Univariate Genetic Analysis of Oxygen Transport Regulation in Children: The Medical College of Virginia Twin Study

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ABSTRACT. We investigated the relative contributions of genetic, individual environmental, and shared environmental effects on 2,3-diphosphoglycerate (DPG) regulation in preadolescent children. In a population of 165 early pubescent boy and girl twin pairs (11.4 y old), of whom 63 were passive smokers, we asked: 1) Are there differences in the control of DPG levels between early pubertal boys and girls? 2) If present, are these differences influenced by exposure to passive cigarette smoke? Non-passive-smoking boys and girls had similar DPG levels. With exposure to passive smoke, DPG levels increased in boys (p = 0.02) but not in girls. Analysis of variance on DPG demonstrated a parental smoking effect (p = 0.008) and suggested an interactive effect between parental smoking and sex of the child (p = 0.08). Univariate genetic analyses suggested that genes operated at different magnitudes in boys (9%)and girls (39%) in explaining a significant portion of the variance in DPG. The magnitude of shared environmental influences was greater in boys (62%) than in girls (34%), whereas individual environmental effects were similar in boys (29%) and girls (26%). Early pubertal boys differ from girls in their regulation of DPG. Environmental stressors such as passive cigarette smoke may elicit different responses in males and females, even at an early age. The use of path analysis may provide important insights into the mechanisms and interactions of genetic and environmental effects that underly the childhood antecedents of atherosclerotic heart disease. (Pediatr Res 33: 645-648, 1993)

Abbreviations

DPG, 2,3-diphosphoglycerate

Systemic oxygen transport is accomplished using many integrated, self-regulated organ systems. Essential components of this system include pulmonary gas exchange, blood flow, Hb concentration, and Hb affinity for oxygen (1). Organic phosphates within the red cell decrease the oxygen affinity of Hb. The most important quantitatively is DPG. The mechanism of the DPG effect on oxygen affinity involves binding of DPG to the two beta chains in the central cavity of deoxy Hb, forcing a conformational shift that drives oxygen off. Oxyhemoglobin (as well as

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carboxyhemoglobin) binds DPG poorly because of a stereochemically smaller central cavity created by changes in the quarternary structure of Hb with oxygenation (2). An increase in red-cell DPG compensates for an impairment in oxygen supply to tissue and is proportional to the increase in deoxyhemoglobin concentration.

Considerable variation in DPG levels, Hb levels, and oxygen affinity occurs in the normal population. A negative correlation has been observed between Hb levels and the levels of DPG in normal people (3, 4). Measurement of red-cell DPG levels has been used as a "functional biopsy" of the adequacy of tissue oxygenation (5). Whereas the correlation between DPG and Hb level may not be causal, feedback mechanisms appear to regulate them in tandem. Therefore, DPG may function as a monitor for environmental influences on systemic oxygen transport. Despite the importance of this organic phosphate, the inheritance of DPG levels in humans has not been studied in detail.

The purpose of the present study was to quantitate the relative contributions in children of both sexes of genetic and environmental effects (including passive cigarette smoking) to the variance of oxygen transport regulation. Specifically, we asked: 1) whether there were differences in the control of DPG levels between early pubertal boys and girls, and 2) if so, whether these differences were influenced by exposure to passive cigarette smoke. We hypothesized that sex differences in oxygen transport regulation exist at an early age and that males demonstrate a greater susceptibility to environmental influences.

MATERIALS AND METHODS

Population. As part of an ongoing longitudinal study of cardiovascular risk factors during adolescence, we recruited families with twins from nearby school systems. Eleven- and 12^{1/2}-y-old twins were ascertained from more than 75 middle schools of central Virginia within a 150-mile radius. Information packets were mailed to the schools for distribution to parents of twins to maintain confidentiality from the investigators. The parents who replied by mail (50%) were invited to participate. From a total population of 341 twin pairs, 165 twin pairs made up the sample for this study based on availability of DPG measurements. The sample population subjects were similar to those not included in weight, height, Tanner stage, and cigarette smoke exposure.

The families participated in a protocol that included the collection of data on family health and smoking histories and collection of blood samples for biochemical assays. The number of cigarettes smoked each day by the parents was recorded. Serum cotinine level was used both as the measure of smoke exposure and to verify non-passive-smoking status. No attempt was made to prescreen enrollees for the presence or absence of cardiovascular risk factors. Informed written consent, which had been approved by the Committee on the Conduct of Human Research of the Virginia Commonwealth University, was obtained from each family before it entered the study.

Procedures. Zygosity. Zygosity was assessed initially by questionnaire and confirmed by blood group and enzyme testing of all twins and their parents. HLA typing was also performed. With this battery of polymorphisms, the probability of dizygosity for concordant pairs typically is <0.001 (6).

Anthropometrics. Height and weight of each subject in stocking feet were measured with a stadiometer and digital scale, respectively. Sexual staging was performed using a five-scale score based upon Tanner's criteria (7).

Blood Samples. Whole blood was obtained by venipuncture, and red-cell DPG level was determined by standard colorimetric methods (8). The error variance for the measurement of DPG was 5% ($R^2 = 0.95$).

Statistical Analysis. Data are presented as mean \pm SD. Statistical differences between group means were assessed by pooled t test. DPG values were found to be normally distributed using the Shapiro-Wilk statistic. For analysis, the subjects were divided into five groups by zygosity and sex. The designation twin 1 or twin 2 was by birth order with the exception of opposite-sex twin-pair groups, in which the male member was consistently designated as twin 1. The methods of path analysis were used to specify the expected contributions of genetic and environmental variables to the variances and covariances of the five groups of twins (9, 10). We have previously described the use of univariate path analysis to quantitate the relative contributions of genes and environment to the variance of coronary risk factors in children (11, 13). This approach provides estimates and tests of significance for the parameters and provides a test of the assumptions implied in the model (14, 15). Path models were specified and fitted using the LISREL VII program for the structural analysis of covariance matrices (16, 17).

RESULTS

Age and anthropometric data are shown in Table 1. No differences were found between the sexes for age, weight, or height. Girls were more sexually mature than boys by Tanner staging.

Mean DPG levels for boys and girls, cross-classified with passive smoking status, are shown in Table 2. In 63 families, one or both parents were cigarette smokers at the time of evaluation. The total number of cigarettes smoked per day by the parents was similar for female and male twin pairs (9.0 \pm 14.2 versus 12.4 \pm 18.2 cigarettes/d, p = 0.22). None of the twins had ever smoked cigarettes.

Non-passive-smoking boys and girls had similar DPG levels. Overall, passive-smoking children demonstrated higher DPG levels than non-passive-smoking children. However, when analyzed by sex, the noted difference in passive-smoking children

Table 1. Age and anthropometric data

		*	
	Female	Male	р
Age (y)	11.4 ± 0.5	11.4 ± 0.5	NS
Wt (kg)	41.0 ± 9.6	41.1 ± 10.3	NS
Ht (cm)	149.1 ± 7.5	148.9 ± 8.0	NS
Tanner stage	2.9 ± 1.2	2.4 ± 1.1	0.0001

Table 2. Mean levels of DPG (μ mol/mL) by sex and passive smoking status*

	Non-PS	PS	р
Both sexes	2.01 ± 0.28 (102)	2.12 ± 0.27 (63)	0.01
Males	2.01 ± 0.28 (62)	2.15 ± 0.27 (38)	0.02
Females	2.01 ± 0.26 (40)†	2.08 ± 0.28 (25)†	NS

* PS, passive smoking.

† Male-female mean difference was not significant (p > 0.05). Analyses are based on one twin per family.

was accounted for by a significant elevation of DPG levels only in the boys. Analysis of variance on DPG demonstrated that there was an effect of the parents' smoking status (p = 0.008) and a suggestion of an interaction effect between parents' smoking and sex of the child (p = 0.08). Multiple linear regression analysis on DPG was performed to evaluate the fraction of the individual variation that was attributable to biologic traits that included: time since last meal, time since exercise, number of workouts per week, height, weight, age, and Tanner stage. None of these variables were found to be significant predictors of DPG levels.

Intraclass correlation coefficients for twin DPG, cross-categorized by sex and zygosity, are shown in Table 3. If family aggregation were totally due to genetic effects, dizygotic twin correlations would be expected to be one-half of monozygotic twin correlations. Conversely, if family aggregation were totally due to environmental effects shared by both twins, dizygotic twin correlations would be identical to monozygotic twin correlations. The data categorized by sex yielded distinct patterns. Femalefemale intrapair correlations were higher in monozygotic than in dizygotic twins (r = 0.76 versus 0.32), consistent with an important genetic effect. Male-male monozygotic and dizygotic twin-pair correlations were similar (r = 0.61 versus 0.73), consistent with an important shared environmental effect with little if any genetic influence. As expected, the opposite-sex dizygotic twin-pair correlation fell between those for same-sex dizygotic twin pairs.

Maximum-likelihood parameter estimates are presented for the full path model for male and female twins in Figure 1. The regression of the phenotypes of the twins, DPG of twin 1 and twin 2, on the latent variables [additive genes (G), shared environment (C), and individual environment (E)] is specified by the path coefficients: h (path from the genotype to DPG), c (path from the shared environment to DPG), and e (path from individual environment to DPG). The model allows for two kinds of sex differences in genetic and environmental effects. The relative magnitudes of the effects may differ between sexes, represented in the model by using h, c, and e for path coefficients in boys and h', c', and e' for the corresponding values in girls. In addition, we allowed for the possibility that different genes and quantitatively different shared environments may contribute in each sex by allowing the correlation between genetic influences on males and on females (r_h) to be less than unity. Similarly, the correlation between shared environmental influences on males and females (r_c) is allowed to be less than unity.

In the full model for female twins, the largest path coefficients were those associated with additive genetic effects and shared environment. Individual environmental effects were also significant. For male twins, the largest path coefficients are seen for shared environment, followed by individual environment with a small genetic effect.

The model-fitting results expressed as estimated percentages of the total phenotypic or DPG variance are presented in Table 4. The main sources of DPG variance are: genes (h²), shared environment (c²), and individual environment (e²). The χ^2 goodness-of-fit statistic tests the agreement between observed and predicted statistics. A large χ^2 indicates poor fit. The full model

 Table 3. Twin-pair intraclass correlation coefficients for DPG

	No. of	
	pairs	DPG1-DPG2*
Monozygotic twins		
Male-male	51	0.61
Female-female	45	0.76
Dizygotic twins		
Male-male	13	0.73
Female-female	20	0.32
Male-female	36	0.65

* DPG1-DPG2, across-twin correlation for DPG.



Fig. 1. Full path model for male and female twins. The phenotypes of twin 1 and twin 2 (*DPG1*, *DPG2*) are modeled as being determined by additive genetic effects (*G1*, *G2*), environmental effects shared by or common to both twins (*C1*, *C2*), and environmental effects unique to each twin (*E1*, *E2*). Maximum-likelihood parameter estimates for path coefficients (\pm SEM) are presented for *h* (path from genotype to DPG), *c* (path from common environment to DPG), and *e* (path from individual environment to DPG). γ , the correlation of genetic effects on twin 1 with those of twin 2; *DZ*, dizygotic pairs; *MZ*, monozygotic pairs.

Table 4. Univariate	analysis (partitioning of	$^{\circ}DPC$	3 variance*,
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	Sources of DPG variance	Females (%)	Males (%)		
Full model [hh	Full model [hh'cc'ee' $r_{\rm h} = 0.5$ (fixed)]				
$\chi^2 = 8.14$	h²	39.4 ± 20.0	9.6 ± 32.7		
df = 9	c ²	34.5 ± 19.6	61.7 ± 33.1		
p > 0.50	e ²	26.1 ± 5.5	28.7 ± 5.8		
Best-fitting mo	del (cc'ee')				
$\chi^2 = 12.44$	h ²	Not in model	Not in model		
df = 11	c ²	62.8 ± 12.6	65.4 ± 12.9		
p > 0.33	e ²	37.1 ± 6.2	34.6 ± 5.9		

* Percentage of variance estimates are \pm SEM; h², genes; c², shared environment; and e², individual environment.

suggests a greater genetic effect in the girls. A significant shared environmental effect predominates in the boys. Individual environmental effects are significant and similar in both sexes. The best-fitting four-parameter model (cc'ee') excluded genetic effects. This model indicates that the same types of effects (shared environment and individual environment) are working in both boys and girls and that the magnitudes of the effects are comparable. The best-fitting five-parameter model (h'cc'ee') would suggest that genetic effects are active in the girls but not in the boys at this age, an interpretation consistent with the pattern of observed correlations. However, this model did not fit significantly better than the best-fitting four-parameter model (χ^2 difference: 3.39, 1 df; p = 0.066). We suspect that a larger sample of like-sexed dizygotic females may provide evidence for a significant genetic effect within females.

DISCUSSION

We found that early pubertal boys differ from girls in their regulation of DPG. Without cigarette-smoke exposure, boys and girls have similar DPG levels. Chronic cigarette-smoke exposure elevates DPG levels in boys but not girls.

Sex differences in oxygen delivery are known to exist in adults, with lower Hb levels, lower oxygen affinity, and higher DPG levels in females (18). The lower blood-oxygen affinity and redcell mass in females is associated with a higher DPG level as well as a younger red blood cell population (19). Higher red-cell glucose consumption, possibly due to greater hexokinase activity, may account for the higher adult female DPG levels (20). Our finding of similar DPG levels in early pubertal boys and girls who are not exposed to cigarette smoke suggests that gene actions, which determine the activity of various erythrocyte glycolytic enzymes, may change during puberty. In another study (21), DPG was found to be significantly higher in sexually mature females than in males of similar age, and no sex differences existed before puberty. These data also suggest a sex hormone-and maturation-induced influence in the development of the red-cell oxygen transport system. The comparison of DPG levels in younger prepubertal and older, mature, non-smoke-exposed boys and girls may provide further insight into the mechanisms of oxygen transport regulation.

Chronic passive cigarette-smoke exposure was associated with an elevation in DPG levels in boys but not girls. We have previously shown (22) that in families with adult smokers, the DPG level in the children correlated directly with the serum thiocyanate level and the total number of cigarettes smoked by the parents. DPG levels in preadolescent boys increased in a dose-dependent relationship with the amount of passive cigarette-smoke exposure. Their serum thiocyanate and cotinine levels correlated with the total number of cigarettes smoked each day by their parents. Passive-smoking boys and girls had similar elevations of serum thiocyanate. The intratwin pair correlation for thiocyanate was also high (r = 0.94, p < 0.0001). This demonstrates that twins from a smoking family have similar exposures to home environmental cigarette smoke.

In hypoxic states, red blood cell DPG levels increase to meet tissue oxygen requirements. Hypoxia has been implicated as the triggering mechanism for DPG synthesis in active cigarette smokers (23, 24). A hypoxia-driven mechanism triggering DPG synthesis is likely in passive-smoking individuals. Although the boys and girls in the present study had similar amounts of passive cigarette-smoke exposure, the discrepant DPG responses suggest sex-specific patterns of reactivity of the systemic oxygen transport system. Boys demonstrated a greater reactivity to this environmental stressor. This may have occurred because environmental factors predominated over genetic factors in the adjustment of DPG levels in boys.

DPG represents an effective, environmentally influenced genetic regulator of systemic oxygen transport. Using the power of path analysis, we have attempted to quantitate the genetic contribution and partition the relative contributions of individual (including laboratory measurement variation) and shared environmental effects on DPG regulation. Despite this large sample population, which included opposite-sex twin pairs, our observations are preliminary. A larger sample will be needed to confirm these findings. The suggested sex differences in the regulation of DPG may be a consequence of the more advanced sexual maturation of the girls at this age. The extent to which these differences persist or whether genetic control will begin to emerge in males with maturity is being examined in ongoing longitudinal studies.

Cigarette smoking is a powerful independent risk factor associated with premature coronary heart disease in men, with the greatest risk occurring in younger age groups (25, 26). We have previously shown (22) that passive-smoking preadolescent boys had lower levels of HDL₂ cholesterol, which were related to the number of cigarettes smoked daily by the parents of the boys. The lowest HDL₂ cholesterol levels were found in boys exposed to the highest number of cigarettes smoked by their mothers. The HDL₃ subfraction was similarly lower in the passive-smoking children with greater differences seen in the girls. Both HDL subfractions have strong inverse and independent associations with myocardial infarction (13, 27). As a consequence of their lowered HDL cholesterol subfractions, children with long-term exposure to passive smoke may be at elevated risk for the development of premature coronary heart disease.

This study estimates the partitioning of the variance in oxygen transport regulation into genetic and environmental influences by use of univariate genetic path analyses. A larger sample of twins will provide more confidence in the tests for sex differences in the genetic or environmental effects on DPG regulation. Environmental factors appear to predominate over genetic factors in the adjustment of DPG levels in boys. Environmental factors other than passive smoke exposure are likely to impact on DPG control. Other lines of investigation, such as gene by environment interaction (i.e. genetic background influence on the response to a given environmental stimulus), may provide some answers. Multivariate genetic path analyses that include sexual maturity as well as cholesterol subfractions may have the potential to explain the genetic and environmental interrelationships of these variables. The use of path analysis may provide important insights into the mechanisms and interactions of genetic and environmental effects that underly the childhood antecedents of atherosclerotic heart disease.

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