

(Continued)

<code>freq(x=variable, w=w, plot=T/F ...); freqC (x=variable, w=w)</code>
<code>imeansC(function1=~y, function2=~x + z, data=design.dataset)</code>
<code>lineType()</code>
<code>logregR2(model=logit.model ...)</code>
<code>orci(model= logit.model ...)</code>
<code>pchisqC(reduced=reduced.logit.model, full=full.logit.model ...)</code>
<code>plotChar()</code>
<code>plotmeans(formula=y ~ x, data=dataset ...)</code>
<code>plotmeansC(data=dataset, formula2=~y, formula3=~x, formula4=y~x, w=~w ...)</code>
<code>printC(objx=table.output)</code>
<code>prop.testC(y=y, x=x, w=w)</code>
<code>scatterplot(formula=y~x, data=dataset ...)</code>
<code>somersD(formula~x+y=, data=design.dataset)</code>
<code>sortC(data=dataset, id=identifier/name, by=sort.criteria, descending=T/F)</code>
<code>spss.get("SPSS.dataset.sav") [import SPSS dataset]</code>
<code>stata.get("Stata.dataset.dta") [import Stata dataset]</code>
<code>svyboxplot(formula=y~x, design=design.dataset ...)</code>
<code>svyby (formula=~y, by=~x, design=design.dataset, FUN=function.applied ...)</code>
<code>svychisq (formula=y~x, design=design.dataset ...)</code>
<code>svychisqC (formula=y~x, design=design.dataset)</code>
<code>svydesign(id=~1, data=data, weights=~w ...) [create design.dataset]</code>
<code>svyglm(formula=binary.y ~ x l... xn, design=design.dataset, family=quasibinomial)</code>
<code>svyglm(formula=y ~ xl ... xn, design=design.dataset ...)</code>
<code>svytable(formula=y~x, design=design.dataset)</code>
<code>welcome()</code>
<code>wtd.boxplot(formula=y ~ x, weights=w ...)</code>
<code>wtd.chi.sq(var1=x, var2=y, weight=w ...)</code>
<code>wtd.cor(x=variable.matrix, weight=w ...)</code>
<code>wtd.hist(x=variable, weight=w ...)</code>
<code>wtd.mean(x=variable, weights=w ...)</code>
<code>wtd.median(x=variable, weights=w)</code>
<code>wtd.mode(x=variable, weights=w)</code>