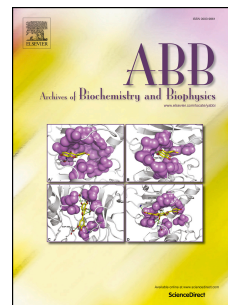


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A single night of sleep curtailment increases plasma acylcarnitines: novel insights in the relationship between sleep and insulin resistance

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1 **A single night of sleep curtailment increases plasma acylcarnitines: novel insights in the**
2 **relationship between sleep and insulin resistance**

3

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22

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25 acylcarnitines

26

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27 **Abstract**

28 We have previously shown that acute sleep curtailment induces insulin resistance, both in
29 healthy individuals as well as in patients with type 1 diabetes, suggesting a causal role for
30 sleep disturbances in pathogenesis of insulin resistance, independent of endogenous insulin
31 production. However, the underlying mechanisms remain unclear. This study aimed to
32 explore the metabolic pathways affected by sleep loss using targeted metabolomics in
33 human fasting plasma samples. Healthy individuals (n = 9) and patients with type 1 diabetes
34 (n = 7) were studied after a single night of short sleep (4 hours) versus normal sleep (8
35 hours) in a cross-over design. Strikingly, one night of short sleep specifically increased the
36 plasma levels of acylcarnitines, essential intermediates in mitochondrial fatty acid oxidation
37 (FAO). Specifically, short sleep increased plasma levels of tetradecenoyl-L-carnitine (C14:1)
38 (+32%, $p=2.67*10^{-4}$), octadecanoyl-L-carnitine (C18:1) (+22%, $p=1.92*10^{-4}$) and
39 octadecadienyl-L-carnitine (C18:2) (+27%, $p=1.32*10^{-4}$). Since increased plasma acylcarnitine
40 levels could be a sign of disturbed FAO, it is possible that sleep curtailment acutely induces
41 inefficient mitochondrial function. Our observations provide a basis for further research into
42 the role of acylcarnitines as a potential mechanistic pathway by which sleep deprivation –
43 even short term – causes adverse metabolic effects, such as insulin resistance.

44

45 Introduction

46 Diabetes mellitus (DM) is characterized by either an absolute (type 1; DM1) or relative (type
47 2; DM2) deficiency of insulin. Both DM1 and DM2 are associated by increased morbidity and
48 increased cardiovascular risk (1;2). Peripheral insulin resistance precedes the development
49 of DM2 and recently it has been recognized that a certain degree of insulin resistance is also
50 present in DM1 (3). Therefore, uncovering modifiable risk factors in an early stage of insulin
51 resistance development is of crucial importance to reduce the number of patients with DM2
52 and improve glycemic control in DM1. Interestingly, the DM2 epidemic coincides with a
53 reduction in the average sleep duration, which has gradually declined with ~1.5 hours per
54 night (4) over the past decades. In fact, large epidemiological cohorts have documented an
55 association between sleep duration and increased insulin resistance (5). Furthermore, short
56 sleep has been associated with poor glycemic control in DM1 (6). Both short and long
57 duration of sleep are associated with an increased risk for insulin resistance, implying that
58 there might be an optimal sleep duration of approximately 8 hours (7-10). Several human
59 intervention studies showed that decreased sleep duration causes insulin resistance.
60 Repeated sleep curtailment during more than 6 nights increased insulin resistance in
61 healthy individuals (11-13). Moreover, we previously published that even one single night
62 with partial sleep loss, i.e. 4 hours sleep allowed, a condition representative for incidental
63 daily life sleep habits, is sufficient to induce peripheral insulin resistance in both healthy
64 young individuals (14) as well as patients with DM1 (15).

65 The mechanism by which acute sleep curtailment induces insulin resistance has not
66 been fully elucidated. Plasma metabolomics is considered a valuable approach to assess
67 underlying biological processes, complementary to genomics and transcriptomics. Strikingly,

68 metabolite levels reflect biological activity of the encoded proteins and are thus closer to
69 the clinical endpoints (16). Indeed, metabolomics has previously been demonstrated to be a
70 powerful tool in investigating insulin resistance and DM2 (17). Thus far, the effects of sleep
71 loss on the human metabolome are poorly characterized. Prolonged sleep deprivation
72 during 5 days has been shown to induce metabolite changes in lipid, carbohydrate, amino
73 acid and protein pathways (18;19). In contrast, Davies et al. (20) subjected healthy
74 individuals to complete sleep restriction of 24 hours. This extreme sleep deprivation
75 resulted in increased plasma levels of glycerophospholipids, acylcarnitines, sphingolipids
76 and amino acids. However, the sleep intervention and control sleep occurred on
77 consecutive days in all individuals. Differences between metabolite levels were also
78 observed between the wake periods, suggesting that the study conditions were not fully
79 comparable. In addition, none of these previous studies included measurements of insulin
80 resistance. Therefore, the aim of the present study was to use metabolomics to explore
81 pathways involved in the relationship between sleep and insulin resistance in a cohort with
82 proven insulin resistance upon short sleep duration (14;15). To this end, we examined 163
83 metabolites in 16 individuals (healthy individuals and individuals with DM1) subjected to a
84 night of normal sleep duration (8 hours) and one night of short sleep duration (4 hours).
85 Here, we report that one night of sleep curtailment specifically increases the metabolic class
86 of acylcarnitines in plasma, suggesting that increased acylcarnitines are associated with the
87 observed relationship between sleep curtailment and induction of insulin resistance.

88

89 **RESEARCH DESIGN AND METHODS**

90 **Protocol**

91 Two studies were previously performed, to study the effect of one night of short sleep
92 duration (4 hours) compared to normal sleep duration (8 hours) on peripheral insulin
93 resistance (14;15). The studies applied the same study design in two different populations,
94 namely healthy individuals and patients (14) with type 1 diabetes (DM1)(15). Healthy
95 individuals were studied to determine the effects of a single night of short sleep duration on
96 insulin resistance. The second study assessed the effects of short sleep duration on insulin
97 resistance in DM1 patients on stable insulin pump therapy. DM1 patients do not have
98 endogenous insulin production and therefore cannot compensate for fluctuations in insulin
99 resistance. We hypothesized that variations in sleep duration could contribute the intra-
100 individual variations in glucoregulation. In both healthy individuals and individuals with
101 DM1, decreased sleep duration induced insulin resistance. Therefore, we reasoned that a
102 single night of short sleep duration may increase peripheral insulin resistance via a common
103 metabolic pathway. To investigate which pathways could be involved, we analyzed
104 metabolites from both studies and pooled the data.

105

106 **Subjects**

107 The study was approved by the medical ethical committee of the Leiden University Medical
108 Center and all subjects gave written informed consent. We recruited a total of 18
109 individuals. Briefly, nine healthy individuals were recruited by advertisement and nine
110 individuals with DM1 with stable continuous subcutaneous insulin pump therapy were
111 included from our outpatient clinic. Exclusion criteria for all individuals were BMI>26 kg/m²,
112 history of sleep disorders, psychiatric disorders and use of sleep medication, β -blocking
113 drugs and prokinetic drugs. All individuals had a stable weight in the past 3 months and had

114 regular and non-extreme sleeping habits. Habitual sleep duration was assessed by 7 days of
115 actigraphy (Actiwatch AW7; Cambridge Neurotechnology, Cambridge, UK) prior to both
116 study days and sleep questionnaires (Epworth Sleepiness Scale, Pittsburg Sleep Quality
117 Index and Berlin Questionnaire). Subjects were instructed to maintain a regular dietary,
118 activity and sleep regiments 3 days prior to both study days, fitting their habits, which they
119 recorded in a diary. DM1 patients were instructed to keep a stable insulin pump setting. Of
120 the 18 recruited individuals, 2 individuals with DM1 were excluded from all analyses, one
121 due to previously undiagnosed sleep apnea and one due to nocturnal hypoglycemia.

122

123 **Experimental design**

124 Subjects were subjected to in-hospital sleep registration for 3 days, of which study day 1
125 was for basal measurements and habituation to hospital conditions. Sleep duration and
126 quality (of parameters) was assessed by polysomnography as described previously (14;15).
127 All subjects underwent both a normal sleep night of at least 8 hours and one night of 4
128 hours sleep, the order of which was determined by balanced assignment, in a cross-over
129 design with at least 3 weeks interval between measurements. In both sleep conditions,
130 subjects spent 8.5 hours (from 23:00 to 7:30) in bed and were fasting from 22:00 onwards.
131 During sleep curtailment, subjects were allowed to sleep from 01:00 to 05:00, the remaining
132 time they were allowed to read or watch movies in upward position in dim light. Their
133 wakefulness was monitored. After the night of normal or short sleep, a fasting plasma
134 sample was obtained at 8:30 am, after which a hyperinsulinemic euglycemic clamp was
135 performed as described in detail previously (14;15) to establish peripheral insulin sensitivity,
136 endogenous glucose production and hepatic insulin sensitivity. Briefly, a primed (17.6

137 $\mu\text{mol}\cdot\text{kg}^{-1}$) continuous ($0.22 \mu\text{mol}\cdot\text{kg}^{-1}\cdot\text{min}^{-1}$) infusion of $[6,6\text{-}^2\text{H}_2]$ glucose (Cambridge
138 Isotope laboratory, Andover, MA) was administered via a catheter. Infusion of insulin
139 (Actrapid, Novo Nordisk, Alphen a/d Rijn) occurred simultaneously according to DeFronzo
140 (21). Blood samples were obtained every 5 minutes from the contralateral arm for glucose
141 measurements to adjust variable infusion of 20% glucose with 3% $[6,6\text{-}^2\text{H}_2]$ glucose to
142 maintain euglycemia (i.e. 5.0 mmol/l), which was started 4 min after start of insulin infusion.
143 Free fatty acids were determined in basal fasting plasma samples as by enzymatic
144 colorimetric assay (14;15).

145

146 **Metabolomics**

147 Metabolomics analysis was performed on fasting plasma samples in all individuals using the
148 Biocrates Absolute/*IDQ*TM p150 kit (Biocrates, Life Science AG, Innsbruck, Austria) in the
149 Genome Analysis Center at the Helmholtz Zentrum, Munich, Germany. The assay
150 procedures of the Absolute/*IDQ*TM p150 kit as well as the metabolite nomenclature have
151 been described in detail previously (22;23). Briefly, 10 μL of each plasma sample was
152 pipetted into a 96 well sandwich plate containing an inserted filter with previously applied
153 stable isotope labeled internal standards. The filters in the wells were dried using a stream
154 of nitrogen. Amino acids were derivatized with 5% phenylisothiocyanate reagent (PITC) and
155 the filters were dried again. Metabolites as well as internal standards were extracted with 5
156 mM ammonium acetate in methanol and the solutions were centrifuged through the filter
157 membrane into the lower deep well plate. The extracts were diluted with MS running
158 solvent and analyzed. Flow injection analysis (FIA) tandem mass spectrometry (MS/MS)
159 method was used to quantify 163 metabolites, including free carnitine, 40 acylcarnitines, 14

160 amino acids (13 proteinogenic + ornithine), hexoses (sum of hexoses), 92
161 glycerophospholipids (15 lysophosphatidylcholines (lysoPC) and 77 phosphatidylcholines
162 (PC), and 15 sphingolipids. Internal standards served as reference for the calculation of
163 metabolite concentrations (μM). The complete list of analyzed metabolites grouped by
164 metabolite class is presented in supplementary material (Table S3).

165

166 **Statistical analysis**

167 For all metabolites, differences between short and normal sleep were calculated by
168 subtracting plasma levels obtained after short sleep from those obtained after normal sleep.
169 Paired Students T-tests for were performed comparing normal and short sleep (SPSS
170 statistical package edition 20) with Bonferroni post-hoc correction for multiple testing. $P <$
171 3.07×10^{-4} ($=0.05/163$; after correction) was considered statistically significant. Calculations
172 for hyperinsulinemic euglycemic clamp analysis were described previously (14;15). Since we
173 aimed to investigate the effect of short sleep on metabolite levels, individuals of both
174 groups (healthy individuals and individuals with DM1) were pooled to determine effects of
175 sleep duration. Two way repeated measure ANOVA was performed to analyze interaction
176 effects of subgroup (healthy vs. DM1) with sleep duration. Data are presented as means \pm
177 SD. Since baseline characteristics and insulin sensitivity data were published for healthy
178 individuals and individuals with DM1 separately, in this paper these data are shown for the
179 two groups together. To allow comparison between subgroups, the baseline characteristics,
180 sleep indices and insulin sensitivity data are included in the supplemental tables and were
181 compared using Student's t-test.

182

183 RESULTS**184 Basal clinical characteristics**

185 Metabolites were measured in sixteen individuals after a night of short sleep (4 hours)
186 versus after a night of normal sleep (8 hours) duration. Subjects had a mean age of 44 ± 14
187 years and included 8 women. Individuals were lean, with an average BMI of $23.7 \pm 2.2 \text{ kg/m}^2$
188 and a waist hip ratio of 0.85 ± 0.08 (Table 1). The study population consisted of nine healthy
189 individuals (56%) and seven individuals with type 1 diabetes mellitus (DM1) (44%). Sleep
190 duration prior to the study days did not differ healthy individuals (mean recorded sleep
191 duration prior to study day 1 and 2: $420 \pm 20 \text{ min}$ vs. $476 \pm 11 \text{ min}$; $p=0.19$) nor in individuals
192 with DM1 (mean recorded sleep duration prior to study day 1 and 2: $475 \pm 8 \text{ min}$ vs. 490 ± 7
193 min; $p=0.12$). Results of healthy individuals and individuals with DM1 were reported
194 previously separately (14;15). Age, sex distribution, BMI and waist-hip ratio were
195 comparable between these two subgroups (Table S1).

196

197 Short sleep increases insulin resistance

198 Short sleep intervention was effective in reducing total sleep time (TST) by -51% (461 ± 25 vs
199 $225 \pm 26 \text{ min}$, $p < 0.001$). The reduction of sleep duration was due to decreased sleep
200 duration of both non-REM (stage 2 and stage 3) and REM sleep (Table 2). Fasting plasma
201 free fatty acids did not differ between sleep conditions (Table 2) or between subgroups
202 (Table S2). Next, the effect of short sleep on insulin resistance was investigated by
203 hyperinsulinemic euglycemic clamp studies. Interestingly, a single night of short sleep
204 increased peripheral insulin resistance, as indicated by a decreased glucose disposal rate
205 (GDR) (34.1 ± 13.8 vs $27.9 \pm 9.8 \mu\text{mol} \cdot \text{kg LBM}^{-1} \cdot \text{min}^{-1}$, $p = 0.001$) and decreased glucose

206 infusion rate (GIR) (29.0 ± 14.7 vs $22.1 \pm 10.7 \mu\text{mol}\cdot\text{kg LBM}^{-1}\cdot\text{min}^{-1}$, $p=0.001$). Short sleep
207 tended to increase endogenous glucose production (EGP) by the liver in all subjects ($4.7 \pm$
208 1.9 vs $5.5 \pm 1.6 \mu\text{mol}\cdot\text{kg LBM}^{-1}\cdot\text{min}^{-1}$, $p=0.08$; Table 2). This was mainly due to increased
209 endogenous glucose production in the subset of healthy individuals (Table S2; previously
210 published in (14)). Expectedly, individuals with DM1 displayed higher baseline insulin
211 resistance than in healthy individuals (3) (EGP 6.2 ± 1.9 vs. 3.6 ± 0.6 , $p=0.003$; GDR 25.5 ± 6.4
212 vs. 40.7 ± 14.3 , $p=0.028$; GIR 19.0 ± 7.0 vs. 36.9 ± 14.4 , $p=0.014$, Table S2). Moreover, short
213 sleep increased peripheral insulin resistance irrespective of this difference in baseline insulin
214 sensitivity, suggesting a that short sleep may induce insulin resistance in healthy individuals
215 and individuals with DM1 via a common pathway. Therefore, the effect of short sleep was
216 investigated for healthy individuals and individuals with DM1 together.

217

218 **Short sleep specifically increases plasma acylcarnitines**

219 To investigate possible pathways which could be involved in the increased of insulin
220 resistance by short sleep duration, we performed metabolomics analysis on fasting morning
221 plasma samples. A total of 163 metabolites representing 5 different classes were measured
222 (Table S3). Short sleep increased thirteen metabolites ($p<0.05$) (Table 3). Strikingly, all of
223 these are acylcarnitines. After stringent post-hoc correction, short sleep significantly
224 increased plasma levels of tetradecenoyl-L-carnitine (C14:1) by +32% (plasma level
225 difference: $+0.017 \mu\text{M}$, $p=2.67\cdot 10^{-4}$), octadecenoyl-L-carnitine (C18:1) by +22% (plasma level
226 difference: $+0.015 \mu\text{M}$, $p=1.92\cdot 10^{-4}$) and octadecadienyl-L-carnitine (C18:2) by +27%
227 (plasma level difference: $+0.005 \mu\text{M}$, $p=1.32\cdot 10^{-4}$). Short sleep duration increased
228 acylcarnitines in both subgroups, indicating that the effect of short sleep on acylcarnitines

229 was not dependent on having DM1 or being healthy. There was no interaction effect of the
230 subgroup (healthy vs. DM1) with the sleep duration (short vs. normal) for the 13 increased
231 acylcarnitines. Baseline acylcarnitine levels (i.e. after normal sleep) did not differ between
232 healthy individuals and DM1, except for a higher level of C:12-DC in DM1 (0.087 ± 0.005 vs
233 $0.101 \pm 0.005 \mu\text{M}$, $p < 0.0001$) (Table S5). Acylcarnitines levels did not differ between healthy
234 individuals and DM1 after short sleep (Table S6). We therefore conclude that a single night
235 of short sleep specifically increased plasma acylcarnitines.

236

237 **DISCUSSION**

238 The present study aimed to explore the metabolic pathways affected by sleep curtailment
239 using targeted plasma metabolomics in individuals (healthy individuals and individuals with
240 type 1 diabetes (DM1)) subjected to both short sleep (4 hours) and normal sleep (8 hours).
241 As part of the same study, we previously reported that this short sleep intervention
242 increased peripheral insulin resistance in both study groups as determined by
243 hyperinsulinemic euglycemic clamp analysis (14;15). We now show that one night of short
244 sleep specifically increases plasma levels of acylcarnitines, in both healthy individuals and
245 DM1 patients.

246 Our study is the first to show that short sleep duration increased plasma
247 acylcarnitines in concert with increased insulin resistance in both healthy individuals and
248 individuals with DM1. This indicates that short sleep duration affects metabolism
249 irrespective of pre-existing insulin producing capacity. The relationship between increased
250 plasma acylcarnitine levels and increased insulin resistance is supported by association
251 studies. Human studies showed increased plasma levels of acylcarnitines in individuals with

252 impaired fasting glucose and with type 2 diabetes (DM2), compared to healthy controls
253 (24;25). The significance of this association is still a matter of debate, since human
254 intervention studies are lacking (26).

255 It is interesting to speculate about the biological relevance of increased plasma levels
256 of acylcarnitines. Acylcarnitines are vital to energy homeostasis. They are esters of fatty
257 acids and carnitine, which are transported over the outer and inner mitochondrial
258 membranes by carnitine palmitoyl transferases (CPTs). Thus, acylcarnitines are essential to
259 shuttle fatty acids from the cytoplasm into mitochondria where they can be oxidized and
260 enter the tricarboxylic acid (TCA) cycle to generate ATP. An excess of acylcarnitines is
261 generally viewed as a result from a mismatch between TCA flux and fatty acid oxidation
262 (FAO) (27). Previously reported causes of this mismatch include prolonged fasting and
263 excessive muscle activity (28-30). The present study, in which subjects participated in a
264 protocol that controlled for food intake and physical activity, adds sleep deprivation as a
265 provoking event. A mismatch between FAO and TCA flux has been related to mitochondrial
266 dysfunction. Patients with inborn errors of FAO have increased plasma levels of especially
267 long chain acylcarnitines (31). Interestingly, altered mitochondrial parameters have been
268 frequently linked to insulin resistance in the context of both DM1 and DM2 (32-38).
269 Moreover, mitochondrial dysfunction in mice induces skeletal muscle insulin resistance (27)
270 while TCA-FAO mismatch predisposes mice to diet-induced obesity and insulin resistance
271 (39). It is therefore tempting to speculate that in our model of insulin resistance due to
272 short sleep deprivation, the increased plasma acylcarnitine levels are a sign of inefficient
273 mitochondrial function.

274 The tissue distribution of acylcarnitines coincides with important targets of insulin,
275 i.e. muscle and liver. The majority of the body's L-carnitine is stored in muscle (~ 97% of the
276 body's L-carnitine), followed by liver which contains 1% of the total L-carnitine pool (40).
277 Acylcarnitine results from the acylation of L-carnitine, and is therefore dependent on the
278 fatty acid pool of the tissue. Interestingly, animal studies demonstrate the distribution of
279 acylcarnitines is different between metabolic organs. In mice, the muscle tissue contains
280 relatively more long-chain acylcarnitines, including C14:1 and C18:1, while liver is richer in
281 free carnitines and short-chain carnitines (41). Collectively, these data suggest that plasma
282 short-chain acyl- and free carnitines are mainly derived from the liver, as indeed
283 demonstrated in pigs (42), while plasma long-chain acylcarnitines in plasma presumably
284 originate from muscle tissue. These data thus suggest that the increase in long-chain
285 acylcarnitine that we observe after a single night of short sleep is likely derived from muscle.

286 Mechanistically, increased acylcarnitine levels after short sleep duration could be a
287 marker of altered metabolic processes: increased fatty acid oxidation (FAO), inefficient
288 mitochondrial function or a disturbed metabolism of the branched-chain amino acids
289 (BCAA) valine, isoleucine or leucine. Although disturbed BCAA metabolism has been
290 associated with insulin resistance in humans (43), our data do not support a role of BCAA
291 metabolism as short sleep duration did not increase BCAA plasma levels or short-chain
292 acylcarnitines. Increased acylcarnitine levels due to increased FAO can be caused by either
293 increased energy demand and/or prolonged fasting. In the present study, the length of
294 fasting was equal; however energy expenditure was not measured. Therefore, we cannot
295 exclude that the increased acylcarnitines after short sleep are due to increased FAO. Sleep is
296 accompanied by lower resting energy expenditure than wakefulness (44) and therefore

297 short sleep duration may increase energy demand. In fact, complete (24 h) sleep deprivation
298 increases energy demand by 7% (45). However, the effects of short sleep duration on
299 energy expenditure are inconclusive (46). A recent study shows that short sleep intervention
300 for five consecutive days increased long-chain plasma acylcarnitines (19). Interestingly, after
301 one night of recovery sleep, plasma acylcarnitines did not normalize. Likely, the increased
302 acylcarnitines were not due to differences in overnight energy expenditure. Besides being a
303 marker of insulin resistance and/or mitochondrial processes, acylcarnitines could also play a
304 causal role in development of insulin resistance. *In vitro* studies have shown that
305 acylcarnitines have bioactive properties and indeed have pro-inflammatory effects (47;48).
306 Of note, treatment of both rodent and human myotubes with acylcarnitines in a
307 physiological concentration caused decreased insulin signaling and glucose uptake in
308 response to insulin (49). Although this finding needs to be confirmed *in vivo*, it provides a
309 putative causal link between acylcarnitines and insulin resistance.

310 Taken all these data together, it is interesting to speculate on a mechanistic model
311 for the relationship between sleep curtailment and insulin resistance. Upon sleep
312 curtailment, the energy demands of peripheral tissues increases at a time conflicting with
313 the physiological circadian rhythm. The energy homeostasis is adapted to anticipate the
314 changing energy need and availability throughout the day. Indeed, muscle tissue is also
315 under circadian control (50). These clock genes are also important in driving rhythmicity in
316 energy producing capacity of the mitochondria, as evidenced by mice studies (51). We
317 hypothesize that the mismatch in energy producing capacity and demand could be the
318 cause of incomplete FAO, leading to accumulation of intermediates of FAO. Acylcarnitine

319 levels increase, which may increase insulin resistance either through direct interaction with
320 insulin signaling or through increased inflammatory pathways.

321 Our findings are supported by three studies which have investigated the effects of
322 sleep on the human metabolome. Davies et al. (20) subjected 12 healthy individuals to an
323 extreme sleep deprivation of 24 hours and reported nine increased short and medium-chain
324 acylcarnitines, including tetradecenoyl-L-carnitine. Bell et al. (18) reported a trend towards
325 increased acylcarnitines after prolonged mild sleep curtailment of 8 consecutive nights of
326 5.5 hours sleep in 11 young individuals with family history of DM2. Weljie et al (19) also
327 reported increased C18:1, C10:0 and C12:0 acylcarnitines upon five consecutive nights of 4
328 hours sleep. Strikingly, despite the difference in study populations and sleep curtailment
329 protocols of the present and previous studies used, the acylcarnitines invariably increase
330 after sleep curtailment.

331 In conclusion, the present study shows that a single night of 4 hours short sleep,
332 which induces insulin resistance (14;15), also increases plasma levels of acylcarnitines, in
333 particular tetradecenoyl-L-carnitine, octadecenoyl-L-carnitine and octadecadienyl-L-
334 carnitine. We propose that sleep curtailment impairs mitochondrial function, which
335 coincides with insulin resistance. Our findings provide a basis for mechanistic studies to
336 further elucidate the role of acylcarnitines in the complex relationship between short sleep
337 and increased insulin resistance.

338

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343 D.O.M.K. and K.W.v.D. analyzed data and supervised metabolome analysis, E.D., M.v.D., and
344 J.G.v.D. performed the initial study, G-J.L. and K.W.v.K. performed polysomnography, C.P.
345 and J.A. performed metabolome analysis and quality assurance, and E.P.M.C, J.A.R, P.C.N.R.
346 and N.R.B designed and supervised the study. All authors have approved final version of the
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513 **TABLES**514 **Table 1: Study population characteristics¹.**

	Subjects (n = 16)
Females (%)	8 (50%)
Age (years)	44 ± 14
BMI (kg/m ²)	23.7 ± 2.2
WHR	0.85 ± 0.08

515 BMI = body mass index. WHR = waist hip ratio. Data is presented as mean (SD or percentage).

516 ¹ Data are pooled from two previously published studies (14;15).

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519 **Table 2: Effects of short sleep on sleep parameters and insulin sensitivity¹.**

	Subjects (n = 16)		p
	Normal sleep	Short sleep	
Sleep parameters			
TST (min)	461 ± 25	225 ± 24	<0.001
Stage 1 (% of TST)	10 ± 3	10 ± 6	0.798
Stage 2 (% of TST)	43 ± 7	37 ± 9	0.002
Stage 3 (% of TST) (SWS)	24 ± 7	34 ± 10	<0.001
REM sleep (% of TST)	23 ± 4	18 ± 8	0.025
Sleep efficiency (%)	93 ± 4	91 ± 7	0.418
Plasma parameter			
Free fatty acids (mmol/l)	0.65 ± 0.24	0.61 ± 0.19	0.24
Insulin sensitivity parameters			
EGP ($\mu\text{mol} \cdot \text{kg LBM}^{-1} \cdot \text{min}^{-1}$)	4.7 ± 1.9	5.5 ± 1.7	0.087
GDR ($\mu\text{mol} \cdot \text{kg LBM}^{-1} \cdot \text{min}^{-1}$)	34.1 ± 13.8	27.9 ± 9.8	0.001
GIR ($\mu\text{mol} \cdot \text{kg LBM}^{-1} \cdot \text{min}^{-1}$)	29.0 ± 14.7	22.1 ± 10.7	0.001

520 Insulin sensitivity parameters were determined by hyperinsulinemic euglycemic clamp.

521 EGP = endogenous glucose production, GDR = glucose disposal rate (glucose Rd), GIR = glucose infusion rate.

522 LBM = lean body mass. Sleep characteristics were determined by polysomnography. TST = total sleep time.

523 SWS = slow wave sleep. Data is presented as means (SD). Effect of sleep intervention was tested with paired

524 Students T-test, significant differences shown in bold.

525 ¹ Data are pooled from two previously published studies (14;15).

Table 3: Difference between short sleep and normal sleep duration in acylcarnitine levels.

Metabolite	<i>All subjects</i>			<i>Healthy</i>			<i>DM1</i>		
	Mean Difference ¹	Change (%) ²	P-Value	Mean Difference ¹	Change (%) ²	P-Value	Mean Difference ¹	Change (%) ²	P-Value
C0	0.191	0.7%	8.81*10 ⁻¹	0.398	1.3%	8.47*10 ⁻¹	-0.075	-0.3%	9.61*10 ⁻¹
C2	0.662[§]	16.8%	2.28*10⁻²	0.184	4.4%	5.40*10⁻¹	1.276[§]	35.6%	1.33*10⁻²
C3	-0.006	-2.6%	6.76*10 ⁻¹	-0.002	-0.7%	9.33*10 ⁻¹	-0.012	-6.2%	6.03*10 ⁻¹
C3:1	-0.001	-5.9%	4.16*10 ⁻¹	0.000	-0.1%	9.93*10 ⁻¹	-0.001	-12.3%	1.58*10 ⁻¹
C3-DC (C4-OH)	0.007	11.3%	4.55*10 ⁻¹	0.006	8.0%	7.20*10 ⁻¹	0.009	17.7%	2.87*10 ⁻¹
C3-OH	0.001	3.3%	5.03*10 ⁻¹	0.002	8.5%	5.91*10 ⁻²	-0.001	-2.1%	8.24*10 ⁻¹
C4	0.011	10.3%	1.22*10 ⁻¹	0.008	7.2%	4.95*10 ⁻¹	0.015	14.4%	8.68*10 ⁻²
C4:1	0.002	6.7%	2.90*10 ⁻¹	0.002	8.7%	2.55*10 ⁻¹	0.001	4.8%	6.58*10 ⁻¹
C5	0.011	10.2%	2.80*10 ⁻¹	0.012	10.4%	4.59*10 ⁻¹	0.010	9.8%	4.30*10 ⁻¹
C5:1	0.000	1.0%	8.74*10 ⁻¹	-0.001	-2.5%	7.86*10 ⁻¹	0.001	5.0%	6.20*10 ⁻¹
C5:1-DC	0.000	2.6%	6.75*10 ⁻¹	-0.001	-4.8%	5.96*10 ⁻¹	0.002	12.5%	1.27*10 ⁻¹
C5-DC (C6-OH)	-0.001	-4.1%	4.87*10 ⁻¹	-0.001	-8.3%	2.56*10 ⁻¹	0.000	0.7%	9.45*10 ⁻¹
C5-M-DC	0.000	0.3%	9.32*10 ⁻¹	0.000	-1.1%	8.45*10 ⁻¹	0.001	1.9%	7.11*10 ⁻¹
C5-OH (C3-DC-M)	0.000	0.3%	9.40*10 ⁻¹	0.000	-1.5%	6.89*10 ⁻¹	0.001	2.4%	7.40*10 ⁻¹
C6 (C4:1-DC)	0.005	8.7%	1.22*10 ⁻¹	0.001	1.9%	7.82*10 ⁻¹	0.011	18.8%	6.20*10 ⁻²
C6:1	0.001	2.6%	3.99*10 ⁻¹	0.001	3.5%	3.35*10 ⁻¹	0.000	1.5%	8.05*10 ⁻¹
C7-DC	0.005[§]	20.0%	5.41*10⁻⁴	0.002	8.4%	8.50*10⁻²	0.008[§]	34.8%	3.21*10⁻⁴
C8	0.003	1.9%	6.06*10 ⁻¹	-0.005	-3.1%	5.56*10 ⁻¹	0.013 [§]	11.0%	4.38*10 ⁻³
C8:1	0.012	15.4%	1.62*10 ⁻¹	0.008	11.8%	3.93*10 ⁻¹	0.017	18.9%	3.09*10 ⁻¹
C9	0.000	-0.4%	9.47*10 ⁻¹	-0.001	-4.7%	5.73*10 ⁻¹	0.001	5.4%	5.51*10 ⁻¹

C10	0.014	5.8%	1.84*10 ⁻¹	0.003	1.1%	8.62*10 ⁻¹	0.027*	15.1%	4.54*10 ⁻³
C10:1	0.008	6.9%	2.68*10 ⁻¹	-0.004	-3.1%	6.68*10 ⁻¹	0.022*	23.9%	2.72*10 ⁻²
C10:2	0.000	-0.7%	8.65*10 ⁻¹	0.000	-0.1%	9.79*10 ⁻¹	0.000	-1.3%	8.51*10 ⁻¹
C12	0.012[§]	17.2%	1.70*10⁻³	0.010	12.0%	9.00*10⁻²	0.015[§]	27.5%	3.30*10⁻³
C12:1	0.017[§]	23.6%	1.74*10⁻³	0.014	17.0%	8.32*10⁻²	0.021[§]	35.2%	5.85*10⁻³
C12-DC	0.002	2.4%	1.53*10 ⁻¹	0.002	2.1%	3.78*10 ⁻¹	0.003	2.8%	3.01*10 ⁻¹
C14	0.004[§]	15.6%	1.21*10⁻²	0.002	8.5%	2.42*10⁻¹	0.006[§]	28.0%	1.97*10⁻²
C14:1	0.020[#]	32.4%	2.67*10⁻⁴	0.017[§]	23.5%	2.34*10⁻²	0.024[§]	49.1%	6.55*10⁻³
C14:1-OH	0.001	6.2%	3.79*10 ⁻¹	0.000	-1.7%	8.10*10 ⁻¹	0.001	16.9%	2.36*10 ⁻¹
C14:2	0.006[§]	26.1%	5.48*10⁻⁴	0.005	17.3%	5.01*10⁻²	0.008[§]	41.5%	4.17*10⁻³
C14:2-OH	0.000	5.0%	3.64*10 ⁻¹	0.000	6.4%	2.91*10 ⁻¹	0.000	3.3%	7.48*10 ⁻¹
C16	0.007[§]	10.8%	1.94*10⁻²	0.005	6.7%	2.05*10⁻¹	0.010	17.1%	5.91*10⁻²
C16:1	0.005[§]	8.8%	9.60*10⁻³	0.005[§]	8.3%	3.06*10⁻²	0.005	9.6%	1.55*10⁻¹
C16:1-OH	0.001[§]	20.8%	5.71*10⁻³	0.001	11.8%	1.90*10⁻¹	0.002[§]	33.8%	1.10*10⁻²
C16:2	0.001[§]	21.3%	9.72*10⁻³	0.001	9.6%	2.39*10⁻¹	0.002[§]	39.5%	1.91*10⁻²
C16:2-OH	0.000	2.1%	6.29*10 ⁻¹	0.000	3.8%	5.45*10 ⁻¹	0.000	-0.1%	9.83*10 ⁻¹
C16-OH	0.000	0.9%	8.89*10 ⁻¹	0.000	-3.4%	5.84*10 ⁻¹	0.000	6.1%	6.49*10 ⁻¹
C18	0.003	10.8%	7.34*10 ⁻²	0.002	9.6%	3.04*10 ⁻¹	0.003	12.5%	1.18*10 ⁻¹
C18:1	0.016[#]	22.3%	1.92*10⁻⁴	0.015*	20.1%	2.96*10⁻³	0.019[§]	25.1%	2.84*10⁻²
C18:1-OH	0.001	12.8%	6.02*10 ⁻²	0.001	5.7%	3.93*10 ⁻¹	0.002	22.3%	1.07*10 ⁻¹
C18:2	0.007[#]	27.0%	1.32*10⁻⁴	0.005[§]	20.7%	9.01*10⁻⁴	0.010[§]	34.4%	1.38*10⁻²

¹Difference in metabolite levels (μM) as measured by Biocrates/DQTM p150 kit between short and normal sleep duration. Positive mean difference indicates an increase after short sleep duration. Negative mean difference indicates a decrease after short sleep duration.

²Change (%) represents percentage of change in metabolite level in short compared to normal sleep (metabolite level (short sleep) – metabolite level (normal sleep)) / metabolite level (normal sleep).

DM1 = individuals with type 1 diabetes. P-values are based on paired Students t-tests. N= 16 (healthy: n=9, DM1: n=7). Full results table is shown in Supplemental Table S2. Abbreviations of acylcarnitines are shown in Supplemental Table S3.

§: Significant difference ($p < 0.05$). #: Significant difference after Bonferroni correction ($p < 3.0 \times 10^{-4}$ ($= 0.05/163$)). Significant differences metabolites in all subjects are displayed in bold.

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Table 4. Interaction effects of diabetes status and short sleep on increased acylcarnitine levels

Metabolite	Interaction P-value	Effect of sleep P-value	Effect of DM1 status P-value
C2	$3.24 * 10^{-2}$	$6.70 * 10^{-3\text{§}}$	$8.33 * 10^{-1}$
C7-DC	$1.10 * 10^{-3}$	$<1.0 * 10^{-4\#\}$	$4.53 * 10^{-1}$
C12	$4.48 * 10^{-1}$	$1.80 * 10^{-3\#\}$	$2.35 * 10^{-1}$
C12:1	$4.43 * 10^{-1}$	$1.90 * 10^{-3\#\}$	$2.25 * 10^{-1}$
C14	$2.24 * 10^{-1}$	$8.70 * 10^{-3\#\}$	$1.01 * 10^{-1}$
C14:1	$4.05 * 10^{-1}$	$3.00 * 10^{-4\#\}$	$1.68 * 10^{-1}$
C14:2	$2.30 * 10^{-1}$	$4.00 * 10^{-4\#\}$	$4.22 * 10^{-1}$
C16	$3.48 * 10^{-1}$	$1.68 * 10^{-2\text{§}}$	$2.09 * 10^{-1}$
C16:1	$8.94 * 10^{-1}$	$1.25 * 10^{-2\text{§}}$	$3.82 * 10^{-1}$
C16:1-OH	$1.42 * 10^{-1}$	$3.30 * 10^{-3\#\}$	$8.79 * 10^{-1}$
C16:2	$8.34 * 10^{-2}$	$4.30 * 10^{-3\text{§}}$	$6.62 * 10^{-1}$
C18:1	$5.97 * 10^{-1}$	$3.00 * 10^{-4\#\}$	$7.64 * 10^{-1}$
C18:2	$1.35 * 10^{-1}$	$<1.00 * 10^{-4\#\}$	$1.81 * 10^{-1}$

DM1 = individuals with type 1 diabetes.

Abbreviations of acylcarnitines are shown in Supplemental Table S3.

[§]p<0.05, #p<0.004 (0.05/13) (two way repeated measure ANOVA).

SUPPLEMENTARY TABLES

Table S1. General population characteristics of healthy individuals and patients with type 1 diabetes¹.

	Healthy	DM1
Females (%)	4 (44%)	4 (57%)
Age (years)	45 ± 14	43 ± 16
BMI (kg/m ²)	23.8 ± 2.2	23.5 ± 2.2
WHR	0.88 ± 0.05	0.81 ± 0.09

DM1 = individuals with type 1 diabetes. BMI = body mass index. WHR = waist hip ratio. Healthy individuals n= 9, DM n= 7. Data are represented as mean ± SD (percentage).

¹Data previously published separately (14;15).

Table S2. Sleep and insulin sensitivity parameters of healthy individuals and patients with type 1 diabetes¹.

	Healthy			DM1			Healthy vs. DM1	
	Normal sleep	Short sleep	Effect of sleep P-value	Normal sleep	Short sleep	Effect of sleep P-value	Normal sleep P-value	Short sleep P-value
Sleep characteristics								
TST (min)	454 ± 26	228 ± 32	<0.0001	469 ± 22	222 ± 19	<0.0001	0.237	0.761
Stage 1 (% of TST)	10 ± 3	11 ± 6	0.490	11 ± 3	10 ± 6	0.868	0.577	0.664
Stage 2 (% of TST)	41 ± 7	35 ± 10	0.006	44 ± 7	41 ± 6	0.160	0.533	0.327
Stage 3 (% of TST) (SWS)	25 ± 5	33 ± 9	0.007	23 ± 9	35 ± 12	0.006	0.570	0.447
REM sleep (% of TST)	24 ± 5	21 ± 8	0.364	22 ± 4	14 ± 6	0.038	0.684	0.069
Sleep efficiency (%)	91 ± 5	90 ± 7	0.699	94 ± 2	93 ± 8	0.484	0.199	0.464
Plasma parameter								
Free fatty acids (mmol/l)	0.63 ± 0.15	0.62 ± 0.14	0.76	0.67 ± 0.31	0.59 ± 0.24	0.23	0.74	0.81
Insulin sensitivity parameters								
EGP ($\mu\text{mol}\cdot\text{kg LBM}^{-1}\cdot\text{min}^{-1}$)	3.57 ± 0.6	4.43 ± 0.8	0.017	6.21 ± 1.9	6.88 ± 1.4	0.505	0.003	0.001
GDR ($\mu\text{mol}\cdot\text{kg LBM}^{-1}\cdot\text{min}^{-1}$)	40.7 ± 14.3	32.5 ± 10.2	0.009	25.5 ± 6.4	22.1 ± 5.1	0.039	0.028	0.035
GIR ($\mu\text{mol}\cdot\text{kg LBM}^{-1}\cdot\text{min}^{-1}$)	36.9 ± 14.4	27.8 ± 10.5	0.006	19.0 ± 7.0	14.9 ± 5.0	0.041	0.014	0.014

DM1= individuals with type 1 diabetes. Insulin sensitivity parameters were determined by hyperinsulinemic euglycemic clamp. EGP = Endogenous glucose production, GDR = glucose disposal rate (glucose Rd), GIR = Glucose infusion rate. Sleep characteristics were determined by polysomnography. TST = total sleep time. SWS = slow wave sleep. Free fatty acids were measured in basal fasting plasma samples. Effect of sleep intervention was tested with paired Students T-test, significant differences shown in bold. Healthy individuals n= 9, DM1 n = 7. Data is presented as means ± SD.

¹Data previously published in separately (14;15).

Table S3. Metabolites determined by Biocrates/iDQ™ p150 kit.

Metabolite Class	Short name	Biochemical Name
Acylcarnitines	C0	DL-Carnitine
	C2	Acetyl-L-carnitine
	C3	Propionyl-L-carnitine
	C3:1	Propenyl-L-carnitine
	C3-DC / C4-OH	Malonyl-L-carnitine / hydroxybutyryl-L-carnitine
	C3-DC-M / C5-OH	Methylmalonyl-L-carnitine / hydroxyvaleryl-L-carnitine
	C3-OH	Hydroxypropionyl-L-carnitine
	C4	Butyryl-L-carnitine
	C4:1	Butenyl-L-carnitine
	C4:1-DC / C6	Fumaryl-L-carnitine/Hexanoyl-L-carnitine
	C5	Valeryl-L-carnitine
	C5:1	Tiglyl-L-carnitine
	C5:1-DC	Glutaconyl-L-carnitine
	C5-DC / C6-OH	Glutaryl-L-carnitine/Hydroxyhexanoyl-L-carnitine
	C5-M-DC	Methylglutaryl-L-carnitine
	C6:1	Hexenoyl-L-carnitine
	C7-DC	Pimelyl-L-carnitine
	C8	Octanoyl-L-carnitine
	C8:1	Octenoyl-L-carnitine
	C9	Nonayl-L-carnitine
	C10	Decanoyl-L-carnitine
	C10:1	Decenoyl-L-carnitine
	C10:2	Decadienyl-L-carnitine
C12	Dodecanoyl-L-carnitine	
C12:1	Dodecenoyl-L-carnitine	

	C12-DC	Dodecanedioyl-L-carnitine
Acylcarnitines	C14	Tetradecanoyl-L-carnitine
	C14:1	Tetradecenoyl-L-carnitine
	C14:1-OH	Hydroxytetradecenoyl-L-carnitine
	C14:2	Tetradecadienyl-L-carnitine
	C14:2-OH	Hydroxytetradecadienyl-L-carnitine
	C16	Hexadecanoyl-L-carnitine
	C16:1	Hexadecenoyl-L-carnitine
	C16:1-OH	Hydroxyhexadecenoyl-L-carnitine
	C16:2	Hexadecadienyl-L-carnitine
	C16:2-OH	Hydroxyhexadecadienyl-L-carnitine
	C16-OH	Hydroxyhexadecanoyl-L-carnitine
	C18	Octadecanoyl-L-carnitine
	C18:1	Octadecenoyl-L-carnitine
	C18:1-OH	Hydroxyoctadecenoyl-L-carnitine
	C18:2	Octadecadienyl-L-carnitine
Sugars	H1	Hexose
Amino acids	Arg	Arginine
	Gln	Glutamine
	Gly	Glycine
	His	Histidine
	Met	Methionine
	Orn	Ornithine
	Phe	Phenylalanine
	Pro	Proline
	Ser	Serine
	Thr	Threonine

	Trp	Tryptophan
Amino acids	Tyr	Tyrosine
	Val	Valine
	xLeu	xLeucine
Glycerophospholipids	lysoPC a C14:0	lysoPhosphatidylcholine acyl C14:0
	lysoPC a C16:0	lysoPhosphatidylcholine acyl C16:0
	lysoPC a C16:1	lysoPhosphatidylcholine acyl C16:1
	lysoPC a C17:0	lysoPhosphatidylcholine acyl C17:0
	lysoPC a C18:0	lysoPhosphatidylcholine acyl C18:0
	lysoPC a C18:1	lysoPhosphatidylcholine acyl C18:1
	lysoPC a C18:2	lysoPhosphatidylcholine acyl C18:2
	lysoPC a C20:3	lysoPhosphatidylcholine acyl C20:3
	lysoPC a C20:4	lysoPhosphatidylcholine acyl C20:4
	lysoPC a C24:0	lysoPhosphatidylcholine acyl C24:0
	lysoPC a C26:0	lysoPhosphatidylcholine acyl C26:0
	lysoPC a C26:1	lysoPhosphatidylcholine acyl C26:1
	lysoPC a C28:0	lysoPhosphatidylcholine acyl C28:0
	lysoPC a C28:1	lysoPhosphatidylcholine acyl C28:1
	lysoPC a C6:0	lysoPhosphatidylcholine acyl C6:0
	PC aa C24:0	Phosphatidylcholine diacyl C 24:0
	PC aa C26:0	Phosphatidylcholine diacyl C 26:0
	PC aa C28:1	Phosphatidylcholine diacyl C 28:1
	PC aa C30:0	Phosphatidylcholine diacyl C 30:0
	PC aa C30:2	Phosphatidylcholine diacyl C 30:2
	PC aa C32:0	Phosphatidylcholine diacyl C 32:0
	PC aa C32:1	Phosphatidylcholine diacyl C 32:1
PC aa C32:2	Phosphatidylcholine diacyl C 32:2	

	PC aa C32:3	Phosphatidylcholine diacyl C 32:3
	PC aa C34:1	Phosphatidylcholine diacyl C 34:1
Glycerophospholipids	PC aa C34:2	Phosphatidylcholine diacyl C 34:2
	PC aa C34:3	Phosphatidylcholine diacyl C 34:3
	PC aa C34:4	Phosphatidylcholine diacyl C 34:4
	PC aa C36:0	Phosphatidylcholine diacyl C 36:0
	PC aa C36:1	Phosphatidylcholine diacyl C 36:1
	PC aa C36:2	Phosphatidylcholine diacyl C 36:2
	PC aa C36:3	Phosphatidylcholine diacyl C 36:3
	PC aa C36:4	Phosphatidylcholine diacyl C 36:4
	PC aa C36:5	Phosphatidylcholine diacyl C 36:5
	PC aa C36:6	Phosphatidylcholine diacyl C 36:6
	PC aa C38:0	Phosphatidylcholine diacyl C 38:0
	PC aa C38:1	Phosphatidylcholine diacyl C 38:1
	PC aa C38:3	Phosphatidylcholine diacyl C 38:3
	PC aa C38:4	Phosphatidylcholine diacyl C 38:4
	PC aa C38:5	Phosphatidylcholine diacyl C 38:5
	PC aa C38:6	Phosphatidylcholine diacyl C 38:6
	PC aa C40:1	Phosphatidylcholine diacyl C 40:1
	PC aa C40:2	Phosphatidylcholine diacyl C 40:2
	PC aa C40:3	Phosphatidylcholine diacyl C 40:3
	PC aa C40:4	Phosphatidylcholine diacyl C 40:4
	PC aa C40:5	Phosphatidylcholine diacyl C 40:5
	PC aa C40:6	Phosphatidylcholine diacyl C 40:6
	PC aa C42:0	Phosphatidylcholine diacyl C 42:0
	PC aa C42:1	Phosphatidylcholine diacyl C 42:1
	PC aa C42:2	Phosphatidylcholine diacyl C 42:2

Glycerophospholipids

PC aa C42:4	Phosphatidylcholine diacyl C 42:4
PC aa C42:5	Phosphatidylcholine diacyl C 42:5
PC aa C42:6	Phosphatidylcholine diacyl C 42:6
PC ae C30:0	Phosphatidylcholine acyl-alkyl C 30:0
PC ae C30:1	Phosphatidylcholine acyl-alkyl C 30:1
PC ae C30:2	Phosphatidylcholine acyl-alkyl C 30:2
PC ae C32:1	Phosphatidylcholine acyl-alkyl C 32:1
PC ae C32:2	Phosphatidylcholine acyl-alkyl C 32:2
PC ae C34:0	Phosphatidylcholine acyl-alkyl C 34:0
PC ae C34:1	Phosphatidylcholine acyl-alkyl C 34:1
PC ae C34:2	Phosphatidylcholine acyl-alkyl C 34:2
PC ae C34:3	Phosphatidylcholine acyl-alkyl C 34:3
PC ae C36:0	Phosphatidylcholine acyl-alkyl C 36:0
PC ae C36:1	Phosphatidylcholine acyl-alkyl C 36:1
PC ae C36:2	Phosphatidylcholine acyl-alkyl C 36:2
PC ae C36:3	Phosphatidylcholine acyl-alkyl C 36:3
PC ae C36:4	Phosphatidylcholine acyl-alkyl C 36:4
PC ae C36:5	Phosphatidylcholine acyl-alkyl C 36:5
PC ae C38:0	Phosphatidylcholine acyl-alkyl C 38:0
PC ae C38:1	Phosphatidylcholine acyl-alkyl C 38:1
PC ae C38:2	Phosphatidylcholine acyl-alkyl C 38:2
PC ae C38:3	Phosphatidylcholine acyl-alkyl C 38:3
PC ae C38:4	Phosphatidylcholine acyl-alkyl C 38:4
PC ae C38:5	Phosphatidylcholine acyl-alkyl C 38:5
PC ae C38:6	Phosphatidylcholine acyl-alkyl C 38:6
PC ae C40:0	Phosphatidylcholine acyl-alkyl C 40:0
PC ae C40:1	Phosphatidylcholine acyl-alkyl C 40:1

	PC ae C40:2	Phosphatidylcholine acyl-alkyl C 40:2
	PC ae C40:3	Phosphatidylcholine acyl-alkyl C 40:3
	PC ae C40:4	Phosphatidylcholine acyl-alkyl C 40:4
	PC ae C40:5	Phosphatidylcholine acyl-alkyl C 40:5
Glycerophospholipids	PC ae C40:6	Phosphatidylcholine acyl-alkyl C 40:6
	PC ae C42:0	Phosphatidylcholine acyl-alkyl C 42:0
	PC ae C42:1	Phosphatidylcholine acyl-alkyl C 42:1
	PC ae C42:2	Phosphatidylcholine acyl-alkyl C 42:2
	PC ae C42:3	Phosphatidylcholine acyl-alkyl C 42:3
	PC ae C42:4	Phosphatidylcholine acyl-alkyl C 42:4
	PC ae C42:5	Phosphatidylcholine acyl-alkyl C 42:5
	PC ae C44:3	Phosphatidylcholine acyl-alkyl C 44:3
	PC ae C44:4	Phosphatidylcholine acyl-alkyl C 44:4
	PC ae C44:5	Phosphatidylcholine acyl-alkyl C 44:5
	PC ae C44:6	Phosphatidylcholine acyl-alkyl C 44:6
Sphingolipids	SM (OH) C14:1	Hydroxysphingomyeline C 14:1
	SM (OH) C16:0	Hydroxysphingomyeline C 16:0
	SM (OH) C22:1	Hydroxysphingomyeline C 22:1
	SM (OH) C22:2	Hydroxysphingomyeline C 22:2
	SM (OH) C24:1	Hydroxysphingomyeline C 24:1
	SM C16:0	Sphingomyeline C 16:0
	SM C16:1	Sphingomyeline C 16:1
	SM C18:0	Sphingomyeline C 18:0
	SM C18:1	Sphingomyeline C 18:1
	SM C20:2	Sphingomyeline C 20:2
	SM C22:3	Sphingomyeline C 22:3
	SM C24:0	Sphingomyeline C 24:0

	SM C24:1	Sphingomyeline C 24:1
	SM C26:0	Sphingomyeline C 26:0
	SM C26:1	Sphingomyeline C 26:1

ACCEPTED MANUSCRIPT

Table S4. Metabolite changes after one night of short sleep duration.

Metabolite	All subjects			Healthy			DM1		
	Mean	Change	P-value	Mean	Change (%) ²	P-value	Mean	Change (%) ²	P-value
C0	0.191	0.6%	8.81×10^{-1}	0.398	1.2%	8.47×10^{-1}	-0.075	-0.3%	9.61×10^{-1}
C2	0.662	16.8%	2.28×10^{-2}	0.184	4.3%	5.40×10^{-1}	1.276	35.6%	1.33×10^{-2}
C3	-0.006	-2.6%	6.76×10^{-1}	-0.002	-0.7%	9.33×10^{-1}	-0.012	-6.2%	6.03×10^{-1}
C3-DC (C4-OH)	0.007	11.3%	4.55×10^{-1}	0.006	8.0%	7.20×10^{-1}	0.009	17.7%	2.87×10^{-1}
C3-OH	0.001	3.3%	5.03×10^{-1}	0.002	8.5%	5.91×10^{-2}	-0.001	-2.1%	8.24×10^{-1}
C3:1	-0.001	-5.9%	4.16×10^{-1}	0.000	-0.1%	9.93×10^{-1}	-0.001	-12.3%	1.58×10^{-1}
C4	0.011	10.3%	1.22×10^{-1}	0.008	7.2%	4.95×10^{-1}	0.015	14.4%	8.68×10^{-2}
C4:1	0.002	6.7%	2.90×10^{-1}	0.002	8.7%	2.55×10^{-1}	0.001	4.8%	6.58×10^{-1}
C5	0.011	10.2%	2.80×10^{-1}	0.012	10.4%	4.59×10^{-1}	0.010	9.8%	4.30×10^{-1}
C5-DC (C6-OH)	-0.001	-4.1%	4.87×10^{-1}	-0.001	-8.3%	2.56×10^{-1}	0.000	0.7%	9.45×10^{-1}
C5-M-DC	0.000	0.3%	9.32×10^{-1}	0.000	-1.0%	8.45×10^{-1}	0.001	1.8%	7.11×10^{-1}
C5-OH (C3-DC-M)	0.000	0.3%	9.40×10^{-1}	0.000	-1.5%	6.89×10^{-1}	0.001	2.4%	7.40×10^{-1}
C5:1	0.000	1.0%	8.74×10^{-1}	-0.001	-2.5%	7.86×10^{-1}	0.001	5.0%	6.20×10^{-1}
C5:1-DC	0.000	2.6%	6.75×10^{-1}	-0.001	-4.8%	5.96×10^{-1}	0.002	12.5%	1.27×10^{-1}
C6 (C4:1-DC)	0.005	8.7%	1.22×10^{-1}	0.001	1.9%	7.82×10^{-1}	0.011	18.8%	6.20×10^{-2}
C6:1	0.001	2.6%	3.99×10^{-1}	0.001	3.5%	3.35×10^{-1}	0.000	1.5%	8.05×10^{-1}
C7-DC	0.005	19.9%	5.41×10^{-4}	0.002	8.4%	8.50×10^{-2}	0.008	34.8%	3.21×10^{-4}
C8	0.003	1.9%	6.06×10^{-1}	-0.005	-3.1%	5.56×10^{-1}	0.013	11.0%	4.38×10^{-3}
C8:1	0.012	15.4%	1.62×10^{-1}	0.008	11.8%	3.93×10^{-1}	0.017	18.9%	3.09×10^{-1}
C9	0.000	-0.4%	9.47×10^{-1}	-0.001	-4.7%	5.73×10^{-1}	0.001	5.3%	5.51×10^{-1}
C10	0.014	5.8%	1.84×10^{-1}	0.003	1.1%	8.62×10^{-1}	0.027	15.1%	4.54×10^{-3}
C10:1	0.008	6.9%	2.68×10^{-1}	-0.004	-3.1%	6.68×10^{-1}	0.022	23.9%	2.72×10^{-2}

C10:2	0.000	-0.7%	8.65×10^{-1}	0.000	-0.1%	9.79×10^{-1}	0.000	-1.3%	8.51×10^{-1}
C12	0.012	17.2%	1.70×10^{-3}	0.010	12.0%	9.00×10^{-2}	0.015	27.5%	3.30×10^{-3}
C12-DC	0.002	2.4%	1.53×10^{-1}	0.002	2.1%	3.78×10^{-1}	0.003	2.8%	3.01×10^{-1}
C12:1	0.017	23.6%	1.74×10^{-3}	0.014	17.0%	8.32×10^{-2}	0.021	35.2%	5.85×10^{-3}
C14	0.004	15.6%	1.21×10^{-2}	0.002	8.5%	2.42×10^{-1}	0.006	28.0%	1.97×10^{-2}
C14:1	0.020	32.4%	2.67×10^{-4}	0.017	23.5%	2.34×10^{-2}	0.024	49.1%	6.55×10^{-3}
C14:1-OH	0.001	6.2%	3.79×10^{-1}	0.000	-1.7%	8.10×10^{-1}	0.001	16.9%	2.36×10^{-1}
C14:2	0.006	26.1%	5.48×10^{-4}	0.005	17.3%	5.01×10^{-2}	0.008	41.5%	4.17×10^{-3}
C14:2-OH	0.000	5.0%	3.64×10^{-1}	0.000	6.4%	2.91×10^{-1}	0.000	3.3%	7.48×10^{-1}
C16	0.007	10.8%	1.94×10^{-2}	0.005	6.7%	2.05×10^{-1}	0.010	17.1%	5.91×10^{-2}
C16-OH	0.000	0.9%	8.89×10^{-1}	0.000	-3.4%	5.84×10^{-1}	0.000	6.0%	6.49×10^{-1}
C16:1	0.005	8.8%	9.60×10^{-3}	0.005	8.3%	3.06×10^{-2}	0.005	9.6%	1.55×10^{-1}
C16:1-OH	0.001	20.8%	5.71×10^{-3}	0.001	11.7%	1.90×10^{-1}	0.002	33.8%	1.10×10^{-2}
C16:2	0.001	21.3%	9.72×10^{-3}	0.001	9.6%	2.39×10^{-1}	0.002	39.5%	1.91×10^{-2}
C16:2-OH	0.000	2.1%	6.29×10^{-1}	0.000	3.8%	5.45×10^{-1}	0.000	-0.1%	9.83×10^{-1}
C18	0.003	10.8%	7.34×10^{-2}	0.002	9.6%	3.04×10^{-1}	0.003	12.5%	1.18×10^{-1}
C18:1	0.016	22.3%	1.92×10^{-4}	0.015	20.1%	2.96×10^{-3}	0.019	25.1%	2.84×10^{-2}
C18:1-OH	0.001	12.8%	6.02×10^{-2}	0.001	5.7%	3.93×10^{-1}	0.002	22.3%	1.07×10^{-1}
C18:2	0.007	27.0%	1.32×10^{-4}	0.005	20.7%	9.01×10^{-4}	0.010	34.4%	1.38×10^{-2}
Arg	4.297	6.0%	2.74×10^{-1}	6.988	10.0%	2.66×10^{-1}	0.838	1.1%	8.57×10^{-1}
Gln	13.186	4.5%	2.52×10^{-1}	10.500	3.6%	4.88×10^{-1}	16.641	5.7%	4.00×10^{-1}
Gly	-9.002	-4.7%	2.84×10^{-1}	-3.264	-1.9%	7.57×10^{-1}	-16.380	-7.4%	2.69×10^{-1}
His	0.533	1.0%	8.27×10^{-1}	-0.174	-0.3%	9.57×10^{-1}	1.441	3.0%	7.32×10^{-1}
Met	0.264	1.3%	7.98×10^{-1}	0.557	2.6%	6.91×10^{-1}	-0.113	-0.6%	9.48×10^{-1}
Orn	3.038	8.3%	1.48×10^{-1}	5.466	15.2%	1.05×10^{-1}	-0.083	-0.2%	9.70×10^{-1}

Phe	2.588	6.2%	1.06×10^{-1}	3.131	7.6%	1.16×10^{-1}	1.890	4.5%	5.13×10^{-1}
Pro	-3.161	-1.9%	6.20×10^{-1}	-0.909	-0.5%	9.10×10^{-1}	-6.057	-4.0%	5.93×10^{-1}
Ser	3.273	3.6%	4.27×10^{-1}	2.253	2.7%	7.18×10^{-1}	4.585	4.6%	4.29×10^{-1}
Thr	4.527	4.7%	5.04×10^{-1}	4.930	5.1%	5.55×10^{-1}	4.010	4.1%	7.47×10^{-1}
Trp	1.178	1.9%	5.56×10^{-1}	2.563	4.2%	3.30×10^{-1}	-0.602	-1.0%	8.58×10^{-1}
Tyr	4.638	9.8%	1.05×10^{-1}	6.966	14.7%	1.30×10^{-1}	1.644	3.5%	6.12×10^{-1}
Val	3.933	2.6%	4.86×10^{-1}	6.183	3.9%	5.06×10^{-1}	1.039	0.7%	8.67×10^{-1}
xLeu	9.916	6.4%	1.56×10^{-1}	9.156	5.7%	3.55×10^{-1}	10.892	7.3%	3.24×10^{-1}
lysoPC a C14:0	-0.047	-1.4%	5.83×10^{-1}	-0.022	-0.6%	8.67×10^{-1}	-0.079	-2.4%	5.04×10^{-1}
lysoPC a C16:0	-1.774	-2.4%	5.84×10^{-1}	0.872	1.3%	8.39×10^{-1}	-5.176	-6.6%	3.34×10^{-1}
lysoPC a C16:1	-0.144	-7.2%	1.58×10^{-1}	-0.061	-3.1%	6.72×10^{-1}	-0.249	-12.3%	1.02×10^{-1}
lysoPC a C17:0	-0.015	-1.3%	7.88×10^{-1}	0.005	0.5%	9.46×10^{-1}	-0.042	-3.4%	6.57×10^{-1}
lysoPC a C18:0	-0.003	0.0%	9.97×10^{-1}	1.077	6.4%	3.22×10^{-1}	-1.392	-6.9%	4.33×10^{-1}
lysoPC a C18:1	-0.7	-2.9%	5.91×10^{-1}	-0.210	-1.6%	7.91×10^{-1}	-0.728	-4.2%	6.67×10^{-1}
lysoPC a C18:2	0.575	1.9%	8.14×10^{-1}	-1.447	-5.5%	4.00×10^{-1}	3.176	8.9%	5.59×10^{-1}
lysoPC a C20:3	0.008	0.4%	9.43×10^{-1}	0.080	4.0%	5.80×10^{-1}	-0.085	-4.5%	6.19×10^{-1}
lysoPC a C20:4	-0.248	-4.7%	3.27×10^{-1}	-0.302	-6.3%	2.31×10^{-1}	-0.179	-3.1%	7.32×10^{-1}
lysoPC a C24:0	-0.003	-0.5%	9.28×10^{-1}	0.014	3.1%	6.56×10^{-1}	-0.024	-4.7%	6.56×10^{-1}
lysoPC a C26:0	-0.009	-0.9%	9.02×10^{-1}	0.014	1.4%	8.63×10^{-1}	-0.039	-4.1%	7.87×10^{-1}
lysoPC a C26:1	-0.004	-1.1%	8.81×10^{-1}	-0.002	-0.6%	9.42×10^{-1}	-0.007	-1.7%	9.03×10^{-1}
lysoPC a C28:0	0.001	0.1%	9.84×10^{-1}	0.012	1.5%	8.49×10^{-1}	-0.013	-1.7%	8.84×10^{-1}
lysoPC a C28:1	-0.006	-0.6%	9.28×10^{-1}	0.010	1.1%	8.84×10^{-1}	-0.025	-2.6%	8.31×10^{-1}
lysoPC a C6:0	-0.002	-8.4%	4.94×10^{-1}	0.002	10.9%	6.26×10^{-1}	-0.007	-26.5%	1.14×10^{-2}
PC aa C24:0	-0.005	-1.8%	7.93×10^{-1}	-0.005	-1.9%	8.42×10^{-1}	-0.005	-1.8%	8.78×10^{-1}
PC aa C26:0	-0.028	-2.4%	6.22×10^{-1}	-0.040	-3.4%	5.37×10^{-1}	-0.013	-1.1%	9.09×10^{-1}

PC aa C28:1	0.061	2.7%	3.74×10^{-1}	0.089	3.9%	2.47×10^{-1}	0.026	1.1%	8.47×10^{-1}
PC aa C30:0	0.081	2.7%	6.62×10^{-1}	0.099	3.1%	6.76×10^{-1}	0.058	2.1%	8.60×10^{-1}
PC aa C30:2	0.002	1.2%	9.33×10^{-1}	0.010	7.0%	7.71×10^{-1}	-0.009	-4.3%	8.40×10^{-1}
PC aa C32:0	0.352	4.0%	2.19×10^{-1}	0.490	5.8%	5.12×10^{-2}	0.174	1.9%	7.76×10^{-1}
PC aa C32:1	0.370	3.7%	6.36×10^{-1}	1.063	9.4%	3.68×10^{-1}	-0.521	-6.5%	6.17×10^{-1}
PC aa C32:2	0.020	0.8%	9.12×10^{-1}	-0.038	-1.3%	8.84×10^{-1}	0.095	4.1%	7.38×10^{-1}
PC aa C32:3	0.001	0.2%	9.68×10^{-1}	-0.005	-1.3%	7.93×10^{-1}	0.007	1.9%	7.90×10^{-1}
PC aa C34:1	2.395	1.9%	5.09×10^{-1}	5.489	4.3%	1.84×10^{-1}	-1.583	-1.3%	8.16×10^{-1}
PC aa C34:2	7.304	3.7%	1.04×10^{-1}	6.544	3.3%	1.31×10^{-1}	8.282	4.1%	3.78×10^{-1}
PC aa C34:3	0.353	3.0%	5.32×10^{-1}	0.435	3.5%	5.90×10^{-1}	0.247	2.2%	7.78×10^{-1}
PC aa C34:4	-0.013	-1.2%	8.47×10^{-1}	-0.032	-2.7%	7.55×10^{-1}	0.012	1.3%	8.98×10^{-1}
PC aa C36:0	0.017	0.7%	9.23×10^{-1}	-0.046	-2.1%	8.05×10^{-1}	0.099	3.3%	7.81×10^{-1}
PC aa C36:1	1.673	5.8%	1.72×10^{-1}	2.555	8.7%	1.40×10^{-1}	0.538	1.9%	7.75×10^{-1}
PC aa C36:2	6.029	4.8%	1.46×10^{-1}	6.133	4.9%	1.96×10^{-1}	5.894	4.7%	4.61×10^{-1}
PC aa C36:3	3.808	4.5%	1.64×10^{-1}	6.521	7.5%	5.27×10^{-2}	0.320	0.4%	9.47×10^{-1}
PC aa C36:4	1.441	1.3%	6.29×10^{-1}	1.110	1.0%	7.39×10^{-1}	1.866	1.9%	7.49×10^{-1}
PC aa C36:5	-0.045	-0.3%	9.65×10^{-1}	0.559	4.1%	7.51×10^{-1}	-0.821	-6.8%	3.50×10^{-1}
PC aa C36:6	-0.010	-1.5%	8.53×10^{-1}	0.004	0.5%	9.69×10^{-1}	-0.028	-4.4%	6.29×10^{-1}
PC aa C38:0	0.050	2.0%	6.68×10^{-1}	0.039	1.7%	8.06×10^{-1}	0.065	2.3%	7.41×10^{-1}
PC aa C38:1	0.041	2.3%	6.99×10^{-1}	0.151	9.3%	2.19×10^{-1}	-0.101	-5.0%	5.98×10^{-1}
PC aa C38:3	2.020	7.5%	6.46×10^{-2}	3.788	12.9%	1.68×10^{-2}	-0.254	-1.1%	8.48×10^{-1}
PC aa C38:4	1.113	2.0%	5.25×10^{-1}	1.760	3.0%	4.88×10^{-1}	0.280	0.6%	9.16×10^{-1}
PC aa C38:5	0.060	0.2%	9.55×10^{-1}	0.609	2.3%	7.31×10^{-1}	-0.645	-2.6%	5.62×10^{-1}
PC aa C38:6	-0.622	-1.6%	7.12×10^{-1}	-0.987	-2.3%	7.03×10^{-1}	-0.154	-0.4%	9.47×10^{-1}
PC aa C40:1	0.020	2.5%	6.90×10^{-1}	0.060	8.0%	3.34×10^{-1}	-0.031	-3.5%	7.38×10^{-1}

PC aa C40:2	0.078	5.4%	4.46×10^{-1}	0.149	11.7%	2.16×10^{-1}	-0.012	-0.7%	9.52×10^{-1}
PC aa C40:3	0.068	4.6%	4.70×10^{-1}	0.136	10.5%	2.55×10^{-1}	-0.020	-1.2%	9.04×10^{-1}
PC aa C40:4	0.154	4.6%	2.35×10^{-1}	0.290	8.9%	1.09×10^{-1}	-0.021	-0.6%	9.16×10^{-1}
PC aa C40:5	0.232	4.1%	3.01×10^{-1}	0.540	9.3%	9.49×10^{-2}	-0.164	-3.0%	5.89×10^{-1}
PC aa C40:6	0.033	0.3%	9.48×10^{-1}	0.413	3.3%	5.51×10^{-1}	-0.454	-4.4%	5.85×10^{-1}
PC aa C42:0	0.028	4.0%	2.80×10^{-1}	0.040	6.2%	2.00×10^{-1}	0.012	1.6%	8.01×10^{-1}
PC aa C42:1	0.009	1.9%	7.04×10^{-1}	0.041	10.0%	1.49×10^{-1}	-0.033	-6.2%	3.74×10^{-1}
PC aa C42:2	0.020	3.1%	6.55×10^{-1}	0.054	9.5%	3.06×10^{-1}	-0.024	-3.1%	7.75×10^{-1}
PC aa C42:4	0.031	5.6%	2.44×10^{-1}	0.053	10.6%	1.88×10^{-1}	0.002	0.4%	9.49×10^{-1}
PC aa C42:5	0.029	5.5%	1.84×10^{-1}	0.047	9.7%	1.34×10^{-1}	0.005	0.8%	8.81×10^{-1}
PC aa C42:6	0.019	3.4%	5.38×10^{-1}	0.049	9.3%	1.60×10^{-1}	-0.020	-3.5%	7.21×10^{-1}
PC ae C30:0	0.002	0.5%	9.20×10^{-1}	0.012	3.5%	6.07×10^{-1}	-0.012	-3.6%	5.95×10^{-1}
PC ae C30:1	-0.003	-1.2%	8.94×10^{-1}	-0.003	-1.2%	9.07×10^{-1}	-0.004	-1.3%	9.43×10^{-1}
PC ae C30:2	0.006	2.5%	6.96×10^{-1}	0.011	5.1%	5.28×10^{-1}	-0.001	-0.3%	9.81×10^{-1}
PC ae C32:1	0.046	2.4%	4.28×10^{-1}	0.035	1.9%	6.30×10^{-1}	0.060	2.9%	5.66×10^{-1}
PC ae C32:2	0.011	2.2%	4.36×10^{-1}	0.017	3.7%	3.50×10^{-1}	0.003	0.6%	8.99×10^{-1}
PC ae C34:0	0.006	0.7%	8.25×10^{-1}	0.015	1.6%	6.90×10^{-1}	-0.005	-0.5%	9.25×10^{-1}
PC ae C34:1	0.159	2.7%	3.75×10^{-1}	0.202	3.3%	4.23×10^{-1}	0.105	1.8%	7.15×10^{-1}
PC ae C34:2	0.274	4.1%	4.06×10^{-1}	0.029	0.4%	9.36×10^{-1}	0.589	8.7%	3.55×10^{-1}
PC ae C34:3	0.213	3.9%	3.04×10^{-1}	0.093	1.8%	7.10×10^{-1}	0.368	6.3%	3.34×10^{-1}
PC ae C36:0	0.048	6.1%	1.66×10^{-1}	0.037	4.6%	3.13×10^{-1}	0.063	8.0%	3.68×10^{-1}
PC ae C36:1	0.832	6.7%	3.11×10^{-1}	0.792	6.9%	3.45×10^{-1}	0.883	6.4%	5.98×10^{-1}
PC ae C36:2	0.308	2.9%	3.51×10^{-1}	0.152	1.4%	7.39×10^{-1}	0.508	4.7%	3.39×10^{-1}
PC ae C36:3	0.143	2.8%	5.30×10^{-1}	0.044	0.9%	8.77×10^{-1}	0.272	5.4%	5.13×10^{-1}
PC ae C36:4	0.257	2.7%	5.65×10^{-1}	-0.112	-1.2%	7.36×10^{-1}	0.732	7.4%	4.57×10^{-1}

PC ae C36:5	0.079	1.2%	7.36×10^{-1}	-0.063	-0.9%	8.10×10^{-1}	0.260	3.7%	5.64×10^{-1}
PC ae C38:0	0.054	2.8%	6.14×10^{-1}	0.049	2.4%	7.92×10^{-1}	0.061	3.4%	5.34×10^{-1}
PC ae C38:1	0.263	6.4%	4.50×10^{-1}	0.411	11.8%	3.06×10^{-1}	0.073	1.5%	9.13×10^{-1}
PC ae C38:2	0.414	6.9%	3.07×10^{-1}	0.631	11.3%	1.77×10^{-1}	0.134	2.1%	8.61×10^{-1}
PC ae C38:3	0.776	7.3%	1.96×10^{-1}	0.799	7.6%	2.04×10^{-1}	0.745	6.9%	5.38×10^{-1}
PC ae C38:4	0.189	2.2%	5.31×10^{-1}	0.199	2.3%	6.11×10^{-1}	0.176	2.0%	7.39×10^{-1}
PC ae C38:5	0.132	1.3%	7.42×10^{-1}	-0.130	-1.3%	7.68×10^{-1}	0.470	4.4%	5.50×10^{-1}
PC ae C38:6	0.096	2.5%	5.50×10^{-1}	-0.033	-0.9%	8.70×10^{-1}	0.261	6.8%	3.48×10^{-1}
PC ae C40:0	0.010	0.3%	9.35×10^{-1}	0.044	1.1%	8.20×10^{-1}	-0.032	-0.9%	8.54×10^{-1}
PC ae C40:1	0.040	1.8%	7.94×10^{-1}	0.031	1.5%	8.49×10^{-1}	0.051	2.2%	8.67×10^{-1}
PC ae C40:2	0.144	5.2%	3.60×10^{-1}	0.233	8.5%	2.19×10^{-1}	0.031	1.1%	9.15×10^{-1}
PC ae C40:3	0.349	7.0%	1.55×10^{-1}	0.441	9.1%	1.16×10^{-1}	0.231	4.5%	6.22×10^{-1}
PC ae C40:4	0.223	4.5%	3.41×10^{-1}	0.332	7.1%	2.64×10^{-1}	0.082	1.6%	8.42×10^{-1}
PC ae C40:5	0.278	4.4%	3.21×10^{-1}	0.241	3.8%	5.41×10^{-1}	0.325	5.2%	4.65×10^{-1}
PC ae C40:6	0.051	1.8%	6.17×10^{-1}	0.057	2.0%	7.26×10^{-1}	0.044	1.5%	7.36×10^{-1}
PC ae C42:0	0.028	4.0%	2.93×10^{-1}	0.049	7.1%	1.08×10^{-1}	0.000	0.1%	9.94×10^{-1}
PC ae C42:1	0.044	4.5%	4.52×10^{-1}	0.075	8.4%	2.37×10^{-1}	0.004	0.3%	9.75×10^{-1}
PC ae C42:2	0.023	2.3%	6.72×10^{-1}	0.057	6.1%	4.38×10^{-1}	-0.020	-1.8%	8.26×10^{-1}
PC ae C42:3	0.047	3.4%	5.37×10^{-1}	0.090	7.0%	3.21×10^{-1}	-0.007	-0.5%	9.59×10^{-1}
PC ae C42:4	0.060	4.2%	2.59×10^{-1}	0.120	8.7%	1.16×10^{-1}	-0.017	-1.1%	8.27×10^{-1}
PC ae C42:5	0.119	4.3%	1.96×10^{-1}	0.149	5.4%	2.06×10^{-1}	0.081	2.9%	6.20×10^{-1}
PC ae C44:3	0.011	3.0%	6.00×10^{-1}	0.022	6.8%	4.11×10^{-1}	-0.003	-0.8%	9.26×10^{-1}
PC ae C44:4	0.016	3.3%	2.93×10^{-1}	0.040	8.1%	5.12×10^{-2}	-0.014	-2.7%	5.62×10^{-1}
PC ae C44:5	0.041	2.8%	3.86×10^{-1}	0.058	3.9%	3.78×10^{-1}	0.018	1.2%	8.06×10^{-1}
PC ae C44:6	0.021	1.9%	6.18×10^{-1}	0.060	5.7%	2.69×10^{-1}	-0.031	-2.7%	6.51×10^{-1}

SM (OH) C14:1	0.146	2.6%	2.81×10^{-1}	0.213	4.0%	1.67×10^{-1}	0.059	1.0%	8.19×10^{-1}
SM (OH) C16:1	0.096	3.7%	1.60×10^{-1}	0.084	3.3%	3.39×10^{-1}	0.111	4.2%	3.54×10^{-1}
SM (OH) C22:1	0.271	2.4%	5.34×10^{-1}	0.730	6.9%	1.15×10^{-1}	-0.320	-2.7%	7.02×10^{-1}
SM (OH) C22:2	-0.036	-0.4%	8.93×10^{-1}	0.084	1.0%	7.57×10^{-1}	-0.191	-2.0%	7.29×10^{-1}
SM (OH) C24:1	0.021	1.8%	7.53×10^{-1}	0.075	6.5%	9.51×10^{-2}	-0.049	-4.0%	7.43×10^{-1}
SM C16:0	3.468	3.9%	1.56×10^{-1}	4.587	5.4%	8.66×10^{-2}	2.030	2.1%	6.71×10^{-1}
SM C16:1	0.585	4.0%	1.77×10^{-1}	0.568	4.3%	1.42×10^{-1}	0.606	3.8%	5.14×10^{-1}
SM C18:0	0.859	5.0%	9.13×10^{-2}	0.375	2.1%	5.47×10^{-1}	1.482	8.9%	9.81×10^{-2}
SM C18:1	0.265	3.2%	3.15×10^{-1}	0.014	0.2%	9.65×10^{-1}	0.588	6.8%	2.07×10^{-1}
SM C20:2	0.024	5.5%	3.80×10^{-1}	0.010	2.3%	7.28×10^{-1}	0.042	9.9%	4.41×10^{-1}
SM C22:3	-0.017	-5.8%	7.54×10^{-1}	-0.021	-7.4%	8.15×10^{-1}	-0.061	-19.9%	2.00×10^{-1}
SM C24:0	1.132	5.0%	1.61×10^{-1}	2.201	9.9%	3.84×10^{-2}	-0.242	-1.0%	8.48×10^{-1}
SM C24:1	2.222	4.7%	6.72×10^{-2}	2.991	6.6%	3.74×10^{-2}	1.234	2.5%	5.82×10^{-1}
SM C26:0	0.032	51.0%	1.31×10^{-1}	0.023	24.4%	3.09×10^{-1}	0.050	139.3%	9.71×10^{-2}
SM C26:1	0.017	5.1%	4.07×10^{-1}	-0.008	-2.5%	7.53×10^{-1}	0.048	14.8%	1.48×10^{-1}
H1	419.618	7.8%	3.58×10^{-1}	126.200	2.8%	3.61×10^{-1}	796.870	12.3%	4.65×10^{-1}

¹ Difference in metabolite levels (μM) as measured by Biocrates*IDQ*TM p150 kit between short and normal sleep duration. Positive mean difference indicates an increase after short sleep duration. Negative mean difference indicates a decrease after short sleep duration.

²Change (%) represents percentage of change in metabolite level in short compared to normal sleep (metabolite level (short sleep) – metabolite level (normal sleep)) / metabolite level (normal sleep).

DM1 = individuals with type 1 diabetes. P-values are based on paired Students t-tests. Abbreviations of all metabolites are shown in Supplemental Table S3. N= 16 (healthy: n=9, DM1: n=7).

Table S5: Acylcarnitine levels after normal sleep duration.

Metabolite	Healthy		DM1		Healthy vs. DM1
	Mean	SD	Mean	SD	P-value
C0	31.9	5.8	26.6	4.8	0.073
C2	4.24	0.57	3.58	1.06	0.133
C3	0.273	0.043	0.192	0.042	0.002
C3:1	0.010	0.002	0.012	0.003	0.139
C3-DC (C4-OH)	0.074	0.062	0.049	0.011	0.304
C3-OH	0.020	0.003	0.024	0.005	0.020
C4	0.105	0.019	0.105	0.057	0.979
C4:1	0.022	0.003	0.031	0.007	0.005
C5	0.120	0.032	0.097	0.027	0.153
C5:1	0.022	0.003	0.024	0.004	0.148
C5:1-DC	0.020	0.004	0.019	0.002	0.628
C5-DC (C6-OH)	0.017	0.003	0.019	0.005	0.273
C5-M-DC	0.035	0.003	0.039	0.004	0.047
C5-OH (C3-DC-M)	0.023	0.003	0.025	0.004	0.405
C6 (C4:1-DC)	0.066	0.022	0.058	0.014	0.385
C6:1	0.027	0.003	0.026	0.002	0.828
C7-DC	0.025	0.008	0.024	0.008	0.975
C8	0.172	0.126	0.121	0.066	0.356
C8:1	0.065	0.009	0.088	0.043	0.151
C9	0.025	0.010	0.024	0.005	0.817
C10	0.274	0.228	0.181	0.073	0.321
C10:1	0.124	0.073	0.093	0.039	0.324
C10:2	0.027	0.006	0.028	0.004	0.593
C12	0.084	0.051	0.055	0.015	0.179
C12:1	0.080	0.035	0.059	0.018	0.163
C12-DC	0.087	0.005	0.101	0.005	0.000#
C14	0.029	0.008	0.022	0.004	0.044
C14:1	0.071	0.028	0.049	0.015	0.080
C14:1-OH	0.009	0.002	0.009	0.001	0.684
C14:2	0.027	0.016	0.020	0.008	0.300
C14:2-OH	0.007	0.001	0.007	0.001	0.277
C16	0.072	0.018	0.060	0.010	0.136
C16:1	0.059	0.004	0.056	0.006	0.231
C16:1-OH	0.006	0.001	0.005	0.001	0.174
C16:2	0.007	0.002	0.006	0.001	0.164
C16:2-OH	0.010	0.001	0.010	0.002	0.684

C16-OH	0.006	0.001	0.007	0.003	0.714
C18	0.025	0.007	0.024	0.007	0.798
C18:1	0.073	0.009	0.074	0.019	0.949
C18:1-OH	0.009	0.001	0.009	0.001	0.668
C18:2	0.025	0.006	0.028	0.006	0.467

Mean = mean plasma metabolite level (μM). DM1 = individuals with type 1 diabetes. [#] $P < 0.001$ (0.05/41). P-values are based on independent Students t-tests. Abbreviations of all metabolites are shown in Supplemental Table S3. Healthy individuals n=9, DM1 n=7.

Table S6: Acylcarnitine levels after short sleep duration.

Metabolite	Healthy		DM1		Healthy vs. DM1
	Mean	SD	Mean	SD	P-value
C0	32.3	7.0	26.6	6.4	0.113
C2	0.287	0.212	0.208	0.078	0.434
C3	0.120	0.057	0.115	0.037	0.839
C3:1	0.027	0.005	0.028	0.005	0.680
C3-DC (C4-OH)	0.094	0.056	0.070	0.018	0.309
C3-OH	0.094	0.034	0.080	0.023	0.356
C4	0.089	0.006	0.104	0.011	0.004
C4:1	0.032	0.008	0.028	0.006	0.300
C5	0.087	0.032	0.073	0.026	0.336
C5:1	0.009	0.001	0.010	0.003	0.261
C5:1-DC	0.032	0.016	0.028	0.008	0.595
C5-DC (C6-OH)	0.007	0.001	0.008	0.002	0.447
C5-M-DC	0.077	0.016	0.070	0.014	0.398
C5-OH (C3-DC-M)	0.063	0.006	0.061	0.011	0.577
C6 (C4:1-DC)	0.007	0.002	0.007	0.002	0.610
C6:1	0.008	0.002	0.008	0.003	0.715
C7-DC	0.010	0.001	0.010	0.001	0.937
C8	0.006	0.001	0.007	0.002	0.213
C8:1	0.027	0.009	0.027	0.006	0.932
C9	0.088	0.014	0.092	0.024	0.664
C10	0.010	0.002	0.011	0.002	0.225
C10:1	0.031	0.006	0.037	0.009	0.103
C10:2	4.421	1.169	4.859	1.478	0.517
C12	0.271	0.065	0.180	0.039	0.006
C12:1	0.010	0.002	0.010	0.003	0.810
C12-DC	0.080	0.047	0.057	0.015	0.241
C14	0.021	0.003	0.024	0.005	0.255
C14:1	0.113	0.028	0.120	0.072	0.793

C14:1-OH	0.024	0.004	0.032	0.007	0.012
C14:2	0.132	0.038	0.106	0.036	0.188
C14:2-OH	0.021	0.005	0.026	0.006	0.121
C16	0.019	0.004	0.021	0.004	0.209
C16:1	0.015	0.003	0.019	0.003	0.028
C16:1-OH	0.034	0.005	0.040	0.007	0.086
C16:2	0.023	0.003	0.025	0.005	0.222
C16:2-OH	0.068	0.016	0.069	0.025	0.919
C16-OH	0.027	0.003	0.027	0.004	0.637
C18	0.027	0.008	0.033	0.008	0.159
C18:1	0.166	0.109	0.135	0.063	0.507
C18:1-OH	0.073	0.028	0.104	0.048	0.128
C18:2	0.024	0.007	0.025	0.005	0.645

Mean = mean plasma metabolite level (μM). DM1 = individuals with type 1 diabetes. P-values are based on independent Students t-tests. Abbreviations of all metabolites are shown in Supplemental Table S3. Healthy individuals n=9, DM1 n=7.

- One night of short sleep acutely increases plasma acylcarnitine levels
- Short sleep increases acylcarnitine in healthy individuals and patients with DM1
- Acylcarnitines reflect fatty acid oxidation and have pro-inflammatory properties
- Acylcarnitines may mediate the relation between short sleep and insulin resistance

ACCEPTED MANUSCRIPT