```
*****start of Matthias's email!
Greetings All,
Here's a short rundown on how the tau analysis works (starting from BEANs).
Potential caveat: you might need dbs access for this... and you should run on earth and the NDCMS condor pool.
1. Create a CMS release, e.g.,
  cmsrel CMSSW_5_3_11_patch3
  cd CMSSW_5_3_11_patch3/src
  cmsenv
        install
                     the
                               BEAN
                                            packages
                                                                              instructions
                                                                                                         [github]
                                                                    per
                                                                                                on
                                                            as
(https://github.com/cms-ttH/BEAN/blob/master/README.md).
3. install grid-control:
    cd ~; svn co https://ekptrac.physik.uni-karlsruhe.de/public/grid-control/trunk/grid-control; cd -
4. install the TauAnalysis package:
    cd $CMSSW BASE/src
    mkdir -p ttH
    git clone https://github.com/matz-e/TTHTauTau-Analysis.git ttH/TauAnalysis
    scram b -i32
5. fix paths (look for 'workdir' and 'se path' in ttH/TauAnalysis/test/grid control.cfg and run the configuration:
    cd ttH/TauAnalysis/test
    vim arid control.cfa
    voms-proxy-init -voms cms -valid 192:00 # this might not be needed?
    ~/grid-control/go.py -c grid control.cfg
 - wait until grid-control finishes
6. Analysis
Perform the installation steps on [github](https://github.com/cms-ttH/ttH-TauRoast), then fix the paths and run the
analysis:
  cd $LOCALRT/src/ttH/TauRoast/
  vim data/generic ttl.yaml
  # look and fix items under paths to where ntuples are stored, and the 'root' under which output will be stored
  roaster -atfpvy data/generic ttl.yaml
This will give you a yield table (how many events passed which cuts for each sample), and plots as specified in the
configuration.
*****end of Matthias's email!
#!/bin/tcsh
#May take X minutes to install and run
date
set CMSSW BASE = (CMSSW 5 3 11 patch3)
#set CMSSW_BASE = (CMSSW_5_3_8_patch1)
set finalOutput = (/afs/<u>crc.nd.edu/user/p/pivie/roast.out</u>)
#set ntupleFolder = ("/hadoop/users/matze/ttH/v53/2012/ttl_{p}/*_ntuple.root") #Use this if the TauRoast gets
updated again
set ntupleFolder = ("/hadoop/users/matze/ttH/v53/2012/ttl_{p}")
module load git
cmsrel $CMSSW BASE
cd ~/$CMSSW BASE
cd src
cmsenv
setenv CVSROOT :pserver:anonymous@cmssw.cvs.cern.ch:/local/reps/CMSSW
cvs login
cvs co -d PhysicsTools/NtupleUtils UserCode/Bicocca/PhysicsTools/NtupleUtils
cvs co -d SusyAnalysis -r V01-02-02 SusyAnalysis/EventSelector/interface/uncorrectionTypeMET.h
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cvs co -r V07-00-01 TopQuarkAnalysis/Configuration
cvs co -r V06-07-11-01 TopQuarkAnalysis/TopTools
cvs co -r V06-05-06-07 DataFormats/PatCandidates
cvs co -r V00-02-14 DataFormats/StdDictionaries
cvs co -r V00-00-70 FWCore/GuiBrowsers
cvs co -r V08-09-52 PhysicsTools/PatAlgos
cvs co -r V03-09-23 PhysicsTools/PatUtils
cvs co -r V00-03-15 CommonTools/ParticleFlow
cvs co -r V01-08-00 RecoBTag/SecondaryVertex
cvs co -r V00-00-08 RecoMET/METAnalyzers
cvs co -r V00-00-07 RecoMET/METFilters
cvs co -r V15-01-11 RecoParticleFlow/PFProducer
cvs co -r V02-02-00 RecoVertex/AdaptiveVertexFinder
cvs co -d Muon/MuonAnalysisTools UserCode/sixie/Muon/MuonAnalysisTools
cvs co -r V00-00-13 -d EGamma/EGammaAnalysisTools UserCode/EGamma/EGammaAnalysisTools
cd EGamma/EGammaAnalysisTools/data
cat download.url | xargs wget
cd -
cd EGamma/EGammaAnalysisTools/interface/
wget -r http://nd.edu/~abrinke1/ElectronEffectiveArea.h -O ElectronEffectiveArea.h
cvs co -r V01-04-23 RecoTauTag/RecoTau
cvs co -r V01-04-10 RecoTauTag/Configuration
cvs co -r V00-04-00 CondFormats/EgammaObjects
cvs co -r V00-02-10 -d CMGTools/External UserCode/CMG/CMGTools/External
git clone https://github.com/cms-ttH/BEAN.git
scram b -j32
wget -O - http://pyyaml.org/download/pyyaml/PyYAML-3.10.tar.gz|tar xzf -
cd PyYAML-3.10/
python setup.py install --user
cd ..
rm -rf PyYAML-3.10
cd ~/$CMSSW_BASE
cd src
mkdir -p ttH
git clone https://github.com/cms-ttH/ttH-TauRoast.git ttH/TauRoast
scram b -j32
git clone https://github.com/cms-ttH/ttHMultileptonAnalysis.git
cd ttH
cd TauRoast
git checkout 351767b174f301eb1e13995a7023b3aad00dfdf5
mv data/generic_ttl.yaml data/generic_ttl.old.yaml
cat data/generic_ttl.old.yaml | sed "s,^\([]*root\: \).*,\1$finalOutput,g" | sed "s,^\([]*ntuples\: \).*,\1$ntupleFolder,g" >
data/generic_ttl.yaml
scram b clean; scram b
# look and fix items under paths to where ntuples are stored, and the 'root' under which output will be stored
echo "Installation complete:"
date
scripts/roaster -atfpvy data/generic_ttl.yaml
```

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The script of the third solution
#!/bin/tcsh
#May take X minutes to install and run
echo "<1>the start of python building:"
date
set CMSSW_BASE = (CMSSW_5_3_11_patch3)
set finalOutput = (/afs/crc.nd.edu/user/p/pivie/roast.out)
set ntupleFolder = ("/hadoop/users/matze/ttH/v53/2012/ttl_{p}")
set PARROT HELPER = '/afs/crc.nd.edu/user/h/hmeng/cctools/lib/$LIB/libparrot helper.so'
setenv
cmsenv
cd ~/$CMSSW BASE
wget -O - http://pyyaml.org/download/pyyaml/PyYAML-3.10.tar.gz|tar xzf -
cd PyYAML-3.10/
cmsenv
python setup.py install --user
cd -
rm -rf PyYAML-3.10
echo "<1>the end of python building:"
echo "<2>the start of analysis"
date
cd ttH
cd TauRoast
scripts/roaster -atfpvy data/generic ttl.yaml
echo "<2>the end of analysis"
date
```

#declaration of environment variables #clear data dependency #clear software dependency #software build and install command #actual analysis program