

Great expectations: views of genetic research participants regarding current and future genetic studies

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Purpose: Recruitment of prior participants in genetic research is one strategy suggested to maximize efficient use of research dollars in gene-environment studies. We explored attitudes toward genetic research participation among people in a case-control genetic epidemiology study of colon cancer, the North Carolina Colorectal Cancer Study (NCCCS). **Methods:** Quantitative and qualitative cross-sectional analysis of 801 NCCCS participants. **Results:** Participants were “very positive” (63%) or “positive” (32%) about genetic research, and “very likely” (49%) or “somewhat likely” (40%) to participate in future genetic research. Variables significantly associated with feeling “very positive” were white race, more education, nonreligious, hearing “a lot” about genetic research, and two measures of trust in medical research. Except for race and education, the same variables were significantly associated with being “very likely” to participate in future studies. Qualitatively, “good things” for self and family included discovering causes and cures for cancer, and the value of genetic information. Many could not list “bad things”; those who did mentioned anxiety, “knowing too much,” losing confidentiality, or abuse of information. **Conclusions:** Despite very positive attitudes of these participants toward genetic research, there is significant variation based on participant characteristics. These findings should encourage and caution researchers attempting to recruit prior participants into genetic studies. *Genet Med* 2008;10(3):193–200.

Key Words: genetic research participants, research ethics, minority recruitment, genetic information, genetic epidemiology studies

A large and diverse group of research participants is critical to the study of genetic variation, gene-environment interaction, and disease expression.¹ Through its extramural and intramural research programs, the National Human Genome Research Institute (NHGRI) has promoted such large-scale genomic studies. Recruiting subjects who participated in previous genetic research studies is one strategy that has been suggested in these initiatives to maximize efficient use of research dollars. People who have participated in research studies or in clinical contexts may be more positive about a request to use previously collected biological samples or may be more willing to participate in future genetic studies than people in the general population²; however, these propensities have not been

the subject of systematic evaluation. Few studies have explored the attitudes of such individuals who now seem to represent an important resource for genetic research. To address this gap in the literature, we interviewed individuals who had participated in a genetic epidemiology study, examining their attitudes, positive as well as negative, toward genetic research, and how likely they would be to participate in a genetic research study in the future.

METHODS

The LeARN Study and North Carolina Colorectal Cancer Study

Learning About Research in North Carolina (LeARN) uses a cross-sectional design to study African American and white individuals who recently participated in a case-control genetic epidemiology study of colon cancer risk factors, the North Carolina Colorectal Cancer Study (NCCCS). In the NCCCS study, cases had an initial diagnosis of invasive colorectal cancer, and controls were drawn from Department of Motor Vehicles (DMV) records and Medicare (HCFA) beneficiary lists. Race was initially obtained from cancer registry records and DMV or HCFA files and further confirmed by self-identification during the interview. In the case of conflicting data, the participants' self-identified race was used. The NCCCS partic-

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Participants completed a 2-hour in-person interview that collected data on demographics; dietary, lifestyle, and environmental exposure; and health care access and utilization. Blood and/or a mouthwash sample were obtained from consenting participants at the conclusion of the interview. Individuals were asked to consent to storage of residual DNA and serum for future analyses. In the final NCCCS study, 22% of the respondents were African American and 51% were cases. Of the respondents interviewed during the time period that LeARN was recruiting from NCCCS, 21% were African American and 45% were cases.

Potential participants for the LeARN telephone interviews were identified through the NCCCS database of subjects interested in hearing about other studies. They were eligible if they met the following criteria: (1) self-reported race of African American or non-Latino white, (2) completed the entire NCCCS interview, (3) agreed to be contacted about future studies, (4) lived in the state of North Carolina at the time of the LeARN study, and (5) cognitive functioning as assessed by the interviewer was sufficient to allow successful completion of the telephone interview.

Potential participants were mailed a letter by the NCCCS investigators that introduced the LeARN study, described the telephone interview, and alerted them to expect a follow-up telephone call. Participants were given a toll-free number to call the NCCCS offices and invited to call with questions or if they did not wish to be contacted. Potential participants were contacted on average 4 months after completing the NCCCS interview. During the initial phone call, the nature and purpose of the LeARN study was explained and verbal consent was sought. Each participant was offered an incentive of \$25, which was mailed after completion of the interview. We contracted with a professional survey group, FGI, Inc., to conduct the 45-minute telephone interviews, which consisted of both closed and open-ended questions, using Computer-Assisted Telephone Interviewing methods. All interviews were audio-taped and transcribed for content analysis of the open-ended questions. All procedures were approved by the UNC Institutional Review Board.

We asked participants questions on a range of topics having to do with research participation. The final LeARN Participant Questionnaire contained 94 items—16 open-ended and 78 closed-ended questions—and took on average 44 minutes to complete. To address external and internal validity, we conducted extensive formative research and pilot testing that employed a range of methods, including cognitive interviews, focus groups, and in person or telephone pilot interviews to refine the questionnaire.

LeARN variables and analytic approaches

This analysis focuses on two main quantitative outcomes, each derived from the answers to a 5-point Likert scale question: (1) “Overall, how positive or negative do you feel about this kind of genetic research, the kind that looks at whether genes put people at risk for disease or illness” (“very positive” to “very negative”), and (2) “How likely would you be to take

part in such a study in the future” (“very likely” to “very unlikely”). Because most of the participants answered positively to both these questions, we dichotomized each response at the highest category, i.e., “very positive” versus the other responses for the first question, and “very likely” versus the other responses for the second question.

We were interested in seeing whether characteristics of the LeARN participants were associated with responses to these two questions. We used means and frequencies to describe demographic information (e.g., age, race, gender, education) and other characteristics, such as case/control status, perceived health status, how much they had heard about genetic research, and two measures of trust in medical researchers. For the analyses, we collapsed years of education into a four-category variable; a three-category variable was created for religion, based on a combination of two questions that asked whether respondents are not at all, somewhat, or very religious and whether they rely on religion when times are tough not at all, somewhat, or a great deal (questions drawn from the North Carolina Breast Cancer Screening Project, an eight-year study examining the effects of mammography in rural North Carolina); and a four-category variable was created for cancer history, which differentiated cases and controls by whether they had a relative who had ever been diagnosed with cancer.

We tested associations between these participant characteristics and each of the dichotomous outcomes, using t-tests to calculate mean age differences, and χ^2 tests for the comparisons of each outcome to all other variables. Finally, we fit separate logistic regression models for each outcome to determine which set of characteristics remained statistically significant ($P \leq 0.05$) after adjustment for all other variables in the model. We included all variables in the initial models, whether significant or not in bivariate comparisons. We did not include income in these models because of a large number of missing values. Also, because there were no differences in people who felt positive about genetic research and the first three levels of education (less than high school, high school grad, and technical school and/or some college), we collapsed these groups into one category and compared them with college graduates. All four education categories were used for the outcome of how likely they were to participate in the future. We used likelihood ratio tests to remove nonsignificant variables ($P > 0.05$) from the models. Variables that remained are reported as adjusted percents (of “very positive” or “very likely”) estimated from the beta coefficients of the final reduced models.

A subsample of 194 of the LeARN population was created to explore respondents’ perceptions of genetic research qualitatively, based on responses to open-ended questions. The primary goal of this subsample was to examine race differences in views of the causes of colorectal cancer and views of research that focuses specifically on race and genetic links to health. At the time of subsample creation, there were 97 African American respondents and we decided to include them all. We added an equal number of white respondents, matched by case status, age, education, and sex. Although this subsample was created

for other analyses, we have used some of these qualitative responses to complement our quantitative results.

In this analysis, we examined responses to four open-ended questions: (1) “List the good things for yourself and your family about taking part in genetic research studies”; (2) “List the good things for society about taking part in genetic research studies”; (3) “List the bad things for yourself and your family . . .”; and (4) “List the bad things for society . . .”. Qualitative codes were initially developed for responses to all open-ended questions by the team of LeARN investigators. They were applied, and validated through an iterative process. Three team members coded all responses to open-ended questions, reviewing codes for 10% of transcripts together, to assure uniformity of application. For the subsample of 194, four investigators worked in pairs to further validate the codes for the four questions used in this article.

RESULTS

The overall response rate in the LeARN study was 73%. The final sample of 801 respondents included 19% African Americans and 81% whites, 45% were cases and 55% controls (Table 1). The majority had at least a high school education, and 28% had a college degree. About half the sample had an annual income of less than \$40,000. The mean age of the respondents was 64 years, and 57% were male. Most were “very” (53%) or “somewhat” (42%) religious. Most said they had heard about genetic research and expressed trust in medical researchers. Compared with the larger sample of 801, the subsample includes more African Americans (by design), and some of the demographic characteristics are consistent with the intentional over-sampling (i.e., lower education and income). Most other variables are similar, however, and there was no difference between the two samples in responses to the two outcome variables: how positive people felt about genetic research and how likely they were to participate in future studies.

Positive views of genetic research

When asked, “Overall, how positive or negative do you feel about this kind of genetic research, the kind that looks at whether genes put people at risk for disease or illness,” most were “very positive” (63%) or “positive” (32%) about genetic research (Table 1). Most reported being “very likely” (49%) or “somewhat likely” (40%) to participate in a genetic research study in the future.

What is behind these positive opinions about genetic research? Qualitative analyses of the subsample of 194 provide some insight. When asked to list “the good things” about taking part in genetic research studies, for self and family and for society, respondents’ answers focused on current and future contributions to improving health. Specifically, “good things” for self and family included the potential for discovering the causes of diseases such as cancer, even “eventually finding a cure for all types of cancer whether it’s colon, breast, or whatever.” Many respondents commented on the value of genetic information. One said, “The more you know about anything

Table 1
Characteristics of the sample (n = 801)

Characteristic	n	Percent or mean (SD)
Race		
African American	153	19
White	648	81
Education		
Less than high school	118	15
High school grad	197	25
Technical or some college	258	32
College degree, graduate school	228	28
Age (yr)	800	64.3 (9.90)
Gender		
Male	457	57
Female	344	43
Income		
Less than \$20,000	147	21
\$20,000 to \$40,000	183	27
More than \$40,000	361	52
How religious		
Not religious	40	5
Somewhat religious	326	42
Very religious	413	53
Perceived health status		
Excellent	105	13
Very good	294	37
Good	235	30
Fair	113	14
Poor	45	6
Case/control status		
Case	363	45
Control	438	55
Cancer history		
Case, family history (relative)	51	6
Case, no family history (relative)	312	39
Control, family history (self or relative)	144	18
Control, no family history (self or relative)	294	37
How much heard about genetic research		
A lot	279	35
Little	460	58
None	56	7
Trust medical researchers		
Agree	740	93
Disagree or don’t know	59	7

(Continued)

Table 1
Continued

Characteristic	n	Percent or mean (SD)
Researchers want to know more than they need to know		
Agree	153	19
Disagree or don't know	647	81
How positive about genetic research		
Very positive	489	63
Positive	253	32
Neutral or not positive	41	5
How likely to take part in genetic research		
Very likely	377	49
Somewhat likely	308	40
Neutral or unlikely	82	11

the better off you are.” They mentioned increased awareness and preventive health activities that might reduce or eliminate the risk of disease. As one noted, “If they find that genes (are) running in their family, they can keep a closer check on it,” and although that individual might not benefit, “. . . it may help someone else in my family that comes behind me” or simply help “prepare to deal with the illness later in life.” A few mentioned genetic testing and counseling as positives, and also gene altering. Although most of the responses regarding “good things” for society were similar, a few additional items emerged. Specifically, some participants spoke of financial benefits for society stemming from reduced medical expenditures, increasing productivity due to reduced illnesses, dropping insurance rates, and overall reduction in the emotional burden and stress of disease. Some respondents focused on their expectation for medical breakthroughs, and belief in the inevitability of medical progress, which will “take a long time.” Researchers received kudos from some as “a dedicated bunch of people that really want to see these cures. . . not for their own glorification.”

When we compared characteristics of the 801 respondents to the views on genetic research and participating in future studies, African Americans were somewhat less likely than whites to say they feel “very positive” about genetic research (52% vs. 65%, $P = 0.004$), but there was no significant difference between these two groups’ responses to being “very likely” to participate in future genetic studies (44% vs. 50%, $P = 0.153$) (Table 2). Other respondent characteristics significantly associated both with feeling “very positive” about genetic research and with being “very likely” to participate in future studies included having a college degree, not being religious, having heard a lot about genetic research, agreeing with the statement that they “trust medical researchers,” and disagreeing with the statement that “researchers want to know more than they need to know.” Other variables, including case-control status and cancer history, were not related to either outcome.

After adjustment for all other variables in the model, the following variables remained significantly associated with feeling “very positive” about genetic research (Table 3): white race (66% vs. 56%, $P = 0.035$), more education (72% vs. 61%, $P = 0.010$), not being religious (84% vs. 58% and 67%, $P = 0.004$), hearing “a lot” about genetic research (76% vs. 57% and 58%, $P < 0.001$), agreeing that they “trust medical researchers” (66% vs. 40%, $P = 0.001$), and disagreeing with the statement that “researchers want to know more than they need to know” (54% vs. 67%, $P = 0.007$). With the exception of race and education level, the same variables remained significantly associated with being “very likely” to participate in future genetic studies.

Negative views of genetic research

When respondents were asked to list the “bad things” for self and family, or for society, half of the qualitative subsample of 194 said “none.” Some respondents were emphatic in asserting “none,” justifying this response by the value of the research, e.g., “None, because anything that helps is worth doing.” For others, “none” meant that they could not think of any bad things: “I don’t know of any bad things, I think the research is very important, I really don’t see any bad parts.” For still others, “none” was followed by qualifications that revealed underlying concerns that the goals of research participation were presented honestly and completely: “None, as long as it’s voluntary and for the purpose stated,” “None, if they are doing the study to find the truth, the causes of disease,” “None unless they lie to you about it,” “None except if it is detected but can’t be prevented,” “None unless you give a needle and give somebody some kind of disease.”

Despite generally positive assessments, respondents did list some concerns. Interestingly, these did not vary in number or type when African Americans responses were compared with those of whites in the subsample. The most common concern was anxiety about the implications of the genetic information. Respondents worried about knowing “too much,” about “knowing what conditions we are facing,” “realizing there is no treatment or prevention,” and “if you take that to heart and start concentrating on it and worrying about it.” There was concern that people may not want to know this information, that “people are not interested in knowing what their fate could be,” “maybe it’s best that you don’t know,” that facing it is “not bad, but hard.” Other concerns included failing to maintain the confidentiality of sensitive information collected; as one participant said, “you are judged if the information is given out to employers or. . . [if the] government actually used it against you and starts categorizing people.” Another worried that “researchers perhaps don’t use this material appropriately.” Similar issues were described when respondents were asked to list “bad things” for society, although a few took the opportunity to focus again on the benefits of genetic research for society, arguing that the “bad” thing is people who will not participate. One noted a citizen’s responsibility to participate because all will benefit from the results. Lastly, two respondents warned that the good will of research participants might

Table 2

Bivariate comparisons of sample characteristics by “how positive or negative you feel about this kind of genetic research” (very positive vs. other), and by “how likely would you be to take part in a genetic research study in the future” (very likely vs. other)

Characteristic	n	Feel very positive		Very likely to participate	
		% or mean	P	% or mean	P
Race					
African American	146	52	0.004	44	0.153
White	637	65		50	
Education					
Less than high school	110	58	0.015	50	0.024
High school grad	193	59		42	
Technical or some college	254	59		48	
College degree, graduate school	226	71		57	
Age (yr) (“very” vs. “other”) ^a	782	64.5/63.8	0.328	64.3/63.9	0.435
Gender					
Male	448	63	0.530	51	0.178
Female	335	61		46	
Income					
Less than \$20,000	143	60	0.096	51	0.850
\$20,000 to \$40,000	180	58		53	
More than \$40,000	358	67		51	
How religious					
Not religious	38	82	0.007	71	0.003
Somewhat religious	322	58		44	
Very religious	402	65		52	
Perceived health status					
Excellent	103	67	0.403	58	0.134
Very good	287	65		51	
Good	232	58		44	
Fair	108	62		45	
Poor	45	58		52	
Case/Control Status					
Case	352	61	0.311	49	0.764
Control	431	64		50	
Cancer history					
Case, family history (relative)	50	56	0.637	48	0.150
Case, no family history (relative)	302	61		49	
Control, family history (self or relative)	142	63		58	
Control, no family history (self or relative)	289	65		46	
How much heard about genetic research					
A lot	277	75	<0.001	58	0.002
Little	449	55		44	
None	52	54		48	

(Continued)

Table 2
Continued

Characteristic	n	Feel very positive		Very likely to participate	
		% or mean	P	% or mean	P
Trust medical researchers					
Agree	725	65	<0.001	51	<0.001
Disagree or don't know	56	38		19	
Researchers want to know more than they need to know					
Agree	148	50	<0.001	36	<0.001
Disagree or don't know	634	65		52	

^aFirst number is mean age for “very positive”; second number is mean age for “other.”

Table 3
Adjusted percents^a of characteristics associated with very positive about genetic research and very likely to participate

Characteristic	n	Feel very positive		Very likely to participate		
		Adjusted %	P	n	Adjusted %	P
Race						NS
African American	140	56	0.035			
White	613	66				
Education						NS
Less than college	537	61	0.023			
College graduate, graduate school	216	72				
Religious						
Not religious	37	84	0.002	37	71	0.003
Somewhat religious	320	58		313	43	
Very religious	396	67		390	52	
How much heard about genetic research						
A lot	267	77	<0.001	262	58	0.002
Little	434	56		424	44	
None	52	60		54	50	
Trust medical researchers						
Agree	700	66	<0.001	691	52	<0.001
Disagree or don't know	53	40		49	20	
Researchers want to know more than they need to know						
Agree	140	54	0.007	138	36	0.001
Disagree or don't know	613	67		602	52	

^aAdjusted percents calculated using the beta estimates from a logistic regression model including statistically significant variables ($P < 0.05$).

have limits. As one said, “I think that the time of study is over and the time of finding out some of the causes and problems and cures would be here, now. We have donated to the research for so many years and talked about it for so many years.”

DISCUSSION

As expected for a sample of individuals who have agreed to participate in genetic epidemiology research, the majority of

NCCCS participants interviewed for the LeARN project were positive or very positive about genetic research and described themselves as somewhat or very likely to participate in future studies. Our interviews revealed a number of positives about genetic research for these study participants. They included the potential to discover the causes of disease, and the value of awareness and information, which might lead to prevention strategies even without a cure. Society would benefit as well, especially if medical progress ultimately resulted in cost savings

associated with a healthier population. Half of the qualitative subsample of respondents saw no negatives of genetic research, although in a few cases, “none” was accompanied by qualifications that revealed underlying concerns. Those who listed negatives mentioned anxiety from “knowing too much,” loss of confidentiality, abuse of information, and possible discrimination. It is important to note that many of those who felt quite positive about genetic research were also able to list some of these negative consequences.

A number of studies have examined responses to the request to provide DNA for research purposes and to provide biological samples for long-term storage, examining perceptions of risks and benefits, and how those might vary depending upon the nature of the request and who is responding. These studies document faith in the potential of genetic research to contribute to improved health^{3,4} and different responses by majority and minority groups.^{3,5–8} For example, Chen and colleagues⁹ analyzed NIH Clinical Center consent forms for 61 studies, and found that although 87% of 1670 subjects authorized future use for any medical condition, fewer African Americans (75%) did so. Sterling et al.² reviewed empirical studies of the willingness of different racial/ethnic groups to participate in genetic research and found significantly lower acceptance related to minority group status and other demographic factors.

Several studies of individuals who were already participating in research demonstrated similar differences in response to the request for genetic samples. McQuillan and colleagues,⁵ for example, conducted interviews with 3201 NHANES (National Health and Nutrition Examination Survey) study participants and found African Americans, women, and those of older age were significantly less likely to consent to donate and store specimens. Moorman and colleagues⁶ interviewed 872 female genetic research participants, and reported that being African American, of older age, lower income, less education, higher occupation category, and having poorer health status were associated with being less likely to consent to enroll in a cancer genetics registry. Other studies of genetic research have identified specific concerns of a variety of groups, including minorities, that might explain different participation rates. These specific concerns include control of DNA, potential for misuse of genetic data, racial discrimination, stigmatization of those at genetic risk, and unequal access to potential benefits, including misuse for ancestry testing.^{3,4,10–13}

In this study, we were able to explore the relationship between attitudes toward genetic research and a number of variables raised in these previous studies. We found that respondents were more likely to be “very positive” about genetic research if they were white, more educated, more knowledgeable about genetic research, and more trusting of medical researchers. Thus, despite the fact that the respondents had participated in a genetic epidemiology study (NCCCS), as well as the LeARN study, variations in their perceptions of genetic research reflect the same demographic factors that have been identified in the studies described earlier. The finding that those who are “not religious” are also more positive about genetic research, compared with those who are “somewhat” or

“very” religious, indicates the need for more attention to the role of religion in attitudes toward genetic research.^{14,15} However, because these respondents represented only 5% of the total sample, this particular finding should be interpreted with caution.

Despite generally positive attitudes, it is clear that all prior participants are not equally receptive to the prospect of recruitment into future studies. Some participants expressed distrust of medical researchers, more than half had heard little about genetic research, and 7% of respondents reported a complete lack of knowledge about genetic research even though they had recently participated in a genetic study (Table 1). Those expressing this distrust or having little knowledge were much less likely to be willing to participate in future studies. These findings, and the concerns described in response to open-ended interview questions, represent potential barriers to recruitment and retention of participants from diverse backgrounds for all genomics studies. Focused interventions to help educate potential study participants about genetic research and efforts to demonstrate the trustworthiness of the research team might help encourage future study participation.

Many LeARN respondents expressed “great expectations” about genetic research studies and the promise of medical progress. These expectations coincide with positive attitudes that Americans hold regarding the potential of medical science to address major health conditions, even making us “better than well.”^{16,17} They are also good news for researchers who hope to enlist current genetic study subjects for additional studies or recruit patients into future studies. Yet, our results demonstrate that positive responses may also be associated with overly high expectations, and that good will toward medical research may not persist without “results.” It is important that researchers and scientific leaders address such expectations through careful explanation of the goals, potential benefits, and limitations of genetic research for participants. Equally important, researchers must not assume that prior experience in genetic research guarantees the ability to articulate participants’ reservations about the risks of this type of research.

Our study has several limitations. First, because LeARN participants were drawn from the North Carolina Colorectal Cancer Study, the generalizability of its findings is limited to individuals in North Carolina and those whom have joined similar genetic research studies. Individuals who have not joined such studies may have different and potentially less positive attitudes toward research participation. Additionally, although the LeARN response rate was quite good (73%), there are biases inherent in the sample that further limit its generalizability. The response rate differed by race: 64.6% of African Americans participated compared with 75.4% of whites. Much of this difference is related to unusable telephone numbers and differences in response to callbacks; the number of African Americans and whites who refused to participate in the LeARN study was similar. However, it may be that differences between African Americans and whites about how positive they felt about genetic research conceivably could have been larger than we

observed had we had been able to recruit both races equally. Finally, the LeARN project asks about participation in future studies, assessing intention, which is one step removed from behavior. Despite these limitations, the contributions of LeARN findings are highly relevant to current goals of recruiting genetic study participants. Similar to other reports in the literature, LeARN participants were very positive about the promise of genetic research. Yet, these participants also demonstrate concerns about genetic research studies that will need to be addressed to ensure future participation. Knowledge of barriers and facilitators gathered from this and other studies provides a roadmap to improve public understanding and acceptance of genomics research.¹⁸

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