were corrected and the installation program was recompiled to support 64-bit Windows installations. We also created an installation program to allow the 2000 Beef NRC to be operated on modern Windows based computers. Both are available on the website. We are working to develop a review of techniques used to evaluate models, an article listing guidelines for reporting dietary information in publications, and an article describing a software platform that could be used across NRC species requirement models. We are also developing a database to store observational data collected by the NRC committees which can be downloaded and used for model development. Future work includes development of a web based mechanism to provide access to the model source code for use by the animal nutrition research community, collection of feedback from model users to help identify software problems and prediction problems, summarization of evaluations of the current NRC models, and an assessment of supply and requirement model subcomponents that could be used across species. These changes will facilitate the use of NRC models in diet formulation, update NRC models more rapidly as new information becomes available and at lower cost, and improve efficiency of feed use and sustainability in animal agriculture.

Key Words: NRC, nutrient requirements, nutrition model

Effects of stage of gestation and diet on maternal fat T56 deposition. P. P. Rotta*1, S. C. Valadares Filho1, T. R. Santos1, L. F. Costa e Silva¹, M. I. Marcondes¹, B. C. Carvalho², A. A. G. Lobo¹, J. V. F. Souza¹, M. A. S. Novaes¹, M. F. L Ferreira¹, and J. S. A. A. Santos¹, ¹Universidade Federal de Vicosa, Vicosa, Brazil, ²Empresa Brasileira de Pesquisa Agropecuaria, Brazil.

The objectives were to evaluate the maternal nutrient intake and stage of gestation on mesentery, kidney, pelvic and heart (KPH) fat and fat thickness growth rate in Holstein × Gyr cows. Sixty-two multiparous cows were inseminated with the same Gyr semen and in the d 30 the gestation was confirmed in 44 cows. Six cows non-pregnant were previously slaughtered as reference group. The 44 pregnant cows were randomly distributed in four time of gestation: 140, 200, 240 and 270 days. Thus, 11 cows were slaughtered in each time. These eleven cows were randomly allocated in two different diets: maintenance level (n = 6; 1.1% of body weight) and ad libitum (n = 5). The cows that achieved the time of gestation were slaughtered and necropsied. The mesentery was removed of the viscera and weighted. The KPH fat was removed and weighted. The fat thickness was measured using an electronic caliper rule 18 h after the slaughter. The values are given in function of final body weight. Data were analyzed using MIXED procedure of SAS. Data of non-pregnant cows were used considering as initial time. The growth rate of mesentery for cows fed ad libitum was exponential and the estimated equation was: $y = 2.7808 + 3.9181 \times [1 - \exp(-0.00726)]$ × days)]. However, the growth rate of mesentery for cows fed at maintenance level was linear and the estimated equation was: y = 2.8351 + (0.00484 × days). In the same way, the KPH fat was exponential to cows fed ad libitum: $y = 1.0812 + 6.5984 \times [1 - \exp(-0.00172 \times$ days)] while the estimated equation to cows fed at maintenance level was linear: $y = 1.1236 + (0.002572 \times days)$. The fat thickness was also exponential to cows fed ad libitum: $y = 0.8033 + 0.0111 \times [exp(0.0186)]$ × days)] in comparison to cows fed at maintenance level, which estimated equation was linear: $y = 0.757 + (0.002986 \times days)$. Thus, cows fed ad libitum present exponential growth rate of fat deposition, while cows fed at maintenance level present linear growth rate. This can be explained by the high dry matter intake observed to this group and the higher average daily gain. In this way, the excess of energy and protein are being deposited as fat.

Key Words: fat thickness, KPH fat, mesentery

Effects of different diets on the gene expression of enzymes related to fatty acid synthesis in the mammary gland of lactating dairy cows. H. Zhang, C. Ao*, L. Song, E. Khas, and X. Zhang, Department of Animal Science of Inner Mongolia Agricultural University, Huhhot, Inner Mongolia, China.

The objectives of this study were to determine the effects of three different diets on the genes expression of key enzymes involved in fatty acid synthesis in the mammary gland of dairy cows. Thirty multiparous Holstein cows (BW = 566 ± 19.6 kg, DIM = 95 ± 24 d) were assigned to a single factor block design with 30 d period. Animals were fed diets with different roughage and same concentrate profiles, the concentrate-roughage ratio in the treatments were different. Diets (on DM base) were: (1) hay (4%), corn silage (27%) and alfalfa (23%) with additional 46% of concentrate (MF; CP: 17.45%, NDF: 38.33% and NEL: 1.6 Mcal/kg); (2) corn straw (35%) and 65% of concentrate (CSA; CP: 17.14%, NDF: 41.58% and NEL: 1.58Mcal/kg); (3) corn straw (54%) and 46% of concentrate (CSB; CP: 13.80%, NDF: 55.59% and NEL: 1.36 Mcal/kg). Mammary gland tissue (500 mg/animal) biopsies were randomly performed for half numbers of cows in each group after milking on the last day of experiment. Gene expressions for key enzymes involved in fatty acids synthesis were determined by RT-PCR. Statistical analysis was performed using the PROC MIXED procedure of SAS 9.0. Results showed that mRNA abundance for ACACA and FASN which involve in de novo fatty acid synthesis was higher (P < 0.05) in MF compared to CSA and CSB. Gene expression for fatty acid desaturase SCD was significantly increased (P < 0.01) by MF treatment and gene expression for fatty acid uptake and intracellular trafficking enzymes LPL, ACSL1 and CD36 were increased (P < 0.05) also by MF. However, no effects were observed for milk fat synthesis regulator genes PPARA, PPARG, SREBF1 (P > 0.05). No effects on the gene expressions for all above enzymes and regulators between CSA and CSB were found. The data indicated that diet with high quality roughage and the same concentrate-roughage can increase the gene expression of enzymes related to fatty acid synthesis and desaturation, but with the same roughage and different concentrateroughage ratio had no effect on milk fat synthesis key genes in the mammary gland of dairy cows.

Key Words: dairy cow, mammary gland, fatty acid synthesis

Ven SP 6368 Effects of stage of gestation and diet on dairy cow placentomes. P. P. Rotta*1, S. C. Valadares Filho1, T. R. Santos1, L. F. Costa e Silva1, M. I. Marcondes1, M. M. Campos2, F. A. S. Silva1, J. R. Oliveira, A. C. B. Menezes, E. C. Martins, and F. A. C. Villadiego, ¹Universidade Federal de Vicosa, Vicosa, Brazil, ²Empresa Brasileira de Pesquisa Agropecuaria, Brazil.

The objectives were to evaluate the maternal nutrient intake and stage of gestation on caruncular, cotyledon and placentoma growth and number rate in Holstein × Gyr cows. Sixty-two multiparous cows were inseminated with the same Gyr semen and in the d 30 the gestation was confirmed in 44 cows. Six/cows non-pregnant were previously slaughtered as reference group. The 44 pregnant cows were randomly distributed in four time of gestation: 140, 200, 240 and 270 days. Thus, 11 cows were slaughtered in each time. These eleven cows were randomly allocated in two different diets: maintenance level (n = 6; 1.1% of body weight) and ad libitum (n = 5). The cows that achieved the time of gestation were slaughtered and necropsied. The gravid uterus was removed and the uterus was separated from placenta. In the uterus, all carunculars were removed, counted and weighted. In the placenta, all cotyledons were removed, counted and weighted. Except for placentomas number, that/consider 140 days of pregnancy at minimum, the

J. Anim. Sci. Vol. 91, E-Suppl. 2/J. Dairy Sci. Vol. 96, E-Suppl. 1

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