

Fig. S 1: Effects of Shp2 knockdown in 3T3L1 cells on Akt and Erk phosphorylation. 3T3L1 cell lines with Shp2 knockdown (KD) and control (scrambled shRNA) were generated. Cells were starved O/N then stimulated with insulin (100nM) for 10, 20 and 30 minutes. Total cell lysates were immunoblotted for pAkt, Akt, pErk, Erk, Shp2 and Tubulin (as a control for loading). Bar graphs represent normalized data for pAkt/Akt and pMAPK/MAPK from two independent experiments and presented as means ± SEM. Statistical analysis was performed using two-tailed Student's *t*-test. (*) indicates significant difference between starved and insulin-stimulated cells, while (#) indicates significant difference between WT and Shp2 KD cells.





Bettaieb, et al., Figure S2



Fig. S2: Immuno-staining of Shp2 in epididymal adipose tissue. (A) Representative images of Shp2 immunofluorescnece in fl/fl (control), fl/+, Cre (heterozygous) and fl/fl, Cre (FSHKO) mice. The middle lane represents brightfield (BF) images of sections in the first lane. The experiment was repeated another day on different set of mice revealing comparable staining pattern (right lane). Please note that staining in KO sections likely reflects Shp2 expression in other cell types in the adipose tissue, such as vascular endothelial cells and macrophages. Differences in Shp2 staining in heterozygous mice reflects differences between mice. **(B)** Higher magnification of Shp2 staining in control and FSHKO mice.



Fig. S3: Expression of macrophage markers in W and CW fractions of FSHKO mice. Immunoblots of F4/80 (a mouse macrophage-specific membrane marker) expression in lysates of white adipose tissue (W) and purified adipocytes from collagenase-treated white adipose tissue (CW) from Adipoq-Cre (Cre), Shp2^{fix/fix} (fl/fl) and Adipoq-Shp2flx/flx (FSHKO) mice on a HFD for 12 weeks. Blot were also probed with another macrophage marker Mac-3 (although it is presumably less specific as suggested by Inoue, *et al.*, Kidney International, 2005). Blots were probed with anti-Tubulin antibodies (bottom panel) as a loading control. The numbers on each lane reflect samples from different mice (W1 and CW1 are from the same mouse; W2 and CW2 are from a different mouse). Notably, FSHKO lysates exhibit increased expression of macrophages markers. This is consistent with our preliminary data suggesting increased inflammation in FSHKO adipose tissue (and presumably increased macrophage infiltration).

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- 1 Adipose-Specific Deletion of Src Homology Phosphatase 2 does not <u>Significantly</u>
- 2 Alter <u>Systemic</u> Glucose Homeostasis
- 3
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22 ABSTRACT

23	<i>Objective</i> - The SH2 domain-containing protein-tyrosine phosphatase Shp2 has been
24	implicated in a variety of growth factor signaling pathways, but its metabolic role in
25	some peripheral insulin-responsive tissues remains unknown. <i>Materials/Methods</i> - To
26	address the metabolic <u>function</u> of Shp2 in adipose tissue, we generated mice with
27	adipose-specific Shp2 deletion using Adiponectin (Adipoq)-Cre transgenic mice. We
28	then analyzed insulin sensitivity, glucose tolerance and body mass in adipose-specific
29	Shp2_deficient and control mice on regular chow and high fat diet (HFD). <i>Results</i>
30	Control mice on a HFD exhibited increased Shp2 expression in various adipose depots
31	compared with those on regular chow, Adipoq-Cre mice enabled efficient and specific
32	deletion of Shp2 in adipose tissue. However, adipose Shp2 deletion did not significantly
33	alter body mass in mice on chow or HFD. In addition, mice with adipose Shp2 deletion
34	exhibited comparable insulin sensitivity and glucose tolerance compared with controls.
35	Consistent with this, basal and insulin-stimulated Erk and Akt phosphorylation were
36	comparable in adipose tissue of Shp2-deficient and control mice. Conclusions - Our
37	findings indicate that adipose-specific Shp2 deletion does not significantly alter systemic
38	insulin sensitivity and glucose homeostasis.
39	
40	Keywords: Protein-tyrosine phosphatases, diabetes, glucose homeostasis, body weight,
41	and adiposity
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- 43 Abbreviations: Shp2; Src homology phosphatase 2, PTP; protein-tyrosine phosphatase,
- 44 ITT; insulin tolerance test, and GTT; glucose tolerance test

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45	INTRODUCTION
46	Metabolic syndrome and type 2 diabetes are complex disorders that are associated
47	with obesity and sedentary life style [1, 2]. The increasing incidence of obesity
48	worldwide has focused attention on adipose tissue function and contribution to whole
49	body metabolic homeostasis. White adipose tissue (WAT) is specialized in lipid storage
50	and adipokine secretion and is a regulator of energy balance and systemic insulin
51	sensitivity [3].
52	Tyrosyl phosphorylation is a major regulator of insulin signaling and is tightly
53	controlled by the opposing actions of protein-tyrosine kinases (PTKs) and protein-
54	tyrosine phosphatases (PTPs) [4]. Src homology phosphatase 2 (Shp2) is a ubiquitously
55	expressed non-transmembrane protein-tyrosine phosphatase that contains two SH2
56	domains, a tyrosine phosphatase domain, a C-terminal region with phosphorylation sites
57	and a proline-rich domain [5]. Multiple studies indicate that Shp2 plays an essential role
58	in most receptor tyrosine kinase signaling pathways [6, 7]. <u>However, its function in</u>
59	regulating glucose homeostasis and energy balance in vivo requires additional
60	investigation.
61	In vivo studies have not completely resolved the physiological role of Shp2 in
62	insulin signaling and glucose homeostasis. Targeted mutation of Shp2 exon 3 in mice
63	leads to embryonic lethality [8], precluding detailed studies of the effects of global Shp2
64	deletion. Hemizygous mice are viable but do not manifest any apparent defects in insulin
65	action [9]. On the other hand, transgenic mice that express a presumptive dominant
66	negative mutant of Shp2 in skeletal muscle, liver and adipose tissue exhibit insulin

67 resistance and impaired insulin-stimulated glucose uptake [10]. Shp2 deletion in striated

Deleted:, where it is required for normal activation of the Erk pathway

Deleted: While multiple studies have addressed the potential role of Shp2 in regulating insulin signaling and glucose homeostasis, its overall function in this pathway in some insulin-responsive tissues remains unresolved.

68	and cardiac muscle results in insulin resistance, impaired glucose uptake in muscle cells,
69	and glucose intolerance [11], although these mice also exhibit marked dilated
70	cardiomyopathy [11, 12]. In addition, Shp2 deletion in the pancreas causes defective
71	glucose-stimulated insulin secretion and impaired glucose tolerance [13]. Moreover, we
72	recently reported that mice lacking Shp2 in the liver exhibit increased hepatic insulin
73	action and enhanced systemic insulin sensitivity [14]. <u>However, the role of adipose Shp2</u>
74	in regulating insulin sensitivity and glucose homeostasis <u>in vivo</u> remains unknown.
75	Shp2 also is implicated in regulating adiposity, body mass and leptin signaling
76	(reviewed in [15, 16]). In vitro biochemical studies identify Shp2 as a positive mediator
77	of leptin signaling through regulating tyrosine 985 site of leptin receptor [17-19]. These
78	findings are supported by <i>in vivo</i> deletion of Shp2 in postmitotic forebrain neurons with
79	the mice developing early onset obesity and leptin resistance [20]. In addition, mice with
80	proopiomelanocortin (POMC) neuron-specific Shp2 deletion exhibit elevated adiposity,
81	decreased leptin sensitivity and reduced energy expenditure [21]. Together, these studies
82	demonstrate a role for Shp2 in regulating energy balance, at least in part, through
83	modulating leptin signaling.
84	In this study we assessed the physiological effects of Shp2 in adipose tissue using
85	tissue-specific knockout approach. We determined the metabolic effects of adipose Shp2
86	deletion on body mass, systemic insulin sensitivity and glucose homeostasis in chow and

87 high fat diet-fed mice.

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88 METHODS

Mouse studies. Shp2-floxed (Shp2^{fl/fl}) mice were generated previously [22]. 89 Adiponectin (Adipoq)-Cre mice were generated and kindly provided by Dr. E. Rosen 90 (BIDMC/Harvard University), Shp2^{fl/fl} mice were on <u>a mixed</u> 129Sv/J x C57Bl/6J 91 background and Adipoq-Cre mice were on a mixed FVB x C57Bl/6J background. All 92 93 mice studied were age-matched and were maintained on a 12-hour light-dark cycle with free access to water and food. Mice were placed on standard lab chow (Purina lab chow, 94 # 5001), and in some experiments, switched to a high fat diet (HFD; 60% kcal from fat, # 95 D12492, Research Diets) at weaning. Genotyping for the Shp2 floxed allele and for the 96 presence of Cre was performed by polymerase chain reaction (PCR), using DNA 97 extracted from tails [14]. Mouse studies were conducted in line with federal regulations 98 and were approved by the Institutional Animal Care and Use Committee at University of 99 100 California Davis. 101

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Metabolic measurements. Glucose was measured in blood collected from the tail using 102 103 a glucometer (Home Aide Diagnostics). Serum insulin was determined by enzyme linked 104 immunosorbent assay (ELISA) using mouse insulin as a standard (Crystal Chem). Serum 105 leptin was assayed by ELISA using rat leptin standard (Crystal Chem). Free fatty acid 106 (FFA) and triglyceride (TG) concentrations were measured by an enzymatic colorimetric 107 method (Wako). Fed glucose measurements were taken between 7-9 am and, where 108 indicated, from mice fasted for 12 hrs. For insulin tolerance tests (ITTs), mice were 109 fasted for 4 hrs and injected intraperitoneally (i.p.) with 1 mU/g body weight human 110 insulin (HumulinR; Eli Lilly). Blood glucose values were measured before and at 15, 30,

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111 45, 60, 90 and 120 min post-injection. For glucose tolerance tests (GTTs), overnight-

fasted mice were injected with 20% D-glucose at 2 mg/g body weight, and glucose was 112

measured before and at 30, 60, 90 and 120 min following injection. 113

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115	Isolation of adipocytes. Three grams of adipose tissue were incubated at 37°C for 60
116	minutes in siliconized tubes containing 20 ml isolation buffer {0.1 M HEPES, pH 7.4,
117	0.12 M NaCl, 0.05 M KCl, 1.2 mM CaCl_2, 0.6 mM, MgSO_4.7H_2O and 1.5% (w/v) bovine
118	serum albumin Fraction V (Fisher)} containing 0.002% (w/v) collagenase (Worthington).
119	Tissue remnants were removed by filtration through a nylon screen (pore size 250 $\mu m)$
120	(Tetko) into a siliconized tube. Adipocytes were allowed to float to the surface for 5 min
121	whereafter the infranatant was aspirated through a siliconized injection needle. Further
122	purification of adipocytes was performed by adding 5 ml isolation buffer and 2 ml
123	dinonylphthalate oil (Fluka). Cells were allowed to float to the surface by centrifugation
124	at 1,000g for 5 min. The supernatant was transferred to Eppendorf tubes and cells were
125	pelleted and lysed in radio-immunoprecipitation assay (RIPA) buffer (150 mM NaCl,
126	1.0% IGEPAL® CA-630, 0.5% sodium deoxycholate, 0.1% SDS, and 50 mM Tris, pH
127	7.4 and I add 5 mM EDTA, 1 mM NaF, 1 mM sodium orthovanadate and protease
128	inhibitors)
129	
130	Biochemical analyses. For insulin signaling experiments, 30 week-old male mice were
131	fasted overnight, injected i.p. with insulin (10 mU/g body weight), and sacrificed 10
132	minutes after injection. Tissues were ground in liquid nitrogen and lysed using RIPA

buffer. Lysates were clarified by centrifugation at 13,000 rpm for 10 min and protein

Deleted: Tissues were dissected and immediately frozen in liquid nitrogen.

- 134 concentrations were determined using bicinchoninic acid protein assay kit (Pierce
- 135 Chemical). Proteins were resolved by SDS-PAGE and transferred to PVDF membranes.
- 136 Immunoblot<u>ting of lysates was</u> performed with antibodies for Shp2 (Santa Cruz;
- 137 1/10,000), PTP1B (Millipore; 1/5<u>,0</u>00), TCPTP (Mediamab; 1/2,000), pAkt (1/5,000),
- 138 Akt (1/5,000), pErk (1/10,000), Erk (1/10,000) (all from Cell Signaling) and Tubulin
- 139 (Santa Cruz; 1/5,000). Proteins were visualized using enhanced chemiluminescence
- 140 (ECL, Amersham Biosciences) and pixel intensities of immuno-reactive bands were
- 141 quantified using FluorChem 8900 (Alpha Innotech).

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- 143 **Statistical analyses.** Data are expressed as means <u>+</u> standard error of the mean (SEM).
- 144 Statistical analyses were performed using the JMP program (SAS Institute). <u>ITTs, GTTs,</u>
- 145 body weight and adiposity data were analyzed by analysis of variance (ANOVA). Post-
- 146 hoc analysis was performed using Tukey-Kramer honestly significant difference test. For
- 147 <u>biochemistry studies, comparisons between groups were performed using unpaired two-</u>
- 148 <u>tailed Student's t test</u>

Deleted: Comparisons between groups were made by unpaired two-tailed Student's t test. ITTs and GTTs were analyzed by repeated measures analysis of variance (ANOVA). Post-hoc analysis was performed using Tukey-Kramer honestly significant difference test.

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157 **RESULTS**

- 158 Generation of adipose-specific, Shp2 knockout mice. To investigate the role of adipose
- 159 Shp2 in regulating body mass and glucose homeostasis, we <u>assessed</u> the <u>physiological</u>
- 160 effects of its deletion in adipose tissue using Cre-LoxP approach. Mice with adipose-
- specific Shp2 deletion were generated by crossing Shp2^{fl/fl} (fl/fl) mice to BAC transgenic
- 162 mice expressing Cre recombinase under the control of the Adiponectin locus (Adipoq-
- 163 Cre) to generate Adipoq-Shp $2^{fl/+}$ mice. These mice were crossed to Shp $2^{fl/fl}$, yielding
- 164 Adipoq-Shp2^{fl/fl} (hereafter termed fat-specific Shp2 KO; FSHKO). FSHKO mice
- survived to adulthood, and were fertile. Efficiency of Shp2 deletion was determined
- using immunoblot analysis of lysates from whole white adipose tissue (W) and purified
- adipocytes from collagenase-treated white adipose tissue (CW) (Fig. 1A, B). Shp2
- 168 protein <u>expression was</u> comparable between Cre and fl/fl mice in white adipose tissue
- and purified adipocytes. On the other hand, <u>FSHKO mice exhibited decreased Shp2</u>
- 170 expression by ~70% in white adipose (W) and ~85% in collagenase-treated white adipose
- 171 (CW) compared with controls (Fig. 1B), These findings are consistent with complete
- 172 deletion of Shp2 in adipocytes; the residual Shp2 in FSHKO white adipose (W) lysates
- 173 <u>likely reflects Shp2 expression in other cell types in the adipose tissue, such as vascular</u>
- 174 <u>endothelial cells and macrophages</u>. Indeed, immuno-staining of Shp2 in WAT sections
- 175 of FSHKO and control mice supports this notion (data not shown). Shp2 levels were
- unchanged in other peripheral insulin-responsive tissues (liver and muscle), pancreas,
- brain and macrophages confirming the specificity of deletion (Fig. 1C, D). The
- 178 expression of other PTPs known to regulate glucose homeostasis, protein-tyrosine
- 179 phosphatase 1B (PTP1B) [23, 24] and its closely related T cell protein-tyrosine

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Deleted: Heterozygous mice expressed about 55% Shp2 compared with controls in collagenase-treated white adipose (data not shown).

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180 phosphatase (TCPTP) [25, 26] was unaltered in FSHKO mice (Fig. 1C). In addition,

- 181 Shp2 deletion also was observed in adipose tissue of old (60 weeks) FSHKO mice on
- 182 regular chow (Fig. 1D). Therefore this approach enables efficient and specific deletion of
- 183 Shp2 in adipose tissue.
- 184

185 Adipose-specific Shp2 deletion does not significantly alter body mass or adiposity.

- 186 <u>Shp2</u> protein expression, was determined in various adipose depots of control mice fed
- regular chow or HFD (for 12 weeks). Immunoblot analysis of lysates revealed that Shp2
- 188 was expressed in subcutaneous (SubQ), epididymal (Epi), retroperitoneal (Ret), visceral
- 189 (Vis) and brown adipose tissue (BAT) depots of mice fed regular chow (Fig. 2A).
- 190 Notably, mice fed a HFD exhibited significantly increased Shp2 expression in all
- 191 examined adipose depots compared with those fed regular chow (Fig. 2A, B). Next, we
- 192 evaluated the effect of adipose Shp2 deletion on body mass and adiposity in mice fed
- 193 <u>regular chow or challenged with a HFD</u>. As expected, on HFD mice gained more weight
- than their counterparts on regular chow, but comparable body weights (females and

195 males) were detected between genotypes on either diet (Fig. 2C-F). Similar data were

obtained in another independent cohort of mice on a HFD for 24 weeks (data not shown).

197 In line with this observation, white adipose tissue weight was similar in FSHKO mice

198 compared with controls on a HFD in both genders (Fig. 2G, J). In addition, adiposity

index (total adipose depot weight (g) \div body weight (g) \times 100), which correlates strongly

- with body fat percentage [27], was comparable between genotypes (Fig. 2H, K). Similar
- 201 head-rump length was also observed in mice of different genotypes (Fig. 2I, L).

202 Moreover, we assayed several parameters of whole-body lipid homeostasis. Leptin is a

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Deleted: Taken together, our findings indicate that

Deleted: To evaluate the effect of adipose Shp2 deletion on body mass regulation, mice were fed regular chow or challenged with a high fat diet. Initially, we determined

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- 203 cytokine that is produced by adipocytes and its levels typically reflect body fat content
- with lean animals normally having low serum leptin [28, 29]. Consistent with their
- 205 comparable adiposity and body weight, FSHKO mice exhibited similar fasted serum
- 206 leptin concentrations compared with controls (Table 1). Furthermore, fasted serum
- 207 triglyceride and free fatty acid concentrations were comparable between FSHKO and
- 208 controls, Together, our data indicate that adipose Shp2 protein expression increases after
- 209 high fat feeding but its deletion does not significantly alter adiposity and body weight
- 210 under the tested conditions.
- 211

212 Adipose-specific Shp2 deletion does not significantly alter systemic glucose

- 213 homesotasis. Body weights of control and FSHKO mice on regular chow and HFD were
- 214 comparable suggesting that any potential differences in glucose homeostasis are primary
- and not caused by body weight alterations. We assayed several metabolic parameters in
- 216 control and FSHKO mice on a HFD (Table 1). FSHKO mice exhibited comparable fed
- and fasted glucose and insulin concentrations compared with controls. In addition,
- insulin/glucose ratio was comparable between genotypes. To directly evaluate insulin

sensitivity *in vivo*, male and female mice on regular chow and a HFD were subjected to

220 ITTs at 14 weeks of age (Fig. 3A-D). On either diet, FSHKO mice exhibited comparable

- 221 insulin sensitivity to controls. In addition, we tested the ability of mice to clear glucose
- 222 from the peripheral circulation during intraperitoneal GTTs (Fig. 3E-H). Similarly,
- 223 FSHKO mice exhibited comparable glucose tolerance to controls on either diet.

224 Additional ITTs and GTTs were performed in another independent cohort of mice on a

HFD (for 24 weeks) revealing comparable results (data not show). Next, we evaluated

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- phosphorylation of Akt (Ser473) and Erk in control and FSHKO mice on a HFD at basal
- and insulin-stimulated (10 min) conditions (Fig. 3I, J). <u>As expected, insulin induced</u>
- 228 significant Akt (Ser473) and Erk phosphorylation in adipose tissue of FSHKO and
- 229 control mice. FSHKO mice exhibited a trend for decreased insulin-induced Akt
- 230 phosphorylation, but it did not reach statistical significance (P=0.4). In addition, no
- 231 significant differences were observed in insulin-induced Erk phosphorylation in FSHKO
- and control mice (P=0.5) (Fig. 3J). Collectively, our data indicate that adipose Shp2
- 233 deletion does not significantly alter systemic insulin sensitivity and glucose tolerance.

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DISCUSSION 234

235	The role of Shp2 in regulating glucose homeostasis, as well as its specific	
236	functions in adipose tissue, has heretofore remained <u>largely</u> unresolved. To begin to	Deleted: the
237	address these issues, we generated mice with adipose-specific Shp2 deletion using the	
238	novel Adipoq-Cre transgenic mice, <u>These</u> mice might provide some advantages	Deleted: with expression
239	compared with <u>the commonly-used</u> fatty acid binding protein <u>2</u> (aP2)-Cre mice that have	Deleted: Adipo
240	been utilized for adipose-specific deletion. Although aP2 is predominantly expressed in	
241	adipocytes postnatally [30], it is also expressed in non-adipose tissues (such as trigeminal	
242	ganglia, dorsal root ganglia and vertebrae) during development [31]. In addition, aP2 is	Deleted: In ac in adipose tissue,
243	expressed in activated macrophages [32], which are implicated in the regulation of	Deleted: are kr
244	adipose tissue inflammation and function [33, 34]. On the other hand, Adipoq-Cre mice	Deleted. Her
245	do not express Cre in <u>bone marrow-derived</u> macrophages (Fig. 1). <u>Although a</u> dditional	Deleted: How
246	studies are required to <u>fully</u> evaluate the utility of Adipoq-Cre <u>mice; they</u> enabled	Deleted: mice
247	efficient and specific deletion of Shp2 in various adipose depots.	deletion. At any Deleted: differ
248	Our studies demonstrated that Shp2 expression was dynamically regulated in	Deleted: and p achieve adipose-
249	adipose tissue <u>depots of mice on a HFD</u> . The regulatory point(s) for <u>adipose</u> .Shp2	Deleted: in adi
250	abundance <u>in response to high fat</u> feeding remains to be determined, and could be	Deleted: IED
251	attributed to increased expression and/or pretranslational alterations (involving mRNA	Deleted: HFD
252	stability or gene transcription). Number of factors can contribute to increased adipose	
253	Shp2 expression including, but not limited to, insulin resistance. Of note, Shp2 (and	
254	PTP1B) expression and activity are increased in liver and muscle of diabetic rats [35]. In	
255	addition, improved insulin sensitivity in obese subjects following weight loss is	

accompanied by decreased PTP1B expression and activity in adipose tissue [36]. Thus, 256

adipocyte-specific Cre oq-Cre genic

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257	in subjects with insulin resistance, reduction of the elevated PTP activity (in one or more
258	tissue) could potentially reduce the risk of developing diabetes and may have beneficial
259	metabolic effects. Additional factors can contribute to increased PTP expression in vivo.
260	Zabolotny et al. report that inflammation underlies PTP1B over-expression in diabetes
261	and obesity [37], Given that Shp2 plays a role in TNF receptor and interleukin 6
262	signaling [38, 39], inflammatory responses might contribute, at least in part, to the
263	regulation of adipose Shp2 expression. Preliminary studies revealed increased
264	inflammatory response in adipose tissue of FSHKO mice on a HFD compared with
265	controls (Bettaieb and Haj, unpublished observations). Since adipose inflammation
266	accompanies obesity in humans [33], additional studies are warranted to address the
267	potential role of adipose Shp2 in inflammation and the metabolic implications of such
268	regulation.
269	Body weights of control and FSHKO mice on regular chow and HFD were
270	comparable indicating that adipose Shp2 deletion did not significantly alter body mass
271	under these experimental conditions. In line with this, FSHKO and control mice
272	exhibited comparable leptin concentrations. <u>Our findings and previous reports on Shp2</u>
273	deletion in muscle [11, 12], suggest that Shp2 deletion in these peripheral insulin-
274	responsive tissues does not significantly alter body mass. On the other hand, neuronal
275	Shp2 has been identified as a regulator of energy balance and adiposity in vivo, at least in
276	part, through modulating leptin signaling [20, 21]. Additional studies are required to
277	fully assess the effects of adipose Shp2 deficiency on energy balance and adipokine
278	secretion under different experimental conditions such as various diets and/or prolonged
279	high fat feeding.

Deleted: At any rate, adipose Shp2 deficiency does not significantly alter insulin sensitivity suggesting that Shp2 over-expression is not regulated simply by obesity or diabetes *per se*.

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280	Shp2 plays diverse roles in peripheral tissues to modulate systemic insulin
281	sensitivity and glucose homeostasis. Transgenic mice that express a presumptive
282	dominant negative mutant of Shp2 to varying levels in liver, skeletal muscle and adipose
283	tissue exhibit insulin resistance, impaired insulin-stimulated glucose uptake and
284	decreased IRS1 phosphorylation in skeletal muscle and liver [10]. In addition, Shp2
285	deletion in muscle leads to insulin resistance and glucose intolerance [11]. The dilated
286	cardiomyopathy that also develops in these mice [11, 12] could lead to secondary
287	changes in muscle cells that affect insulin sensitivity. However, the similarity between
288	the muscle-specific KO and transgenic mutant mice suggest that Shp2 is a positive
289	modulator of insulin signaling in muscle. On the other hand, we recently reported that
290	mice lacking Shp2 in the liver exhibit increased hepatic insulin action and enhanced
291	systemic insulin sensitivity, indicating that Shp2 is a negative regulator of insulin
292	signaling in the liver. Our current findings indicate that adipose-specific Shp2 deletion
293	does not significantly alter systemic insulin sensitivity and glucose homeostasis.
294	Consistent with their insulin sensitivity, FSHKO and control mice exhibited comparable
295	Akt and Erk phosphorylation. However, we cannot exclude alterations in Akt and/or Erk
296	signaling at later times post stimulation. The mechanism(s) through which Shp2 plays
297	distinct roles in individual peripheral insulin-responsive tissues remains unclear.
298	Conceivably, Shp2 has distinct substrates in different tissues. Alternatively, Shp2 may
299	affect the same pathways in different tissues, but the effects of those pathways and/or the
300	feedback regulatory pathways may differ in a tissue-specific manner. At any rate, these
301	studies highlight the need to dissect the tissue-specific roles of Shp2.

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302	In summary, our studies indicated that adipose Shp2 deletion did not significantly
303	alter body mass and systemic glucose homeostasis. It would be of interest to examine if
304	enhanced/prolonged metabolic challenge(s) will lead to the manifestation of metabolic
305	alterations in FSHKO mice. Finally, we do not exclude other biologically relevant
306	effects of adipose Shp2 deletion such as nonshivering thermogenesis, endoplasmic
307	reticulum stress, adipokine secretion and inflammation. Indeed, Shp2 plays a role in TNF
308	receptor and interleukin 6 signaling, thus it will be important to explore these and other
309	signaling systems, in FSHKO mice.
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313	(BIDMC/Harvard University) for Shp2 ^{fl/fl} and Adipoq-Cre mice, respectively. We thank
314	members of the laboratory for critical comments on the manuscript.
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322	DISCLOSURE STATEMENT
323	The authors declare no conflict of interest

325 AUTHOR CONTRIBUTIONS

- A. B., performed research, collected and analyzed data and co-wrote manuscript, K. M.,
- 327 performed research, I.M., performed research, N.N., performed research, S.C., performed
- 328 research, S.L., performed research, F.G.H., designed study, data interpretation and
- 329 manuscript writing.

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Genotype	flx/flx	FSHKO	Cre
Metabolic parameters			
Glucose (mg/dl) Fed	178 ± 14	197 ± 19	175 ± 13
Fasted	128 ± 10	139 ± 5	129 ± 9
Insulin (ng/ml) Fed	10.5 ± 1.6	11.2 ± 1.2	10.7 ± 1.7
Fasted	3.2 ± 0.6	2.6 ± 0.5	2.9 ± 0.5
Insulin/Glucose ratio Fed	0.05 ± 0.01	0.06 ± 0.02	0.06 ± 0.01
Fasted	0.025 ± 0.01	0.018 ± 0.01	0.022 ± 0.02
Leptin (ng/ml) Fasted	8.16 ± 0.62	8.96 ± 1.90	9.71 ± 1.10
TG (mg/dl) Fasted	15.9 ± 1.6	13.4 ± 0.8	18.3 ± 4.6 *
FFA (mM) Fasted	1.86 ± 0.3	1.41 ± 0.2	1.45 ± 0.2

- 423 **Table 1: Metabolic variables in mice with adipose-specific Shp2 deletion**. Male
- 424 flx/flx, FSHKO and Cre mice were fed a HFD upon weaning. Serum was collected from
- 425 fed or fasted mice at 10 weeks of age (6 weeks on HFD) and the indicated metabolic
- 426 parameters were measured. Values are expressed as the mean \pm SEM of measurements
- 427 obtained for 6-8 animals per genotype. * indicates statistically significant difference
- 428 between Cre and FSHKO.

429 FIGURE LEGENDS

- 430
- 431 Figure 1: Adipose-specific Shp2 deletion. (A) Immunoblots of Shp2 expression in
- 432 lysates of white adipose tissue (W) and purified adipocytes from collagenase-treated
- 433 white adipose tissue (CW) from Adipoq-Cre (Cre), Shp2^{flx/flx} (fl/fl) and Adipoq-Shp2^{flx/flx}
- 434 (FSHKO) mice on a HFD for 12 weeks. <u>Blots were probed with anti-</u>Tubulin <u>antibodies</u>
- 435 (bottom panel) as a loading control, <u>Numbers reflect samples from different mice (W1</u>

436 and CW1 are from the same mouse; W2 and CW2 are from a different mouse). (B)

- 437 Quantitative determination of Shp2 protein expression (normalized to Tubulin) from six
- 438 mice per genotype. <u>Note that compared with control mice, adipose Shp2 protein</u>
- 439 expression was decreased by ~70% and 85% in W and CW, respectively. (C) Shp2
- 440 protein expression in lysates from white adipose tissue, collagenase-treated white adipose
- 441 tissue, brown adipose tissue (BAT), liver (L), muscle (M), pancreas (P) and brain (Br).
- 442 Blots were probed for PTP1B, TCPTP and Tubulin. (D) Shp2 expression in W, CW and
- 443 bone marrow-derived macrophages (Mac) from mice on regular chow for 60 weeks. **
- 444 Indicates statistically significant difference ($P \le 0.01$) between FSHKO and fl/fl<u>mice</u>.
- 445

446 Figure 2: Effects of adipose-specific Shp2 deletion on body weight and adiposity.

- 447 (A) Shp2 protein expression in different adipose depots (SubQ: subcutaneous; Epi:
- 448 epididymal; Ret: retroperitoneal; Vis: visceral; BAT: brown adipose tissue) of wild type
- 449 male mice on regular chow or a HFD (for 12 weeks). Each lane represents sample from a
- 450 different mouse. (B) Quantitative determination of Shp2 protein expression, normalized
- to Tubulin, from 3 mice per genotype. Body weight of male (C, E) and female (D, F)

Deleted: Shp2 deletion efficiency was evaluated in

1	Deleted:	Lysates were blotted			
-{	Deleted: for				
1	Deleted:				

452 Cre (n= 9), flx/flx (n= 9), and FSHKO (n= 9) mice on a HFD (\mathbf{C} , \mathbf{D}) and chow (\mathbf{E} , \mathbf{F}).

453 Total white adipose tissue weight (G, J), adiposity index (H, K), and head-rump length

454 (cm) (I, L) of male (G-I) and female (J-L) mice on HFD for 12 weeks. Adiposity index

455 (H) in FSHKO and fl/fl mice (H) has a P value of 0.052. * Indicates statistically

456 significant difference in Shp2 expression between chow and HFD fed mice (*, $P \le 0.05$;

457 **, $P \le 0.01$).

458

Figure 3: Insulin sensitivity and glucose tolerance in mice with adipose-specific Shp2 459 deletion. (A-D) Insulin tolerance tests (ITTs) in male (A, C) and female (B, D) Cre (n= 460 9), flx/flx (n= 9), and FSHKO (n= 9) mice on a HFD (A, B) and chow (C, D) at 14 weeks 461 of age (insulin 1 mU/g B.W.). (E-H) Glucose tolerance tests (GTTs) in male (E, G) and 462 females (F, H) Cre (n= 9), flx/flx (n= 9), and FSHKO (n= 9) mice on a HFD (E, F) and 463 chow (G, H) at 15 weeks of age (glucose dose, 2 mg/g B.W.). (I, J) Male mice (30 464 weeks old) were injected intraperitoneally with saline or insulin (10 mU/g B.W.) and 465 466 sacrificed after 10 minutes. Total adipose tissue lysates were immunoblotted for pAkt (S473) (I) and pErk (J) and the corresponding total proteins. Bar graph indicates 467 quantitation of Akt and Erk phosphorylation (adjusted to protein level) from at least 4 468 mice per group. All blots were scanned and quantified using FluorChem 8900 and 469 470 statistical analysis was performed using two-tailed Student's t-test. * Indicates statistically significant difference between basal and insulin stimulated conditions for 471 each genotype (*, $P \le 0.05$; **, $P \le 0.01$). 472