

Cognition and Purpose

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Preparation

make composites

```
### SENSE OF PURPOSE
# make purpose mean
let$purpose <- composite(let[,c("LE_1", "LE_2", "LE_3", "LE_4",
                                "LE_5", "LE_6")],
                           nomiss = .75)
# exclude people with missing data
let <- let %>%
  filter(!is.na(purpose))

### DEPRESSION
# make depression mean
bdi$depressionM <- composite(bdi[,c("BDI_1", "BDI_2", "BDI_3", "BDI_4", "BDI_5",
                                       "BDI_6", "BDI_7", "BDI_8", "BDI_9", "BDI_10",
                                       "BDI_11", "BDI_12", "BDI_13", "BDI_14", "BDI_15",
                                       "BDI_16", "BDI_17", "BDI_18", "BDI_19", "BDI_20",
                                       "BDI_21")],
                               nomiss = .75)
bdi$depression <- rowSums(bdi[,c("BDI_1", "BDI_2", "BDI_3", "BDI_4", "BDI_5",
                                   "BDI_6", "BDI_7", "BDI_8", "BDI_9", "BDI_10",
                                   "BDI_11", "BDI_12", "BDI_13", "BDI_14", "BDI_15",
                                   "BDI_16", "BDI_17", "BDI_18", "BDI_19", "BDI_20",
                                   "BDI_21")],
                           na.rm = FALSE) # use based on CESD
# exclude people with missing data
bdi <- bdi %>%
  filter(!is.na(depression))

### LONELINESS
# make loneliness mean
ucla$loneliness <- composite(ucla[,c("UCLA_1", "UCLA_2", "UCLA_3", "UCLA_4", "UCLA_5",
                                         "UCLA_6", "UCLA_7", "UCLA_8", "UCLA_9", "UCLA_10",
                                         "UCLA_11", "UCLA_12", "UCLA_13", "UCLA_14", "UCLA_15",
                                         "UCLA_16", "UCLA_17", "UCLA_18", "UCLA_19", "UCLA_20")],
                                 nomiss = .75)
# exclude people with missing data
ucla <- ucla %>%
  filter(!is.na(loneliness))

### COGNITIVE DECLINE
# sum excludes participants with any NAs
ad8$decline <- rowSums(ad8[,c("AD8_1", "AD8_2", "AD8_3", "AD8_4",
                             "AD8_5", "AD8_6", "AD8_7", "AD8_8")],
                           na.rm = FALSE)
# dummy coded variable based on score of 2+
ad8$dementia <- ifelse(ad8$decline < 2, yes = 0, no = 1)
# exclude people with missing data
ad8 <- ad8 %>%
  filter(!is.na(decline))
```

```

# dummy coded Black and white race variable
demo$raceDc <- ifelse(demo$PRACE == 1, yes = 0,
                      no = ifelse(demo$PRACE == 2, yes = 1,
                                  no = is.na(demo$raceDc)))

## Warning: Unknown or uninitialized column: `raceDc`.

# dummy coded gender: male = 0, female = 1
demo$genderDc <- ifelse(demo$PGENDER == 1, 0,
                        ifelse(demo$PGENDER == 2, 1,
                               is.na(demo$PGENDER)))

```

merge files

```

dataA <- merge(demo, let, by = "PARTID", all = FALSE)
dataB <- merge(dataA, ucla, by = "PARTID", all = FALSE)
dataC <- merge(dataB, bdi, by = "PARTID", all = FALSE)
dataCf <- merge(dataC, ad8, by = "PARTID", all = FALSE)

```

standardize necessary variables

```

# standardize
dataCf$age_z <- as.numeric(scale(dataCf$age, center = TRUE, scale = TRUE))
dataCf$purpose_z <- as.numeric(scale(dataCf$purpose, center = TRUE, scale = TRUE))
dataCf$depression_z <- as.numeric(scale(dataCf$depression, center = TRUE, scale = TRUE))
dataCf$loneliness_z <- as.numeric(scale(dataCf$loneliness, center = TRUE, scale = TRUE))

```

Descriptive Statistics

Cronbach's α

```
# sense of purpose
psych::alpha(let[,c("LE_1", "LE_2", "LE_3", "LE_4", "LE_5", "LE_6")])

##
## Reliability analysis
## Call: psych::alpha(x = let[, c("LE_1", "LE_2", "LE_3", "LE_4", "LE_5",
##     "LE_6")])
##
##   raw_alpha std.alpha G6(smc) average_r S/N      ase mean    sd median_r
##       0.86      0.87      0.85      0.52 6.5 0.0076  4.2 0.68      0.51
##
##   lower alpha upper      95% confidence boundaries
## 0.85 0.86 0.88
##
## Reliability if an item is dropped:
##   raw_alpha std.alpha G6(smc) average_r S/N alpha se var.r med.r
## LE_1      0.85      0.85      0.82      0.53 5.6  0.0085 0.0060  0.53
## LE_2      0.84      0.85      0.83      0.53 5.6  0.0088 0.0057  0.51
## LE_3      0.82      0.83      0.80      0.49 4.8  0.0100 0.0033  0.49
## LE_4      0.83      0.83      0.81      0.50 4.9  0.0096 0.0052  0.47
## LE_5      0.83      0.84      0.81      0.51 5.2  0.0092 0.0050  0.51
## LE_6      0.85      0.86      0.84      0.55 6.1  0.0083 0.0037  0.55
##
## Item statistics
##   n raw.r std.r r.cor r.drop mean    sd
## LE_1 774  0.78  0.75  0.68  0.63  3.9 1.10
## LE_2 774  0.75  0.75  0.68  0.63  4.0 0.87
## LE_3 775  0.83  0.83  0.80  0.74  4.1 0.89
## LE_4 775  0.81  0.82  0.78  0.72  4.2 0.77
## LE_5 775  0.78  0.79  0.74  0.68  4.3 0.83
## LE_6 775  0.69  0.70  0.61  0.56  4.5 0.77
##
## Non missing response frequency for each item
##   1   2   3   4   5 miss
## LE_1 0.03 0.11 0.17 0.35 0.34   0
## LE_2 0.01 0.05 0.16 0.50 0.28   0
## LE_3 0.01 0.05 0.13 0.44 0.37   0
## LE_4 0.01 0.02 0.14 0.48 0.36   0
## LE_5 0.01 0.04 0.08 0.41 0.47   0
## LE_6 0.01 0.02 0.06 0.26 0.65   0

# loneliness
psych::alpha(ucla[,c("UCLA_1", "UCLA_2", "UCLA_3", "UCLA_4", "UCLA_5",
                     "UCLA_6", "UCLA_7", "UCLA_8", "UCLA_9", "UCLA_10",
                     "UCLA_11", "UCLA_12", "UCLA_13", "UCLA_14", "UCLA_15",
                     "UCLA_16", "UCLA_17", "UCLA_18", "UCLA_19", "UCLA_20")])

##
## Reliability analysis
## Call: psych::alpha(x = ucla[, c("UCLA_1", "UCLA_2", "UCLA_3", "UCLA_4",
##     "UCLA_5", "UCLA_6", "UCLA_7", "UCLA_8", "UCLA_9", "UCLA_10",
```

```

##      "UCLA_11", "UCLA_12", "UCLA_13", "UCLA_14", "UCLA_15", "UCLA_16",
##      "UCLA_17", "UCLA_18", "UCLA_19", "UCLA_20"))
##
##      raw_alpha std.alpha G6(smc) average_r S/N      ase mean     sd median_r
##          0.94      0.94    0.95      0.43 15 0.0033  1.8 0.51      0.45
##
##      lower alpha upper      95% confidence boundaries
## 0.93 0.94 0.94
##
##      Reliability if an item is dropped:
##      raw_alpha std.alpha G6(smc) average_r S/N alpha se var.r med.r
## UCLA_1      0.93      0.94    0.95      0.44 15 0.0034 0.0126 0.45
## UCLA_2      0.93      0.94    0.95      0.44 15 0.0034 0.0119 0.46
## UCLA_3      0.93      0.93    0.94      0.43 14 0.0035 0.0117 0.44
## UCLA_4      0.93      0.94    0.94      0.43 15 0.0035 0.0119 0.45
## UCLA_5      0.93      0.93    0.94      0.43 14 0.0035 0.0123 0.45
## UCLA_6      0.93      0.93    0.94      0.43 14 0.0035 0.0123 0.44
## UCLA_7      0.93      0.93    0.94      0.43 14 0.0035 0.0120 0.44
## UCLA_8      0.93      0.94    0.95      0.44 15 0.0034 0.0123 0.45
## UCLA_9      0.93      0.94    0.95      0.44 15 0.0034 0.0120 0.46
## UCLA_10     0.93      0.93    0.94      0.43 14 0.0035 0.0122 0.44
## UCLA_11     0.93      0.93    0.94      0.43 14 0.0035 0.0128 0.45
## UCLA_12     0.93      0.94    0.95      0.43 15 0.0034 0.0126 0.45
## UCLA_13     0.93      0.93    0.94      0.43 14 0.0035 0.0124 0.44
## UCLA_14     0.93      0.93    0.94      0.43 14 0.0035 0.0124 0.44
## UCLA_15     0.93      0.94    0.95      0.44 15 0.0034 0.0122 0.45
## UCLA_16     0.93      0.93    0.94      0.43 14 0.0035 0.0121 0.45
## UCLA_17     0.94      0.94    0.95      0.46 16 0.0031 0.0069 0.46
## UCLA_18     0.93      0.94    0.95      0.43 15 0.0034 0.0128 0.45
## UCLA_19     0.93      0.93    0.94      0.43 14 0.0035 0.0116 0.45
## UCLA_20     0.93      0.93    0.94      0.43 14 0.0035 0.0113 0.45
##
##      Item statistics
##      n raw.r std.r r.cor r.drop mean     sd
## UCLA_1   774 0.63 0.65 0.62  0.60 1.6 0.58
## UCLA_2   774 0.64 0.63 0.60  0.59 2.1 0.91
## UCLA_3   774 0.76 0.75 0.74  0.72 1.6 0.82
## UCLA_4   772 0.69 0.67 0.66  0.64 1.9 0.83
## UCLA_5   774 0.72 0.72 0.71  0.68 1.7 0.78
## UCLA_6   774 0.72 0.74 0.72  0.69 1.7 0.63
## UCLA_7   774 0.76 0.75 0.74  0.72 1.6 0.82
## UCLA_8   774 0.64 0.64 0.61  0.59 2.1 0.82
## UCLA_9   774 0.57 0.59 0.56  0.53 1.6 0.62
## UCLA_10  774 0.74 0.75 0.74  0.71 1.6 0.62
## UCLA_11  774 0.73 0.73 0.71  0.69 2.0 0.78
## UCLA_12  774 0.68 0.67 0.65  0.63 1.7 0.80
## UCLA_13  774 0.72 0.72 0.70  0.68 1.9 0.84
## UCLA_14  774 0.74 0.73 0.72  0.70 1.9 0.86
## UCLA_15  774 0.64 0.64 0.61  0.59 1.6 0.78
## UCLA_16  773 0.71 0.72 0.70  0.67 1.7 0.70
## UCLA_17  774 0.38 0.37 0.32  0.30 2.2 0.84
## UCLA_18  773 0.66 0.66 0.64  0.62 2.1 0.79
## UCLA_19  774 0.71 0.72 0.72  0.68 1.4 0.65
## UCLA_20  774 0.72 0.74 0.73  0.69 1.4 0.64

```

```

##  

## Non missing response frequency for each item  

##      1   2   3   4 miss  

## UCLA_1  0.44 0.52 0.03 0.01    0  

## UCLA_2  0.33 0.32 0.31 0.05    0  

## UCLA_3  0.60 0.23 0.16 0.02    0  

## UCLA_4  0.36 0.39 0.22 0.03    0  

## UCLA_5  0.49 0.37 0.11 0.03    0  

## UCLA_6  0.38 0.55 0.06 0.01    0  

## UCLA_7  0.61 0.22 0.15 0.02    0  

## UCLA_8  0.28 0.39 0.32 0.02    0  

## UCLA_9  0.49 0.46 0.05 0.01    0  

## UCLA_10 0.43 0.51 0.05 0.01    0  

## UCLA_11 0.26 0.48 0.24 0.03    0  

## UCLA_12 0.46 0.36 0.16 0.02    0  

## UCLA_13 0.36 0.38 0.23 0.03    0  

## UCLA_14 0.39 0.34 0.24 0.03    0  

## UCLA_15 0.56 0.32 0.08 0.04    0  

## UCLA_16 0.44 0.46 0.07 0.02    0  

## UCLA_17 0.24 0.35 0.38 0.03    0  

## UCLA_18 0.24 0.43 0.31 0.02    0  

## UCLA_19 0.69 0.25 0.05 0.01    0  

## UCLA_20 0.71 0.23 0.05 0.01    0  

# depressive symptoms  

psych::alpha(dataCf[,c("BDI_1", "BDI_2", "BDI_3", "BDI_4", "BDI_5",
                      "BDI_6", "BDI_7", "BDI_8", "BDI_9", "BDI_10",
                      "BDI_11", "BDI_12", "BDI_13", "BDI_14", "BDI_15",
                      "BDI_16", "BDI_17", "BDI_18", "BDI_19", "BDI_20",
                      "BDI_21")])  

##  

## Reliability analysis  

## Call: psych::alpha(x = dataCf[, c("BDI_1", "BDI_2", "BDI_3", "BDI_4",
##      "BDI_5", "BDI_6", "BDI_7", "BDI_8", "BDI_9", "BDI_10", "BDI_11",
##      "BDI_12", "BDI_13", "BDI_14", "BDI_15", "BDI_16", "BDI_17",
##      "BDI_18", "BDI_19", "BDI_20", "BDI_21")])  

##  

##      raw_alpha std.alpha G6(smc) average_r S/N      ase mean     sd median_r
##            0.89      0.91      0.92      0.33    10 0.0063 0.25 0.28      0.32  

##  

##      lower alpha upper      95% confidence boundaries
## 0.88 0.89 0.9  

##  

## Reliability if an item is dropped:  

##      raw_alpha std.alpha G6(smc) average_r  S/N alpha se var.r med.r
## BDI_1        0.89      0.90      0.92      0.32  9.4  0.0067 0.013  0.31
## BDI_2        0.89      0.90      0.92      0.32  9.4  0.0067 0.013  0.31
## BDI_3        0.89      0.91      0.92      0.32  9.5  0.0066 0.013  0.32
## BDI_4        0.88      0.90      0.92      0.32  9.3  0.0069 0.013  0.31
## BDI_5        0.89      0.91      0.92      0.33  9.7  0.0066 0.013  0.32
## BDI_6        0.89      0.91      0.92      0.33 10.1  0.0064 0.013  0.33
## BDI_7        0.88      0.90      0.91      0.32  9.2  0.0068 0.012  0.31
## BDI_8        0.89      0.91      0.92      0.32  9.5  0.0066 0.013  0.32
## BDI_9        0.89      0.91      0.92      0.33  9.9  0.0065 0.013  0.33

```

```

## BDI_10      0.89      0.91      0.92      0.32    9.6    0.0066  0.014   0.31
## BDI_11      0.89      0.91      0.92      0.33    9.8    0.0065  0.013   0.32
## BDI_12      0.88      0.90      0.91      0.32    9.3    0.0068  0.013   0.31
## BDI_13      0.89      0.91      0.92      0.33    9.6    0.0066  0.014   0.31
## BDI_14      0.88      0.90      0.91      0.32    9.2    0.0068  0.012   0.31
## BDI_15      0.89      0.91      0.92      0.33    9.7    0.0067  0.014   0.33
## BDI_16      0.89      0.91      0.92      0.33    9.9    0.0064  0.014   0.34
## BDI_17      0.89      0.90      0.92      0.32    9.4    0.0067  0.013   0.31
## BDI_18      0.89      0.91      0.92      0.33   10.0    0.0064  0.013   0.34
## BDI_19      0.89      0.91      0.92      0.33    9.7    0.0067  0.014   0.32
## BDI_20      0.88      0.90      0.92      0.32    9.5    0.0068  0.014   0.31
## BDI_21      0.90      0.91      0.92      0.34   10.4    0.0056  0.011   0.34
##
## Item statistics
##          n raw.r std.r r.cor r.drop mean   sd
## BDI_1  595  0.66  0.68  0.67   0.63  0.121  0.34
## BDI_2  595  0.65  0.66  0.64   0.60  0.229  0.47
## BDI_3  595  0.60  0.63  0.61   0.54  0.193  0.50
## BDI_4  595  0.70  0.70  0.69   0.66  0.281  0.52
## BDI_5  595  0.54  0.57  0.54   0.49  0.151  0.39
## BDI_6  595  0.43  0.45  0.41   0.38  0.066  0.35
## BDI_7  595  0.70  0.73  0.74   0.66  0.124  0.44
## BDI_8  595  0.61  0.63  0.61   0.56  0.136  0.43
## BDI_9  595  0.46  0.51  0.47   0.43  0.054  0.23
## BDI_10 595  0.60  0.62  0.59   0.54  0.139  0.49
## BDI_11 595  0.51  0.54  0.50   0.46  0.126  0.36
## BDI_12 595  0.71  0.70  0.70   0.66  0.198  0.47
## BDI_13 595  0.58  0.59  0.57   0.53  0.106  0.37
## BDI_14 595  0.70  0.74  0.74   0.66  0.106  0.39
## BDI_15 595  0.60  0.58  0.55   0.54  0.689  0.55
## BDI_16 595  0.54  0.49  0.45   0.44  0.565  0.74
## BDI_17 595  0.67  0.69  0.67   0.63  0.134  0.39
## BDI_18 595  0.50  0.48  0.44   0.42  0.324  0.59
## BDI_19 595  0.60  0.59  0.56   0.54  0.291  0.50
## BDI_20 595  0.67  0.64  0.62   0.61  0.466  0.60
## BDI_21 595  0.42  0.34  0.29   0.29  0.697  0.90
##
## Non missing response frequency for each item
##          0    1    2    3 miss
## BDI_1  0.88 0.11 0.01 0.00    0
## BDI_2  0.79 0.20 0.01 0.01    0
## BDI_3  0.85 0.10 0.04 0.00    0
## BDI_4  0.75 0.23 0.02 0.01    0
## BDI_5  0.86 0.13 0.00 0.00    0
## BDI_6  0.95 0.04 0.00 0.01    0
## BDI_7  0.91 0.05 0.03 0.00    0
## BDI_8  0.89 0.09 0.01 0.01    0
## BDI_9  0.95 0.05 0.00 0.00    0
## BDI_10 0.90 0.08 0.00 0.02    0
## BDI_11 0.88 0.11 0.01 0.00    0
## BDI_12 0.83 0.16 0.01 0.01    0
## BDI_13 0.91 0.07 0.02 0.00    0
## BDI_14 0.92 0.06 0.02 0.00    0
## BDI_15 0.35 0.61 0.03 0.00    0

```

```
## BDI_16 0.55 0.36 0.05 0.03      0
## BDI_17 0.88 0.11 0.01 0.00      0
## BDI_18 0.73 0.22 0.04 0.01      0
## BDI_19 0.73 0.26 0.01 0.00      0
## BDI_20 0.58 0.39 0.02 0.01      0
## BDI_21 0.54 0.30 0.10 0.07      0
```

Demographics

```
Hmisc::describe(dataCf[,c("age", "raceDc", "PRACE", "genderDc", "PGENDER",
                         "purpose", "loneliness", "depression",
                         "decline", "dementia")])

## Warning in all.is.numeric(names(weights), "vector"): NAs introduced by coercion
## dataCf[, c("age", "raceDc", "PRACE", "genderDc", "PGENDER", "purpose", "loneliness", "depression", "decline", "dementia")]
## 
## 10 Variables      595 Observations
## -----
## age
##      n    missing   distinct     Info      Mean      Gmd     .05     .10
##      595        0       14    0.991    71.43    3.516    67      67
##      .25       .50       .75     .90      .95
##      69        71       74      76      76
## 
## lowest : 65 66 67 68 69, highest: 74 75 76 77 78
## 
## Value      65     66     67     68     69     70     71     72     73     74     75
## Frequency  2      23     36     58     73     57     59     54     57     68     38
## Proportion 0.003 0.039 0.061 0.097 0.123 0.096 0.099 0.091 0.096 0.114 0.064
## 
## Value      76     77     78
## Frequency  46     18      6
## Proportion 0.077 0.030 0.010
## -----
## raceDc
##      n    missing   distinct     Info      Sum      Mean      Gmd
##      584       11       2    0.455     109    0.1866    0.3041
## 
## PRACE
##      n    missing   distinct
##      595        0       7
## 
## lowest : . 1 2 3 5, highest: 2 3 5 7 8
## 
## Value      NA     1     2     3     5     7     8
## Frequency  2    475   109     1     2     2     4
## Proportion 0.003 0.798 0.183 0.002 0.003 0.003 0.007
## 
## genderDc
##      n    missing   distinct     Info      Sum      Mean      Gmd
##      595        0       2    0.739     333    0.5597    0.4937
## 
## PGENDER
##      n    missing   distinct
##      595        0       2
## 
## Value      1     2
## Frequency  262   333
## Proportion 0.44 0.56
```

```

## -----
## purpose
##      n  missing distinct      Info      Mean      Gmd      .05      .10
##      595       0       20     0.989     4.212    0.7115    3.000    3.400
##      .25       .50       .75       .90       .95
##      3.833     4.167     4.833     5.000     5.000
##
## lowest : 1.500000 2.000000 2.166667 2.333333 2.500000
## highest: 4.333333 4.500000 4.666667 4.833333 5.000000
##
## 1.5 (1, 0.002), 2 (2, 0.003), 2.16666666666667 (3, 0.005), 2.33333333333333 (5,
## 0.008), 2.5 (6, 0.010), 2.66666666666667 (5, 0.008), 2.83333333333333 (5,
## 0.008), 3 (10, 0.017), 3.16666666666667 (9, 0.015), 3.33333333333333 (14,
## 0.024), 3.5 (26, 0.044), 3.66666666666667 (27, 0.045), 3.83333333333333 (51,
## 0.086), 4 (69, 0.116), 4.16666666666667 (68, 0.114), 4.33333333333333 (57,
## 0.096), 4.5 (41, 0.069), 4.66666666666667 (36, 0.061), 4.83333333333333 (57,
## 0.096), 5 (103, 0.173)
##
## -----
## loneliness
##      n  missing distinct      Info      Mean      Gmd      .05      .10
##      595       0       48     0.999     1.705    0.5345    1.10     1.15
##      .25       .50       .75       .90       .95
##      1.35     1.65     1.95      2.40      2.65
##
## lowest : 1.00 1.05 1.10 1.15 1.20, highest: 3.20 3.40 3.60 3.65 3.75
## -----
## depression
##      n  missing distinct      Info      Mean      Gmd      .05      .10
##      595       0       32     0.99     5.197    5.567      0       0
##      .25       .50       .75       .90       .95
##      1         4        7       12       17
##
## lowest : 0 1 2 3 4, highest: 30 33 34 44 50
## -----
## decline
##      n  missing distinct      Info      Mean      Gmd
##      595       0       9     0.709     0.684    1.066
##
## lowest : 0 1 2 3 4, highest: 4 5 6 7 8
## -----
## Value      0      1      2      3      4      5      6      7      8
## Frequency 391   111    45    20    10     8     5     4     1
## Proportion 0.657 0.187 0.076 0.034 0.017 0.013 0.008 0.007 0.002
## -----
## dementia
##      n  missing distinct      Info      Sum      Mean      Gmd
##      595       0       2     0.396      93    0.1563    0.2642
##
## -----
library(gmodels)
CrossTable( dataCf$raceDc, dataCf$dementia)

##
##

```

```

##      Cell Contents
## |-----|
## |           N |
## | Chi-square contribution |
## |           N / Row Total |
## |           N / Col Total |
## |           N / Table Total |
## |-----|
## 
## 
## Total Observations in Table:  584
##
## 
##          | dataCf$dementia
## dataCf$raceDc |      0 |      1 | Row Total |
## -----|-----|-----|-----|
##      0 |    401 |     74 |    475 |
##      | 0.000 | 0.000 |      |
##      | 0.844 | 0.156 | 0.813 |
##      | 0.813 | 0.813 |      |
##      | 0.687 | 0.127 |      |
## -----|-----|-----|-----|
##      1 |    92 |     17 |    109 |
##      | 0.000 | 0.000 |      |
##      | 0.844 | 0.156 | 0.187 |
##      | 0.187 | 0.187 |      |
##      | 0.158 | 0.029 |      |
## -----|-----|-----|-----|
## Column Total |    493 |     91 |    584 |
##      | 0.844 | 0.156 |      |
## -----|-----|-----|-----|
## 
## 
## 
```

Race Differences

```

# mean differences in groups
t.test(purpose ~ raceDc, data = dataCf)

## 
## Welch Two Sample t-test
## 
## data: purpose by raceDc
## t = -2.8165, df = 177.01, p-value = 0.005406
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -0.30633930 -0.05391823
## sample estimates:
## mean in group 0 mean in group 1
##        4.176140        4.356269

t.test(depression ~ raceDc, data = dataCf)

## 
## Welch Two Sample t-test

```

```

##  

## data: depression by raceDc  

## t = -0.67189, df = 144.18, p-value = 0.5027  

## alternative hypothesis: true difference in means is not equal to 0  

## 95 percent confidence interval:  

## -1.860992 0.916752  

## sample estimates:  

## mean in group 0 mean in group 1  

## 5.124211 5.596330  

t.test(loneliness ~ raceDc, data = dataCf)  

##  

## Welch Two Sample t-test  

##  

## data: loneliness by raceDc  

## t = 0.041501, df = 148.9, p-value = 0.967  

## alternative hypothesis: true difference in means is not equal to 0  

## 95 percent confidence interval:  

## -0.1085737 0.1132321  

## sample estimates:  

## mean in group 0 mean in group 1  

## 1.707834 1.705505  

t.test(decline ~ raceDc, data = dataCf)  

##  

## Welch Two Sample t-test  

##  

## data: decline by raceDc  

## t = -0.74192, df = 139.62, p-value = 0.4594  

## alternative hypothesis: true difference in means is not equal to 0  

## 95 percent confidence interval:  

## -0.4352512 0.1977234  

## sample estimates:  

## mean in group 0 mean in group 1  

## 0.6610526 0.7798165  

# summary stats  

dataCf %>%  

  group_by(raceDc) %>%  

  summarise_at(vars(purpose, depression,  

                    loneliness, decline),  

    list(name = mean))  

## # A tibble: 3 x 5  

##   raceDc purpose_name depression_name loneliness_name decline_name  

##   <dbl>        <dbl>          <dbl>           <dbl>         <dbl>  

## 1     0        4.18          5.12          1.71        0.661  

## 2     1        4.36          5.60          1.71        0.780  

## 3    NA        4.33          4.36          1.59        0.727  

dataCf %>%  

  group_by(raceDc) %>%  

  summarise_at(vars(purpose, depression,  

                    loneliness, decline),  

    list(name = sd))
```

```

## # A tibble: 3 x 5
##   raceDc purpose_name depression_name loneliness_name decline_name
##   <dbl>      <dbl>          <dbl>          <dbl>          <dbl>
## 1     0       0.663        5.67        0.476        1.22
## 2     1       0.587        6.82        0.540        1.57
## 3     NA      0.483        6.23        0.459        1.42

# group differences in groups
chisq.test(dataCf$dementia, dataCf$raceDc)

## 
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: dataCf$dementia and dataCf$raceDc
## X-squared = 0.0000000000000000000000000000000000033918, df = 1, p-value = 1

dataCf %>%
  group_by(raceDc) %>%
  summarise_at(vars(dementia),
               list(name = sum))

## # A tibble: 3 x 2
##   raceDc name
##   <dbl> <dbl>
## 1     0    74
## 2     1    17
## 3     NA     2

```

Correlations

```

cor <- cor.ci(dataCf[,c("age", "purpose", "depression", "loneliness",
                     "decline", "dementia")])
print(cor, short=FALSE)

## Call:corCi(x = x, keys = keys, n.iter = n.iter, p = p, overlap = overlap,
##             poly = poly, method = method, plot = plot, minlength = minlength,
##             n = n)
##
## Coefficients and bootstrapped confidence intervals
##          age   purps dprss llnls decln demnt
## age     1.00
## purpose -0.05  1.00
## depression -0.01 -0.50  1.00
## loneliness -0.06 -0.54  0.58  1.00
## decline    0.04 -0.30  0.55  0.37  1.00
## dementia   0.03 -0.27  0.46  0.31  0.83  1.00
##
## scale correlations and bootstrapped confidence intervals
##          lower.emp lower.norm estimate upper.norm upper.emp      p
## age-purps    -0.12     -0.12    -0.05      0.03    0.04  0.25
## age-dprss    -0.08     -0.09    -0.01      0.06    0.07  0.69
## age-llnls    -0.14     -0.14    -0.06      0.02    0.02  0.14
## age-decln    -0.03     -0.03    0.04       0.10    0.10  0.34
## age-demnt    -0.06     -0.06    0.03       0.10    0.09  0.58
## purps-dprss   -0.56     -0.56    -0.50     -0.43   -0.43  0.00

```

```

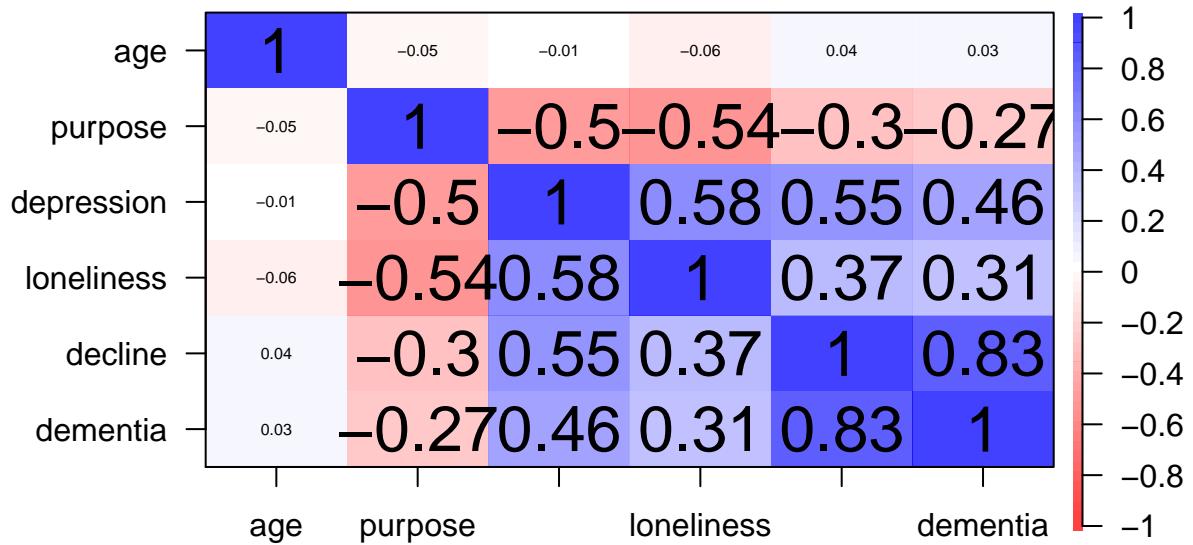
## purps-lnlns -0.61 -0.61 -0.54 -0.46 -0.48 0.00
## purps-decln -0.38 -0.39 -0.30 -0.21 -0.20 0.00
## purps-demnt -0.37 -0.36 -0.27 -0.18 -0.18 0.00
## dprss-lnlns 0.51 0.50 0.58 0.65 0.65 0.00
## dprss-decln 0.46 0.45 0.55 0.64 0.63 0.00
## dprss-demnt 0.38 0.39 0.46 0.54 0.53 0.00
## lnlns-decln 0.26 0.26 0.37 0.46 0.45 0.00
## lnlns-demnt 0.20 0.21 0.31 0.41 0.42 0.00
## decln-demnt 0.81 0.81 0.83 0.85 0.85 0.00

cor$ci$p

## [1] 0.2477374626545241209641 0.6864974683795088683524 0.1441234648695293163456
## [4] 0.3422836957427735082859 0.5764681402453248892925 0.00000000000000000000000000000000
## [7] 0.00000000000000000000000000000000 0.0000000008978784382663 0.0000000925825858200113
## [10] 0.00000000000000000000000000000000 0.000000000000000000000000000000002220446 0.00000000000000000000000000000000
## [13] 0.0000000003915472390759 0.0000000168997051908093 0.00000000000000000000000000000000

round(cor.ci(dataCf[,c("age", "purpose", "depression", "loneliness",
"decline", "dementia")])$ci, digits = 4)

```



```

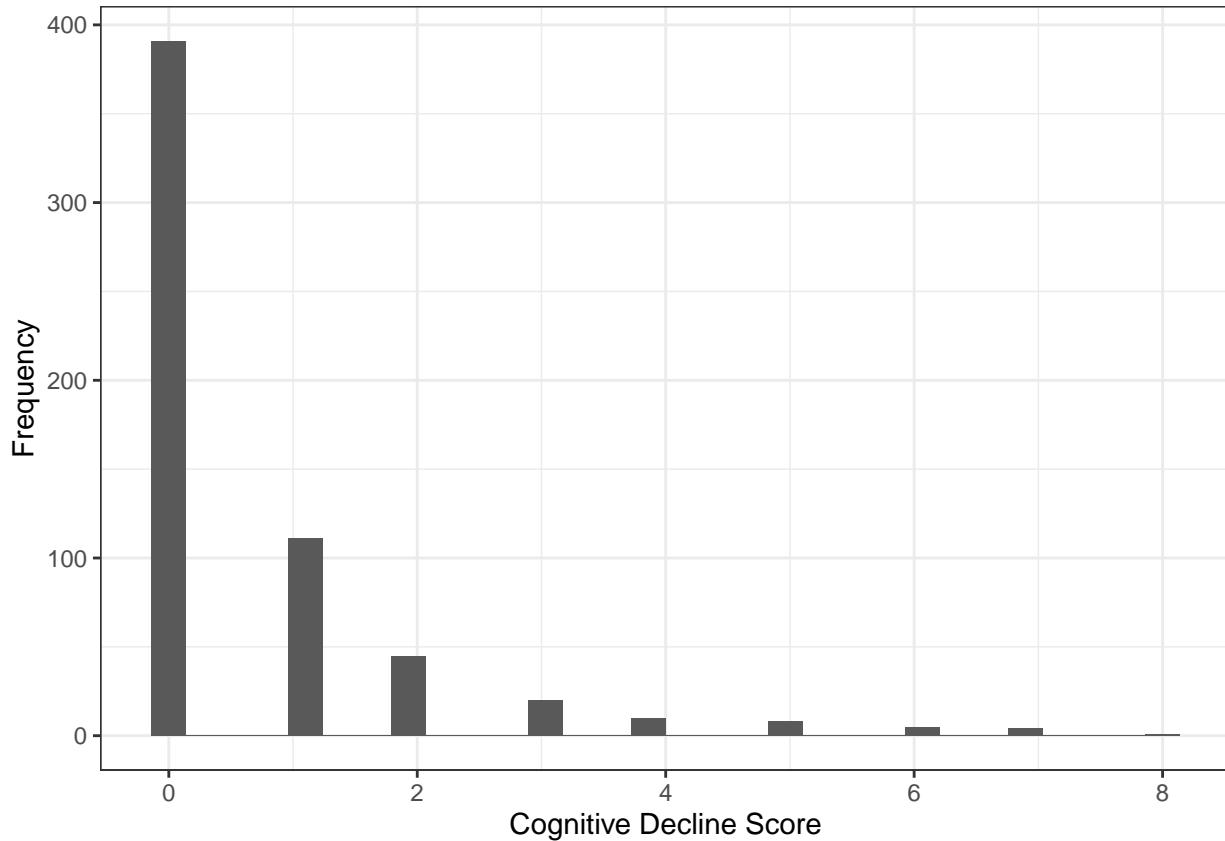
## lower low.e upper up.e p
## age-purps -0.1216 -0.1189 0.0310 0.0201 0.2437
## age-dprss -0.1005 -0.1079 0.0666 0.0657 0.6894
## age-lnlns -0.1361 -0.1356 0.0154 0.0079 0.1184
## age-decln -0.0394 -0.0436 0.1146 0.1067 0.3369
## age-demnt -0.0602 -0.0569 0.1168 0.1144 0.5288
## purps-dprss -0.5674 -0.5638 -0.4396 -0.4415 0.0000
## purps-lnlns -0.6132 -0.6050 -0.4717 -0.4805 0.0000
## purps-decln -0.3871 -0.3854 -0.2031 -0.1997 0.0000
## purps-demnt -0.3546 -0.3508 -0.1795 -0.1728 0.0000
## dprss-lnlns 0.5005 0.5025 0.6504 0.6577 0.0000
## dprss-decln 0.4442 0.4427 0.6251 0.6211 0.0000
## dprss-demnt 0.3919 0.3825 0.5218 0.5123 0.0000
## lnlns-decln 0.2767 0.2910 0.4468 0.4353 0.0000
## lnlns-demnt 0.2313 0.2262 0.3872 0.3946 0.0000
## decln-demnt 0.8081 0.8060 0.8530 0.8516 0.0000

```

Distributions

cognitive decline

```
ggplot(dataCf, aes(x = decline)) +  
  geom_histogram() +  
  theme_bw() +  
  labs(x = "Cognitive Decline Score",  
       y = "Frequency")  
  
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

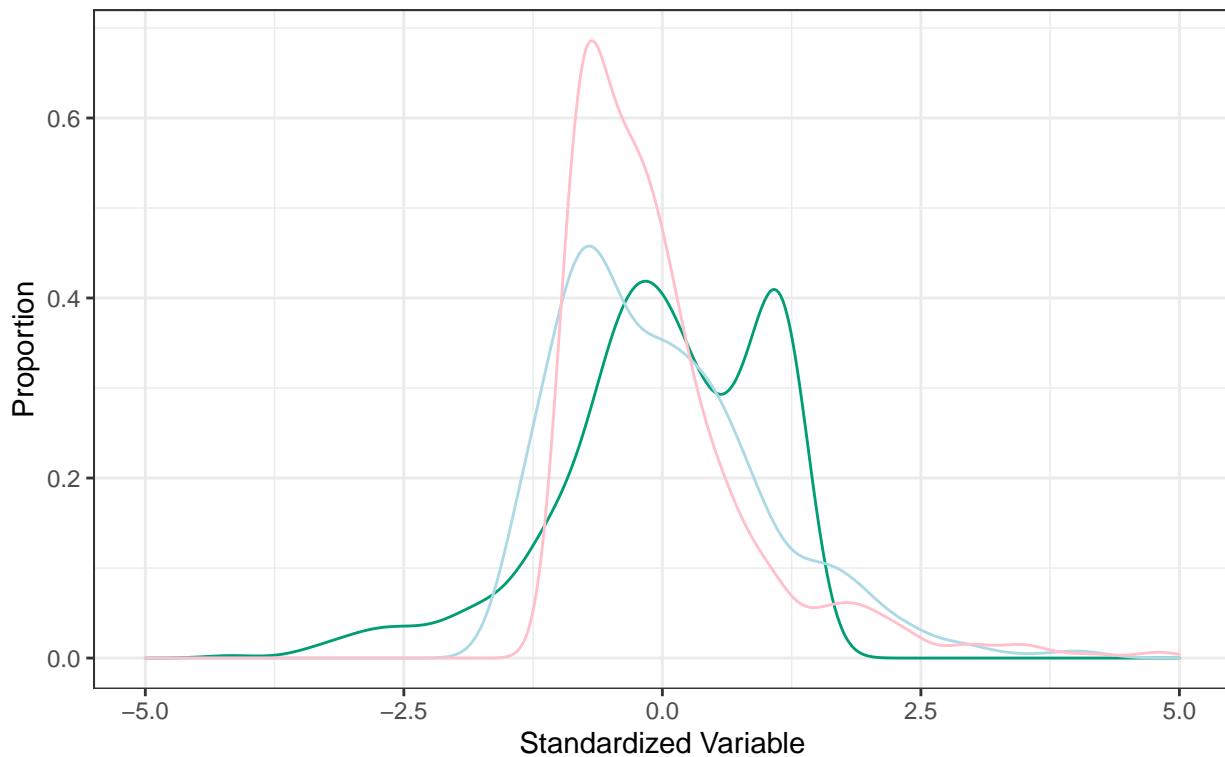


standardized psychological variables

```
ggplot(dataCf) +
  geom_density(mapping = aes(x = purpose_z), color = "#009E73") +
  geom_density(mapping = aes(x = loneliness_z), color = "lightblue") +
  geom_density(mapping = aes(x = depression_z), color = "pink") +
  theme_bw() +
  xlim(-5,5) +
  labs(x = "Standardized Variable",
       y = "Proportion",
       title = "Distribution of Purpose (green), Loneliness (light blue),\nand Depression (pink)")
```

Warning: Removed 2 rows containing non-finite values (stat_density).

Distribution of Purpose (green), Loneliness (light blue),
and Depression (pink)



Hypothesis Testing

Only depression is a unique predictor of cognitive decline both continuously and dichotomously when standardized purpose, loneliness, and depression are entered simultaneously alongside age.

```
mod1cdA <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z + depression_z,  
               data = dataCf)  
summary(mod1cdA)
```

```
##  
## Call:  
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z +  
##       depression_z, data = dataCf)  
##  
## Residuals:  
##      Min      1Q Median      3Q      Max  
## -2.6247 -0.5368 -0.1843  0.2063  7.1727  
##  
## Coefficients:  
##             Estimate Std. Error t value     Pr(>|t|)  
## (Intercept)  0.775262  0.066941 11.581 <0.0000000000000002 ***  
## age_z        0.061144  0.044630   1.370     0.1712  
## genderDc     -0.163005  0.089673  -1.818     0.0696 .  
## purpose_z    -0.005646  0.055163  -0.102     0.9185  
## loneliness_z 0.087766  0.058826   1.492     0.1362  
## depression_z  0.655391  0.056942  11.510 <0.0000000000000002 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 1.081 on 589 degrees of freedom  
## Multiple R-squared:  0.3072, Adjusted R-squared:  0.3013  
## F-statistic: 52.23 on 5 and 589 DF,  p-value: < 0.0000000000000022
```

```
mod1cd <- lm(decline ~ age + genderDc + purpose_z + loneliness_z + depression_z,  
               data = dataCf)  
summary(mod1cd)
```

```
##  
## Call:  
## lm(formula = decline ~ age + genderDc + purpose_z + loneliness_z +  
##       depression_z, data = dataCf)  
##  
## Residuals:  
##      Min      1Q Median      3Q      Max  
## -2.6247 -0.5368 -0.1843  0.2063  7.1727  
##  
## Coefficients:  
##             Estimate Std. Error t value     Pr(>|t|)  
## (Intercept) -0.650004  1.042881  -0.623     0.5333  
## age          0.019953  0.014564   1.370     0.1712  
## genderDc     -0.163005  0.089673  -1.818     0.0696 .  
## purpose_z    -0.005646  0.055163  -0.102     0.9185  
## loneliness_z 0.087766  0.058826   1.492     0.1362  
## depression_z  0.655391  0.056942  11.510 <0.0000000000000002 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```

##
## Residual standard error: 1.081 on 589 degrees of freedom
## Multiple R-squared:  0.3072, Adjusted R-squared:  0.3013
## F-statistic: 52.23 on 5 and 589 DF,  p-value: < 0.00000000000000022
mod1dem <- glm(dementia ~ age + genderDc + purpose_z + loneliness_z + depression_z,
                 data = dataCf)
summary(mod1dem)

##
## Call:
## glm(formula = dementia ~ age + genderDc + purpose_z + loneliness_z +
##     depression_z, data = dataCf)
##
## Deviance Residuals:
##      Min        1Q     Median        3Q       Max
## -0.71718  -0.15404  -0.07727  -0.00423   0.94536
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.147089  0.311268 -0.473   0.637
## age          0.004282  0.004347  0.985   0.325
## genderDc    -0.004409  0.026765 -0.165   0.869
## purpose_z   -0.010867  0.016464 -0.660   0.509
## loneliness_z  0.019542  0.017558  1.113   0.266
## depression_z  0.151745  0.016996  8.929 <0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.104013)
##
## Null deviance: 78.464 on 594 degrees of freedom
## Residual deviance: 61.264 on 589 degrees of freedom
## AIC: 349.88
##
## Number of Fisher Scoring iterations: 2
mod1demA <- glm(dementia ~ age_z + genderDc + purpose_z + loneliness_z + depression_z,
                  data = dataCf)
summary(mod1demA)

##
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + loneliness_z +
##     depression_z, data = dataCf)
##
## Deviance Residuals:
##      Min        1Q     Median        3Q       Max
## -0.71718  -0.15404  -0.07727  -0.00423   0.94536
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept)  0.158770  0.019980  7.947  0.0000000000098 ***
## age_z        0.013121  0.013321  0.985   0.325
## genderDc    -0.004409  0.026765 -0.165   0.869

```

```
## purpose_z     -0.010867   0.016464   -0.660          0.509
## loneliness_z  0.019542   0.017558    1.113          0.266
## depression_z   0.151745   0.016996   8.929 < 0.0000000000000002 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.104013)
##
## Null deviance: 78.464  on 594  degrees of freedom
## Residual deviance: 61.264  on 589  degrees of freedom
## AIC: 349.88
##
## Number of Fisher Scoring iterations: 2
```

Purpose X Loneliness

Results

Higher sense of purpose and lower loneliness are associated with less cognitive decline when treating cognitive decline both continuously and dichotomously when accounting for age. Furthermore, there is an interaction between loneliness and purpose, wherein the association between loneliness and cognitive decline gets weaker as purpose levels get higher. In other words, sense of purpose could be a buffer against the effects of loneliness on cognitive decline.

```
# for graphing
mod2cd <- lm(decline ~ age + genderDc + purpose + loneliness + purpose * loneliness,
               data = dataCf)
summary(mod2cd)

##
## Call:
## lm(formula = decline ~ age + genderDc + purpose + loneliness +
##     purpose * loneliness, data = dataCf)
##
## Residuals:
##    Min      1Q  Median      3Q     Max 
## -3.2032 -0.5674 -0.3246  0.3702  7.0913 
## 
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) -4.29753   1.66112 -2.587  0.009917 **  
## age          0.02077   0.01597  1.300  0.193971    
## genderDc     -0.05990   0.09802 -0.611  0.541407    
## purpose       0.54196   0.26257  2.064  0.039447 *   
## loneliness   2.48823   0.53128  4.683  0.00000351 *** 
## purpose:loneliness -0.42711   0.12905 -3.310  0.000991 *** 
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 
## Residual standard error: 1.185 on 589 degrees of freedom
## Multiple R-squared:  0.1668, Adjusted R-squared:  0.1598 
## F-statistic: 23.59 on 5 and 589 DF,  p-value: < 0.0000000000000022

mod2dem <- glm(dementia ~ age + genderDc + purpose_z + loneliness_z + purpose_z * loneliness_z,
                 data = dataCf)
summary(mod2dem)

##
## Call:
## glm(formula = dementia ~ age + genderDc + purpose_z + loneliness_z +
##     purpose_z * loneliness_z, data = dataCf)
##
## Deviance Residuals:
##    Min      1Q  Median      3Q     Max 
## -0.78984 -0.15335 -0.09322 -0.05363  0.95329 
## 
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) -0.195174   0.328991 -0.593  0.55324    
## age          0.004482   0.004593  0.976  0.32960
```

```

## genderDc          0.019990  0.028190  0.709   0.47855
## purpose_z         -0.035139  0.017470 -2.011   0.04474 *
## loneliness_z     0.075203  0.017153  4.384  0.00000138 ***
## purpose_z:loneliness_z -0.037151  0.011760 -3.159   0.00166 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1161233)
##
## Null deviance: 78.464  on 594  degrees of freedom
## Residual deviance: 68.397  on 589  degrees of freedom
## AIC: 415.41
##
## Number of Fisher Scoring iterations: 2
# standardized age for tables
mod2cdA <- lm(decline ~ age_z + genderDc + purpose + loneliness + purpose * loneliness,
               data = dataCf)
summary(mod2cdA)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose + loneliness +
##     purpose * loneliness, data = dataCf)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3.2032 -0.5674 -0.3246  0.3702  7.0913
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -2.81398   1.14645 -2.455  0.014396 *
## age_z        0.06364   0.04894  1.300  0.193971
## genderDc    -0.05990   0.09802 -0.611  0.541407
## purpose      0.54196   0.26257  2.064  0.039447 *
## loneliness   2.48823   0.53128  4.683  0.00000351 ***
## purpose:loneliness -0.42711   0.12905 -3.310  0.000991 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.185 on 589 degrees of freedom
## Multiple R-squared:  0.1668, Adjusted R-squared:  0.1598
## F-statistic: 23.59 on 5 and 589 DF,  p-value: < 0.0000000000000022
confint(mod2cdA)

##
##                  2.5 %    97.5 %
## (Intercept) -5.06561660 -0.5623434
## age_z        -0.03247768  0.1597662
## genderDc     -0.25241705  0.1326229
## purpose       0.02627964  1.0576360
## loneliness   1.44479530  3.5316600
## purpose:loneliness -0.68055115 -0.1736608
mod2demA <- glm(dementia ~ age_z + genderDc + purpose_z + loneliness_z + purpose_z * loneliness_z,
                 data = dataCf)

```

```

summary(mod2demA)

##
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + loneliness_z +
##       purpose_z * loneliness_z, data = dataCf)
##
## Deviance Residuals:
##      Min        1Q     Median        3Q       Max
## -0.78984 -0.15335 -0.09322 -0.05363  0.95329
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)             0.12495   0.02219   5.631 0.0000000278 ***
## age_z                  0.01373   0.01407   0.976   0.32960
## genderDc                0.01999   0.02819   0.709   0.47855
## purpose_z               -0.03514   0.01747  -2.011   0.04474 *
## loneliness_z            0.07520   0.01715   4.384 0.0000137914 ***
## purpose_z:loneliness_z -0.03715   0.01176  -3.159   0.00166 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1161233)
##
## Null deviance: 78.464 on 594 degrees of freedom
## Residual deviance: 68.397 on 589 degrees of freedom
## AIC: 415.41
##
## Number of Fisher Scoring iterations: 2

confint(mod2demA)

## Waiting for profiling to be done...

##                               2.5 %      97.5 %
## (Intercept)             0.08145663  0.1684424319
## age_z                  -0.01385300  0.0413196580
## genderDc                -0.03526234  0.0752414567
## purpose_z               -0.06937979 -0.0008978967
## loneliness_z            0.04158280  0.1088226524
## purpose_z:loneliness_z -0.06020002 -0.0141014549

# odds ratio effects
exp(coef(mod2demA))

##                               (Intercept)           age_z           genderDc
##                         1.1330913          1.0138281          1.0201907
##                         purpose_z        loneliness_z purpose_z:loneliness_z
##                         0.9654714          1.0781027          0.9635309

exp(confint(mod2demA))

## Waiting for profiling to be done...

##                               2.5 %      97.5 %
## (Intercept)             1.0848662  1.1834601
## age_z                  0.9862425  1.0421852

```

```
## genderDc          0.9653521 1.0781444
## purpose_z          0.9329723 0.9991025
## loneliness_z      1.0424595 1.1149646
## purpose_z:loneliness_z 0.9415762 0.9859975
```

Simple Slopes Visualization

```

# creating data with low purpose and mean age
predictDataLow1 <- with(dataCf,
                        data.frame(age = mean(dataCf$age),
                                   genderDc = .5,
                                   purpose = mean(dataCf$purpose, na.rm=T)-
                                       sd(dataCf$purpose, na.rm=T),
                                   loneliness = seq(from = mean(dataCf$loneliness, na.rm=T)-
                                       (sd(dataCf$loneliness, na.rm=T)),
                                       to = mean(dataCf$loneliness, na.rm=T)+
                                           (sd(dataCf$loneliness, na.rm=T)),
                                       0.01)))

# creating data with mean purpose and mean age
predictDataMean1 <- with(dataCf,
                        data.frame(age = mean(dataCf$age),
                                   genderDc = .5,
                                   purpose = mean(dataCf$purpose, na.rm=T),
                                   loneliness = seq(mean(dataCf$loneliness, na.rm=T)-
                                       (sd(dataCf$loneliness, na.rm=T)),
                                       mean(dataCf$loneliness, na.rm=T)+
                                           (sd(dataCf$loneliness, na.rm=T)),
                                       0.01)))

# creating data with high purpose and mean age
predictDataHigh1 <- with(dataCf,
                        data.frame(age = mean(dataCf$age),
                                   genderDc = .5,
                                   purpose = mean(dataCf$purpose, na.rm=T)+
                                       sd(dataCf$purpose, na.rm=T),
                                   loneliness = seq(mean(dataCf$loneliness, na.rm=T)-
                                       (sd(dataCf$loneliness, na.rm=T)),
                                       mean(dataCf$loneliness, na.rm=T)+
                                           (sd(dataCf$loneliness, na.rm=T)),
                                       0.01)))

# get predicted values for pa based on three conditions
predictedLow1 <- predict(mod2cd,
                           newdata = predictDataLow1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
predictedMean1 <- predict(mod2cd, newdata = predictDataMean1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
predictedHigh1 <- predict(mod2cd, newdata = predictDataHigh1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")

# combine data frames
predictDataAll1 <- as.data.frame(rbind(predictDataLow1,
                                         predictDataMean1,
                                         predictDataHigh1))
predictions1 <- rbind(as.matrix(predictedLow1),
                      as.matrix(predictedMean1),

```

```

    as.matrix(predictedHigh1))
predictions1 <- cbind(predictDataAll1, predictions1)

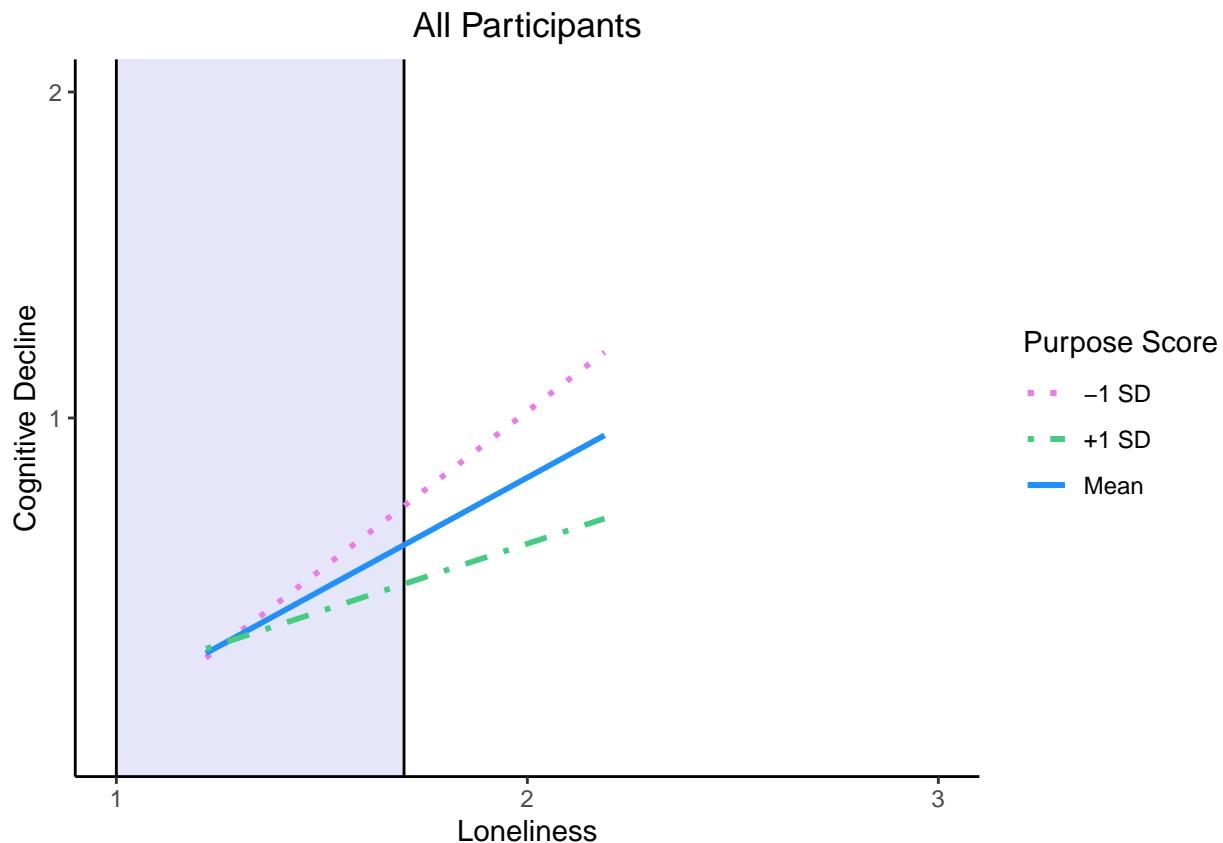
predictionsWide1 <- cbind(predictDataLow1, predictDataMean1, predictDataHigh1,
                           as.matrix(predictedLow1),
                           as.matrix(predictedMean1),
                           as.matrix(predictedHigh1))

names(predictionsWide1) <- c("ageL", "genderDcL", "purpL", "lonelyL",
                            "ageM", "genderDcM", "purpM", "lonelyM",
                            "ageH", "genderDcH", "purpH", "lonelyH",
                            "predictedCdL", "predictedCdM", "predictedCdH")

# getting groupings
colors <- c("-1 SD" = "orchid2", "+1 SD" = "seagreen3", "Mean" = "dodgerblue")
linetype <- c("-1 SD" = "dotted", "+1 SD" = "dotdash", "Mean" = "solid")

# graphing
library(ggplot2)
ggplot(predictionsWide1) +
  geom_rect(aes(xmin = 1, xmax = 1.70, ymin = -Inf, ymax = Inf),
            fill = "lavender") +
  geom_vline(xintercept = 1) +
  geom_vline(xintercept = 1.70) +
  geom_line(aes(x = lonelyL, y = predictedCdL, color = "-1 SD", linetype = "-1 SD"), size = 1) +
  geom_line(aes(x = lonelyM, y = predictedCdM, color = "Mean", linetype = "Mean"), size = 1) +
  geom_line(aes(x = lonelyH, y = predictedCdH, color = "+1 SD", linetype = "+1 SD"), size = 1) +
  expand_limits(x=c(1,3), y=c(0,2)) +
  scale_x_continuous(breaks = c(1,2,3,4,5)) +
  scale_y_continuous(breaks = c(1,2,3)) +
  theme_classic() +
  theme(plot.title = element_text(hjust = .5)) +
  labs(x = "Loneliness",
       y = "Cognitive Decline",
       color = "Purpose Score",
       linetype = "Purpose Score",
       title = "All Participants") +
  scale_color_manual(values = colors) +
  scale_linetype_manual(values = linetype)

```



```

# filtering based on values of loneliness
lowPurp <- dataCf %>% filter(purpose_z < -1)
meanPurp <- dataCf %>% filter(purpose_z < 1 & purpose_z > -1)
highPurp <- dataCf %>% filter(purpose_z > 1)

# models based on simple slope levels
modL <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z,
           data = lowPurp)
summary(modL)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z,
##      data = lowPurp)
##
## Residuals:
##      Min       1Q   Median       3Q      Max 
## -2.9931 -1.2572 -0.3184  0.7977  5.1261 
## 
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 1.2306    0.5626   2.187   0.0316 *  
## age_z       -0.2029    0.2021  -1.004   0.3183    
## genderDc    -0.4138    0.3804  -1.088   0.2799    
## purpose_z     0.1631    0.2890   0.564   0.5740    
## loneliness_z  0.7816    0.1905   4.103  0.0000966 ***
## ---
```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.721 on 81 degrees of freedom
## Multiple R-squared:  0.2299, Adjusted R-squared:  0.1918
## F-statistic: 6.044 on 4 and 81 DF,  p-value: 0.0002624
modM <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z,
            data = meanPurp)
summary(modM)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z,
##      data = meanPurp)
##
## Residuals:
##    Min      1Q  Median      3Q     Max 
## -1.7512 -0.6440 -0.3764  0.3038  7.0910 
##
## Coefficients:
##             Estimate Std. Error t value     Pr(>|t|)    
## (Intercept)  0.66518   0.08678   7.665 0.000000000000136 ***
## age_z        0.12742   0.05753   2.215     0.0273 *  
## genderDc     -0.04360  0.11471  -0.380     0.7041    
## purpose_z    -0.12766  0.10919  -1.169     0.2430    
## loneliness_z 0.29856   0.07045   4.238 0.000028032256489 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.144 on 401 degrees of freedom
## Multiple R-squared:  0.06433, Adjusted R-squared:  0.055 
## F-statistic: 6.893 on 4 and 401 DF,  p-value: 0.00002253
modH <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z,
            data = highPurp)
summary(modH)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z,
##      data = highPurp)
##
## Residuals:
##    Min      1Q  Median      3Q     Max 
## -0.5711 -0.3510 -0.2303 -0.0577  3.4740 
##
## Coefficients: (1 not defined because of singularities)
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept)  0.272819   0.102148   2.671  0.00885 ** 
## age_z       -0.009002   0.059788  -0.151  0.88063    
## genderDc     0.211720   0.131093   1.615  0.10948    
## purpose_z      NA         NA         NA         NA        
## loneliness_z  0.176594   0.083626   2.112  0.03723 *  
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

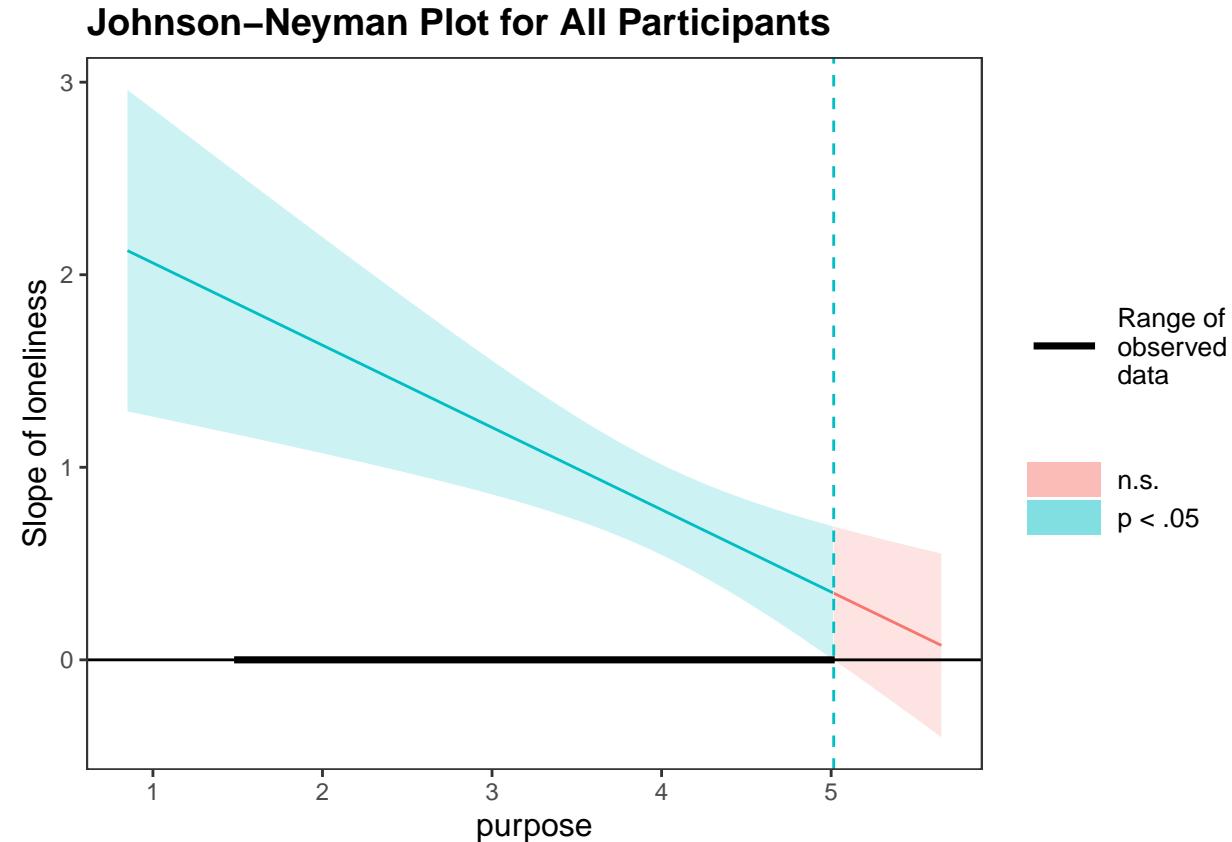
```

```
##  
## Residual standard error: 0.6423 on 99 degrees of freedom  
## Multiple R-squared:  0.05605,   Adjusted R-squared:  0.02745  
## F-statistic:  1.96 on 3 and 99 DF,  p-value: 0.125
```

Johnson-Neyman Technique

```
johnson_neyman(model = mod2cd, pred = loneliness, modx = purpose,
                 title = "Johnson-Neyman Plot for All Participants")

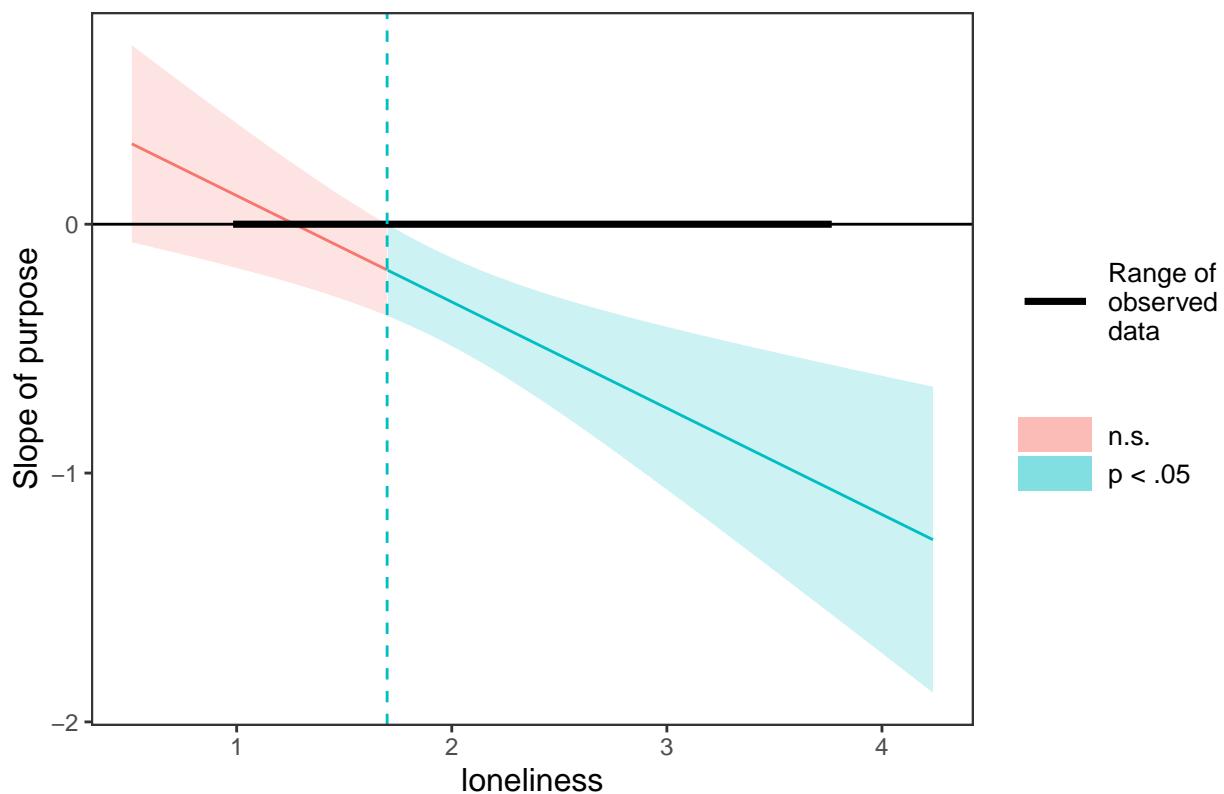
## JOHNSON-NEYMAN INTERVAL
##
## When purpose is OUTSIDE the interval [5.02, 8.61], the slope of loneliness
## is p < .05.
##
## Note: The range of observed values of purpose is [1.50, 5.00]
```



```
johnson_neyman(model = mod2cd, pred = purpose, modx = loneliness,
                 title = "Johnson-Neyman Plot for All Participants")

## JOHNSON-NEYMAN INTERVAL
##
## When loneliness is OUTSIDE the interval [0.14, 1.70], the slope of purpose
## is p < .05.
##
## Note: The range of observed values of loneliness is [1.00, 3.75]
```

Johnson–Neyman Plot for All Participants



Purpose X Depression

```
# for graphing
mod3cd <- lm(decline ~ age + genderDc + purpose_z + depression_z +
               data = dataCf)
summary(mod3cd)

##
## Call:
## lm(formula = decline ~ age + genderDc + purpose_z + depression_z +
##     purpose_z * depression_z, data = dataCf)
##
## Residuals:
##      Min    1Q   Median    3Q   Max 
## -2.6073 -0.5435 -0.1778  0.1910  7.2184 
##
## Coefficients:
##             Estimate Std. Error t value    Pr(>|t|)    
## (Intercept) -0.43678   1.04666  -0.417   0.6766    
## age          0.01721   0.01458   1.180   0.2384    
## genderDc     -0.17867  0.09043  -1.976   0.0487 *  
## purpose_z     -0.03825  0.05195  -0.736   0.4618    
## depression_z  0.70959   0.06369  11.141 <0.0000000000000002 *** 
## purpose_z:depression_z 0.01741   0.03657   0.476   0.6341  
## --- 
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
##
## Residual standard error: 1.082 on 589 degrees of freedom
## Multiple R-squared:  0.3048, Adjusted R-squared:  0.2989 
## F-statistic: 51.65 on 5 and 589 DF,  p-value: < 0.0000000000000022

mod3dem <- glm(dementia ~ age + genderDc + purpose_z + depression_z +
                 purpose_z * depression_z, data = dataCf)
summary(mod3dem)

##
## Call:
## glm(formula = dementia ~ age + genderDc + purpose_z + depression_z +
##     purpose_z * depression_z, data = dataCf)
##
## Deviance Residuals:
##      Min    1Q   Median    3Q   Max 
## -0.72295 -0.15684 -0.07288  0.00289  0.95844 
##
## Coefficients:
##             Estimate Std. Error t value    Pr(>|t|)    
## (Intercept) -0.081633  0.312006  -0.262   0.794    
## age          0.003472  0.004348   0.799   0.425    
## genderDc     -0.009707  0.026957  -0.360   0.719    
## purpose_z     -0.019055  0.015485  -1.231   0.219    
## depression_z  0.169301  0.018986   8.917 <0.0000000000000002 *** 
## purpose_z:depression_z 0.009242  0.010902   0.848   0.397  
## --- 
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
##
```

```

## (Dispersion parameter for gaussian family taken to be 0.1041048)
##
## Null deviance: 78.464 on 594 degrees of freedom
## Residual deviance: 61.318 on 589 degrees of freedom
## AIC: 350.4
##
## Number of Fisher Scoring iterations: 2
# standardized age for tables
mod3cdA <- lm(decline ~ age_z + genderDc + purpose_z + depression_z + purpose_z * depression_z,
               data = dataCf)
summary(mod3cdA)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + depression_z +
##     purpose_z * depression_z, data = dataCf)
##
## Residuals:
##    Min      1Q  Median      3Q      Max
## -2.6073 -0.5435 -0.1778  0.1910  7.2184
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.79278   0.07157 11.077 <0.0000000000000002 ***
## age_z       0.05275   0.04469  1.180   0.2384
## genderDc   -0.17867   0.09043 -1.976   0.0487 *
## purpose_z   -0.03825   0.05195 -0.736   0.4618
## depression_z 0.70959   0.06369 11.141 <0.0000000000000002 ***
## purpose_z:depression_z 0.01741   0.03657  0.476   0.6341
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.082 on 589 degrees of freedom
## Multiple R-squared:  0.3048, Adjusted R-squared:  0.2989
## F-statistic: 51.65 on 5 and 589 DF,  p-value: < 0.0000000000000022
confint(mod3cdA)

##
##                  2.5 %      97.5 %
## (Intercept) 0.65221631  0.933339024
## age_z       -0.03502957  0.140526260
## genderDc   -0.35627347 -0.001058991
## purpose_z   -0.14027551  0.063765851
## depression_z 0.58449690  0.834679364
## purpose_z:depression_z -0.05441096  0.089240667

mod3demA <- glm(dementia ~ age_z + genderDc + purpose_z + depression_z + purpose_z * depression_z,
                  data = dataCf)
summary(mod3demA)

##
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + depression_z +
##     purpose_z * depression_z, data = dataCf)
##

```

```

## Deviance Residuals:
##      Min       1Q   Median       3Q      Max
## -0.72295 -0.15684 -0.07288  0.00289  0.95844
##
## Coefficients:
##                               Estimate Std. Error t value     Pr(>|t|)
## (Intercept)               0.166380  0.021334  7.799 0.000000000000286 ***
## age_z                     0.010640  0.013323  0.799  0.425
## genderDc                 -0.009707 0.026957 -0.360  0.719
## purpose_z                -0.019055 0.015485 -1.231  0.219
## depression_z              0.169301  0.018986  8.917 < 0.0000000000000002 ***
## purpose_z:depression_z   0.009242  0.010902  0.848  0.397
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1041048)
##
## Null deviance: 78.464 on 594 degrees of freedom
## Residual deviance: 61.318 on 589 degrees of freedom
## AIC: 350.4
##
## Number of Fisher Scoring iterations: 2
confint(mod3demA)

## Waiting for profiling to be done...
##                               2.5 %    97.5 %
## (Intercept)           0.12456489 0.20819429
## age_z                 -0.01547275 0.03675224
## genderDc              -0.06254230 0.04312819
## purpose_z              -0.04940452 0.01129445
## depression_z           0.13208886 0.20651405
## purpose_z:depression_z -0.01212452 0.03060949
# odds ratio effects
exp(coef(mod3demA))

##                               (Intercept)          age_z          genderDc
##                         1.1810213        1.0106965        0.9903399
##                         purpose_z      depression_z purpose_z:depression_z
##                         0.9811254        1.1844771        1.0092853
exp(confint(mod3demA))

## Waiting for profiling to be done...
##                               2.5 %    97.5 %
## (Intercept)           1.1326555 1.231452
## age_z                 0.9846463 1.037436
## genderDc              0.9393733 1.044072
## purpose_z              0.9517960 1.011358
## depression_z           1.1412097 1.229385
## purpose_z:depression_z 0.9879487 1.031083

```

Exploratory Race Analyses

loneliness

The association between sense of purpose and loneliness is moderated by race (i.e., three-way interaction), wherein the influence of sense of purpose one the connection between loneliness on cognitive decline in stronger for Black participants than white participants. In other words, sense of purpose appears to provide an even more powerful buffering effect for Black participants than white participants when it comes to the negative influence of loneliness on cognitive decline.

```
### STANDARDIZING AGE FOR TABLES
modExCdLa <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z + raceDc +
                  purpose_z * loneliness_z +
                  raceDc * purpose_z +
                  raceDc * loneliness_z +
                  purpose_z * loneliness_z * raceDc,
                  data = dataCf)
summary(modExCdLa)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z +
##     raceDc + purpose_z * loneliness_z + raceDc * purpose_z +
##     raceDc * loneliness_z + purpose_z * loneliness_z * raceDc,
##     data = dataCf)
##
## Residuals:
##      Min      1Q  Median      3Q     Max 
## -2.8693 -0.5712 -0.3189  0.3807  7.0188 
##
## Coefficients:
##              Estimate Std. Error t value    Pr(>|t|)    
## (Intercept) 0.64333   0.08180   7.865 0.0000000000000184
## age_z        0.07121   0.04920   1.447  0.1483    
## genderDc     -0.09052   0.09841  -0.920  0.3581    
## purpose_z     -0.09604   0.06786  -1.415  0.1575    
## loneliness_z 0.33022   0.06852   4.819 0.0000018461304584
## raceDc        0.10830   0.14088   0.769  0.4423    
## purpose_z:loneliness_z -0.10625   0.04519  -2.351  0.0191    
## purpose_z:raceDc -0.29883   0.15975  -1.871  0.0619    
## loneliness_z:raceDc -0.08840   0.13957  -0.633  0.5267    
## purpose_z:loneliness_z:raceDc -0.25157   0.10872  -2.314  0.0210  
##
## (Intercept) ***
## age_z
## genderDc
## purpose_z
## loneliness_z ***
## raceDc
## purpose_z:loneliness_z *
## purpose_z:raceDc .
## loneliness_z:raceDc
## purpose_z:loneliness_z:raceDc *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```

## Residual standard error: 1.174 on 574 degrees of freedom
##   (11 observations deleted due to missingness)
## Multiple R-squared:  0.1868, Adjusted R-squared:  0.174
## F-statistic: 14.65 on 9 and 574 DF,  p-value: < 0.00000000000000022

confint(modExCdLa)

##                               2.5 %      97.5 %
## (Intercept)          0.48267049  0.80399519
## age_z              -0.02542291  0.16783326
## genderDc           -0.28380283  0.10276857
## purpose_z            -0.22932203  0.03724403
## loneliness_z        0.19564253  0.46480419
## raceDc              -0.16839573  0.38499865
## purpose_z:loneliness_z  -0.19501977 -0.01748824
## purpose_z:raceDc     -0.61259709  0.01494205
## loneliness_z:raceDc -0.36253072  0.18573281
## purpose_z:loneliness_z:raceDc -0.46510428 -0.03804421

# dichotomous with all participants
modExDemLa <- glm(dementia ~ age_z + genderDc + purpose_z + loneliness_z + raceDc +
                    purpose_z * loneliness_z +
                    raceDc * purpose_z +
                    raceDc * loneliness_z +
                    purpose_z * loneliness_z * raceDc,
                    data = dataCf)
summary(modExDemLa)

## 
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + loneliness_z +
##     raceDc + purpose_z * loneliness_z + raceDc * purpose_z +
##     raceDc * loneliness_z + purpose_z * loneliness_z * raceDc,
##     data = dataCf)
## 
## Deviance Residuals:
##      Min        1Q        Median         3Q        Max 
## -0.75694  -0.15340  -0.09564  -0.04178   0.95643 
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)          0.129858   0.023743  5.469 0.000000675 ***
## age_z                0.013847   0.014280  0.970  0.3326    
## genderDc             0.010502   0.028564  0.368  0.7132    
## purpose_z            -0.033867   0.019697 -1.719  0.0861 .  
## loneliness_z         0.084062   0.019889  4.227 0.0000275920 ***
## raceDc               -0.001718   0.040891 -0.042  0.9665    
## purpose_z:loneliness_z -0.030990   0.013118 -2.362  0.0185 *  
## purpose_z:raceDc      -0.022288   0.046369 -0.481  0.6309    
## loneliness_z:raceDc -0.056815   0.040512 -1.402  0.1613    
## purpose_z:loneliness_z:raceDc -0.042275   0.031556 -1.340  0.1809    
## ---                
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 
## (Dispersion parameter for gaussian family taken to be 0.1160656)
## 
```

```

##      Null deviance: 76.820  on 583  degrees of freedom
## Residual deviance: 66.622  on 574  degrees of freedom
##   (11 observations deleted due to missingness)
## AIC: 411.53
##
## Number of Fisher Scoring iterations: 2
confint(modExDemLa)

## Waiting for profiling to be done...

##                               2.5 %      97.5 %
## (Intercept)          0.08332226  0.176392820
## age_z              -0.01414060  0.041835371
## genderDc           -0.04548212  0.066486914
## purpose_z            -0.07247189  0.004738023
## loneliness_z        0.04508105  0.123042770
## raceDc              -0.08186204  0.078426677
## purpose_z:loneliness_z  -0.05670051 -0.005279129
## purpose_z:raceDc     -0.11317070  0.068593787
## loneliness_z:raceDc -0.13621620  0.022586387
## purpose_z:loneliness_z:raceDc -0.10412292  0.019573513

# odds ratios
exp(coef(modExDemLa))

##                               (Intercept)          age_z
##                         1.1386662          1.0139437
##                         genderDc          purpose_z
##                         1.0105577          0.9667001
##                         loneliness_z          raceDc
##                         1.0876962          0.9982838
##                         purpose_z:loneliness_z          purpose_z:raceDc
##                         0.9694854          0.9779581
##                         loneliness_z:raceDc purpose_z:loneliness_z:raceDc
##                         0.9447689          0.9586064

exp(confint(modExDemLa))

## Waiting for profiling to be done...

##                               2.5 %      97.5 %
## (Intercept)          1.0868920  1.1929066
## age_z              0.9859589  1.0427228
## genderDc           0.9555367  1.0687470
## purpose_z            0.9300919  1.0047493
## loneliness_z        1.0461126  1.1309328
## raceDc              0.9213991  1.0815840
## purpose_z:loneliness_z  0.9448770  0.9947348
## purpose_z:raceDc     0.8929982  1.0710011
## loneliness_z:raceDc 0.8726539  1.0228434
## purpose_z:loneliness_z:raceDc 0.9011145  1.0197663

### FOR GRAPHING
# continuous with all participants
modExCdL <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z + raceDc +
                 purpose_z * loneliness_z +
                 raceDc * purpose_z +

```

```

            raceDc * loneliness_z +
            purpose_z * loneliness_z * raceDc,
            data = dataCf)
summary(modExCdL)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z +
##     raceDc + purpose_z * loneliness_z + raceDc * purpose_z +
##     raceDc * loneliness_z + purpose_z * loneliness_z * raceDc,
##     data = dataCf)
##
## Residuals:
##    Min      1Q  Median      3Q     Max 
## -2.8693 -0.5712 -0.3189  0.3807  7.0188 
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)    
## (Intercept)               0.64333   0.08180  7.865 0.000000000000184
## age_z                     0.07121   0.04920  1.447  0.1483    
## genderDc                  -0.09052   0.09841 -0.920  0.3581    
## purpose_z                 -0.09604   0.06786 -1.415  0.1575    
## loneliness_z                0.33022   0.06852  4.819 0.0000018461304584
## raceDc                      0.10830   0.14088  0.769  0.4423    
## purpose_z:loneliness_z      -0.10625   0.04519 -2.351  0.0191    
## purpose_z:raceDc              -0.29883   0.15975 -1.871  0.0619    
## loneliness_z:raceDc          -0.08840   0.13957 -0.633  0.5267    
## purpose_z:loneliness_z:raceDc -0.25157   0.10872 -2.314  0.0210    
##
## (Intercept) ***                                 
## age_z                                 ***      
## genderDc                                ***      
## purpose_z                                *        
## loneliness_z                            ***      
## raceDc                                 .        
## purpose_z:loneliness_z                   *        
## purpose_z:raceDc                         .        
## loneliness_z:raceDc                    *        
## purpose_z:loneliness_z:raceDc           *        
## ---                                 
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.174 on 574 degrees of freedom
##   (11 observations deleted due to missingness)
## Multiple R-squared:  0.1868, Adjusted R-squared:  0.174 
## F-statistic: 14.65 on 9 and 574 DF,  p-value: < 0.0000000000000022
# dichotomous with all participants
modExDemL <- glm(dementia ~ age_z + genderDc + purpose_z + loneliness_z + raceDc +
                  purpose_z * loneliness_z +
                  raceDc * purpose_z +
                  raceDc * loneliness_z +
                  purpose_z * loneliness_z * raceDc,
                  data = dataCf)

```

```

summary(modExDemL)

##
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + loneliness_z +
##      raceDc + purpose_z * loneliness_z + raceDc * purpose_z +
##      raceDc * loneliness_z + purpose_z * loneliness_z * raceDc,
##      data = dataCf)
##
## Deviance Residuals:
##    Min      1Q   Median      3Q     Max
## -0.75694 -0.15340 -0.09564 -0.04178  0.95643
##
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)                0.129858   0.023743   5.469 0.0000000675 ***
## age_z                      0.013847   0.014280   0.970  0.3326
## genderDc                   0.010502   0.028564   0.368  0.7132
## purpose_z                  -0.033867  0.019697  -1.719  0.0861 .
## loneliness_z                0.084062  0.019889   4.227 0.0000275920 ***
## raceDc                     -0.001718  0.040891  -0.042  0.9665
## purpose_z:loneliness_z    -0.030990  0.013118  -2.362  0.0185 *
## purpose_z:raceDc            -0.022288  0.046369  -0.481  0.6309
## loneliness_z:raceDc       -0.056815  0.040512  -1.402  0.1613
## purpose_z:loneliness_z:raceDc -0.042275  0.031556  -1.340  0.1809
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1160656)
##
## Null deviance: 76.820 on 583 degrees of freedom
## Residual deviance: 66.622 on 574 degrees of freedom
##   (11 observations deleted due to missingness)
## AIC: 411.53
##
## Number of Fisher Scoring iterations: 2

#### STRATIFICATION
# making dataframes separated by race
white <- dataCf %>%
  filter(raceDc == 0)
black <- dataCf %>%
  filter(raceDc == 1)

# continuous with white people only (standardized)
modExCdLz <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z +
                  purpose_z * loneliness_z,
                  data = white)
summary(modExCdLz)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z +
##      purpose_z * loneliness_z, data = white)

```

```

## 
## Residuals:
##   Min     1Q Median     3Q    Max
## -2.7986 -0.5489 -0.3174  0.4157  6.7306
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)               0.57769   0.08269   6.986 0.0000000000974 ***
## age_z                     0.05163   0.05212   0.990   0.3225
## genderDc                  0.02478   0.10457   0.237   0.8128
## purpose_z                 -0.09590   0.06525  -1.470   0.1423
## loneliness_z                0.33073   0.06586   5.022 0.00000072891379 ***
## purpose_z:loneliness_z   -0.10898   0.04345  -2.508   0.0125 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 
## Residual standard error: 1.128 on 469 degrees of freedom
## Multiple R-squared:  0.1553, Adjusted R-squared:  0.1463
## F-statistic: 17.24 on 5 and 469 DF,  p-value: 0.000000000000001127
confint(modExCdLz)

##                               2.5 %      97.5 %
## (Intercept)               0.4151982  0.74018437
## age_z                   -0.0507985  0.15405384
## genderDc                  -0.1807110  0.23026668
## purpose_z                 -0.2241276  0.03231945
## loneliness_z                0.2013154  0.46015146
## purpose_z:loneliness_z   -0.1943711 -0.02359474

# continuous with black people only (standardized)
modExCdBz <- lm(decline ~ age_z + genderDc + purpose_z + loneliness_z +
                  purpose_z * loneliness_z,
                  data = black)
summary(modExCdBz)

## 
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + loneliness_z +
##     purpose_z * loneliness_z, data = black)
## 
## Residuals:
##   Min     1Q Median     3Q    Max
## -2.6277 -0.7133 -0.3139  0.1616  6.7154
## 
## Coefficients:
##                               Estimate Std. Error t value Pr(>|t|)
## (Intercept)               1.0332    0.2018   5.119 0.00000143 ***
## age_z                     0.1627    0.1333   1.221   0.22494
## genderDc                  -0.6088   0.2630  -2.315   0.02258 *
## purpose_z                 -0.3497   0.1674  -2.089   0.03916 *
## loneliness_z                0.2514   0.1459   1.723   0.08787 .
## purpose_z:loneliness_z   -0.3831   0.1134  -3.377   0.00104 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

## 
## Residual standard error: 1.338 on 103 degrees of freedom
## Multiple R-squared:  0.303, Adjusted R-squared:  0.2692
## F-statistic: 8.957 on 5 and 103 DF, p-value: 0.0000004316
confint(modExCdBz)

##          2.5 %      97.5 %
## (Intercept) 0.63289163 1.43345027
## age_z       -0.10159588 0.42694552
## genderDc    -1.13039358 -0.08729402
## purpose_z   -0.68167662 -0.01772676
## loneliness_z -0.03795525 0.54074116
## purpose_z:loneliness_z -0.60803260 -0.15807620

# continuous with white people only (plotting)
modExCdL <- lm(decline ~ age + genderDc + purpose + loneliness +
                 purpose * loneliness,
                 data = white)
summary(modExCdL)

## 
## Call:
## lm(formula = decline ~ age + genderDc + purpose + loneliness +
##     purpose * loneliness, data = white)
## 
## Residuals:
##      Min      1Q Median      3Q      Max 
## -2.7986 -0.5489 -0.3174  0.4157  6.7306 
## 
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) -3.63186   1.78197 -2.038 0.042099 *  
## age          0.01685   0.01701  0.990 0.322455    
## genderDc     0.02478   0.10457  0.237 0.812804    
## purpose       0.43896   0.28284  1.552 0.121346    
## loneliness   2.12720   0.56495  3.765 0.000187 *** 
## purpose:loneliness -0.34392  0.13713 -2.508 0.012478 * 
## --- 
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## 
## Residual standard error: 1.128 on 469 degrees of freedom
## Multiple R-squared:  0.1553, Adjusted R-squared:  0.1463
## F-statistic: 17.24 on 5 and 469 DF, p-value: 0.00000000000001127
confint(modExCdL)

##          2.5 %      97.5 %
## (Intercept) -7.13349841 -0.13022613
## age         -0.01657717  0.05027269
## genderDc    -0.18071102  0.23026668
## purpose     -0.11683519  0.99476311
## loneliness  1.01705642  3.23734801
## purpose:loneliness -0.61338009 -0.07445831

# continuous with black people only (plotting)
modExCdB <- lm(decline ~ age + genderDc + purpose + loneliness +

```

```

        purpose * loneliness,
      data = black)
summary(modExCdB)

##
## Call:
## lm(formula = decline ~ age + genderDc + purpose + loneliness +
##     purpose * loneliness, data = black)
##
## Residuals:
##    Min     1Q Median     3Q    Max
## -2.6277 -0.7133 -0.3139  0.1616  6.7154
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -10.05528   4.44406 -2.263 0.025757 *
## age          0.05309   0.04348  1.221 0.224941
## genderDc    -0.60884   0.26298 -2.315 0.022584 *
## purpose       1.52347   0.68862  2.212 0.029150 *
## loneliness    5.60738   1.46165  3.836 0.000216 ***
## purpose:loneliness -1.20881   0.35798 -3.377 0.001036 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.338 on 103 degrees of freedom
## Multiple R-squared:  0.303, Adjusted R-squared:  0.2692
## F-statistic: 8.957 on 5 and 103 DF,  p-value: 0.0000004316
confint(modExCdB)

##
##              2.5 %    97.5 %
## (Intercept) -18.86901679 -1.24154567
## age          -0.03315398  0.13932597
## genderDc    -1.13039358 -0.08729402
## purpose       0.15775541  2.88918589
## loneliness    2.70853157  8.50621929
## purpose:loneliness -1.91877876 -0.49884375

```

Simple Slopes Visualization

```
### Graphing for white participants

# creating data with low purpose and mean age
predictDataLow1 <- with(white,
                       data.frame(age = mean(white$age),
                                  genderDc = .5,
                                  purpose = mean(white$purpose, na.rm=T)-
                                  sd(dataCf$purpose, na.rm=T),
                                  loneliness = seq(from = mean(white$loneliness, na.rm=T)-
                                  (sd(white$loneliness, na.rm=T)),
                                  to = mean(white$loneliness, na.rm=T)+
                                  (sd(white$loneliness, na.rm=T)),
                                  0.01)))

# creating data with mean purpose and mean age
predictDataMean1 <- with(white,
                         data.frame(age = mean(white$age),
                                    genderDc = .5,
                                    purpose = mean(white$purpose, na.rm=T),
                                    loneliness = seq(mean(white$loneliness, na.rm=T)-
                                    (sd(white$loneliness, na.rm=T)),
                                    mean(white$loneliness, na.rm=T)+
                                    (sd(white$loneliness, na.rm=T)),
                                    0.01)))

# creating data with high purpose and mean age
predictDataHigh1 <- with(white,
                         data.frame(age = mean(white$age),
                                    genderDc = .5,
                                    purpose = mean(white$purpose, na.rm=T)+
                                    sd(white$purpose, na.rm=T),
                                    loneliness = seq(mean(white$loneliness, na.rm=T)-
                                    (sd(white$loneliness, na.rm=T)),
                                    mean(white$loneliness, na.rm=T)+
                                    (sd(white$loneliness, na.rm=T)),
                                    0.01)))

# get predicted values for pa based on three conditions
predictedLow1 <- predict(modExCdL,
                          newdata = predictDataLow1,
                          n.sims = 1000, level = 0.95,
                          type = "response",
                          include.resid.var = FALSE, which = "fixed")
predictedMean1 <- predict(modExCdL, newdata = predictDataMean1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
predictedHigh1 <- predict(modExCdL, newdata = predictDataHigh1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")

# combine data frames
predictDataAll1 <- as.data.frame(rbind(predictDataLow1,
```

```

                                predictDataHigh1))
predictions1 <- rbind(as.matrix(predictedLow1),
                      as.matrix(predictedMean1),
                      as.matrix(predictedHigh1))
predictions1 <- cbind(predictDataAll1, predictions1)

predictionsWide1 <- cbind(predictDataLow1, predictDataMean1, predictDataHigh1,
                           as.matrix(predictedLow1),
                           as.matrix(predictedMean1),
                           as.matrix(predictedHigh1))

names(predictionsWide1) <- c("ageL", "genderL", "purpL", "lonelyL",
                            "ageM", "genderM", "purpM", "lonelyM",
                            "ageH", "genderH", "purpH", "lonelyH",
                            "predictedCdL", "predictedCdM", "predictedCdH")

# getting groupings
colors <- c("-1 SD" = "orchid2", "+1 SD" = "seagreen3", "Mean" = "dodgerblue")
linetype <- c("-1 SD" = "dotted", "+1 SD" = "dotdash", "Mean" = "solid")

# graphing
whiteGraph <- ggplot(predictionsWide1, aes(x = lonelyL, y = predictedCdL, color = "-1 SD", linetype = "-1 SD"),
                       geom_rect(aes(xmin = 1, xmax = 1.68, ymin = -Inf, ymax = Inf),
                                 fill = "lavender") +
                         geom_vline(xintercept = 1) +
                         geom_vline(xintercept = 1.68) +
                         geom_line(size = 1) +
                         geom_line(aes(x = lonelyM, y = predictedCdM, color = "Mean", linetype = "Mean"), size = 1) +
                         geom_line(aes(x = lonelyH, y = predictedCdH, color = "+1 SD", linetype = "+1 SD"), size = 1) +
                         expand_limits(x=c(1,3), y=c(0,2)) +
                         scale_x_continuous(breaks = c(1,2,3,4,5)) +
                         scale_y_continuous(breaks = c(1,2,3)) +
                         theme_classic() +
                         theme(plot.title = element_text(hjust = .5)) +
                         labs(x = "Loneliness",
                              y = "Cognitive Decline",
                              color = "Purpose Score",
                              linetype = "Purpose Score",
                              title = "White Participants") +
                         scale_color_manual(values = colors) +
                         scale_linetype_manual(values = linetype)

### Graphing for Black participants

# creating data with low purpose and mean age
predictDataLow1 <- with(black,
                        data.frame(age = mean(dataCf$age),
                                   genderDc = .5,
                                   purpose = mean(black$purpose, na.rm=T) -
                                     sd(black$purpose, na.rm=T),
                                   loneliness = seq(from = mean(black$loneliness, na.rm=T) -
                                                     (sd(black$loneliness, na.rm=T)),
                                                     to = mean(black$loneliness, na.rm=T) +
```

```

        (sd(black$loneliness, na.rm=T)),
        0.01)))
# creating data with mean purpose and mean age
predictDataMean1 <- with(black,
                         data.frame(age = mean(black$age),
                                     genderDc = .5,
                                     purpose = mean(black$purpose, na.rm=T),
                                     loneliness = seq(mean(black$loneliness, na.rm=T)-
                                         (sd(black$loneliness, na.rm=T)),
                                         mean(black$loneliness, na.rm=T)+
                                         (sd(black$loneliness, na.rm=T)),
                                         0.01)))
# creating data with high purpose and mean age
predictDataHigh1 <- with(black,
                         data.frame(age = mean(black$age),
                                     genderDc = .5,
                                     purpose = mean(black$purpose, na.rm=T)+
                                         sd(black$purpose, na.rm=T),
                                     loneliness = seq(mean(black$loneliness, na.rm=T)-
                                         (sd(black$loneliness, na.rm=T)),
                                         mean(black$loneliness, na.rm=T)+
                                         (sd(black$loneliness, na.rm=T)),
                                         0.01)))
# get predicted values for pa based on three conditions
predictedLow1 <- predict(modExCdB,
                           newdata = predictDataLow1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
predictedMean1 <- predict(modExCdB, newdata = predictDataMean1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
predictedHigh1 <- predict(modExCdB, newdata = predictDataHigh1,
                           n.sims = 1000, level = 0.95,
                           type = "response",
                           include.resid.var = FALSE, which = "fixed")
# combine data frames
predictDataAll1 <- as.data.frame(rbind(predictDataLow1,
                                         predictDataMean1,
                                         predictDataHigh1))
predictions1 <- rbind(as.matrix(predictedLow1),
                      as.matrix(predictedMean1),
                      as.matrix(predictedHigh1))
predictions1 <- cbind(predictDataAll1, predictions1)
names(predictions1) <- c("purpose", "loneliness", "age", "predictedCog")

predictionsWide1 <- cbind(predictDataLow1, predictDataMean1, predictDataHigh1,
                           as.matrix(predictedLow1),
                           as.matrix(predictedMean1),
                           as.matrix(predictedHigh1))

names(predictionsWide1) <- c("ageL", "genderL", "purpL", "lonelyL",

```

```

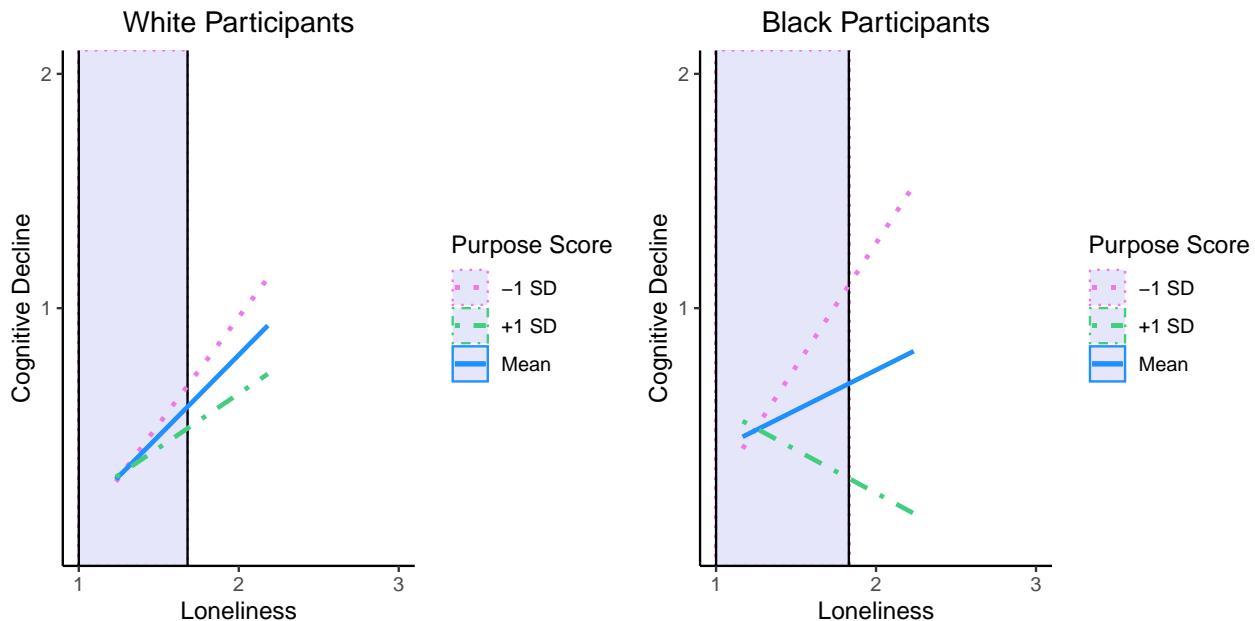
"ageM", "genderM", "purpM", "lonelyM",
"ageH", "genderH", "purpH", "lonelyH",
"predictedCdL", "predictedCdM", "predictedCdH")

# getting groupings
colors <- c("-1 SD" = "orchid2", "+1 SD" = "seagreen3", "Mean" = "dodgerblue")
linetype <- c("-1 SD" = "dotted", "+1 SD" = "dotdash", "Mean" = "solid")

# graphing
blackGraph <- ggplot(predictionsWide1, aes(x = lonelyL, y = predictedCdL, color = "-1 SD", linetype = "-1 SD"),
  geom_rect(aes(xmin = 1, xmax = 1.83, ymin = -Inf, ymax = Inf),
    fill = "lavender") +
  geom_vline(xintercept = 1) +
  geom_vline(xintercept = 1.83) +
  geom_line(size = 1) +
  geom_line(aes(x = lonelyM, y = predictedCdM, color = "Mean", linetype = "Mean"), size = 1) +
  geom_line(aes(x = lonelyH, y = predictedCdH, color = "+1 SD", linetype = "+1 SD"), size = 1) +
  expand_limits(x=c(1,3), y=c(0,2)) +
  scale_x_continuous(breaks = c(1,2,3,4,5)) +
  scale_y_continuous(breaks = c(1,2,3)) +
  theme_classic() +
  theme(plot.title = element_text(hjust = .5)) +
  labs(x = "Loneliness",
       y = "Cognitive Decline",
       color = "Purpose Score",
       linetype = "Purpose Score",
       title = "Black Participants") +
  scale_color_manual(values = colors) +
  scale_linetype_manual(values = linetype)

# make combined graph
ggarrange(whiteGraph, blackGraph)

```



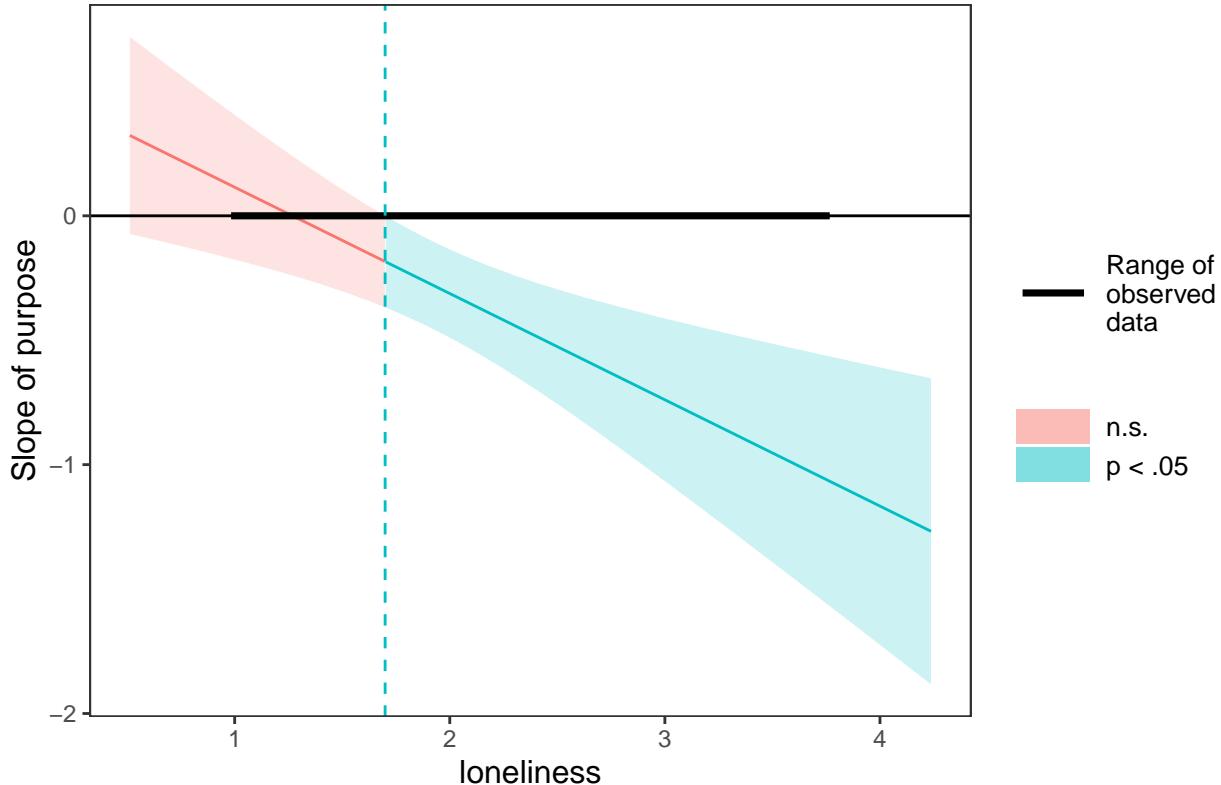
Johnson-Neyman Technique

```
# participant groups
jnAll <- lm(decline ~ age + genderDc + purpose + loneliness +
            purpose * loneliness,
            data = dataCf)
jnWhite <- lm(decline ~ age + genderDc + purpose + loneliness +
            purpose * loneliness,
            data = white)
jnBlack <- lm(decline ~ age + genderDc + purpose + loneliness +
            purpose * loneliness,
            data = black)

# johnson_neyman analyses
johnson_neyman(model = jnAll, pred = purpose, modx = loneliness,
                 title = "Johnson-Neyman Plot for White Participants")

## JOHNSON-NEYMAN INTERVAL
##
## When loneliness is OUTSIDE the interval [0.14, 1.70], the slope of purpose
## is p < .05.
##
## Note: The range of observed values of loneliness is [1.00, 3.75]
```

Johnson-Neyman Plot for White Participants



```
johnson_neyman(model = jnWhite, pred = purpose, modx = loneliness,
                 title = "Johnson-Neyman Plot for White Participants")
```

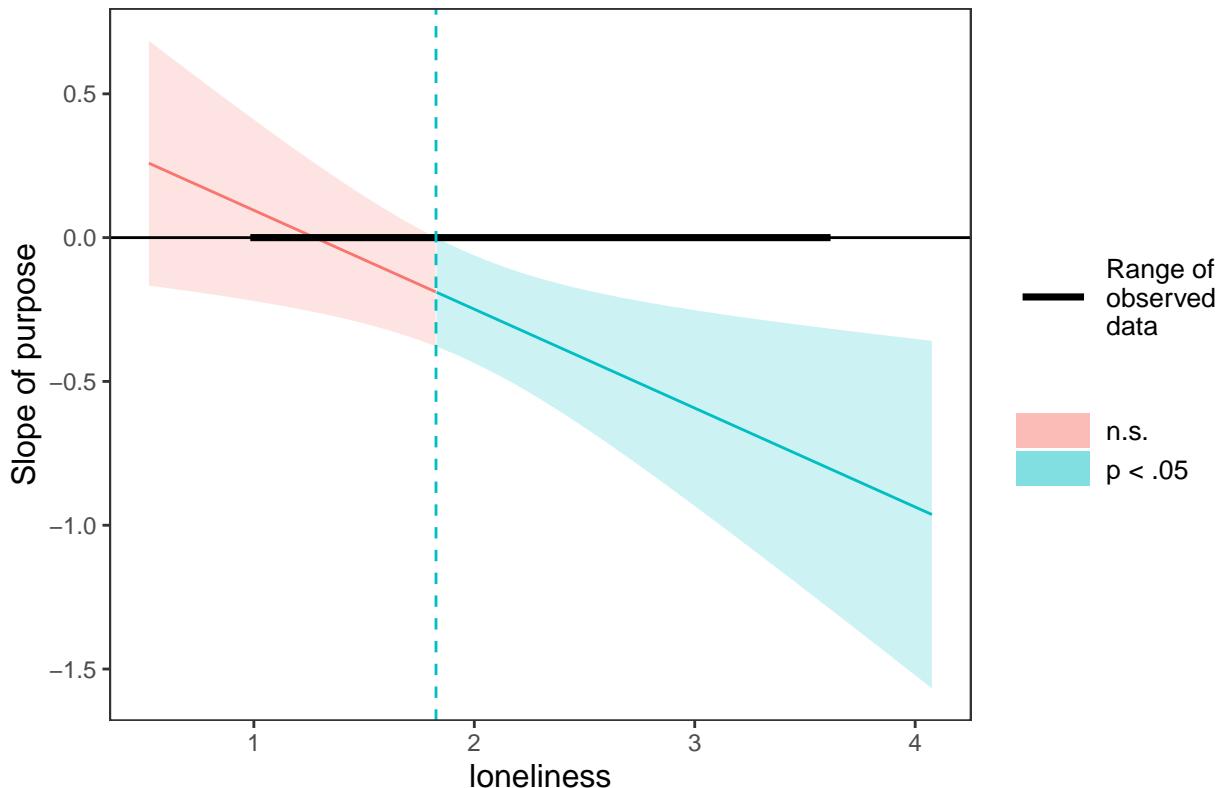
```
## JOHNSON-NEYMAN INTERVAL
```

```

## 
## When loneliness is OUTSIDE the interval [-1.39, 1.83], the slope of purpose
## is p < .05.
## 
## Note: The range of observed values of loneliness is [1.00, 3.60]

```

Johnson–Neyman Plot for White Participants



```

johnson_neyman(model = jnBlack, pred = purpose, modx = loneliness,
                 title = "Johnson–Neyman Plot for Black Participants")

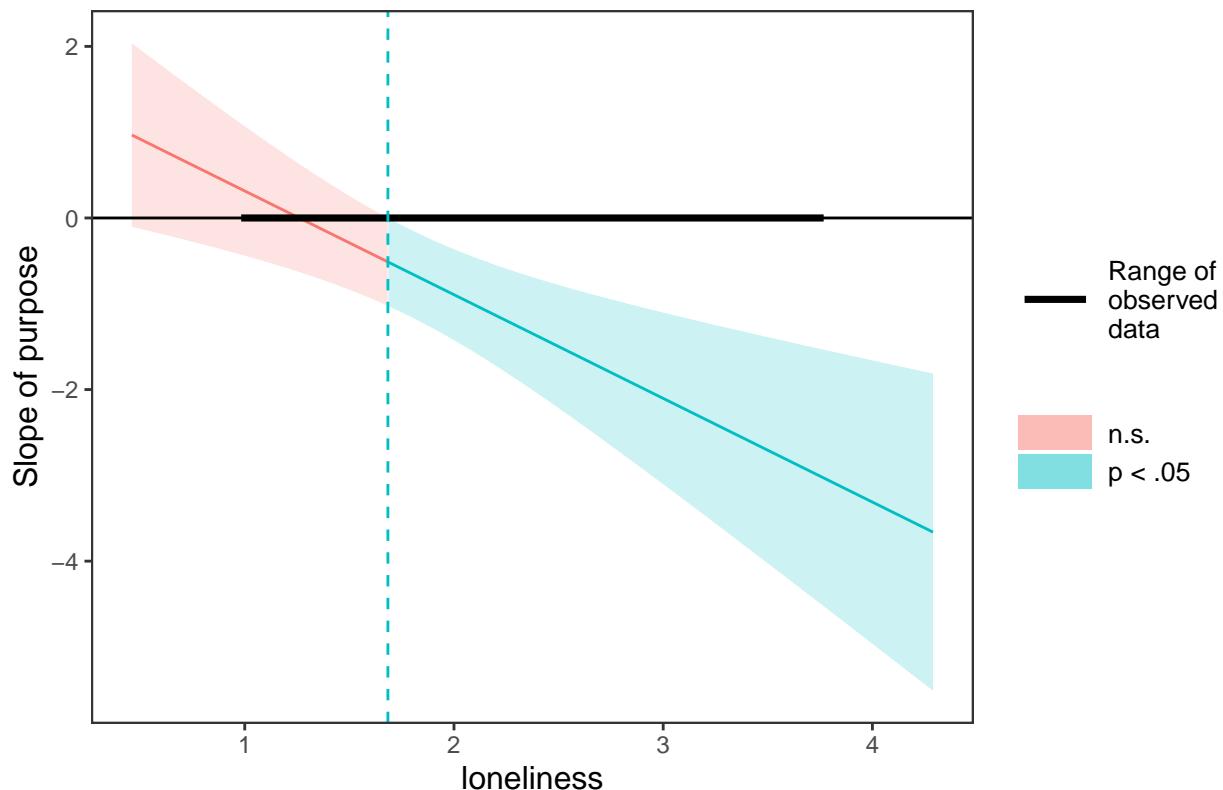
```

```

## JOHNSON–NEYMAN INTERVAL
## 
## When loneliness is OUTSIDE the interval [0.28, 1.68], the slope of purpose
## is p < .05.
## 
## Note: The range of observed values of loneliness is [1.00, 3.75]

```

Johnson–Neyman Plot for Black Participants



```
# proportion above value
table(dataCf$loneliness >= 1.70)

##
## FALSE TRUE
## 322 273
273/(322+273)

## [1] 0.4588235
table(white$loneliness >= 1.68)

##
## FALSE TRUE
## 255 220
220/(220+255)

## [1] 0.4631579
table(black$loneliness >= 1.83)

##
## FALSE TRUE
## 69 40
40/(69+40)

## [1] 0.3669725
```

depression

There is no interaction between the association for depression and loneliness with race when predicting cognitive decline.

```
modExCdD <- lm(decline ~ age_z + genderDc + purpose_z + depression_z + raceDc +
                  purpose_z * depression_z +
                  raceDc * purpose_z +
                  raceDc * depression_z +
                  purpose_z * depression_z * raceDc,
                  data = dataCf)
summary(modExCdD)

##
## Call:
## lm(formula = decline ~ age_z + genderDc + purpose_z + depression_z +
##     raceDc + purpose_z * depression_z + raceDc * purpose_z +
##     raceDc * depression_z + purpose_z * depression_z * raceDc,
##     data = dataCf)
##
## Residuals:
##    Min      1Q  Median      3Q     Max 
## -2.8424 -0.5355 -0.1833  0.2166  7.1292 
##
## Coefficients:
##              Estimate Std. Error t value     Pr(>|t|)    
## (Intercept) 0.78384   0.07638 10.262 <0.0000000000000002
## age_z        0.05934   0.04535  1.308   0.1913    
## genderDc     -0.18930   0.09184 -2.061   0.0397    
## purpose_z    -0.02821   0.05725 -0.493   0.6225    
## depression_z  0.70742   0.07202  9.822 <0.0000000000000002
## raceDc        0.08126   0.13100  0.620   0.5353    
## purpose_z:depression_z  0.02270   0.04136  0.549   0.5833    
## purpose_z:raceDc    -0.15638   0.15609 -1.002   0.3168    
## depression_z:raceDc -0.16426   0.17455 -0.941   0.3471    
## purpose_z:depression_z:raceDc -0.07603   0.09463 -0.804   0.4220  
##
## (Intercept) ***
## age_z        
## genderDc      
## purpose_z      
## depression_z ***
## raceDc        
## purpose_z:depression_z
## purpose_z:raceDc
## depression_z:raceDc
## purpose_z:depression_z:raceDc
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.088 on 574 degrees of freedom
##   (11 observations deleted due to missingness)
## Multiple R-squared:  0.301,  Adjusted R-squared:  0.29
## F-statistic: 27.46 on 9 and 574 DF,  p-value: < 0.0000000000000022
```

```

confint(modExCcdD)

##                               2.5 %      97.5 %
## (Intercept)          0.63382026  0.933858014
## age_z                -0.02973739  0.148420788
## genderDc             -0.36967425 -0.008916318
## purpose_z              -0.14065700  0.084246298
## depression_z           0.56596207  0.848873738
## raceDc                -0.17603496  0.338546285
## purpose_z:depression_z -0.05852740  0.103929804
## purpose_z:raceDc       -0.46296913  0.150199936
## depression_z:raceDc   -0.50709611  0.178580193
## purpose_z:depression_z:raceDc -0.26188769  0.109820661

modExDemD <- glm(dementia ~ age_z + genderDc + purpose_z + depression_z + raceDc +
                  purpose_z * depression_z +
                  raceDc * purpose_z +
                  raceDc * depression_z +
                  purpose_z * depression_z * raceDc,
                  data = dataCf)
summary(modExDemD)

```

```

##
## Call:
## glm(formula = dementia ~ age_z + genderDc + purpose_z + depression_z +
##       raceDc + purpose_z * depression_z + raceDc * purpose_z +
##       raceDc * depression_z + purpose_z * depression_z * raceDc,
##       data = dataCf)
##
## Deviance Residuals:
##      Min        1Q     Median        3Q       Max
## -0.63455 -0.15757 -0.07457 -0.00010  0.96477
##
## Coefficients:
##                               Estimate Std. Error t value
## (Intercept)          0.165257117 0.022757358 7.262
## age_z                 0.012227510 0.013512997 0.905
## genderDc             -0.014417954 0.027362881 -0.527
## purpose_z              -0.023821607 0.017058536 -1.396
## depression_z            0.164740904 0.021458373 7.677
## raceDc                0.0000006052 0.039030120 0.000
## purpose_z:depression_z 0.001994520 0.012322105 0.162
## purpose_z:raceDc       0.036758475 0.046507840 0.790
## depression_z:raceDc   0.008773503 0.052007392 0.169
## purpose_z:depression_z:raceDc 0.021931976 0.028193452 0.778
##                               Pr(>|t|)
## (Intercept)          0.0000000000012506 ***
## age_z                   0.366
## genderDc                0.598
## purpose_z                  0.163
## depression_z          0.0000000000000704 ***
## raceDc                   1.000
## purpose_z:depression_z    0.871
## purpose_z:raceDc          0.430

```

```

## depression_z:raceDc          0.866
## purpose_z:depression_z:raceDc 0.437
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1051231)
##
## Null deviance: 76.820  on 583  degrees of freedom
## Residual deviance: 60.341  on 574  degrees of freedom
##   (11 observations deleted due to missingness)
## AIC: 353.7
##
## Number of Fisher Scoring iterations: 2
confint(modExDemD)

## Waiting for profiling to be done...

##                               2.5 %    97.5 %
## (Intercept)            0.12065352 0.20986072
## age_z                 -0.01425748 0.03871250
## genderDc              -0.06804822 0.03921231
## purpose_z              -0.05725572 0.00961251
## depression_z           0.12268327 0.20679854
## raceDc                -0.07649158 0.07650368
## purpose_z:depression_z -0.02215636 0.02614540
## purpose_z:raceDc       -0.05439522 0.12791217
## depression_z:raceDc   -0.09315911 0.11070612
## purpose_z:depression_z:raceDc -0.03332617 0.07719013

# odds ratios
exp(coef(modExDemD))

##                               (Intercept)      age_z
##                         1.1796964 1.0123026
##                         genderDc     purpose_z
##                         0.9856855 0.9764599
##                         depression_z   raceDc
##                         1.1790876 1.0000061
##                         purpose_z:depression_z purpose_z:raceDc
##                         1.0019965 1.0374424
##                         depression_z:raceDc purpose_z:depression_z:raceDc
##                         1.0088121 1.0221742

exp(confint(modExDemD))

## Waiting for profiling to be done...

##                               2.5 %    97.5 %
## (Intercept)            1.1282339 1.233506
## age_z                 0.9858437 1.039472
## genderDc              0.9342154 1.039991
## purpose_z              0.9443525 1.009659
## depression_z           1.1305263 1.229735
## raceDc                0.9263607 1.079506
## purpose_z:depression_z 0.9780873 1.026490
## purpose_z:raceDc       0.9470577 1.136453

```

```
## depression_z:raceDc          0.9110485 1.117067  
## purpose_z:depression_z:raceDc 0.9672230 1.080247
```