



Accepted Article

Title: Metabolic inhibitors of O-GlcNAc transferase (OGT) that act in vivo implicate decreased O-GlcNAc levels in leptin-mediated nutrient sensing.

Authors: Ta-Wei Liu, Wesley F. Zandberg, Tracey M. Gloster, Lehua Deng, Kelsey D Murray, Xiaoyang Shan, and David Jaro Vocadlo

This manuscript has been accepted after peer review and appears as an Accepted Article online prior to editing, proofing, and formal publication of the final Version of Record (VoR). This work is currently citable by using the Digital Object Identifier (DOI) given below. The VoR will be published online in Early View as soon as possible and may be different to this Accepted Article as a result of editing. Readers should obtain the VoR from the journal website shown below when it is published to ensure accuracy of information. The authors are responsible for the content of this Accepted Article.

To be cited as: Angew. Chem. Int. Ed. 10.1002/anie.201803254 Angew. Chem. 10.1002/ange.201803254

Link to VoR: http://dx.doi.org/10.1002/anie.201803254 http://dx.doi.org/10.1002/ange.201803254

WILEY-VCH

COMMUNICATION

Metabolic inhibitors of *O*-GlcNAc transferase (OGT) that act *in vivo* implicate decreased *O*-GlcNAc levels in leptin-mediated nutrient sensing.¹

Tai-Wei Liu,[†] Wesley F. Zandberg,[†] Tracey M. Gloster,[†] Lehua Deng, Kelsey D. Murray, Xiaoyang Shan and David J. Vocadlo*

Abstract: O-linked glycosylation of serine and threonine residues of nucleocytoplasmic proteins with N-acetylglucosamine (O-GlcNAc) residues is catalyzed by O-GlcNAc transferase (OGT). O-GlcNAc is conserved within mammals and is implicated in a wide range of physiological processes. Here we describe metabolic precursor inhibitors of OGT suitable for use both in cells and in vivo in mice. These 5-thiosugar analogues of N-acetylglucosamine are assimilated through a convergent metabolic pathway, most likely involving Nacetylglucosamine-6-phosphate de-N-acetylase (NAGA), to generate a common OGT inhibitor within cells. Of these inhibitors, we show that 2-deoxy-2-N-hexanamide-5-thio-D-glucopyranoside acts in vivo to induce dose- and time-dependent decreases in O-GlcNAc levels in various tissues. Decreased O-GlcNAc correlates, both in vitro within adipocytes and in vivo within mice, with lower levels of transcription factor Sp1 and the satiety-inducing hormone leptin revealing a link between decreased O-GlcNAc levels and nutrient sensing in peripheral tissues of mammals.

β-O-linked *N*-acetylglucosamine (*O*-GlcNAc) is a dynamic modification of nucleocytoplasmic proteins that is present in all multicellular eukaryotes.^[1] The attachment of GlcNAc (1, Figure 1) to hydroxyl groups of serine or threonine residues of hundreds of target proteins is catalyzed by the glycosyltransferase *O*-GlcNAc transferase (OGT), which uses uridine diphospho-*N*-acetylglucosamine (UDP-GlcNAc, 2) as a donor sugar substrate. The glycosidase *O*-GlcNAcase (OGA) cleaves *O*-GlcNAc off from proteins.

O-GlcNAc is implicated in various diseases including cancer, neurodegeneration, cardiovascular disease, and obesity. [2] Notably, O-GlcNAc levels respond to nutrient availability both within cells and *in vivo* in animal models. [1b] Transgenic mice overexpressing OGT in fat or muscle tissue exhibit elevated serum leptin and insulin levels in addition to insulin resistance. [3] Further, deletion of the gene encoding OGT from neurons of the paraventricular nucleus (PVN) within the hypothalamus of mice results in uncontrolled eating. [2b]

Notably, OGA inhibitors that are active *in vivo* have helped gain insights into the roles of increased *O*-GlcNAc levels in mammals. Strikingly, OGA inhibitors sometimes yield different results from those made using genetic approaches to increase *O*-GlcNAc levels^[4]; perhaps because OGT and OGA also have non-

catalytic roles. Unfortunately, similar studies regarding the roles of decreased O-GlcNAc on mammalian physiology are lacking because there are no OGT inhibitors suitable for use *in vivo*. Given the emerging roles of OGT in nutrient sensing and other processes, inhibitors of OGT that can be used as research tools *in vivo* are of high interest.^[5]

One approach to decreasing O-GlcNAc levels using small molecules has been to use broad-spectrum amidotransferase inhibitors such as 6-diazo-5-oxo-L-norleucine (DON), which promiscuously blocks all amidotransferases that biosynthesize many cellular metabolites including UDP-GlcNAc. throughput screening has been pursued to deliver hits that can be improved upon.^[5b, 5d] These leads, however, show modest cellular activity and limited solubility. [5b] They also exhibit off-target cellular toxicity. [5b] Another approach [5c] has been to generate a GlcNAc analogue, 2-acetamido-2-deoxy-5-thio- α -D-glucopyranose (5SGlcNAc, 3), which in its per-O-acetylated form (Ac₄5SGlcNAc, 3-OAc) can diffuse across the plasma membrane. Within cells, 3-OAc is de-O-acetylated and assimilated via the GlcNAc salvage pathway (Figure 1) to generate UDP-5SGlcNAc (4), which is a competitive OGT inhibitor ($K_i = 8 \mu M$). Ac₄5SGlcNAc (3-OAc) lowers cellular O-GlcNAc levels (EC₅₀ = 0.8 to 5 μ M). This metabolic OGT inhibitor is not toxic but suffers from poor aqueous solubility. Indeed, while often used in cell studies, [6] solubilizing 3-OAc requires high concentrations of DMSO, making incompatible for dosing of mammals.

De novo hexosamine biosynthetic pathway

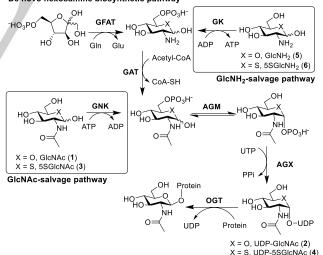


Figure 1. The hexosamine biosynthetic pathway (HBP) and the GlcNAc-salvage pathway yield UDP-GlcNAc (2). Glutamine fructose amidotransferase (GFAT) catalyzes the conversion of glucosamine (5) to glucosamine-6-phosphate (GlcNH₂-6PO₄; 5-6PO₄). Acetylation of 5-6PO₄ by glucosamine acetyltransferase (GAT) yields GlcNAc-6PO₄, an intermediate also produced by GlcNAc kinase (GNK) in the GlcNAc-salvage pathway. The sequential action of phosphoglucosamine mutase (AGM) and GlcNAc pyrophosphorylase (AGX) on GlcNAc-6PO₄ leads to the formation of UDP-GlcNAc (2). 5SGlcNAc (3) can be similarly salvaged and converted into UDP-5SGlcNAc (4).

[*] Current address: Department of Chemistry, University of British Columbia, 3247 University Way, Kelowna, BC, Canada [**] Current address: School of Biology, Biomedical Sciences Research Complex, University of St Andrews, North Haugh, St

Andrews, Fife, UK.

†These autors contributed equally.

Supporting information for this article is given via a link at the end of the document.

[[]a] Dr. T.-W. Liu, Dr. W. F. Zandberg, [+] Dr. T. M Gloster, [++] Dr. L. Deng, K. D. Murray, Dr. X. Shan, Prof. D. J. Vocadlo*, Departments of Chemistry & Molecular Biology and Biochemistry Simon Fraser University, 8888 University Dr., Burnaby, BC, Canada E-mail: dvocadlo@sfu.ca
[1] Current address: Department of Chemistry University of British

COMMUNICATION

Accordingly, to explore the roles of decreased O-GlcNAc levels in organismal physiology, compounds that act *in vivo* are of high interest.

To generate a tool compound for inhibiting OGT in vivo, we synthesized a panel of water soluble analogues of 5SGlcNAc (3) possessing various N-acyl substituents (7-17, Figure 2a,b Scheme S1). We reasoned these hydrophobic N-acyl groups would confer a balance between hydrophilicity and lipophilicity, making them water soluble yet able to diffuse into cells. Furthermore, although the substrate specificity of acetylglucosamine-6-phosphate de-N-acetylase (NAGA) is not known,[7] we speculated that compounds 7-17, once phosphorylated, might be substrates for this recently identified enzyme (Figure 2b). Action of NAGA on phosphorylated 7-17 would lead to formation of a common intermediate, 5-thioglucosamine-6-phosphate (5-6PO₄), which can be assimilated by the hexosamine biosynthetic pathway (HBP). However, direct entry of 7-17 into the GlcNAc-salvage pathway cannot be ruled out since the specificity of NAGA is unknown and bulkier Nglycolyl (GlcNGc) and N-azidoacetyl (GlcNAz) containing analogues enter the HBP.[8]

To measure the potential of **7–17** and their per-*O*-acetylated congeners to block OGT activity in cells, we treated cells with these compounds. Analysis of *O*-GlcNAc levels of cell lysates showed only some deprotected analogues were effective in reducing *O*-GlcNAc levels (Figure 2c, Figure S1). Of these the promising 2-hexanamide derivative (5SGlcNHex, **12**) showed dose-dependent decreases in *O*-GlcNAc comparable to Ac₄5SGlcNAc (**3**-OAc) (Figure S2).

We next tested whether these compounds were directly activated as UDP-linked analogues or if they were metabolized by NAGA. We therefore analyzed the pool of nucleotide sugars from cells treated with compounds 7, 8, and 12 by capillary electrophoresis (CE) (Figure 3). Electropherograms for nucleotide sugars from cells treated with 7, 8, and 12 revealed two new peaks that had CE mobilities matching those of the chemoenzymatically-prepared standards UDP-5SGlcNAc (4) and

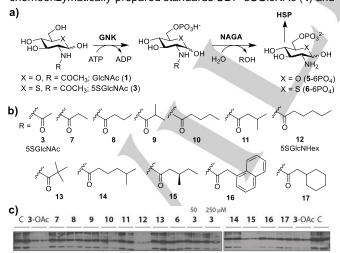
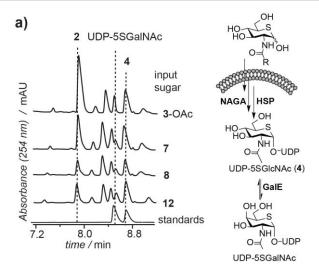


Figure 2. Synthetic analogues of 5SGlcNAc (3) are metabolized within cells, leading to decreased O-GlcNAc levels in cells. a) Possible metabolism of 5SGlcNR analogues (7 – 17) by NAGA prior to their entry into the HBP. b) Synthetic analogues (7 – 17) bearing various N-acyl R groups. c) Effects of analogues of O-GlcNAc levels of HEK293 cells. C = vehicle (PBS alone).



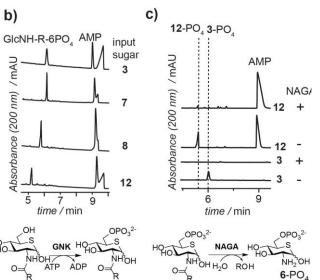


Figure 3. 5SGIcNR analogues are sequentially processed in cells by GIcNAc kinase (GNK) and, likely, N-acetylglucosamine-6-phosphate-de-N-acetylase (NAGA) to generate 5SGIcNH₂-6PO₄ (6-PO₄). a) Nucleotide sugar analysis of cells treated with selected inhibitors indicated that all the 5SGIcNR analogues tested (3-OAc, 7, 8, and 12) were converted within cells into two new nucleotide sugars, indistinguishable from synthetically prepared UDP-5SGIcNAc (4) and UDP-5SGaINAc. b) 5SGIcNR analogues were phosphorylated by GNK in vitro. c) 5SGIcNR-6-phosphates were not substrates for AGM or AGX but were readily hydrolyzed in vitro by NAGA to produce 5SGIcNH2-6PO₄ (6-PO₄).

its epimer UDP-5SGalNAc (Figure 3a). Extracts from cells treated with per-O-acetylated 5SGlcNH $_2$ (6-OAc, Figure S3) also yielded the same two unnatural nucleotide sugars in cells. These data suggest that these 5SGlcNR analogues are processed within cells to the common intermediate 2-amino-2-deoxy-5-thio-glucopyranose 6-phosphate (6-PO $_4$) and then assimilated by the HBP to form UDP-5SGlcNAc (4).

To clarify the metabolic processing of compounds **7**, **8**, and **12**, we used recombinant enzymes of the mammalian HBP (Figure 1). **7**, **8**, and **12** were all phosphorylated by GNK (Figure 3b), yet none of the phosphorylated products were substrates for AGM except for **7**. However, the nucleotide sugar produced *in*

COMMUNICATION

vitro from 7 by the combined HBP enzymes had a different mobility than that of nucleotide sugar UDP-5SGlcNAc (4) that we detected in cells treated with compound 7 (Figure S4). We accordingly tested whether NAGA converts the corresponding 5thiosugar-6-PO₄ derivatives of **7**, **8**, and **12** into 5SGlcNH₂-6-PO₄ (6-6PO₄). We confirmed this scenario by digestion of 12-PO₄ with recombinant human NAGA. Electropherograms obtained by CE analysis of the processing of 12-PO4 by NAGA showed it was converted to 5SGlcNH₂-6-PO₄ (6-PO₄) (Figure 3c). These data suggest that 5SGlcNR analogues 7-17, including 7, 8, and the most potent derivative 5SGlcNHex (12), are phosphorylated within cells by GNK and then deacylated by NAGA, which we found has remarkable substrate tolerance. The resulting common intermediate, 5SGlcNH2-6-PO4 (6-PO4), is then assimilated via the HBP to form UDP-5SGIcNAc (4), which leads to metabolic inhibition of OGT and decreased O-GlcNAc levels in tissues.

Previous efforts to use Ac45SGlcNAc (3-OAc) in vivo in mammals failed because of its poor aqueous solubility. We therefore evaluated if these new water-soluble metabolic OGT inhibitors could be used in vivo. Accordingly, we dosed mice by intraperitoneal (IP) delivery with our most cell active compound 12. A concentration-dependent decrease in spleen O-GlcNAc levels was observed upon treatment with 12, with apparent effects at even 3 mg kg⁻¹ (Figure S5a). Compound 12 (300 mg kg⁻¹) decreased spleen O-GlcNAc levels over time with maximal inhibition by 8 h that was maintained for at least 48 h (Figure S5b). 12 reduced O-GlcNAc levels in the kidneys, lungs, fat, pancreas, heart, spleen and muscle tissue but not in the blood or brain (Figure S5c). Mice injected with high doses of 12 (300 mg kg⁻¹), became lethargic, which is a known consequence of low leptin levels. [9] A second injection of 300 mg kg-1 on the second day caused mice to be moribund, and we discontinued treatment. We therefore dosed mice IP with a lower dose of 12 (50 mg kg⁻¹, n=3) and found reduced O-GlcNAc levels in various tissues 16 h after dosing but no effect in brain, liver, pancreas, and kidney (Figure 4a,b and Figure S6a-c). O-GlcNAc levels returned to baseline levels after 16 h (Figure 4a). Metabolic inhibitor 12 therefore acts in vivo to reversibly lower O-GlcNAcylation in various tissue types. Notably, others have shown Ac₄5SGlcNAc (3-OAc) appears able to inhibit other glycosyltransferases in cell lines.[5b] We find, however, by lectin blot assessment that 12 induced no apparent changes in other forms of protein glycosylation in all tissues tested (Figure S7), even at a dose of 300 mg kg⁻¹ after 48 h. Notably, these observations are consistent with a recent report using parent compound Ac₄5SGlcNAc (3-OAc) in cells.^[10]

Curiously, we observed that mice treated with either low or high doses (50-300 mg kg⁻¹) of **12** exhibited engorged stomachs that were full of rodent chow after 16 h (Figure S8). Because lethargy and excessive food consumption (hyperphagia) are seen in mice deficient in leptin signaling,^[11] we measured the serum leptin concentration of animals treated with 50 mg kg⁻¹ **12** (Figure 4b). Their leptin levels dropped transiently to a minimum at 16 h (Figure 4b), consistent with the engorged stomachs^[9,12]. Leptin levels were also lower in mice treated with a high dose of **12** (300 mg kg⁻¹) (Figure S9). We therefore hypothesized that OGT inhibition might impair leptin production (Figure S10).

To test this hypothesis we used 3T3-L1 adipocytes and found these cells secreted less leptin when treated with either 12 (Figure 4d) or Ac₄5SGlcNAc (3-OAc) (Figure S11) without affecting

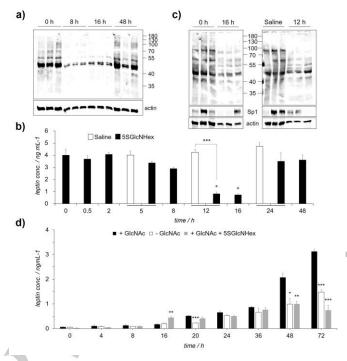


Figure 4. Dosing of mice with 50 mg kg 1 5SGlcNHex (12) reduced O-GlcNAc levels and impaired the secretion of the hormone leptin. **a)** Skeletal muscle from mice dosed with 50 mg kg 1 of 12 showed a transient reduction in O-GlcNAc levels by immunoblot analysis (CTD110.6). **b)** Leptin levels as measured by ELISA decreased in mice dosed with 12 to a minimum at 16 h. **c)** Dosing with 12 decreased O-GlcNAc and Sp1 levels to a minimum at 16 h in fat pad tissue. **d)** Compound 12 lowered GlcNAc-induced leptin secretion from 3T3-L1 adipocytes. Results are given as the means \pm SEM of three independent samples (n = 3). Each independent sample was tested in duplicate; symbols denote statistical significance (*p < 0.05, **p < 0.01, ***p < 0.001 compared to control. Student's t-test).

glucose uptake. Notably, transcription factors C/EBP- α , - β , and Sp1 all regulate leptin production. Compound **12** lowered levels of these known *O*-GlcNAcylated proteins in adipocytes (Figure S9) and in mice (Figure S6d). These collective data are consistent with OGT acting as a nutrient sensor in a process coupling HBP flux to leptin levels (Figure S10).

We envision that future studies to detail dosing regimens with 12 will be important to explore the effects of decreased *O*-GlcNAc on varied physiological processes. Experiments will also be needed to detail the mechanistic links between reduced *O*-GlcNAc levels due to *in vivo* OGT inhibition, impaired leptin secretion, and apparent hyperphagia. However, these data provide the first direct correlation between decreased *O*-GlcNAc levels and impaired leptin production *in vivo*. Our findings are in accord with studies showing that reduced *O*-GlcNAc lowers levels of Sp1 in cells^[6f,13] and OGT having a pivitol role in fat tissues^[14]. Notably, these data also suggest that regulation of leptin by *O*-GlcNAc is bidirectional *in vivo*, since overexpression of OGT^[3], knock out of OGA, ^[15], and metabolic upregulation of UDP-GlcNAc levels^[16] all increase leptin expression.

In summary, we describe convenient new tools to inhibit OGT in cells and *in vivo*. Strikingly, inhibition of OGT with 5SGIcNHex (12) provides support for the hypothesis that reduced *O*-GlcNAc levels signal impaired nutrient supply in mammals. We expect these observations will stimulate activity in the creation of

WILEY-VCH

COMMUNICATION

additional metabolic OGT inhibitors, as well as metabolic inhibitors of other glycosyltransferases for use *in vivo*. Moreover, we envision that this chemical strategy of metabolic OGT inhibition will serve as a valuable complement to using genetic methods in various rodent models to accelerate studies to understand the *in vivo* roles of *O*-GlcNAc in mammals.

Experimental Section: Please see supporting information for full experimental procedures. All experiments carried out on animals were approved by the university animal care committee of SFU.

Acknowledgments: K. Buettner and M. Dearden from the SFU animal care facility are thanked for their expert advice and assistance. S. Yuzwa is thanked for early studies on leptin levels. This work was supported by a grant from the Natural Sciences and Engineering Research Council of Canada (NSERC) and Canadian Glycomics Network (GlycoNet). D.J.V. is a scholar of the Michael Smith Foundation of Health Research (MSFHR) and a Tier II Canada Research Chair in Chemical Glycobiology. T.M.G was supported by a Sir Henry Wellcome Post-doctoral Fellowship and a Research Career Development Fellowship from the Wellcome Trust.

Keywords: glycoprotein • inhibitors • thiosugar • nucleotidesugars • leptin

References:

- a) C. R. Torres, G. W. Hart, J. Biol. Chem. 1984, 259, 3308-3317;
 b) G. W. Hart, M. P. Housley, C. Slawson, Nature 2007, 446, 1017-1022
- a) C. M. Ferrer, V. L. Sodi, M. J. Reginato, J Mol Biol 2016, 428, 3282-3294; b) O. Lagerlof, J. E. Slocomb, I. Hong, Y. Aponte, S. Blackshaw, G. W. Hart, R. L. Huganir, Science 2016, 351, 1293-1296; c) S. Dassanayaka, S. P. Jones, Pharmacol. Therapeut. 2014, 142, 62-71.
- [3] D. A. McClain, W. A. Lubas, R. C. Cooksey, M. Hazel, G. J. Parker, D. C. Love, J. A. Hanover, *Proc. Natl. Acad. Sci., USA* 2002, 99, 10695-10699.
- [4] a) C. F. Teo, E. G. El-Karim, L. Wells, *Glycobiology* 2016, 26, 1198-1208; b) M. S. Macauley, Y. He, T. M. Gloster, K. A. Stubbs, G. J. Davies, D. J. Vocadlo, *Chem. Biol.* 2011, 17, 937-948; c) M. S. Macauley, X. Shan, S. A. Yuzwa, T. M. Gloster, D. J. Vocadlo, *Chem. Biol.* 2011, 17, 949-958; d) P. J. Stivers, L. Harmonay, A. Hicks, H. Mehmet, M. Morris, G. M. Robinson, P. R. Strack, M. J. Savage, D. M. Zaller, I. Zwierzynski, P. E. Brandish, *PLoS One* 2015, 10, e0145151.
- [5] a) J. Jiang, M. B. Lazarus, L. Pasquina, P. Sliz, S. Walker, Nat Chem. Biol. 2011, 8, 72-77; b) R. F. Ortiz-Meoz, J. Jiang, M. B. Lazarus, M. Orman, J. Janetzko, C. Fan, D. Y. Duveau, Z. W. Tan, C. J. Thomas, S. Walker, ACS Chem. Biol. 2015, 10, 1392-1397; c) T. M. Gloster, W. F. Zandberg, J. E. Heinonen, D. L. Shen, L. Deng, D. J. Vocadlo, Nat Chem Biol 2011, 7, 174-181; d) B. J. Gross, B. C. Kraybill, S. Walker, J, J. Am. Chem. Soc. 2005, 127, 14588-14589; e) R. Trapannone, K. Rafie, D. M. van Aalten, Biochem. Soc. Trans. 2016, 44, 88-93.
- [6] a) S. Olivier-Van Stichelen, L. Drougat, V. Dehennaut, I. El Yazidi-Belkoura, C. Guinez, A. M. Mir, J. C. Michalski, A. S. Vercoutter-Edouart, T. Lefebvre, Oncogenesis 2012, 1, e36; b) P. J. Lund, J. E. Elias, M. M. Davis, J. Immunol. 2016, 197, 3086-3098; c) M. Swamy, S. Pathak, K. M. Grzes, S. Damerow, L. V. Sinclair, D. M. van Aalten, D. A. Cantrell, Nat. Immunol. 2016, 17, 712-720; d) F. Doll, A. Buntz, A. K. Spate, V. F. Schart, A. Timper, W. Schrimpf, C. R. Hauck, A. Zumbusch, V. Wittmann, Angew. Chem. Int. Ed. Engl. 2016, 55, 2262-2266; e) C. M. Speakman, T. C. Domke, W. Wongpaiboonwattana, K. Sanders, M. Mudaliar, D. M. van Aalten, G. J. Barton, M. P. Stavridis, Stem Cells 2014, 32, 2605-2615; f) Y. Zhu, T. W. Liu, S. Cecioni, R. Eskandari, W. F. Zandberg, D. J. Vocadlo, Nat. Chem. Biol. 2015, 11, 319-325; g) Y. Zhu, T. W. Liu, Z. Madden, S. A. Yuzwa, K. Murray, S. Cecioni, N. Zachara, D. J. Vocadlo, J. Mol. Cell Biol. 2016, 8, 2-16; h) W. Lin, L. Gao, X.

Chen, Chembiochem 2015, 16, 2571-2575; i) P. H. Chen, T. J. Smith, J. Wu, P. F. Siesser, B. J. Bisnett, F. Khan, M. Hogue, E. Soderblom, F. Tang, J. R. Marks, M. B. Major, B. M. Swarts, M. Boyce, J. T. Chi, EMBO J 2017, 36, 2233-2250; j) P. S. Banerjee, J. Ma, G.W. Hart, Proc. Natl. Acad. Sci., USA 2015, 112, 6050-6055.

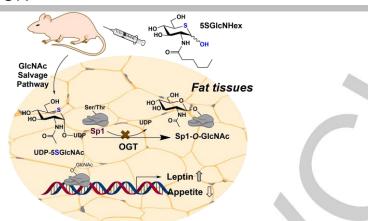
- [7] A. K. Bergfeld, O. M. Pearce, S. L. Diaz, R. Lawrence, D. J. Vocadlo, B. Choudhury, J. D. Esko, A. Varki, J. Biol. Chem. 2012, 287, 28898-28916.
- [8] a) M. S. Macauley, J. Chan, W. F. Zandberg, Y. He, G. E. Whitworth, K. A. Stubbs, S. A. Yuzwa, A. J. Bennet, A. Varki, G. J. Davies, D. J. Vocadlo, J. Biol Chem 2012, 287, 28882-28897; bD. J. Vocadlo, H. C. Hang, E. J. Kim, J. A. Hanover, C. R. Bertozzi, Proc. Natl. Acad. Sci., USA 2003, 100, 9116-9121.
 [9] M. A. Pelleymounter, M. J. Cullen, M. B. Baker, R. Hecht, D.
- [9] M. A. Pelleymounter, M. J. Cullen, M. B. Baker, R. Hecht, D. Winters, T. Boone, F. Collins, Science 1995, 269, 540-543.
 [10] L. M. Andres, I. W. Blong, A. C. Evans, N. G. Rumachik, T. Yamaguchi, N. D. Pham, P. Thompson, J. J. Kohler, C. R. Bertozzi, ACS Chem. Biol. 2017, 12, 2030-2039.
- [11] Y. Zhang, R. Proenca, M. Maffei, M. Barone, L. Leopold, J. M. Friedman, *Nature* 1994, 372, 425-432.
- J. L. Halaas, K. S. Gajiwala, M. Maffei, S. L. Cohen, B. T. Chait, D. Rabinowitz, R. L. Lallone, S. K. Burley, J. M. Friedman, *Science* 1995, 269, 543-546.
- I. Han, J. E. Kudlow, *Mol. Cell Biol.* 1997, 17, 2550-2558.
 H. B. Ruan, M. O. Dietrich, Z. W. Liu, M. R. Zimmer, M. D. Li, J. P. Singh, K. Zhang, R. Yin, J. Wu, T. L. Horvath, X. Yang, *Cell* 2014, 159, 306-317.
- [15] C. Keembiyehetty, D. C. Love, K. R. Harwood, O. Gavrilova, M. E. Comly, J. A. Hanover, *J. Biol. Chem.* 2015, 290, 7097-7113.
 [16] a) J. Wang, R. Liu, M. Hawkins, N. Barzilai, L. Rossetti, *Nature* 1998, 393, 684-688; b) E. E. Wollaston-Hayden, R. B. Harris, B. Liu, R. Bridger, Y. Xu, L. Wells, *Front Endocrinol.* 2014, 5, 223.

WILEY-VCH

COMMUNICATION

COMMUNICATION

New metabolic inhibitors of OGT enable chemical control of *O*-GlcNAc *in vivo* in mice and link decreased *O*-GlcNAc modification in fat with leptin-mediated nutrient sensing in vivo.



Tai-Wei Liu,† Wesley F. Zandberg,† Tracey M. Gloster,† Lehua Deng, Kelsey D. Murray, Xiaoyang Shan and David J Vocadlo*

Page No. - Page No.

Metabolic inhibitors of O-GlcNAc transferase (OGT) that act in vivo implicate decreased O-GlcNAc levels in leptinmediated nutrient sensing.