



CLINICAL RESEARCH ARTICLE

Clinical features, genetic background, and outcome in infants with urinary tract infection and type IV renal tubular acidosis

Min-Hua Tseng¹, Jing-Long Huang², Shih-Ming Huang³, Jeng-Daw Tsai^{4,5,6,7}, Tai-Wei Wu⁸, Wen-Lang Fan⁹, Jhao-Jhuang Ding^{1,10} and Shih-Hua Lin¹¹

BACKGROUND: Type IV renal tubular acidosis (RTA) is a severe complication of urinary tract infection (UTI) in infants. A detailed clinical and molecular analysis is still lacking.

METHODS: Infants with UTI who exhibited features of type IV RTA were prospectively enrolled. Clinical, laboratory, and image characteristics and sequencing of genes responsible for phenotype were determined with follow-up.

RESULTS: The study cohort included 12 infants (9 males, age 1–8 months). All exhibited typical type IV RTA such as hyperkalemia with low transtubular potassium gradient, hyperchloremic metabolic acidosis with positive urine anion gap, hypovolemic hyponatremia with renal salt wasting, and high plasma renin and aldosterone levels. Seven had hyperkalemia-related arrhythmia and two of them developed life-threatening ventricular tachycardia. With prompt therapy, all clinical and biochemical abnormalities resolved within 1 week. Five had normal urinary tract anatomy, and three of them carried genetic variants on *NR3C2*. Three variants, c.1645T>G (S549A), c.538G>A (V180I), and c.1-2C>G, on *NR3C2* were identified in four patients. During follow-up, none of them had recurrent type IV RTA, but four developed renal scarring.

CONCLUSIONS: Genetic mutation on *NR3C2* may contribute to the development of type IV RTA as a complication of UTI in infants without identifiable risk factors, such as urinary tract anomalies.

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INTRODUCTION

Renal tubular acidosis (RTA) is traditionally classified into three forms: distal RTA (type I), proximal RTA (type II), and hyperkalemic RTA (type IV). Unlike type I and type II RTA that presents commonly with hypokalemia, type IV RTA is characterized by hyponatremia, hyperkalemia, and non-anion gap metabolic acidosis associated with reduced distal hydrogen secretion. The etiologies of type IV RTA are diverse, including inherited mutations in genes encoding mineralocorticoid receptor (MR) and renal epithelial sodium channel (ENaC) or acquired disorders, such as tubulointerstitial nephritis, lupus nephritis, and chronic kidney disease. Its course can be permanent or transient in nature. Type IV RTA has been also reported as a severe complication of urinary tract infection (UTI). Although the clinical features of type IV RTA in infants with UTI are usually non-specific and asymptomatic, cases with life-threatening hyperkalemia have been reported.¹ Therefore, type IV RTA should not be considered as a benign complication of UTI. A comprehensive analysis of clinical outcomes in infants with UTI complicated by type IV RTA remains lacking.

Young age and urinary tract anomalies have been identified as two important factors for the development of type IV RTA in cases

with UTI.^{1–5} Nevertheless, many do not exhibit the known structure anomalies.^{6–8} Recently, a pathogenic mutation on *NR3C2* has been identified in an infant with UTI complicated by type IV RTA, which suggests the possible overlap between primary and secondary type IV RTA.⁹ To date, the genetic sequencing of ENaC and MR has not been fully evaluated in these infants. The aim of this study was to investigate clinical features, genetic background, and outcome in infants with UTI and type IV RTA.

MATERIALS AND METHODS

Subjects and diagnosis of type IV RTA

The study protocol was approved by the Ethics Committee on Human Studies at Chang Gung Memorial Hospital in Taiwan, R.O.C. (IRB 00481A3). Informed consent was obtained from the parents after a detailed description of the study. Infants with UTI admitted to Chang Gung Memorial Hospital were prospectively enrolled from January 2015 to December 2018. Laboratory tests, which includes serum creatinine, Na⁺, K⁺, Cl⁻, HCO₃⁻, osmolality, plasma aldosterone and renin, and urine creatinine, were measured at presentation and followed up during hospitalization. The

¹Division of Pediatric Nephrology, Department of Pediatrics, Chang Gung Memorial Hospital and Chang Gung University, Taoyuan, Taiwan; ²Division of Pediatric Allergy, Asthma, and Rheumatology, Department of Pediatrics, Chang Gung Memorial Hospital and Chang Gung University, Taoyuan, Taiwan; ³Department of Biochemistry, National Defense Medical Center, Taipei, Taiwan; ⁴Division of Nephrology, Department of Pediatrics, MacKay Children's Hospital, Taipei, Taiwan; ⁵Department of Medicine, MacKay Medical College, New Taipei City, Taiwan; ⁶Department of Pediatrics, Taipei Medical University Hospital, Taipei, Taiwan; ⁷Department of Pediatrics, School of Medicine, College of Medicine, Taipei Medical University, Taipei, Taiwan; ⁸Fetal and Neonatal Institute, Division of Neonatology, Children's Hospital Los Angeles, Department of Pediatrics, University of Southern California Keck School of Medicine, Los Angeles, CA, USA; ⁹Genomic Medicine Core Laboratory, Chang Gung Memorial Hospital, Linkou, Taiwan; ¹⁰Department of Pediatrics, Department of Medicine, Tri-Service General Hospital, National Defense Medical Center, Taipei, Taiwan and ¹¹Division of Nephrology, Department of Medicine, Tri-Service General Hospital, National Defense Medical Center, Taipei, Taiwan
Correspondence: Shih-Hua Lin (l521116@ndmctsg.h.edu.tw)

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diagnosis of type IV RTA included the findings of hyperkalemia (serum K^+ >6.0 mmol/L) with impaired renal K^+ excretion, non-anion gap metabolic acidosis (anion gap <12 with $[HCO_3^-]$ <22 mmol/L) with positive urine anion gap (low urine ammonium excretion), relatively low urine pH (<6.0), and normal renal function. To exclude the other causes of type IV RTA such as obstructive nephropathy, diabetic nephropathy, etc., patients who had hyporeninemia or hypoaldosteronism were excluded. As shown in Fig. 1, patients who had (1) impaired renal function

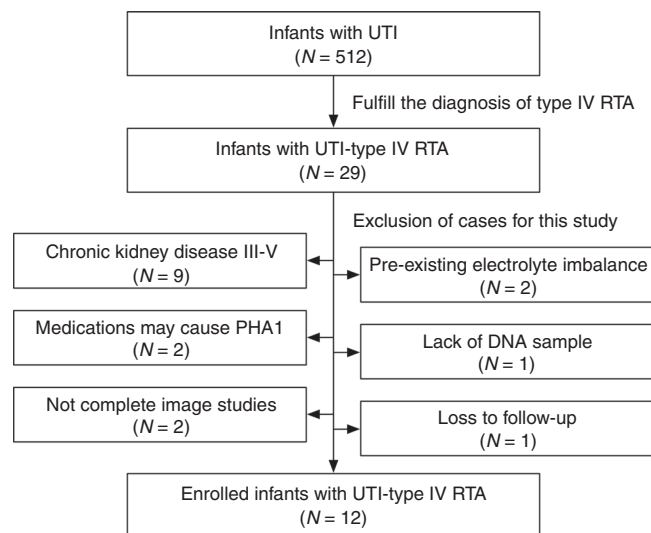


Fig. 1 Patient enrollment.

defined as chronic kidney disease stage III–V, (2) pre-existing electrolyte imbalance, (3) pharmacological history that may cause type IV RTA, (4) lack of DNA for analysis, (5) no imaging studies, and (6) lost to follow-up were excluded. Finally, 12 infants with UTI who exhibited features of type IV RTA were enrolled. UTI was defined by a positive urine culture with at least 100,000 colony-forming units per mL of a single urinary tract pathogen from a specimen obtained by catheter or urine bag or any pathogen from a suprapubic puncture.

Image studies

Technetium-99m-dimercaptosuccinic acid (DMSA) renal scanning and voiding cystourethrography (VCUG) were performed following the diagnosis of UTI. Acute pyelonephritis and renal scar detected by Technetium-99m-DMSA were defined according to a previous study.¹¹ Grading of vesicoureteral reflux (VUR) was based on the criteria previously proposed by International Reflux Study in Children.¹² An experienced nuclear medicine consultant and an experienced radiologist blinded to the patients' clinical condition interpreted the DMSA and VCUG studies, respectively.

Clinical characteristics, treatment, and outcome

Patient demographics and clinical features, treatment, and outcome data were collected in all patients. The clinical features consisted of gender, age at onset, underlying disease, symptoms at presentation. Clinical outcomes, including arrhythmia, unstable hemodynamics, renal scar, and sequelae of vital organs, were determined at follow-up.

Molecular analysis

Genomic DNA was extracted from peripheral leukocyte with DNA isolation kits (QIAamp Blood Kit; Qiagen, Dusseldorf, Germany). We

Table 1. Clinical characteristics of patients with transient type IV renal tubular acidosis.

Patient	1	2	3	4	5	6	7	8	9	10	11	12
Sex	M	M	M	M	M	M	F	M	M	M	F	F
Age (month/s)	2	1.5	1	6	2	8	1	1	1.5	3.5	2	2
Onset of PHA ^a	1	4	3	2	1	4	2	3	5	3	4	3
Symptoms at onset												
Poor feeding	Yes	Yes	Yes	—	—	—	—	—	—	Yes	Yes	Yes
Vomiting	—	Yes	Yes	Yes	Yes	—	—	Yes	Yes	—	—	—
Poor activity	—	—	—	—	—	Yes	Yes	—	—	—	—	—
Fever	Yes	—	—	Yes	Yes	—	Yes	Yes	—	—	—	Yes
Laboratory												
Na ⁺ (133–146), mmol/L	129	125	129	121	109	124	129	129	128	127	129	125
K ⁺ (3.5–5.6), mmol/L	6.5	6.1	6.1	6.2	6.7	7.2	6.1	6.0	6.7	6.5	6.5	6.4
Cl ⁻ (102–112)	104	102	104	98	91	108	102	104	100	101	102	100
HCO ₃ ⁻ (22–26), mmol/L	17	12	15	18	12	10	16	17	14	16	18	16
Serum AG ^c (7–16)	8	11	10	5	6	6	11	11	11	10	9	9
Urine AG (<0)	9	5	35	8	13	33	10	12	9	15	14	7
UpH	5.4	5.0	5.2	5.4	5.1	5.0	5.2	5.3	5.2	5.4	5.0	5.2
Creatinine (0.1–0.5), mg/dL	0.20	0.30	0.25	0.21	0.35	0.34	0.20	0.23	0.24	0.18	0.20	0.30
FENa (%)	1.5	2.0	2.2	2.4	3.0	2.3	2.1	2.3	2.6	3.4	2.8	3.3
TTKG	3.0	4.2	4.8	2.8	5.0	2.0	2.5	3.0	3.0	2.5	3.2	3.5
Aldosterone (38–313), pg/mL	355	420	465	530	755	630	2,101	1,014	980	1,022	846	1,320
Renin (3.0–28.4), pg/mL	806	243	63	94	138	1,433	216	245	1,550	366	244	316
CRP (<0.5), mg/dL	6.0	15.3	4.0	12.1	5.5	7.5	6.0	12.0	5.5	4.5	7.0	21.2
Urine culture	EC	K.P	EC	E. coli	E. coli	K.P	E. coli	P.A	E. coli	E. coli	E. coli	E. coli

AG anion gap (mEq/L), EC *Enterococcus* spp., K.P *Klebsiella pneumoniae*, P.A *Pseudomonas aeruginosa*.

^aDays after onset of urinary tract infection.

sequenced the complete coding regions and intron–exon junctions of the *NR3C2*, *SCNN1A*, *SCNN1B*, and *SCNN1G* genes. The identified mutations were predicted by using eight pathogenicity computation score that was calculated using PolyPhen2, MutationTaster, Mutation Assessor, FATHMM, M-CAP, CADD, DANN, and GERP++. Score 8 from all of pathogenicity computation is a pathogenic mutation.

RESULTS

Demographic and clinical manifestations

The overall incidence of type IV RTA in infants with UTI was around 0.57% (29/512) in this study. There were 9 males and 3 females, who presented at mean age 2.6 ± 1.2 months (ranging from 1 to 8 months). Three-fourths (75%) of patients had disease onset at <3 months of age. As shown in Table 1, poor oral feeding and vomiting were the most common symptoms at presentation. The duration from the onset of UTI to the diagnosis of type IV RTA ranged from 1 to 5 days. Of note, half of them developed type IV RTA after resolution of fever.

Laboratory characteristics

All patients manifested moderate-to-severe hyperkalemia ($K^+ 6.4 \pm 0.2$ mmol/L) with inappropriately low renal K^+ excretion (transtubular potassium gradient 3.3 ± 0.5), non-anion gap metabolic acidosis ($HCO_3^- 15.1 \pm 1.5$ mmol/L), and positive urine anion gap (14.2 ± 5.6 mEq/L). They also had hypovolemic hyponatremia ($Na^+ 125.3 \pm 3.3$ mmol/L) with renal Na^+ wasting (fractional excretion of sodium 2.4 ± 0.2 %) and inappropriately high plasma renin (476.2 ± 295.2 pg/mL) and aldosterone levels (869.8 ± 280.3 pg/mL) (Table 1), excluding the hyporeninemia or hypoaldosteronism causes for type IV RTA. The serum creatinine levels ranged from 0.18 to 0.30 mg/dL. There were no leukocytosis, thrombocytosis, or traumatic blood sampling. The most common pathogens cultured from the urine are *Escherichia coli* followed by *Enterococcus* spp. and *Klebsiella pneumoniae*.

Molecular analysis of corresponding genes

Sanger sequence of *NR3C2*, *SCNN1A*, *SNN1B*, and *SCNN1G* were conducted in all patients. Three genetic variants, c.1-2C>G (slicing site), c.538G>A, and c.1645T>G, on *NR3C2* were detected in four patients (Table 2). The genetic variant c.1645T>G, but not c.1-2C>G or c.538G>A, was predicted to be pathogenic by in silico analysis. This novel and heterozygous thymine-to-guanine substitution at position 1645 (c.1645T>G) in exon 2 resulted in amino acid substitution of serine for alanine (S549A) in patient 9. This specific genetic variant was not found in 200 healthy subjects and was inherited from his father (Fig. 2).

Kidney and urinary tract anomalies

Nine patients had pyelonephritis diagnosed by DMSA renal scan, and two of them had bilateral pyelonephritis (Table 2). Urinary tract anomalies, including VUR, ureteropelvic junction obstruction, ectopic ureter, and unilateral renal agenesis, were identified in seven patients. Of note, VUR was the most common anomaly. Seven patients had both pyelonephritis and urinary tract anomalies. Five patients did not have urinary tract anomalies and one of them (patient 9) harbored heterozygous mutation on *NR3C2* gene.

Complications, treatment, and follow-up outcome

Seven patients exhibited arrhythmia with predominant tented T wave and two of them developed ventricular tachycardia (Table 2) requiring intravenous calcium and insulin administration. Intravenous antibiotics, salt and fluid hydration, and furosemide were administered to all patients for the treatment of the underlying UTI, hyponatremia, and hyperkalemia, respectively. Four patients received $NaHCO_3$ supplement for severe metabolic acidosis. All

Table 2. Images, genetic variants, complications, treatment, and outcome of UTI infants with transient type IV renal tubular acidosis.

Patient no.	APN	UTA	Genetic variant	PS	Complications	Treatment	Outcome
1	Left	Negative	Negative	—	Tent T wave	Antibiotics, intravenous hydration, furosemide	Recovery
2	Bilateral	Right VUR, grade III, Left VUR, grade V	Negative	—	—	Antibiotics, intravenous hydration, furosemide sodium bicarbonate	Recovery, renal scar
3	Negative	Negative	<i>NR3C2</i> , c.1-2C>G, homo	Not applicable (benign)	Tent T wave	Antibiotics, intravenous hydration, furosemide	Recovery
4	Left	Negative	Negative	—	—	Antibiotics, intravenous hydration, furosemide	Recovery
5	Negative	Negative	<i>NR3C2</i> , c.538G>A (V180I), homo	3 (Benign)	Ventricular tachycardia	Antibiotics, intravenous hydration, furosemide, sodium bicarbonate, calcium polystyrene sulfonate, calcium gluconate, insulin	Recovery
6	Left	Left UPJO	<i>NR3C2</i> , c.538G>A (V180I), hetero	3 (Benign)	Ventricular fibrillation	Antibiotics, intravenous hydration, furosemide, sodium bicarbonate, calcium polystyrene sulfonate, calcium gluconate, insulin	Recovery, prolonged hospital stay
7	Right	Right ectopic ureter	Negative	—	—	Antibiotics, intravenous hydration, furosemide	Recovery
8	Left	Left renal duplication with ectopic ureter	Negative	—	—	Antibiotics, intravenous hydration, furosemide	Recovery, renal scar
9	Negative	Negative	<i>NR3C2</i> , c.1645T>G (S549A), hetero.	8 (Pathogenic)	Tent T wave	Antibiotics, intravenous hydration, furosemide sodium bicarbonate	Recovery
10	Bilateral	Bilateral VUR, grade V	Negative	—	Tent T wave	Antibiotics, intravenous hydration, furosemide	Recovery, renal scar
11	Left	Right renal agenesis, left VUR, grade V	Negative	—	Tent T wave	Antibiotics, intravenous hydration, furosemide	Recovery, renal scar
12	Right	Left renal agenesis, right VUR, grade IV	Negative	—	—	Antibiotics, intravenous hydration, furosemide	Recovery

APN acute pyelonephritis, UTA urinary tract anomaly, PS pathogenicity score, VUR vesicoureteral reflux, UPJO ureteropelvic junction obstruction, homo homozygous mutation, hetero heterozygous mutation.

patients achieved full recovery with resolution of type IV RTA within the first week of hospitalization. Prolonged hospital stay was noted in one patient (patient 6) due to complications from seizure and aspiration pneumonia. During their 6-month follow-up, four patients had renal scar detected by DMSA renal scan. None of the patients had recurrence of type IV RTA. Patient 9 who harbored genetic defect on *NR3C2* had a normal high serum K^+ (5.2 mmol/L) at the age of 1 year during follow-up.

DISCUSSION

In this study, we found that nearly half of the patients with UTI-associated type IV RTA did not have urinary tract anomalies. Over half of the patients developed arrhythmia, and two patients had ventricular tachycardia. All patients had full resolution of type IV RTA within 1 week after treatment and did not have recurrence during follow-up. In the investigation of genetic background of these infants, we identified one novel missense mutation at the phosphorylation site of *NR3C2*.

The cardinal presentations for UTI in this cohort, such as vomiting and poor oral feeding, were non-specific, thus increasing the difficulty of diagnosing type IV RTA correctly and promptly without measurement of blood electrolytes and acid–base status. Previous studies have pointed out that young age is a necessary factor for the development of UTI-associated type IV RTA. All our patients were aged <8 months. The relatively immature renal

tubules in infants have limited compensatory capacity, resulting from both/either physiological partial aldosterone resistance and/or low renal MR expression associated with UTIs.^{6,10,13–15} Similar to the previous reports, we found that VUR was the leading anomaly of urinary tract in our patients with UTI-associated type IV RTA.⁶ Urinary tract anomaly is also shown to be associated with the decreased expression of MR receptor and increased MR resistance.^{16,17} However, the percentage of urinary tract anomalies in this study was approximately 60% (7/12), which was less than that previously reported (90%).⁶ Furthermore, recent studies reveal that bacterial endotoxins and inflammatory cytokines, such as transforming growth factor- β , interferon- α , interleukin (IL)-1, and IL-6, were involved in the pathogenesis of transient type IV RTA in infants with UTI.^{16–21} Accordingly, both bacterial and host factors may predispose the development of UTI-associated type IV RTA.

Three genetic variants, c.1-2C>G (slicing site), c.538G>A, and c.1645T>G, on *NR3C2* encoding the MR were identified in four patients. The c.1-2C>G and c.538G>A genetic variants were predicted as benign. However, c.538G>A (V180I) has been reported to cause defective MR function in in-vitro functional study.^{22,23} It remains to be determined whether UTI alone or a combination of genetic defect and UTI was responsible for the development of transient type IV RTA in infants with UTI carrying this mutation.^{9,20} A novel and heterozygous mutation, S549A, was identified at a phosphorylation site for MR in patient 9, who had no urinary tract anomalies. The mutation was found in the father, which is consistent with the diagnosis of autosomal-dominant pseudohypoaldosteronism type I affecting the MR. Based on the absence of this variant in 200 healthy subjects and predictions from different mutation softwares, we believe this mutation to be pathogenic. A functional study is still warranted to confirm the pathogenicity of the phosphorylation site mutant S549A on MR. In summary, we proposed that bacterial factor (endotoxins) and/or host factors (inflammatory cytokines, urinary tract anomalies, genetic background) can contribute to the development of UTI-associated type IV RTA (Fig. 3).

Although the outcome of UTI-associated type IV RTA is usually favorable, some patients may present with life-threatening hyperkalemia and severe metabolic acidosis.^{10,24–26} Half of our patients developed hyperkalemia-related arrhythmia and two of them presented with ventricular arrhythmia requiring emergent management. Early recognition and timely treatment to normalize electrolyte status is crucial to avoid the complications of transient type IV RTA in infants with UTI. Furthermore, we emphasize that any form of stress, especially infections in infancy not limited to UTI, could trigger the clinical manifestations of type IV RTA in infants harboring the genetic defect in *NR3C2*. Close monitoring of

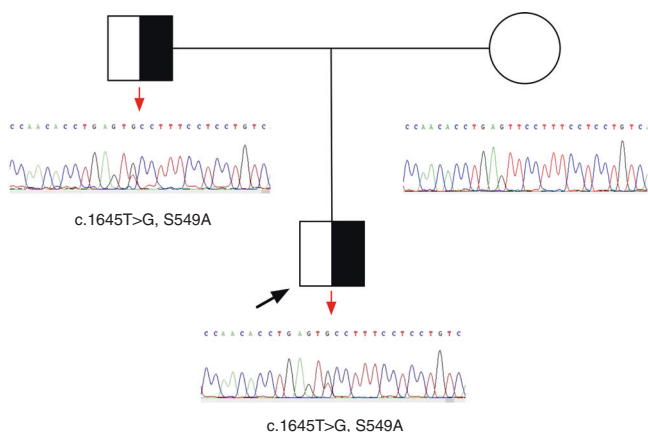


Fig. 2 Direct sequencing of *NR3C2* in patient 9 and their parents. Arrow indicates the mutant base and left and right upper rows indicate the father and mother, respectively. Male and female indicated by squares and circles, respectively. Filled symbols represent patient 9.

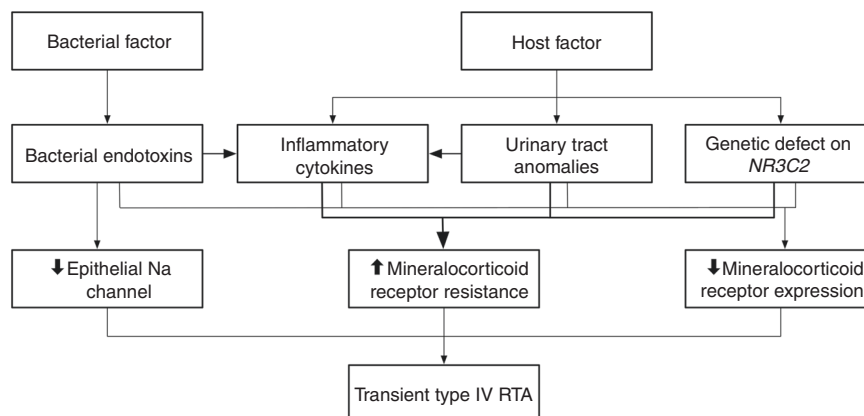


Fig. 3 Proposed mechanism of transient type IV renal tubular acidosis in infants with urinary tract infection.

recurrence of type IV RTA should be considered in these infants during their acute illness.

CONCLUSION

Although UTI-associated type IV RTA is usually transient with a benign outcome, rapid-onset and life-threatening complications such as hyperkalemia-related arrhythmia and severe metabolic acidosis can occur and require emergent therapy. Aside from young age and urinary tract anomalies, underlying genetic defects on *NR3C2* could be a contributing factor for UTI-associated type IV RTA without other identifiable risk factors.

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AUTHOR CONTRIBUTIONS

M.-H.T. and S.-H.L. interpreted the data analyses and reviewed and revised the manuscript. J.-L.H., S.-M.H., and J.-D.T. carried out the data analysis, assisted with interpretation of the data analyses, and reviewed and revised the manuscript for important intellectual content. T.-W.W. assisted with acquisition of the data, interpreted the data analyses, and reviewed and revised the manuscript for important intellectual content. W.-L.F. and J.-J.D. conceptualized and designed the study, assisted with acquisition of data, and provided analysis and interpretation of the data. All authors approved the final manuscript as submitted and agree to be accountable for all aspects of the work.

ADDITIONAL INFORMATION

Competing interests: The authors declare no competing interests.

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