

SUPPLEMENTARY TABLE S1. MICROARRAY RESULTS
FOR *RXR*, *NCOA*, *DIO*, AND *DGAT* ISOFORMS
AFTER T3 TREATMENT AND WITHDRAWAL

<i>Gene</i>	<i>Group</i>	<i>FC</i> ^a	<i>Z-score</i> ^b
<i>Rxb</i>	Acute	1.17	1.12
	Chronic	1.12	0.52
	Recovery	0.99	-0.09
<i>Rrg</i>	Acute	1.00	0.03
	Chronic	1.04	0.21
	Recovery	1.00	0.01
<i>Ncoa2</i>	Acute	1.00	0.05
	Chronic	1.06	0.35
	Recovery	1.05	0.51
<i>Ncoa3</i>	Acute	0.94	-0.57
	Chronic	0.87	-0.80
	Recovery	0.92	-0.79
<i>Dio3</i>	Acute	0.96	-0.37
	Chronic	0.98	-0.72
	Recovery	0.99	-0.13
<i>Dgat1</i>	Acute	1.04	0.33
	Chronic	0.93	-0.38
	Recovery	0.97	-0.30
<i>Dgat2</i>	Acute	1.30	1.30
	Chronic	1.18	0.50
	Recovery	1.07	0.38

^aFold changes (FCs) are presented for treated mice relative to non-treated mice.

^bZ-scores were determined as described previously (9). |Z-score| >2.58 was considered statistically significant ($n = 3/\text{group}$).