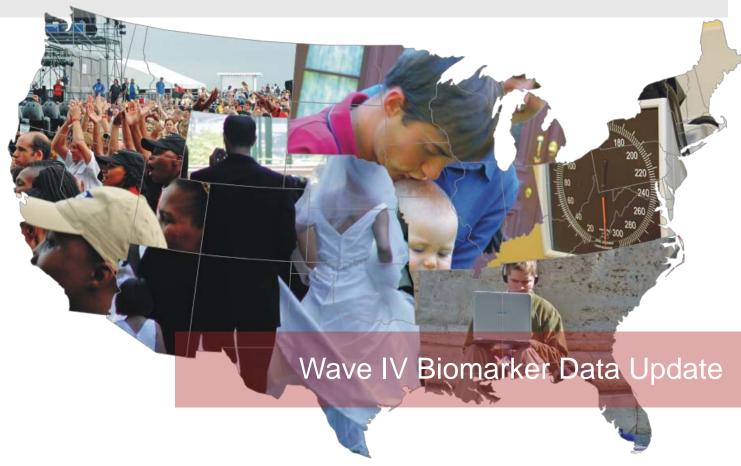


2010 Add Health Users Conference







Wave IV Biomarkers – Overview

- Types + updates
- Measures & Classification + updates
- Collection Sequence
- Collection Methods
- Central Labs
- QA/QC + updates
- Discussion





Wave IV Biomarkers – Types

- Cardiovascular
- Anthropometric
- Metabolic
- Immune
- Inflammatory
- Genetic
- Hematologic
- [Pharmacologic]





- Cardiovascular
 - Primary
 - systolic blood pressure (SBP)
 - diastolic blood pressure (DBP)
 - pulse rate (PR)
 - Secondary
 - pulse pressure (PP = SBP DBP)
 - mean arterial pressure (MAP = [SBP + 2 DBP] 3)





- Cardiovascular
 - Classified according to Joint National Committee VII guidelines*

< 120/80 mm Hg</p>

120-139/80-89 mm Hg

140-159/90-99 mm Hg

• \geq 160/100 mm Hg

Normal

Pre-Hypertension

Stage 1 Hypertension

Stage 2 Hypertension





- Anthropometric
 - Primary
 - weight
 - height
 - waist circumference (waist)
 - Secondary
 - body mass index (BMI = weight in kg / height in m²)





- Anthropometric
 - Classified according to NHLBI Evidence Report*

< 18.5 kg/m²

• 18.5-24.9 kg/m²

• 25.0-25.9 kg/m²

• 30.0-34.9 kg/m²

35.0-39.9 kg/m²

• \geq 40.0 kg/m²

Underweight

Normal

Overweight

Obesity, Stage I

Obesity, Stage II

Obesity, Stage III





- Metabolic, lipids*
 - Primary
 - total cholesterol (τc)
 - high density lipoprotein cholesterol (нос-с)
 - triglycerides (тд)
 - Secondary
 - low density lipoprotein cholesterol (LDL-C = TC HDL-C TG 5)[†]
 - TC:HDL-C ratio (TC:HDL-C = TC / HDL-C)
 - non-HDL-C (= TC HDL-C)





- Metabolic, lipids*
 - Classified according to National Cholesterol Education
 Panel Adult Treatment Panel III guidelines[†]

TC (mg/dL)	< 200 200-239	desirable borderline high
 HDL-C (mg/dL) 	≥ 240 < 40 ≥ 60	high Iow high
 LDL-C (mg/dL) 	< 100 100-129	optimal near optimal
• TG‡ (mg/dL)	130-123 130-159 160-189 ≥ 190 < 150 150-199 200-499 ≥ 500	borderline high high very high normal borderline high high very high





- Metabolic, glucose homeostasis*
 - Primary
 - glucose
 - glycosylated hemoglobin (HbA1c)
 - Secondary
 - mean plasma glucose (MPG = 35.6 HbA1c 77.3)†
 - estimated average glucose (EAG = 28.7 HbA1c 46.7)[‡]





- Metabolic, glucose homeostasis*
 - Classified according to American Diabetes Association and Endocrine Society guidelines[†]
 - fasting glucose

≥ 126 mg/dL diabetes

100-125 mg/dL impaired

≤ 99 mg/dL normal

random glucose

≥ 200 mg/dL diabetes

HbA1c

≥ 6.5% diabetes

5.7-6.4% increased risk for diabetes





- Immune / Inflammatory
 - Epstein-Barr virus antibody titer (EBV)
 - high sensitivity C-reactive protein (hscrp)
 - classified according to Centers for Disease Control / American Heart Association guidelines*
 - low < 1 mg/L
 - average 1-3 mg/L
 - high > 3 mg/L





- Hematologic
 - Hemoglobin (нь)
 - gender-specific classification

	<u>Females</u>	<u>Males</u>
- low	< 12.0 g/dL	13.5 g/dL
average	12.0-16.0 g/dL	13.5-17.5 g/dL
high	> 16.0 g/dL	17.5 g/dL

above, if conversion of OD → g/dL possible





- Genetic
 - candidate gene loci & SNP panels }
 - ancestry informative markers (AIM)

focusing on DA, 5-HT & NA neurotransmission systems with known roles in behavior and health

- 10⁶ SNPs
- Pharmacologic
 - prescription medications
 - select over-the-counter medications (salicylates & NSAIDS)
 - classification according to Multum therapeutic category
 - examples lisinopril ACEI metoprolol BB

nifedipine CCB hctz DIURETIC

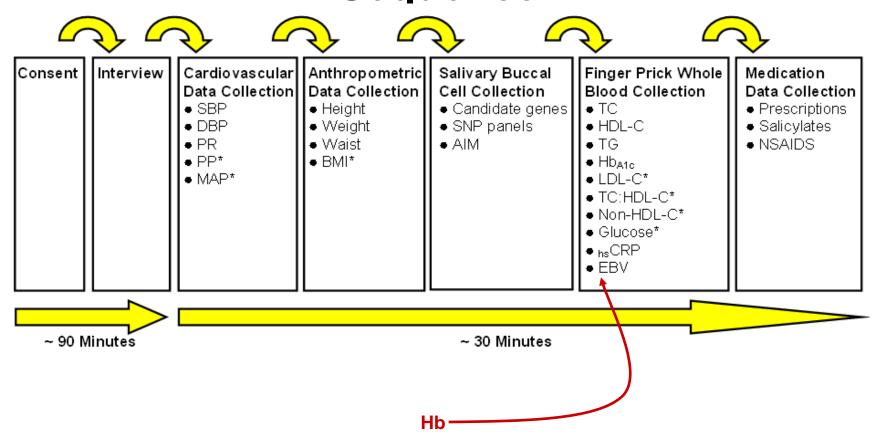
clonidine CASM

hydralazine VASODILATOR





Wave IV Biomarkers – Collection Sequence







Cardiovascular Data Collection Equipment

<u>Measures</u>	Equipment	Picture	Specifications
SBP	MicroLife		30-280/40-200 BP/PR range
DBP	3MC1-PC_IB		1-unit graduations
PR	Oscillometric	Micro I s	198 measure recall
PP*	BP	microlife 70 mm.	3 mmHg accuracy
MAP*	Monitor	On/OH START B	5 beat pulse accuracy
			BHS-approved
			4 "AA" battery-powered
			w/ AC adapter + USB cable
			2 cuffs (24-41 cm)
			160 x 140 x 98 mm
			735 g (w/ batteries)
			< \$65

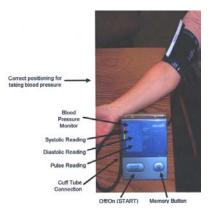




- Cardiovascular Data Collection Protocol
 - trained & certified staff
 - resting & seated respondents
 - cuff matched to arm circumference
 - SBP, DBP & PR
 - measure 3X
 - @ 30-sec intervals
 - double enter
 - automatically average over last 2





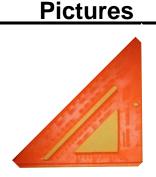






Anthropometric Data Collection Equipment

<u>Measure</u>	Equipment
Height	Carpenter's Square



Specifications portable light weight inexpensive true 90 angle

Tape Measure







10 ft steel tape

adherent pre-labeled





Anthropometric Data Collection Equipment

Measure	Equipment	Pictures	Specifications
Weight	Health-O-Meter	1	4-point load cell
	844KL High	Emn 1	digital display in lb / kg
	Capacity		440 lb maximum
	Digital		0.1 lb graduations
	Scale	-	long life Li++ battery
			low battery warning
		↑ Measure to top ↑	1-year warranty
		Measure to top	12.6 x 12.6 in
		Height:cm	4.5 lb
		Weight:kg	< \$70





Anthropometric Data Collection Equipment

<u>Measure</u>	Equipment
Waist	Seca
	200
	Circumference
	Tape
	Measure



Pictures

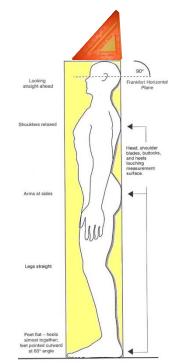


Specifications 200 cm maximum 2-sided cm scaling 1 mm graduations fiberglass tape plastic case automatic roll-up end-peg positioner 90 x 25 x 65 mm 50 g < \$13





- Anthropometric Data Collection Protocol
 - trained & certified staff
 - dressed & unshoed respondents
 - standing on uncarpeted floor
 - measure
 - height to nearest 0.5 cm
 - weight to nearest 0.1 kg
 - waist to nearest 0.5 cm
 - @ superior border of iliac crest
 - @ end expiration
 - horizontal to floor
 - hair/shoe height to nearest
 0.5 cm, as needed











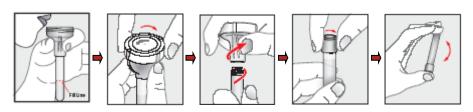
Salivary Buccal Cell DNA Collection Equipment

Meas	sures	Equipment	Sc	chematic	Specifications
Gen	es	DNAGenotek		⋖ Big cap	collection funnel
SNP	panels	Oragene•DNA			preservative-containing cap
AIM		Collection		⋖ Funnel	tube for storage
		Kit		⋖ — Tube	small cap for shipping





- Salivary Buccal Cell DNA Collection Protocol
 - trained & certified staff
 - respondents spit into funnel
 - remove funnel
 - turn big cap clockwise to add preservative
 - turn big cap counter-clockwise to remove it
 - replace big cap with small cap
 - invert 5X to mix saliva and preservative
 - package & FedEx to lab







Wave IV Biomarkers - Central Labs

- Salivary Buccal Cell DNA
 - Andy Smolen, Ph.D. (PI)
 Director, Genotyping Laboratory
 Institute for Behavioral Genetics
 University of Colorado (Boulder, CO)





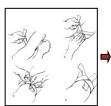
Dried Whole Blood Spot Collection Equipment

<u>Measures</u>	Equipment	<u>Workstation</u>
TC	Gloves	DNA Collection Gloves Device
HDL-C	Band-aids	2 Band-aids
TG	Gauze	Biohazard Container
Hb _{A1c}	Alcohol prep pads	2 Sets of Gauze – 2
LDL-C*	Tourniquet	each
TC:HDL-C*	Lancets	2 Alcohol Chux
Non-HDL-C*	7-spot cards	Prep Pads Pad
Glucose*	Chux pad	7-Spot Blood Card
_{hs} CRP	Biohazard container	
EBV		Rubber 2 Lancets
Hb		Strap





- Dried Whole Blood Spot Collection Protocol
 - trained & certified staff
 - non-fasting respondents
 - clean middle or ring finger w/ alcohol prep pad
 - apply tourniquet to arm
 - prick finger & firmly wipe away 1st drop
 - drop up to 7 blood spots onto card
 - repeat prick X1, if necessary
 - air dry over desiccant
 - package & FedEx to lab















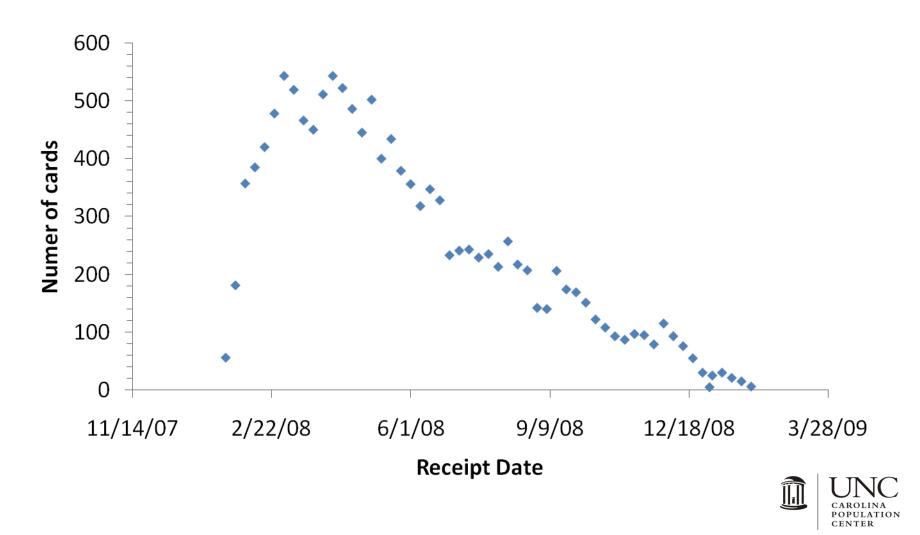
Wave IV Biomarkers – Central Labs

- Dried Whole Blood Spots
 - Mark H. Wener, M.D. (PI)
 - Director, UW Medical Center Laboratories Department of Laboratory Medicine University of Washington (Seattle, WA)
 - Robert Ray, Ph.D.
 Director, FlexSite Diagnostics, Inc. (Palm City, FL)



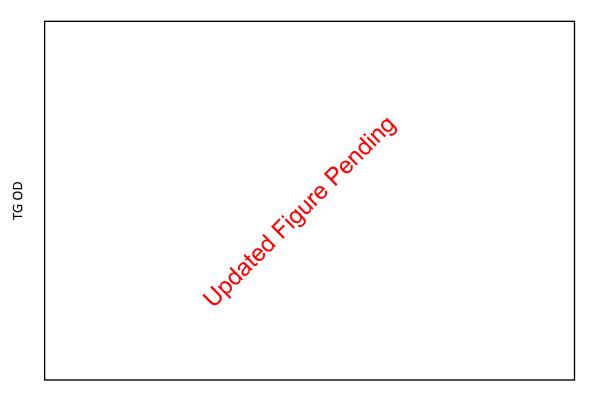


Cards Received Per Week





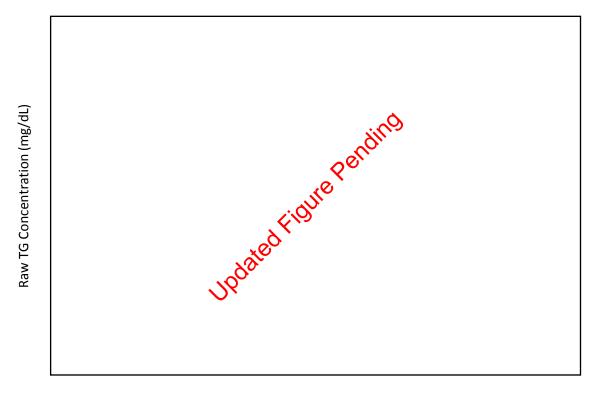
Weekly Mean TG Optical Density



Receipt Date



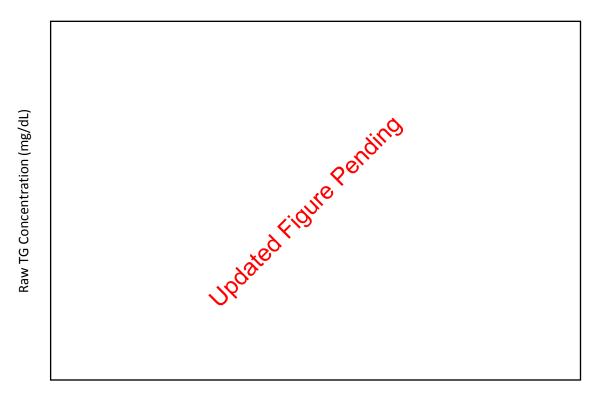




Receipt Date



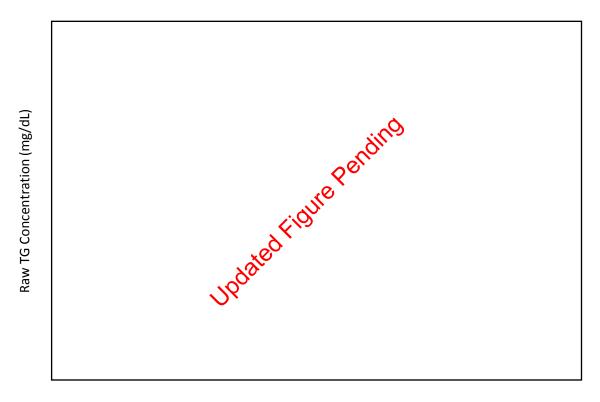




Assay Date





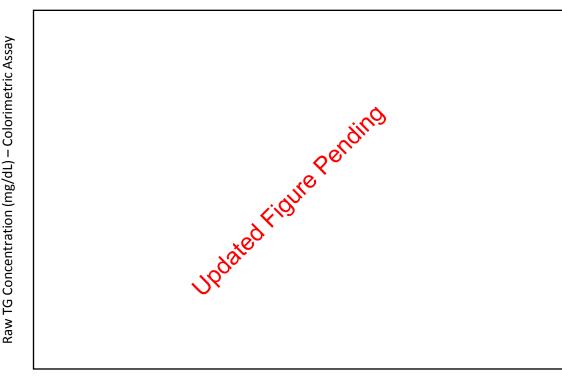


Assay-Receipt Date Interval (days)





Raw TG Concentration (mg/dL) – Colorimetric Assay



Raw TG Concentration (mg/dL) – Fluorometric Assay

Assay Date





Wave IV Biomarkers – QA/QC

- Issues
 - Conversion of Concentrations
 - OD → Raw → Plasma Equivalent
 - Colorimetric → Fluorometric
 - Trends in Concentrations Over
 - Receipt Dates
 - Assay Dates
 - Assay-Receipt Date Intervals
- Other Issues
 - Assay validity (colorimetric vs. criterion standard)
 - Intra-individual variation (visit 1 vs. 2)
 - Intra-sample variation (split sample 1 vs. 2)





Wave IV Biomarkers – QA/QC

- Routine components
 - automated range checks
 - duplicate entry requirements
 - post-encounter verification interviews
- External validation studies
 - to assess digit preference
 - to assess accuracy
 - cardiovascular: BP monitor calibration
 - anthropometric: weight scale calibration
 - metabolic: lipid calibration
- Inter-assay variation study
 - to inform Colorimetric → Fluorometric conversion
- Intra-individual variation study
 - to assess short-term reliability





Digit Preference

DIGIT PREFERENCE IN ANTHROPOMETRIC VARIABLES ADD HEALTH WAVE IV, MAIN STUDY (2008-2009)

Variable	k	n	ChiSq	р	DPS
WEIGHT (kg)	10	15204	608.4	0.000	6.7
HEIGHT (cm)	10	15373	63.4	0.000	2.1
WAIST (cm)	10	15357	26.3	0.002	1.4

Sorted from high to low DPS. DPS = Hense (1991) digit preference score = 100*(Chisq/(n*(k-1)))**0.5. Range = 0-100. Chisq = goodness of fit test stat, where ni = observed cell freq & sum(ni/k) = expected cell freq in cell i. k = number of possible terminal digits, 0-9 by 1 for weight. k = number of possible penultimate digits, 0-9 by 1 for height and waist. p = p value.





Distribution of terminal digits, Add Health Wave IV, Main Study (2008-2009)

Weight	n	%
x.0	2369	(15.6)
x.1	1338	(8.8)
x.2	1426	(9.4)
x.3	1358	(8.9)
x.4	1331	(8.8)
x.5	1717	(11.3)
x.6	1482	(9.8)
x.7	1366	(9.0)
x.8	1484	(9.8)
x.9	1333	(8.8)





DIGIT PREFERENCE IN CARDIOVASCULAR VARIABLES ADD HEALTH WAVE IV, MAIN STUDY (2008-2009)

Variable	k	n	ChiSq	р	DPS
SBP2 (mm Hg)	10	15298	19.3	0.023	1.2
SBP1 (mm Hg)	10	15347	12.5	0.187	1.0
SBP3 (mm Hg)	10	15222	11.7	0.230	0.9
DBP1 (mm Hg)	10	15347	67.9	0.000	2.2
DBP3 (mm Hg)	10	15222	61.8	0.000	2.1
DBP2 (mm Hg)	10	15298	45.0	0.000	1.8
PR1 (beat/min)	10	15262	18.7	0.028	1.2
PR3 (beat/min)	10	15137	10.4	0.319	0.9
PR2 (beat/min)	10	15210	10.0	0.349	0.9

Sorted from high to low DPS. DPS = Hense (1991) digit preference score = 100*(Chisq/(n*(k-1)))**0.5. Range = 0-100. Chisq = goodness of fit test stat, where ni = observed cell freq & sum(ni/k) = expected cell freq in cell i. k = number of possible terminal digits, 0-9 by 1 for weight. p = p value.





Terminal Digit Preference of Blood Pressure, Add Health Wave IV

		SBP Measure			DBP Measure	
Terminal Digit	1	2	3	1	2	3
	n (%)b					
0	1606 (10.5)	1630 (10.7)	1626 (10.7)	1591 (10.4)	1637 (10.7)	1599 (10.5)
1	1545 (10.1)	1517 (9.9)	1519 (10.0)	1445 (9.4)	1442 (9.4)	1411 (9.3)
2	1560 (10.2)	1500 (9.8)	1492 (9.8)	1667 (10.9)	1685 (11.0)	1700 (11.2)
3	1590 (10.4)	1554 (10.2)	1462 (9.6)	1753 (11.4)	1628 (10.6)	1682 (11.1)
4	1538 (10.0)	1598 (10.5)	1560 (10.3)	1558 (10.2)	1468 (9.6)	1490 (9.8)
5	1491 (9.7)	1453 (9.5)	1502 (9.9)	1476 (9.6)	1467 (9.6)	1476 (9.7)
6	1509 (9.8)	1533 (10.0)	1525 (10.0)	1419 (9.3)	1529 (10.0)	1508 (9.9)
7	1500 (9.8)	1445 (9.5)	1527 (10.0)	1506 (9.8)	1478 (9.7)	1445 (9.5)
8	1552 (10.1)	1533 (10.0)	1507 (9.9)	1476 (9.6)	1503 (9.8)	1442 (9.5)
9	1456 (9.5)	1535 (10.0)	1502 (9.9)	1456 (9.5)	1461 (9.6)	1469 (9.7)
Pearson χ²	12.49	19.26	11.71	67.90	44.99	61.76
P value	0.187	0.023	0.230	<0.001	<0.001	<.0001
DPS ^a	0.95	1.2	0.9	2.2	1.8	2.1

^aDPS = digit preference score (see methods). ^bUnweighted sample size and percent.





	FI 122					
SBP3	n	%				
95	1	(1)				
99	1	(1)				
100	2	(2)				
101	1	(1)				
105	1	(1)				
108	3	(3)				
110	9	(8)				
112	4	(4)				
113	1	(1)				
114	2	(2)				
115	3	(3)				
116	5	(5)				
118	15	(14)				
119	2	(2)				
120	36	(32)				
122	5	(5)				
124	6	(5)				
128	6	(5)				
130	5	(5)				
132	1	(1)				
134	2	(2)				

FI 122					
DBP3	n	%			
60	5	(5)			
61	1	(1)			
62	4	(4)			
64	2	(2)			
65	4	(4)			
67	2	(2)			
68	2	(2)			
69	6	(5)			
70	32	(29)			
72	9	(8)			
74	5	(5)			
74 75	5 11	(5) (10)			
	11				
75	11 2 4	(10)			
75 76	11	(10)			
75 76 78	11 2 4	(10) (2) (4)			
75 76 78 79	11 2 4 1 16 2	(10) (2) (4) (1)			
75 76 78 79 80	11 2 4 1 16	(10) (2) (4) (1) (14)			





- Summary
 - affects relatively few FIs & participants
 - yet some FI / participant data
 - consistently affected
 - severely affected
 - highly suspect FI data has been deleted & flagged













- Standard protocol (Dec 2008 Jul 2009)
 - involved two technicians
 - inspected 292 monitor/cuff pairs returned from the field
 - damage
 - missing parts
 - electronic malfunction
 - attached large adult cuff to 37 cm rigid cylinder
 - connected in tandem with
 - BP monitor (Microlife 3MC1-PC_IB)
 - Pressure meter (Netech DigiMano, Model 2000)
 - recorded pressures over 280-40 mm Hg in 20-unit dec
 - attached adult cuff to 28.5 cm rigid cylinder
 - repeated pressure recordings





	MICE	olite Biood Pressu	re Moi	iitor	Calibration
Tech ID:	FI Cuff:	FI Monitor:	_ Test	Date: _	Cuff: ADULT or LARGE ADULT
		<u>Visual (</u>	<u>Check</u>		
◆ TUBE	HAS CRACKING?.			Y N	NO MATCHING TUBING
◆ TUBE	HAS HOLES?			Y N	NO MATCHING TUBING
+ CUFF	HAS WORN OUTER	CLOTH OR VELCRO?		Y N	NO MATCHING CUFF
◆ TUBE	:LEAKS?			Y N	NO MATCHING TUBING
• CUFF	HAS LEAKAGE OF	CUFF BLADDER?		Y N	NO MATCHING CUFF
COMMENTS:					
	Ca	libration Check with F	raccur	0-\/201	uum Matar
	from 280 t	re values on the Digimani o 40 (± 2) mmHg in appro	ximate (ire-vac lecrem	uum Meter and the Microlife ents of 20 (± 2) mmHg.
MEASUR	EMENT NUMBER	DIGIMANO			MICROLIFE
	1 (280)		łg		□□□ mmHg
	2 (260)		łg		□□□ mmHg
	3 (240)		łg		□□□ mmHg
	4 (220)		łg		□□□ mmHg
	5 (200)		łg		□□□ mmHg
	6 (180)		łg		□□□ mmHg
	7 (160)		łg		□□□ mmHg
	8 (140)		łg		□□□ mmHg
	9 (120)		łg		□□□ mmHg
1	0 (100)		łg		□□□ mmHg
•	11 (80)		łg		□□□ mmHg
•	12 (60)		łg		mmHg
•	13 (40)		łg		□ □ □ mmHg

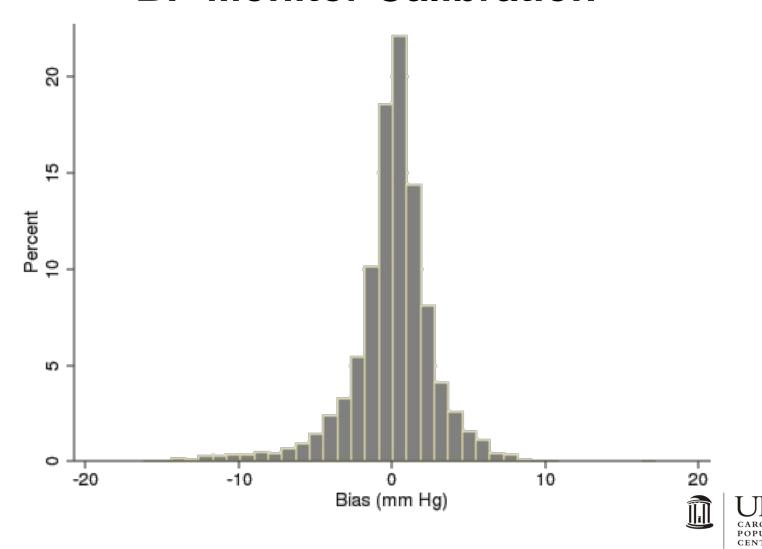




- Analysis
 - bias (mm Hg) = meter monitor difference
 - relative bias (%) = $100 \times \text{bias} / \text{meter pressure}$









Accuracy of Monitor Pressure, by Meter Pressure (mm Hg), Add Health Wave IV (2008-2009)

	Bias	s (mm Hg) ^a	Relat	ive Bias	(%)b
Meter Pressure (mm Hg)	Median	Mean	SD	Median	Mean	SD
280	1.70	1.79	1.78	-0.60%	0.64%	0.64%
260	1.30	1.34	1.89	0.50%	0.51%	0.73%
240	1.10	1.20	1.89	0.46%	0.50%	0.79%
220	1.10	1.26	2.15	0.50%	0.57%	0.98%
200	1.10	1.38	2.16	0.55%	0.69%	1.08%
180	0.70	0.57	2.66	0.39%	0.31%	1.48%
160	0.30	0.06	2.69	0.19%	0.03%	1.68%
140	0.00	-0.28	2.65	0.00%	-0.20%	1.89%
120	-0.40	-0.71	2.59	-0.33%	-0.60%	2.16%
100	-0.60	-1.30	3.27	-0.59%	-1.30%	3.27%
80	-0.75	-1.23	2.74	-0.95%	-1.53%	3.42%
60	-0.70	-1.30	2.73	-1.16%	-2.17%	4.53%
40	-0.90	-1.55	2.53	-2.24%	-3.86%	6.28%

 aBias = meter pressure - monitor pressure. $^bRelative\ bias$ = 100 \times (bias \div meter pressure). SD = standard deviation.





Model: multi-level, random-intercept

```
- Y_{ijk} = \beta_0 + \beta_1 P_{ijk} + \beta_2 C_{ijk} + \gamma_{1k} + \gamma_{2ijk} + \varepsilon_{ijk}
```

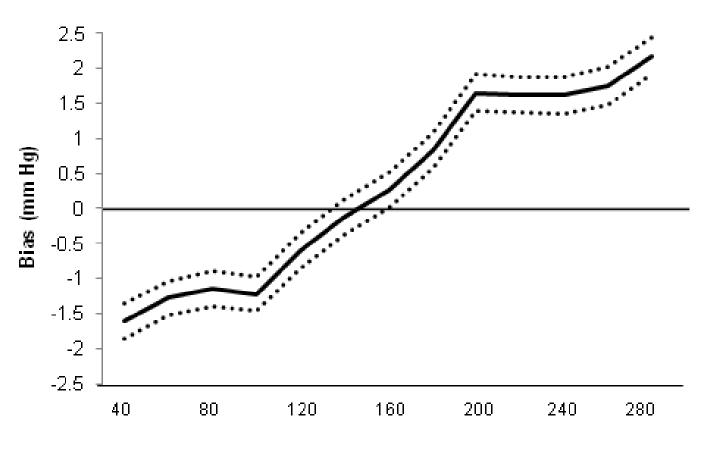
- where
 - i = i^{th} cuff pressure (level 1)
 - $j = j^{th}$ cuff (level 2)
 - $k = k^{th}$ monitor (level 3)
- and
 - Y_{ijk} = bias (mm Hg)

 - β_0^{ijk} = intercept P_{ijk}^{ijk} = vector of meter pressure categories (1-13) C_{ijk} = vector of covariates, e.g. technician ID

 - $\gamma_{1k} + \gamma_{2i(k)}$ = random intercepts @ levels 3 & 2
 - = random error @ level 1 ullet ${\cal E}_{iik}$







Meter Pressure (mm Hg)





Summary

– bias: < 2 mm Hg</pre>

- relative bias: < 4%</p>

- both: > 0 @ BP < 140</pre>

@ BP > 140 mm Hg





- Standard Protocol (Apr 2008 Jul 2008)
 - procured 3 CDC LSP plasma pools, together representing

low
medium
high
TC, HDL-C & TG concentrations

- mixed each pool with washed RBCs
- spotted 20 cards with each plasma-RBC mixture

 - pool 1
 pool 2
 pool 3
 x 20 cards = 60 cards
- sent 1 card/pool to the lab, 2x/wk x 10 wk
- masked lab & technicians to origin of card
- processed the spots per Add Health protocol





- Analysis
 - lab | CDC correlations (Pearson; Spearman)
 - bias (mg/dL) = lab CDC difference
 - relative bias (%) = 100 x bias / CDC
 - coefficient of variation = sd (lab) / mean (lab)





Correlation of lipid concentrations, lab vs. CDC

Variable		Pearson*	Spearman*
Primary	TC		
	HDL-C	Updated tabl	oendir
	TG	(180)	()
Secondary	LDL-C	aated	
	NON - HDL - C	760	

LDL-C = TC - HDL-C - TG \div 5 when TG < 400 (Friedwald). NON-HDL-C = TC - HDL-C. *P values < 0.0001 for all tests of H_0 : r = 0.0.





Difference in lipid concentrations (mg/dL), lab vs. CDC

			Mean (SD)		Relative		_
Variable		Lab	CDC	Bias	Bias	CV	р
Primary	TC		ري.		్యర్తు		رم.
	HDL - C		Sendin		oendir		oendir
	TG	, xi	jobe Pendinos	Updatedtal	le t		Lable pending
Secondary	LDL-C	agtedi		agied !		kaje ^c	`
	NON-HDL-C	760		760		760	

LDL-C = TC - HDL-C - TG \div 5 when TG < 400 (Friedwald). Bias = lab - CDC. Relative bias = (100 * bias) / CDC. NON-HDL-C = TC - HDL-C. P = p value for student's t test of H_0 : Bias = 0.





- Summary
 - pending completion of trend & conversion analyses





Inter-Assay Variation Study

- Standard Protocol
 - identified a race/ethnicity-, gender- & TG-stratified random sample of 96 Main Study participants
 - four race/ethnic groups: NHW, NHB, Hispanic & other
 - three TG concentrations: low, medium & high
 - two genders: male & female
 - 4 x 3 x 2 = 24 groups
 - 4 participants / group
 - resubmitted calorimetrically assayed cards for fluorometric assay
 - masked labs & technicians to participant identity
 - processed the lipids





Inter-Assay Variation Study

- Summary
 - pending completion of trend analysis





- Standard Protocol (2007 2009)
 - computed n needed to estimate the reliability assuming
 - measures are interval-scale
 - reliability estimated as an ICC
 - underlying ICC = 0.7 & precise, 95%CI = 0.2
 - identified a race/ethnicity- & gender-stratified random sample
 - ~ 50 participants in Pre-Test & ~ 50 in Main Study
 - excluded siblings and pregnant women
 - examined participants 2x, 1-2 wk apart
 - visit 1: full interview + biomarkers
 - visit 2: abbreviated interview + biomarkers
 - masked labs & technicians to respondent identity
 - processed the biomarkers





Characteristics of the IIV population*

		Mean (sd) or n (%)
Characteristic		Visit 1 Visit 2
Age (y)		28.5 (1.9)
Female		50 (50%)
Race/Ethnicity	Non-Hispanic White	64 (64%)
	Non-Hispanic Black	16 (16%)
	Hispanic	12 (12%)
	Other	8 (8%)
Pre-Test Partic	ipant	42 (42%)
Time of Day		x:xx pm x:xx pm
Fasting duratio	X.X $(X.X)$ $X.X$ $(X.X)$	
Fasting ≥ 8 hr	XX (XX%)	
V1-V2 Interval	8.6 (3.0)	
Same Field Inte	rviewer Both Visits	84 (84%)

^{*}Based on a final sample of 100 IIV participants.





- Analysis
 - Nested, random-effects model partitioning biomarker variance into its components:

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$$Y_{ijk} = \mu + P_i + V_j(P_i) + \varepsilon_{ijk}$$

- where:
 - Y_{ijk} = value @ jth visit of ith participant μ = intercept

 - P_i = ith participant $V_i(P_i)$ = jth visit nested within ith participant
 - \mathcal{E}_{ijk} = random error
- assuming:
 - P_i , $V_i(P_i)$ and \mathcal{E}_{iik} are independent and ~ $N(0,\sigma^2)$
- under this assumption:
 - $\sigma_T^2 = \sigma_{RP}^2 + \sigma_{RV}^2 + \sigma_{WV}^2$ and ICC = $\sigma_{RP}^2 / \sigma_T^2$





Reliability of available biomarkers Add Health, Wave IV (2007-2009)

Туре	Measure	ICC (95% CI)*
Anthropometric	Weight	1.00 (1.00-1.00)
	Height	0.98 (0.98-0.99)
	BMI	0.99 (0.99-1.00)
	Waist	0.98 (0.97-0.99)
Cardiovascular	SBP	0.81 (0.74-0.88)
	DBP	0.68 (0.57-0.79)
	PR	0.47 (0.31-0.63)
Metabolic	HbA1C	
	Non-HDL-C	
	LDL-C	^
	HDL-C	andinis
	TC	We be.
	Glucose	adial
	TG	vodate
Inflammatory	hsCRP	Updated table pending
Immune	EBV	
Hematologic	Hb	

^{*}ICC (95% CI) = intra-class correlation coefficient, 95% confidence interval.





- Summary of reliability
 - anthropometric measures
 - excellent, near unity
 - cardiovascular measures
 - SBP / DBP acceptable
 - PR marginal
 - other measures
 - pending completion of trend & conversion analyses





To Learn More

- Resources
 - General
 - http://www.cpc.unc.edu/projects/addhealth
 - User Guides
 - http://www.cpc.unc.edu/projects/addhealth/data/guides
 - cardiovascular & anthropometric measures
 - BP monitor calibration form & protocol
 - FI verification script
 - Wave IV Data
 - http://www.icpsr.umich.edu/icpsrweb/DSDR/access/add-health.jsp





Wave IV Biomarkers – Discussion

