

How it works in principle

The idea is to reduce risks when changing AAPS settings. I first test the impact of such changes offline on a PC by emulating in a python script what the loop would do. That python is a translation of the original AAPS „determine-basal“ java version.

The code scans an existing AAPS logfile and extracts the results of the original run. Also the inputs for "determine_basal" like last glucose, IOB array or profile definition are extracted. Here profile is not the usual daily pattern of basals etc. but a collection of settings like which SMB feature is ON or OFF; for details go to the „SMB“ tab in AAPS and scan the section named „Profile“.

After collecting all these inputs the emulator will also read your alternative AAPS settings like

- different temp target
- different SMB settings
- different basal safety factors
- different ISF (here named „sens“ !) or IC values

Settings like those related to meals already eaten or insulin already dosed are handled in other modules of AAPS and are therefore out of reach here. For the same reason changes to Autosense cannot be handled and the only meaningful change you can do is set it to „1“ which means switch it off.

Then determine_basal.py is called to rerun the analysis and produce results in the working folder for comparing the original situation with the emulated scenario.

Comments and current limitations

This is is not a complete virtual loop! Please keep in mind that the variant shows a new and independent decision at each timestep. It is not a sequence of decisions building on each other because there is no forecasting what a different decision would have lead to at the next time step. Therefore you need to carefully consider the results at each timestep separately and consider whether it is safe enough to be activated in AAPS. Once this emulator gives me promising hints and the parameter is tuned I go to quasi "open loop", e.g. for a few days I manually apply an increase in SMB as a correction bolus before going live.

The code still contains dead sections because it is a spinoff from scanAPSlog.py, a more complete analysis of the logfiles. I use that to populate my own local database for treatments, sgvs, etc. independant of Nightscout.

The code contains many (in)active print statements for debugging purposes. I am still learning python and have not yet taken the lesson on proper debugging.

If the logfile starts in the middle of preparing for the loop decision this event lacks required json definitions and may lead to a crash. If you absolutely need that logfile append it to its predecessor logfile and use the combined one.

Be aware that maxBolus=0 in the tabular output file means maxBolus was high enough to not limit the SMB and was therefore not listed explicitly in the logfile by AAPS.

Glucose values are assumed to be in mg/dl.

Tested system settings

- Windows10, 64 bit
- python 3.7, 3.8
- some python libs like matplotlib (see the import statements in vary_settings_core.py)
- AAPS versions for logfile: orefl & SMB for 2.3 / 2.5 / 2.5.1 / 2.6.1
- AAPS versions for determine_basal: orefl & SMB for 2.3 / 2.5 / 2.5.1 / 2.6.1

Installation

Download the Github repo as a zip archive and extract it to a folder containing

- "determine_basal.py" as the emulation of the original java version
- "vary_settings_batch.py" which manages the process in a DOS window
- "vary_settings_GUI.py" which manages the process in an interactive window
- create a shortcut on your desktop to point to this GUI file.
- "vary_settings_core.py" which contains the core programme
- "<AAPS logfile>" (zipped or unzipped) for the period to be investigated
- "<variant_label>.dat" which contains the definition of the settings you want to change

Define your variant

Create or edit "<variant_label>.dat" to define the changes for your what-if scenario. You have access to the relevant lists or dictionaries as shown in the tab „SMB“, sections „Glucose-data“ through to „Autosense-data“. The general format per row is:

```
.  
.   
<array> <item in array> <new value or formula> [<optionally: ### any comment>]  
.   
.
```

The entries may be separated by several BLANKS and aligned for better readability. Their meaning is as follows:

- *<array>*
name of python dictionary; the available names are
 - autosens_data do not change; gets calculated inside AAPS elsewhere
 - glucose_status not really meaningful
 - currenttemp not really meaningful
 - iob_data do not change; gets calculated inside AAPS elsewhere
 - meal_data do not change; gets calculated inside AAPS elsewhere
 - profile most variations will happen here

If array is not a recognised name the row will be skipped.

Must not contain BLANKS

- *<item in array>*
Item in <array>
Must not contain BLANKS
- *<new value or formula>*
Boolean or numeric value or expression that evaluates accordingly
May contain BLANKS but must not contain '###'

optionally also:

- *<### any comment>* an optional comment after a block of '###'

For detailed content of the arrays and their correct spelling see the original logfile and look for rows containing "[DetermineBasalAdapterSMBJS.invoke():" or check the appropriate sections in the AAPS “SMB“ tab.

CAUTION:

Whereas AAPS will check for limits and deactivate some SMB settings this emulator does not know about them and just goes ahead. Therefore it is your responsibility to check whether AAPS would accept the settings from your variant definition file.

Example: This repo contains the variant definition file „Demo_Sports_Adaptations.dat“. It will be used for a logfile covering 2-3 hours of playing badminton. I like to stay below 140mg/dl because otherwise I lose concentration and react too slow. In this case it worked quite well but what if I tweak things a bit?

```
profile    min_bg          101                ### just above the temp target threshold for SMB
profile    max_bg          profile['min_bg']      ### redefine target
#profile   target_bg       int((profile['min_bg'] + profile['max_bg'])/2)  ### example of numerical expression
profile    tempTargetset   True               ### spelling error
profile    temptargetSet   True               ### i.e. not using NS profile target
profile    allowSMB_with_high_temptarget False      ### was True at the time
```

1. The first row changes the upper target from 100 to 101 to move it just outside of the „enableSMB_with_temptarget“ range.
2. The second row sets the upper target value the same as the lower one with a simple expression. It needs to be the same because I want to simulate a temp target which does not allow for a finite bandwidth if defined in the AAPS GUI.
3. The third row is ignored because „#rofile“ is not a recognised dictionary. But just look at the rather complex expression assignment possible. As far as targets are concerned the „target_bg“ is defined as the average of „min_bg“ and „max_bg“ anyway inside determine_basal meaning this row is not required.
4. The fourth row tries to redefine the flag for „tempTargetset“ as True. However, that spelling is not really correct because there are 2 swaps in lower case and upper case spelling.
5. The fifth row keeps the flag for „temptargetSet“ as True and this time in correct spelling. There is no check to verify the target bandwidth is zero in such a case.
6. Finally the sixth row disables SMB for higher TempTargets, i.e. above 100.

In summary the target increases from 100 to 101 and following a general recommendation for sport SMB is now disabled.

For first time use I recommend to leave things unchanged and verify that the results agree with the original. Minor numerical deviations are probably due to roundings in the original logfile.

There are two special cases of quasy arrays that were introduced to handle time varying assignments:

- STAIR <UTC-date/time> < value from this time onwards>
This is like defining a basal profile
- INTERPOL <UTC-date/time 1> < value at this time>
INTERPOL <UTC-date/time 2> < value at this time>
This uses linear interpolation and extrapolation to define a time dependent function

An example shows possible usage of how to define and how to reference the functions:

```
STAIR      2020-04-14T00:00:00Z    111
STAIR      2020-04-14T12:50:00Z    125
STAIR      2020-04-14T13:10:00Z    131
INTERPOL   2020-04-14T13          100
INTERPOL   2020-04-14T13:30:00Z    140

profile    min_bg          round(INTERPOL) - 20      ### time varying value
profile    max_bg          STAIR - 10                ### time varying value
```

Execute the analysis

For those who are less fit in DOS commands I created a GUI front end which collects the inputs in a form (see appendix). Otherwise, open a DOS command window, navigate to the working folder containing "<variant_label>.dat" and enter

```
DOS-prompt>python vary_settings_batch.py <logfile(s)> <options> <variant> [<from> [<to>]]
```

The meaning of the arguments is as follows:

- *<logfile(s)>*
the AAPS logfile (still zipped or unzipped) downloaded from Android.
Optionally you may use the wild characters (? for matching any single character; * for matching any string) to match several logfiles in the directory which provides a longer time window to evaluate. An example could be „AndroidAPS._2020-04-02_“ to handle all logfiles of that day.

- *<options>*
"All" or a slash separated list of things to be shown so it does not overload the plot.
The list is any subset from "insReq/maxBolos/SMB/basal/target/bg/pred/flowchart" where
 - insReq is the total insulin required at the current time step
 - maxBolos is the upper limit of allowed SMB
 - SMB is the actual SMB to be delivered
 - basal is the actual basal insulin to be delivered
 - target is the lower and upper target
 - bg is the original glucose value
 - cob is the original COB value to help understanding the situation
 - iob is the original IOB value to help understanding the situation
 - activity is the original insulin activity to help understanding the situation
 - as_ratio is the Autosense ratio
 - pred plots the prediction curves per timestep of the emulated scenario including hints if SMB needed to be disabled; you get a quasi animation by using forward/backward keys in the pdf-file created with one page per loop execution
 - flowchart in the graphics output only it plots a flowchart of how determine-basal was transversed with all the if-branches executed.

With the special flag „-“ you can suppress individual <plot-options>. I normally use „All/-pred/-flowchart“ which results in everything apart from „pred“ and „flowchart“ in order to speed up the programme significantly in routine analyses.

- *<variant_label>*
A label for the result file names and plot. It is also the base name of the variant definition file "<variant_label>.dat".

or alternatively:

- *<variant_label>.dat*
This alternative method of giving the filename directly may be easier especially when using the DOS type ahead method

The optional parameters may be used to define a time window to be evaluated, whether for a single logfile or a list of them:

- *<from>*
Define the beginning of the time window like „2020-04-02T22:00:00Z“ or for short „2020-04-02T22“. If you want to skip the start time then just use „2000“ as a very early year.
- *<to>*
Optionally also define the end of the time window, again in the readable UTC format used in the logfile system.

The process will create the following files:

- `<logfile>.<variant>.log`
echo of parameter changes as requested by `<variant>.dat`
- `<logfile>.orig.txt`
extract of original logfile with entries relevant for the loop result; for the same logfile this will always be the same name and be independent of the variant; therefore there is no harm when a different variant is emulated and it gets overwritten.
- `<logfile>.<variant>.txt`
equivalent, emulated subset for the variant loop result
- `<logfile>.<variant>.tab`
comparison of key values showing original versus emulated run; a summary line at the bottom compares total basal and total SMB delivered
- `<logfile>.<variant>.pdf`
plot of the tabular results; if the previous file with that name is still open in pdf-viewer a warning will be printed until it is clear to proceed.

The comparison table file is also output to the screen and always contains all columns.

The plot is saved in a predefined resolution(200dpi) and size(9 x up to 12 inches, portrait). Without the „pred“ option is is also displayed interactively and things like aspect ratio or margins can be modified. After such modifications the graph can be saved manually by clicking the disk symbol, preferably in jpeg or png format which makes it easier to later include it in other documents. With the “pred“ option you can inspect it in a pdf-viewer and transverse the history of the logfile with the forward and backward keys.

A file compare between the two txt-files is possible, but mainly the formating is still different and flags too many lines although numerically equivalent. Personally I find the table and plot more useful.

Example: What do the result files look like for the example from above?

First let us check „AndroidAPS. 2019-11-13 00-00-00 .7.Demo Sports Adaptations.log“:

```
===== Echo of what-if definitions actioned for variant Demo_Sports_Adaptations
===== created on Wed, 22 Apr 2020 12:02:11 +0200
===== for loop events found in logfile AndroidAPS._2019-11-13_00-00-00_.7

loop execution in row=2050 of logfile AndroidAPS._2019-11-13_00-00-00_.7 at= 2019-11-13T18:23:30Z
edited old value of 100 in profile with min_bg=101
edited old value of 100 in profile with max_bg=101
not actioned: [#rofile], [target_bg], [int((profile['min_bg'] + profile['max_bg'])/2)]
appended new entry to profile with tempTargetset=True
edited old value of True in profile with temptargetSet=True
edited old value of True in profile with allowSMB_with_high_temptarget=False

loop execution in row=5288 of logfile AndroidAPS._2019-11-13_00-00-00_.7 at= 2019-11-13T18:28:30Z
edited old value of 100 in profile with min_bg=101
edited old value of 100 in profile with max_bg=101
not actioned: [#rofile], [target_bg], [int((profile['min_bg'] + profile['max_bg'])/2)]
appended new entry to profile with tempTargetset=True
edited old value of True in profile with temptargetSet=True
edited old value of True in profile with allowSMB_with_high_temptarget=False
...
```

This is an echo of the changes assigned at each timestep. It is useful to check results of complex expressions for values and verify that there were no spelling errors like in the „appended ...“ rows

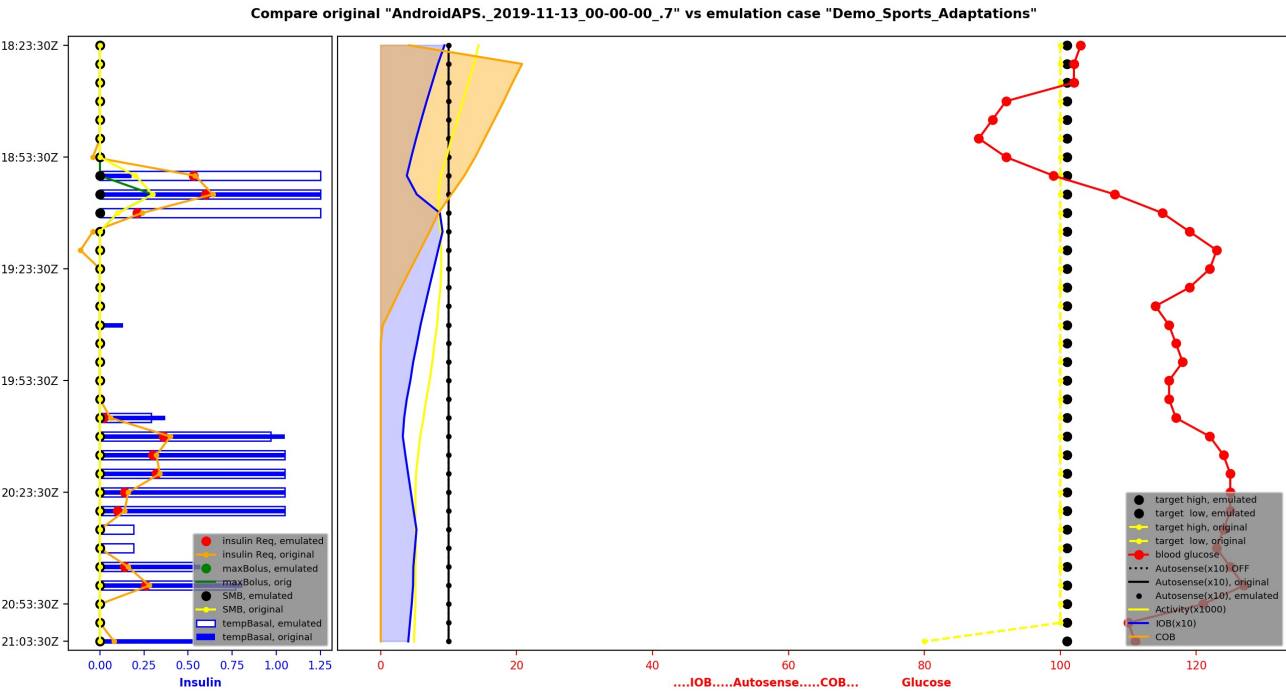
Next check „AndroidAPS. 2019-11-13 00-00-00 .7.Demo Sports Adaptations.tab“:

id	time	--UNIXtime--	bg	cob	iob	act	-Autosens-		----target----		insulin Req		-maxBolus-		---SMB---		---tmpBasal---			
							orig	emul	orig	emul	orig	emul	orig	emul	orig	emul	orig	emul		
0	18:23:30Z	1573669410.3	103	4.2	0.94	0.014	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
1	18:28:30Z	1573669710.2	102	20.8	0.84	0.014	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
2	18:33:30Z	1573670010.3	102	19.4	0.76	0.013	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
3	18:38:30Z	1573670310.3	92	18.1	0.68	0.012	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
4	18:43:30Z	1573670610.3	90	16.7	0.6	0.011	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
5	18:48:30Z	1573670910.3	88	15.3	0.52	0.01	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
6	18:53:30Z	1573671210.3	92	13.9	0.45	0.01	1.0	1.0	100-100	101-101	-0.04	0	0	0	0	0	0	0		
7	18:58:30Z	1573671510.3	99	12.3	0.38	0.009	1.0	1.0	100-100	101-101	0.54	0.53	0	0	0.2	0	0.18	1.25		
8	19:03:30Z	1573671810.6	108	10.4	0.53	0.009	1.0	1.0	100-100	101-101	0.64	0.6	0.3	0	0.3	0	1.25	1.25		
9	19:08:30Z	1573672110.3	115	8.5	0.87	0.009	1.0	1.0	100-100	101-101	0.24	0.21	0.1	0	0.1	0	0	1.25		
10	19:13:30Z	1573672410.3	119	7.2	0.91	0.009	1.0	1.0	100-100	101-101	-0.04	0	0	0	0	0	0	0		
11	19:18:30Z	1573672710.3	123	5.8	0.84	0.009	1.0	1.0	100-100	101-101	-0.11	0	0	0	0	0	0	0		
12	19:23:30Z	1573673010.3	122	4.4	0.78	0.009	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
13	19:28:30Z	1573673310.3	119	3.0	0.71	0.009	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
14	19:33:30Z	1573673610.2	114	1.7	0.65	0.009	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
15	19:38:40Z	1573673920.6	116	0.3	0.58	0.008	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0.13	0		
16	19:43:30Z	1573674210.4	117	0	0.53	0.008	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0.03	0.01		
17	19:48:30Z	1573674510.4	118	0	0.48	0.008	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
18	19:53:30Z	1573674810.3	116	0	0.44	0.007	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
19	19:58:30Z	1573675110.4	116	0	0.38	0.007	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
20	20:03:30Z	1573675410.6	117	0	0.35	0.006	1.0	1.0	100-100	101-101	0.06	0.02	0	0	0	0	0.37	0.29		
21	20:08:30Z	1573675710.4	122	0	0.33	0.006	1.0	1.0	100-100	101-101	0.4	0.36	0	0	0	0	1.05	0.97		
22	20:13:30Z	1573676010.3	124	0	0.36	0.006	1.0	1.0	100-100	101-101	0.32	0.3	0	0	0	0	1.05	1.05		
23	20:18:30Z	1573676310.4	125	0	0.4	0.005	1.0	1.0	100-100	101-101	0.34	0.32	0	0	0	0	1.05	1.05		
24	20:23:30Z	1573676610.3	125	0	0.44	0.005	1.0	1.0	100-100	101-101	0.16	0.14	0	0	0	0	1.05	1.05		
25	20:28:30Z	1573676910.8	125	0	0.48	0.005	1.0	1.0	100-100	101-101	0.14	0.1	0	0	0	0	1.05	1.05		
26	20:33:30Z	1573677210.3	124	0	0.53	0.005	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0.19		
27	20:38:30Z	1573677510.4	123	0	0.5	0.005	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0.19		
28	20:43:30Z	1573677810.3	125	0	0.48	0.005	1.0	1.0	100-100	101-101	0.16	0.14	0	0	0	0	0.57	0.53		
29	20:48:30Z	1573678110.7	127	0	0.48	0.005	1.0	1.0	100-100	101-101	0.28	0.26	0	0	0	0	0.81	0.77		
30	20:53:30Z	1573678410.3	121	0	0.46	0.005	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
31	20:58:30Z	1573678710.3	110	0	0.43	0.005	1.0	1.0	100-100	101-101	0	0	0	0	0	0	0	0		
32	21:03:30Z	1573679010.7	111	0	0.41	0.005	1.0	1.0	80- 80	101-101	0.08	0	0	0	0	0	0.53	0		
Totals:																	0.6	0.0	0.76	0.91

This table lists the main comparisons between the original case and the emulated case. In the target columns the last row shows that the original target was reduced from 100 to 80 because sport was over. As the variant continued with 101 just ignore that last timestep in the final judgement. The Insulin required is slightly lower as expected due to a slightly higher target. The SMB columns confirm that all of the original SMBs were disabled. The tempBasal now has to offset that lack of insulin around those timesteps and is otherwise slightly reduced because of reduced requirements. The totals row at the bottom gives an indication of the changes but the timing of the major differences is important. While the SMBs of 0.6U were delivered the basals added up to 0.12U in those 15 minutes giving a total of 0.72U. In the variant case the total basal insulin adds up to 0.31U in that short period. So in summary this difference is too small to justify a change in settings which worked well on that day and the next week will be different anyway because the body will be in a different condition. But the process could be repeated for the logfiles of the equivalent activities on other days.

The graphical representation of the comparison tells the same story. The best evaluation method is to look at both, the graph and the table when assessing the comparison. The graph for the example

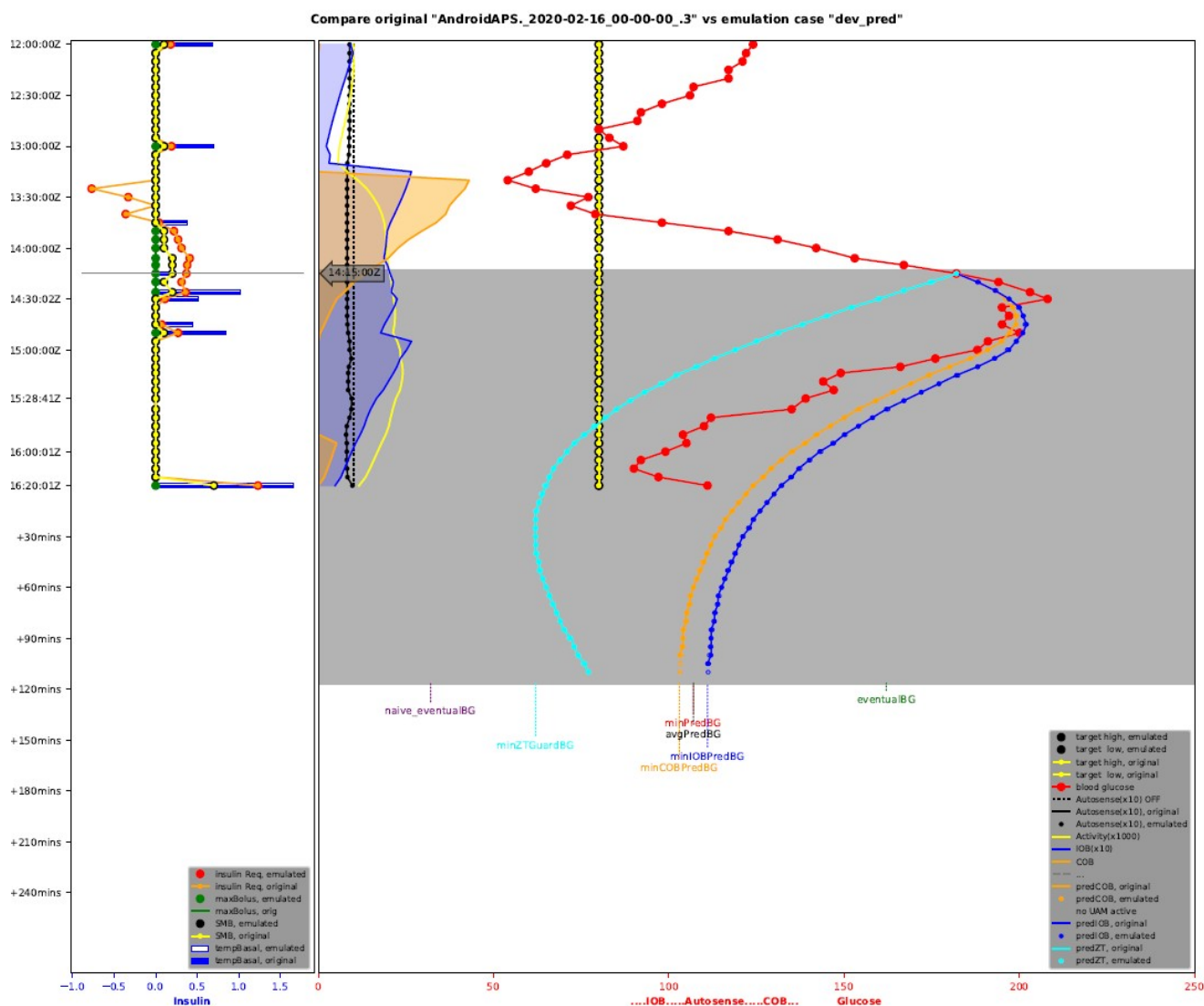
looks like this:



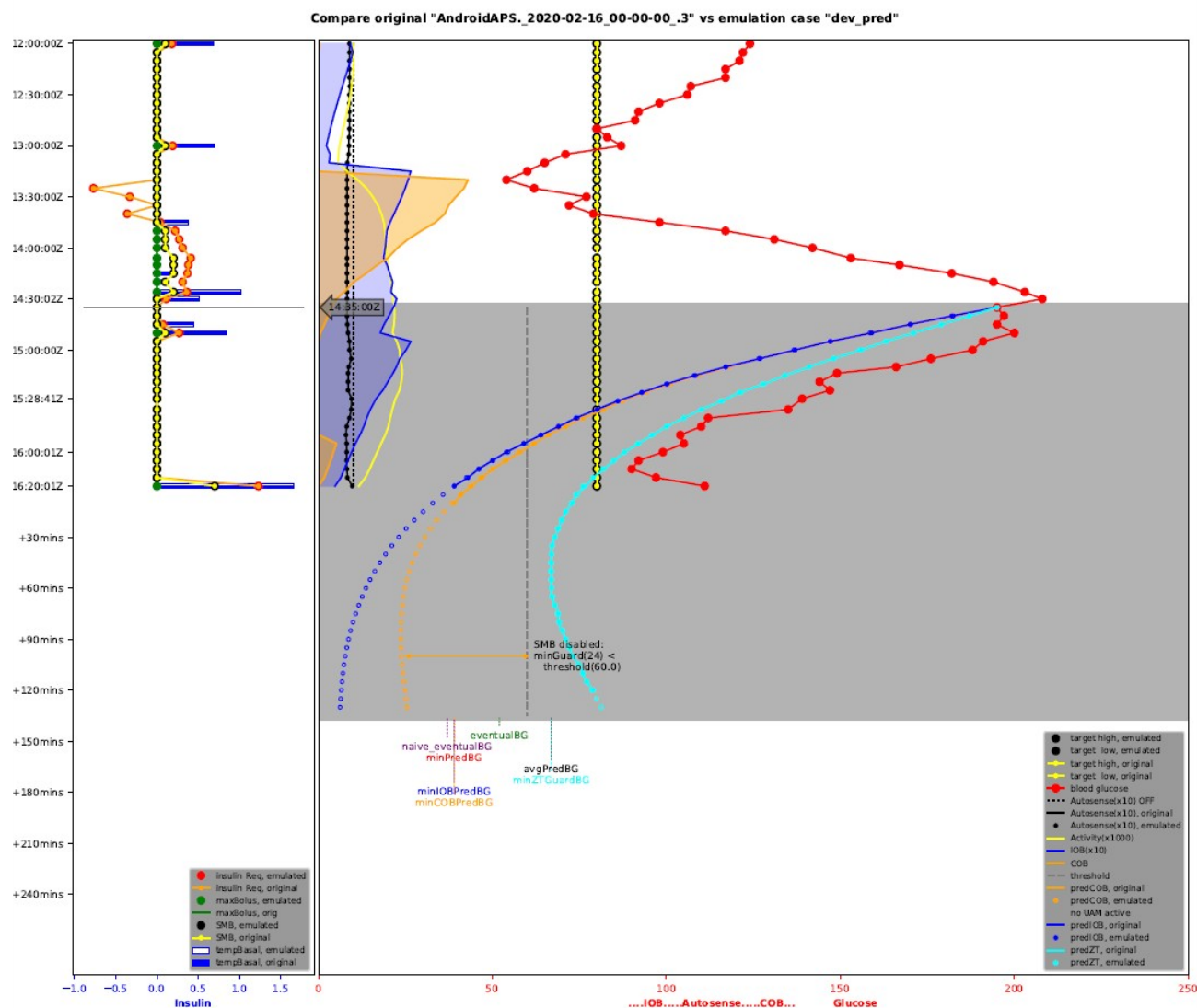
The new „pred“ output option - an early release

It overlays the graph with the prediction lines known from the AAPS “Home“ tab. With the backward and forward arrows in the pdf-viewer you can get a slow motion animation of the history. The hope is that this will improve the understanding of how the loop works.

The example below is from a different logfile with no variation active and shows a moment when the predictions matched the future glucose quite well. This means the settings matched real life quite well for about one hour and that there was no outside disturbance like new carbs or target changes.



Some timesteps later the loop reports in its „SMB“ tab a reason why SMB was disabled. Such a situation is included in the graph whenever it applies to help understanding of the reasoning. What I find interesting is that the loop compares the predictions against the threshold rather than the lower target. That threshold is defined as 50% of the lower target plus 20mg/dl, i.e. always below target!



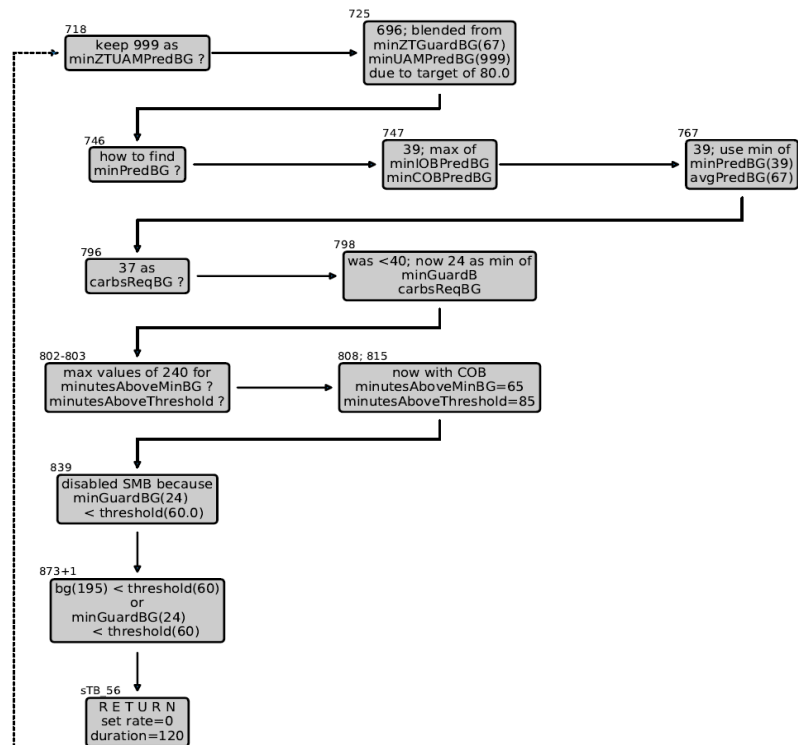
This graphical sketch of special situations is still under development and needs more debugging.

By the way, open circles in the predictions show the initial calculations which later get capped, truncated and rounded in AAPS and those final values are shown as filled circles.

The new „flowchart“ output option - an early release

In order to better understand the logic and decisions in determine-basal I added this extra graphic. As for the „pred“ option a new flowchart is created at each time step. As an example here is a zoomed in region for the same timestep as before, i.e. the disabling of SMB at 14:35:00Z;

00-00-00.3" vs emulation case "dev_pred"



Flowchart and decision logic at time 14:35:00Z

It shows the source of the decision in row 839 of the original determine-basal.js

Appendix – Using the GUI front end instead of the DOS window

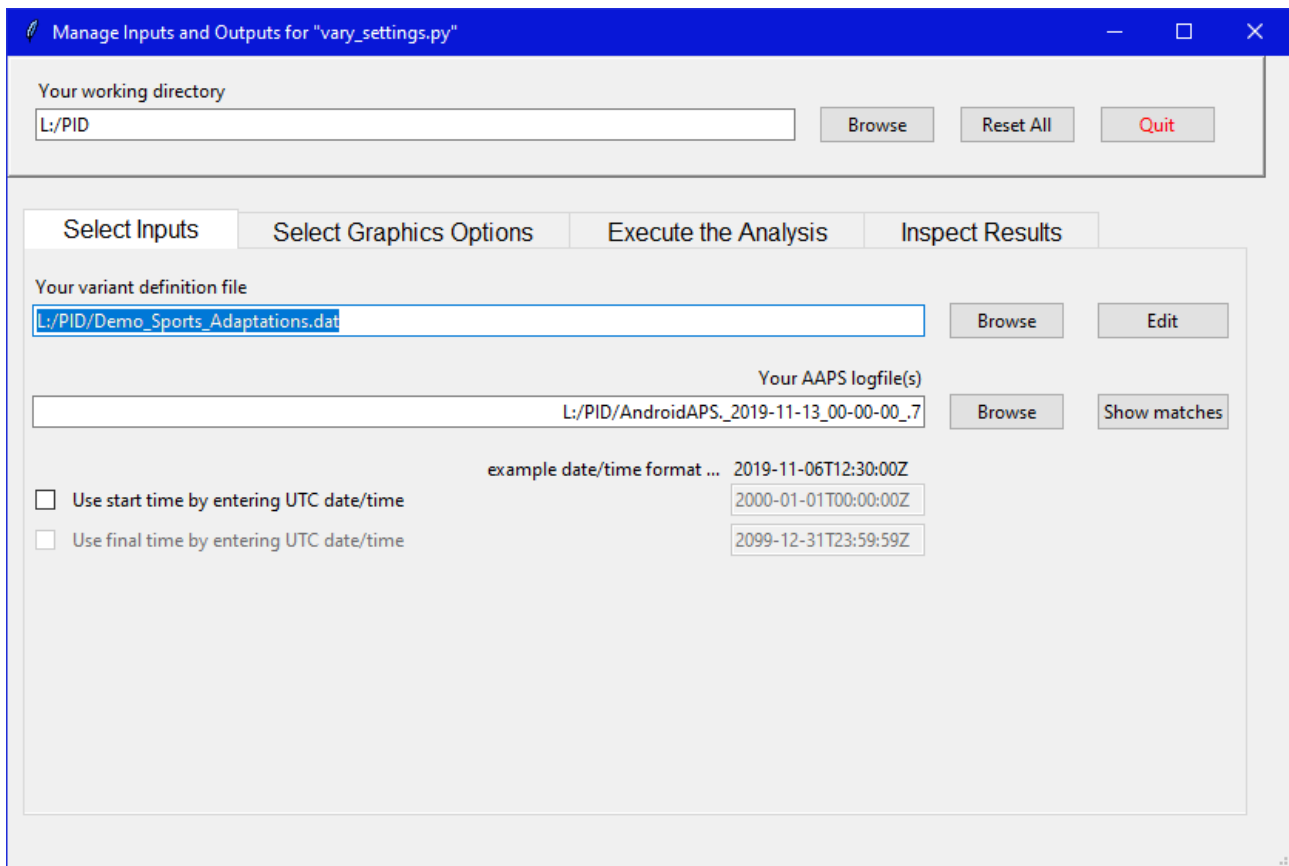
Start by clicking on the shortcut created on the desktop during the installation process. This opens a DOS window. The DOS window should remain empty but may be error messages will show up and I recommend to check for those. I tried to catch such messages in the code for display in the form itself but you never know.

After opening that DOS window another window opens which contains a form itself to manage the whole process. The size of the window may be adjusted so longer filenames are displayed without scrolling. In general, boxes with white background are used to enter text, mostly filenames. Boxes with grey background are command buttons to start certain actions:

- Browse – will start the standard file selection dialog to find the respective file
- Edit - loads that file into your standard Editor (as long as your system has assigned a standard app for that file type)
- Show – similar to „Edit“ apart from the result PDF where it uses your standard PDF viewer

The process goes from top to bottom and from left to right but of course you can jump to wherever. First, you select the working folder, i.e. the one containing your logfiles, the variant definition file and later the result files. This input is not really required but it preselects the folder for all later file browser actions.

Click the „**Select Inputs**“ tab if not already active. The two required inputs are the names of variant definition file and the logfile.



If you need to scan several logfiles for longer time windows you edit the filename by using „*“ or „?“ in the suitable position of the filename. In such cases you can check which files will match the wild card specification by clicking the button „Show matches“.

For specifying an optional start date/time you activate tick box. The time field will no longer be greyed out and you can enter/edit the start date and time. Above the field you see the format of how

to specify the date and time. Remember AAPS uses the internal UTC format which is Greenwich time without daylight savings offset., e.g. in MESZ (Middle-European-SummerTime) it is numerically 2 hours behind. Normaly I use copy and paste from that part of the logfile field und then do the detailed adjustments. If final time time is also wanted, again tick the box, copy and paste from the start time and adapt it.

Clicking „Reset All“ will clear all input fields and return to the defaults.

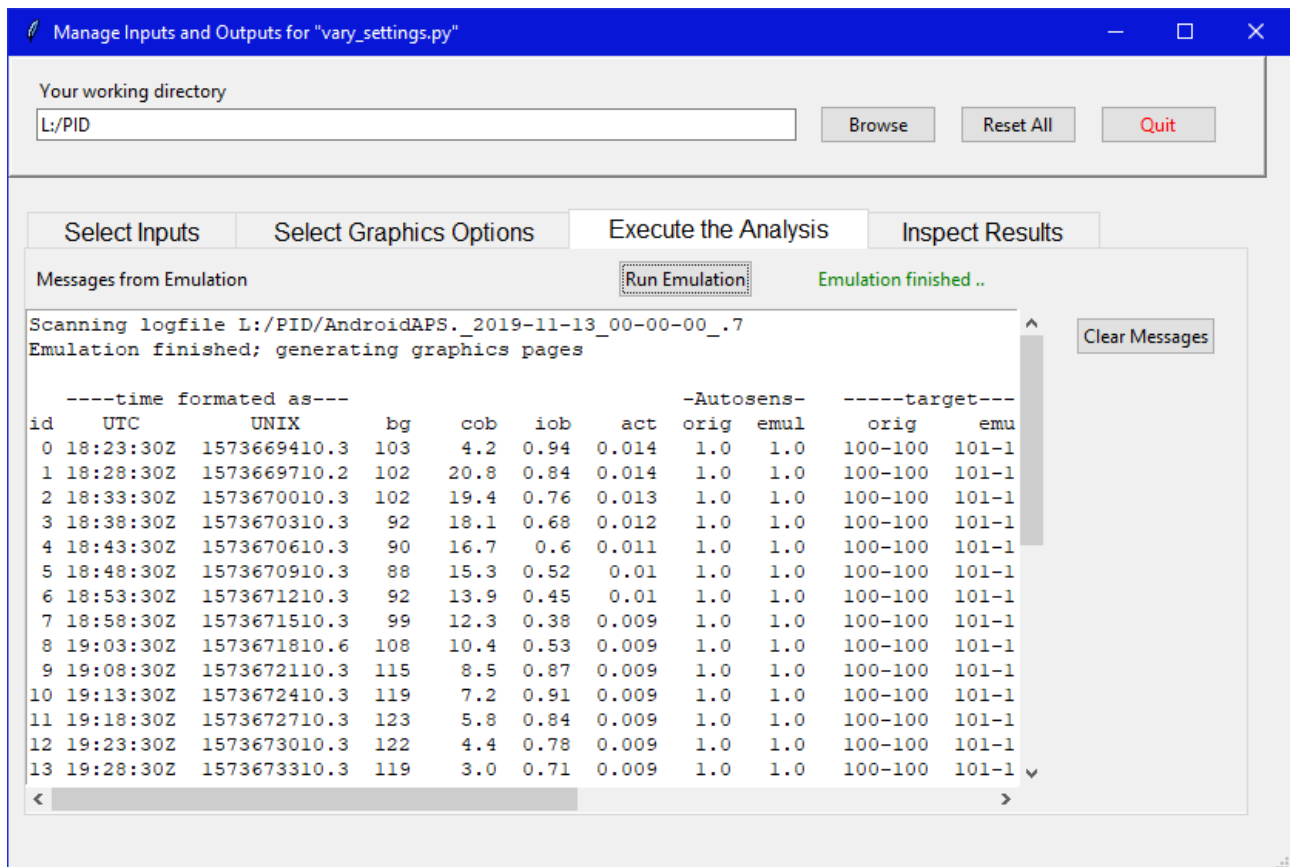
Click the „**Select Graphics Options**“ tab if not already active. This form selects what information to show in the graphic output. It uses radio buttons for the coarse selection and where appropriate shows tick boxes for details to be included or excluded. The resulting string as used in the DOS method is dynamically updated to reflect the tick actions but could also be edited manually.

The initial selection is „most“ which is the most frequent option. In mode „most“ you tick those options you are not inntereted in.

In mode „just a few“ it is the opposite, i.e. you select those options you are specifically interested in.

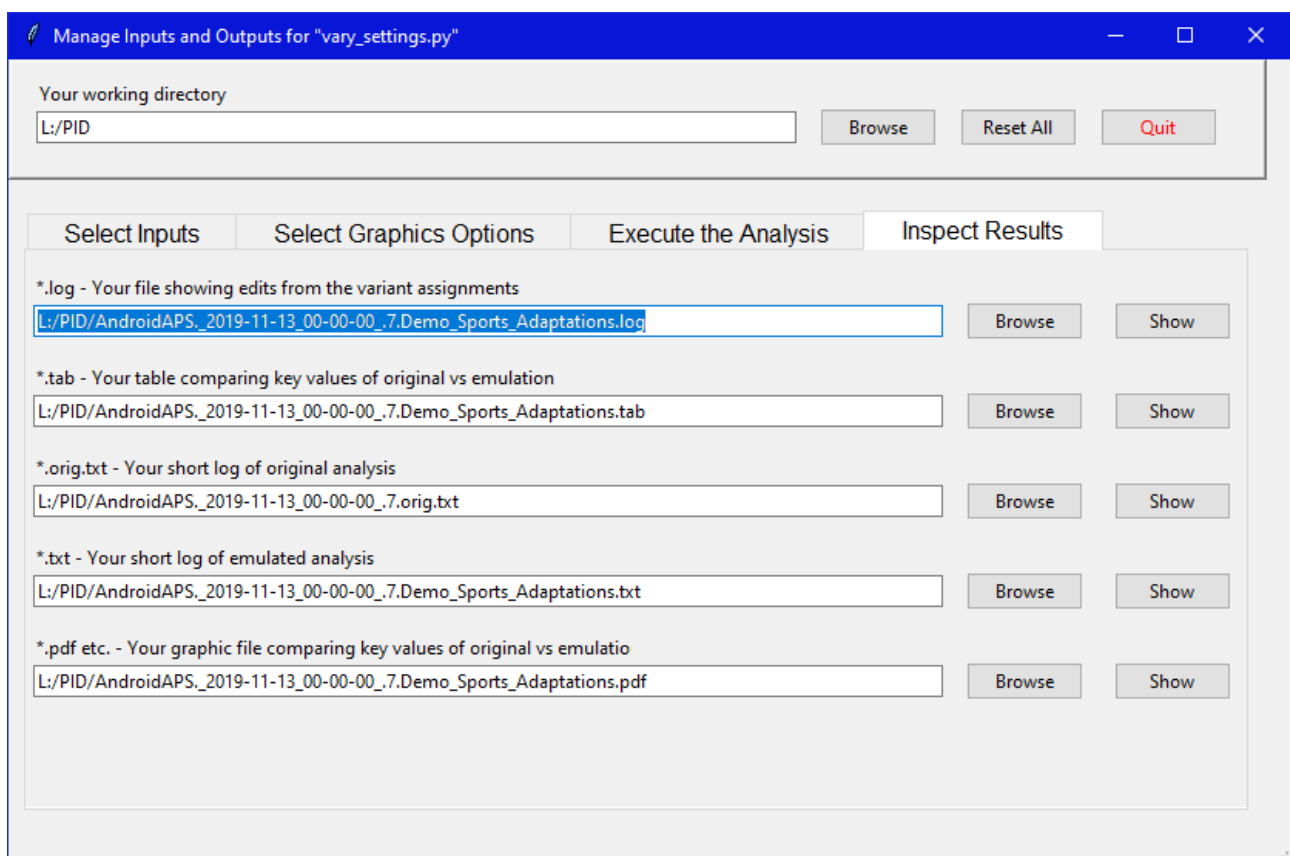
In mode „All“ the fine grained selctions in the bottom are disabled and hidden because they make no sense in that context.

Click the „**Execute the Analysis**“ tab if not already active. The button „Run Emulation“ will kick off the analysis. The large white box shows messages otherwise displayed in the DOS window. Here it may male sense to adjust the window size the show more content without scrolling. The vertical resizing is limited to 30 lines so the window fits on a normal monitor.



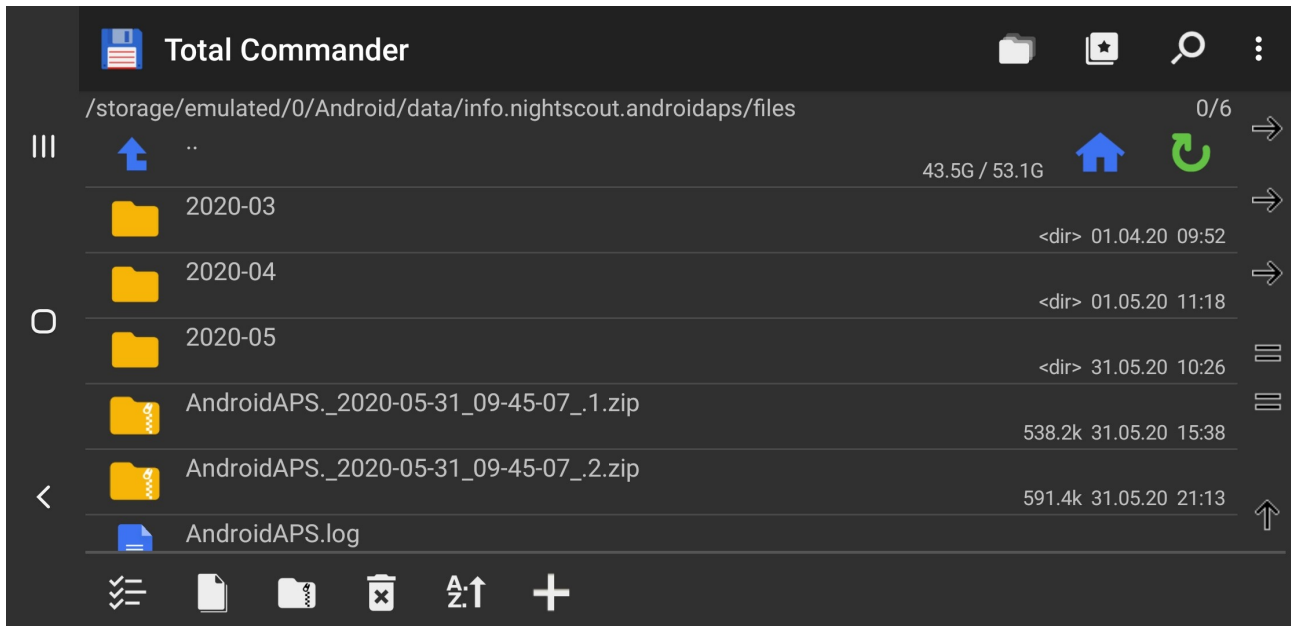
Error messages or hints about missing inputs are displayed in red. The button „Clear Messages“ obviously will clear all contents in the rmessage box. This is useful before starting the next analysis.

Click the „Inspect Results“ tab if not already active. This tab will also be activated automatically once the analysis is finished. Also the filenames are already prefilled after the analysis. However, you may use the „Browse“ buttons to select and see content of analyses run before.



Appendix – the AAPS logfile system

Where can you find those logfiles? They are stored on the phone in the logfolder „storage/emulated/0/Android/data/info.nightscout.androidaps/files“.



The current logfile is named „AndroidAPS.log“. All messages are appended until it reaches a size of about 5MB. Once that happens it is converted into a zip archive and a fresh logfile is started.

Those zip archives accumulate on the phone in the logfolder and occupy about 0.5MB each. The only time they are deleted is when installing a new release of AAPS. So if you want to keep them for later reference you should copy them elsewhere, e.g. to a USB drive.

The zip archives are named like „AndroidAPS._2019-11-13_00-00-00_.7.zip“, the logfile contained within the zip would be named „AndroidAPS._2019-11-13_00-00-00_.7“. Obviously the names help to find the correct day. Within a day the files are numbered sequentially, here „7“. Some events like rebooting the phone trigger a restart of the naming and the time part of the filename reflects that time and may look like „..._15-30-45_.0“ which by the way also restarts the counter field at the end.

Depending on the activity in the loop there may be more or less messages. Normally I end up with 5 to 12 logfiles per day and in my first year of looping I collected 2500 logfiles. The folder would therefore become very messy and I create subfolders for each month, both on the phone and in the offline backup.

The logfiles are plain text files which can be inspected with a standard editor. They contain more information than stored in Nightscout. You need to get used to the code words for the various messages. For example you may look for words like „alarm“ or „pump“ or „ruffy“ when chasing for a pump problem.