

## ANEVIS supplementation optimizes many health factors in growing beef cattle

### Background

Niacin plays important roles as a cofactor in lipid, protein, and carbohydrate synthesis. Therefore, supplementation may be justified when high forage levels result in lower niacin concentrations. Rumen-protected niacin (Anevis) has been widely used in dairy cattle due to its numerous metabolic advantages, such as its vasodilatory, antioxidative, and antilipolytic effect. In this study, we investigated the effect of supplementation with Anevis on beef cattle liver transcriptome (i.e., genes present in the DNA).

### Design

A group of 6 Angus × Simmental, weaned steers (n = 4) and heifers (n = 2) with average body weight (BW; 299 ± 7 kg) and age of 7–9 months old were randomly allocated in two groups based on dietary treatment: Anevis (n = 3), and Control (n = 3). After a training period of ~10 d, animals were successfully adapted to Calan gates. The diet offered was *ad libitum* Bermudagrass hay combined with a nutritional supplement composed of 1.61 kg of endophyte-free tall fescue seeds, and 1.61 kg of pellets composed of 46.5% ground corn, 46.5% soybean meal, 5% wheat middlings, and 2% soybean oil; and 0.1 kg of molasses per animal per day. Anevis was supplemented at 6 g/h/day (4.2 g niacin). Liver samples were obtained 30 days after the beginning of the supplementation with Anevis. The total RNA of liver samples was extracted and analyzed through RNA-sequencing; a technique that determines the level of expression of differentially expressed genes (DEG) in the whole genome (DNA). DEG were clustered based on their related metabolic pathways (KEGG pathway; Figure 1).

### Results

A list of 1,192 genes were differentially expressed between treatments. These DEG were significantly altered by administration Anevis as compared with the Control. They resulted in alterations to the following metabolic pathways:

#### Downregulation of arachidonic acid metabolism pathway

- Inhibitory effect of niacin on lipolysis.
- Anevis supplementation may improve liver health status by preventing hepatic lipid accumulation.

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#### Downregulation of Cholesterol metabolism

- Inhibitory effect of niacin on total cholesterol decreasing metabolic pathways.
- Anevis supplementation downregulates genes that encode for proteins of glycan biosynthesis, which might suggest that glycation could potentially be inhibited, leading to decreased VLDL and LDL synthesis, which is beneficial to liver health.

#### Increased the expression of genes that are critical regulators of liver size

- The decrease in organ size could be explained also by the lipolytic effect of niacin in liver adipose tissue content.

#### Promotion of anti-inflammatory effects

- Mediated via HCA2-dependent mechanisms in monocytes and macrophages.
- Inhibition of adhesion and accumulation in adipose tissue by oxidized LDL.
- Inhibition of angiotensin II-induced ROS production in vascular endothelium.

#### Promotion of a vasodilator effect

- Niacin derived activation of genes that attenuate the increment of hepatic vascular resistance.
- Anevis supplementation produced an inhibition of the Phospholipase D pathway, which causes vasoconstriction and regulates blood pressure.

#### Causing an inhibition on cell migration through actin reorganization and cell proliferation

- Niacin interferes with the signaling cascade of chemo-attractants in macrophages.
- This leads to the macrophage proinflammatory responses of niacin.

#### Inhibition of inflammation and increased activation of angiogenesis

- Observed in the HIF1 pathway (an oxygen-sensing metabolic pathway).
- Anevis activates an early response gene involved in hepatic tissue proliferation and liver regeneration.
- The FOXO signaling pathway was downregulated, potentially inhibiting oxidative stress resistance and DNA repair, glucose metabolism and immunoregulation.



# ANEVIS™

Rumen-protected niacin delivered to where the cow needs it most.



## Potential antifibrotic properties

- Anevis also inhibited the conversion of mechanical forces into biochemical signals through the RAP1 pathway, leading to attenuation of collagen accumulation exerting of niacin.

## Downregulation of Insulin signaling suggests an improved insulin sensitivity

- Downregulation of G6PD, which encodes for a key enzyme involved in the last step of gluconeogenic and glycogenolytic pathways, suggesting gluconeogenesis inhibition.
- Anevis treatment is associated with an increase of adiponectin, an important insulin-sensing hormone.

## **Conclusion**

Anevis supplementation presented a significant list of potential benefits observed at the liver transcriptomics level. Several metabolic pathways revealed positive effects of Anevis. The most impacted pathways showed that rumen-protected niacin had a down-regulatory effect on the expression of genes related to lipolysis, inflammatory responses, atherosclerosis, oxidative stress, and fibrosis, and enhancing vasodilation. Therefore, results from our study could potentially promote supplementation of rumen-protected niacin on beef cattle backgrounding operations or new arrivals to a feedlot, especially during the acclimation period when health status is often compromised. Finally, it is important to remark that our study seeks to bring light on the specific role of niacin in growing beef cattle, and caution must be exercised when translating our findings to other species or cattle breeds (i.e., transition dairy cows).

## **Reference**

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**Figure 1.** Results of flux and impact uncovered by Kyoto Encyclopedia of Genes and Genomes (KEGG) database analysis of the bovine liver transcriptome of growing beef cattle supplemented with ANEVIS.

| KEGG category  | KEGG subcategory                                    | KEGG pathway  | Impact   | Flux       |
|--|---|---|----------|------------|
| <a href="#">Metabolism</a>                           | <a href="#">Lipid Metabolism</a>                    | <a href="#">Arachidonic acid metabolism</a>             | Blue bar | Green bar  |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">VEGF signaling pathway</a>                  | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">HIF-1 signaling pathway</a>                 | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">TNF signaling pathway</a>                   | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">Notch signaling pathway</a>                 | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">FoxO signaling pathway</a>                  | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">Rap1 signaling pathway</a>                  | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signal Transduction</a>                 | <a href="#">Phospholipase D signaling pathway</a>       | Blue bar | Yellow bar |
| <a href="#">Environmental Information Processing</a> | <a href="#">Signaling Molecules and Interaction</a> | <a href="#">ECM-receptor interaction</a>                | Blue bar | Green bar  |
| <a href="#">Cellular Processes</a>                   | <a href="#">Cellular community - eukaryotes</a>     | <a href="#">Focal adhesion</a>                          | Blue bar | Green bar  |
| <a href="#">Cellular Processes</a>                   | <a href="#">Cell Motility</a>                       | <a href="#">Regulation of actin cytoskeleton</a>        | Blue bar | Yellow bar |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">Fc epsilon RI signaling pathway</a>         | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">B cell receptor signaling pathway</a>       | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">T cell receptor signaling pathway</a>       | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">Toll-like receptor signaling pathway</a>    | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">Chemokine signaling pathway</a>             | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Immune System</a>                       | <a href="#">Neutrophil extracellular trap formation</a> | Blue bar | Yellow bar |
| <a href="#">Organismal Systems</a>                   | <a href="#">Endocrine System</a>                    | <a href="#">Thyroid hormone signaling pathway</a>       | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Endocrine System</a>                    | <a href="#">Regulation of lipolysis in adipocytes</a>   | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Endocrine System</a>                    | <a href="#">Adipocytokine signaling pathway</a>         | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Endocrine System</a>                    | <a href="#">Insulin signaling pathway</a>               | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Digestive System</a>                    | <a href="#">Carbohydrate digestion and absorption</a>   | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Digestive System</a>                    | <a href="#">Protein digestion and absorption</a>        | Blue bar | Green bar  |
| <a href="#">Organismal Systems</a>                   | <a href="#">Digestive System</a>                    | <a href="#">Cholesterol metabolism</a>                  | Blue bar | Green bar  |

'Impact' represents the change in the expression of the genes belonging to a specific pathway due to the supplementation of ANEVIS; and 'flux' as the report of the average direction in the expression as downregulation, upregulation, or neutral or no change. Flux represents the direction of each subcategory belonging to 'Metabolism' KEGG category: green color represents inhibition, yellow neutrality, whereas red color shows activation. Color intensity depicts flux level. Blue lines show the impact of each category and the corresponding subcategory (P value < 0.05).

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