

```

/*****
* Malaria early morbidity and mortality
* Preliminary analysis - generating tables and graphs - file last edited - 28 May 2015 *
* - first run do files 'la. Setup', and optionally 'lb stvary diagnostics'
*****/

capture log close
version 13.1
set linesize 100
set more off
cd "C:\Users\Carl\Google Drive\MPH\Projects\Malaria project\Data\results"
loc today = c(current_date)
log using "malariaproject_log_`today'.txt", append text

*TIES
loc ties efron

*prepare folder for results
local T = c(current_time)
local T = subinstr("`T'", ":", "_", .)
mkdir "`ties' `today' `T'"
cd "`ties' `today' `T'"
mkdir figures
cd figures
mkdir PH
cd ..

keep obsno hrn Age Sex Ethnic AGR4 AdmNext14 DiedNext14 YearCat sexPreg hrnmal AdmFU14m
DiedFU14m whiteCat tgn SpeciesX AdmFU15 DiedFU15

*timer Start

timer clear 1
timer on 1

*** use "C:\Users\Carl\Google Drive\MPH\Internships\Malaria project\Data\File
archive\MalariaEpisodes_vs 1.9.3.dta", clear /*FIRST RUN SET UP */

*Outcome / Failure variables
loc outcome Adm Died /* list of outcomes of interest to be analysed
separately */
loc Adm_f early admission /* full title of 'admission' outcome for graph
display */
loc Died_f early death /* full title of 'death' outcome for graph display */

*Exposures
loc commonExp SpeciesX ///
AGR4 ///
sexPreg ///
Ethnic ///
whiteCat ///
tgn

*Exposures
loc iExp i.SpeciesX ///

```

```

b4.AGR4          ///
i.sexPreg        ///
i.Ethnic         ///
i.whiteCat       ///
i.tgn

```

*Model specifications

```

loc modell ", cluster(hrn)"
loc model2 "i.SpeciesX b4.AGR4 i.sexPreg i.Ethnic i.whiteCat i.tgn, cluster(hrn) `ties'"

```

*save graph style in local macro

```

loc graph_style graphregion(fcolor(white) lcolor(white)) scheme(s2color)
di "`graph_style'"

```

```

foreach outc of loc outcome {

```

*Initiate outcome timer

```

timer clear 2
timer on 2

```

*Copy template result sheet for each outcome

```

copy ..\Template_resTable_v2.xlsx `outc'_resTable.xlsx

```

*Set up for export of data to Excel results worksheet

```

putexcel set "`outc'_resTable.xlsx", modify keepcellformat
putexcel A1 = ("Model 2: stcox `model2' ") /* Title for excel results sheet */ ///
C2 = (`:di proper("`outc'_f'"))')

```

*Create macro references for excel export columns

```

loc nN_Cell      = "C"
loc uHRCCell     = "D"
loc uHRpvalCell  = "E"
loc M2_HRCCell   = "G"
loc M2_HRpvalCell = "H"
loc mfp_HRCCell  = "J"
loc mfp_HRpvalCell = "K"

```

*Start cell for input values

```

loc vcell = 5
loc varcell = `vcell'

```

*Set up for survival analysis (AdmFU14m and DiedFU14m are currently specified - recoding of follow up time through 0.5 to 14.5)

```

stset `outc'FU15, fail(`outc'Next14) id(obsno)
loc axismax 15

```

* *Graph example risk set for outcome (need to fix outpoints and legend to be closed dot, not arrow)

```

loc Adm_hlstart = 72745
loc Adm_hlend = 72752

```

```

loc Died_hlstart = 72745
loc Died_hlend = 72752
loc `outc'rs = ``outc'_hlstart'+1
loc `outc're = ``outc'_hlend'-1
loc hlimit = "if hrnmal>``outc'_hlstart' & hrnmal <``outc'_hlend'"
loc mcols = "black"
loc ts = "_t0"
loc te = "_t"
loc yvar = "hrnmal"
loc textv = ``outc'rs' -.2

twoway sc `yvar' `ts' `hlimit', mc(`mcols') ms(o)
|| ///
    pcspike `yvar' `ts' `yvar' `te' `hlimit' & _d==0, mc(`mcols') lcolor(`mcols')
    || ///
    pcspike `yvar' `ts' `yvar' `te' `hlimit' & _d==1, lcolor(`mcols')
    || ///
    sc `yvar' `te' `hlimit' & _d==1, mc(red) ms(X)
    || ///
    sc `yvar' `te' `hlimit' & _d==0, mc(`mcols') ms(o)
    ///
    text(72746.1 0 "(not malaria patient)", size(small) placement(e) color(gray))
    ///
    text(72748.1 0 "(not malaria patient)", size(small) placement(e) color(gray))
    ///
    text(72751.1 0 "(not malaria patient)", size(small) placement(e) color(gray))
    ///
    name("``outc'_egRiskset``outc'rs'to``outc're",replace)
    ///
    title("HRN clusters (``outc'_f' riskset example)",size(medsmall)
    ///
    placement(west) margin(-10 0 0 -3) justification(left))
    ///
    ylabel(72746.1(1)72751.1, format(%9.0f)
    ///
    nogrid angle(horizontal) labsize(small))
    ///
    ytitle("Hospital Record Number clusters")
    ///
    xlabel(0(1)`axismax', labsize(small)) yscale(rev)
    ///
    xtitle("Time (days) from entry (_t0) until ``outc'_f' or censoring (_t)"
    ///
    , margin(medsmall)) /***/
    ///
    xline(15, lpattern(shortdash) lc(edkblue) noextend)
    ///
    text(72746 15 "End of two weeks' follow up", size(small) placement(w))
    ///
    legend(on order(1 "entry / exit (censored)" 2 "time at risk" 4 "``outc'_f'")
    ///
    colfirst notextfirst nostack cols(6) size(small) nobox
    ///
    region(fcolor(white) margin(zero) lcolor(white)) position(12) ring(1))
    ///
    `graph_style' xscale(nofextend)
graph export figures/`outc'_egRiskset``outc'rs'to``outc're'.png, as(png) replace

```

***Loop code over explanatory variables for descriptive statistics

```
foreach v of varlist `commonExp' {
```

*Export variable name

```
putexcel A`varcell' = ("`': var label `v'")
```

*Macros for key aspects (min max, n etc)

```
su `v', meanonly
loc vmax = r(max)
loc vmin = r(min)
loc vcat = (`vmax' - `vmin') + 1
if `vcat' < 6 {
```

/*alternate spacing for extra categories*/

```
    loc alt=""
}
else {
    loc alt="alt"
}
loc labname = "`': val label `v'"" /****/
qui: levelsof `v', loc(vl)
```

*Macros for Kaplan-Meier curve and other graphs

```
loc labname = "`': val label `v'"" /****/
```

```
tempvar `v'_S /* generating temporary survivor function variable by explanatory
variable to establish scaling */
```

```
sts gen ``v'_S' = s, by(`v')
```

```
tempvar `v'_F
```

```
gen ``v'_F' = 1 - ``v'_S'
```

```
su ``v'_F', meanonly
```

```
loc fmax = r(max)
```

```
loc fmin = r(min)
```

```
loc failmax = round(trunc((r(max) * 10)) / 10, .25)
```

```
loc gap = round(`failmax' / 5, .05)
```

```
loc roundmax = `failmax' - `gap'
```

```
loc mindif = r(max) - `roundmax'
```

```
loc med = ""
```

```
loc call = ""
```

```
if `mindif' > .14 {
    loc med = `roundmax' + .1
}
```

```
loc ordnum = 1
```

```
foreach j of loc vl { /* establishing labels for value categories */
```

```
    local call `call' `ordnum' "`': label `labname' `j'""
```

```
    loc ++ordnum
```

*KM survival curve (automatic y axis scaling)

```

sts graph, by(`v`) failure                                     ///

    name("`outc'_"`v'_KM", replace)                               ///

    title("Probability of failure: "`outc'_f", by `: var label `v'", size(medsmall)
    ///
    placement(west) margin(-8 0 0 -3 ) justification(left))
    ///
    xlab(0(1)`axismax', labsize(small)) xmtick(0(1)15)
    ///
    xtitle("Days since presentation with malaria", margin(medsmall))
    ///
    ylab(minmax `fmin' `fmax' 0(`gap')`roundmax' `med',
    ///
    add format(%5.3f) nogrid labsize(small) angle(horizontal))
    ///
    legend(on order(`call') colfirst notextfirst nostack cols(6) size(small)
    ///
    nobox region(fcolor(white) margin(zero) lcolor(white)) position(12) ring(1))
    ///
    `graph_style' xscale(nofextend) yscale(nofextend)
    ///

    note(" ")
graph export figures/`outc'_"`v'_KM.png, as(png) replace

loc adjvar = subinstr("`commonExp'", "`v'", " ", 1)
loc adjcall
foreach av of varlist `adjvar' {
loc adjcall `adjcall' "`:var label `av'", "

}

loc adjcall = subinstr("`adjcall'", char(34), "", . )          /***/
loc adjcall = substr("`adjcall'",1,length("`adjcall'")-2)

di "Adjusting for `adjcall'"

sts graph, by(`v`) failure adjustfor("`adjvar'")              /***/
///
    name("`outc'_"`v'_KM_adj", replace)
    ///
    title("Probability of failure: "`outc'_f", by `: var label `v'", adjusted*",
    size(medsmall) ///
    placement(west) margin(-8 0 0 -3 ) justification(left))
    ///
    xlab(0(1)`axismax', labsize(small)) xmtick(0(1)15)
    ///
    xtitle("Days since presentation with malaria event", margin(medsmall))
    ///
    ylab(minmax `fmin' `fmax' 0(`gap')`roundmax' `med',
    ///
    add format(%5.3f) nogrid labsize(small) angle(horizontal))
    ///
    legend(on order(`call') colfirst notextfirst nostack cols(6) size(small)
    ///
    nobox region(fcolor(white) margin(zero) lcolor(white)) position(12) ring(1))
    ///

```

```

`graph_style' xscale(nofextend) yscale(nofextend)
                                     ///

note(" *adjusted for: `adjcall'")
graph export figures/`outc'_'v'_KM_adj.png, as(png) replace

graph combine `outc'_'v'_KM `outc'_'v'_KM_adj, name("`outc'_'v'_KM_combo", replace)
xsize(20) ysize(10.4) `graph_style'

*Export cumulative incidence to Excel
tab `v' `outc' Next14, row matcell(`v'_'outc'_tab)
mata : st_matrix("`v'_'outc'_N", rowsum(st_matrix("`v'_'outc'_tab"))) /*sums columns
for total N*/
loc r = 1

foreach i of loc vl {
    putexcel B`varcell' = (`": label `labname' `i'') /**/
    putexcel `nN_Cell'`varcell' = (`":di %6.0fc `v'_'outc'_tab[`r',2]'/`":di %6.0fc (`v'_'
`outc'_N[`r',1])' (`":di%-5.2f ((`v'_'outc'_tab[`r',2]/(`v'_'outc'_N[`r',1]))*100)')")

    loc ++r /*increments the row number in stored matrix results */
    loc ++varcell /*increments the row number for output to Excel */
}
}

```

***Loop code over explanatory variables - Hazard ratios

```

loc varcell = `vcell'
foreach v in `iExp' {
    ***Univariable unadjusted model (results for each outcome output to excel worksheets
per variable in folder 'Results')
    stcox `v', `ties'
    matrix vHR = r(table)'
    local names: rownames vHR
    loc r = 1
    foreach n of loc names {
        loc vr `=substr("`n'",1,1)' /**/
        loc br `=substr("`n'",2,1)' /**/

        di "`vr'"
        di "`br'"

        if "`br'" == "b" {
            putexcel `uHRCcell'`varcell' = ("1.00 (reference)")
            ///
            `uHRpvalCell'`varcell' = ("-")

        }

        if "`br'" == "o" {
            putexcel `uHRCcell'`varcell' = ("(omitted)")
            ///
            `uHRpvalCell'`varcell' = ("-")

        }

        if "`br'" == "." {
            putexcel `uHRCcell'`varcell' = (`":di%3.2f vHR[`r',1]' (
`":di%3.2f vHR[`r',5]', `":di%3.2f vHR[`r',6]')")
            ///
            `uHRpvalCell'`varcell' = (`": di subinword("`": di
%4.3f vHR[`r',4]'", "0.000", "< 0.001", 1)')")

        }
    }
}

```

```

loc ++r
loc ++varcell
di "r = `r`; varcell = `varcell`"
}

}

/* end of univariable loop */

***Plot failure rate for Year with Era marker
loc `outc'_var YearCat /*define list of explanatory variables*/
foreach v of varlist ``outc'_var' {
  su `v', meanonly
  loc vmax = r(max)
  loc eramarker = ""
  set varabbrev off
  loc Adm_era_mark = 22.75
  loc Died_era_mark = 0.55

  strate `v', per(10000) graph cluster(hrn)

  name("`outc'_YearEra_strate",replace)

  title("Rate of ``outc'_f' per 10,000 patient-days, by Year & ACT Era",
    size(medsmall) placement(west) margin(-10 0 0 -3) justification(left))

  m(o) mc(black) ciopts(lc(black) ls(p2other))

  xlabel(#`vmax',value label labsize(small))

  xtitle("`": var label `v'", margin(medsmall)) /****/

  ylabel(, nogrid angle(horizontal) labsize(small)) ytitle("")

  xline(3.4, lpattern(shortdash) lc(blue) noextend)

  text("`outc'_era_mark' 5.7 "ACT usage commences in April 2006", size(small)
    justification(left))

  addplot(pcarrowi ``outc'_era_mark' 4 ``outc'_era_mark' 3.6 (3), mc(black)
    msize(medsmall) mfc(black) lc(black))

  legend(off)

  xscale(nofextend) yscale(nofextend) `graph_style'

graph export figures/`outc'_Year-Era_strate10k.png, as(png) replace
}

timer off 2
timer list 2
di "Time to process data for `outc': " r(t1)/60 "minutes"
} /* end of outcome loop */

```

```
timer off 1
timer list 1
di "Time to process complete do-file: " r(t1)/60 "minutes"
```