

Appendix II: Implementation in Stata

```

use melanoma if year8594 == 1
stset surv_mm, failure(status=1,2) scale(12) exit(time 120) id(id)

// Pohar Perme estimate
stpp R_pp using popmort.dta, agediag(age) datediag(dx) pmother(sex)

// conditional model (no covariates)
gen _age = floor(min(age + _t,99))
gen _year = floor(year(dx + _t*365.24))

merge m:1 _age _year sex using popmort, keep(match master)
stpm2, scale(hazard) df (5) bhazard(rate)
range tt_cond 0 10 101
predict s_cond, surv timevar(tt) ci

preserve
rename _t PP_t
keep tt_cond s_cond* R_pp* PP_t
save cond, replace
restore

// marginal model
// setup data
mrsprep using popmort.dta, pmother(sex) agediag(age) datediag(dx) ///
    pmmaxyear(2000) /// 
    verbose breaks(0(.2)10)

// fit model
stset tstop [iw=wt], enter(tstart) failure(event==1)
stpm2, scale(hazard) df (5) bhazard(meanhazard_wt) vce(cluster id)
range tt_mrs 0 10 101
predict s_mrs, surv timevar(tt_mrs) ci

// merge in PP and conditional model
merge 1:1 _n using cond, nogenerate

// Plot results
twoway (rarea R_pp_lci R_pp_uci PP_t, sort connect(stairstep) color(${C1ci}))    ///
    (line R_pp PP_t, sort connect(stairstep) color(${C1}))    ///
    (line s_mrs* tt_mrs, color(${C2} ${C2} ${C2}) lpattern(solid dash dash))    ///
    (line s_cond* tt_cond, color(${C3} ${C3} ${C3}) lpattern(solid dash dash))    ///
    , legend(order(2 "Pohar Perme" 3 "Marginal model" 6 "Conditional model (no covariates)")    ///
    ring(0) cols(1) pos(7))    ///
    ylabel(0.6(0.1)1, format(%3.1f))    ///
    title("(a)")    ///
    ytitle("Marginal relative survival")    ///
    xtitle("Years from diagnosis")    ///
    name(int_stand, replace)

///////////////////////////////
// External age standardization
/////////////////////////////
frame change default
drop _age _year rate prob

// change age groups to those defined in ICSS
drop agegrp
egen agegrp=cut(age), at(0 45 55 65 75 200) icodes
replace agegrp = agegrp + 1
label variable agegrp "Age group"
label define agegrplab 1 "0-44" 2 "45-54" 3 "55-64" 4 "65-74" 5 "75+", replace
label values agegrp agegrplab

recode agegrp (1=0.28) (2=0.17) (3=0.21) (4=0.20) (5=0.14), gen(ICSSwt)

```

```

// Pohar Perme
stpp R_pp2 using popmort.dta, agediag(age) datediag(dx) pmother(sex) ///
standstrata(agegrp) /// standweights(0.28 0.17 0.21 0.20 0.14)

preserve
rename _t PP_t
keep PP_t R_pp2*
save stand, replace
restore

// Proportion within each age group
local total= _N
bysort agegrp: gen a_age = _N/`total'
gen double wt_age = ICSSwt/a_age

// Prepare data for marginal model
mrsprep using popmort.dta, pmother(sex) agediag(age) datediag(dx) ///
pmmaxyear(2000) ///
verbose breaks(0(.2)10) ///
indweights(wt_age) ///
newframe(mrs_stand, replace)

// Fit model
stset tstop [iw=wt], enter(tstart) failure(event==1)
stpm2, scale(hazard) df(5) bhazard(meanhazard_wt) vce(cluster id)
range tt_stand 0 10 101
predict s_extstand, surv timevar(tt_stand) ci

// merge in PP
merge 1:1 _n using stand.dta

// Plot results
twoway (rarea R_pp2_lci R_pp2_uci PP_t, connect(stairstep) color(${C1ci})) ///
(line R_pp2 PP_t, sort connect(stairstep) color(${C1ci})) ///
(line s_extstand* tt_stand, sort color(${C2} ${C2} ${C2}) lpattern(solid dash dash)) ///
, legend(order(2 "Pohar Perme" 3 "Marginal Model") ///
ring(0) cols(1) pos(7)) ///
ylabel(0(.6(0.1)1, angle(h) format(%3.1f)) ///
xtitle("Years from diagnosis") ///
ytitle("Marginal Relative Survival") ///
title("(b)") ///
name(ext_stand, replace)

graph combine int_stand ext_stand, nocopies

```