Ubqln3, a Testis-Specific Gene, is Dispensable for Embryonic Development and Spermatogenesis in Mice

Increasing lines of evidence suggest that the ubiquitin-proteasome pathway (UPP) plays a key role in spermatogenesis (Hou and Yang, 2013). Ubiquilins are ubiquitin-like proteins, all of which contain an amino-terminal UBL domain and a carboxy-terminal ubiquitin-associated (UBA) domain in its structure. Among all ubiquilin proteins identified so far, Ubiquilin 3 is the only one that has been reported to be testis-specific (Conklin et al., 2000). By examining the expression of *Ubqln3* mRNAs and protein in eight different mouse organs using quantitative PCR (qPCR) and Western blot, we found that *Ubqln3* was indeed expressed exclusively in the testis, with both transcript and protein detectable in the testes starting at postnatal Day 28 (P28); the highest levels were detected in adult testes (Fig. 1A–D).

The onset of testicular *Ubgln3* expression at P28 coincides with the elongation steps during late spermiogenesis, suggesting that Ubgln3 is mainly expressed in elongating/elongated spermatids. To define its physiological role in the testis, we generated Ubgln3 global knockout mice (on the C57BL/6J background) using a targeted embryonic stem cell line (cell-line number 11472C-G12) obtained from the Knockout Mouse Project repository; in this case, a gene-trap cassette (LacZ-pA) had recombined with exon 2 of Ubqln3, leading to a "gene trap" allele (Fig. 1E). Neither Ubgln3 mRNAs (Fig. 1F) nor UBQLN3 protein (Fig. 1B) was detected in homozygous knockout mice, suggesting that these animals are truly *Ubgln3*-null (herein called Ubqln3 KO mice). Both male and female Ubgln3 KO mice were viable and displayed normal development, with no discernible differences in growth compared to wild-type (WT) mice. Together, these data indicate that Ubgln3 is dispensable for embryonic and postnatal development in mice.

To determine the fertility of *Ubqln3* KO males, we performed a fecundity test using *Ubqln3* KO males bred with fertility-proven adult WT females. Our breeding data showed that no significant difference in either litter size or litter interval compared to those WT breeding pairs over a 6-month period (data not shown), suggesting that *Ubqln3* KO males are fertile. Consistent with their normal fertility, both sperm count and sperm motility are comparable between gametes isolated from *Ubqln3* KO and WT males (Fig. 1G–H) and histological analyses of *Ubqln3* KO testes and sperm demonstrated normal spermatogenesis and sperm morphology (Fig. 1I–J). Taken together, these

data indicate that *Ubqln3* is not required for male germ-cell development or spermatogenesis in mice. Interestingly, *Ubqln1*, *Ubqln2*, and *Ubqln4* transcript abundance was increased significantly in *Ubqln3* KO compared to WT testes (Fig. 1F), suggesting that other members of the ubiquilin family may compensate for the loss of *Ubqln3*. Therefore, the testis-specific expression of *Ubqln3* is dispensable for both embryonic development and spermatogenesis in mice.

SHUIQIAO YUAN, ¹ WEIBING QIN, ² CONNOR R. RIORDAN, ¹ HAYDEN MCSWIGGIN, ¹ HUILI ZHENG, ¹ AND WEI YAN ^{1,*} ¹Department of Physiology and Cell Biology, University of Nevada Reno School of Medicine, Reno, Nevada ²Family Planning Research Institute of Guangdong, Guangzhou, China

*Corresponding author:
Department of Physiology and
Cell Biology,
University of Nevada Reno
School of Medicine,
Reno, NV 89557.
Email: wyan@medicine.nevada.edu
Grant sponsor: National Institutes of Health (NIH);
Grant numbers: HD060858, HD071736,



Published online 16 March 2015 in Wiley Online Library (wileyonlinelibrary.com).

DOI 10.1002/mrd.22475

Mol. Reprod. Dev. 82: 266–267, 2015. © 2015 Wiley Periodicals, Inc.

Received 1 February 2015; Accepted 21 February 2015

ACKNOWLEDGMENT

This work was supported, in part, by National Institutes of Health (NIH) Grants HD060858, HD071736, and HD074573 (to W.Y.).

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HD074573

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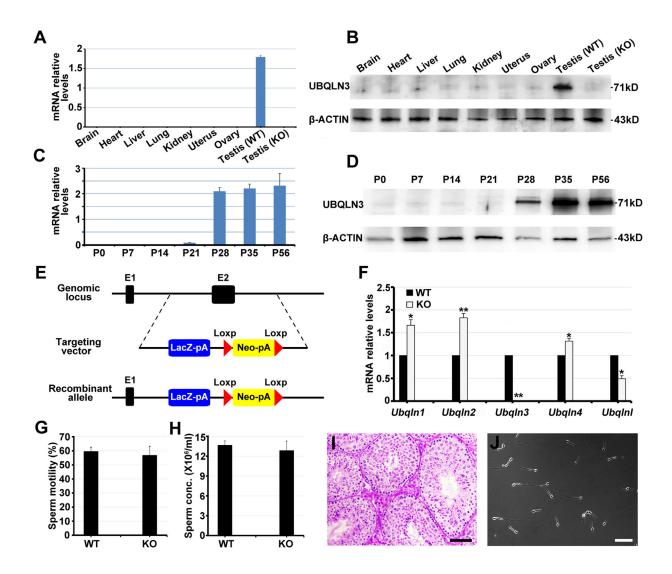


Figure 1. Ubqln3 is a testis-specific gene dispensable for spermatogenesis. A: qPCR analyses of Ubqln3 mRNA levels in eight organs of adult mice. B: A representative Western blot showing that UBQLN3 is exclusively detected in wild-type (WT) but not in Ubqln3 KO testes. (The rabbit polyclonal UBQLN3 antibody (1:1000) was purchased from ProteinTech Group, Inc. [Chicago, IL], catalog number 13568-1AP.) C: qPCR analyses of Ubqln3 mRNA levels in developing testes. Testes at postnatal Day 0 (P0, newborn), P7, P14, P21, P28, P35, and P56 were analyzed. D: A representative Western blot showing UBQLN3 expression in developing mouse testes. E: Schematic illustration of the targeting strategy for generating a Ubqln3-null allele in mouse embryonic stem cells. F: qPCR assays showing mRNA levels of five ubiquilin-family genes in Ubqln3 KO testes. P < 0.05, P < 0.05, P < 0.01 compared to WT (n = 3, Student's t-test). G and H: Normal sperm concentration and sperm motility in Ubqln3 KO males. I: Normal histology in Ubqln3 KO testes. J: Representative phase-contrast micrograph showing Ubqln3 KO sperm. Scale bars, 100 μm (I) or 50 μm (J).