

A simple metric to assess the influence of recruitment bias in dementia research-related autopsy cohorts:

Putting it to the test with NACC data

Pete Nelson





**Kathryn
Gauthreaux**

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RESEARCH ARTICLE

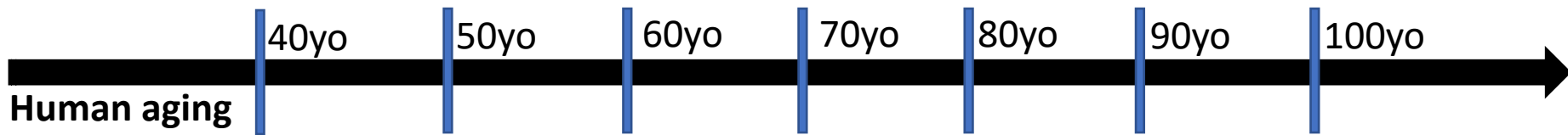
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**Different cohort, disparate results: Selection bias is a key factor
in autopsy cohorts**

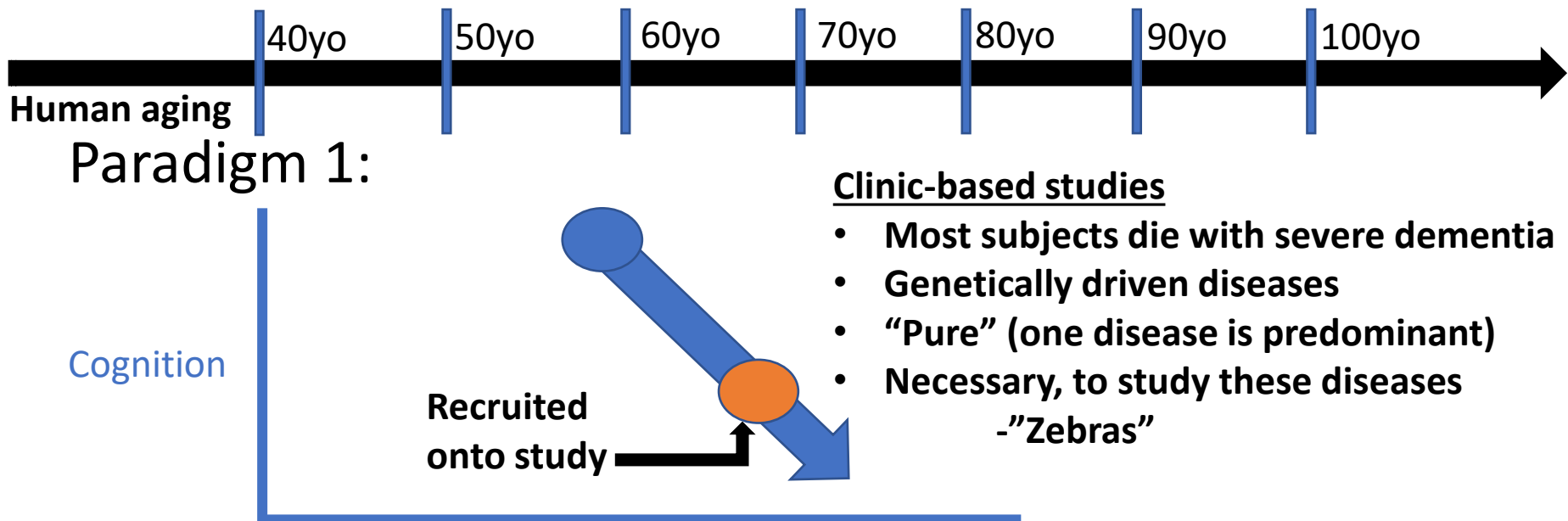
**Kathryn Gauthreaux¹ | Walter A. Kukull¹ | Karin B. Nelson² | Charles Mock¹ |
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Erin L. Abner^{5,6,7} | Peter T. Nelson^{5,8}**

VERY DIFFERENT STUDY DESIGNS...

...can be conceptualized along human aging spectrum



VERY DIFFERENT STUDY DESIGNS...

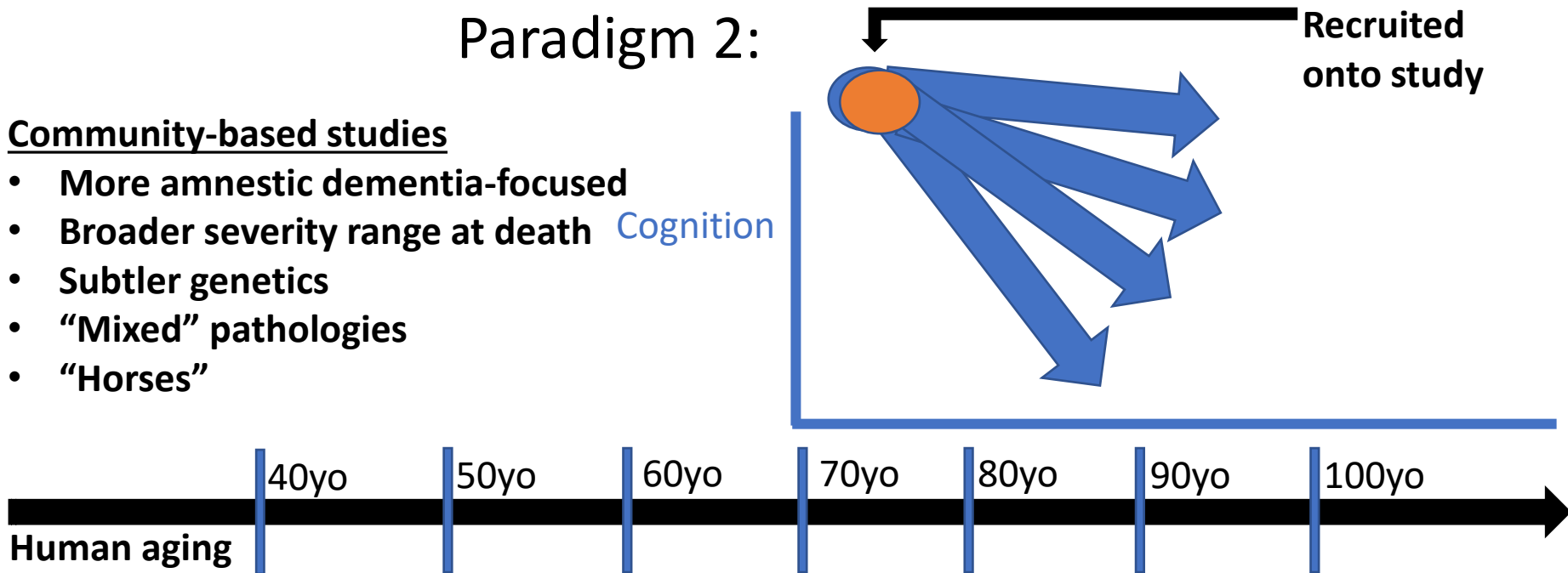


VERY DIFFERENT STUDY DESIGNS...

Paradigm 2:

Community-based studies

- More amnesic dementia-focused
- Broader severity range at death **Cognition**
- Subtler genetics
- “Mixed” pathologies
- “Horses”

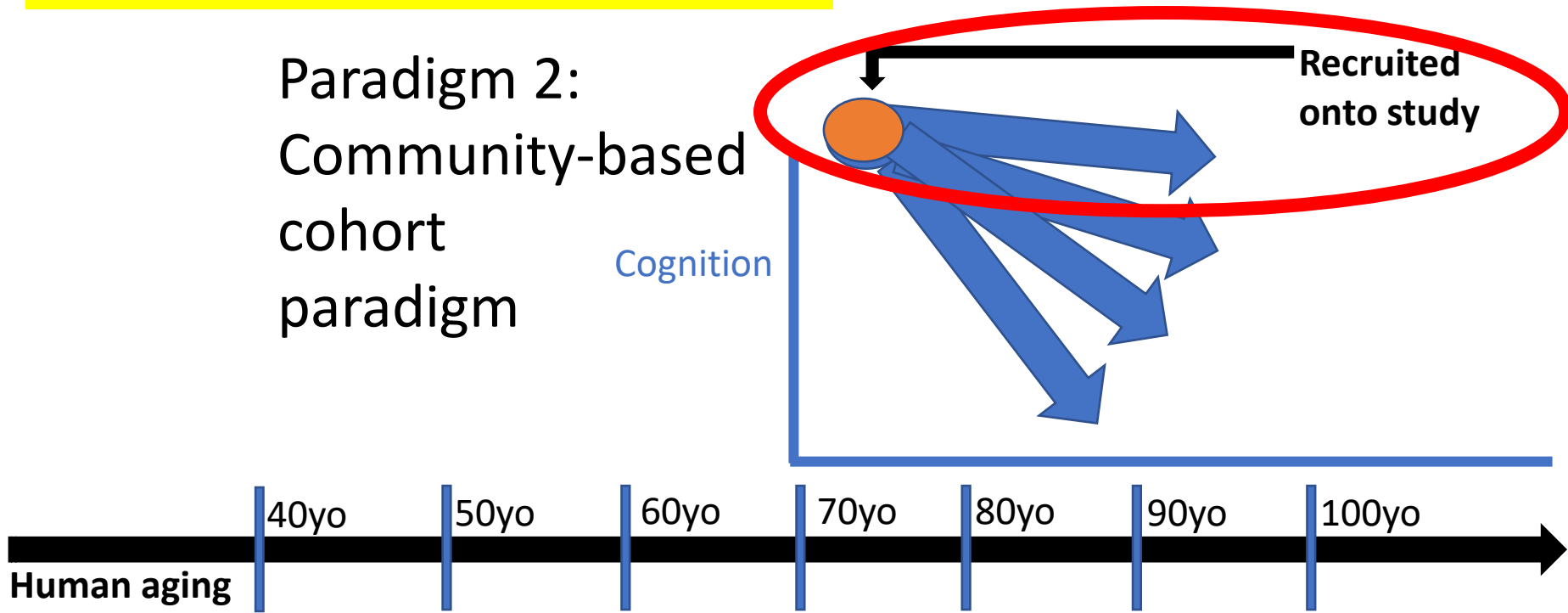


Despite these “extreme” examples,
many study cohorts fall somewhere
betwixt/between

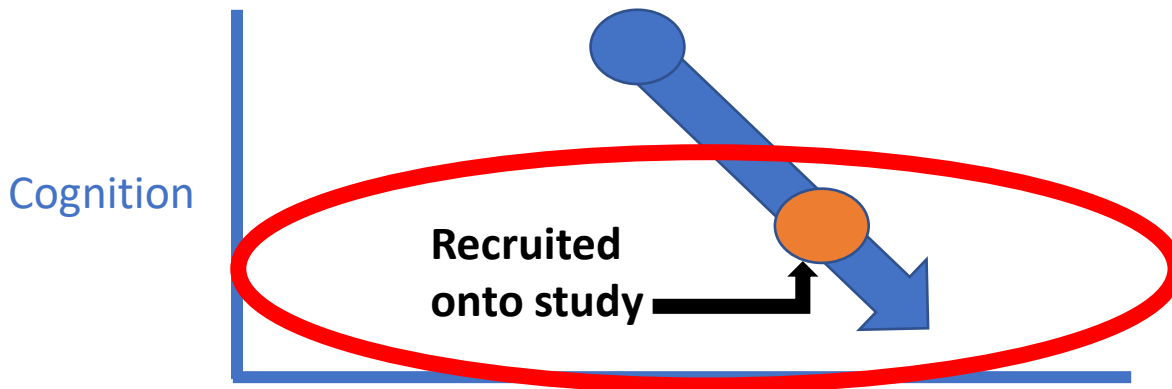


HOW CAN YOU COMPARE?

Paradigm 2:
Community-based
cohort
paradigm



Paradigm 1: Clinic-based cohort paradigm



Different cohort, disparate results: Selection bias is a key factor in autopsy cohorts

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NACC:

**A great context to study
different cohort types**

28 ADRCs met inclusion criteria
>30 study participants each

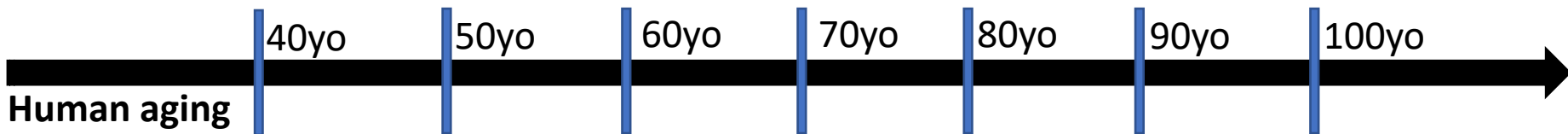
- Followed longitudinally
- Came to autopsy
- Detailed data available

HOW TO COMPARE THE PARADIGMS?

Paradigm 2:

Cognition

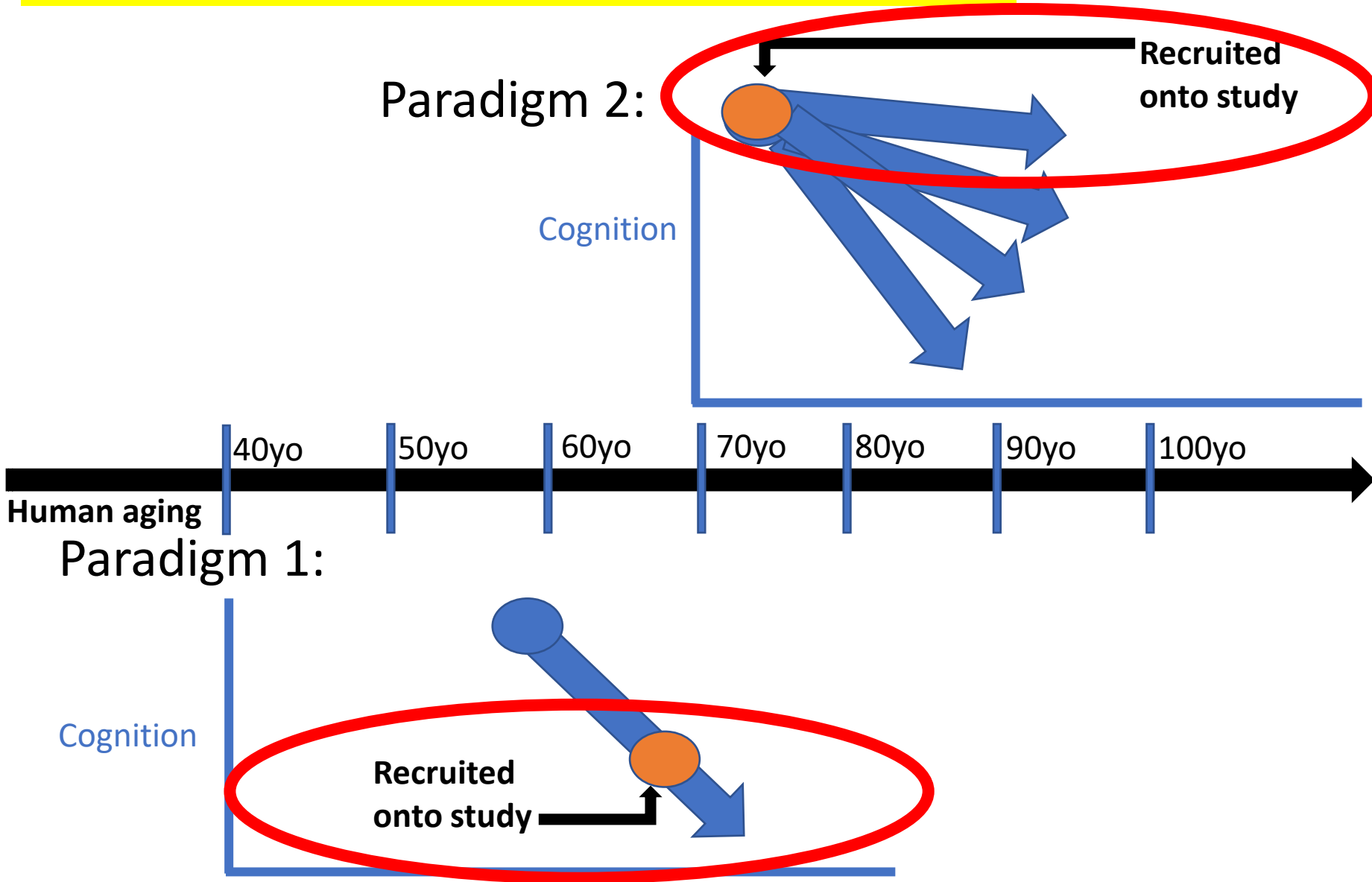
Recruited onto study



Paradigm 1:

Cognition

Recruited onto study



NACC:

28 ADRCs met inclusion criteria
(Avg~150 cases each)

- 8 ADRCs: More normal (>30% normal) at recruitment
- 10 ADRCs: Middle (15-30% normal) at recruitment
- 10 ADRCs: More impaired (<15% normal) at recruitment

NACC:

28 ADRCs met inclusion criteria (>30 cases each)

- 8 ADRCs: More normal (>30% normal at recruitment)
- 10 ADRCs: Middle (15-30% normal at recruitment)
- 10 ADRCs: More impaired (<15% normal at recruitment)



NACC:

28 ADRCs met inclusion criteria (>30 cases each)

- 8 ADRCs: More normal (>30% normal at recruitment)
- 10 ADRCs: Middle (15-30% normal at recruitment)
- 10 ADRCs: More impaired (<15% normal at recruitment)



- Predominantly Caucasian
- Predominantly high-SES
- AD-oriented

Clinic-
based



Community-
based

28 ADRCs, and over 4000 participants, included

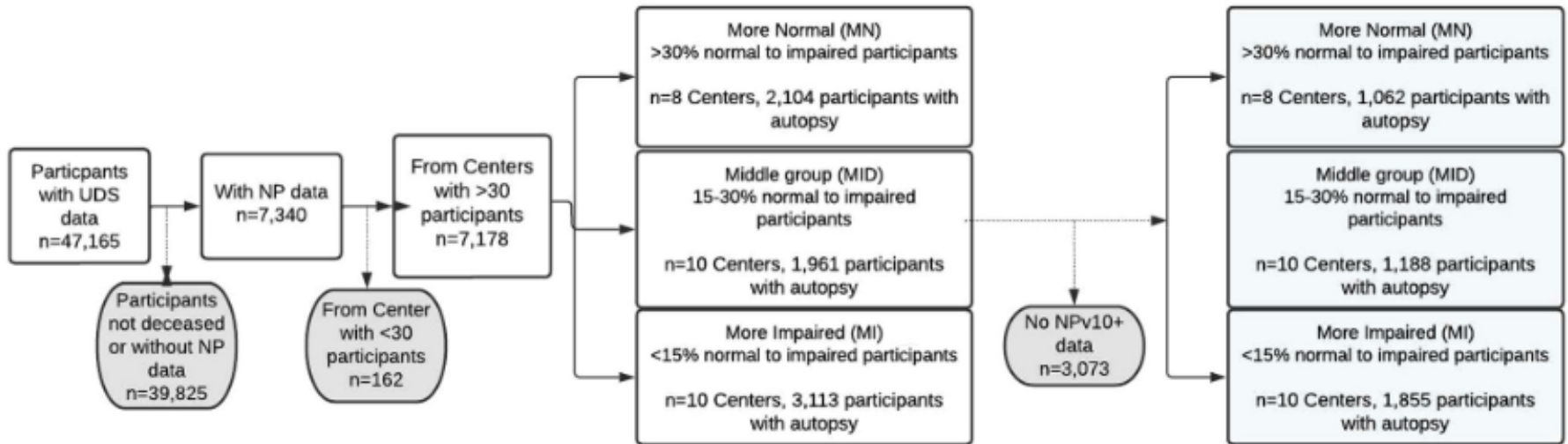


FIGURE 1 Study cohort exclusions, inclusions, sample sizes, and delineation of study groups. NP, National Alzheimer's Coordinating Center Neuropathology form; UDS, Uniform Data Set

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2.2 | Statistical analyses

To compare the demographic characteristics, clinical profiles, and neuropathologic features between the M-Norm, M-Imp, and Mid groups, we report both descriptive statistics (mean proportion, min, max) and *P*-values from trend test using regression models. The analysis was done at the center level; thus, all statistics are reported for the ADRCs within each group. Continuous measures were compared with linear models, while counts were compared using Poisson models with a log offset for the number of participants with autopsy in each ADRC. Robust standard errors were used for the calculation of *P*-values, which allows non-constant variance across groups and overdispersion of count outcomes. Participant characteristics examined included

Thanks to Drs Yen-Chi Chen, Gary Chan, Dave Fardo, Yuri Katsumata, & Erin Abner

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TABLE 1 Demographic characteristics among participants with NP data available.

| Center group, n | More impairment (N/MCI&D < 15%) n = 10 centers, 3113 participants with autopsy | Middle group (N/MCI&D 15%–30%) n = 10 centers, 1961 participants with autopsy | More normal (N/MCI&D > 30%) n = 8 centers, 2104 participants with autopsy | P-values |
|---|--|---|---|----------|
| Number of unique 3-digit zip codes in ADRC, mean (min, max) | 34.6 (0, 101) | 16.8 (0, 26) | 23.1 (0, 66) | 0.22 |
| Average number of UDS visits, mean (min, max) | 3.6 (2.5, 4.7) | 4.5 (3.4, 5.7) | 4.5 (3, 5.5) | 0.01 |
| Average follow-up time (in years), mean (min, max) | 3.0 (1.5, 4.2) | 4.1 (2.9, 5) | 4.1 (2.5, 5.5) | <0.01 |
| % deceased participants with autopsy, mean % (min%, max%) | 60.0 (33.7, 86.3) | 60.5 (42.2, 76.0) | 62.8 (47.3, 82.2) | 0.68 |
| Age at baseline, mean (min, max) | 72.1 (66.2, 78.3) | 74.9 (69.6, 78.5) | 79.3 (75.1, 83.9) | <0.01 |
| Age at death, mean (min, max) | 77.1 (70.7, 83.8) | 80.3 (75.4, 84.4) | 85.1 (80.0, 89.8) | <0.01 |
| Sex, mean % (min%, max%) | | | | |
| Male | 58.9 (51.2, 66.7) | 54.7 (47.3, 66.7) | 46.7 (38.8, 56.9) | <0.01 |
| Female | 41.1 (33.3, 48.9) | 45.4 (33.3, 52.7) | 53.3 (43.1, 61.2) | <0.01 |
| Race, mean % (min%, max%) | | | | |
| White | 94.4 (89.1, 97.4) | 91.6 (85.4, 96.6) | 88.0 (43.9, 97.8) | 0.46 |
| Black or African American | 2.4 (0.6, 4.8) | 4.4 (0.5, 10.2) | 8.2 (0.0, 45.9) | 0.09 |
| Other | 2.1 (0.7, 3.4) | 3.5 (1.2, 8.8) | 3.7 (1.6, 9.2) | 0.32 |
| APOE ε4, mean % (min%, max%) | | | | |
| ε4 carrier | 39.7 (22.2, 53.8) | 40.4 (16.7, 54.7) | 35.7 (26, 46.3) | 0.36 |
| No ε4 allele | 42.4 (14.3, 58) | 46.3 (22.5, 56.9) | 55.7 (40.8, 71.8) | 0.01 |
| Missing/unknown/ not assessed | 17.9 (1.2, 63.5) | 13.3 (0.0, 60.9) | 8.7 (0.5, 21.7) | 0.09 |

Demographics and genetics:

ADRCs which recruited higher % normal:

Longer followup, older at recruitment and death, more female

Lower APOE ε4 allele

Clinical findings:

ADRCs which recruited
higher % **impaired**:
More DLB, FTD

ADRCs which recruited
higher % **normal**:
More CVD

AD—not different!

TABLE 2 Primary clinical diagnoses at initial and last standardized Uniform Data Set visit among participants with NP data available.

| Center group, n | More impairment (N/MCI&D < 15%) n = 10 centers, 3113 participants with autopsy | Middle group (N/MCI&D 15-30%) n = 10 centers, 1961 participants with autopsy | More normal (N/MCI&D > 30%) n = 8 centers, 2104 participants with autopsy | P-values |
|---|--|--|---|----------|
| Initial UDS visit | | | | |
| Primary clinical diagnoses, mean % (min%, max%) | | | | |
| AD | 51 (30.7, 69.2) | 54.5 (28.2, 82.8) | 41.3 (22.3, 57.1) | 0.16 |
| DLB | 8.8 (2.3, 20.1) | 4.2 (1.2, 7.4) | 1.6 (0.0, 3.8) | <0.01 |
| CVD | 0.8 (0.0, 3.3) | 1.5 (0.0, 5.5) | 1.3 (0.0, 3.4) | 0.12 |
| FTD ^a | 17.5 (3.1, 42.2) | 9.5 (0.0, 38.9) | 3.4 (0.0, 8.2) | <0.01 |
| Clinical diagnoses, ^b mean % (min%, max%) | | | | |
| AD | 54.8 (37.9, 72.8) | 59.0 (44.7, 82.8) | 43.1 (23.5, 60.2) | 0.08 |
| DLB | 11.5 (3.1, 24.3) | 6.0 (1.2, 10.1) | 4.5 (2.1, 6.0) | <0.01 |
| CVD | 3.3 (0.0, 8.0) | 4.6 (0.9, 10.7) | 3.7 (1.3, 6.6) | 0.90 |
| FTD | | | | |
| Any FTD ^a | 21.4 (3.6, 53.7) | 12.7 (1.2, 53.2) | 5.0 (1.9, 10.2) | <0.01 |
| bvFTD | 10.7 (2.6, 20.7) | 6.6 (0.0, 23.3) | 3.5 (1.3, 9.2) | <0.01 |
| PPA | 6.8 (1.0, 15.4) | 5.2 (0.0, 30.2) | 1.4 (0.0, 2.7) | <0.05 |
| Last UDS visit | | | | |
| Primary clinical diagnoses, mean % (min%, max%) | | | | |
| AD | 57.9 (34.3, 76.2) | 64.1 (37.1, 85.1) | 58.7 (43.2, 66.8) | 0.56 |
| DLB | 10.2 (3, 22.5) | 6.0 (2.2, 11.6) | 2.9 (0.0, 5.8) | <0.01 |
| CVD | 1.2 (0.0, 3.1) | 2.3 (0.0, 10.2) | 4.3 (1.0, 7.4) | <0.01 |
| FTD ^a | 16.9 (1.5, 40.7) | 10.1 (0.0, 37.6) | 4.0 (0.9, 8.2) | <0.01 |
| Clinical diagnoses, ^b mean % (min%, max%) | | | | |
| AD | 62.3 (41.2, 80) | 69.2 (48.8, 88.5) | 61.9 (45.3, 69.8) | 0.65 |
| DLB | 15.0 (4.1, 30.8) | 10.2 (6.8, 15.4) | 6.6 (3.2, 9.8) | <0.01 |
| CVD | 5.0 (0.0, 11.9) | 8.9 (4.0, 19.1) | 10.0 (4.1, 16.7) | <0.01 |
| FTD | | | | |
| Any FTD ^a | 21.6 (3.1, 53.9) | 13.2 (1.2, 53.5) | 5.0 (1.9, 9.6) | <0.01 |
| bvFTD | 10.7 (1.5, 22.0) | 6.4 (0.0, 20.1) | 3.3 (0.4, 8.2) | <0.01 |
| PPA | 7.3 (1.0, 16) | 4.8 (0.0, 29.7) | 1.1 (0.0, 2.9) | 0.03 |

TABLE 3 Neuropathological features.

| Center group, n | More impairment (N/MCI&D < 15%) n = 10 centers, 3113 participants | Middle group (N/MCI&D 15-30%) n = 10 centers, 1961 participants | More normal (N/MCI&D > 30%) n = 8 centers, 2104 participants | P-values |
|--|--|--|---|----------|
| Alzheimer's disease pathology, mean % (min%, max%) | | | | |
| Pathological NIA-AA criteria ^b | | | | |
| No ADNC | 14.2 (6.6, 25.7) | 11.1 (4.3, 17.4) | 13.0 (6.9, 19.5) | 0.47 |
| Low ADNC | 14.5 (0.0, 30.0) | 14.9 (7.5, 26.3) | 18.5 (10.0, 28.5) | 0.48 |
| Intermediate ADNC | 16.2 (10, 30.7) | 23.1 (10.1, 52.5) | 28.3 (9.8, 40) | <0.01 |
| High ADNC | 50.0 (34.7, 68.2) | 48.9 (23.2, 67.5) | 39.7 (28.8, 54.1) | <0.05 |
| Braak stage | | | | |
| 0 | 9.5 (1.3, 27.1) | 3.7 (0.0, 12.6) | 4.2 (0.5, 10.4) | <0.01 |
| I | 7.7 (3.1, 17.7) | 6.6 (3.5, 11.4) | 6.6 (1.9, 13.2) | 0.49 |
| II | 10.1 (3.3, 18.9) | 10.6 (6.5, 14.8) | 11.6 (2.2, 19.2) | 0.27 |
| III | 9.5 (5.5, 20.6) | 9.4 (4.9, 16.4) | 13.4 (7.6, 31.6) | 0.01 |
| IV | 10.6 (7.3, 19.1) | 10.3 (7.2, 13.5) | 18.5 (4.7, 35.3) | 0.01 |
| V | 15.6 (5.8, 25.4) | 21.4 (12.7, 37) | 20.4 (5.7, 36.2) | 0.06 |
| VI | 34.5 (12.7, 51.8) | 36.7 (17.1, 51.8) | 24.6 (12.2, 56.6) | <0.01 |
| CERAD score | | | | |
| None | 23.9 (9.5, 37.3) | 17.6 (12.2, 26.2) | 23.9 (12.1, 33.9) | 0.89 |
| Sparse | 10.3 (2.6, 19.5) | 8.8 (2.8, 18.0) | 17.4 (4.2, 45.4) | 0.01 |
| Moderate | 18.3 (9.5, 34.9) | 19.1 (3.8, 35.1) | 19.0 (5.8, 42.4) | 0.71 |
| Frequent | 46.7 (23.8, 73.4) | 54.2 (34.3, 69.8) | 39.7 (6.6, 59.0) | 0.16 |

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2023*

Pathology findings (AD neuropathologic changes/ADNC):

ADRCs which recruited higher % **impaired**:

More Braak NFT stages 0, VI

ADRCs which recruited higher % **normal**:

More Braak NFT stages III, IV

TABLE 3 Neuropathological features.

| Center group, n | | | | |
|---|--|--|---|----------|
| Characteristic, mean % (min%, max%) | More impairment (N/MCI&D < 15%) n = 10 centers, 3113 participants | Middle group (N/MCI&D 15-30%) n = 10 centers, 1961 participants | More normal (N/MCI&D > 30%) n = 8 centers, 2104 participants | P-values |
| FTLD-TDP-43 ^b | 7.9 (0.0, 20.8) | 4.3 (0.0, 14.6) | 2.5 (0.0, 5.0) | 0.01 |
| FTLD-tau ^b | 14.4 (0.0, 39.0) | 10.2 (1.0, 29.5) | 15.9 (0.0, 54.6) | 0.97 |
| Argyrophilic grains ^b | 2.4 (0.0, 7.89) | 2.4 (0.0, 11.74) | 6.1 (0.0, 24.68) | 0.72 |
| Cerebrovascular pathology, mean % (min%, max%) | | | | |
| Infarcts or lacunes | 15.2 (8.5, 30.1) | 15.6 (9.0, 31.6) | 23.2 (10.4, 41.8) | 0.03 |
| Microinfarcts | 17.2 (3.1, 32.1) | 19.6 (8.4, 37.8) | 21.4 (5.3, 31.7) | 0.13 |
| Mod/severe arteriolosclerosis | 46.5 (15.9, 76.7) | 41.0 (8.7, 72.4) | 22.0 (0.0, 45.7) | 0.09 |

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Pathology findings (non-ADNC):

ADRCs which recruited higher % **impaired**:
More FTLD-TDP

ADRCs which recruited higher % **normal**:
More infarcts or lacunes

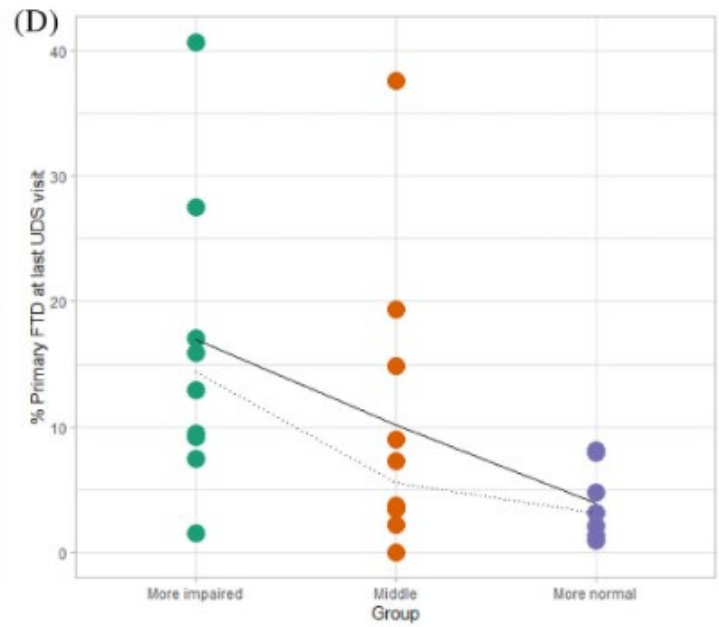
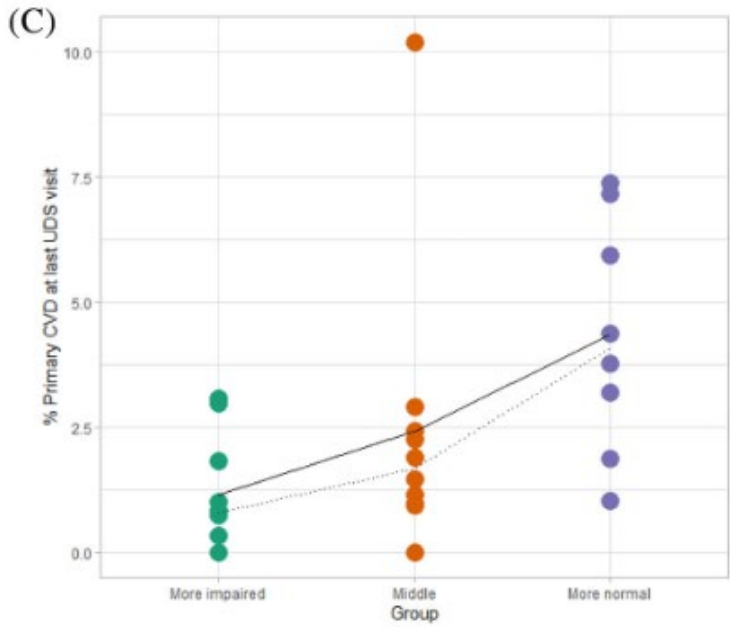
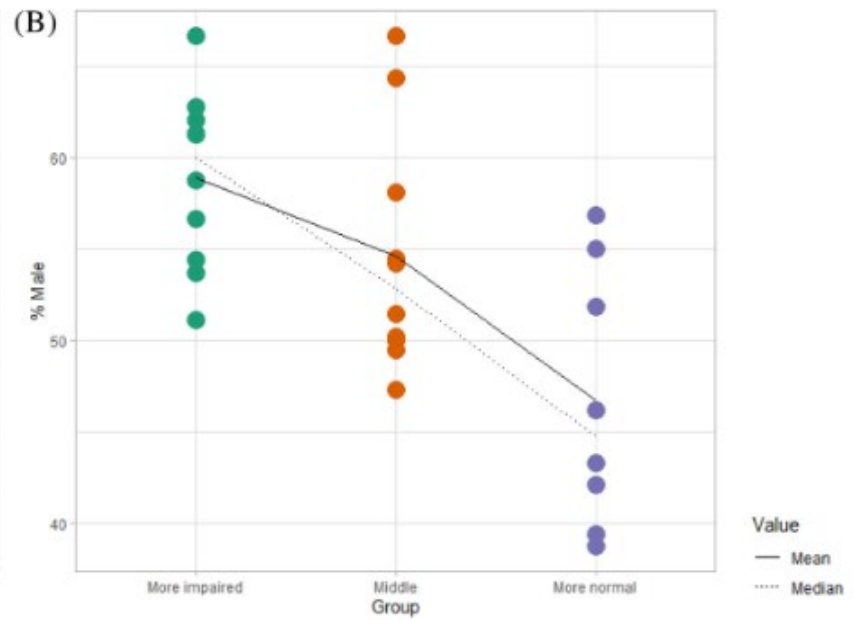
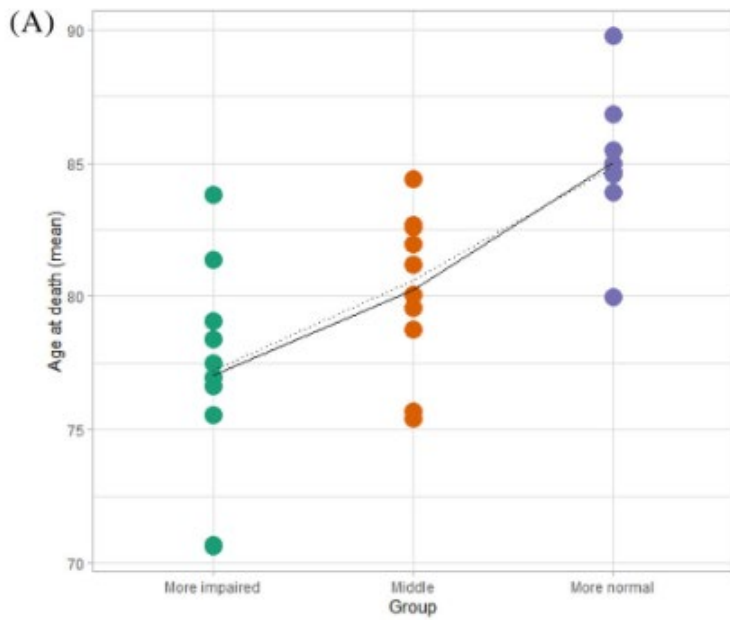


TABLE 4 Summary of findings.

| Cohort recruitment criteria | Findings |
|---|---|
| More participants recruited while cognitively normal | Older participants (at intake and death) |
| | Longer study follow-up |
| | More females |
| | More CVD clinically |
| | More infarcts at autopsy |
| | More intermediate ADNC |
| More participants recruited while cognitively impaired | Younger participants |
| | More males |
| | More APOE ϵ 4 (though not statistically significant at the $p=0.05$ level) |
| | More severe ADNC |
| | More clinical DLB at last UDS visit |
| More clinical FTD at last UDS visit and FTLD-TDP at autopsy | |

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2023*

Conclusion:

For an autopsy cohort used in dementia research, the % of subjects cognitively normal at recruitment into the study is a measure (related to ascertainment bias) associated with a set of clinical and pathological observations.

This parameter may also provide a proxy for where a cohort falls along the clinic/community-based spectrum of study designs.

THANK
YOU!

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RESEARCH ARTICLE

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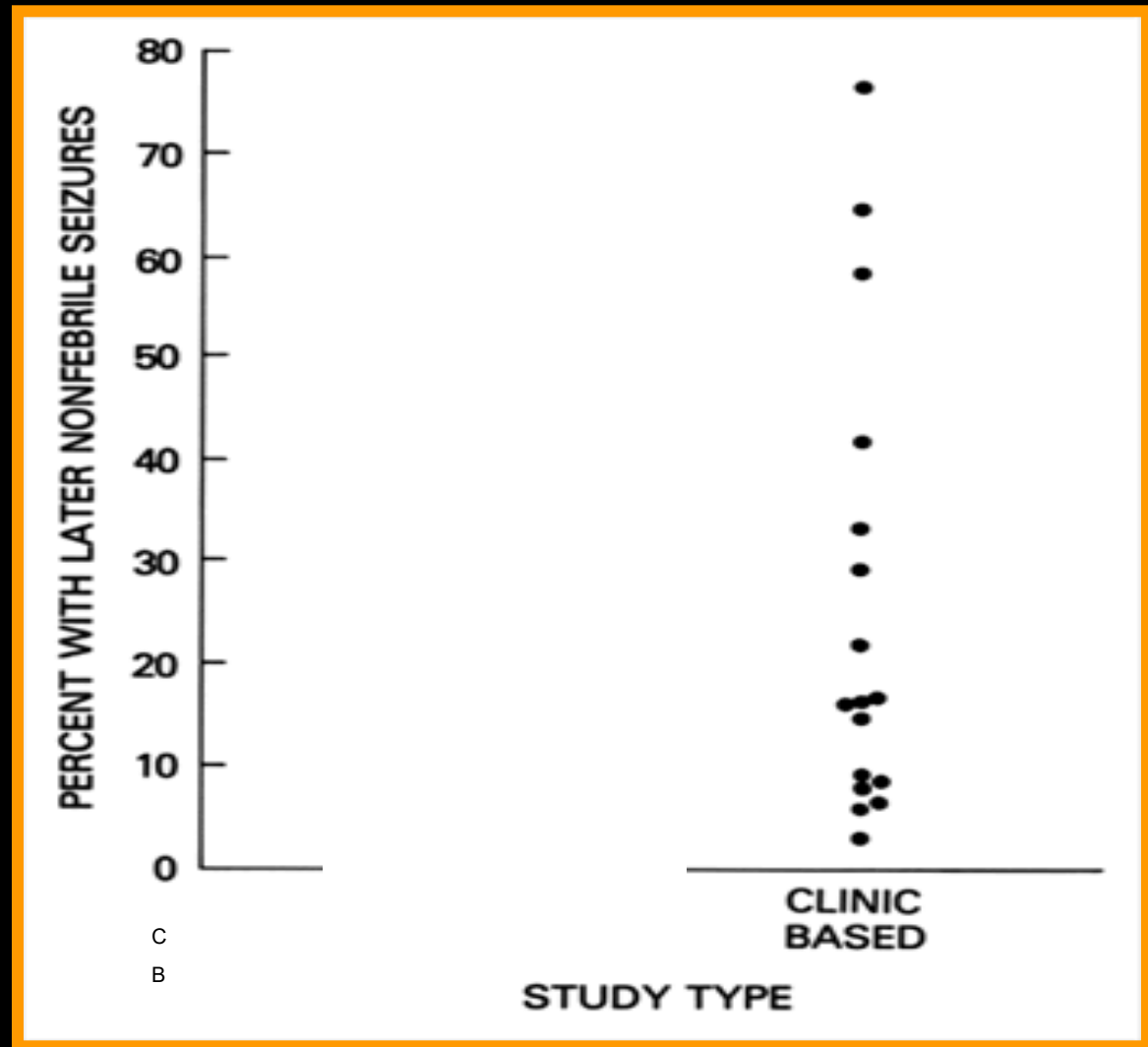
Different cohort, disparate results: Selection bias is a key factor in autopsy cohorts

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Population based vs. clinical samples

% Febrile →
Nonfebrile
seizures:

Give
phenobarbital?



Average Stanford–Binet IQ Scores at the Two-Year Visit, According to Expected IQ Level and Treatment Group.

Table 2. Average Stanford–Binet IQ Scores at the Two-Year Visit, According to Expected IQ Level and Treatment Group.

| EXPECTED IQ LEVEL* | PHENOBARBITAL GROUP | | | PLACEBO GROUP | | |
|---------------------------|---------------------|-------------|------------|------------------|-------------|------------|
| | NO. AT BASE LINE | NO. AT 2 YR | AVERAGE IQ | NO. AT BASE LINE | NO. AT 2 YR | AVERAGE IQ |
| 1 | 24 | 11 | 85.64 | 19 | 11 | 97.73 |
| 2 | 17 | 11 | 92.00 | 26 | 21 | 98.00 |
| 3 | 22 | 17 | 97.47 | 21 | 17 | 105.35 |
| 4 | 21 | 18 | 97.33 | 22 | 20 | 107.65 |
| 5 | 24 | 20 | 114.30 | 21 | 20 | 115.70 |
| Total | 108 | 77 | — | 109 | 89 | — |
| Horvitz–Thompson average† | — | — | 95.54 | — | — | 103.95 |

*The higher the number, the higher the expected Stanford–Binet IQ score of the children at that level.

†The Horvitz–Thompson average is the weighted average of the averages for each level. The weight is the inverse of the proportion of the number of observations. For example, the weight for the third level of the placebo group is 21 divided by 17. The Horvitz–Thompson average of the placebo group is greater by 8.41, and its t statistic is 3.48 (P adjusted for three interim analyses, 0.0057).



