

Caregiver burden data preparation

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Overview

Our analyses focus on caregiver burden as the *outcome* from a variety of predictors, such as demographics, disorder type, cognitive function, and neuropsychological profiles.

Load data and construct data set

The observations we work with require at least summary scores from Zarit's (caregiver) burden interview (ZBI; CITE). Conditional to the ZBI, we include participants who classify as:

- Dementia (community and institution),
- Alzheimer's,
- Parkinson's disease,
- Amyotrophic lateral sclerosis,
- Stroke,
- Epilepsy, and
- Multiple sclerosis.

Other conditions to consider (perhaps for later): Depression (depressn==1), Learning Disability (learning==1), Psychiatric Illness (psychiat==1), or Migraines (migraine==1).

The following lines of code will read in the data and include only the subjects outlined in the above conditions.

```
## data exist one directory above and in a separate folder
### (in part to avoid committing it to Github for now!)
CSHA.1991.full <- read.csv("../IPN2017_Case_Comp/Dataset.csv")

## Conditionals to extract specific participants
### start with primary outcome of interest: caregiver burden
CSHA.1991.zarit <- CSHA.1991.full[CSHA.1991.full$zarscore < 99,]
dim(CSHA.1991.zarit)

## [1] 1086 1724

### Now get all data for above conditions
conditions.of.interest <- (
  CSHA.1991.zarit$cgcc==1 |      # Dementia: COMMUNITY CASE
  CSHA.1991.zarit$cgcc==3 |      # Dementia: INSTITUTION CASE
  CSHA.1991.zarit$adcc==1 |      # Alzheimer's: YES
  CSHA.1991.zarit$parkin==1 |    # Parkinson's: YES
  CSHA.1991.zarit$prstroke==1 |  # Stroke: YES
  CSHA.1991.zarit$als==1 |       # ALS: YES
  CSHA.1991.zarit$epilepsy==1 |  # Epilepsy: YES
  CSHA.1991.zarit$ms==1         # Multiple sclerosis: YES
)
CSHA.1991.zarit_disorders <- CSHA.1991.zarit[conditions.of.interest,]
dim(CSHA.1991.zarit_disorders)
```

```
## [1] 861 1724
```

We now have a subset of observations we should make a pseudo-design matrix of the conditionals above. We want to denote which of the aforementioned categories each individual belongs to:

```
cols.for.design <- c("cgcc", "adcc", "parkin", "prstroke",
  "als", "epilepsy", "ms")
CSHA.1991.zarit_disorders_design <- CSHA.1991.zarit_disorders[,
  cols.for.design]

dim(CSHA.1991.zarit_disorders_design)
```

```
## [1] 861 7
```

```
head(CSHA.1991.zarit_disorders_design)
```

```
##      cgcc adcc parkin prstroke als epilepsy ms
## 1      3     1      2         8  2         2  2
## 2      3     0      2         1  2         2  2
## 4      1     0      2         2  2         2  2
## 11     3     0      2         2  2         2  2
## 26     3     0      8         8  2         2  2
## 35     3     0      1         9  9         9  9
```

We can recode some of the values in these columns to something more sensible with a design matrix. The code below is hidden from the output.

```
summary(as.factor(CSHA.1991.zarit_disorders_design$adcc))
```

```
##      0      1      2      9
## 473 314     10     64
```

```
head(CSHA.1991.zarit_disorders_design)
```

```
##      DEMENTIA  ALZ  PD  STROKE      ALS      EPI      MS
## 1  INS.CASE CASE  NO      DNK      NO      NO      NO
## 2  INS.CASE N/A  NO      YES      NO      NO      NO
## 4  COMM.CASE N/A  NO      NO      NO      NO      NO
## 11 INS.CASE N/A  NO      NO      NO      NO      NO
## 26 INS.CASE N/A DNK      DNK      NO      NO      NO
## 35 INS.CASE N/A YES MISSING MISSING MISSING MISSING
```

```
head(CSHA.design)
```

```
##      DEMENTIA.INS.CASE DEMENTIA.COMM.CASE DEMENTIA.INS.CON ALZ.CASE ALZ.N/A
## 1                      1                      0                      0          1      0
## 2                      1                      0                      0          0      1
## 4                      0                      1                      0          0      1
## 11                     1                      0                      0          0      1
## 26                     1                      0                      0          0      1
## 35                     1                      0                      0          0      1
##      ALZ.MISSING ALZ.CON PD.NO PD.DNK PD.YES PD.MISSING STROKE.DNK
## 1              0      0      1      0      0              0          1
## 2              0      0      1      0      0              0          0
## 4              0      0      1      0      0              0          0
## 11             0      0      1      0      0              0          0
## 26             0      0      0      1      0              0          1
## 35             0      0      0      0      1              0          0
##      STROKE.YES STROKE.NO STROKE.MISSING ALS.NO ALS.MISSING ALS.DNK ALS.YES
```

## 1	0	0	0	1	0	0	0	
## 2	1	0	0	1	0	0	0	
## 4	0	1	0	1	0	0	0	
## 11	0	1	0	1	0	0	0	
## 26	0	0	0	1	0	0	0	
## 35	0	0	1	0	1	0	0	
##	EPI.NO	EPI.MISSING	EPI.DNK	EPI.YES	MS.NO	MS.MISSING	MS.DNK	MS.YES
## 1	1	0	0	0	1	0	0	0
## 2	1	0	0	0	1	0	0	0
## 4	1	0	0	0	1	0	0	0
## 11	1	0	0	0	1	0	0	0
## 26	1	0	0	0	1	0	0	0
## 35	0	1	0	0	0	1	0	0

colSums(CSHA.design)

##	DEMENTIA.INS.CASE	DEMENTIA.COMM.CASE	DEMENTIA.INS.CON
##	539	312	10
##	ALZ.CASE	ALZ.N/A	ALZ.MISSING
##	314	473	64
##	ALZ.CON	PD.NO	PD.DNK
##	10	620	27
##	PD.YES	PD.MISSING	STROKE.DNK
##	48	166	45
##	STROKE.YES	STROKE.NO	STROKE.MISSING
##	241	433	142
##	ALS.NO	ALS.MISSING	ALS.DNK
##	631	204	25
##	ALS.YES	EPI.NO	EPI.MISSING
##	1	644	177
##	EPI.DNK	EPI.YES	MS.NO
##	20	20	665
##	MS.MISSING	MS.DNK	MS.YES
##	176	18	2