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| Download | Instructions |
| Command line tool *(hyper link for download)* | Software that you can download to run the pipline on your linux system.  *Example: $Rscript geneExpFile doseResFile –p 1*  *(Detail see ReadMe.txt in the download file)* |
| R package *(hyper link to git hub repository maybe)* | R packge that provide the function that you can work in R console.  Library(DIGRE\_drugSyn)  Res <- DIGRE.pred(geneFile = geneFile, doseFile = doseFile, pathway = 1) |