

IN	inputnode	OUT
		flair
		roi
	utility	subject_id
		subjects_dir
		t1w
		t2w

IN	anat_template_wf	OUT
		outputnode.out_report
	inputnode.t1w	outputnode.t1w_realign_xfm
		outputnode.t1w_ref
		outputnode.t1w_valid_list

IN	anat_validate	OUT
	images	
	in_file	out_file

IN	brain_extraction_wf	OUT
		outputnode.bias_corrected
	inputnode.in_files	outputnode.out_file
		outputnode.out_segm

IN	surface_recon_wf	OUT
		outputnode.fsnative2t1w_xfm
		outputnode.out_aparc
		outputnode.out_aseg
		outputnode.out_brainmask
		outputnode.subject_id
		outputnode.subjects_dir
		outputnode.surfaces
		outputnode.t1w2fsnative_xfm

IN	applyrefined	OUT
	in_file	fsl
	mask_file	out_file

IN	buffernode	OUT
	t1w_brain	utility
	t1w_mask	t1w_brain
		t1w_mask

IN	t1w_dseg	OUT
	fsl	probability_maps
	in_files	tissue_class_map

IN	anat_norm_wf	OUT
		outputnode.anat2std_xfm
		outputnode.std2anat_xfm
		outputnode.template
		poutputnode.anat2std_xfm
		poutputnode.standardized
		poutputnode.std2anat_xfm
		poutputnode.std_dseg
		poutputnode.std_mask
		poutputnode.std_tpms
		poutputnode.template

IN	outputnode	OUT
		anat2std_xfm
		fsnative2t1w_xfm
		std2anat_xfm
		std_dseg
		std_mask
		std_preproc
		std_tpms
	utility	std_preproc
		std_tpms
		surfaces
		t1w2fsnative_xfm
		t1w_aparc
		t1w_aseg
		t1w_brain
		t1w_dseg
		t1w_mask
		t1w_preproc
		t1w_ref_xfms
		t1w_tpms
		template

IN	anat_derivatives_wf		OUT

IN	anat_reports_wf	OUT