

* Encoding: UTF-8.

/*2012 - remove those not eligible for LBQ*/

```
DATASET ACTIVATE DataSet1.  
FILTER OFF.  
USE ALL.  
SELECT IF (NLBELIG = 1).  
EXECUTE.
```

/*2014 - remove those not eligible for LBQ*/

```
DATASET ACTIVATE DataSet2.  
FILTER OFF.  
USE ALL.  
SELECT IF (OLBELIG = 1).  
EXECUTE.
```

/*run the following for both 2012 and 2014 cohorts*/

```
COMPUTE COUNT=1.  
EXECUTE.
```

```
AGGREGATE  
/OUTFILE=* MODE=ADDVARIABLES  
/BREAK=HHID  
/COUNT_sum=SUM(COUNT).
```

```
FILTER OFF.  
USE ALL.  
SELECT IF (COUNT_sum >= 2).  
EXECUTE.
```

/* assign a value of 1 to everyone with xPN_SP*/

```
DATASET ACTIVATE DataSet1.  
FILTER OFF.  
USE ALL.  
SELECT IF (VAR00001 = 1).  
EXECUTE.
```

/*removed 4,006 participants, leaving 6,073 in the dataset*/

```
DATASET ACTIVATE DataSet1.  
COMPUTE COUNT2=1.  
EXECUTE.
```

```
AGGREGATE
/OUTFILE=* MODE=ADDVARIABLES
/BREAK=HHID
/COUNT2_sum=SUM(COUNT2).
```

1 - 25
2 - 6,004
3 - 12
4 - 32

remove the 25 participants and double check the 12 participants for the value 3.

```
FILTER OFF.
USE ALL.
SELECT IF (COUNT2_sum >= 2).
EXECUTE.
```

for households with 3 people, remove:

014427/010
208867/021
501283/011
535283/020

32 participants belong to 8 households. use xSUBHH.

/*removed 3853 participants, leaving 5,696 in the dataset*/

```
DATASET ACTIVATE DataSet2.
COMPUTE COUNT2=1.
EXECUTE.
```

```
AGGREGATE
/OUTFILE=* MODE=ADDVARIABLES
/BREAK=HHID
/COUNT2_sum=SUM(COUNT2).
```

1 - 24
2 - 5,622
3 - 6
4 - 44

remove the 24 participants and double check the 6 participants for the value 3.

```
FILTER OFF.
USE ALL.
SELECT IF (COUNT2_sum >= 2).
```

EXECUTE.

for households with 3 people, remove:

050125/021

140836/011

44 participants belong to 11 households. use xSUBHH.

/*did participant report on spa?*/

/*2012*/

DATASET ACTIVATE DataSet1.

RECODE NNSPA NPSPA (MISSING=0) (ELSE=1) INTO NNSPA_count NPSPA_count.

VARIABLE LABELS NNSPA_count 'NNSPA data yes/no' /NPSPA_count 'NPSPA data yes/no'.

EXECUTE.

AGGREGATE

/OUTFILE=* MODE=ADDVARIABLES

/BREAK=HHID

/NNSPA_count_sum=SUM(NNSPA_count)

/NPSPA_count_sum=SUM(NPSPA_count).

NNSPA:

0 - 902

1 - 1,114

2 - 4,002

3 - 16

4 - 12

NPSPA:

0 - 906

1 - 1,108

2 - 4,004

3 - 16

4 - 12

FILTER OFF.

USE ALL.

SELECT IF (NNSPA_count_sum >= 2).

EXECUTE.

FILTER OFF.

USE ALL.

SELECT IF (NPSPA_count_sum >= 2).

EXECUTE.

4016 participants/2010 couples with both PSPA and NSPA data in 2012.

/*2014*/

COMPUTE ONSPA=MEAN(OLB028B01,OLB028B3,OLB028B7,OLB028B8).

VARIABLE LABELS ONSPA 'NEGATIVE SPA 2014'.

EXECUTE.

COMPUTE OPSPA=MEAN(OLB028B02,OLB028B4,OLB028B5,OLB028B6).

VARIABLE LABELS OPSPA 'POSITIVE SPA 2014'.

EXECUTE.

DATASET ACTIVATE DataSet2.

RECODE ONSPA OPSPA (MISSING=0) (ELSE=1) INTO ONSPA_count OPSPA_count.

VARIABLE LABELS ONSPA_count 'NSPA data yes/no' /OPSPA_count 'PSPA data yes/no'.

EXECUTE.

AGGREGATE

/OUTFILE=* MODE=ADDVARIABLES

/BREAK=HHID

/ONSPA_count_sum=SUM(ONSPA_count)

/OPSPA_count_sum=SUM(OPSPA_count).

NSPA:

0 - 674

1 - 776

2 - 4,186

4 - 32

PSPA:

0 - 672

1 - 780

2 - 4,186

4 - 32

FILTER OFF.

USE ALL.

SELECT IF (ONSPA_count_sum >= 2).

EXECUTE.

FILTER OFF.

USE ALL.

SELECT IF (OPSPA_count_sum >= 2).

EXECUTE.

/*ended up with 4,210 participants with both PSPA and NSPA data in 2014*/

/*combine 2012 and 2014 cohorts*/

```
GET FILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\H12LB_R_eligible_cleaned_SPA.sav'
/KEEP = hhid pn NLBELIG NNSPA NPSPA NSUBHH.
SORT CASES BY hhid (a).
SAVE OUTFILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\H12_tobecombined.sav'.
```

```
GET FILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\H14LB_R_eligible_cleaned_SPA.sav'
/KEEP = hhid pn OLBELIG ONSPA OPSPA OSUBHH.
SORT CASES BY hhid (a).
SAVE OUTFILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\H14_tobecombined.sav'.
```

In the 2012 dataset, create a new variable LB and assign a value 0.
In the 2014 dataset, rename OLBELIG as LB.
in both datasets, remove x (N/O) in variable names.

/*extract age from tracker*/

```
GET FILE ='C:\Users\mmhuo\Box Sync\Research\Papers in progress\HRS\HRS
data\trk2018tr_r.sav'
/KEEP = hhid pn NAGE OAGE
gender race hispanic schlyrs .
SORT CASES BY hhid (a).
SAVE OUTFILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\trk_cov.sav'.
```

```
RECODE hhid pn (CONVERT) INTO hhid1 pn1.
COMPUTE hhidpn = (hhid1*1000)+pn1.
SORT CASES BY hhidpn (A).
FORMATS hhid1 pn1 hhidpn (f10.0).
```

/*Age*/

```
IF (LB = 0) AGE_a=NAGE.
EXECUTE.
IF (LB = 1) AGE_b=OAGE.
EXECUTE.
```

```
COMPUTE AGE=MAX(AGE_a, AGE_b).
FORMATS AGE (f6.0).
EXECUTE.
```

```
IF (AGE >= 50) age50=1.  
EXECUTE.
```

```
RECODE age50 (MISSING=0).  
EXECUTE.
```

among the sample of 8,226 participants, 245 age under 50 (478 participants removed due to at least one partner < 50).

```
AGGREGATE  
  /OUTFILE=* MODE=ADDVARIABLES  
  /BREAK=HHID_r  
  /age50_sum=SUM(age50).
```

```
FILTER OFF.  
USE ALL.  
SELECT IF (age50_sum = 2).  
EXECUTE.
```

```
SORT CASES BY HHID_r .  
CASESTOVARS  
  /ID=HHID_r  
  /GROUPBY=VARIABLE.
```

```
/*pre-restructure dataset*/
```

```
DATASET ACTIVATE DataSet2.  
DATASET DECLARE aggr_gender.  
AGGREGATE  
  /OUTFILE='aggr_gender'  
  /BREAK=HHID_r  
  /GENDER_sum=SUM(GENDER).
```

2 - 9 gay couples
4 - 15 lesbian couples

```
AGGREGATE  
  /OUTFILE=* MODE=ADDVARIABLES  
  /BREAK=HHID_r  
  /GENDER_sum=SUM(GENDER).
```

```
FILTER OFF.  
USE ALL.
```

```
SELECT IF (GENDER_sum = 3).  
EXECUTE.
```

removed 24 same-sex couples. the final sample includes 7,700 participants, thus 3,850 couples.

```
/*recode gender*/
```

```
RECODE GENDER (2=0) (1=1) (MISSING=SYSMIS).  
EXECUTE.
```

```
RECODE GENDER (ELSE=Copy) INTO hw.  
VARIABLE LABELS hw 'husband 1 wife 0'.  
EXECUTE.
```

```
/*extract RQ variables*/
```

```
C  
GET FILE ='C:\Users\mmhuo\Box Sync\Research\Papers in progress\HRS\HRS  
data\2014\H14LB_R.sav'  
/KEEP = hhid pn OLB004A  
OLB004B  
OLB004C  
OLB004D  
OLB004E  
OLB004F  
OLB004G.  
SORT CASES BY hhid (a).  
SAVE OUTFILE ='C:\Users\mmhuo\Box Sync\Research\Papers in  
progress\HRS\HRS_Profiles\H14_marital quality.sav'.
```

```
/*marital quality in 2012 and 2014*/  
/*reverse code all items*/
```

```
RECODE NLB005A NLB005B NLB005C OLB004A OLB004B OLB004C (1=4) (2=3) (3=2)  
(4=1)  
(MISSING=SYSMIS) INTO NLB005Ar NLB005Br NLB005Cr OLB004Ar OLB004Br  
OLB004Cr.  
VARIABLE LABELS NLB005Ar 'UNDERSTAND REVERSE' /NLB005Br 'RELY ON  
REVERSE' /NLB005Cr  
'OPEN UP REVERSE' /OLB004Ar 'UNDERSTAND REVERSE'  
/OLB004Br 'RELY ON REVERSE' /OLB004Cr 'OPEN UP REVERSE'.  
EXECUTE.
```

```

RECODE NLB005D NLB005E NLB005F NLB005G OLB004D OLB004E OLB004F
OLB004G (1=4) (2=3) (3=2) (4=1)
(MISSING=SYSMIS) INTO NLB005Dr NLB005Er NLB005Fr NLB005Gr OLB004Dr
OLB004Er OLB004Fr OLB004Gr.
VARIABLE LABELS NLB005Dr 'TOO MANY DEMANDS REVERSE' /NLB005Er
'CRITICIZE REVERSE' /NLB005Fr
'LET YOU DOWN REVERSE' /NLB005Gr 'GETS ON YOUR NERVES REVERSE'
/OLB004Dr 'TOO MANY DEMANDS REVERSE'
/OLB004Er 'CRITICIZE REVERSE' /OLB004Fr 'LET YOU DOWN REVERSE' /OLB004Gr
'GETS ON YOUR NERVES '+
'REVERSE'.
EXECUTE.

```

```

COMPUTE
RQ12=MEAN(NLB005Ar,NLB005Br,NLB005Cr,NLB005D,NLB005E,NLB005F,NLB005G).
EXECUTE.

```

```

COMPUTE RQ14=MEAN(OLB004Ar, OLB004Br, OLB004Cr, OLB004D, OLB004E,
OLB004F, OLB004G).
EXECUTE.

```

```

/*marital quality*/

```

```

IF (LB = 0) RQ_a = RQ12.
EXECUTE.
IF (LB = 1) RQ_b = RQ14.
EXECUTE.

```

```

COMPUTE RQ=MAX(RQ_a, RQ_b).
FORMATS RQ (f8.2).
EXECUTE.

```

```

/*covariates from rand*/

```

```

GET FILE ='C:\Users\mmhuo\Box Sync\Research\Papers in progress\HRS\HRS
data\randhrs1992_2018v1.sav'
/KEEP = hhidpn H11ITOT H12ITOT R11CONDE R12CONDE R11MCURLN R12MCURLN
S11MCURLN S12MCURLN.
SORT CASES BY hhid (a).
SAVE OUTFILE ='C:\Users\mmhuo\Box Sync\Research\Papers in
progress\HRS\HRS_Profiles\rand_cov.sav'.

```



```
/*income*/
```

```
IF (LB = 0) ITOT_a=H11ITOT.  
VARIABLE LABELS ITOT_a 'ITOT 12'.  
EXECUTE.  
IF (LB = 1) ITOT_b=H12ITOT.  
VARIABLE LABELS ITOT_b 'ITOT 14'.  
EXECUTE.
```

```
COMPUTE ITOT=MAX(ITOT_a,ITOT_b).  
VARIABLE LABELS ITOT 'ITOT'.  
EXECUTE.
```

```
/*log transform income*/
```

```
COMPUTE Income=LN(ITOT+1).  
EXECUTE.
```

```
/*chronic conditions*/
```

```
IF (LB = 0) CONDE_a=R11CONDE.  
VARIABLE LABELS CONDE_a 'CONDE 12'.  
EXECUTE.  
IF (LB = 1) CONDE_b=R12CONDE.  
VARIABLE LABELS CONDE_b 'CONDE 14'.  
EXECUTE.
```

```
COMPUTE CONDE=MAX(CONDE_a,CONDE_b).  
VARIABLE LABELS CONDE 'CONDE'.  
EXECUTE.
```

```
/*marital duration*/
```

```
IF (LB = 0) MDUR_a=R11MCURLN.  
VARIABLE LABELS MDUR_a 'marital duration 12'.  
EXECUTE.  
IF (LB = 1) MDUR_b=R12MCURLN.  
VARIABLE LABELS MDUR_b 'marital duration 14'.  
EXECUTE.
```

```
COMPUTE MDUR=MAX(MDUR_a,MDUR_b).  
VARIABLE LABELS MDUR 'marital duartion'.  
EXECUTE.
```

/*functional limitations*/

```
IF (LB = 0) FL_a=NG013.  
VARIABLE LABELS FL_a 'FL 12'.  
EXECUTE.  
IF (LB = 1) FL_b=OG013.  
VARIABLE LABELS FL_b 'FL 14'.  
EXECUTE.
```

```
COMPUTE FL=MAX(FL_a,FL_b).  
VARIABLE LABELS FL 'FL'.  
EXECUTE.
```

/*self-rated health*/

```
IF (LB = 0) HEALTH_a=NC001.  
VARIABLE LABELS HEALTH_a 'HEALTH 12'.  
EXECUTE.  
IF (LB = 1) HEALTH_b=OC001.  
VARIABLE LABELS HEALTH_b 'HEALTH 14'.  
EXECUTE.
```

```
COMPUTE HEALTH=MAX(HEALTH_a,HEALTH_b).  
VARIABLE LABELS HEALTH 'HEALTH'.  
EXECUTE.
```

```
RECODE HEALTH (8=SYSMIS) (MISSING=SYSMIS) (5=1) (4=2) (3=3) (2=4) (1=5) INTO  
Healthr.  
EXECUTE.
```

/*recode minority status*/

```
IF (HISPANIC = 5 and race = 1) Minority=0.  
EXECUTE.
```

```
DO IF (HISPANIC = 1 OR HISPANIC = 2 OR HISPANIC = 3 OR RACE =2 OR RACE = 7).  
RECODE Minority (MISSING=1).  
END IF.  
EXECUTE.
```

/*recode work status*/

```
IF (LB = 0) work_a=NJ020.  
VARIABLE LABELS work_a 'work 12'.  
EXECUTE.  
IF (LB = 1) work_b=OJ020.  
VARIABLE LABELS work_b 'work 14'.  
EXECUTE.
```

```
COMPUTE work=MAX(work_a,work_b).  
VARIABLE LABELS work 'work'.  
EXECUTE.
```

```
RECODE work (9=SYSMIS) (5=0) (ELSE=Copy) INTO workr.  
EXECUTE.
```

```
/*generate depressive symptom variables*/
```

```
RECODE ND113 ND115 OD113 OD115 PD113 PD115 (1=0) (0=1) (MISSING=SYSMIS)  
INTO ND113r ND115r OD113r OD115r PD113r PD115r .  
EXECUTE.
```

```
COMPUTE DEP12=SUM(ND110, ND111, ND112, ND113r, ND114, ND115r, ND116,  
ND117).  
EXECUTE.  
COMPUTE DEP14=SUM(OD110, OD111, OD112, OD113r, OD114, OD115r, OD116,  
OD117).  
EXECUTE.  
COMPUTE DEP16=SUM(PD110, PD111, PD112, PD113r, PD114, PD115r, PD116, PD117).  
EXECUTE.
```

```
IF (LB = 0) w1dep_a=dep12.  
EXECUTE.  
IF (LB = 0) w2dep_a=dep14.  
EXECUTE.
```

```
IF (LB = 1) w1dep_b=dep14.  
EXECUTE.  
IF (LB = 1) w2dep_b=dep16.  
EXECUTE.
```

```
COMPUTE w1dep=max(w1dep_a, w1dep_b).  
EXECUTE.  
COMPUTE w2dep=max(w2dep_a, w2dep_b).  
EXECUTE.
```

```
/*center continuous covariates*/
```

```
COMPUTE cage=AGE - 67.44.  
EXECUTE.  
COMPUTE cedu=SCHLYRS - 13.02.  
EXECUTE.  
COMPUTE cinc=income - 10.89.  
EXECUTE.  
COMPUTE chea=healthr - 3.22.  
EXECUTE.  
COMPUTE cfl=fl - 1.90.  
EXECUTE.  
COMPUTE ccon=conde - 2.12.  
EXECUTE.  
COMPUTE crq=rq - 3.26.  
EXECUTE.  
COMPUTE cmdu=mdur - 37.18.  
EXECUTE.
```

```
/*reorder the 5 profiles*/
```

```
RECODE Profile_5 (1=1) (4=2) (5=3) (2=4) (3=5) INTO Profile5r.  
VARIABLE LABELS Profile5r 'reordered profiles 5 new model'.  
EXECUTE.
```

```
IF (Profile5r = 1) pro5_1=1.  
VARIABLE LABELS Pro5_1 'similarly positive'.  
EXECUTE.
```

```
IF (Profile5r = 2) pro5_2=1.  
VARIABLE LABELS Pro5_2 'similarly negative'.  
EXECUTE.
```

```
IF (Profile5r = 3) pro5_3=1.  
VARIABLE LABELS Pro5_3 'similarly average'.  
EXECUTE.
```

```
IF (Profile5r = 4) pro5_4=1.  
VARIABLE LABELS Pro5_4 'husband negative'.  
EXECUTE.
```

```
IF (Profile5r = 5) pro5_5=1.  
VARIABLE LABELS Pro5_5 'wife negative'.  
EXECUTE.
```

```
RECODE pro5_1 pro5_2 pro5_3 pro5_4 pro5_5 (MISSING=0).  
EXECUTE.
```