

```

/*****

```

```

*Malaria early morbidity and mortality

```

```

* Tables do file - last modified 5 October 2015

```

```

*

```

```

*****/

```

```

capture log close

```

```

version 13.1

```

```

set linesize 100

```

```

set more off

```

```

cd "C:\Users\Carl\Google Drive\MPH\Projects\Malaria project\Data\excel"

```

```

loc today = c(current_date)

```

```

log using "malariaproject_log_`today'.txt", append text

```

```

*prepare folder for results

```

```

local T = c(current_time)

```

```

local T = subinstr("`T'", ":", "_", .)

```

```

mkdir "table `today' `T'"

```

```

cd "table `today' `T'"

```

```

***Table 1a: Characteristics of Patients by Species(export csv for placement in maltab.xlsx)***

```

```

loc dem constant AGR4 sexPreg Ethnic tgn whiteCat HbAdmGr5 pltCat MalNut Malaria_Last63Gr
MalPres_cat Year Era

```

```

/*explanatory variables*/

```

```

tabout `dem' Species using tables\workfiles\maltable1_Species.csv, replace c(freq row)
f(1) ptotal(none) style(csv)

```

```

***Table 1b: Characteristics of Patients by early outcomestatus (14 days or less) following a
malaria episode (export csv for placement in maltab.xlsx)***

```

```

loc group AdmNext14 DiedNext14 /*group variables*/

```

```

loc dem constant Species AGR4 sexPreg Ethnic tgn whiteCat HbAdmGr5 pltCat MalNut
Malaria_Last63Gr MalPres_cat Year Era

```

```

/*explanatory variables*/

```

```

foreach var of varlist `group' {

```

```

tabout `dem' `var' using tables\workfiles\maltable1_`var'.csv, replace c(freq row) f(1)
ptotal(none) style(csv) npos(lab)

```

```

}

```

```

foreach var of varlist `dem' {

```

```

tab `var' AdmNext14, row chi

```

```

}

```

```

***Table 2Adm: Characteristics of Patients with early re-admission (14 days or less) following a
malaria episode (export csv for placement in maltab.xlsx)***

```

```

*AdmNext14==0 (not re-admitted within 14 days)*

```

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin

```

```

tokenize `cont'

```

```

local counter = 0

```

```

local filemethod = "replace"

```

```

local heading = "h1(nil) h2(nil) h3(|Count | Mean | SD | Median |Min | Max )"

```

```

foreach v of varlist `cont' {

```

```

if `counter' > 0 {

```

```

local filemethod = "append"

```

```

local heading = "h1(nil) h2(nil) h3(nil)"

```

```

}

```

```

label define dummy 1 "`1'", modify
label val dummy dummy
tabout dummy if AdmNext14==0 using tables\workfiles\table2adm0.csv, ///
`filemethod' c(count `v' mean `v' sd `v' median `v' min `v' max `v' ) ///
f(0 lc lc lc lc lc) sum `heading' ///
lines(none) ptotal(none) style(csv)
mac shift
local counter = `counter' + 1
}

```

### \*AdmNext14==1 (re-admitted within 14 days)\*

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin
tokenize `cont'
local counter = 0
local filemethod = "replace"
local heading = "h1(nil) h2(nil) h3(|Count | Mean | SD | Median |Min | Max )"
foreach v of varlist `cont' {
if `counter' > 0 {
local filemethod = "append"
local heading = "h1(nil) h2(nil) h3(nil)"
}
label define dummy 1 "`1'", modify
label val dummy dummy
tabout dummy if AdmNext14==1 using tables\workfiles\table2adm1.csv, ///
`filemethod' c(count `v' mean `v' sd `v' median `v' min `v' max `v' ) ///
f(0 lc lc lc lc lc) sum `heading' ///
lines(none) ptotal(none) style(csv)
mac shift
local counter = `counter' + 1
}

```

### \*ttest vars by AdmNext14\*

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin
loc groupv AdmNext14
foreach v of varlist `cont' {
qui ttest `v', by(`groupv')
di as text "`v' Mean difference by `groupv':"
di %9.1f r(mu_1)-r(mu_2) as text " (95% CI" %9.1f (r(mu_1)-r(mu_2)-(1.96*r(se))) as
text"," as result %9.1f (r(mu_1)-r(mu_2)+(1.96*r(se))) as text "; {it:P-value}:" %9.3f r(p)
as text")"
}

```

### \*Mann-Whitney U Test

```

loc cont PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin log10pltmin
wbcmin log2wbcmin log10wbcmin
loc groupv AdmNext14
foreach v of varlist `cont' {
ranksum `v', by(`groupv')
}

```

\*\*\*Ask Julie for help with this - which to use with?

\*\*\*Table 1 Died: Characteristics of Patients by early death status (14 days or less) following a malaria episode (export csv for placement in maltab.xlsxs)\*\*\*

```

loc dem constant Species AGR4 Sex pregWom Ethnic MalNut Malaria_Last63Gr HbAdmGr5
sevThrom q5_wbcmin q5_pltmin Source Year Era TreatGr_First q5_PQmgkg_Sum

```

```

about `dem' DiedNext14 using tables\workfiles\maltable1_Died.csv, replace c(freq row)
f(1) ptotal(none) style(csv) npos(lab)

```

```

foreach var of varlist `dem' {
  tab `var' DiedNext14, row chi
}

```

### \*\*\*Table 2 Died: Characteristics of Patients with early death (14 days or less) following a malaria episode (export csv for placement in maltab.xlsxs)\*\*\*

#### \*DiedNext14==0 (did not die within 14 days)\*

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin
tokenize `cont'
local counter = 0
local filemethod = "replace"
local heading = "h1(nil) h2(nil) h3(|Count | Mean | SD | Median |Min | Max )"
foreach v of varlist `cont' {
  if `counter' > 0 {
    local filemethod = "append"
    local heading = "h1(nil) h2(nil) h3(nil)"
  }
  label define dummy 1 "`1'", modify
  label val dummy dummy
  tabout dummy if DiedNext14==0 using tables\workfiles\table2d0.csv, ///
  `filemethod' c(count `v' mean `v' sd `v' median `v' min `v' max `v' ) ///
  f(0 lc lc lc lc lc) sum `heading' ///
  lines(none) ptotal(none) style(csv)
  mac shift
  local counter = `counter' + 1
}

```

#### \*DiedNext14==1 (died within 14 days)\*

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin
tokenize `cont'
local counter = 0
local filemethod = "replace"
local heading = "h1(nil) h2(nil) h3(|Count | Mean | SD | Median |Min | Max )"
foreach v of varlist `cont' {
  if `counter' > 0 {
    local filemethod = "append"
    local heading = "h1(nil) h2(nil) h3(nil)"
  }
  label define dummy 1 "`1'", modify
  label val dummy dummy
  tabout dummy if DiedNext14==1 using tables\workfiles\table2d1.csv, ///
  `filemethod' c(count `v' mean `v' sd `v' median `v' min `v' max `v' ) ///
  f(0 lc lc lc lc lc) sum `heading' ///
  lines(none) ptotal(none) style(csv)
  mac shift
  local counter = `counter' + 1
}

```

#### \*ttest vars by DiedNext14\*

```

loc cont Age hbmin_Min PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin
log10pltmin wbcmin log2wbcmin log10wbcmin
loc groupv DiedNext14

```

```
foreach v of varlist `cont' {
  qui ttest `v', by(`groupv')
  di as text "`v' Mean difference by `groupv':"
  di %9.1f r(mu_1)-r(mu_2) as text " (95% CI" %9.1f (r(mu_1)-r(mu_2)-(1.96*r(se))) as
  text"," as result %9.1f (r(mu_1)-r(mu_2)+(1.96*r(se))) as text "; {it:P-value}:" %9.3f r(p)
  as text")"
}
```

### \*Mann-Whitney U Test

```
loc cont PQmgkg_Sum log2PQmgkg_Sum log10PQmgkg_Sum pltmin log2pltmin log10pltmin
wbcmin log2wbcmin log10wbcmin
loc groupv AdmNext14
foreach v of varlist `cont' {
  ranksum `v', by(`groupv')
}
```

### \*\*\*Graph histogram density plots for continuous variables (or is kernel density better? pretty similar)

```
/*(hbmin_Min is normally distributed; PQmgkg_Sum has an odd distribution suggestive of
two distinct treatment groupings - this is especially noticeable if restricted to
AGR4==4, and has some relationship with white blood cell count (eg. second peak is more
prominent in wbcmin>8000))*/
loc cont Age Age_log10 Age_log2 predwt hbmin_Min pltmin log10pltmin log2pltmin wbcmin
log10wbcmin log2wbcmin
foreach var of varlist `cont' {
  histogram `var', normal dens name("`var'_histo_density",replace) scheme(tufte)
}
```

```
/*****
```

```
*
```

### \*EXTRA GRAPHS

```
*
```

```
*****/
```

### \*Graph of initial species by year - note both absolute increase in Pv and mixed diagnoses across study period, and relative to Pf

```
use "C:\data\malaria\MalEps_v1.9.3_r9aug2015.dta", clear
```

```
cd\data\malaria
```

```
tempfile original
```

```
save `original'
```

### /\*Generate indicators of variables for plotting frequencies \*/

```
g pres_pf = SpeciesX if SpeciesX==1
```

```
g pres_pv = SpeciesX if SpeciesX==2
```

```
g pres_all = constant
```

```
g adm_pf = SpeciesX if AdmNext14==1 & SpeciesX==1
```

```
g adm_pv = SpeciesX if AdmNext14==1 & SpeciesX==2
```

```
g adm_all = constant
```

```
g died_pf = SpeciesX if DiedNext14==1 & SpeciesX==1
```

```
g died_pv = SpeciesX if DiedNext14==1 & SpeciesX==2
```

```
g died_all = constant
```

### /\*Collapse for frequencies of each variable \*/

```
collapse (count) pres_pf ///
```

```

pres_pv      ///
pres_all     ///
adm_pf       ///
adm_pv       ///
adm_all      ///
died_pf      ///
died_pv      ///
died_all     ///
, by(YearCat)

/*Colours from colorbrewer2.org */
loc farb1 = "27 158 119"
loc farb2 = "217 95 2"
loc farb3 = "102 194 165"
loc farb4 = "252 141 98"
loc farb5 = "179 226 205"
loc farb6 = "253 205 172"

tw line pres_pf Year, sort lc("`farb5'") || ///
line pres_pv Year, lc("`farb6'") || ///
line adm_pf Year, sort lc("`farb3'") lpattern(dash) || ///
line adm_pv Year, lc("`farb4'") lpattern(dash) || ///
line died_pf Year, sort lc("`farb1'") lpattern(.) || ///
line died_pv Year, lc("`farb2'") lpattern(.) ///
legend(order(1 "{it:P.falciparum}" 2 "{it: P.vivax}" {bf: malaria episodes}" ///
3 "{it:P.falciparum}" 4 "{it: P.vivax}" {bf: early admissions}" ///
5 "{it:P.falciparum}" 6 "{it: P.vivax}" {bf: early deaths}" ) ///
col(2) pos(6) ) ///
xlab(2004(1)2013, labsize(small)) xsca(nofextend) ///
xtitle("Year", margin(medsmall)) ///
ytitle(, margin(medsmall)) ylab(,angle(h) nogrid) ///
graphr(color(white) lcolor(white)) plotr(color(white) ) ///
title("Yearly frequency of malaria episodes, early admissions and early deaths", ///
size(medsmall) placement(west) margin(-8 0 0 -3 ) justification(left)) xsize(12)
ysize(10)

```