### DELETION OF ERF AND CIC CAUSES ABNORMAL SKULL MORPHOLOGY AND GLOBAL **DEVELOPMENTAL DELAY**

Ram Singh<sup>1,2\*</sup>, Ana S. A. Cohen<sup>1,2,a\*</sup>, Cathryn Poulton<sup>3</sup>, Tina Duelund Hjortshøj<sup>4</sup>, Moe Akahira-Azuma<sup>1,b</sup>, Geetu Mendiratta<sup>1,2</sup>, Wahab A. Khan<sup>1,2,c</sup>, Dimitar N. Azmanov<sup>5,6</sup>, Karen J. Woodward<sup>5,6</sup>, Maria Kirchhoff<sup>4</sup>, Lisong Shi<sup>1,2</sup>, Lisa Edelmann<sup>1,2</sup>, Gareth Baynam<sup>7,8,9</sup>, Stuart A. Scott<sup>1,2,d†</sup>, and Ethylin Wang Jabs<sup>1†</sup>

#### **Current affiliations:**

#### Running Title: ERF and CIC deletion and abnormal skull morphology

#### †CORRESPONDENCE TO:

Stuart A. Scott, PhD Ethylin Wang Jabs, MD

Adjunct Associate Professor Professor

Department of Genetics and Genomic Sciences Department of Genetics and Genomic Sciences

Icahn School of Medicine at Mount Sinai Icahn School of Medicine at Mount Sinai

New York, NY 10029 New York, NY 10029 Tel. 212-241-3780 Tel. 212-241-3504 Fax. 212-241-0139 Fax. 212-426-9065

E-mail: sascott@stanford.edu E-mail: ethylin.jabs@mssm.edu

#### **SUPPLEMENTAL MATERIAL:**

**Supplemental Table S1:** Page 2 **Supplemental Table S2:** Page 5

<sup>&</sup>lt;sup>1</sup> Department of Genetics and Genomic Sciences, Icahn School of Medicine at Mount Sinai, New York, NY.

<sup>&</sup>lt;sup>2</sup> Sema4, Stamford, CT.

<sup>&</sup>lt;sup>3</sup> Genetic Service of Western Australia, King Edward Memorial Hospital, Perth.

<sup>&</sup>lt;sup>4</sup> Department of Clinical Genetics, Copenhagen University Hospital, Rigshospitalet, Denmark.

<sup>&</sup>lt;sup>5</sup> Department of Diagnostic Genomics, PathWest Laboratory Medicine, QEII Medical Centre, Nedlands 6009, Western Australia.

<sup>&</sup>lt;sup>6</sup> Pathology and Laboratory Medicine, Medical School, Faculty of Health and Medical Sciences, The University of Western Australia, Crawley 6009, Western Australia.

<sup>&</sup>lt;sup>7</sup> Western Australian Register of Developmental Anomalies and Genetic Services of Western Australia, King Edward Memorial Hospital, Perth, Western Australia.

<sup>&</sup>lt;sup>8</sup> Faculty of Health and Medical Sciences, Division of Paediatrics and Telethon Kids Institute, University of Western Australia, Perth, Western Australia.

<sup>&</sup>lt;sup>9</sup> Faculty of Medicine, University of Notre Dame, Australia, Perth, Western Australia.

<sup>\*</sup> These authors contributed equally.

<sup>&</sup>lt;sup>a</sup> Department of Pathology and Laboratory Medicine, Children's Mercy - Kansas City, Kansas City, MO; Center for Pediatric Genomic Medicine, Children's Mercy - Kansas City, Kansas City, MO; University of Missouri-Kansas City School of Medicine, Kansas City, MO.

<sup>&</sup>lt;sup>b</sup> National Center for Global Health and Medicine, Tokyo, Japan.

<sup>&</sup>lt;sup>c</sup> Department of Pathology and Laboratory Medicine, Dartmouth-Hitchcock Medical Center, Lebanon, NH.

<sup>&</sup>lt;sup>d</sup> Department of Pathology, Stanford University, Stanford, CA; Stanford Medicine Clinical Genomics Laboratory, Stanford Health Care, Palo Alto, CA.

# Supplemental Table S1. Previously reported *ERF* variants implicated in variable craniosynostosis.

Nucleotide change	Protein change	Craniosynostosis type / Phenotype	Population frequency (gnomAD)	Reference
c.1A>T	p.Met1?	P + OHT, mild exorbitism, malar hypoplasia, broad thumbs/first toes, early CM (1 pt)	absent	Glass (2019) PMID: 30758909
c.1A>G	p.Met1?	MS + proptosis, hypertelorism, malar hypoplasia, depressed nasal bridge, ectopic posterior pituitary, CM (1 pt)	absent	Chaudhry (2015) PMID: 26097063
c.3G>A	p.Met1?	MS (4 pts, 1 family)	absent	Twigg (2013) PMID: 23354439
c.21A>T	p.(Thr7=)	SS (3 pts, 1 family)	absent	Twigg (2013) PMID: 23354439
c.23-2A>G	-	BI-ME + proptosis, hypertelorism, depressed nasal bridge, speech delay (1 pt)	absent	Chaudhry (2015) PMID: 26097063
c.90_99del	p.Trp30Cysfs*44	SS + developmental delay (1 pt)	absent	Lee (2018) PMID: 29215649
c.157G>A	p.Gly53Arg	SS + OHT, exorbitism, malar hypoplasia, low-set ears, open mouth appearance (1 pt)	absent	Glass (2019) PMID: 30758909
c.161A>G	p.Glu54Gly	UL + medial epicanthic folds, short upturned nose (1 pt); CC-SS, OHT, exorbitism (1 pt, same family)	absent	Glass (2019) PMID: 30758909
c.194G>A	p.Arg65Gln	SS (2 pts, 1 family); BI (1 pt)	absent	Twigg (2013) PMID: 23354439; Lee (2018) PMID: 29215649
c.202G>C	p.Gly68Arg	SS-UL + hydrocephalus, macrocephaly, dysplastic auricles (1 pt); ME + dysplastic auricles (1 pt, same family); SS (1 pt, same family)	absent	Glass (2019) PMID: 30758909
c.247C>T	p.Arg83Trp	CC variable (10 pts, 1 family, with reduced penetrance in 4 pts); SB + OHT, exorbitism (1 pt, different family)	absent	Glass (2019) PMID: 30758909
c.248G>A	p.Arg83GIn	CC variable + OHT, exorbitism, malar hypoplasia, medial epicanthic folds, CM (2 pts, 1 family)	1/250822 =0.0004%	Glass (2019) PMID: 30758909
c.256C>T	p.Arg86Cys	MS (1 pt)	absent	Twigg (2013) PMID: 23354439
c.257G>A	p.Arg86His	SB + hypertelorism, vertical nasal displacement (2 pts, 1 family)	1/250504 =0.0004%	Yoon (2019) PMID: 31754721

204 C: T	. A 404T	B. OUT	4/254472	Class (2040) BN41B
c.301C>T	p.Arg101Trp	P + OHT, exorbitism, malar hypoplasia,	1/251472	Glass (2019) PMID:
		early CM (3 pts, 1 family, with reduced	=0.0004%	30758909
		penetrance in other 2 pts)		
c.311A>G	p.Tyr104Cys	Multiple congenital anomalies	1/251472	Retterer (2016)
			=0.0004%	PMID: 26633542
c.373+1G>C	-	Crouzon syndrome	absent	Lee (2018) PMID: 29215649
c.383T>G	p.Val128Gly	UC + low anterior hairline, asymmetric	7/241336	Topa (2020) PMID:
		face, flat malar regions, low-set and	=0.0029%	31837199
		posteriorly rotated right ear, normal		
		development (1 pt)		
c.547C>T	p.Arg183*	MS (4 pts, 1 family); SS + OHT, mild	absent	Twigg (2013)
		exorbitism, malar hypoplasia (1 pt,		PMID: 23354439;
		different family); SB + OHT, mild		Glass (2019) PMID:
		exorbitism, malar hypoplasia, short		30758909
		lateral metatarsals (3 pts, 1 family,		
		with reduced penetrance in 1 pt)		
c.652C>T	p.Arg218*	SB + OHT, mild exorbitism	0%	Glass (2019) PMID:
	h9==0	(2 pts, 1 family, reduced penetrance in		30758909
		other pt)		30730303
c.886G>A	p.Gly296Ser	SS, hypertelorism, vertical nasal	5/245366	Yoon (2019) PMID:
C.880G2A	p.diy2505ei	displacement (1pt)	=0.002%;	31754721
		displacement (1pt)	1	31/34/21
			homozygo	
c.891_892del	p.Gly299Argfs*9	P (1 pt); M (4 pts, 2 families); B-UL +	us absent	Twigg (2013)
C.091_092UEI	p.diy233Aigis 3		absent	PMID: 23354439;
		OHT, exorbitism, malar hypoplasia,		-
		early CM (1pt)		Glass (2019) PMID:
. 005 402744	. A 220C C . * E	CC has a talled as for stall have in	-1	30758909
c.985_1027del	p.Arg329Serfs*5	SS, hypertelorism, frontal bossing,	absent	Yoon (2019) PMID:
	4	sunken midface, short stature (2 pts, 1		31754721
		family)		
c.1103C>T	p.Ser368Phe	UC (1 pt who also carries POR	absent	Clarke (2018)
		p.Arg453His)		PMID: 29168297
c.1201_1202del	p.Lys401Glufs*10	P (1 pt); P + OHT, malar hypoplasia,	1/236032	Twigg (2013)
		CM, frontal bossing, long philtrum,	=0.0004%	PMID: 23354439;
		high arched palate, low-set ears,		Glass (2019) PMID:
		inverted nipples, bilateral 5th finger		30758909; Korberg
		clinodactyly, broad DP of		(2020) PMID:
		halluces, sacral dimple (1 pt, different		32370745
		family); UC + long philtrum, short up-		
		turned nose, joint hypermobility (2		
		pts, 1 family, with reduced penetrance		
		in other pt); BLSS and SS.		
c.1214G>A	p.Ser405Asn	ME (1 pt)	absent	Yoon (2019) PMID:
				31754721

c.1270C>T	p.Gln424*	P (3 pts, 1 family); SB + OHT (1 pt, different family)	absent	Twigg (2013) PMID: 23354439; Glass (2019) PMID: 30758909
c.1300G>A	p.Glu434Lys	UC + high forehead, tendency toward turricephaly preoperative, prominent scalp veins, shallow orbits, mild right proptosis, flat malar regions, small mouth, hypotelorism, asymmetric facial midline, right epicanthus, flat malar regions, midface hypoplasia, low-set and anteriorly rotated ears and cup-shaped right helix, psychomotor delay, strabismus, tapering fingers, mild skin syndactyly of fingers, short proximal phalanges 2-5, bilateral syndactyly 2nd-3rd toes, overlapping 2nd and 4th toes right foot, clinodactyly 3rd toe, diarrhea, gastroesophageal reflux, food allergy, growth delay, short stature, sparse hair (1 pt)	3/245440 =0.0012%	Topa (2020) PMID: 31837199
c.1390_1391dup	p.Lys465Leufs*6 7	P + OHT, exorbitism, malar hypoplasia, left proximal radio-ulnar synostosis, C5/6 vertebral fusion, broad thumbs (1pt); SS + OHT, exorbitism (1 pt, same family)	absent	Glass (2019) PMID: 30758909
c.1512del	p.Phe504Leufs*2 7	CC (1 pt)	absent	Twigg (2013) PMID: 23354439
c.1593_1594insC AGA	p.Ser532Glnfs*3	ME (3 pts, 1 fam)	absent	Timberlake (2017) PMID: 28808027

BI: bicoronal; BLSS: bilateral lambdoid and sagittal synostosis; CC: complex craniosynostosis; CM: Chiari-1 malformation; ME: metopic; MS: multiple-suture synostosis; OHT: orbital hypertelorism; P: pansynostosis; pt(s): patient(s); SB: Sagittal bilambdoid; SS: sagittal synostosis; UC: unicoronal; UL: unilambdoid. All variants described for *ERF* (NM\_006494.3) in HGMD® Professional 2021.1 with the exception of c.266A>G, c.288G>T and c.1460G>A associated with distinct phenotypes; variants frequencies from https://gnomad.broadinstitute.org/.

## Supplemental Table S2. Previously reported *CIC* sequence variants implicated in neurodevelopmental phenotypes (mostly *de novo*).

Nucleotide change	Protein change	Phenotype	Population frequency (gnomAD)	Reference
c.457C>T	p.Leu153Phe	ASD (1/2,508 pts)	absent	lossifov (2014) PMID: 25363768
c.673C>T	p.Gln225*	ASD + CLP (1/2,620 pts)	absent	Yuen (2017) PMID: 28263302
c.680A>G	p.Asn227Ser	nsID (1/236 pts)	absent	Athanasakis (2014) PMID: 24307393
c.791A>G	p.His264Arg	Schizophrenia (1/623 pts)	absent	Fromer (2014) PMID: 24463507
c.820C>T	p.Arg274*	ASD (1/2,508 pts)	1/246948 =0.0004%	lossifov (2014) PMID: 25363768; PMID: 28191890 uses same data set
c.1057C>T	p.Arg353*	DD, ID, ASD, seizures, macrocephaly, abnormal MRI, mild telangiectasia (1 pt)	absent	Lu (2017) PMID: 28288114
c.1474C>T	p.Arg492Trp	nsID (1/10 pts)	1/250214 =0.0004%	Vissers (2010) PMID: 21076407
c.1801_1808 dup	p.Glu604Argfs*127	DD, ID, seizures (2 siblings) + pulmonary stenosis (sib A) + heart murmur, mild telangiectasia (sib B)	absent	Lu (2017) PMID: 28288114
c.1927G>C	p.Gly643Arg	ASD, DD (3 affected siblings, variable severity)	absent	Mahfouz (2020) PMID: 32382396
c.2571_2578 delinsC	p.Thr859Alafs*63	DD, ID, ASD, abnormal MRI, mild hypotonia (1 pt)	absent	Lu (2017) PMID: 28288114
c.2974C>T	p.Gln992*	DD, ID, ASD, ADHD, abnormal MRI, marfanoid habitus, stereotypic movements (1 pt)	absent	Lu (2017) PMID: 28288114
c.3795+1G>C	-	nsID (1/108 pts)	absent	Kim (2019) PMID: 30952489

ASD: autism spectrum disorder; CLP: cleft lip and palate; DD: developmental delay; ID: intellectual disability; MRI: magnetic resonance imaging; nsID: nonsyndromic intellectual disability; pt(s): patient(s). All variants described for CIC (NM\_015125.4) in HGMD® Professional 2021.1; variants frequencies from <a href="https://gnomad.broadinstitute.org/">https://gnomad.broadinstitute.org/</a>.