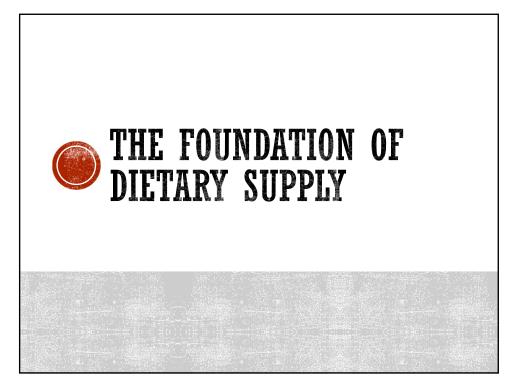
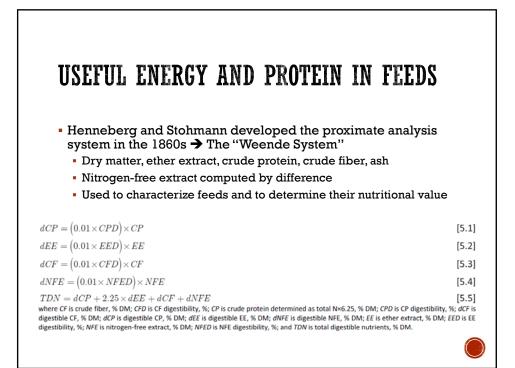
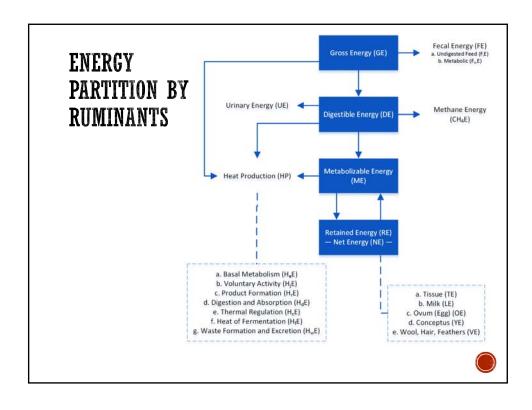


## **APPLICATIONS OF THE RNS**

- extending and refining the use of conventional diet formulation programs
- estimating biological values for feeds for which no nutritive values have been determined
- predicting requirements and balances for nutrients that need more detailed accounting systems
- pragmatically extrapolating research results to varying farm production conditions
- evaluating and diagnosing feeding programs to account for more of the variation in performance under specific production settings
- delivering information that can be used to improve whole-farm nutrient management planning
- providing feedback on the usability and adequacy of the use of applied models (e.g., RNS) so they can be enhanced and reengineered to fit new production situations
- serving as a meta-modeling tool to generate data to be used in other modeling applications



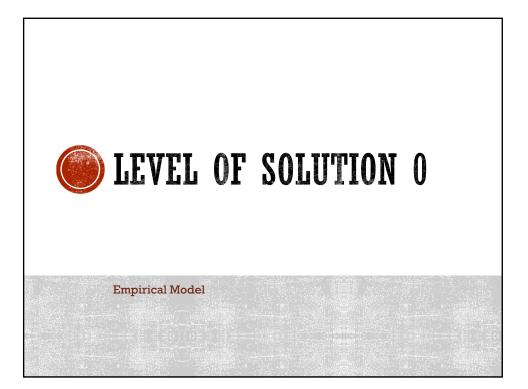




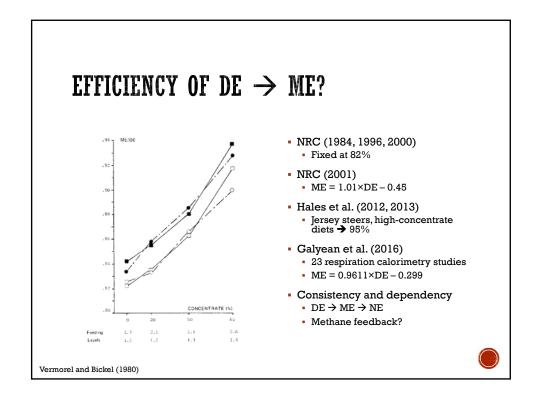
## PREDICTING TOTAL DIGESTIBLE NUTRIENTS

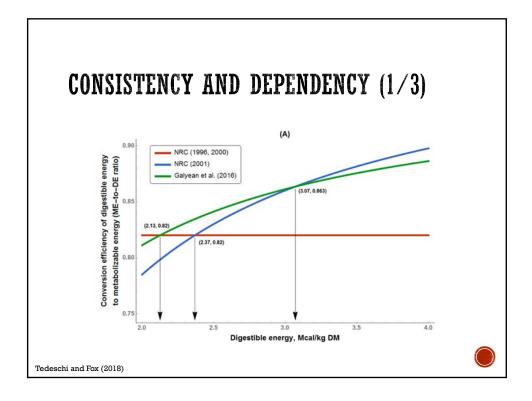
- Digestibility trials using the digestible EE, CP, CF, and NFE
- Empirical equations
- Theoretical equations
- Mechanistic modeling

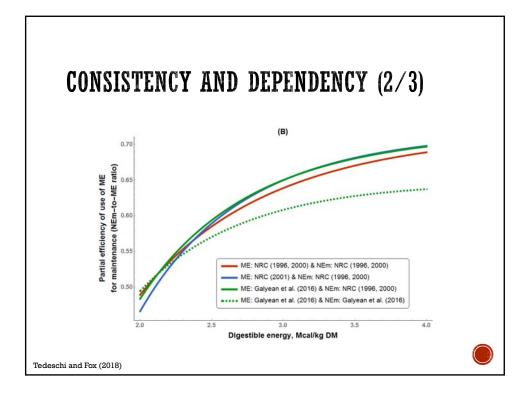
$$\begin{split} dCP &= \begin{pmatrix} 0.01 \times CPD \end{pmatrix} \times CP & [5.1] \\ dEE &= \begin{pmatrix} 0.01 \times EED \end{pmatrix} \times EE & [5.2] \\ dCF &= \begin{pmatrix} 0.01 \times CFD \end{pmatrix} \times CF & [5.3] \\ dNFE &= \begin{pmatrix} 0.01 \times OFD \end{pmatrix} \times NFE & [5.3] \\ dNFE &= \begin{pmatrix} 0.01 \times NFED \end{pmatrix} \times NFE & [5.4] \\ TDN &= dCP + 2.25 \times dEE + dCF + dNFE & [5.5] \\ where CF is crude fiber, \% DM; CFD is CF digestibility, \%; CP is crude protein determined as total Nx6.25, \% DM; CPD is CP digestibility, \%; dCF is digestible CF, % DM; dCF is digestible EE, % DM; dNFE is digestible NFE, % DM; EE is ether extract, % DM; EED is EE digestibility, %; NFE is nitrogen-free extract, % DM; NFE digestibility, %; and TDN is total digestible nutrients, % DM. \end{split}$$

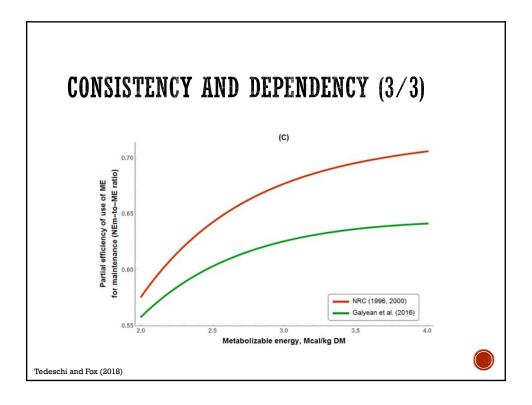


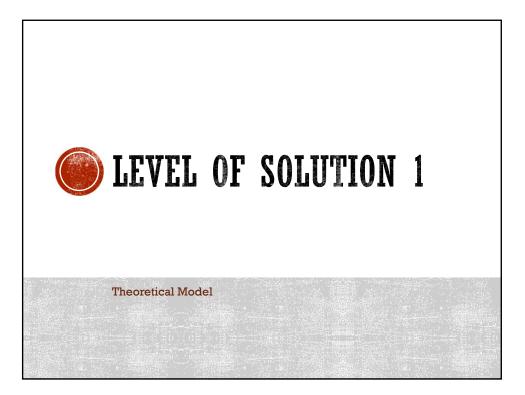
Feed class <sup>(1)</sup>	Total digestible nutrients, %
Dry forages	
(H)	92.464 - 3.338×CF - 6.945×EE - 0.762×NFE + 1.115×CP + 0.031×CF <sup>2</sup> - 0.133×EE <sup>2</sup> + 0.036×CF×NFE + 0.027×EE×NFE + 0.100×EE×CP - 0.022×EE2×CP
(K)	-17.2649 + 1.2120×CP + 0.8352×NFE + 2.4637×EE + 0.4475×CF
Pasture	
(H)	-54.572 + 6.769×CF - 51.083×EE + 1.851×NFE - 0.334×CP - 0.049×CF <sup>2</sup> + 3.384×EE <sup>2</sup> - 0.086×CF×NFE + 0.687×EE×NFE + 0.942×EE×CP - 0.112×EE <sup>2</sup> ×CP
(K)	-21.7656 + 1.4284×CP + 1.0277×NFE + 1.2321×EE + 0.4867×CF
Silages	
(H)	-72.943 + 4.675×CF - 1.280×EE + 1.611×NFE + 0.497×CP - 0.044×CF <sup>2</sup> - 0.760×EE2 - 0.039×CF×NFE + 0.087×EE×NFE - 0.152×EE×CP + 0.074×EE <sup>2</sup> ×CP
(K)	-21.9391 + 1.0538×CP + 0.9736×NFE + 3.0016×EE + 0.4590×CF
Energy feeds	
(H)	-202.686 - 1.357×CF + 2.638×EE + 3.003×NFE + 2.347×CP + 0.046×CF <sup>2</sup> + 0.647×EE <sup>2</sup> + 0.041×CF×NFE - 0.081×EE×NFE + 0.553×EE×CP - 0.046×EE <sup>2</sup> ×CP
(K)	40.2625 + 0.1969×CP + 0.4228×NFE + 1.1903×EE - 0.1379×CF
Protein feeds	
(H)	-133.726 - 0.254×CF + 19.593×EE + 2.784×NFE + 2.315×CP + 0.028×CF <sup>2</sup> - 0.341×EE <sup>2</sup> - 0.008×CF×NFE - 0.215×EE×NFE - 0.193×EE×CP + 0.004×EE±×CP
(K)	40.3227 + 0.5398×CP + 0.4448×NFE + 1.4218×EE - 0.7007×CF

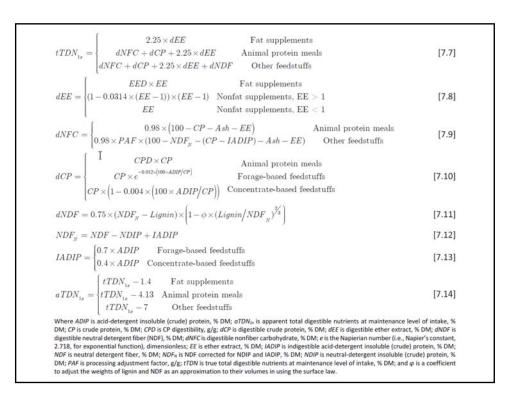


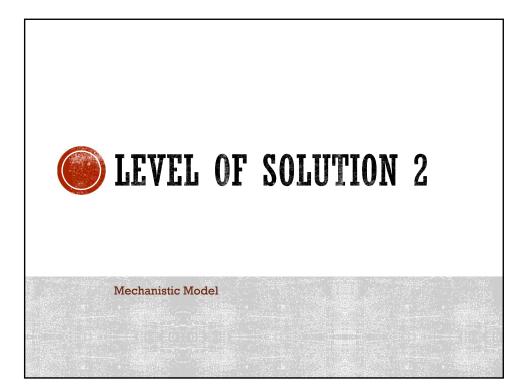






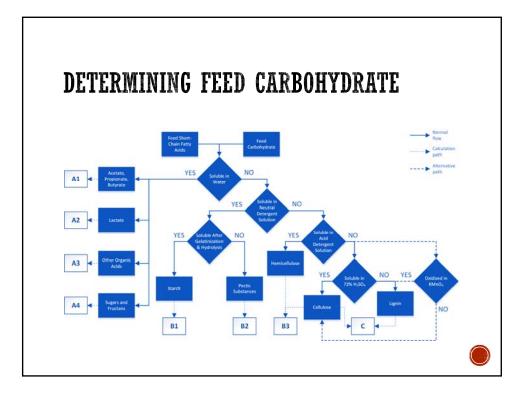


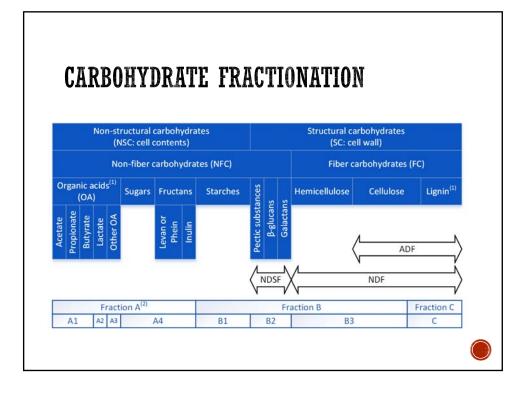


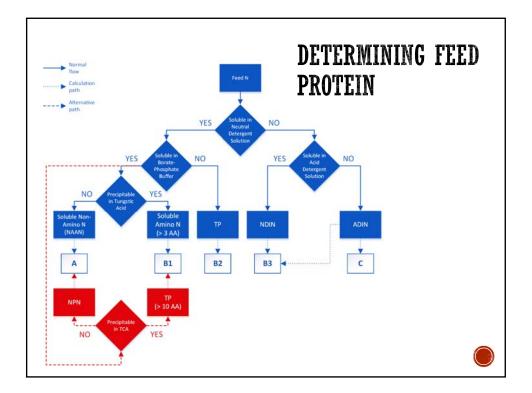


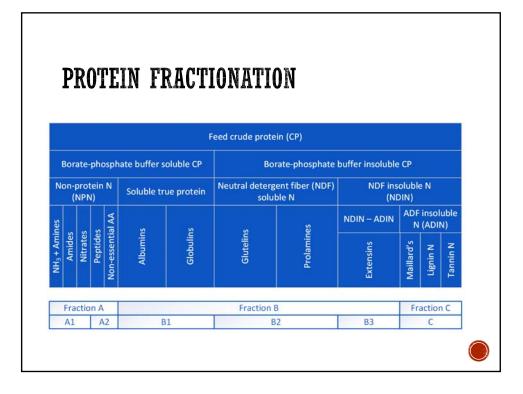
## **MECHANISTIC MODELING**

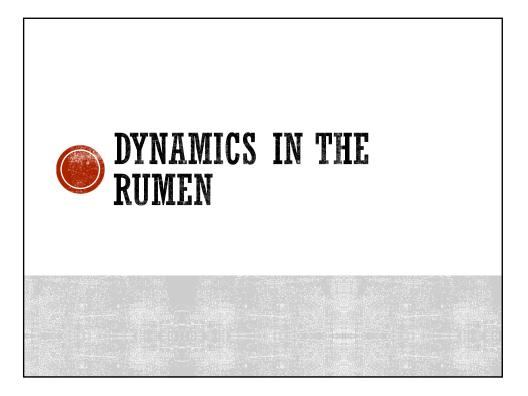
- Nutrient fractionation
  - Carbohydrate
  - Protein
- Bacteria submodel
  Deficiency of N or Branched-Chain AA
- Ruminal degradation and passage fractional rates
- Ruminal pH submodel
- Intestinal digestibility
  - Midgut (small intestine)
  - Hindgut (large intestine)
- Fecal matter
- Digestibility and TDN

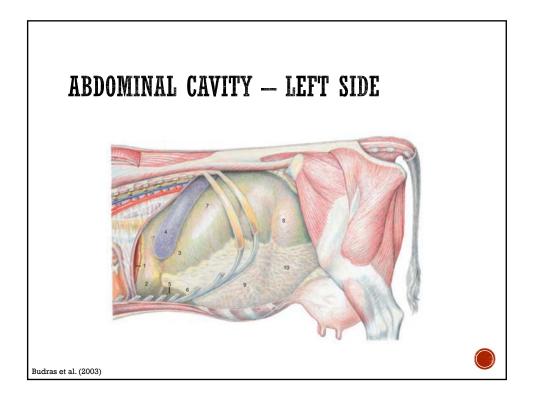


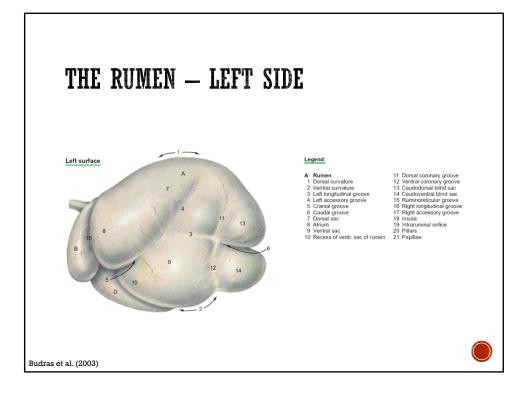


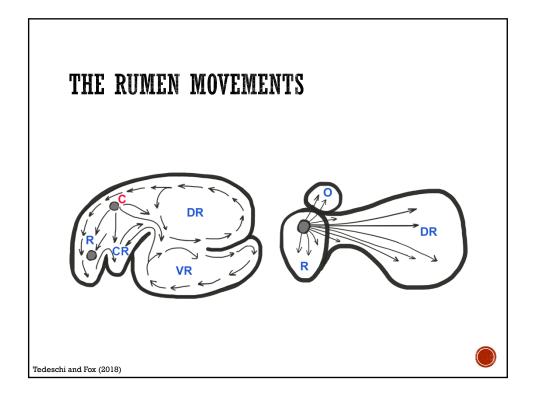


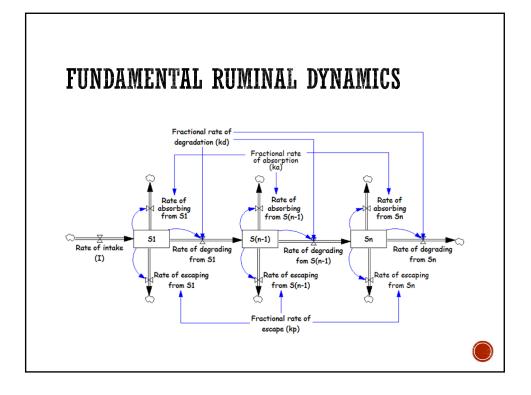


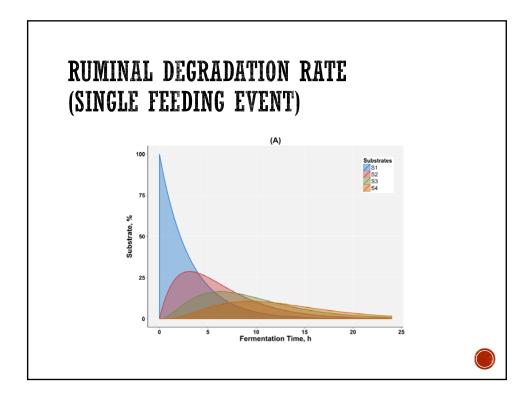


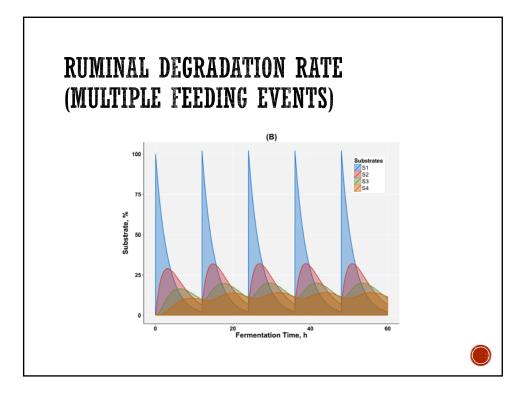






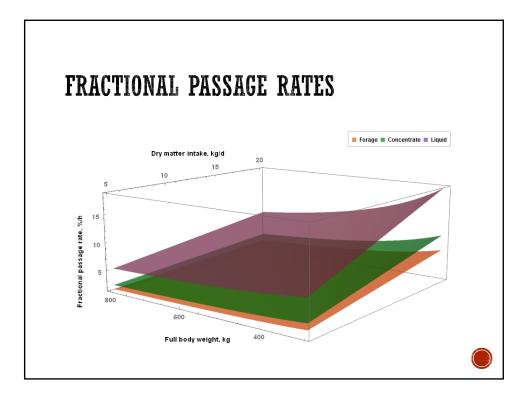


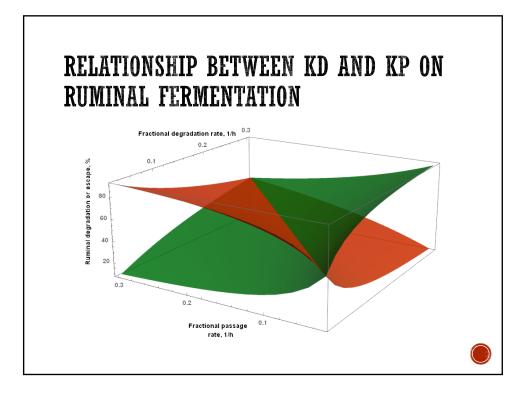


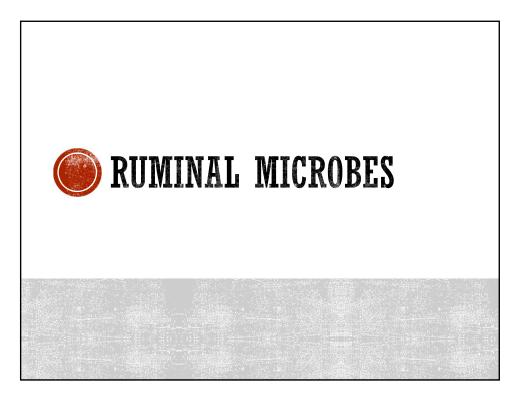


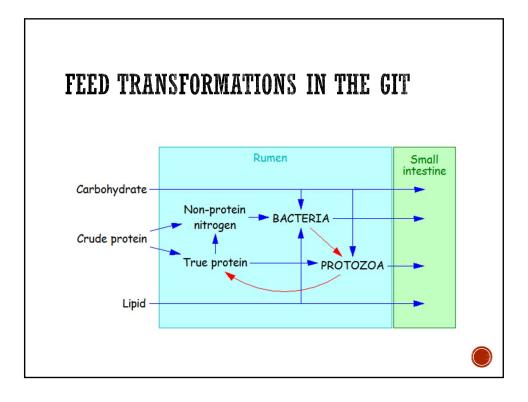
## FRACTIONAL PASSAGE RATES

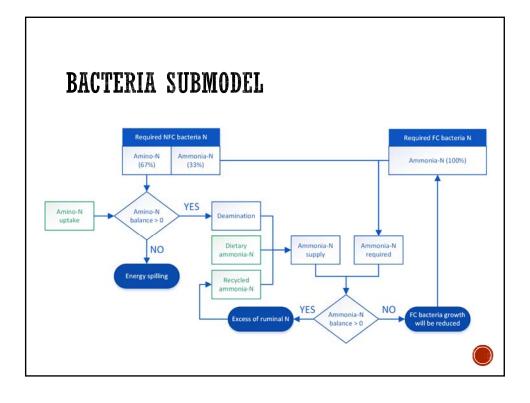
$kp_{\textit{Forage}} = 2.365 + 21.4 \times \textit{Ff} \times \textit{DMI} / \textit{FBW} + 73.4 \times \left(1 - \textit{Ff}\right) \times \textit{DMI} / \textit{FBW} + 0.069 \times \textit{Ff} \times \textit{DMI}$	[8.140]
$kp_{\textit{Liq}} = 1.169 + 137.5 \times \textit{Ff} \times \textit{DMI} \big/ \textit{FBW} + 172.1 \times \big(1 - \textit{Ff}\big) \times \textit{DMI} \big/ \textit{FBW}$	[8.141]
$kp_{\textit{Lig}} = 4.524 + 22.3 \times \textit{Ff} \times \textit{DMI} / \textit{FBW} + 204.6 \times (1 - \textit{Ff}) \times \textit{DMI} / \textit{FBW} + 0.344 \times \textit{Ff} \times \textit{DMI}$	[8.142]
Where <i>DMI</i> is dry matter intake, kg/d; <i>FBW</i> is full body weight, kg; <i>Ff</i> is the fraction of diet forage, dimensionless; and kp <sub>rompe</sub> , kp fractional passage rates, %/h.	

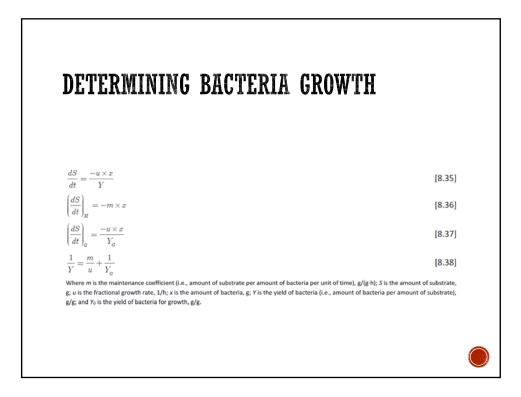


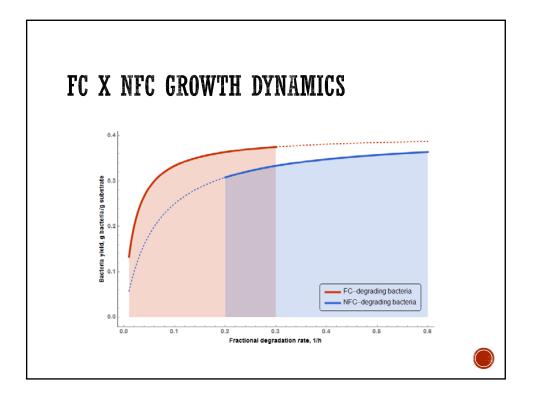


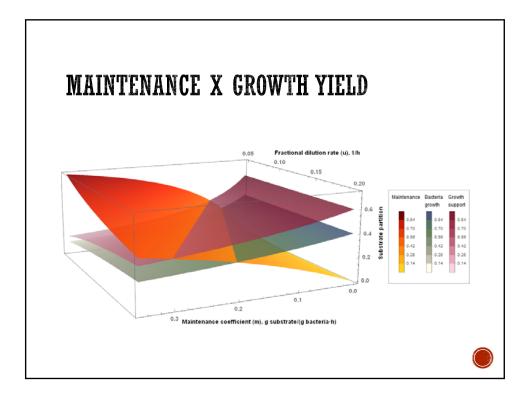


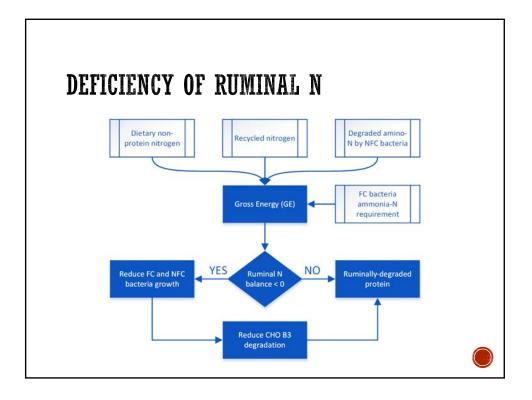


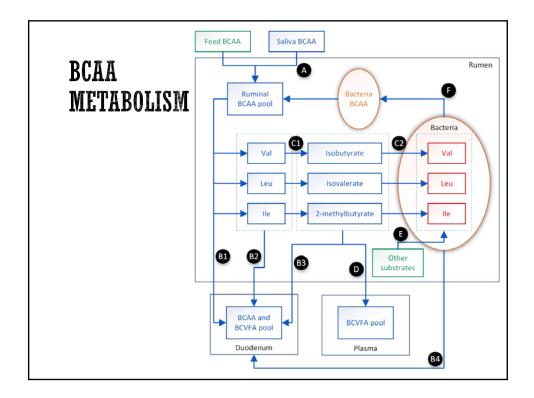


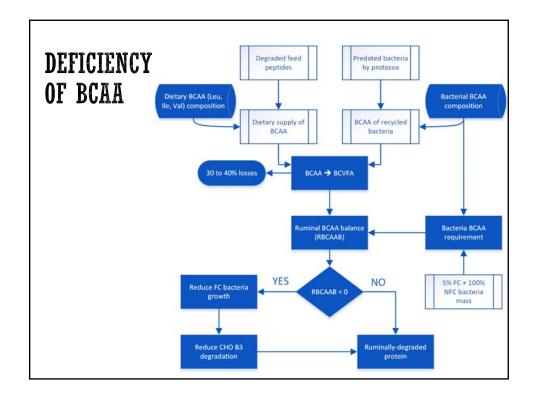


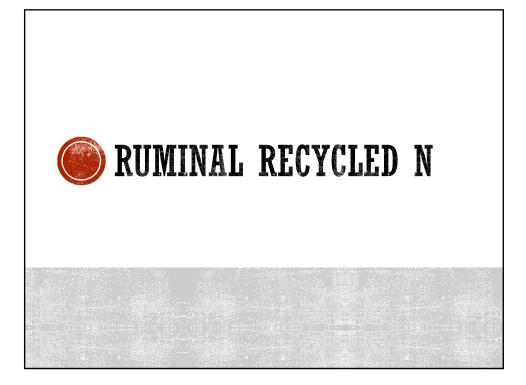


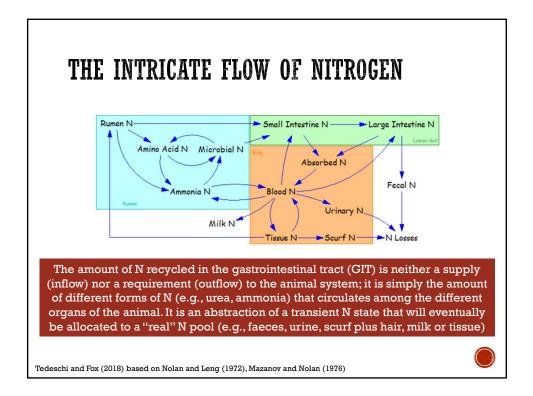














- UUA is the amount of recycled N that is effectively used by the ruminal microbes, but two sources of N retention (Lobley et al. 2000):
  - synthesis of nitrogenous compounds by the GIT microbes and
  - anabolic reactions of ammonia within the body of the ruminant animal
- The amount of recycled N to the GIT increases as N intake decreases (Eisemann and Tedeschi 2016), and about 54% to 57% of the recycled N is used for the synthesis of microbial protein (Marini and Van Amburgh 2003; Reynolds and Kristensen 2008)

