Corrections

COMMENTARY. For the article “Extracting functional information from microarrays: A challenge for functional genomics,” by Michael Q. Zhang, which appeared in number 20, October 1, 2002, of Proc. Natl. Acad. Sci. USA (99, 12509–12511; First Published September 23, 2002; 10.1073/pnas.212532499), Fig. 1 appeared incorrectly. The locants for B and C were reversed. In addition, the locant for the description of D was omitted from the legend. The corrected figure and its legend appear below.

![Figure 1](https://www.pnas.org/cgi/doi/10.1073/pnas.242607999)

Fig. 1. Relations among different concepts in the SP-analysis method. (A) Expression profile matrix (table). \( t = (t_1, t_2, \ldots) \) is the experimental condition index; in this example it indicates a set of time points. (B) Expression profiles (patterns). \( g_1 \) and \( g_4 \) are not strongly correlated directly, but both are strongly correlated with the correlated set \( g_x, g_j \). \( g_1, g_2 \) are the transitive genes interpolating the two terminal genes along SP1 (see C and D); similarly, \( g_y \) is the transitive gene interpolating \( g_1, g_5 \) along SP2. (C) GO biological process tree. The Ps are process annotations for genes at a particular node. A gene may belong to more than one node (“multiple-function,” such as \( g_2 \)). (D) Expression profile space. \( g_3 \) is on the short path SP1 terminated by the known genes \( g_1, g_2, \) and \( g_4 \) and hence is assigned a function of \( P_{1,1,1} \) (level L0) according to the GO tree in C; \( g_y \) is on SP2 terminated by \( g_1, g_5 \) and is assigned a function of \( P_{1,1,1} \) (level L1). \( g_1 \) is shared by both SPs and may be involved in both processes, which means the processes represented by SP1 and SP2 actually crosstalk to each other. The linked gene network can be formed by the subgraph SP1 + SP2.
CHEMISTRY. For the article “Creating nanocavities of tunable sizes: Hollow helices,” by Bing Gong, Huaiqiang Zeng, Jin Zhu, Lihua Yu, Yaohua Han, Shizhi Cheng, Mako Furukawa, Ruben D. Parra, Andrey Y. Kovalevsky, Jeffrey L. Mills, Ewa Skrzypczak-Jankun, Suzana Martinovic, Richard D. Smith, Chong Zheng, Thomas Szyperski, and Xiao Cheng Zeng, which appeared in number 18, September 3, 2002, of Proc. Natl. Acad. Sci. USA (99, 11583–11588; First Published August 12, 2002; 10.1073/pnas.162277099), the accession numbers for the crystal structure data were omitted from the paper. The omitted footnote appears below.

Data deposition: The atomic coordinates have been deposited in the Cambridge Structural Database, Cambridge Crystallographic Data Centre, Cambridge CB2 1EZ, United Kingdom, www.ccdc.cam.ac.uk [reference nos. 188141 (2b), 188142 (4c), 188143 (5b), 188144 (8), and 188145 (9)].

In addition, the author name Lihua Yua should have appeared as Lihua Yuan. The online version has been corrected. The corrected author line appears below.

Bing Gong, Huaiqiang Zeng, Jin Zhu, Lihua Yuan, Yaohua Han, Shizhi Cheng, Mako Furukawa, Ruben D. Parra, Andrey Y. Kovalevsky, Jeffrey L. Mills, Ewa Skrzypczak-Jankun, Suzana Martinovic, Richard D. Smith, Chong Zheng, Thomas Szyperski, and Xiao Cheng Zeng

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IMMUNOLOGY. For the article “Genomic expression programs and the integration of the CD28 costimulatory signal in T cell activation,” by Maximilian Diehn, Ash A. Alizadeh, Oliver J. Rando, Chih Long Liu, Kryn Stankunas, David Botstein, Gerald R. Crabtree, and Patrick O. Brown, which appeared in number 18, September 3, 2002, of Proc. Natl. Acad. Sci. USA (99, 11796–11801; First Published August 23, 2002; 10.1073/pnas.092284399), the figures in the printed article are of low resolution. The online article contains high-resolution figures (www.pnas.org/cgi/reprint/99/18/11796.pdf).

www.pnas.org/cgi/doi/10.1073/pnas.222520199

BIOCHEMISTRY. For the article “Generation and characterization of androgen receptor knockout (ARKO) mice: An in vivo model for the study of androgen functions in selective tissues,” by Shuyuan Yeh, Meng-Yin Tsai, Qingquan Xu, Xiaoming Mu, Henry Lardy, Ko-En Huang, Hank Lin, Shahud-Der Yeh, Saleh Altuwaijri, Xinchang Zhou, Lianping Xing, Brendan F. Boyle, Min-Chie Hung, Su Zhang, Lin Gan, and Chawnshang Chang, which appeared in number 21, October 15, 2002, of Proc. Natl. Acad. Sci. USA (99, 13498–13503; First Published October 7, 2002; 10.1073/pnas.212474399), The author name Min-Chie Hung appeared incorrectly. The correct name is Mien-Chie Hung. The online version has been corrected. The corrected author line appears below.

Shuyuan Yeh, Meng-Yin Tsai, Qingquan Xu, Xiaoming Mu, Henry Lardy, Ko-En Huang, Hank Lin, Shahud-Der Yeh, Saleh Altuwaijri, Xinchang Zhou, Lianping Xing, Brendan F. Boyle, Mien-Chie Hung, Su Zhang, Lin Gan, and Chawnshang Chang

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CELL BIOLOGY. For the article “Untangling the wires: A strategy to trace functional interactions in signaling and gene networks,” by Boris N. Kholodenko, Anatoly Kiyatkin, Frank J. Bruggeman, Eduardo Sontag, Hans V. Westerhoff, and Jan B. Hoek, which appeared in number 20, October 1, 2002, of Proc. Natl. Acad. Sci. USA (99, 12841–12846; First Published September 19, 2002; 10.1073/pnas.192442699), the affiliation of the communicating member appeared incorrectly. The correct “Communicated by” line appears below.

Communicated by Rudolf E. Kalman, Swiss Federal Institute of Technology, Zurich, Switzerland, July 25, 2002 (received for review March 30, 2002)

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Generation and characterization of androgen receptor knockout (ARKO) mice: An in vivo model for the study of androgen functions in selective tissues


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Contributed by Henry Lardy, July 29, 2002

By using a cre-lox conditional knockout strategy, we report here the generation of androgen receptor knockout (ARKO) mice. Phenotype analysis shows that ARKO male mice have a female-like appearance and body weight. Their testes are 80% smaller and serum testosterone concentrations are lower than in wild-type (wt) mice. Spermatogenesis is arrested at pachytene spermatoocytes. The number and size of adipocytes are also different between the wt and ARKO mice. Cancellous bone volumes of ARKO male mice are reduced compared with wt littermates. In addition, we found the average number of pups per litter in homologous and heterozygous ARKO female mice is lower than in wt female mice, suggesting potential defects in female fertility and/or ovulation. The cre-lox ARKO mouse provides a much-needed in vivo animal model to study androgen functions in the selective androgen target tissues in female or male mice.

Androgen receptor (AR), a member of the nuclear receptor superfamily, was first cloned in 1988 (1–3). It contains an N-terminal transactivation domain, a central DNA-binding domain, and a C-terminal ligand-binding domain (4). AR may form a dimer and interact with many coregulators to modulate androgen target genes (5). In addition to its natural ligands, testosterone (T) and dihydrotestosterone, 17β-estradiol can also induce AR transactivation in the presence of some selective coregulators in some selective tissues (6, 7). The increasing evidence shows that androgen and AR may also play important roles in female physiological processes, including folliculogenesis (8), bone metabolism (9), autoimmune diseases (10), maintenance of brain functions (11), and several female cancers, including breast, ovary, and endometrium (12–14).

In the cartilage and bone, AR is expressed in chondrocytes, osteoblasts, osteocytes (15), and osteoclasts (16). Clinical studies suggested that combined therapy of estrogens plus androgens may enhance bone mineral density and bone mass to a more significant degree than estrogen therapy alone in postmenopausal women (17, 18). However, the mechanism of androgen action on bone metabolism remains controversial. Some studies suggest that the effect is mainly through the effects of aromatase that converts androgens to estrogens (19). Other studies show that administration of anti-androgens, including flutamide and casodex, to female mice resulted in osteopenia, suggesting the direct role of AR in bone metabolism (20, 21). As more evidence suggests that estrogens and androgens may have some crosstalk effects, whether estrogen can go through AR, or androgens may go through the estrogen receptor for their physiological functions, remains an interesting question for future studies (22, 23).

Transgenic and knockout (KO) mice have long been used as animal models to study the function of the genes in vivo. For AR, the testicular feminization (Tfm) male mice and the patients with the androgen-insensitive syndrome are the natural models for the study of the loss of androgen function in males (24). However, female models have been lacking, and the classical KO strategy does not work because AR is located in the X chromosome, which is critical for male fertility. To generate tissue-specific ARKO mice or female ARKO mice, a cre-lox strategy for conditional KO is necessary. The cre-lox system utilizes the expression of P1 phage cre recombinase (Cre) to catalyze the excision of DNA located between flanking lox sites (25). This strategy differs from the standard targeted disruption procedure in that embryonic stem (ES) cells are generated in which the targeted segment is not disrupted but flanked by lox sites (floxed). The target gene thus functions normally and mice can be bred to homozygosity for the targeted locus. Here we describe the generation and characterization of ARKO mice. The potential in vivo application of this model for the study of androgen functions in selective tissues is also discussed.

Experimental Procedures

The steps leading to the birth of the female mice carrying the Cre and the homozygous floxed AR genes in both X chromosomes are illustrated in Figs. 1–3. We first constructed the targeting vector followed by generation of the founder mice carrying the floxed AR fragment. The founder mice were then mated with the female Cre mice to generate the F3 offspring that carries ARKO with Cre expression.

Construction of Targeting Vectors. Two genomic clones containing exon 2 of mouse AR were isolated from an ES129/SVJ bacteriophage λ genomic library (Stratagene) by using the mouse AR exon 2 sequence as the probe. The flanking region was sequenced and cloned into the PKI vector (26). Fig. 1 details the procedure for the construction of targeting vectors.

Generation of the Chimera Founder Mice. The ES cell line 129/SEVE was grown according to the conditions described (27). For electroporation, 40 μg of the targeting vector was linearized by NotI and suspended together with 109 ES cells in 1 ml of Dulbecco’s modified Eagle’s medium. They were electropolarized at 300 F, 0.4 ms in the Gene Pulsar II system (Bio-Rad). The neo+ colonies were selected in the presence of 300 μg/ml G418. Homologous recombinations were identified by genomic Southern blot hybridization. The clones with homologous recombination were amplified and reelectropolarized to introduce pCMV-Cre vector into the cells. The transient expression of the Cre in the cells resulted in three types of recombination, which was checked by Southern blot hybridization (27). The ES cells with

Abbreviations: AR, androgen receptor; KO, knockout; ES cell, embryonic stem cell; wt, wild type; T, testosterone; Tfm, testicular feminization; Cre, cre recombinase.

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an artificial Kpn site at the 5’ end of intron 2 sequences into 5’ cloning site (R1), followed by a fragment containing intron 1, exon 2, and a small fragment of intron 2 sequences into 5’ XbaI site (X). A lox sequence plus an artificial Kpn site were finally inserted into the XhoI site shortly 5’ to the beginning of exon 2. The constructed plasmid was linearized by NotI before being electroporated into ES cells. The vector was modified from the pBluescript vector and a thymidine kinase selective marker (MCT-TK) was inserted at the 5’ end. Two multiple cloning sites (MCS), two lox sites (loxP), a positive Neo selective marker (PKG-Neor), and a negative thymidine kinase-selective marker (MCT-TK) were inserted at the 5’ I site at the 5’ end primer is 5’-GTTGATACCTTAACCTCTGC-3’, and the 3’ end primer is 2-9, which is located in intron 2 and its sequence is 5’-CCTACATGTACTTGAGAGG-3’. The primer design and PCR conditions of Cre and IL2 follow the Jackson Laboratory’s suggestions.

**Histologic Analysis of Bones.** Preparation and analysis of bones were performed, as described (28). In brief, bones were fixed in 10% buffered formalin, decalcified in 10% EDTA, and embedded in paraffin. Sections were stained with hematoxylin and eosin for assessment of bone volume and for tartrate-resistant acid phosphatase activity for quantitation of osteoclasts. Bone volume (amount of bone matrix expressed as a percentage of the cancellous space) and osteoclast numbers (expressed per square millimeter of the cancellous bone volume) in the cancellous space of the long bones were measured. Bone samples were examined from a minimum of four wt and KO mice. Double labeling of bone-forming surfaces was performed by s.c. injection of calcine (20 mg/kg) given 7 days and 1 day before killing. Bone formation rate was measured, as described (29).

**Generation of Osteoclasts.** Splenocytes from ARKO mice and their wt littermates were used to generate osteoclasts in the absence of osteoblast/stromal cells. These cells were incubated with ammonium chloride solution for 10 min on ice to lyse red blood cells, and then cultured (1.75 × 10^6 cells per well in 96-well plates) in α-MEM (GIBCO/BRL) supplemented with 10% FCS (HyClone) in the presence of RANKL (100 ng/ml) and M-CSF (10 ng/ml). Cultures were maintained for 5–6 days at 37°C in an atmosphere of 5% CO₂/air and the media were changed every 2 days by replacing half of the spent media with fresh media supplemented with RANKL/M-CSF. Cells were then fixed and stained for tartrate-resistant acid phosphatase (TRAP) to identify osteoclasts, as described (30).

**Results and Discussion**

**Construction of Targeting Vectors and Electroporation of the ARKO Plasmid in ES Cells.** To disrupt the AR gene, exon 2 is targeted for loxP-Cre-mediated excision. Exon 2 encodes the second zinc finger of the DNA-binding domain, and deletion of the DNA-binding domain has been reported to result in the complete androgen-insensitive syndrome (31). As shown in Fig. 1, the PKI vector was modified from the pBluescript vector and a thymidine kinase-selective marker (MCT-TK) was inserted at the 5’ end of the multiple cloning site. The other neomycin-resistant marker (PKG-NeoⅠ′) flanked by two loxP sequences was inserted at the middle of the multiple cloning site separating it into two parts. The XhoI-cloning site was filled in by using Klenow fragment. Two fragments, one was a 3-kb fragment with the intron 2 sequence and the other was a 5-kb fragment including the 3’ end of intron 1, exon 2, and the 5’ end of intron 2 sequence, were generated by using the extended high fidelity PCR system. The loxP-Cre system utilizes the expression of the P1 phage Cre recombinase to catalyze the excision of DNA located between flanking lox sites. This strategy differs from the standard targeted disruption procedure because the ES cells were generated in which the targeted segment is not disrupted but flanked by lox sites (flxed). The arrangement of loxp sites located in intron 1 and intron 2 can preserve the AR function before Cre introduction. After selection and screening for homologous recombinants, a pCMV-Cre expression plasmid was then transiently transfected into the ES cell clones to induce recombination between any two loxp sites.

**PCR product with 580 bp. For examining the floxed AR X chromosome, primer “select” and primer “2-3” are used. 2-3 primer is the 3’ end primer which is located in the exon 2 with the sequence 5’-TTTACGCCGCTTTTGGAG-3’. This pair of primers will amplify a type 1 recombination were then injected into the inner cell mass of blastocysts, which were then implanted into the uterus of foster mothers for further development.

**Mating of the Chimera Founder Mice with the Homozygous Cre Mice.** The mating strategy is illustrated (see Fig. 3). The strain of the mosaic founder was C57BL/6J–129/SEV (B6/129). The mating between the founder and the female B6 mice create agouti female offspring carrying the heterozygous floxed AR (F1). The F1 offspring mate with the B6 male mice to create male mice carrying the floxed AR in the X chromosome (F2), or mate with the homozygous ACTB/Cre male that are carrying the Cre under the control of β-actin promoter to generate female mice carrying both the heterozygous floxed AR and Cre (F2). Mating these two genotypes of the F2 mice finally generate female mice carrying the homozygous ARKO and Cre. β-Actin is a housekeeping gene and is universally expressed in every tissue. Therefore, the β-actin promoter-driven Cre will express and delete the floxed AR fragment in all of the cells.

**Primer Design and Genotyping of ARKO Mice.** Based on the sequence information we obtained for the AR genomic DNA, two pairs of primers have been designed to distinguish the wild-type AR (wtAR), ARKO, and floxed AR X chromosome on mice. For examining the ARKO on the X chromosome, the 5’ primer named “select” is located in the intron 1 and its sequence is 5’-GTTGATACCTTAACCTCTGC-3’, and the 3’ end primer is 2-9, which is located in intron 2 and its sequence is 5’-CCTACATGTACTTGAGAGG-3’. If the mice contain wtAR, this pair of primers will amplify a PCR product with 238 bp. If the mouse contains wtAR, this pair of primers will amplify a PCR product with 580 bp. For examining the floxed AR X chromosome, primer “select” and primer “2-3” are used. 2-3 primer is the 3’ end primer which is located in the exon 2 with the sequence 5’-TTTACGCCGCTTTTGGAG-3’.
Screening of ARKO ES Cells and Genotype Screening of Floxed AR Chimera Male Mice. For the screening of ES cell clones with ARKO, we designed two pairs of primers to distinguish between the wt and floxed AR locus (Fig. 2A). Southern blotting was applied to verify the floxed AR construct in ES cells (Fig. 2B). The transient transfection of pCMV-Cre would result in type 1, type 2, and type 3 recombinations. As determined by Southern blot screening of ES cell clones containing wtAR, we were able to obtain 6 and 12 floxed AR clones, A9 and K7, respectively.

Generation of ARKO Mice. To generate mice with the disruption of AR, the floxed AR male mice were crossed to female mice carrying Cre under the control of the β-actin (ACTB) promoter. ACTB-Cre expression was activated in all tissues to generate mice lacking functional AR. As shown in Fig. 3, after two matings, we were able to identify ARKO male mice by using select and exon 2–9 primers to PCR amplify the 238-bp DNA. In contrast, wt and floxed AR male mice were distinguished by using the wt and floxed AR locus (Fig. 2A).

Genotype Screening of ARKO Mice. Alternatively, we were able to obtain ARKO male mice by mating floxed AR female and wild-type male mice. This strategy will generate pups of four possible genotypes (ar/Y-ACTB/Cre-male, ar/AR-ACTB/Cre-female, AR/Y-ACTB/Cre-male, AR/AR ACTB/Cre-female) with ratio of 1:1:1:1. Three primers, select, exon 2–3, and exon 2–9 (for the relative position of each primer in the AR gene see Fig. 4A), were synthesized to amplify mice genomic DNA to distinguish the floxed AR, ARKO, and wt mice. As shown in Fig. 4, we were able to identify ARKO male mice by using select and exon 2–9 primers to PCR amplify the 238-bp DNA. In contrast, wt mice can produce 580-bp PCR-amplified DNA fragments by using select and 2–9 primers (Fig. 4B).

Phenotype Characterization of ARKO Male Mice. Six 5- and 8-week-old ARKO mice were killed for the comparison of their phenotype with wt male and female mice. We first noticed that ARKO male mice have female-like appearance and body weight. In the 5-week-old ARKO male mice, the genito-anal distance is 0.55 cm, similar to female mice, yet is shorter than the wt male mice of 1.05 cm. In the 8-week-old ARKO male mice, the genito-anal distance is 0.59 cm, which is shorter than that of wt male sibling of 1.12 cm and similar to their female siblings (Fig. 5A). The ARKO male mice have testes, yet the size is much smaller, only 20% as compared with the same age of wt male mice (Figs. 5B–D).

Male Reproductive Organs. The external genitalia in male ARKO mice showed ambiguous or feminized appearance. The penis seems microphallus and the urethra shows hypospadias. The scrotum is poorly developed and looks like the labia major in the female. All of vas deferens, epididymis, seminal vesicle, and prostate are agenesis (Fig. 5B and C). Testes are small in size and cryptorchid in the low abdominal area close to the internal inguinal ring, and are similar in Tfm mice or humans with complete androgen-insensitive syndrome (Fig. 5B–D). No vaginal opening is found, and also no fallopian tubes and uterus can be observed.

Analysis of Testes. To dissect further the testes morphology in the ARKO mice, immunostaining was performed. As shown in Fig.
6A and B, we could observe obvious AR positive signals in testes from 8-week-old wt mice (Fig. 6A). In contrast, no AR signal could be detected from 8-week-old ARKO mice (Fig. 6B).

We then compared the histology of testes from ARKO mice and wt mice. As shown in Fig. 6 C and D, in the testes from wt mice, spermatogenesis sequentially developed round spermatids and elongated spermatids. In addition, mature spermatocytes can be clearly observed (Fig. 6C). However, testes from ARKO mice are less cellular and thin in the seminiferous tubules. Some tubules contain Sertoli cells only and the others contain a few germ cells (Fig. 6D). In the tubules with germ cells, the spermatogenesis is hypoplastic. Occasionally, some spermatocytes at the pachytene stage are found (Fig. 6E). No round spermatids, elongated spermatids, or mature spermatocytes occur in the whole testis, suggesting spermatogenesis may be arrested at pachytene spermatocyte. The Sertoli cells show fibrillary degeneration (Fig. 6F). Many pyknotic cells contain apoptotic-like bodies in tubules (Fig. 6G). In the interstitial space, Leydig cells are hypertrophic (Fig. 6H).

Early studies showed that testes in Tfm mice are smaller (20% of normal), cryptorchid, and composed of immature tubular elements surrounded by several layers of peritubular cells and enlarged Leydig cells. The numbers of Leydig cells are normal or slightly reduced. The reduced seminiferous tubules contain only Sertoli cells, spermatogonia, and primary spermatocytes. Spermatogenesis was arrested at the spermatocyte stages or earlier. In older Tfm mice, the Leydig cells seemed to be hypertrophied. In the pachytene spermatocyte stage, homologues are fully paired and exchange of genetic material occurs through crossing over between chromatids of homologous chromosomes (32). Together, our results indicate AR may play some important roles during these processes.

**Testosterone Concentration in Serum.** Previous in vivo animal models clearly established that initiation, maintenance, and reinitiation of spermatogenesis are androgen-dependent processes (33). We also compared the serum T concentration in 8-week-old mice in wt and ARKO mice. The results from three individual mice indicate that the T concentration in ARKO mice is lower than wt mice.

**Bones.** Histologic analysis of bone sections from 8-week-old ARKO mice shows that cancellous bone volumes are lower in the ARKO mice than in both female and male wt littermates. The decrease in bone volume is seen readily in the metaphyses of the femora (Fig. 7A and C) and tibiae (data not shown). Osteoclast numbers in the femoral metaphyses are higher in ARKO than in wt mice (Fig. 7C). However, the number of osteoclasts formed in vitro from spleen cells from ARKO mice is similar to that in wt mice (data not shown). Mean values for mineral apposition and bone formation rates in the femoral

Fig. 5. Phenotype of 8-week-old male ARKO mice. Six 8-week-old male ARKO mice were killed. The results were always compared among siblings. (A) The external genitalia of male ARKO, wt male, and wt female. (B and C) The internal genitalia of male ARKO and wt male mice. Arrows in B identify the testis. (D) The testes of wt male and ARKO mice.

Fig. 4. Genotyping of ARKO mice. We have applied the primers "select" and "2–9" to identify wt and ARKO male mice in our study. (A) Schematic presentation of the DNA construct and primer location in exon 2 area of wt, KO, and floxed AR genes. (B) List of the sizes of PCR product amplified by designed primer pairs. (C) The identification of wt and ARKO mice, by using select and 2–9. We amplified a DNA fragment with 380 bp, which represents wtAR, and with 238 bp, which represents ARKO exon 2. The expression of Cre and internal control IL-2 were confirmed by PCR (Bottom).
metaphyses are higher in the ARKO mice (Fig. 7 B and D) than in the wt mice. The difference in the values for bone formation rates did not reach statistical significance, presumably because of the small number of animals studied (three per group). These data indicate that in the absence of the AR, mice develop osteopenia. Because osteoclast numbers and bone formation are both increased in these mice, the osteopenic phenotype most likely is because of bone resorption being increased beyond that of the increase in bone formation. This increase in osteoclast function could be the result of an imbalance in the production of RANKL (the major osteoclastogenic cytokine) and OPG (the major inhibitor of RANKL, which inhibits RANKL by acting as its decoy receptor) by osteoblast/stromal cells (34), because estrogen regulates OPG production by these cells (35). We have demonstrated previously that testosterone and estrogen promote apoptosis of osteoclasts and their precursors in vitro and in vivo (34, 36). Thus, it is possible that osteoclasts are not only increased in number in ARKO mice, but that they also live longer than normal because of reduced testosterone concentrations, and therefore resorb more bone than wt cells. These possibilities are currently under investigation. The increased bone mineral apposition and bone formation rates coupled with increased osteoclast numbers and low bone volumes observed in the mice is a phenotype that is seen in sex steroid deficiency (37) and is consistent with the low T concentrations in the mice. Taken together, the osteopenic phenotype of ARKO mice strongly supports an important role of AR signaling in bone metabolism.

Body Weight and Adipose Tissue. The body weight of ARKO male mice is similar to wt female mice with an average of 18.8 g at 5 weeks and 20.1 g at 8 weeks. In contrast, the average body weight of wt male mice is 24.3 g at 5 weeks and 26.7 g at 8 weeks. Histologic analysis of adipose tissue indicated that the size and number of adipocytes start to be different between wt male and ARKO male mice at 5 weeks. These differences become more obvious in 8-week-old mice (Fig. 8). In contrast, infrarenal adipocytes were well-developed in both 5- and 8-week-old mice, and show no obvious difference between wt and ARKO mice (Fig. 8B). These results suggest that AR may play roles in adipogenesis. Early reports also suggest that androgens may play some roles in the site specificity of adipose metabolism (38). Women with abdominal fat distribution may have increased percentage of free T in the peripheral blood. In contrast, obesity in men may be characterized by reduced T. However, the detailed mechanisms of how T-AR influence adipose tissue and obesity remain unclear. Because peroxisome-proliferator-activated receptors play important roles in adipogenesis (39, 40) and the AR coregulator, ARA70, can modulate both AR and peroxisome-proliferator-activated receptors’ transcriptional activities (41, 42), it will be interesting to see whether AR can crosstalk to peroxisome-proliferator-activated receptors’ pathway by sharing a common coregulator, such as ARA70, in adipose tissue.

ARKO Female Mice. Because of the infertility of Tfm male mice, it is difficult to generate ARKO female mice to study the roles of AR in female tissues. With floxed AR male mice, we now are able to mate them with ACTB-Cre ar/AR-female to generate ARKO in female mice. By using this strategy, we were able to obtain a few ARKO female mice, and we are in the process of obtaining more ARKO female mice and analyzing their phenotypes with focus on the ovary and breast.

Early analysis showed that although the female ARKO were fertile, the average number of the pups per litter from heterozygous and homologous ARKO females are reduced as compared to wt female mice. Even though we obtained homologous ARKO female mice, we are still in the process of obtaining enough mice for further analysis. Previous studies have revealed that androgens played a positive role in the early stage of folliculogenesis.
(43, 44). However, it is unclear whether the androgen effect is mediated by AR. For example, Vendola et al. suggested that this effect could be mediated by way of growth factor signal pathways instead of the AR-mediated pathway (44). With ARKO female mice, we are in the process of dissecting potential roles of AR in the folliculogenesis.

The morphology of breast in our ARKO mice at this early stage show no major differences as compared with wt mice. Early studies suggested that androgen-AR may play some roles in the progression of breast cancer (45). The detailed mechanisms, however, remain unclear. Our ARKO female mice may provide a nice in vivo model to study how carcinogens induce breast tumors in the presence versus absence of AR in breast tissue.


**Tissue-Specific Knockout of AR.** One advantage of creating the floxed AR mice is to provide a base to generate tissue-specific ARKO in selective tissues such as breast (mating with MMTV-Cre mice), prostate (mating with PSA-Cre or probasin-Cre), and liver (mating with α-fetal protein-Cre or albumin-Cre). We are in the process of generating these tissue-specific ARKO mice to study the roles of AR in these tissues.

In summary, the generation of ARKO male and female mice provides us with a valuable in vivo model to study androgen functions in selective tissues.

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