R§Medicine 2025

An R Consortium Conference

★June 9-13, 2025

Virtual Conference

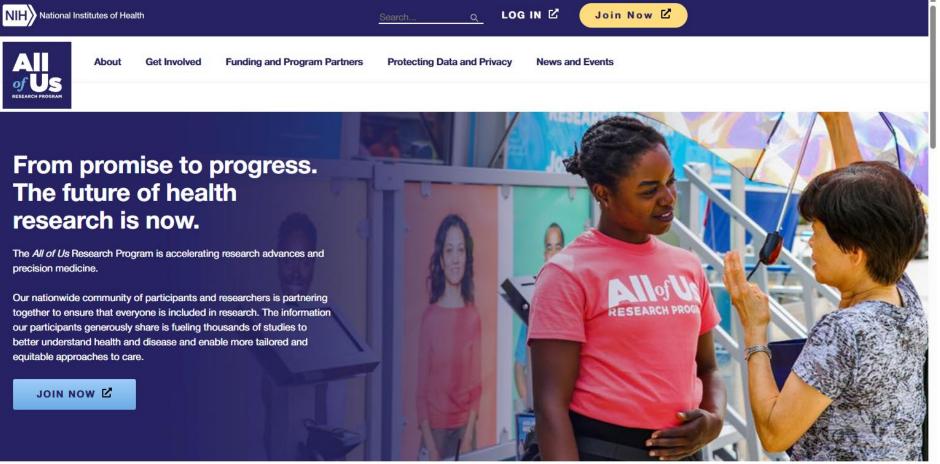
#RMED25

AOUSDOHtools: An R Package for Social Determinants of Health Survey data in the *All of Us* Research Program

Presenter: Zhirui Deng, Biostatistician University of Pittsburgh, School of Nursing R/Medicine 2025 · June 13, 2025

Authors: Zhirui Deng, MS | University of Pittsburgh, School of Nursing
Theresa A. Koleck, PhD, RN | University of Pittsburgh, School of Nursing
Chen Zhang, PhD, MPH, MS | University of Rochester, School of Nursing
Peter D.R. Higgins, MD, PhD, MSc | University of Michigan, School of Medicine
Caitlin Dreisbach, PhD, RN | University of Rochester, School of Nursing

All of Us (AoU) Research Program



- All of Us:
- >1 million participant cohort
- Data:
 EHR, surveys, biospecimens,
 wearables

https://allofus.nih.gov/

Why Do We Need AOUSDOHtools?

- SDOH Survey: rich data but not analysis-ready
- Highly disaggregated responses require transformation
- No built-in tools for SDOH scoring on AoU platform
- Manual scoring is time-consuming and inconsistent
- Need for reproducibility and accessibility

People around here are willing to help their neighbors. (40192463)

See			

Answer	Concept Code ()		
Agree	40192455		
Strongly agree	40192514		
Neutral (neither agree nor disagree)	40192524		
Disagree	40192408		
Strongly disagree	40192422		
Skip	903096		

People around here are willing to help their neighbors. (40192463)

See Answers >

People in my neighborhood generally get along with each other. (40192411)

See Answers >

People in my neighborhood can be trusted. (40192499)

See Answers >

People in my neighborhood share the same values. (40192417)

See Answers >

There is a lot of graffiti in my neighborhood. (40192420)

See Answers >

My neighborhood is noisy. (40192522)

See Answers >

259,200 80

Participants completed this survey

Questions Available

What Does AOUSDOHtools Do?

- An R package designed for AoU SDOH Survey users
- Scoring SDOH constructs based on a published user guide
- Koleck et al. (2024) https://pmc.ncbi.nlm.nih.gov/articles/PMC11631056/





► J Am Med Inform Assoc. 2024 Aug 27;31(12):3032–3041. doi: 10.1093/jamia/ocae214 🗷

User guide for Social Determinants of Health Survey data in the All of Us Research Program

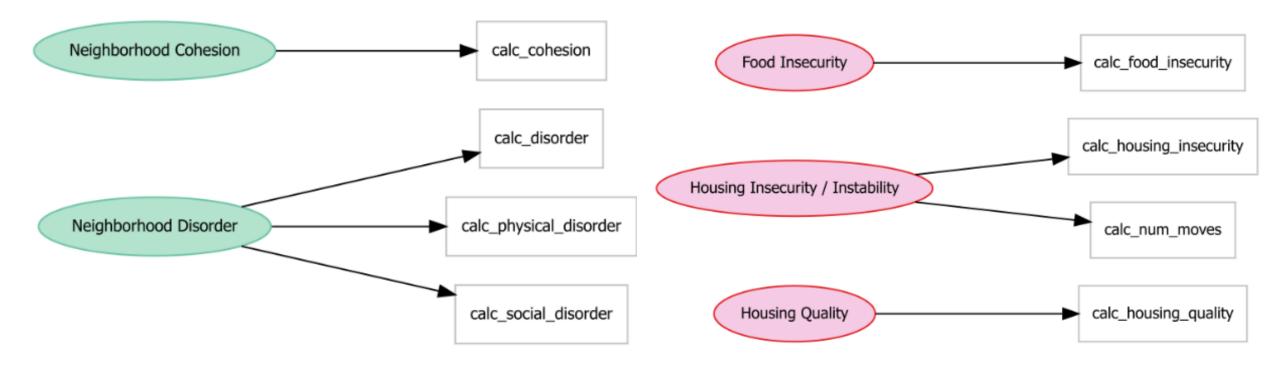
Theresa A Koleck ^{1,⊠}, Caitlin Dreisbach ^{2,3}, Chen Zhang ⁴, Susan Grayson ⁵, Maichou Lor ⁶, Zhirui Deng ⁷, Alex Conway ⁸, Peter D R Higgins ⁹, Suzanne Bakken ^{10,11,12}

► Author information ► Article notes ► Copyright and License information

PMCID: PMC11631056 PMID: 39190874

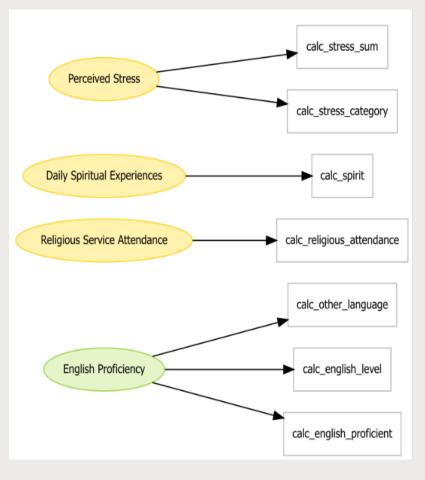
Key Features and SDOH Constructs

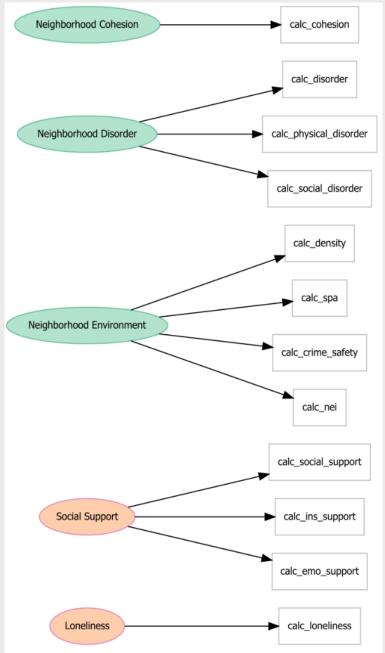
- 30 functions across 14 SDOH constructs
- Examples: neighborhood cohesion and neighborhood disorder
- Built for All of Us Researcher Workbench (Jupyter/RStudio environments)

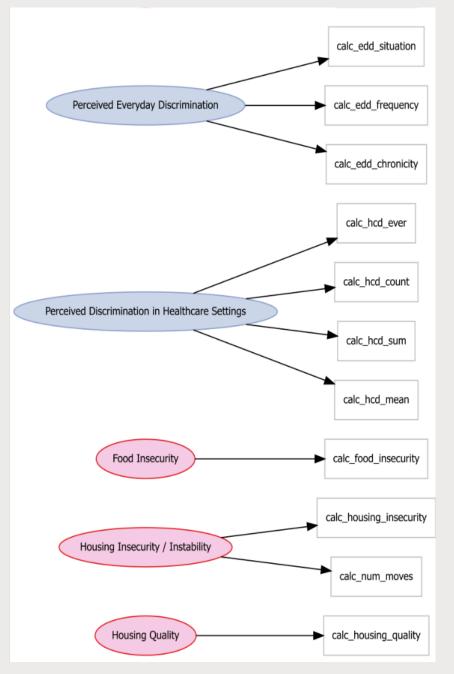


14 SDOH Constructs

Constructs definitions & details can be found in Koleck et al. (2024).







DiagrammeR package (lannone & Roy, 2024)

All of Us Researcher Workbench

RESEARCHER LOGIN





ABOUT

DATA & TOOLS

DISCOVER

SUPPORT

Q

REGISTER

lome > Data & Tools > Researcher Workbench

Researcher Workbench

The Researcher Workbench is a cloud-based platform where registered researchers can access Registered and Controlled Tier data. Its powerful tools support data analysis and collaboration. Integrated help and educational resources are provided through the Workbench User Support Hub.





WORKSPACES

Registered researchers use workspaces to access, store, and analyze data for specific research projects. Workspaces are collaborative and can be shared among other registered researchers within a project team.

USES: Organizing research projects, collaboration



ANALYSIS TOOLS

Researchers with R, Python, or SAS experience can perform queries and analyses within the *All of Us* datasets using our integrated cloud-based analysis tools, including Jupyter Notebook, RStudio, and SAS Studio.

USES: Analysis, queries

How to Use AOUSDOHtools

library(AOUSDOHtools)
survey_df <- your_sdoh_survey_data
cohesion_scores <- calc_cohesion(survey_df)
disorder scores <- calc disorder(survey df)</pre>

Input:

• SDOH Survey data includes person_id, question_concept_id, answer_concept_id

Output:

Scored datasets indexed by person_id, one row per participant

Next steps:

- Merge multiple scores (dplyr::full_join)
- Ready for downstream analysis

calc_cohesion()

```
calc_cohesion <- function(survey_df) {</pre>
  if (!is.null(survey_df)){
    df_cohesion <- survey_df >>
     dplyr::filter(question concept id %in% c(40192463, 40192411, 40192499, 40192417)) > # 4 specific items
     # 40192463 = How much you agree or disagree that people around here are willing to help their neighbor?
     # 40192411 = How much you agree or disagree that people in your neighborhood generally get along with each other?
     # 40192499 = How much you agree or disagree that people in your neighborhood can be trusted?
     # 40192417 = How much you agree or disagree that people in your neighborhood share the same values?
     dplyr::select(person_id, question_concept_id, answer_concept_id) > # map answer_concept_id to value
     dplyr::mutate(value = dplyr::case_when(
        answer_concept_id == 40192514 ~ 5, # Strongly agree
        answer_concept_id == 40192455 ~ 4, # Agree
        answer concept id == 40192524 ~ 3, # Neutral (neither agree nor disagree)
        answer concept id == 40192408 ~ 2, # Disagree
        answer_concept_id == 40192422 ~ 1, # Strongly disagree
       TRUE ~ 999)) |>
     dplyr::filter(value != 999) |> # remove skips
     dplyr::group_by(person_id) |> # group by person_id and calculate mean score
     dplyr::mutate(cohesion = round(mean(value, na.rm = TRUE), 2), # rounded to 2 decimals
                    nrows = length(value)) > # how many questions did the participant answer?
     dplyr::filter(nrows == 4) > # include only participants who answered all 4 questions
     dplyr::select(person_id, cohesion) > # only 2 columns in the final result
     dplyr::distinct(person id, .keep all = TRUE) |> # remove duplicate rows
     dplyr::right_join(survey_df |> # include participants without scores as NA
                          dplyr::group_by(person_id) |>
                         dplyr::summarize(),
                       by = 'person id') >
     dplyr::ungroup() # ungroup
  df_cohesion
```

Availability & Development

Package released by CRAN on 03/26/2025

https://r-packages.io/packages/AOUSDOHtools

Version: 1.0.4

Depends: $R (\geq 4.1.0)$

Imports: <u>dplyr</u>

Suggests: $\underline{\text{testthat}} \ (\geq 3.0.0), \underline{\text{knitr, rmarkdown, covr, devtools, usethis}}$

Published: 2025-03-26

DOI: 10.32614/CRAN.package.AOUSDOHtools

Paper submitted to JOSS on 05/13/2025, currently pending

Actively updating on Github

https://github.com/zhd52/AOUSDOHtools

Impact and Contributions

Standardizes SDOH score generation

Simplifies data transformation

Enables equity-driven research at scale



Get Involved

1. Register for an All of Us account

https://allofus.nih.gov/

2. Have access to All of Us Researcher Workbench and SDOH Survey data

https://www.researchallofus.org/data-tools/workbench/https://www.researchallofus.org/data-tools/survey-explorer/

3. Install AOUSDOHtools

CRAN: install.packages("AOUSDOHtools") GitHub: github.com/zhd52/AOUSDOHtools

4. Apply AOUSDOHtools to your research

Feedback and collaboration welcome!

Acknowledgements

- We thank R/Medicine 2025 for the opportunity to present our work.
- We are grateful to the All of Us Research Program and its participants for making this work possible.
- We also acknowledge all the authors of the published user guide (Koleck et al., 2024),
 whose work provided the conceptual and methodological foundation for scoring the Social
 Determinants of Health constructs implemented in AOUSDOHtools.
- We thank the **CRAN team** for supporting open-source distribution and **GitHub** for providing the platform used to develop and share the package.

References

- All of Us Research Program. All of Us Research Program. National Institutes of Health. Retrieved April 3, 2025, from https://allofus.nih.gov/
- All of Us Research Program Investigators. (2019). The 'All of Us' Research Program. *New England Journal of Medicine*, 381(7), 668–676. https://doi.org/10.1056/NEJMsr1809937
- All of Us Researcher Workbench. All of Us Research Hub. Retrieved April 3, 2025, from https://www.researchallofus.org/
- Deng, Z., Koleck, T. A., Dreisbach, C., Zhang, C., & Higgins, P. D. R. (2025). *AOUSDOHtools: Analyzing AOU SDOH Survey Data*. R package version 1.0.4. https://doi.org/10.32614/cran.package.aousdohtools
- Iannone, R., & Roy, O. (2024). DiagrammeR: Graph/Network Visualization. R package version 1.0.11. https://CRAN.R-project.org/package=DiagrammeR
- Koleck, T. A., Dreisbach, C., Zhang, C., Grayson, S., Lor, M., Deng, Z., Conway, A., Higgins, P. D. R., & Bakken, S. (2024). User guide for Social Determinants of Health Survey data in the All of Us Research Program. *Journal of the American Medical Informatics Association*, 31(12), 3032–3041.
 https://doi.org/10.1093/jamia/ocae214
- Survey Explorer Social Determinants of Health. All of Us Research Hub. Retrieved April 3, 2025, from https://www.researchallofus.org/data-tools/survey-explorer/
- Wickham, H., Francois, R., Henry, L., Muller, K., & Vaughan, D. (2023). *dplyr: A Grammar of Data Manipulation*. R package version 1.1.4. https://CRAN.R-project.org/package=dplyr
- Yu, G., Gatto, L., Rainer, J., Gibb, S., Corty, R., & Neitmann, T. (2020). hexSticker: Create Hexagon Sticker in R. R package version 0.4.9. https://CRAN.R-project.org/package=hexSticker

Useful Links

- All of Us: https://allofus.nih.gov/
- SDOH Survey: https://www.researchallofus.org/data-tools/survey-explorer/
- All of Us Researcher Workbench: https://www.researchallofus.org/data-tools/workbench/
- User Guide by Koleck et al. (2024): https://pmc.ncbi.nlm.nih.gov/articles/PMC11631056/
- CRAN: https://r-packages.io/packages/AOUSDOHtools
- Github: https://github.com/zhd52/AOUSDOHtools

Q & A

