- ·	9	•	10		6.3+0.9	7.1 ±0.97
Solenomonas Gr (-)	~	5		3	1	1	3	9	, -	2.2.0 6	1 0				-				•	7	-		0	10.0-6.0	0.81+0.2
Oecillospira Gr (-)	8	2	9	1				-		3000			'		4		~	•	\$	2	1	*	2	2.3+0.5	2.29+0.37
The second secon			-							2000		•													

Table (3) The percentage of different types of bacteria in the rumen of goats fod green (Barseem) and dry concentrate (wheat and grain sorghums) ration-

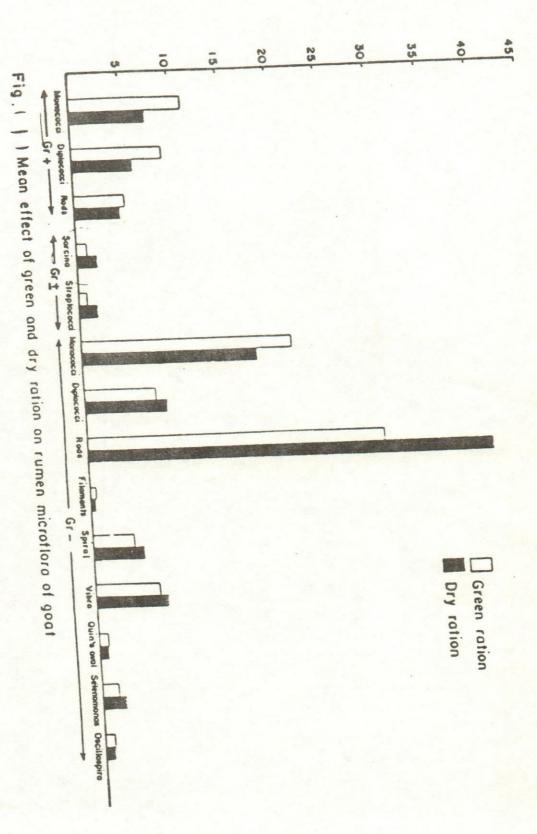
The percentage of different types of bacteria in the rumen of goats fed on green (Barseem) and dry concentrate (wheat bran and grain sorghums) ration Table (3)

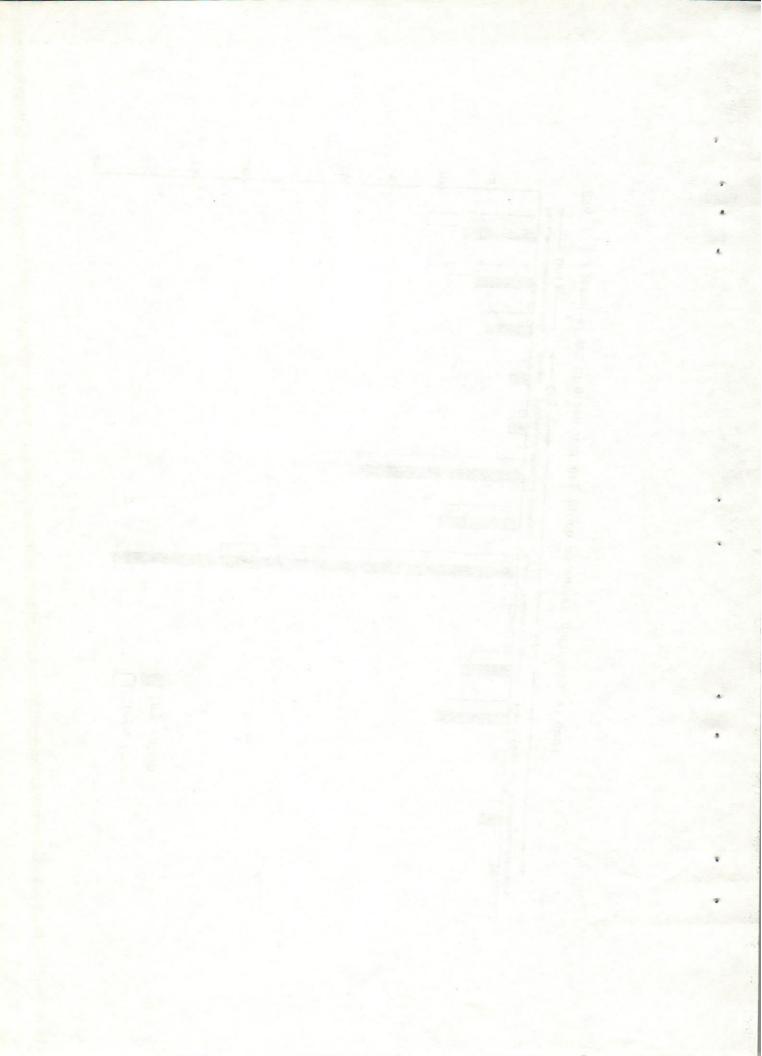
1		1		Gree	ate	Concentr	Dry
-		i m	Bar see m)	and the later of t	2 5	ration	
		Early at morning	Afternoon	Mean	Early	Afternoon	Mean
Mono	Gr+	13.9	11.2	12.5 **	7.6 ±1.45	7.3 ±0.83	7.4 ±0.74
Monococci	Gr-	22.7 ±1.6	20.1	21.4	18.2 ±1.37	18.8	18.2 +0.96
Diplo	GI+	10.2	8.0	9.1 *	6.1 +0.96	6.0 ±1.14	6.1 ±0.73
Diplococci	Gr-	8.7 ±1.0	6.6 ±0.97	7.7 ±0.55	7.7 ±0.83	+0.89	+0.60
Sarcina	Gr+	1.3 ±0.94	1.1	1.1	0.7 ±0.29	1.0	0.8
Strept.	Gr+	0.5 ±0.26	0.8	0.7 ±0.27	2.2	1.2 +0.32	1.7 ±0.51
Rods	Gr+	5.0 ±1.13	4.5	4.8 ±0.77	5.1 +1.42	2.5	3.6 ±0.73
ds	GI-	24.3 ±1.15	35.5**	29.8 +1.91	36.6 +1.99	38.5	37.7* ±1.74
Filament	Gr-	1.1 +0.47	0.2	0.6 +0.22	0.4	0.3 ±0.18	0.4 ±0.13
Spiral	Gr-	3.9 +0.6	3.4 ±0.75	3.7 ±0.47	4.2 ±0.88	5.6 +0.91	5.0 ±0.62
Vibrio	-15	7.6 ±0.8	5.4 ±0.53	6.5 +54	8.2	+0.95	7.1 ±0.95
oval	Gr-	0.3 +0.18	1.4 *	0.8 +0.29	0.7 +0.29	0.9	0.8 ±0.24
monas	Gr-	1.9	2.3 ±0.57	2.12 ±0.39	2.2 ±0.6	+0.5	2.3 ±0.36
spira	Gr-	0.1	0.6 ±0.29	0.4 ±0.15	1.2 ±0.66	0.4 ±0.34	0.8 ±0.34

Afternoon: 6 h. after feeding

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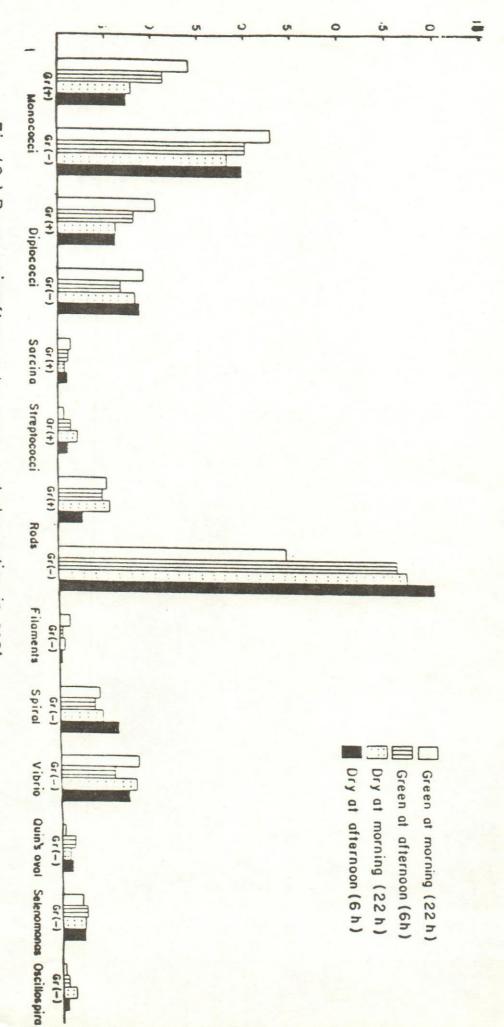


Fig. (2) Rumen microflora under green and dry ration in goat

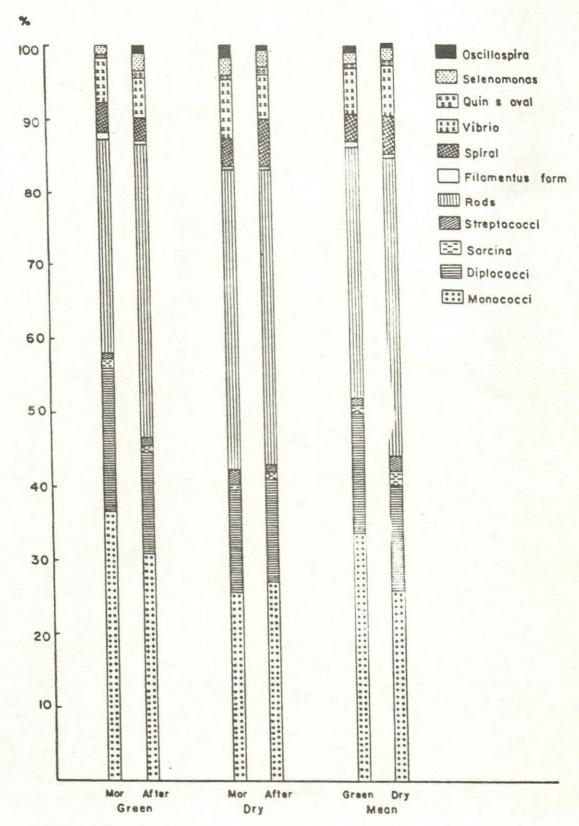


Fig.(3) Changes in the average percent of different types of microflora (bacteria) in rumen content of goat.