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MAMLD1 mutations seem not sufficient to explain a 46,XY DSD phenotype. What else?

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Background

The *MAMLD1* gene (Xp28) is thought to cause disorder of sex development (DSD) in 46,XY patients, mostly presenting with hypospadias, and, recently, also gonadal dysgenesis. However, there is some controversy about the role of *MAMLD1* in sex development.

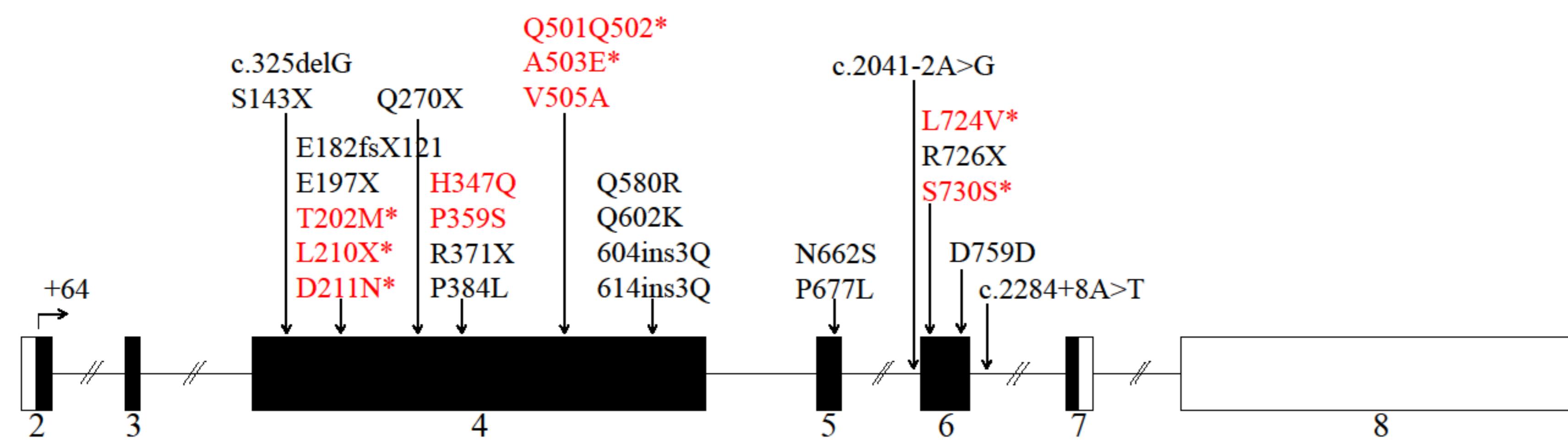
Aims and Objectives

We searched for *MAMLD1* sequence variations in 108 46,XY DSD individuals presenting with a wide spectrum of DSD phenotypes. Identified variations were functionally tested *in vitro*, and findings were compared with reported cases and the literature of *MAMLD1* focusing on sex development.

Methods and Results I

Nine *MAMLD1* mutations (7 novel) in 9 46,XY DSD patients were detected by Sanger sequencing.

All the mutations and polymorphisms detected in the *MAMLD1* gene



In red: present study; *: novel

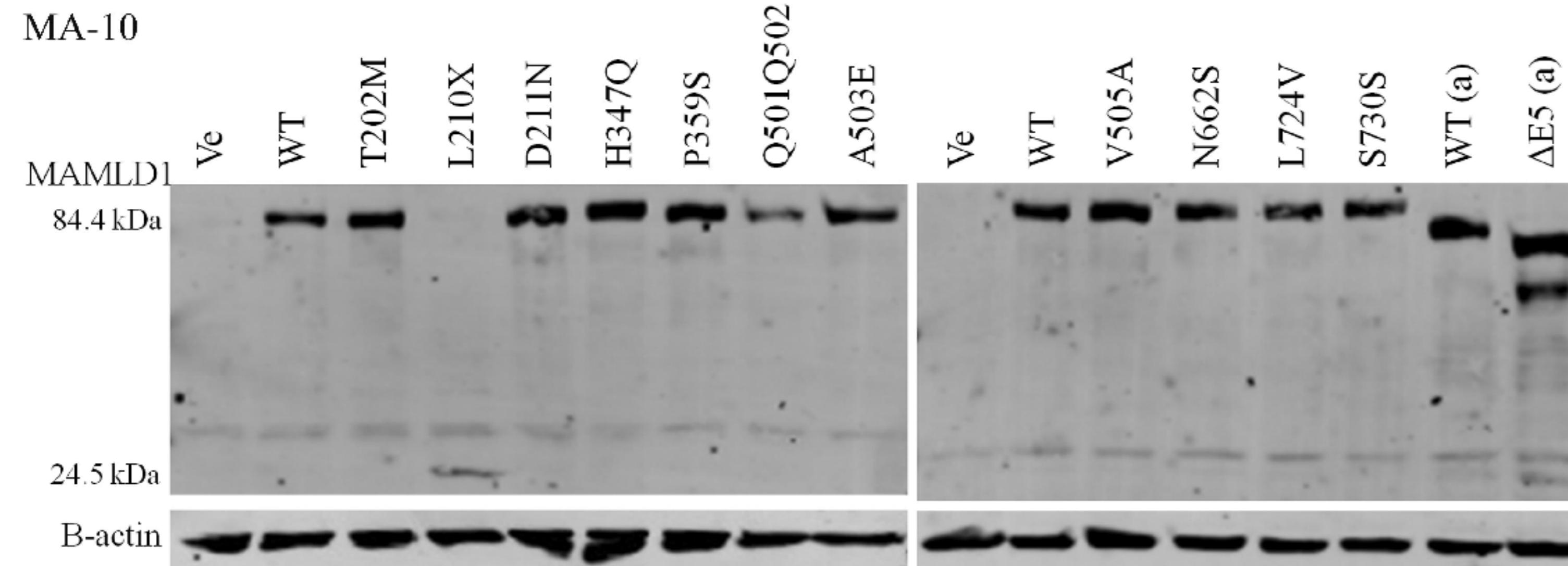
Characteristics of the 9 patients with *MAMLD1* mutations and polymorphisms

| Pat. | Karyotype | MAMLD1 gene mutation | Genital anatomy. <i>Testes histology</i> (age) | Gonadal function (age) | Adrenal function (age) | Remarks |
|------|-----------------|--------------------------------------|---|--|--------------------------|--|
| | | | | | | |
| 1 | 46,XY Male | T202M c.605C>T | Penoscrotal hypospadias. Small penis. Unilateral cryptorchidia. | Normal T (minipuberty). Normal gonadotropins. No hCG test. | ND | SGA. Short stature. Low implantation thumbs. |
| 2 | 46,XY Female | L210X c.626delT D211N c.631G>A | Penoscrotal hypospadias. Small penis. Testes 0.5 ml. <i>Normal for age</i> (15 d). | Normal T (baseline) and hCG test (10 days). | Normal baseline (15 d). | Abnormal GGN repeat in AR. Mother non-carrier; norfloxacin treatment during pregnancy. |
| 3 | 46,XY Female | H347Q c.1041C>A rs62641609 | Female genitalia. Gonads in labia. | Normal hCG test (2 y). | Normal baseline (2 y). | |
| 4 | 46,XY Male | H347Q c.1041C>A rs62641609 | Penoscrotal hypospadias. Testes 2 ml. | Normal hCG test. Normal AMH (2.5 y). | Normal baseline (2.5 y). | |
| 5 | 46,XY Male | Q501Q502 c.1503_1504dupCAGCAG | Hypospadias. Short penis. Delayed puberty. Testes 8 ml. | Baseline T and gonadotropins normal (70 y). Fathered a boy. | Normal baseline (70 y). | |
| 6 | 46,XY Male | A503E c.1508C>A | Penoscrotal hypospadias. Small penis. Testes 2 ml. | Normal baseline T (3 m). Normal hCG test (9 m). | Normal baseline (3 d). | |
| 7 | 46,XY Female | V505A c.1514T>C rs61740566 | Penoscrotal hypospadias. Small penis. Unilateral cryptorchidia. <i>Normal for age</i> (2 y). | Normal hCG test. | Normal Synacthen test. | |
| 8 | 46,XY Male | L724V c.2170C>G | Penoscrotal hypospadias. Small penis. Testes 1 ml. | Normal prepubertal baseline T (15 m). Normal AMH. | Normal baseline (15 m). | Esophageal atresia. Right aortic arch. |
| 9 | 46,XY Female | S730S c.2190G>A | Penoscrotal hypospadias. Small penis. <i>Normal for age</i> (nests of Normal Leydig cells; normal fertility index (1 y)). | Normal baseline (12 m). No hCG test. | | Müllerian ducts. |

ND: not done. d: day(s), m: month(s), y: year(s)

Methods and Results III

WT and mutant *MAMLD1* expression was assessed in mouse Leydig MA-10 cells.

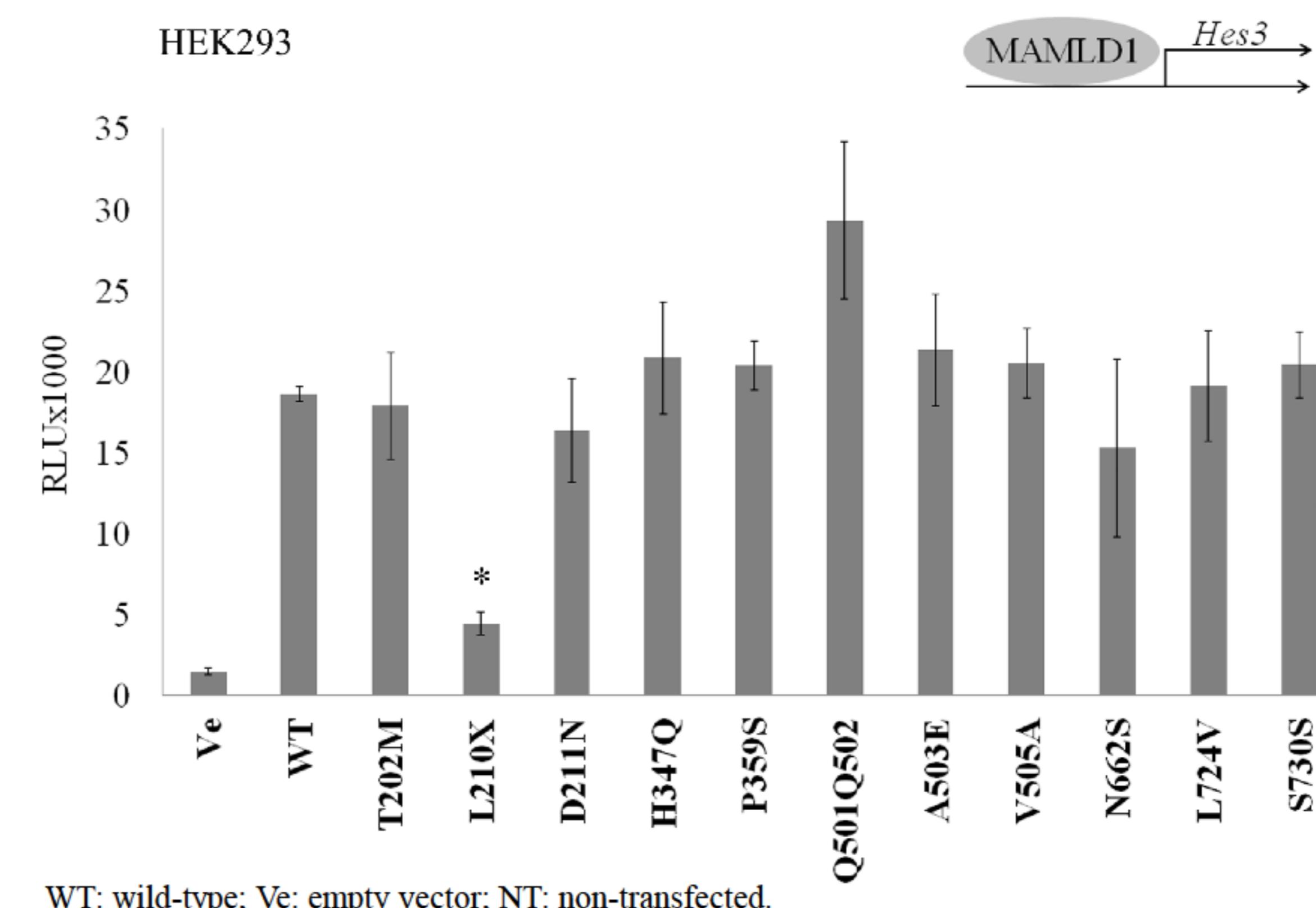


No difference for *MAMLD1* protein expression was found, except for a shorter L210X.

Methods and Results II

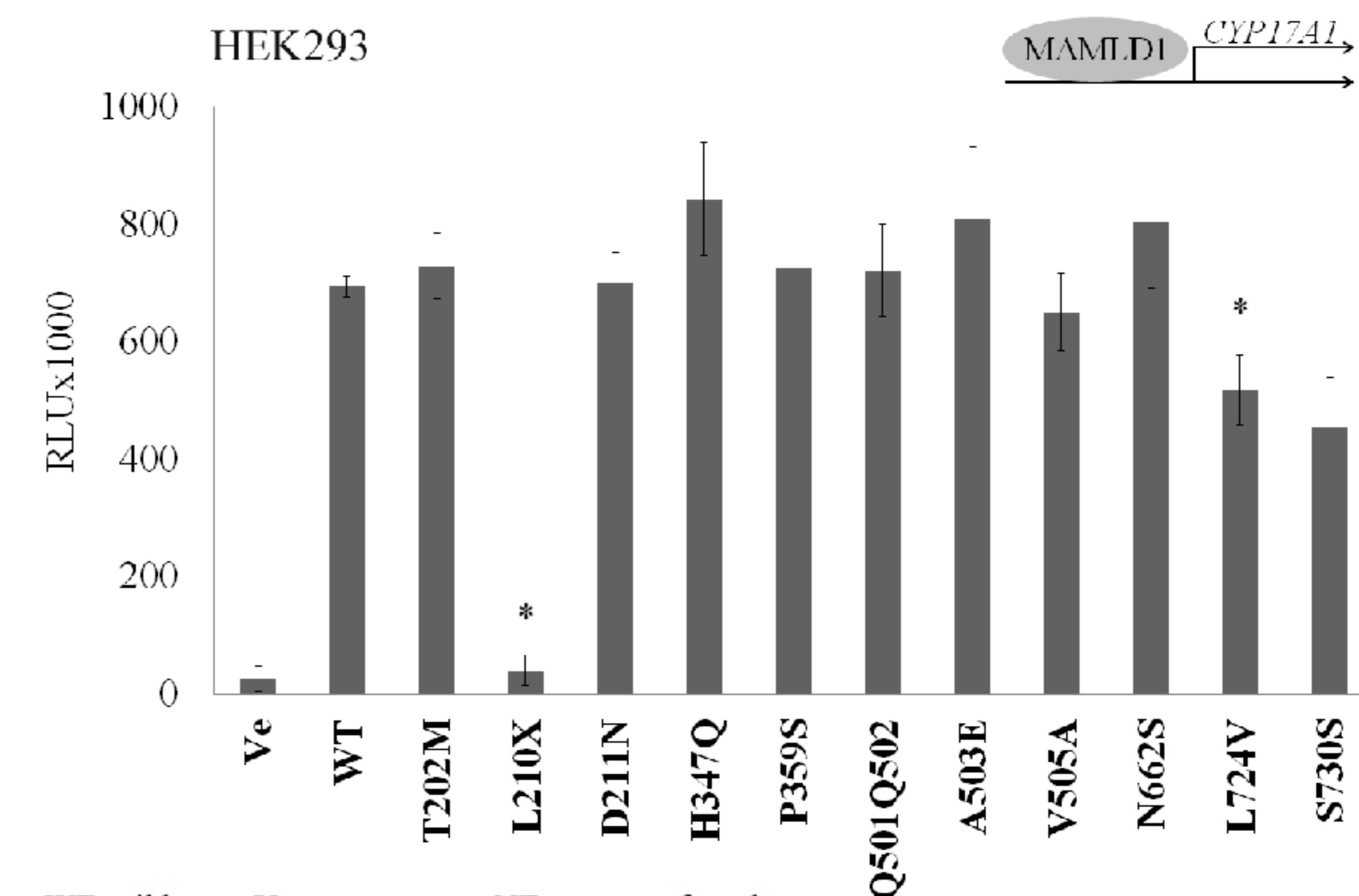
Functional studies were completed in non-steroidogenic HEK293, adrenal NCI-H295R and Leydig MA-10 cells.

MAMLD1 transcriptional activity tested on *Hes3* promoter by luc assays



WT: wild-type; Ve: empty vector; NT: non-transfected.

MAMLD1 transcriptional activity tested on *CYP17A1* promoter by luc assays

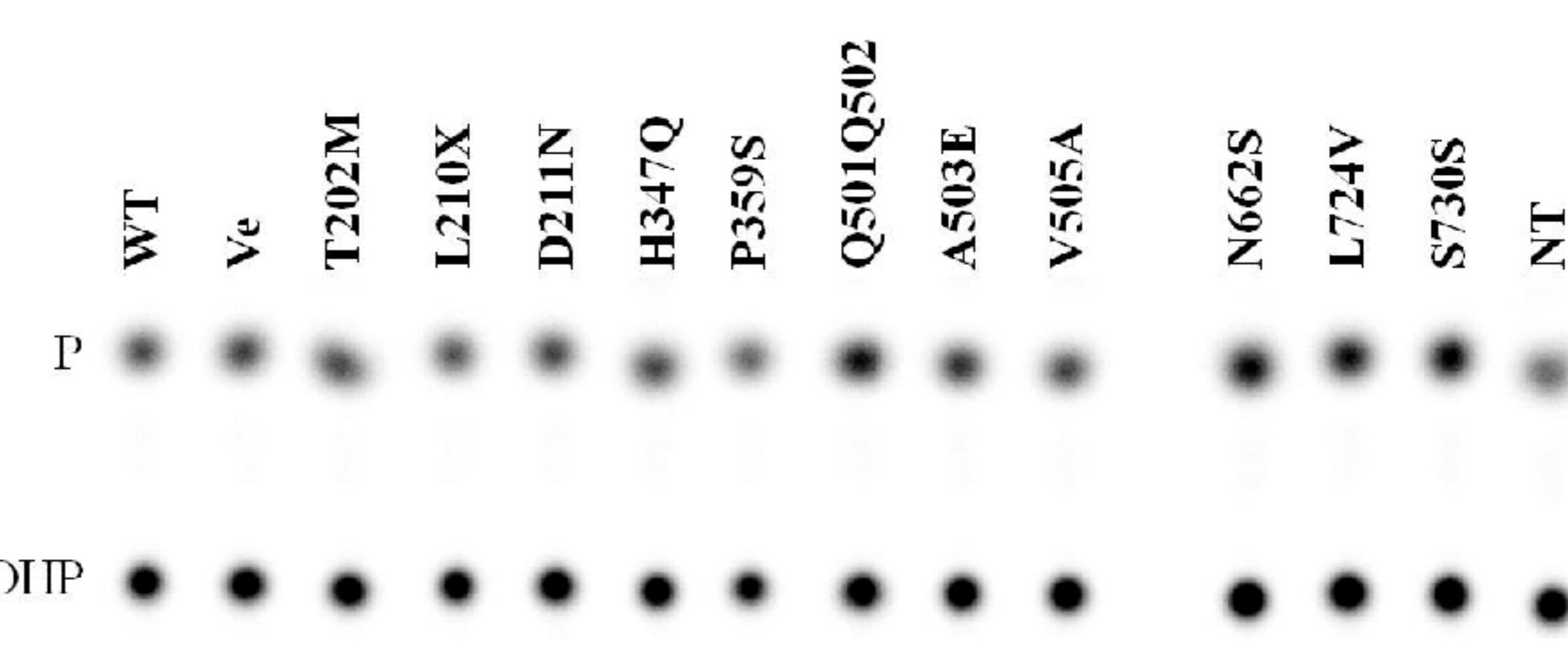


WT: wild-type; Ve: empty vector; NT: non-transfected.

Most *MAMLD1* variants acted similarly to the WT.

Only the L210X mutation showed loss of function in all tests, while variants L724V and S730S showed a decrease in *CYP17A1* promoter activation.

Effect of *MAMLD1* on *CYP17A1* activity in NCI-H295R cells



P: progesterone; 17OHP: 17-hydroxyprogesterone; WT: wild-type; Ve: empty vector; NT: non-transfected.

Effect of *MAMLD1* on androgen production assessed by testing the *CYP17A1* activity in the 3 cell lines. No effect of either WT or any *MAMLD1* variant on *CYP17A1* enzyme activity was found.

Conclusion

Our data support the notion that *MAMLD1* sequence variations may not suffice to explain the DSD phenotype in carriers.

