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## Normal and Abnormal Epithelial Differentiation in the Female Reproductive Tract

**Takeshi Kurita**

Division of Reproductive Biology Research, Department of Obstetrics and Gynecology, Northwestern University Feinberg School of Medicine, Chicago, Illinois 60611

### Abstract

In mammals, the female reproductive tract (FRT) develops from a pair of paramesonephric or Müllerian ducts (MDs), which arise from coelomic epithelial cells of mesodermal origin. During development, the MDs undergo a dynamic morphogenetic transformation from simple tubes consisting of homogeneous epithelium and surrounding mesenchyme into several distinct organs namely the oviduct, uterus, cervix and vagina. Following the formation of anatomically distinctive organs, the uniform MD epithelium (MDE) differentiates into diverse epithelial cell types with unique morphology and functions in each organ. Classic tissue recombination studies, in which the epithelium and mesenchyme isolated from the newborn mouse FRT were recombined, have established that the organ specific epithelial cell fate of MDE is dictated by the underlying mesenchyme. The tissue recombination studies have also demonstrated that there is a narrow developmental window for the epithelial cell fate determination in MD-derived organs. Accordingly, the developmental plasticity of epithelial cells is mostly lost in mature FRT. If the signaling that controls epithelial differentiation is disrupted at the critical developmental stage, the cell fate of MD-derived epithelial tissues will be permanently altered and can result in epithelial lesions in adult life. A disruption of signaling that maintains epithelial cell fate can also cause epithelial lesions in the FRT. In this review, the pathogenesis of cervical/vaginal adenoses and uterine squamous metaplasia is discussed as examples of such incidences.

### Keywords

diethylstilbestrol (DES); adenosis; clear cell adenocarcinoma; squamous metaplasia; p63; Wnt

### Development of Müllerian duct derived organs

In mammals, the majority of the FRT develops from a pair of paramesonephric or Müllerian ducts (MD) of mesodermal origin. The MDs arise as cranio-caudal invaginations of thickened coelomic epithelium (Müllerian plaque) at the upper end of the urogenital ridge on the lateral aspect of the corresponding mesonephric or Wolffian duct (WD) (Fig. 1A). The site of the invagination remains open throughout the development of the MDs and becomes the abdominal ostium of the oviduct (Fig. 1A insert). The initial segment of the MD grows

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To whom correspondence is addressed: Takeshi Kurita, Ph.D. Department of Obstetrics and Gynecology, Division of Reproductive Biology Research, Northwestern University Feinberg School of Medicine, 7th Floor, Suite 7-127 303, East Superior Street, Chicago, Illinois 60611, Phone: 312-503-0525, Fax: 312-503-0095, t-kurita@northwestern.edu.

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caudally through urogenital ridge mesenchyme using the WD as a guide (Fig. 1B) (Grünwald, 1941). The cellular origin of the MD during caudal growth, particularly the presence/absence of cellular contribution of the WD to the MD, has been debated for decades. Some have argued that the WD simply acts as a guide for caudal MD growth (O’Rahilly, 1973), whereas others have proposed that the MD partially or completely splits off from the WD after they make intimate contact with each other (Grünwald, 1941). Recent cell fate tracing experiments in the chick and the mouse have settled the debate by demonstrating that both the epithelium and mesenchyme of the Müllerian duct arise from coelomic epithelium (Guioli et al., 2007; Orvis and Behringer, 2007), as previously suggested by the expression pattern of *Amhr2* (Zhan et al., 2006). In the cell fate tracing studies, the long debated cellular contribution of the WD to the MD (Del Vecchio, 1982; Frutiger, 1969) was not detected.

As the right and left MDs grow caudally, they cross the WDs ventrally to join (Fig. 1B’) and fuse with each other in the midline. Subsequently, the medial walls of the MDs degenerate forming a single canal (Fig. 2A). In females, the caudal tip of the MDs reaches the urogenital sinus (UGS), which is derived from the endoderm. The fusion of the MDs, the WDs and the UGS forms the sinovaginal bulbs, which are projections of solid epithelial cords on the dorsal wall of the UGS (Koff, 1933; Bloomfield and Frazer, 1927) (Fig. 2A). At the same time, the majority of WDs in female embryos regress due to the absence of testicular androgen, leaving fragments of WDE at the junction with the UGS. In human and mouse female embryos, a flat epithelial cord, which is called the vaginal plate in humans, is formed following the union of fused MDs with the UGS (Koff, 1933). Most textbooks indicate that the lower portion of vagina forms via the simultaneous growth and canalization of the vaginal plate (Russell, 1989; Moore and Persaud, 2002; Sadler, 2004; Gilbert, 2003; Carlson, 1999; Forsberg, 1978).

### **Classic problem; Developmental origin of the lower female reproductive tract**

As described above, the MDs, WDs, and UGS are present at the site of organogenesis of the lower FRT (Fig. 2A), and the degree of contribution of these structures to the formation of the vaginal bulb and plate, as well as the adult vagina have been debated for decades. There have been four major theories for the developmental origin of vaginal epithelium. The most widely accepted has been the “UGS + MD origin” theory, in which the upper two-thirds of the vagina (Müllerian vagina) develops from the caudal portion of the MDs, and the lower portion (sinus vagina) develops from the UGS (Moore and Persaud, 2002; Sadler, 2004; Koff, 1933; Forsberg, 1963; 1973; Kobayashi and Behringer, 2003; Yin and Ma, 2005; Shapiro et al., 2000; Gilbert, 2003; Del Vecchio, 1982; Sajjad, 2010). In this theory, the vaginal bulb/plate consists solely of epithelium of UGS origin, and is thus referred to as the “sinovaginal bulb” (SVB). Accordingly, the lower vagina develops through the simultaneous growth and canalization of the UGS epithelium (UGE) (Koff, 1933; Forsberg, 1963). The shallow vagina that is present in mice and humans with complete androgen insensitivity syndrome (CAIS) or testicular feminization mutation (*Tfm*) provides strong support for this theory (Morris, 1953; Cunha, 1975). CAIS arises from a spontaneous mutation of the androgen receptor gene, which causes insensitivity to testosterone; as a result, the male reproductive tract, excluding the testes, does not develop normally in affected XY individuals (Quigley et al., 1995). At the same time, the MDs regress due to the production of the anti Müllerian hormone (AMH)/Müllerian inhibiting substance (MIS) by the fetal testis (Teixeira et al., 2001; Josso et al., 2001). Therefore, the short vagina present in *Tfm* males is believed to arise from the sinus vagina of the UGS origin (Cunha, 1975).

The alternative “MD origin” and “MD + WD origin” theories describe how the vaginal bulb/plate is derived from the MDs (Bloomfield and Frazer, 1927; Cai, 2009) or the WDs (Forsberg, 1963; Witschi, 1970; Drews, 2007), and thus, the vagina develops from either the MDs alone or the MDs plus the WDs (Hart, 1901; Bloomfield and Frazer, 1927; Mauch et al., 1985; Sánchez-Ferrer et al., 2006). In the MD + WD origin theory, the WDs contribute to the epithelium of the vagina (Hart, 1896; 1901; Forsberg, 1963; Ación, 1992; Sánchez-Ferrer et al., 2006). In contrast, the MD origin theory proposes that the WDs play a role in the downward growth of the Müllerian-derived vagina and/or in the formation of the hymen, but they do not contribute to the vaginal epithelium (Bloomfield and Frazer, 1927; Drews, 2007). In these theories, the presence of short vaginae in CAIS-affected XY individuals is considered a derivative of residual MDs and/or WDs (Forsberg, 1978; Drews et al., 2002).

Finally, the UGS origin theory suggests that the entire squamous epithelium of the cervix and vagina are originated solely from the UGE of endodermic origin (Arey, 1954; Bulmer, 1957; 1959; Ferris, 2004; Fliegner, 1994; Ulfelder and Robboy, 1976). In this view, the squamous epithelium derived from the UGS grows upward and replaces the original columnar epithelium of MD origin. However, mouse studies demonstrated that the squamous vaginal epithelium might develop from MD epithelium (MDE) (Forsberg, 1965; Cunha, 1976; Kurita et al., 2004), which questions the observations supporting the UGS origin theory. All of the theories are based upon anatomical/histological observations, with the boundaries of structures drawn subjectively. Thus, none of these theories were definitively proven or disproven until recently.

The origin of vaginal epithelium has been finally determined by a recent cell lineage tracing experiment (Kurita, 2010). In the experiment, mouse epithelial cells of embryonic WD, MD, or UGS origin were permanently labeled to express enhanced green fluorescent protein (EGFP), and the cellular origin of vaginal epithelium was determined by following the developmental cell fate of EGFP-positive cells from the embryo to the adult. The cellular origin and organogenesis process of vagina revealed by the study have refuted the four theories described above. The cell lineage tracing study agreed with the MD+UGS theory in regards to the origin of the vaginal bulb in that the vaginal bulb solely consisted of UGE. Thus the embryonic and newborn vaginae consisted of both the “Müllerian vagina” and “sinus vagina.” However, the Müllerian and sinus vaginae were never “unified” as described in the majority of textbooks. In mice, the epithelium in the sinus vagina remained solid throughout development, and it never became a part of the true vagina. Moreover, the solid epithelium of the sinus vagina never elongated. Instead, the sinus vagina connected the Müllerian vagina and urethra and migrated towards the posterior end of the body leading the caudal growth of the Müllerian vagina (Fig. 2B and C). The solid epithelial cord in the sinus vagina canalized only during the formation of the vaginal orifice at puberty (~ four-weeks-old), and thus the UGE in the sinus vagina became a part of the vulvar epithelium. Although a critical role of the WD was proposed in the MD origin and MD+WD origin theories, the WD did not appear to play a role in the development of the vagina. WDE cells were not detected in the vaginal epithelium throughout development (Kurita, 2010).

## Differentiation of vaginal and uterine epithelial cells

As explained in the previous section, the entire FRT, from the oviduct to the vagina, develops from the MD. As each MD-derived organ becomes anatomically distinctive, the uniform MDE differentiates into epithelial cell types with unique morphology and functions in each organ. While the differentiation of the MDE occurs in the fetal stage of the human, mouse MDE is mostly undifferentiated at the time of birth (Kurita et al., 2005; Kurita and Nakamura, 2008). For example, ciliation of the epithelial cells in the mouse oviduct is first observed on postnatal day 5 (P5) (Komatsu and Fujita, 1978). The mouse uterine epithelium

displays the first signs of morphogenesis around P4 as a complex luminal shape, and the rudimentary glands appear on ~P7 (Brody and Cunha, 1989; Kurita et al., 2001). Since it can be studied postnatally, the mechanism of MDE differentiation has been studied mainly by using the mouse as the model organism. Accordingly, this and the following sections mainly discuss the mouse studies.

Although epithelia in the uterus, cervix and vagina are derived from MDE, the simple columnar epithelium of the uterus and the stratified squamous epithelium of the cervix/vagina are profoundly different in their morphology and gene expression. For example, cytokeratin 14 (K14) is expressed in vaginal and cervical but not in uterine epithelium. Progesterone receptor (PR) is also differentially expressed in the epithelial cells of the mouse uterus versus the cervix/vagina. In mouse cervical/vaginal epithelial cells, PR is upregulated by estrogen through the estrogen receptor  $\alpha$  (ER $\alpha$ ) as in most progesterone target cells/tissues (Kurita et al., 2000). In contrast, PR is expressed independently of estrogen and ER $\alpha$  in the mouse uterine epithelium (Kurita et al., 2000). In fact, PR is down-regulated, not up-regulated, by estrogen in the mouse uterine epithelium (Kurita et al., 2000). Since this unusual pattern of PR regulation is unique for the uterine epithelial cells of rodents, estrogen-independent PR expression is an excellent marker for mouse uterine epithelial differentiation (Kurita et al., 2001).

The critical role of epithelial-mesenchymal tissue interaction in the differentiation of MDE has been established by a series of tissue recombination studies, in which the mesenchyme and epithelium isolated from the uterus and vagina of newborn mice were recombined and grafted under the subrenal capsule of syngeneic or immunodeficient mouse hosts (Cunha, 1976; Boutin et al., 1991b; Kurita et al., 2001; Kurita et al., 2004). In 1976, Cunha demonstrated for the first time that the differentiation fate of uterine and vaginal epithelial cells was exchangeable at birth, and that their organ-specific phenotypes are dictated by regional inductive cues from the surrounding mesenchyme (arrows in Fig. 2A' and B') (Cunha, 1976). In response to the induction by the uterine mesenchyme, vaginal epithelium differentiates into uterine epithelium containing both luminal epithelial cells and glands (Cunha, 1976; Boutin et al., 1991a). The induced uterine epithelium expresses a spectrum of differentiation markers characteristic of normal uterine epithelium such as uterine-type syndecans (Boutin et al., 1991b) and PR (Kurita et al., 2001). In the reciprocal tissue recombinants, vaginal mesenchyme induced the uterine epithelium to differentiate into a stratified squamous vaginal epithelium, which showed a mucification and cornification in response to the estrogen and progesterone (Cunha, 1976; Boutin et al., 1991a; 1992). The induced vaginal epithelium expressed vaginal differentiation markers such as the vaginal isoform of syndecan (Boutin et al., 1991b), involucrin and keratins 1, 10, 14 and 19 (Kurita et al., 2001). In the induced vaginal epithelium, PR was also expressed in the vaginal pattern (Kurita et al., 2001). Though the differentiation markers are already expressed, most cells in the vaginal and uterine epithelia from P5 mice can still trans-differentiate into either uterine or vaginal epithelium when induced by heterotypic mesenchyme (Cunha, 1976). However, the developmental plasticity of the epithelial cells is then gradually lost thereafter, and the uterine and vaginal epithelial phenotypes become irreversibly fixed. Accordingly, uterine and vaginal epithelium from seven-day-old or older mice are induced only partially to become the alternative cell type by heterotypic mesenchyme, and most adult uterine and vaginal epithelial cells cannot be re-programmed by the newborn uterine or vaginal mesenchyme to express alternative epithelial phenotypes (Kurita et al., 2001; Kurita et al., 2004). Hence, the organ specific epithelial phenotype of mouse uterine and vaginal epithelial cells is "induced" and then "stabilized" during the first week of postnatal development. As a result, a boundary known as the squamocolumnar junction (SCJ) is formed between the columnar and squamous epithelia of the uterus and cervix. In the mouse, the SCJ is located in the caudal portion of each uterine horn where the uterine and

cervical epithelia meet (black arrow Fig. 2C and C'). Since cervical and vaginal epithelial cells of the mouse are indistinguishable, there is no epithelial boundary between the cervix and vagina.

## p63 transcription factor in FRT development

The transcription factor p63 is the product of the mouse *Trp63* or human *TP63* gene, which are transcribed into isoforms containing or lacking the N-terminal transactivation domain, TA and  $\Delta N$  forms, respectively. In addition, alternative splicing generates three different C-terminal sequences corresponding to the  $\alpha$ ,  $\beta$  and  $\gamma$  forms (Yang et al., 1998). p63 is highly expressed in germ cells (Petre-Lazar et al., 2007; Nakamuta and Kobayashi, 2004b; Kurita et al., 2005; Saito et al., 2006; Nakamuta and Kobayashi, 2004a; Petre-Lazar et al., 2006) and the basal cells of many epithelial tissues (Mills et al., 1999; Yang et al., 1999; Signoretti et al., 2000; Kurita and Cunha, 2001; Daniely et al., 2004). In the adult FRT, p63 is expressed in cervical and vaginal but not uterine epithelium (Kurita and Cunha, 2001; Kurita et al., 2004; Kurita et al., 2005), and the  $\Delta N$ p63 is the dominant form in the cervix and vagina (Kurita et al., 2005). The MDE and its precursor coelomic cells are negative for p63, but p63 expression is induced by the cervical/vaginal mesenchyme in the early fetal stage of the human and the perinatal stage of the mouse (Fig. 2B') (Kurita et al., 2005; Kurita and Nakamura, 2008). In developing mouse and human FRT, p63 is the first marker that is differentially expressed between uterine and vaginal/cervical epithelial cells, suggesting that the key role of p63 lies in the epithelial cell fate determination in the cervix and vagina. Indeed, the cervical/vaginal epithelium of p63 null mice differentiated into columnar uterine-like epithelium (Kurita and Cunha, 2001). The p63 null vaginal epithelium expressed PR in the absence of estrogen indicating uterine epithelial identity, demonstrating that the expression of p63 determines the developmental cell fate of MDE to become uterine epithelium (p63-negative) or cervical/vaginal epithelium (p63-positive) (Kurita et al., 2004).

When p63 is induced in the MDE by cervical and vaginal mesenchyme, the p63-positive MDE cells subsequently change their morphology and eventually transform into cervical/vaginal epithelium (Kurita et al., 2004; Kurita and Cunha, 2001). Consequently, a layer of p63 positive basal cells forms in the cervix and vagina. In the mouse, a continuous basal epithelial cell layer of cervix/vagina forms by P5 and the SCJ becomes distinctive by P14 (Kurita et al., 2005). As described above, uterine and vaginal epithelia gradually lose their ability to change in the p63-expression patterns in response to induction by heterotypic mesenchyme. Accordingly, adult vaginal epithelium maintains its original squamous phenotype and p63 expression, even when it is combined with inductive uterine mesenchyme from newborn mice (Kurita et al., 2004). This result indicates that the cervical/vaginal mesenchymal factor(s) is required only for activation and not for the maintenance of p63 expression. The developmental plasticity of uterine epithelial cells is gradually lost during postnatal development, and most uterine epithelial cells do not respond to induction by the vaginal mesenchyme (Kurita et al., 2004). However, there is a small number of epithelial cells that maintain developmental plasticity in the adult uterus as assessed by the induction of p63 expression. When the uterine epithelial cells from two-month-old virgin mice were combined with vaginal mesenchyme, p63 expression was detected in a small subpopulation of epithelial cells (< 5%) (Fig. 4C). These rare developmentally plastic cells may be stem cells and the targets of uterine squamous metaplasia (see below).

## DES-induced cervical/vaginal adenosis and clear cell adenocarcinoma

### (a) Adenosis

Cervical and vaginal adenoses are congenital anomalies defined as the development of glandular (columnar) epithelium in cervical/vaginal epithelium, which is normally stratified

squamous. Cervical/vaginal adenosis is known to be associated with *in utero* exposure to diethylstilbestrol (DES), a synthetic non-steroidal estrogen. DES was prescribed during the 1938–1971 for pregnant women to help prevent miscarriages. It has been estimated that more than 2 million mothers, daughters and sons were exposed to DES in the United States alone (Giusti et al., 1995). Female fetuses exposed to DES *in utero* (DES daughters) are at increased risk of developing clear cell adenocarcinoma (CCAC) of the cervix and vagina (Herbst and Scully, 1970; Herbst et al., 1971; Melnick et al., 1987; Verloop et al., 2010), and adenosis is believed to be the precursor of CCAC. Particularly, atypical adenosis of the tuboendometrial type has been considered as the immediate precursors of CCACs (Robboy et al., 1984). The incidences of adenosis and CCACs are directly related to the starting time of DES-treatment during pregnancy. A study demonstrated that more than a third of DES-daughters developed adenosis if DES was given before gestational week eight (Robboy et al., 1984). In the same study, adenosis was essentially absent if DES-treatment was started at week 22 or later (Robboy et al., 1984). Perinatal exposure of mice to DES also generates a spectrum of reproductive tract lesions similar to those observed in DES daughters, including cervical/vaginal adenosis (Forsberg, 1976; Plapinger and Bern, 1979; McLachlan et al., 1980). As in DES-daughters, the frequency of cervical/vaginal adenosis in the mouse is also directly related to the timing of DES exposure. When female neonatal mice were exposed to DES from P1 to 5, vaginal adenosis was detected in 75% of mice by P35, while it was detected only in 15% of mice with prenatal DES-exposure (gestation day 9 to 16) (Newbold and McLachlan, 1982). In the mouse, the first seven days of postnatal development are crucial for the MDE to establish p63 expression patterns, and thus the differentiation cell fate (Kurita et al., 2004). The developmental stage in which the human fetus is susceptible to DES-induced adenosis/CCACs also corresponds to the time period in which p63 expression is induced in the cervical/vaginal epithelium (Kurita et al., 2005). These observations suggest that DES induces cervical and vaginal adenosis by altering the expression pattern of p63, and thus the differentiation cell fate of MDE. Indeed, DES-exposure repressed expression of p63 in developing mouse cervix and vagina (Kurita et al., 2004). This inhibitory effect of DES on p63 expression is usually transient. However, when DES is present throughout the critical period for cell fate determination, the MDE cells in the cervix and vagina miss the developmental window to express p63 and consequently, the p63-negative MDE cells differentiate into uterine epithelium within cervix and vagina forming adenosis (Kurita et al., 2004).

The inhibitory effect of DES on p63 in developing cervical/vaginal epithelium was mediated via ER $\alpha$  (Kurita et al., 2004). A tissue recombination study with ER $\alpha$  null mice determined definitively that DES blocks the expression of p63 via ER $\alpha$  within the MDE (Kurita et al., 2004). In the study, DES blocked p63 expression in wild-type but not ER $\alpha$  null MDE. DES actions via ER $\alpha$  in the mesenchymal cells did not affect the expression of p63 in the epithelial cells. Our recent study indicates that DES via ER $\alpha$  blocks p63 expression in MDE cells autonomously (Fig. 3). In the experiment, uterine epithelial cells from P1 ER $\alpha$  null and wild-type mice were mixed and combined with vaginal mesenchyme, which was then grafted under the subrenal capsule of nude mice (Fig. 3). In intact hosts, uterine epithelium was induced to express p63 in both ER $\alpha$  positive and negative epithelial cells (Fig. 3A–C). In contrast, when hosts were supplemented with DES, p63 expression was detected only in the ER $\alpha$  negative cells (Fig. 3D – F). These experiments clearly demonstrate that DES does not block expression of p63 in the MDE through the down-regulation of mesenchymal factors (red arrows in Fig. 2A' and B'). Cell-autonomous inhibition of p63 expression by DES via epithelial ER $\alpha$  also refutes the involvement of autocrine factors produced by the MDE in the pathogenesis of DES-induced adenosis.

### (b). Clear Cell Adenocarcinoma

The relative risk of CCAC in DES Daughters has been estimated to be greater than 40 times compared to the general population (Hatch et al., 1998). Therefore, it has been considered that *in utero* exposure to DES is the cause of CCACs of the cervix and vagina. However, the incidence of CCAC has been estimated to be 1 to 1.5 per 1,000 DES daughters (Melnick et al., 1987; Herbst, 2000), whereas the prevalence of cervical/vaginal adenosis in DES daughters was as high as ~90% (Johnson et al., 1979; Sherman et al., 1974; Robboy et al., 1984). The discrepancy in the incidences of DES-induced cervical/vaginal adenosis versus CCACs indicates that oncogenesis of cervical/vaginal CCACs requires factors other than the developmental loss of p63 expression. Moreover, the prevalence of adenosis in the general population has been estimated to be 1 to 10 % (Herbst et al., 1975; Chattopadhyay et al., 2001), which is 5 – 90 times lower than that of DES daughters. Thus, the > 40-fold increase in CCAC incidence in DES daughters appears to be proportional to the increased rate of adenosis lesions. These numbers strongly suggest that the *in utero* DES exposure increased the risk of cervical/vaginal CCACs by increasing the prevalence of cervical/vaginal adenosis lesions; however the chance of the malignant transformation of adenosis to adenocarcinoma may be identical between DES-daughters and general population. Progression of adenosis to CCACs may be independent of DES-exposure.

The molecular mechanism underlying the transformation of the adenosis lesion to CCAC is not known. For example, a screening for common mutations to cancers, such as Ras, p53 and WT1, failed to detect mutations in all 16 cervical/vaginal CCAC cases (Boyd et al., 1996). While infection of high-risk types of the human papillomavirus (HPV) is the major cause of both cervical squamous cell carcinomas (SCC) and non-clear cell adenocarcinomas (Clifford et al., 2003; Castellsague et al., 2006), there is no association between this infection and CCACs (Pirog et al., 2000; Goto et al., 2005; Stewart et al., 2006; Liebrich et al., 2009). Even in a case of synchronous SCC and CCAC of the cervix (co-existing cervical SCC and CCAC in the same patient), HPV was found only in SCC but not in CCAC, suggesting different etiologies in these two types of cervical cancers (Goto et al., 2005). The low prevalence of this cancer makes research difficult. Although the etiology of cervical/vaginal CCAC remains unknown, it should be noted that in a study, loss of function mutations in PTEN (phosphatase and tensin homolog deleted on chromosome 10) were detected in 36% (4/11) of HPV-negative cervical adenocarcinomas (Minaguchi et al., 2004), suggesting a link between alterations in the phosphatidylinositol 3,4,5-trisphosphate (PIP<sub>3</sub>) signaling pathway and CCACs of the cervix.

### Cervical/vaginal adenosis and CCAC in post-DES era

Although DES is no longer prescribed to pregnant women, cervical/vaginal adenosis and CCACs are still relevant to current women's health. There have been cases of cervical/vaginal adenoses/CCACs reported in women who have no history of DES exposure (Kurman and Scully, 1974; Robboy et al., 1986; Accetta et al., 2001; Herbst et al., 1975; Chattopadhyay et al., 2001). The persisting incidence of cervical/vaginal adenosis in the general population even after the ban of DES usage suggests the existence of other causal factors in our environment. *In utero* exposure to environmental chemicals may be the cause of adenosis in the cervix and vagina. In this regard, there are a number of environmental chemicals that disrupt the endocrine systems of animals and humans by mimicking or antagonizing the biological activities of natural hormones, thus known as endocrine disruptors (Colborn, 1995; Diamanti-Kandarakis et al., 2009). Some environmental chemicals such as bisphenol A (BPA) have been reported to elicit estrogenic activities, which may explain the underlying cause of non DES-associated adenosis (Newbold et al., 2009).

## Uterine Squamous Metaplasia

In addition to cervical/vaginal adenosis, perinatal DES exposure of mice also induced uterine squamous metaplasia, development of squamous cells in normally columnar uterine epithelium (McLachlan et al., 1980). When pregnant mice were treated with 100  $\mu\text{g}$  DES/kg/day from gestational day 14 to day 17, a small number of p63 positive cells were observed in the uterine epithelium at P7 (Fig. 4A), which were considered to be the seed of squamous metaplasia (Kurita et al., 2004). The metaplastic squamous epithelial cells in the adult uterus expressed vaginal epithelial markers such as K14, and PR was also expressed in the vaginal pattern (Kurita et al., 2004). These observations indicate that uterine squamous metaplasia is the formation of cervical/vaginal epithelium in the uterus. The uterus of p63 null mice was resistant to DES-induced squamous metaplasia, confirming the essential role of p63 in cervical/vaginal epithelial differentiation (Kurita et al., 2004). While cervical and vaginal adenoses are congenital, squamous metaplasia can develop in the adult uterus. For example, uterine squamous metaplasia is a classic histological change caused by chronic estrogen exposure (McEuen, 1936; Gitlin, 1954; Selye et al., 1935) or vitamin A deficiency (VAD) (Wolbach and Howe, 1925; Moll et al., 1983; Darwiche et al., 1993) in mature animals. When CD-1 mice were fed a vitamin A deficient diet from weaning, expression of p63 was detected in the uterine epithelium at four months old (Fig. 4E). This observation is in agreement with the tissue recombination study that demonstrated the presence of developmental plastic cells in adult uterine epithelium (Fig. 3C).

## Canonical Wnt signaling pathway maintains uterine epithelial identity

As described above, epithelial differentiation in MDE was first “induced” and then “stabilized” during development. The Wnt (wingless-type MMTV integration site family member) pathway plays an important role in the “stabilization” of the epithelial cell fate in the uterus. In the *Wnt7a* null mutant mice, uterine epithelial cells are columnar with proper differentiation markers by puberty (Parr and McMahon, 1998; Carta and Sassoon, 2004; Kurita and Nakamura, 2008), thus *Wnt7a* is dispensable for the cell fate determination of MDE in the neonatal uterus and cervix/vagina (Fig. 5A). However, *Wnt7a* is essential for the “stabilization” of the uterine epithelial cell fate, and thus uterine epithelium of *Wnt7a* null mice gradually transforms into a squamous epithelium. The uterine squamous metaplasia becomes prominent in fully mature *Wnt7a* null mice (Miller and Sassoon, 1998), and the post-pubertal onset of uterine squamous metaplasia in *Wnt7a* null mice suggests the involvement of ovarian steroid hormones in its pathogenesis. Indeed, our study demonstrated the essential role of estrogen signaling in the formation of uterine squamous metaplasia in the *postaxial hemimelia* mice (Kurita and Nakamura, 2008), a spontaneous *Wnt7a* null mutant strain (Parr et al., 1998). When *Wnt7a* null mice were ovariectomized at P35, the uterine epithelium remained columnar, negative for p63 (Fig. 5B) and positive for PR in the absence of estrogen at P60 (Kurita and Nakamura, 2008) (Fig. 5B'). However, 17 $\beta$ -estradiol treatment at P60 induced p63 expression and squamous transformation in the uterine epithelium of ovariectomized *Wnt7a* null mice (Fig. 5C) (Kurita and Nakamura, 2008).

The Canonical Wnt pathway is initiated by the binding of the Wnt protein to the membrane receptors, followed by downstream cytoplasmic events that lead to stabilization and nuclear transportation of  $\beta$ -catenin. In the nucleus,  $\beta$ -catenin binds to the T-cell factor/lymphocyte enhancer factor (Tcf/Lef) transcription factor, resulting in the activation of Wnt target genes (Cadigan and Peifer, 2009). The conditional inactivation of  $\beta$ -catenin in the uterus induced expression of p63 in the uterine epithelium of adult mice (Jeong et al., 2009), confirming the importance of the canonical Wnt/ $\beta$ -catenin signaling pathway in the maintenance of the uterine epithelial identity. The regulation of p63 by Wnts in the uterine epithelium, however,



appears to involve more than just Wnt7a and  $\beta$ -catenin. In the uterus, transcripts for Wnts 4, 5a, 7a, 7b, 10b 11 and 16 have been detected (Wang and Shackleford, 1996; Hayashi et al., 2009; Miller et al., 1998), thus the loss of  $\beta$ -catenin in the uterus can affect the downstream signaling of many Wnt family members. Indeed, the conditional inactivation of Wnt4 in the uterus also induced p63 expression in the adult uterine epithelium (Franco et al., 2010). Since Wnt4 was also down-regulated in the uterus of Wnt7a null mutant mice (Miller and Sassoan, 1998), it is possible that the development of uterine squamous metaplasia in the Wnt7a null mutant mice was mediated by the down-regulation of Wnt4. In addition, the loss of Wnt7a affects expression of Wnt5a, Hoxa10 and Hoxa11 in the adult uterus (Miller and Sassoan, 1998). Therefore, in the adult uterus, epithelial differentiation is maintained by a complex signaling network in which canonical Wnt signaling pathway plays the pivotal role. Wnt/ $\beta$ -catenin signaling pathway controls maintenance and differentiation cell fate of epithelial stem cells in the epidermis and gastrointestinal tract (DasGupta and Fuchs, 1999; Huelsken et al., 2001; Korinek et al., 1998; Grigoryan et al., 2008). The same signaling pathway may also control the number and cell fate of stem cells in the FRT, and uterine squamous metaplasia may be a result of the dysregulation of the stem cell maintenance in the adult uterine epithelium.

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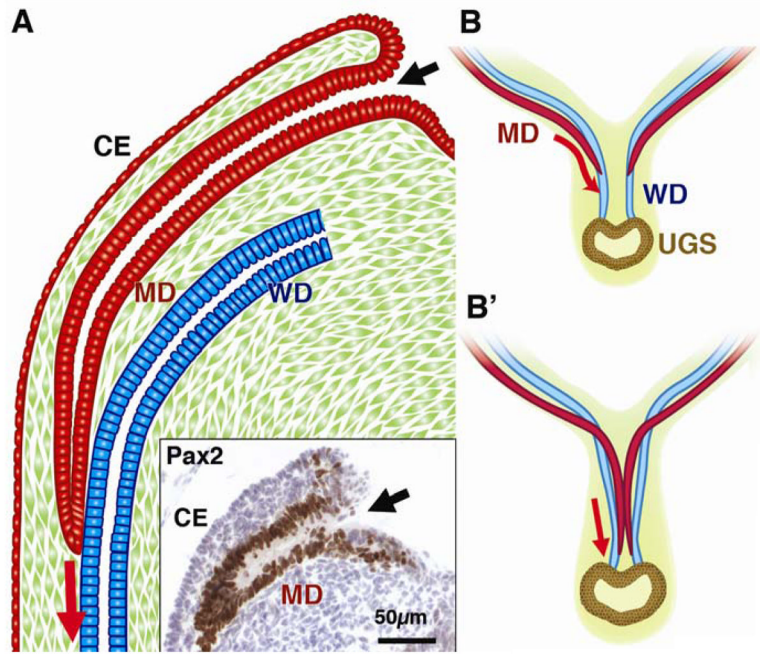
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**Figure 1. Development of the MD**

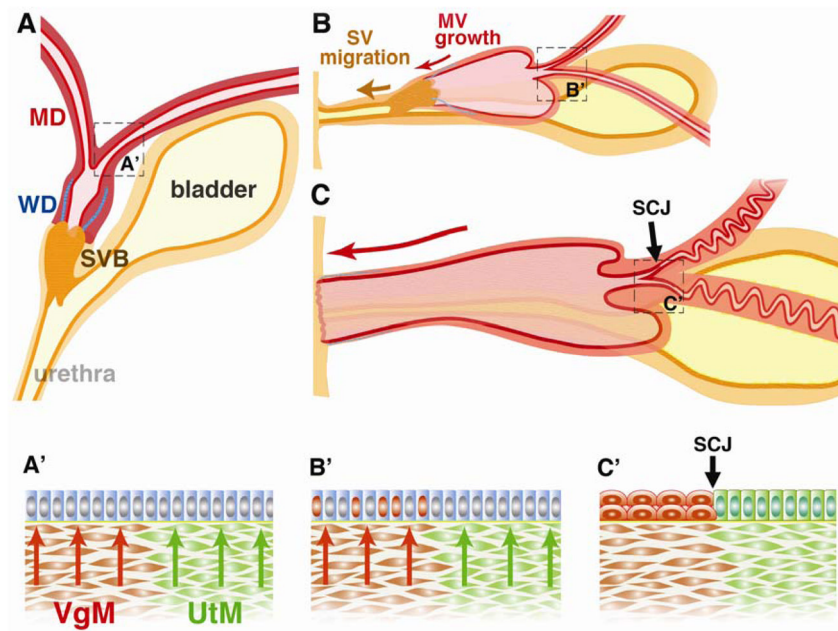
CE; coelomic epithelium, MD; Müllerian duct, WD; Wolffian duct, UGS; urogenital sinus

A; Formation of MD

The MD arises as an invagination of CE at the cranial end of the urogenital ridge. The insert demonstrates Pax2 immunohistochemistry on the E13.5 female mouse embryo. The Pax2 is essential for the development of the MD (Torres et al., 1995) and its expression differentiates the MDE (brown cells in the insert) from the CE. The site of infolding remains open throughout development (black arrows). The MD grows caudally through the urogenital ridge mesenchyme and the tip comes into contact with the WD within a common basement membrane. Afterwards, the tip of the growing MD maintains close contact with WD while the cranial portion is separated from the WD by intervening mesenchyme.

B; Caudal growth and fusion of the MDs

The MDs remain in contact with the WDs and use them as a guide during their caudal growth. As the MDs grow caudally, they cross over the WDs and meet in the midline to fuse with each other (B'). The caudal tips of the MDs remain separated to keep contact with the WDs (Hashimoto, 2003). Right before the MD tips reach the urogenital sinus, they finally become united and fuse with the UGS.



**Figure 2. Formation of mouse cervix and vagina**

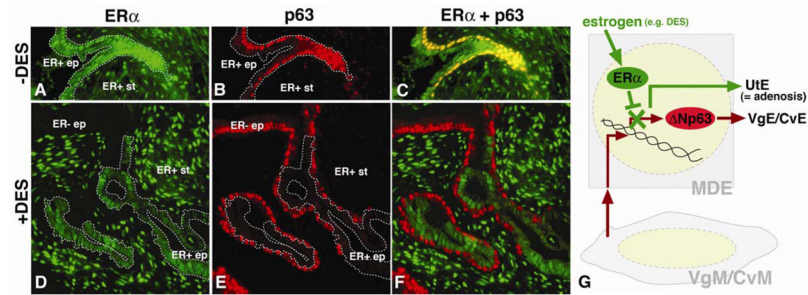
MD; Müllerian duct, WD; Wolffian duct, UGS; urogenital sinus, SVB; sinovaginal bulb. SV; sinus vagina, MV; Müllerian vagina, SJC; squamocolumnar junction, VgM; vaginal mesenchyme, UtM; uterine mesenchyme.

A. Late embryonic state (~E16). The MD, WD and UGS are present. The cranial portion of WD is regressed by this stage. At this stage, the MDE is uniformly undifferentiated (A').

B. Perinatal stage. The SV moves caudally as the MV elongates caudally. During this caudal migration, the SV maintains its connection to the urethra. Around this time,  $\Delta Np63$  expression is induced in epithelial cells (red nuclei in B') of the cervix and vagina in response to the mesenchymal induction (red arrows). By this time, the majority of the WD (blue lines) is regressed.

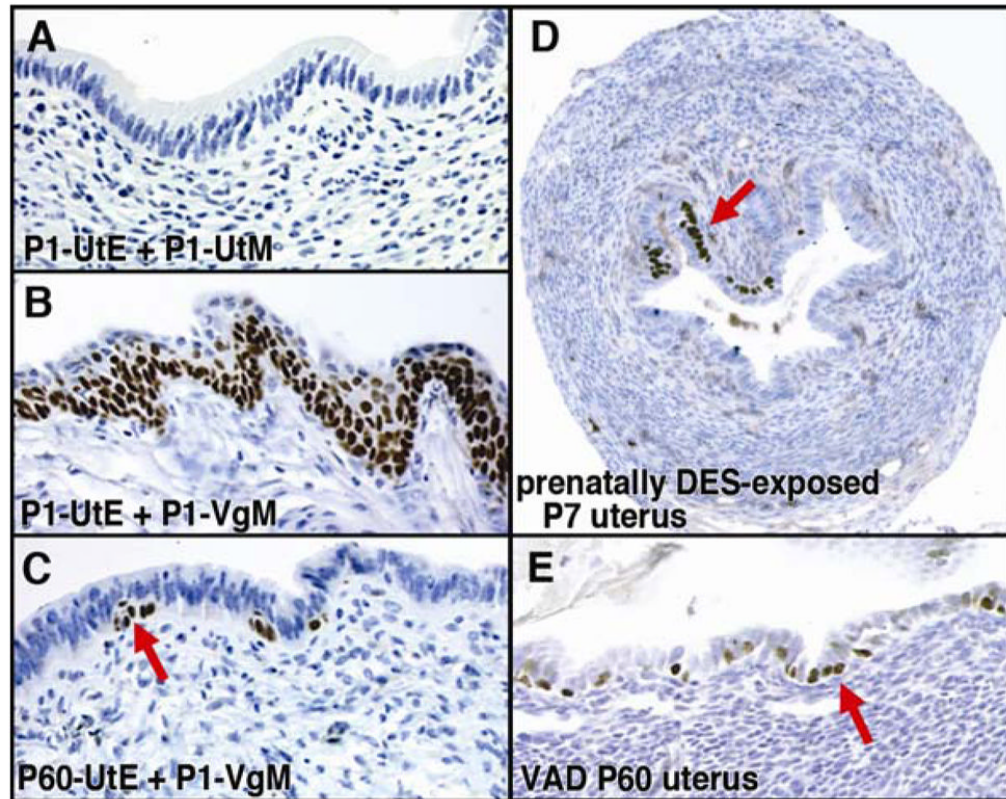
C. Pubertal stage. The MV reaches the posterior body wall in the neonatal stage. At puberty, the solid epithelial cord of the SV is canalized and the vaginal orifice is formed. The entire vagina is lined by epithelial cells derived from MDE. In the cervix, the SCJ is formed as a result of mesenchymal induction (C'). The residual segments of WDs may still be present in the vaginal stroma, and the amount of WDE remnants varies among adult female mice.





**Figure 3. Cell-autonomous inhibition of p63 expression by DES/ER $\alpha$  in MDE**

ER $\alpha$ ; estrogen receptor  $\alpha$ , UtE; uterine epithelium, VgE; vaginal epithelium, CvE; cervical epithelium, VgM; vaginal mesenchyme, CvM; cervical mesenchyme, ep; epithelium, st; stroma, ER+; ER $\alpha$  positive ER-; ER $\alpha$  negative. Dotted line indicates ER+ epithelial cells. The uterine epithelial cells from P1 ER $\alpha$  null and wild-type mice were mixed, combined with P1 rat VgM and grafted under the subrenal capsule of female host nude mice with/ without subcutaneous implantation of a 25 $\mu$ g DES pellet (Kurita et al., 2004). In the absence of DES, p63 was induced in the entire epithelium, which contained both ER $\alpha$  positive and negative epithelial cells. The panels A –C show the area with double positive epithelial cells for ER $\alpha$  and p63. In contrast, in the presence of DES, p63 was induced only in the ER $\alpha$  negative cells. The panels D– F show exclusive expression of ER $\alpha$  and p63 in the epithelium in the DES-treated host. These data confirm the conclusion of our previous study that DES inhibits expression of p63 through ER $\alpha$  in the epithelial cells. ER $\alpha$  in the VgM/CvM does not inhibit induction of p63 in the MDE. Furthermore, co-localization of ER+/p63-negative and ER-/p63-positive epithelial cells indicates that the inhibitory effect of DES/ER $\alpha$  on p63 expression is cell-autonomous, as illustrated in G.



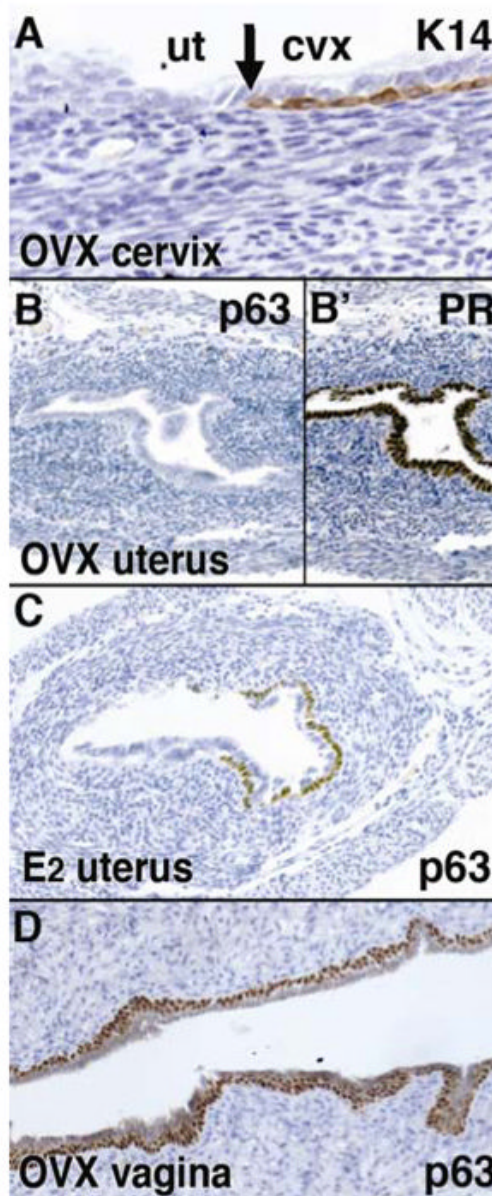
**Figure 4. Uterine squamous metaplasia**

UtE; uterine epithelium, VgE; vaginal epithelium, UtM; uterine mesenchyme, VgM; vaginal mesenchyme

A– C; Tissue recombinants composed with P1 UtE + P1 UtM (A), P1 UtE + P1 VgM (B), and P60 UtE + P1 VgM (C). When the UtE from the P1 mouse is associated with a P1 UtM, it develops into simple columnar UtE, which is negative for p63 (A). The same P1 UtE can be induced to be stratified squamous VgE by P1 VgM (B). In contrast, UtE from mature mice have limited potential to transdifferentiate into VgE. UtE from two month old virgin mice remains mostly simple columnar when it is combined with P1 VgM (C). However, the bipotency of UtE is not completely lost even in the adult mice, and a small number of UtE cells can be induced to express p63 in response to the P1 VgM (C, red arrow). These developmentally plastic cells are likely to be the target of VAD-induced squamous metaplasia (E).

D. Expression of p63 in the uterus of prenatally (gestational day 14 – 17) DES-exposed mouse. The ectopic expression of p63 in the uterus was detected at P7 (red arrow). These cells are believed to develop into squamous metaplasia in a mature animal.

E. Vitamin A deficiency induces expression of p63 in the adult uterus. When female mice were fed a vitamin A deficient diet from P21, p63 positive cells were detected in the UtE by four-month-old.



**Figure 5.**

Uterine and vaginal epithelial differentiation phenotypes of *Wnt7a* null mutant, *postaxial hemimelia* (*px*) mice. Female *px* mice were ovariectomized (OVX) at P35 and the expressions of K14 (A), PR (B') and p63 (B, C and D) were analyzed at P60. The SCJ was normally formed at cervix (A). UtE was columnar, positive for PR (B') and negative for p63 and K14 (B), whereas CvE/VgE were stratified squamous with the expression of p63 (D). However, IP injection of 125ng 17 $\beta$ -estradiol (E<sub>2</sub>)/day for 3 days from P60 induced p63 expression in the uterus (C). Therefore, *Wnt7a* is required for stabilization of the epithelial cell fate in the uterus, but is dispensable for induction of the uterine/vaginal epithelial cell fate.