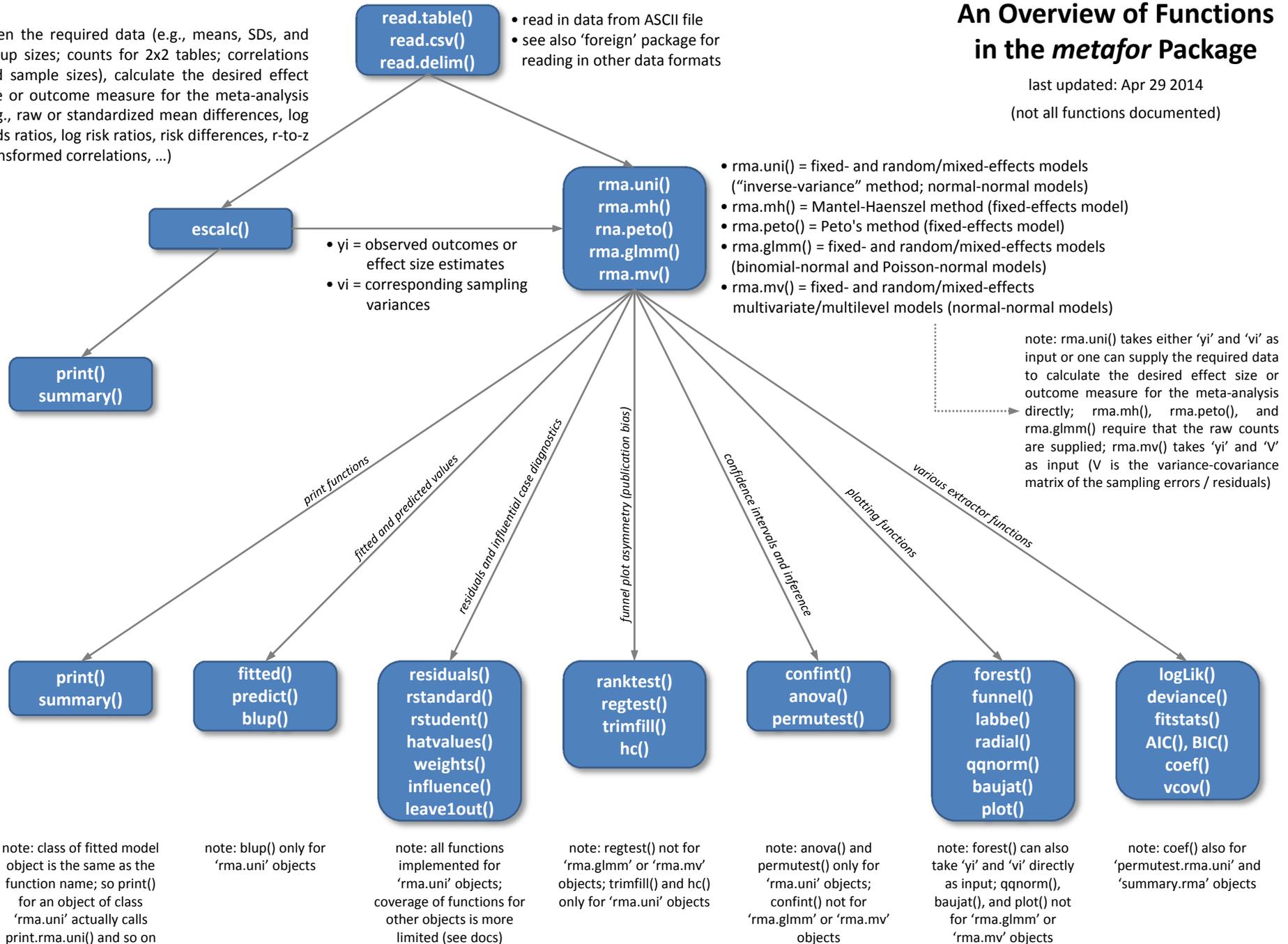


An Overview of Functions in the *metafor* Package

last updated: Apr 29 2014
(not all functions documented)

given the required data (e.g., means, SDs, and group sizes; counts for 2x2 tables; correlations and sample sizes), calculate the desired effect size or outcome measure for the meta-analysis (e.g., raw or standardized mean differences, log odds ratios, log risk ratios, risk differences, r-to-z transformed correlations, ...)



note: class of fitted model object is the same as the function name; so print() for an object of class 'rma.uni' actually calls print.rma.uni() and so on

note: blup() only for 'rma.uni' objects

note: all functions implemented for 'rma.uni' objects; coverage of functions for other objects is more limited (see docs)

note: regtest() not for 'rma.glmm' or 'rma.mv' objects; trimfill() and hc() only for 'rma.uni' objects

note: anova() and permutest() only for 'rma.uni' objects; confint() not for 'rma.glmm' or 'rma.mv' objects

note: forest() can also take 'yi' and 'vi' directly as input; qqnorm(), baujat(), and plot() not for 'rma.glmm' or 'rma.mv' objects

note: coef() also for 'permutest.rma.uni' and 'summary.rma' objects

note: restrictions may change with future updates