model2rjfun.R

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```
model2rjfun <- function(modelformula, pvec, data = NULL, jacobian = TRUE,
                           testresult = TRUE) {
    cat("model2rjfun: modelformula = ")
    print(modelformula)
    print(class(modelformula))
    stopifnot(inherits(modelformula, "formula"))
    if (length(modelformula) == 2) {
        residexpr <- modelformula[[2]]</pre>
    } else if (length(modelformula) == 3) {
        residexpr <- call("-", modelformula[[3]], modelformula[[2]])</pre>
    } else stop("Unrecognized formula")
    if (is.null(names(pvec)))
    names(pvec) <- paste0("p", seq_along(pvec))</pre>
    if (jacobian)
        residexpr <- deriv(residexpr, names(pvec))</pre>
## SHOULD TRY:
## residexpr <- fnDeriv(residexpr, names(pvec))</pre>
    if (is.null(data))
    data <- environment(modelformula)</pre>
    else if (is.list(data))
    data <- list2env(data, parent = environment(modelformula))</pre>
    else if (!is.environment(data))
    stop("'data' must be a dataframe, list, or environment")
## ??140730JN: Why do we have no data=something here? We cannot change data
   for the function.
    rjfun <- function(prm) {</pre>
        if (is.null(names(prm)))
        names(prm) <- names(pvec)</pre>
    localdata <- list2env(as.list(prm), parent = data)</pre>
    eval(residexpr, envir = localdata)
        # Saves Jacobian matrix as "gradient" attribute (consistent with deriv())
    }
    if (testresult) {
    resids <- rjfun(pvec)
    if (any(bad <- !is.finite(resids)))</pre>
        stop("residuals contain ", unique(resids[bad]))
    if (jacobian && any(bad <- !is.finite(attr(resids, "gradient"))))
        stop("Jacobian contains ", unique(attr(resids, "gradient")[bad]))
    rm(resids, bad) # Don't want to capture these in the environment of rjfun
```

```
rjfun
model2ssgrfun <- function(modelformula, pvec, data = NULL, gradient = TRUE,</pre>
                             testresult = TRUE) {
    rjfun <- model2rjfun(modelformula, pvec, data = data, jacobian = gradient,</pre>
                            testresult = testresult)
    function(prm) {
    resids <- rjfun(prm)</pre>
    ss <- as.numeric(crossprod(resids))</pre>
    if (gradient) {
         jacval <- attr(resids, "gradient")</pre>
         grval <- 2*as.numeric(crossprod(jacval, resids))</pre>
         attr(ss, "gradient") <- grval</pre>
    }
    SS
    }
}
modelexpr <- function(fun) {</pre>
    env <- environment(fun)</pre>
    if (exists("rjfun", env))
    env <- environment(env$rjfun)</pre>
    env$residexpr
}
```