

REDUCING CLINICAL NOISE FOR BODY MASS INDEX MEASURES DUE TO UNIT AND TRANSCRIPTION ERRORS IN THE ELECTRONIC HEALTH RECORD



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March 28, 2017

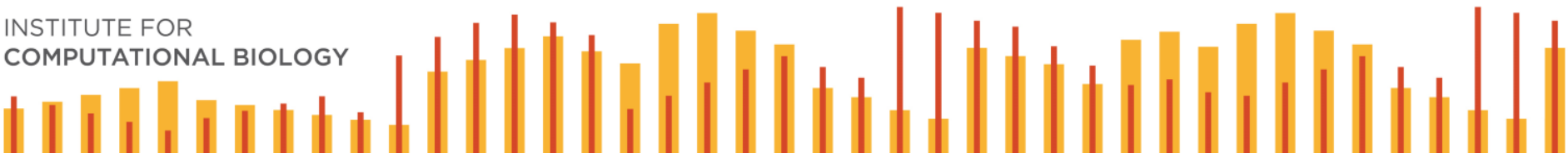
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Associate Professor
Epidemiology and Biostatistics
Institute for Computational Biology

BODY MASS INDEX, THE VARIABLE

Calculated using height and weight:

$$\text{weight} / (\text{height})^2$$

$$\text{kg/m}^2$$



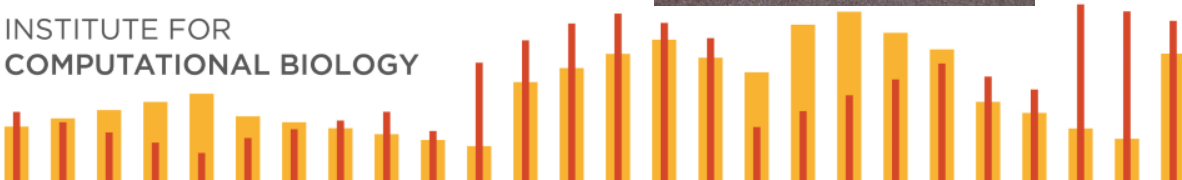
BODY MASS INDEX, THE VARIABLE

BMI is important

Height and weight measured* at most clinic visits

Stadiometers

Scales



BODY MASS INDEX AND EHR

There are problems with EHR-calculated BMI

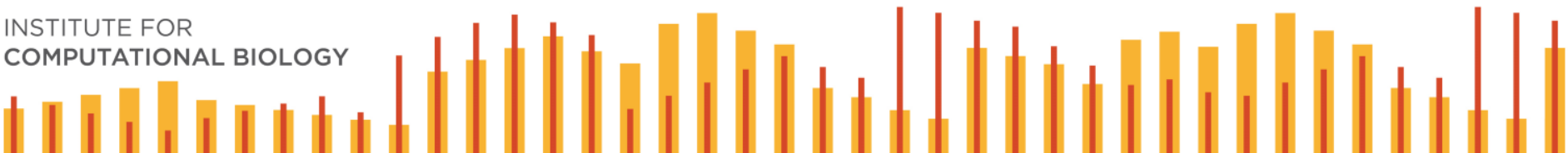
Units (lb v kg; in or ft v m)

Just plain wrong numbers

Age disassociated from weight

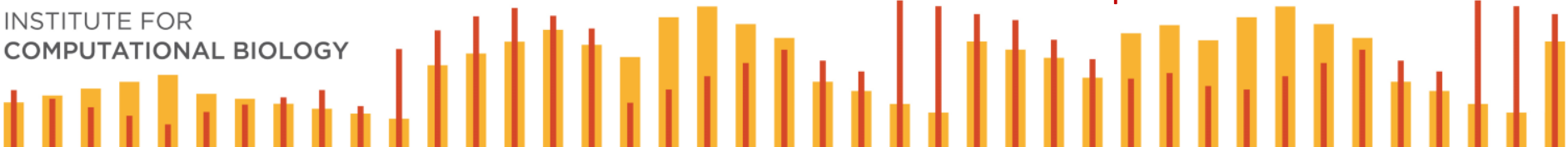
Weight but missing heights

Etc.



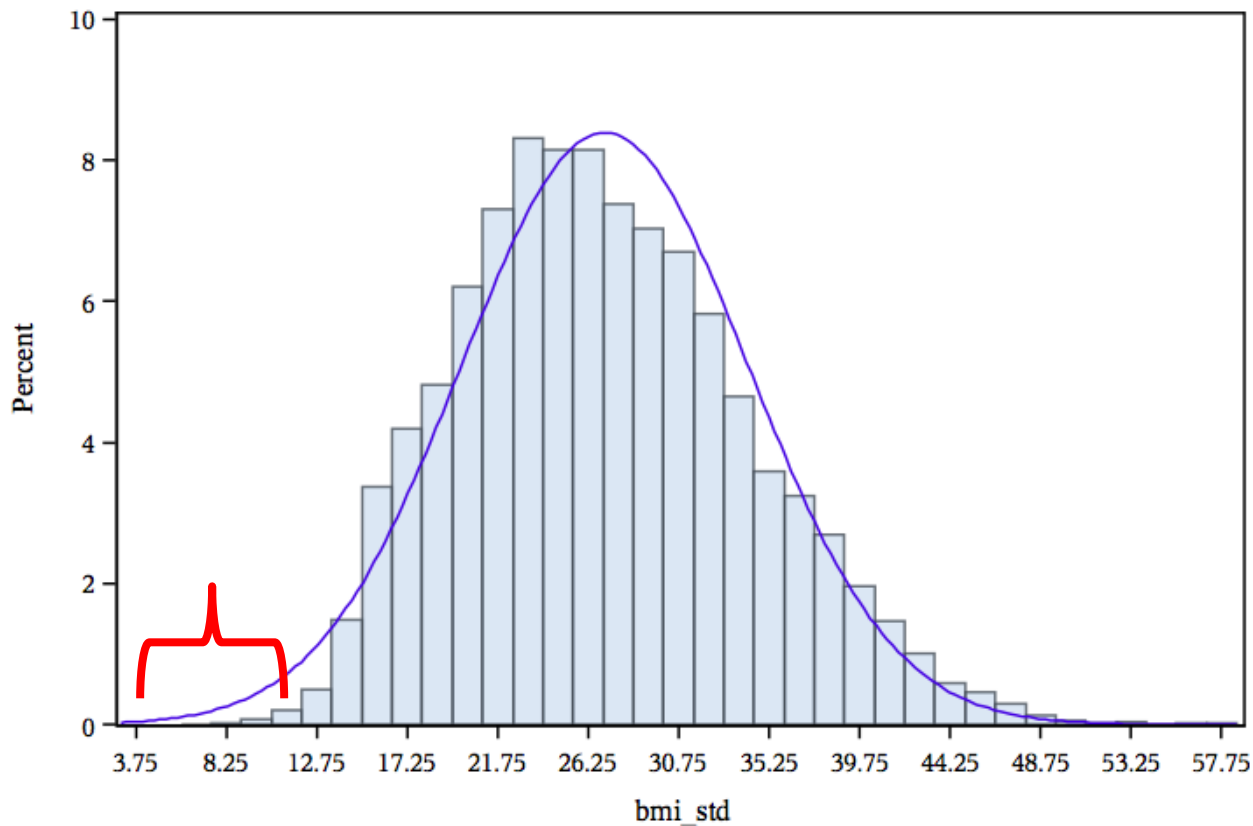
BODY MASS INDEX AND EHR

- In clinical practice, records and values are examined individually
 - Errors are easy to spot and ignore
 - Aggregate data (on a single subject) is synthesized by experts in the context of a patient encounter
- In research, records and values are examined in aggregate over many thousands of individuals
 - Errors are not so easy to identify
 - Data points are not considered in the context of the patient



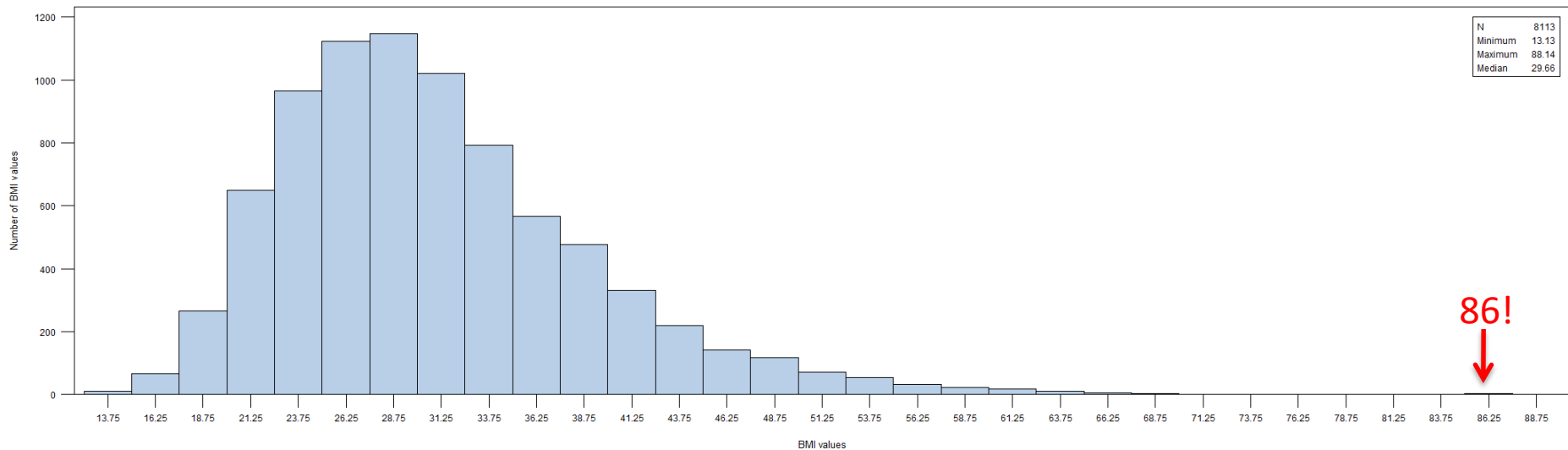
BODY MASS INDEX AND EHR

Implausible low BMIs



BODY MASS INDEX AND EHR

Questionable high BMIs

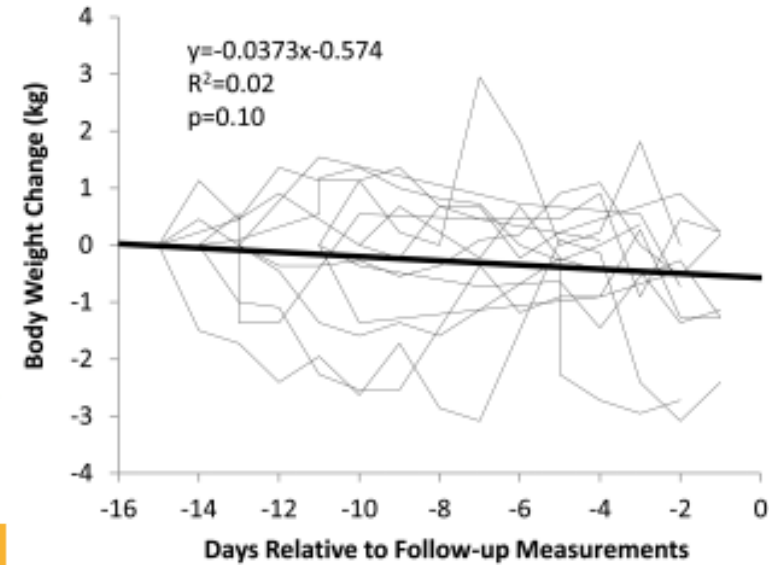


BODY MASS INDEX AND EHR

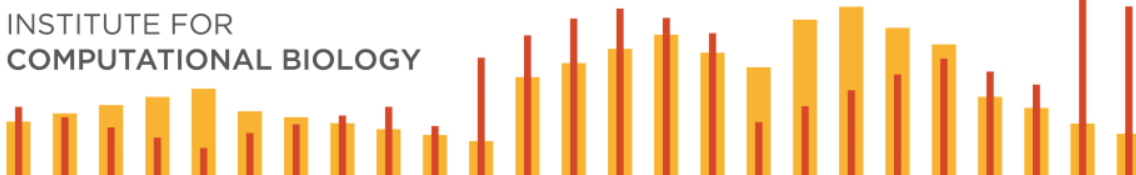
Can we use multiple measures per patient to identify these errors?

Biggest Loser contestants

Fothergill et al (2016) *Obesity* 24(8):1612-1619

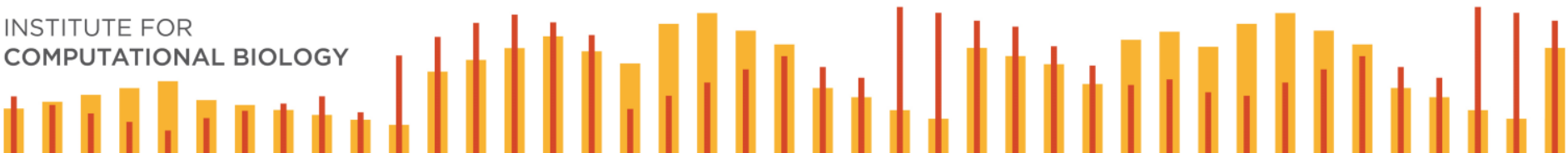
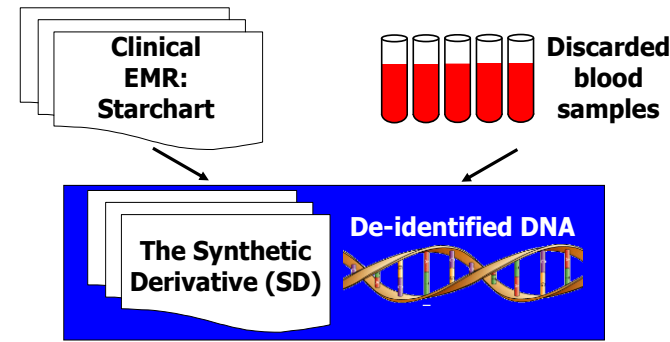


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VUMC BIOVU

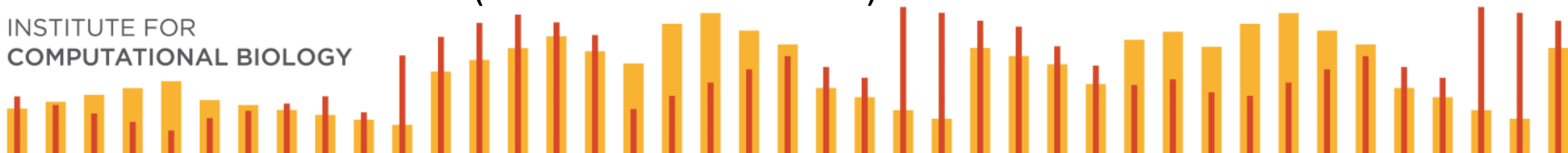
- Opt-out model (2007-2015)
- DNA collected from discarded blood after routine clinical testing has been completed
 - Matched with clinical and demographic data within de-identified EHR (“Synthetic Derivative” database)
 - >225,000 DNA samples



VUMC EHR

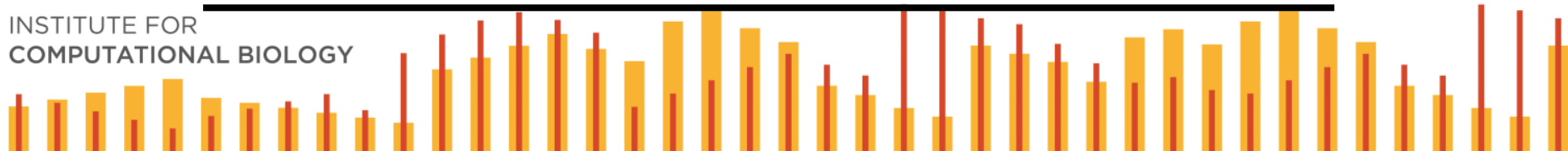
- StarChart
 - Designed, built, and maintained by faculty-led teams
 - Being replaced with EPIC (2018)
- >2 million records, including order entry data on inpatients since 1994
- A document-centric architecture
 - Structured (ex. ICD-9-CM and ICD-10-CM codes)
 - Unstructured (ex. clinical notes)

VUMC in Nashville, TN



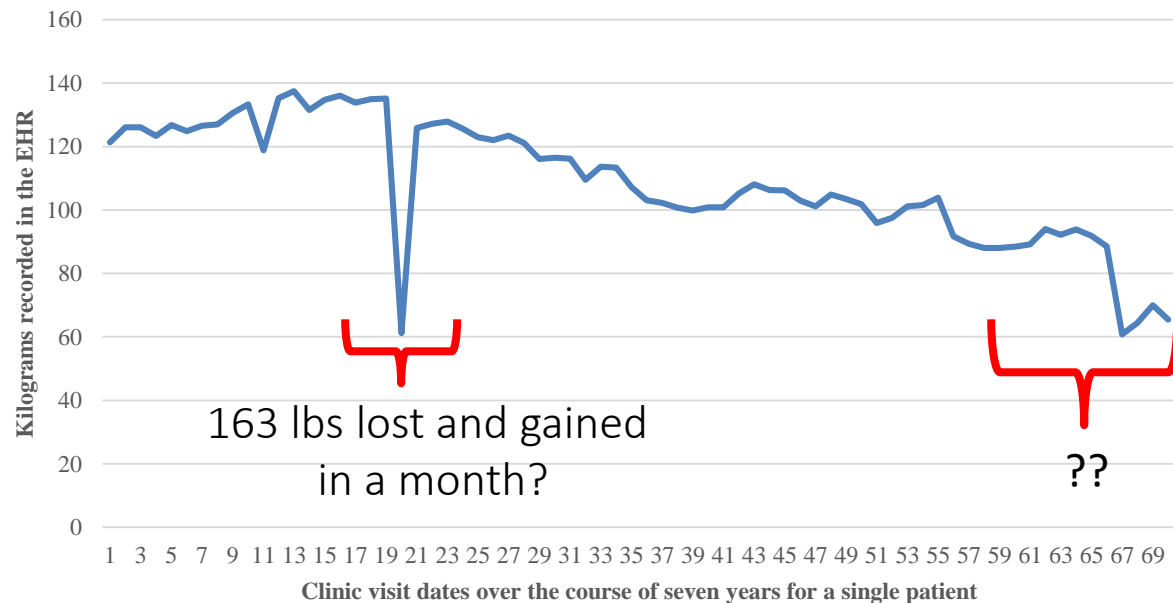
n=15,863

Female (%)	10,050 (63.35)
Median (SD) age	37 (20.46)
African American (%)	11,521 (73.06)
Hispanic (%)	1,714 (10.87)
Asian (%)	1,122 (7.12)
Other (%)	1,412 (8.95)

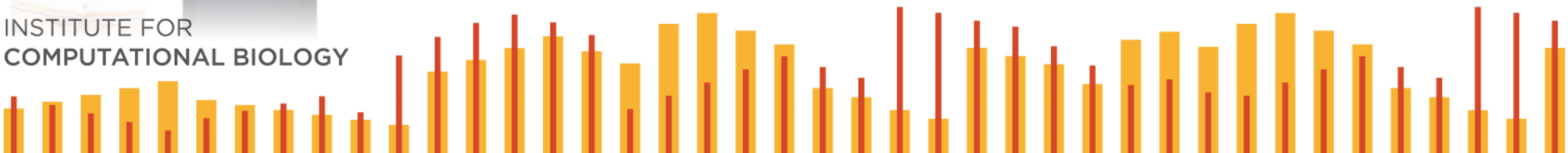


EAGLE BioVU CASE STUDY

Single patient
70 clinic visit dates



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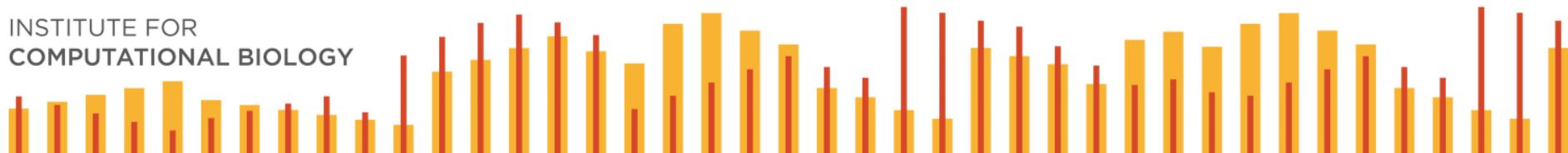
ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

- Step 1: Initial Outlier Detection and Characterization

Calculate and plot raw BMI distributions

Stratify obese from non-obese using ICD-9-CM or ICD-10-CM codes

Extreme outliers in non-obese individuals can be manually inspected



ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

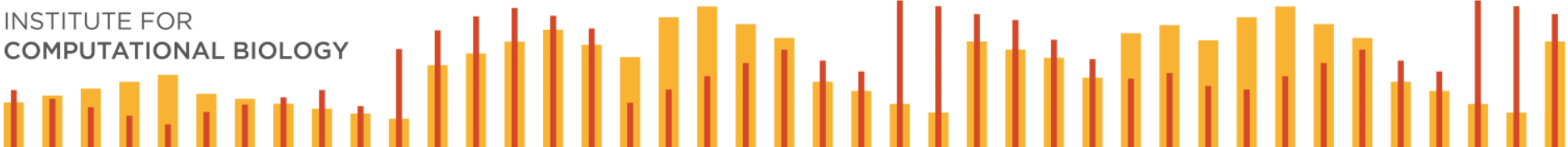
- Step 2: Temporal Partitioning

Divide the smallest two weights of the first three values
by the fourth weight

121.28 kg

126.10 kg

123.23 kg



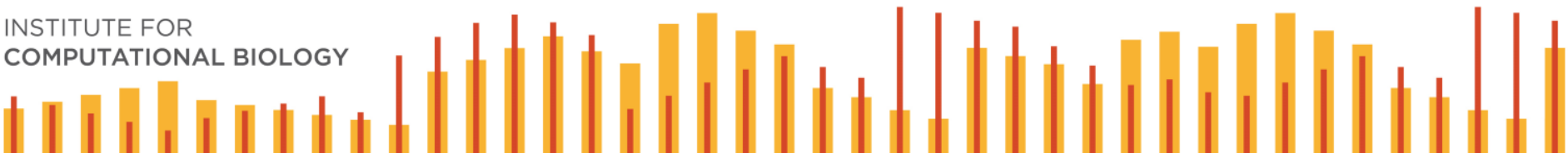
ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

- Step 2: Temporal Partitioning

Establish the “weight index value” closest to 1

$$121.28 \text{ kg} / 123.23 \text{ kg} = 0.984$$

$$126.10 \text{ kg} / 123.23 \text{ kg} = 1.023$$

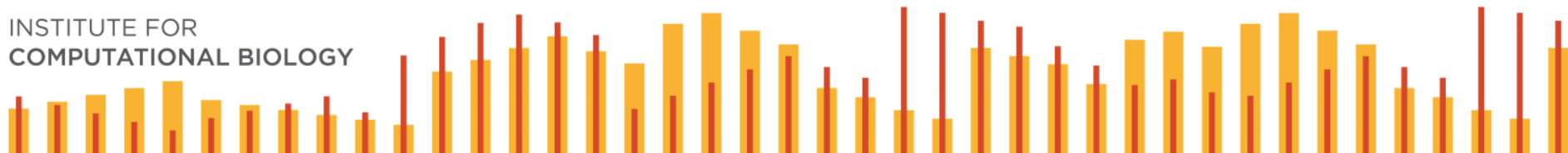


ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

- Step 3: Unit mismatch identification

Divide all measurements with “weight index value”

Generate change ratio distribution



ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

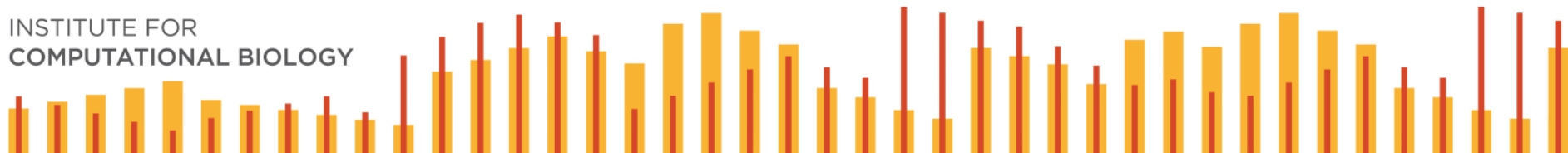
- Step 3: Unit mismatch identification

For weight, if observed value is

within 0.20 SD, value is in kilograms

within 0.45 SD, value is in pounds

within .10 of 0.45, value is in kilograms but assumed pounds and
converted to kilograms (kgx2)

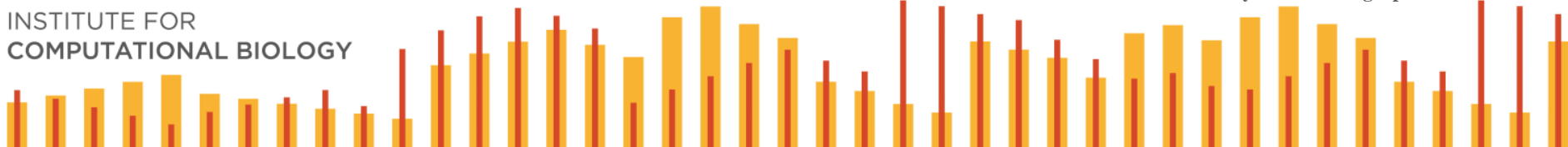
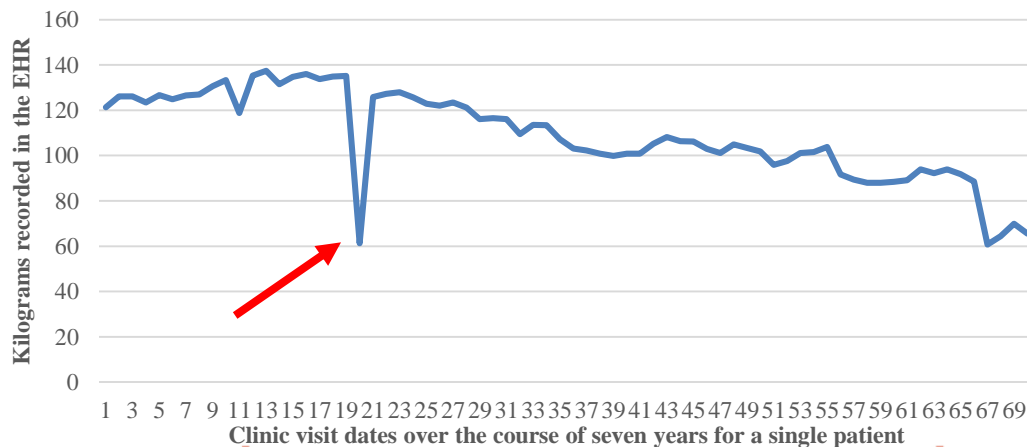


ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

- Step 3: Unit mismatch identification

$$61.23 / 121.28 = 0.50$$

Kg thought to be in lbs
and converted to kg
(kgx2)



ADJACENCY-BASED LONGITUDINAL OUTLIER EXTRACTION (ALOE)

- Step 3: Unit mismatch identification

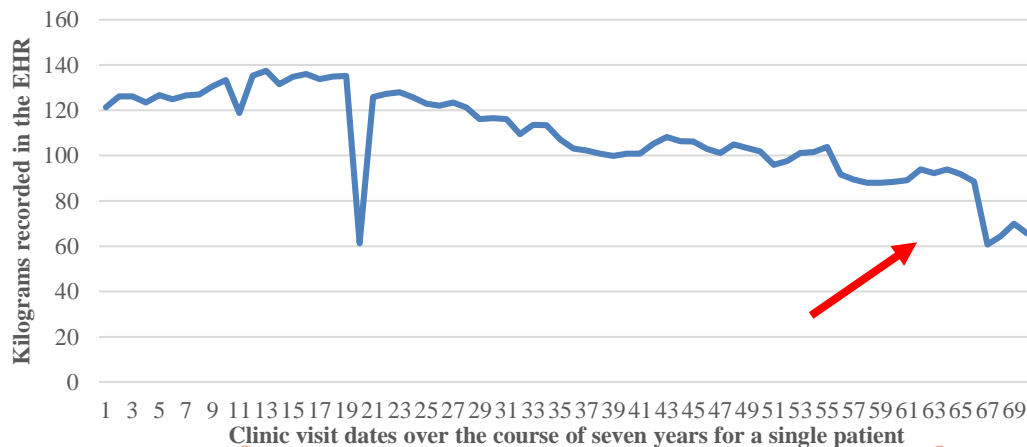
$$60.78 / 121.28 = 0.50$$

$$64.41 / 121.28 = 0.53$$

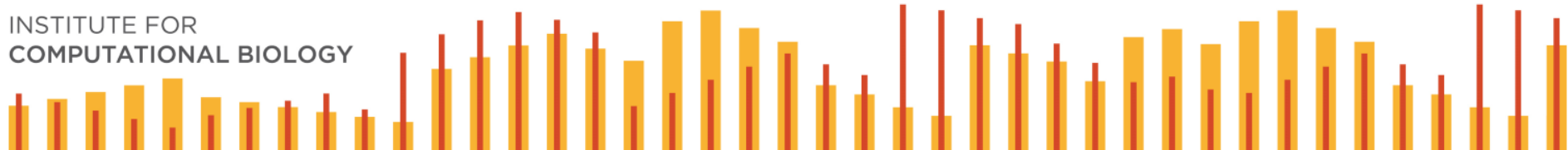
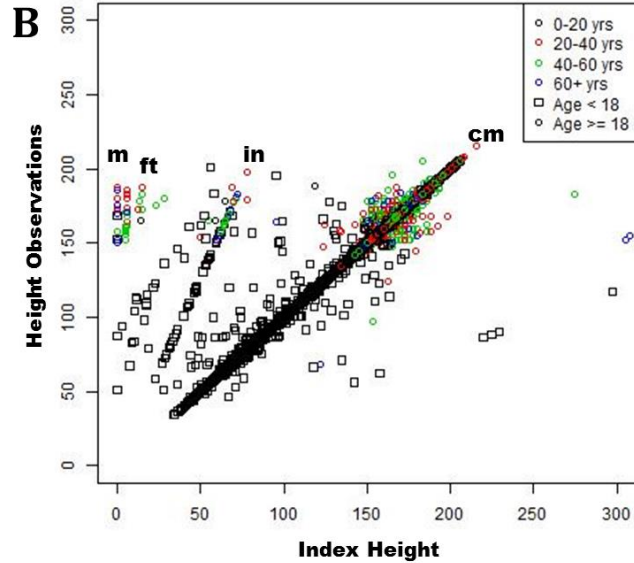
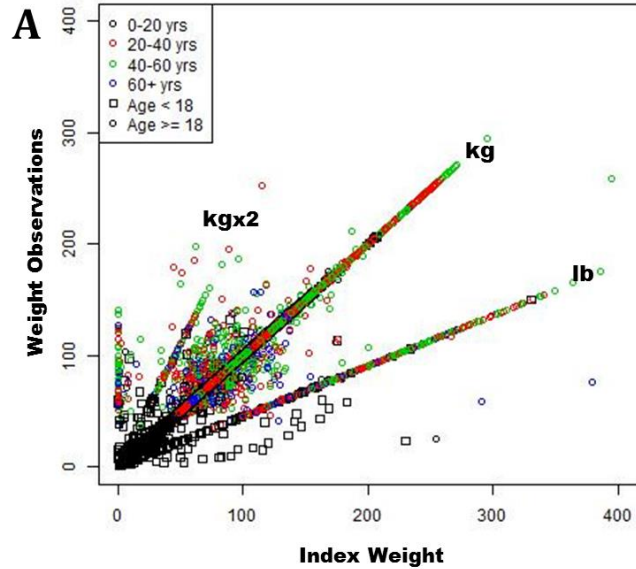
$$69.94 / 121.28 = 0.58$$

$$65.49 / 121.28 = 0.54$$

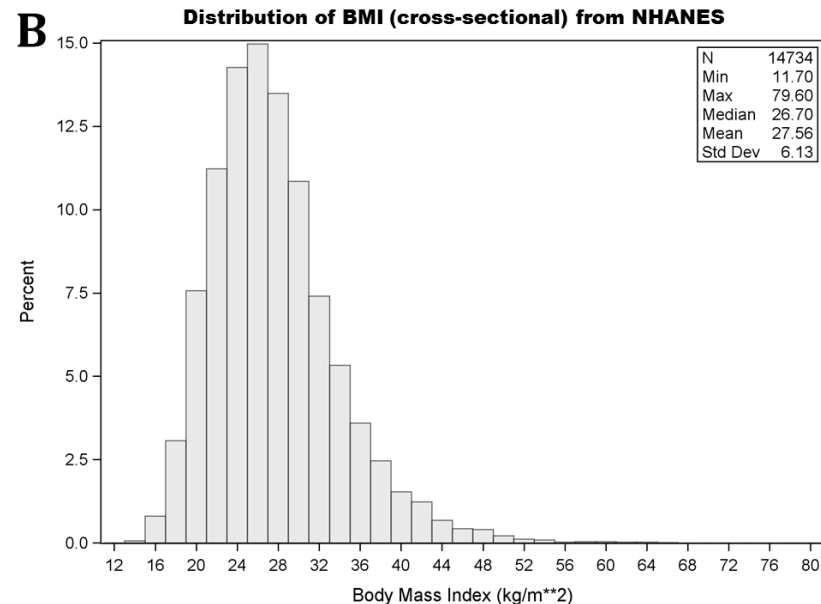
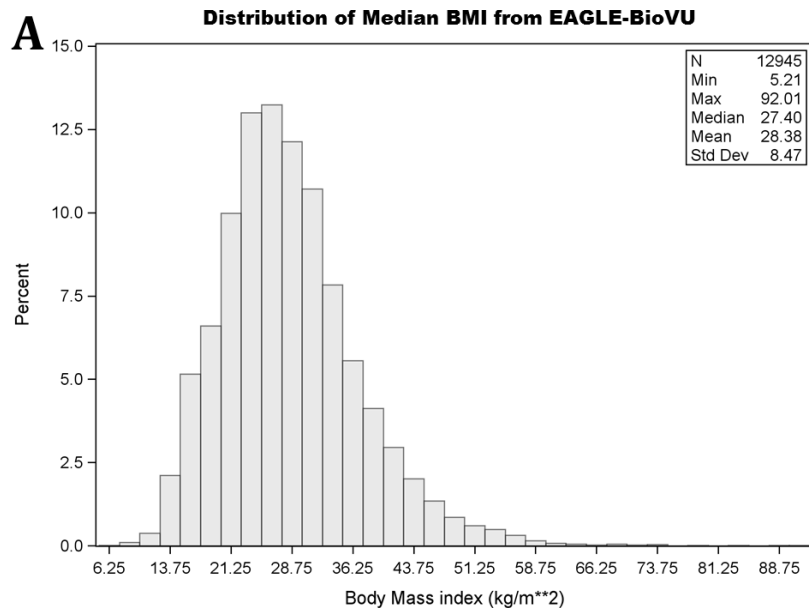
Kg thought to be in lbs and
converted to kg (kgx2)



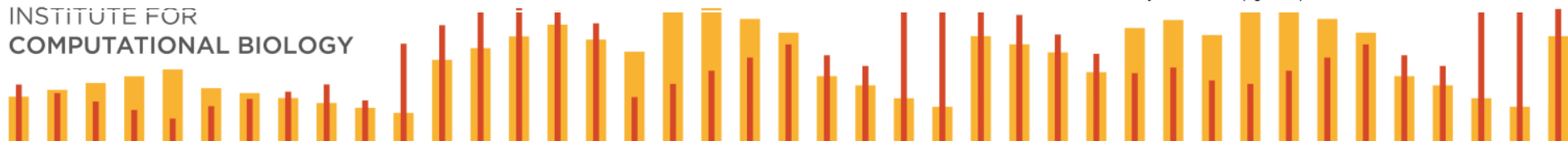
ALOE APPLIED TO EAGLE BIOVU



COMPARISON OF EAGLE BioVU WITH NHANES

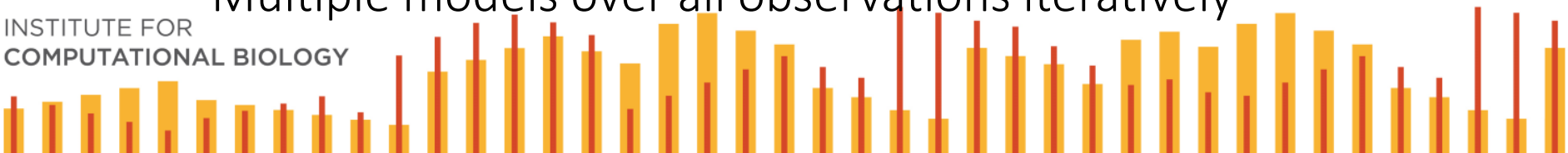


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RESIDUAL MODELING

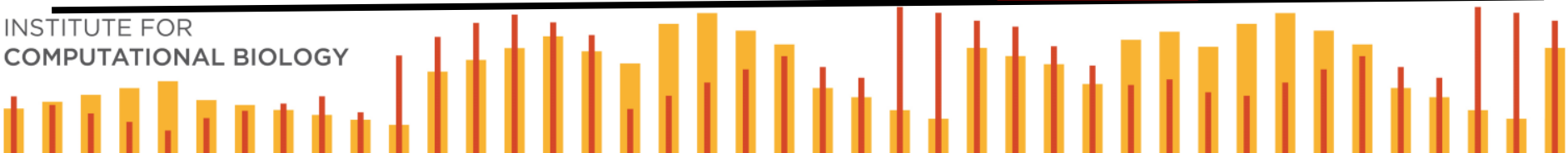
- Exploits relationship between height, weight, and age
 - Regress age onto height and weight, respectively
 - Calculate deviation from predicted value (Cook's Distance, Leverage, DFFits, Studentized residuals, Covariance Ratio)
 - If modeled data has three positive tests, data set to missing
- Model executed two different ways
 - A single model over all observations for an individual
 - Multiple models over all observations iteratively



ALOE VERSUS RESIDUAL MODELING

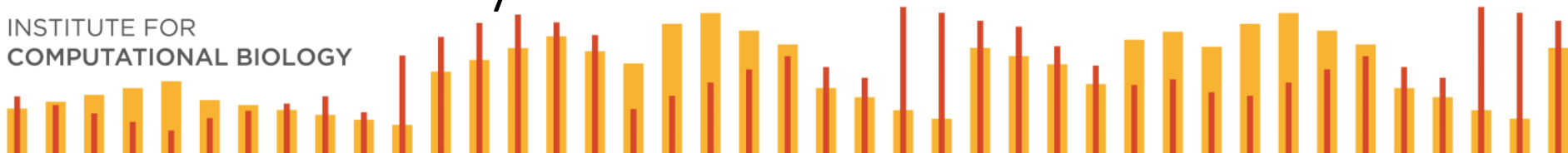
- ALOE retains more data

	RM (all)	RM (individual)	ALOE	Raw Data Total
Weight	155,781 (66%)	226,685 (96%)	230,701 (98%)	235,624
Height	57,707 (51%)	106,424 (94%)	111,536 (99%)	112,862

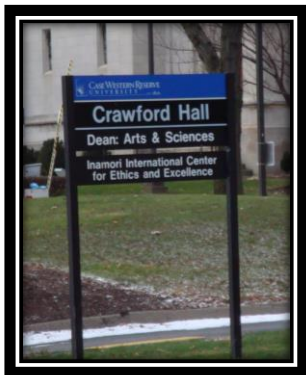


CONCLUSIONS AND DIRECTIONS

- ALOE cleaned BMI data and retained data points
- But relies on dense temporal data with multiple measures
- Manual corrections still necessary and are discretionary



CRAWFORD LAB AND COLLABORATORS

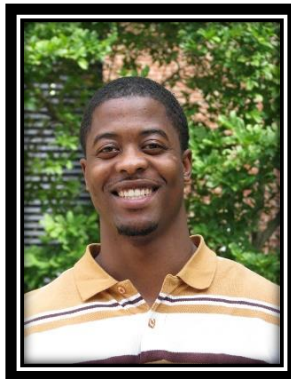


Dana C. Crawford, PhD

Brittany Hollister, PhD candidate

NIH/NHGRI HG004798 (EAGLE)

NCATS 2 UL1 TR000445-06 (BioVU)



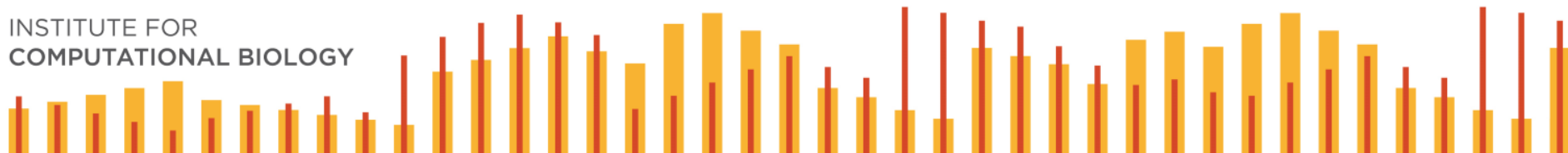
Robert Goodloe,
MS

Williams S. Bush,
PhD, MS



Eric Farber-Eger Jonathan Boston

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CWRU AND CLEVELAND

Biomedical Data Science open rank position available!

<http://epbiwww.case.edu/>



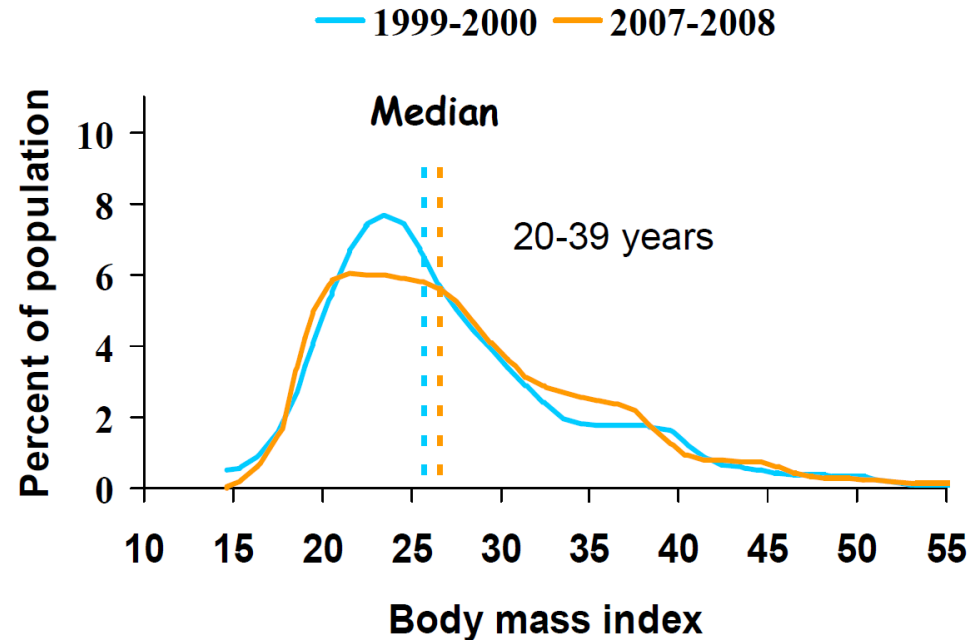
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BODY MASS INDEX, THE VARIABLE

BMI is variable in human populations

US women from NHANES

Flegal et al (2010) *JAMA* 303(3):235-241

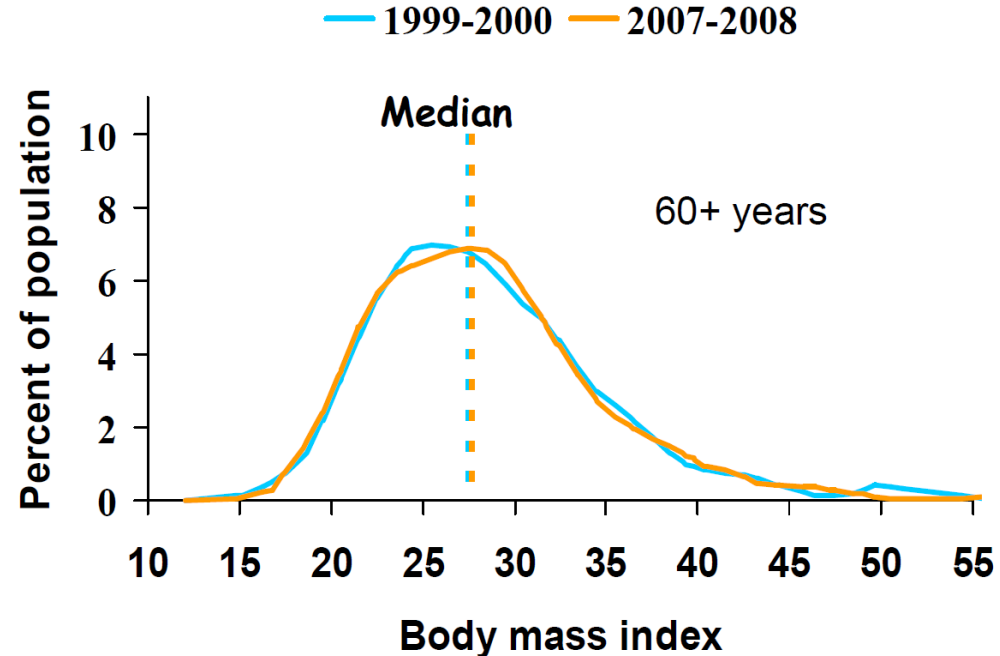


BODY MASS INDEX, THE VARIABLE

BMI changes over the lifecourse

US women from NHANES

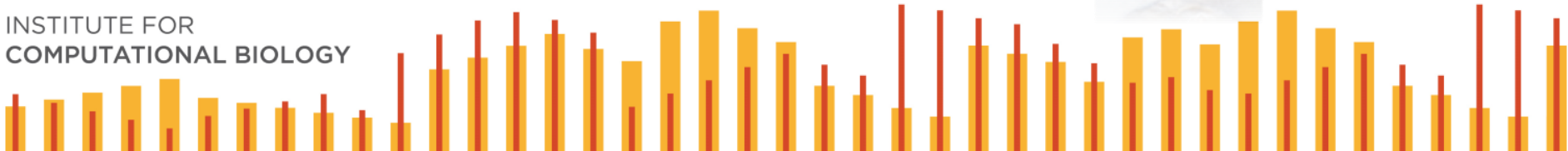
Flegal et al (2010) *JAMA* 303(3):235-241



BODY MASS INDEX, THE VARIABLE

BMI is an important health variable

BMI	WHO classification
<18.5	underweight
18.5-24.9	normal weight
25.0-29.9	overweight
30.0-34.9	class I obesity
35.0-39.9	class II obesity
≥40.0	class III obesity

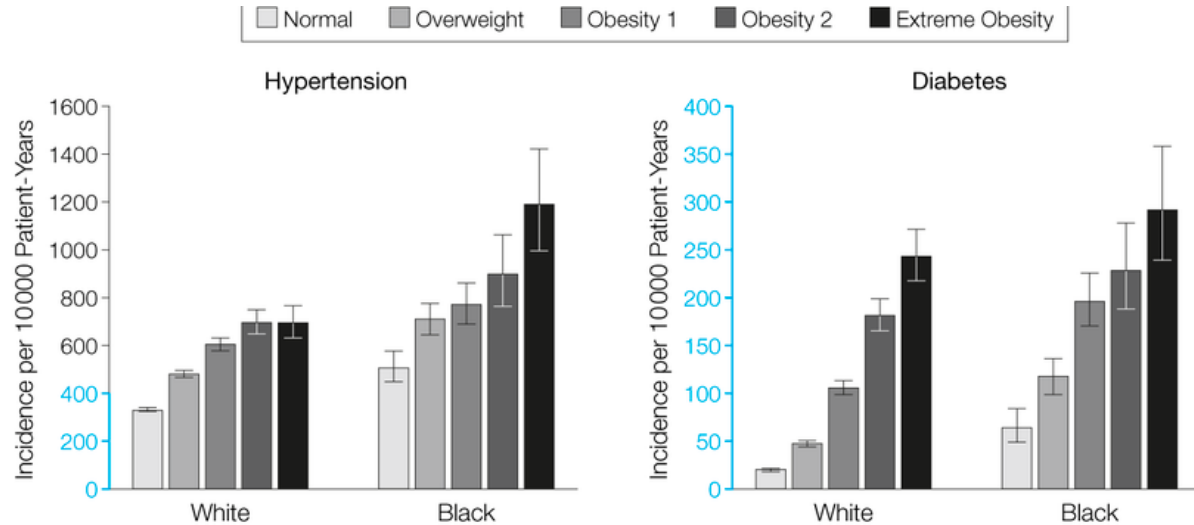


BODY MASS INDEX, THE VARIABLE

BMI is associated with health outcomes

WHI Observational Study

McTigue et al (2006) *JAMA* 296(1):79-86



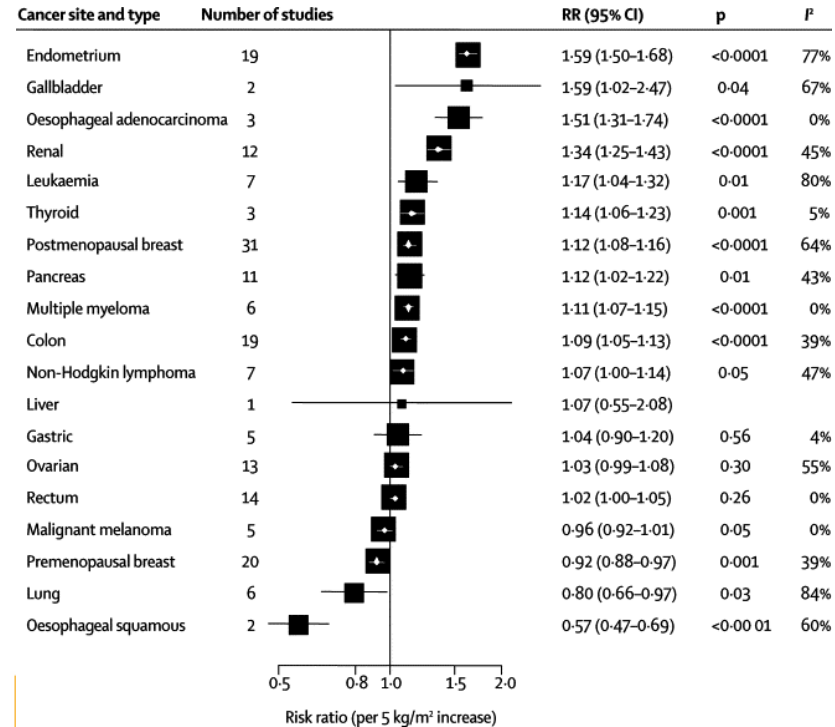
Error bars represent 95% confidence intervals.

BODY MASS INDEX - THE VARIABLE

BMI is associated with health outcomes

Meta-analysis of observational studies

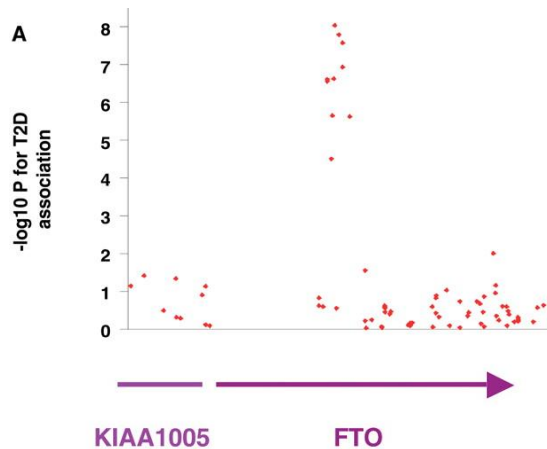
Renehan et al (2008) *Lancet* 371(9612):16-22



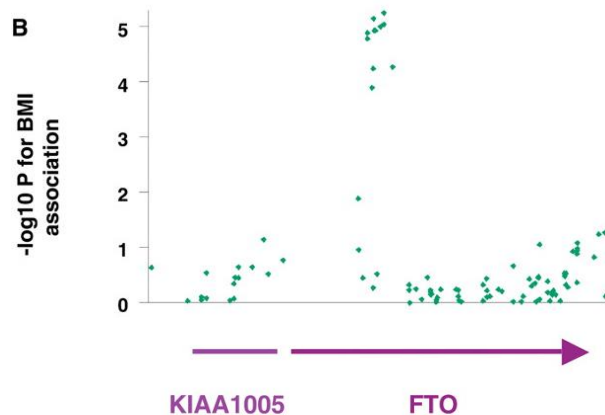
BODY MASS INDEX, THE VARIABLE

BMI is a known mediator

FTO associated with T2D



FTO associated with BMI



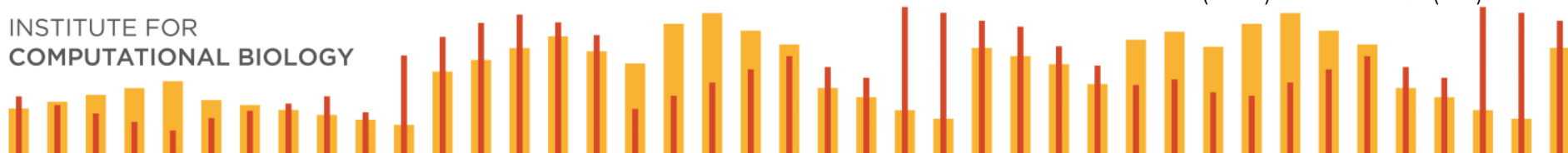
Frayling et al (2007) *Science* 316(5826):89-94

VUMC BioVU IS CLINIC-BASED

	Davidson County (n=626,684)	BioVU (n=162,716)
% female	51.55	51.93
% adults 18-64 years	68.06	57.66
% adults ≥65 years	10.23	24.83
% European American	60.48	81.07
% African American	28.43	8.65
% Hispanic	10.04	1.32
% Asian	3.10	0.83

} Over-represents
European-descent
and elderly

Crawford et al (2015) *Hum Hered* 79(3-4):137-46



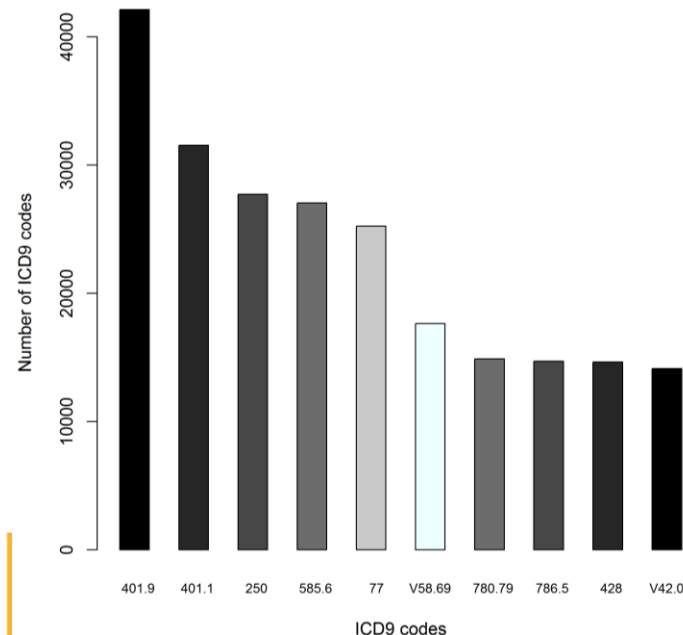
EAGLE BioVU Common Codes

Top 10 codes for African American adults

Hypertension (401.9, 401.1)

Diabetes Mellitus (250)

End-stage renal disease (585.6)



Crawford et al (2015) *Hum Hered* 79(3-4):137-46

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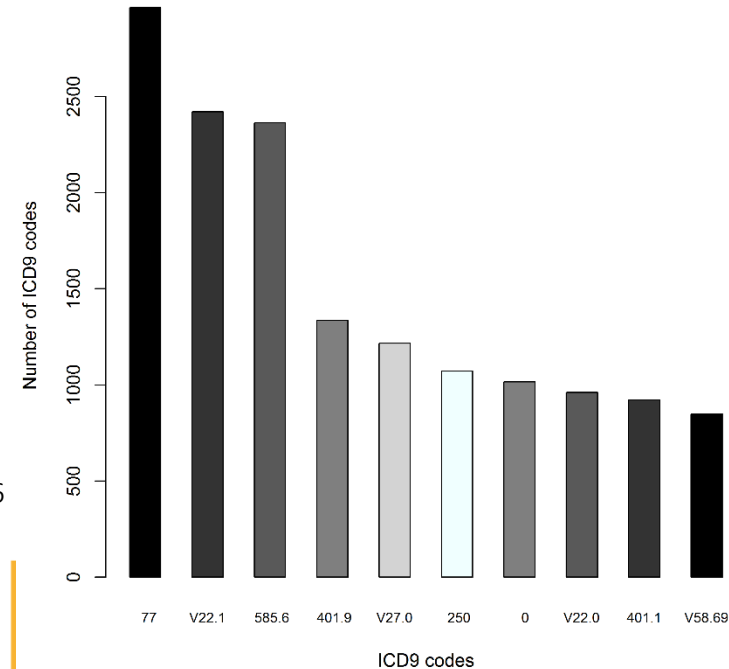
EAGLE BioVU Common Codes

Top 10 codes for Mexican American adults

Sequestrectomy (77)

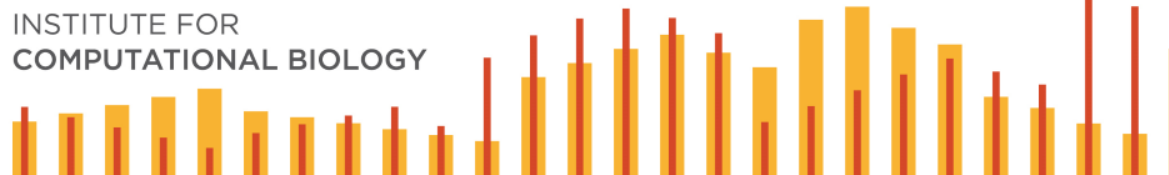
Supervision of other normal
pregnancy (v22.1)

End-stage renal disease (585.6)



Crawford et al (2015) *Hum Hered* 79(3-4):137-46

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RAW EAGLE BIOVU BMI

