Some tools in bioimagetools

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readTIFF() and writeTIFF() in the tiff package do not handle 3d stacks the right way. bioimagetools uses readTIF().

- > library(bioimagetools)
- > download.file("http://ex.volkerschmid.de/cell.tif",destfile="cell3d.tif",quiet=TRUE)
- > img <- readTIF("cell3d.tif")</pre>
- > file.remove("cell3d.tif")
- [1] TRUE
- > print(dim(img))
- [1] 512 512 3 52