

**Importance of Monocarboxylate transporter 8 (Mct8) for the Blood-Brain Barrier
Dependent Availability of 3,5,3'-Triiodo-L-Thyronine (T3)**

Short title: Thyroid hormone action in *Mct8* mutant mice

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PRECIS: The main restriction for T3 entry into the neural target cells of the mouse deficient in the thyroid hormone transporter *Mct8* is at the blood brain barrier.

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Abstract

Mutations of the gene expressing plasma membrane transporter for thyroid hormones *MCT8* (*SLC16A2*) in humans lead to altered thyroid hormone levels and a severe neurodevelopmental disorder. Genetically engineered defect of the *Mct8* gene in mice leads to similar thyroid hormone abnormalities, but no obvious impairment of brain development or function. In this work we studied the relative role of the blood-brain barrier, and the neuronal plasma cell membrane in the restricted access of T3 to the target neurons. To this end we compared the effects of low doses of T4 and T3 on cerebellar structure and gene expression in wild type (Wt) and *Mct8* null male mice (*Mct8*^{-/-}, KO) made hypothyroid during the neonatal period. We found that compared to Wt animals, T4 was considerably more potent than T3 in the *Mct8* KO mice, indicating a restricted access of T3, but not T4, to neurons after systemic administration *in vivo*. In contrast, T3 action in cultured cerebellar neurons was similar in Wt cells as in *Mct8* KO cells. The results suggest that the main restriction for T3 entry into the neural target cells of the mouse deficient in *Mct8* is at the blood-brain barrier.

Introduction

The importance of plasma membrane transporters for the transfer of thyroid hormones from the extracellular milieu to the interior of the cell is now widely recognized. For many years it was thought that thyroid hormones enter the target cells by passive or facilitated diffusion. However mutations in a specific T4 and T3 transporter, the monocarboxylate transporter 8 (*MCT8*, *SIC16A2*), were found in patients with a severe neurodevelopmental defect and abnormal levels of iodothyronines in blood, consisting of decreased T4 and rT3 and increased T3 (1, 2). These and subsequent findings revealed the physiological role of transporters in thyroid hormone action, and their relevance to the brain (3-5).

The generation of *Mct8* knock out (KO) mice demonstrated that absence of *Mct8* impairs brain thyroid hormone uptake and metabolism possibly due to a primary decreased uptake and degradation of T3 in target neurons (6, 7). As a consequence, T3 concentrations increase in serum, with stimulation of *Dio1* expression in liver and other tissues. It is postulated that the increased *Dio1* activity increases conversion of T4 to T3, thereby decreasing T4 and further increasing T3 in serum. On the other hand, circulating rT3 is also decreased, which might be due to increased degradation by *Dio1*, and / or decreased formation from T4 by inner ring deiodination.

However, while the absence of *Mct8* in mice reproduces the endocrine changes characteristic for humans with *MCT8* gene mutations, the mutant mice do not show signs of neurological impairment, which contrasts with the observations in humans. It is logical to think that the neurological syndrome is due to impaired T3 action in neurons, as a consequence of restricted uptake. However no histological changes suggestive of cerebral hypothyroidism in the mutant mice have been found, and only a moderately decreased expression of thyroid hormone regulated genes such as *Nrgn* (also known as RC3) could be related to the decreased T3 uptake (6, 7). At least in part, this could be interpreted as if the mice brains were in a state of locally

compensated hypothyroidism, since *Dio2* activity is increased in the brain, due to the decreased concentration of circulating T4 (6, 7).

Early studies on *Mct8* gene expression in rodents indicated that the gene is expressed predominantly in the choroid plexuses and in neurons (8). Recent studies have shown that *Mct8* is also expressed in the blood-brain barrier (9). Other thyroid hormone transporters are expressed in the blood-brain barrier, such as organic anion transporters and L-type aminoacid transporters (10, 11). In the absence of *Mct8* the restriction to T3 transport through the blood-brain barrier or through the neuronal plasma membrane would depend on the presence of alternative transporters.

In this work we have studied the relevance of *Mct8* gene expression in neurons for T3 action. We have analyzed the relative effects of low doses of T4 and T3 on two T3 target genes, expressed in the striatum (*Nrgn*) and the cerebellum (*Hr*, or *Hairless*). We found that in male *Mct8KO* mice, when compared with wild type (Wt) mice, these genes are less responsive to T3 than to T4, indicating a restricted entry of plasma T3, but not of T3 derived from T4. On the other hand, the action of T3 in primary cultures of cerebellar granular cells was little affected in the absence of *Mct8*. The data suggest that the critical restriction to T3 transport in the absence of *Mct8* is located at the blood-brain barrier rather than at the plasma membrane of individual neurons.

Methods

Animals

Protocols for animal handling were approved by the local institutional Animal Care Committee, and followed the rules of the European Union. Animals were housed in temperature (22 ± 2 °C) and light (12:12 light-dark cycle; lights on at 7 a.m.) controlled conditions and had free access to food and water. *Mct8KO* mice, were originally produced by Dumitrescu et al (6) by homologous recombination. Experiments were carried out on Wt (*Mct8+/y*) and KO (*Mct8-/y*) male litter mates derived from the third and fourth back crossing of heterozygous females (*Mct8-/+*)

with Wt (*Mct8^{+/-}*) males of the C57BL/6J strain. The genotype was confirmed by PCR of tail DNA (38 cycles at 55°C annealing temperature) using the following primers: Forward common: 5'-ACAACAAA AGCCAAGCATT-3'; reverse Wt specific: 3'-GAGAGCAGCGTAAGGACAAA-5'; reverse knockout specific: 3'-CTCCA AGCCTGATTTCTAT-5'. Using this procedure the Wt allele generated a 476 bp product and the null allele a 239 bp PCR product.

Induction of hypothyroidism, and thyroid hormone treatments

After crossing with Wt male mice, *Mct8^{+/-}* pregnant dams were given either drinking water or a solution containing 0.02% 1-methyl-2-mercapto-imidazol (MMI, Sigma Chemical Co, St Louis, MO) plus 1% KClO₄ *ad libitum*. These antithyroid drugs were given from gestational day 17, and throughout the lactating period, until the end of the experiment on postnatal (P) day 21. The pups were genotyped on P11, to select for *Mct8^{+/-}* and *Mct8^{-/-}* mice from the same litters. For simplicity, these animals will be referred to as wild type (Wt) and knock out (KO) mice respectively throughout this paper. The hypothyroid pups were then divided into three groups receiving no hormonal treatment, 20 ng T₄/g bw, or 3 ng T₃/g body weight respectively. The hormones were administered in PBS containing 0.1% BSA, as daily single intra peritoneal injections from P16 to P20. The following groups were thus prepared: euthyroid (n = 7) and hypothyroid (n = 6) Wt mice; euthyroid (n = 8) and hypothyroid (n = 6) KO mice; hypothyroid Wt mice treated with either T₄ (n = 5) or T₃ (n = 6), and hypothyroid KO mice treated with either T₄ (n = 6) or T₃ (n = 6). The pups were sacrificed by decapitation 24 hours after the last T₄ or T₃ injection, on P21. The striatum and cerebellum were rapidly dissected out, frozen on dry ice, and kept at -80°C until RNA isolation.

Histological methods

Examination of stained sections of the cerebellum, and *in situ* mRNA

hybridization analysis were performed on pups perfused with paraformaldehyde under anesthesia. Methods for perfusion, sectioning, staining, and *in situ* hybridization have been previously described in detail (12, 13)

Polymerase chain reaction

Total RNA was extracted using the TRIZOL reagent (Invitrogen, Carlsbad, CA). Complementary DNA was prepared from 250 ng of RNA using the High Capacity cDNA Reverse Transcription kit (Applied Biosystems, Foster City, CA). For quantitative PCR a cDNA aliquot corresponding to 5 ng of the starting RNA was used, with Taqman Assay-on-Demand primers and the Taqman Universal PCR Master Mix, No Amp Erase UNG (Applied Biosystems) on a 7900HT Fast Real-Time PCR System (Applied Biosystems). The PCR program consisted in a hot start of 95°C for 10 minutes, followed by 40 cycles of 15 seconds at 95°C and 1 minute at 60°C. PCRs were performed in triplicates, using the 18S gene as internal standard and the 2-Ct method for analysis. For quantitative assays a standard curve was generated after amplification of known amounts of specific templates for each gene including 18S, to calculate the number of mRNA copies in each sample.

Primary granular cell cultures

All media were purchased from Invitrogen. The cerebella were dissected from P6-P7 Wt and *Mct8*KO mice in Hank's balanced sodium salt solution (HBSS), without Ca²⁺ and Mg²⁺, supplemented with 1 mM Na Pyruvate and 10 mM HEPES pH 7.4. The tissue was disaggregated by passing through a 0.9 mm syringe, rinsed in HBSS/Pyruvate/HEPES and resuspended in serum-free culture medium (Neurobasal medium supplemented with 2% B27, 0.5 mM Glutamine, 10 U/ml penicillin, and 10 U/ml streptomycin), before seeding on poly-L-ornithine (Sigma) coated 12-well multiwells (Sigma; 2.5x10⁵ cells/well). After 4 days, the granular cells were incubated for 24h in the absence or presence of T₃ (Sigma) (from 0.2 to 5 nM), in the same medium containing 0.1% newborn calf serum deprived of thyroid hormones. Astrocyte

contamination of the cultures was 3% as determined by immunofluorescence. Cells plated on glass cover slips were fixed with 4% paraformaldehyde for 5 min and permeabilized with 0.2% Triton-X100 in PBS for 5 min, and then with methanol at -20°C for 2 min. After blocking with 5% non immune serum (Vector Laboratories, Burlingame, CA), the cells were doubly stained by overnight incubation at 4°C with the following combination of primary antibodies diluted 1/2000: rabbit polyclonal anti-GFAP (Dako, Glostrup, DK) for astrocytes, and mouse monoclonal anti-NeuN (Chemicon, Temecula, CA) for neurons. Nuclei were labeled with the nuclear stain 4', 6-diamidino-2-phenylindole dihydrochloride (DAPI).

Statistical calculations

Differences between means were obtained by 2-way ANOVA, with the two factors being genotype and thyroidal state. As post-hoc test we used the Bonferroni tests using the Graph-Pad Prism software (<http://www.graphpad.com/prism/>).

Results

In this study we have examined the relative effects of low doses of T4, and T3, administered to hypothyroid Wt and Mct8-deficient mice. The goal of this study was to evaluate the relative role of Mct8 in the transport of T4 and T3 in the brain in vivo, by determining the effects of the hormones on cerebellar structure and on the expression of thyroid hormone-regulated genes. Preliminary studies using morphological techniques failed to reveal consistent differences between age-matched, Wt and KO mice during development that could be related to deficient thyroid hormone transport into the brain. Although not shown in this paper, we examined the laminar structure of the cerebral cortex, myelin protein expression, maturation of glial cells and of different classes of interneurons and Purkinje cells, and number of interneuron precursors in the cerebellum, and found no consistent deficits in the KO mice. These observations agree with previous studies reporting no obvious phenotype of cerebral hypothyroidism in these mice (6, 7).

The next question we addressed is whether the absence of the Mct8 transporter impaired the biological activities of exogenous T4 and T3 selectively. Wt and KO mice were made hypothyroid by administration of antithyroid drugs, and received 20 ng/g T4 or 3 ng/g T3 daily for 5 days before sacrifice on P21. This dosage schedule was sufficient to completely correct the delayed migration of granular cells in the cerebellum of hypothyroid mice, as shown in Fig. 1: Cerebellar sections from each group of mice were stained and examined by optical microscopy. Migration of granular cells was already completed in the euthyroid Wt animals by P21, so that the external germinal layer (EGL) was absent. As expected, the EGL was still present in the hypothyroid Wt mice at this age. Both T4 and T3 treatments were equally able to prevent the effects of hypothyroidism in the Wt mice.

In the absence of Mct8, the structure of the cerebellar cortex in the untreated KO mice was identical to that of the Wt mice, with no EGL remaining, illustrating the lack of morphological developmental abnormalities. As in the hypothyroid Wt mice, the EGL was still present in the hypothyroid KO P21 mice. T4 treatment prevented the effects of hypothyroidism. However, in contrast to the effect on Wt mice, T3 treatment did not correct the migration abnormality.

To examine the effects of T4, and T3 on gene expression, two well known thyroid hormone target genes, *Nrgn* in the striatum, and *Hr* in the cerebellum, were examined by real time PCR. The results are shown in Fig 2. Interestingly, and despite the lack of morphological impairment, *Hr* expression was decreased in the cerebellum of untreated KO mice with respect to the Wt mice, with levels similar to those present in WT hypothyroid mice. Induction of hypothyroidism decreased *Hr* expression further in the KO mice. T4 treatment significantly increased *Hr* expression in the hypothyroid Wt mice to levels that were similar to the untreated Wt mice. The effect of T4 treatment in the hypothyroid KO mice was not different from the hypothyroid Wt mice. The response to T3 was similar to that of T4 in the Wt mice, but was significantly different when the

hypothyroid Wt and KO mice were compared. T3 was without effect in the hypothyroid KO mice.

Nrgn expression was also lower in the untreated KO than in the Wt mice, and decreased further with hypothyroidism. In contrast to *Hr*, the hypothyroid Wt and KO mice had similar *Nrgn* mRNA levels. Although neither T4 nor T3 treatment were able to fully normalize *Nrgn* expression, again there was a significant difference between the responses to T3 between the Wt and the KO mice, but no difference in the responses to T4.

The liver mRNA *Gpd2* (encoding mitochondrial alpha-glycerol phosphate dehydrogenase) was increased in the untreated KO mice relative to the untreated Wt mice. Hypothyroidism decreased *Gpd2* expression in both genotypes ($P < 0.001$). T4 and T3 significantly increased the *Gpd2* mRNA level compared to hypothyroid mice ($P < 0.05$). This increase was of similar magnitude in both genotypes and with both hormones.

To correlate the effects of thyroid hormones on *Hr* and *Nrgn* gene expression with *Mct8* gene expression we performed *in situ* mRNA hybridization (Fig 3). The *Mct8* gene was heavily expressed in the choroid plexus (panels A-C) and in the ependymal lining of the third ventricle (panel B). Other sites of expression were the upper layers of the cerebral cortex, especially the cingulate, visceral, and piriform cortices, the pyramidal and granular layers of the hippocampus, and the amygdala (panel B). In the cerebellum, besides expression in the choroid plexus, *Mct8* mRNA had low but detectable abundance in the cerebellar cortex. In the striatum *Mct8* was poorly expressed (panel A). Panel D shows *Nrgn* mRNA which is abundantly present following a lateral-medial gradient, contrasting with the poor expression of *Mct8*. The effect of *Mct8* gene deletion (panel E) did not affect the *Nrgn* mRNA signal gradient, in contrast to the effect of hypothyroidism (not shown, but see (14)) which results in a total suppression of the gradient. Interestingly, *Dio2* mRNA distribution in the striatum (panel F) also followed a similar gradient, with no changes in the pattern of distribution in the *Mct8* KO mice (not shown). The lack of

correlation between the sites of expression of *Mct8* with that of the T3-target genes, *Nrgn* and *Hr*, indicate that *Mct8* might be playing a minor role in thyroid hormone transport through the plasma membrane of cerebellar granular cells, and of striatal neurons *in vivo*.

To address this question more directly, we studied transporter expression and *Hr* induction by T3 in primary cultures of neurons. Granular cells from newborn mice cerebella were cultured. To analyze the effect of *Mct8* deficit on the effect of T3 on *Hr* gene induction, T3 was added to granular cells from Wt and KO mice and *Hr* mRNA was measured by quantitative PCR. One representative experiment using different concentrations of T3 is shown in figure 4. Starting at the lowest concentration used, 0.2 nM, all T3 concentrations of T3 gave a significant stimulation of *Hr* expression ($P < 0.001$) both in the Wt and the KO cells. There were no significant differences in the effect of T3 in the KO mice as compared with the Wt except for the 1.25 nM T3 concentration in this particular experiment.

We also examined the profile of transporter expression in the same cultures used to analyze the effect of T3 on *Hr*. We measured the amounts of mRNA of *Oatp2* (*Slco1a4*), *Oatp14* (*Slc1c1*), and *Mct8* (table 1). Granular cells from Wt cells expressed predominantly *Mct8* (591 ± 130 mRNA copies, relative to 18S RNA), which was undetectable in the KO cells. *Oatp2* and *Oatp14* were expressed at much lower levels (27.0 ± 9.1 and 7.4 ± 3.4 , respectively). There were no changes in the KO compared to the Wt, and T3 treatment had no effects on transporter expression, except for the higher dose, that decreased *Oatp14* mRNA in the KO mice.

Discussion

The main finding of the present work is that the brain of animals lacking the thyroid hormone transporter *Mct8* do not readily respond to a low dose of T3, while the sensitivity to an equally low dose of T4 is similar to that of Wt animals. Five ng/g T3 were previously shown to normalize circulating TSH and hypothalamic TRH transcripts in hypothyroid Wt, but not in hypothyroid *Mct8KO* mice (7). We used

low doses of T4 and T3, to avoid inhibition of Dio2 by T4, and the use of low affinity transporters by T3. Although, based on *Gpd2* gene expression, the doses used did not fully restore euthyroidism in the Wt mice liver, they were sufficient to normalize cerebellar development, EGL migration and *Hr* expression in the Wt hypothyroid mice. In contrast, they were insufficient for Nrgn mRNA normalization.

In normal animals, T3 reaches the extracellular fluid of the brain parenchyma from the circulation through the blood-brain barrier, and acts directly on the neurons. T4 may exert some extranuclear actions, but the bulk of genomic responses are mediated by its conversion to T3 by Dio2. In the brain, this reaction takes place predominantly in glial cells, namely astrocytes, and IV ventricle tanocytes, although *Dio2* expression has also been observed in some cerebral cortex interneurons as a response to hypothyroidism (13, 15).

In the developing cerebellum, T4 to T3 conversion takes place in the protoplasmic astrocytes located within the granular layer in close association to the granular cells. Since there was no difference in the effects of T4 in KO versus Wt mice, the results suggest that, in the doses used, T4 could reach the cerebellar astrocytes of *Mct8*-deficient mice in sufficient amount to produce the effects observed in Wt mice. This event is further facilitated by the great increase of Dio2 activity in the brain of *Mct8* KO mice (6,7). In addition, the results also suggest that the T3 produced in astrocytes can access the granular cell nuclear receptors with little restriction at the neuronal cell membrane.

The effects of T3 on granular cells in culture agree with the above conclusion. Trajkovic et al (7) also showed that T3 was effective in inducing Purkinje cell differentiation in vitro in the presence or absence of *Mct8*. In the context of these findings it was surprising that in isolated granular cells the *Mct8* gene was by far the more abundantly expressed transporter. However its absence in *Mct8*-/- cells caused only a minimal impairment of T3 action at the nuclear level, as evidenced by *Hr* gene expression, with a trend toward a lower effect at intermediate doses in the KO cells.

Although expression of other transporters was much lower, it was enough to elicit almost identical responses to T3 in the absence as in the presence of *Mct8*. The presence of other transporters is also likely the cause for the similar effect of T4 in vivo in Wt and in KO mice. The relative effects of T4 and T3 on the expression of the *Nrgn* gene in the striatum suggests a similar conclusion.

These results agree with the preferential accumulation of administered T4, relative to the restricted accumulation of administered T3 in the brain of *Mct8* KO mice (6, 7). Brain T3 concentrations in the KO mice were about 2/3 of normal. Given the restriction to T3 entry, most T3 in the brain of these animals must be derived from T4.

The main site of *Mct8* expression is the choroid plexus. The consequences of the absence of *Mct8* in the choroid plexus are not known. Intrathecally administered T4 and T3 can access brain structures (16). However most studies on the routes of thyroid hormone entry to the brain agree that the cerebrospinal fluid allows only limited access of thyroid hormone to the brain parenchyma, preferentially reaching cells located near the surface of the ventricles (17, 18). Therefore the main access of thyroid hormone to the brain parenchyma is through the blood-brain barrier. In keeping with this concept, our data suggest that the restriction of T3 entry in *Mct8*-deficient mice is at the blood-brain barrier. Indeed *Mct8* has been recently demonstrated in the membrane of the brain parenchyma capillaries (9). The presence of other transporters such as Oatp14 and Oatp2, with more affinity for T4 than for T3 may explain the different sensitivities to T4 and T3. While normal T4 uptake may preserve the compensated phenotype in mice, the lack of alternative transporters in the human BBB would be the reason for the neurological impairment (9).

In conclusion, the data show that the main restriction to T3 action in the absence of *Mct8* is at the level of the blood-brain barrier. The thyroid hormone transport role of *Mct8* in the plasma membrane of neurons, at least in the striatum and the cerebellum seems to be minimal.

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Figures

Fig 1: Effects of hypothyroidism and thyroid hormone treatment on the structure of the cerebellar cortex of Wt and KO mice. The figure shows photomicrographs of lobule 7 from toluidine-stained sagittal slices of P21 mice. Igl: internal granular layer; Mol: molecular layer; the arrows show the external germinal layer. Scale bar, 100 microns.

Fig 2: Influence of *Mct8* deficit on the response to T4 and T3. Wt and KO mice were made hypothyroid from the late fetal period up to P21, and treated with either, vehicle, or with T4 or T3 for five days before sacrifice. Data are means \pm SD. Hairless (*Hr*) and RC3/Neurogranin (*Nrgn*) expression was analyzed by real time PCR in the cerebellum and striatum, respectively. The data were analyzed by two-way ANOVA and the Bonferroni post test to compare the data from each KO condition with the corresponding Wt (** = $P < 0.01$; *** = $P < 0.001$). For *Hr* expression there was a significant effect of genotype ($F_{1,41} = 116.01$, $P < 0.0001$), and thyroidal status ($F_{3,41} = 37.84$, $P < 0.0001$), with a significant interaction ($F_{3,41} = 3.88$, $P = 0.0156$). Also for *Nrgn* there was a significant effect of genotype ($F_{1,35} = 24.68$, $P < 0.0001$), and thyroidal status ($F_{3,35} = 41.16$, $P < 0.0001$), with a significant interaction ($F_{3,35} = 10.30$, $P < 0.0001$). The lower panel shows *Gpd2* (mitochondrial alpha-glycerol phosphate dehydrogenase) expression in the liver. There was a significant effect of thyroidal status ($F_{3,41} = 23.67$, $P < 0.0001$), but not genotype ($F_{3,41} = 1.065$, $P = 0.308$), with a significant interaction ($F_{3,41} = 3.936$, $P < 0.0148$).

Fig 3: ^{35}S in situ hybridization for *Mct8* (A-C), *Nrgn* (D,E) and *Dio2* (F) mRNAs. The slices are from coronal sections at the level of the caudate (A, B, D-F) and sagittal section of the cerebellum (C). All slices are from P21 Wt mice except for panel E which shows the typical *Nrgn* expression in a P21 KO mouse. The arrows in panels A-C show heavy *Mct8* expression in the choroid plexus. The asterisks show the caudate nucleus in A, with low hybridization signal, and the faint, but detectable hybridization in the cerebellar cortex in C.

Fig 4. Hairless (*Hr*) expression in primary cultures of granular cells from Wt mice or *Mct8*-deficient mice, as a function of T3 added to the cultures. Differences in *Hr* expression between the cells without T3 added and the 0.2 nM T3 concentration were $P < 0.001$ (a). Differences between the Wt cells and the KO cells at each T3 concentration, by two-way ANOVA were NS, except for the 0.25 nM T3, with $P < 0.05$ (b).

Table 1: Effect of T3 treatment on transporter expression in cultured granular cells

Transporter mRNA	No T3		2.5 nM T3		5.0 nM T3	
	Wt	KO	Wt	KO	Wt	KO
Mct8	591 ± 130	-	596 ± 48	-	621 ± 109	-
Oatp2	27.0 ± 9.1	28.0 ± 8.5	29.3 ± 5.4	23.9 ± 6.1	28.3 ± 1.6	19.5 ± 4.7
Oatp14	7.4 ± 3.4	6.8 ± 2.4	4.9 ± 0.9	2.8 ± 0.4	5.9 ± 1.7	2.5 ± 0.3*

Primary cultures of granular cells from the cerebella of wild type (Wt) and Mct8 KO mice were incubated in the presence of 0, 0.2, 0.5, 1.25, 2.5, and 5.0 nM T3 for 24 hours. Expression of *Mct8* (*Slc16a2*), *Oatp2* (*Slc1a4*), and *Oatp14* (*Slc1c1*) was quantified by real time PCR using TaqMan probes. Shown are the data (mean number of RNA copies relative to 18S RNA ± SD) from cells incubated without added T3, or in the presence of 2.5 and 5.0 nM only. Mct8 mRNA was not detected in the KO cells. The cells used in this experiment are the same as for Hr mRNA quantification shown in figure 4. Two-way ANOVA using the data from all T3 concentrations revealed that there was no difference of genotype or treatment, except for the highest T3 dose that decreased Oatp14 mRNA in the KO cells. *: P < 0.05.

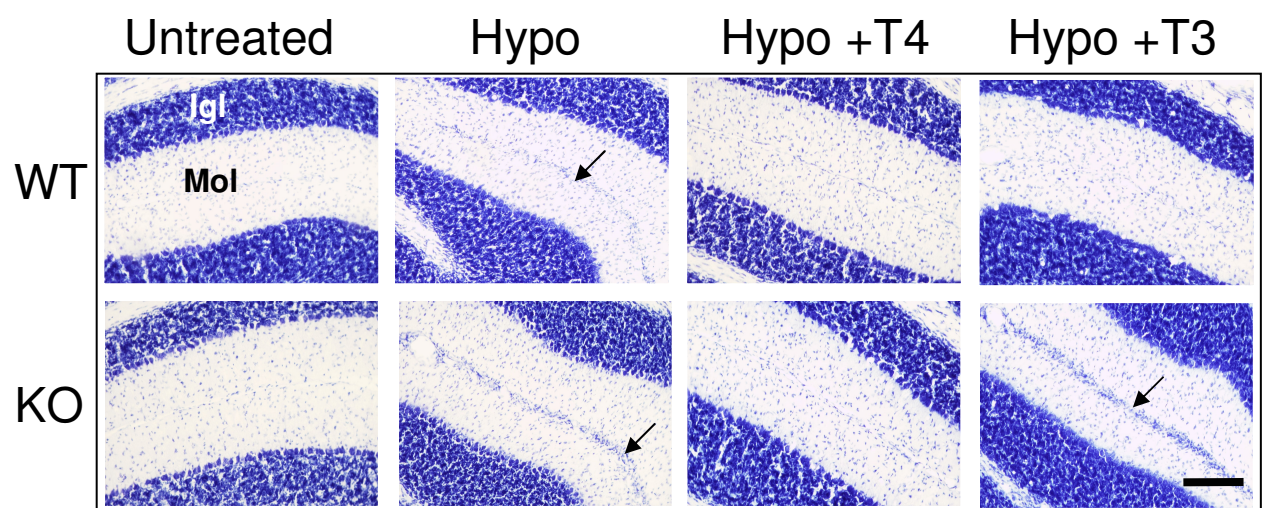


Figure 1

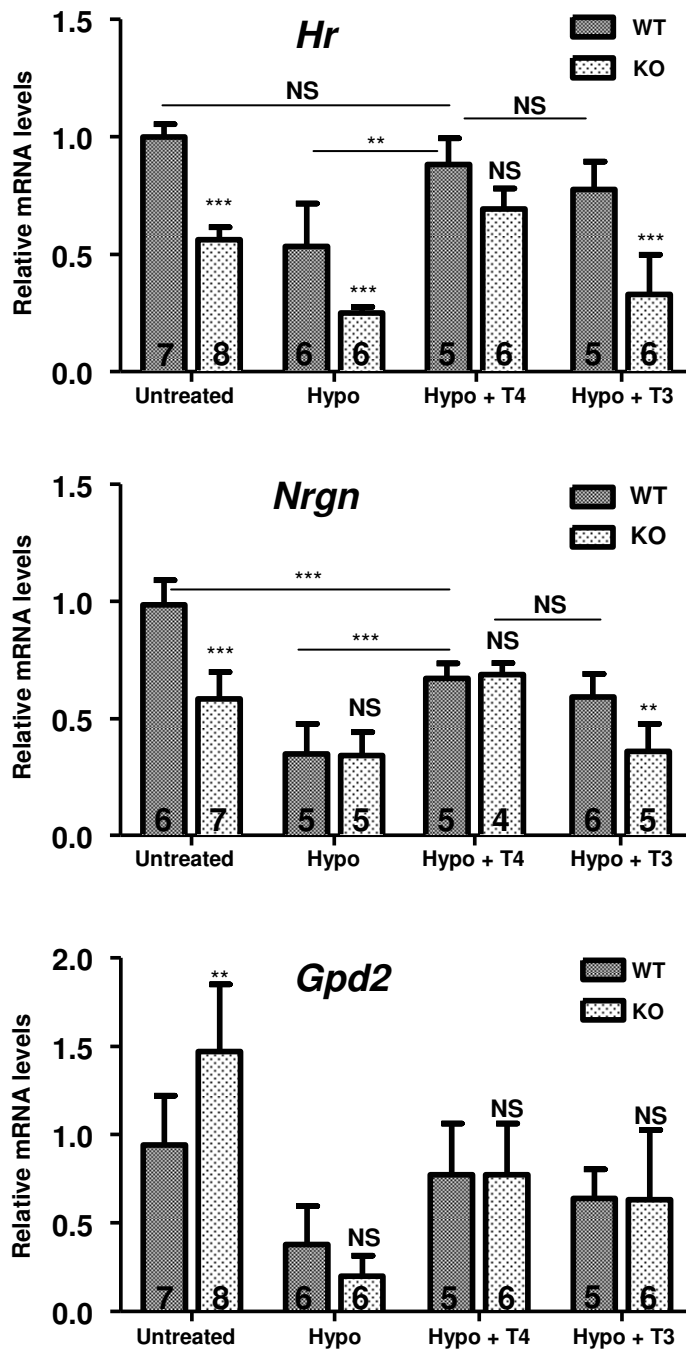


Figure 2

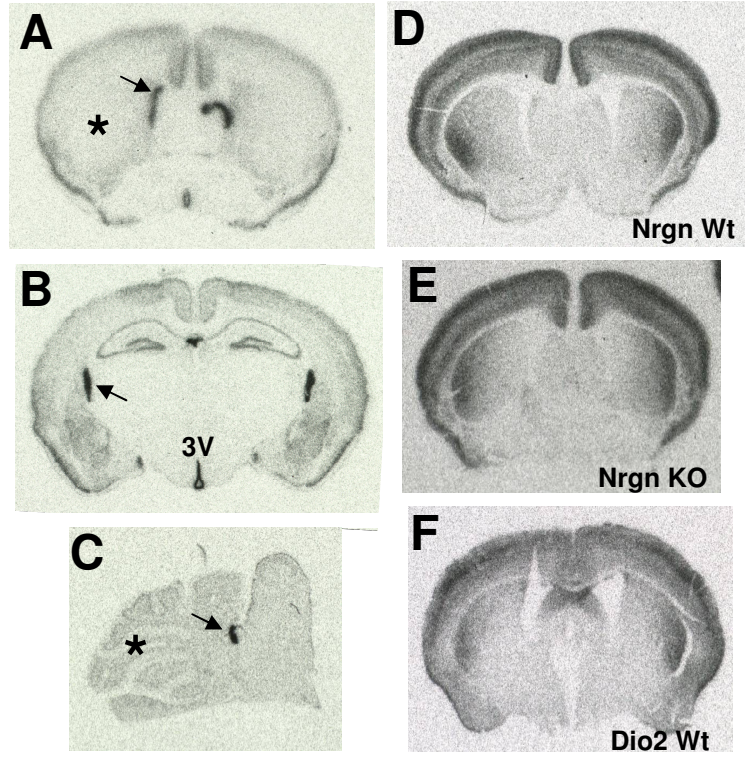


Figure 3

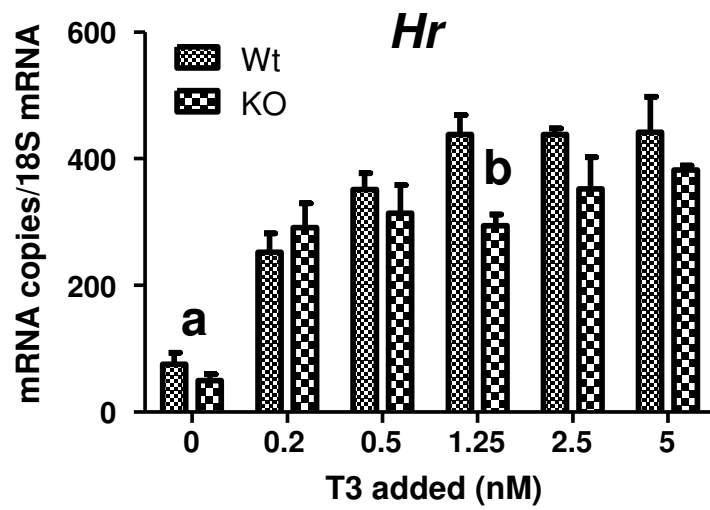


Figure 4