



Perspective

# The Two $\beta$ -Arrestins Regulate Distinct Metabolic Processes: Studies with Novel Mutant Mouse Models

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**Abstract:** The two  $\beta$ -arrestins ( $\beta$ -arrestin-1 and -2; alternative names: arrestin-2 and -3, respectively) are well known for their ability to inhibit signaling via G protein-coupled receptors. However,  $\beta$ -arrestins can also act as signaling molecules in their own right. Although the two proteins share a high degree of sequence and structural homology, early studies with cultured cells indicated that  $\beta$ -arrestin-1 and -2 are not functionally redundant. Recently, the *in vivo* metabolic roles of the two  $\beta$ -arrestins have been studied using mutant mice selectively lacking either  $\beta$ -arrestin-1 or -2 in cell types that are of particular relevance for regulating glucose and energy homeostasis. These studies demonstrated that the  $\beta$ -arrestin-1 and -2 mutant mice displayed distinct metabolic phenotypes *in vivo*, providing further evidence for the functional heterogeneity of these two highly versatile signaling proteins.

**Keywords:**  $\beta$ -arrestins; G protein-coupled receptors; diabetes; obesity; metabolism; mutant mice



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## 1. Introduction

The two  $\beta$ -arrestins,  $\beta$ -arrestin-1 and -2 (alternative nomenclature: arrestin-2 and -3, respectively), are intracellular proteins that are best known for their ability to regulate the activity of G protein-coupled receptors (GPCRs) [1,2]. In contrast to rod and cone arrestin (arrestin-1 and -4, respectively), which are mainly found in the eye,  $\beta$ -arrestin-1 ( $\beta$ arr1) and  $\beta$ -arrestin-2 ( $\beta$ arr2) are found in virtually every cell type [1,2]. Following GPCR occupation by agonist ligands, including hormones, neurotransmitters, metabolites, or sensory stimuli, most GPCRs are subject to phosphorylation by GPCR kinases (GRKs). This structural modification enables the two  $\beta$ -arrestins to bind to the intracellular surface of the receptors, thus interfering with productive receptor/G protein coupling via steric hindrance [1,2]. Moreover, due to the ability of receptor-associated  $\beta$ -arrestins to bind to clathrin and adaptor protein 2 (AP-2),  $\beta$ -arrestins play a key role in GPCR internalization via clathrin-coated pits [2,3].

Beyond these “traditional roles” of  $\beta$ -arrestins, a large body of evidence indicates that  $\beta$ -arrestins can act as signaling molecules in their own right, often by serving as scaffolding proteins for various intracellular signal transduction cascades [4–6]. The best-known example is the ability of  $\beta$ -arrestins to stimulate signaling via different mitogen-activated protein kinase (MAPK) signaling pathways [4–6]. While many of these non-canonical  $\beta$ -arrestin activities are predicted to require the prior recruitment of  $\beta$ -arrestins by GPCRs, GPCR-independent  $\beta$ -arrestin functions have also been reported [4–6]. In addition, recent studies suggest that at least some of the cellular functions of  $\beta$ -arrestins require the presence of functional G proteins [7–9].

$\beta$ arr1 and  $\beta$ arr2 are found in all mammals, suggesting that the existence of the two  $\beta$ -arrestin isoforms is advantageous from an evolutionary point of view [10]. The two  $\beta$ -arrestins share more than 70% identity at the amino acid level and have very similar three-dimensional structures [11]. For this reason, it is not surprising that the two proteins share many similar functions. However, early studies with cultured cells clearly indicated

that  $\beta$ arr1 and  $\beta$ arr2 are not functionally redundant [1,11]. For example,  $\beta$ arr2 has higher affinity for many GPCRs than  $\beta$ arr1, although some GPCRs preferentially recruit  $\beta$ arr1 [10]. Another striking example highlighting this functional heterogeneity is the observation that  $\beta$ arr2, but not  $\beta$ arr1, can promote the activation of c-jun N-terminal kinase 3 (JNK3) [10]. In agreement with these findings, an early global proteomics study using cultured HEK293 cells showed that the two  $\beta$ -arrestins are endowed with distinct protein interaction profiles, both under basal conditions and after stimulation of angiotensin II type 1a receptor signaling [12].

One possibility is that minor local conformational differences between  $\beta$ arr1 and  $\beta$ arr2 contribute to the ability of the two proteins to affect cellular functions in an isoform-specific fashion. In agreement with this notion, subtle structural differences have been observed in the inter-domain hinge region of activated  $\beta$ arr1 and  $\beta$ arr2 [11].

While the two  $\beta$ -arrestins are primarily found in the cytoplasm, both  $\beta$ arr1 and  $\beta$ arr2 contain a nuclear localization sequence [13]. However, since  $\beta$ arr2 also harbors a nuclear export signal domain,  $\beta$ arr2, but not  $\beta$ arr1, is predicted to be rapidly exported back to the cytoplasm [11]. In agreement with this concept, accumulating evidence indicates that nuclear  $\beta$ arr1 can regulate several important transcriptional processes [13]. These findings suggest that differences in subcellular localization can also contribute to the functional divergence of the two  $\beta$ -arrestins [11].

## 2. Studies with Whole-Body $\beta$ -Arrestin Knockout (KO) Mice

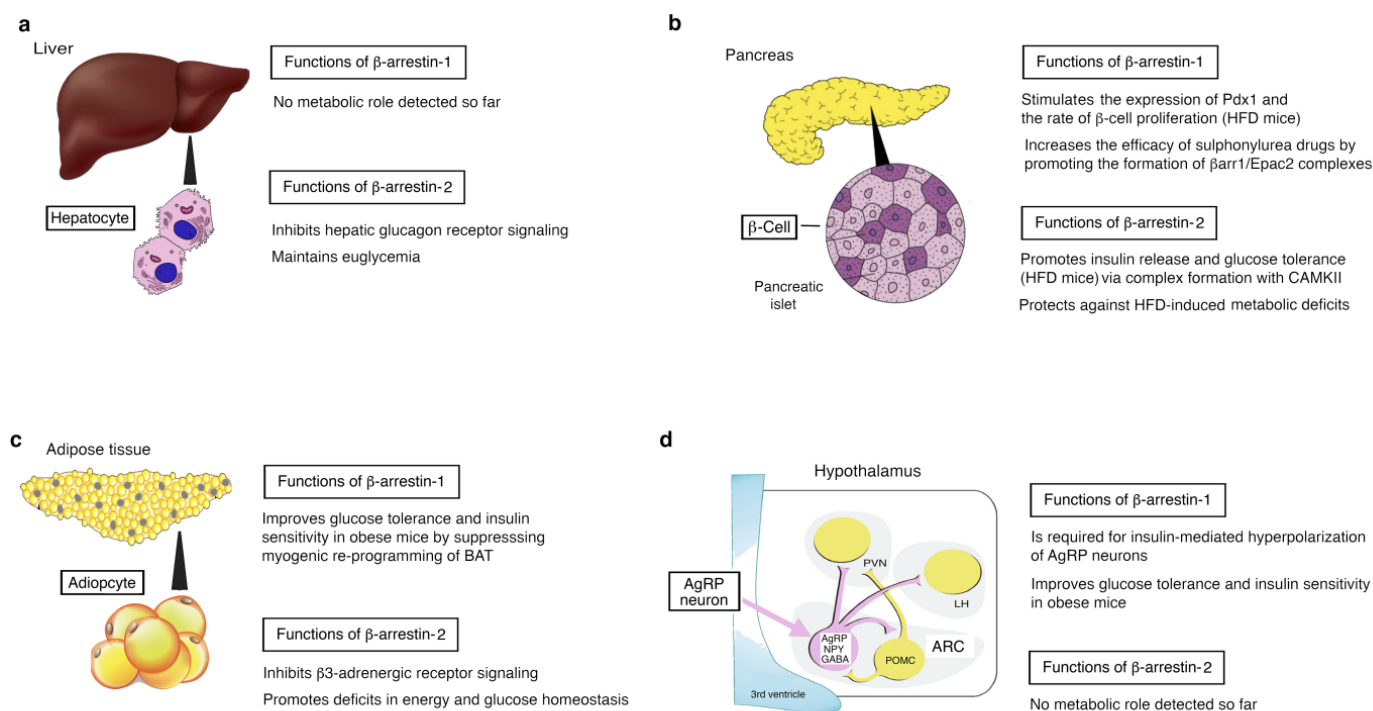
In agreement with published in vitro data, studies with whole-body  $\beta$ -arrestin KO mice confirmed that  $\beta$ arr1 and  $\beta$ arr2 differ in the physiological processes they regulate in vivo [14]. For example, nicotinic acid (niacin), an FDA-approved drug, lowers the plasma lipid levels by activation of the  $G_i$ -coupled hydrocarboxylic acid 2 (HCA<sub>2</sub>) receptors (alternative name: GPR109A) expressed by adipocytes [15]. A major side effect caused by nicotinic acid is the “niacin flush”, a flush of red on the skin that is frequently accompanied by an itching or burning sensation. This response is greatly reduced in  $\beta$ arr1 KO but not  $\beta$ arr2 KO mice [16]. Another striking example highlighting the different in vivo functions of the two  $\beta$ -arrestins are the different metabolic phenotypes displayed by whole-body  $\beta$ arr1 and  $\beta$ arr2 KO mice [17].

## 3. Analysis of Cell-Type Specific $\beta$ -Arrestin KO Mice

The recent availability of floxed  $\beta$ arr1 and  $\beta$ arr2 mice has made it possible to delete either of the two  $\beta$ -arrestin isoforms in a cell type-specific fashion [18,19]. As a result, it is now possible to compare the in vivo physiological importance of  $\beta$ arr1 and  $\beta$ arr2 expressed by a particular cell type. Specifically, recent work has targeted cell types that play critical roles in regulating glucose and energy homeostasis [20]. In the following, I will briefly summarize the outcome of studies carried out with cell type-specific  $\beta$ -arrestin KO mice that provide additional in vivo evidence for the functional heterogeneity of the two  $\beta$ -arrestin isoforms.

### 3.1. Hepatocytes

Mice that selectively lack  $\beta$ arr1 in hepatocytes do not show any impairments in glucose homeostasis [21]. In contrast, hepatocyte-specific  $\beta$ arr2 KO mice display striking metabolic deficits, primarily due to increased hepatic glucagon receptor (GCGR) signaling [21] (Figure 1a). While glucagon-induced GCGR internalization remains intact in hepatocytes lacking  $\beta$ arr1, this process is severely disrupted in  $\beta$ arr2-deficient hepatocytes. Since receptor internalization contributes to GPCR desensitization, the most likely scenario is that the lack of GCGR internalization caused by  $\beta$ arr2 deficiency plays a key role in promoting GCGR signaling in  $\beta$ arr2-deficient hepatocytes [21].



**Figure 1.** The two  $\beta$ -arrestins regulate different functions in metabolically important cell types in vivo. (a–d) Summary of the outcome of metabolic studies with mutant mice lacking  $\beta$ arr1 or  $\beta$ arr2 selectively in mouse hepatocytes (a), pancreatic  $\beta$ -cells (b), adipocytes (c), and AgRP neurons of the arcuate nucleus of the hypothalamus (d) (for a review, see [20]). See text for details. HFD, high-fat diet; CAMKII, calcium/calmodulin-dependent protein kinase II; BAT, brown adipose tissue; AgRP, agouti-related peptide; NPY, neuropeptide Y; POMC, proopiomelanocortin; ARC, arcuate nucleus; PVN, paraventricular nucleus; LH, lateral hypothalamus.

### 3.2. Pancreatic $\beta$ -Cells

The selective inactivation of  $\beta$ arr1 or  $\beta$ arr2 in pancreatic  $\beta$ -cells also results in well-defined metabolic phenotypes [22–24] (Figure 1b). Mice that selectively lack  $\beta$ -cell  $\beta$ arr2 show significantly impaired insulin release when the mutant mice are maintained on a calorie-rich diet [23]. Studies with isolated islets showed that glucose-induced insulin secretion is greatly reduced in  $\beta$ arr2-deficient  $\beta$ -cells, most likely due to impaired function of calcium/calmodulin-dependent protein kinase type II (CAMKII), a multi-functional Ser/Thr protein kinase that plays an important role in promoting insulin exocytosis [23]. Biochemical studies indicated that  $\beta$ arr2 can interact with CAMKII in a protein complex that is critical for the proper function of CAMKII [23]. It remains unknown at present whether this  $\beta$ arr2 function is regulated by the activity of  $\beta$ -cell GPCRs.

Like the  $\beta$ -cell  $\beta$ arr2 KO mice,  $\beta$ -cell  $\beta$ arr1 KO mice display significant impairments in glucose tolerance and glucose-dependent insulin secretion when maintained on an obesogenic diet [24]. Interestingly, Barella et al. [24] reported that obese  $\beta$ -cell  $\beta$ arr1 KO mice exhibit a striking decrease in  $\beta$ -cell mass and rate of  $\beta$ -cell proliferation (Figure 1b). Additional studies showed that  $\beta$ -cell  $\beta$ arr1 is required for the proper expression of the transcription factor Pdx1, the master regulator of  $\beta$ -cell function and  $\beta$ -cell mass expansion [25]. Barella et al. [24] concluded that the lack of nuclear  $\beta$ arr1 leads to reduced Pdx1 expression and that this deficit underlies the metabolic impairments displayed by obese  $\beta$ -cell  $\beta$ arr1 KO mice [24].

Somewhat surprisingly, a related study [22] showed that the presence of  $\beta$ -cell  $\beta$ arr1 is required for most antidiabetic drugs of the sulphonylurea (SU) family to simulate insulin release with high efficacy (Figure 1b). Mechanistic studies revealed that  $\beta$ arr1 enhances

SU-induced insulin release by promoting SU-dependent activation of Epac2 via formation of a  $\beta$ arr1/Epac2 complex that triggers Rap1 activation and insulin secretion [22].

### 3.3. Adipocytes

Mice that selectively lack  $\beta$ arr2 in adipocytes are protected against high-fat diet-induced weight gain and the associated metabolic deficits, including impaired glucose tolerance and insulin resistance [26]. Pydi et al. [26] showed that the metabolic improvements caused by adipocyte  $\beta$ arr2 deficiency are mediated by the browning/beiging of white adipose tissue. At the cellular level,  $\beta$ arr2 acts as a strong negative regulator of adipocyte  $\beta$ 3-adrenergic receptor ( $\beta$ 3-AR) signaling by promoting the internalization of this receptor subtype [26] (Figure 1c). In mice,  $\beta$ 3-ARs are known to mediate the browning/beiging of white adipose tissue caused by activation of the sympathetic nervous system [27].

Strikingly, mutant mice that selectively lack  $\beta$ arr1 in adipocytes exhibit metabolic phenotypes that are opposite to those caused by adipocyte  $\beta$ arr2 deficiency [28]. The absence of  $\beta$ arr1 in adipocytes results in greatly impaired glucose tolerance and insulin resistance when mice are maintained on an obesogenic diet. Pydi et al. [28] demonstrated that  $\beta$ arr1 deficiency promotes the expression of myostatin in brown adipose tissue, resulting in elevated plasma myostatin levels that eventually trigger peripheral insulin resistance. These and other findings indicate that  $\beta$ arr1-mediated suppression of myostatin expression by brown adipose tissue is required for maintaining proper insulin responsiveness and blood glucose homeostasis [28] (Figure 1c).

### 3.4. Agouti-Related Protein (AgRP) Neurons

AgRP neurons, located in the arcuate nucleus of the hypothalamus, play a key role in the central regulation of food intake, energy expenditure, and glucose homeostasis [29]. Interestingly, mutant mice selectively lacking  $\beta$ arr1 in AgRP neurons mice show significant impairments in glucose tolerance and insulin sensitivity when consuming an obesogenic diet [30] (Figure 1d). This phenotype was not observed with mice in which  $\beta$ arr2 was selectively inactivated in AgRP neurons [30]. Electrophysiological studies indicated that  $\beta$ arr1 is required for the ability of insulin to 'silence' AgRP neurons, resulting in multiple beneficial metabolic effects. One possible mechanism underlying this finding is the ability of  $\beta$ arr1 to stabilize insulin receptor substrate 1 (IRS-1), a key transducer of insulin receptor signaling, via complex formation [30].

### 3.5. Concluding Remarks

In summary, studies with cell type-specific  $\beta$ arr1 and  $\beta$ arr2 mutant mice strongly support the concept that the two  $\beta$ -arrestins regulate distinct physiological processes. While some of these effects can be explained by the traditional roles of  $\beta$ -arrestins as inhibitors of GPCR signaling, many of the phenotypes observed with the newly developed  $\beta$ -arrestin mutant mice are consistent with alternative  $\beta$ -arrestin functions. It remains to be determined to which extent these novel  $\beta$ -arrestin functions are regulated by GPCR signaling and/or GPCR/ $\beta$ -arrestin interactions. The outcome of the phenotyping studies summarized in this short article may guide the development of novel drugs capable of modulating the  $\beta$ arr1 or  $\beta$ arr2 activity or expression levels for the treatment of various pathophysiological conditions, including type 2 diabetes and related metabolic disorders (for a detailed review of potential therapeutic opportunities, see [20]).

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## References

1. DeWire, S.M.; Ahn, S.; Lefkowitz, R.J.; Shenoy, S.K. Beta-arrestins and cell signaling. *Annu. Rev. Physiol.* **2007**, *69*, 483–510. [[CrossRef](#)]
2. Pierce, K.L.; Lefkowitz, R.J. Classical and new roles of beta-arrestins in the regulation of G-protein-coupled receptors. *Nat. Rev. Neurosci.* **2001**, *2*, 727–733. [[CrossRef](#)]
3. Tian, X.; Kang, D.S.; Benovic, J.L.  $\beta$ -arrestins and G protein-coupled receptor trafficking. *Handb. Exp. Pharm.* **2014**, *219*, 173–186.
4. Gurevich, V.V.; Gurevich, E.V. Plethora of functions packed into 45 kDa arrestins: Biological implications and possible therapeutic strategies. *Cell. Mol. Life Sci.* **2019**, *76*, 4413–4421. [[CrossRef](#)]
5. Peterson, Y.K.; Luttrell, L.M. The Diverse Roles of Arrestin Scaffolds in G Protein-Coupled Receptor Signaling. *Pharm. Rev.* **2017**, *69*, 256–297. [[CrossRef](#)]
6. Ahn, S.; Shenoy, S.K.; Luttrell, L.M.; Lefkowitz, R.J. SnapShot:  $\beta$ -Arrestin Functions. *Cell* **2020**, *182*, 1362–1362.e1. [[CrossRef](#)]
7. Grundmann, M.; Merten, N.; Malfacini, D.; Inoue, A.; Preis, P.; Simon, K.; Ruttiger, N.; Ziegler, N.; Benkel, T.; Schmitt, N.K.; et al. Lack of beta-arrestin signaling in the absence of active G proteins. *Nat. Commun.* **2018**, *9*, 341. [[CrossRef](#)]
8. O'Hayre, M.; Eichel, K.; Avino, S.; Zhao, X.; Steffen, D.J.; Feng, X.; Kawakami, K.; Aoki, J.; Messer, K.; Sunahara, R.; et al. Genetic evidence that  $\beta$ -arrestins are dispensable for the initiation of  $\beta(2)$ -adrenergic receptor signaling to ERK. *Sci. Signal.* **2017**, *10*, eaal3395. [[CrossRef](#)]
9. Smith, J.S.; Pack, T.F.; Inoue, A.; Lee, C.; Zheng, K.; Choi, I.; Eiger, D.S.; Warman, A.; Xiong, X.; Ma, Z.; et al. Noncanonical scaffolding of G(xi) and  $\beta$ -arrestin by G protein-coupled receptors. *Science* **2021**, *371*, eaay1833. [[CrossRef](#)]
10. Gurevich, V.V.; Gurevich, E.V. Biased GPCR signaling: Possible mechanisms and inherent limitations. *Pharm. Ther.* **2020**, *211*, 107540. [[CrossRef](#)] [[PubMed](#)]
11. Srivastava, A.; Gupta, B.; Gupta, C.; Shukla, A.K. Emerging Functional Divergence of beta-Arrestin Isoforms in GPCR Function. *Trends Endocrinol. Metab.* **2015**, *26*, 628–642. [[CrossRef](#)] [[PubMed](#)]
12. Xiao, K.; McClatchy, D.B.; Shukla, A.K.; Zhao, Y.; Chen, M.; Shenoy, S.K.; Yates, J.R., 3rd; Lefkowitz, R.J. Functional specialization of beta-arrestin interactions revealed by proteomic analysis. *Proc. Natl. Acad. Sci. USA* **2007**, *104*, 12011–12016. [[CrossRef](#)] [[PubMed](#)]
13. Ma, L.; Pei, G. Beta-arrestin signaling and regulation of transcription. *J. Cell Sci.* **2007**, *120*, 213–218. [[CrossRef](#)] [[PubMed](#)]
14. Schmid, C.L.; Bohn, L.M. Physiological and pharmacological implications of beta-arrestin regulation. *Pharmacology* **2009**, *121*, 285–293. [[CrossRef](#)]
15. Offermanns, S. Hydroxy-Carboxylic Acid Receptor Actions in Metabolism. *Trends Endocrinol. Metab.* **2017**, *28*, 227–236. [[CrossRef](#)]
16. Walters, R.W.; Shukla, A.K.; Kovacs, J.J.; Violin, J.D.; DeWire, S.M.; Lam, C.M.; Chen, J.R.; Muehlbauer, M.J.; Whalen, E.J.; Lefkowitz, R.J. beta-Arrestin1 mediates nicotinic acid-induced flushing, but not its antilipolytic effect, in mice. *J. Clin. Investig.* **2009**, *119*, 1312–1321. [[CrossRef](#)]
17. Zhao, J.; Pei, G. Arrestins in metabolic regulation. *Prog. Mol. Biol. Transl. Sci.* **2013**, *118*, 413–427.
18. Urs, N.M.; Gee, S.M.; Pack, T.F.; McCorvy, J.D.; Evron, T.; Snyder, J.C.; Yang, X.; Rodriguiz, R.M.; Borrelli, E.; Wetsel, W.C.; et al. Distinct cortical and striatal actions of a beta-arrestin-biased dopamine D2 receptor ligand reveal unique antipsychotic-like properties. *Proc. Natl. Acad. Sci. USA* **2016**, *113*, E8178–E8186. [[CrossRef](#)]
19. Kim, J.; Grotegut, C.A.; Wisler, J.W.; Li, T.; Mao, L.; Chen, M.; Chen, W.; Rosenberg, P.B.; Rockman, H.A.; Lefkowitz, R.J. beta-arrestin 1 regulates beta2-adrenergic receptor-mediated skeletal muscle hypertrophy and contractility. *Skelet. Muscle* **2018**, *8*, 39. [[CrossRef](#)]
20. Pydi, S.P.; Barella, L.F.; Zhu, L.; Meister, J.; Rossi, M.; Wess, J.  $\beta$ -Arrestins as Important Regulators of Glucose and Energy Homeostasis. *Annu. Rev. Physiol.* **2021**, *84*. [[CrossRef](#)]
21. Zhu, L.; Rossi, M.; Cui, Y.; Lee, R.J.; Sakamoto, W.; Perry, N.A.; Urs, N.M.; Caron, M.G.; Gurevich, V.V.; Godlewski, G.; et al. Hepatic beta-arrestin 2 is essential for maintaining euglycemia. *J. Clin. Investig.* **2017**, *127*, 2941–2945. [[CrossRef](#)]
22. Barella, L.F.; Rossi, M.; Zhu, L.; Cui, Y.; Mei, F.C.; Cheng, X.; Chen, W.; Gurevich, V.V.; Wess, J.  $\beta$ -Cell-intrinsic  $\beta$ -arrestin 1 signaling enhances sulfonylurea-induced insulin secretion. *J. Clin. Investig.* **2019**, *129*, 3732–3737. [[CrossRef](#)]
23. Zhu, L.; Almaca, J.; Dadi, P.K.; Hong, H.; Sakamoto, W.; Rossi, M.; Lee, R.J.; Vierra, N.C.; Lu, H.; Cui, Y.; et al. beta-arrestin-2 is an essential regulator of pancreatic beta-cell function under physiological and pathophysiological conditions. *Nat. Commun.* **2017**, *8*, 14295. [[CrossRef](#)]
24. Barella, L.F.; Rossi, M.; Pydi, S.P.; Meister, J.; Jain, S.; Cui, Y.; Gavrilova, O.; Fulgenzi, G.; Tessarollo, L.; Wess, J.  $\beta$ -Arrestin-1 is required for adaptive  $\beta$ -cell mass expansion during obesity. *Nat. Commun.* **2021**, *12*, 3385. [[CrossRef](#)]
25. Spaeth, J.M.; Walker, E.M.; Stein, R. Impact of Pdx1-associated chromatin modifiers on islet  $\beta$ -cells. *Diabetes Obes. Metab.* **2016**, *18*, 123–127. [[CrossRef](#)]
26. Pydi, S.P.; Jain, S.; Tung, W.; Cui, Y.; Zhu, L.; Sakamoto, W.; Jain, S.; Abel, B.S.; Skarulis, M.C.; Liu, J.; et al. Adipocyte beta-arrestin-2 is essential for maintaining whole body glucose and energy homeostasis. *Nat. Commun.* **2019**, *10*, 2936. [[CrossRef](#)] [[PubMed](#)]
27. Harms, M.; Seale, P. Brown and beige fat: Development, function and therapeutic potential. *Nat. Med.* **2013**, *19*, 1252–1263. [[CrossRef](#)]

28. Pydi, S.P.; Jain, S.; Barella, L.F.; Zhu, L.; Sakamoto, W.; Meister, J.; Wang, L.; Lu, H.; Cui, Y.; Gavrilova, O.; et al. Beta-arrestin-1 suppresses myogenic reprogramming of brown fat to maintain euglycemia. *Sci. Adv.* **2020**, *6*, eaba1733. [[CrossRef](#)]
29. Deem, J.D.; Faber, C.L.; Morton, G.J. AgRP neurons: Regulators of feeding, energy expenditure, and behavior. *FEBS J.* **2021**. [[CrossRef](#)] [[PubMed](#)]
30. Pydi, S.P.; Cui, Z.; He, Z.; Barella, L.F.; Pham, J.; Cui, Y.; Oberlin, D.J.; Egritag, H.E.; Urs, N.; Gavrilova, O.; et al. Beneficial metabolic role of  $\beta$ -arrestin-1 expressed by AgRP neurons. *Sci. Adv.* **2020**, *6*, eaaz1341. [[CrossRef](#)] [[PubMed](#)]