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Epidemiological study of congenital and hereditary anomalies in Sialkot District of Pakistan revealed a high incidence of limb and neurological disorders

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Abstract

Background: Pakistan faces high incidence of congenital anomalies (CA) and hereditary anomalies due to various factors, including a high rate of consanguinity, early marriages, and predominance of extended families. There is a paucity of epidemiological studies that could provide a baseline for management strategies for these anomalies. **Objectives:** We aimed to elucidate the pattern, as well as the clinical and genetic aspects, of CA prevalence among the general population in Sialkot District of Pakistan.

Methods: In a cross-sectional sampling design, subjects and families with a certain type of CA were recruited from hospitals and medical centers in Sialkot District. Subjects were also selected from various towns and remote villages by visiting public places. Phenotypic and descriptive data were obtained, pedigrees were constructed, and parental and demographic attributes were recorded.

Results: A total of 241 independent subjects and/or families with CA were recruited. The malformations were classified into five major and 56 minor categories. Limb defects had the highest representation (n = 113; proportion = 0.469; 95% confidence interval (CI) = 0.406–0.532), followed by neurological anomalies (n = 76; proportion = 0.315; 95% CI = 0.257–0.374). Among the limb defects, polydactyly and talipes were most prevalent while, among neurological disorders, intellectual disability and cerebral palsy were more frequent. In this cohort, sporadic occurrence was customary compared to the familial presentation (n = 144 vs 97). Analyses of various attributes, such as gender differences, parental consanguinity, and paternal ages, as well as pedigree analyses, revealed marked heterogeneity among the major and minor categories of CA.

Conclusion: The pattern of anomalies witnessed in this cohort and a high occurrence of sporadic cases point to a substantial role of nongenetic etiological factors, which could be minimized by strengthening the health-care system.

Keywords: consanguinity; genetic diseases; prevalence

Congenital anomalies (CA) are a heterogeneous group of disorders that are present at or before birth. CA could be due to genetic factors, such as single gene defects and chromosomal abnormalities, or could be caused by environmental influences, which include micronutrient deficiencies, teratogen exposure, and infections [1]. A large number of CA with genetic etiology

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may be hereditary in nature. Every year, 303,000 children die within 4 weeks of birth due to CA, and others with long-term disabilities negatively affect families, health-care systems, and societies [2].

In developed countries, there is systematic registration of CA [1, 3, 4]. In many developing countries, however, birth defect registries are not maintained. Monitoring the malformations and establishing their sociodemographic correlates is not an easy task and is hampered by a number of factors, including inadequate infrastructure for surveillance, ascertainment biases – particularly in rare anomalies, and invasive methods of diagnosis in certain cases. In the absence of detailed epidemiological data on CA, it is difficult to evaluate possible risk factors and to implement effective prevention and care services [5, 6].

The health-care system of Pakistan is facing various challenges in service delivery to the population. The majority of the population resides in rural areas, where the infrastructure of health care is present, but it is poorly maintained, in addition to lacking modern equipment and trained professionals [7]. Despite significant improvements over the past two decades in Pakistan, the infant and neonatal mortality rates remain high. In 2015, Pakistan was ranked 149th among 179 countries on the Maternal Mortality Ratio Index, which is very alarming [8]. The allocation of health-care resources, such as finance and transport, is not a need-based process in Pakistan. Furthermore, owing to the high rate of population growth, the number of health-care professionals is inadequate and the existing medical staff is untrained, underpaid, and deprived of the latest facilities for medical practice. Nonetheless, the health-care system mainly relies on private organizations, which provide more advanced facilities at high prices, and the masses cannot afford these facilities. There are massive inequalities in the accessibility of health-care services to the low-socioeconomic-status population [7-9].

According to a recent estimate, CA caused 2.34% of total deaths in Pakistan [3]. Here, the high incidence of CA has been attributed to various factors, such as maternal malnutrition, inadequate prenatal care, poor socioeconomic setup, rural origin, and a high rate of consanguinity [10–12]. In a study carried out in the Combined Military Hospital, Kharian, Hussain et al. [13] showed that CA affected 7% of the 3,210 cases. Previously, Perveen and Tyyab [14] showed that neural tube defects were the most prevalent type of anomalies in a tertiary care hospital in Karachi and the commonest associated risk factor was consanguinity. A detailed survey of the medical literature revealed that the majority of the reports on CA were hospital based [10–13]. Such studies may have an underrepresentation of subjects from rural areas, where most of the deliveries take place at home and with the help of traditional birth attendants [15]. Due to lack of proper surveillance and documentation, the countrywide picture on the epidemiology of CA is fragmented.

The present clinico-epidemiological study on CA was carried out in the Sialkot District of Pakistan, where the majority of the population resides in rural areas.

Methods

Study population

A descriptive clinical and genetic epidemiological study on CA was carried out in the Sialkot District, which is located in the northeast of Pakistan. The district has four administrative units called "tehsils". The rural population comprises 74% of the total population, and the literacy rate is 59%. Major ethnicities are the Jatts and Gujjars, while there is also a representation of Rajput, Arain, and Kashmiris. Major languages are Punjabi and Kashmiri [16]. According to the Pakistan Census of 2017, it had a population of 3.9 million. It is an industrial city and is famous for its leather, sports, and surgical products [17].

Ethical considerations and sampling strategy

The study was approved by the Ethical Review Committee of Quaid-i-Azam University, Islamabad (DFBS/2014–3278). All the data were acquired and documented in writing in the presence of family head/guardian after participants had provided informed written or formally documented verbal consent when illiterate. When a participant was below the legal age of providing consent or was incapable of providing it because of disability (deaf-mute, blind, neuromuscular defect), or otherwise incapable, a parent/guardian or literate elder provided written informed consent. They were assured before the start of the study that there would be no breaches of an ethical nature or confidentiality.

In a cross-sectional sampling design, subjects and families with CA and/or hereditary anomalies were recruited from June 2013 to December 2014. The subjects/families were recruited from District Headquarters Hospitals and Basic Health Units in different rural and urban areas of Sialkot District. Since there is no systematic record keeping in the public sector hospitals, all the cases were retrospectively recruited by the data collection team. The subjects were also recruited from public places such as community centers and rural gathering areas and were brought to the nearest medical center for examination. Each subject was physically examined by a local physician. The previous health record of the subjects when available was assessed by two expert health workers. The fieldwork and data collection were performed by one of the coauthors, who had had formal training in medical genetics, along with a local resource person who guided in subject recruitment. The resource persons were male nurses or paramedical staff who were familiar with the local population. The data were not collected from any rehabilitation or special education center.

Classification of anomalies

For the primary diagnosis, we relied on the assessment provided by the specialist resident doctor, and only those individuals with confirmed congenital and/or hereditary anomalies were recruited. The cases were identified with respect to the index male or female subject and were categorized as follows: (i) familial or sporadic, and (ii) isolated or syndromic. Syndromic cases were identified with respect to the more severe symptoms in the following order: neurological disorders, neuromuscular defects, musculoskeletal defects, eye/visual impairments, sensorineural/ear anomalies, and limb defects. Secondary symptoms, when present, were separately scored as associated malformations. A pedigree comprising three or more generations was drawn for each case; however, only the index subject was considered in the data analyses. Malformations of infectious or traumatic nature and subjects with poliomyelitis were not included. All the information was recorded on a structured questionnaire that was divided into three sections: the first section included demographic data; the second section dealt with various risk factors such as parental medical history (including paternal ages) and consanguinity; and the third section documented the phenotypic details of the anomalies.

The definition of CA was based on a standard coding system of the International Classification of Diseases, Tenth Revision (ICD-10), according to the primary diagnosis [18]. The corresponding definitions of each entity were searched in the Online Mendelian Inheritance in Man (OMIM) and Orphanet databases [19, 20]. Limb defects, such as syndactyly and polydactyly, were further characterized into well-described entities [21–23]. For each CA, the proportions and 95% confidence intervals (95% CIs) were estimated from the total number of anomalies. Statistical analyses were performed using MS-Excel and GraphPad Prism.

Results

Sample characteristics

A total of 241 independent subjects/families with certain types of CA were identified in this study. There were 181 (75%) index males and 60 (25%) index females (**Table 1**). The CA were classified into five major and 56 minor categories (**Table 2**). Among the major categories, limb defects had the highest representation (n = 113; proportion = 0.337; CI = 0.278–0.397), followed by neurological disorders (n = 76), musculoskeletal defects (n = 23), and neuromuscular anomalies (n = 10). Seven categories with <6 cases were lumped into "others". The sporadic occurrence was customary compared to the familial presentation (n = 144 vs 97). Across all families, the total number of affected individuals was 497, with a preponderance of affected males compared to affected females (324 vs 173; **Table 1**).

Table 1. Major categories of CA, familial/sporadic nature, and total number of affected family members

Congenital anomalies	Index subject		Proportion	95% CI	Familial/sporadic nature*		Total no. of people affected in all families*			
-	Male	Female	Total			Familial	Sporadic	Male	Female	Total
Limb defects	89	24	113	0.469	0.406-0.532	48	65	153	69	222
Neurological disorders	57	19	76	0.315	0.257-0.374	20	56	83	35	118
Musculoskeletal defects	16	7	23	0.095	0.058-0.133	8	15	23	18	41
Neuromuscular anomalies	9	1	10	0.042	0.016-0.067	5	5	21	2	23
Others (<i>n</i> < 6)	10	9	19	0.079	0.045-0.113	16	3	44	49	93
Total	181	60	241	1.000	-	97	144	324	173	497

*Chi-test statistics were statistically highly significant.



Limb defects

Limb defects were further resolved into 16 distinct entities (**Tables 2 and 3**). Clubfoot had the highest representation (n = 26), followed by postaxial polydactyly type A (n = 23), preaxial polydactyly type I (n = 21), and arthrogryposis (n = 10). Certain defects, such as arthrogryposis, clubfoot, preaxial polydactyly type I, and reduction defects of upper limbs, mostly had sporadic presentation, whereas cases with postaxial polydactyly type A were more often familial. In all families with limb anomalies, a total of 222 subjects were observed to be affected.

Among the index cases with limb defects (n = 113), a total of 196 limbs were involved (**Table 4**). There was relatively

higher involvement of the lower limbs compared to the upper limbs (109 vs 87) and of the right leg compared to the left leg (61 vs 48). Detailed analyses of the phenotypic variability and the combination of involved limbs are presented in **Table 4**.

Neurological disorders

Intellectual disability types (mild, severe, Down syndrome, and microcephaly) were in a sizable number and appeared in 35 subjects (**Table 2**). Cerebral palsy (CP) was the primary malformation in at least 32 subjects (and an associated malformation in further 18 individuals with intellectual disabilities; **Table 5**).

Table 2. Major and minor categories of congenital/hereditary malformations observed in the studied population

Malformation (major/minor)	No. of cases	s Proportion	95% CI	ICD-10*	OMIM**	ORPHA#
Limb defects	113	0.469	0.406-0.532			
Clubfoot	26	0.108	0.069–0.147	Q66	119800	293150, 199315
Polydactyly, postaxial type A	23	0.095	0.058-0.133	Q69.0;Q69.2	174200	93335
Polydactyly, preaxial type I	20	0.083	0.048-0.118	Q69.1	174400	93339
Arthrogryposis	10	0.042	0.016-0.067	Q74.3	108120	1146
Polydactyly, postaxial type B	6	0.025	0.005-0.045	Q69.0;Q69.2	174200	93335
Reduction defects of upper limb	6	0.025	0.005-0.045	Q71	217100	93457
Brachydactyly, 4th toe; brachymetatarsus IV	4	0.017	0.001-0.033	Q72.8	113475	294998
Congenital shortening of lower limb	4	0.017	0.001-0.033	Q72.8		295057
Syndactyly, type 1a	4	0.017	0.001-0.033	Q70.3	609815	93402, 295187
Brachydactyly, all fingers short	2	0.008	-0.003 to 0.020	Q71.8		295130
Reduction defects of lower limb	2	0.008	-0.003 to 0.020	Q72		93457
Syndactyly, type 1c	2	0.008	-0.003 to 0.020	Q70.1		295191
Camptodactyly	1	0.004	-0.004 to 0.012		114200	295016
Polydactyly, preaxial type II	1	0.004	-0.004 to 0.012	Q74.0	174500	2950, 93336
Split-hand/foot	1	0.004	-0.004 to 0.012	Q72.7	183600	2440
Syndactyly, type II	1	0.004	-0.004 to 0.012	Q70.4	186000	295195
Neurological disorders	76	0.315	0.257-0.374			
Intellectual disability: mild	21	0.087	0.052-0.123	F70	249500	88616
Spastic diplegic cerebral palsy	20	0.083	0.048-0.118	G80.1		
Intellectual disability: severe	11	0.046	0.019-0.072	F72	611091	88616
Hereditary neuropathy	6	0.025	0.005-0.045	G60	201300	970
Cerebral palsy, unspecified	4	0.017	0.001-0.033	G80.9	605388	
Spastic hemiplegic cerebral palsy	3	0.012	-0.002 to 0.026	G80.2		
Spastic quadriplegic cerebral palsy	3	0.012	-0.002 to 0.026	G80.0	603513	210141

(Continued)

Table 2. Major and minor categories of congenital/hereditary malformations observed in the studied population (Continued)

Malformation (major/minor)	No. of cases	Proportion	95% CI	ICD-10*	OMIM**	ORPHA#
Ataxic cerebral palsy	2	0.008	-0.003 to 0.020	G80.4	605388	
Amyotrophic lateral sclerosis	1	0.004	-0.004 to 0.012	G12.2	105400	803
Down syndrome	1	0.004	-0.004 to 0.012	Q90.9	190685	870
Hereditary ataxia	1	0.004	-0.004 to 0.012	G11	229300	95
Hereditary spastic paraplegia	1	0.004	-0.004 to 0.012	G11.4	270800	
Intellectual disability; succinic semialdehyde dehydro- genase deficiency	1	0.004	-0.004 to 0.012		271980	22
Microcephaly, low IQ^	1	0.004	-0.004 to 0.012	Q02	251200	2512, 52183
Musculoskeletal defects	23	0.095	0.058-0.133			
Achondroplasia	12	0.050	0.022-0.077	Q77.4	100800	15
Kyphosis	2	0.008	-0.003 to 0.020	Q76.4	192900	
Congenital deformity of forehead	1	0.004	-0.004 to 0.012	Q75.8		
Congenital dislocation of hip	1	0.004	-0.004 to 0.012	Q65.2	142700	
Congenital malformation of bony thorax; genu valgum	1	0.004	-0.004 to 0.012	Q76.9; Q74.1		
Congenital scoliosis	1	0.004	-0.004 to 0.012	Q76.3		
Disorder of ligament (leg)	1	0.004	-0.004 to 0.012	M24.2		
Diastrophic dysplasia	1	0.004	-0.004 to 0.012	Q77.5	222600	628
Genu valgum	1	0.004	-0.004 to 0.012	Q44.1	137370	
Multiple congenital exostoses	1	0.004	-0.004 to 0.012	Q78.6	133700	321
Osteochondrodysplasia, unspecified	1	0.004	-0.004 to 0.012	Q78.9		
Neuromuscular anomalies	10	0.042	0.016-0.067			
Muscular dystrophy	8	0.033	0.011-0.056	G71.0	310200	98896
Spinal muscular atrophy	2	0.008	-0.003 to 0.020	G12.1	253300	70, 83330
Others	19	0.079	0.045-0.113			
Oculocutaneous albinism	3	0.012	-0.002 to 0.026	E70.3	203100	79431, 352731
Allergic asthma	3	0.012	-0.002 to 0.026	J45.9		
Deaf-mute	3	0.012	-0.002 to 0.026	H91.3.	220290	90636
Dysphasia	1	0.004	-0.004 to 0.012	R47.0	600117	
Amblyopia	1	0.004	-0.004 to 0.012	H53.0		
Congenital malformation of cardiac septum	1	0.004	-0.004 to 0.012	Q21.9	600001	
Deformed ear pinna	1	0.004	-0.004 to 0.012	H61.9		156243
Disorders of fluid, electrolyte, and acid-base balance	1	0.004	-0.004 to 0.012	E87		
High myopia	1	0.004	-0.004 to 0.012	H52.1	160700	
Hypertensive heart disease	1	0.004	-0.004 to 0.012	111		
Intestinal cancer	1	0.004	-0.004 to 0.012	C17.9	26106	
Night blindness	1	0.004	-0.004 to 0.012	H53.6	257270	215
Vitiligo	1	0.004	-0.004 to 0.012	L80		

*ICD-10, International Classification of Diseases, Tenth Revision; AIQ, intelligence quotient; **OMIM, Online Mendelian Inheritance in Man; ORPHA, #Orphanet database identifiers/Entrez.



Table 3. Spectrum of limb defects and their distribution with respect to familial/sporadic nature and total number of affected family members

		Index cases	;	Familial	/sporadic	Total no. of p	eople affected	cted in all families
LIMD defects*	Male	Female	Total	Familial	Sporadic	Male	Female	Total
Arthrogryposis	8	2	10	2	8	10	8	18
Brachydactyly, brachymetatarsus IV	4	0	4	3	1	8	4	12
Brachydactyly, all fingers	0	2	2	1	1	1	2	3
Camptodactyly	1	0	1	1	0	2	1	3
Clubfoot	19	7	26	8	18	30	11	41
Congenital shortening of lower limb	4	0	4	1	3	4	1	5
Polydactyly, postaxial type A	19	4	23	13	10	35	16	51
Polydactyly, postaxial type B	5	0	5	2	3	7	0	7
Polydactyly, preaxial type I	17	4	21	8	13	25	13	38
Polydactyly, preaxial type ll	0	1	1	0	1	0	1	1
Reduction defects of lower limb	2	0	2	1	1	2	1	3
Reduction defects of upper limb	5	1	6	1	5	7	3	10
Split-hand/foot	1	0	1	1	0	6	2	8
Syndactyly, type ll	0	1	1	1	0	1	2	3
Syndactyly, type 1a	4	0	4	3	1	9	1	10
Syndactyly, type 1c	0	2	2	2	0	6	3	9
Total	89	24	113	48	65	153	69	222

*Presented in alphabetical order.

 Table 4. Phenotypic manifestation of limb defects (n = 113)

Limb defects	No. of	Total no. of	Uppe (<i>n</i> =	r limb 87)	Lowe (n =	r limb 109)	No. o invol	of cases wit vement of	th 		No. of invo	limbs lved	
	cases	affected limbs	RA	LA	RL	LL	Arms only	Legs only	Both	Any 1	Any 2	Any 3	All 4
Arthrogryposis	10	22	6	7	5	4	5	1	4	3	4	1	2
Brachydactyly, brachyme- tatarsus IV	4	5	0	0	3	2	0	4	0	3	1	0	0
Brachydactyly, all fingers	2	4	2	2	0	0	2	0	0	0	2	0	0
Camptodactyly	1	4	1	1	1	1	0	0	1	0	0	0	1
Clubfoot	26	44	0	0	23	21	0	26	0	8	18	0	0
Congenital shortening of lower limb	4	5	0	0	4	1	0	4	0	3	1	0	0
Polydactyly, postaxial type A	23	50	12	12	16	10	3	9	11	9	6	3	5
Polydactyly, postaxial type B	5	6	2	3	1	0	4	1	0	4	1	0	0
Polydactyly, preaxial type	l 21	25	10	13	1	1	20	0	1	19	1	0	1
Polydactyly, preaxial type ll	1	1	1	0	0	0	1	0	0	1	0	0	0
Reduction defects of lower limb	2	3	0	0	1	2	0	2	0	1	1	0	0
Reduction defects of upper limb	6	7	4	3	0	0	6	0	0	5	1	0	0
Split-hand/foot	1	4	1	1	1	1	0	0	1	0	0	0	1
Syndactyly, type II	1	4	1	1	1	1	0	0	1	0	0	0	1
Syndactyly, type 1a	4	8	0	0	4	4	0	4	0	0	4	0	0
Syndactyly, type 1c	2	4	2	2	0	0	2	0	0	0	2	0	0
Total	113	196	42	45	61	48	43	51	19	56	42	4	11

LA, left arm; LL, left leg; RA, right arm; RL, right leg

Associated malformations

The secondary symptoms appearing with the primary presentation in the index subjects were scored as associated malformations. The combinations of associated malformations are depicted in **Table 5**. Limb anomalies were in the majority (n = 23), followed by CP (n = 18) and deaf-mute (n = 8) cases.

Familial cases

The pedigrees with at least two affected subjects with a similar phenotypic presentation were considered as familial (n = 97). The analyses of pedigree structures revealed that the anomalies segregated in one or two generations in most of the cases (n = 47 and 34, respectively; data not shown). In the majority of pedigrees, the malformations appeared in two or one independent sibships (n = 45 and 26, respectively).

Parental consanguinity and parental ages

Parental consanguinity was estimated to be 17% in the overall sample. The differences in the distribution of consanguineous and nonconsanguineous unions among the major categories of CA were statistically significant (**Table 6**). Consanguinity was relatively higher in subjects with neuro-muscular anomalies and neurological disorders (30 and 21%, respectively). Furthermore, consanguinity was significantly higher among familial cases compared to sporadic anomalies (29 vs 9%, respectively; P < 0.0001) (**Table 6**).

The differences in the mean paternal and maternal ages were statistically significant in all categories of sporadic and familial samples (**Table 7**). For the subjects with neuromuscular anomalies, the mean paternal ages were higher than for those with other anomalies.

The distribution of index subjects with respect to the demographic variables is presented in **Table 8**.

Table 5. Combinations of malformations associated with CA

Association CA category	Brachydactyly	Campto- dactyly	Syndactyly type	Clubfoot	Arthro- gryposis	Deaf- mute	Cerebral palsy	Others
Achondroplasia		1				1		High myopia
Arthrogryposis			111					
Camptodactyly	First toe							
Clubfoot	Fourth toe	3	1a					
Deaf-mute								Polydactyly, postaxial type A
Down syndrome	Fourth toe							Squint eye
Dysphasia					1			
Hereditary neuropathy	Third toe							Squint eye
Intellectual disability: mild					3	2	10	Ataxia
Intellectual disability: severe	Fourth toe			1		3	8	Cleft lip, overriding toe; polydactyly postaxial type B
Kyphosis	Fourth toe			1				
Muscular dystrophy								Cleft lip
Polydactyly, postaxial type A			1a					Clinodactyly
Polydactyly, preaxial type l		1						Hypoplastic thumb
Reduction defects of lower limb				1	1			
Spastic diplegic cerebral palsy	Third toe					1		Cleft palate
Spinal muscular atrophy	,					1		
Total	7	5	3	3	5	8	18	



Table 6. Relationship between consanguinity and different sample types

C-1	Parental mar	Total and a function of	
Category	Consanguineous [#]	Nonconsanguineous	- Iotal no. of marriages
Malformation			
Limb defects	10 (9)	103 (91)	113
Neurological disorders	16 (21)	60 (79)	76
Musculoskeletal defects	4 (17)	19 (83)	23
Neuromuscular anomalies	3 (30)	7 (70)	10
Others	8 (42)	11 (58)	19
		<i>P</i> = 0.003	
Familial/sporadic nature			
Familial	28 (29)	69 (71)	97
Sporadic	13 (9)	131 (91)	144
Total	41 (17)	200 (83)	241
		<i>P</i> < 0.0001	

[#]Inbreeding coefficient *F*³ 0.0313

Table 7. Mean parental ages at index subject's birth

Category	N	Paternal age (mean ± SD)	Maternal age (mean ± SD)	Significance level (P)*
Sporadic cases (<i>n</i> = 137)				
Limb defects	62	33.4 ± 5.0	27.0 ± 5.2	<0.0001
Neurological disorders	56	32.8 ± 4.8	26.3 ± 4.6	<0.0001
Musculoskeletal defects	14	35.8 ± 4.9	29.3 ± 4.7	0.0014
Neuromuscular anomalies	5	40.0 ± 3.7	32.8 ± 3.6	0.0143
Total	137	33.7 ± 5.1	27.1 ± 5.0	<0.0001
		ANOVA, <i>P</i> = 0.910	ANOVA, <i>P</i> = 0.726	
Familial cases (<i>n</i> = 70)				
Limb defects	43	34.7 ± 7.4	27.7 ± 6.2	<0.0001
Neurological disorders	16	31.8 ± 5.6	25.4 ± 4.8	0.0018
Musculoskeletal defects	7	31.4 ± 5.6	25.3 ± 5.9	0.0702
Neuromuscular anomalies	4	36.0 ± 6.8	29.0 ± 5.0	0.1469
Total	70	$\textbf{33.8} \pm \textbf{6.9}$	$\textbf{27.0} \pm \textbf{5.9}$	<0.0001
		ANOVA, <i>P</i> = 0.561	ANOVA, <i>P</i> = 0.689	

*T-test statistics were used for comparison between paternal and maternal ages.

One-way analysis of variance (ANOVA) was used for analysis among the paternal ages or the maternal ages.

Table 8. Demographic characteristics of index subjects

Demonwouhiowawiahlaa	Ge	nder	Familial	Total	
Demographic variables	Male	Female	Familial	Sporadic	
Age category, years					
≤9	26	19	16	29	45
>9–19	67	28	28	67	95
>19–29	40	5	22	23	45
>29–39	22	4	15	11	26
≥39	26	4	16	14	30
Total	181	60	97	144	241
	<i>P</i> = 0.0027		<i>P</i> = 0.0187		

Table 8. Demographic characteristics of index subjects (Continued)

	Ge	ender	Familial	/sporadic	Total
Demographic variables	Male	Female	Familial	Sporadic	
Rural/urban origin					
Rural	148	45	72	121	193
Urban	33	15	25	23	48
		<i>P</i> = 0.2553		<i>P</i> = 0.0617	
Caste system					
Jatt	51	12	32	31	63
Rajput	32	7	13	26	39
Arain	18	7	9	16	25
Malik	13	9	10	12	22
Meher	14	4	10	8	18
Others	53	21	23	51	74
		<i>P</i> = 0.3207		<i>P</i> = 0.1325	
Migratory origin [#]					
Native	150	42	76	116	192
Migrated	31	18	21	28	49
		<i>P</i> = 0.0318		<i>P</i> = 0.6766	
Literacy (age >5 years)					
lliterate	98	31	49	80	129
Literate	71	21	41	51	92
		<i>P</i> = 0.8351		<i>P</i> = 0.3263	
Family type					
Nuclear	107	30	67	70	137
Extended	74	30	30	74	104
		<i>P</i> = 0.2166		<i>P</i> = 0.0017	

*During the 1947 partition.

Discussion

Pakistan bears a high burden of congenital and hereditary anomalies. The health-care system is not able to provide management and support to the subjects/families afflicted with such anomalies, resulting in a great social, economic, and psychological impact on the involved families and the society at large. In Pakistan, 6%–9% perinatal deaths are due to CA [24]. Many of the known risk factors, such as advanced maternal age, exposure to teratogens and radiation, maternal illnesses, and smoking, could be substantially minimized by educating pregnant women and providing them timely antenatal care. Furthermore, various screening methods, such as determination of maternal serum markers, ultrasonography, amniocentesis, and chorionic villus sampling, can be utilized to detect the at-risk pregnancies and their subsequent management. In this context, basic epidemiological data provide useful grounds for assigning priorities, allocating resources, and establishing monitoring and management systems for these disorders.

Limb defects had the highest representation in the present study, 113 cases with primary presentation and at least 28 cases

with associated defects. Polydactyly types were most prevalent, followed by clubfoot, arthrogryposis, reduction defects, and syndactyly types. Most of the limb/digit defects do not cause severe disability and, owing to their minor nature, such anomalies remain less reported in epidemiological studies [12, 25, 26]. As witnessed in the present cohort, the majority of limb defects were of milder nature and did not result in any disability. However, there were at least 45 cases that were the potential sources of disability (including clubfoot = 26; arthrogryposis = 10; reduction defects of upper limb = 6; reduction defects of lower limb = 2; split-hand/foot = 1). In a study carried out in a tribal area of Pakistan, Zahra et al. [27] demonstrated that limb defects were the third most common anomalies (21%), after neurological disorders (34%) and musculoskeletal defects (23%). Curiously, amputations/reduction defects were the most common types among limb anomalies. Polydactyly was also witnessed to be the most common type of limb anomaly in other studies [25, 28].

Among the neurological disorders, CP was a major source of severe disability in our cohort. CP is a chronic motor disorder that is nonprogressive in nature. Subjects with CP may have several problems, including walking disability, hearing and eye problems, seizures, feeding problems, and intellectual disability [29, 30]. In our data, different clinical types of CP were recognized, including hemiplegia, diplegia, quadriplegia, and ataxia. There are several other conditions that should be considered in the differential diagnosis of CP. For instance, diplegic CP (OMIM-605388) resembles hereditary spastic paraplegia (HSP). However, CP is nonprogressive, whereas HSP is characterized by a steady weakness of the lower limbs.³¹ On the other hand, in muscular dystrophies, there is no spasticity, yet the patients can develop contractures. In primary dystonia (OMIM-128100), which is another movement disorder, the onset of muscular deformity occurs after several years of normal development. The patients have sustained episodes of muscle contraction and dystonia but without the development of contractures. Hereditary spastic paraparesis type 4 (OMIM-182601) is the single most common dominantly inherited paraparesis, representing approximately 40% of all cases [32].

Among the musculoskeletal anomalies, achondroplasia was predominant (n = 12) in the present cohort, followed by other disorders, including kyphosis, deformities of the hip, and disorders of the ligament, which were also the causes of physical disability. Another source of disability among the subjects was the occurrence of neuromuscular disorders, represented by muscular dystrophy (n = 8) and spinal muscular atrophy (n = 2). Azhar et al [33]. carried out an interventional study on disabled subjects in Sialkot District. Among the 644 individuals with disabilities, poliomyelitis was a predominant disorder, followed by CP, skeletal dysplasias, muscular dystrophy, congenital dislocated hip, and talipes equinovarus. The cases of poliomyelitis were excluded from our sample (n = 7). Nonetheless, the primary focus of Azhar et al. [33] was on the recovery of disabled individuals by surgical interventions, physiotherapy, and bracing, and no attempt was made to describe the nature and clinical spectrum of anomalies. The current study, however, presents a range of phenotypes for these anomalies.

Parental consanguinity was calculated to be 17% in the present cohort. Among the subjects with neuromuscular anomalies, the rate of consanguinity was 30%. This is rather surprising as several studies have advocated a high incidence of consanguineous marriages, that is, 57–62%, in various Pakistani populations [34–36]. Nonetheless, the increased incidence of CA is generally attributed to a high rate of parental consanguinity in Pakistani society [11]. The low rate of consanguinity may indicate the involvement of nongenetic factors in the etiology of these anomalies.

In this study, the high incidence of limb and neurological disorders, the preponderance of cases with sporadic nature,



and the relatively low level of parental consanguinity may indicate a substantial contribution of environmental factors in the etiology of CA. Gao et al. [37] in a case–control study demonstrated that several factors, including advanced maternal age, alcohol consumption during pregnancy, rural residence, father's occupational exposure to harmful substances, and multiple births, were important risk factors for CP in Chinese children. Prenatal exposure to environmental factors was reported to cause congenital limb defects [38]. Furthermore, agricultural compounds in water were observed to be the source of birth defects, which include limb anomalies [39].

There could be several potential nongenetic causes of the high prevalence of limb and neurological anomalies in the study region, for instance, poor prenatal care, lack of basic health facilities, and maternal exposures. In the rural areas of Pakistan, primary maternity care is provided by traditional midwives/birth attendants, who are not professionally trained and cannot handle birth complications [15]. Birth defects such as CP may appear in newborns due to birth asphyxia. Second, Sialkot is an industrial city that is famous for its leather, sports, and surgical products. With 117 operational tanning units, Sialkot is the second largest tanning cluster of Punjab Province. Only 3% of industrial plants meet international waste treatment standards [17]. A large number of industries discharge their toxic waste into natural watercourses and open streams [40, 41]. The industrial waste is used to irrigate vegetable and fruit farms [42]. Third, due to the adjoining boundary with India, the rural population is continuously threatened by cross-border shelling, which may add heavy metals in the human food chain. Thus, Sialkot population is constantly exposed to teratogens, such as heavy metals, which could be a potential cause for the high incidence of CA in the region. However, a direct association between environmental exposures and CA incidence remains to be established in this population.

Similar to other epidemiological studies, the current study has several limitations [25, 27, 28]. For instance, the exact prevalence rate of CA was not established in this study. This study presents CA cases that were rather explicit in nature and could be easily phenotyped with the physical examination, notwithstanding the fact that detailed clinical data, including X-rays and other laboratory investigations, were available for many of the cases/families. Maladies of biochemical or metabolic nature or those that require invasive diagnostic methods (such as structural brain defects) could be underrepresented in our cohort. Furthermore, there was no exhaustive coverage of areas of the whole district due to sociocultural limitations and unavailability of resource persons.

Conclusion

This preliminary study in Sialkot District presents a detailed clinical and descriptive account of CA prevalent in this population. High incidence of limb and neurological anomalies, low level of parental consanguinity, and the preponderance of sporadic cases in this cohort suggest a significant role of nongenetic etiological factors, which could be minimized by strengthening the health-care system.

Author contributions. Sajid M contributed to the conception and design of this study. NAB collected the data and drafted the initial version of the manuscript. Sara M and Sajid M critically revised it and completed the write-up. All the authors approved the final version submitted for publication and take responsibility for statements made in the published article.

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Conflict of interest statement. The authors have completed and submitted the International Committee of Medical Journal Editors Uniform Disclosure Form for Potential Conflicts of Interest. None of the authors disclose any conflict of interest.

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