1) Specific path to ENIGMA folders

```
addpath('ENIGMA/scripts/surface_viewer/')
Warning: Name is nonexistent or not a directory: ENIGMA/scripts/surface_viewer
addpath(genpath('ENIGMA/shared/'))
```

2) Load subcortical templates

```
sctx_template_L = SurfStatReadSurf('sctx_template_L')

sctx_template_L = struct with fields:
    tri: [51952×3 int32]
    coord: [3×25910 double]

sctx_template_R = SurfStatReadSurf('sctx_template_R')

sctx_template_R = struct with fields:
    tri: [50884×3 int32]
    coord: [3×25368 double]
```

3) Plot subcortical data



this is an example

2 4 6 8 10 12 14