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The prevalence of patients with ocular genetic disorders attending a general paediatric ophthalmology clinic in the East End of London

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Abstract

Aims To document the prevalence of ocular genetic disorders in a general paediatric ophthalmology clinic in a deprived, ethnically diverse inner city area of London, and to assess the consanguineous relationships and ethnicity among families with these conditions. Methods A prospective audit documented all ocular genetic conditions (excluding strabismus and amblyopia) presenting for 16 consecutive weeks to the paediatric ophthalmology clinic. Information regarding ethnicity and consanguinity were sought. The disorders were divided according to the mode of inheritance if known.

Results Thirteen per cent of patients (45/342) had an ocular genetic disorder or were being examined for one. Of them, 22% (10/45) had a history of consanguinity with an inheritance pattern of 30% autosomal recessive (3/10), 20% autosomal dominant, 50% X-linked/unknown/ isolated cases. In the remaining nonconsanguineous families (35/45), 22% were autosomal recessive, 17% autosomal dominant, and 60% X-linked/unknown/isolated cases. The vast majority of cases (9/10) with a history of consanguineous marriage had South Asian ancestry. Variable ethnic backgrounds were documented for patients with genetic disease and no consanguinity.

Conclusion Ocular genetic disorders are common in secondary care. Less than a third of patients with such disorders had a family history of consanguinity. The proportion of patients with proven autosomal recessive disease was similar irrespective of

consanguinity within family. The proportion of children of South Asian ancestry was high in this clinic population.

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Introduction

Providing a service in paediatric ophthalmology in a deprived, ethnically diverse inner city area can prove challenging. Ocular genetic disorders involve the use of resources and require a multidisciplinary team approach to diagnose and counsel families appropriately. Although amblyopia and squint have a genetic predisposition, these conditions were not considered as ocular genetic disorders for this study. It is believed that high levels of ocular genetic disease are present in East London and social customs of the local communities contribute to this burden.

Consanguinity is defined by marriage between second cousins or closer relatives and is prevalent in the South Asian community in Britain. Over 50% of marriages in the Pakistani community in the 1980s were consanguineous compared to 1% in the general population.² Pakistani children of consanguineous unions have increased post-neonatal mortality and childhood morbidity compared with other ethnic groups.3 The predominant ethnic community in East London is Bangladeshi and consanguineous marriages do occur.

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Some ocular genetic disorders are characterised by only one inheritance pattern. Autosomal recessive disease occurs by the inheritance of a copy of a recessive allele from each parent. An increased frequency of proven autosomal recessive disorders is expected to be present in children of consanguineous parents compared to non-consanguineous parents. In addition, the predominant inheritance pattern of ocular genetic disorders in patients with consanguineous parents is expected to be autosomal recessive.

Materials and methods

A prospective audit documented all ocular genetic conditions (excluding strabismus and amblyopia) presenting for 16 consecutive weeks to the general paediatric ophthalmology clinic from March 2006. Details regarding the ethnicity of all patients entering clinic were also sought. Consanguinity was recorded in patients with or suspected of having an ocular genetic disorder and was restricted to second cousins or closer. No patient was documented more than once during the 16-week audit period.

Co-variates such as autosomal recessive inheritance, consanguinity and ethnicity were individually assessed using the Fisher's exact test and P < 0.05 was considered statistically significant. Institutional approval for this work was acquired before the start of the project.

Results

During the 16 weeks of this audit, 342 patients presented to the clinic at least once. The most common ethnic group was Bangladeshi followed by white children of English/ Scottish and Welsh descent (see Table 1).

Overall, 45 children were identified with an ocular genetic disorder or possible ocular genetic disorder. Ten children had parents who were first or second cousins. Thirty-five children had parents who were nonconsanguineous (including a family with a third cousin marriage). The conditions and known inheritance patterns are listed in Table 2. The most common entity was Down's syndrome (7) followed by non-syndromic retinitis pigmentosa (4) and sickle cell disease (4). There were 17 conditions that presented in a single patient during this period.

In patients with consanguineous parents, 9 of 10 had South Asian ancestry. Three children had documented autosomal recessive disease (two with syndromic pulverulent cataracts and one with global developmental delay and myopia). The ethnic makeup of the consanguineous, non-consanguineous, and non-ocular genetic disorder groups was assessed. The proportion of children of South Asian ancestry in the consanguineous

Table 1 Patients who presented during the 16-week study period categorised by ethnicity (n = 342)

Ethnicity	No.	Percentage	
Bangladeshi (including British)	126	37	
England/Scot/wales	76	22	
African (non-Somali)	27	8	
Pakistani (including British)	12	3	
Not stated	14	4	
Mixed	19	5	
Indian (including British)	9	3	
Somali	6	2	
Black British	11	3	
Caribbean	7	2	
Any other group	7	2	
Other Asian	6	2	
Other black	6	2	
Turkish/Turkish Cypriot	4	1	
Eastern European	2	<1	
Greek/Greek Cypriot	2	<1	
Arab or Middle East	2	<1	
Irish	2	<1	
Jewish	2	<1	
Other white European	2	<1	

group was not statistically different from the non-consanguineous group (Fisher's exact test two-sided P-value P = 0.68) nor the group without ocular genetic disorders (P = 0.26).

Similar proportions of patients had autosomal recessive disease and autosomal dominant disease in both consanguineous and non-consanguineous groups (Table 3). A large number of patients in both groups have conditions affecting one individual. The inheritance pattern in such cases is unknown and there may be more than one possible mode of inheritance.

Discussion

There have been few studies looking at the prevalence of ocular genetic disorders in general paediatric ophthalmology clinics in the UK and as a result it is difficult to state if the proportion that is seen in East London is comparable with other areas of the country. As the examination and discussion with a family with an ocular genetic disorder is necessarily longer and more detailed than a child with more straightforward conditions, such cross-sectional studies are important when considering the allocation of resources to paediatric ophthalmology clinics and future planning. This study included children who were suspected of having ocular genetic disorders, as this reflects the burden on the workload in clinic.

We restricted consanguineous marriage between second cousins and closer and it is possible that



Table 2 Features of patients presenting with ocular genetic disorders

Ethnicity	Consanguinity	Genetic disorder	Inheritance pattern	
BANGLADESHI	No	Retinitis pigmentosa	Unknown	
NOT STATED	No	Retinitis pigmentosa	Unknown	
GREEK	No	Usher	AR	
PAKISTAN	No	22q11 deletion	Sporadic	
BANGLADESHI	No	X-linked retinoschisis	X-linked	
PAKISTAN	No	Craniosynostosis	AD	
ENGLISH	No	Microcephaly and muscular disorder	Unknown	
BANGLADESHI	No	Down	Sporadic	
MOROCCAN	No	Down	Sporadic	
BANGLADESHI	No	Down	Sporadic	
AFRICAN	No	Down	Sporadic	
ENGLISH	No	Retinitis pigmentosa	AD	
RWANDA	No	High myopia (>-10)	Unknown	
TURKISH	No	Louis-Bar	AR	
BANGLADESHI	No	High myopia (>-10)	Unknown	
ENGLISH	No	Retinitis pigmentosa	Unknown	
AFRICAN	No	Sickle cell disease	AR	
NIGERIAN	No	Sickle cell disease	AR	
CARIBBEAN	No	Sickle cell disease	AR	
BANGLADESHI	No	Tuberous sclerosis	AD	
ENGLISH	No	Ataxic telengiectasia	AR	
CARIBBEAN	No	Sickle cell disease	AR	
ALBANIAN	No	Idiopathic motor nystagmus	Unknown	
ENGLISH	No	CHARGE	Sporadic	
GREEK	No	Williams-like syndrome	Unknown	
ENGLISH	No	Trisomy 3p	Sporadic	
BANGLADESHI	No	Non-syndromic cataract	Unknown	
BANGLADESHI	No	Peter's anomaly	Unknown	
NIGERIAN	No	Craniosynostosis	AD	
BANGLADESHI	No	Tuberous sclerosis	AD	
BANGLADESHI	No	Pierre-robin	AD	
AFRICAN	No	Down	Sporadic	
BANGLADESHI	No	Knobloch	AR	
BANGLADESHI	No	Coloboma	Unknown	
BANGLADESHI	No	Down	Sporadic	
INDIA	Yes		AR-brother	
BANGLADESHI	Yes	Global developmental delay and myopia NF-1	AR-brother AD	
		NF-1 NF-1		
BANGLADESHI CARIBBEAN	Yes Yes		AD Unknown	
		Non-syndromic cataract		
BANGLADESHI	Yes	Familial exudative vitreoretinopathy	Unknown	
PAKISTAN	Yes	Down	Sporadic	
PAKISTAN	Yes	Agammaglobulinaemia	X-linked	
BANGLADESHI	Yes	Non-syndromic cataract	Unknown	
BANGLADESHI	Yes	Syndromic cataract	AR	
BANGLADESHI	Yes	Syndromic cataract	AR	

Table 3 Identified inheritance patterns of patients with consanguineous and non-consanguineous parents

	Autosomal recessive	Autosomal dominant	X-linked recessive	Sporadic	Unknown
Consanguineous	3/10 (30%)	2/10 (20%)	1/10 (10%)	1/10 (10%)	3/10 (30%)
Non-consanguineous	8/35 (23%)	6/35 (17%)	1/35 (3%)	9/35 (26%)	11/35 (31%)

'unrelated' parents who originally come from the same village in South Asia may share similar haplotypes and therefore be at risk of recessive disease. There have been concerns that the health problems exhibited by socially and economically disadvantaged migrant groups are

blamed on consanguinity⁴ without understanding the nature of inheritance pattern of the phenotype. The assumption that consanguinity automatically entails a high level of autosomal recessive disease was assessed in this study.



The sample size is relatively small in this paper; larger studies have been conducted in South India, where consanguinity is endemic. A population-based cross-sectional epidemiological study from Andhra Pradesh has found that of 10 290 children with consanguineous parents, only 80 had a possible ocular genetic disorder. The inheritance pattern of the ocular genetic disorders was not discussed. Another study from Chennai demonstrated that among 430 consanguineous families with retinitis pigmentosa, 39% of the offspring had definite autosomal recessive disease.

One criticism of our paper is that the sporadic cases of ocular genetic disorder could in fact be autosomal recessive in nature. This is difficult to prove, as sporadic cases may also represent *de novo* autosomal dominant mutations, X-linked recessive mutations (in boys), and mitochondrial inheritance. Only large empirical studies will allow us to differentiate between these inheritance patterns for phenotypes that have more than one inheritance pattern.

Another drawback of this paper is that the prevalence of consanguinity in families without ocular genetic disorders was not analysed. It is likely that consanguinity is high in the South Asian community (specifically Bangladeshi population) and in those families who have children without ocular genetic disorders.

This paper confirms the findings of previous studies that have shown that consanguinity *per se* does not translate to high levels of autosomal recessive disorders. It is important that clinicians who discuss children with ocular genetic disorders with their families do not assume that the phenotype is autosomal recessive unless the condition is characterised as such. This is because the use of assumptions can have devastating effects on the

family and may be incorrect. This study has not assessed the age of parents⁷ nor the number of offspring and both these factors may contribute to the burden of ocular genetic disorders in the most deprived area of London.

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