

Editing the 'automatic_MAP18.ini' file

This text file (which is found in the directory '...\MAP18_for_SPM12\MAP18_Program') controls the automated conversion of DICOM images into ANALYZE format, the detection of different MR sequences, the naming rules for image files, and the subsequent image processing with MAP18.

The following overview shall explain which items have at least to be adjusted for each user site.

The screenshot shows the 'automatic_MAP15.ini' file in Notepad++ with several sections and annotations:

- General Information:**
 - Name:** automatic_MAP15.ini
 - Purpose:** Parameter file used by map15 for automated image analysis (pipeline)
 - Note:** This file can be used as a template for creating different *.ini files.
- [Directories]** # paths to important directories: DICOM imagebox, directory of MAP15 analysis, and backup directory for images converted to ANALYZE format
 - DICOMDirectory:** '\\fsmri01\MRI\KFacsV16\Imagebox' (Annotated: Path to the Imagebox of the PACS system)
 - Destination:** 'E:\MR-Daten_Zuerich\3D-MRT-Analyse' (Annotated: Directory where all result files shall be created)
 - BackupDirectory:** 'E:\MR-Daten_Zuerich\ANALYZE-Daten'
- [TimeStamp]** # defines the time point from that on new images in the DICOMDirectory shall be analyzed with MAP15
 - StartingTime:** '26-Oct-2017 09:12:19'
 - alternatively, use a distinct time point (e.g., StartingTime = '17-Jul-2012 09:47:27' or StartingTime = 'now')**
- [Diverse Items]**
 - SolidStateDisk:** 'd:' (if there is no SSD, leave empty (i.e. ''))
 - SkipExistingData:** 'no' (if 'yes' >> if subdirectory with MRI data already exists, don't create alternative subdirectory but skip)
 - Start_MATLAB_in_new_window:** 'yes' (if 'no' >> process images consecutively within the same MATLAB window; 'yes' >> start each processing in new window)
 - Delay:** 5 (wait for x minutes before proceeding with next data set (shall prevent crash due to multiple open windows))
 - CropImage:** 'yes' (if 'yes' >> try to crop input images to accelerate image processing; 'no' >> don't try cropping)
- [InstitutionNames]** # determine institution from item 'InstitutionName' in DICOM header
 - Basel Bilddiagnostik = 'Bild Diagnostik Basel', 'BILDDIAGNOSTIK BASEL'
 - Berlin = 'Radiologische Praxis am KEH'
 - Bethel = 'EvKB-Mara'
 - Bochum = 'Bochum', 'Borad'
 - Brussel = 'Brussels'
 - CHUV Lausanne = 'Trio CHUV Lausanne'
 - Cleveland = 'Cleveland', 'CCF', 'MELLEN CENTER CLEVELAND CLINIC', 'Cleveland Clinic L10 3T'
 - Epilepsiezentrum Kork = 'Klinikum Offenburg Radiologie', 'Olghospital Stuttgart', 'Epilepsie Zentrum Kork', 'Ortenau Klinikum Offenburg'
 - Genf = 'Genf', 'HUG', 'Hopitaux Universitaires Geneve 3T'
 - Heidelberg = 'Heidelberg'
 - Hirslanden Zurich = 'Klinik Hirslanden Zurich', 'Klinik Hirslanden Zurich', 'KLINIK HIRSlanden', 'Klinik Hirslanden Zurich', 'KLINIK HIRSlanden'
 - Hirslanden Park = 'neuroradiologie schanze', 'KLINIK IM PARK ZUERICH', 'Klinik im Park Zuerich'
 - HUG = 'HUG', 'Genf', 'Hopitaux Universitaires Geneve 3T'
- [Tokens]** # define token for each institution
 - BBD:** 'Basel Bilddiagnostik'
 - B:** 'Berlin'
 - BE:** 'Bethel'
 - BO:** 'Bochum'
 - BRU:** 'Brussel'
 - CHUV:** 'CHUV Lausanne'
 - CCF:** 'Cleveland'
 - Kork:** 'Epilepsiezentrum Kork'
 - HUG:** 'Genf'
 - HD:** 'Heidelberg'
- [SeriesDescriptions]** # define series descriptions of volume data sets
 - T1:** 'T1', 'sT1/3D/TFE', 'sT1/3D/TFE', 't1_mpr_ns_sag', '3DT1 FSPGR', 't1_mpr_ns_sag', '3DT1 FSPGR', '3DT1 FSPGR mit Asset', 'T1W 3D'
 - T2:** 't2_TSE_TE100_cor_thin', 'T2W.tra.Str', 'T2 Ax 1mm^3 1 NEX', 'T2W_VISTA_HR', 'T2 SENSE', 't2_tse3dsvfl_ns_sag_st2_p2', 't2_spc'
- [SeriesDescriptions for Images to Coregister]** # define sequences which are to coregister and to normalize together with the T1 or T2 image
 - FLAIR_WBA:** 'T2W_FLAIR cor', 'T2W_FLAIR c', 'flair_tse3dsvfl_opt_fov_big', 'FLAIR 3mm Ep', 't2_tirm_cor_da-fl_2mm', 'Sag FLAIR CUBE', 'T2W_FLAIR'
 - FLAIR:** 'FLAIR', 'T2W_FLAIR', 't2flair3dwip_ns_tra_1.2mm', 'T2 Axial FLAIR', 'T2 Cor FLAIR', 'T2W_FLAIR', 't2_blade_tra_dark-fl_32'
 - Hemoflash:** 't2_fl2d_tra_hemo', 'T2W_Haemo', 'cor T2 fl2d hemo 320_FIL', 'T2 Ax GRE'
 - T2TSE:** 'T2W_TSE', 't2_tse_tra_2mm', 'cor T2 tse1 1mm 384_FIL', 'T2 TSE 3mm', 't2_tse_cor_512_2mm', 'T2 frFSE cor', 'STIR_tra_3mm'
 - T2:** '3D_T2_32chSHC', '3D_T2_32chSHC', 'e3D_T2_32chSHC SENSE', 'e3D_T2_32chSHC SENSE'
 - IR:** 'spc_ir_ns_cor_p2_iso', 't1_tir_cor'
 - T1KM:** 'T1W_FFE_KM', 'T1 TFE 3D KM NEU', 'T1W_FFE_KM'

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268
269 # determine several input arguments for MAP07 (i.e., mode,norm,space,sensitivity,ROI_mode,viewer)
270 # from a combination of different image items (i.e., 'Institution', 'Manufacturer', 'SeriesDescription' / 'ProtocolName', Manufacturer
271 #
272 # Parameters left of the identity sign (= input arguments for MAP07):
273 # first item: mode (of action; please, cf. map07('help'))
274 # second item: norm (database; please, cf. map07('help'))
275 # third item: space (i.e., 'standard', 'native', or 'both')
276 # fourth item: sensitivity (for detection of FCDs; i.e., 'high', 'medium', or 'low')
277 # fifth item: ROI_mode (i.e., 'full', 'closed', or 'dotted')
278 # sixth item: viewer (i.e., 'MRicro', 'MRicron', or 'none')
279 #
280 # Parameters right of the identity sign (= items defining the image type):
281 # first item: 'Institution'
282 # second item: 'Manufacturer'
283 # third item: 'SeriesDescription' / 'ProtocolName' (one expression for both items)
284 # fourth item: 'ManufacturersModelName' (optionally)
285 # Combinational logic 'Institution' AND 'Manufacturer' AND ('SeriesDescription' OR 'ProtocolName') AND ManufacturersModelName (opt.
286
287 [MAP07 Parameters]
288 'all','B_Symphony_T1','standard','medium','closed','MRicro'      = 'Berlin','Siemens','T1'
289 'all','BE_Symphony_T1','standard','medium','closed','MRicro'    = 'Bethel','Siemens','T1'
290 'all','AVG_T1','standard','medium','closed','MRicro'             = 'Bochum','Siemens','T1'
291 'all','BRU_Achieva_T1','standard','medium','closed','MRicro'     = 'Bruessel','Philips','T1'
292 'all','AVG_T1','standard','medium','closed','MRicro'             = 'CHUV Lausanne','Siemens','T1'
293 'all','BO_Trio_T2','standard','medium','closed','MRicro'         = 'CHUV Lausanne','Siemens','T2'
294 'all','CC_Trio_T1','standard','medium','closed','MRicro'         = 'Cleveland','Siemens','T1'
295 'all','AVG_T1','standard','medium','closed','MRicro'             = 'Epilepsiezentrum Kork','Siemens','T1','Symphony'

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Define on the left side the parameters (i.e. mode of action, viewer etc) with which MAP18 shall be started when an image as defined on the right side of the equation sign is found.

MS ini file 23164 chars 23882 bytes 360 lines Ln: 267 Col:1 Sel: 0 (0 bytes) in 0 ranges Dos/Windows ANSI INS

To start the automatic mode of MAP18 call **automatic_MAP18('automatic_MAP18.ini')** from the MATLAB command window. You may define different 'ini' files for different modes of action. In this case the command **map18('automatic')** opens a menu for the selection of the desired 'ini' file.

After calling the automatic mode of MAP18 the program waits for the arrival of new images in the image directory of the local PACS system. If an appropriate MRI with a 3D T1 image arrives, the DICOM images are converted to ANALYZE format and transferred to the destination directory. Then, MAP18 is automatically started and does the analysis on the new image(s).