NAD Metabolism in Male Reproductive Aging

Mirella L. Meyer-Ficca
Utah State University, Mirella.Meyer@usu.edu

Follow this and additional works at: https://digitalcommons.usu.edu/funded_research_data

Part of the Dairy Science Commons, and the Veterinary Medicine Commons

Recommended Citation

This Grant Record is brought to you for free and open access by the Data Services at DigitalCommons@USU. It has been accepted for inclusion in Funded Research Records by an authorized administrator of DigitalCommons@USU. For more information, please contact digitalcommons@usu.edu.
Resource Sharing Plan

This proposal does not involve clinical data sets. When resources have been developed with NIH funds and the associated research findings published or provided to NIH, they will be made available for research purposes to qualified individuals (e.g., other NIH-funded researchers) within the scientific research community.

1. Data Sharing Plan
Data generated by the proposed research will be shared via the conventional mechanisms of presentations at scientific meetings and publication in peer-reviewed scientific journals according to the NIH Public Access Policy, as described in the NIH Data Sharing Policy and Implementation Guidance. Peer-reviewed journal articles produced by the proposed research will be submitted to the digital archive PubMed Central.

2. Sharing Model Organisms
Model organisms and materials developed with NIH funds and the associated research findings published or provided to NIH will be made available to qualified individuals (e.g., all other NIH-funded researchers) within the scientific community via applicable Utah State University Material Transfer Agreements and/or licensing agreements through the Office for Technology Commercialization, making them available to the broader research community.

3. Functional genomics data
Functional genomics data and other large data sets generated with NIH funds and their associated activities will be made available to the scientific research community. This will be accomplished by submission of such data sets to the appropriate NIH-designated data repositories such as the NCBI Gene Expression Omnibus (GEO) for public access.