

SUPPLEMENTARY DATA

The BBSome in POMC and AgRP Neurons is Necessary for Body Weight Regulation and Sorting of Metabolic Receptors

Deng-Fu Guo¹, Zhihong Lin¹, Yuanming Wu², Charles Searby³, Daniel R. Thedens⁴, George B. Richerson², Yuriy M. Usachev^{1,7}, Justin L. Grobe^{1,6,7}, Val C. Sheffield^{3,6,7} and Kamal Rahmouni^{1,5,6,7,*}

Department of Pharmacology¹, Department of Neurology², Department of Pediatrics³, Department of Radiology⁴, Department of Internal Medicine⁵, Obesity Education and Research Initiative⁶ and Fraternal Order of Eagles Diabetes Research Center⁷, University of Iowa, Iowa City, Iowa

Short title: Neuronal BBSome and Energy Homeostasis.

Contain:

Supplemental figures 1-8

Supplemental table 1

***Corresponding author:**

Kamal Rahmouni, Ph.D.

Department of Pharmacology

University of Iowa Carver College of Medicine

Iowa City, IA, 52242, USA

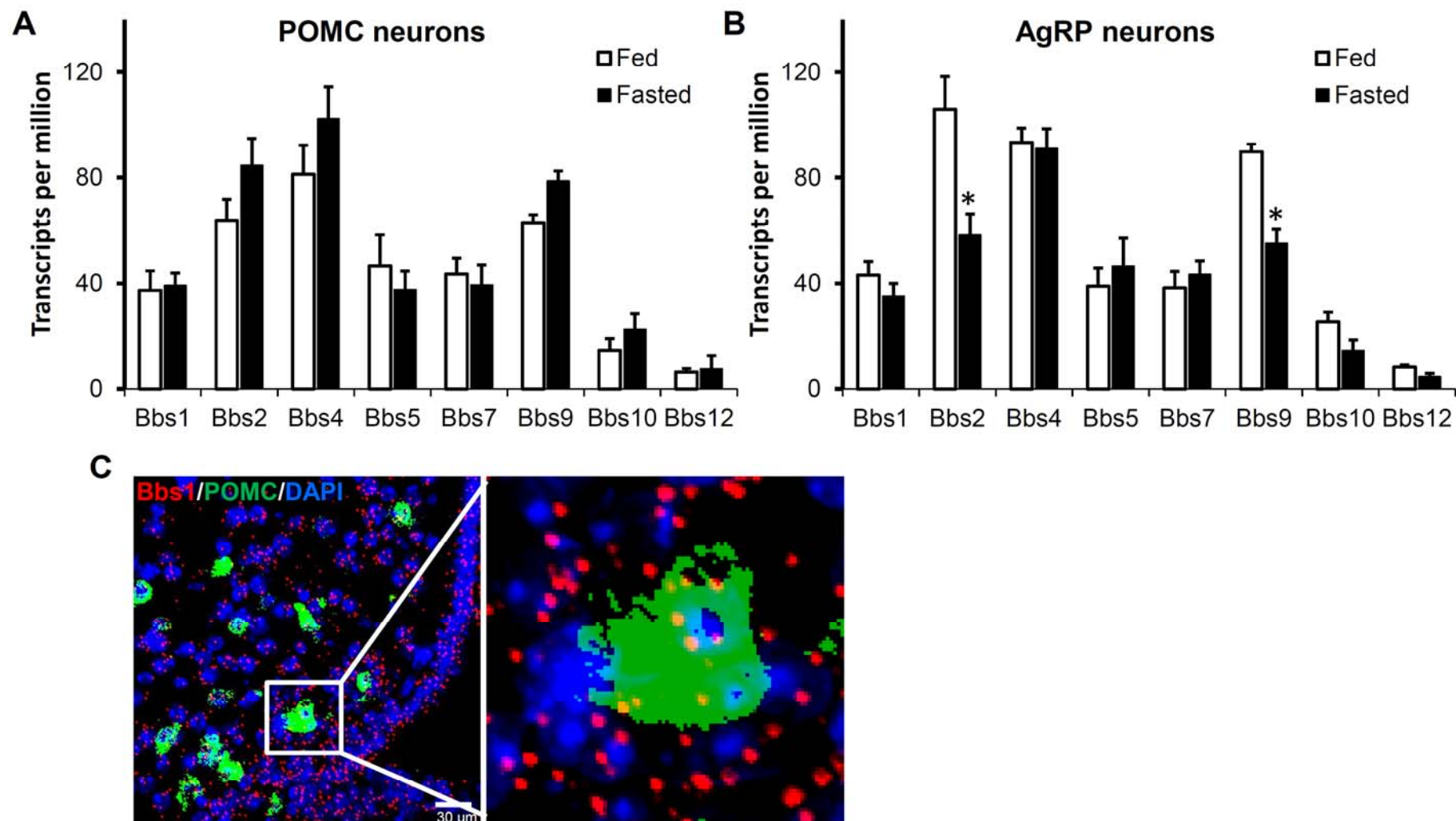
e-mail: kamal-rahmouni@uiowa.edu

Tel: 319 353 5256

Fax: 319 353 5350

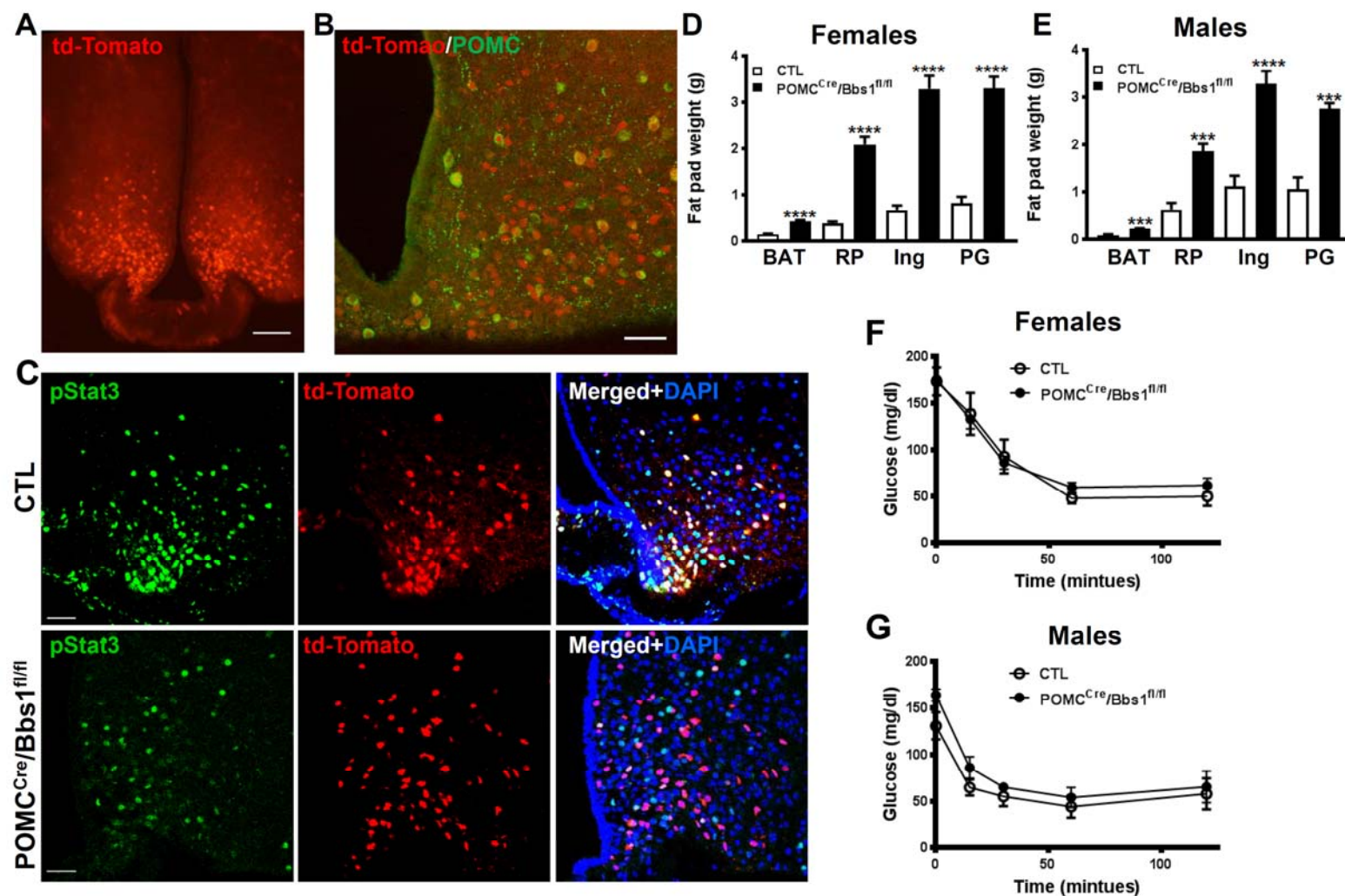
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Supplementary Figure S1. Expression of BBS genes in POMC and AgRP neurons is modulated by energy status. (A-B) In silico reanalysis of available data (ref 23) showing BBS gene expression in POMC and AgRP neurons in fed versus fasted mice (cells obtained from 6 fed mice and 5 fasted mice were used for single-cell RNA sequencing). (C) Co-localization of Bbs1 mRNA and POMC mRNA by RNAscope. * $P < 0.05$, vs. the fed group.



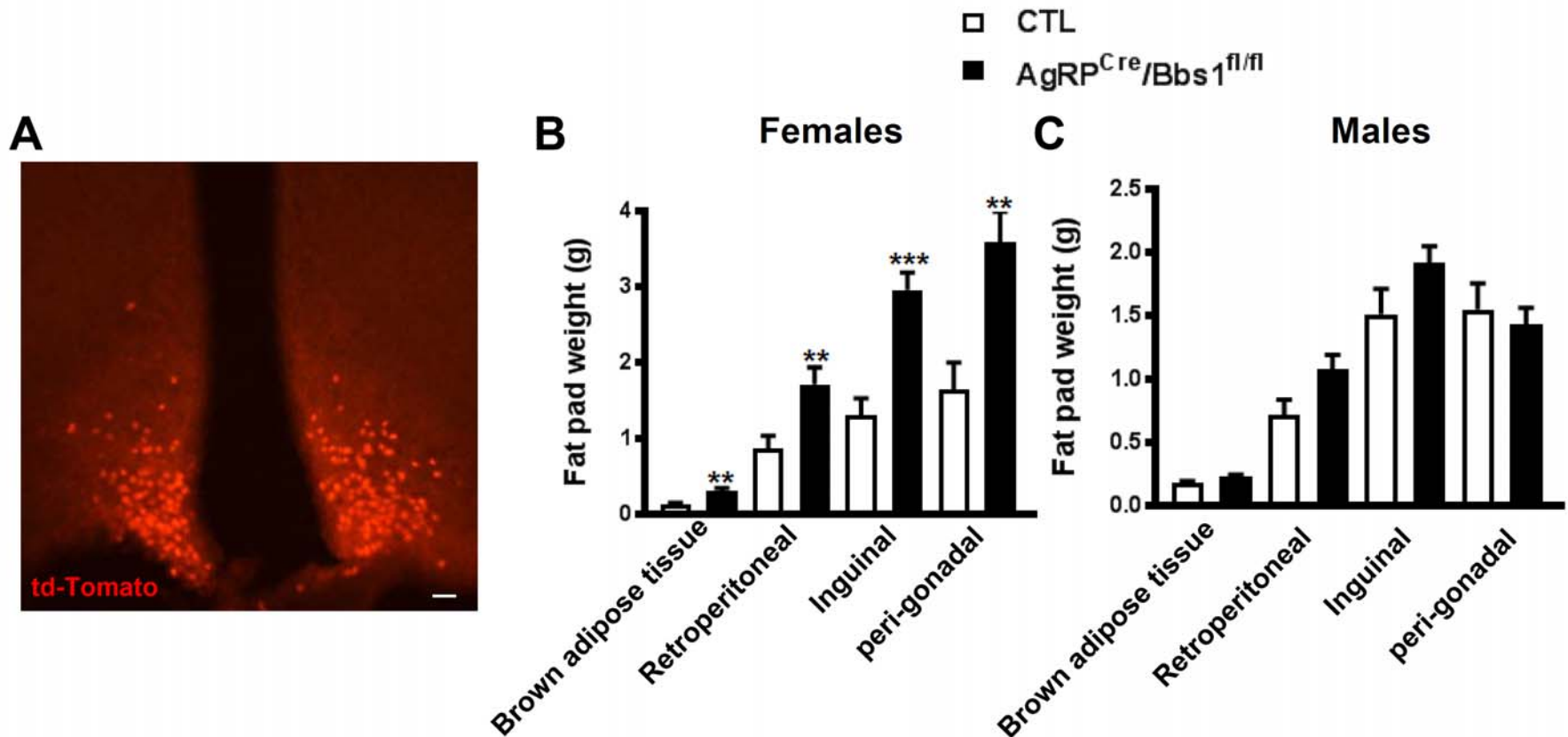
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Supplementary Figure S2. Evidence of Cre recombinase (presence of td-Tomato) (A) and its co-localization with POMC (B) in the arcuate nucleus of POMC^{Cre}/Td-Tomato mice. (C) Intraperitoneal leptin increased p-Stat3 in POMC neurons of control mice, but not POMC^{Cre}/Bbs1^{fl/fl} mice. (D-E) Weight of fat pads of POMC^{Cre}/Bbs1^{fl/fl} mice compared to control littermates (n=6-8 per group: 7 males and 7 females for controls and 8 males and 6 females for POMC^{Cre}/Bbs1^{fl/fl} mice). (F-G) Insulin tolerance test (n=10-12/group: 5 males and 5 females for controls and 6 males and 6 females for POMC^{Cre}/Bbs1^{fl/fl} mice). BAT: brown adipose tissue, RP: retroperitoneal fat, Ing: inguinal fat, PG: perigonadal fat. ***P<0.001 and ****P<0.0001 vs. control group.



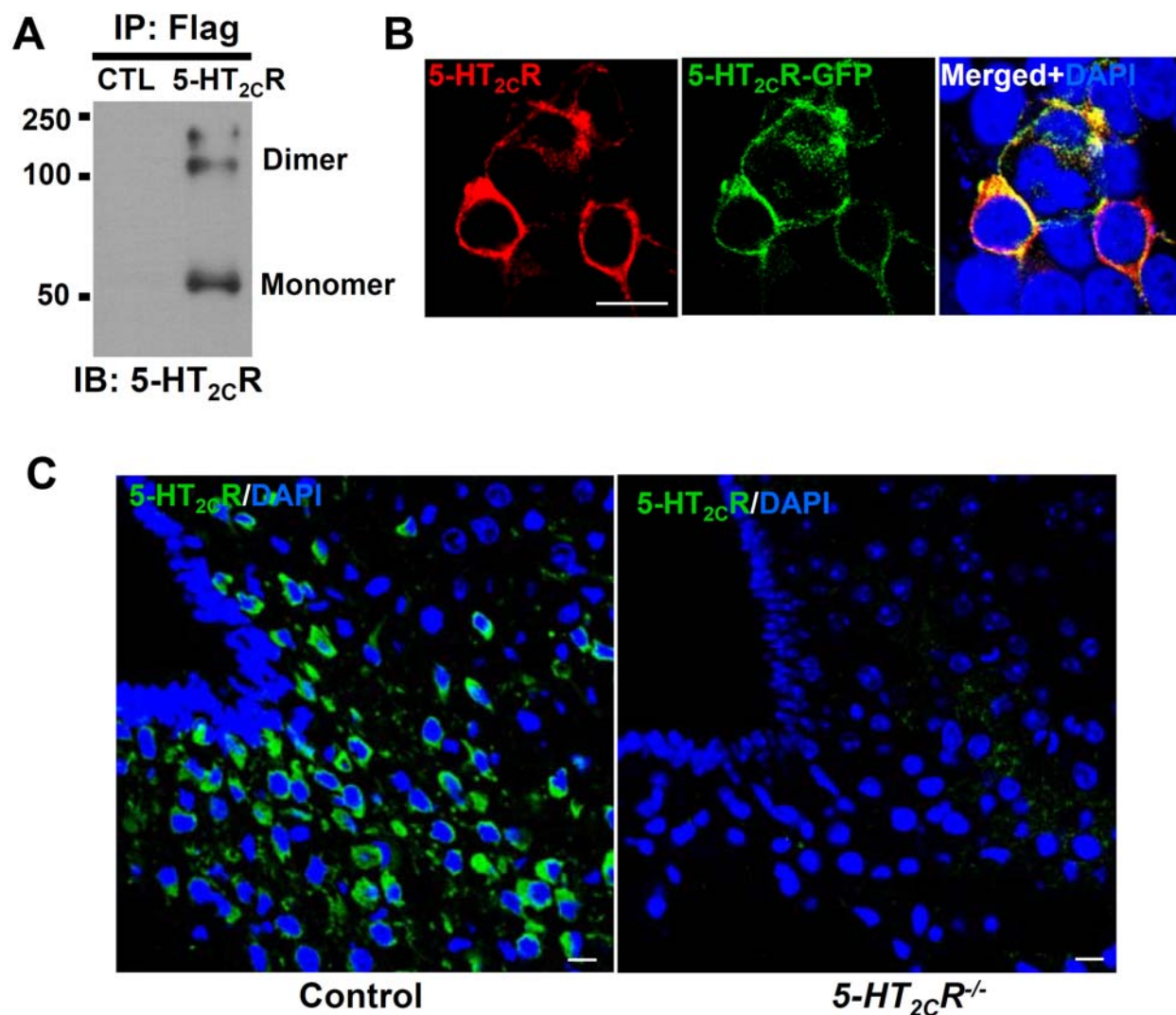
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Supplementary Figure S3. BBSome disruption in AgRP neurons causes obesity. (A) Evidence of Cre recombinase (presence of td-Tomato) in the arcuate nucleus of the hypothalamus (where AgRP neurons reside) in $\text{AgRP}^{\text{Cre}}/\text{td-Tomato}$ mice. (B-C) Weight of fat pads in female (B) and male (C) $\text{AgRP}^{\text{Cre}}/\text{Bbs1}^{\text{fl/fl}}$ mice compared to control littermates (n=6-8 per group: 7 males and 9 females for control and 7 males and 8 females for $\text{AgRP}^{\text{Cre}}/\text{Bbs1}^{\text{fl/fl}}$ mice). **P< 0.01 and ***P<0.001 vs. control group. Scale bar: 100 μm .



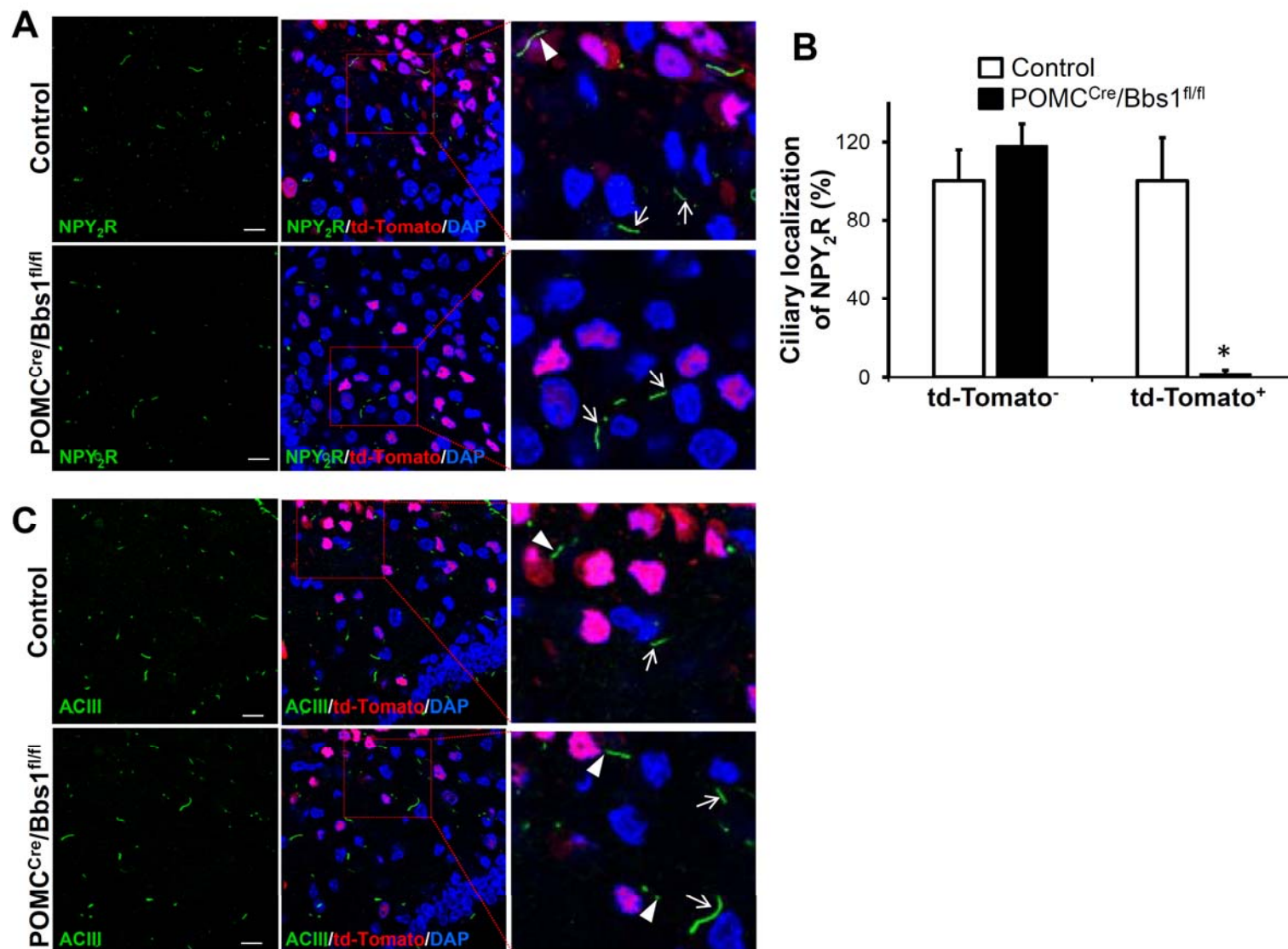
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Supplementary Figure S4. Validation of 5-HT_{2C}R antibody. (A) The anti-5-HT_{2C}R antibody detects monomer, dimer and oligomer forms of the Flag-tagged 5-HT_{2C}R in HEK 293 cells. (B) Staining with the anti-5-HT_{2C}R antibody co-localized with GFP in HEK 293 cells transfected with GFP tagged 5-HT_{2C}R (C) The anti-5-HT_{2C}R antibody detects the endogenous 5-HT_{2C}R in the hypothalamic arcuate nucleus of control mice. This staining is absent in the arcuate nucleus of 5-HT_{2C}R knockout mice. Scale bar: 10 μm



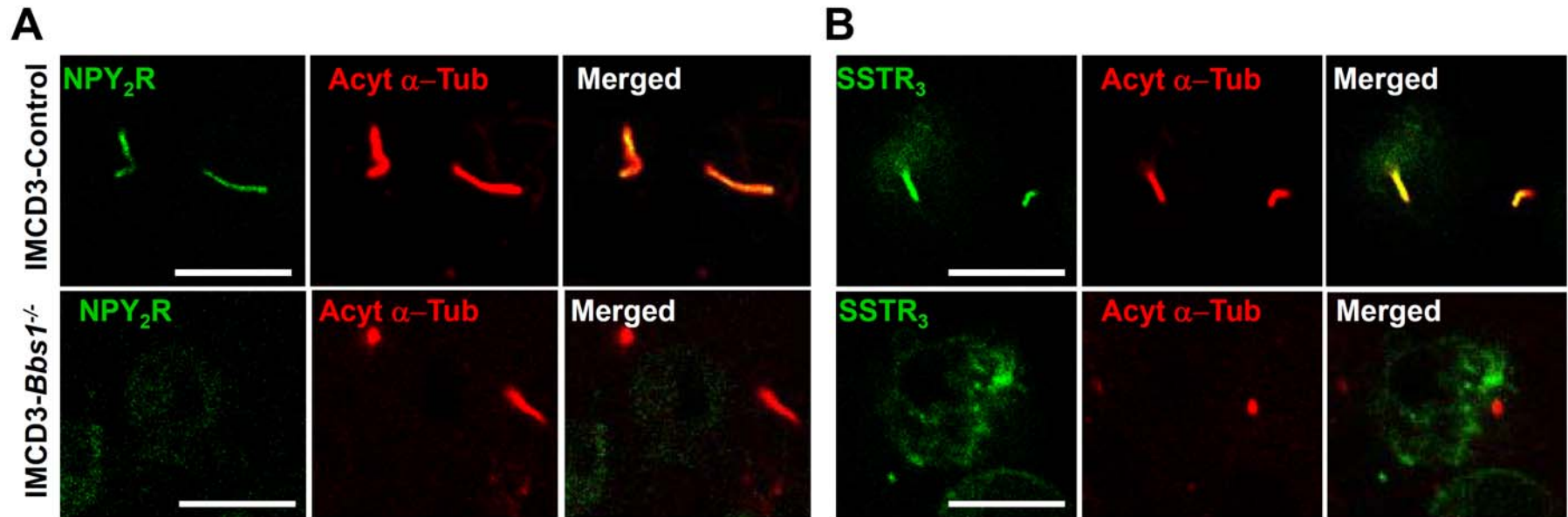
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Supplementary Figure S5. The BBSome is required for the ciliary localization of NPY₂R in POMC neurons. (A-B) Loss of NPY₂R in cilia of td-Tomato⁺ (red, arrow head), but not in td-Tomato⁻ (non-red, arrow), cells in the arcuate nucleus of POMC^{Cre}/Bbs1^{fl/fl} mice (n=3) vs controls (n=4). (C) Presence of cilia in td-Tomato⁺ (arrow head) and td-Tomato⁻ (arrow) cells of POMC^{Cre}/Bbs1^{fl/fl} mice. *P<0.05 vs Control. Scale bar: 10 μm.



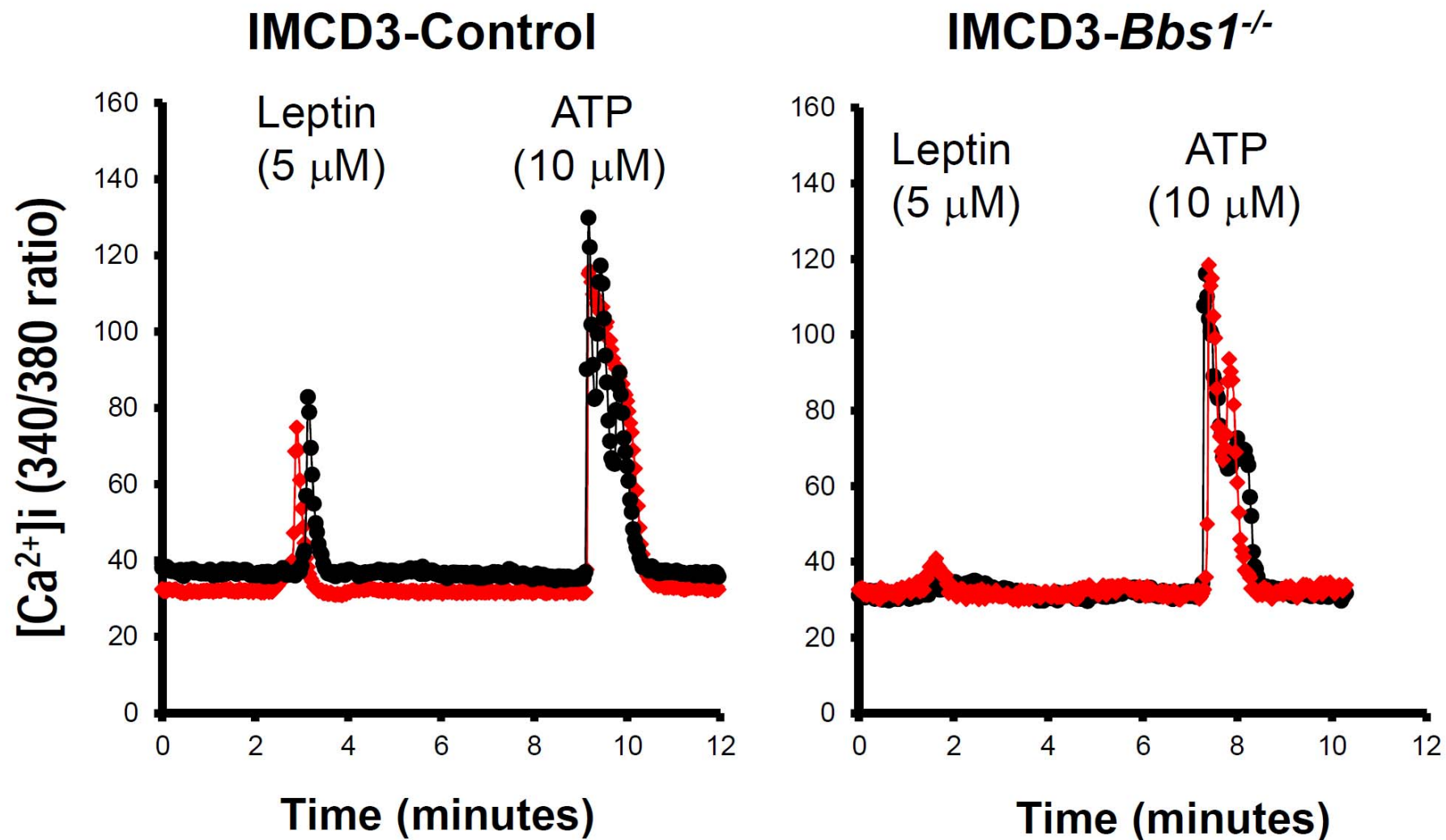
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Supplementary Figure S6. The BBSome is required for the ciliary localization of NPY₂R and SSTR₃. (A-B) Loss of localization in cilia (stained with Acetylated α -Tub) of NPY₂R (A) and SSTR₃ (B) in IMCD3-*Bbs1*^{-/-} cells. Scale bar: 10 μ m.



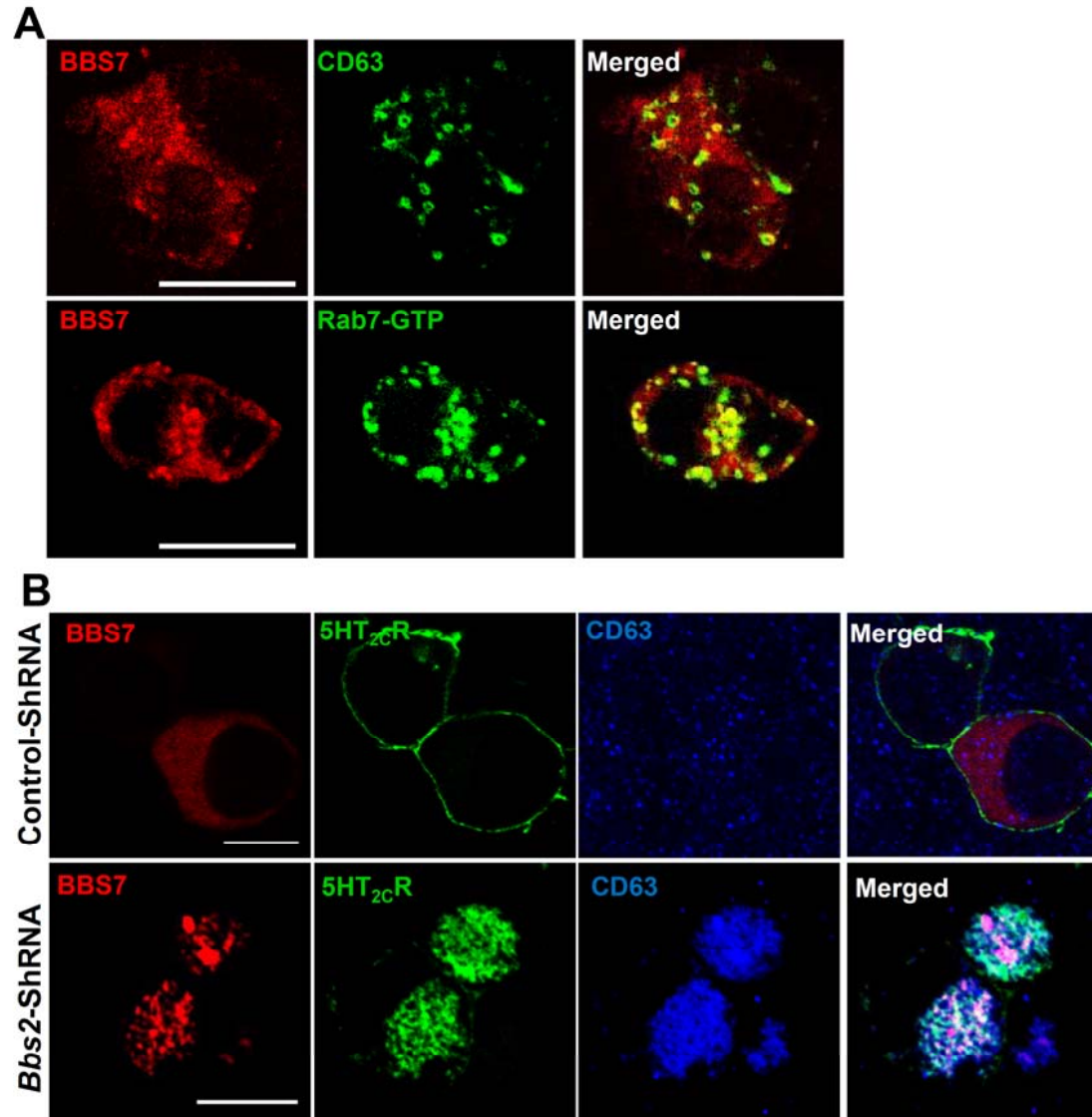
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Supplementary Figure S7. Representative tracings showing reduced $[Ca^{2+}]_i$ response to leptin in control and $Bbs1^{-/-}$ IMCD3 cells stably expressing the LepRb.



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Supplementary Figure S8. BBSome proteins interacts with 5-HT_{2C}R in the late endosome. (A) BBS7 (red) colocalizes with markers of the late endosome: CD63 (green) or Rab7-GTP (green). (B) Silencing *Bbs2* gene expression promotes BBS7 (red) and 5-HT_{2C}R (green) interaction in late endosome marked with CD63 (blue) in HEK 293 cells. Scale bar: 10 μm.



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Supplementary Table S1. Primer sequences used.

Name	Sense (5'-3')	Anti-sense (5'-3')
<u>Genotyping</u>		
<i>Cre</i>	ACCTGAAGATGTTGCGCATTATCT	ACCGTCAGTACGTGAGATATCTT
<i>Bbs1</i>	ACATCCACACCTTCTCCTCCT	GCCTACCTTGTACTIONCCCATC
<u>Hypothalamic gene expression</u>		
S18	ACTGCCATTAAGGGCGTGG	CCATCCTTCACATCCTTCTG
AgRP	CAGAAGCTTTGGCGGAGGT	AGGACTCGTGCAGCCTTACAC
NPY	TCAGACCTCTTAATGAAGGAAAGCA	GAGAACAAGTTTCATTTCCCATCA
POMC	CTGCTTCAGACCTCCATAGATGTG	CAGCGAGAGGTGAGTTTTGC
<u>Generation of <i>Bbs1</i> knock-out cells</u>		
sgRNA	CCTTTGAGCACCTTCAGGCG	
<i>Bbs1</i>	ACTGCCATTAAGGGTGTGG	CCATCCTTTACATCCTTCTG