All:
1. Identifiers can be merged to de-identified data using the variable "instanceID."
Household:
1. **"FH_HOUSEHOLD_15 Dec 2014_correct_eacodes.dta"** -- this is the "main" data set. It is a single observation per Household and includes all of the non-roster data. It also includes (variable names in parentheses): region (region), health facility ID (h_facility), community (settlement), and national eacode developed (ea_code). There are 130 unique eacodes.
2. **"FH_HOUSEHOLD-Flap_15 Dec 2014.dta"** This is the roster data, and has 1 observation for every household member (11,306). It can be merged to the main data file using the variable "instanceID" (m:1 if the main file is the using file).
3. **"FH_HOUSEHOLD-fh_08_04c_1_15 Dec 2014.dta"** This is section 8, the module on deaths. It is a long data set with one observation for every reported death (N=1,547). It can be merged to the main data file using the variable "instanceID" (m:1 if the main file is the using file).
4. **"FH_HOUSEHOLD-Module9_15 Dec 2014.dta"** This is the illness and health seeking module and has 1 observation for every HH member (N=11,324). It can be merged to the roster data 1:1 using pid instanceID: (merge 1:1 instanceID pid using "/FH_HOUSEHOLD-Module9_15 Dec 2014.dta"), but results in 18 unmatched observations -- all of these observations are empty except for pid and instanceID. As before, can also be merged to the main data file using the variable "instanceID" (m:1 if the main file is the using file).
5. **"FH_HOUSEHOLD-Module16_15 Dec 2014.dta"** This is the child anthro and vax section, and contains one observation for each HH member (N=11,306), but data is missing unless the member is <5. It can be merged to the roster data 1:1 using pid instanceID: (merge 1:1 instanceID pid using "/FH_HOUSEHOLD-Module9_15 Dec 2014.dta"). As before, can also be merged to the main data file using the variable "instanceID" (m:1 if the main file is the using file).
VDC:
This is in the Village/VDC data folder.
1. **"vdc_main.dta"** is the main data and contains f_facility and Region for 108 settlements.
2. **"VDC_COMMUNITY_vdc_08_ 2 Dec 2014.dta"** contains 207 observations which includes a single observation for each community program. It can be merged m:1 using "instanceID" if vdcmain is the using file.
VSG
This is in the Village/VSG data folder
1. **"vsgmain.dta"** is the main data and contains f_facility and Region for 108 settlements.
2. **"VSG_VILLAGE_vsg_01_13_ 2 Dec 2014.dta"** contains data on people who left and has 24 unique observations in 17 unique settlement, one for each member who left. It can be merged m:1 using "instanceID" if vsgmain is the using file.
3. "**VSG_VILLAGE_vsg_03_01_ 2 Dec 2014.dta**" contains data on activities and contains 140 observations in 84 communities.
Health Facility Data
1. **"F1_HEALTH_FACILITY_22 Dec 2014.dta"** is the main data set, and contains 24 observations. It contains region and facility name (region, h_facility).
2. **"F1_HEALTH_FACILITY_GEN1-f1_04_01_22 Dec 2014.dta"** contains all of the staff details. This includes 207 observations in the 24 facilities. It can be merged m:1 using "instanceID" if F1_HEALTH_FACILITY_GEN1-f1_04_01_22 Dec 2014.dta is the using file.
Health Worker Data
1. **"F2_HEALTH_WORKER_ 2 Dec 2014.dta"** includes all of the health worker interview data. 128 observations in 24 facilities, region and h_facility are included in the file.
U5 Exit
1. **"F4_EXIT_INTERVIEW_ 2 Dec 2014.dta"** is the main data file. It contains 160 observations at 24 facilities. It contains region and facility name (region, h_facility).
2. **"F4_EXIT_INTERVIEW_GEN1-f4_02_18_ 2 Dec 2014.dta"** is the data on any medicines perscribed, 86 observations across 43 patients. It can be merged m:1 using "instanceID" if "F4_EXIT_INTERVIEW_ 2 Dec 2014.dta" is the using file.
ANC Exit
1. **"F3_EXIT_INTERVIEW_ 2 Dec 2014.dta"** includes all of the ANC exit interview data. It contains region and facility name (region, h_facility).