

structural_preproc_local_STANDARD.m

/home/michele/Documents/FastTrack_fullSample/SCRIPTS/structural_preproc_local_STANDARD.m/home/michele/Documents/FastTrack_fullSample/orig_sof
Thu Feb 20 15:47:28 GMT 2020

structural_preproc.m

/home/michele/Documents/FastTrack_fullSample/orig_sof
Thu Aug 15 10:37:12 BST 2019

35 differences found. Use the toolstrip buttons to navigate to them.

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6 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%% [5 unmodified lines hidden]
7 . %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
8 %% setup . %% setup
9 [~,hostname] = system('hostname'); x [~,hostname] = system('hostname');
10 switch deblank(hostname) x switch deblank(hostname)
11     case 'michele-Precision-7530' x     case 'fslvm6.localdomain'
12         setenv('FOLDER','/home/michele/Documents/FastTrack_fullSample') x         bpath = 'xxx';
13         setenv('FSLDIR','/usr/local/fsl') x         case 'xxx'
14         setenv('FREESURFER_HOME','/usr/local/freesurfer') x         bpath = 'xxx';
15         setenv('SUBJLIST', 'subj_list_39x1.txt') % <
16         setenv('SGE_TASK_ID','1') <
17         setenv('NSLOTS','8') <
18     end x end
19 .
20 wrk_path = getenv('FOLDER'); x vps = [3:24 26 28:38 40 41 43:45];
21 bpath = fullfile(wrk_path, 'processing'); <
22 .
23 do_dcm2nii = 0; . do_dcm2nii = 0;
24 do_t1uniXinv2 = 0; %mp2rage: multiply t1_uni with t1_inv2 for better brain extra . do_t1uniXinv2 = 0; %mp2rage: multiply t1_uni with
[10 unmodified lines hidden]
35 do_fsseg = 0; . do_fsseg = 0;
36 do_overlap = 0; . do_overlap = 0;
37 .
38 fsl_path = getenv('FSLDIR'); x fsl_path = '/usr/local/fsl';
39 fs_path = getenv('FREESURFER_HOME'); x fs_path = '/usr/local/freesurfer';
40 fssub_path = fullfile(bpath,'freesurfer_subjects'); . fssub_path = fullfile(bpath,'freesurfer_subjects');
41 dicom_path = fullfile(bpath, 'raw'); %rawdata . dicom_path = fullfile(bpath, 'raw'); %rawdata
42 nii_path = fullfile(bpath, 'nifti'); . nii_path = fullfile(bpath, 'nifti');
[3 unmodified lines hidden]
46 datasets = {'t1_mp2rage_uni', 't2'}; . datasets = {'t1_mp2rage_uni', 't2'};
47 direcs = {'t2w2t1wReg', 'ACPCAlignment', 'BrainExtraction_FNIRTbased', 'xfms', ' . direcs = {'t2w2t1wReg', 'ACPCAlignment', 'BrainExtraction_FNIRTbased', 'xfms', '
48 .
49 % ADDED BY ME %%%%%%%%%% <
50 subj_list = getenv('SUBJLIST'); <
51 <
52 <
53 <
54 downloaded_data = fullfile(wrk_path,'data_mri'); <
55 n_cores = getenv('NSLOTS'); <
56 %%%%%%%%%% <
57 <
58 <
59 ref_t1_lmm = fullfile(fsl_path,'data','standard','MNI152_T1_lmm.nii.gz'); . ref_t1_lmm = fullfile(fsl_path,'data','standard',
60 ref_t1_lmm_brain = fullfile(fsl_path,'data','standard','MNI152_T1_lmm_brain.nii . ref_t1_lmm_brain = fullfile(fsl_path,'data','stand
61 ref_t1_lmm_mask = fullfile(fsl_path,'data','standard','MNI152_T1_lmm_brain_mask . ref_t1_lmm_mask = fullfile(fsl_path,'data','stand
[8 unmodified lines hidden]
70 '--match 9 --match 10 --match 11 --match 12 --match 13 --match 17 --match 18 . '--match 9 --match 10 --match 11 --match 12 --
71 '--match 1025 --match 2025', '--match 1008', '--match 1007', '--match 1013 --m . '--match 1025 --match 2025', '--match 1008', '--
72 .
73 <
74 % import subj_list for parallelization <
75 %subj_list = fullfile(wrk_path,'subj_list_2x1.txt'); %fullfile(wrk_path,'subj_l <
76 fid = fopen(subj_list,'r'); <
77 linenum = str2double(getenv('SGE_TASK_ID')); <
78 format_spec = '%d'; % This must reflect the columns in the file e.g.: format spe <
79 vps = cell2mat(textscan(fid,format_spec, 39,'delimiter','\n', 'headerlines',int3 <
80 <
81 <
82 %% do work . %% do work
83 for n_vp = 1:length(vps) . for n_vp = 1:length(vps)
84     vp_path = fullfile(nii_path,sprintf('VP%03d',vps(n_vp))); .     vp_path = fullfile(nii_path,sprintf('VP%03d',vps(n_vp)));
[16 unmodified lines hidden]
101     mkdir(mnifsfms_path) .     mkdir(mnifsfms_path)
102     mkdir(mnifs_path) .     mkdir(mnifs_path)
103     end .     end
104     %%% ADDED BY ME %%% <
105     subj_fold = fullfile(downloaded_data,sprintf('ID%03d',vps(n_vp))); <
106     struct_data = fullfile(subj_fold,'highres'); <
107     t1_uni_defaced = fullfile(struct_data,'t1_mp2rage_sag_p3_iso_uni_defaced.nii <
108     t1_inv_defaced = fullfile(struct_data,'t1_mp2rage_sag_p3_iso_inv2_defaced.ni <
109     t2_defaced = fullfile(struct_data,'t2w_spc_sag_p2_iso0.8_defaced.nii.gz'); <
110     % this is to obtain the T2 MNI152 that should be compatible with the script <
111     % system('flirt -in /usr/local/fsl/data/standard/mni_icbm152_t2_tal_nlin_asym <
112     % system('flirt -in /usr/local/fsl/data/standard/mni_icbm152_t2_tal_nlin_asym <
113     %%%%%%%%%% <
114     .
115     for set=1:length(datasets) .     for set=1:length(datasets)
116         set_path = fullfile(sess_path,datasets{set}); .         set_path = fullfile(sess_path,datasets{set});
117     .
118     %%% ADDED BY ME %%% <
119     if ~exist(set_path,'dir') <
120         mkdir(set_path) <
121     end <
122     if set == 1 <
123         t1_uni = fullfile(set_path, 't1_uni.nii.gz'); <
124         t1_inv = fullfile(set_path, 't1_inv2.nii.gz'); <
125         system(['cp ' t1_uni_defaced ' ' t1_uni ], '-echo'); <
126         system(['cp ' t1_inv_defaced ' ' t1_inv ], '-echo'); <
127     elseif set == 2 <
128         t2 = fullfile(set_path, 'T2.nii.gz'); <
129         system(['cp ' t2_defaced ' ' t2 ], '-echo'); <
130     end <
131     %%%%%%%%%% <
132     <

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133 %dcm2nii . %dcm2nii
134 if do_dcm2nii . if do_dcm2nii
135 if ~exist(set_path,'dir') . if ~exist(set_path,'dir')
[80 unmodified lines hidden]
216 acpc_wd = fullfile(set_path,'ACPCAlignment'); . acpc_wd = fullfile(set_path,'ACPCAlignment');
217 acpc_in = fullfile(set_path,sprintf('%sw_reoriented.nii.gz',datasets{set})); . acpc_in = fullfile(set_path,sprintf('%sw_reoriented.nii.gz',datasets{set}));
218 if set ==1 . if set ==1
219 acpc_tmp_in = fullfile(set_path,'t1mult_reoriented.nii.gz'); % ADDED < acpc_tmp_in = fullfile(set_path,'t1mult_reoriented.nii.gz');
220 acpc_ref = ref_t1_lmm; . acpc_ref = ref_t1_lmm;
221 elseif set ==2 . elseif set ==2
222 acpc_tmp_in = acpc_in; % ADDED BY ME !!!!! < acpc_tmp_in = acpc_in;
223 acpc_ref = fullfile(bpath,'analysis','MNI_t2w_templates','MNI152_T2_ . acpc_ref = fullfile(bpath,'analysis','MNI152_T2_2001.nii.gz');
224 end . end
225 acpc_out = fullfile(set_path,sprintf('%sw_acpc.nii.gz',datasets{set})(1:2 . acpc_out = fullfile(set_path,sprintf('%sw_acpc.nii.gz',datasets{set})(1:2));
[4 unmodified lines hidden]
230 robustroi=fullfile(acpc_wd,'robustroi.nii.gz'); . robustroi=fullfile(acpc_wd,'robustroi.nii.gz');
231 roi2full = fullfile(acpc_wd,'roi2full.mat'); . roi2full = fullfile(acpc_wd,'roi2full.mat');
232 cropcmd=sprintf('robustfov -i %s -m %s -r %s -b 170 -v',... . cropcmd=sprintf('robustfov -i %s -m %s -r %s -b 170 -v',...);
233 acpc_tmp_in, roi2full, robustroi); % MODIFIED BY ME (acpc_in -> x acpc_in, roi2full, robustroi);
234 fprintf('%s\n',cropcmd); . fprintf('%s\n',cropcmd);
235 system(cropcmd, '-echo'); . system(cropcmd, '-echo');
236 full2roi = fullfile(acpc_wd,'full2roi.mat'); . full2roi = fullfile(acpc_wd,'full2roi.mat');
[33 unmodified lines hidden]
270 system(multwarp, '-echo'); . system(multwarp, '-echo');
271 end . end
272 %QA . %QA
273 % zcmd = sprintf('fslview %s', robustroi); % MODIFIED BY ME (fslview x zcmd = sprintf('fslview %s', robustroi);
274 % system(zcmd); . system(zcmd);
275 % zcmd = sprintf('fsleyes %s %s %s', acpc_ref, acpc_final, acpc_out); x zcmd = sprintf('fsleyes %s %s %s', acpc_ref, acpc_final, acpc_out);
276 % system(zcmd); . system(zcmd);
277 end . end
278 .
279 %% initial brain extraction . %% initial brain extraction
[47 unmodified lines hidden]
327 fprintf('%s\n',mathcmd) . fprintf('%s\n',mathcmd);
328 system(mathcmd, '-echo'); . system(mathcmd, '-echo');
329 %QA . %QA
330 % zcmd = sprintf('fsleyes %s %s -l Red -t 0.5', acpc_out, be_outbrain); x zcmd = sprintf('fsleyes %s %s -l Red -t 0.5', acpc_out, be_outbrain);
331 % system(zcmd); . system(zcmd);
332 % zcmd = sprintf('fsleyes %s %s -t 0.5', be_ref2mm, acpc2MNI_lin); x zcmd = sprintf('fsleyes %s %s -t 0.5', be_ref2mm, acpc2MNI_lin);
333 % system(zcmd); . system(zcmd);
334 % zcmd = sprintf('fsleyes %s %s -t 0.5', acpc_ref, acpc2MNI_nonlin); x zcmd = sprintf('fsleyes %s %s -t 0.5', acpc_ref, acpc2MNI_nonlin);
335 % system(zcmd); . system(zcmd);
336 end . end
337 end . end
338 .
[43 unmodified lines hidden]
382 fprintf('%s\n',copy3cmd) . fprintf('%s\n',copy3cmd);
383 system(copy3cmd, '-echo'); . system(copy3cmd, '-echo');
384 conv3cmd=sprintf('convertwarp --relout --rel -r %s -w %s --postmat=%s -- . conv3cmd=sprintf('convertwarp --relout --rel -r %s -w %s --postmat=%s --');
385 t2_acpc_dc, t1_dc_transform, t2w2t1w_mat, t2_dc_transform); x t2_acpc_dc, t1_dc_transform, t2w2t1w_mat, t2_dc_transform);
386 fprintf('%s\n',conv3cmd) . fprintf('%s\n',conv3cmd);
387 system(conv3cmd, '-echo'); . system(conv3cmd, '-echo');
388 %QA ?????????????????? . %QA
389 % if n_vp==1 . if n_vp==1
390 % sl1 = [t1_acpc_dc ' ' t2_acpc_dc]; x sl1 = [t1_acpc_dc ' ' t2_acpc_dc];
391 % else . else
392 % sl1 = [sl1 ' ' t1_acpc_dc ' ' t2_acpc_dc]; x sl1 = [sl1 ' ' t1_acpc_dc ' ' t2_acpc_dc];
393 % end . end
394 % if n_vp==length(vps) . if n_vp==length(vps)
395 % cd(qa_path); . cd(qa_path);
396 % slname = 'slicesdir_t2t1reg'; . slname = 'slicesdir_t2t1reg';
397 % sl = sprintf('slicesdir -o %s', sl1); x sl = sprintf('slicesdir -o %s', sl1);
398 % fprintf('%s\n', sl); . fprintf('%s\n', sl);
399 % system(sl, '-echo'); . system(sl, '-echo');
400 % movefile(fullfile(qa_path,'slicesdir'), fullfile(qa_path,slname)); x movefile(fullfile(qa_path,'slicesdir'), fullfile(qa_path,slname));
401 % end . end
402 end . end
403 .
404 %% bias field correction based on sqrt T1w*T2w . %% bias field correction based on sqrt T1w*T2w
[91 unmodified lines hidden]
496 fprintf('%s\n',bfc4cmd) . fprintf('%s\n',bfc4cmd);
497 system(bfc4cmd, '-echo'); . system(bfc4cmd, '-echo');
498 %QA . %QA
499 % if n_vp == 1 . if n_vp == 1
500 % sl1 = [t1_acpc_dc_brain ' ' t1_acpc_dc_rest_brain]; x sl1 = [t1_acpc_dc_brain ' ' t1_acpc_dc_rest_brain];
501 % sl2 = [t2_acpc_dc ' ' t2_acpc_dc_rest]; x sl2 = [t2_acpc_dc ' ' t2_acpc_dc_rest];
502 % else . else
503 % sl1 = [sl1 ' ' t1_acpc_dc_brain ' ' t1_acpc_dc_rest_brain]; x sl1 = [sl1 ' ' t1_acpc_dc_brain ' ' t1_acpc_dc_rest_brain];
504 % sl2 = [sl2 ' ' t2_acpc_dc ' ' t2_acpc_dc_rest]; x sl2 = [sl2 ' ' t2_acpc_dc ' ' t2_acpc_dc_rest];
505 % end . end
506 % if n_vp==length(vps) . if n_vp==length(vps)
507 % cd(qa_path); . cd(qa_path);
508 % slname = {'slicesdir_t1_bfc', 'slicesdir_t2_bfc'}; x slname = {'slicesdir_t1_bfc', 'slicesdir_t2_bfc'};
509 % for slx = 1:length(slname) . for slx = 1:length(slname)
510 % if slx ==1 . if slx ==1
511 % sl = sprintf('slicesdir %s', sl1); x sl = sprintf('slicesdir %s', sl1);
512 % else . else
513 % sl = sprintf('slicesdir %s', sl2); x sl = sprintf('slicesdir %s', sl2);
514 % end . end
515 % system(sl, '-echo'); . system(sl, '-echo');
516 % movefile(fullfile(qa_path,'slicesdir'), fullfile(qa_path,slname)); x movefile(fullfile(qa_path,'slicesdir'), fullfile(qa_path,slname));
517 % end . end
518 % end . end
519 end . end
520 .
521 %% reg to MNI . %% reg to MNI
[24 unmodified lines hidden]
546 t1_acpc_dc_rest, ref_t1_2mm, acpc2mnilin, ref_t1_2mm_mask, acpc_dc2s . t1_acpc_dc_rest, ref_t1_2mm, acpc2mnilin, ref_t1_2mm_mask, acpc_dc2s;
547 fprintf('%s\n',mnireg2cmd) . fprintf('%s\n',mnireg2cmd);
548 system(mnireg2cmd, '-echo'); . system(mnireg2cmd, '-echo');
549 inv2cmd=sprintf('invwarp -w %s -o %s -r %s -v', ... . inv2cmd=sprintf('invwarp -w %s -o %s -r %s -v', ...);
550 acpc_dc2std, std2acpc_dc, ref_t1_2mm); . acpc_dc2std, std2acpc_dc, ref_t1_2mm);
551 fprintf('%s\n',inv2cmd) . fprintf('%s\n',inv2cmd);

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552      system(inv2cmd, '-echo');
553
554      [33 unmodified lines hidden]
555      system(mnimask2cmd, '-echo');
556
557      .
558      .
559      %QA
560      zcmd = sprintf('fslview %s %s', ref_t1_lmm, mni_t1_rest);
561      system(zcmd);
562      zcmd = sprintf('fslview %s %s', ref_t1_lmm, mni_t2_rest);
563      system(zcmd);
564
565      end
566
567      .
568      .
569      %% freesurfer
570
571      [5 unmodified lines hidden]
572      fs_nu = fullfile(fsmri_path, 'nu.mgz');
573      fs_t1 = fullfile(fsmri_path, 'T1.mgz');
574      fs_talairachskull = fullfile(fsmri_path, 'transforms', 'talairach_with_skull.lta');
575      % MODIFIED BY ME !!!!!!!
576      % {
577      fs_gca1 = fullfile(fs_path, 'average', 'RB_all_2008-03-26.gca');
578      fs_gca2 = fullfile(fs_path, 'average', 'RB_all_withskull_2008-03-26.gca');
579      %}
580      fs_gca1 = fullfile(fs_path, 'average', 'RB_all_2016-05-10.vc700.gca');
581      fs_gca2 = fullfile(fs_path, 'average', 'RB_all_withskull_2016-05-10.vc700.gca');
582
583      % ADDED BY ME !!!!!!!!!!!!! I'm trying to export the SUBJDIR path but it
584      subjDir_path = fullfile(bpath, 'freesurfer_subjects');
585      system(['mkdir -p ' subjDir_path], '-echo');
586      setenv('SUBJECTS_DIR', subjDir_path)
587      system('echo $SUBJECTS_DIR');
588      % ADDED BY ME !!!!!!!!!!!!!
589
590      .
591      .
592      if do_freesurfer1 % It takes ~200 seconds no parallelization
593      zcmd = sprintf('recon-all -i %s -subjID %s -motioncor -talairach -nuinte
594      t1_acpc_dc_rest, fs_subjID);
595      fprintf('%s\n', zcmd);
596      system(zcmd);
597
598      .
599      .
600      %if talairach transform fails: register t1 to mni manually
601      zcmd=sprintf('recon-all -i %s -subjID %s -motioncor -talairach -notalc
602      t1_acpc_dc_rest, fs_subjID);
603
604      [16 unmodified lines hidden]
605      system(zcmd);
606
607      .
608      end
609
610      .
611      if do_prefreesurfer2 % It takes ~ 400 seconds
612      %generate brain mask
613      tic
614      zcmd=sprintf('mri_convert %s %s --conform', ...
615      t1_acpc_dc_rest_brain, fs_brainmask); %--conform to lmm voxel size i
616      fprintf('%s\n', zcmd)
617      system(zcmd);
618
619      .
620      .
621      zcmd=sprintf('mri_em_register -mask %s %s %s %s', ...
622      fs_brainmask, fs_nu, fs_gca1, fs_talairachskull); % (added comment) cre
623      fprintf('%s\n', zcmd)
624      system(zcmd);
625
626      .
627      .
628      zcmd=sprintf('mri_watershed -T1 -brain atlas %s %s %s %s', ...
629      fs_gca2, fs_talairachskull, fs_t1, fs_brainmaskauto); % (added comment) x
630      fprintf('%s\n', zcmd)
631      system(zcmd);
632
633      .
634      .
635      toc
636      %if watershed error: T1 does not have enough wm values of 110, then pro
637      %-- add control points to nu.mgz and rerun normalization and watershed
638      controlDir = fullfile(fssub_path, fs_subjID, 'tmp');
639
640      [18 unmodified lines hidden]
641      fprintf('%s\n', zcmd)
642      system(zcmd);
643
644      .
645      .
646      .
647      % I UNCOMMENTED THIS BECAUSE THERE WHERE DURA LEFT IN THE BRAIN MASK AND
648      % THE SECOND STEP OF FREESURFER WAS FAILING (IT SEEMS TO WORK NOW)
649      %then if still parts of dura left can try
650      %option 1 gcut
651      %old brainmask file is saved in subj/trash
652      tic
653      zcmd=sprintf('recon-all -skullstrip -clean-bm -gcut -subjID %s', ...
654      fs_subjID);
655      fprintf('%s\n', zcmd)
656      system(zcmd);
657      %and check
658      zcmd=sprintf('fsleyes -v %s $SUBJECTS_DIR/%s/mri/brainmask.gcuts.mgz:co
659      old brainmaskauto, fs_subjID);
660      system(zcmd);
661      %option 2 if gcut fails then lower preflooding height of watershed
662      % (default 25) to 15
663      zcmd=sprintf('recon-all -skullstrip -wsthresh 15 -clean-bm -subjID %s'
664      fs_subjID);
665      fprintf('%s\n', zcmd)
666      system(zcmd);
667      %and check
668      zcmd=sprintf('freeview -v %s $SUBJECTS_DIR/%s/mri/brainmask.mgz:colorma
669      system(zcmd);
670      toc
671      % ADDED BY ME !!!
672      % if the segmentation is too aggressive use the following command
673      % to manually adjust the brain mask (see also
674      % https://surfer.nmr.mgh.harvard.edu/fswiki/FsTutorial/SkullStripFix_t
675      zcmd=sprintf('tkmedit %s brainmask.mgz -aux T1.mgz', ...
676      fs_subjID);
677      fprintf('%s\n', zcmd)
678      system(zcmd);
679
680      .
681      .
682      .
683      .
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719      .
720      .
721      .

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722                                     <
723                                     <
724     end                             .     end
725                                     .
726 if do_freesurfer2                   x   if do_freesurfer2
727     recon2cmd=sprintf('recon-all -subjID %s -autorecon2 -parallel -openmp %s' x       recon2cmd=sprintf('recon-all -subjID %s -a
728         fs_subjID, n_cores);          x       fs_subjID);
729     fprintf('%s\n',recon2cmd);         x       fprintf('%s\n',recon2cmd);
730     system(recon2cmd);                 x       system(recon2cmd);
731     %if big defects during fix topology check unfixed surfaces to see if sku x       %if big defects during fix topology check
732     hemi = 'xx';                      . %       hemi = 'xx';
733     zcmd=sprintf('freeview -v $SUBJECTS_DIR/%s/mri/T1.mgz -f $SUBJECTS_DIR, . %       zcmd=sprintf('freeview -v $SUBJECTS_DIR,
734     fs_subjID, fs_subjID);            . %       fs_subjID, fs_subjID);
735     system(zcmd);                     . %       system(zcmd);
736 end                                   x   end
737                                     .
738 if do_freesurfer3                   x   if do_freesurfer3
739     recon3cmd=sprintf('recon-all -subjID %s -T2 %s -T2pial -autorecon3 -paral x       recon3cmd=sprintf('recon-all -subjID %s -T
740         fs_subjID, t2_acpc_dc_rest, n_cores); x       fs_subjID, t2_acpc_dc_rest);
741     fprintf('%s\n',recon3cmd);         x       fprintf('%s\n',recon3cmd);
742     system(recon3cmd);                 x       system(recon3cmd);
743 end                                   x   end
744                                     .
745 %% segmentation and brain mask based on freesurfer output                    . %% segmentation and brain mask based on freesu
746 fsnatmask_path = fullfile(mask_path,'native','fs');                          . fsnatmask_path = fullfile(mask_path,'native',
[178 unmodified lines hidden]
925     fprintf('%s\n',zcmd)              .     fprintf('%s\n',zcmd)
926     system(zcmd,'-echo')               .     system(zcmd,'-echo')
927     %QA                                .     %QA
928     if n_vp==1                         x     if n_vp==1
929     sl = [t1_acpc_dc_rest_fsbrain ' ' bin_out]; x     sl = [t1_acpc_dc_rest_fsbrain ' '
930     else                                x     else
931     sl = [sl ' ' t1_acpc_dc_rest_fsbrain ' ' bin_out]; x     sl = [sl ' ' t1_acpc_dc_rest_fsbra
932     end                                  x     end
933     if n_vp==length(vps)                x     if n_vp==length(vps)
934     cd(qa_path);                        x     cd(qa_path);
935     slice = sprintf('slicesdir -o %s', sl); x     slice = sprintf('slicesdir -o %s',
936     fprintf('%s\n', slice);             x     fprintf('%s\n', slice);
937     system(slice, '-echo');              x     system(slice, '-echo');
938     movefile(fullfile(qa_path,'slicesdir'), fullfile(qa_path,sprint x     movefile(fullfile(qa_path,'slicesd
939     end                                  x     end
940 end                                     .     end
941                                     .
942 %create total_gm mask in 2mm native and mni space                          . %create total_gm mask in 2mm native and mri
[10 unmodified lines hidden]
953 %combine total gm and wm in 2mm mni space                                . %combine total gm and wm in 2mm mni space
954 allm = fullfile(fsmni2mmmask_path, 'allm.nii.gz');                      .     allm = fullfile(fsmni2mmmask_path, 'allm.n
955 totgm = fullfile(fsmni2mmmask_path, 'total_gm.nii.gz');                  .     totgm = fullfile(fsmni2mmmask_path, 'total
956 %wm = fullfile(fsmni2mmmask_path, 'wm_aseg.nii.gz');                    x     wm = fullfile(fsmni2mmmask_path, 'wm_aseg
957 wm = fullfile(fsmni2mmmask_path, 'wm.nii.gz'); % Modified by me !! wm_as <
958 zcmd = sprintf('fslmaths %s -add %s -bin %s', ...                        .     zcmd = sprintf('fslmaths %s -add %s -bin %
959 totgm, wm, allm);                                                         .     totgm, wm, allm);
960 fprintf('%s\n',zcmd)                                                       .     fprintf('%s\n',zcmd)
961 system(zcmd,'-echo')                                                       .     system(zcmd,'-echo')
962 end                                                                         .     end
963 end                                                                         .     end
964                                     .
965 <
966 <
967 %% create overlap mask for allm                                           . %% create overlap mask for allm
968 if do_overlap                                                              . if do_overlap
969     over_path = fullfile(nii_path,'overlap_masks');                        .     over_path = fullfile(nii_path,'overlap_masks'
[7 unmodified lines hidden]
977     if n_vp == 1                                                           .     if n_vp == 1
978     addmasks = vpmask;                                                     .     addmasks = vpmask;
979     else                                                                    .     else
980     addmasks = [addmasks ' -add ' vpmask];                                x     addmasks = [addmasks30 ' -add ' vpmask
981     end                                                                    .     end
982 end                                                                         .     end
983 over_out = fullfile(over_path, '30overlap_allm_erolvox.nii.gz');          .     over_out = fullfile(over_path, '30overlap_all
[8 unmodified lines hidden]

```

Number of matching lines: 798

Number of unmatched lines in left file: 193

Number of unmatched lines in right file: 109