

List of Abnormal Hemoglobins Recorded in the World

Susumu SHIBATA and Satoshi UEDA
Department of Medicine, Yamaguchi University
School of Medicine, Ube, Japan
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Hemoglobinopathy is a congenital abnormality of hemoglobin synthesis, in which the α or the non- α polypeptide chains of abnormal amino acid sequence are produced owing to the mutational change of the structural gene of the hemoglobin polypeptides. Hereditary suppression of the production of the α or the non- α polypeptide chains due to mutation of their operator or regulator gene, namely thalassemia, is also included into the category of hemoglobinopathy, although it is not accompanied by abnormal hemoglobins.

Pauling and his associates are credited for the pioneer work of exploring the vast area of hemoglobinopathy. In 1949¹⁾ they discovered a slow-moving hemoglobin, which was named Hb S later, in the hemolysate of a patient with sickle cell anemia by moving boundary electrophoresis, and established that Hb S which is different from the normal adult hemoglobin in molecular structure was responsible for "sickling" and clinical manifestations of this disease. They coined the word "molecular disease" for sickle cell anemia in order to characterize it properly. Sickle cell anemia was the first instance of hemoglobinopathy. Pauling's discovery gave a great impetus to the survey of hemoglobinopathy in the world, and within a few years a considerable number of electrophoretically abnormal hemoglobins, such as Hbs C, D, E, etc., were reported from various countries.

Schroeder and his associates²⁾ demonstrated two different sorts of polypeptides, the α and the β chains, in the molecule of hemoglobin. In 1958 Ingram³⁾ invented an ingenious method called finger-printing for the analysis of polypeptides, which enabled him to reveal the characteristic abnormality in Hb S molecule, i.e. the substitution of valine for the glutamic acid at the sixth residue of the β chain. The abnormal β chain of Hb S, namely the β^S chain, is discriminated from the β^A chain of normal adult hemoglobin (Hb A) only by this amino acid substitution. The remainder of the amino acid sequence is entirely the same between the β^S and the β^A chains. However, the presence of valine in the place of glutamic acid residue (β 6) leads to build an intramolecular ring (Val-His-Leu-Thr-Pro-Val) between the sixth amino acid residues and the N-terminal end of the β^S chain. In deoxygenated state a pair of these rings of a Hb S molecule are fitted to a pair of pockets belonging to the α chains of another Hb S molecule. In this way the connection of deoxygenated Hb S molecule extends endlessly to form a filament. Several (six) of the filaments snuggle together to form a bundle and the

bundles thus produced create crystal-like structure called "tactoid" which is extremely insoluble. The tactoids push and deform the stroma of an erythrocyte from inside to give rise to fish-fin or foliate spicules on the surface. This is the phenomenon of "sickling" which is characteristic of the erythrocytes containing Hb S.⁴⁾

The basic study on the chemistry of hemoglobinopathy was almost completely accomplished in 1965, by which year the amino acid sequences had been established for all of the polypeptide chains composing the molecules of the normal human hemoglobins, namely the α^A ⁵⁾, the β^A ⁵⁾, the γ^F ⁶⁾ and the δ^{A2} ⁷⁾ chains. The erythrocytes of the normal adults contain three kinds of hemoglobins, (1) Hb A = $\alpha_2^A \beta_2^A$, the major component occupying about 95 per cent, (2) Hb F = $\alpha_2^A \gamma_2^F$, a minor component which is alkali-resistant, comprising about 1 per cent and (3) Hb A₂ = $\alpha_2^A \delta_2^{A2}$, a slow moving hemoglobin forming about 2.5 per cent of the total hemoglobin respectively, in the foetal period Hb F makes the principal component of hemoglobins.

As many as 199 sorts of abnormal hemoglobins have hitherto been recorded in the world. They are listed in Table I. Most of them are not associated with clinical symptoms (symptomless hemoglobinopathies). However, some abnormal hemoglobins, which are represented by Hb S, give rise to diseases showing characteristic clinical manifestations. They are classified into the following major groups: ... (1) Sick cell hemoglobins, (2) unstable hemoglobins producing congenital Heinz body hemolytic anemia, (3) hemoglobin M's causing congenital cyanosis, (4) hemoglobins with low oxygen affinity, and (5) hemoglobins with high oxygen affinity entailing polycythemia.

1. Sick cell hemoglobins. This is represented by Hb S (β 6 Glu \rightarrow Val). Sick cell hemoglobins are listed in Table II. They give positive result to sickling test of blood drop with reducing agent (2 per cent sodium metabisulfite solution). Hb I was once said to cause sickling, but recent study revealed it to be an artifact⁸⁾.

2. Unstable hemoglobins. These hemoglobins are so unstable that they undergo irreversible oxidation and denaturation spontaneously even while they are contained in living erythrocytes, yielding intraerythrocytic inclusion bodies (Heinz bodies). Occasionally denaturation becomes manifest only when drugs (sulfonamide) which enhance the oxidation process in the erythrocytes are given to the carrier. Hb Zürich (β 63 His \rightarrow Arg) is such an example. The hemoglobins are precipitated when they are warmed at 50–60°C. Accordingly, heat denaturation test of the hemolysate containing these abnormal hemoglobins are used for their detection. A few hemoglobins belonging to this group such as Hb Hammersmith, Hb Torino, Hb Mie, Hb Kanazawa and Hb Chiba are not demonstrable by electrophoresis. Table III presents the list of unstable hemoglobins. It is worth while to mention that almost all the unstable hemoglobins have β chain anomaly.

3. Hb M's. Hb M is the oldest example of abnormal hemoglobins that have

ever been detected from the Caucasians and the Asian races. They were for the first time reported from Germany by Hörlein and Weber in 1948⁹⁾, a year before the discovery of Hb S by Pauling¹⁾ and his associates. Hb M's are characterized by their chocolate brown color reminiscent of methemoglobin. So, they are demonstrable by the spectroscopic examination of the hemolysates (of O₂ Hb and metHb types). Irreversible oxidation of heme iron as a result of the amino acid substitution in the α or β chains at a residue which is in intimate relation to heme is responsible for the causation of this peculiar color. It is thought that half a moiety of the molecule of this hemoglobin relevant to the abnormal polypeptide chains (the α or the β) is methemoglobinized. Substitution of tyrosine for the proximal (α 87, β 92) or the distal (α 58, β 63) histidine is commonly found in Hb M's. Agar gel or starch block electrophoresis (pH 7.0) of the methemoglobin type hemolysate is convenient for the demonstration of the Hb M's. Hb M's are listed in Table IV.

4. Hemoglobins with low O₂ affinity. Hb Kansas is the representative of this group (Table V). Like Hb M's it causes cyanosis. However, another hemoglobin belonging to this group, i.e. Hb E is not related to cyanosis.

5. Hemoglobins with high O₂ affinity. The molecules of these hemoglobins are difficult in releasing O₂ when they are exposed to lowered oxygen tension. Presumably the amino acid substitution in the abnormal chains seen in these hemoglobins result in altered conformation of the α or β subunit pair which hampers their interrelated to-and-fro shift inherent in oxygenation and deoxygenation of hemoglobin molecule. Chronic oxygen deficit in the tissues will stimulate production of erythropoietin, thus causing polycythemia. Hemoglobins with high O₂ affinity are seen in Table VI.

Theoretically 2151 sorts of abnormal hemoglobins are expected from the consideration of possible amino acid substitution on the α or the β chains.¹⁵⁾ Only 687 kinds of these are demonstrable by electrophoresis which is, at present, the most useful measure for their detection. The remaining variants are thought to be buried until new procedures except electrophoresis which enable us to disclose electrically inert abnormality of hemoglobins, because the amino acid substitutions in their abnormal polypeptide chains are of the one which does not entail the positive or negative change in electrical charge. Perhaps, heat denaturation test of the hemolysate may be mentioned as one of such procedures, although it has only a limited usefulness. Further efforts should be made along this line to promote the study of abnormal hemoglobins.

Table I. List of known hemoglobin substitutions and deletions

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
A ₂ (B ₂) ¹⁰⁾	delta	16	A 13	Gly→Ala
A ₂ Babinga ¹¹⁾	delta	136	H 14	Gly→Asp
A ₂ Flatbush ¹²⁾	delta	22	B 4	Ala→Glu
A ₂ N.Y.U. ¹³⁾	delta	12	A 9	Asn→Lys
A ₂ Sphakia ¹⁴⁾	delta	2	NA 2	His→Arg
Abiko ¹⁵⁾	beta			
Aegina ¹⁶⁾	gamma ?			
Agenogi = Izumo ¹⁷⁾	beta	90	F 6	Glu→Lys
Alexandra ¹⁸⁾	gamma ?			
Ann Arbor ¹⁹⁾	alpha	80	F 1	Leu→Arg
Atago ¹⁹⁹⁾	alpha	85	F 6	Asp→Tyr
Augusta I ²⁰⁾	beta			β_4^S (tetramer)
Augusta II ²⁰⁾	beta			β_4^C (tetramer)
Bart's ²¹⁾				γ_4 (tetramer)
Bibba ²²⁾	alpha	136	H 19	Leu→Pro
Boras ²³⁾	beta	88	F 4	Leu→Arg
Broussais ²⁴⁾	alpha	90	FG 2	Lys→Asn
C ²⁵⁾	beta	6	A 3	Glu→Lys
C Georgetown ²⁶⁾	beta	6 core	A 3	Glu→Val
C Harlem ²⁷⁾	beta	6 73	A 3 E 17	Glu→Val Asp→Asn
Cardeza ²⁸⁾	alpha			
Caserto ²⁹⁾	beta			
Chad ³⁰⁾	alpha	23	B 4	Glu→Lys
Chesapeake ³¹⁾	alpha	92	FG 4	Arg→Leu
Chiapas ³²⁾	alpha	114	GH 2	Pro→Arg
Chiba ³³⁾				
Chicago I ³⁴⁾				
Cyprus I ³⁵⁾	gamma			
D Bushman ³⁶⁾	beta	16	A 13	Gly→Arg
D Frankfurt ³⁷⁾	beta			
D Hollywood ³⁸⁾	beta			
D Ibadan ³⁹⁾	beta	87	F 3	Thr→Lys
D Punjab ⁴⁰⁾ =D Caucasian =D Chicago =D Cyprus =D Los Angeles =D North Carolina =D Portugal	beta	121	GH 4	Glu→Gln

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
Dakar ⁴¹⁾	alpha	112	G 19	His→Gln
Dhofar ⁴²⁾	beta	58	E 2	Pro→Arg
Douglas et al ⁴³⁾	beta			
E ⁴⁴⁾	beta	26	B 8	Glu→Lys
E Saskatoon ⁴⁵⁾	beta	22	B 4	Glu→Lys
Etobicoke ⁴⁶⁾	alpha	84	F 5	Ser→Arg
F Akashi ¹⁵⁾	gamma			
F Alexandra ¹⁹⁾	gamma	12	A 9	Thr→Lys
F Houston ⁴⁷⁾	gamma			
F Hull ⁴⁸⁾	gamma	121	GH 4	Glu→Lys
F Jamaica ²⁰⁰⁾	gamma	61	E 5	Lys→Glu
	gamma	136	H 14	Ala
F Roma ⁴⁹⁾	gamma			
F Texas I ⁵⁰⁾ = F Galveston	gamma	5	A 2	Glu→Lys
F Texas II ⁵¹⁾	gamma	6	A 3	Glu→Lys
Fessas-Papaspyrou ⁵²⁾ = Bart's				γ_4 (tetramer)
Freiburg ⁵³⁾	beta	23	B 5	Val deleted
Fukuoka ⁵⁴⁾				
G Accra ⁵⁵⁾	beta	79	EF 3	Asp→Asn
G Audhali ⁵⁶⁾	alpha	23	B 4	Glu→Val
G Baltimore ⁵⁷⁾	alpha	Tp-9		Asp→His
G Chinese ⁵⁸⁾ = G Hong Kong = G Honolulu = G Shingapore	alpha	30	B 11	Glu→Gln
G Copenhagen ⁵⁹⁾	beta	47	CD 6	Asp→Asn
G Coughatta ⁶⁰⁾ = G Saskatoon = G Hsin-Chu	beta	22	B 4	Glu→Ala
G Galveston ⁶¹⁾ = G Port Arther = G Texas	beta	43	CD 2	Glu→Ala
G Georgia ²⁰¹⁾	alpha	95	G 2	Pro→Leu
G Ibadan ⁶²⁾	alpha			
G Makassar ²⁰²⁾	beta	6	A 3	Glu→Ala
G Norfolk ⁶³⁾	alpha	85	F 6	Asp→Asn
G Paris ⁶⁴⁾	alpha	Tp-7		
G Philadelphia ⁶⁵⁾ = D Azuokoli = D Baltimore = D St. Lous = D Washington = G Bristol = Stanleyville I	alpha	68	E 17	Asn→Lys

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
G San Jose ⁶⁶⁾	beta	7	A 4	Glu→Gly
G Taegu ⁶⁷⁾				
G Taichung ²⁰⁸⁾ =Mahidol	alpha	74	EF 3	Asp→His
G Taipei ⁶⁸⁾	beta	22	B 4	Glu→Gly
G Taiwan Ami ⁶⁹⁾	beta	25	B 7	Gly→Arg
Genova ⁷⁰⁾	beta	28	B 10	Leu→Pro
Gifu ⁷¹⁾ =G Szuhu ⁷²⁾ =Ichnomiya	beta	80	EF 4	Asn→Lys
Gower I ⁷³⁾	epsilon			ϵ_4 (tetramer)
Gower II ⁷⁴⁾				$\alpha_2 \epsilon_2$
Gun Hill ⁷⁵⁾	beta	91-95 or 92-96 or 93-97	F7-FG2 or F8-FG3 or F9-FG4	(Leu, His, Cys, Asp, Lys) deletion
H ⁷⁶⁾				β_4 (tetramer)
Hammersmith ⁷⁷⁾	beta	42	CD 1	Phe→Ser
Hasharon ⁷⁸⁾	alpha	47	CD 6	Asp→His
Hikairi ⁷⁹⁾	beta	61	E 5	Lys→Asn
Hirose ⁸⁰⁾	beta	37	C 3	Try→Ser
Hiroshima ⁸¹⁾	beta	146	H 21	His→Asp
Hijiyama ⁸²⁾	beta	120	GH 3	Lys→Gln
Hofu ⁸³⁾	beta	126	H 4	Val→Glu
Honolulu ⁸⁴⁾				
Hope ⁸⁵⁾	beta	136	H 14	Gly→Asp
Hopkins I ⁸⁶⁾	beta			
Hopkins II ⁸⁷⁾	alpha			
I ⁸⁸⁾	alpha	16	A 14	Lys→Glu
I Burlington ⁸⁹⁾	alpha			
I Toulouse ⁹⁰⁾	beta	66	E 10	Lys→Glu
Ichinomiya ⁹¹⁾ =Gifu	beta	80	EF 4	Asn→Lys
J ⁹²⁾ =Mexico	alpha	54	E 3	Gln→Glu
J Baltimore ⁹³⁾ =New Heaven No. 2 =J Trinidad =J Ireland	beta	16	A 13	Gly→Asp
J Bangkok ⁹⁴⁾ =J Korat =J Meining	beta	56	D 7	Gly→Asp
J Cambridge ⁹⁵⁾	beta	69	E 13	Gly→Asp
J Cape Town ⁹⁶⁾	alpha	92	FG 4	Arg→Gln

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
J Georgia ⁹⁶⁾	beta			
J India ⁹⁷⁾	alpha			
J Iran ⁹⁸⁾	beta	77	EF 1	His→Asp
J Jamaica ³⁸⁾	beta			
J Kaohsiung ²⁰⁴⁾	beta	59	E 3	Lys→Thr
J Korat ⁹⁹⁾	beta	56	D 7	Gly→Asp
J Malaya ⁹⁷⁾	alpha			
J Medellin ¹⁰⁰⁾	alpha	22	B 3	Gly→Asp
J Meining ⁹⁹⁾	beta	56	D 7	Gly→Asp
J Oxford ¹⁰¹⁾ =I Interlaken	alpha	15	A 13	Gly→Asp
J Paris ¹⁰²⁾	alpha	12	A 10	Ala→Asp
J Sardegna ¹⁰³⁾	alpha	50	CD 8	His→Asp
J Taichung ¹⁹⁸⁾	beta	129	H 7	Ala→Asp
J Tongariki ¹⁰⁴⁾	alpha	115	GH 3	Ala→Asp
J Toronto ¹⁰⁵⁾	alpha	5	A 3	Ala→Asp
K Calcutta ⁹⁷⁾	alpha			
K Cameroon ¹⁰⁶⁾	beta	Tp-8		Ala→Glu or Asp
K Ibadan ¹⁰⁶⁾	beta	46	CD 5	Gly→Glu
K Liberia ³⁸⁾	beta			
K Madras ³⁸⁾	alpha			
K Woolwich ¹⁰⁶⁾	beta	132	H 10	Lys→Gln
Kanazawa ¹⁰⁷⁾				
Kansas ¹⁰⁸⁾	beta	102	G 4	Asn→Thr
Kempsey ¹⁰⁹⁾	beta	99	FG 6	Asp→Asn
Khartoum ¹¹⁰⁾	beta	124	H 2	Pro→Arg
Koelliker ¹¹¹⁾	alpha	141	HC 3	Arg deleted
Köln ¹¹²⁾	beta	98	FG 5	Val→Met
Kokura ¹¹³⁾ =L Ferrara =Beilinson =Umi =Tagawa II =Yukuhashi II =L Gaslini	alpha	47	CD 6	Asp→Gly
Korle Bu ¹¹⁴⁾	beta	73	E 17	Asp→Asn
Kumamoto ¹¹⁵⁾	alpha			
L ¹¹⁶⁾	beta			
L Bombay ¹¹⁷⁾	alpha	Tp-8,9		
L Ferrara ¹¹⁸⁾ =Kokura	alpha	47	CD 6	Asp→Gly
L Persian Gulf ²⁰⁵⁾	alpha	57	E 6	Gly→Arg

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
Leiden ¹¹⁹⁾	beta	6 or 7	A3 or A4	Glu deleted
Lepore Boston ¹²⁰⁾				$\alpha_2(\delta-\delta)_2$
Lepore Hollandia ¹²¹⁾				$\alpha_2(\delta-\beta)_2$
M Boston ¹²²⁾ (Osaka ¹²³⁾ =M Gothenberg =M Kiskunhalas ¹²⁴⁾ =M Leipzig II =M Norin =M Osaka	alpha	58	E 7	His→Tyr
M Hyde Park ¹²⁵⁾ (Akita ¹²⁶⁾ =M Akita	beta	92	F 8	His→Tyr
M Iwate ¹²⁷⁾ (Kankakee ¹²⁸⁾ =M Kankakee =M Oldenburg	alpha	87	F 8	His→Tyr
M Leipzig I ¹²⁹⁾				
M Leipzig II ¹³⁰⁾				
M Milwaukee I ¹²²⁾ =M Mineapolis	beta	67	E 11	Val→Glu
M Milwaukee II ¹²²⁾	beta			
M Reserve ¹³¹⁾	alpha			
M Saskatoon ¹²²⁾ (Kurume ¹³²⁾ =M Chicago =M Emory =M Elberfeld =M Hamburg =M Kurume =M Radom =M Arhus =M Yonago	beta	63	E 7	His→Tyr
Malmö ²¹⁰⁾	bata	97	FG 4	His→Gln
Manitoba ¹³³⁾	alpha	102	G 9	Ser→Arg
Matsue ¹³⁴⁾	alpha	Tp-9		
Memphis ¹³⁵⁾	alpha	23	B 4	Glu→Gln
Mexico ³²⁾ =J	alpha	54	E 3	Gln→Glu
Mie ¹³⁶⁾				
Miyada ¹¹⁵⁾				$\alpha_2(\beta-\delta)_2$
N Baltimore ¹³⁷⁾ =N Memphis =N New Haven =Jenkins =Kenwood	beta	95	FG 2	Lys→Glu
N New Haven ¹³⁸⁾	bata	95	FG 2	Lys→Glu
N Seattle ¹³⁹⁾	beta	61	E 5	Lys→Glu
Nagasaki ¹⁵⁾	beta	17	A 14	Lys→Glu
New York ¹⁴⁰⁾	beta	113	G 15	Val→Glu
Nishiki IV ¹⁴¹⁾ =Waziro ?	beta	Tp-3		

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
Norfolk ¹⁴²⁾ =Kagoshima =Nishiki I, II, III	alpha	57	E 6	Gly→Asp
O Arab ¹⁴³⁾ =O Bulgaria =O New York	beta	121	GH 4	Glu→Lys
O Indonesia ¹⁴³⁾	alpha	116	GH 4	Glu→Lys
Oak Ridge ¹⁴⁴⁾	beta	94	FG 1	Asp→Asn
Okayama ¹⁵⁾	alpha			
Oki ¹⁵⁾	alpha			
Olmsted ^{145, 210)}	beta	141	H 19	Leu→Arg
Omori ¹⁵⁾	beta	6 or 7	A3 or A4	Glu→Gly
Otsu ¹⁴⁶⁾ =E	beta	26	B 8	Glu→Lys
P ¹⁴⁷⁾	beta	117	G 19	His→Arg
Philly ¹⁴⁸⁾	beta	35	C 1	Tyr→Phe
Picardy ¹⁴⁹⁾				
Port Alegre ¹⁵⁰⁾	beta	9	A 6	Ser→Cys
Portland I ¹⁵¹⁾				$\gamma_2 \chi_2$
Pylos ¹⁵²⁾				$\alpha_2(\delta-\beta)_2$
Q Chinese ¹⁵³⁾	alpha			
Q Iran ²¹¹⁾	alpha	75	EF 4	Asp→His
Q Thailand ²¹²⁾	alpha	74	EF 3	Asp→His
R ¹⁵⁴⁾ =Durham No. 1	beta	Tp-1		
Rainier ¹⁵⁵⁾	beta	145	HC 2	Tyr→Cys
Rambam ¹⁵⁶⁾	beta	69 or 74	E13 or E18	Gly→Asp
Riverdale-Bronx ¹⁵⁷⁾	beta	24	B 6	Gly→Arg
Russ ¹⁵⁸⁾	alpha	51	CD 9	Gly→Arg
S ¹⁵⁹⁾	beta	6	A 3	Glu→Val
Santa Ana ¹⁶⁰⁾	beta	88	F 4	Leu→Pro
Sabine ¹⁶¹⁾	beta	91	F 7	Leu→Pro
Salt Lake City ¹⁶²⁾				
Sawara ²⁰⁶⁾	alpha	6	A 4	Asp→Ala
Sealy ¹⁶³⁾ =Sinai =Hasharon	alpha	47	CD 6	Asp→His
Seattle ¹⁶⁴⁾	beta	76	E 20	Ala→Glu
Shepherds Bush ²⁰⁷⁾	beta	74	E 18	Gly→Asp
Shimonoseki ^{165,166)} =Hikoshima =Mobara	alpha	54	E 3	Gln→Arg

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
Singapore ¹¹⁰⁾	alpha	141	HC 3	Arg→Pro
Singapore-Bristol ¹⁶⁷⁾	gamma			
Siriraj ¹⁶⁸⁾	beta	7	A 4	Glu→Lys
Sogn ¹⁶⁹⁾	beta	14	A 11	Leu→Arg
Stanleyville II ¹⁷⁰⁾	alpha	78	EF 7	Asn→Lys
St. Mary ¹⁷¹⁾	beta	core ?		
Sud-Vietnam ¹⁷²⁾				
Sydney ¹⁷³⁾	beta	67	E 11	Val→Ala
Tacoma ¹³³⁾	beta	30	B 12	Arg→Ser
Tagawa I ¹¹⁵⁾	alpha	Tp-10		
Ta-Li ²⁰⁸⁾	beta	83	EF 7	Gly→Cys
Tochigi ¹⁷⁴⁾	beta	56-59	D7-E3	deleted
Tokuchi ¹¹⁵⁾	beta			
Tokyo ¹¹⁵⁾	beta			
Torino ¹⁷⁵⁾	alpha	43	CD 1	Phe→Val
Tsukiji ¹⁷⁶⁾	beta			
Ube I ¹⁷⁷⁾	beta	93	F 9	Cys SH blocked
Ube II ¹⁷⁸⁾	alpha	68	E 17	Asn→Asp
Warren ¹⁷⁹⁾	gamma			
Wajiro ¹⁸⁰⁾	alpha			
Wien ¹³³⁾	beta	130	H 8	Tyr→Asp
Yakima ¹⁸¹⁾	beta	99	FG 6	Asp→His
Yatsushiro ¹⁵⁾	beta			
Yoshizuka ¹⁵⁾	beta	108	G 10	Asn→Asp
Ypsilanti ¹⁸²⁾				
Yukhashi ¹⁵⁾	beta	58	E 2	Pro→Arg
Zambia ²⁰⁹⁾	alpha	60	E 9	Lys→Asn
Zurich ¹⁸³⁾	beta	63	E 7	His→Arg

Table II. List of sickling hemoglobins

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
S ¹⁵⁹⁾	beta	6	A 3	Glu→Val
C Georgetown ²⁶⁾	beta	6 core	A 3	Glu→Val
C Harlem ²⁷⁾	beta	6	A 3	Glu→Val
		73	E 17	Asp→Asn
G Philadelphia/S ¹⁸⁴⁾	alpha	68	E 17	Asn→Lys
	beta	6	A 3	Glu→Val

Memphis/S ¹⁸⁵)	alpha	23	B	4	Glu→Gln
	beta	6	A	3	Glu→Val
Stanleyville II/S ¹⁸⁶)	alpha	78	EF	7	Asn→Lys
	beta	6	A	3	Glu→Val

Table III. List of unstable hemoglobins

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
Bibba ²²)	alpha	136	H 19	Leu→Pro
Chiba ³³)				
Genova ⁷⁰)	beta	28	B 10	Leu→Pro
Gun Hill ⁷⁵)	beta	91-95 or 92-96 or 93-97	F7-FG2 or F8-FG3 or F9-FG4	(Leu, His, Cys, Asp, Lys) deleted
Hammersmith ⁷⁷)	beta	42	CD 1	Phe→Ser
Kansas ¹⁰⁸)	beta	102	G 4	Asn→Thr
Köln ¹¹²)	beta	98	FG 5	Val→Met
L Ferrara ¹¹⁸)	alpha	47	CD 6	Asp→Gly
Mie ¹³⁶)				
Olmsted ¹⁴⁵)				
Philly ¹⁴⁸)	beta	35	C 1	Tyr→Phe
Sabine ¹⁶¹)	beta	91	F 7	Leu→Pro
Salt Lake City ¹⁶²)				
Santa Ana ¹⁶⁰)	beta	88	F 4	Leu→Pro
Seattle ¹⁶⁴)	beta	76	E 20	Ala→Glu
Sealy ¹⁶³)	alpha	47	CD 6	Asp→His
Shepherds Bush ²⁰⁷)	beta	74	E 18	Gly→Asp
Sogn ¹⁶⁹)	beta	14	A 11	Leu→Arg
St. Mary's ¹⁷¹)	beta	core ?		
Sydney ¹⁷³)	beta	67	E 11	Val→Ala
Tacoma ¹³³)	beta	30	B 12	Arg→Ser
Tochigi ¹⁷⁴)	beta	56-59	D7-E3	deleted
Torino ¹⁷⁵)	alpha	43	CD 1	Phe→Val
Ube 1 ¹⁷⁷)	beta	93	F 9	Cys SH blocked
Wien ¹³³)	beta	130	H 8	Tyr→Asp
Zurich ¹⁸³)	beta	63	E 7	His→Arg

Table IV. List of hemoglobin M disgenes

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
M Boston ¹²²⁾	alpha	58	E 7	His→Tyr
M Hyde Park ¹²⁵⁾	beta	92	F 8	His→Tyr
M Iwate ¹²⁷⁾	alpha	87	F 8	His→Tyr
M Milwaukee ¹²²⁾	beta	67	E 11	Val→Glu
M Saskatoon ¹²²⁾	beta	63	E 7	His→Tyr

Table V. List of hemoglobins with low oxygen affinity

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
E ^{44, 187)}	beta	26	B 8	Glu→Lys
Kansas ¹⁰⁸⁾	beta	102	G 4	Asn→Thr

(Hb M's of α chain anomaly will also be classified into this group.)²⁰³⁾

Table VI. List of hemoglobins with high oxygen affinity

Name	Chain anomaly	Residue no.	Helical no.	Amino acid substitution
A ₂ ^{7, 188)}				
A' ₂ (B ₂) ^{10, 188)}	delta	16	A 13	Gly→Ala
Bart's ^{21, 189)}				γ_4 (tetramer)
C Harlem ²⁷⁾	beta	6 73	A 3 E 17	Glu→Val Asp→Asn
Chesapeake ^{31, 190)}	alpha	92	FG 4	Arg→Leu
D Punjab ^{40, 191)}	beta	121	GH 4	Glu→Gln
Freiburg ⁵³⁾	beta	23	B 5	Val deleted
Gun Hill ⁷⁵⁾	beta	91-95 or 92-96 or 93-97	F7-FG2 or F8-FG3 or F9-FG4	(Leu, His, Cys, Asp, Lys) deleted
H ^{76, 192)}				β_4 (tetramer)
Hiroshima ⁸¹⁾	beta	143	H 21	His→Asp
J Cape Town ^{95, 193)}	alpha	92	FG 4	Arg→Gln
Kempsey ¹⁰⁹⁾	beta	99	FG 6	Asp→Asn
Köln ^{112, 194)}	beta	98	FG 5	Val→Met
Lepore ^{120, 121, 195)}				$\alpha_2(\delta-\beta)_2$
*M Hyde Park ^{125, 126, 196)}	beta	92	F 8	His→Tyr
Rainier ¹⁵⁵⁾	beta	145	HC 2	Tyr→His
Tacoma ¹³³⁾	beta	30	B 12	Arg→Ser
Yakima ^{181, 197)}	beta	99	FG 6	Asp→His
Zurich ¹⁸³⁾	beta	63	E 7	His→Arg

*Hb M's of β chain anomaly are inferior to normal adult hemoglobin in the capacity of transporting oxygen in spite of slightly increased oxygen affinity of their normal α chain, because their abnormal β chains can not carry oxygen.

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