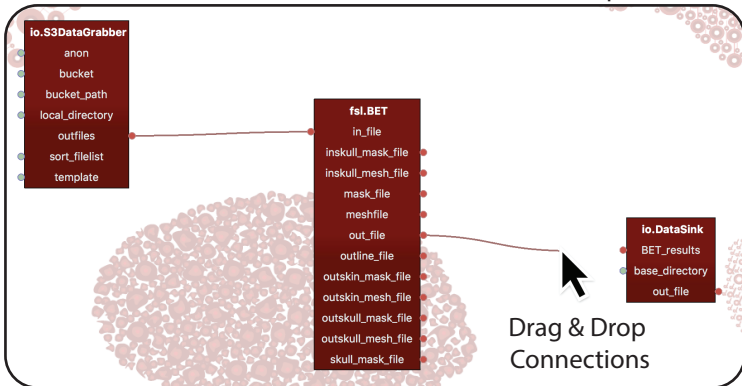


Workflow Editor

.pork file



Generated Code

.py file

```
#This is a NiType generator. Warning, here be dragons.
import nipype
import nipype.pipeline as pe
import nipype.interfaces.io as io
import nipype.interfaces.fsl as fsl

WorkingDirectory = "~/Porcupipelines/ThisStudy"

#Generic datagrabber module that wraps around glob in an
NodeHash_610000684510 = pe.Node(interface = io.S3DataGrabber(), name = 'NodeName_610000684510')
NodeHash_610000684510.inputs.anon = True
NodeHash_610000684510.inputs.bucket = 'openneuro'
NodeHash_610000684510.inputs.bucket_path = 'ds000101/ds000101_R2.0.0/uncompressed/'
NodeHash_610000684510.inputs.local_directory = '/tmp'
NodeHash_610000684510.inputs.sort_filelist = True
NodeHash_610000684510.inputs.template = 'sub-01/anat/sub-01_T1w.nii.gz'

#Wraps command **bet**
NodeHash_60800049f220 = pe.Node(interface = fsl.BET(), name = 'NodeName_60800049f220')

#Generic datasink module to store structured outputs
NodeHash_610000695fe0 = pe.Node(interface = io.DataSink(), name = 'NodeName_610000695fe0')
NodeHash_610000695fe0.inputs.base_directory = '/tmp'

#Create a workflow to connect all those nodes
analysisflow = nipype.Workflow('MyWorkflow')
analysisflow.connect(NodeHash_60800049f220, 'out_file', NodeHash_610000695fe0, 'BET_results')
analysisflow.connect(NodeHash_610000684510, 'outfiles', NodeHash_60800049f220, 'in_file')

#Run the workflow
analysisflow.run()
```