

Changes in Neuropeptide Y Receptors and Pro-Opiomelanocortin in the Anorexia (*anx/anx*) Mouse Hypothalamus

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The pro-opiomelanocortinergic (POMCergic) system originating in the hypothalamic arcuate nucleus extends projections widely over the brain and has been shown to be intricately linked and parallel to the arcuate neuropeptide Y (NPY) system. Both NPY and POMC-derived peptides (melanocortins) have been strongly implicated in the control of feeding behavior, with the former exerting orexigenic effects and the latter having anorexigenic properties. Mice homozygous for the lethal anorexia (*anx*) mutation are hypophagic, emaciated, and exhibit anomalous processing of NPY exclusively in the arcuate nucleus, providing an interesting model to study NPY–POMC interactions. In the present study, several morphological markers were used to investigate the histochemistry and morphology of the POMC system in *anx/anx* mice. *In situ* hybridization demonstrated decreased numbers of POMC mRNA-expressing neurons in the *anx/anx* arcuate nucleus. In parallel, mRNA levels for both the

NPY Y1 and Y5 receptors, which are expressed in POMC neurons, were decreased. Also, expression of the NPY Y2 autoreceptor was attenuated. Immunohistochemistry using antibodies against adrenocorticotrophic hormone to demonstrate POMC cell bodies, against α -melanocyte-stimulating hormone to demonstrate axonal projections and against the NPY Y1 receptor to demonstrate dendritic arborizations, showed strikingly decreased immunoreactivities for all these markers. The present data suggest that degeneration of the arcuate POMC system is a feature characteristic of the *anx/anx* mouse. The possible relationship to the NPYergic phenotype of this animal is discussed.

Key words: adrenocorticotrophic hormone; α -melanocyte-stimulating hormone; arcuate nucleus; feeding; immunohistochemistry; *in situ* hybridization; paraventricular nucleus of the hypothalamus; trophism

The neurochemical basis for the central control of food intake has been illuminated by recent investigations (Leibowitz, 1995; Sawchenko, 1998; Woods et al., 1998; Elmquist et al., 1999; Kalra et al., 1999). Two messenger molecules expressed in discrete populations of the hypothalamic arcuate nucleus appear to play antagonistic roles in energy balance control, namely neuropeptide Y (NPY) (Tatemoto, 1982; Tatemoto et al., 1982) and pro-opiomelanocortin (POMC) (Mains et al., 1977; Roberts and Herbert, 1977; Nakanishi et al., 1979), the precursor protein for the melanocortin peptides adrenocorticotrophic hormone (ACTH) and α -melanocyte-stimulating hormone (α -MSH). Intracerebral injections of NPY stimulate food intake (Clark et al., 1984; Stanley and Leibowitz, 1985), whereas melanocortin treatment induces anorexia (Poggioli et al., 1986; Fan et al., 1997; Grill et al., 1998), and mice deficient for the melanocortin-4 receptor develop obesity (Huszar et al., 1997). Furthermore, fasting increases arcuate NPY mRNA levels and decreases POMC mRNA (Brady et al., 1990).

The projections of the arcuate NPY and POMC neurons follow seemingly parallel routes (Broberger et al., 1998b) and may terminate on common postsynaptic targets. NPY neurons also innervate the POMC cell bodies, which express the inhibitory NPY Y1 receptor (Y1R) (Csiffáry et al., 1990; Zhang et al., 1994; Broberger et al., 1997b; Fuxe et al., 1997). A similar expression pattern has been suggested for the Y5 receptor (Y5R) (Gerald et al., 1996; Naveilhan et al., 1998a). Activation of the Y1R and the Y5R (Flood and Morley, 1989; Stanley et al., 1992; Gerald et al., 1996) stimulates food intake, possibly through inhibition of POMC signaling. In addition, NPY neurons also express agouti gene-related protein (AGRP) (Shutter et al., 1997; Broberger et al., 1998a,b; Hahn et al., 1998), an antagonist of the melanocortins (Ollmann et al., 1997). Thus, both anatomical and pharmacological evidence suggest that the balance between NPY and POMC transmission may be correlated to the level of food intake.

The anorexia (*anx*) mutation is a recessive autosomal mutation in mice that phenotypically results in decreased food intake, emaciation, and premature death during the fourth postnatal week (Maltais et al., 1984). A possible explanation for this phenotype was provided by the recent demonstration of NPY abnormalities in *anx/anx* mice that appear to be restricted to the arcuate system, characterized by a decrease in the number of its NPY-immunoreactive (-IR) nerve terminals and a strongly increased perikaryal staining in the arcuate nucleus, a pattern suggestive of cell body accumulation and impaired transport and/or release (Broberger et al., 1997a). The same pattern was also noted with AGRP histochemistry (Broberger et al., 1998b). In accordance with the orexigenic function of arcuate NPY neurons described

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above, diminished NPY-AGRP signaling could be causally related to the anorectic phenotype of *anx/anx* mice.

The present study investigated the histochemistry of POMC neurons in *anx/anx* mice, using several markers expressed by this population, to gain understanding of how the melanocortinergic system functions in a potentially “hypoNPYergic” state. These markers include mRNAs for POMC, the Y1R and the Y5R studied by *in situ* hybridization, as well as ACTH, α -MSH, and the Y1R studied by immunohistochemistry. We have also investigated the expression of the autoinhibitory Y2 receptor coexpressed with NPY in arcuate neurons (Broberger et al., 1997b) in *anx/anx* mice.

Part of these results have been presented previously in abstract form (Broberger et al., 1998c).

MATERIALS AND METHODS

Animals. Animals were housed together in ventilated cages in an animal room maintained on a 12 hr light/dark schedule with lights on at 7.00 A.M., at a temperature of +25°C. The *anx/anx* mice were characterized genotypically using nearby located simple sequence length polymorphisms and phenotypically on the basis of body weight and neurological abnormalities determined before dissection–perfusion. Both male and female mice were analyzed. For *in situ* hybridization, 10 *anx/anx* mice and eight wild-type littermates were used. For immunohistochemistry, six *anx/anx* mice and seven wild-type littermates were used. Dissection was performed on postnatal days 20 or 21. For *in situ* hybridization experiments, brains were dissected from decapitated animals and rapidly frozen. For immunohistochemistry, mice were anesthetized with 0.1 ml of sodium pentobarbital (Mebumal; 6 mg/ml, i.p.; Apoteksbolaget, Umeå, Sweden) and perfused via the ascending aorta with 10 ml of Ca²⁺-free Tyrode's solution (37°C), followed by 10 ml of a mixture of formalin and picric acid (4% paraformaldehyde and 0.4% picric acid in 0.16 M phosphate buffer, pH 6.9, 37°C) according to Zamboni and de Martino (1967), followed by 50 ml of ice-cold fixative (as above). The brains were rapidly dissected out, immersed in the same fixative for 3 hr, and rinsed for at least 24 hr in 0.1 M phosphate buffer, pH 7.4, containing 10% sucrose, 0.02% bacitracin (Sigma, St. Louis, MO), and 0.01% sodium azide (Merck, Darmstadt, Germany).

In situ hybridization. Brains from all animal groups were mounted together on the same chuck, frozen, and coronally sectioned at 14 μ m in a cryostat (Microm, Heidelberg, Germany). The sections were thaw-mounted onto ProbeOn slides (Fisher Scientific, Pittsburgh, PA) and then stored at –20°C until processing. Oligonucleotide probes complementary to nucleotides 266–319 of the rat POMC mRNA (Drouin and Goodman, 1980), nucleotides 546–586 of the rat Y1R (Eva et al., 1990), nucleotides 737–775, 1040–1081, and 1166–1203 of the mouse Y2R (Nakamura et al., 1996), and nucleotides 755–802, 898–945, and 1176–1223 of the mouse Y5R (Nakamura et al., 1997) were synthesized (Scandinavian Gene Synthesis, Köping, Sweden), 3' end-labeled with ³⁵S- α -dATP (NEN, Boston, MA) using terminal deoxynucleotidyl transferase (Amersham, Buckinghamshire, UK), and purified using Qiaquick Nucleotide Removal Kit (Qiagen, Hilden, Germany). The probes had specific activities ranging between 2.5–3.8 $\times 10^6$ cpm/ng oligonucleotide. *In situ* hybridization was performed essentially as described previously (Schalling et al., 1988; Young, 1990; Dagerlind et al., 1992). In brief, tissue sections were air-dried and incubated for 16 hr at 42° with 10⁶ cpm of the labeled probe in a hybridization solution containing 50% deionized formamide (Baker, Deventer, The Netherlands), 4 \times SSC (1 \times SSC: 0.15 M NaCl and 0.015 M Na sodium citrate), 1 \times Denhardt's solution, 0.02% bovine serum albumin, 0.02% Ficoll (Pharmacia, Uppsala, Sweden), 0.02% polyvinylpyrrolidone (Sigma), 0.02 M NaPO₄ buffer, pH 7.0, 1% N-lauroylsarcosine, 10% dextran sulfate (Pharmacia), 500 μ g/ml denatured salmon testis DNA (Sigma), and 200 mM dithiothreitol (LKB, Stockholm, Sweden). After hybridization, the sections were rinsed in 1 \times SSC at 55°C for 60 min, including four changes of SSC, and for 60 min at room temperature, transferred through distilled water, dehydrated through 60 and 95% ethanol (15 sec each), and apposed to β -max autoradiography film (Amersham) at –20°C. After 1 day (POMC probe) or 5 weeks (Y1R, Y2R, and Y5R probes) of exposure, the films were developed with Kodak LX 24 (Eastman Kodak, Rochester, NY) for 4 min and fixed for 10 min in Kodak AL4. In addition to film autoradiography, sections were dipped in Kodak NTB2 autoradiography emulsion

in distilled water, exposed for 1 (POMC probe) or 8 (Y1R, Y2R, and Y5R probes) weeks in light-free metal containers at –20°C, developed in Kodak D19 for 3 min, and fixed in Kodak 3000 for 6–7 min. Sections were rinsed in distilled water, counterstained with cresyl violet, and coverslipped with Entellan (Merck). All sections were examined under bright- and dark-field illumination using a Nikon (Tokyo, Japan) Microphot-FX microscope. Photomicrographs were taken with Kodak T-max 100 film.

Quantification of autoradiograms. Measurements were performed on a MacIntosh IIX (Apple Computer Inc., Cupertino, CA) equipped with a Quick Capture frame grabber card (Data Translation, Marlboro, MA), a Northern Light precision illuminator (Imaging Research, St. Catharines, Ontario, Canada), and a Dage-MTI CCD-72 series camera (Dage-MTI, Michigan City, IN) equipped with a Nikon 55 mm lens. To process images, NIH Image software (courtesy of Wayne Rasband, National Institute of Mental Health, Bethesda, MD) was used. Each image was an average of eight video frames digitized to a 512 \times 512 matrix with 256 gray levels for each picture element. To adjust for possible defect in the illumination or optical pathway, an image of the empty illumination screen was taken through a neutral filter and used for background shading correction. The gray levels of eight ¹⁴C-plastic standards (Amersham) (Miller, 1991) exposed to the autoradiography film together with the tissue sections were determined and used in a third-degree polynomial approximation to construct a gray level to activity transfer function. The borders of the measuring fields were interactively defined, and the average gray level was calculated using the Macintosh computer and Microsoft Excel (Apple). The average gray level of control tissue was set to 100%, and changes in gray levels in the arcuate nucleus from sections of *anx/anx* mice were expressed as percentage of controls. To quantify the relative numbers of POMC mRNA-expressing neurons, one section from each animal at the level of bregma –1.7 mm was chosen. Under bright-field illumination combined with epi-illumination, the total number of labeled cells in the arcuate nucleus on both sides of the third ventricle was counted. Cells were considered labeled when the number of silver grains overlying the cytoplasm exceeded five times the background levels.

Immunohistochemistry. Coronal brain sections of 14 μ m thickness were cut on a cryostat (Microm) and thaw-mounted onto gelatin-coated glass slides. The sections were then processed according to the catalyzed reporter deposition method, by incubating overnight with primary polyclonal antisera to α -MSH (1:400; Peninsula, Belmont, CA) or Y1R (1:2000; Broberger et al., 1997), both raised in rabbits, or ACTH (1:400; Peninsula), raised in guinea pig, at 4°C, rinsing in PBS, followed by catalyzed reporter deposition using the Renaissance kit (DuPont NEN, Boston, MA). After 10 min rinsing in TNT buffer (0.1 M Tris-HCl, pH 7.5, 0.15 M NaCl, and 0.05% Tween 20), the sections were preincubated for 30 min at room temperature in TNB buffer (0.1 M Tris-HCl, pH 7.5, 0.15 M NaCl, and 0.05% DuPont Blocking Reagent) and incubated for 30 min at room temperature with horseradish peroxidase-conjugated swine anti-rabbit immunoglobulins (Dako A/S, Copenhagen, Denmark) for α -MSH and Y1R antisera or horseradish peroxidase-conjugated rabbit anti-guinea pig immunoglobulins (Dako A/S) for ACTH antiserum, diluted at 1:100 in TNT buffer. The sections were washed three times for 10 min, twice in Tris buffer (0.1 M Tris-HCl, pH 7.5, and 0.15 M NaCl) and once in TNT buffer. The tyramide signal amplification involved deposition of biotinylated tyramide and was performed by incubating the sections for 10 min at room temperature with a 1:50 dilution of biotinyl tyramide (NEN) in 1 \times amplification diluent (Renaissance kit). After three washes, in Tris buffer (twice) and TNT buffer (once), the chromogenic detection of the deposited biotin was performed by incubation for 30 min at room temperature with fluorescein-conjugated streptavidin (Amersham) diluted at 1:200 in TNB buffer. Finally, the sections were rinsed extensively in Tris buffer and then mounted in a mixture of glycerol and PBS (3:1) containing 0.1% *p*-phenylenediamine (Sigma) (Johnson and Nogueira Araujo, 1981; Platt and Michael, 1983).

After processing, the sections were examined in a Nikon fluorescence microscope equipped with an oil dark-field condenser, a KP500 excitation filter, and an LP 520 stop filter. Photographs were taken on black-and-white Kodak Tri-X film (Kodak) or color film (Kodak Ektachrome 400 X). For confocal microscopy, samples were recorded using a Zeiss (Oberkochen, Germany) LSM410 with a 63 \times /1.4 NA objective using 488 nm excitation. Approximately 30 optical slices of 0.5 μ m thickness were sampled and combined to produce composite confocal images, which were printed on a Codonics NP-1600 photographic network printer (Codonics Inc., Middleburg Heights, OH).

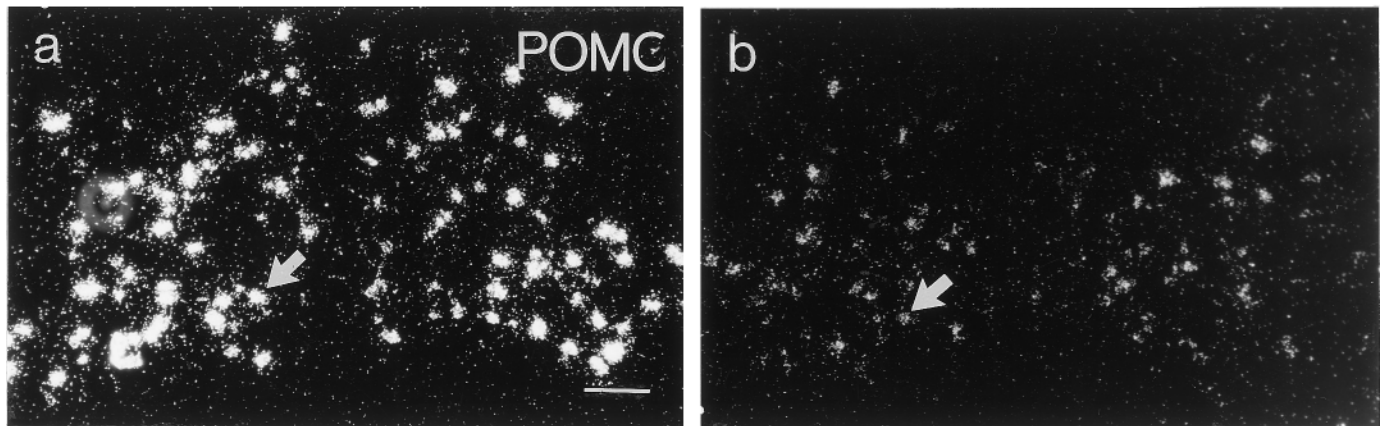


Figure 1. Dark-field micrograph of the arcuate nucleus from a wild-type (*a*) and an *anx/anx* (*b*) mouse after hybridization for POMC mRNA. Note fewer silver grains over individual cell profiles (*arrows*) in the mutant animal. Scale bar, 50 μ m.

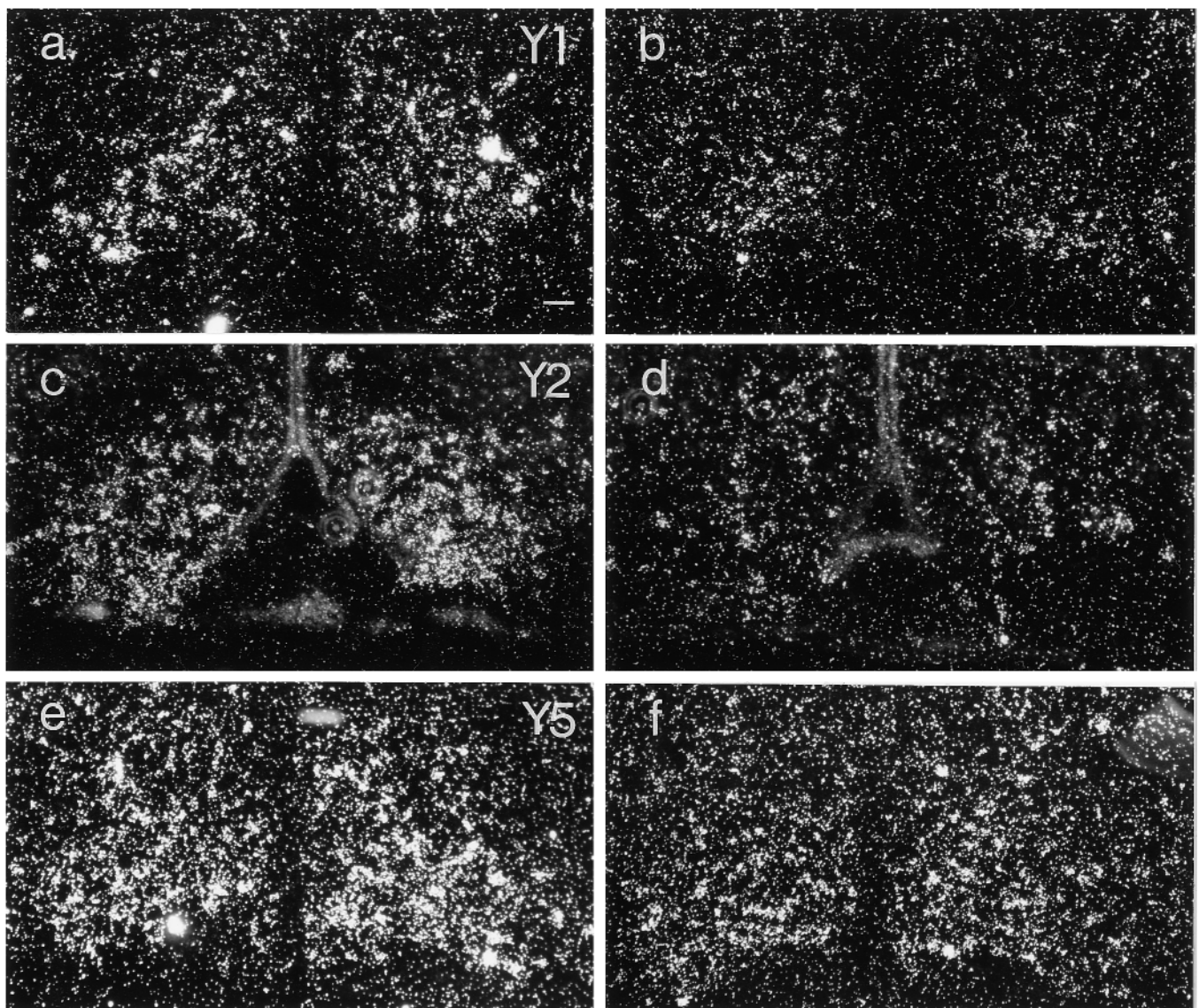


Figure 2. Dark-field micrographs of arcuate nuclei from wild-type (*a, c, e*) and *anx/anx* (*b, d, f*) mice after hybridization for Y1R (*a, b*), Y2R (*c, d*), and Y5R (*e, f*) mRNAs. There is an attenuation of signal for all three mRNAs in mutant animals. Scale bar, 50 μ m.

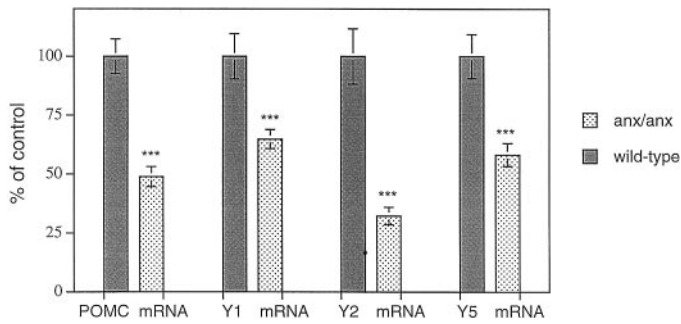


Figure 3. Relative POMC, Y1R, Y2R, and Y5R mRNA levels in wild-type (filled bars) and *anx/anx* (stippled bars) mice, as measured by quantitative autoradiography. Levels of all four mRNA transcripts are significantly lower in mutant mice. Statistical analysis was performed by using ANOVA and Mann–Whitney *U* test. *** $p < 0.001$.

Quantification of immunohistochemically labeled structures. For quantification of ACTH- and Y1R-IR cell bodies, coronal brain sections from both *anx/anx* mice and wild-type littermates at the level of bregma -1.9 mm were selected. The total number of stained cell bodies within the arcuate nucleus was counted at $20\times$ magnification. For quantification of α -MSH-IR nerve terminal arborizations, the periventricular area at the level of the suprachiasmatic nucleus (bregma -0.3) was selected. A grid (Graticules Ltd, Tonbridge, UK), with a 1.0 mm side length of individual squares, was placed in the eyepiece. The total number of crossings between fibers and grid lines within the field of view were counted in sections from both *anx/anx* mice and wild-type littermates through a $20\times$ objective.

Statistical test. Statistical analysis was performed using ANOVA and Mann–Whitney *U* test. All data are presented as mean \pm SD.

Controls. For the control for specificity in *in situ* hybridization, an excess ($100\times$) of unlabeled probe was added to the hybridization cocktail.

For immunohistochemistry, antiserum was preabsorbed with cognate peptide (10^{-6} M) before tissue incubation.

RESULTS

The *anx/anx* mice used in this study could be clearly distinguished from their control littermates phenotypically (also confirmed with genotyping), characterized by emaciation, particularly apparent on the tail where individual vertebrae could easily be delineated, hyperactivity, tremor, and uncoordinated gait. The average body weight of these animals was reduced by $>50\%$ (wild-type, 10.2 ± 0.2 gm; *anx/anx*, 4.5 ± 0.1 gm; $p < 0.001$), and at dissection, an almost complete absence of abdominal adipose tissue was noted.

A distinct autoradiographic POMC mRNA signal could be visualized in the arcuate nucleus of wild-type mice (Fig. 1*a*). In *anx/anx* mice, this labeling was markedly decreased ($48.9 \pm 4.2\%$ of wild-type mRNA levels; $p < 0.001$) (Fig. 1, compare *a*, *b*; Fig. 3). The decrease in POMC expression was at least partly caused by decreased expression within individual cells. A reduction in the number of POMC mRNA-labeled cell bodies in the arcuate nucleus at the level of bregma -1.7 mm was also seen in *anx/anx* mice ($54.8 \pm 11.2\%$ of wild-type; $p < 0.001$).

Hybridization signal for both Y1R, Y2R, and Y5R mRNAs was observed in the arcuate nucleus (Fig. 2*a–f*). Y1R and Y5R mRNAs had similar distributions, predominantly in the ventrolateral portion of the nucleus, whereas Y2R mRNA was restricted to the ventromedial portion. Decreases in autoradiographic signal could be observed and quantified for all three receptor mRNAs in *anx/anx* mice, by approximately two-thirds for Y2R mRNA ($32.9 \pm 3.7\%$ of wild-type mRNA levels; $p < 0.001$) (Fig. 2, compare *c*, *d*; Fig. 3) and approximately one-third for Y1R ($64.9 \pm 4.0\%$ of wild-type mRNA levels; $p < 0.001$) (Fig. 2,

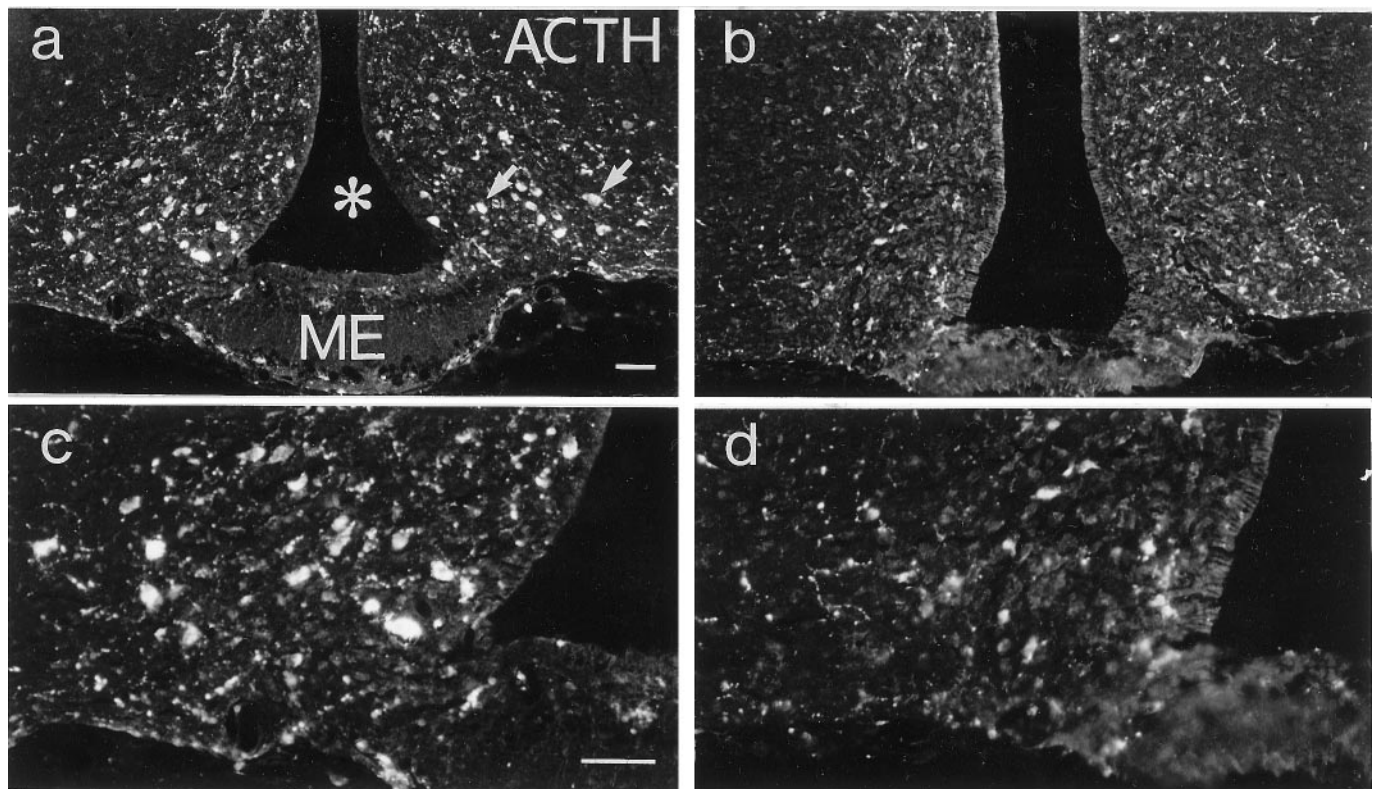


Figure 4. Fluorescence micrographs of sections from the arcuate nucleus of a wild-type (*a*, *c*) and an *anx/anx* (*b*, *d*) mouse stained with antiserum against ACTH. The number of ACTH-IR cell bodies (*a*, arrows) is reduced in mutant mice. *c* and *d* represent a higher magnification of *a* and *b*, respectively. Asterisk indicates third ventricle. ME, Median eminence. Scale bars: (in *a*), *a*, *b*, $50 \mu\text{m}$; (in *c*) *c*, *d*, $50 \mu\text{m}$.

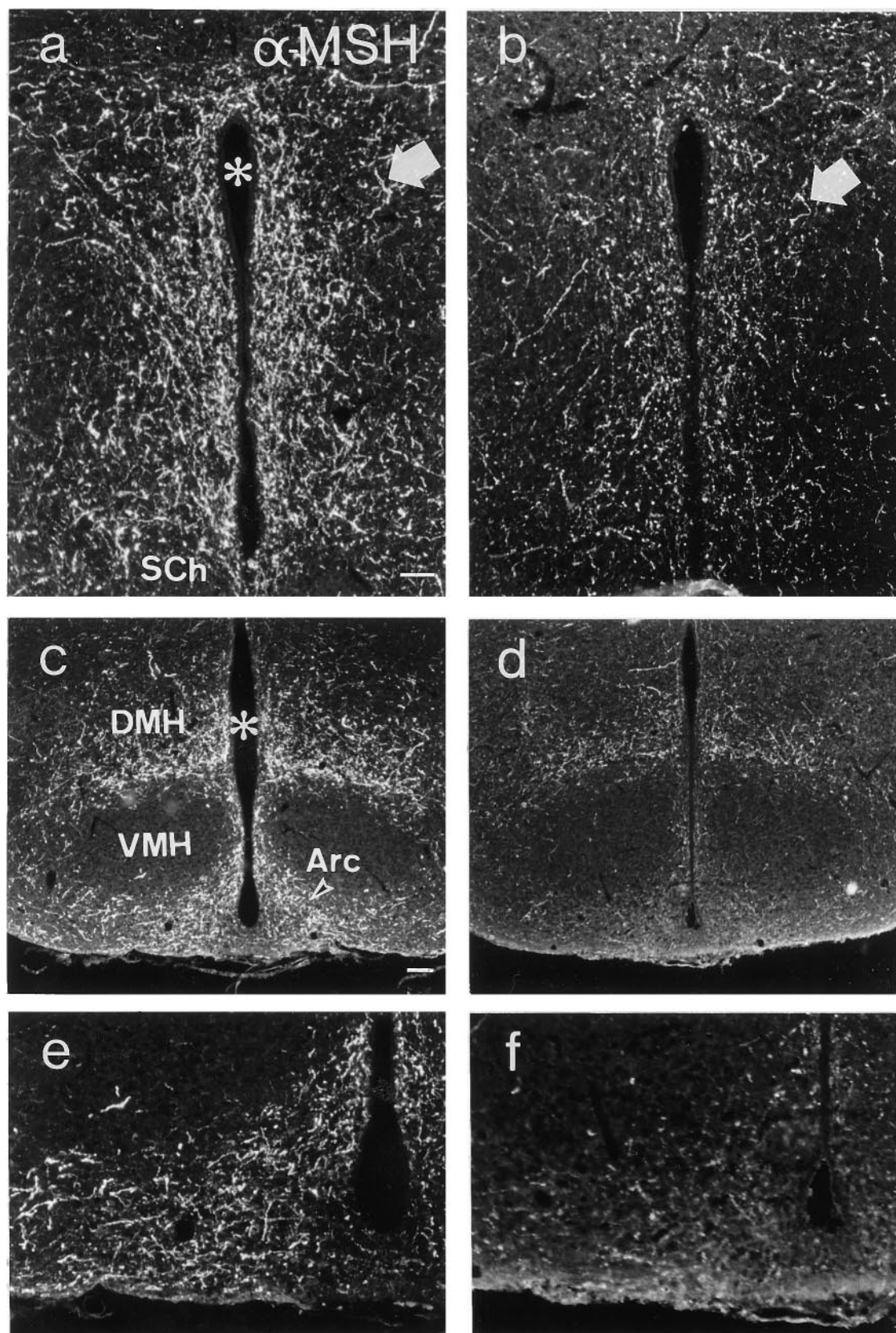


Figure 5. Fluorescence micrographs of sections from the periventricular area (*a, b*) and arcuate nucleus (*c-f*) of wild-type (*a, c, e*) and *anx/anx* (*b, d, f*) mice stained with antiserum against α -MSH. The number of α -MSH-IR terminals (*arrows*) are reduced in mutant mice. *e* and *f* represent a higher magnification of *c* and *d*, respectively. *Asterisk* indicates third ventricle. *Arc*, Arcuate nucleus; *DMH*, dorsomedial hypothalamic nucleus; *SCh*, suprachiasmatic nucleus; *VMH*, ventromedial hypothalamic nucleus. Scale bars: (in *a, b, e, f*, 50 μ m; (in *c, d*, 100 μ m).

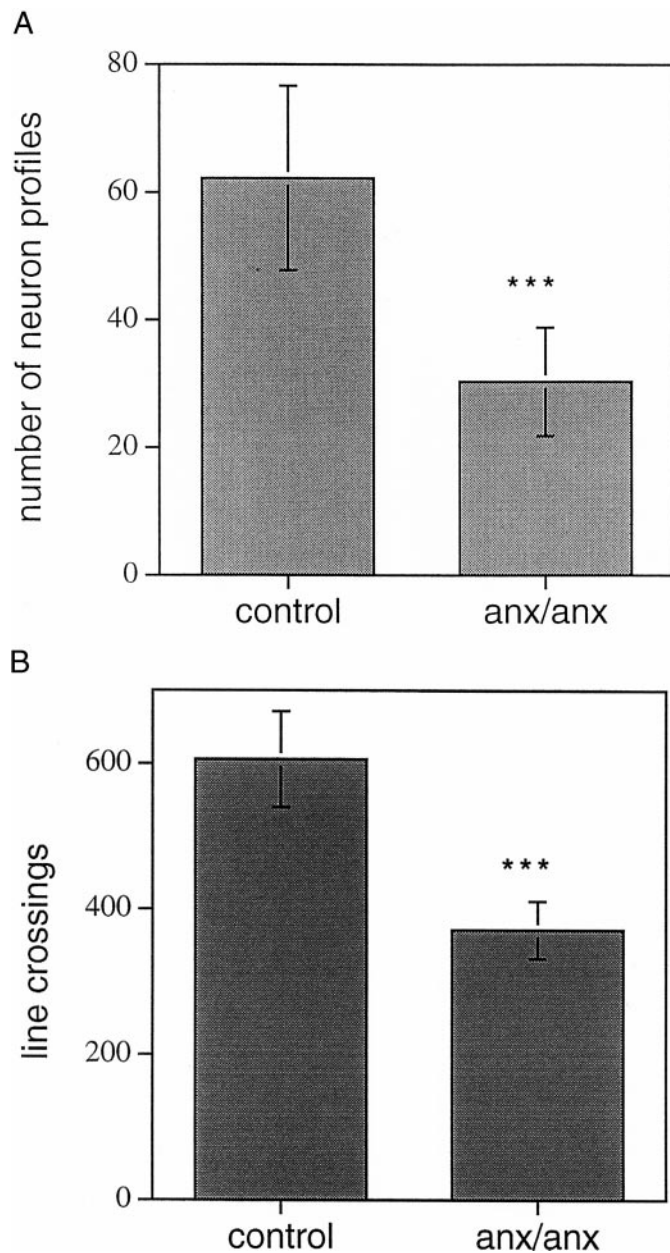


Figure 6. *a*, Quantification of the number of ACTH-IR neuron profiles in the arcuate nucleus. There is a significant reduction in the number of stained profiles in *anx/anx* mice compared with wild-type controls. *b*, Density of α -MSH-IR terminals in the periventricular area of *anx/anx* compared with wild-type mice. Density was quantified as the number of crossings between the lines of a grid inserted in the microscope eyepiece and stained terminals. The number of line crossings is significantly reduced in mutant mice. Statistical analysis was performed by using ANOVA and Mann–Whitney *U* test. *** $p < 0.001$.

compare *a*, *b*; Fig. 3) and Y5R ($58.3 \pm 4.9\%$ of wild-type mRNA levels; $p < 0.001$) (Fig. 2, compare *e*, *f*; Fig. 3).

ACTH-like immunoreactivity (-LI) was seen in cell bodies of the ventrolateral portion of the arcuate nucleus of both *anx/anx* and wild-type mice (Fig. 4*a–d*) and in a limited number of varicose axons in areas in which α -MSH-LI could be detected (see below). The number of stained cell bodies was decreased in *anx/anx* mice ($48.8 \pm 3.8\%$ of wild-type; $p < 0.001$) (Figs. 4, compare *a*, *c*, with *b*, *d*, respectively; Fig. 6).

α -MSH-IR terminals could be observed in many nuclei, includ-

ing the preoptic area, periventricular zone, paraventricular thalamic and hypothalamic nuclei, dorsomedial hypothalamic nucleus, and the arcuate itself (Fig. 5*a–f*); however, no arcuate cell bodies exhibiting α -MSH-LI were seen. In *anx/anx* mice, the density of terminal arborizations was notably attenuated (Fig. 5, compare *a*, *c*, *e* with *b*, *d*, *f*, respectively). Fiber density was quantified in the periventricular zone at the level of the supra-chiasmatic nucleus and was found to be significantly decreased in *anx/anx* mice ($61.3 \pm 6.5\%$ of wild-type; $p < 0.001$) (Fig. 6*b*).

In wild-type animals, Y1R-LI was observed in the ventrolateral arcuate nucleus, decorating the cytoplasm and dendrites of large polygonal cells (Fig. 7*a*, *c*). The rich dendritic arborization observed in control mice was distinctly less pronounced in *anx/anx* mice and was almost completely absent in approximately half of the mutant mice (Fig. 7, compare *a*, *c* with *b*, *d*). Furthermore, the number of stained cell bodies also appeared to be decreased, although this reduction was more variable ($60.7 \pm 30.8\%$ of wild-type). Similar changes could also be observed in Y1R-IR cells of the paraventricular hypothalamic nucleus (Fig. 8, compare *a*, *c* with *b*, *d*).

Autoradiographic signal after *in situ* hybridization was abolished if an excess of the cognate unlabeled oligonucleotide probe was added to the hybridization cocktail. The histochemical staining patterns described above could not be observed after incubation of antisera with corresponding peptide before tissue incubation.

DISCUSSION

The present study investigated the morphology and histochemistry of arcuate POMC neurons in a genetic model of hypophagia, the *anx/anx* mouse. Mutant mice displayed decreased levels, not only of POMC mRNA, but also of Y1R and Y5R mRNAs, which are expressed in the POMC neurons. In addition, morphological characteristics of POMC neurons were also affected, as seen with immunohistochemical markers: thus, the number of ACTH-IR cell bodies, as well as α -MSH-IR terminals was decreased, and, furthermore, a dramatic reduction of dendritic extensions of the arcuate Y1R-IR cells [primarily corresponding to POMC cells (Broberger et al., 1997*b*; Fuxe et al., 1997)] was seen, as determined by staining with Y1R antiserum. In the arcuate neurons, ACTH, after being cleaved from the POMC precursor, is further processed to α -MSH (Watson and Akil, 1980; Khachaturian et al., 1985). The ratio of α -MSH/ACTH in the brain gradually increases with the distance from the arcuate cell bodies (Barnea et al., 1979). Thus, it could be expected that antiserum directed against the intact ACTH segment would predominantly stain cell bodies and, only to a lesser extent, terminals. In contrast, antiserum directed against α -MSH would be more likely to stain terminals, all in accordance with the results shown here. It has been demonstrated that the concentrations of α -MSH-LI in the brain are 4–15 times higher than those of ACTH-LI (Mezey et al., 1985).

Our previous studies on the *anx/anx* mouse have demonstrated altered histochemistry of arcuate NPY-AGRP coexpressing neurons (Broberger et al., 1997*a*, 1998*b*). However, it is important to note that the changes occurring in the NPY-AGRP population are distinct from those seen in POMC cells (present paper). Thus, the decreases in NPY-AGRP terminals are paired with increased cell body staining for these peptides, and NPY-AGRP mRNA levels remain unaffected in *anx/anx* mice. In contrast, the histochemical features of arcuate POMC neurons in *anx/anx* mice were characterized by decreased mRNA levels for all markers

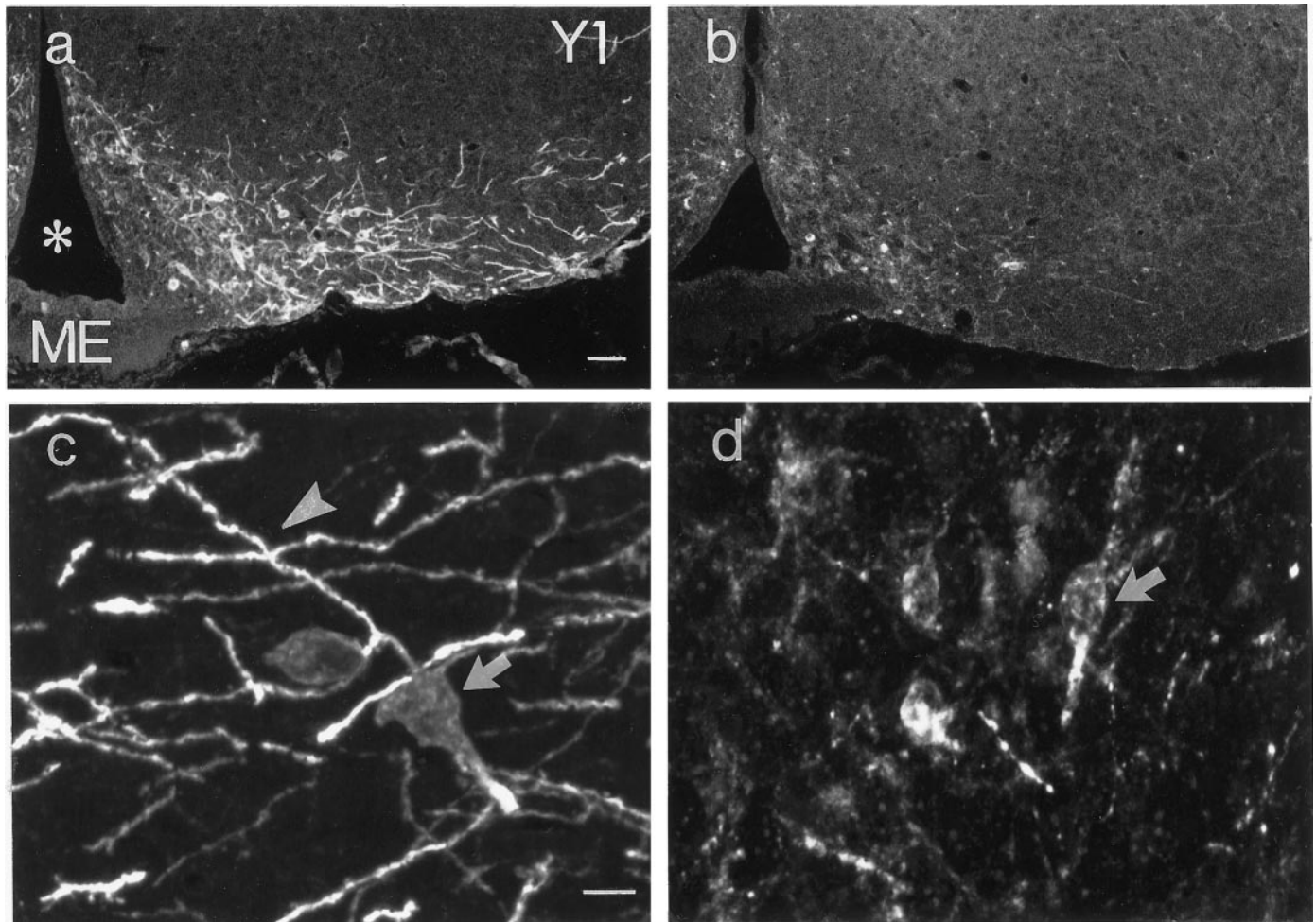


Figure 7. Fluorescence (*a, b*) and confocal (*c, d*) micrographs of sections from the arcuate nucleus of a wild-type (*a, c*) and an *anx/anx* (*b, d*) mouse stained with antiserum against the Y1R. Note dramatic reduction in dendritic arborizations (*arrowheads*), as well as attenuation in fluorescence and number of cell bodies (*arrows*) in *anx/anx* mice. *Asterisk* indicates third ventricle. *ME*, Median eminence. Scale bars: (in *a, b*) 50 μm ; (in *c, d*) 10 μm .

studied, as well as decreased markers for both cell body and dendritic and axonal arborizations, suggestive of atrophy. One speculative explanation to account for these changes is that some molecule(s) secreted from the NPY cells may act as a trophic signal for POMC neurons and that POMC cells deteriorate–degenerate in the absence of this signal. Alternatively, synaptic signaling contributed by NPY terminals may be necessary for the survival of POMC neurons.

There is good evidence for an antagonistic relationship between the arcuate NPY and POMC populations. Recent studies suggest that NPY neurons may regulate POMC signaling in several ways. POMC cell bodies and dendrites are lined by Y1Rs (Broberger et al., 1997b; Fuxe et al., 1997) and possibly also Y5Rs (Gerald et al., 1996; Naveilhan et al., 1998a) [which may be explained by the proximity of the two genes within the same transcriptional region (Herzog et al., 1997)], both of which couple to inhibitory G-proteins (Herzog et al., 1992; Larhammar et al., 1992; Wahlestedt and Reis, 1993). Also, NPY neurons coexpress AGRP (Shutter et al., 1997; Broberger et al., 1998a,b; Hahn et al., 1998), which acts as an antagonist of melanocortins at the MC4 receptor (Ollmann et al., 1997) and may be released from terminals in close proximity to POMC terminals (Broberger et al., 1998b). Thus, through regulation on both the cell body and terminal levels, NPY may have the “upper hand” in its antago-

nistic relationship with POMC neurons, with POMC derivatives serving primarily to counterbalance the orexigenic NPY signals in states of positive energy balance. It may be speculated that the presence of inflow from the NPY neurons, although inhibitory on POMC signaling itself, is necessary for the development of the POMC system. It is noteworthy that changes in Y1R immunostaining, similar to those observed in the arcuate, could also be seen in cells in the paraventricular hypothalamic nucleus (present data). These Y1R-expressing cells also receive innervation from arcuate NPY neurons (Bai et al., 1985; Broberger et al., 1998b) but do not express POMC. This suggests that changes in the tone exerted by arcuate-derived NPY terminals, rather than the histochemical phenotype of the target cell, are of relevance for the morphological changes observed in hypothalamic Y1R-IR cells in *anx/anx* mice. This hypothesis remains to be validated experimentally, and to fully determine whether the NPY and POMC changes are interrelated may have to await molecular characterization of the *anx* gene. Furthermore, we have not in the present study investigated whether the brainstem POMC neurons, which also appear to modulate food intake (Grill et al., 1998), are affected in the *anx/anx* mouse.

The argument made above is based on the assumption that there is a *de facto* impairment in release of secreted molecules from the NPY neurons. It should be kept in mind, however, that

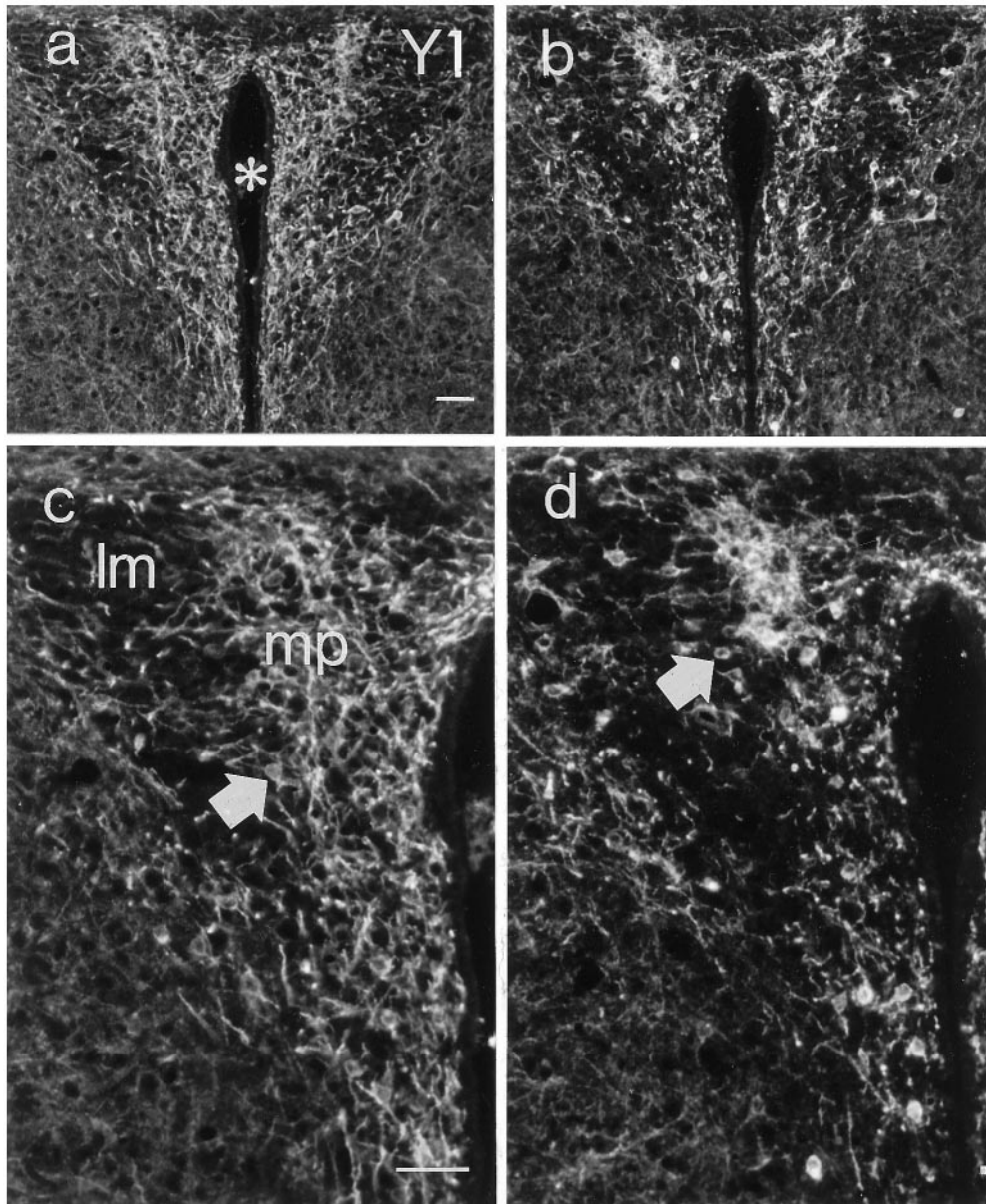


Figure 8. Fluorescence micrographs of sections from the paraventricular hypothalamic nucleus of a wild-type (*a, c*) and an *anx/anx* (*b, d*) mouse stained with antiserum against the Y1R. (Micrographs in *a* and *b* are magnified in *c* and *d*, respectively). There is a reduction in both cell bodies (*arrows*) and dendritic arborizations. *Asterisk* indicates third ventricle. *mp*, Medial parvocellular nucleus; *lm*, lateral magnocellular nucleus. Scale bar, 50 μm .

the changes in NPY histochemistry (Broberger et al., 1997a) are partly compatible also with a hypersecretion of NPY as an effect secondary to energy depletion, wherein the decreased terminal staining could be the result of increased release and the increased cell body staining the result of increased production of NPY. Yet, this view does not appear compatible with the normal arcuate NPY mRNA levels in the *anx/anx* mouse (Broberger et al., 1997a) (although it cannot be ruled out that translational and post-translational mechanisms are affected in mutant mice). Starvation per se increases arcuate NPY mRNA (Brady et al., 1990) and also increases NPY staining in arcuate terminal fields (Calzà et al., 1989), contrary to what is seen in *anx/anx* mice.

In this context, it is also notable that, in the arcuate nucleus of mice lacking the NPY gene (Erickson et al., 1996), POMC expression is comparable with normal animals (Hohmann et al., 1998), and Y1R-LI is distributed in a normal pattern with richly arborized dendrites (C. Broberger, J. Erickson, R. Palmiter, and T. Hökfelt, unpublished observations). This would suggest that NPY itself is not necessary for the development of the POMC

system. However, as mentioned above, the NPY neurons express several other messenger molecules, including AGRP (Broberger et al., 1998a,b; Hahn et al., 1998) and, in a subpopulation of NPY neurons, the GABA-synthesizing enzyme glutamic acid decarboxylase (Horvath et al., 1997). Likely, several other coexpressed molecules remain to be discovered, which may further explain how NPY neurons regulate the POMC cells. Finally, the possible involvement of β -endorphin, a POMC-derived peptide that, in contrast to the melanocortins, stimulates food intake when administered centrally (Grandison and Guidotti, 1977), remains to be investigated in the *anx* phenotype.

Microinjection of the Y2R agonist NPY(13–36) into the hypothalamic paraventricular nucleus inhibits food intake (Leibowitz and Alexander, 1991), and mice rendered deficient of the Y2R gene develop hyperphagia when fed a high-fat diet (Naveilhan et al., 1998b), indicating that the Y2R exerts an inhibitory tone on food intake. The Y2R is expressed in arcuate NPY neurons (Broberger et al., 1997b). Pharmacologically, this receptor has inhibitory properties, activating potassium currents and inhibiting

calcium channels (Ewald et al., 1988; Chen and van den Pol, 1996; McQuiston and Colmers, 1996; Sun et al., 1998). Together, this suggests that the Y2R may act as a presynaptic auto-inhibitory receptor (Wahlestedt et al., 1986). Combined with our previous demonstration of decreased hypothalamic NPY terminal staining in the *anx/anx* mouse (Broberger et al., 1997a), the present data may be interpreted as a compensatory downregulation of an auto-inhibitory mechanism in the absence of neurotransmitter release.

In summary, the results presented here demonstrate decreased labeling for all markers of arcuate POMC neurons so far studied, suggesting atrophy–degeneration of this population. In addition, we also demonstrate decreased mRNA levels for the Y2 receptor, which may be a response to decreased NPY transmission in this mutant. These data further delineate the hypothalamic abnormalities that may underlie the *anx/anx* phenotype.

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