A STUDY OF HAEMOGLOBINOPATHIES AND HAEMOGLOBIN VARIANTS USING HIGH PERFORMANCE LIQUID CHROMATOGRAPHY (HPLC) IN A TEACHING HOSPITAL OF ODISHA

Sarojini Raman¹, Nageswar Sahu², Urmila Senapati³

¹Assistant Professor, Department of Pathology, Kalinga Institute of Medical Sciences (KIMS), KIIT University, Bhubaneswar, Odisha. ²Assistant Professor, Department of Pathology, Kalinga Institute of Medical Sciences (KIMS), KIIT University, Bhubaneswar, Odisha. ³Professor and HOD, Department of Pathology, Kalinga Institute of Medical Sciences (KIMS), KIIT University, Bhubaneswar, Odisha.

ABSTRACT

BACKGROUND

Thalassaemias and haemoglobinopathies are highly prevalent in India. Identification of these disorders is important for epidemiologic purposes and for prevention of thalassaemia major and clinically severe haemoglobinopathies. The use of high performance liquid chromatography (HPLC) as a screening method for detection of these groups of disorder is increasing in last two decades.

The aim is to study of thalassaemias and haemoglobinopathies using high performance liquid chromatography as a diagnostic tool in patients of a tertiary care hospital of Odisha.

METHODS AND MATERIALS

A total of 788 cases were included in the study. Samples were analysed on the BIO-RAD D-10TM dual mode HPLC system. Personal history, family history, peripheral blood findings and sickling test result were correlated. Family studies were done whenever required and possible. However, secondary confirmatory tests were not done. Statistical analysis was done on Microsoft Excel. Continuous variables were expressed as mean ± SD. Categorical variables were expressed as frequencies and percentages.

RESULTS

Abnormal HPLC patterns were seen in 37.18% of cases. Sickle cell heterozygous (15.10%), sickle cell homozygous (9.90%) and β -thalassaemia trait (6.10%) were the most common abnormalities found. Other patterns detected include β -thalassaemia major and intermedia, compound heterozygous state of HbS and β -thalassaemia, HbE trait, HbE disease, double heterozygous state of HbS and HbE, HbE and β -thalassaemia and HbS and hereditary persistent of foetal haemoglobin, HbD Punjab trait, Hb Lepore trait and HbH disease.

CONCLUSION

Sickle cell along with β -thalassaemias are the major abnormal haemoglobins in Odisha. Premarital and antenatal screenings are important measures to prevent birth of children with severe haemoglobin disorders. HPLC is a rapid and reliable technique for identification of various haemoglobin fractions.

KEYWORDS

Haemoglobinopathies, Haemoglobin Variants, High Performance Liquid Chromatography, Prevalence.

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BACKGROUND

Haemoglobinopathies and thalassaemias are the most common single gene disorders in the world.^[1] The prevalence of thalassaemias and haemoglobinopathies varies with geographic locations.^[2] World Health Organization figures estimate that 5% of the world population are carriers of a potentially pathological haemoglobin gene.^[3] The general incidence of thalassaemia trait and sickle cell haemoglobinopathy in India varies between 3-17% and 1-44% respectively.

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Corresponding Author:
Dr. Nageswar Sahu,
Assistant Professor,
Department of Pathology,
KIMS & PBMH, KIIT University,
Patia, Bhubaneswar-751024, Odisha.
E-mail: nageswar.sahu@yahoo.in
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But because of consanguinity and caste and area endogamy, some communities show a very high incidence, making this group of diseases a major public health problem in our country.^[4,5,6]

Diagnosis of haemoglobinopathies in most centres in India relies upon conventional methods like, clinical and family history, haemoglobin concentration, red cell indices, HbA2, HbF estimation, sickling test, and Hb electrophoresis. Various limitations of these methods have been felt in recent years. One of the most important is the difficulty in the identification of Hb variants with same electrophoretic mobility, such as in A2/E/C/O-Arab and S/D/G/Q/Lepore. Other issues come up in diagnosis of HbS traits where low quantity of HbS is associated with negative sickling test and while diagnosing certain compound heterozygous states such as, for HbD and HbE, HbS and β -thalassaemia, HbS and HbD.[7,8]

High performance liquid chromatography (HPLC) being an automated instrument is highly sensitive and specific, has high resolution and helps in quantification of various haemoglobins.

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It acts as a good alternative to electrophoresis.^[9,10] With the help of basic haematological parameters like Hb% and RBC indices along with HPLC, a laboratory can identify around 75% of the common variants encountered, without the need for confirmatory studies such as alkaline and acid electrophoresis.^[11]

The knowledge of the common Hb variants encountered in a particular area is important for the formulation of specific diagnostic, preventive and therapeutic strategies. As the exact data pertaining to the prevalence of haemoglobinopathies in this region is scarce, we considered it important to find out the extent of burden of the common Hb disorders in patients of a tertiary care hospital of Odisha.

MATERIALS AND METHODS

This was a prospective study conducted over a period of 2 years. All the cases received in our laboratory for haemoglobin HPLC were considered which included patients opting for premarital or antenatal haemoglobin analysis, patients having positive family history and cases suspected of haemoglobinopathy on the basis of red cell indices. However, patients with a history of blood transfusion within the last 1 month and infants with inconclusive HPLC findings were excluded. A total of 788 cases were included in the study. The geographical distribution included whole Odisha, along with cases from West Bengal, Chhattisgarh and Jharkhand.

A detailed clinical history, family history and history of blood transfusion were obtained from each patient. Blood samples were collected in vacutainer tubes containing dipotassium EDTA and analysed in Beckman Coulter (LH-750) automated cell counter for complete blood count. For each patient, a peripheral blood smear (PBS) was prepared and stained with Leishman stain. Keeping in mind that HbS prevalence is high in Odisha^[4,12] sickling test (Using sodium meta-bisulphite) was done in all the samples.

HPLC was performed on BIO-RAD D- 10^{TM} dual mode instrument manufactured by Bio-Rad laboratories, which utilizes the principle of cation-exchange high performance liquid chromatography (CE-HPLC). Before every run, the instrument was calibrated with the help of D- 10^{TM} HbA₂/F/A_{1c} calibrator and two levels of controls. The sample

were injected into the analysis stream and separated by cation exchange cartridge using a phosphate ion gradient generated by mixing two buffers of different ionic strengths elute the different haemoglobins. The separated haemoglobin fractions pass through a flow cell, where absorbance is measured at 415 nm with simultaneous use of secondary wavelength of 690 nm to reduce background noise. A printed chromatogram is delivered by the software. where the different peaks are identified in different windows with relevant information like retention time, relative percentage and area. The total area acceptable was between one to four million. Haemoglobins were identified on the basis of retention time. Quantification of the haemoglobin was done by determining the area under the corresponding peak in the elution profile. If a peak eluted at a retention time that is not predefined, it was labelled as unknown. In these cases the identification of Hb variant was done by manufacturer assigned retention time in the manual.

Diagnosis was made by seeing the chromatogram finding in correlation with age, Hb%, RBC indices, PBS findings and the result of sickling test. The presence of HbH was confirmed by using brilliant cresyl blue test for HbH inclusions. A second confirmatory method like Hb electrophoresis at alkaline pH, isoelectric focusing and molecular studies were not done though recommended to many patients.

Statistics

Statistical analysis was done on Microsoft Excel. Continuous variables were expressed as mean±SD. Categorical variables were expressed as frequencies and percentages.

RESULTS

Out of 788 cases 495 (62.82%) were normal and 293(37.18%) cases showed abnormal haemoglobin fractions. Male to female ratio in the abnormal group was 4:3. Age group ranged from 6 months to 84 years with a mean of 19.18 years and median of 17 years. Normal chromatogram showed primarily HbA, a small percentage of A_2 (<3.9%) and traces of HbF. P2 and P3 are normal associates of HbA. Table-1&2 shows the number, age group affected, Hb%, RBC indices and chromatogram finding of all abnormal cases.

Diagnosis	No. of	Age (Years)			Hb(g/dL) TRBC(106/μL)		MCV(fl)	MCH (pg)	MCHC (g/dL)	RDW-CV(%)	
	Cases (%)	Range	Mean	Median	Mean±SD	Mean±SD	Mean±SD	Mean± SD	Mean±SD	Mean±SD	
AS	119 (15.1)	0.5-84	19.61	15	10.14±2.59	4.31±1.14	76.87±10.56	23.86±3.85	30.94±1.39	18.14±5.55	
SS	78 (9.90)	0.7-48	15.32	12.5	7.81±1.89	3.02±0.87	84.14±12.00	26.50±4.43	31.46±3.04	20.44±3.57	
BTT	48 (6.10)	2-64	28.22	27	10.19±2.00	5.12±1.11	66.23±5.69	19.74±1.85	29.85±1.00	17.98±2.72	
BTI	01 (0.12)	7			5.8	2.88	83.6	24.6	29.4	41.9	
BTM	07 (0.89)	0.6-1	0.7	0.6	5.2±0.9	2.04±0.36	74.36±5.46	25.71±3.2	33.28±2.99	32.31±3.96	
Sβ	22 (2.79)	1-46	17.77	12.5	7.11±2.11	3.28±0.92	73.30±7.38	22.23±2.54	30.31±0.73	22.84±4.13	
AE	06 (0.76)	1-21	10.66	11.5	9.76±3.03	4.52±1.28	71.25±9.81	21.55±2.95	30.27±0.46	17.98±5.75	
EE	01 (0.12)	16			11.8	5.73	64.7	20.6	31.9	15.9	
Εβ	05 (0.63)	1-53	13.8	5	5.35±0.9	2.50±0.56	66.96±8.07	20.18±2.95	30.11±0.98	29.45±6.13	
SE	01 (0.12)	27		10.3	3.07	69.2	23.4	33.8	14.2		
SF	02 (0.25)	28, 40		9.2, 9.9	3.3, 3.4	84.2, 91.7	27.9, 28.8	31.4, 33.1	14.4, 19.5		
AD	01 (0.12)	24		9.9	4.3	92.2	32.4	33.8	16.8		
HbH disease	01 (0.12)	41		6.9	3.17	78.8	21.7	27.6	24.7		
Lepore trait	01 (0.12)	24			6.3	2.09	63.8	19.1	29.9	22.1	
Table 1. Number, Age and RBC indices of all abnormal cases											

Hb=Haemoglobin, TRBC=Total RBC count, MCV=Mean corpuscular volume, MCH=Mean corpuscular haemoglobin, MCHC=Mean corpuscular haemoglobin concentration, RDW=Red cell distribution width, SD=Standard deviation

Diagnosis	A ₀ ([%]	A ₂ /E	(%)	F	(%)	S (%)		Other (%)		
	Range	Mean±SD	Range	Mean±SD	Range	Mean±SD	Range	Mean±SD	(RT in Minutes)		
AS	32.5-77.5	59.7± 6.98	1.6-4.9	3.23±0.63	0.0-18.5	2.43±3.60	13.2-58.9	26.50±6.83			
SS	2.2-28.4	6.38±4.17	1.1-4.9	2.77±0.85	7.9-45.2	22.03±8.36	37.4-82.3	64.7±9.41			
BTT	75.8-85.8	81.7±2.17	4.1-8.2	5.03±0.85	0.0-6.4	1.28±1.36					
BTI	6.	6	1.	8		81.7					
BTM	1.6-22.6	10.67±8.11	0.6-3.3	1.95±0.95	61.8-85.9	76.25±8.55					
Sβ	3.9-24.7	11.06±5.71	4.5-8.2	5.93±0.99	3.0-30.0	13.89±8.20	29.6-81.0	64.86±11.82			
AE	61.8-73.4	66.0±4.31	20.6-33.4	27.1±4.1	0.0-1.2	0.63±0.51					
EE	6.	3	100	0.1	5.2				6.9(3.78)		
Εβ	5.0-29.9	15.28±9.95	39.8-88.0	60.9±21.3	17.1-37.3	26.26±9.2					
SE	6.3		46	.3		5.3	50.8				
SF	2.8-5.6	4.2±1.97	1.4-1.9	1.65±0.35	26.7-35.2	30.95±6.01	50.4-61.0	55.7±7.49			
AD	52	9	3.	2	1.0				33.3(3.88)		
HbH disease	66	.2	0.	8	0.0				15.2(0.39)		
Lepore trait	75	.9	17	7.2	0.8						
Table 2. Haemoglobin Fractions of all Abnormal Cases											

AS=Sickle cell heterozygous, SS=Sickle cell homozygous, BTT= β -thalassaemia trait, BTI= β -thalassaemia intermedia, BTM= β -thalassaemia major, S β =Compound heterozygous for Hb S and β -thalassaemia, AE=Hb E heterozygous, EE=Hb E homozygous, E β =Compound heterozygous for Hb E and β -thalassaemia, SE=Compound heterozygous for Hb S and Hb E, SF=Compound heterozygous for Hb S and hereditary persistent of foetal haemoglobin, AD=Hb D Punjab heterozygous, RT=Retention time.

In our study, most common abnormal haemoglobin fraction observed was HbS, seen in 222(28.17%) patients. Sickling test was positive in all of these cases. Hb concentration in sickle cell heterozygous (AS) group ranged from 2.6-14.86 g/dL. Anaemia (<11 g/dL) was seen in 68 (57.14%) patients. Red cell morphology was mostly normocytic normochromic and microcytic hypochromic. HbF was mostly normal, with 13(10.92%) patients having a raised HbF (>5.0%). In sickle cell homozygous (SS) group Hb ranged from 2.9-12.1 g/dL. Only four (5.1%) patients had Hb>11 g/dL. Most of the patients had either normocytic normochromic or microcytic hypochromic blood picture with anisopoikilocytosis [Fig-1a].

All patients of compound heterozygous for HbS and β -thalassaemia (S β) were anaemic (Hb=3.5-10.22 g/dL). PBS showed moderate to marked anisopoikilocytosis [Fig-1b]. Two adults with mild anaemia were diagnosed to have compound heterozygous for HbS and hereditary persistent of foetal haemoglobin (SF) [Fig-1c]. Parenteral study of one case confirmed the diagnosis, while the other case parents did not turn around. The single case of compound heterozygous for HbS and HbE (SE) had mild anaemia with microcytosis. Both HbS and HbE (eluted in A_2 window) were around 50% [Fig-1d].

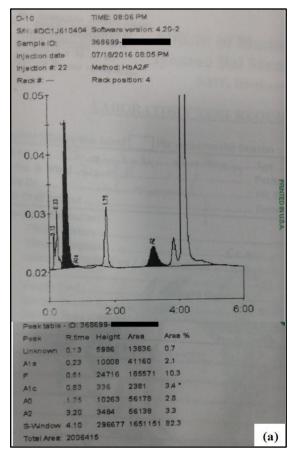
HbA $_2$ levels of 4-9% are diagnostic of β -thalassaemia trait (BTT) in an asymptomatic individual with no or mild anaemia. We got this range of increased HbA $_2$ in 55 patients without any other significant abnormality in chromatogram.

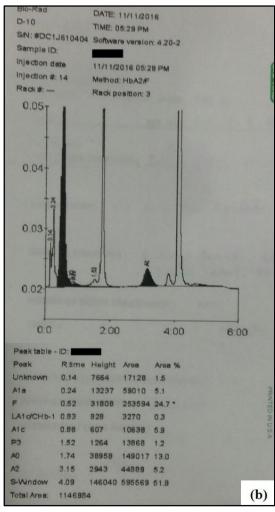
Most of them had microcytic hypochromic blood picture. However, 9 patients were suspected to have either megaloblastic or dimorphic anaemia. HPLC was repeated in 7 of them after nutritional supplements. Repeat HbA_2 was normal in 5 of them and were included in normal category. Two cases with persistent raised HbA2 (>4%) were diagnosed as BTT. Two patients lost to follow up, were excluded from the study. Total 48 patients were diagnosed as BTT [Fig-2a]. 11 (22.91%) patients had Hb < 9 g%.

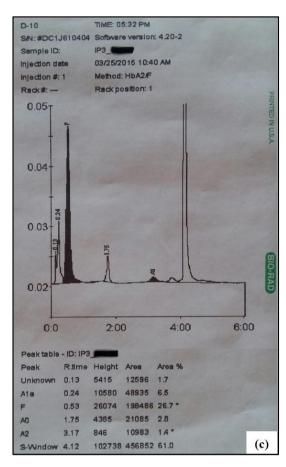
Seven cases were diagnosed as β -thalassaemia major (BTM) [Fig-2b]. All presented in their 1^{st} year of life with Hb level \leq 6.5 g%. PBS showed marked degree of anisopoikilocytosis with raised RDW, hypochromia, target cells, polychromasia and NRBC. HbF levels were high. One patient had similar presentation at the age of 7 years with 81.7% HbF was diagnosed as β -thalassaemia intermedia (BTI) on the basis of clinical data.

HbE elutes in the A_2 window. We got six cases of HbE heterozygous (AE) [Fig-2c], one case of HbE homozygous (EE) [Fig-2d] and five cases of compound heterozygous for HbE and β -thalassaemia (E β) [Fig-3a]. All except one AE cases had microcytic hypochromic picture. The EE case was advised HPLC based on severe microcytic hypochromic blood picture with normal Hb level. All E β patients were of paediatric age group except one adult case. Hb value was 4.0-6.13 g%. The PBS showed anisopoikilocytosis with microcytosis, hypochromia and target cells.

One case having mild normocytic normochromic anaemia showed decreased A₀ with an unknown peak of 33.3% at retention time of 3.88 minutes [Fig-3b]. We diagnosed it as HbD Punjab heterozygous (AD) based on the retention time. One adult patient having an unknown peak of 15.2% in the 1st minute (RT=0.39 minutes) was diagnosed as HbH disease [Fig-3c]. Presence of HbH inclusion on brilliant cresyl blue stain confirmed the diagnosis. Hb Lepore trait constituted one case. HbA₂ was raised to 17.2% with microcytic hypochromic anaemia [Fig-3d].







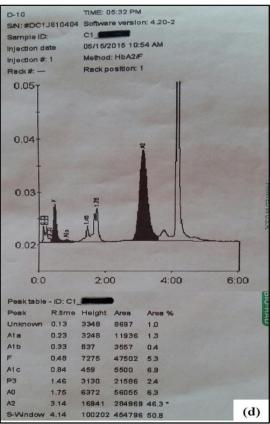
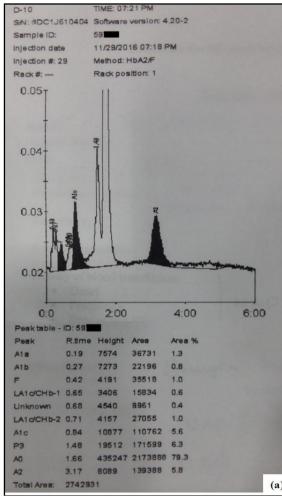
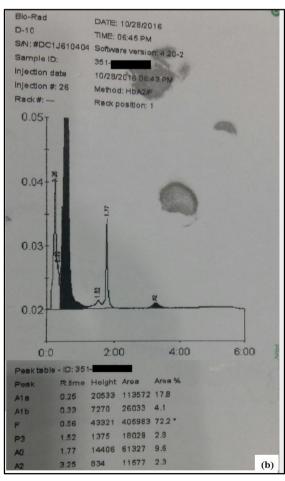
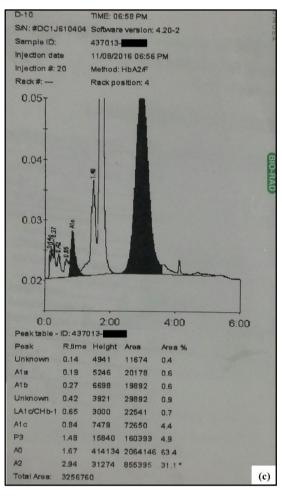


Figure 1. Chromatogram of (a) Sickle Cell Homozygous (SS), (b) Compound Heterozygous for HbS and β-Thalassaemia (Sβ), (c) Compound Heterozygous for HbS and Hereditary Persistent of Foetal Haemoglobin (SF), (d) Compound Heterozygous for HbS and HbE (SE)







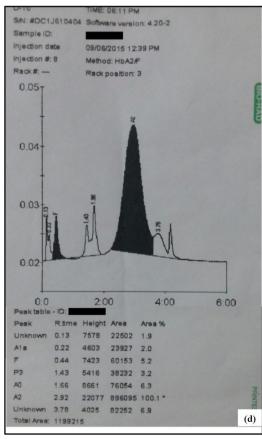
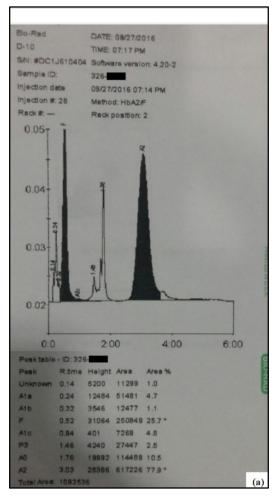
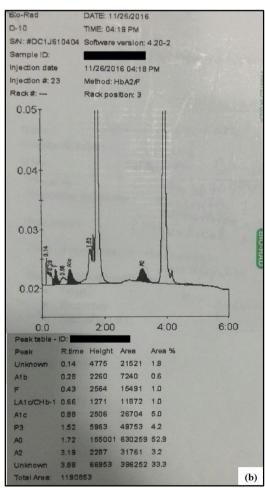
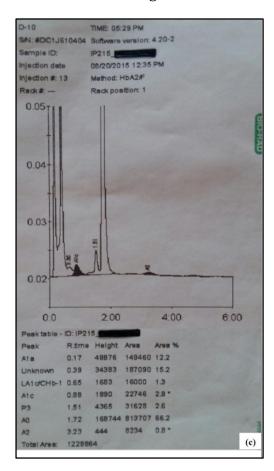


Figure 2. Chromatogram of (a) β-thalassaemia Trait (BTT), (b) β-thalassaemia Major (BTM), (c) HbE Heterozygous (AE), (d) HbE Homozygous (EE)







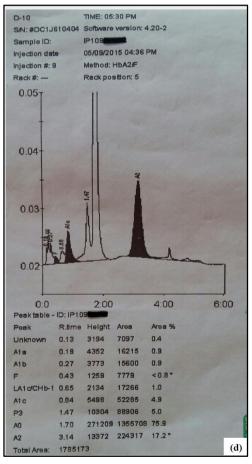


Figure 3. Chromatogram of (a) Compound Heterozygous for HbE and β-thalassaemia (Eβ), (b) HbD Punjab Heterozygous, (c) HbH Disease, (d) Hb Lepore Trait

DISCUSSION

In the present study, the prevalence of Hb disorders was found to be 37.26%. Previous institutional based studies from Odisha reported higher prevalence (44.2% and 65.7%) of Hb disorder. [4,12] However, there is a wide range of prevalence of Hb disorder in different parts of India. Madan et al reported a prevalence of 11.43% in a study from West Bengal. [2] Two hospital based studies from Western India got 8.6% and 11.36% of abnormal haemoglobin variants. [7,13] In the North Indian population, incidence of haemoglobinopathies was found to be 12.5%. [14] Baruah et al noted a high prevalence (59.11%) in Nort-East region. [1]

HbS was detected in 28.17% patients, which was lower than reported by Balgir et al $(39.3\%)^{[4]}$ but was similar to that got by Dash et al $(29.0\%)^{[12]}$ from Odisha. Most common abnormal Hb pattern detected in this study was AS (15.10%) followed by SS (9.90%) and BTT (6.10%). The two above said studies from Odisha also got AS as the most common variant followed by BTT and SS. $^{[4,12]}$ However, most of the other studies across the country reported BTT as the most common variant. $^{[2,7,8,13,14,15,16]}$ In spite of these variations, the high incidence of both AS and BTT highlights the need for antenatal screening for prevention of more severe form like BTM, SS and Sβ in offspring.

The microcytic hypochromic blood picture in AS group could be due to associated iron deficiency. High incidence of iron deficiency has been reported in patients with sickle cell disease from India.^[5,17] The finding of a raised HbF in some SCT patients was difficult to explain.

Many sickle cell homozygous patients had HbF >25%. In Indian subcontinent HbSS patients have slightly higher HbF level, than the other parts of the world. The reason for this is that the haplotype of HbS gene, which is prevalent in India, is the Saudi Arabia/Indian haplotype, that reduces the clinical severity of the disease.[7,18]

BTT was the 3rd most common abnormal haemoglobin variant we got. Most of the cases had characteristic microcytic hypochromic red cells with normal or slightly reduced Hb and raised RBC count. Raised A2 level is the most important abnormal chromatogram finding helpful for its diagnosis. However, conditions with borderline HbA2 need careful interpretation. Nutritional anaemia must always be taken into account. A low level of HbA2 may be induced by iron deficiency. Similarly, cobalamine or folate deficiency may raise HbA2 level. However, in previous studies no significant difference was found in HbA2 level in patients of BTT with or without concomitant iron deficiency. Thus elevation in HbA2 level can be used with reliability for diagnosis of BTT even in the presence of iron deficiency.[10,19] In our study, the peripheral red cell morphology was of help in cases of borderline elevation of HbA2. In cases of cobalamine and folate deficiency HPLC was repeated after nutritional supplement whenever feasible. Cases with normal A₂ level on repeat HPLC were taken as normal and cases still having raised A2 were diagnosed as BTT. Similarly, milder forms of thalassaemia or a co-inheritance of delta thalassaemia may also lead to borderline A2 levels. Genetic studies should be advised in cases of dilemma for a conclusive opinion.[14]

BTM was seen in only 0.89% of patients. This low incidence of the disease may be due to decrease incidence of the disease due to effective prenatal diagnosis.

Two patients had received blood transfusion more than 1 month before diagnosis. Thalassaemia intermedia is suspected when a patient presents after 3 years of age or needs fewer blood transfusion.^[17] One of our patients presented like major at the age of 7 years and diagnosed as BTI.

HbE is the most frequent variant Hb in Asia, with a significant prevalence in North-East India and Bangladesh. It is a β-chain variant that tends to elute in A_2 window on HPLC. In our study, the prevalence of HbE gene was 1.63% which is slightly lower than the finding observed by Balgir et al (1.90%) in a study from Odisha. HbE homozygous usually presents with HbE values >70-75% and heterozygous with HbE values <40%. Clinical effects are more severe when HbE is coinherited with β-thalassaemia (Eβ). Γ7.14 All Eβ cases in our study had severe anaemia (Hb≤6.13 g%).

Though cases of HPFH heterozygous cases have been reported, we could not find much literature about SF. We diagnosed two cases of adult onset mild anaemia as SF based on raised HbF and HbS with normal HbA₂. The age of presentation and degree of anaemia helped to exclude SS. Parenteral study confirmed the diagnosis in one. Molecular study was advised in the other case for confirmation. The prevalence of SE in our study was 0.12% as compared to 0.004% in a study of more than 90,000 cases from West Bengal.^[2]

Other variants detected in the present study included one case each of HbD-Punjab trait, Hb Lepore and HbH.

The gene frequency of HbD Punjab is relatively low with highest prevalence in North-Western India. Balgir et al[4] got only two cases (0.2%) of HbD Punjab trait in a series of more than 1000 cases. We also got only one case (0.12%) indicating a very low prevalence of HbD in Odisha. On CE-HPLC, it gets eluted in the D-window, which is separate from HbS peak. Homozygous HbD disease usually presents with mild haemolytic anaemia and mild to moderate splenomegaly. Heterozygous HbD is a clinically silent condition, unless coinherited with HbS.[20] HPLC is useful to distinguish between HbD-Iran and HbD-Punjab. HbD-Iran gets eluted in A2 window and HbD-Punjab in the D window. Both the situations are clinically important because HbD-Punjab produces a significant sickling disorder when coinherited with HbS but with HbD-Iran, the combination is clinically benign.[7,8,21]

Like HbD-Iran Hb Lepore also elutes in A_2 window. However, the HbA $_2$ % in HbD-Iran (40-50%) is significantly more than that of Hb Lepore (10-18%).[11] A sharp peak in the first minute of elution indicates HbH. HbH disease shows considerable variability in clinical and haematological severity.[22]

Due to socio-cultural practices, marriages in India are usually among individuals of the same caste or ethnic group and this makes it important to know the prevalence of different abnormal Hb variants in different ethnic groups. [23] In our country premarital screening is still considered taboo. So the best approach would be to target those patients attending the haematology OPD, the antenatal population and extended family members. The couple at risk should be counselled regarding the nature of the disease and the implications of being carriers. [13,16]

CONCLUSION

In India, where β -thalassaemia trait is so rampant, premarital and antenatal screening should be mandatory to prevent birth of children with β-thalassaemia major. Moreover, in Odisha where HbS is so common, the occurrence of sickle cell homozygous and compound heterozygous like Sβ or SE can be prevented by these screening protocols. HPLC is an excellent and powerful diagnostic tool for direct identification of Hb variants. The simplicity and rapidity of sample preparation, accurate quantification of Hb concentration combined with complete automation, makes HPLC an ideal methodology for the routine diagnosis of Hb disorders. Hb variants that are eluted in the same window can be differentiated according to the retention time and the percentage of different haemoglobins obtained. However, the chromatograms must be interpreted only in the light of other relevant investigations, family studies and if necessary molecular studies.

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