README.md

QLC = Quick Look CAMS/IFS results

Caution: under development

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Purpose

- QLC provides a standardized quick look to CAMS / IFS data.
- It combines data retrieval, post processing, plotting and compilation into tex pdf presentation.
- · Some flexibility in complexity is given by adopting the configuration file to user specific needs.
- Don't expect too much, as we follow the KISS principle "Keep it simple, stupid!" ;-) ...

Quick start

• execute the following (on ATOS/ECMWF):

ln -s ~cysm/bin/qlc \$HOME/bin/qlc
cp -rp ~cysm/.qlc \$HOME/.qlc

Quick usage ¶

- qlc takes the following command line input: qlc exp1 exp2 startDate endDate
- start qlc with, e.g.:

```
qlc i5la i59h 2018-12-01 2018-12-07
```

Usage (after installing qlc)

• start qlc by writing all command line output into a user defined log file, type e.g.:

```
$HOME/qlc/start.sh i89e i7rs 2018-12-16 2018-12-22 log001
```

• submit qlc with sbatch, type e.g.:

```
$HOME/qlc/start_batch.sh i89e i7rs 2018-12-16 2018-12-22
```

• make an alias or shortcut for sbatch, type e.g.:

```
ln -s $HOME/qlc/start_batch.sh $HOME/bin/sqlc
```

• export bin PATH and/or source your .profile

```
export PATH="$HOME/bin:$PATH"
```

· source your .profile

```
source ~/.profile
```

submit a qlc batch job with sqlc from anywhere, type e.g.:

```
sqlc i89e i87n 2018-12-16 2018-12-22
```

• submit sqlc batch job with dependency for plotting, type e.g.:

```
sqlc i89e i7rs 2018-12-16 2018-12-22 all
```

• view your outcome (single pdf) of this example using, e.g. xdg-open:

```
qlclastpdf="`ls -t $HOME/qlc/Presentations/CAMS2_35_i89e-i7rs_20181216-20181222_qlc_Z1-png_*.pdf | head -n 1`"
```

/usr/bin/xdg-open \$qlclastpdf

How to get qlc

• get your qlc (beta) version from the developer (until available on GitHub)

Installation

- · change to your installation directory
- unzip latest version (currently qlc_v01.zip)
- link "qlc_version" to qlc in your home directory
- change to the qlc directory, copy and adopt the example configure file and save it as qlc.conf
- make sure all paths point to the right directories, especially scratch and working directory
- caution: qlc may require temporarily huge amount of space (depending in the data to be processed)
- · caution: all paths defined in qlc.conf will be created automatically, in case they do not exist
- you may use symbolic links for "scratch" and "working directory" to match the example configure file
- hint: you may simply use the settings of the example configure file and execute the following commands:

```
cd $PERM
unzip glc v01.zip
ln -s $PERM/qlc v01
                                 $HOME/alc
                                 $HOME/.alc
mkdir
cp -p $HOME/qlc/qlc example.conf $HOME/.qlc/qlc.conf
cp -p $HOME/glc/README.md
                                 $HOME/.glc/
cp -p $HOME/qlc/LICENSE
                                 $HOME/.qlc/
cp -p $HOME/qlc/nml/*
                                 $HOME/.glc/
mkdir -p
                                 $HOME/bin
ln -s ~cysm/bin/qlc src
                                 $HOME/bin/qlc src
ln -s ~cysm/bin/qlc
                                 $HOME/bin/glc
ln -s $HOME/qlc/start_batch.sh
                                 $HOME/bin/sqlc
export PATH="$HOME/bin:$PATH"
source ~/.profile
```

configure

• a user specific configuration file is used throughout all qlc scripts:

```
CONFIG_DIR="$HOME/.qlc"
CONFIG FILE="$CONFIG DIR/qlc.conf"
```

• the following entries are required:

```
# ~/.qlc/qlc.conf - config file for QLC
SCRIPTS_PATH="$HOME/../cysm/bin/qlc_src"
QLC_DIRECTORY="$HOME/qlc"
WORKING_DIRECTORY="$SCRATCH/qlc"
MARS_RETRIEVAL_DIRECTORY="$SCRATCH/qlc/Results"
ANALYSIS_DIRECTORY="$PERM/qlc/Analysis"
PLOTS_DIRECTORY="$PERM/qlc/Plots"
```

```
TEX DIRECTORY="$PERM/glc/Presentations"
TEAM PREFIX="CAMS2 35"
EVALUATION PREFIX="EVAL"
MODEL RESOLUTION="T511L137"
TIME RESOLUTION="03hh"
#PLOTEXTENSION="pdf"
PLOTEXTENSION="png"
#SUBSCRIPT NAMES=("A1" "B1" "B2" "C1" "D1" "E1" "F1" "G1" "Z1")
SUBSCRIPT_NAMES=("A1" "B1" "B2" "C1" "71")
#MARS RETRIEVALS=("A" "B" "C" "D" "E" "F" "G")
#MARS_RETRIEVALS=("A" "C" "F")
MARS_RETRIEVALS=("F" "G")
param A=("207.210" "72.210" "73.210" "74.210")
ncvar_A=("var207" "var72"
                            "var73"
                                      "var74")
                  "PM1"
                            "PM2.5"
                                      "PM10")
myvar_A=("AOD"
param_B=("19.217" "6.217"
                            "122.210" "203.210")
#ncvar B=("var19"
                  "var6"
                             "var122" "var203")
ncvar_B=("nh3"
                  "hno3"
                            "so2"
                                      "qo3")
                            "S02"
myvar_B=("NH3"
                  "HN03"
                                      "03")
param_C=("210247" "210248"
                            "210249"
                                      "210011")
ncvar_C=("aermr16" "aermr17" "aermr18" "aermr11")
                  "N03b"
                            "NH4"
                                      "S04")
myvar_C=("N03a"
param_D=("51.212" "52.212")
ncvar D=("var51"
                  "var52")
myvar_D=("pHcloud" "pHrain")
param_E=("210025")
ncvar_E=("aergn10")
myvar_E=("EQdiag")
param_F=("22.210")
ncvar F=("aergn07")
```

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```
myvar_F=("pHaer")
param_G=("22.210")
ncvar_G=("aergn07")
myvar_G=("AW")
```

- SCRIPTS_PATH <- you may keep this (cysm) as script source directory
- QLC_DIRECTORY <- keep default, or change / link path to your qlc installation directory
- WORKING_DIRECTORY <- keep default, or change / link path to your working directory
- MARS_RETRIEVAL_DIRECTORY <- keep default, or change / link path to your mars retrieval directory
- ANALYSIS_DIRECTORY <- keep default, or change / link path to your post processing directory
- PLOTS_DIRECTORY <- keep default, or change / link path to your plots output directory
- TEX DIRECTORY <- keep default, or change / link path to your tex plots directory
- TEAM_PREFIX <- keep default, or change to your evaluation identifier (tex prefix)
- EVALUATION_PREFIX <- keep default, or change to your evaluation identifier (tex prefix)
- MODEL_RESOLUTION <- keep default, or define your model resolution
- TIME_RESOLUTION <- keep default, or define your IFS data output time step resolution
- PLOTEXTENSION <- keep default (png), or use pdf (for pyFerret 10 times larger file size)
- MARS_RETRIEVALS=("A" "B" "C") <- define / add your additional MARS retrieval namelists
 - MARS namelists are located in \$H0ME/.qlc and named e.g.: mars_A.nml, mars_B.nml, mars_C.nml, etc
 - where A,B,C, etc may denote different retrieval types (e.g., A=AOD(surface), B=NH3(3D), C=NH4, etc)
 - add additional namelists accordingly and/or adopt to your needs, except for:
 - EXP, SDATE, EDATE, MYPATH, MYFILE, which will be auto-replaced depending on your command line input
 - add your param_, ncvar_ and myvar_ to your \$HOME/qlc.conf file
 - variable renaming will be done automatically by subscript glc_C1.sh
 - make sure your MARS entries fit your exp type, e.g. check settings and redefine fixed entries in mars_?.nml:

```
CLASS = rd,
TYPE = fc,
STREAM = oper,
EXPVER = EXP.
```

• SUBSCRIPT_NAMES=("A1" "B1" "C1" "Z1") <- keep default, or choose the subscripts to be considered

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- subscripts are currently to be provided by the developer (request any extensions)
- subscripts are located in \$HOME/bin/qlc_src and named e.g.: qlc_A1.sh
- where the qlc_extensions denote:
 - -- A1,2,... = mars retrieval scripts (user specific, except placeholders)
 - -- B1,2,... = post processing scripts (grb2nc, chname, timavg, ...)
 - -- C1,2,... = plot scripts using ferret (https://ferret.pmel.noaa.gov/Ferret/))
 - -- D1,2,... = plot scripts using python (own developments, and/or user provided)
 - -- E1,2,... = plot scripts using evaltools (https://opensource.umr-cnrm.fr/projects/evaltools (<a href="https://opensource.umr-cnrm.fr/project
 - -- F1,2,... = plot scripts using ver0D (ECMWF, Luke Jones)
 - -- G1,2,... = plot scripts using Metview (ECMWF, https://metview.readthedocs.io/en/latest)
 - -- Z1,2,... = tex scripts to assemble standardized plots in a single pdf presentation (with \$USER name)

