



# Expression of mRNA Mastermind-like Domain-containing 1. Androgen Receptor, and Estrogen Receptor in Patients with **Hypospadias**

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#### Abstract

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BACKGROUND: Androgen (AR) and Estrogen (ER) hormones play an important role in the prenatal and postnatal development of the urogenital tract and especially the penis. Growth factors also influence the development of genital structures. Little is known about the exact role of Mastermind Like Domain 1 (MAMLD1) in sexual development. A role in sex differentiation through supporting testosterone production in critical periods of male development has been suggested. MAMLD1 mutations result in hypospadias with and without a disorder of sexual development (DSD) primarily because of compromised testosterone production around the critical period for fetal sex development, but the underlying etiology remains unclear.

AIM: The objective of this study was to investigate the correlation between gene factor MAMLD1, AR, ER1, and ER2 with the incidence of hypospadias

OBJECTIVE: The objective of this study was to investigate the correlation between gene factor MAMLD1, AR, ER1, and ER2 with the incidence of hypospadias.

METHODS: From 2017-2018, peri-urethral dartos were harvested from 46 patients with proximal hypospadias, 24 patients with distal hypospadias and 10 patients with normal penile were used as controls. The expressions of MAMLD1, AR, ER1, and ER2 were investigated by one-step quantitative polymerase chain reaction.

RESULTS: Median age was 5 years old in the 70 patients with hypospadias and 6 years old in the control subjects. Total specimens taken included 24 distal penile, 46 proximal penile, and 10 normal penile specimens. We found decreasing MAMLD1 and AR expressions, but ER2 expression increased in patients with hypospadias compared to controls, which was statistically significant (p < 0.001). A positive correlation between MAMLD1 and AR was found in patients with hypospadias (r = 0.062; p = 0.038).

CONCLUSIONS: Decreasing of MAMLD1 and AR expression was followed by increasing ER2 expression in patients with hypospadias. MAMLD1 had a positive correlation with AR so the defect of MAMLD1 may influence AR and increase the incidence of hypospadias.

# Introduction

Hypospadias is a relatively common genital anomaly in humans in which the urethral opening is not correctly positioned at the tip of the penis. Its prevalence in many countries ranges from 4 to 43 cases per 10,000 births [1]. This congenital disorder is the second most common in boys after cryptorchidism and the most frequent malformation of the penis [2]. Hypospadias is defined as a malformation of the penis due to incomplete development of the male urethra leading to the localization of the urinary meatus on the ventral side of the penis in a variable position from the glans to the perineum [3], [4]. The precise etiology remains unclear in most disorder of sexual development (DSD) cases including hypospadias. Several candidate genes and their response to environmental agents have been analyzed both in human and mouse tissues, opening many promising areas of research concerning the etiology of hypospadias [5].

Mastermind-like domain-containing 1 (MAMLD1, previously known as CXORF6) is an X-linked transcriptional coactivator [6]. It is expressed in Sertoli and Leydig cells of fetal mice around the critical period for sexual development [7]. MAMLD1 is known to be implicated in various human DSDs, including hypospadias. MAMLD appears to regulate testosterone production around the critical period for sex development [8]. However, the underlying mechanism of hypospadias' pathogenesis due to MAMLD1 is poorly understood.

Several genes may affect the anomalies of male development such as androgen receptors (AR) and estrogen receptors (ER). The previous studies report that several mutations specifically affect AR synthesis in the testis which can lead to hypospadias. Several studies reported hemizygous mutations (as AR is X-linked) in patients with hypospadias [9]. ER dysfunction may also contribute to the pathogenesis of hypospadias. Various homozygous mutations have been reported to cause hypospadias and cryptorchidism in the 46 XY sibling [10]. As previously reported, there are two ER receptors, ER1 and ER2, which interact to form homo-heterodimer ER complexes. ER bound by ER translocates to the nucleus to activate ER-responsive genes. ER1 and ER2 are expressed in most cells of the male urethra [11]. This study examined the interaction between MAMLD1, AR, ER1, and ER2 in the development of hypospadias. We compared the expression levels of MAMLD1, AR, ER1, and ER2 between normal boys and those with hypospadias.

The objective of this study was to investigate the correlation between gene factor MAMLD1, AR, ER1, and ER2 with the incidence of hypospadias.

## Methods

#### Patients

Seventy patients underwent repair of hypospadias without DSD and 10 normal patients who underwent circumcision as controls in four pediatric urology centers from 2017 to 2018. We divided the samples into distal hypospadias, proximal hypospadias, and controls. We used periurethral dartos obtained during chordee excision following the standard procedure using samples from RNAlater fixation. The surgical approach of each patient was decided depending on the patient's condition with distal hypospadias treated by tubularized incision plate, or Mathieu onlay island flap urethroplasty, and proximal hypospadias using Duckett, two-stage urethroplasty.

#### Polymerase chain reaction (PCR) analysis

All foreskin tissues of the patients were pooled in a single tube. Total RNA was extracted from a homogenized sample using the Hybrid- $R^{TM}$  Isolation Kit and cDNA was synthesized from 200 ng of total RNA using NEXpro<sup>TM</sup> quantitative reverse transcription (qRT)-PCR kit. Quantitative PCR was performed for MAMLD1, AR, ER1, and ER2. We used GAPDH as internal control, Table 1. Amplification conditions consisted of an initial denaturation step at 95°C for 10 min, followed by 40 cycles at 95°C for 20 s, at 56°C for 40 s, and at 72°C for 60 s. An extension was carried out at 72°C for 5 min. The quantitative qPCR amplified samples were analyzed by the BiONEER Exicycle<sup>TM</sup> 96 (BioNEER, Daejeon, South Korea).

#### Statistical analysis

Differences with statistical significance were determined using one-way ANOVA test or Kruskal–Wallis test, and Mann–Whitney U-test was used to compare differences between groups. P values < 0.05 were considered as statistically significant.

Table 1: MAMLD1, AR, ER1, ER2, and GADPH primers

Gen	Forward primer	Reverse primer
MAMLD1	TTTGCAACTCCAGCATCATC	CCTAACTCCTTCCCCTGGAC
AR	CCTGGCTTCCGCAACTTACAC	GGACTTGTGCATGCGGTACTCA
ER1	ATTTCTCTGCGCCCCTAGAC	CATAGTGGTACCCAGACGCA
ER2	TGAACTGGGCCCAGACTATG	ATGGTACCCTGAGGCGTAGT
GADPH	5'-GCATCCTGGGCTACACTGAG-3'	5'-TCCACCACCCTGTTGCTGTA-3'
	D. Colorado MAMI D4: Masterial like dee	

AR: Androgen, ER: Estrogen, MAMLD1: Mastermind-like domain 1

#### **Results**

Periurethral dartos was harvested from 46 patients with proximal hypospadias, 24 patients with distal hypospadias during urethroplasty, and 10 patients with normal penile during circumcision as controls. Characteristic data of patients are presented in Table 2.

#### Table 2: Variable characteristics

<u> </u>	N / 1
Variables	Value
Age, year	5 (1-14)
Hypospadias type, n (%)	
Distal	24 (30)
Glandular	5 (6.3)
Subcoronal	2 (2.5)
Distal penile	2 (2.5)
Midshaft	14 (17.5)
Proximal penile	1 (1.3)
Proximal	46 (57.5)
Penoscrotal	33 (41.3)
Scrotal	12 (15.0)
Perineal	1 (1.3)
Control	10 (12.5)
MAMLD1, median (min-max)	4.59 (1.86-10.55)
AR, Mean (SD)*	6.84 ± 2.57
ER1, Median (min-max)	14.92 (4.59-39.39)
ER2, Mean (SD)*	21.89 (8.14)
Hypospadias characteristics penile torsion, n (%)	· · ·
Yes	4 (5.7)
No	66 (94.3)
Penile curvature, n (%)	
Mild (<30 degrees)	22 (31.4)
Moderate (30-60 degrees)	14 (20)
Severe (>60 degrees)	34 (48.6)
Urethral plate, n (%)	
Narrow (<8mm)	36 (51.4)
Wide (≥8 mm)	34 (48.6)
Penoscrotal transposition	
Yes	9 (12.9)
No	61 (87.1)
Bifid scrotum	
Yes	10 (14.3)
No	60 (85.7)
Urethroplastv	
First urethroplasty	47 (67.1)
Redo-urethroplasty	23 (32.9)

\*Kolmogorov–Smirnov p>0.05; AR: Androgen, ER: Estrogen, MAMLD1: Mastermind-like domain 1, SD: Standard deviation, VEGF: Vascular endothelial growth factor. The quantitative PCR analyses showed significantly decreased MAMLD1 expression in both distal (5.33  $\pm$  2.21) and proximal (4.82  $\pm$  2.42) hypospadias compared with the control group (9.55  $\pm$  0.93) (p < 0.05). Meanwhile, the expression of the AR showed decreases in both distal (5.33  $\pm$  2.21) and proximal (4.82  $\pm$  2.42) hypospadias compared to the control group (9.55  $\pm$  0.93) with significant difference (p < 0.05). In line with the result of MAMLD1 expression, ER1 showed decreasing

Table 3: Compared mean analysis among the groups

Variable	Proximal hypospa	adias Distal hypospac	dias Control	р		
MAMLD1 <sup>#</sup>	4.82 (± 2.42)	5.33 (± 2.21)	9.55 (± 0.93)	< 0.001		
AR*	6.52 (± 2.61)	6.27 (± 2.18)	9.69 (± 1.10)	< 0.001		
ER1 <sup>#</sup>	15.31 (± 6.49)	17.93 (± 8.22)	12.82 (± 2.23)	0.158		
ER2*	24.44 (± 8.41)	21.22 (± 5.38)	11.79 (± 2.49)	< 0.001		
"Kruskal Wallis *one way ANOVA AP: Androgen EP: Estrogen MAMI D1: Mastermind like domain 1						

expression in both hypospadias types, with proximal  $(15.31 \pm 6.49)$  and distal  $(17.93 \pm 8.22)$  compared to the control group  $(12.82 \pm 2.23)$ . Reciprocally with ER1 and ER2 showed increasing expression in both hypospadias types, with proximal  $(24.44 \pm 8.41)$  and distal  $(21.22 \pm 5.38)$  compared to the control group  $(11.79 \pm 2.49)$  (Table 3 and Figure 1).



Figure 1: Expression of MAMLD1, androgen receptor, estrogen receptor 1, and estrogen receptor 2 in patients with hypospadias compared to the control group. AR: Androgen, ER: Estrogen, MAMLD1: Mastermind-like domain 1,\*statistically significant

*Post hoc* analyses were used to explore which group was significantly related. It was revealed that there were significant differences in the MAMLD1, AR, and ER2 hypospadia groups (both distal and proximal) compared with the control group, while among hypospadias types, there were no significant differences (Table 4).

	Table	4:	Post	hoc	analysis	;
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Variable dependent	Hypospadias type		р	95% CI	95% CI	
				Lower	Upper	
MAMLD1 <sup>#</sup>	Distal	Proximal	0.199	N/A	N/A	
	Distal	Control	<0.001	N/A	N/A	
	Proximal	Control	< 0.001	N/A	N/A	
AR*	Distal	Proximal	1.00	-1.69	1.21	
	Distal	Control	<0.001	-5.59	-1.26	
	Proximal	Control	<0.001	-5.19	-1.17	
ER1 <sup>#</sup>	Distal	Proximal	0.195	N/A	N/A	
	Distal	Control	0.059	N/A	N/A	
	Proximal	Control	0.121	N/A	N/A	
ER2*	Distal	Proximal	0.230	-7.60	1.17	
	Distal	Control	0.002	2.87	15.98	
	Proximal	Control	<0.001	6.56	18.72	

like domain 1.

Linear regression analysis of each of the mRNA expressions showed the AR expression and MAMLD1 expression had strong correlation (r = 0.062) and the positive pattern indicates that the more MAMLD1 expression is reduced, the AR expression will also

decrease. Furthermore, the *post hoc* results showed a significant association between AR expression and MAMLD1 expression with p = 0.038 (Table 5).

## Discussion

The etiology of hypospadias cases, however, is not yet solved. Genes or environmental factors may play a principal role in the pathogenesis of hypospadias. Hypospadias can be equally transmitted through the maternal and paternal sides of the family and recurrence risks for brothers and sons of hypospadias cases are similar [12]. Various genes are known to be implicated in testes development, masculinization, and hypospadias such as MAMLD1, AR, ER1, and ER2. Mutations that affect the ability of the gonads to produce these hormones or the cell's sensitivity to these hormones can result in anomalies of male development including hypospadias [6].

MAMLD1 (previously known as CXorf6) located on human chromosome Xq28 is a causative gene for hypospadias, a mild form of 46, XY DSD [13]. At present, multiple mutations have been reported in patients with various types of hypospadias [13], [14], [15]. In this regard, the mouse homologous gene MAMLD1 is transiently expressed in fetal Sertoli and Leydig cells around the critical period for sex development [13], and transient MAMLD1 knockdown using small interfering RNAs (siRNAs) reduces testosterone production in cultured mouse Leydig tumor cells (MLTCs) [13]. Various mutations affect the expressions of MAMLD1 in patients with hypospadias. They were downregulated compared to the control group. These findings are similar with other studies which stated that the mRNA level of MAMLD1 was severely reduced (25-30%) in the MLTCs with reduction in testosterone production (50-60%) [8]. This association between testosterone production and expression of MAMLD1 is consistent with the hypospadias phenotype in the affected patients [16].

ARs affect and play an important role in the prenatal and postnatal development of the urogenital tract, especially the penis [17]. The present study showed a decrease in the expression of the AR in patients with hypospadias compared to the control group. Decreased AR expression may be affected by elevation in the number of CAG repeats. The relationship between CAG repeats and AR transcriptional activity has not been well explored. The AR gene is composed of eight exons and is located on the X chromosome at q11-q12. Exon 1 of the AR gene contains a polymorphic sequence of CAG repeats, which usually varies in number from 10 to 35, which encodes polyglutamine stretches of AR transactivation domain [18], [19]. Many findings suggest that the CAG number is negatively correlated with AR transcriptional activity [20]. In fact, patients affected by Kennedy

Table 5: Linear regression analysis between MAMLD1 and AR, ER1,	and ER2 in hypospadias group
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Model		Unstandardize	d coefficients	Standardized coefficients	r	t	р	95% confidence interval for B	
		В	SE	Beta				Lower bound	Upper bound
AR	(Constant)	5.129	0.678			7.567	0.000	3.776	6.481
	MAMLD1	0.261	0.123	0.249	0.062	2.120	0.038	.015	0.506
ER1	(Constant)	17.990	2.029			8.867	0.000	13.942	22.039
	MAMLD1	-0.357	0.368	-0.117	0.014	-0.971	0.335	-1.092	0.377
ER2	(Constant)	26.075	2.137			12.199	0.000	21.810	30.340
	`MAMLD1	-0.548	0.388	-0.169	0.029	-1.415	0.162	-1.322	0.225

AR: Androgen, ER: Estrogen, MAMLD1: Mastermind-like domain 1, SD: Standard deviation.

syndrome have a number of CAG repeats >40 together with decreased virilization, reduced sperm production. testicular atrophy, and infertility [18]. Repeat length is positively affecting AR transcriptional activity in skeletal muscle, while the opposite appears in prostate cells. It is possible that alterations of the repeat length induce conformational changes in the AR N-terminal domain that enhance or suppress transcriptional cofactor and AR-associated protein interactions [21]. Other studies have shown that ethnic differences in CAG repeat length of the AR gene exist between African, Caucasian, and Asian populations. The allele expansion found in Africans is between 18 and 20 repeats, although some African subpopulations seem to have a shorter track [22]. In contrast, Caucasian and Asian populations have a longer expansion, where the mean number of CAG repeats is 21-22 in Caucasians [20] and 23 in Asians [23].

This research also found a positive correlation between MAMLD1 and AR. The result of our study showed a decreased in expression MAMLD1 similar to the expression AR which has decreased. It is known that defects in the AR function and its regulatory mechanisms contribute to the development of hypospadias [24].

Several studies showed that the expressions of ER and AR can also influence the development of male external genitalia [11], [25], [26], [27], [28]. Our study found that both ERs, including ER1 and ER2, have elevated expression compared to the control group. These findings show a counterbalance between AR and ER. Dominant expression of ER receptors in the penile tissues of children with hypospadias and the theory of disrupted AR and ER receptors interaction and/or balance could play a role during the development of external genitalia in patients with hypospadias [17]. Reduced testosterone levels and elevated ER levels may also be a risk factor for hypospadias [6], [29]. Interestingly, there were no studies that definitively stated that endocrine disruptors caused hypospadias. However, there were multiple studies that implicated endocrine disruptors as one component of a multifactorial model for hypospadias.

One limitation of this study is the nature of the cross-sectional study that does not include following the possibility of changes of the expression ER2 overtime. However, as far as we know, this is the first study that discussed about ER2 and its role in hypospadias in an Asian population.

#### Conclusions

Decreasing MAMLD1 and AR expressions were followed by increasing ER2 expression in hypospadias patients. MAMLD1 had a positive correlation with AR so the defect of MAMLD1 may influence AR and increase the incidence of hypospadias.

# Ethics Approval and Consent to Participate

The Ethical Committee of Faculty of Medicine, Universitas Gadjah Mada/Dr. Sardjito Hospital gave approval for this study (KE/1049/09). Written informed consent was obtained from all parents for participating this study.

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