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The authors have nothing to disclose

Background

Insulin/insulin like-growth factor-I (IGF-I) signaling plays a critical role in central glucose bioavailability and lipid metabolism. An increase in glucose disposal can generate reducing agents through the pentose-phosphate pathway necessary for the synthesis of free fatty acids (FFA) and recent evidences suggest that nicotinamide adenine dinucleotide (NAD⁺) depletion leads to abnormal lipid metabolism. Disturbances in lipid synthesis are related to the appearance of insulin resistance and diabetes. The insulin receptor substrate (IRS) 2-deficient mice (IRS2^{-/-}) is an excellent model to study the development of diabetes as a high proportion of them present an abrupt increase in glycemia, showing similarities with type 1 diabetes. However, the molecular mechanisms involved in the onset of diabetes in IRS2^{-/-} mice remain to be elucidated.

Hypothesis and objective

We hypothesized that alterations in insulin/IGF-I signaling could be related to the incidence of diabetes in IRS2^{-/-} mice. Thus, we analyzed the changes in hypothalamic signaling and its relationship with fatty acid metabolism in this model.

Animals and methods

Animals

Eighteen male mice:

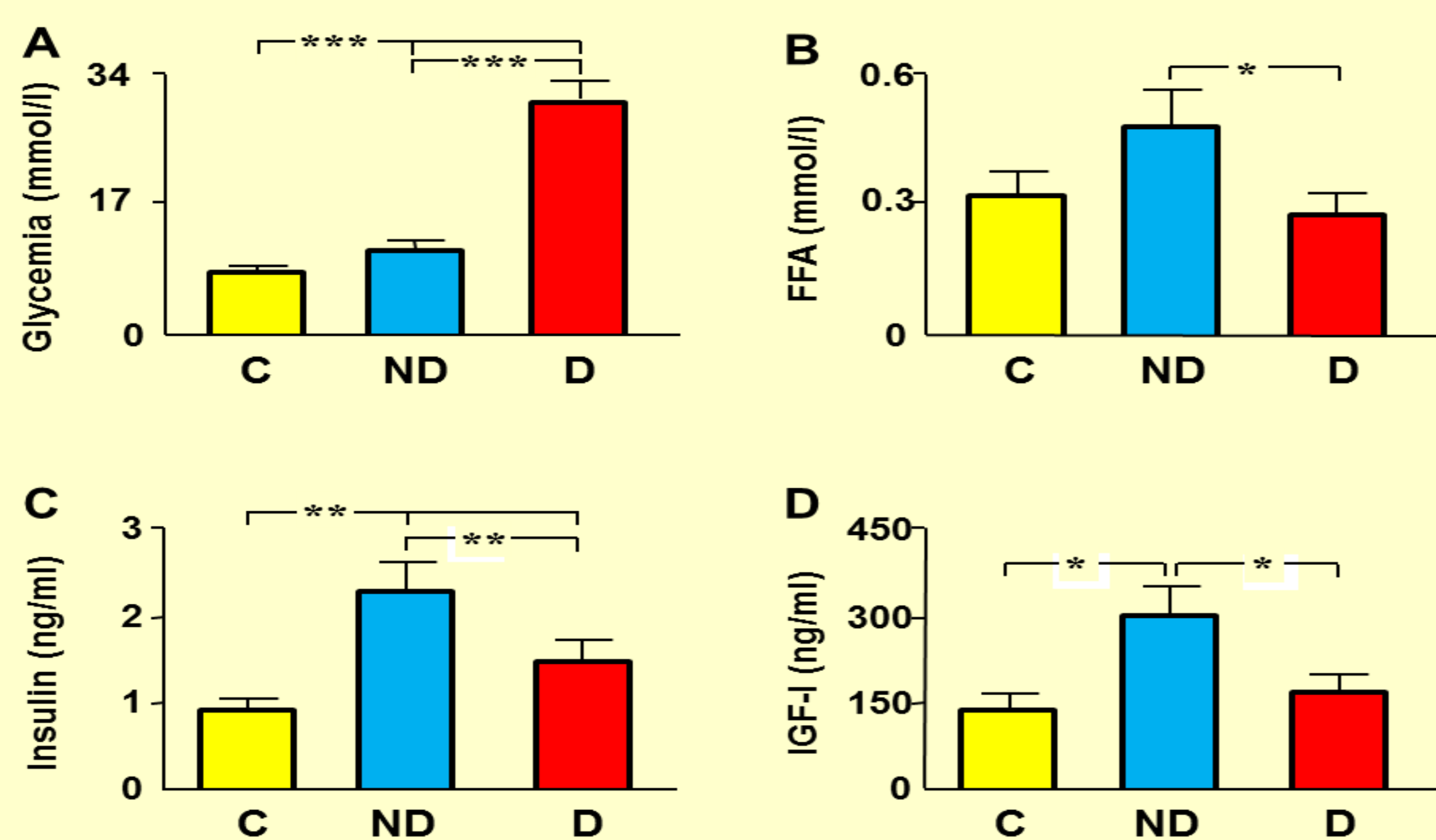
- Controls (C, wild type mice)
- Non-diabetic IRS2^{-/-} (ND, glycemia < 11.10 mmol/l)
- Diabetic IRS2^{-/-} (D, glycemia > 27.75 mmol/l)

Methods

- Serum insulin and IGF-I: ELISA
- Serum glucose and free fatty acids: colorimetric methods
- Hypothalamic insulin/IGF-I signaling: multiplexed bead immunoassay and immunoprecipitation/Western blot
- Enzymes: Western blot in hypothalamic extracts
- Metabolomic studies: nuclear magnetic resonance in hypothalamic extracts

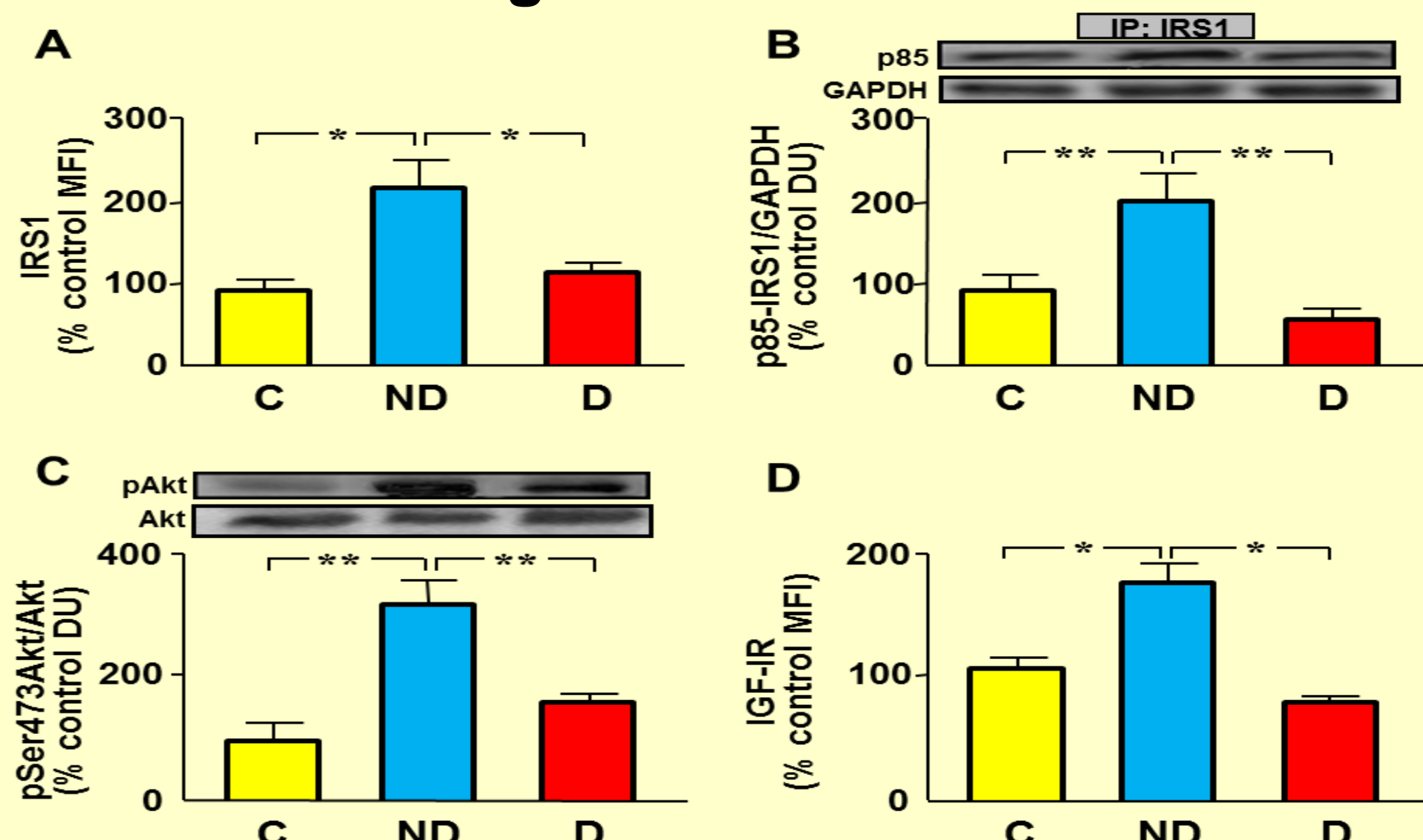
Results

1 Serum metabolite and hormone levels are modified in IRS2^{-/-} mice



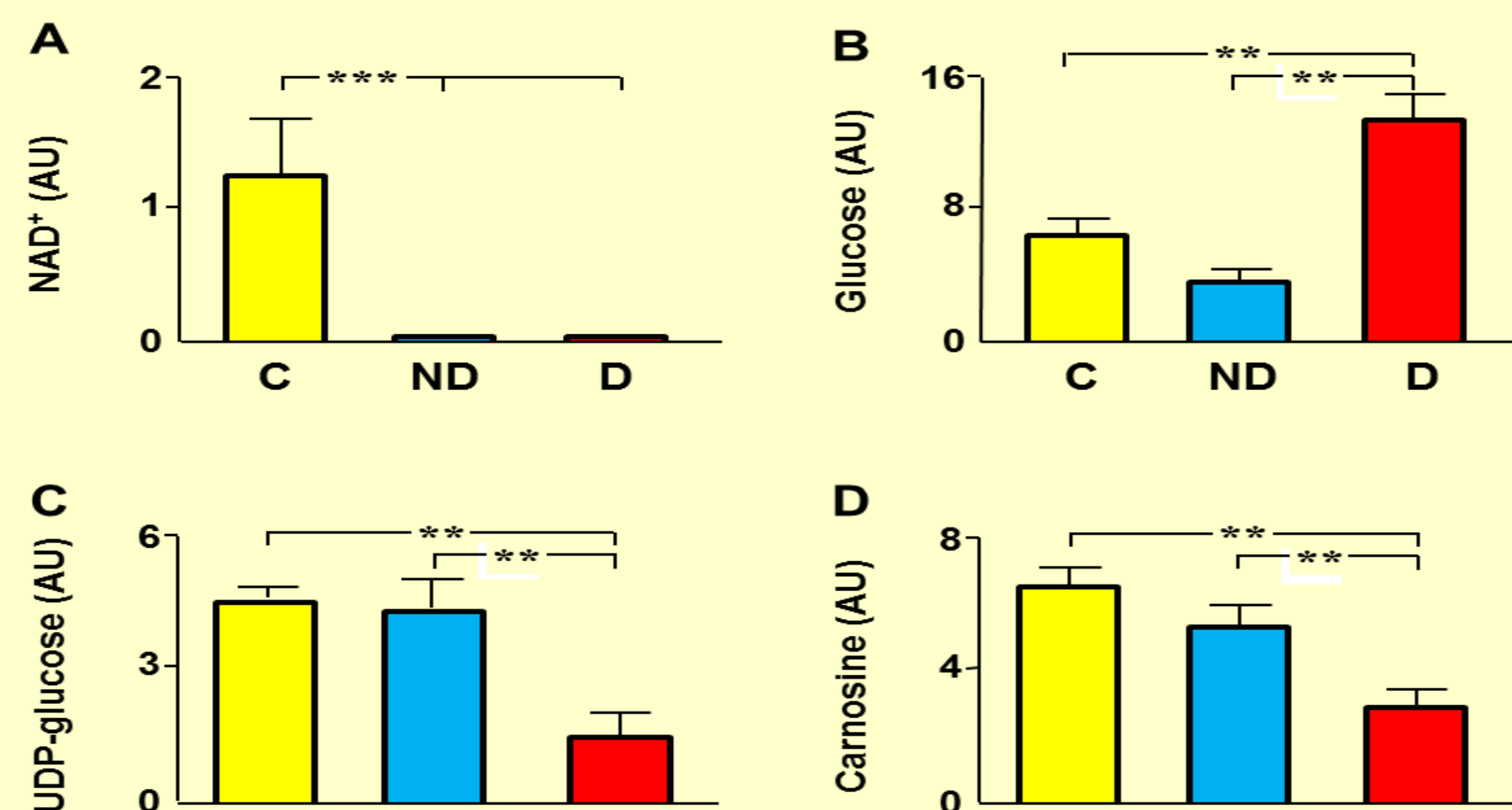
A. Glycemia in controls (C), non-diabetic IRS2 deficient (ND) and diabetic IRS2 deficient mice (D). B. Serum free fatty acid concentrations (FFA) in the same groups. C. Serum insulin levels in the same groups. D. Serum insulin-like growth factor-I (IGF-I) in the same groups. **p*<0.05, ***p*<0.01, ****p*<0.001.

2 Activation of hypothalamic insulin/IGF-I signaling is augmented in ND mice



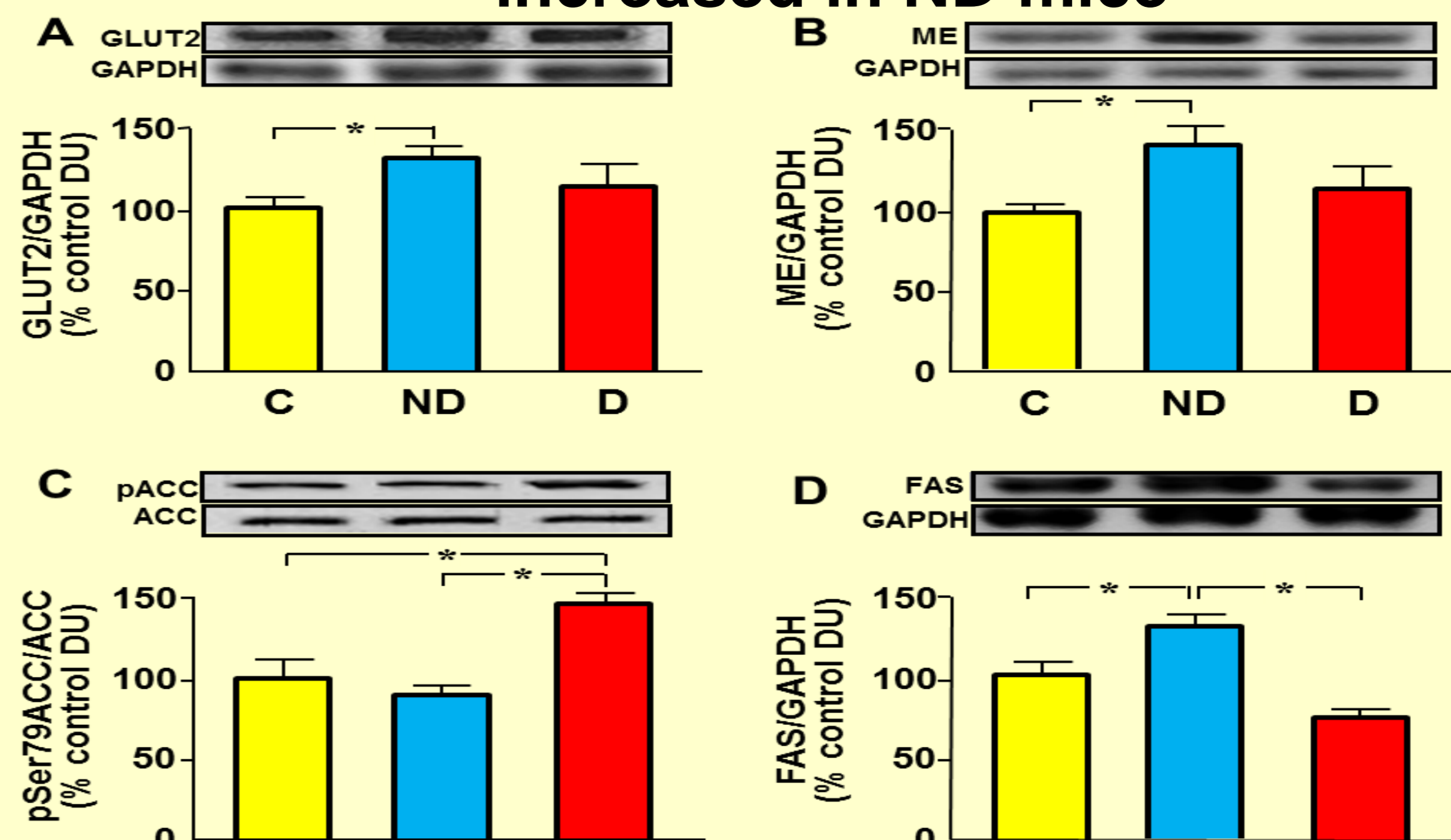
A. Relative levels of insulin receptor substrate 1 (IRS1) in controls (C), non-diabetic IRS2 deficient (ND) and diabetic IRS2 deficient mice (D). B. Relative levels of the regulatory subunit of PI3K (p85) associated to IRS1 in the same groups. C. Relative protein levels of Akt phosphorylated on serine 473 (pSer473-Akt) in the same groups. D. Relative protein levels of insulin-like growth factor-I receptor (IGF-IR) in the same groups. **p*<0.05, ***p*<0.01. DU, densitometry units; MFI, median fluorescent intensity.

3 Metabolite profiling in the hypothalamus of ND mice is related to deregulation of lipid homeostasis



A. Levels of hypothalamic nicotinamide adenine dinucleotide (NAD⁺) in controls (C), non-diabetic IRS2 deficient (ND) and diabetic IRS2 deficient mice (D). B. Levels of glucose in the same groups. C. Levels of uridine diphosphate (UDP)-glucose in the same groups. D. Levels of carnosine in the same groups. ***p*<0.01, ****p*<0.001. AU, arbitrary units.

4 Reduction capacity and fatty acid anabolism are increased in ND mice



A. Relative levels of glucose transporter 2 (GLUT2) in controls (C), non-diabetic IRS2 deficient (ND) and diabetic IRS2 deficient mice (D). B. Relative protein levels of malic enzyme (ME) in the same groups. C. Relative protein levels of acetyl CoA carboxylase phosphorylated on Ser79 (pSer79/ACC) in the same groups. D. Relative protein levels of fatty acid synthase (FAS) in the same groups. **p*<0.05. DU, densitometry units.

Conclusion

Increased hypothalamic insulin/IGF-I signaling might be a triggering factor for the occurrence of diabetes in IRS2^{-/-} mice.