

Original Paper

Luteinizing Hormone and GATA4 Action in the Adrenocortical Tumorigenesis of Gonadectomized Female Mice

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Key Words

Lhcgr • GATA4 • Adrenocortical tumors • Molecular mechanisms • Transgenic mice

Abstract

Background/Aims: Physiological role of luteinizing hormone (LH) and its receptor (LHCGR) in adrenal remains unknown. In inhibin- α /Simian Virus 40 T antigen (SV40Tag) (inh α /Tag) mice, gonadectomy-induced (OVX) elevated LH triggers the growth of transcription factor GATA4 (GATA4)-positive adrenocortical tumors in a hyperplasia-adenoma-adenocarcinoma sequence. **Methods:** We investigated the role of LHCGR in tumor induction, by crossbreeding inh α /Tag with *Lhcgr* knockout (LuRKO) mice. By knocking out *Lhcgr* and *Gata4* in C α 1 adrenocortical cells (*Lhcgr-ko*, *Gata4-ko*) we tested their role in tumor progression. **Results:** Adrenal tumors of OVX inh α /Tag mice develop from the hyperplastic cells localized in the topmost layer of zona fasciculata. OVX inh α /Tag/LuRKO only developed SV40Tag positive hyperplastic cells that were GATA4 negative, cleaved caspase-3 positive and did not progress into adenoma. In contrast to *Lhcgr-ko*, *Gata4-ko* C α 1 cells presented decreased proliferation, increased apoptosis, decreased expression of *Inha*, *SV40Tag* and *Lhcgr* tumor markers, as well as up-regulated adrenal- and down-regulated sex steroid gene expression. Both *Gata4-ko* and *Lhcgr-ko* C α 1 cells had decreased expression of steroidogenic genes resulting in decreased basal progesterone production. **Conclusion:** Our data indicate that LH/LHCGR signaling is critical for the adrenal cell reprogramming by GATA4 induction prompting adenoma formation and gonadal-like phenotype of the adrenocortical tumors in inh α /Tag mice.

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Introduction

The embryonic adrenogonadal primordium (AGP) is a source of progenitors for three major steroidogenic organs: ovaries, testes and adrenal cortex [1]. AGP cells differentiate into gonadal and adrenal primordium and later into independent organs [1] although, recently AGP-like cells were found in the capsule of the mouse adrenal cortex [2]. Therefore disrupted endocrine homeostasis, e.g. due to gonadectomy (GDX) may promote the differentiation of AGP progenitors into neoplastic cells bearing gonadal-like phenotype [2]. Most of the wild-type (WT) mouse strains develop sub-capsular, non-steroidogenic, spindle-shaped A cells when aging [3, 4]. Conversely, adrenal glands of gonadectomized certain inbred (DBA/2J, NU/J) and transgenic mice (21-OH-GATA-4, *Inha*^{-/-}) develop steroidogenically active adrenocortical tumors [3, 5, 6]. Therefore, they are referred to as mouse models that are genetically susceptible to gonadectomy-induced adrenocortical tumorigenesis (GIACT) [3, 5-8] and this phenotype is stronger in females than in males [3, 6, 9].

The hallmark of neoplastic A cells and adrenocortical tumor cells is the expression of transcription factor GATA4 which does not occur in healthy adrenal cells [3, 6]. On the other hand, GATA6, another member of the GATA transcription factor family, regulates adrenal steroidogenic cell differentiation [10]. Interestingly, the interplay between GATA6 and GATA4 transcription factors in adrenal pathophysiology has been suggested to determine the cell fate [10-13]. The lack of GATA6 in the adrenal gland promotes A cell formation [10], whereas GATA4 is pivotal for adrenal tumor onset in DBA/2J and 21-OH-GATA-4 mice [6, 14, 15]. Additionally, in *Inha*^{-/-} mouse model, it has been proposed that the development of tumorous and gonadal-like phenotypes of adrenal tumors is a consequence of the switch from GATA6 to GATA4 expression triggered by elevated gonadotropins [9].

Transgenic mice expressing the Simian Virus (SV40Tag) T-antigen under the inhibin- α promoter (*inh α /Tag*) [8] belong to the group of GIACT models. Both male and female transgenic mice present with gonadal tumors [16], but when prepubertally GDX they develop adrenocortical tumors in a hyperplasia-adenoma-carcinoma sequence by the age of 5-8 months [8, 17, 18]. In this model, genetic or chemical gonadotropin ablation prevents tumor formation [19], whereas transgenic elevation of LH results in the occurrence of gonadal and adrenocortical tumors [20]. An adrenocortical tumor cell line (α 1) has also been established from an *inh α /Tag* female founder [8] and extensively characterized [8, 17, 19]. Similar to other GIACT models, *inh α /Tag* adrenocortical tumors and α 1 cells express an array of gonadal factors including *Lhcgr* and transcription factor GATA4 (*Gata4*) [3, 8, 17, 21]. However, the direct involvement of LHCGR and GATA4 in the adrenocortical tumor induction, progression and phenotype, as well as their inter-relationship in these processes, remains unknown.

In the present study, we characterized the functional role of LHCGR and GATA4 expression in adrenocortical tumorigenesis in ovariectomized (OVX) *inh α /Tag* mice *in vivo* by crossbreeding them with LHCGR knockout mice (LuRKO), as well as *in vitro* by knocking out these genes in α 1 cells.

Materials and Methods

Experimental animals

Inh α /Tag females and control C57Bl/6N (WT) littermates were ovariectomized (OVX) at 21-24 days of age under isoflurane (2-4%) anesthesia (Isoflo, Orion Pharma, Turku, Finland). Temgesic (buprenorphine, 0.1 mg/kg/8 h) (Schering-Plough, Brussels, Belgium) and Comforion® (ketoprofen, 5 mg/kg/24h) (Orion, Turku, Finland) were administered subcutaneously as pre-operative and a post-operative analgesia respectively. 2-, 4-, 6-mo old intact and OVX, WT and *inh α /Tag* animals (n=8-14/group/for all the groups) were sacrificed by terminal cardiac puncture blood collection under isoflurane anesthesia. Tissues were weighted, adrenals snap-frozen in liquid nitrogen and/or fixed with 4% paraformaldehyde (PFA). Blood was collected into a tube consisting 0.5 M sterile EDTA solution, centrifuged at 3000 RPM for 10 minutes in

4 °C and plasma was stored in -80 °C. To elucidate LHCGR involvement in the adrenal tumorigenesis *in vivo*, we crossbreed *inhα/Tag* mice with Luteinizing hormone Receptor Knockout (LuRKO) mice [22]. Samples from OVX: WT, *inhα/Tag*, LuRKO and *inhα/Tag/LuRKO* mice were collected at the age of 7-mo as described above. Experimental mice were housed in a room with controlled light (12 h light and 12 h darkness) and temperature (21±1 °C) in a specific pathogen-free surrounding. Mice were fed with commercial mouse chow SDS RM-3 (Witham, Essex, UK) and tap water *ad libitum*. The State Provincial Office of Southern Finland approved all the animal experiments.

Gene expression analysis

Total RNA from cells and tissues was isolated using phenol/chloroform method with TRIsure (Bioline Reagents Ltd., London, UK) reagent according to the manufacturer's protocol. Prior to qPCR, 900 ng of total RNA was DNase I treated (#18068015, Thermo Fisher Scientific, Waltham, MA) and transcribed (1 h in 48 °C) using SensiFAST™ cDNA Synthesis Kit (#BIO-65053, Bioline Reagents Ltd.). qPCR was carried out on a CFX Real Time PCR Detection System (Bio-Rad, Vienna, Austria), using the DyNAmo™ Flash SYBR® Green qPCR Kit (#F415, Thermo Fisher Scientific) with 7.5ng of cDNA template in total reaction volume of 10 µl in duplicates. Conditions were as follows: 95 °C for 7 min, [95 °C for 15 s, 54-62 °C for 15 s, 72 °C for 15s]x40, 72 °C for 3 min, 65-95 °C melt curve. Primer sequences are presented (for all online suppl. material, see www.karger.com/doi/10.1159/000481718) in supplementary Table S1. Gene expression was calculated using qBase MExcel VBA applet [23]. In brief, Ct of gene of interest was normalized by geometric mean of 2-3 reference genes: peptidylprolyl isomerase A (*Ppia*), β-glucuronidase (*Gusb*), hypoxanthine phosphoribosyltransferase (*Hprt1*), peptidylprolyl isomerase B (*Ppib*) and hydroxymethylbilane synthase (*Hmbs*) validated separately for each experiment using Bio-Rad CFX Manager Software (Bio-Rad).

Immunohistochemistry

PFA fixed paraffin embedded adrenal glands from all experimental group females (n=4-6/ group) were cut into 4±1 µm sections and immunohistochemically stained with antibodies and reagents listed (see supplementary material) in supplementary Table S2. Antigens were retrieved using high-temperature antigen retrieval method in citrate buffer (pH6), washed in TBS buffer with 0.1% Tween20 (#P1379, Sigma-Aldrich, Saint Louis, MO) and visualized using Liquid DAB+ Substrate Chromogen System (Dako, Glostrup, Denmark). Slides were scanned by Panoramic 250 Slide Scanner (3DHISTECH Ltd., Budapest, Hungary).

Semi-quantitative densitometric analysis

Optical density of the antibody staining was calculated from minimum 5 images of good-quality representative areas of OVX *inhα/Tag* and OVX WT adrenals (n=4/group). Images were acquired with Panoramic Viewer (3DHISTECH Ltd.) at 50X magnification and analyzed by Fiji [24] using Hematoxylin-Diaminobenzidine (H-DAB) color deconvolution and DAB units of intensity (color 2) were counted. Optical density (OD) values were calculated using formula $OD = \log(255/\text{Mean intensity})$.

Hormone measurement

Plasma concentration of LH was measured by immunofluorometric (Delfia; Perkin Elmer, Turku, Finland) assay as described previously [25]. The intra- and inter- assay coefficients for LH were below 10%. Plasma and cell culture medium progesterone were analyzed by Elecsys® Progesterone II assay (Roche Diagnostics, Basel, Switzerland), using Cobas e411 immunoanalyzer (Roche Diagnostics). Detection limit for progesterone was 0.10 nmol/l.

CRISPR/Cas9 mediated knockout

All-in-one pLV-U6gRNA-Ef1aPuroCas9GFP plasmids with targeting *Lhcgr* (MM0000334875, MM0000334894), *Gata4* (MM0000234246, MM0000234282) and non-targeting guide sequence (#CRISPR12-1EA, control) were obtained as custom-made plasmids from Sigma- Aldrich. Packaging cell line 293FT was transfected with a mixture of 2.6 µg of plasmid and 26 µl of MISSION® Lentiviral Packaging Mix (Sigma- Aldrich) using FuGENE® 6 Transfection Reagent (#E2691, Promega) according to the provided protocol. After 72 h, viral particles were concentrated for 2 h in 25.000 rpm and snap frozen in liquid nitrogen. Cα1 cells were seeded 50000 cells/well on 24-well plate and let to attach over-night. Next day, lentiviral stocks were added in low volume of full media and incubated for 16 h. Single GFP positive cells were isolated by fluorescence-activated cell sorter (FACS) (BD FACSAria II, Becton Dickinson, Franklin Lakes, NJ) into microtiter plates to establish clonal cell populations.

Genotyping of knockout clones

Clones were cultured in 10% FCS medium, and genotyped using qPCR (primer sequences, see supplementary material, Table S1). Amplicons were separated by electrophoresis on 4% agarose gels (#17856, Thermo Fischer Scientific) and amplicons with significant size difference were isolated from the gel (#28704, QIAquick Gel Extraction Kit, Qiagen) and Sanger sequenced (The Finnish Microarray and Sequencing Centre, Turku, Finland). Three mutated clones for *Lhcgr* and *Gata4* were selected for further experiments (see supplementary material, sequences in Suppl. Fig. S3) and analyzed separately.

In vitro experiments

Mycoplasma-free Cα1 murine adrenocortical cell line [8], and 3 mutated clones/gene were cultured in phenol red-free DMEM F-12 medium (Sigma- Aldrich) supplemented with 10% heat-inactivated fetal calf serum (FCS) and 5 U/ml of penicillin/streptomycin (#15140-122, Gibco) (full culture medium). Results from 3 mutated clones for *Lhcgr* (*Lhcgr*-ko) or *Gata4* (*Gata4*-ko) were pooled and compared with the Cas9 expressing Cα1 cells (mock).

Cell proliferation rate was analyzed using CyQUANT® Cell Proliferation Assay kit (C7026, Life Technologies) in 8-plicates according to manufacturer's protocol. In brief, Cα1 (passages 7, 11, 20), mock and ko cells were seeded on 96 well plates, 5000 cells/well in DMEM/F12 supplemented with 0.5% steroid stripped serum (SFS). One million cells of each cell line was harvested to prepare a cell line-specific DNA standard curve. Cells were grown in medium containing 2.5% SFS without or supplemented with human recombinant hCG (50ng/ml) for 48 h (Serono, Geneva, Switzerland). Fluorescence signal was read using Wallac 1420 Victor2 Microplate Reader (Perkin Elmer, Turku, Finland). Standard curves and interpolated values were obtained using GraphPad Prism.

Caspase 3/7 activity was assessed using Caspase Glo 3/7 kit (Promega) according to the provided protocol. In brief, cells were seeded 8000/well on 96 well plate and assay was performed after 16 h of incubation. Absolute luminescence was normalized by Cα1-mock values.

For progesterone production measurements cells were seeded on 24-well plates (80000/well) in phenol red-free DMEM F-12 medium supplemented with 0.5% SFS. Medium was collected after 24h and stored at -8 °C until the analysis.

For the measurement of cAMP production cells were seeded on 24-well plates (120000/well) in medium containing 0.5% SFS and allowed to attach over-night. Next day, medium was changed for medium containing 0.5% SFS without (C) or with 1, 10, 50 ng/ml of recombinant hCG or 1 μM forskolin (FRK) (Sigma-Aldrich). Medium was collected after 1 h stimulation, mixed 1:1 with 2 mM teophiline (Sigma- Aldrich), boiled for 5 minutes in a water bath and frozen in -20 °C. Extracellular cAMP concentration was assessed by a standard radioimmunoassay method after acetylation according to the method of Harper and Brooker [26, 27]. Pellet radiation was read using γ-counter and automatically subtracted to a standard curve (Wallac 1470 Wizard Gamma Counter, Perkin Elmer).

Statistical analysis

To test the statistical differences between 2 or more experimental groups Mann-Whitney U or Kruskal-Wallis with multiple comparison of mean range as post hoc tests were used respectively. All numerical data were presented as mean ± SEM. Graphs and statistical analysis were done with Graph Pad Prism 5 (GraphPad Software, San Diego California USA), values of P<0.05 were considered to be significant.

Results

Formation of the adrenocortical tumors in $Inh\alpha/Tag$ mice

Total adrenal (Fig. 1A) and tumor weights of OVX $inh\alpha/Tag$ females (Fig. 1B; calculated by subtraction of the average total adrenal weight of age-matched OVX WT from total adrenal weights of $inh\alpha/Tag$ mice) were significantly increased from 4-mo and continued at 6-mo vs. age-matched controls. Using histological and immunohistochemical staining criteria, two distinct types of neoplastic cells expressing SV40Tag were identified at 2-mo of age (Fig. 1C and see supplementary material Fig. S1A). The first abnormal cells in the intact and OVX $inh\alpha/Tag$ mouse adrenals were the spindle-shaped neoplastic A cells (Fig.

1D and see supplementary material Fig. S1C), originating from the sub-capsular region of the adrenal gland, and migrated towards the peri-medullary region. As observed in WT, these cells did not form tumors (see supplementary material, Fig. S1B, S1D). The second type of neoplastic cells observed had round nuclei (Fig. 1E), and was exclusively found in OVX *inhα*/Tag already at 2-mo, formed hyperplasia in the topmost layer of zona fasciculata, and resulted in a subsequent tumor formation. Both A and hyperplastic cells were SV40Tag, GATA4 and MKI67 positive (Fig. 1D-E). Representative pictures were taken from adjacent slides of the same specimen.

Tumor cells lose GATA6 and gain GATA4 expression during their ontogeny

The reciprocal expression of GATA6 and GATA4 in mouse models for adrenocortical tumorigenesis has been previously reported [9, 17], and was suggested to underpin the transition from adrenal-to-gonadal phenotype observed in these tumors [9]. We therefore examined the age-related GATA6 and GATA4 localization during the ontogeny of the adrenocortical tumors. GATA6 expression was abundant in the normal adrenal cortex of OVX

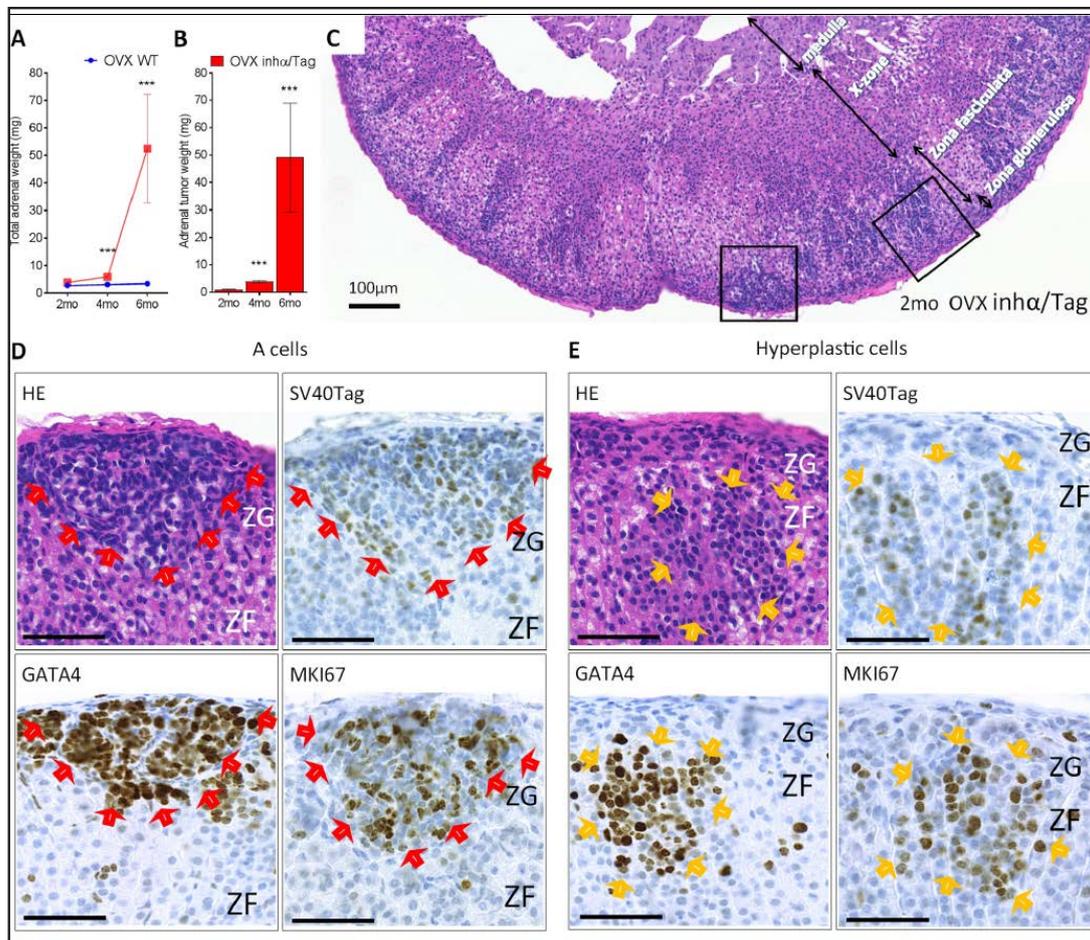


Fig. 1. Onset of adrenocortical tumors in *inhα*/Tag mice. (A) Age dependent adrenal gland mass of intact and OVX WT and *inhα*/Tag females (mean ± SEM; ***. $P \leq 0.001$; $n=8-14$ /group). (B) Approximate adrenal tumor weight calculated by average total adrenal weight of age-matched OVX WT subtracted from total adrenal weights of *inhα*/Tag mice (mean ± SEM; ***. $P \leq 0.001$; $n=8-14$ /group). (C) Histopathology of the 2-month-old OVX female with the foci of (D) sub-capsular A cells and (E) hyperplastic cells stained for adrenal neoplastic cell biomarkers SV40Tag, GATA4 and proliferation marker MKI67 ($n=4$ /group). Bar= 50μm; ZG- zona glomerulosa, ZF- zona fasciculata, red arrows- A cells; yellow arrows- hyperplastic cells; OVX, subjected to prepubertal ovariectomy; WT, wild-type C57Bl/6N; *inhα*/Tag, transgenic mice expressing SV40Tag oncogene under inhibin α promoter.

WT mice (Fig. 2A) and hyperplastic cells of 2-mo OVX inh α /Tag (Fig. 2B), and persisted in some cells within the tumor foci of 4-mo (Fig. 2C) and 6-mo (Fig. 2D) OVX inh α /Tag mice. In turn, GATA4 expression was absent in the normal adrenal cortex of OVX WT mice (Fig. 2F), with expression of GATA4 observed in hyperplastic cells of 2-mo OVX inh α /Tag (Fig. 2G), increasing in abundance within tumor foci of 4-mo (Fig. 2H) and 6-mo (Fig. 2I) OVX inh α /Tag mice. Densitometric analysis of the same region from consecutive sections (5 images/specimen, n=4/ group), showed a significant decrease in GATA6 staining (Fig. 2E) in 4-mo and 6-mo OVX inh α /Tag vs. 6-mo OVX WT adrenals. Moreover, a concomitant increase in GATA4 staining was already present in hyperplastic cells at 2-mo, and continued to increase at 4-mo and 6-mo OVX inh α /Tag vs. 6-mo OVX WT (Fig. 2J).

Lack of LHCGR prevents GATA4 induction in SV40Tag-positive cells and adrenocortical tumor formation

To evaluate the role of LHCGR in the adrenocortical tumor induction in the OVX inh α /Tag model, we crossbred inh α /Tag mice with *Lhcgr* knockout (LuRKO) [22] mice (inh α /Tag/LuRKO). At 7-mo of age, OVX inh α /Tag/LuRKO mice had significantly decreased total adrenal

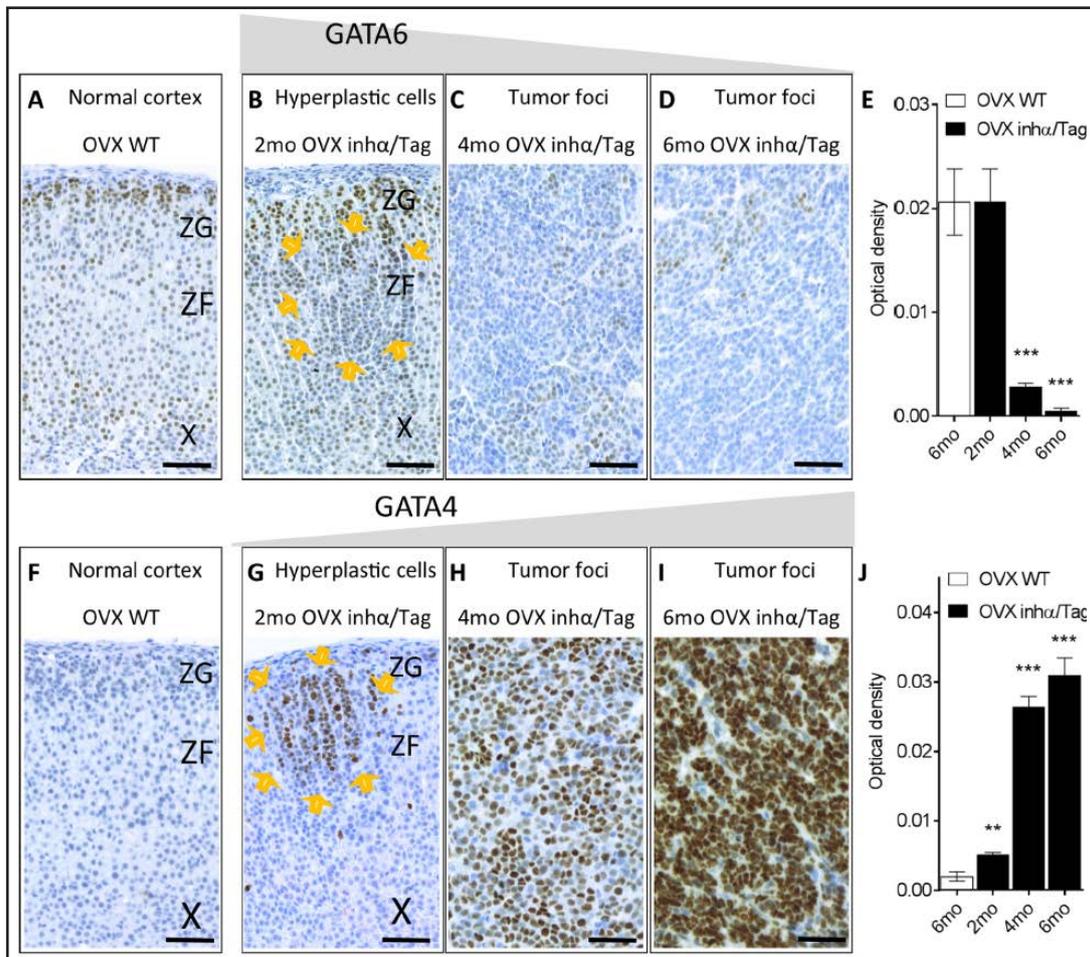


Fig. 2. GATA6 and GATA4 expression during the ontogeny of adrenocortical tumors in inh α /Tag mice. GATA6 localization in adrenals of 6mo OVX WT (A), hyperplastic cells of 2-mo (B), and in tumor foci of 4-mo (C) and 6-mo (D) OVX inh α /Tag mice (n=4/group). GATA4 localization in adrenals of 6-mo OVX WT (F), hyperplastic cells of 2-mo (G), and in tumor foci of 4-mo (H) and 6-mo (I) OVX inh α /Tag mice (n=4/group). Average optical density of GATA6 (E) and (GATA4) (J) staining in 6-mo OVX WT and 2-4-6-mo OVX inh α /Tag adrenals (mean \pm SEM; 5 images/ specimen, n=4/ group; **, P \leq 0.01. ***, P \leq 0.001). Bar= 50 μ m; ZG- zona glomerulosa, ZF- zona fasciculata, X- x-zone, yellow arrows- hyperplastic cells.

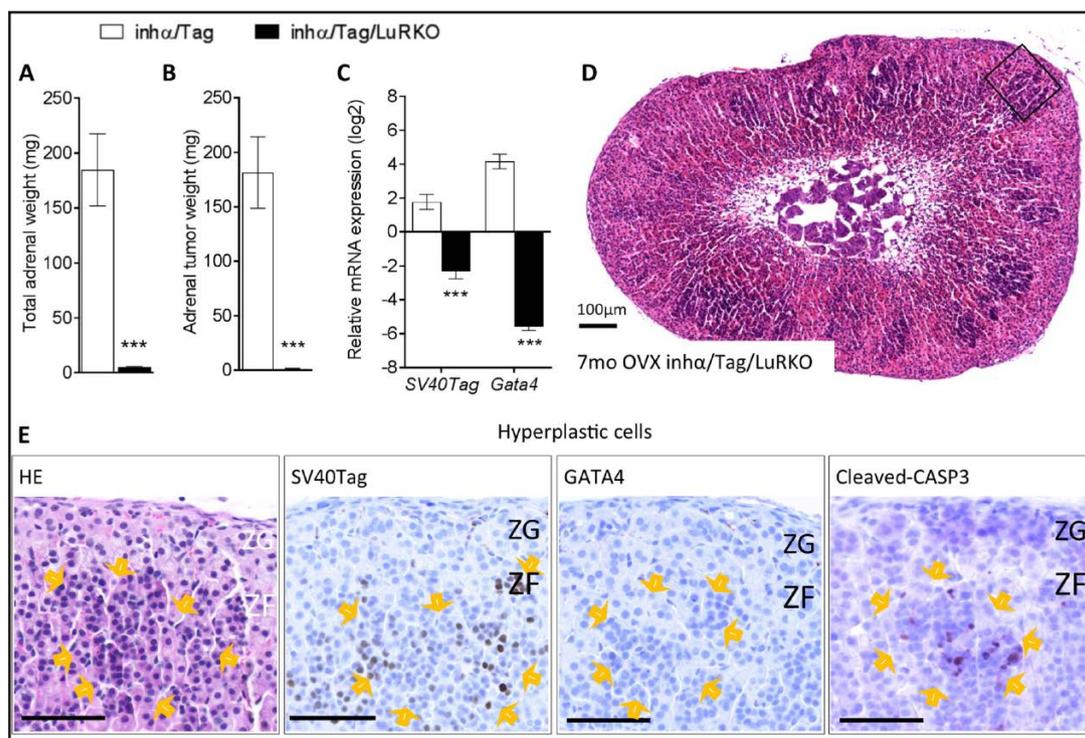


Fig. 3. Analysis of LH/LHCGR signaling in the onset of adrenal tumors in inhα/Tag mice. Comparison of adrenal (A), approximate adrenal tumor weights (B) and tumor marker *SV40Tag*, *Gata4* expression (C) in 7mo-old inhα/Tag and inhα/Tag/LuRKO females (mean ± SEM; n=8/group; ***. P ≤ 0.001). Hematoxylin and eosin staining of 7mo-old OVX inhα/Tag/LuRKO adrenal gland (D), and localization of adrenocortical tumor biomarkers SV40Tag, GATA4 and Cleaved-CASP3 in hyperplastic cells (E) (n=6/ group). Bar= 50μm; ZG- zona glomerulosa, ZF- zona fasciculata, yellow arrows- hyperplastic cells; OVX, subjected to prepubertal ovariectomy; inhα/Tag, transgenic mice expressing SV40Tag oncogene under inhibin α promoter; inhα/Tag/LuRKO, inhα/Tag mice crossbred with LH deficient mice (LuRKO).

(Fig. 3A) and adrenal tumor weights (Fig. 3B), as well as down-regulated expression of the tumor markers *SV40Tag* and *Gata4*, in comparison to OVX inhα/Tag mice (Fig. 3C). Plasma concentrations of LH were similar (see supplementary material, Fig. S2A) in all the groups however, progesterone was increased in OVX inhα/Tag, and decreased in both OVX inhα/Tag/LuRKO and OVX LuRKO, when compared with OVX WT females (see supplementary material, Fig. S2B). The expression of neoplastic cell biomarker *Gata4* was significantly decreased in OVX LuRKO mice vs. OVX WT (see supplementary material, Fig. S2C). As *SV40Tag* transcripts were detected in the adrenal glands of OVX inhα/Tag/LuRKO mice, we further analyzed their histopathology. The hematoxylin and eosin staining of 7-mo inhα/Tag/LuRKO adrenals (Fig. 3D) revealed the presence of hyperplastic cells (Fig. 3E), similar to those previously found in 2-mo inhα/Tag adrenals (Fig. 1E). We immunolocalized tumor biomarkers SV40Tag and GATA4, and to detect potential apoptotic cells, active caspase-3 (Cleaved-CASP3). Hyperplastic cells were SV40Tag and Cleaved-CASP3 positive, but GATA4 negative (Fig. 3E). Taken together, the absence of GATA4 in hyperplastic cells (Fig. 3E) could result in the decreased *Gata4* expression in the adrenals (Fig. 3C) of 7-mo inhα/Tag/LuRKO mice.

GATA4 induces the tumorous phenotype, but both LHCGR and GATA4 regulate steroidogenesis in Cα1 cells

We analyzed next the gene expression profile of *Gata4* and *Lhcgr* markers, and steroidogenic enzymes in 6-mo OVX inhα/Tag and Cα1 cells. *Lhcgr* and *Gata4* expression were significantly increased in both 6-mo OVX inhα/Tag and Cα1 cells when compared

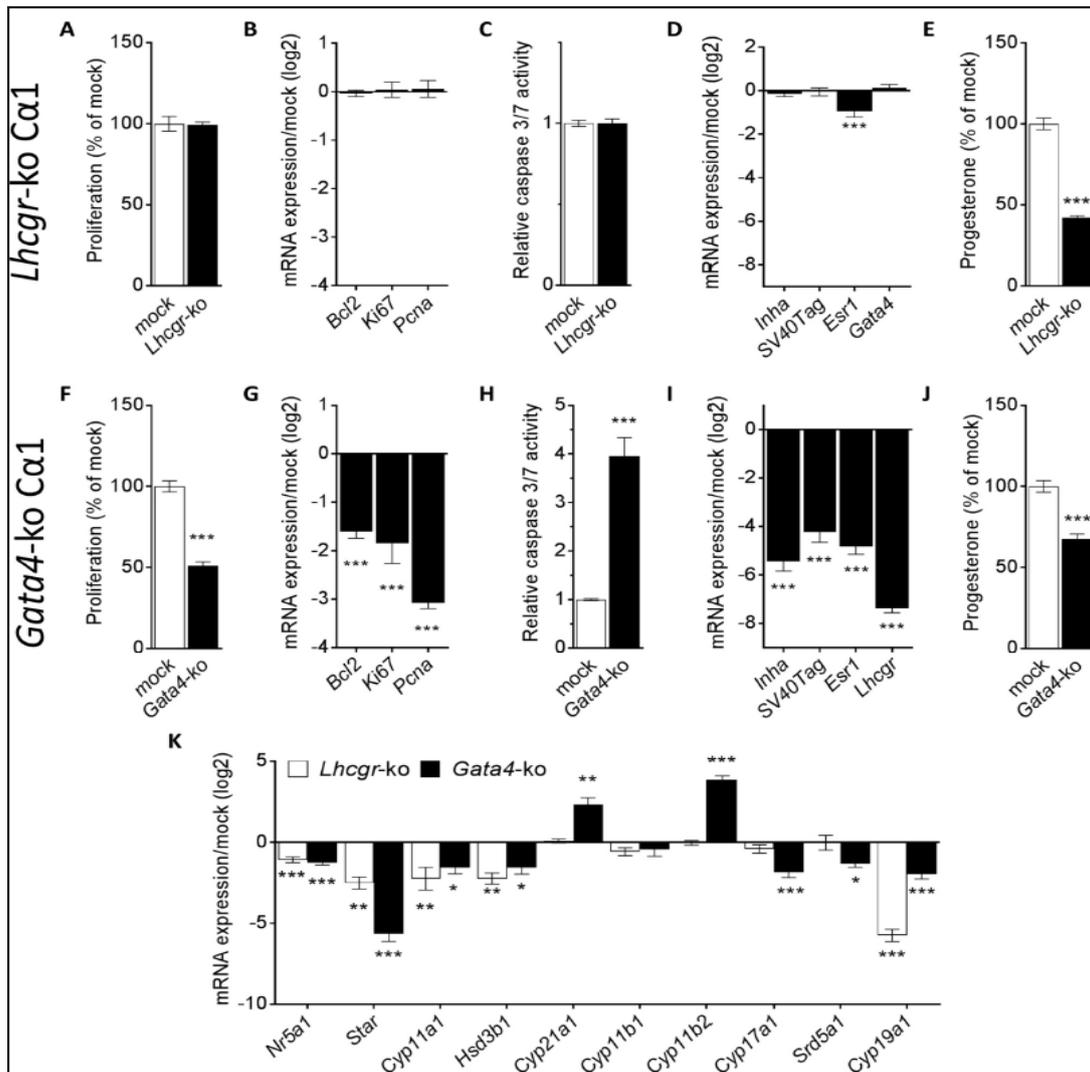


Fig. 4. Effects of *Lhcgr* and *Gata4* knockout in Ca1 cell line. Characterization of *Lhcgr-ko* and *Gata4-ko* cells by proliferation (A, F) gene expression of pro-survival *Bcl2*, proliferation markers *Mki67* and *Pcna* (B, G), active caspase 3/7 levels (C, H), tumor cell markers *Inha*, *SV40Tag*, *Lhcgr* and *Esr1* expression (D, I) and progesterone production (E, J) (mean \pm SEM; n=5-8/group). Comparison of steroidogenic enzymes gene expression in *Lhcgr-ko* and *Gata4-ko* cells vs. mock (K) (mean \pm SEM; n=9/ group). *. $P \leq 0.05$. **. $P \leq 0.01$. ***. $P \leq 0.001$.

to 6-mo OVX WT (see supplementary material, Fig. S3A). Moreover, Ca1 cells expressed functional LHCGR as shown *via* the dose dependent hCG stimulation of cAMP production (see supplementary material, Fig. S3B). However, hCG stimulation did not affect Ca1 cell proliferation (see supplementary material, Fig. S3C). Steroidogenic gene expression profiling showed up-regulation of the proximal *Star* (only in Ca1 cells) and *Cyp11a*, up-regulation of gonad-specific *Srd5a1* and *Cyp19a1*, and down-regulation of adrenal-specific *Cyp21a1*, *Cyp11b1* and *Cyp11b2* in comparison to OVX WT (see supplementary material, Fig. S3D).

To analyze the impact of LHCGR and GATA4 on tumor progression, we used CRISPR/Cas9-mediated mutagenesis to knockout these genes in Ca1 cells. Representative amplicon lengths of mutated clones as well as the actual sequences of *Lhcgr* and *Gata4* mutants used in the experiments are shown in (see supplementary material) Fig. S4A-B and Fig. S4C-D, respectively. Functional LHCGR knockout was confirmed by abrogation of hCG-mediated cAMP production in all 3 independent *Lhcgr-ko* clones (see supplementary material, Fig.

S4E). For all the experiments 3 different clones (see supplementary material, with mutations indicated in Fig. S4B and Fig. S4D) of *Lhcgr*-ko or *Gata4*-ko were used, results were pooled and compared with a mock-transfected cell line (control).

Lhcgr-knockout had no effect on either the proliferation of α 1 cells (Fig. 4A), expression of their anti-apoptotic *Bcl2* and proliferation markers *Pcna* and *Mki67* (Fig. 4B) or caspase 3/7 activity (Fig. 4C) in comparison to control. Gene expression analysis of the adrenocortical tumor biomarkers in *Lhcgr*-ko α 1 cells showed down-regulation of *Esr1*. In contrast, expression of *Inha*, *SV40Tag* and *Gata4* remained unaffected (Fig. 4D). *Gata4*-ko α 1 cells had a significantly reduced proliferation rate vs. control (Fig. 4F) reflecting the decreased expression of anti-apoptotic *Bcl2* and proliferation markers *Mki67* and *Pcna* (Fig. 4G). Activity of caspase 3/7 was significantly increased in *Gata4*-ko α 1 cells (Fig. 4H) indicating that these cells undergo apoptosis. Gene expression of tumor cell markers *Inha*, *SV40Tag*, *Lhcgr* and *Esr1* was also decreased in *Gata4*-ko α 1 cells (Fig. 4I).

Basal progesterone production was significantly decreased in both *Lhcgr*-ko (Fig. 4E) and *Gata4*-ko (Fig. 4J) cells vs. mock. Moreover, lack of *Lhcgr* and *Gata4* in α 1 cells significantly down-regulated *Nr5a1*, *Star*, *Cyp11a1*, *Hsd3 β 1* and *Cyp19a1* steroidogenic genes (Fig. 4K). Additionally, in *Gata4*-ko cells, adrenal steroidogenic enzymes *Cyp21a1* and *Cyp11b2* were up-regulated, while gonadal steroidogenic enzymes *Cyp17a1* and *Srd5a* were down-regulated (Fig. 4K), showing enhanced adrenal steroidogenic phenotype.

Discussion

Adrenocortical tumor cells in *inh α /Tag* and *Inha*^{-/-} mice [7, 9] or large lipid-laden steroidogenic neoplastic B-cells in NU/J [5], DBA/2J [3] and 21-OH-GATA4 [6] appear after gonadectomy-induced elevated gonadotropins. It was previously considered that adrenocortical tumors in *inh α /Tag* mice, similarly to those of a complementary *Inha*^{-/-} model, originate from the x-zone [7, 8]. However, a recent report suggests that adrenocortical tumors in *Inha*^{-/-} mice, as well as in other GACT models, may originate from the sub-capsular pluripotent progenitors [9]. To revisit the origin of the *inh α /Tag* adrenocortical tumors we analyzed spatiotemporal expression of *SV40Tag* oncogene and identified the presence of two distinct types of *SV40Tag* positive cells, sub-capsular A and hyperplastic cells in adrenal parenchyma.

The tumor constituting *SV40Tag* expressing hyperplastic cells, at the age of 2-mo, were localized in the zona fasciculata, where a 6kb promoter of *Inha* was shown to be most active in young and adult mouse adrenals [28]. Therefore, we suggest that the adrenocortical tumors in *inh α /Tag* model originate from zona fasciculata, not x-zone [8], nor from the stem/progenitor cell niche [9] as previously suggested. Failure in tumor development in OVX *inh α /Tag/LuRKO* mice, indicate that *SV40Tag* expression alone is able to cause the hyperplasia but is not enough to drive tumor formation. Generally, *SV40Tag* expression inhibits p53 action and enables cells to pass the cell cycle despite DNA damage [29, 30], but may also require additional co-factors to maintain tumor formation [31]. An analogous situation was reported in a pancreatic tumor model expressing *SV40Tag* (RIP1-Tag2), where less than 2% of the islets expressing *SV40Tag* progressed into tumors [32]. In addition, anti-apoptotic BCL2 family members were identified as *SV40Tag* partners for tumor formation [31]. A recent study has shown [33], that expression of *SV40Tag* under a zona fasciculata-specific *Akr1b7* promoter (AdTAG mice), resulted in a development of metastatic adrenal tumors in intact mice by the age of 8 months. Similarly to our model, *SV40Tag* expression was implicated mostly into tumor initiation, whereas spontaneously activated WNT/ β -catenin signaling promoted the malignant progression of adrenocortical tumors in intact mice [33]. In our OVX *inh α /Tag/LuRKO* model, lack of LHCGR and GATA4 expression in the hyperplastic cells resulted in a failure in hyperplasia to adenoma progression, suggesting these proteins as a potential *SV40Tag* partners during tumorigenesis.

Based on the *in vitro* studies, lack of GATA4 had more severe effects on the α 1 tumor cell phenotype, than the lack of LHCGR. In α 1 cell line, we found that GATA4 regulated

proliferation and cell survival, as well as the expression of *Inha*, *SV40Tag* and *Lhcgr*. A recent report from our group showed that transgenic GATA4 expression under adrenal-specific 21-hydroxylase (21-OH) promoter (21-OH-GATA-4 model) was able to trigger tumor formation in C57Bl/6 strain mice, which is non-susceptible to G1ACT [6]. However, similar to the *inh α /Tag* model, gonadectomy was a prerequisite for the tumor formation in 21-OH-GATA-4 mice [6]. On the other hand, knock out of GATA4 in mice susceptible to G1ACT prevented tumor formation [14]. Moreover, GATA4 has been shown to be a potent driver of the tumorous and gonadal phenotype in the G1ATC models [6, 9, 14], as well as a pro-survival factor in gonadal Leydig and granulosa cells [34, 35]. Therefore, GATA4 could be a potent partner for SV40Tag in tumor formation, regulator of anti-apoptotic pathways and gonad-related gene expression. The failure in hyperplasia-adenoma progression in OVX *inh α /Tag/LuRKO* mouse adrenals, supports the involvement of LH as an upstream factor for tumorigenesis in *inh α /Tag* mice [9, 20]. In the complementary *Inha*^{-/-} model, follicle-stimulating hormone (FSH) was proposed to trigger the tumorigenesis by induction of the GATA6 to GATA4 transcription factor switch [9]. In *inh α /Tag* model we see a gradual decrease of GATA6 expression with concomitant increase in GATA4 expression during the adrenal hyperplasia-adenoma transformation, suggesting that the tumorigenesis mechanism could be similar to that observed in *Inha*^{-/-} model.

The lack of effects of hCG or *Lhcgr* knock out on C α 1 cell proliferation suggests that tumor progression is independent on LH/LHCGR signaling in this cell line model. Conversely, the tumor progression and survival was clearly regulated by GATA4. Moreover, no change was observed in *Gata4* expression in *Lhcgr*-ko cells, suggesting that *Gata4* expression is not regulated by LH/LHCGR, in the tumor cells. Therefore the high levels of *Gata4* transcripts in adenoma and C α 1 cells might be an effect of GATA4 auto-regulation as previously shown [36]. Thus, LH/LHCGR signaling appears to be a prerequisite for the adrenocortical cell reprogramming probably through inducing the GATA6 to GATA4 switch, and thereafter GATA4 takes over the transcriptional control of the tumor cell progression.

Localization of SV40Tag in A cells of intact and OVX *inh α /Tag* mice supported the notion that oncogene expression is not sufficient for tumor formation. Moreover, the presence of A cells in OVX LuRKO and OVX *inh α /Tag/LuRKO*, confirmed an earlier report on their gonadotropin-independent appearance [6]. In contrast, a significant decrease of A cell biomarker *Gata4* in OVX LuRKO and OVX *inh α /Tag/LuRKO* when compared to OVX WT adrenals, indicated a decrease in their population when LH/LHCGR signaling was lacking. An increase in A cell population after GDX [37], and *Lhcgr* expression in a sub-population of A cells [21] suggest that LHCGR signaling may affect their proliferation.

Gene expression of gonadal-type steroidogenic enzymes was up-regulated at the expense of reduced adrenal steroidogenesis in the adrenal tumors of OVX *inh α /Tag* females and C α 1 cells. This corresponds to the impaired expression of steroidogenic enzymes in *inh α /Tag* males bearing adrenocortical tumors [21]. *In vitro* studies revealed that both *Lhcgr* and *Gata4* are important regulators of general steroidogenic genes resulting in decreased progesterone production in *Lhcgr*-ko and *Gata4*-ko cells. Also, *in vivo* the absence of functional LHCGR resulted in a decreased plasma progesterone levels supporting the potent role of LHCGR in the steroidogenic activity of OVX mouse adrenal glands [38]. We and others [17], have shown that GATA4 regulates *Lhcgr* expression, therefore it is impossible to excise the direct and indirect (through *Lhcgr* regulation) effects of GATA4 on the general steroidogenic gene expression. In contrast, only in *Gata4*-ko cells, we saw a clear inversion of gonadal-like (*Cyp17a1*, *Srd5a1*, *Cyp19a1*) to adrenal (*Cyp21a1*, *Cyp11b2*) steroidogenic phenotype, consistent with the previous reports that gonadal steroidogenic phenotype in G1ACT models is an effect of GATA4 expression [9, 14]. Our data confirms that LHCGR is a potent steroidogenesis enhancer in the adrenal gland [38], however, GATA4 controls the gonadal steroidogenic phenotype in the adrenocortical tumor cells [9].

In conclusion, the gonadotropin-dependent tumorigenic mechanism in the adrenal gland of *inh α /Tag* mice is activated in the cells of zona fasciculata that express GATA6. Increased LH/LHCGR signaling in definitive adrenocortical cells causes a transcription factor switch

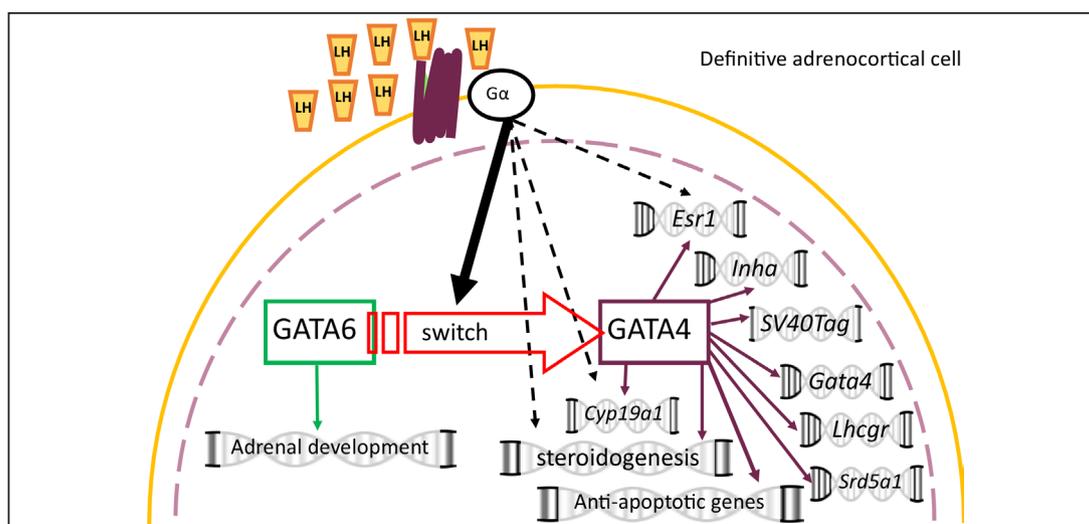


Fig. 5. Possible mechanism of SV40Tag oncogene activation in *inha/Tag* adrenals. Oncogene expression in our *inha/Tag* model is dependent on the activation of inhibin α promoter. Gonadectomy-induced chronically elevated LH levels up-regulate the expression of LHCGR, and LH/LHCGR signaling induces expression of GATA4, which independently of gonadotropins, takes control on the regulation of inhibin α promoter expression and cell survival.

from GATA6 to GATA4. GATA4 enhances strongly *Inha/SV40Tag* expression, the gonadal steroidogenic phenotype and most likely also stimulates the pro-survival pathways resulting in hyperplasia-adenoma transition. Once the transcription factor switch is induced, tumor progression becomes dependent on GATA4 and its downstream targets, but independent of LHCGR signaling (Fig. 5). Whether the process of GATA4 induction in human ACTs is similar to our *inha/Tag* model, is an important question to address in future studies. Furthermore, identification of the direct GATA4 downstream targets in both *inha/Tag* and human ACTs could be of great importance to find potential therapeutic targets for human ACTs.

Abbreviations

A cells (sub-capsular spindle shaped neoplastic non-steroidogenic type 'A' cells); B cells (lipid laden neoplastic steroidogenic type 'B' cells); SV40Tag (Simian Virus 40T antigen; oncogene); GIACT (Gonadectomy-induced adrenocortical tumorigenesis); LHCGR (luteinizing hormone receptor);

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Disclosure Statement

The authors have nothing to disclose.

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