

Supplementary Data

Mitochondrial DNA Sequence Variation Associated with Dementia and Cognitive Function in the Elderly

Gregory J. Tranah^{a,*}, Michael A. Nalls^b, Shana M. Katzman^c, Jennifer S. Yokoyama^d, Ernest T. Lam^e, Yiqiang Zhao^c, Sean Mooney^c, Fridtjof Thomas^f, Anne B. Newman^g, Yongmei Liu^h, Steven R. Cummings^a, Tamara B. Harrisⁱ, Kristine Yaffe^j and for the Health, Aging and Body Composition Study

^a*California Pacific Medical Center Research Institute, San Francisco, CA, USA*

^b*Laboratory of Neurogenetics, Intramural Research Program, National Institute on Aging, Bethesda, MD, USA*

^c*Buck Institute for Research on Aging, Novato, CA, USA*

^d*Memory and Aging Center, Department of Neurology, University of California, San Francisco, CA, USA*

^e*Institute for Human Genetics, University of California, San Francisco, CA, USA*

^f*The University of Tennessee Health Science Center, Memphis, TN, USA*

^g*Department of Epidemiology, University of Pittsburgh, Pittsburgh, PA, USA*

^h*Division of Public Health Sciences, Wake Forest School of Medicine, Winston-Salem, NC, USA*

ⁱ*Laboratory of Epidemiology, Demography, and Biometry, National Institute on Aging, Bethesda, MD, USA*

^j*Departments of Psychiatry, Neurology, and Epidemiology, University of California, and the San Francisco VA Medical Center, San Francisco, CA, USA*

Handling Associate Editor: Aleksandra Maruszak

Accepted 14 June 2012

*Correspondence to: Gregory J. Tranah, PhD, California Pacific Medical Center Research Institute, San Francisco Coordinating Center, UCSF, 185 Berry Street, Lobby 5, Suite 5700, San Francisco, CA 94107-1728, USA. E-mail: gtranah@sfcc-cpmc.edu.

Supplementary Table 1

Complex	Gene	No Dementia (n=113)		Dementia (n=22)				
		Nucleotide	Protein	Nucleotide	Protein			
I	ND1	m.3308, T>G	p.M1R	m.3943, A>G	p.I213V			
		m.3337, G>C	p.V11L					
		m.3388, C>A	p.L28M					
		m.3547, A>G	p.I81V					
		m.3593, T>C	p.V96A					
		m.3866, T>C	p.I187T					
		m.4021, A>G	p.T239A					
		m.4172, T>A	p.L289Q					
		ND2	m.4501, C>T			p.S11F	m.4890, A>G m.5461, C>A	p.I141V p.A331D
			m.4640, A>C			p.M57I		
m.4735, A>C	p.N89T							
m.5046, G>A	p.V193I							
m.5277, T>C	p.F270P							
ND3	m.10289, A>G	p.W77C						
	m.10345, T>C	p.I96T						
ND4L			m.10750, A>G	p.N94S				
ND4	m.10775, G>A	p.V6I						
	m.10907, T>C	p.F50L						
	m.11025, T>C	p.L89P						
	m.11065, T>G	p.L102R						
	m.11172, A>G	p.N138S						
	m.11204, T>C	p.F149L						
	m.11928, A>G	p.N390S						
	m.11969, G>A	p.A404T						
	m.12092, C>T	p.L445F						
	ND5	m.12557, C>T			p.T74I	m.12811, T>C m.13676, A>G	p.Y159H p.N447S	
m.12634, A>G		p.I100Y						
m.12952, G>A		p.A206T						
m.13105, A>G		p.I257V						
m.13928, G>A		p.S531N						
m.13934, C>T		p.T533M						
ND6	m.14180, AT>G	p.Y165C						
III	CytB	m.14861, G>A	p.A39T	m.15317, G>A m.15326, A>G	p.A191T p.T194A			
		m.14927, A>G	p.T61A					
		m.15122, A>G	p.T126A					
		m.15257, G>A	p.D171N					
		m.15674, T>C	p.S310P					
		m.15693, T>C	p.M316T					
		m.15758, A>G	p.I338V					
		m.15866, A>G	p.N374D					
		m.15884, G>A	p.A380T					
		IV	COI			m.5913, G>A	p.D4N	m.7080, T>C
m.6261, G>A	p.A120T							
m.6267, G>A	p.A122T							
m.6367, T>C	p.V155A							
m.6465, G>A	p.V188I							
m.6489, C>A	p.L196I							
m.7775, G>A	p.V64I							
m.7805, G>A	p.V74I							
COIII	m.9214, A>G			p.H3R				
	m.9300, G>A			p.A32T				
	m.9621, G>A	p.A139T						
	m.9667, A>G	p.N154S						
	m.9966, G>A	p.V254I						
V	ATP6	m.8705, T>C	p.M60T					
		m.8836, A>G	p.M104V					
		m.8951, T>C	p.V142A					
		m.9070, T>G	p.S182A					
		m.9088, T>C	p.S188P					
		m.9110, T>C	p.I195T					
ATP8			m.8519, G>A	p.E52K				