

N-acetylgalactosamine

N-acetylglucosamine

ABO Blood and Human Origins

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Many people know what their blood type is and understand that blood types must be matched in a medical emergency. The ABO blood group is the most significant blood factor in clinical applications involving blood transfusions. Understanding the importance of the ABO blood group is not limited to clinical applications, however. With our recent ability to rapidly sequence genes, the ABO blood group is also proving to be a valuable asset for determining human migration patterns and origins.

What Determines Blood Type?

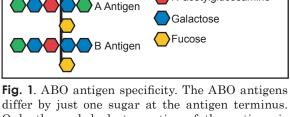
ABO blood types are determined by a cell surface marker that identifies the cell as belonging to "self" or to that individual. These cell surface markers are characterized by a protein or lipid that has an extension of a particular arrangement of sugars. Fig. 1 shows the arrangement of sugars that determines each of the A, B, and O blood types.¹ Note that each is identical, except that types A and B have an additional sugar: N-acetylgalactosamine for A, and galactose for B.

These sugar arrangements are part of an antigen capable of stimulating an immune response that produces antibodies to identify and destroy foreign antigens. People with blood type A produce antibody B when exposed to

antigen B, and those with blood type B produce antibody A when exposed to antigen A. Blood type AB, however, produces no antibodies because both antigens present on the cells are recognized as "self." Blood type O

Blood Type	Cell Antigen	Serum Antibodies	Donor
А	А	В	A or O
В	В	А	B or O
AB	AB	none	all
0	none	A and B	0

Table	1. ABO	blood	groups.
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differ by just one sugar at the antigen terminus. Only the carbohydrate portion of the antigen is illustrated.

produces antibodies A and B, because neither antigen A nor B is present on the cells of type O individuals (Table 1). Antibodies A and B belong to the "M" class of immunoglobins and are expressed from the immunoglobin genes of B-cell lymphocytes upon exposure to foreign antigens. Immunoglobin genes are capable of producing an essentially infinite number of

antibodies through a complex editing and selective process.² Consequently, there isn't a specific "antibody A" gene or "antibody B" gene inherited with a complementary A or B antigen.

A gene for the specification of antigens A or B or type O determines the blood type. An enzyme, glycosyltransferase, is the product of this gene,³ and differences in the sequence of this enzyme (polymorphisms)

determine whether the enzyme attaches N-acetylgalactosamine (antigen A), galactose (antigen B), or no sugar (type O) (Fig. 1). People inherit two genes for blood type; or, more accurately, two alleles, one from each parent. These alleles are represented as IA for type A, IB for type B, and i for type O. Both glycosyltransferase alleles for antigens A and B are expressed when inherited together, producing both antigens and resulting in blood type AB. When the allele for blood type A or B is inherited with type O, the individual will be either type A or B. This is not necessarily because the type O allele is silenced or recessive, but is instead a result of the activity of the A or B glycosyltransferase, while the glycosyltransferase for the O allele is inactive.⁴ A type O individual has both alleles for the inactive glycosyltransferase.

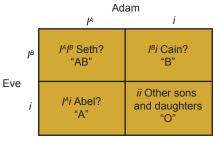


Fig. 2. The possible inheritance of four blood types from Adam and Eve. Alleles for blood type IA = A, IB = B, i = O.

Blood Types and Human Origins

So what light does this shed on human origins? Is it possible for the two people of the Creation account (Adam and Eve) or the eight people on Noah's Ark to give rise to all of the ABO blood types present in humans today? If Adam and Eve were heterozygous for blood types A and B, respectively (one allele for type O and one allele for either type A or B), they could have produced children that had any of the ABO blood types, as illustrated in Fig. 2. The Punnett square simply predicts what the possible phenotypes would be for a given couple's children. From the number of children that Adam and Eve likely produced, it is not difficult to envision all of the ABO blood types being passed down to their offspring.

If Adam and Eve were heterozygous for the ABO blood type gene locus, then the allele frequency for the type O allele is 50 percent (2 of 4 alleles), the allele frequency for type A is 25 percent (1 of 4 alleles), and the allele frequency for type B is 25 percent (Fig. 2). If there are no selective pressures or genetic drift for these alleles, then the allele frequency will remain constant through all of their descendants. The overall allele frequency in the Punnett square is actually the same for the children as it might have been for Adam and Eve. This scenario would also be true for Noah's family and their descendants.

Modern Allele Frequencies

Do human populations today reflect these allele frequencies? The answer is yes. Table 2 shows the allele frequencies for several populations. (Note that these are not blood type frequencies.) There is a general increase in the frequency of the type O allele, and in many populations a drop in the type B allele. But as expected, the frequencies for each allele are close to what they could have been at the start of human history or with Noah's family. The shift in frequency (the increase in type O and decrease in type B) can be caused by migration of people groups that had a higher or lower frequency for one of the alleles at the time

Table	2.	The	allele	frequencies	for	several
popula	tions	$5.^{5,6}$				

Population	Number	Allele Frequency			
		0	А	В	
American	20,000	67	26	7	
French	10,433	64	30	6	
Japanese	29,799	55	28	17	
African	1,538	57	22	21	
Hindu	2,357	55	18	26	

of migration. It could also result from random genetic drift, or from a mutation that renders glycosyltransferase inactive—which would result in blood type O from type A and is likely one cause for the increase in the frequency of the O allele.

Unfortunately, the origin of the ABO alleles gets more complicated when examining the actual gene for glycosyltransferase. There are more than 180 variations (polymorphisms) for the ABO gene listed on the National Center for Biotechnology Information (NCBI) website,⁷ and each one of these polymorphisms can be assigned to one of the three ABO alleles. Most of these polymorphisms do not change glycosyltransferase activity or blood type, but can identify ethnic groups that formed after humans migrated across the globe. Mutation and chromosome crossing-over events are the most plausible cause of these variants.⁸

There are DNA differences, or polymorphisms, that determine the function of glycosyltransferase, resulting in different ABO blood types. These differences are few, but not trivial. The glycosyltransferase specific for antigen A synthesis differs from the antigen B-specific enzyme by just four amino acid residues (out of 354), and there are several DNA sequence differences in the alleles that code for the A- and O-specific enzyme.⁹ The four differences between the A and B glycosyltransferase are enough to allow the enzyme to specify the characteristic terminal sugar that distinguishes antigens A and B. A single DNA deletion in the A-specific allele results in a truncated version of the glycosyltransferase gene product, eliminating enzymatic activity and effectively resulting in blood type O.

Origin Implications of Blood Type O

It can be argued that one of the three alleles is ancestral to the other two. For example, the origin of the O allele, and subsequently blood type O, is simply the result of the deletion resulting in a loss of function of glycosyltransferase activity for the A antigen. A mutation resulting in the loss of function in a protein, at best, would be a "nearly neutral" mutation since blood type O does not appear to have any deleterious effects or selective advantage over the other two blood types. Because neutral or nearly neutral mutations have no selective advantage, it is likely impossible to fix these mutations in a large population of organisms (fixation=100% O alleles) in a reasonable length of time. For example, if a mutation that gave blood type O were actually 1% more beneficial than type A, it would take 100,000 generations to fix this mutation in the modern human population from a beginning population of 10,000 people.^{10, 11} The larger the population at the time of the mutation, the longer it will take for fixation and the less likely the mutation will ever be fixed.

ABO Blood and Human Origins

Molecular evolutionary time scales place modern humans at roughly 200,000 years ago,¹² a timeframe too short to increase the O allele frequency to 60 percent of all people alive today within a population of 10,000. Certainly a biblical timeframe would be far too short for such fixation. The deletion responsible for converting an A allele to an O allele is not present in chimpanzees, and sequence comparisons between humans and chimps indicate this allele is unique to the human lineage,^{13,14} further complicating an evolutionary scenario for the origin of blood type O. This scenario would fit better if the O allele was rare in the population today and appeared in a specific people group. However, the O allele is by far the most common allele globally, indicating that if it did originate via a mutational event, it had to occur when the human population was extremely small and before humans divided into ethnic groups and spread across the globe.

It is possible to achieve the current O allele frequency via a mutation if it occurred at the time of Noah's Flood and was passed on by one of Noah's family members. Noah or Mrs. Noah could have had the O allele and passed it on to each one of their sons, or the alleles could have mutated in one son's offspring. The population of the human race at the time of the Flood and immediately afterward certainly qualifies as a population size that would enable a mutated allele to become common as the population grew. With a starting population of only eight people, the O allele could easily have increased in frequency through random genetic drift in the post-Flood population, reflecting the present levels that are observed today and consistent with computer simulations modeling fixation.¹⁵

Conclusion

If Adam and Eve did not have all three blood type alleles, then there must have been a mutation creating the O allele while the human race was still very small and before humans dispersed across the globe. Whether the origin of blood type O was in Adam and Eve at Creation or whether it arose as a mutational event that took place shortly before or after the Flood, it strongly supports that all humans today are descendants of two individuals or a small group of people that eventually populated the globe. Both scenarios are consistent with the biblical model of human origins.

Footnotes

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