

## Supplemental File 2

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*****  
* Supplemental File 2 – GBTM Example Do-File (Stata)  
*****
```

```
** download traj package for Stata (Jones & Nagin, 2013)  
net from http://www.andrew.cmu.edu/user/bjones/traj  
net install traj, force  
help traj
```

```
** Square Root Transform Outcome (Step 2)
```

```
* Alcohol
```

```
gen alcsq1 = sqrt(alcohol1)  
gen alcsq2 = sqrt(alcohol2)  
gen alcsq3 = sqrt(alcohol3)  
gen alcsq4 = sqrt(alcohol4)  
gen alcsq5 = sqrt(alcohol5)  
gen alcsq6 = sqrt(alcohol6)  
gen alcsq7 = sqrt(alcohol7)  
gen alcsq8 = sqrt(alcohol8)  
gen alcsq9 = sqrt(alcohol9)  
gen alcsq10 = sqrt(alcohol10)  
gen alcsq11 = sqrt(alcohol11)  
gen alcsq12 = sqrt(alcohol12)  
gen alcsq13 = sqrt(alcohol13)  
gen alcsq14 = sqrt(alcohol14)
```

```
* Marijuana
```

```
gen mjsq1 = sqrt(marijuana1)  
gen mjsq2 = sqrt(marijuana2)  
gen mjsq3 = sqrt(marijuana3)  
gen mjsq4 = sqrt(marijuana4)  
gen mjsq5 = sqrt(marijuana5)  
gen mjsq6 = sqrt(marijuana6)  
gen mjsq7 = sqrt(marijuana7)  
gen mjsq8 = sqrt(marijuana8)  
gen mjsq9 = sqrt(marijuana9)  
gen mjsq10 = sqrt(marijuana10)  
gen mjsq11 = sqrt(marijuana11)  
gen mjsq12 = sqrt(marijuana12)  
gen mjsq13 = sqrt(marijuana13)  
gen mjsq14 = sqrt(marijuana14)
```

```
**** Censored Normal Models **** (Step 3)
```

```
** Alcohol - 4 Trajectories
```

```
* All cubic
```

```
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 3 3) min(0) max(6)  
    * Plotting  
    trajplot, ci
```

```
* Mixed
```

```
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 3 2) min(0) max(6)
```

```

* Mixed
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 2 2) min(0) max(6)

** Alcohol - 5 Trajectories
* All cubic
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 3 3) min(0) max(6)
* Mixed
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 3 2) min(0) max(6)
* Mixed
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 2 2) min(0) max(6)

** Marijuana - 4 Trajectories
* All cubic
traj, model(cnorm) var(mjsq*) indep(t_*) order(3 3 3 3) min(0) max(6)
* All quadratic
traj, model(cnorm) var(mjsq*) indep(t_*) order(2 2 2 2) min(0) max(6)
* Mixed
traj, model(cnorm) var(mjsq*) indep(t_*) order(1 2 2 2) min(0) max(6)

** Marijuana - 5 Trajectories
* All cubic
traj, model(cnorm) var(mjsq*) indep(t_*) order(3 3 3 3 3) min(0) max(6)
* All quadratic
traj, model(cnorm) var(mjsq*) indep(t_*) order(2 2 2 2 2) min(0) max(6)
* Mixed
traj, model(cnorm) var(mjsq*) indep(t_*) order(1 2 2 2 2) min(0) max(6)

*** ZIP Models *** (Step 3)
** Alcohol - 4 Trajectories
* All cubic
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 3) iorder(1 1 1 1)
* Mixed
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 2) iorder(1 1 1 1)
* Mixed
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 2 2) iorder(1 1 1 1)

** Alcohol - 5 Trajectories
* All cubic
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 3 3) iorder(1 1 1 1 1)
* Mixed
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 3 2) iorder(1 1 1 1 1)
* Mixed
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 2 2) iorder(1 1 1 1 1)

** Marijuana - 4 Trajectories
* All cubic
traj, model(zip) var(marijuana*) indep(t_*) order(3 3 3 3) iorder (1 1 1 1)
* Mixed
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 3 2) iorder (1 1 1 1)
* Mixed

```

```
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 2 2) iorder (1 1 1 1)
```

```
** Marijuana - 5 Trajectories
```

```
* All quadratic
```

```
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 3 3 3) iorder (1 1 1 1 1)
```

```
* Mixed
```

```
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 2 2 3) iorder (1 1 1 1 1)
```

```
* Mixed
```

```
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 2 2 2) iorder (1 1 1 1 1)
```

```
*** Addition of Covariates to "Best" Models (Step 4) ***
```

```
** Alcohol
```

```
* Censored Normal
```

```
traj, model(cnorm) var(alcsq*) indep(t_*) order(3 3 3 3 3) min(0) max(6) risk(black hispanic other)
```

```
** ZIP
```

```
traj, model(zip) var(alcohol*) indep(t_*) order(3 3 3 3 2) iorder(1 1 1 1 1) risk(black hispanic other)
```

```
** Marijuana
```

```
* Censored Normal
```

```
traj, model(cnorm) var(mjsq*) indep(t_*) order(1 2 3 2 2) min(0) max(6) risk(black hispanic other)
```

```
** ZIP
```

```
traj, model(zip) var(marijuana*) indep(t_*) order(1 2 2 2 2) iorder(1 1 1 1 1) risk(black hispanic other)
```