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## PERIPHERAL ADIPOSE TISSUE INSULIN RESISTANCE ALTERS LIPID COMPOSITION AND FUNCTION OF HIPPOCAMPAL SYNAPSES

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### Abstract

Compelling evidence indicates that type 2 diabetes mellitus (T2D), insulin resistance (IR), and metabolic syndrome are often accompanied by cognitive impairment. However, the mechanistic link between these metabolic abnormalities and CNS dysfunction requires further investigations. Here, we evaluated whether adipose tissue (AT) IR and related metabolic alterations resulted in CNS changes by studying synapse lipid composition and function in the adipocyte-specific ectonucleotide pyrophosphate phosphodiesterase overexpressing transgenic (*AtENPP1*-Tg) mouse, a model characterized by white adipocyte IR, systemic IR, and ectopic fat deposition. When fed a high-fat diet (HFD), *AtENPP1*-Tg mice recapitulate essential features of the human metabolic syndrome, making them an ideal model to characterize peripherally induced CNS deficits. Using a combination of gas chromatography and western blot analysis, we found evidence of altered lipid composition, including decreased phospholipids and increased triglycerides (TG) and fatty acid (FFA) in hippocampal synaptosomes isolated from HFD-fed *AtENPP1*-Tg mice. These changes were associated with impaired basal synaptic transmission at the Schaffer collaterals to hippocampal cornu ammonis 1 (CA1) synapses, decreased phosphorylation of the GluN1 glutamate receptor subunit, down-regulation of insulin receptor expression and up-regulation of the FFA receptor 1.

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#### CONFLICT OF INTEREST

The authors have no conflicts of interest to declare.

## Keywords

Cognitive dysfunction; ENPPI; Glutamate receptors; Insulin Resistance; Lipids; synaptic transmission

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## INTRODUCTION

A growing body of evidence indicates that cognitive impairment is associated with obesity, metabolic syndrome, and type 2 diabetes (T2D) (Calvo-Ochoa and Arias 2014; Elias *et al.* 2003; Hassing *et al.* 2004), suggesting common yet undefined pathogenic mechanisms among the conditions. However, the cellular and molecular mechanisms driving these peripherally induced CNS deficits remain elusive in this emerging area of interest. This critical knowledge gap precludes the development of therapeutic interventions to prevent and/or treat cognitive impairment in the growing population affected by the metabolic complications of obesity.

Altered production of adipokines and lipid metabolism regulation in adipose tissue (AT) (*i.e.*, AT dysfunction) affect the plasma concentrations of circulating adipokines and free fatty acids (FFA). This aberrant endocrine signaling triggers numerous functional changes throughout the body that are part of the complex cluster of metabolic abnormalities that increase the risks for T2D and cardiovascular diseases in obese patients.

We have developed an adipocyte-specific ecto-nucleotide pyrophosphate phosphodiesterase overexpressing transgenic (*AtENPPI-Tg*) animal model of metabolic syndrome and systemic IR, which is highly suitable for evaluating peripherally driven changes in central synapses triggered by adipocyte IR. The *AtENPPI-Tg* mouse is characterized by adipocyte and systemic IR, increased circulating FFA, and TG deposition in the liver, all of which are induced by the adipocyte-specific overexpression of ecto-nucleotide pyrophosphate phosphodiesterase (*ENPPI*), a negative modulator of the insulin receptor (Pan *et al.* 2011). We have shown translational validity of our model by identifying association between higher AT *ENPPI* expression and systemic abnormalities of glucose and lipid metabolism in humans (Chandalia *et al.* 2012). In the present study, we used the *AtENPPI-Tg* mouse model to investigate the effects of AT-induced metabolic alterations of systemic lipid and glucose metabolism on lipid composition and functional activity of hippocampal synapses, and to explore involved candidate molecular mechanisms.

## MATERIALS AND METHODS

### Animals

Eighty-one adult male *AtENPPI-Tg* (transgenic, Tg) and C57Bl/6J (wild type, WT) mice were randomized to receive dietary intervention. Animals were generated through a breeding program at UTMB and were individually housed in their filter-top cages in a temperature-controlled environment at 22°C, humidity 40%, and a 12:12-h light-dark cycle. Understanding the 3 R's of Russell and Birch (Replacement, Refinement and Reduction), the number of mice required for each experiment was calculated based on the minimal number of animal required to detect a significant difference based on power analysis. The

study was approved by the Animal Care and Use Committee of the University of Texas Medical Branch (UTMB, Galveston, TX, USA) and was performed according to the guidelines of the National Institutes of Health Guidelines on the use of laboratory animals.

### Experimental Procedure

Following weaning at 8 weeks of age, a random group of Tg and WT siblings were pair-fed with high-fat chow (60% calories from fat - 37.1% saturated; Research Diets D12492, New Brunswick, NJ, USA). Another random group of Tg and WT siblings were fed a regular chow (RC) diet (4% calories from fat; Teklad 7001; Teklad, Madison, WI). Animal weights were recorded weekly. In the morning following 12 weeks of the respective diets, the animals were brought into the lab. and were either intracardially perfused and decapitated for electrophysiological studies, under general anesthesia (see below), or were immediately sacrificed by anesthetic overdose (5% isoflurane inhalation confirmed by chest opening) and their brains collected, flash-frozen in liquid nitrogen, and stored at  $-80^{\circ}\text{C}$  for future analyses.

### Synaptosome isolation

Hippocampi were dissected and homogenized from frozen brains. Synaptosomes were isolated using a sucrose gradient and ultracentrifugation, as previously described (Bjorklund *et al.* 2012). We employed samples with synaptosome yields 25 mg for lipid measurements.

### Lipid composition in hippocampal synaptosomes

Synaptosome lipids were extracted with the chloroform:methanol (2:1) Folch extraction (n=4-12 per group; a total of 38). TG, diacylglycerols (DAG), and phospholipids were isolated using TLC, as previously described (Pan *et al.* 2011). Briefly, TG, DAG, and phospholipid bands on TLC plates were identified, scraped off, and processed to separate fatty acids and glycerol. The fatty acids were derivitized and quantified with a gas chromatography-flame ionization detector (GC-FID 6890N model – Agilent Technologies, Santa Clara, CA, USA) using a Supelco SP-2330 fused silica capillary column (30 m  $\times$  0.25 mm  $\times$  0.2  $\mu\text{m}$  film thickness). The concentrations of TG, DAG, and phospholipids were calculated using an internal standard method as described previously (Zhang *et al.* 2011).

### Extracellular recording of hippocampal synaptic responses

The mice (n=4-5 per group; a total of 18) were anesthetized with 2-2-2-tribromoethanol (Sigma-Aldrich - 250 mg/Kg IP) and intracardially perfused with an ice-cold sucrose-based solution containing (in mM) 56 NaCl, 100 sucrose, 2.5 KCl, 20 glucose, 5  $\text{MgCl}_2$ , 1  $\text{CaCl}_2$ , 30  $\text{NaHCO}_3$ , 1.25  $\text{NaH}_2\text{PO}_4$ , and 1 kynurenic acid). Following decapitation and harvesting hippocampi, horizontal hippocampal slices (350  $\mu\text{m}$ ) were cut with a vibratome VT1200S (Leica, IL) in the sucrose-based solution and transferred to a recovery chamber with 95%  $\text{O}_2$  and 5%  $\text{CO}_2$  bubbled artificial CSF (ACSF) containing (in mM) 125 NaCl, 2.5 KCl, 2  $\text{MgCl}_2$ , 2.5  $\text{CaCl}_2$ , 1.25  $\text{NaH}_2\text{PO}_4$ , 26  $\text{NaHCO}_3$ , and 10 glucose) at  $31.5^{\circ}\text{C}$ . After at least 90 min of recovery, recordings were performed in ACSF in a submerged chamber, at  $30.5 \pm 0.5^{\circ}\text{C}$ . Recordings of field excitatory postsynaptic potentials (fEPSPs) were performed in

the cornu ammonis 1 (CA1) subfield with a tungsten electrode connected to an A-M Model 1800 Differential AC Amplifier (A-M Systems, Carlsborg, WA, USA). Cornu ammonis 3 (CA3) Schaffer collaterals were stimulated by a bipolar tungsten electrode, with 0.1-ms pulses of constant current. The traces were digitized by a Digidata 1200 interface using Clampex 7, and the slopes of the fEPSPs were measured offline with Clampfit 9.0 (PClamp software, Molecular Devices, Union City, CA, USA).

### Hippocampal receptor expression

To evaluate whether ENPP1-induced adipocyte insulin resistance associates with changes in key receptor proteins involved in cognitive function, we studied the insulin and glutamatergic neuronal transmission. To this goal, content of insulin and FFA1 receptors in hippocampal tissue and levels of glutamate receptors (namely, NMDA receptor 1 subunit (GluN1) and  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptor 1 subunit (GluA1)) in synaptosomes were determined by western blot (n=4-5 per group; a total of 18). Tissues were homogenized in cell lysis buffer (Cell Signaling Technology Inc. Danvers, MA), and mixed 1 mM PMSF, 1 $\times$  protease inhibitor cocktail, and phosphatase inhibitor cocktail (Sigma-Aldrich, Saint Louis, MO). Protein concentration of hippocampal tissue lysate was measured using the Bio-Rad protein assay kit (Bio-Rad Laboratories, Inc. Hercules, CA). Protein concentrations of synaptosome preparations were determined using a bicinchoninic acid protein assay kit (Thermo Scientific, Waltham, MA, USA). SDS-PAGE was used to separate proteins which were then transferred to a nitrocellulose membrane (Bio-Rad). Primary antibodies used were anti- $\beta$ -actin antibody (A1978, Sigma-Aldrich, Saint Louis, MO), anti-Insulin Receptor antibody- $\alpha$  (sc710 – Santa Cruz Biotechnology Inc., Santa Cruz, CA), anti-Insulin Receptor antibody- $\beta$  (Cat. # 610109 - BD Biosciences, San Jose, CA), anti-total GluN1 (D65B7 - 5704 - Cell Signaling Technology Inc. Danvers, MA), anti-phosphorylated-GluN1 (ser897, 3385; Cell Signaling Technology Inc., Danvers, MA), anti-total GluA1 (G12 - SC-55509; Santa Cruz Biotechnology Inc., Santa Cruz, CA), anti-phosphorylated-GluA1 (ser831 - SC-16313; Santa Cruz Biotechnology Inc., Santa Cruz, CA), and anti-FFA1 (ab109257, Abcam, Cambridge, MA). Secondary antibodies were from Southern Biotechnology Associates (Birmingham, AL). Immunoblots were detected using the ECL Western Blotting Detection Reagents (Amersham Biosciences, Piscataway, NJ). Expression levels were evaluated by quantification of relative density of each band normalized to that of corresponding  $\beta$ -actin band density using the NIH ImageJ software version 1.46r (NIH, Bethesda, MD).

### Brain response to insulin

In order to understand alterations of insulin responses in the brain, synaptosomes were isolated from different brain regions of RC-fed WT and *AtENPP1*-Tg (n=2-5 each; a total of 7), thirty minutes following the intra-peritoneal administration of insulin (0.5 U/Kg). Regional expressions of protein kinase B (Akt) and glucagon synthase kinase 3 $\beta$  (GSK-3 $\beta$ ) were determined by western blot, as detailed above. All antibodies used were from Cell Signaling (anti-pAKT Cat. # 2965 – anti-AKT Cat. # 4691 – anti-pGSK-3 $\beta$  Cat. #9323 – anti-GSK-3 $\beta$  Cat. #9315).

## Statistical Analysis

All data were analyzed using SAS version 9.2 (SAS Institute, Cary, NC, USA) and are presented as means  $\pm$  SEM. Two-way ANOVA was used for multiple comparisons between all four groups of animals with post-hoc Bonferroni or Scheffé test was used. Significance was set at  $p < 0.05$ .

## RESULTS

To determine the effect of peripherally-induced changes in insulin and lipid handling in the CNS, we included four study groups ( $n = 4-12$  per group). The weights of the RC-fed groups were  $31.8 \pm 1$  and  $30.7 \pm 1$  g for WT and *AtENPPI*-Tg, respectively; and the weight of high fat fed animals were  $37.3 \pm 3$  and  $36.6 \pm 3$  g (WT and *AtENPPI*-Tg, respectively –  $p > 0.05$  each).

### Lipid composition in hippocampal synaptosomes

The concentrations of TG, FFA, DAG, and phospholipids in the hippocampal synaptosomes are summarized in Fig. 1. TG content (Fig. 1A) was significantly different in the four groups ( $p = 0.004$  - ANOVA), and highest in the *AtENPPI*-Tg mice ( $*p < 0.05$  compared to the corresponding RC group), suggesting an additive effect of *ENPPI*-induced adipocyte IR and HFD on hippocampal lipid levels. Fig. 1B shows a non-significant trend towards increased synaptosomal FFA content in HFD groups. Fig. 1C shows that DAG content was increased in hippocampal synaptosomes isolated from HFD-fed WT animals, suggesting independent effects of diet and adipocyte IR on these lipid compositional changes in hippocampal synapses ( $p = 0.03$  - ANOVA;  $*p < 0.01$  vs. WT-RC). Fig. 1D shows that the synaptosomal phospholipid content was lower in *AtENPPI*-Tg mice and was further decreased by HFD ( $p = 0.001$  - ANOVA;  $*p < 0.01$  vs. WT-RC).

### Extracellular recording of hippocampal synaptic responses

To determine whether changes in lipid synaptic content were associated with functional deficits in hippocampal circuits, we measured CA3 Schaffer collateral axons synaptic responses. Extracellular stimulation of the elicited postsynaptic responses in the CA1 (Fig. 2), but fEPSP magnitude and slope were diet and genotype dependent. For slices from mice fed a RC diet, the input-output curves of fEPSP slopes in WT mice were indistinguishable from those recorded in *AtENPPI*-Tg mice ( $n = 4-5$  each,  $p > 0.05$  - ANOVA), indicating that overexpression of the *ENPPI* gene in adipose tissue was not sufficient to elicit changes in basal synaptic transmission in the CA1 hippocampal area. However, when challenged by the HFD, the *AtENPPI*-Tg group exhibited synaptic responses that were greatly diminished compared to HFD-fed WT littermates ( $n=4-5$  each,  $p < 10^{-8}$  - ANOVA). Notably, the HFD alone had a significant effect on synapses in WT animals, with HFD-fed mice exhibiting suppressed postsynaptic responses compared to RC-fed WT mice ( $n=5$  each,  $p < 10^{-10}$  - ANOVA). Indeed, HFD-fed WT animals exhibited fEPSPs with intermediate amplitudes between those from HFD-fed *AtENPPI*-Tg and RC-fed WT mice. This phenotype, along with the lack of effect of the *AtENPPI* genotype alone on synaptic responses, indicated an interaction between genotype and diet. Two-way ANOVA analysis of fEPSPs curves across the four experimental groups confirmed that the HFD and *AtENPPI* genotype additively

contribute to cause the greatest degree of synaptic suppression in HFD-fed *AtENPPI*-Tg animals.

### Hippocampal receptor expression

Next we addressed potential molecular mechanisms responsible for the observed suppression of synaptic transmission. As shown in Fig. 3A, there was a significant decrease in GluN1 phosphorylation in *AtENPPI*-Tg mice, particularly the HFD-fed group ( $p = 0.01$ , ANOVA-  $*p < 0.05$  vs. corresponding WT). Conversely, there were no changes in the levels of phosphorylated or total GluA1 in response to adipocyte IR and/or HFD (Fig. 3B), suggesting that synaptosomal lipid changes specifically affected GluN1 phosphorylation ( $p > 0.05$ , ANOVA).

In addition, we explored the insulin and FFA1 receptor expressions in the hippocampal tissue. Only in the HFD-fed *AtENPPI*-Tg hippocampus, insulin receptor- $\alpha$  and - $\beta$  subunits were down-regulated ( $p = 0.001$  and  $0.01$  for  $\alpha$ - and  $\beta$ -subunits, respectively, ANOVA -  $*p < 0.01$  vs. *AtENPPI*-Tg RC – Fig. 4A and B), while FFA1 receptors were up-regulated ( $p = 0.005$ , ANOVA –  $*p = 0.001$  vs. *AtENPPI*-Tg RC – Fig. 4C). These data confirm that a combination of the transgene and HFD is necessary to alter insulin receptor expression and is linked to the elevated FFA levels in this *AtENPPI*-Tg animal model (Pan *et al.* 2011).

### Brain response to insulin

To gain initial knowledge on post-receptor insulin signaling activation of *AtENPPI*-Tg mice CNS, we studied the regional expression of Akt and GSK-3 $\beta$  in the brain of RC-fed animals in response to acute systemic insulin elevation. After 30 min from intra-peritoneal insulin injection, we observed a differential regional response to insulin that was blunted in *AtENPPI*-Tg compared to their WT littermates (Figure 1S). It should be noted that these animals did not show a significant decrease in insulin receptor in absence of HFD (refer to Fig. 4 A and B).

## DISCUSSION

Our results provide new evidence for AT IR-induced biochemical and functional changes in hippocampal synapses. We found that HFD-fed *AtENPPI*-Tg animals exhibited alterations in synaptic lipid composition, decreased basal synaptic transmission at the Schaffer collaterals to CA1 synapses, decreased GluN1 receptor phosphorylation, decreased insulin receptor expression and increased FFA1 receptor expression compared to WT littermates. These results extend the impact of AT dysfunction-induced lipotoxicity from peripheral organs to the brain, thus providing a novel mechanistic link between the increased risk for cognitive impairment and the metabolic complications of obesity, such as metabolic syndrome and T2D.

One innovative aspect of the present work is in the use of the *AtENPPI*-Tg mouse, a model of adipocyte IR and consequent manifestations of metabolic syndrome (Pan *et al.* 2011). Due to adipocyte-specific overexpression of *ENPPI*, a negative modulator of the insulin receptor, *AtENPPI*-Tg mice exhibit specific adipocyte IR/AT dysfunction (decreased adipocyte insulin signaling activation and defective adipocyte maturation/TG storage and



adiponectin production) upon excessive caloric intake (HFD). In humans, high adipocyte lipolysis and the inability to store lipids in AT leads to FFA spillover and ectopic fat deposition in the liver and muscle, resulting in lipid and glucose metabolic alterations that recapitulate essential features of IR and metabolic syndrome (Brassard *et al.* 2008; Chandalia *et al.* 2012; Lomonaco *et al.* 2013; Pan *et al.* 2011; Ravussin and Smith 2002). Thus, *AtENPP1*-Tg mice are an ideal model to determine the dietary contribution of IR to the clinical manifestations of humans with metabolic syndrome and T2D.

The lipid composition of hippocampal synaptosomes from the brains of WT and *AtENPP1*-Tg mice was influenced by AT dysfunction. TG contents were clearly affected by the combination of diet and gene overexpression in the HFD-fed *AtENPP1*-Tg mouse, which are known to have high levels of AT dysfunction, circulating FFA, and IR (Pan *et al.* 2011). The increased plasma-derived FFA and higher glucose availability in the *AtENPP1*-Tg mouse brain would increase substrates for TG synthesis. Such TG excess, outside of AT, is known to cause cellular dysfunction and apoptosis (*e.g.*, lipotoxicity), manifested as cellular IR. Indeed, we have shown evidence of elevated cellular TG content in both the liver (Pan *et al.* 2011) and hippocampal synapses that would potentially affect their integrity and function. A mechanistic interpretation includes the view that a recent human study showed that acute FFA elevation induced by intralipid infusion results in decreased hippocampal glucose utilization (Emmanuel *et al.* 2013), similar to the well-established effect of acute FFA elevation on muscle insulin-mediated glucose disposal (Boden and Chen 1995; Boden *et al.* 1994; Homko *et al.* 2003). A plausible mediator of these intracellular effects downstream of increased circulating FFA is increased levels of the lipid metabolism intermediate DAG (Yu *et al.* 2002; Samuel and Shulman 2012; Itani *et al.* 2002; Nowotny *et al.* 2013). Interestingly, the pattern of hippocampal synaptosome DAG content was similar to the FFA content.

Neural phospholipid content was decreased in the hippocampal synaptosomes of mice with AT dysfunction. This change was most pronounced in HFD-fed *At-ENPP1*-Tg mice, a combination that we previously demonstrated to lead to severe systemic IR (Pan *et al.* 2011). The result is of interest given the previous findings by other investigators who reported decreased phospholipids in the hippocampal region of Alzheimer's disease patients (Prasad *et al.* 1998; Soderberg *et al.* 1991). Indeed, plasma phospholipids were recently identified as useful biomarkers for cognitive impairment (Mapstone *et al.* 2014). Such a decrease in phospholipids may result from hippocampal IR, similar to that reported in mice with streptozotocin-induced brain IR (Muller *et al.* 1998).

Because of the association between IR and cognitive impairment, next we studied whether that peripherally-induced IR could have a negative impact on synaptic transmission in the hippocampus. Extracellular field recordings revealed that the synaptic strength of the CA3 to CA1 inputs was significantly suppressed in HFD-fed *AtENPP1*-Tg mice. Given the role of the CA1 region and of synaptic GluN1 in cognitive function and memory formation (Morris 2013; Tsien *et al.* 1996), our findings might suggest reduced synaptic plasticity along with learning and memory deficit, a reduced capability of the hippocampus to decode memory engrams and store new memories; however future studies are required to directly test such hypothesis. This result is in line with the essential role of GluN1 (which is less

phosphorylated and therefore less active in these mice) in hippocampal synaptic plasticity and in learning and remembering novel places in spatial navigation (Tsien et al., 1996). Furthermore, we demonstrated that the HFD was sufficient to induce synaptic deficits in the CA3-CA1 pathway, although this effect was lower in WT animals compared to *AtENPPI*-Tg mice.

These functional manifestations occurred concomitantly with lipid profile alterations, indicating that brain circuitry changes might explain the relationship between cognitive impairment and insulin-dependent metabolic dysfunction. While previous animal studies have reported HFD-induced memory deficits (Granholm *et al.* 2008; Molteni *et al.* 2002; Winocur and Greenwood 2005) associated with IR (McNay *et al.* 2010), the CNS synaptic deficits described in the present study can be clearly causally linked to peripherally induced adipocyte IR.

Next, we evaluated hippocampal expression of both  $\alpha$ - and  $\beta$ - subunits of the insulin receptor and found that they were down-regulated in the HFD-fed *AtENPPI*-Tg mice, suggesting an interaction between gene and diet. On the other hand, FFA1 receptor expression was increased in these animals, suggesting increased neuronal FFA uptake, which may contribute to defective post-receptor CNS insulin signaling (Ruddock *et al.* 2008). Decreased insulin receptor expression could have a negative effect on hippocampal synaptic plasticity, resulting in memory and learning dysfunction (Calvo-Ochoa and Arias 2014), variables that remain to be evaluated in the future. The role of insulin receptor down regulation and neuronal IR in neurodegeneration is further supported by changes observed in the NIRKO (neuron-specific disruption of the IR gene) mouse (Schubert *et al.* 2004), in which complete knockout of brain insulin receptors mimics neuronal changes typically found in Alzheimer's disease.

The synapse between Schaffer collaterals and CA1 pyramidal neurons is under the control of NMDA-dependent plasticity (Morris, 2013; Tsien et al., 1996). At the same time, neuronal insulin is known to control NMDA-dependent synaptic plasticity in the hippocampus, a mechanism responsible for long-term potentiation and/or depression, i.e. memory formation (Liao and Leonard 1999; Liu *et al.* 1995; van der Heide *et al.* 2005; Costello DA. 2012). For these reasons, the NMDA receptor is an excellent candidate as a molecular mechanism of the decreased synaptic transmission in *AtENPPI*-Tg mice. In particular, we evaluated GluN1 phosphorylation at S897, a site that has been linked to neurophatologies (Emamian *et al.* 2004; Hei *et al.* 2012). The reduction in GluN1 phosphorylation in synaptosomes isolated from *AtENPPI*-Tg mice supports the view that AT dysfunction affects both synaptic lipid composition and post-translational modification of a glutamate receptor subtype critical for synaptic plasticity and memory formation (Morris 2013). Our studies indicate a decrease in GluN1 phosphorylation at S897, a residue that controls NMDA receptor permeation through protein kinase A phosphorylation (Aman *et al.* 2014). Although the functional role of S897 phosphorylation at synapses under normal conditions is still poorly understood, severe impairment in NMDA trafficking, AMPA and NMDA-mediated synaptic transmission, and long-term potentiation along with aberrant social interaction and sensorimotor gating have been reported in GluN1 S897A knock-in phosphomutant mice (Li *et al.* 2009). Furthermore, evidence indicates that the S897 site is downregulated in ischemic or NMDA-induced brain



damage rodent model (Hei *et al.* 2012) as well as in schizophrenia in humans (<http://www.ncbi.nlm.nih.gov/pubmed/14973229>Emamian *et al.* 2004). Thus, lack of phosphorylation of GluN1 at S897 is an absolute requirement for synaptic function and a causative link to neuropathologies. A reduction of the AMPA GluA1 receptor, similar to that observed in GluN1 S897A knock-in phosphomutant animals, was not detected in our current study. This observation requires further investigation, but suggests that S897 might be a point of convergence of intracellular signaling pathways, carrying specialized functions. This result provides insight into one of the possible mechanisms through which neuronal IR contributes to cognitive dysfunction by altering brain circuitry via effects on specific glutamate receptor subtypes.

## CONCLUSIONS

Our results suggest that peripheral lipids and IR alter hippocampal molecular and functional integrity, which could be a pivotal mechanism responsible for the reported cognitive deficits associated with metabolic syndrome and T2D. It is therefore tempting to speculate that the mechanisms described in the present work could underscore, at least in part, the reported epidemiological link between T2D and heightened risk of developing Alzheimer's Disease in humans. It is of interest to study the potential detrimental consequences on learning, memory, and other cognitive functions related to our current findings. Further ongoing studies in our laboratories are necessary to ultimately address this important issue.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## Abbreviations used

<b>AT</b>	adipose tissue
<b>CA</b>	cornu ammonis
<b>DAG</b>	diacylglycerol
<b>ENPP1</b>	ecto-nucleotide pyrophosphate phosphodiesterase 1 gene
<b>fEPSPs</b>	field excitatory postsynaptic potentials
<b>FFA</b>	free fatty acids
<b>FFA1</b>	free fatty acid receptor 1
<b>GC-FID</b>	gas chromatography-flame ionization detector

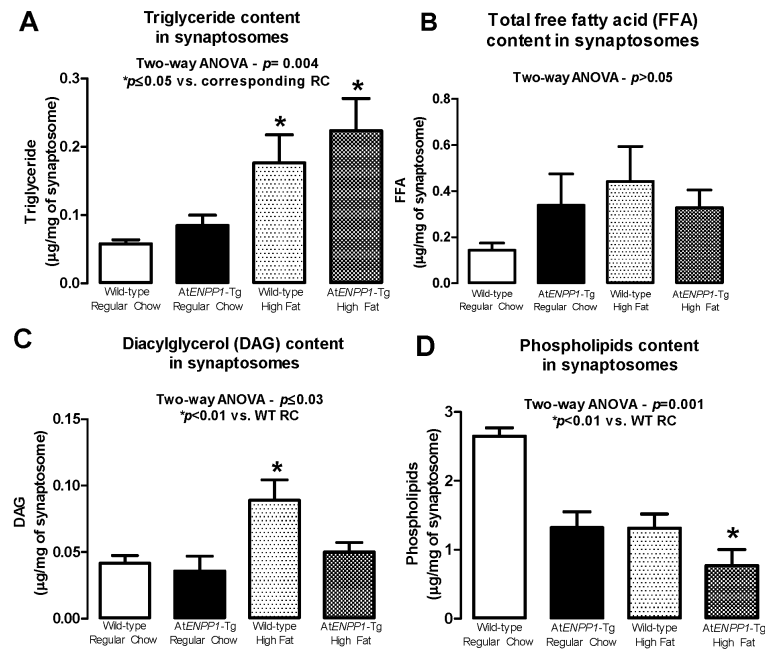
<b>GluA1</b>	$\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) receptor 1 subunit
<b>HFD</b>	high-fat diet
<b>IR</b>	insulin resistance
<b>GluN1</b>	NMDA receptor 1 subunit
<b>RC</b>	regular chow
<b>T2D</b>	type 2 diabetes mellitus
<b>Tg</b>	transgenic
<b>TG</b>	triglyceride
<b>WT</b>	wild type

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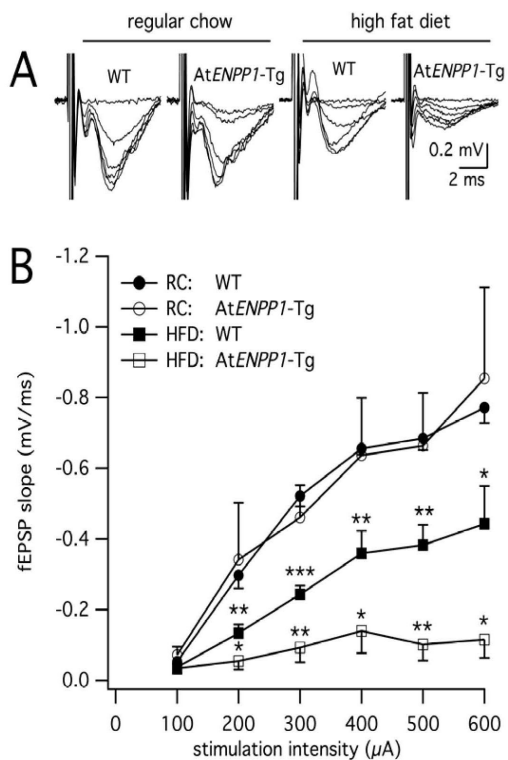
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**Figure 1.**

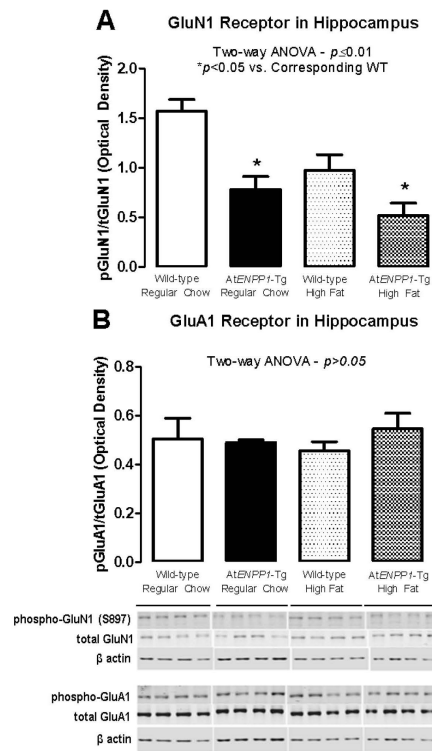
A. Neuronal triglyceride (TG) content in hippocampal synaptosomes. TG were significantly increased in response to HFD in both WT and *AtENPP1*-Tg mice ( $p = 0.004$ , ANOVA –  $*p < 0.05$  vs. corresponding RC littermates). B. Neuronal FFA content in hippocampal synaptosomes. An insignificant trend toward higher FFA concentrations was detected after HFD ( $p > 0.05$ , ANOVA). C. Neuronal DAG content in hippocampal synaptosomes. DAG increased in response to a HFD in the WT mice ( $p = 0.03$  – ANOVA;  $*p < 0.01$  vs. WT-RC). D. Neuronal phospholipid content in hippocampal synaptosomes. Phospholipid content was decreased in response to *ENPP1* overexpression in *AtENPP1*-Tg and in response to HFD. Consumption of a HFD further decreased phospholipid levels ( $p = 0.001$  – ANOVA;  $*p < 0.01$  vs. WT-RC).



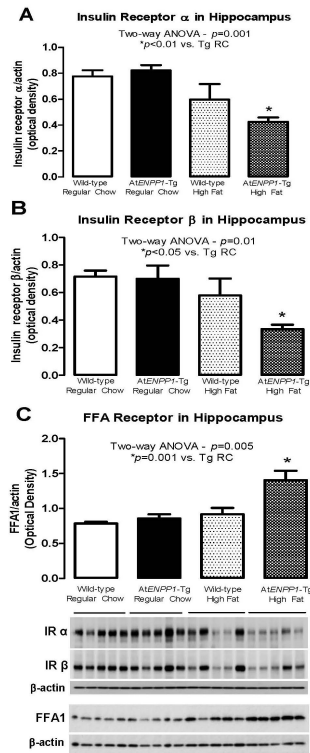
**Figure 2. Basal synaptic transmission in the hippocampal CA1 area is greatly suppressed in HFD-fed AtENPP1-Tg animals**

A. Representative traces of extracellular field recordings in the CA1. Notice the depressed signal in the HFD-fed AtENPP1-Tg mice. B. Input/output curves of fEPSP slopes versus presynaptic stimulation intensity. fEPSP slopes were diminished in HFD-fed animals ( $p < 10^{-8}$  for AtENPP1-Tg HFD vs. WT HFD and  $p < 10^{-10}$  for WT HFD vs. WT RC, ANOVA). Asterisks signify significant decreases at a specific stimulation intensity between HFD-fed WT relative to RC-fed WT and between HFD-fed AtENPP1-Tg relative to HFD-fed WT (\*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$ ).



**Figure 3.**

A. pGluN1/GluN1 ratio decreases in hippocampal synaptosomes. The ratio between phosphorylated GluN1 and total GluN1 was lower in the presence of AT-specific *ENPP1* overexpression, particularly after exposure to HFD ( $p = 0.01$ , ANOVA- \* $p < 0.05$  vs. corresponding WT). B. The pGluA1/GluA1 ratio is unchanged in hippocampal synaptosomes. The ratio between phosphorylated and total GluA1 was not affected by HFD or AT-specific *ENPP1* overexpression ( $p > 0.05$ , ANOVA).

**Figure 4.**

A & B Insulin receptor  $\alpha$  and  $\beta$  subunits 1 levels in hippocampal tissue. Only in the HFD-fed *AtENPPI-Tg* hippocampus, insulin receptor- $\alpha$  and - $\beta$  subunits were down-regulated ( $p = 0.001$  and  $0.01$  for  $\alpha$ - and  $\beta$ -subunits, respectively, ANOVA - \* $p<0.01$  vs. *AtENPPI-Tg* RC). C. Free fatty acid receptor levels in hippocampal tissue. ( $p = 0.005$ , ANOVA - \* $p = 0.001$  vs. *AtENPPI-Tg* RC).