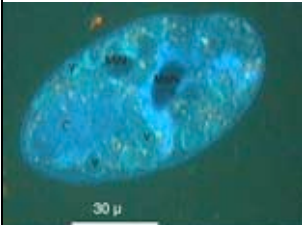
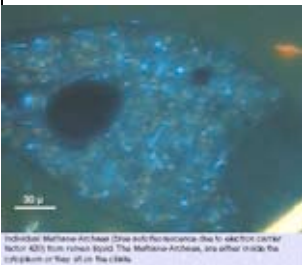


Experiment	Microbial Diversity in the Rumen
Advisor	Kurt Hanselmann, hanselma@botinst.unizh.ch
Reading	Chapters in BBOM 10 th : 19.10, 19.11, 13.4, 14.8 BBOM: Madigan M.T., J.M. Martinko and J. Parker: "Brock - Biology of Microorganisms", 10 th Edition 2003, Prentice Hall.
Objectives	<ul style="list-style-type: none"> • Sampling of microbes from a natural ecosystem • Studying microbial diversity by microscopy • Learning about syntrophic interactions • Determining habitat conditions • Evaluating experiments quantitatively • Linking microbial physiology with biochemistry
Background	<p>Ruminants, e.g. cattle, sheep, camels, llamas, deer etc. are cloven-hoofed, mammals which feed on plant materials. They were domesticated early by humans for meat, milk and other products and they play important roles as mediators in the global carbon cycle. Although they are herbivorous, they lack the glycolytic hydrolases needed to cleave the major plant polysaccharides like cellulose, pectins, hemicellulose and starch. They rely on the glycolytic hydrolase enzymes produced by microorganisms (cellulases, amylases, pectinases etc.), which they harbor in a particular digestive compartment. Digestion in ruminants is achieved by one of the most fascinating but also extremely complex microbial ecosystems, the rumen. The rumen is a large pregastric fermentation chamber present in the digestive tracts of all ruminants. It has a volume of up to 250 l in an adult cow and contains a microbial community consisting of about 10¹¹ microbes per ml of rumen fluid. The microbiota comprises mostly anaerobic bacteria and archaea (about 10⁸ – 10¹¹ ml⁻¹, belonging to more than 200 species), anaerobic ciliated protozoa (10⁴ to 10⁶ ml⁻¹) and anaerobic fungi (10² to 10⁴ zoospores ml⁻¹). They are kept in the host's rumen under more-or-less constant anoxic conditions at redox potentials between -350 and -400 mV, a pH of 6.7 to 6.9 and a temperature of 39°C. Substrates for the rumen microbes are supplied by the feed, which is collected, prepared, conditioned and reconditioned by the host animal while the microbes hydrolyze the plant polymers and ferment the hydrolysis products. Microbial digestion products are removed by diffusion into the blood stream of the host, gases are belched up through the esophagus. Solids, including 500 to 700 g/day of microbial biomass, are further digested in the gastric stomach (the abomasum) and in the small and large intestines. Undigested material leaves the digestive system through the rectum in fecal shapes and consistencies, which are characteristic for particular ruminants, their diets and their digestive systems, mostly the size of the caecum and the large intestine.</p> <p>Digestion in the rumen is carried out by synergistically interacting microbes in an anaerobic food-web symbiosis (figures 1 and 3). First, the plant polysaccharides are enzymatically broken down into di- and monosaccharides, which are then fermented to short chain volatile fatty acids (vFA) like formic, acetic, propionic, butyric acid etc. as well as CO₂, H₂ and CH₄. Depending on the diet, small amounts of lactate and succinate can be produced. Ammonia, branched chain fatty acids and other essential growth factors are produced by several groups of prokaryotes from plant proteins and nucleic acids. Some of the nitrogen is recycled through the ureohepatic cycle and added as urea to the rumen ecosystem with the saliva. While the bulk vFA, which serve as energy sources for the ruminant are taken up and distributed in the body by the blood stream, H₂ and some of the CO₂ are converted by methanogens into CH₄. The metabolic gases are mostly belched into the atmosphere.</p> <p>Through many years of ingenious microbiological research, the prokaryotic composition of the rumen community has been able to be analyzed in great detail. Some of the prokaryotes present and their contribution to the digestion process are listed in table 1.</p>



Rumen ciliate as viewed in the fluorescence microscope. Blue/green is chlorophyll autofluorescence due to the presence of electron carrier compounds (FAD) known to be present in Muller-like ciliates, some monoxenous, some xenotrophic, if present, it is chlorophyll.



Individual Methanogenic Archaea (blue) with fluorescence due to electron carrier factor 420 from rumen fluid. The Methanogenic Archaea, are either inside the ciliophore or free off on the ciliate.

Table 1. Some prokaryotes of the rumen microbiota and selected catabolic abilities

Organisms	Catabolic Abilities												
	1	2	3	4	5	6	7	8	9	10	11	12	13
	Starch and/or Glycogen degradation	Cellulose degradation	Pectin degradation	Hemicellulose degradation	Sugar metabolism, Glycolysis	Acetogenesis	Butyrogenesis	Lactate formation	Succinate formation	Lactate conversion to propionate	Succinate conversion to propionate	Formate formation	Methanogenesis from H ₂ , CO ₂ and formate
<i>Bacteroides amylophilus</i>	X				X	X			X				
<i>Ruminobacter amylophilum</i>	X				X	X			X			X	
<i>Bacteroides ruminicola</i>	X		X		X	X			X	(X)		X	
<i>Succinimonas amylolytica</i>	X				X	X			X	X			
<i>Selenomonas ruminantium</i>	X				X	X		X			(X)		
<i>Selenomonas ruminantium subsp. lactilytica</i>	X				X	X			(X)	X			
<i>Streptococcus bovis</i>	X		X		X			X					
<i>Ruminococcus flavefaciens</i>		X		X	X	X			X			X	
<i>Ruminococcus albus</i>		X		X	X	X						X	
<i>Fibrobacter succinogenes</i>		X		X	X	X			X			X	
<i>Butyrivibrio fibrisolvens</i>		X		X	X	X	X	X				X	
<i>Clostridium lochheadii</i>		X			X	X	X					X	
<i>Lachnospira multiparus</i>			X	X	X	X		X				X	
<i>Lactobacillus spp.</i>					X			X					
<i>Schwartzia succinivorans</i>											X		
<i>Veillonella parvula</i>						X				X	(X)		
<i>Megasphaera elsdenii</i>					X	X	X			X	(X)		
<i>Methanobrevibacter ruminantium</i>													X
<i>Methanomicrobium mobile</i>													X
() some strains only or only by resting cells													

In a cow's rumen **multispecies communities** colonizes grass and hay in a biofilm arrangement which provides an example of the intricate relations between the organisms in a microbial community.

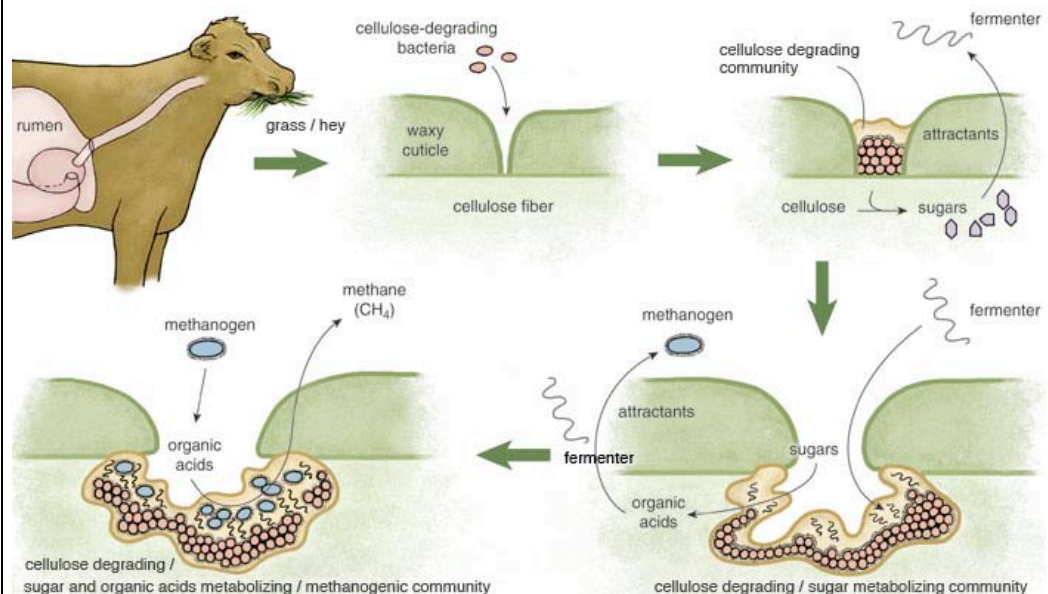


Figure 1 Colonization of fibrous plant polymeric material by hydrolytic, fermenting and methanogenic microorganisms (from: Joe J. Harrison, Raymond J. Turner, Lyriam L. R. Marques and Howard Ceri, 2005, "Biofilms" in American Scientist, Volume 93, 508-515)

The **establishment of a colony** begins with **cellulose-degrading bacteria**, which hydrolyze and partially digest the plant polymers. The mono- and disaccharide sugars produced by these hydrolytic bacteria attract more **fermenting microorganisms**, which convert the sugars into organic acids. In turn, the organic acids attract other fermenters and **methanogenic microbes** that join the community. The organic acids not neutralized by the cow's saliva would normally inhibit further growth in the community, but the methanogens convert some of these molecules (e.g. acetic acid, formic acid) into methane. The entire process produces **a protein-rich microbial cell mass** that can be digested by the cow, providing the bulk of the animal's proteinaceous nutrients.

A cow's cud can be passed between its mouth and the rumen several times before the products are passed to its remaining stomachs and intestines.

Growth in dense communities is slowed down by several factors (figure 2). Bacteria near the center of a microcolony grow slowly because they are exposed to lower concentrations of oxygen and nutrients (1). Intercellular signals (2) can alter the physiology of the entire biofilm community, causing members to expel metabolites from the cells and allow the community to grow on each others metabolites. The **biofilm matrix** is negatively charged (3) and so binds to positively charged ions, metabolites and antimicrobials, slowing their diffusion to other cells within the colony. Populations of **persister cells** (4) do not grow when certain nutrients are lacking or in the presence of an antibiotic, but neither do they die. When the stress is removed, or when the diet changes (e.g. from hay to grass) the persisters (resting cells) can give rise to a normal bacterial colony. Finally, population diversity (5), genetic as well as physiological, improves the chance that some cells will survive any challenge. This guarantees communities with a great **robustness** towards environmental alterations.

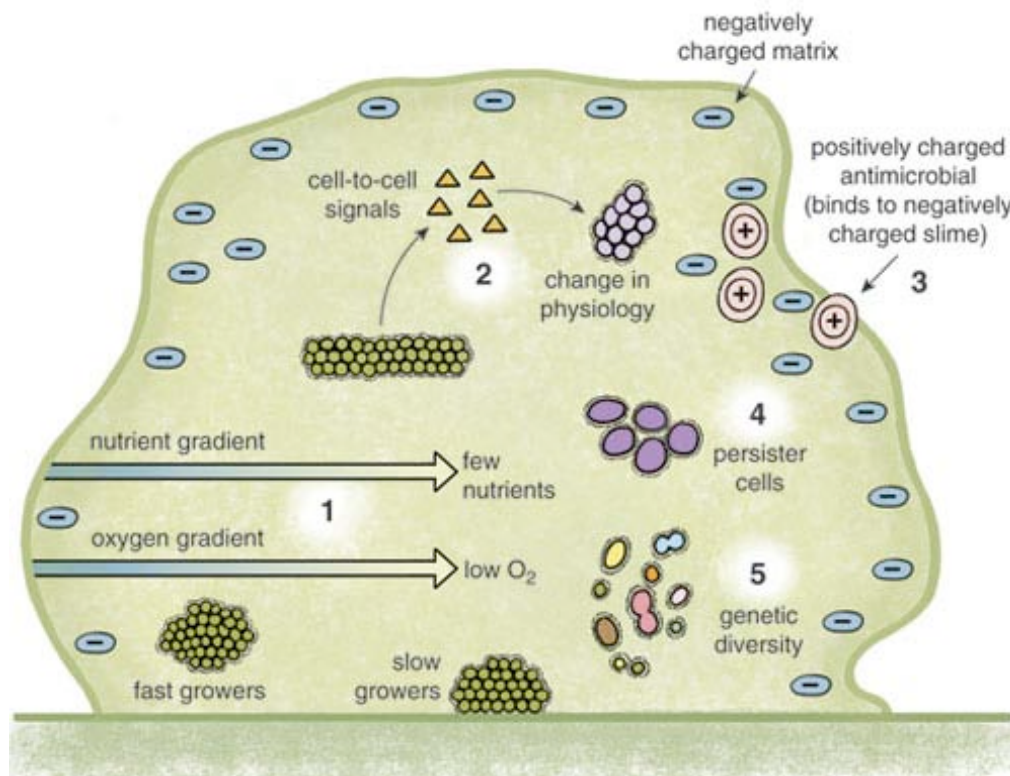


Figure 2 The entire digestion process is achieved by an anaerobic multispecies community which forms biofilm-like colonies embedded in a matrix of exopolymers (EPS) (from: Joe J. Harrison, Raymond J. Turner, Lyriam L. R. Marques and Howard Ceri, 2005, "Biofilms" in American Scientist, Volume 93, 508-515)

Literature

- The chapters of BBOM mentioned above
- Hungate, Robert E., 1966. The Rumen and its Microbes, AP, New York. 533 pgs. A classic describing all aspects of rumen microbiology.
- Dirksen G., 1969. Ist die Methylenblauprobe als Schnelltest für die klinische Pansenuntersuchung geeignet? Deutsche Tierärztliche Wochenschrift 76/12, 305-309.

www. Links

- Images of the exterior and the interior of calf rumens (Penn. State): <http://www.das.psu.edu/dcn/calfmgt/rumen/index.html>
- Anaerobic zoospore fungi of the rumen: http://www.towson.edu/~wubah/Research/Rumen_fungi/rumen_fungi.html
- Rumen physiology and rumination (Colorado State): <http://arbl.cvmbs.colostate.edu/hbooks/pathophys/digestion/herbivores/rumination.html>
- The microbe zoo (MSU): <http://commtechlab.msu.edu/sites/dlc-me/zoo/zacmain.html>

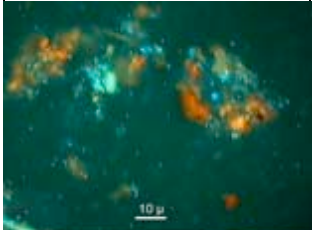
Practical work



Sampling rumen fluid from a fistulated cow at the Veterinary School. The liquid is drawn from the bottom of the cow's rumen and kept as much as possible in a sterile, which must be done in the rumen area. Experiment 1

We will collect rumen fluid from a **fistulated cow** at the Animal Hospital of the Veterinary School, observe the organisms of the **microbiota** by phase contrast microscopy and make a few physiological tests, which will give a qualitative impression about the **habitat conditions** and the **activity** of the rumen microbes.

Material and Experimental Protocols



1. Sampling rumen fluid from a fistulated cow: **sampling technique**

Insert a stiff plastic tubing all the way to the **bottom of the rumen chamber**. Remove rumen fluid by suction and fill it into a sterile 250 ml bottle with a wide neck. Fill the bottle almost completely, leaving only a 2 ml gas space and close tightly. Keep sample at 35 to 40°C.

2. Organismic composition of rumen microbiota: **microscopy**

Prepare a wet mount with the rumen fluid and observe the microbiota in the **phase contrast microscope** beginning with the lowest magnification. You will be able to observe the **ciliates** best at low magnification while you need oil immersion objectives to clearly identify shapes of **bacteria** and **archaea**. Methanogens containing the blue **autofluorescent F₄₂₀** coenzyme are best observed in the fluorescence microscope with filter block A or D.

What happens to the organisms if you aerate a small aliquote of the rumen fluid for a few seconds?

3. Activity of rumen microbiota: **reduction**

Prepare 3 tubes containing 9 ml of freshly collected rumen fluid each. Add 1 ml of glucose stock solution (100mM) to tubes #2 and #3 and the same volume of sterilized water to tube #1. Close the tubes airtight and keep tubes #1 and #2 in the waterbath at 39°C. Tube #3 (loosen cap) is kept in boiling water for 5 minutes before it is transferred to the 39°C water bath. Incubate for 5 minutes. Mix 0.05 ml of a 5 mM **methylene blue** solution to each of the tubes, note the time of the addition and keep in the waterbath at 39°C. Record the time it takes for the methylene blue to lose its color. Compare with the color change in assay #3.

Alternatively, you might use **resazurin** or **phenosafranin** instead of the methylene blue. Why is the color change fastest with methylene blue? And why is it slower with **resazurin** and **phenosafranin**? See also “5. Habitat conditions”.

Methylene blue ($C_{16}H_{18}ClN_3S \cdot 3H_2O$, MW aq. 373.9, Eh +10mV). A 5 mM (approximately) stock solution is prepared by dissolving 14.5 mg methylene blue powder in 10 ml 20 mM phosphate buffer pH 7. This stock solution is 200x concentrated. 50 μl dye stock solution are added to 10 ml culture.

Resazurin ($C_{12}H_7NO_4^- Na^+$, MW 251.2 for the sodium salt, Eh -45 mV for the resorufin / dihydroresorufin couple): 12.56 mg powder dissolved in 10 ml 20 mM phosphate buffer, pH 7 will give a 5 mM dye solution which is 1000x concentrated. The solution is sterilized by filtration and kept in a brown glass bottle.

Phenosafranin Chloride ($C_{18}H_{15}N_4^+ Cl^-$, MW 322.80, Eh -270 mV): 16.14 mg powder dissolved in 10 ml 20 mM phosphate buffer, pH 7 will give a 5 mM dye solution which is 1000x concentrated. The solution is sterilized by filtration and kept in a brown glass bottle.

Glucose monohydrate ($C_6H_{12}O_6 \cdot H_2O$, MW 198): To prepare a 100mM stock solution dissolve 1.98g / 100ml distilled water and autoclave. Add 1 ml to 9 ml rumen culture.

4. Changes of habitat conditions: **acid production**

Prepare 3 tubes containing 9 ml of freshly collected rumen fluid each. Add 1 ml of glucose stock solution to tubes #2 and #3 and the same volume of sterilized water to tube #1. Close the tubes airtight and keep tubes #1 and #2 in the waterbath at 39°C. Tube #3 (loosen cap) is kept in boiling water for 5 minutes before it is transferred to the 39°C water bath. After another 5 minutes, remove a 100 μl aliquote from the culture assay #1 and add to 100 μl bromthymol blue indicator prepared in a well of a white ceramic plate. Record the time it takes for the bromthymol blue to change its color from blue (pH 7.2) to green (pH 6.6) or yellow (pH 6). Repeat with assays #2 and #3. Why is the pH changing during the incubation? How can the color change from green or yellow back to blue after a few minutes be understood?

Bromthymol blue ($C_{27}H_{28}Br_2O_5S$, MW 624.39, pH 7.0, pH 6 yellow pH 7.6 blue) To make a 0.04% (w/v) stock solution dissolve 4 mg of bromthymol blue powder in 0.6 ml 0.01N NaOH and add distilled water to a final volume of 10 ml.

5. Habitat conditions: **redox potential of rumen fluid**

Oxygen, which enters the rumen with the feed, is reduced immediately by reducing

	<p>compounds or through consumption by facultative aerobes. The rumen thus has a constantly low redox potential of -350 to -400 mV, which can be illustrated qualitatively by adding $10\text{ }\mu\text{l}$ of resazurin solution (1000x) to 10 ml of rumen fluid in a completely filled and tightly closed tube. At the concentration used, resazurin ($\text{C}_{12}\text{H}_7\text{NO}_4$) changes its color from purple to pink (resorufin, $\text{C}_{12}\text{H}_7\text{NO}_3$) to the colorless dihydroresorufin, ($\text{C}_{12}\text{H}_9\text{NO}_3$) at a redox potential for the last step of approximately -45 mV. As soon as the redox potential of the rumen fluid becomes low enough, the color of the indicator dye will disappear. Actively metabolizing microbes are able to maintain the low redox potential even in the test tube culture. Phenosafranin is a redox indicator dye with a midpoint potential of -270 mV.</p> <p>You will obtain 5mM solutions of resazurin and phenosafranin with which you should design your redox experiments. Carry out the experiment with 10x diluted rumen contents. Dilute with organism-free rumen fluid which is obtained as supernatant after centrifugation at $20'000g$ for 10 minutes. Observe redox changes and add glucose to stimulate microbial activity if you cannot observe a color change within a few minutes. Incubation temperature is 39°C. How can you check whether or not the redox dye actually responds to reducing and oxidizing conditions?</p> <p>6. Volatile fatty acids (vFA): gas chromatography of the volatile metabolites Since many of the short chain fatty acids are volatile when protonated they can easily be separated, detected and identified by gas chromatography coupled to flame ionization detection (FID). Acidify a 1 ml aliquote of each tube from the “pH-experiment” and inject $100\text{ }\mu\text{l}$ via the injection loop into the gas chromatograph. Follow separation and identify the peaks with vFA standards. Compare with a sample taken from the original rumen fluid.</p> <p>7. Gas production: Design an experiment in which you can follow the production of metabolic gases by the rumen community.</p>
Laboratory Rules & Precautions	Use good laboratory practice ! Do not contaminate yourself others or the laboratory environment. All waste must be sterilized before disposal. It is necessary to work cautiously and, where necessary, aseptically. Wash your hands before you leave the room and disinfect bench surfaces with 70 % ethanol.
Experiences gained	<ul style="list-style-type: none"> • Handling anaerobic microbes • Using the phase contrast microscope and the epi-fluorescence microscope • Learning how to design experiments • Detect microbial activities through product analysis and changes in habitat conditions • Learning how to interpret color changes of redox and pH dyes • Learning how to formulate stoichiometrically correct equations
Timing	90 minutes
Reporting	<ul style="list-style-type: none"> • Make drawings, digital images and video recordings of a few of the microbes observed in the microscope. Do not forget to note the magnification. • Report conclusions from the redox and the pH-indicator experiments. • What vFA are produced during glucose fermentation?

Questions to be answered

(after the lab sessions)

The following exercises will allow you to quantitatively describe some of the microbiological processes, which happen in the rumen.

1. Balancing fermentation

(Consult Fig. 19.28 in BBOM 10th (pg 660) before you attempt to solve this problem)

Hungate observed the production of the following fermentation products in bovine rumen fluid (all values are given in μ moles per hour per gram of rumen contents): acetic acid 20.3, propionic acid 7.1, butyric acid 5.3, carbon dioxide 18.6, methane 7.8

- Determine how much glucose must have been fermented in order to arrive at the quantities of fermentation products observed. Try to reconstruct a balanced stoichiometric equation for this mixed acid glucose fermentation. Electrons, charges and masses of all atoms involved must be balanced.
- Which organisms of the ones listed in table 1 might have been present in the rumen fluid from which the products under (a) were determined?
- What is the fate of the fatty acids produced, what happens to the methane?

2. Fermentation patterns

During digestion of 1044g of hexose monomer ($C_6H_{12}O_6$, MW 180 D) *in vitro* by a rumen microbe culture the following products were detected: 356.3g of acetic acid (MW 60), 155.9g propionic acid (MW 74), and 134.9g butyric acid (MW 88). The gas phase contained predominantly methane and carbon dioxide.

- Derive the balanced stoichiometry for this hexose fermentation process.
- When lactate was added to the rumen fluid one observed an increase in the concentrations of propionate and acetate and the number of *Veillonella alcalescens* cells dramatically increased. How was lactate metabolized? Use the metabolic summary (figure 1) to answer the question.
- One observes elevated concentrations of succinate and propionate but less methane in the rumen microbe culture when *Wolinella succinogenes* cells from a pure culture are added, and the medium is supplemented with fumarate. How can the result of this experiment be explained? (see pg. 619 BBOM 9th or pg 589 BBOM 10th for a short note on the metabolic abilities of *W. succinogenes*)
- When Monensin[®] (an ionophoric antibiotic acting as a growth promotor, sometimes added to the feed of cows), was added to the complete experimental rumen microbe culture, one observed a decrease in the amount of methane formed but an increase in the concentrations of butyrate and propionate. What is the role of Monensin[®] as a feed additive?

3. Syntrophism in fermentation

- Ruminococcus flavefaciens*, a cellulose degrader, produces in an axenic batch culture acetate, formate and succinate in millimolar ratios of 107:62:93 from 100 mMol/l glucose. Hydrogen and carbon dioxide were found in the gas phase, but no other organic metabolites could be detected. Reconstruct and comment the fermentation balance. (Hint: Have a quick look at question 6a below for the enzymes involved in pyruvate conversion)
- Cont. from (3a): If *Methanobrevibacter ruminantium*, a hydrogenotrophic and formatotrophic methanogen (see BBOM 10th pg 454 tab. 13.5. and fig 13.7a), and *Ruminococcus flavefaciens* are cultured together, succinate and acetate are produced in millimolar ratios of 11:189. The gas phase contained CH_4 and CO_2 but no H_2 . Reconstruct the fermentation balance of this mutualistic bacterial community for the degradation of 100 mMol/l glucose.
- Predict the outcome of the experiment in which Monensin[®] will be added to the two-membered community *in vitro*.

	<p>4. The rumen food web</p> <p>a) Which enzymes are needed to hydrolyze the plant polymers, cellulose (beta-1,4-glycosidically linked glucose monomers), starch (alpha-1,4-glycosidically linked glucose monomers), and pectin (a galacturonic acid polymer)?</p> <p>b) What is the role of the ciliates in the rumen ecosystem?</p> <p>5. Host microbe interactions</p> <p>The production of acids during fermentation requires constant neutralization in order to maintain pH-homeostasis. Degradation would quickly be inhibited at pH values below 6. How do ruminants buffer their rumen ecosystem?</p> <p>6. Biochemistry of Pyruvate conversion (these aspects will be studied in detail during the course "Bio-146: Biochemistry and Physiology of Prokaryotes")</p> <p>a) Pyruvate is one of the key intermediates between glycolysis and the different fermentation pathways. Different organisms employ different routes for pyruvate conversion, which leads to the various fermentation patterns. Pyruvate can be converted by anaerobes ...</p> <ul style="list-style-type: none"> • to oxaloacetate by pyruvate carboxylase, • to acetyl~CoA by pyruvate-ferredoxin oxidoreductase, • to acetyl~CoA and formate by pyruvate-formate lyase or • to lactate by lactate dehydrogenase. <p>Which of the organisms listed in tab.1 must contain which of these enzymes?</p> <p>b) What can you find out about propionate production from lactate or succinate from the text book ? (BBOM 10th fig 12.68)</p> <p>c) <i>Ruminococcus flavefaciens</i> excretes succinate, which it can apparently not convert further to propionate. Diagnose this inability from the point of view of the enzymes, which must be lacking.</p> <p>d) By which enzyme do organisms which produce succinate from pyruvate carboxylate pyruvate? How is this carboxylation step achieved enzymatically by organisms which produce propionate from pyruvate via succinate?</p>
<p>Outlook</p>	<p>Experiment 1 will be treated from a bio-thermodynamic point of view in exercise 18. In exercise 13 we will analyzed a few organisms of the rumen community genotypically</p> <p>Exercise 17 will introduce you to system biology by showing you a number of computational models describing interactions between microbes.</p>

Figure 3. Degradation of plant polymers in the rumen (the numbers refer to the catabolic abilities listed in table 1)