Loss of $G_{q/11}$ Family G Proteins in the Nervous System Causes Pituitary Somatotroph Hypoplasia and Dwarfism in Mice

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Heterotrimeric G proteins of the Gq/11 family transduce signals from a variety of neurotransmitter and hormone receptors and have therefore been implicated in various functions of the nervous system. Using the Cre/*loxP* system, we generated mice which lack the genes coding for the α subunits of the two main members **of the Gq/11 family,** *gnaq* **and** *gna11***, selectively in neuronal and glial precursor cells. Mice with defective** *gnaq* **and** *gna11* **genes were morphologically normal, but they died shortly after birth. Mice carrying a single** *gna11* **allele survived the early postnatal period but died within 3 to 6 weeks as anorectic dwarfs. In these mice, postnatal proliferation of pituitary somatotroph cells was strongly impaired, and plasma growth hormone (GH) levels were reduced to 15%. Hypothalamic levels of GH-releasing hormone (GHRH), an important stimulator of somatotroph proliferation, were strongly decreased, and exogenous administration of GHRH restored normal proliferation. The hypothalamic effects of ghrelin, a regulator of GHRH production and food intake, were reduced in these mice, suggesting that an impairment of ghrelin receptor signaling might** contribute to GHRH deficiency and abnormal eating behavior. Taken together, our findings show that $G_{q/11}$ **signaling is required for normal hypothalamic function and that impairment of this signaling pathway causes somatotroph hypoplasia, dwarfism, and anorexia.**

The $G_{q/11}$ family of heterotrimeric G proteins mediates the cellular effects of numerous neurotransmitter receptors, e.g., the metabotropic glutamate receptor subtypes 1 and 5, the M_1 muscarinic-acetylcholine receptor, the $5-\text{HT}_2$ serotonin receptors, or the α_1 adrenergic receptor. $G_{q/11}$ -coupled receptors are also involved in the signal transduction of several hypothalamic peptide hormone receptors, such as the receptors for thyrotropin-releasing hormone (39), gonadotropin-releasing hormone (13), and prolactin-releasing hormones (3). There is an increasing amount of evidence that the release of hypothalamic hormones themselves is also controlled by $G_{q/11}$ -coupled receptors. Kisspeptins, for example, have been suggested to control the release of gonadotropin-releasing hormone via the $G_{q/11}$ -coupled receptor GPR54 (14, 20), and the gastrointestinal peptide hormone ghrelin regulates the production of growth hormone-releasing hormone (GHRH) via the hypothalamic growth hormone secretagogue receptor (GHS-R) (12, 22). G_{q/11} family G proteins mediate the activation of β isoforms of phospholipase C, resulting in the activation of protein kinase C and intracellular calcium mobilization (2). The $G_{q/11}$ family consists of four members, two of which, G_q and $G₁₁$, are expressed almost ubiquitously in the central nervous system (26). Genetic inactivation of the *gnaq* gene, which codes for the α subunit of G_q ($G\alpha_q$), leads to a defect in primary hemostasis (16) and cerebellar ataxia (15). In contrast, mice homozygous for a null allele of the gene coding for $G\alpha_{11}$, *gna11*, did not show any phenotypic abnormalities (17). These defects were relatively mild when one considers the number of potential

transmitter systems affected, and this fact is probably due to the high functional redundancy of Ga_q and Ga_{11} , which share 88% amino acid sequence identity (24). Indeed, mice lacking both G α_{q} and G α_{11} die at day 10.5 of embryonic development (17). To circumvent this embryonic lethality, we used the Cre/ *loxP* system (18) to generate a mouse line which allows conditional, tissue-specific inactivation of *gnaq* in constitutively Ga_{11} -deficient mice (36). Postnatal inactivation of $G_{q/11}$ -mediated signaling in the forebrain by use of the *Camkcre4* mouse line was shown to inhibit maternal behavior (35). To investigate the role of Ga_{q} and Ga_{11} in pre- and postnatal development and the function of the whole nervous system, we used in this study the *Nestin*Cre mouse line (31), which causes Cremediated recombination in neuronal and glial precursor cells starting at embryonic day 9.5 (40). Interestingly, the $G\alpha_q/G\alpha_{11}$ deficiency in descendants of neuronal and glial precursors did not cause gross morphological abnormalities of the developing nervous system but was incompatible with postnatal survival. In addition, we provide evidence that $G_{q/11}$ -mediated signaling is critically involved in the regulation of hypothalamic GHRH production and consecutive control of somatotroph proliferation.

MATERIALS AND METHODS

Generation of mice lacking $G\alpha_q$ and $G\alpha_{11}$ in neuronal precursor cells. Mice in which the gene coding for G_{α_q} , *gnaq*, is flanked by *loxP* sites (*gnaq*^{flox}) (36) were crossed to the constitutively $G\alpha_{11}$ -deficient mouse line (17) and to mice which express the recombinase Cre under the control of the *Nestin* promoter (31, 40). This action led to the generation of mice which lacked one, two, three, or all of the four *gnaq*/*gna11* alleles in derivatives of neuronal and glial precursor cells. Genotyping for the *gnaq*flox allele, for wild-type and knockout *gna11* alleles, and for the Cre transgene was performed as described previously (36).

Hormone levels. Plasma samples were obtained from 15-day-old mice at the beginning of the dark period. Levels of insulin-like growth factor I (IGF-I) were determined with an OCTEIA rat and mouse IGF-I kit (IDS Inc., Boldon, United

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Kingdom), ghrelin levels were determined with a rat and mouse ghrelin radioimmunoassay (RIA) kit (Phoenix, Belmont, Calif.), thyroxine levels were determined with a Micro enzyme immunoassay kit (Leinco, St. Louis, Mo.), and adrenocorticotropic hormone (ACTH) levels were determined with an immunometric assay kit (Nichols Institute Diagnostics, San Clemente, Calif.). Corticosterone was measured by RIA in the Steroid Laboratory of the Pharmacology Institute, Heidelberg, Germany. GH and thyroid-stimulating hormone (TSH) were measured by RIA in the laboratory of the National Hormone and Pituitary Program, Torrance, Calif.

Histology. Mice were deeply anesthetized with pentobarbital (100 mg/kg of body weight) intraperitoneally (i.p.) and perfused with 4% paraformaldehyde (PFA) via the left cardiac ventricle. Brains were postfixed in 4% PFA overnight and then stored in 0.5% PFA at 4°C. For immunohistochemistry, 50- μ m-thick vibratome sections were cut and incubated at 4°C with the following antibodies: rabbit anti-c-*fos* antibody (sc-52; Santa Cruz Biotechnology, Santa Cruz, Calif.) at a 1:20,000 dilution for 3 days, rabbit anti-G $\alpha_{q/11}$ antibody (sc-392; Santa Cruz Biotechnology) at a 1:1,000 dilution for 16 h, rabbit anti-GHRH antibody (obtained from F. Talamantes, University of California, Santa Cruz, Santa Cruz, Calif.) at a 1:20,000 dilution for 16 h, and mouse anti-NeuN antibody (Chemicon, Hofheim, Germany) at a dilution of 1:2,000 for 16 h. For staining, we used the Vectastain Elite ABC kit (Vector Laboratories, Burlingame, Calif.). Internal organs and brains were weighed after preparation, and hematoxylin-eosin staining was performed on 5 - μ m-thick paraffin sections according to standard proto c ols. For immunofluorescence staining of pituitary somatotroph cells, 5- μ m-thick paraffin sections were incubated for 16 h with a rabbit anti-GH antibody (dilution, 1:20,000; obtained from A. F. Parlow) and then with fluorescein isothiocyanate (FITC)-labeled goat anti-rabbit antibody (dilution, 1:200; Dianova, Hamburg, Germany) for 2 h. For BrdU labeling, 5-day-old mice were injected intraperitoneally on three consecutive days with 10μ g of GHRH (Phoenix)/kg or with a comparable volume of saline. In the afternoon of the third day, mice were intraperitoneally injected with 10 μ g of BrdU/kg of body weight and sacrificed after 2 h. Paraffin sections of the pituitary were stained with anti-BrdU antibody (dilution, 1:100; BD Biosciences, San Diego, Calif.) and FITC-labeled goat anti-rabbit antibody (dilution, 1:200; Jackson ImmunoResearch) for 2 h. For ghrelin immunohistochemistry, 16 - μ m-thick cryotome sections of the stomach were stained with goat anti-ghrelin antibody (dilution, 1:300; Santa Cruz Biotechnology) and by using a Vectastain Elite ABC kit (Vector Labs). The numbers of c-*fos*-, NeuN-, ghrelin-, or BrdU-positive cells, as well as the area of GHRH immunoreactivity, were determined with the CellExplorer 2003 program (BioSciTec, Frankfurt, Germany).

Food uptake. Food uptake during the dark phase was determined in 3-to 4-week-old mice by weighing standard chow before and after the test period. Water was accessible ad libitum. To determine the effect of ghrelin on food intake, mice were injected intraperitoneally with 1 mg of rat ghrelin (Bachem, Heidelberg, Germany) or saline/kg of body weight, and food uptake during the following 3 h was determined every 30 min. In some cases, mice were sacrificed 1 h after ghrelin injection for c-*fos* immunohistochemistry of their arcuate nuclei (arcN).

Statistics. Values were expressed as means \pm standard errors of the means. Differences between two groups were statistically analyzed by using the unpaired Student's *t* test. Statistical significance was accepted at a P value of ≤ 0.05 .

RESULTS

We used the *Nestin*Cre mouse line (31) to generate mice which lacked the G protein α subunit Ga_q selectively in neuronal and glial precursor cells, and we then mated these mice with the constitutively $G\alpha_{11}$ -deficient mouse line (17). This procedure resulted in the generation of mice which lacked one, two, three, or all four *gnaq*/*gna11* alleles in neuronal and glial precursor cells. To prove that Cre expression leads to the recombination of the *gnaq*flox allele and thereby to a loss of Ga_{q} protein, we performed Western blot experiments with extracts from the brains of newborn mice by using an antibody directed against G $\alpha_{q/11}$ (Fig. 1a). In the brains of *NestinCre*^{+/-} $\frac{g}{\text{max}}$ *flox*/flox $\frac{g}{\text{max}}$ *gna11* –/² mice (i.e., no *gnaq* or *gna11* alleles were left), neither Ga_{q} nor Ga_{11} protein was detectable, whereas *NestinCre*^{+/-} *gnaq*^{flox/wt} *gna11*^{-/-} (i.e., one *gnaq* allele is intact) or *NestinCre*^{+/-} $gnaq^{\text{flow/flox}}$ $gna11^{+/-}$ mouse brains (i.e.,

FIG. 1. The survival of *Nestin*-G $\alpha_{q/11}$ -deficient mice depends on the number of remaining *gnaq*/*gna11* alleles. (a) Western blot for G $\alpha_{q/11}$ proteins in brain extracts (10 μ g per lane) from newborn mice with different genotypes. $+$, wild-type allele; fl, recombined floxed allele; $-$, null allele. (b) Reduced body length of a mouse with only one remaining *gna11* allele (bottom) compared to a control littermate (top) at postnatal day 20. (c and d) Survival rate (c) and postnatal weight gain (d) of *NestinCre^{+/ –} gnaq*^{flox/flox} *gna11^{-/ –}* mice (asterisks), *NestinCre^{+/ –}* $gnaq^{\text{flow/flox}}$ *gna11^{+/* $\overline{}$ mice (circles), *NestinCre^{+/-} gnaq*^{flox/wt} *gna11^{-/-}*} mice (triangles), or control mice (squares) (24 to 30 mice per group). p, postnatal day. (e) Representative hematoxylin-eosin stainings of frontal brain sections at embryonic day 18.5 from a control animal (left) and a *NestinCre*^{+/-} *gnaq*^{flox/flox} *gna11*^{-/-} mouse (right) (magnification, \times 2.5). (f) Tenfold magnification of the mouse sensory cortices shown in panel e.

one gna11 allele is intact) lacked only Ga_{11} or Ga_q , respectively. These results show that the *Nestin*Cre-mediated recombination of gnaq^{flox} alleles indeed leads to a loss of $G\alpha_q$ protein in the nervous system.

Matings between triple-heterozygous mice (NestinCre^{+/-} *gnaq*^{flox/wt} *gna11*^{+/-}) showed that mice of all genotypic combinations were born at expected frequencies (data not shown). However, the postnatal survival of a newborn strongly depended on the number of intact *gnaq*/*gna11* alleles (Fig. 1c). Normal numbers of newborns in which all four *gnaq*/*gna11* alleles were inactivated were born without obvious malformations, but they did not take up rhythmic breathing after delivery. Accordingly, the lungs of these animals were only partially inflated (data not shown). Neither histological nor basic immunohistochemical analysis of the brains of these animals revealed major structural abnormalities (Fig. 1e and f and data not shown). The peripheral nervous system, including the phrenic nerve, was histologically normal in each mouse. Initial reflexes towards tactile stimuli and basal cardiovascular function seemed to be normal, suggesting that inefficient respiration caused the deaths of these newborns.

FIG. 2. Organ weights and food uptake of wild-type mice and mice with only one intact *gna11* allele (*NestinCre^{+/-} gnaq*^{flox/flox} *gna11*^{+/-}). (a to d, f) Organ weights as percentages of body weight $($ %bw) for kidneys (a), brains (b), stomachs (c), retrorenal fat deposits (d), and pituitaries (f) at postnatal days 15 (p15) and 25 (p25) (five to six mice per group). (e) Food uptake in 3-week-old mice (five to six mice per group). $\hat{P} < 0.05$; **, $P < 0.01$; ***, $P < 0.005$. (g) Pituitaries from a control mouse (left) and an animal with only one intact *gna11* allele (right) at postnatal day 15. Borders of the posterior pituitary are indicated by hatched ovals.

Mice with at least one intact *gnaq* allele or two intact *gna11* alleles were viable and fertile. Of note, mice lacking both *gnaq* alleles showed an ataxic gait comparable to the one seen in constitutively Ga_q -deficient mice (16), whereas animals with a single intact *gnaq* allele were normal in this respect. In contrast, mice with only one *gna11* allele were strongly retarded in their somatic growth (Fig. 1b and d) and died as underweight dwarfs between the third and sixth postnatal weeks (Fig. 1c).

Growth retardation in mice with a single *gna11* allele was first detectable between postnatal days 5 and 9 (Fig. 1d). During the following weeks, the body weights increased only slowly, peaked between the second and third weeks, and finally decreased again (Fig. 1d), usually leading to the deaths of the animals between the third and sixth weeks (Fig. 1c). Except for their small size and slightly retarded motor development, these animals appeared to be normal, i.e., they did not show obvious behavioral abnormalities or signs of illness. Neither the removal of normal littermates to decrease competition for the dam's milk nor the presence of easily accessible extra food changed the final outcome. To determine whether the growth retardation and premature deaths of mice with only one intact *gna11* allele could be attributed to the failure of a specific organ system, we performed macroscopic and microscopic analyses of their central nervous systems and internal organs. Brains, hearts, lungs, kidneys, livers, thyroids, and adrenal glands did not show major morphological abnormalities (data not shown). Analyses of organ weight/body weight ratios showed that kidneys and hearts were of normal sizes relative to the body weights of the mice (Fig. 2a and data not shown), whereas the relative brain weights were increased (Fig. 2b). In contrast, the relative weights of the stomach and of retrorenal fat deposits were decreased (Fig. 2c and d), and this was probably due to reduced food uptake (Fig. 2e). Most strikingly, the pituitaries were much too small in relation to body weights, especially at older ages, and this circumstance was due mainly to the reduced size of the anterior pituitary in each mouse, since the posterior pituitary was of normal size (Fig. 2f and g).

As somatic growth is regulated mainly through secretion of GH from somatotroph cells of the anterior pituitary, we reasoned that the observed dwarfism might be due to impaired somatotroph function. Immunofluorescence staining showed that the densities of GH-positive cells were reduced to about 50% of that of the control cells in pituitaries from 2-week-old mice (Fig. 3a and b). This reduced density of GH-positive cells, together with the massive reduction of anterior pituitary volume (Fig. 2f and g), leads to GH deficiency, with plasma GH levels at 15% of the normal level (Fig. 3c). Consistent with this, IGF-I levels were reduced to 30% of normal levels in 2-weekold animals (Fig. 3d). Interestingly, other anterior pituitary functions, as determined by levels of TSH, thyroxine, ACTH, or the mouse glucocorticoid analogue corticosterone in plasma, were normal on postnatal day 15 (Fig. 3e through h). These findings show that mice with only one intact *gna11* allele

FIG. 3. Reduced numbers of somatotrophs in pituitaries of mice with only one intact *gna11* allele (*NestinCre*^{+/-} *gnaq*^{flox/flox} *gna11*^{+/-}). (a) Immunofluorescence staining for GH in the anterior pituitary of 15-day-old control mice (left) and mice with one remaining *gna11* allele (right). (b) Statistical evaluation of immunofluorescence staining (two mice per group, three sections each). (c to h) Levels in plasma, at postnatal day 15, of GH (15 mice per group) (c), IGF-I (6 to 8 mice per group) (d), ACTH (8 mice per group) (e), mouse cortisol analogue corticosterone (8 mice per group) (f), TSH (g), and thyroxine, T4 (h) (6 to 8 mice per group). ***, $P < 0.001$.

FIG. 4. Impaired postnatal proliferation of somatotroph cells can be rescued by administration of GHRH. (a) Hematoxylin-eosin staining of coronal head sections from a control animal (left) and a mouse with only one intact *gna11* allele (right) at embryonic day 18.5. (b) BrdU-labeled cells in the anterior pituitary of 7-day-old control animals (left) and mice with only one intact *gna11* allele (right) after 3 days of intraperitoneal administration of saline (top) or GHRH (bottom). (c) Statistical analysis of BrdU-positive (pos.) cells (two mice per group, three sections each). Black bars, control mice; white bars, mice with one intact *gna11* allele; ***, $P < 0.001$. (d) β -Galactosidase staining of a pituitary from a mouse carrying both the *Rosa26*-*lacZ* reporter construct and the *Nestin*Cre transgene. (a and d) ap, anterior pituitary; ht, hypothalamus; il, intermediate lobe; pp, posterior pituitary; sb, sphenoid bone.

in the nervous system suffer from selective somatotroph hypoplasia, leading to strongly reduced plasma GH and IGF-I levels and, consecutively, proportionate dwarfism.

To test whether hypoplasia of the anterior pituitary was due to a defect in prenatal development, we histologically analyzed pituitaries from mice at embryonic day 18.5. At this age, pituitary sizes and development were normal for mice of all genotypes (Fig. 4a), indicating that somatotroph hypoplasia is due to a defect in postnatal proliferation. To test this concept, we performed BrdU labeling of pituitaries in the first postnatal week and found that the number of proliferating cells was reduced in mice with only one intact *gna11* allele to 38% of that of the control littermates (Fig. 4b and c). An important stimulator of somatotroph proliferation is GHRH (4), which is produced in neurons of the hypothalamic arcN and which is released into the pituitary portal system at the median eminence. To test whether pituitaries from mice with only one intact *gna11* allele would be responsive to exogenously administered GHRH, we treated pups of different genotypes with daily intraperitoneal injections of saline or GHRH from postnatal days 5 through 7. This treatment strongly enhanced anterior pituitary proliferation in mice with only one intact *gna11* allele, which almost reached the proliferation levels of salinetreated control littermates (Fig. 4b and c). These findings suggest that somatotroph cells are responsive to GHRH in these mice and that somatotroph hypoplasia is not due to a defect of the pituitary itself but rather to inappropriate GHRH release from the hypothalamus. Consistent with this, β -galactosidase staining of mice carrying both the *Nestin*Cre transgene and a *Rosa26*-*lacZ* reporter construct (23) did not show any recombination in the anterior pituitary, whereas the posterior pituitary and hypothalamus were recombined (Fig. 4d and data not shown).

Immunohistochemical analyses of the hypothalamic arcN and median eminence, in which GHRH-producing neurons are located, revealed a considerable immunoreactivity for $Ga_q/$ Ga_{11} in wild-type controls, which was strongly reduced in mice with only one intact gna11 allele (Fig. 5a). To quantify $Ga_q/$ Ga_{11} protein expression in the hypothalami of mice with only one intact Ga_q or Ga_{11} allele, we performed Western blotting experiments using extracts from hypothalami of wild-type or Ga_q/Ga_{11} -deficient mice. We found that in wild-type mouse hypothalami, Ga_q levels were severalfold higher than Ga_{11} levels and that the reduction of Ga_q/Ga_{11} immunoreactivity was much stronger in hypothalami from animals with one intact Ga_{11} allele than in hypothalami from mice with one intact Ga_{q} allele (Fig. 5b).

Immunohistochemistry using an antibody directed against GHRH revealed that the number of GHRH-immunoreactive terminals was markedly decreased in mice with only one intact

FIG. 5. Immunohistochemical changes in the hypothalami of mice with only one gnall allele. (a) Staining with antibodies against $Ga_{q/11}$ in the arcN of a control mouse (left) and a mouse with one intact *gna11* allele (right). (b) Western blot of extracts from wild-type and mutant mouse hypothalami (10 μ g per lane) with antibodies against G $\alpha_{q/11}$ and total extracellular signal-regulated kinase (ERK) as the loading control. $+$, wild-type allele; fl, recombined floxed allele; $-$, null allele. (c) Staining with antibodies against GHRH in the arcN of a control mouse (left) and a mouse with one intact *gna11* allele (right). (d) Statistical evaluation of GHRH immunoreactivity (GHRH-IR) in the median eminence (two mice per group, three sections each). Black bars, control mice; white bars, mice with only one intact *gna11* allele. ***, $P < 0.001$.

FIG. 6. Impaired ghrelin actions in mice with only one intact *gna11* allele (*Nestin*Cre^{+/-} *gnaq*^{flox/flox} *gna11*^{+/-}). (a) c-*fos* immunohisto- $\frac{1}{\pi}$ gnaq^{flox/flox} $\frac{1}{\pi}$ gna11^{+/-}). (a) c-*fos* immunohistochemistry of the arcN in control mice (left) or mice with one intact *gna11* allele (right) after the i.p. administration of saline (top) or ghrelin (bottom). (b) Statistical evaluation of c-*fos* immunohistochemistry of mouse arcN (two mice per group, three sections each). (c) Relative increase in plasma IGF-I levels after i.p. administration of ghrelin (six mice per group). (d) Immunohistochemistry for ghrelinpositive cells in the stomachs of control mice (left) and mice with only one intact *gna11* allele (right). (e) Statistical evaluation of immunochemistry for ghrelin-positive cells (two mice per group, three sections each). (f) Plasma ghrelin levels on postnatal day 20 (eight mice per group). (b, c, e, and f) **, $P < 0.005$; ***, $P < 0.001$.

 Ga_{11} allele (Fig. 5c and d) but that the total number of arcN neurons, as determined by NeuN immunohistochemistry, was not changed (not shown). These findings suggest that the loss of Ga_{q}/Ga_{11} -mediated signaling in the arcN leads to impaired GHRH production, which in turn causes impaired somatotroph proliferation and, consecutively, dwarfism.

The peptide hormone ghrelin, in addition to having a regulatory function in food uptake and energy homeostasis, is a potent inducer of hypothalamic GHRH release, and the ghrelin receptor GHS-R was reported to couple to $G_{q/11}$ family G proteins (12, 22). We therefore tested whether impaired GHS-R signaling might contribute to reduced GHRH release and weight loss in mice with only one intact *gna11* allele. Since ghrelin administration is known to cause c-*fos* expression in the arcN of rodents (8, 30, 33), we performed c-*fos* immunohistochemistry after intraperitoneal ghrelin application and found that the number of activated neurons in these mice was reduced to 65% of the number in controls (Fig. 6a and b). In contrast to its effect on wild-type animals, ghrelin administration did not increase plasma IGF-I levels in mice with only one *gna11* allele (Fig. 6c), indicating that ghrelin-induced GH release is inhibited in these mice. We then tested whether impaired GHS-R signaling led to a compensatory up-regulation of ghrelin production and found both the number of ghrelinimmunoreactive cells in the stomach (Fig. 6d and e) and the plasma ghrelin levels (Fig. 6f) to be increased. These findings suggest that defective signaling via the ghrelin receptor GHS-R contributes to deregulation of somatic growth and food uptake in mice with only one intact *gna11* allele.

DISCUSSION

We show in this study that conditional inactivation of $G_{q/11}$ mediated signaling in the nervous system leads to different degrees of impairment, depending on the number of intact *gnaq*/*gna11* alleles. Newborn mice which lacked all four alleles did not show any gross morphological abnormalities but did not take up rhythmic breathing activity. This outcome suggests that Ga_{q}/Ga_{11} -mediated signaling is crucial for the postnatal function of the nervous system but not for its prenatal development. Pups carrying one intact *gna11* allele survived the early postnatal period but died as underweight dwarfs between the third and sixth postnatal weeks. In contrast, mice which inherited one intact *gnaq* allele were viable and fertile. *gnaq* and *gna11* are obviously equally able to guarantee vital functions in neonates, but normal postnatal development and weight gain depend on the presence of an intact *gnaq* allele. Two factors might account for these differences between Ga_q and Ga_{11} : functional differences and differential expression patterns of the two proteins. Potential functional differences between $G\alpha_{q}$ and $G\alpha_{11}$ have been intensively studied, but up to now, neither differential receptor coupling (34, 38) nor differences with regard to effector preferences (7, 29, 38) have been described. Our data show that total $G\alpha_q/G\alpha_{11}$ immunoreactivity is much lower in hypothalami from mice with only one intact Ga_{11} allele than in hypothalami from mice with one intact G α_{q} allele. Though both G α_{q} and G α_{11} are basically ubiquitously expressed in the central nervous system (26), differences in expression levels have also been described for other brain regions, e.g., in the cerebellum or the hippocampus, in which expression levels of $G\alpha_{11}$ are also lower than those of Ga_{q} (6, 11, 15). Thus, a single *gna11* allele, but not a single *gnaq* allele, might be unable to compensate for the loss of the other alleles. This finding is in line with observations for constitutively *gnaq*- and/or *gna11*-deficient mice, in which the presence of either two *gna11* alleles or one *gnaq* allele ensures normal cardiomyocyte development (17), craniofacial development (10, 17), and melanoblast migration and proliferation (32).

We showed that growth retardation in mice with only one intact *gna11* allele is due to a massive hypoplasia of anterior pituitary somatotroph cells, resulting in GH deficiency. The anterior pituitary is a derivative of the oral ectoderm and is not of neuronal origin; hence, it does not undergo *Nestin*Cre-mediated recombination (Fig. 4d) (31). The defect leading to somatotroph hypoplasia is therefore unlikely to reside in the pituitary but rather involves brain regions which regulate somatotroph proliferation, e.g., the hypothalamus. In rats, somatotroph cells are first detected at embryonic day 18.5 and rapidly increase in number during the first 10 days after birth (27). An important stimulator of somatotroph proliferation is GHRH, and the overexpression of GHRH leads to somatotroph hyperplasia, increased growth, and tumorigenesis, whereas inactivating mutations cause somatotroph hypoplasia and dwarfism (4). Our data show that impaired somatotroph proliferation in mice with only one *gna11* allele is accompanied

by decreased production of GHRH in the hypothalamus and that exogenous application of GHRH can restore normal proliferation.

The synthesis and release of GHRH from arcN neurons are modulated by a variety of neuropeptides and neurotransmitters, e.g., somatostatin, galanin, neuropeptide Y, and the peptide hormone ghrelin (1). Ghrelin enhances GHRH release via the activation of the GHS-R, which was shown to activate protein kinase C and to release calcium from intracellular stores via the pathway mediated by phospholipase C - β and IP₃ (12). However, several other effectors seem to contribute to GHS-R function, like adenylyl cyclase-mediated cyclic AMP production or L-type Ca^{2+} channels (12, 22). We showed that impaired hypothalamic $G_{q/11}$ signaling led to a significant reduction of ghrelin-mediated c-*fos* activation in arcN neurons, suggesting that $G_{q/11}$ proteins are indeed crucial for the signal transduction of the GHS-R. This notion is corroborated by the finding that ghrelin-induced IGF-I surges are inhibited in mice with only one intact *gna11* allele. The latter finding also suggests that the pituitary GHS-R (5, 9), which should signal normally because *gnaq* is not inactivated in the anterior pituitary of these mice, does not play a major role in ghrelinmediated GH release. This finding is in line with the observation that GHRH antiserum inhibits the GH response to ghrelin in rats (28) and that patients with hypothalamopituitary disconnection do not respond with GH release to synthetic GHS-R ligands (19).

Impaired proliferation of the somatotroph cell line explains very well the dwarf phenotype but not the premature deaths of these mice. Dwarfism itself is not expected to be lethal, but it might handicap the pup in competing for the dam's milk or in reaching the food tray. However, even in very small litters and when food and water were put directly into the cage, the survival of dwarf mice was not improved. In addition to being of short stature, these mice are also impaired with regard to motor development, i.e., they develop an ataxia resembling the one seen in constitutively Ga_q -deficient mice (15). However, constitutively Ga_q -deficient mice have, except for an increased perinatal mortality due to defective primary hemostasis (16), a normal life expectancy (15), and therefore, ataxia cannot account for the premature deaths of mice with only one intact *gna11* allele. We hypothesized that additional defects must exist in these mice, but neither levels of other pituitary hormones in plasma (Fig. 3) nor histological analyses of internal organs revealed any abnormalities. However, stomach weight, size of retrorenal fat deposits, and food uptake were reduced in mice with only one intact *gna11* allele (Fig. 2), suggesting that they suffer from a deregulation of energy homeostasis and/or appetite control. Ghrelin, besides having a role as a growthhormone secretagogue, is a well-known regulator of food intake and energy expenditure (12), and impaired hypothalamic ghrelin signaling is expected to result in dwarfism and reduced food intake. Surprisingly, genetic inactivation of the ghrelin gene did not affect growth or food intake in mice (25, 37). On the other hand, a small interfering RNA-based knockdown of the GHS-R in rats impaired somatic growth, weight gain, and food uptake (21). These studies suggest that the GHS-R, but not ghrelin itself, is indispensable for normal regulation of growth and energy homeostasis. It is quite possible that parallel hormonal systems, known or unknown, are involved in the

regulation of these vital processes, and our findings strongly suggest that they converge on Ga_q/Ga_{11} -coupled receptors, including the GHS-R.

Taken together, our data show that the complete loss of Ga_q/Ga_{11} -mediated signaling in the developing nervous system does not cause obvious developmental abnormalities but leads to functional defects incompatible with extrauterine life. The presence of a single *gnaq* allele or a single *gna11* allele is sufficient to maintain vital functions after birth, but only the presence of a *gnaq* allele, not of a *gna11* allele, allows normal postnatal proliferation of the somatotroph cell line. In addition to showing somatotroph hypoplasia and consecutive dwarfism, mice with only one intact *gna11* allele show abnormal eating behavior, and our findings suggest that both defects involve impaired signaling via the GHS-R.

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