

Propuesta de registro de pacientes España

Dra. V. Seidel – Hospital Universitario Gregorio Marañón

Dra. A. Cueto – Hospital Universitario Vall d'Hebron

Dr. F. Santos – Hospital Universitario La Paz

Base de datos conjunta HULP - HGUG

[illegible]

Kabuki syndrome: a Spanish case series

Verónica Seidel¹, Sixto García-Miñaur², Elena Vallespin², Víctor Martínez-Glez², Gema Gordo³, Andrea Naughton³, Geraldine Malone³, Pablo Lapunzina³, Fernando Santos-Simarro³

¹Clinical Genetics/ Pediatrics Service, Hospital General Universitario Gregorio Marañón, Madrid, Spain

²INGEMM-CIBERER, Hospital Universitario La Paz, Madrid, Spain

³Manchester Centre for Genomic Medicine, Manchester, UK

Introduction:

Kabuki syndrome is a clinically heterogeneous multiple congenital malformation syndrome with distinctive facial features and developmental delay. Two histone methyltransferase genes have been identified as causes, *KMT2D* and *KDM6A* (X chromosome).

Material and methods:

We present 20 cases diagnosed with Kabuki syndrome, seen at two large hospitals in Madrid, Spain.

ID	Sex	Age (years)	Height (percentile)	DOI/ ID	Heart	Renal	Gastro-Intestinal	Neurologic	Skeletal	Palate	Teeth	Eyes	Hearing	Endocrine / Immunology	Other
LP1	M	12	p10	+	VSD	N	N	N	N	N	hypodontia	Strabismus	bilateral SNHL	N	
GM1	F	3	SS	+	Acromioclavicular, VSD, ASD	N	Feeding difficulties	Hypotonia, microcephaly	hirsutism, hyperlacty	VPI	N	retinal coloboma (unilateral)	transient hypothyroidism	N	premature telarche
LP2	F	14	p10	+	Ao, MV stenosis	N	Volvulus; short bowel syndrome	stroke	N	N	hypodontia	N	N	N	genetic mosaicism
LP3	F	10	FTT	++	CHD	N	N	N	N	N	N	N	N	N	Decreased
GM2	F	8	SS	++	N	unilateral agenesis; ectopic dysplastic	GER	Hypotonia	-	cleft	hypoplastic	N	Transmissive HL	Low IgA	
LP4	M	6	+	N	N	N	N	N	N	N	N	N	N	N	
GM3	F	3	p25	++	VSD	N	-	Hypotonia, microcephaly	Trigonocephaly; hip dysplasia	N	N	NI duct obstruction	N	N	Abbreviations: FTT: Failure to thrive; DOI: Developmental delay; ID: Intellectual disability; R: Normal; S: Short stature; PFO: Patent Foramen Ovale; VPI: Velopharyngeal insufficiency; SNHL: Sensorineural Hearing Loss
GM4	F	4	p25	+	PFO	N	-	Hypotonia, microcephaly	hip dysplasia	VPI	N	N	N	N	
LP5	M	10	SS	+	MV dysfunction	N	N	N	N	N	N	N	N	N	
LP6	M	16	SS	+	N	N	N	N	N	N	N	N	N	N	
GM5	M	3	SS	+	VSD	Horseshoe	-	Hypotonia	-	N	N	Strabismus	N	neonatal hypoglycemia	stereotypes
LP7	M	10	p3	+	N	unilateral preauricular stenosis	feeding difficulties	N	N	cleft	N	N	N	N	hypoplasia
LP8	F	7	++	VSD	ectopic kidney	PEG	N	Hypotonia, EEG anomalies	Vth finger dystrophy	cleft	N	N	N	N	
GM6	M	10	p2-10	+	PFO	duplex renal pelvis (unilateral)	GER, PEG	N	N	high arched	Hypodontia	iris coloboma; corneal clouding (unilateral)	Transmissive HL	neonatal hypoglycemia	cryptorchidism
GM7	F	7	SS	++	Ao coarctation	N	feeding difficulties	Hypotonia, EEG anomalies	-	high arched	Hypodontia	Strabismus	N	N	premature telarche, umbilical hernia, behavioural problems
LP9	F	5	+	+	N	unilat R agnathia	ant placed anus	-	hip dysplasia	N	NA	Myopia	N	N	
LP10	F	5	SS	+	N	N	N	N	hip dysplasia	N	N	bulb's eye	N	N	
GM8	M	5m	FTT	+	Tetralogy of Fallot	N	feeding difficulties	N	clavicle hypoplasia	cleft	NA	microphthalmia; retinal coloboma	presenile pitted	N	
GM9	F	14	p25	+	N	N	-	Macrocephaly	-	N	N	Strabismus	presenile pitted	N	AI steering cholangitis
LP11	F	9	SS	+	N	N	feeding difficulties	N	N	N	hypodontia	Autism	N	N	ADHD

Table 1: Clinical data

ID	Gene	Variant	Protein	Change	Other
LP1	KMT2D	c.12764G>A	p.S427N	missense	
GM1	KMT2D	c.12764G>A	p.S427N	missense	
LP2	KMT2D	c.12764G>A	p.S427N	missense	
LP3	KMT2D	c.12764G>A	p.S427N	missense	
GM2	KMT2D	c.12764G>A	p.S427N	missense	
LP4	KMT2D	c.12764G>A	p.S427N	missense	
GM3	KMT2D	c.12764G>A	p.S427N	missense	
LP5	KMT2D	c.12764G>A	p.S427N	missense	
LP6	KMT2D	c.12764G>A	p.S427N	missense	
GM5	KMT2D	c.12764G>A	p.S427N	missense	
LP7	KMT2D	c.12764G>A	p.S427N	missense	
LP8	KMT2D	c.12764G>A	p.S427N	missense	
GM6	KMT2D	c.12764G>A	p.S427N	missense	
GM7	KMT2D	c.12764G>A	p.S427N	missense	
LP9	KMT2D	c.12764G>A	p.S427N	missense	
LP10	KMT2D	c.12764G>A	p.S427N	missense	
GM8	KMT2D	c.12764G>A	p.S427N	missense	
GM9	KMT2D	c.12764G>A	p.S427N	missense	
LP11	KMT2D	c.12764G>A	p.S427N	missense	
GM10	KMT2D	c.12764G>A	p.S427N	missense	
GM11	KMT2D	c.12764G>A	p.S427N	missense	
GM12	KMT2D	c.12764G>A	p.S427N	missense	
GM13	KMT2D	c.12764G>A	p.S427N	missense	
GM14	KMT2D	c.12764G>A	p.S427N	missense	
GM15	KMT2D	c.12764G>A	p.S427N	missense	
GM16	KMT2D	c.12764G>A	p.S427N	missense	
GM17	KMT2D	c.12764G>A	p.S427N	missense	
GM18	KMT2D	c.12764G>A	p.S427N	missense	
GM19	KMT2D	c.12764G>A	p.S427N	missense	
GM20	KMT2D	c.12764G>A	p.S427N	missense	
GM21	KMT2D	c.12764G>A	p.S427N	missense	
GM22	KMT2D	c.12764G>A	p.S427N	missense	
GM23	KMT2D	c.12764G>A	p.S427N	missense	
GM24	KMT2D	c.12764G>A	p.S427N	missense	
GM25	KMT2D	c.12764G>A	p.S427N	missense	
GM26	KMT2D	c.12764G>A	p.S427N	missense	
GM27	KMT2D	c.12764G>A	p.S427N	missense	
GM28	KMT2D	c.12764G>A	p.S427N	missense	
GM29	KMT2D	c.12764G>A	p.S427N	missense	
GM30	KMT2D	c.12764G>A	p.S427N	missense	
GM31	KMT2D	c.12764G>A	p.S427N	missense	
GM32	KMT2D	c.12764G>A	p.S427N	missense	
GM33	KMT2D	c.12764G>A	p.S427N	missense	
GM34	KMT2D	c.12764G>A	p.S427N	missense	
GM35	KMT2D	c.12764G>A	p.S427N	missense	
GM36	KMT2D	c.12764G>A	p.S427N	missense	
GM37	KMT2D	c.12764G>A	p.S427N	missense	
GM38	KMT2D	c.12764G>A	p.S427N	missense	
GM39	KMT2D	c.12764G>A	p.S427N	missense	
GM40	KMT2D	c.12764G>A	p.S427N	missense	
GM41	KMT2D	c.12764G>A	p.S427N	missense	
GM42	KMT2D	c.12764G>A	p.S427N	missense	
GM43	KMT2D	c.12764G>A	p.S427N	missense	
GM44	KMT2D	c.12764G>A	p.S427N	missense	
GM45	KMT2D	c.12764G>A	p.S427N	missense	
GM46	KMT2D	c.12764G>A	p.S427N	missense	
GM47	KMT2D	c.12764G>A	p.S427N	missense	
GM48	KMT2D	c.12764G>A	p.S427N	missense	
GM49	KMT2D	c.12764G>A	p.S427N	missense	
GM50	KMT2D	c.12764G>A	p.S427N	missense	
GM51	KMT2D	c.12764G>A	p.S427N	missense	
GM52	KMT2D	c.12764G>A	p.S427N	missense	
GM53	KMT2D	c.12764G>A	p.S427N	missense	
GM54	KMT2D	c.12764G>A	p.S427N	missense	
GM55	KMT2D	c.12764G>A	p.S427N	missense	
GM56	KMT2D	c.12764G>A	p.S427N	missense	
GM57	KMT2D	c.12764G>A	p.S427N	missense	
GM58	KMT2D	c.12764G>A	p.S427N	missense	
GM59	KMT2D	c.12764G>A	p.S427N	missense	
GM60	KMT2D	c.12764G>A	p.S427N	missense	
GM61	KMT2D	c.12764G>A	p.S427N	missense	
GM62	KMT2D	c.12764G>A	p.S427N	missense	
GM63	KMT2D	c.12764G>A	p.S427N	missense	
GM64	KMT2D	c.12764G>A	p.S427N	missense	
GM65	KMT2D	c.12764G>A	p.S427N	missense	
GM66	KMT2D	c.12764G>A	p.S427N	missense	
GM67	KMT2D	c.12764G>A	p.S427N	missense	
GM68	KMT2D	c.12764G>A	p.S427N	missense	
GM69	KMT2D	c.12764G>A	p.S427N	missense	
GM70	KMT2D	c.12764G>A	p.S427N	missense	
GM71	KMT2D	c.12764G>A	p.S427N	missense	
GM72	KMT2D	c.12764G>A	p.S427N	missense	
GM73	KMT2D	c.12764G>A	p.S427N	missense	
GM74	KMT2D	c.12764G>A	p.S427N	missense	
GM75	KMT2D	c.12764G>A	p.S427N	missense	
GM76	KMT2D	c.12764G>A	p.S427N	missense	
GM77	KMT2D	c.12764G>A	p.S427N	missense	
GM78	KMT2D	c.12764G>A	p.S427N	missense	
GM79	KMT2D	c.12764G>A	p.S427N	missense	
GM80	KMT2D	c.12764G>A	p.S427N	missense	
GM81	KMT2D	c.12764G>A	p.S427N	missense	
GM82	KMT2D	c.12764G>A	p.S427N	missense	
GM83	KMT2D	c.12764G>A	p.S427N	missense	
GM84	KMT2D	c.12764G>A	p.S427N	missense	
GM85	KMT2D	c.12764G>A	p.S427N	missense	
GM86	KMT2D	c.12764G>A	p.S427N	missense	
GM87	KMT2D	c.12764G>A	p.S427N	missense	
GM88	KMT2D	c.12764G>A	p.S427N	missense	
GM89	KMT2D	c.12764G>A	p.S427N	missense	
GM90	KMT2D	c.12764G>A	p.S427N	missense	
GM91	KMT2D	c.12764G>A	p.S427N	missense	
GM92	KMT2D	c.12764G>A	p.S427N	missense	
GM93	KMT2D	c.12764G>A	p.S427N	missense	
GM94	KMT2D	c.12764G>A	p.S427N	missense	
GM95	KMT2D	c.12764G>A	p.S427N	missense	
GM96	KMT2D	c.12764G>A	p.S427N	missense	
GM97	KMT2D	c.12764G>A	p.S427N	missense	
GM98	KMT2D	c.12764G>A	p.S427N	missense	
GM99	KMT2D	c.12764G>A	p.S427N	missense	
GM100	KMT2D	c.12764G>A	p.S427N	missense	

Table 2: Genetic testing results: 18 confirmations of 19 tested

Results:

Age at diagnosis ranged from 5 months to 16 years. Female:male ratio was 12:8.

Eighteen cases had a molecular confirmation (17 *KMT2D* and 1 *KDM6A* mutation), one test is ongoing and one result was negative for both genes. Of the 17 *KMT2D* mutations 8 (47%) were truncating (either nonsense or frameshift producing a downstream stop codon).

One case was found to carry 3 variants in *KMT2D*: 1 *de novo* nonsense (clearly pathogenic) and 2 missense variants inherited from an unaffected parent.

The combination of congenital anomalies, developmental delay and characteristic facial features was variable but the distinctive eyebrows and long palpebral fissures reminiscent of the Kabuki make-up of Japanese theatre were present in all.

Additional features found in our series, rarely or not previously described as part of the syndrome, were trigonocephaly in one case, unilateral corneal opacity and iris coloboma in another, and autoimmune cholangitis in the girl with *KDM6A* mutation. One patient died at 10 years from complications of her heart anomalies. Interestingly, one case with a milder phenotype and less characteristic facial dysmorphism was found to be mosaic for a *KMT2D* frameshift mutation.

Conclusions:

This is the first series of cases with Kabuki syndrome in Spain. Additional cranial, ophthalmological and hepatic features are shown. A case with mosaicism for Kabuki syndrome is presented.

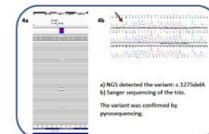


Figure 4: Mosaicism for a *KMT2D* mutation in blood.

REDCap – Programa disponible en IISGM



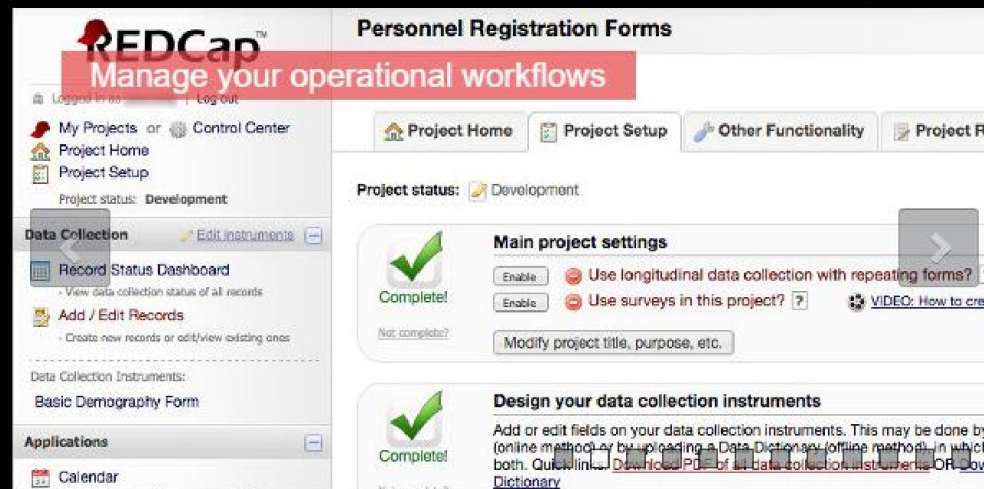
Institutions **2553** Countries **116** Projects **464k** Users **608k** Articles **4110**

[ABOUT](#)

[PARTNERS](#)

[RESOURCES](#)

[SOFTWARE](#)



REDCap is a secure web application for building and managing online surveys and databases. While REDCap can be used to collect virtually any type of data (including 21 CFR Part 11, FISMA, and HIPAA-compliant environments), it is specifically geared to support online or offline data capture for research studies and operations. The REDCap Consortium, a vast support network of collaborators, is composed of thousands of active institutional partners in over one hundred countries who utilize and support REDCap in various ways.