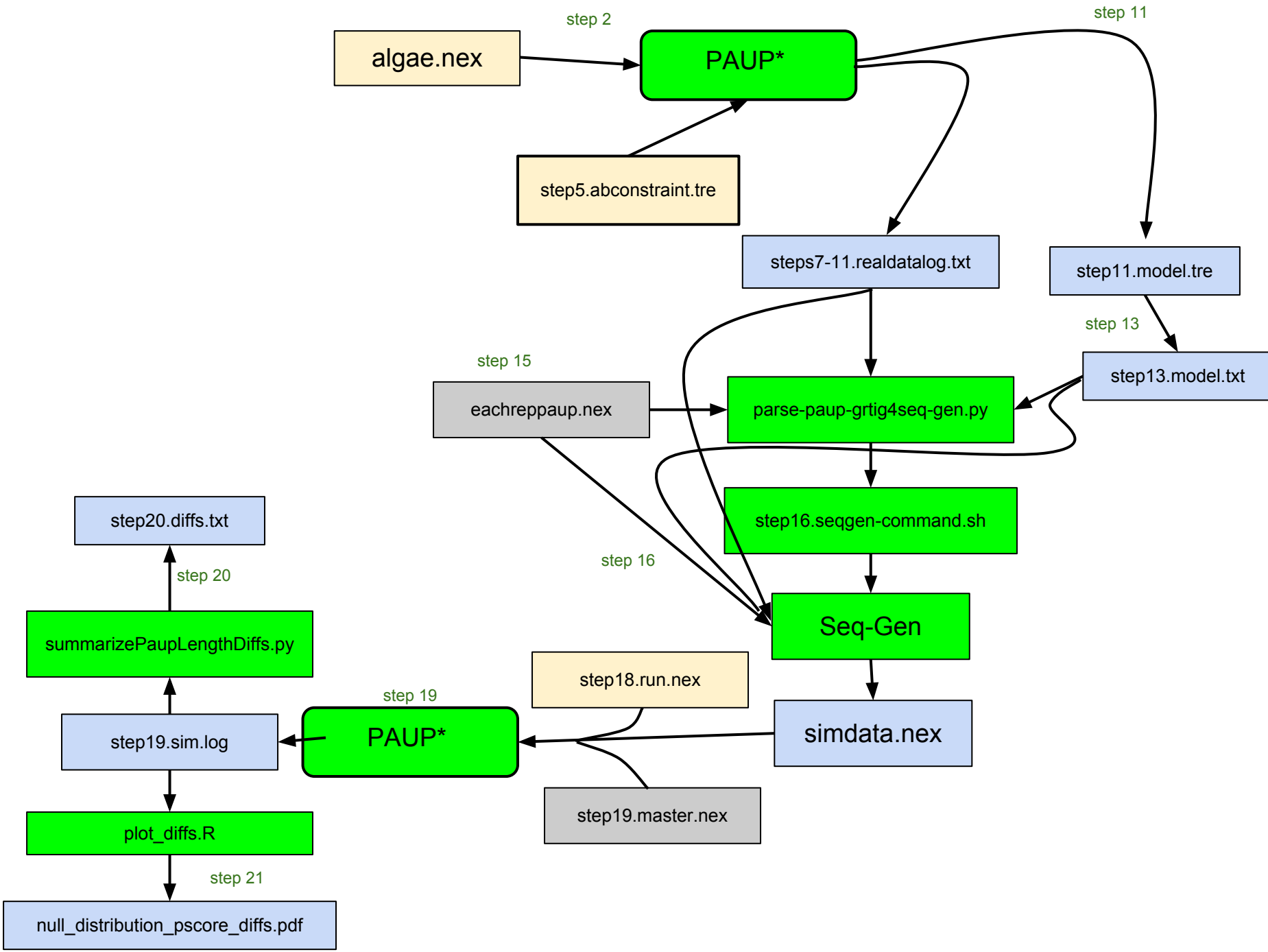
step 2
algae.nex
PAUP*
step 11
step5.abconstraint.tre
steps7-11.realdatalog.txt
step11.model.tre
step 13
step13.model.txt
step 15
eachreppaup.nex
parse-paup-grtig4seq-gen.py
step16.seqgen-command.sh
step 16
Seq-Gen
step20.diffs.txt
step 20
summarizePaupLengthDiffs.py
step18.run.nex
step 19
PAUP*
simdata.nex
step19.sim.log
step19.master.nex
plot_diffs.R
step 21
null_distribution_pscore_diffs.pdf