[11pt] article [] graphicx [] color

 $fgcolorrgb0.345, 0.345, 0.345 \\ [1][rgb]0.686, 0.059, 0.5691 \\ [1][rgb]0.192, 0.494, 0.81 \\ [1][rgb]0.678, 0.584, 0.6861 \\ [1][rgb]0, 0, 01[10] \\ [10][rgb]0.192, 0.494, 0.81 \\ [10][rgb]0.678, 0.584, 0.6861 \\ [10][rgb]0.0, 0.1[10][rgb]0.0, 0.1$

@end@of@kframe

shade colorrgb.97, .97, .97 message colorrgb0, 0, 0 warning colorrgb1, 0, 1 error colorrgb1, 0, 0 knitrout alltt

hyperref [hmargin=2cm, vmargin=3cm]geometry

 $The \ LumiR \ User \ Guide \ Renan \ Sauteraudrs autera@fhcrc.org \ upquote.styupquote$

document

Introduction Technology Luminex xMAPhttp://www.luminexcorp.com/TechnologiesScience/xMAPTechnology/technology is a multiplex assay using flow cytometry to concurrently measure up to 500 analytes in a single reaction volume.

LumiR LumiR is an R package that provides data structure and functions to read, store and analyze Luminex xMAP experiment.

As with any R package, it should first be loaded in the session knitrout shadecolorrgb0.969, 0.969, 0.969fgcolorkframe alltt library(LumiR)

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