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Distributions of Genetic Markers in United States Populations: I. Blood Group and Secretor Systems

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ABSTRACT: All published and unpublished population frequency data that could be located for U.S. populations are tabulated and presented for the blood group and secretor systems. Results obtained by combining data for comparable racial/ethnic groups are also presented. The results obtained with combined data may give better information on frequencies for the U.S. population at large than is obtainable from studies conducted in restricted geographic areas.

KEYWORDS: forensic sciences, genetic typing, demography, population genetics, genetic markers, genotypic frequencies, phenotypic frequencies, ABO blood group system, Rh blood group system, MN blood group system, MNSs blood group system, Kell blood group system, Duffy blood group system, Kidd blood group system, Lewis blood group system, secretor system

Forensic serology has witnessed extraordinary growth and development in the past 25 years, reflective of the significant advances in immunology and human genetics. Important aspects of this progress have been the unraveling of many human polymorphic genetic marker systems and the discovery of new ones. Workers involved in disputed parentage testing now have a substantial number of genetic marker systems from which to construct routine testing protocols. The choice of systems having relatively high exclusion probabilities, especially the human leukocyte antigens (HLA) system, has enabled calculations of the probability of paternity in nonexclusion cases [1-3]. Moreover, a wider range of genetic marker systems than ever before is available to criminalists for the partial individualization of blood and physiological fluid stains [4-10]. To interpret and appreciate better the significance of typing results, both parentage and criminalistics applications of genetic marker testing often require knowledge of genotypic and phenotypic frequencies in applicable populations.

Thousands of frequency studies on various genetic marker systems have been carried out on many populations throughout the world, the most complete compilation of them being

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the extraordinary work by Mourant et al. [11]. A number of papers and studies have appeared which compile or analyze or both frequency data for many different genetic marker systems for the populations of entire countries, such as Japan [12-14], Ireland [15], and England [16-18]. Genetic marker frequency data from the many different studies of United States populations, however, have not to our knowledge been thoroughly compiled.

In this and two planned following papers, we summarize all the published and some unpublished population frequency data that could be located for U.S. populations for 22 genetic marker systems, along with the results obtained by combining data from different studies. In this paper, we summarize the sources and methods employed and report data for the classical blood group and secretor systems.

Methods

In connection with the preparation of a lengthy review book [10], all population frequency studies of U.S. populations published since 1950 that could be located were collected, in addition to data from papers presented at professional meetings and unpublished studies. The data from a number of additional published and unpublished studies that have since appeared are included in the present work as well.

Within the separate tables, each representing a different genetic marker system, data are tabulated separately for Caucasian, Negro, Hispanic, Chinese, and Asian populations. Each population studied is identified by location (city and state; two-letter abbreviations have been used for the states) and by any description provided by the original author(s). Each row (and in a few cases, several related rows) of each table represents a different study, and a reference is provided. References to population studies are given a "T" (for "Table") prefix in the tables, and are separately compiled at the end of the paper. Data for each phenotype within each system are reported using a $NNN(\%.%)$ format, where NNN represents the number of individuals who possessed the phenotype and $%.%$ represents the percentage rounded to one decimal place.

The total number of people studied is also given, and is not always the sum of the major phenotypes because rare phenotypes were observed. The notes in each table provide data for rarer types, unusual or descriptive features of a population, or explanations about the calculations.

Data from studies of the ABO system in which anti- A_1 was used, that is, the data in Table 3, were used to compute simple ABO frequencies which are included in Table 1. Similarly, data from studies of the MNSs system (Table 8) were used to compute data for the MN system and the computed data included in Table 7.

Where sufficient data were available, two types of calculations were used to combine all the data for a particular racial/ethnic class within a genetic marker system. Some authors provided the actual numbers of individuals possessing the different phenotypes, whereas others expressed the distributions in percentages. We calculated percentage values for the data sets that provided numbers of individuals. For data sets that provided percentages, only the percentages given by the original author(s) are tabulated. As a result, every data set has a set of percentage distributions, but some sets do not have numbers of individuals. The first calculation sums the numbers of individuals for all data sets showing numbers, and a percentage value for each phenotype is computed from the resulting totals. This result is referred to as the "numerical total." The second calculation was carried out on every group of data for a particular racial/ethnic group for which at least two independent percentage distributions were available. In the second calculation, the percentage distributions for each phenotype are weighted according to the number of individuals typed and what is referred to as the "weighted mean of proportions," or "WMP," is computed according to: $WMP = \sum np/N$, where n is the number of individuals in a particular data set, p is the corresponding phenotypic proportion, and N is the total number of people in the racial/ethnic group. A

weighted standard deviation of proportions (WSDP) is calculated for each WMP according to $\sqrt{\Sigma np^2/N - (\text{WMP})^2}$, where n , p , N , and WMP have the meanings defined above.

All available data were included in the tables for completeness, but in some circumstances a data set was not used in the calculations. These circumstances include cases in which: (a) one study included the data from another study by the same author(s); (b) a data set was incomplete for all phenotypes (as, for example, an Rh study in which anti-e was not employed and Rh₂Rh₂ was indistinguishable from Rh₂rh); and (c) one data set was provided for a group of individuals related to those in another data set, such as mothers and their infant children.

In the Kell, Duffy, and Kidd data, separate calculations were carried out on data from studies in which only one of the antisera was used in testing.

Gene frequencies were calculated for data sets in which there was sufficient information to enable the calculation. Those for the ABO and A₁A₂BO systems were calculated using the maximum likelihood methods of estimation described by Stevens [19]. The method applicable to ABO was also used for calculation of Fy^a, Fy^b, and Fy in populations that showed four Duffy types. Rh and MNSs gene frequencies were calculated by the methods described by Mourant et al. [11], which give estimates nearly as good as true maximum likelihood methods [20, 21]. Gene frequencies for two-allele three-phenotype systems (MN, some Kell, Kidd, and Duffy data) were calculated by gene counting. Gene frequencies for two-allele two-phenotype systems (Kell, using only anti-K; Duffy, using only anti-Fy^a; Kidd, using only anti-Jk^a; Secretor) were estimated by the square root method. This method assumes that all K⁻, Fy(a⁻), and Jk(a⁻) people are, respectively, kk , Fy^bFy^b, and Jk^bJk^b homozygotes, and that nonsecretors are *sese*. The square root of the phenotypic distribution of the assumed homozygotes provides an estimate of the frequency of one gene. An estimate of the other is obtained by difference from unity. No gene frequency calculations were attempted for the Lewis system, as the data are for red cell Lewis types [11].

A chi-square value was calculated for every data set for which gene frequencies could be meaningfully calculated and for the corresponding numerical totals. Chi-square values were not calculated for data sets in which gene frequencies were estimated by the square root method. It is common practice in human genetic studies to regard data with some concern if χ^2 values corresponding to $P < 0.05$ are obtained. Observed frequencies in such cases are said to represent "significant" deviations from expectation under Hardy Weinberg equilibrium assumptions. Data yielding χ^2 values corresponding to $P < 0.01$ are said to represent "highly significant" deviations. Chi-square values must be interpreted in context, however. In some data sets, there are phenotypic classes with small numbers and small differences between observed and expected values can yield disproportionately large contributions to the chi-square statistic. In the tables where gene frequency data are presented (except in those where gene frequencies were obtained by the square root method, as noted above), data sets having χ^2 values corresponding to $P < 0.05$ or $P < 0.01$ are indicated.

In the Rh system, tabulation and WMP calculations were carried out for seven major phenotypes. Less common phenotypes are reported in the table notes. Phenotypes having C^w or D^u were not included with the corresponding major phenotypes (that is, Rh₁^wRh₁ were not included with Rh₁Rh₁, and so forth) but are reported in table notes. For data sets containing sufficient data, frequencies were estimated for all the Rh genes except r^y, and in these cases, all the phenotypes except those having r^y, C^w, or D^u had to be taken into account. In a few data sets, the original authors' gene frequencies are cited. These are indicated in table notes, and in those cases C^w or D^u types may be included within the major Rh phenotypes. In some cases, frequencies of the four major genes were estimated from data for the seven major phenotypes.

Most computations were carried out on a Data General MV 8000 mainframe computer with programs written in FORTRAN.

Results and Discussion

In Tables 1 through 14 are reported the phenotypic distribution and, where applicable, the estimates of gene frequencies for the ABO, A_1A_2BO , Rh, MN, MNSs, Kell, Duffy, Kidd, Lewis, and Secretor systems, respectively. Tables 2, 4, 6, and 9 show the gene frequency estimates for applicable data in Tables 1, 3, 5, and 8, respectively.

In addition to tabulating all the data, phenotypic proportions for all individuals typed as well as a WMP were calculated for each phenotype within each system for each major racial/ethnic group as described under **Methods**. Populations are defined by the size of the effective interbreeding gene pool rather than by geographical boundaries, especially in a country like the United States where mobility is very high. According to this reasoning, which we have discussed elsewhere [22,23], it might be useful in certain circumstances to have frequency estimates for larger and presumably better randomized samples of the population at large than would result from local population studies. Computation of WMP for systems in which a number of different studies have been done and in which fairly large numbers of people have been typed provides a possible approach to obtaining such an estimate.

The results suggest that the approach may be a useful one. Although there are no large studies on randomly sampled individuals throughout the United States in every racial/ethnic group against which to compare the WMP values, a recent study has been carried out on a completely ascertained, randomly selected, country-wide population of Caucasian and Negro teenagers [T30]. Table 15 shows a comparison between WMPs obtained by the method in the present work with the values obtained in the U.S. study. The WMP values used for comparison in Table 15 were obtained from all available data except that the data from the U.S. study was omitted. If the U.S. data are regarded as "expected" and the WMP data as "observed" in a chi-square goodness-of-fit test, only one system shows a significant deviation, the A_1A_2BO set for Negroes. The greatest contribution to the chi-square statistic in this instance is the A_2 class, which is very small. Although only a few of the blood group systems could be compared with the U.S. study, results obtained with those which could indicate that WMP may give a reasonably good estimate of the phenotypic proportions based upon the population data which has been gathered.

Our expectation that the aggregation of appropriate data for a given ethnic/racial category for a particular system would yield larger, better randomized data sets for gene frequency calculations was not altogether realized, at least to the extent that in some cases the numerical total values showed noteworthy deviations from expectation. In some cases, the results are understandable because of the small numbers in certain phenotypic classes. In other cases, there were not very many population studies amenable to aggregation. Another factor to be considered is the difficulty of defining ethnic/racial groups which can properly be regarded as belonging to an effective interbreeding gene pool.

In the studies on the ABO system for which gene frequencies were calculated (Table 2), several data sets and the Caucasian and Hispanic totals showed deviations from expectation based on Hardy Weinberg equilibrium assumptions. In every case, however, the greatest contribution to the chi-square statistic results from Group AB, the smallest class. Similarly, in the A_1A_2BO studies (Table 4) data sets showing significant deviations from expectation were accounted for by the AB phenotypes. A number of Rh data sets (Table 6) did not show goodness of fit on the basis of chi-square, which was computed separately for the major types and for all types. MN data (Table 7) including the aggregate totals showed good fit for the most part. Highly significant deviations from expectation are seen in two studies.

Caucasian data sets for MNSs (Table 9) were mostly within expectation. The Negro data sets, however, uniformly showed highly significant deviations from expectation. The largest Hispanic data set showed highly significant deviations, and so did the Hispanic total as a result. Kell (Table 10) three-phenotype data, including the totals, showed good fit through-

TABLE 1—Phenotypic frequencies of ABO blood groups in U.S. populations.

Population	Frequency—Number (Percent)					Reference
	Total	O	A	B	AB	
CAUCASIAN						
New York, NY	3 268	1 470(45.0)	1 199(36.7)	425(13.0)	166(5.1)	T1
Minnesota	300	(40.0)	(45.7)	(11.7)	(2.6)	T2
Iowa City, IA	49 979	22 392(44.8)	21 144(42.3)	4 695(9.4)	1748(3.5)	T3
Iowa City and Des Moines, IA	6 313	(45.8)	(41.6)	(9.0)	(3.6)	T4
Brooklyn, NY						
Leukemia patients						
Jewish	665	226(34.0)	261(39.2)	120(18.0)	58(8.7)	
Non-Jewish	639	291(45.5)	235(36.8)	90(14.1)	23(3.6)	
Blood donors						
Jewish	375	142(37.9)	165(44.0)	46(12.3)	22(5.9)	T5
Non-Jewish	548	281(51.3)	184(33.6)	62(11.3)	21(3.8)	
Seattle, WA						
University students	5 657	2 399(42.4)	2 458(43.5)	599(10.6)	201(3.6)	T6
St. Louis, MO						
Veteran's Hospital	359	169(47.1)	149(41.5)	29(8.1)	12(3.3)	
Control donors	32 945	14 918(45.3)	13 611(41.3)	3 248(9.9)	1168(3.5)	T7
Iowa City, IA						
Control donors	1 261	563(44.6)	512(40.6)	144(11.4)	42(3.3)	T8
New Haven, CT						
Students	1 000	431(43.1)	422(42.2)	110(11.0)	37(3.7)	T9
New York, NY						
Memorial Hospital						
Donors	4 738	2 029(42.8)	1 828(38.6)	636(13.4)	245(5.2)	
Transfused patients	2 332	976(41.9)	932(40.0)	291(12.5)	133(5.7)	
Tumor patients	525	192(36.6)	233(44.4)	70(13.3)	30(5.7)	T10
Southeastern GA	333	172(51.7)	136(40.8)	20(6.0)	5(1.5)	T11
Boston, MA						
Rheumatoid	608	276(45.4)	246(40.5)	55(9.0)	31(5.1)	
Nonrheumatoid	605	281(46.4)	225(37.2)	76(12.6)	23(3.8)	T12

TABLE 2—Gene frequencies for the ABO system in U.S. populations.

Population	Total Number Typed	Gene Frequency			χ^2	Reference to Population Study
		A	B	O		
CAUCASIAN						
New York, NY	3 268	0.2371	0.0950	0.6680		T1
Iowa City, IA	49 979	0.2638	0.0667	0.6695		T3
Brooklyn, NY						
Leukemia patients						
Jewish	665	0.2779	0.1438	0.5738		
Non-Jewish	639	0.2284	0.0929	0.6133		
Blood donors						
Jewish	375	0.2916	0.0951	0.6133		
Non-Jewish	548	0.2085	0.0787	0.7129		T5
Seattle, WA						
University students	5 657	0.2725	0.0735	0.6540	*	T6
St. Louis, MO						
Veteran's Hospital	359	0.2571	0.0588	0.6842		
Control donors	32 945	0.2575	0.0694	0.6731		T7
Iowa City, IA						
Controls	1 261	0.2517	0.0768	0.6714		T8
New Haven, CT						
Students	1 000	0.2648	0.0765	0.6587		T9
New York, NY						
Memorial Hospital						
Donors	4 738	0.2497	0.0976	0.6526		
Transfused patients	2 332	0.2622	0.0952	0.6426		
Tumor patients	525	0.2938	0.1003	0.6059		T10
Southeastern GA	333	0.2410	0.0383	0.7207		T11
Boston, MA						
Rheumatoid	608	0.2640	0.0738	0.6621		
Nonrheumatoid	605	0.2320	0.0855	0.6825		T12
Pittsburgh, PA						
School population	1 578	0.2625	0.1411	0.6234		
County fair sample	3 871	0.2677	0.0934	0.6389		
Male blood donors	1 959	0.2519	0.1009	0.6472	**	T13
Salt Lake City, UT						
College age	247	0.2645	0.0586	0.6769		T14
San Francisco Bay Area, CA						
Caucasians	8 962	0.2512	0.0727	0.6761	*	
"Caucasians of Western European Origin"	5 056	0.2471	0.0678	0.6851		T16
New York, NY	500	0.2471	0.1012	0.6666	**	T17
Denver, CO	3 648	0.2516	0.0648	0.6836		T18
Tecumseh, MI	8 965	0.2738	0.0658	0.6604		T20
Brooklyn, NY						
Kings County Hospital						
Adults	599	0.2442	0.0996	0.6562		
Newborns	253	0.2488	0.0762	0.6751		
Patients	68	0.2354	0.0613	0.7033		T21
Los Angeles, CA	205	0.2694	0.0981	0.6325		T22
Greater Philadelphia, PA including part of NJ	223	0.2367	0.1029	0.6605	**	T26
Miami/Dade County, FL	336	0.2006	0.0939	0.7055	*	T28
Los Angeles, CA						
Case material	419	0.2415	0.0748	0.6837	*	T29
CA, TX, HI, and Mexico City	914	0.2615	0.0741	0.6644	*	T31
Philadelphia, PA	396	0.2515	0.0844	0.6641		T32
Los Angeles County, CA						
Case material	516	0.2454	0.0749	0.6796	*	T33
North Carolina	773	0.2581	0.0601	0.6818		T34

TABLE 2—(Continued).

Population	Total Number Typed	Gene Frequency			χ^2	Reference to Population Study
		A	B	O		
Connecticut						
Case material	382	0.2578	0.0708	0.6714		T35
Total Caucasian	140 681	0.2605	0.0721	0.6674	*	
NEGRO						
Miami, FL	502	0.1523	0.1465	0.7012		T38
St. Louis, MO						
Veteran's Hospital	99	0.1474	0.1183	0.7343		
Control donors	1 395	0.1608	0.1257	0.7134		T7
Southeastern GA	300	0.1443	0.1135	0.7422		T11
San Francisco Bay Area, CA	3 146	0.1744	0.1248	0.7008		T16
Birmingham, AL						
School children	610	0.1665	0.1271	0.7063		T40
New York, NY	500	0.1569	0.1415	0.7016		T17
Brooklyn, NY						
Kings County Hospital						
Adults	1 150	0.1523	0.1396	0.7081		
Newborns	2 933	0.1678	0.1281	0.7042		
Patients	628	0.1648	0.1363	0.6953		T21
Greater Philadelphia, PA						
including part of NJ	176	0.1679	0.1443	0.6877		T26
Miami/Dade County, FL	346	0.1397	0.1413	0.7190		T28
Los Angeles, CA						
Case material	185	0.1884	0.1211	0.6904		T29
CA, TX, HI, and Mexico City	713	0.1659	0.1271	0.7072		T31
Philadelphia, PA	1 347	0.1710	0.1393	0.6897		T32
Los Angeles County, CA						
Case material	224	0.1959	0.1317	0.6723		T33
North Carolina	642	0.1586	0.1377	0.7037		T34
Connecticut						
Case material	67	0.1445	0.0775	0.7781		T35
Total Negro	14 963	0.1654	0.1306	0.7040	*	
HISPANIC						
Southern TX						
Mexican surnames	962	0.1676	0.0584	0.7735		T41
San Francisco Bay Area, CA						
"Mexicans"	335	0.1708	0.0904	0.7389		T16
Brooklyn, NY						
Kings County Hospital—Puerto Rican						
Adults	228	0.1915	0.0662	0.7432		
Newborns	928	0.1776	0.0668	0.7566		
Patients	248	0.1962	0.0818	0.7220		T21
Miami/Dade County, FL	357	0.1921	0.0518	0.7561		T28
Los Angeles, CA						
Case material	215	0.1593	0.0526	0.7881		T29
CA, TX, HI, and Mexico City	1 212	0.1315	0.0564	0.8120		T31
Los Angeles County, CA						
Case material	310	0.1671	0.0598	0.7731		T33
Total Hispanic	5 511	0.1648	0.0624	0.7728	*	
CHINESE						
New York, NY	400	0.1745	0.1639	0.6616		T17
New York, NY	946	0.1808	0.1724	0.6468		T43
Total Chinese	946	0.1808	0.1724	0.6468		
ASIAN						
CA, TX, HI, and Mexico City	1 342	0.2449	0.1705	0.5847	*	T31

Notes:

* $\chi^2 > 3.841$.

** $\chi^2 > 6.635$.

Detroit, MI	507	(53.1)	(16.0)	(0.2)	(26.2)	(2.8)	(1.8)	2,5	T25
Greater Philadelphia, PA including part of NJ	176	82(46.6)	41(23.3)	6(3.4)	40(22.7)	6(3.4)	1(0.6)	6	T26
U.S. national sample	999	(47.5)	(23.4)	(3.4)	(21.2)	(2.2)	(2.3)	2	T30
CA, TX, HI, and Mexico City Connecticut	713	359(50.4)	147(20.6)	37(5.2)	137(19.2)	20(2.8)	13(1.8)		T31
Case material	66	41(62.1)	14(21.2)	1(1.5)	8(12.1)	0(0.0)	2(3.0)	2	T35
Southeastern MO	79	(50.6)	(31.7)	(1.3)	(15.2)	(1.3)	(0.0)	2	T36
TOTAL NEGRO									
Numerical total	4 901	2 426(49.5)	955(19.5)	358(7.3)	958(19.5)	121(2.5)	76(1.6)		
WMP		49.5	19.9	6.2	20.3	2.5	1.5		
WSDP		2.013	2.394	2.572	2.120	0.589	0.519		
HISPANIC									
San Francisco Bay Area, CA	335	186(55.5)	74(22.1)	17(5.1)	44(13.1)	13(3.9)	1(0.3)		T16
CA, TX, HI, and Mexico City Connecticut	1 212	801(66.1)	239(19.7)	39(3.2)	173(14.3)	15(1.2)	5(0.4)		T31
Case material	21	12(57.1)	4(19.0)	2(9.5)	3(14.3)	0	0		T35
TOTAL HISPANIC									
Numerical total	1 568	999(63.7)	317(20.2)	58(3.7)	160(10.2)	28(1.8)	6(0.4)		
WMP		63.7	20.2	3.7	10.2	1.8	0.4		
WSDP		4.390	0.979	1.019	1.630	1.101	0.065		
CHINESE									
New York, NY	103	(45.6)	(27.2)	(0)	(22.3)	(4.9)	(0)	2	T37
Blood donors	817	(40.4)	(28.6)	(0)	(25.9)	(5.5)	(0)	2	T42
New York, NY	400	172(43.0)	108(27.0)	0	101(25.2)	19(4.8)	0		T17
TOTAL CHINESE									
Numerical total	400	172(43.0)	108(27.0)	0	101(25.2)	19(4.8)	0		
WMP		41.6	28.0	0	25.4	5.2	0		
WSDP		1.653	0.759	0	0.954	0.351	0		
ASIAN									
CA, TX, HI, and Mexico City	1 342	475(35.4)	442(32.9)	4(0.3)	287(21.4)	125(9.3)	9(0.7)		T31

Notes:

1. Six were A₁₂, 1 was A₃, and 1 was A₄.
2. Distributions given in percentages in original data; not used in calculating numerical totals.
3. Includes the 5056 "Caucasians of Western European Origin" in next row; only data for the 8962 used in calculations.
4. Two A₁ and 2 A₂ found in Caucasians.
5. Data of Stolow and collaborators.
6. Identical twin study; data for one member of each twin pair tabulated and used in calculations.
7. Four A₁, 2 A₂, and 1 A₃ found in Negroes.

TABLE 4—Gene frequencies for the A_1A_2BO system in U.S. populations.

Population	Total Number Typed	Gene Frequency			χ^2	Reference to Population Study
		A1	A2	B		
New York, NY	3 268	0.1819	0.0551	0.0950	0.6681	T1
Iowa City, IA						
Controls	1 261	0.2010	0.0509	0.0768	0.6713	T8
Southeastern GA	333	0.1653	0.0755	0.0384	0.7208	T11
Boston, MA						
Rheumatoid	608	0.1880	0.0725	0.0732	0.6663	
Nonrheumatoid	605	0.1710	0.0612	0.0854	0.6824	T12
San Francisco Bay Area, CA						
Caucasians	8 962	0.1903	0.0612	0.0729	0.6757	
"Caucasians of Western European Origin"						
New York, NY	5 056	0.1823	0.647	0.0679	0.6851	T16
Tecumseh, MI	500	0.1793	0.0525	0.1013	0.6670	T17
Los Angeles, CA	8 965	0.2039	0.0700	0.0658	0.6603	T20
Greater Philadelphia, PA	205	0.1778	0.0924	0.0979	0.6319	T22
including part of NJ	223	0.1674	0.0688	0.1028	0.6610	T26
CA, TX, HI, and Mexico City	914	0.1959	0.0655	0.0741	0.6646	T31
Connecticut	376	0.1987	0.0537	0.0691	0.6784	T35
Total Caucasian	26 220	0.1933	0.0635	0.0742	0.6690	

CAUCASIAN

TABLE 5—Phenotypic frequencies of Rh blood groups in U.S. populations.

Population	Total	Frequency—Number (Percent)								rh	Note	Reference
		Rh ₀	Rh ₁ rh	Rh ₁ Rh ₁	Rh ₂ rh	Rh ₂ Rh ₂	Rh ₁ Rh ₂	Rh ₁ Rh ₂ '	rh			
CAUCASIAN												
New York, NY	2 390	69(2.9)	798(33.4)	489(20.5)	349(14.6)	330(13.8)	321(13.4)				1,2,3	T1
Minnesota	300	(4.7)	(39.6)	(15.3)	(11.0)	(13.7)	(14.7)				1,2,4	T2
University of Iowa	2 181	91(4.2)	723(33.2)	382(17.5)	319(14.6)	303(13.9)	338(15.5)				1,5	T43
Southeastern GA	331	6(1.8)	112(33.8)	64(19.3)	31(9.4)	44(13.3)	46(13.9)				6	T11
San Francisco Bay Area, CA	4 928	(2.6)	(34.6)	(19.6)	(11.8)	(0)	(13.8)					
Mothers	4 928	(2.3)	(34.8)	(20.4)	(11.8)	(0)	(13.8)				2,7	T15
Children												
San Francisco Bay Area, CA	8 962	206(2.3)	3086(34.4)	1728(19.3)	1033(11.5)	212(2.4)	1189(13.3)	1365(15.2)				
Caucasians												
"Caucasians of Western	5 056	99(2.0)	1746(34.5)	912(18.0)	594(11.7)	109(2.2)	661(13.1)	853(16.9)			8	T16
European Origin"	500	11(2.2)	162(32.4)	111(22.2)	47(9.4)	9(1.8)	80(16.0)	60(12.0)			9	T17
New York, NY	1 412	(2.1)	(33.4)	(17.4)	(12.8)	(3.9)	(14.4)	(14.3)			2,10	T19
South Central WV	8 963	188(2.1)	3051(34.0)	1452(16.2)	1124(12.5)	208(2.3)	1124(12.5)	1412(15.8)			11	T20
Tecumseh, MI	505	(2.6)	(32.3)	(13.5)	(9.3)	(2.4)	(13.1)	(23.2)			2,12,13	T25
Detroit, MI												
Greater Philadelphia, PA	200	6(3.0)	74(37.0)	38(19.0)	29(14.5)	4(2.0)	26(13.0)	23(11.5)			14	T26
including part of NJ	370	10(2.7)	138(37.3)	64(17.3)	41(11.1)	12(3.2)	43(11.6)	51(13.8)			15,16	T28
Miami/Dade County, FL												
Los Angeles, CA	256	11(4.4)	80(32.0)	56(22.4)	24(9.6)	14(5.6)	34(13.6)	31(12.4)			15,17	T29
Case material	5 735	(2.4)	(35.4)	(17.6)	(12.1)	(3.1)	(14.2)	(14.1)			18	T30
U.S. national sample	914	37(4.1)	284(31.1)	150(16.4)	113(12.4)	13(1.4)	122(13.3)	165(18.1)			19	T31
CA, TX, HI, and Mexico City												
Connecticut	164	6(3.7)	50(30.5)	27(16.5)	14(8.5)	6(3.7)	35(21.3)	22(13.4)			20	T35
Case material												
TOTAL CAUCASIAN												
Numerical total	22 841	671(2.9)	7760(34.0)	4072(17.8)	2775(12.1)	491(2.1)	3000(13.1)	3513(15.4)				
WMP		2.8	34.2	18.0	12.1	2.1	13.4	15.0				
WSDP		2.372	0.994	1.547	1.000	1.166	0.880	1.420				

TABLE 5—(Continued).

Population	Total	Frequency—Number (Percent)							rh	Note	Reference
		Rh _o	Rh ₁ rh	Rh ₁ Rh ₁	Rh ₂ rh	Rh ₂ Rh ₂	Rh ₁ Rh ₂	Rh ₁ Rh ₂ ^a			
TOTAL HISPANIC											
Numerical total	2 042	54(2.6)	481(23.6)	488(23.9)	181(8.9)	113(5.5)	457(22.4)	107(5.2)			
WMP		2.6	23.6	23.9	8.9	5.5	22.4	5.2			
WSDP		2.050	7.214	4.448	1.765	1.870	5.157	4.391			
CHINESE											
New York, NY	103	(1.0)	(11.7)	(53.4)	(8.7)	(8.7)	(24.2)	(1.0)	1,2	T37	
New York, NY	400	1(0.3)	30(7.5)	213(53.2)	6(1.5)	19(4.8)	126(31.5)	0	35	T17	
New York, NY	946	3(0.3)	70(7.4)	506(53.5)	21(2.2)	42(4.4)	284(30.0)	2(0.2)	36	T43	
TOTAL CHINESE											
Numerical total	946	3(0.3)	70(7.4)	506(53.5)	21(2.2)	42(4.4)	284(30.0)	2(0.2)			
WMP		0.4	7.8	53.5	2.0	4.0	29.4	0.3			
WSDP		0.203	1.280	0.041	0.661	1.321	1.732	0.235			
ASIAN											
CA, TX, HI, and Mexico City	1 342	23(1.7)	112(8.3)	560(41.7)	46(3.4)	95(7.1)	466(34.7)	2(0.1)	37	T31	

^aIncludes Rh₁Rh₂, Rh₁rh^o, Rh₂rh^o, Rh₂Rh₂, Rh₂Rh_o, and Rh_orh_o.

Notes:

1. Anti-hr^o (anti-c) not used; Rh₂Rh₂ and Rh₂rh indistinguishable.
2. Data not complete for the major phenotypes or given in percentages not used in calculating numerical totals; data collected not using one or more of the five major antisera were not used in calculating WMPs.
3. 22(0.9) were rh^orh, 11(0.5) were rh^orh^o, and 1 was Rh₂Rh₂.
4. 0.3% were Rh₂Rh₁ and 0.7% were rh^orh.
5. 25(1.2) were rh^orh^o (or rh_orh).
6. One was Rh₂Rh₁, 1 was Rh₂Rh₂, 7 were rh^orh, and 2 were rh^orh^o; 3 of the 9 Rh_o had D^o, 2 of the 112 Rh₁rh had C^o, and 1 of the 65 Rh₁Rh₁ had C^o (Rh^o).
7. Mothers and their infants (children) studied; "mothers" phenotypic percentages used in calculating the WMPs. 4.3% children and 3.9% mothers among Caucasians and 2.8% children and 2.7% mothers among Negroes reported to be "other" Rh types.

8. 143(1.6) Caucasians, 82(1.6) "Caucasians of Western European Origin," 50(1.6) Negroes, and 18(5.4) Hispanic ("Mexicans") had "other" Rh types. The 8962 Caucasians include the 5056 "Caucasians of Western European Origin" in next row; the data on the 5056 "Caucasians of Western European Origin" was not used in calculating numerical totals nor WMPs.
9. One was Rh₂Rh₁, 5 were Rh₁^wRh₁, 5 were Rh₁^wRh₁, 4 were rh'rh, 4 were rh"rh, and 1 was rh_yrh.
10. 0.2% were Rh₂Rh₁, 0.3% were Rh₂Rh₂, 0.3% were rh'rh, 0.6% were rh"rh, 0.3% were CcD^{ee}, and 0.1% were ccD^{ee}.
11. 5 were Rh₂Rh₁, 10(0.1) were Rh₂Rh₂, there were one each rh'rh', rh"rh", and rh'rh" (or rh₁rh), 54(0.6) were rh"rh, 38(0.4) were rh'rh, 79(0.9) were Rh₁^wRh₁, 37(0.4) were Rh₁^wRh₂, 109(1.2) were Rh₁^wRh₂, 15(0.2) were ccD^{ee}, 33(0.4) were CcD^{ee}, 16(0.2) were ccD^{ee}, 4(0.04) were CCD^{ee}, and 1 was ccD^{ee}.
12. Data of Stolorow and collaborators.
13. 0.2% were Rh₂Rh₁, 0.4% were Rh₂Rh₂, 0.2% were Rh₂Rh₂, 0.6% were rh'rh', 1.2% were rh"rh, 1.0% were rh'rh, and 0.2% were CcD^{ee}.
14. Identical twin study; data for one member of each twin pair tabulated and used in calculations. Among Caucasians, 2 were Rh₂Rh₁, 1 was rh'rh, 4 were Rh₁^wRh₁, 6 were Rh₁Rh₁, C^{w+}, and 5 were Rh₁Rh₂, C^{w+}.
15. And see Shaler, 1978 (Ref T24).
16. 2 were Rh₂Rh₁, 1 was Rh₂Rh₂, 3 were rh'rh, 3 were rh"rh, 1 was rh'rh', and 1 was rh_yrh".
17. 2 were RzR2, 1 was RzRz, 2 were rh'rh, and 1 was rh"rh.
18. 0.2% were Rh₂Rh₁, 0.5% were rh'rh, and 0.4% were rh"rh; Rh₁Rh₂, Rh₂rh" and Rh₂rh_y were included to the Rh₁Rh₂ category, and Rh₀ includes those with D^u.
19. 4 were Rh₂Rh₁, 3 were Rh₂Rh₂, 7 were rh'rh, 5 were rh"rh, 2 were rh"rh", 6 were CcD^{ee}, and there were one each of ccD^{ee}, CcD^{ee}, and CCD^{ee}.
20. There were one each Rh₂Rh₁, Rh₂Rh₂, rh"rh, and rh'rh".
21. Anti-hr' (anti-c) not used; Rh₁Rh₁ and Rh₁rh indistinguishable. 1.4% were rh'rh'.
22. Anti-hr' (anti-c) not used; Rh₁Rh₁ and Rh₁rh indistinguishable. 1.2% were rh'rh' or rh'rh.
23. 3 were rh'rh.
24. 1 was Rh₂Rh₂, 9 were rh'rh, and 1 was ccD^{ee}.
25. 5 were rh'rh, 3 were ccD^{ee}, and 1 was Rh₁^wRh₁.
26. 1% were rh'rh.
27. 1.2% were rh'rh.
28. 2 were rh'rh.
29. 0.9% were rh'rh and 0.4% were rh"rh; Rh₂Rh₂, Rh₂rh", and Rh₂rh_y were included in the Rh₁Rh₂ category, and Rh₀ contains those with D^u.
30. 2 were Rh₂Rh₁, 12 were rh'rh, 12 were ccD^{ee}, 7 were CcD^{ee}, and 4 were ccD^{ee}.
31. 2 were Rh₂Rh₁, 1 was Rh₂Rh₂, and 3 were rh'rh.
32. Primarily Mexican.
33. 2 were Rh₂Rh₂, 2 were rh'rh, and 1 was rh"rh.
34. 46 were Rh₂Rh₁, 47 were Rh₂Rh₂, 28 were R₂R₂, 2 were rh'rh, 2 were rh"rh, 2 were rh"rh", and 4 were rh_yrh_y.
35. 4 were Rh₂Rh₁ and 1 was Rh₂Rh₂.
36. 15(1.6) were Rh₂Rh₁, 2(0.2) were Rh₂Rh₂, and 1(0.1) was rh'rh. Includes the 400 people on the line immediately above; data on the 946 people used for calculations.
37. 14 were Rh₂Rh₁, 6 were Rh₂Rh₂, 3 were rh'rh, 3 were rh"rh, 1 was rh'rh_y, 4 were CCD^{ee}, 1 was CcD^{ee}, 2 were CcD^{ee}, 1 was ccD^{ee}, and 3 were CcD^{ee}.

TABLE 6—Gene frequencies for the Rh blood group system in U.S. populations.

Population	Total	Gene Frequency							Reference to Population Study	
		R_0	R_1	R_2	R_z	r	r'	r''		
CAUCASIAN										
Southeastern GA	327	0.0235	0.4274	0.1428	0.0034	0.3714	0.0080	0.0236		T11
San Francisco, CA	8 819	0.0279	0.4383	0.1500	...	0.3837	1,3	T16
New York, NY	489	0.0311	0.4595	0.1382	0.0022	0.3545	0.0116	0.0130		T17
Tecumseh, MI	8 669	0.0261	0.4050	0.1498	0.0007	0.4051	0.0054	0.0078	5	T20
Greater Philadelphia, PA including part of NJ	218	0.0433	0.4371	0.1505	0.0101	0.3521	0.0070	0.0000	6,7	T26
Miami/Dade County, FL	369	0.0365	0.4066	0.1403	0.0065	0.3900	0.0110	0.0090	5	T28
Los Angeles, CA										
Case material CA, TX, HI, and Mexico City	255	0.0522	0.4402	0.1742	0.0000	0.3184	0.0108	0.0043	2	T29
	905	0.044	0.382	0.136	0.008	0.408	0.010	0.012	6	T31
Connecticut	164	0.0428	0.4221	0.1850	0.0078	0.3339	0.0000	0.0084		T35
TOTAL CAUCASIAN										
Major types	20 042	0.0286	0.4275	0.1532	...	0.3908	1,3	
All types	11 396	0.0291	0.4080	0.1492	0.0017	0.3974	0.0063	0.0083	5	
NEGRO										
Southeastern GA	304	0.5337	0.1046	0.1086	0.0000	0.2294	0.0236	0.0000		T11
Birmingham, AL	613	0.5110	0.0736	0.1003	0.0000	0.2549	0.0601	0.0000	3,5	T40
San Francisco, CA	3 096	0.4887	0.1549	0.1021	...	0.2544	1	
New York, NY	496	0.4700	0.1300	0.1159	0.0000	0.2639	0.0202	0.0000		T17
Greater Philadelphia, PA including part of NJ	173	0.539	0.168	0.092	0.0000	0.201	0.0000	0.0000	1,6	T26
Miami/Dade County, FL	350	0.5019	0.1353	0.1029	0.0000	0.2481	0.0118	0.0000		T28
Los Angeles, CA										
Case material CA, TX, HI, and Mexico City	125	0.4964	0.1840	0.1120	...	0.2076	1,2	T29
	690	0.397	0.163	0.114	0.005	0.292	0.029	0.0000	6	T31

	TOTAL NEGRO									
Major types	2 777	0.4764	0.1338	0.1063	0.0024	0.2592	0.0219	0.0000	3	
All types	5 812	0.4838	0.1542	0.1055	...	0.2565	3	
	HISPANIC									
San Francisco, CA	317	0.0469	0.5110	0.1861	...	0.2559	1	T16
Miami/Dade County, FL	364	0.0808	0.3792	0.1496	0.0070	0.3726	0.0108	0.0000		T28
Los Angeles, CA										
Case material	116	0.0312	0.4514	0.2306	0.0000	0.2016	0.0572	0.0281		T29
CA, TX, HI, and Mexico City	1 212	0.040	0.490	0.235	0.061	0.138	0.005	0.016	4,6,8	T31
	TOTAL HISPANIC									
Major types	1 867	0.0475	0.5094	0.2298	...	0.2133	3	
All types	1 660	0.0499	0.4563	0.2382	0.0275	0.2155	0.0087	0.0039	3,5	
	CHINESE									
New York, NY	946	0.0262	0.7018	0.2051	0.0106	0.0451	0.0112	0.0000	2	T43
	ASIAN									
CA, TX, HI, and Mexico City	1 330	0.020	0.631	0.262	0.000	0.062	0.013	0.003	3,5,6,8	T31

Notes:

1. Frequencies of R_0 , R_1 , R_2 , and r estimated from the seven major phenotypes.
2. χ^2 major types > 7.815.
3. χ^2 major types > 11.345.
4. χ^2 all types > 14.067.
5. χ^2 all types > 18.475.
6. Gene frequencies are those quoted by original author(s).
7. Frequency of C^w estimated to be 0.035.
8. Frequency of r^d estimated to be 0.015 for Hispanic and 0.009 for Asian.

TABLE 7—Genotypic and phenotypic frequencies of MN blood groups in U.S. populations.

Population	Total	Frequency—Number (Percent)			Gene Frequency M	Note	Reference
		M	MN	N			
CAUCASIAN							
New York, NY	3 263	1037(31.8)	1 621(49.7)	603(18.5)	0.5662	1	T1
New York, NY	954	287(30.1)	481(50.4)	186(19.5)	0.5529		T46
Western Alaska	784	(36.6)	(45.0)	(18.4)		2	T47
Minnesota	300	(35.0)	(49.7)	(15.3)		2	T2
University of Iowa							
Controls	2 186	587(26.9)	1 208(55.3)	391(17.9)	0.5448	**	T44
Southeastern GA	333	86(25.8)	166(49.8)	81(24.3)	0.5075		T11
Boston, MA							
Rheumatic	606	200(33.0)	293(48.3)	113(18.6)	0.5718		
Nonrheumatic	600	182(30.3)	305(50.8)	113(18.8)	0.5575		T12
New York, NY	332	95(28.6)	159(47.9)	78(23.5)	0.5256		T48
Eastern San Francisco Bay Area, CA							
Mothers	4 928	(31.3)	(48.9)	(19.9)			
Children	4 928	(28.4)	(52.4)	(19.2)			
New York, NY	900	280(31.1)	429(47.7)	191(21.2)	0.5494	2,3	T15 T49
San Francisco Bay Area, CA							
Caucasians	8 962	2755(30.7)	4 411(49.2)	1796(20.0)	0.5535		
"Caucasians of Western European Origin"							
New York, NY	5 056	1522(30.1)	2 514(49.7)	1020(20.2)	0.5496	4	T16
South Central WV	500	158(31.6)	249(49.8)	93(18.6)	0.5650		T17
Tecumseh, MI	1 051	(29.8)	(51.8)	(18.5)		2	T19
Detroit, MI	8 447	2546(30.1)	4 191(49.6)	1710(20.2)	0.5495		T20
Philadelphia, PA	461	(37.3)	(49.0)	(14.7)		2,5	T25
including part of NJ	218	55(25.2)	119(54.6)	44(20.2)	0.5252	6	T26
Miami/Dade County, FL	366	108(29.5)	181(49.5)	77(21.0)	0.5423	7	T28
CA, HI, TX, and Mexico City	751	241(32.1)	363(48.3)	147(19.6)	0.5626		T31
Connecticut							
Case material	361	87(24.1)	192(53.2)	82(22.7)	0.5069		T35
Southeastern MO	328	(14.9)	(57.9)	(27.1)		2	T36

TABLE 7—(Continued).

Population	Total	Frequency—Number (Percent)			Gene Frequency M	Note	Reference
		M	MN	N			
CA, HI, TX, and Mexico City Connecticut Case material	1 189 19	593(49.9) 4(21.1)	472(39.7) 11(57.9)	124(10.4) 4(21.1)	0.6972 0.5000	* T35	T31
Numerical total	1 902	839(44.1)	823(43.3)	240(12.6)	0.6575		
WMP		44.1	43.3	12.6			
WSDP		7.732	4.945	4.258			
		TOTAL HISPANIC					
		CHINESE					
New York, NY	103	(38.9)	(43.7)	(17.5)		2	T37
New York, NY	400	141(35.2)	201(50.3)	58(14.5)	0.6037	8	T17
New York, NY	946	321(33.9)	477(50.4)	148(15.6)	0.5914	9	T43
		TOTAL CHINESE					
Numerical total	946	321(33.9)	477(50.4)	148(15.6)	0.5914		
WMP		34.4	49.8	15.8			
WSDP		1.478	2.000	0.552			
		ASIAN					
CA, HI, TX, and Mexico City	1 342	432(32.3)	641(47.8)	267(19.9)	0.5622		T31

Notes:

1. Two were MN₂.
 2. Data given in percentages; not used to calculate numerical totals.
 3. Mothers and their infants (children) studied; "mothers" phenotypic percentages used in calculating WMPs.
 4. Included in the 8962 "Caucasians" on line above; data for the 8962 people used for calculations.
 5. Data of Stolorow and collaborators.
 6. Identical twin study; data for one member of each twin pair tabulated and used for calculations.
 7. And see Shaler (1978), Ref T24.
 8. MN class includes nine MN₂.
 9. MN class includes 17 MN₁; includes the 400 reported by Wiener, 1969 (Ref T16); data for the 946 people used for calculations.
- * $\chi^2 > 3.841$.

TABLE 8—Phenotypic frequencies of MNSs blood groups in U.S. populations.

Population	Frequency—Number (Percent)											Note	Reference
	Total	MS	MSs	Ms	MNS	MNSs	MNs	NS	NSs	Ns			
CAUCASIAN													
New York, NY	394	88(22.3)		37(9.4)	106(26.9)		87(22.1)	27(6.9)		49(12.4)		1	T46
Southeastern GA	333	13(3.9)	39(11.7)	34(10.2)	11(3.3)	76(22.8)	79(23.7)	1(0.3)	21(6.3)	59(17.7)			T47
New York, NY	332	12(3.6)	44(13.3)	39(11.7)	7(2.1)	75(22.6)	77(23.2)	4(1.2)	26(7.8)	48(14.5)			T48
New York, NY	900	50(5.6)	121(13.4)	109(12.1)	29(3.2)	194(21.6)	206(22.9)	11(1.2)	63(5.9)	127(14.1)			T49
San Francisco Bay Area, CA	8 962	575(6.4)	1327(14.8)	853(9.5)	331(3.7)	1992(22.2)	2088(23.3)	41(0.5)	457(5.1)	1298(14.5)			
Caucasians of "Caucasians of Western European Origin"	5 056	335(6.6)	719(14.2)	468(9.3)	183(3.6)	1144(22.6)	1187(23.5)	24(0.5)	239(4.7)	757(15.0)		2	T16
South Central WV	1 051	(6.4)	(13.8)	(9.6)	(3.6)	(24.1)	(24.1)	(0.6)	(4.5)	(13.4)		3	T19
Tecumseh, MI	8 447	587(6.9)	1262(14.9)	697(8.3)	278(3.3)	1959(23.2)	1954(23.1)	29(0.3)	423(5.0)	1258(14.9)			T20
Detroit, MI	461	(6.3)	(17.6)	(13.4)	(2.4)	(24.3)	(22.3)	(1.7)	(3.0)	(10.0)		3,4	T25
Philadelphia, PA including part of NJ	218	13(6.0)	25(11.5)	17(7.8)	14(6.4)	41(18.8)	64(29.4)	2(0.9)	11(5.0)	31(14.2)		5	T26
Miami/Dade County, FL	370	28(7.6)	50(13.5)	30(8.1)	19(5.1)	82(22.2)	80(21.6)	3(0.8)	19(5.1)	59(15.9)		6	T28
CA, TX, HI, and Mexico City	751	49(6.5)	111(14.8)	81(10.8)	28(3.7)	158(21.0)	177(23.6)	6(0.8)	22(2.9)	119(15.8)			T31
Southeastern MO	328	(2.1)	(7.9)	(4.9)	(6.4)	(30.5)	(21.0)	(2.7)	(8.8)	(15.6)		3	T36
TOTAL CAUCASIAN													
Numerical total	20 313	1327(6.5)	2979(14.7)	1860(9.2)	717(3.5)	4577(22.5)	4725(23.3)	97(0.5)	1032(5.1)	2999(14.8)			
WMP		6.5	14.6	9.2	3.6	22.8	23.2	0.5	5.1	14.6			
WSDP		0.806	1.126	1.270	0.585	1.236	0.747	0.382	0.795	0.896			

TABLE 8—(Continued).

Population	Frequency—Number (Percent)											Note	Refer- ence
	Total	MS	MSs	Ms	MNS	MNSs	MNs	NS	NSs	Ns	Ns		
NEGRO													
New York, NY	580	(6.9)	(6.9)	(17.2)	(16.6)	(33.1)			(26.0)	(20.2)		1	T37
Ann Arbor, MI	96	10(10.4)	10(10.4)	18(18.7)	12(12.5)	31(32.3)			8(8.3)	17(17.7)		1	T50
Southeastern GA	304	6(2.0)	22(7.2)	42(13.8)	9(3.0)	31(10.2)	113(37.2)	4(1.3)	15(4.9)	58(19.1)		7	T11
New York, NY	206	10(4.9)	16(7.8)	28(13.7)	4(2.0)	33(16.2)	64(31.4)	5(2.5)	10(4.9)	34(16.7)		8	T48
Houston, TX	263	0	12(4.6)	48(18.3)	4(1.5)	48(18.6)	79(30.0)	0	15(5.7)	56(21.3)		9	T51
New York, NY	493	10(2.0)	32(6.5)	78(15.8)	8(1.6)	71(14.4)	163(33.1)	9(1.8)	23(4.7)	99(20.1)		10	T49
San Francisco													
Bay Area, CA	3 146	57(1.8)	201(6.4)	482(15.3)	98(3.1)	349(11.1)	1132(36.0)	33(1.0)	167(5.3)	627(19.9)			T16
South Central WV	106	(2.0)	(6.0)	(16.0)	(4.0)	(10.0)	(34.0)	(1.0)	(4.0)	(24.0)		3	T19
Detroit, MI	485	(3.1)	(6.2)	(13.4)	(2.7)	(11.0)	(38.1)	(2.3)	(4.0)	(19.4)		3,4	T25
Philadelphia, PA													
including part of NJ	173	4(2.3)	4(2.3)	35(20.2)	8(4.6)	14(8.1)	51(29.5)	4(2.3)	13(7.5)	38(22.0)		5,11	T26
Miami/Dade County, FL	337	7(2.1)	23(6.8)	49(14.5)	11(3.3)	40(11.9)	122(36.2)	6(1.8)	11(3.3)	68(20.2)		6	T28
CA, HI, TX, and Mexico City	710	16(2.3)	68(9.6)	128(18.0)	21(3.0)	75(10.6)	198(27.9)	11(1.5)	48(6.8)	145(20.4)			T31
Southeastern MO	64	(0.0)	(3.1)	(9.4)	(1.6)	(10.9)	(20.3)	(6.3)	(9.4)	(39.1)		3	T36
TOTAL NEGRO													
Numerical total	5 632	110(2.0)	378(6.7)	890(15.8)	163(2.9)	662(11.8)	1922(34.1)	72(1.3)	302(5.4)	1125(20.0)			
WMP		2.0	6.6	15.6	2.9	11.7	34.3	1.4	5.3	20.2			
WSDP		0.772	1.383	1.644	0.625	2.021	3.410	0.732	0.989	2.149			
HISPANIC													
San Francisco Bay Area, CA	335	23(6.9)	62(18.5)	41(12.2)	18(5.4)	77(23.0)	78(23.3)	1(0.3)	12(3.6)	23(6.9)			T16
"Mexican" Miami/Dade County, FL	363	24(6.6)	57(15.7)	36(9.9)	12(3.3)	77(21.2)	81(22.3)	2(0.6)	12(3.3)	62(17.1)			T28
CA, HI, TX, and Mexico City	1 189	133(11.2)	277(23.3)	183(15.4)	52(4.4)	205(17.2)	215(18.1)	8(0.7)	31(2.6)	85(7.1)			T31

		TOTAL HISPANIC									
Numerical total	1 887	180(9.5)	396(21.0)	260(13.8)	82(4.3)	359(19.0)	374(19.8)	11(0.6)	55(2.9)	170(9.0)	
WMP		9.5	21.0	13.8	4.3	19.0	19.8	0.6	2.9	9.0	
WSDP		2.151	3.135	2.220	0.629	2.389	2.287	0.140	0.410	3.940	
		CHINESE									
New York, NY	103	(3.9)	(35.0)	(5.8)	(37.9)	(1.0)	(16.5)	1	737		
		ASIAN									
CA, HI, TX, and Mexico City	1 342	8(0.6)	79(5.9)	347(25.9)	8(0.6)	80(6.0)	553(41.2)	3(0.2)	30(2.2)	234(17.4)	737

Notes:

1. Anti-s not used in testing. MS is indistinguishable from MSs, MNS indistinguishable from MNSs, and NS indistinguishable from NSs. Data not used in calculations.
2. Included in the 8962 Caucasians above; data for the 8962 people used for calculations.
3. Data given in percentages; not used in calculating numerical totals.
4. Data of Stolorow and collaborators.
5. Identical twin study; data on one member of each twin pair tabulated and used for calculations.
6. And see Shaler, 1978 (Ref T24).
7. Four were S-s.
8. 1 was MS-s-U-; 1 was MNS-s-U-.
9. Anti-M and anti-M₁ used; M and M₁ combined in tabular data and calculations.
10. 2 were Mu, 1 was MNu, and 4 were Nu.
11. 1 was MNS^u and 1 was NS^u.

TABLE 9—Haplotype frequencies for the MNSs system in U.S. populations.

Population	Total Number Typed	Haplotype Frequency			χ ²	Reference to Population Study
		MS	M _s	NS		
CAUCASIAN						
Southeastern GA	333	0.1965	0.3110	0.0828	0.4097	T11
New York, NY	332	0.1872	0.3384	0.1004	0.3740	T48
New York, NY	900	0.2137	0.3357	0.0907	0.3598	T49
San Francisco Bay Area, CA						
Caucasians	8 962	0.2473	0.3062	0.0691	0.3774	T16
"Caucasians of Western European Origin"	5 056	0.2483	0.3013	0.0668	0.3836	T16
Tecumseh, MI	8 447	0.2600	0.2894	0.0615	0.3890	T20
Philadelphia, PA including part of NJ	218	0.2333	0.2919	0.0763	0.3984	T26
Miami/Dade County, FL	370	0.2610	0.2754	0.0781	0.3854	T28
CA, HI, TX, and Mexico City	751	0.2473	0.3153	0.0570	0.3805	T31
Total Caucasian	20 313	0.2490	0.3011	0.0677	0.3821	
NEGRO						
Southeastern GA	300	0.1096	0.3787	0.0671	0.4446	T11
New York, NY	204	0.1495	0.3627	0.0882	0.3995	T48
New York, NY	493	0.1046	0.3842	0.0779	0.4332	T49
San Francisco Bay Area, CA	3 146	0.1001	0.3861	0.0736	0.4402	T16
Philadelphia, PA including part of NJ	171	0.0690	0.3959	0.1152	0.4198	T26
Miami/Dade County, FL	337	0.1149	0.3762	0.0661	0.4428	T28
CA, TX, HI, and Mexico City	710	0.1185	0.3872	0.0836	0.4107	T31
Total Negro	5 624	0.1037	0.3855	0.4338	0.0769	
HISPANIC						
San Francisco Bay Area, CA						
"Mexicans"	335	0.2752	0.3591	0.0756	0.2901	T16
Miami/Dade County, FL	363	0.2544	0.3021	0.0514	0.3921	T28
CA, TX, HI, and Mexico City	1 189	0.3196	0.3776	0.0584	0.2444	T31
Total Hispanic	1 887	0.3006	0.3584	0.0587	0.2823	
ASIAN						
CA, TX, HI, and Mexico City	1 342	0.0579	0.5043	0.0267	0.4111	T31

Notes:

*χ² > 11.070.

**χ² > 15.086.

TABLE 10—Genotypic and phenotypic frequencies of Kell blood groups in U.S. populations.

Population	Total	Frequency—Number (Percent)			Gene Frequency K	Note	Reference
		KK	Kk	kk			
Boston, MA	210	CAUCASIAN			0.0513	1	T52
Minnesota	300	21(10.0)		189(90.0)	0.0566	1,2	T2
Boston, MA	1 925	(11.0)	(9.5)	(89.0)		2	T55
Southeastern GA	333	28(8.4)		305(91.6)	0.0429	1	T11
San Francisco Bay Area, CA	4 928	(8.3)		(91.7)	0.0424		
Mothers	4 928	(8.6)		(91.4)	0.0440	1,2,3	T15
Children							
NC and VA							
Mongoloids	585	44(7.5)		541(92.5)	0.0382		
Controls	585	36(6.2)		549(93.8)	0.0315		
Donors	253	19(7.5)		234(92.5)	0.0382	1	T53
San Francisco Bay Area, CA	8 962	768(8.6)		8 194(91.4)	0.0440	1	
Caucasians							
“Caucasians of Western	5 056	452(8.9)		4 604(91.1)	0.0455	1,4	T16
European Origin”	500	42(8.4)		458(91.6)	0.0429	1	T17
New York, NY	1 412	(0.4)	(8.1)	(91.4)		2	T19
South Central WV	8 442	6(0.1)	594(7.0)	7 842(92.9)	0.0359		T20
Tecumseh, MI							
Philadelphia, PA	222	9(4.1)		213(95.9)	0.0207	1,5	T26
including part of NJ	911	3(0.1)	84(7.2)	824(90.5)	0.0494		T31
CA, TX, HI, and Mexico City							
TOTAL CAUCASIAN							
Two phenotype (K +, K -):	11 650	967(8.3)		10 683(91.7)	0.0424		
Numerical total		8.3		91.7			
WMP		0.793		0.791			
WSDP							
Three phenotype:	9 353	9(0.1)	678(7.2)	8 666(92.7)	0.0372		
Numerical total		0.2	7.7	92.1			
WMP		0.129	0.984	1.100			
WSDP							

TABLE 10—(Continued).

Population	Total	Frequency—Number (Percent)			Gene Frequency K	Note	Reference
		KK	Kk	kk			
New York, NY	200			(96.5)	0.0177	1,2	T37
New York, NY	126			(98.4)	0.0080	1,2	T54
Southeastern GA	303			300(99.0)	0.0050	1	T11
San Francisco Bay Area, CA							
Mothers	1 453			(98.4)	0.0080		
Children	1 453			(97.9)	0.0106	1,2,3	T15
San Francisco Bay Area, CA	3 146			3 094(98.3)	0.0085	1	T16
New York, NY	500			495(99.0)	0.0050	1	T17
South Central WV	133	(0)	(1.0)	(99.0)		2	T19
Philadelphia, PA							
including part of NJ	176			173(98.3)	0.0085	1,5	T26
CA, TX, HI, and Mexico City	713	0(0)	19(2.7)	694(97.3)	0.0133		T31
TOTAL NEGRO							
Two phenotype (K +, K -):							
Numerical total	4 251			4 186(98.5)	0.0075		
WMP				98.4			
WSDP				0.417			
Three phenotype:							
Numerical total	713	0(0)	19(2.7)	694(97.3)	0.0133		
WMP		0.0	2.4	97.6			
WSDP		0.000	0.606	0.606			
HISPANIC							
San Francisco Bay Area, CA	335			321(95.8)	0.0212	1	T16
"Mexican"							
CA, TX, HI, and Mexico City	1 212	0(0)	23(1.9)	1 189(98.1)	0.0095		T31
CHINESE							
New York, NY	103			(100.0)	0.0000	1,2	T37

New York, NY	160	0(0)	160(100.0)	0.0000	1	T42
New York, NY	946	2(0.2)	944(99.8)	0.0010	1	T43
TOTAL CHINESE						
Two phenotype (K+, K-):	1 106	2(0.2)	1 104(99.8)	0.0010		
Numerical total		0.2	99.8			
WMP		0.087	0.088			
WSDP						
ASIAN						
CA, TX, HI, and Mexico City	1 342	0(0)	1 339(99.8)	0.0011		T31

Notes:

1. Tests done with anti-K; KK and Kk indistinguishable.
2. Distributions given in percentages; data not used in calculating numerical totals.
3. Mothers and their newborns (children) studied; "mothers" percent distribution used in calculating WMPs.
4. The S056 "Caucasians of Western European Origin" are included in the 8962 Caucasians; data for the 8962 Caucasians used in calculating numerical totals and WMPs.
5. Identical twin study; data for one member of each twin pair tabulated and used in calculations.

TABLE 11—Genotypic and phenotypic frequencies of Duffy blood groups in U.S. populations.

Population	Frequency—Number (Percent)				Gene Frequencies			Note	Reference
	Fy(a+b-)	Fy(a+b+)	Fy(a-b+)	Fy(a-b-)	Fy ^a	Fy ^b	Fy		
CAUCASIAN									
Minnesota	100	68(68.0)	32(32.0)		0.4343	0.5657	...	1	T2
Southeastern GA	333	221(66.4)	112(33.6)		0.4203	0.5797	...	1	T11
San Francisco Bay Area, CA									
Mothers	4 928	(66.5)	(33.4)		0.4221	0.5779	...		
Children	4 928	(65.9)	(34.1)		0.4160	0.5840	...	1,2,3	T15
San Francisco Bay Area, CA									
Caucasians	8 962	6 007(67.0)	2955(33.0)		0.4255	0.5745	...		
"Caucasians of Western European Origin"	5 056	3 405(67.4)	1651(32.6)		0.4286	0.5714	...	1,4	T16
South Central WV	1 016	(15.4)	(46.2)					2	T19
Tecumseh, MI	8 946	6 009(67.2)	2937(32.8)		0.4273	0.5727	...	1	T20
Philadelphia, PA including part of NJ	223	49(22.0)	81(36.3)	2(0.9)	0.3887	0.5201	0.0912	5	T26
CA, TX, HI, and Mexico City	913	201(22.0)	408(44.7)	291(31.9)	0.4100	0.5001	0.0900	*	T31
TOTAL CAUCASIAN									
Two Phenotype [Fy(a+), (a-)]:									
Numerical total	18 341	12 305(67.1)	6036(32.9)		0.4263	0.5737	...		
WMP		67.0	33.0						
WSDP		0.265	0.234						
All phenotype:									
Numerical total	1 136	250(22.0)	499(43.9)	372(32.7)	0.4508	0.5039	0.0903	*	
WMP		18.9	45.0	35.4					
WSDP		3.298	1.595	3.055			0.677		
NEGRO									
New York, NY	125	11(8.8)	2(1.6)	27(21.6)	0.0534	0.1235	0.8231		T56
Southeastern GA	304	27(8.9)	85(68.0)	277(91.1)				1	T11
San Francisco Bay Area, CA									
Mothers	1 453	(17.4)	(82.6)						
Children	1 453	(17.1)	(82.9)					1,2,3	T15

New York, NY	179	19(10.6)	9(5.0)	43(24.0)	108(60.3)	0.0809	0.1564	0.7627	*	T57
San Francisco Bay Area, CA	3 146	564(17.9)		2582(82.1)					1	T16
Detroit, MI	404	85(21.0)		319(79.0)					1	T58
South Central WV	103	(17.0)	(5.0)	(78.0)	(0.0)				2	T19
Philadelphia, PA	176	22(12.5)	5(2.8)	30(17.0)	119(67.6)	0.0796	0.1046	0.8158	**	T26
including part of NJ	713	117(16.4)	33(4.6)	149(20.9)	414(58.1)	0.1108	0.1363	0.7529	**	T31
CA, TX, HI, and Mexico City										
TOTAL NEGRO										
Two phenotype [Fy(a+), (a-)]:										
Numerical total	3 854	676(17.5)		3178(82.5)						
WMP		17.5		82.5						
WSDP		2.306		2.306						
All phenotype:										
Numerical total	1 193	169(14.2)	49(4.1)	249(20.9)	726(60.9)	0.0955	0.1332	0.7713	**	
WMP		14.2	4.1	20.9	60.9				6	
WSDP		2.886	1.088	1.925	4.097					
HISPANIC										
San Francisco Bay Area, CA	335	268(80.0)		67(20.0)					1	T16
CA, TX, HI, and Mexico City	1 212	493(40.7)	517(42.7)	182(15.0)	20(1.7)	0.5697	0.3363	0.0937	7,**	T31
ASIAN										
CA, TX, HI, and Mexico City	1 342	1 093(81.4)	232(17.3)	15(1.1)	2(0.1)	0.8763	0.0955	0.0281		T31

Notes:

1. Tests with anti-Fy^a only; Fy(a + b -) indistinguishable from Fy(a + b +) and Fy(a - b +) indistinguishable from Fy(a - b -).
2. Distributions given in percentages; not used in calculation of numerical totals.
3. Mothers and newborns (children) studied; "mothers" phenotypic distributions used in calculation of WMPs.
4. The 5056 "Caucasians of Western European Origin" are included in the 8962 Caucasians; data for the 8962 Caucasians used in calculation of WMPs.
5. Identical twin study; data for one member of each twin pair tabulated and used in calculations.
6. South Central WV data excluded from the calculations.
7. "Mexican."

* $\chi^2 > 3.841$.

** $\chi^2 > 6.635$.

TABLE 12—Genotypic and phenotypic frequencies of Kidd blood groups in U.S. populations.

Population	Total	Frequency—Number (Percent)			Gene Frequencies			Note	Reference
		Jk(a+b ⁻)	Jk(a+b ⁺)	Jk(a-b ⁺)	Jk ^a	Jk ^b			
CAUCASIAN									
Boston, MA	189	146(77.2)		43(22.8)	0.5230	0.4770		1	T52
New York, NY	726	557(76.7)		169(23.3)	0.5175	0.4825		1	T59
Southeastern GA	333	92(27.6)	173(52.0)	68(20.4)	0.5360	0.4640			T11
South Central WV	1016	(77.4)		(22.6)	0.5246	0.4754		2	T19
Philadelphia, PA	177	48(27.1)	84(47.5)	45(25.4)	0.5085	0.4915		3	T26
including part of NJ	514	144(28.0)	244(47.5)	126(24.5)	0.5175	0.4825			T31
CA, TX, HI, and Mexico City									
TOTAL CAUCASIAN									
Two phenotype [Jk(a ⁺), Jk(a ⁻)]:									
Numerical total	915	703(76.8)		212(23.2)	0.5187	0.4813			
WMP		77.1		22.9					
WSDP		0.325		0.320					
All phenotype:									
Numerical total	1024	284(27.7)	501(48.9)	239(23.3)	0.5220	0.4780			
WMP		27.7	48.9	23.3					
WSDP		0.330	2.101	2.053					
NEGRO									
New York, NY	305	283(92.8)		22(7.2)	0.7314	0.2686		1	T59
Southeastern GA	303	166(54.6)	118(38.8)	19(6.3)	0.7401	0.2566			T11
New York, NY	67	38(56.7)	23(34.3)	6(9.0)	0.7388	0.2612			T57
South Central WV	103	(86.0)		(14.0)	0.6258	0.3742		1,2	T19

Philadelphia, PA including part of NJ	119	68(57.1)	39(32.8)	12(10.1)	0.7353	0.2647	3	T26
CA, TX, HI, and Mexico City	697	303(43.5)	353(50.6)	41(5.9)	0.6879	0.3121	**	T31
TOTAL NEGRO								
Two phenotype [Jk(a +), Jk(a -)]:								
Numerical total	305	283(92.8)		22(7.2)	0.7314	0.2686		
WMP		91.1		8.9				
WSDP		2.949		2.948				
All phenotype								
Numerical total	1186	575(48.5)	533(44.9)	78(6.6)	0.7095	0.2905	**	
WMP		48.5	44.9	6.6				
WSDP		6.022	7.038	1.364				
HISPANIC								
CA, TX, HI, and Mexico City	1143	288(25.2)	649(56.8)	206(18.0)	0.5359	0.4641	4,**	T31
CHINESE								
New York, NY	103	54(52.4)		49(47.6)			1	T59
ASIAN								
CA, TX, HI, and Mexico City	1340	262(19.6)	767(57.2)	311(23.2)	0.4817	0.5183	**	T31

Notes:

1. Tests with anti-Jk^a only; Jk(a + b -) indistinguishable from Jk(a + b +).
 2. Distributions given in percentages, not used in calculation of numerical totals.
 3. Identical twin study; data for one member of each twin pair tabulated and used in calculations.
 4. "Mexican."
- * $\chi^2 > 3.841$.
 ** $\chi^2 > 6.635$.

TABLE 13—Phenotypic frequencies of red cell Lewis groups in U.S. populations.

Population	Total	Frequency—Number (Percent)			Note	Reference
		Le(a+b-)	Le(a-b+)	Le(a-b-)		
CAUCASIAN						
New York, NY	460	(22.8)	(71.5)	(5.7)	1	T37
Boston, MA	1194	307(25.7)	666(55.8)	221(18.5)		T12
South Central WV	1412	(25.9)	(74.1)		2	T19
Southeastern PA, southern NJ and DE	935	(20.1)	(70.2)	(9.7)	1	T60
Connecticut	228	46(20.2)	149(65.4)	32(14.5)		T35
TOTAL CAUCASIAN						
Numerical total	1422	353(24.8)	815(57.3)	254(17.9)		
WMP		22.9	63.9	13.2		
WSDP		2.56	7.122	5.033		
NEGRO						
New York, NY	211	(23.2)	(54.5)	(22.3)	1	T61
New York, NY	236	44(11.1)	138(34.8)	54(13.6)		T62
South Central WV	133	(20.0)	(80.0)		2	T19
Southeastern PA, southern NJ and DE	883	(19.6)	(51.9)	(28.5)	1	T60
Connecticut	45	4(8.9)	28(62.2)	13(28.9)		T35
TOTAL NEGRO						
Numerical total	281	48(17.1)	166(59.1)	67(23.8)		
WMP		19.6	53.8	26.6		
WSDP		2.429	2.909	2.775		
CHINESE						
New York, NY	85	(23.5)	(70.6)	(5.9)	1	T37

Notes:

1. Distributions given in percentages; data not used in calculation of numerical totals.
2. Only anti-Le^a used in testing; Le(a-b+) and Le(a-b-) indistinguishable; data not used in calculation of WMPs.

out. Duffy (Table 11) all-phenotype data for total Caucasians and total Negroes did not show good fit. Only a few different studies had sufficient data to contribute to these totals, and some of them exhibited significant or highly significant deviations from expectations. In the Kidd system (Table 12), all Caucasian and most Negro data showed good fit. The Negro all-phenotype total deviates from expectation because of the highly significant deviations in the largest data set contributing to it. The single data sets for Hispanic and Asians showed highly significant deviations as well.

Some but not all of the deviations can be accounted for by inordinately large contributions from small phenotypic classes to the chi-square statistic. Other factors contributing to deviations in the totals values probably include the limited number of data sets available, and the difficulty in deciding which subpopulations belong to the same effective gene pool.

The limited comparisons of WMPs with phenotypic proportions obtained in a fairly large, full ascertained U.S. population sample suggest that these values are relatively representative of the larger population, at least where a sufficiently large data base exists. There may be some value in the combined population gene frequency estimates for parentage probability

TABLE 14—Genotypic and phenotypic frequencies of secretor-nonsecretor groups in U.S. populations.

Population	Total	Frequency—Number (Percent)		Gene Frequencies		Note	Reference
		Secretors	Nonsecretors	Se	se		
Iowa City, IA	1 261	971(77.0)	290(23.0)	0.5204	0.4796		T8
New Haven, CT		CAUCASIAN					
Yale University students	1 000	773(77.3)	227(22.7)	0.5235	0.4765		T9
Boston, MA	1 194	857(71.8)	337(28.2)	0.4687	0.5313		T12
Tecumseh, MI	8 664	6461(74.6)	2203(25.4)	0.4957	0.5043		T20
Los Angeles, CA	205	141(68.8)	64(31.2)	0.4413	0.5587		T22
U.S. national sample	5 735	(77.6)	(22.4)	0.5267	0.4733	1	T30
Los Angeles, CA	38	28(73.7)	10(26.3)	0.4870	0.5130		T33
Numerical total	12 362	TOTAL CAUCASIAN					
WMP		9231(74.7)	3131(25.3)	0.4967	0.5033		
WSDP		75.6	24.4				
		1.890	1.889				
U.S. national sample	999	NEGRO					
Los Angeles, CA	18	(74.2)	(25.8)	0.4921	0.5079	1	T30
		14(77.8)	4(22.2)	0.5286	0.4714		T33

Note:

1. Distributions given in percentages; not used in calculation of numerical totals.

TABLE 15—Comparison of WMP and U.S. phenotypic frequency values^a for the ABO, A₁A₂BO, Rh, and Secretor systems.

System	Phenotype	Proportion, %			
		Caucasian		Negro	
		U.S.	WMP	U.S.	WMP
ABO	O	44.3	44.7	47.5	49.4
	A	42.1	41.1	26.8	26.2
	B	10.2	10.3	21.2	21.3
	AB	3.6	3.8	4.5	4.1
		$\chi^2 = 0.0395$		$\chi^2 = 0.1632$	
A ₁ A ₂ BO	O	44.3	44.7	47.5	49.8
	A ₁	35.2	32.1	23.4	19.4
	A ₂	6.9	8.7	3.4	6.6
	B	10.2	10.6	21.2	20.2
	A ₁ B	2.5	2.8	2.2	2.5
	A ₂ B	1.1	1.0	2.3	1.4
		$\chi^2 = 0.8070$		$\chi^2 = 4.247$	
Rh	R ₀	2.4	2.8	51.6	48.0
	R ₁ r	35.4	34.0	22.7	22.2
	R ₁ R ₁	17.6	18.0	1.7	2.3
	R ₂ r	12.1	12.1	14.8	14.2
	R ₂ R ₂	3.1	1.9	1.2	1.1
	R ₁ R ₂	14.2	13.3	1.9	3.8
	rr	14.1	15.2	4.7	6.5
		$\chi^2 = 0.7385$		$\chi^2 = 3.0960$	
Secretor					
Secretors		77.6	74.7		
Nonsecretors		22.4	25.3		
		$\chi^2 = 0.4838$			

^aRef T30.

calculations as well. It has been shown that differences in the gene frequency values employed in the calculations can have significant effects on the outcomes [24,25].

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