

1 **Presence and regulation of D1 and D2 deiodinases in rat white adipose tissue**

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27 **Abstract**

28 Thyroid hormones regulate adipogenic differentiation, lipogenic and lipolytic metabolism and
29 mitochondrial activity in adipose tissue. T3 levels in tissues are regulated by the deiodinase
30 enzymes.

31 Objective: To study the activity and mRNA expression of the 5' outer-ring deiodinases (D1
32 and D2) and thyroid hormone concentrations in rat white adipose tissue (WAT), where only
33 D1 activity had been described.

34 Methods: Control, thyroidectomized and thyroid hormone treated rats were used. D1 and D2
35 mRNA were determined in WAT by qRT-PCR using Taqman probes, D1 and D2 activities
36 were determined using rT3 and T4 as substrates. T4 and T3 were measured by
37 radioimmunoassay in plasma, liver and adipose tissue.

38 Results: D1 and D2 mRNAs are present in epididymal rat WAT with similar abundance,
39 which are 7% of the D2 mRNA levels in BAT and 1% that of D1 in liver. The Kms in WAT
40 are 40 nM T4 for D2 and 0.35 μ M rT3 for D1. Both D1 and D2 are regulated in rat
41 epididymal WAT by thyroidal status. T4 and T3 concentrations in plasma, liver and WAT
42 decreased after thyroidectomy (Tx) and recovered after treatment with T4+T3. Both D1 and
43 D2 mRNA increased in WAT from Tx rats and T4+T3 treatment inhibited them, especially
44 D2 mRNA. D1 activity did not changed with thyroidal status, while D2 activity was inhibited
45 by T4+T3.

46 Conclusions: the presence of both deiodinases in WAT suggests important roles in regulating
47 T3 bioavailability for adipose tissue function and regulation of lipid metabolism and
48 thermogenesis.

49

50 **Keywords:** epididymal adipose tissue, thyroid hormones, outer-ring deiodinases.

51

52 **List of abbreviations:**

53 BAT: brown adipose tissue; D1: type1 deiodinase; D2: type 2 deiodinase; D3: type3
54 deiodinase; DTT: dithiothreitol; HEPES: 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid;
55 Km: Michaelis-Menten constant; PCR: polymerase chain reaction; PTU: 6-propyl-2-
56 thiouracil; qRT-PCR: quantitative Real-time PCR; rT3: reverse triiodothyronine; T3:
57 triiodothyronine; T4: thyroxine; Tx: thyroidectomized; UCP1: uncoupling protein 1; Vmax:
58 maximal velocity; WAT: white adipose tissue.

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64 **Introduction**

65 Deiodinases are selenoenzymes that regulate thyroid hormone concentrations in tissues.
66 Two isoenzymes, D1 and D2, catalyze 5' deiodination producing T3 from T4. D1 is present in
67 liver, kidney and thyroid. D1 Km is 0.2-0.5 μM [1], is inhibited by PTU, increases in
68 hyperthyroidism and decreases in hypothyroidism, except thyroid D1 which increases in
69 hypothyroidism, by thyrotropin stimulation [2]. D2 is present in pituitary, brain, BAT and
70 other tissues, is not inhibited by PTU, its Km in nanomolar range. D2 increases in
71 hypothyroidism and produces T3 for local needs. D3, the main T3 degradation pathway,
72 catalyzes inner ring deiodination of T4 and T3, leading to inactive metabolites.

73 D2 is essential for the regulation of T3 availability during specific events of development,
74 in the ear, retina, brain or BAT. T3 is required for the differentiation program of adipocytes
75 [3], regulating the expression of genes involved in the differentiation and metabolism of
76 adipose tissue. Most studies demonstrating the importance of deiodinases, specifically D2,
77 have been performed in BAT, where T3, locally produced by D2, is important for full
78 thermogenesis, UCP1 expression and lipogenesis [4-7]. So far, no studies have been
79 performed on D1 or D2 abundance in WAT, its hormonal regulation or potential role in
80 different metabolic processes, though D1 activity in WAT was found when D2 was reported
81 in BAT [8]. Some studies have shown induction of brown adipocytes in inguinal WAT
82 depots, using UCP1 and D2 as markers [9, 10]. In this respect epididymal WAT is considered
83 "pure" WAT, never converted into BAT even under extreme cold exposure, as opposed to
84 inguinal WAT considered "convertible" adipose tissue [11]. In humans, WAT represents 15-
85 20% of the body weight in lean subjects and this percentage is much higher in obese people
86 (50%). Thus, WAT might represent one of the largest pools of thyroid hormones. In addition,
87 if WAT can be converted into BAT (inducing UCP1), WAT represents a potential therapeutic

88 target to increase energy expenditure in obesity. Here we studied the presence and regulation
89 by thyroid status of D1 and D2 activities and mRNA levels in rat WAT.

90

91 **Materials and Methods**

92 *Animals and treatments*

93 Protocols following the European Community guidelines were approved by our ethic
94 Committee. Male Wistar rats were divided in 3 groups: Control, Tx and Tx rats treated with
95 T4+T3 (2.4 µg T4+0.4 µg T3/day/100 g body weight) as described [12]. Plasma and organs
96 were dissected out and frozen on dry ice. Epididymal WAT was carefully dissected avoiding
97 the reproductive male organs.

98 *Analytical procedures*

99 *Determination of D1 and D2 activities.* D1 and D2 activities were assayed in WAT
100 homogenates (1:8, weight/volume) using 0.32 M sucrose, 10 mM HEPES and 10 mM DTT.
101 Homogenates were centrifuged 5 minutes at 1000 rpm to separate the upper lipid cake, the
102 infranatant was used to measure deiodinase activities.

103 D1 activity: 60.000 cpm [¹²⁵I]-rT₃/sample, 100 nM rT₃ and 5 mM DTT, for 1 h at 37 C
104 using 20-30 µg protein/ 100 µl [13]. D2 activity: 100.000 cpm [¹²⁵I]-T₄/tube, 2 nM T₄+ 1
105 µM T₃, 20 mM DTT and 1 mM PTU/100 µl, using the same conditions and protein [14]. For
106 kinetic analysis: 2-500 nM rT₃ and 5 mM DTT were used for D1 and 1-50 nM T₄ and 20
107 mM DTT for D2.

108 *D1 and D2 mRNA*

109 Total RNA was extracted using the RNeasy lipid tissue (Quiagen). D1 and D2 mRNAs
110 were measured by qRT-PCR, using Taqman probes (Rn00572183m1, Rn00581867m1,
111 Applied Biosystems). After normalization to 18S rRNA, the fold-change in mRNA
112 expression was calculated by the $2^{-\Delta\Delta Ct}$ method.

113 *Determination of thyroid hormone concentrations.*

114 Thyroid hormone concentrations were determined by RIA in plasma, liver and WAT, after
115 extraction and purification [15]. High specific activity T4, rT3 and T3 labelled with [¹²⁵I] were
116 synthesized as described [15].

117 *Statistical analysis*

118 Mean values (\pm SEM) are given. Significant differences were achieved by one-way
119 ANOVA analysis.

120

121 **Results**

122 *Presence and characterization of D1 and D2 in epididymal rat WAT*

123 We first identified D1 in WAT, using PCR amplification and gel separation (not shown).
124 Then, we did titration curves using qRT-PCR using Taqman probes (**Figure 1A**), confirming
125 that D1 and D2 mRNA were present in rat epididymal WAT being D2 mRNA abundance
126 higher than D1 mRNA (approx. double). WAT D1 mRNA abundance was less than 1% of
127 liver D1 and WAT D2 mRNA abundance was about 7% of BAT D2 (Figure 1B, 1C).

128 We then analyzed D1 and D2 activities in WAT. First, the kinetic characteristics were
129 determined; for D2: $K_m=40$ nM T4 and $V_{max}= 3$ pmol/h/mg protein and for D1: $K_m= 0.35$
130 μ M rT3 and $V_{max}= 6-18$ pmol/h/mg protein (C and Tx rats, respectively). D1 activity was
131 higher in perirenal WAT; D2 activity was similar in perirenal and epididymal WAT and
132 lower in subcutaneous WAT (results not shown).

133 *Regulation of D1 and D2 in rat WAT by thyroid status*

134 Deiodinases are regulated by thyroid status in most tissues. To study this regulation we
135 used control, Tx and T4+T3-treated rats. The thyroidal status was checked measuring T4 and
136 T3 concentrations in plasma, liver and WAT. **Figure 2** shows that T4 and T3 decreased after
137 thyroidectomy in all samples. Treatment of Tx rats with T4+T3 reverted T4 to control values

138 in plasma and liver, while WAT T4 doubled the values in control rats, pointing to a
139 preferential T4 uptake in WAT. The treatment also increased T3 in all cases, reaching control
140 values in WAT and plasma, while liver T3 concentrations did not fully recovered.

141 D1 and D2 mRNA increased in WAT from Tx rats (**Figure 1D, 1E**), and T4+T3 treatment
142 inhibited them to control values for D1, and to less than 50% of control values for D2 mRNA.
143 Opposite to the mRNA responses no variation was observed in D1 and D2 activities, except
144 for D2 activity that was inhibited in the T4+T3 group.

145

146 **Discussion**

147 D1 activity was identified in WAT in 1983 together with D2 activity in BAT [8],
148 establishing clear differences between BAT and WAT. Then, a full UCP1 expression was
149 associated to T3 produced by D2 in BAT [4] and this was recently confirmed in the D2
150 knockout mice [7].

151 Herein, we describe that D2 is also present in WAT, both at mRNA and activity levels. We
152 used epididymal WAT, the purest of all WAT locations, never converted into BAT under cold
153 exposure; in contrast, inguinal fat, is considered “convertible” adipose tissue, that is
154 transformed into BAT under several stimuli [11, 16]. Other WAT locations (perirenal,
155 periuterine) are considered a mix of BAT and WAT in terms of UCP1 induction [17]. UCP1
156 levels in epididymal WAT are near detection limits by qRT-PCR and less than 5% those in
157 inguinal WAT (unpublished). Therefore it is unlikely that D2 could come from residual BAT
158 cells present in epididymal WAT.

159 Despite the 1983 report, WAT D1 had not been further studied, possibly because lipids
160 cause technical difficulties, which we solved by improving the RNA isolation using a specific
161 kit to avoid lipids and specific Taqman probes. Regarding D1 and D2 activities, we

162 characterized them in terms of their kinetic characteristics which were in the same ranges as
163 in other tissues [1].
164 Thyroid status, a main regulator of deiodinases, was studied in this work. Both deiodinases
165 mRNAs were up-regulated in hypothyroidism. WAT D2 increased as expected in
166 hypothyroidism. WAT D1 also increased, responding to hypothyroidism as thyroid D1 [2].
167 Regarding deiodinase activities, WAT D2 followed a pattern similar to D2 mRNA, as it
168 tended to increase in hypothyroidism and decreased in T4+T3-treated rats, reflecting the high
169 WAT T4 concentrations. We showed recently increased D2 mRNA in parallel to D2 activity
170 levels in human subcutaneous fat [18]. D1 activity did not change among groups in our
171 study. Other studies have shown discrepancies between D1 and/or D2 mRNA and activity
172 levels [19], probably due to posttranscriptional modifications.

173 Recently, D1 activity in WAT was related to adiposity, increasing by high-fat diets and by
174 leptin, and decreasing under caloric restriction [20]. Other possible regulators (hormones,
175 nutrients) remain to be identified, as we have only explored the role of thyroid hormones. D1
176 and D2 may have distinct functions in WAT, regulating lipogenesis and lipolysis or the
177 expression of genes and providing the T3 required for specific functions, including the
178 conversion of WAT into BAT under specific conditions, which would increase energy
179 expenditure. Each enzyme may have specific functions.

180 *Conclusions*

181 D1 and D2 activities and mRNAs are present in rat WAT. Thyroidal status regulates D1
182 and D2 mRNA in vivo, both increasing in hypothyroid rats. No changes are found in
183 deiodinase activities, except for the D2 inhibition by T4+T3.

184

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189

190 **Authors contribution**

191 Both authors contributed equally to the study. MJO wrote the manuscript.

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194 **Legends to Figures**

195

196 **Figure 1.** Ct values for 18S rRNA, D1 and D2 mRNAs from epididymal rat WAT (panel
197 A). Relative expression of D1 (panel B) and D2 mRNAs (panel C) versus liver D1 and BAT
198 D2 mRNAs, respectively. Panels D to G show D1 and D2 mRNAs and activities in WAT
199 from control (C), thyroidectomized (Tx) and T4+T3 treated thyroidectomized rats. Values
200 are means \pm SEM. * $P < 0.05$ vs C, # $P < 0.05$ vs Tx. (n=4-5/ group).

201

202 **Figure 2.** T4 and T3 concentrations in plasma, liver and epididymal WAT from control
203 (C), thyroidectomized (Tx) and T4+T3 treated thyroidectomized rats. Values are means \pm
204 SEM. * $P < 0.05$ vs C, # $P < 0.05$ vs Tx. (n=4-5 /group)

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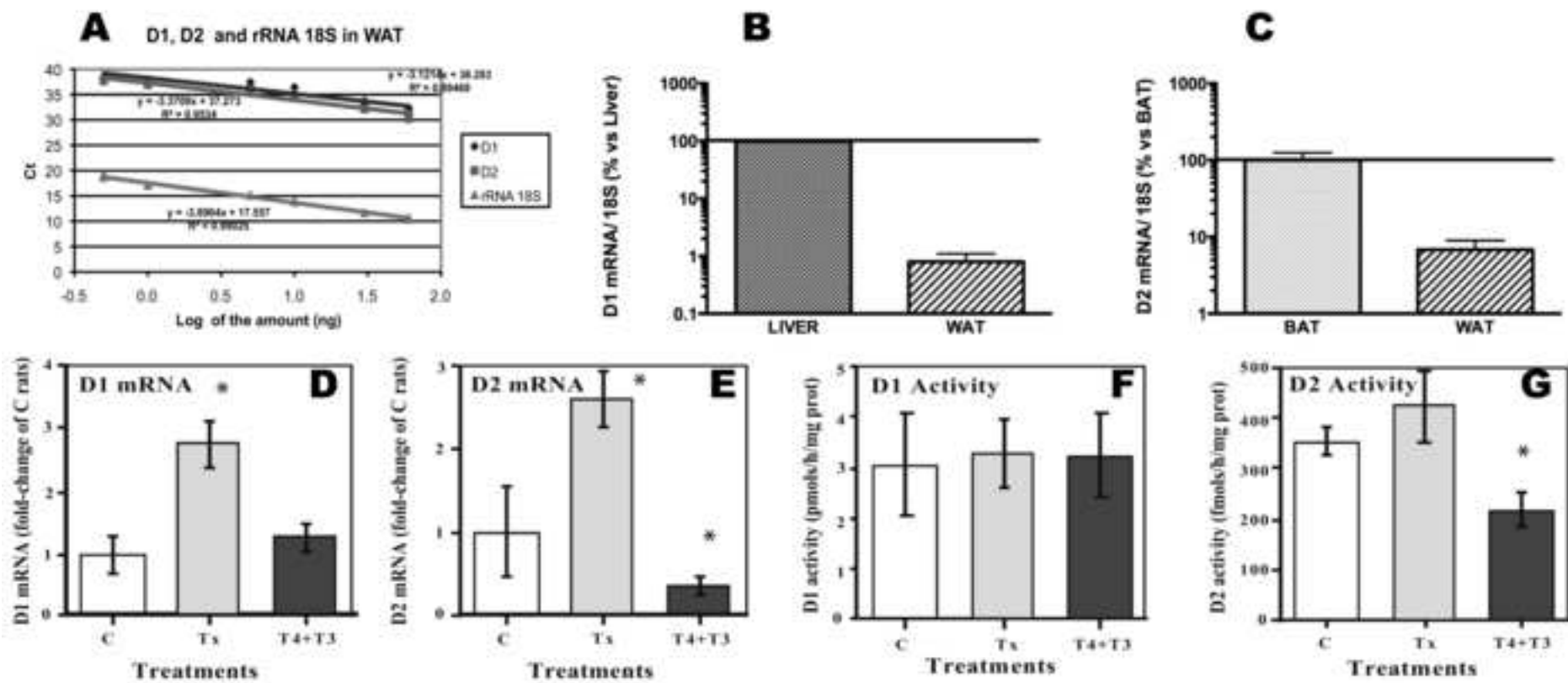
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