



# **The BIOCARD Study**

Biomarkers of Cognitive Decline  
Among Normal Individuals

## **Genetics**

Updated: September  
2016

## Glossary of Terms

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Term	Description
Allowable Codes	codes (and their meanings) allowed to be values for that variable
Audit Findings	error rates based on BIOCARD or NIH phase audits error rates are calculated as number of errors / total number of variables examined
Baseline visit	date admitted to NIH phase of BIOCARD study <i>[Note: some data may have been collected prior to this date]</i>
Collection	when the variable information was collected (i.e., Baseline, Follow-up)
Comments	further information about the variable not covered in the above fields
Data Type	numeric or character <i>[Note: Dates are numeric data]</i> numeric or character classifications are strictly related to how the data are stored and not how the data should be analyzed
JHU phase	the study phase at JHU from 2009 - present
Missing OK If	instances (such as skips) or reasons why a blank or missing value is acceptable
NA	not applicable for this variable
NIH / NIH phase	the study phase that was performed at the NIH from 1995-2005
Question Text	the question as it appears on the NACC or BIOCARD data collection forms
Short Description	a short explanation of what the variable means
Source	the name of the NACC form, BIOCARD form, or NIH dataset containing the variable information (or "DERIVED" if the variable was derived) and the variable question number located on the form or in the dataset, if applicable
Unknown Code	the codes for the "unknown", "don't know", or missing values for the variable
Variable Name	the name of the variable in the provided dataset <i>[Note: Variables will follow the NACC naming scheme as closely as possible]</i>

## Acronyms and Definitions

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AD	Alzheimer's Disease	JHU	The Johns Hopkins University
CDR	Clinical Dementia Rating	MCI	Mild Cognitive Impairment
CERAD	Consortium to Establish a Registry for Alzheimer's Disease	MMSE	Mini-Mental State Examination
CNS	Central Nervous System	NACC	National Alzheimer's Coordinating Center
CSF	Cerebrospinal Fluid	NIA	National Institute on Aging
CVD	Cardiovascular Disease	NINDS	National Institute of Neurological Disorders and Stroke
CVLT	California Verbal Learning Test	NPI-Q	Neuropsychiatric Inventory Questionnaire
FAQ	Functional Assessment Questionnaire	UPDRS	Unified Parkinson's Disease Rating Scale
FTD	Frontotemporal Degenerations	WAIS	Wechsler Adult Intelligence Scale
GDS	Geriatric Depression Scale	WMS	Wechsler Memory Scale

# Genetics Dataset Characteristics

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Number of variables: 5

Order of variables:

- |              |                                  |
|--------------|----------------------------------|
| 1) JHUANONID | Participant ID Anonymized by JHU |
| 2) CODENAME  | Participant Codename             |
| 3) NIHID     | Participant NIH ID               |
| 4) STUDY_ID  | Participant Study ID             |
| 5) APOECODE  | Apolipoprotein E (APOE) genotype |

<b>1)</b>	Variable Name	<b>JHUANONID</b>
	Short Description	Participant ID Anonymized by JHU
	Source	NA
	Question Text	NA
	Time of Collection	Baseline
	Data Type	Character
	Allowable Codes	JHU + 6 numbers
	Missing OK If	NA
	Audit Findings	NA
	Comments	None

<b>2)</b>	Variable Name	<b>CODENAME</b>
	Short Description	Participant Codename
	Source	NA
	Question Text	NA
	Time of Collection	Baseline
	Data Type	Character
	Allowable Codes	
	Missing OK If	NA
	Audit Findings	NA
	Comments	None

<b>3)</b>	Variable Name	<b>NIHID</b>
	Short Description	Participant NIH ID
	Source	NA
	Question Text	NA
	Time of Collection	Baseline
	Data Type	Numeric
	Allowable Codes	
	Missing OK If	NA
	Audit Findings	NA
	Comments	None

4)	Variable Name	<b>STUDY_ID</b>
	Short Description	Participant Study ID
	Source	NA
	Question Text	NA
	Time of Collection	Baseline
	Data Type	Numeric
	Allowable Codes	
	Missing OK If	NA
	Audit Findings	NA
	Comments	None

5)	Variable Name	<b>APOECODE</b>
	Short Description	Apolipoprotein E (APOE2, APOE3 and APOE4) genotype
	Source	APOE
	Question Text	NA
	Time of Collection	NA
	Data Type	Numeric
	Allowable Codes	Formatted as X.X where each X is an allelic variant (2, 3, or 4) Possible Values = 2.2, 2.3, 2.4, 3.3, 3.4, 4.4
	Missing OK If	NA
	Audit Findings	No NIH or JHU audit
	Comments	None