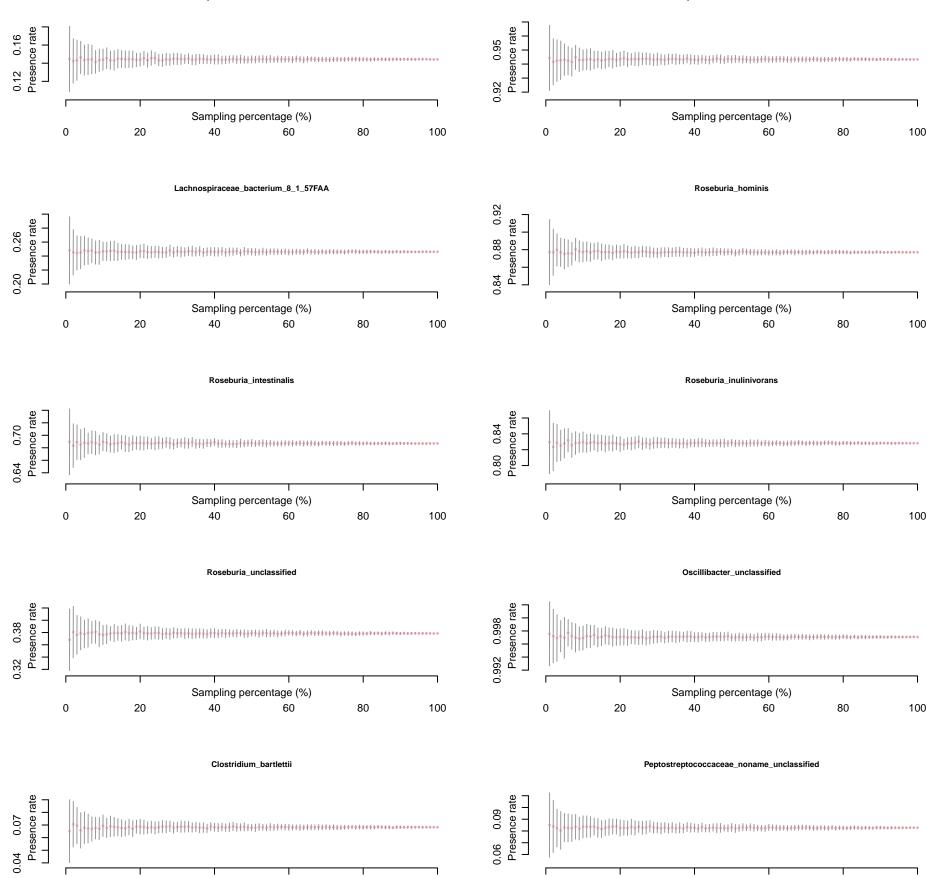
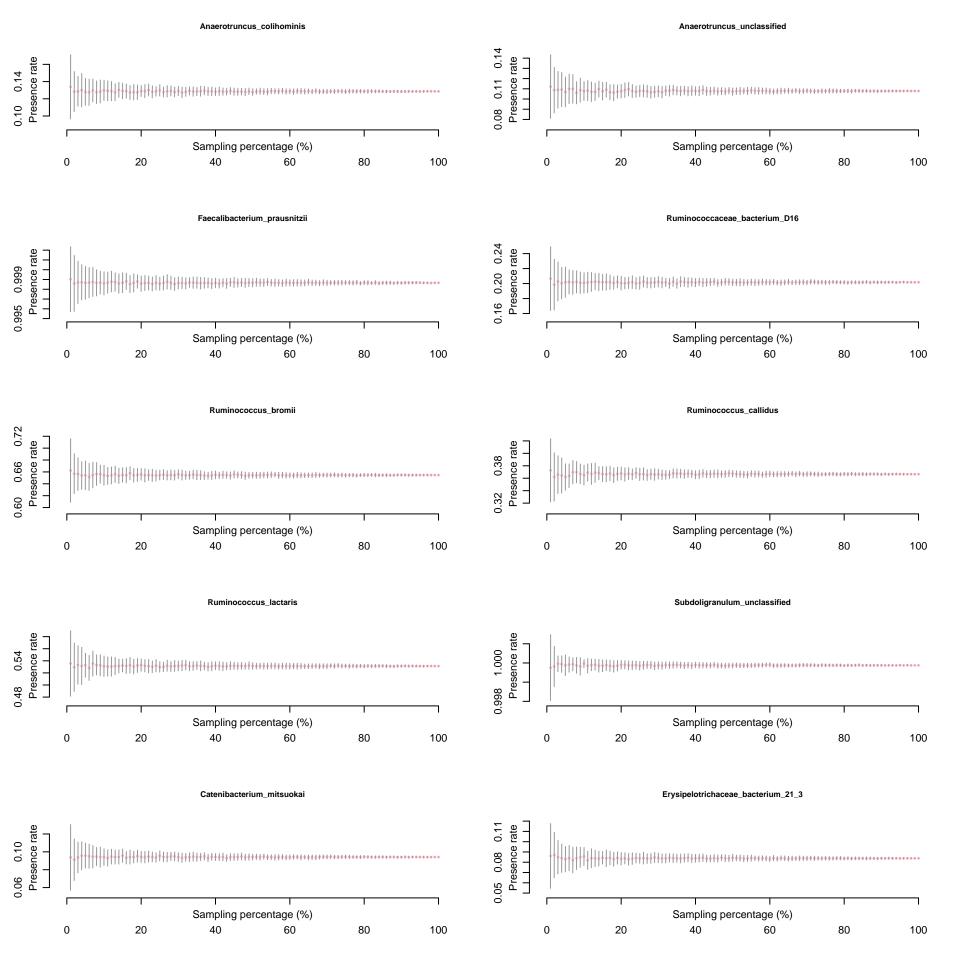
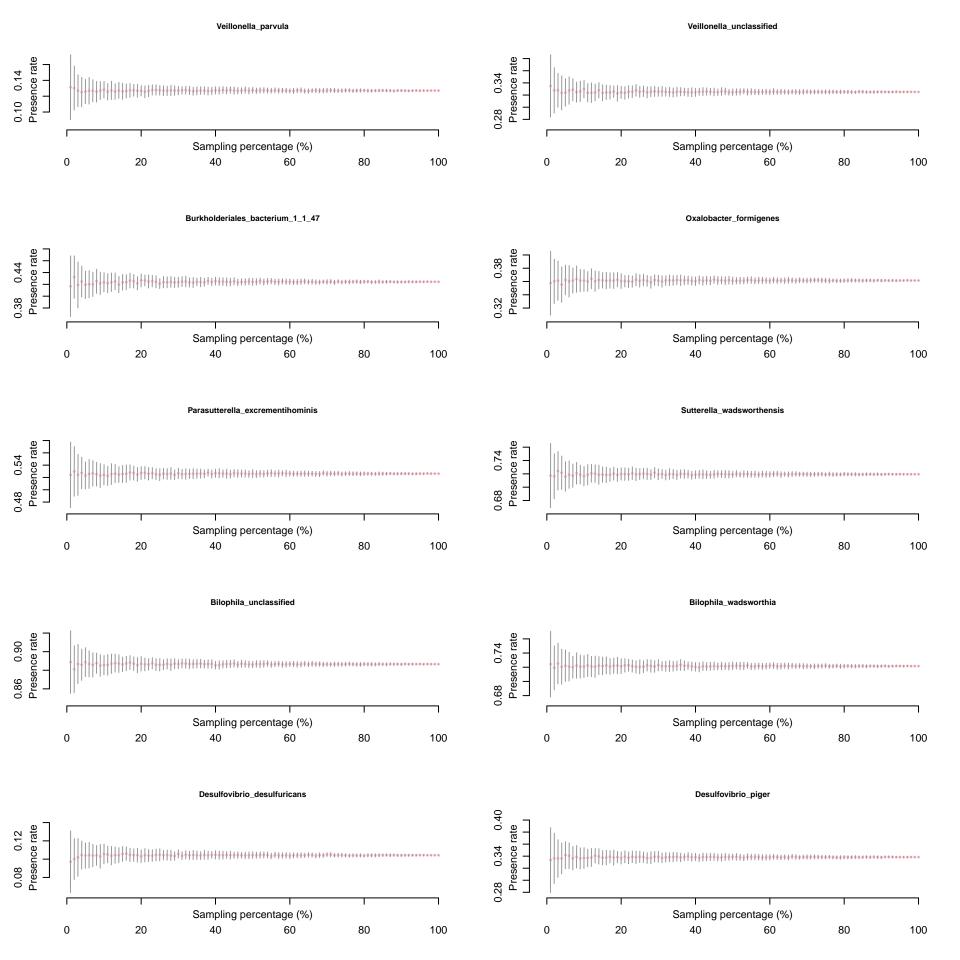


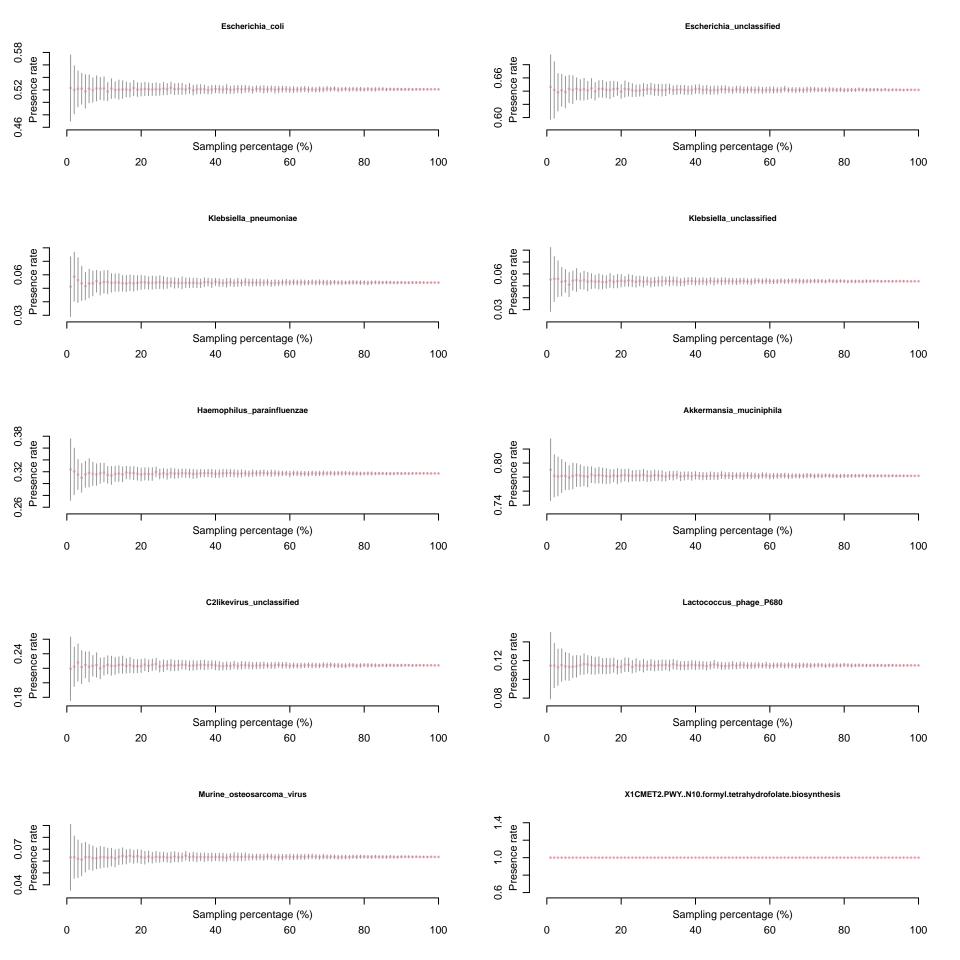
Sampling percentage (%)

Sampling percentage (%)

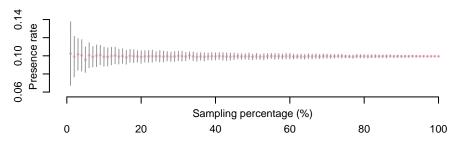


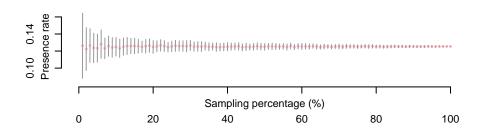




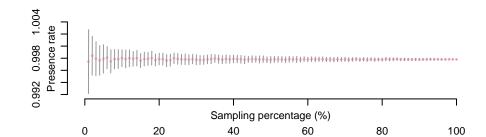


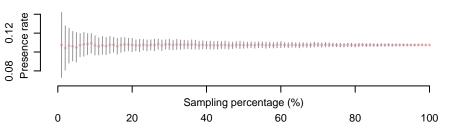
ANAEROFRUCAT.PWY..homolactic.fermentation



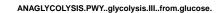


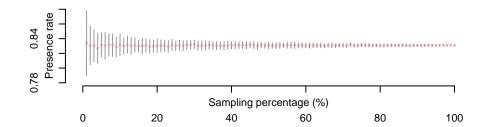
ALL.CHORISMATE.PWY..superpathway.of.chorismate.metabolism

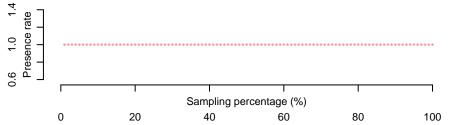






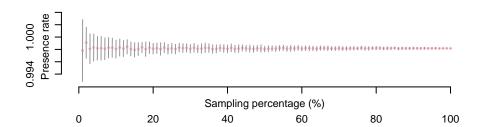


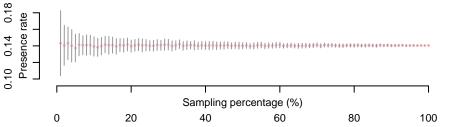






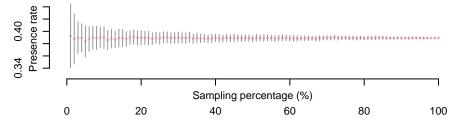
ARGDEG.PWY..superpathway.of.L.arginine..putrescine..and.4.aminobutanoate.degradation

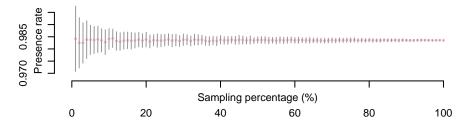


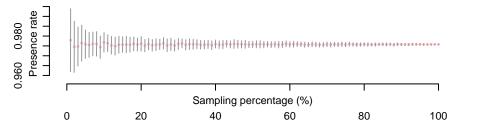


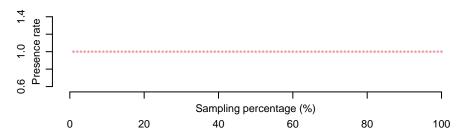
ARGORNPROST.PWY..arginine..ornithine.and.proline.interconversion

ARGSYN.PWY..L.arginine.biosynthesis.l..via.L.ornithine.



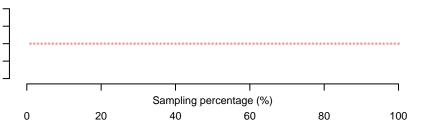




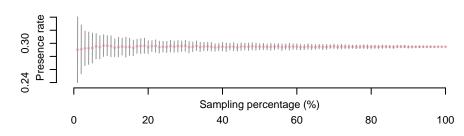


ASPASN.PWY..superpathway.of.L.aspartate.and.L.asparagine.biosynthesis

AST.PWY..L.arginine.degradation.II..AST.pathway.

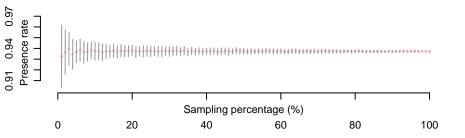


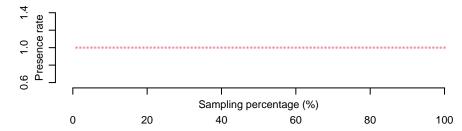
0.6 1.0 1.4 Presence rate



BIOTIN.BIOSYNTHESIS.PWY..biotin.biosynthesis.I

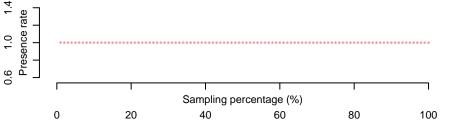
BRANCHED.CHAIN.AA.SYN.PWY..superpathway.of.branched.amino.acid.biosynthesis

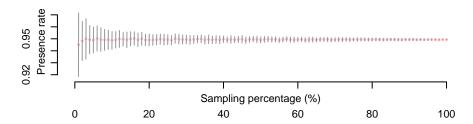




CALVIN.PWY..Calvin.Benson.Bassham.cycle

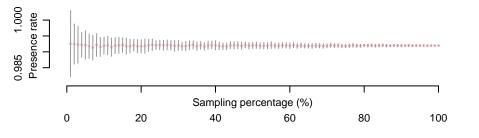
CENTFERM.PWY..pyruvate.fermentation.to.butanoate

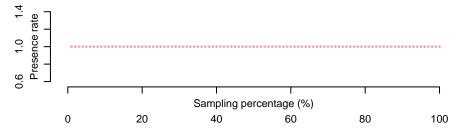


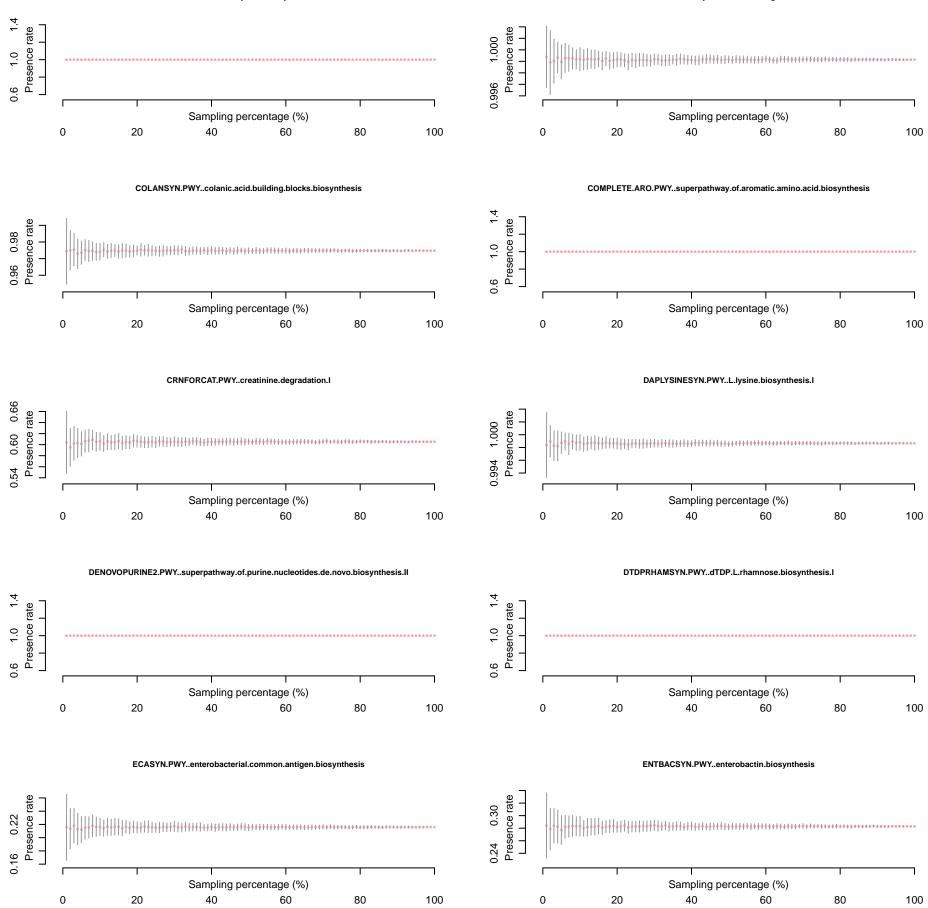


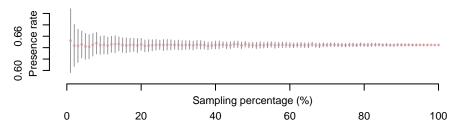
CITRULBIO.PWY..L.citrulline.biosynthesis

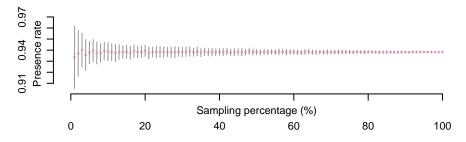
COA.PWY.1..coenzyme.A.biosynthesis.II..mammalian.





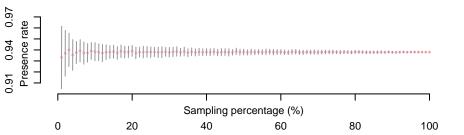


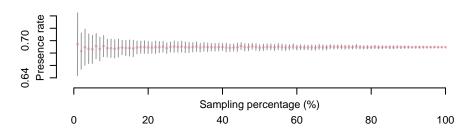




FASYN.INITIAL.PWY..superpathway.of.fatty.acid.biosynthesis.initiation..E..coli.

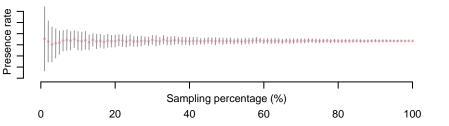
FERMENTATION.PWY..mixed.acid.fermentation



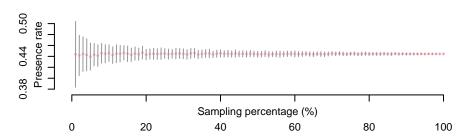


FOLSYN.PWY..superpathway.of.tetrahydrofolate.biosynthesis.and.salvage

FUC.RHAMCAT.PWY..superpathway.of.fucose.and.rhamnose.degradation

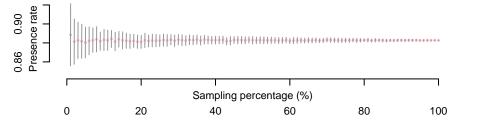


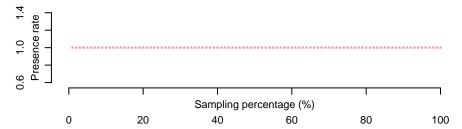
0.58 0.64 0.70



FUCCAT.PWY..fucose.degradation

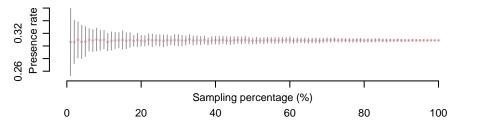
GALACT.GLUCUROCAT.PWY..superpathway.of.hexuronide.and.hexuronate.degradation

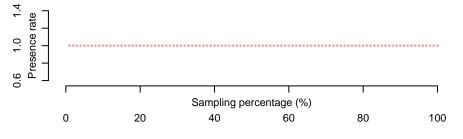


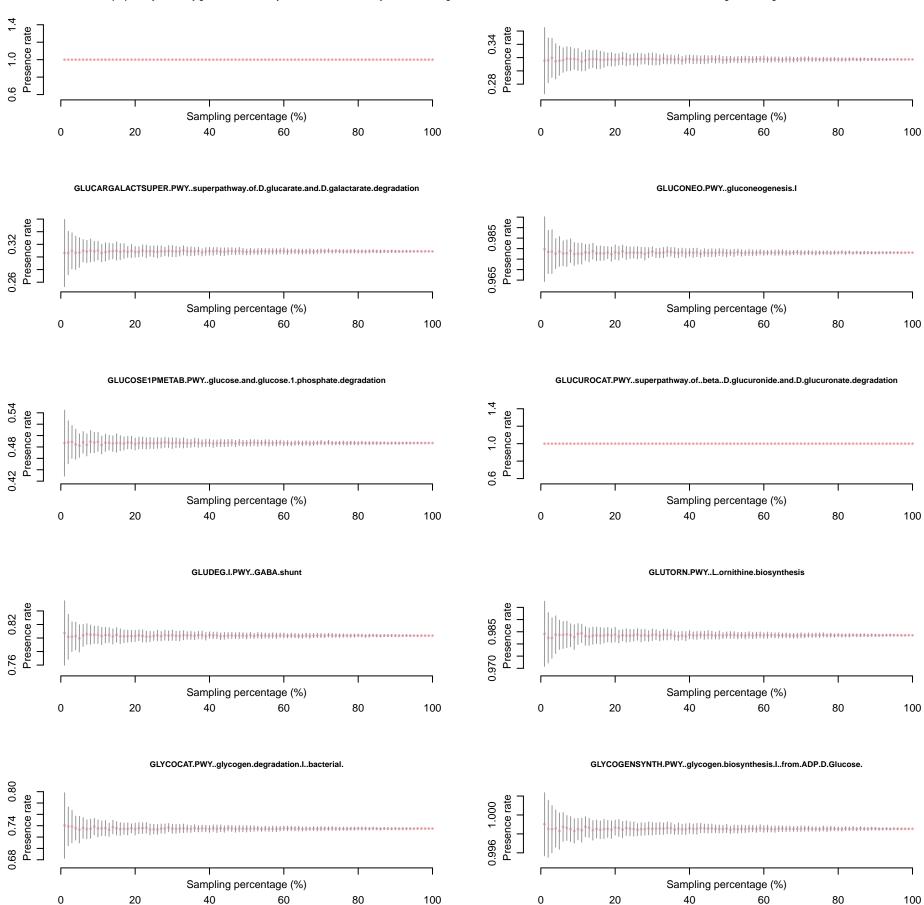


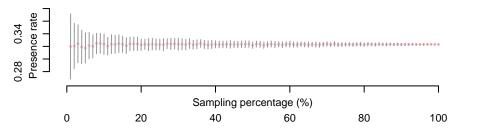
GALACTARDEG.PWY..D.galactarate.degradation.I

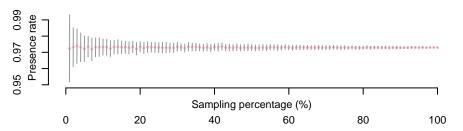
GALACTUROCAT.PWY..D.galacturonate.degradation.l



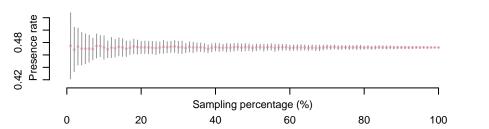


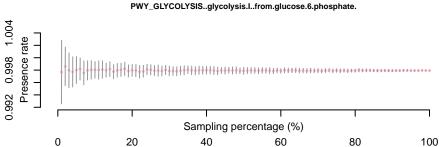




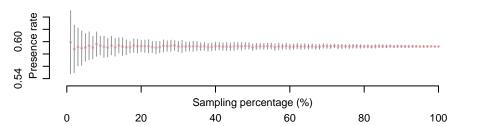


 $PWY_GLYCOLYSIS.TCA.GLYOX.BYPASS..superpathway.of.glycolysis..pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.dehydrogenase..TCA..and.glyoxylate.bypacture.pyruvate.pyr$

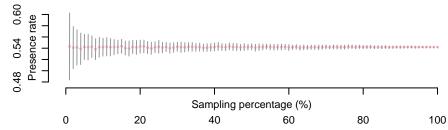




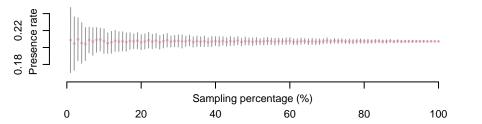
PWY_GLYOXYLATE.BYPASS..glyoxylate.cycle



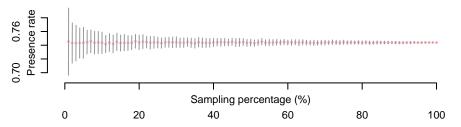
GOLPDLCAT.PWY..superpathway.of.glycerol.degradation.to.1.3.propanediol



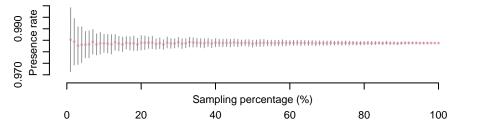
HCAMHPDEG. PWY..3. phenyl propanoate. and. 3..3. hydroxyphenyl. propanoate. degradation. to. 2. oxopent. 4. enoate the propanoate of the



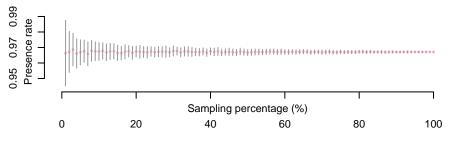
PWY_HEME.BIOSYNTHESIS.II..heme.biosynthesis.I..aerobic.



HEMESYN2.PWY..heme.biosynthesis.II..anaerobic.

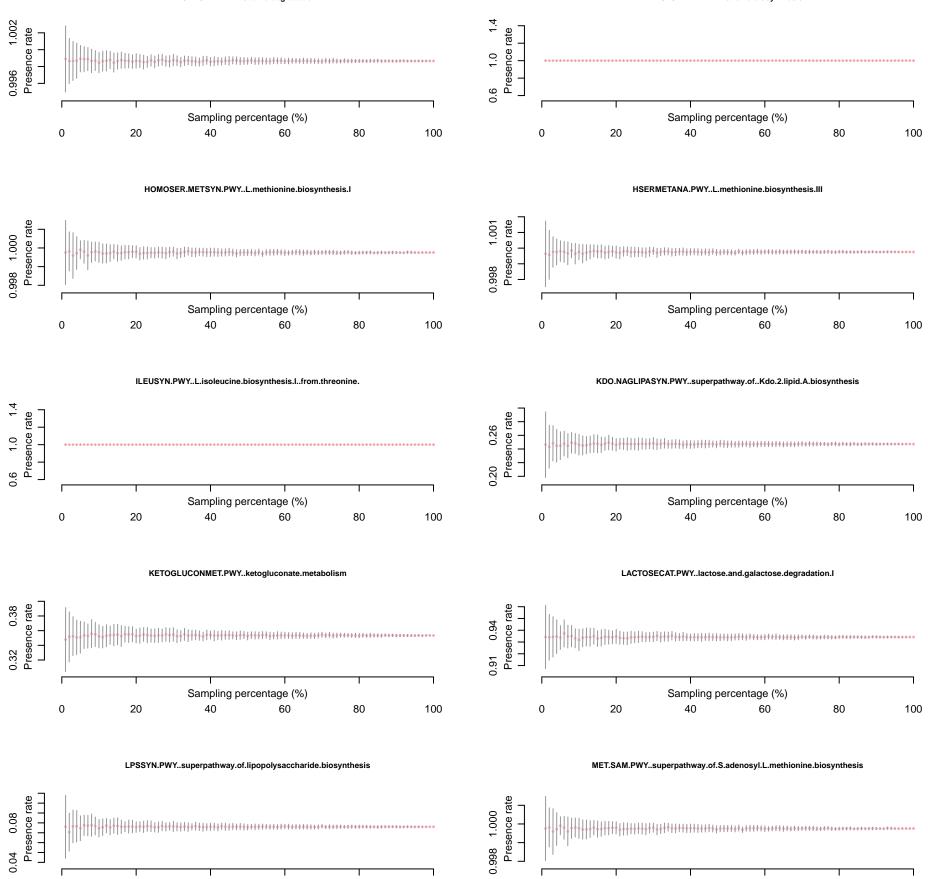


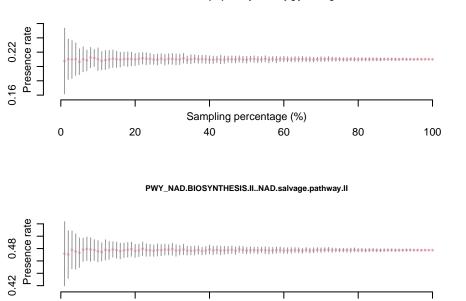
 $\label{lem:hexitoldegsuper.PWY} HEXITOLDEGSUPER.PWY..superpathway.of. hexitol. degradation.. bacteria. A superpathway of the superpathway of the$

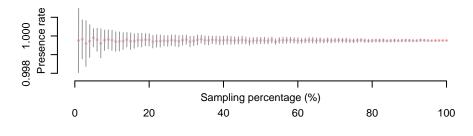


Sampling percentage (%)

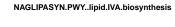
Sampling percentage (%)

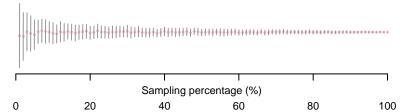


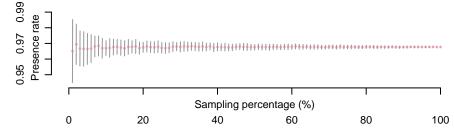




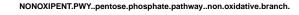


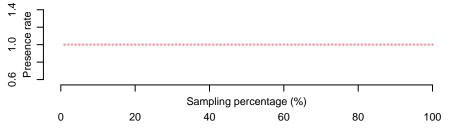


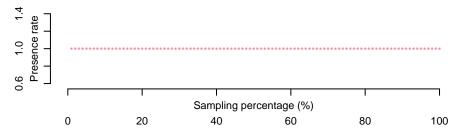




${\bf NONMEVIPP.PWY..} methylery thritol.phosphate.pathway. I$

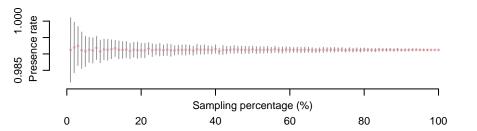


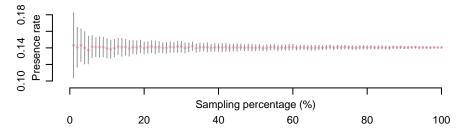




OANTIGEN.PWY..O.antigen.building.blocks.biosynthesis..E..coli.

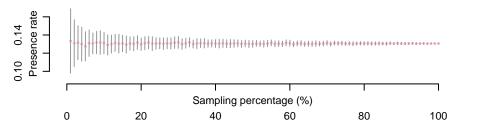
ORNARGDEG.PWY..superpathway.of.L.arginine.and.L.ornithine.degradation

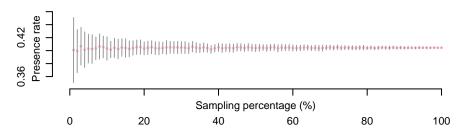


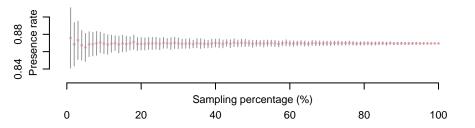


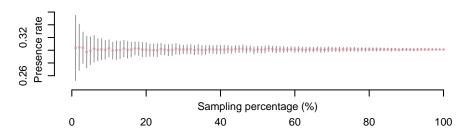
ORNDEG.PWY..superpathway.of.ornithine.degradation

P105.PWY..TCA.cycle.IV..2.oxoglutarate.decarboxylase



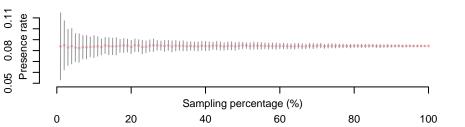


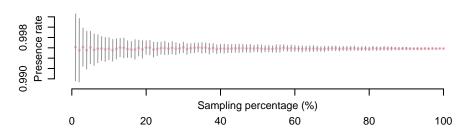






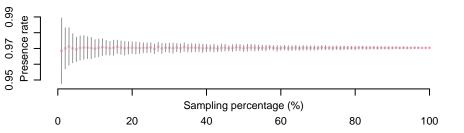
P161.PWY..acetylene.degradation

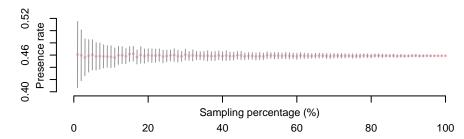




P162.PWY..L.glutamate.degradation.V..via.hydroxyglutarate.

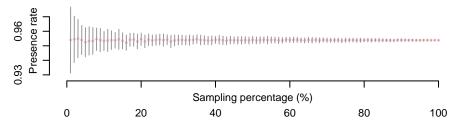
P164.PWY..purine.nucleobases.degradation.l..anaerobic.

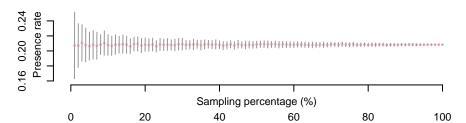




P185.PWY..formaldehyde.assimilation.lll..dihydroxyacetone.cycle.

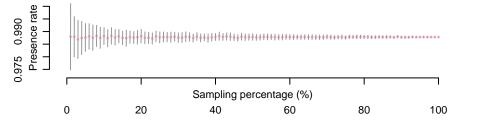
P23.PWY..reductive.TCA.cycle.l

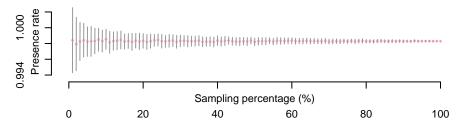


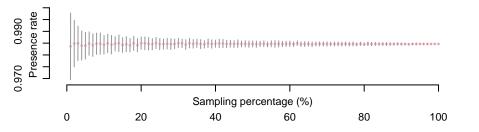


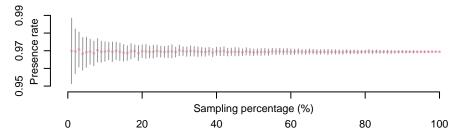
P4.PWY..superpathway.of.L.lysine..L.threonine.and.L.methionine.biosynthesis.l

P42.PWY..incomplete.reductive.TCA.cycle



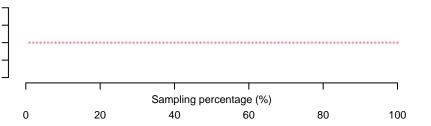






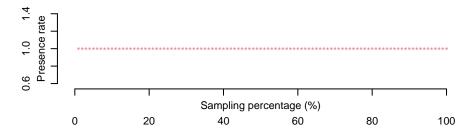


PANTOSYN.PWY..pantothenate.and.coenzyme.A.biosynthesis.I



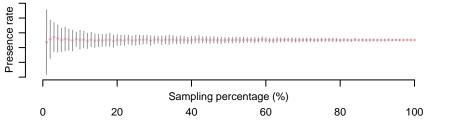
0.6 1.0 1.4 Presence rate

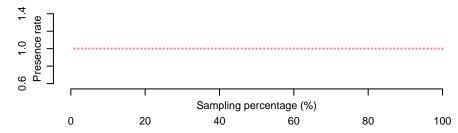
0.980 0.995



PENTOSE.P.PWY..pentose.phosphate.pathway

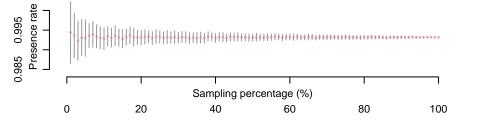
PEPTIDOGLYCANSYN.PWY..peptidoglycan.biosynthesis.l..meso.diaminopimelate.containing.

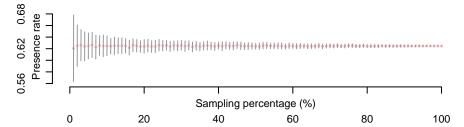




PHOSLIPSYN.PWY..superpathway.of.phospholipid.biosynthesis.l..bacteria.

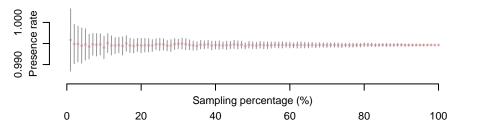
POLYAMINSYN3.PWY..superpathway.of.polyamine.biosynthesis.ll

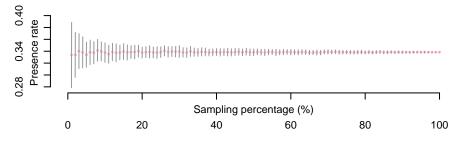


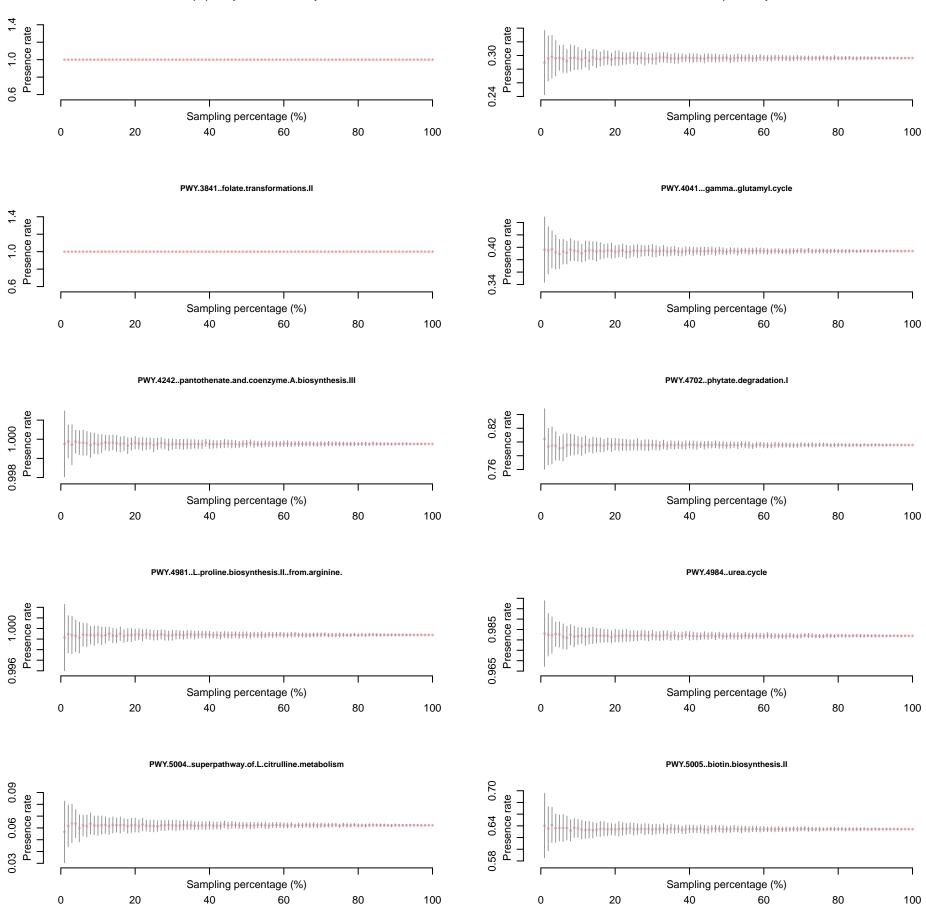


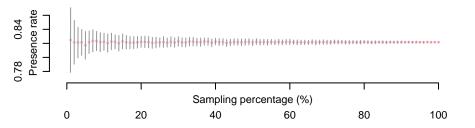
POLYAMSYN.PWY..superpathway.of.polyamine.biosynthesis.I

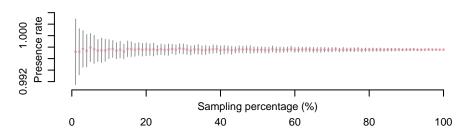
POLYISOPRENSYN.PWY..polyisoprenoid.biosynthesis..E..coli.





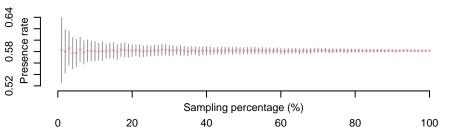


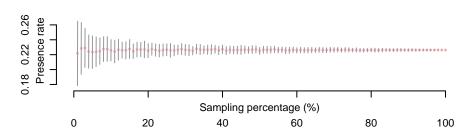






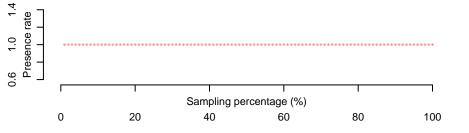
PWY.5088..L.glutamate.degradation.VIII..to.propanoate.

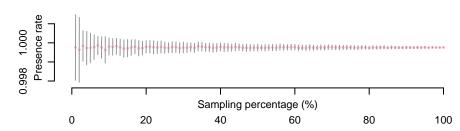




PWY.5097..L.lysine.biosynthesis.VI

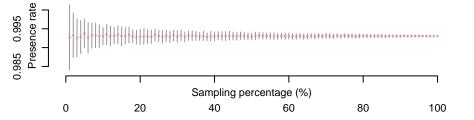
PWY.5100..pyruvate.fermentation.to.acetate.and.lactate.ll

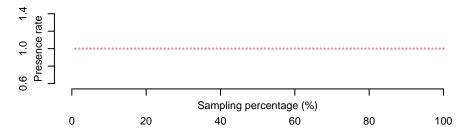




PWY.5101..L.isoleucine.biosynthesis.ll

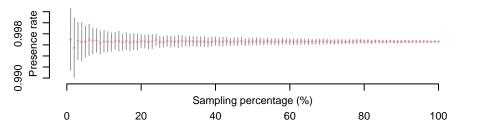
PWY.5103..L.isoleucine.biosynthesis.III

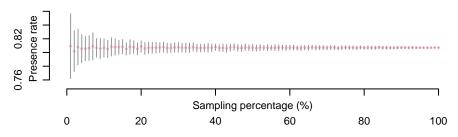


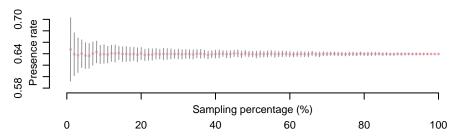


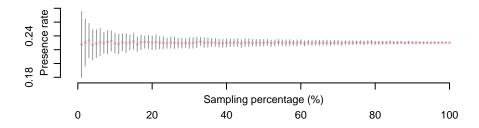
PWY.5104..L.isoleucine.biosynthesis.IV

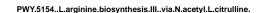
PWY.5121..superpathway.of.geranylgeranyl.diphosphate.biosynthesis.ll..via.MEP.

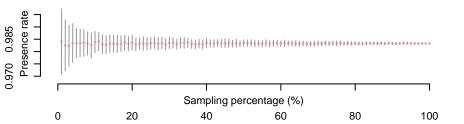




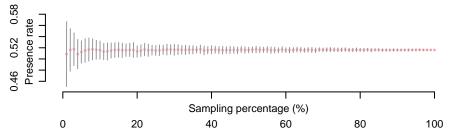




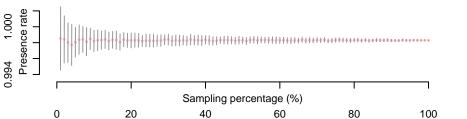






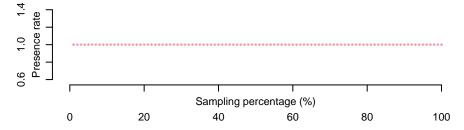


PWY.5177..glutaryl.CoA.degradation

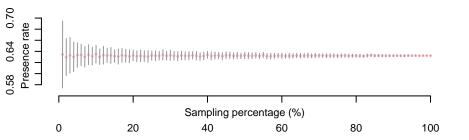


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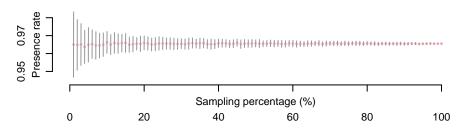
PWY.5188..tetrapyrrole.biosynthesis.l..from.glutamate.



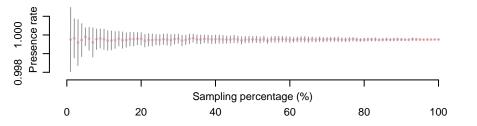
PWY.5189..tetrapyrrole.biosynthesis.ll..from.glycine.



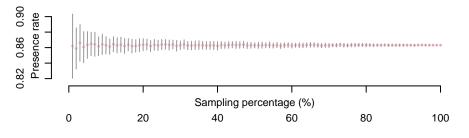
PWY.5345...superpathway.of.L.methionine.biosynthesis...by.sulfhydrylation.

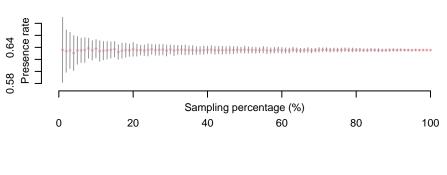


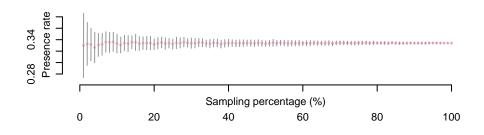
PWY.5347..superpathway.of.L.methionine.biosynthesis..transsulfuration.



PWY.5367..petroselinate.biosynthesis

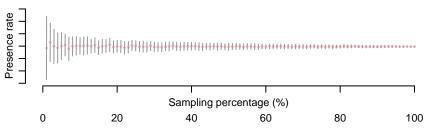




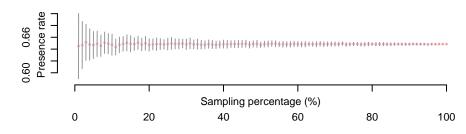




PWY.5505..L.glutamate.and.L.glutamine.biosynthesis

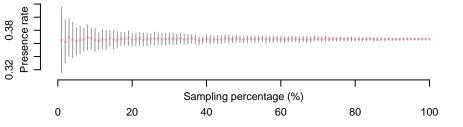


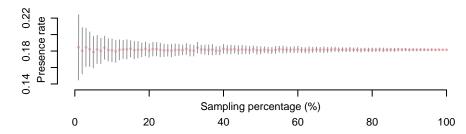
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PWY.561..superpathway.of.glyoxylate.cycle.and.fatty.acid.degradation

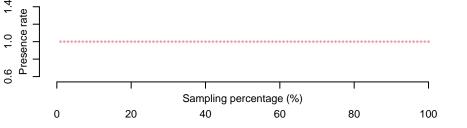
PWY.5656..mannosylglycerate.biosynthesis.l

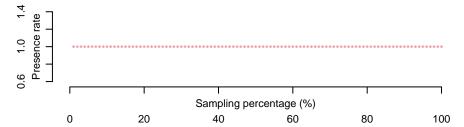




PWY.5659..GDP.mannose.biosynthesis

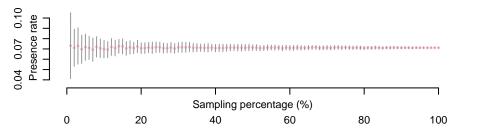
PWY.5667..CDP.diacylglycerol.biosynthesis.l

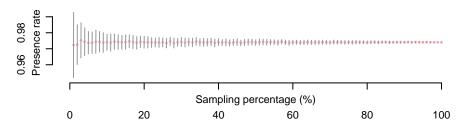


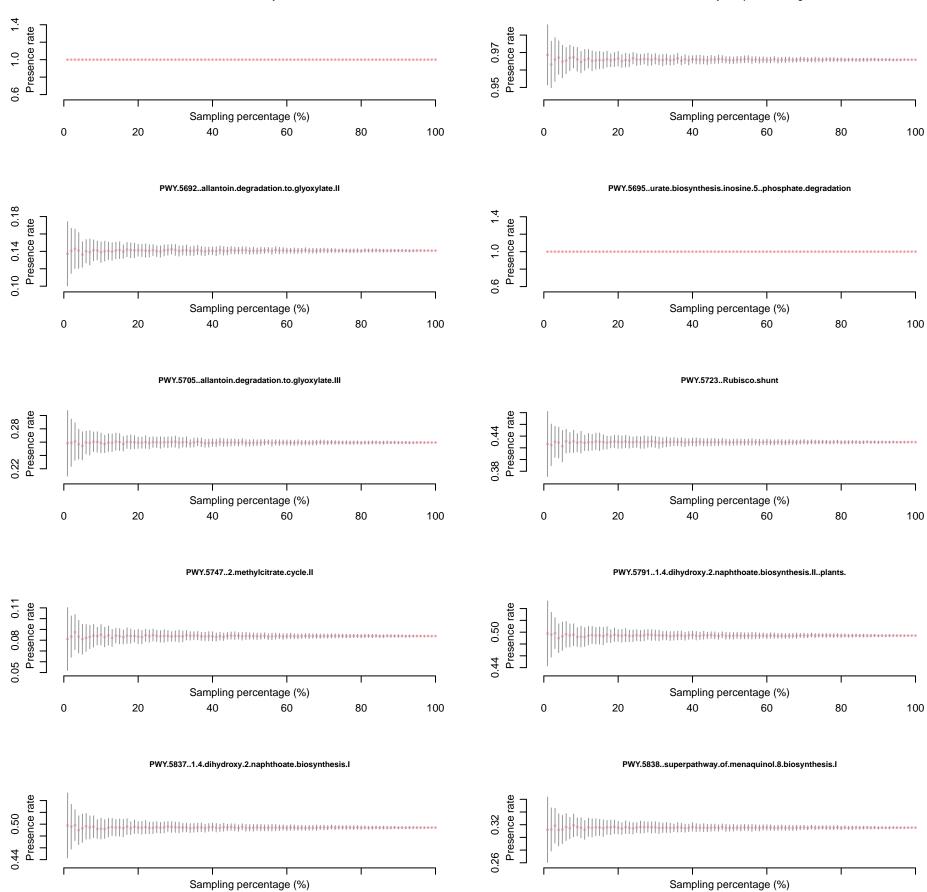


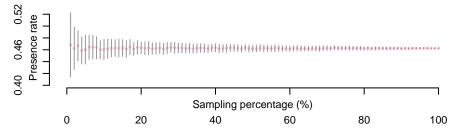
PWY.5675..nitrate.reduction.V..assimilatory

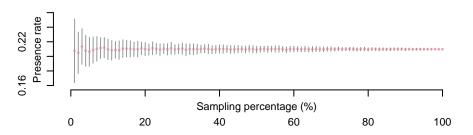
PWY.5676..acetyl.CoA.fermentation.to.butanoate.II





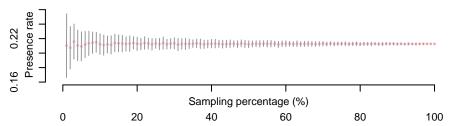


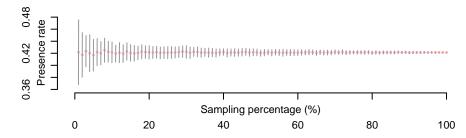






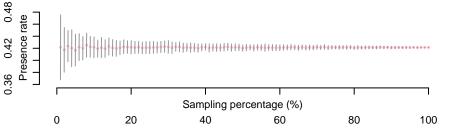
PWY.5855..ubiquinol.7.biosynthesis..prokaryotic.

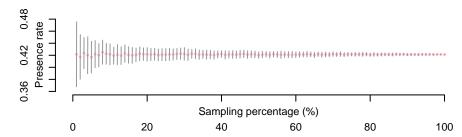




PWY.5856..ubiquinol.9.biosynthesis..prokaryotic.

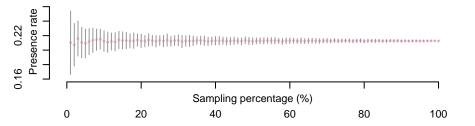
PWY.5857..ubiquinol.10.biosynthesis..prokaryotic.

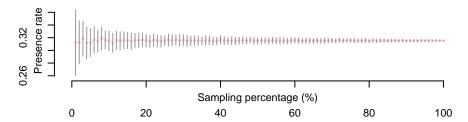




PWY.5860..superpathway.of.demethylmenaquinol.6.biosynthesis.l

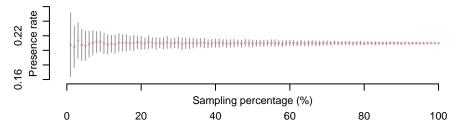
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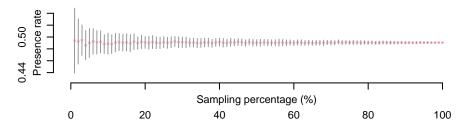


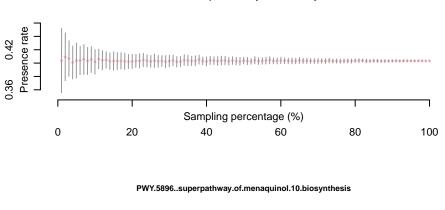


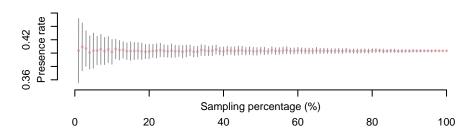
PWY.5862..superpathway.of.demethylmenaquinol.9.biosynthesis

PWY.5863..superpathway.of.phylloquinol.biosynthesis

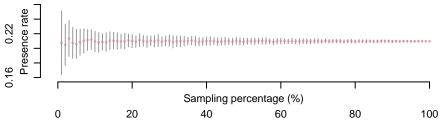


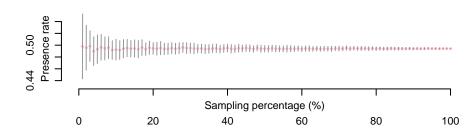






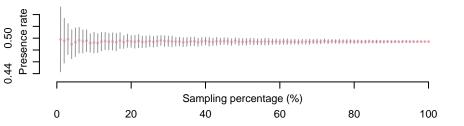
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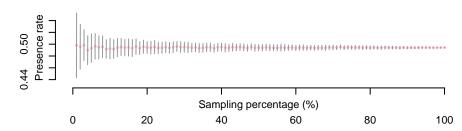




PWY.5898..superpathway.of.menaquinol.12.biosynthesis

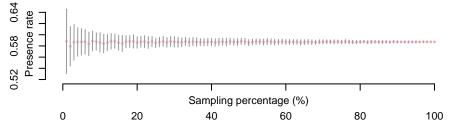
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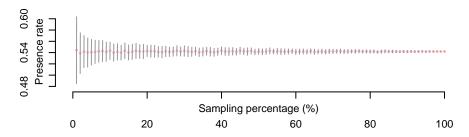




PWY.5913..TCA.cycle.VI..obligate.autotrophs.

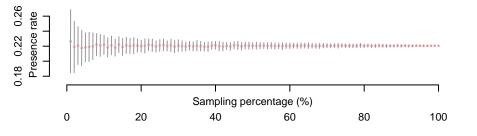
PWY.5918..superpathay.of.heme.biosynthesis.from.glutamate

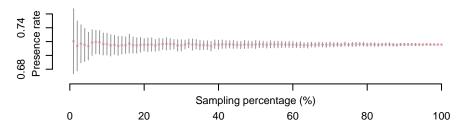


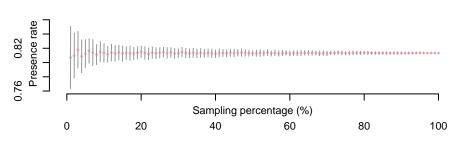


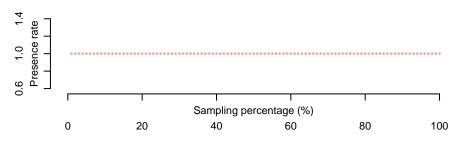
PWY.5920..superpathway.of.heme.biosynthesis.from.glycine

PWY.5941..glycogen.degradation.ll..eukaryotic.



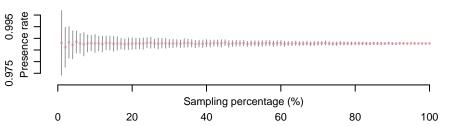


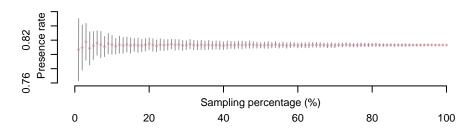




PWY.5989..stearate.biosynthesis.ll..bacteria.and.plants.

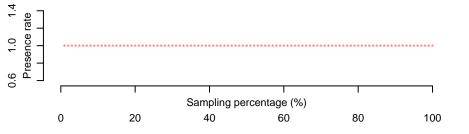
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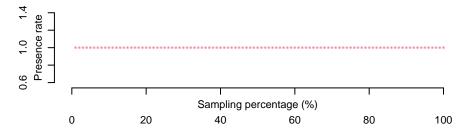




PWY.6121..5.aminoimidazole.ribonucleotide.biosynthesis.l

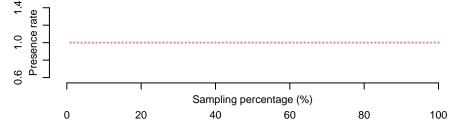
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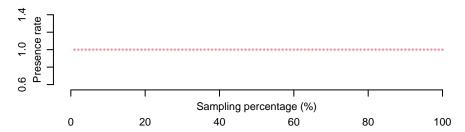




PWY.6123..inosine.5..phosphate.biosynthesis.l

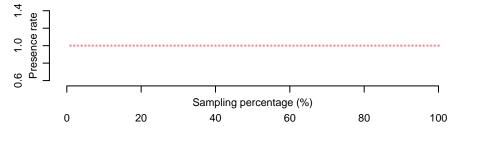
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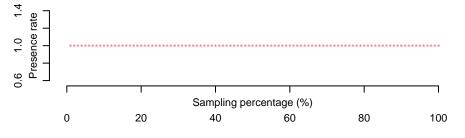


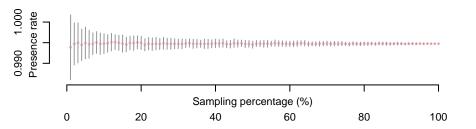


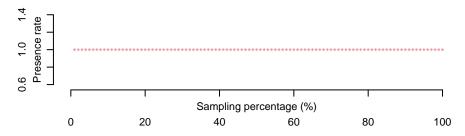
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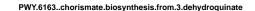
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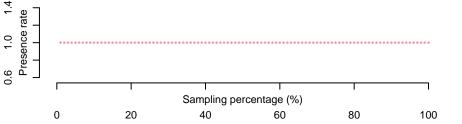


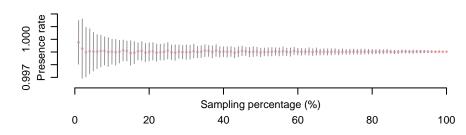






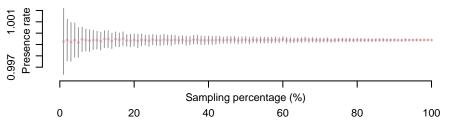
PWY.6168..flavin.biosynthesis.lll..fungi.

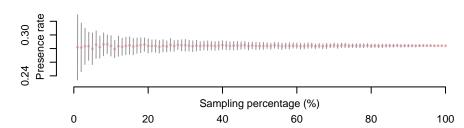




PWY.621..sucrose.degradation.III..sucrose.invertase.

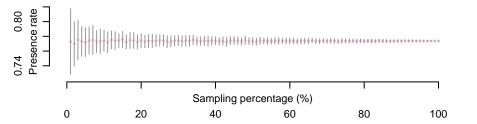
PWY.6263..superpathway.of.menaquinol.8.biosynthesis.II

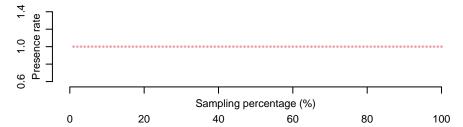




PWY.6270..isoprene.biosynthesis.I

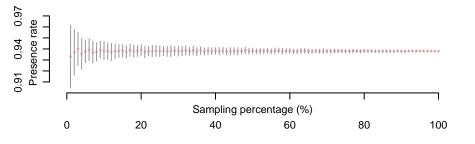
PWY.6277..superpathway.of.5.aminoimidazole.ribonucleotide.biosynthesis

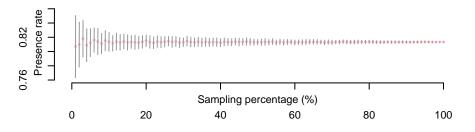


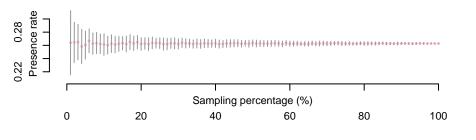


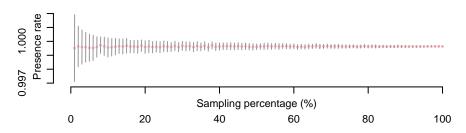
PWY.6282..palmitoleate.biosynthesis.l..from..5Z..dodec.5.enoate.

PWY.6284..superpathway.of.unsaturated.fatty.acids.biosynthesis..E..coli.



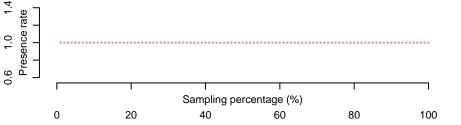


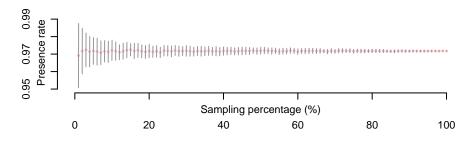




PWY.6317..galactose.degradation.l..Leloir.pathway.

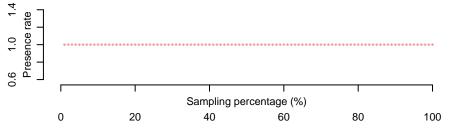
PWY.6353..purine.nucleotides.degradation.ll..aerobic.

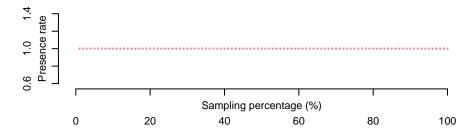




PWY.6385..peptidoglycan.biosynthesis.lll..mycobacteria.

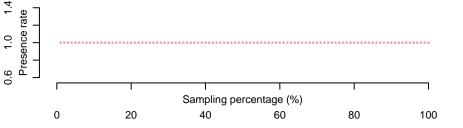
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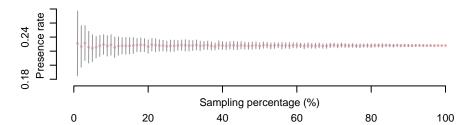




PWY.6387..UDP.N.acetylmuramoyl.pentapeptide.biosynthesis.l..meso.diaminopimelate.containing.

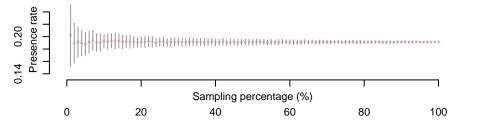
PWY.6415..L.ascorbate.biosynthesis.V

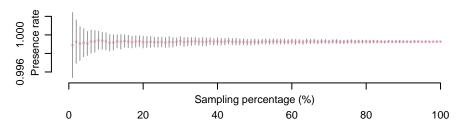


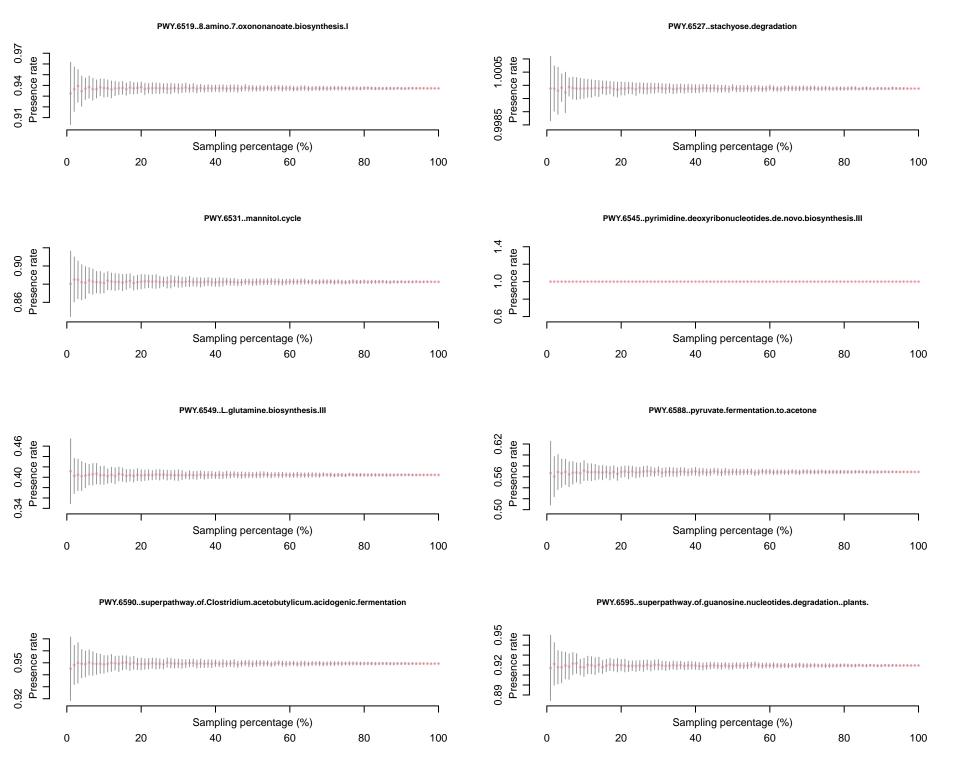


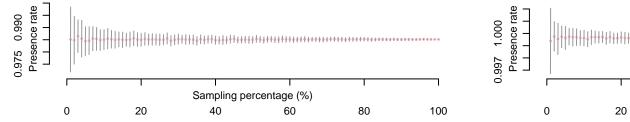
PWY.6471..peptidoglycan.biosynthesis.IV..Enterococcus.faecium.

PWY.6507..4.deoxy.L.threo.hex.4.enopyranuronate.degradation

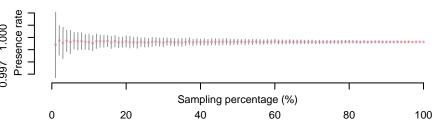






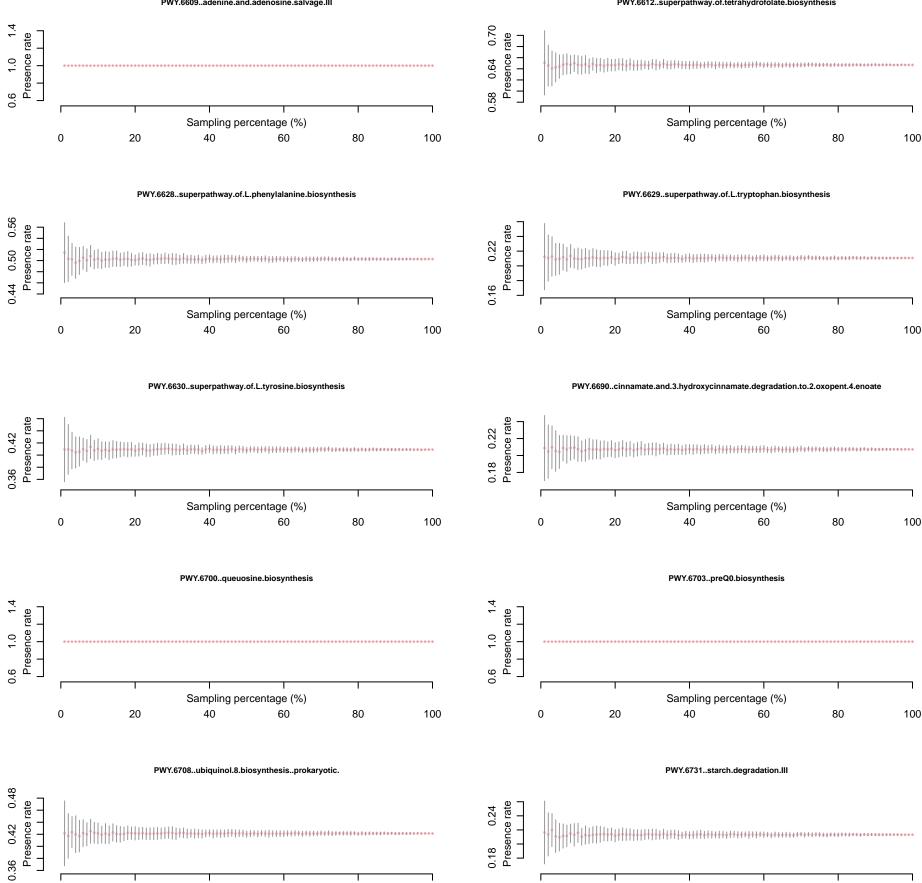


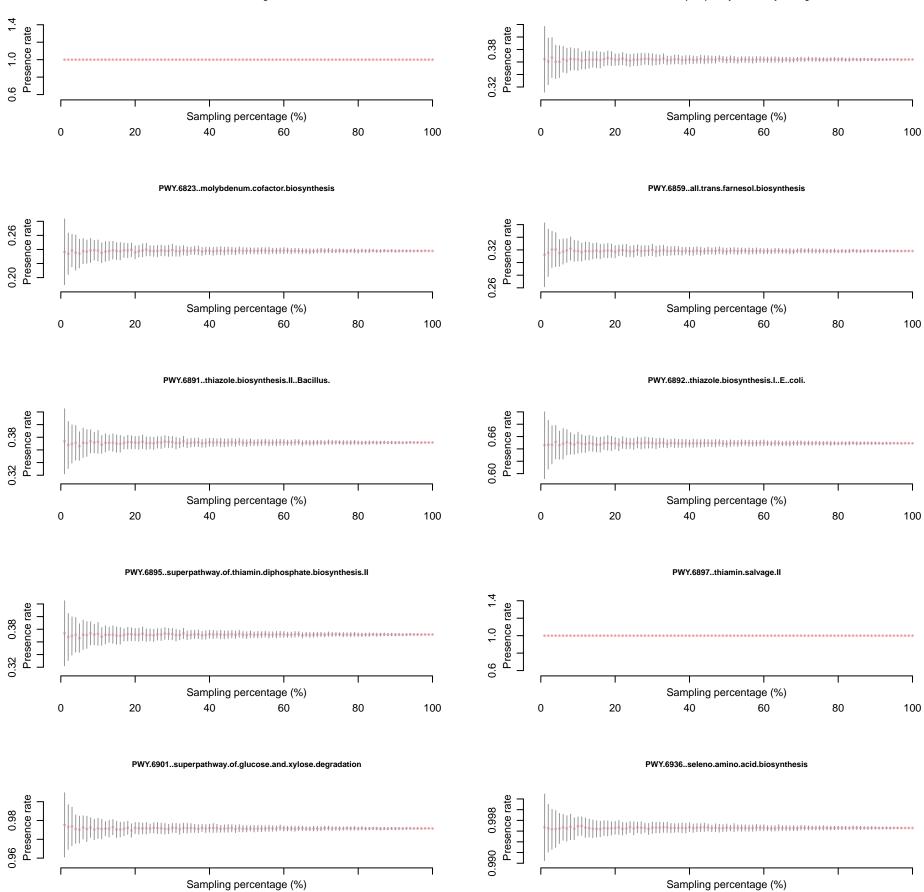
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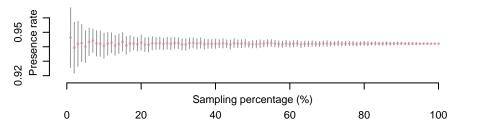


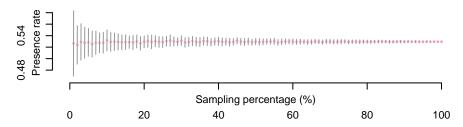
PWY.6608..guanosine.nucleotides.degradation.III

Sampling percentage (%)



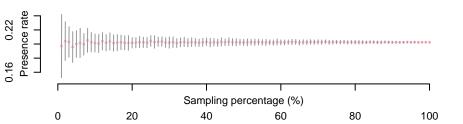


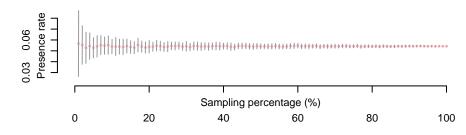






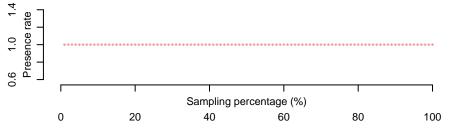
PWY.7039..phosphatidate.metabolism..as.a.signaling.molecule

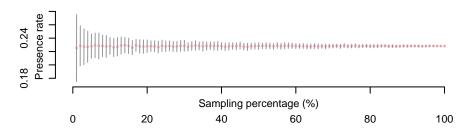




PWY.7111..pyruvate.fermentation.to.isobutanol..engineered.

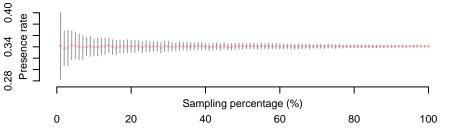
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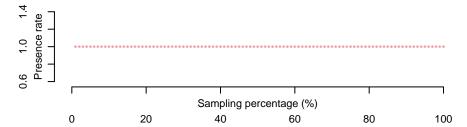




PWY.7117..C4.photosynthetic.carbon.assimilation.cycle..PEPCK.type

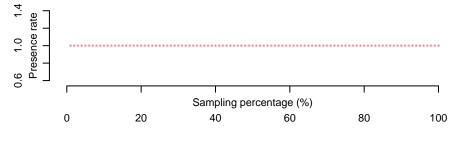
PWY.7184..pyrimidine.deoxyribonucleotides.de.novo.biosynthesis.l

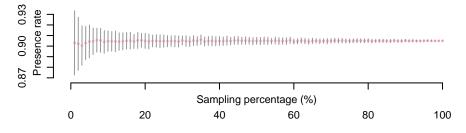




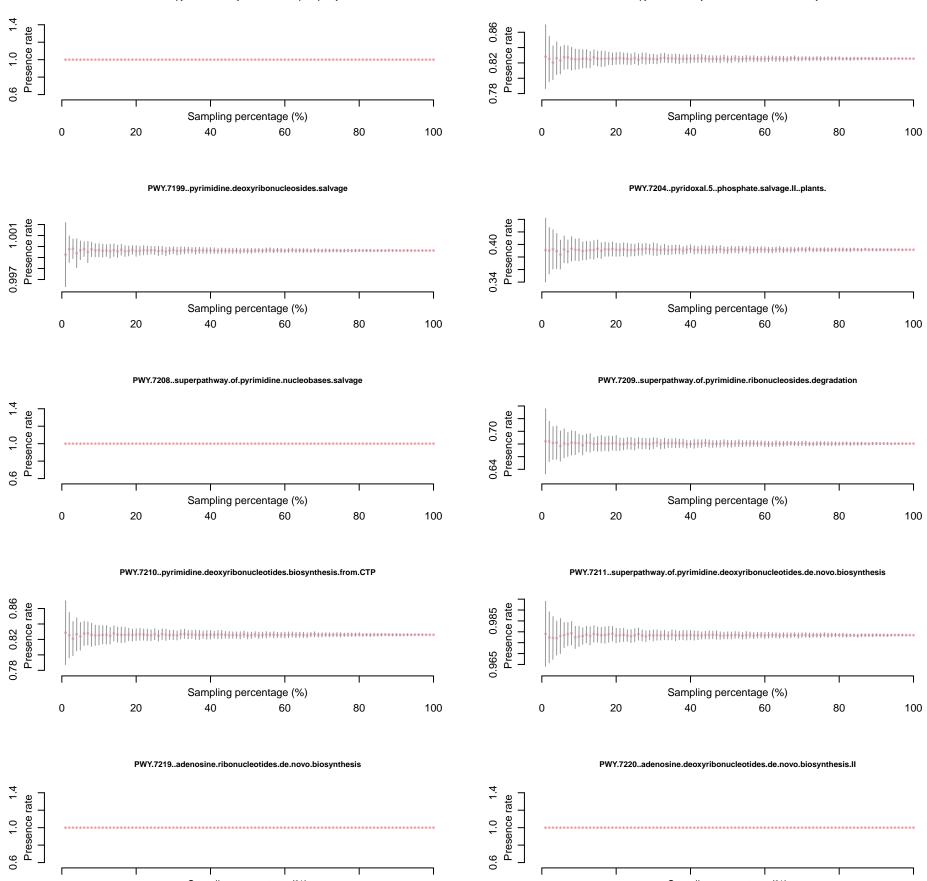
PWY.7187..pyrimidine.deoxyribonucleotides.de.novo.biosynthesis.ll

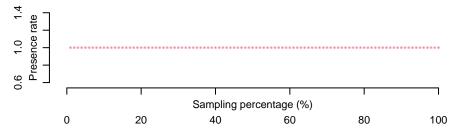
PWY.7196..superpathway.of.pyrimidine.ribonucleosides.salvage

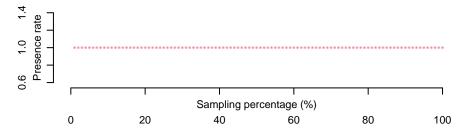




Sampling percentage (%)

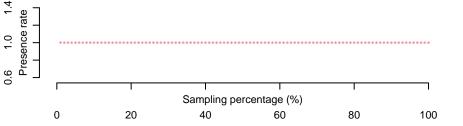


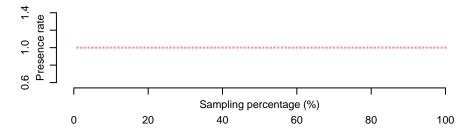




PWY.7228..superpathway.of.guanosine.nucleotides.de.novo.biosynthesis.l

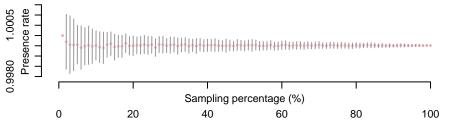
PWY.7229..superpathway.of.adenosine.nucleotides.de.novo.biosynthesis.l

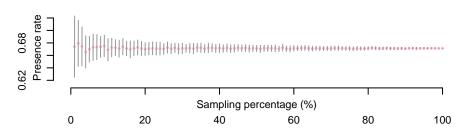




PWY.7234..inosine.5..phosphate.biosynthesis.III

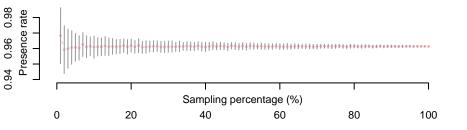
PWY.7235..superpathway.of.ubiquinol.6.biosynthesis..eukaryotic.

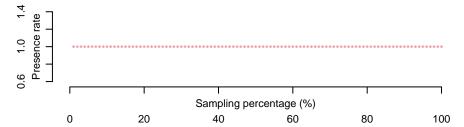




PWY.7237..myo...chiro..and.scillo.inositol.degradation

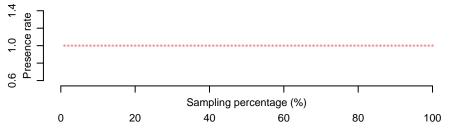
PWY.7242..D.fructuronate.degradation

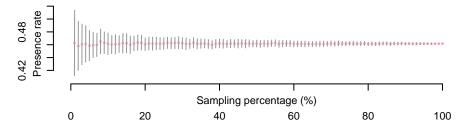


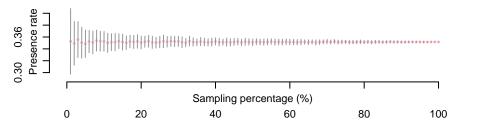


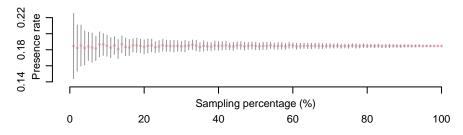
PWY.724..superpathway.of.L.lysine..L.threonine.and.L.methionine.biosynthesis.ll

PWY.7254..TCA.cycle.VII..acetate.producers.



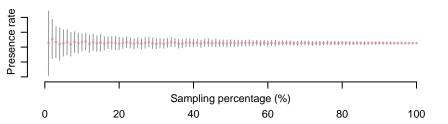




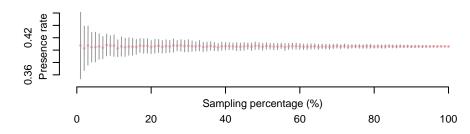


PWY.7282..4.amino.2.methyl.5.phosphomethylpyrimidine.biosynthesis..yeast.

PWY.7315..dTDP.N.acetylthomosamine.biosynthesis

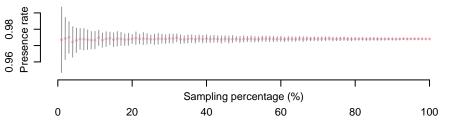


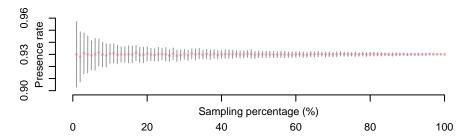
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PWY.7323..superpathway.of.GDP.mannose.derived.O.antigen.building.blocks.biosynthesis

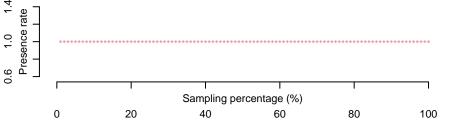
PWY.7328..superpathway.of.UDP.glucose.derived.O.antigen.building.blocks.biosynthesis

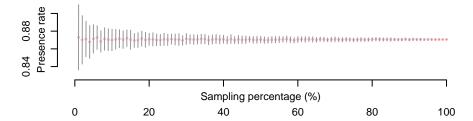




PWY.7357..thiamin.formation.from.pyrithiamine.and.oxythiamine..yeast.

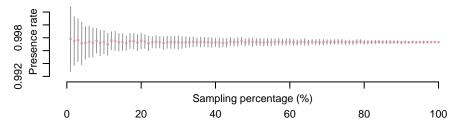
PWY.7371..1.4.dihydroxy.6.naphthoate.biosynthesis.ll

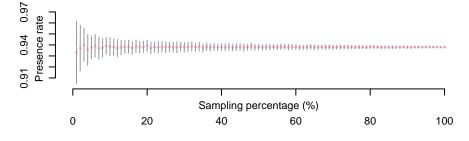


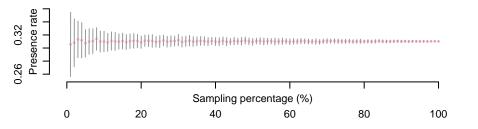


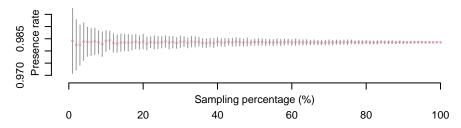
PWY.7383..anaerobic.energy.metabolism..invertebrates..cytosol.

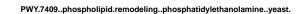
PWY.7388..octanoyl..acyl.carrier.protein..biosynthesis..mitochondria..yeast.



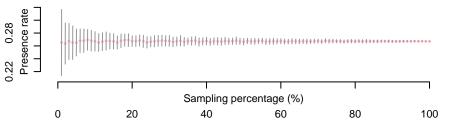


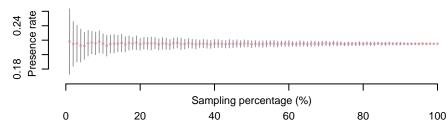






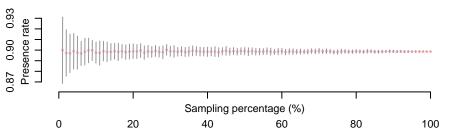


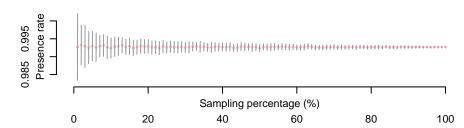




PWY.7456..mannan.degradation

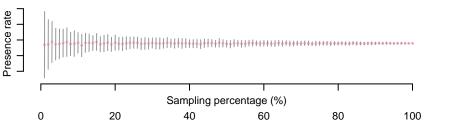
PWY.7539..6.hydroxymethyl.dihydropterin.diphosphate.biosynthesis.lll..Chlamydia.



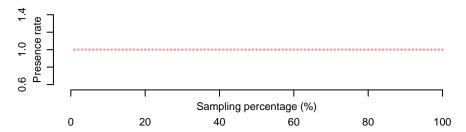


PWY.7560..methylerythritol.phosphate.pathway.ll

PWY.7663..gondoate.biosynthesis..anaerobic.

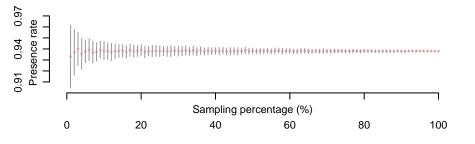


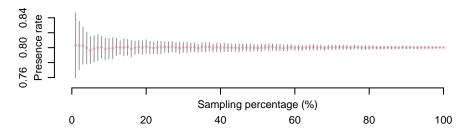
0.76 0.80 0.84

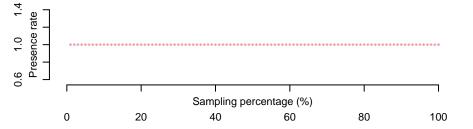


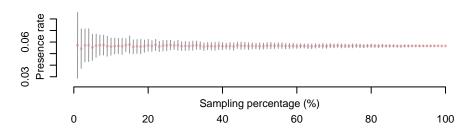
PWY.7664..oleate.biosynthesis.IV..anaerobic

PWY.821.. superpathway. of. sulfur. a mino. acid. biosynthesis.. Saccharomyces. cerevisiae.



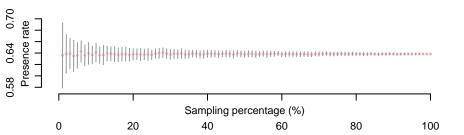


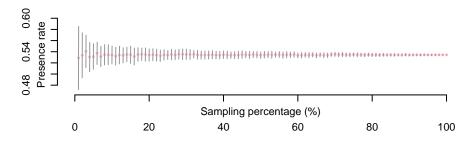






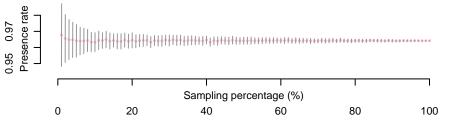
PWY0.1241..ADP.L.glycero..beta..D.manno.heptose.biosynthesis

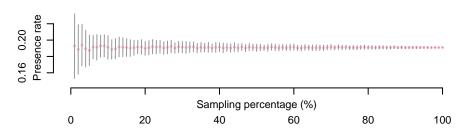




PWY0.1261..anhydromuropeptides.recycling

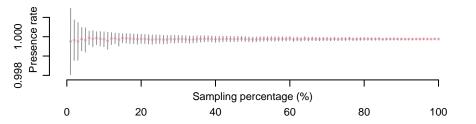
PWY0.1277..3.phenylpropanoate.and.3..3.hydroxyphenyl.propanoate.degradation

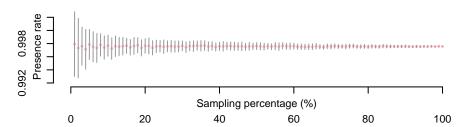




PWY0.1296..purine.ribonucleosides.degradation

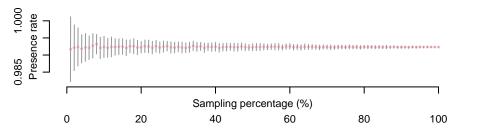
PWY0.1297..superpathway.of.purine.deoxyribonucleosides.degradation

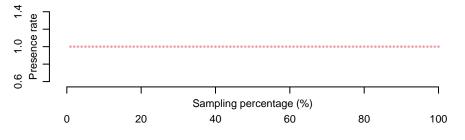




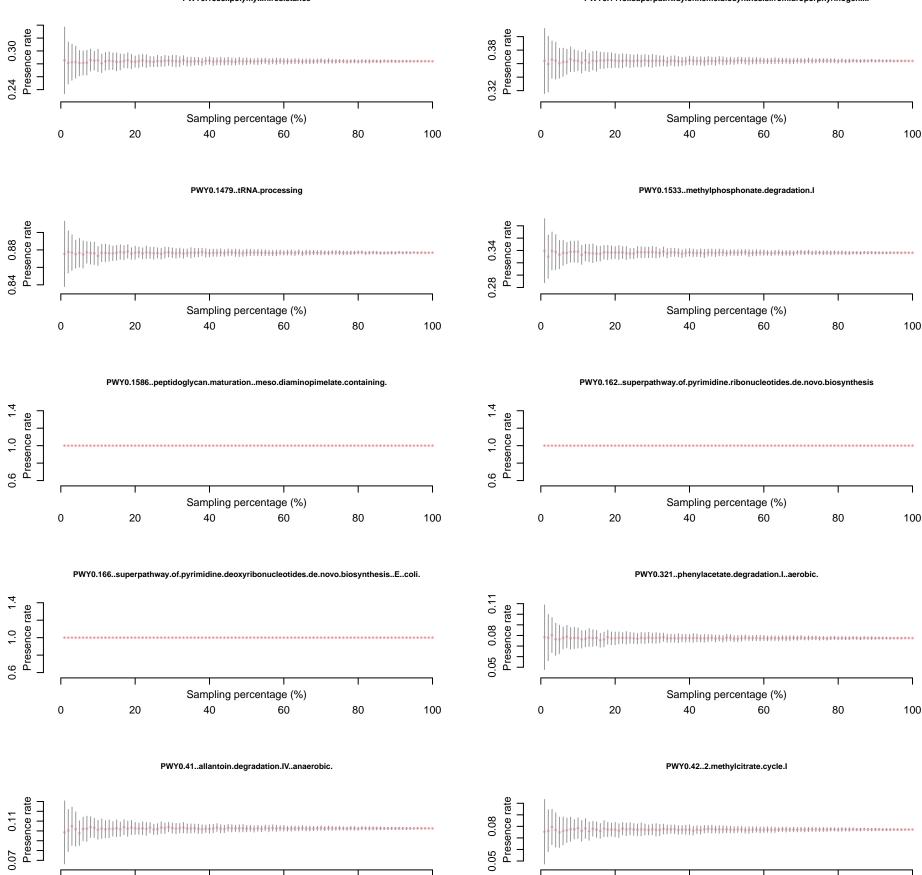
PWY0.1298..superpathway.of.pyrimidine.deoxyribonucleosides.degradation

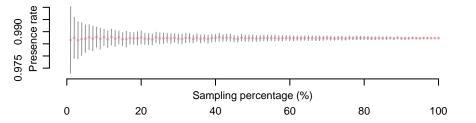
PWY0.1319..CDP.diacylglycerol.biosynthesis.ll

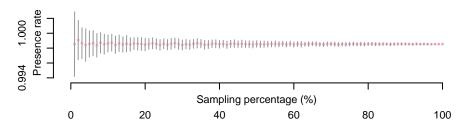




Sampling percentage (%)

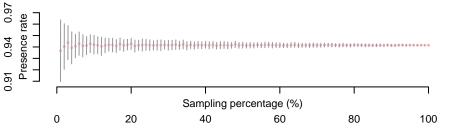


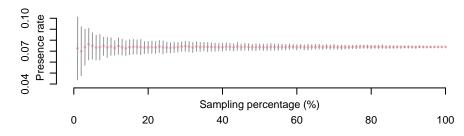






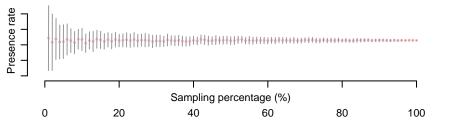
PWY3O.19..ubiquinol.6.biosynthesis.from.4.hydroxybenzoate..eukaryotic.



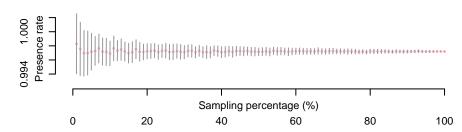


PWY490.3..nitrate.reduction.VI..assimilatory.

PWY4FS.7..phosphatidylglycerol.biosynthesis.l..plastidic.

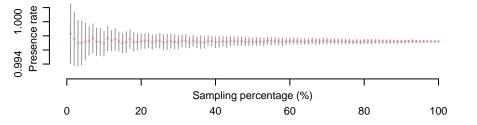


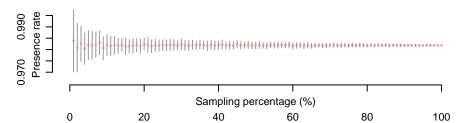
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PWY4FS.8..phosphatidylglycerol.biosynthesis.ll..non.plastidic.

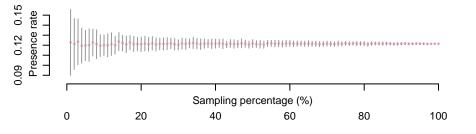
PWY4LZ.257..superpathway.of.fermentation..Chlamydomonas.reinhardtii.

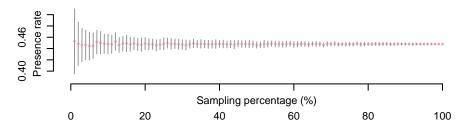


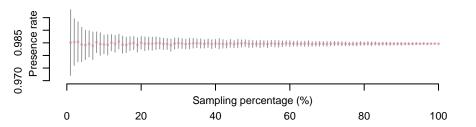


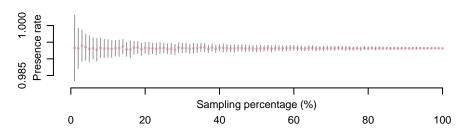
PWY66.389..phytol.degradation

PWY66.398..TCA.cycle.III..animals.



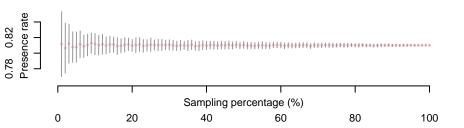


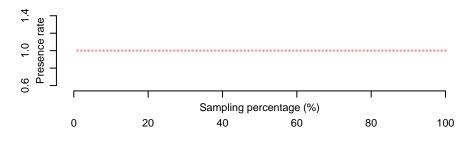




PWY66.409..superpathway.of.purine.nucleotide.salvage

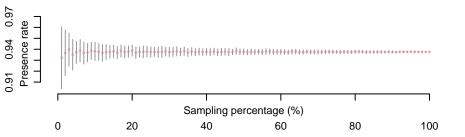
PWY66.422..D.galactose.degradation.V..Leloir.pathway.

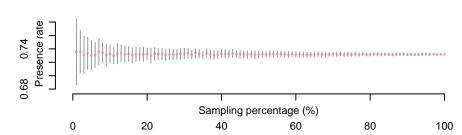




PWYG.321..mycolate.biosynthesis

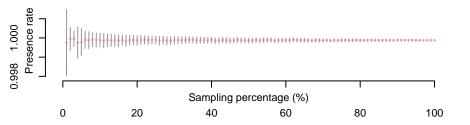
PYRIDNUCSAL.PWY..NAD.salvage.pathway.l

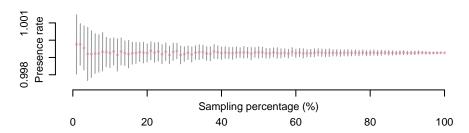




PYRIDNUCSYN.PWY..NAD.biosynthesis.l..from.aspartate.

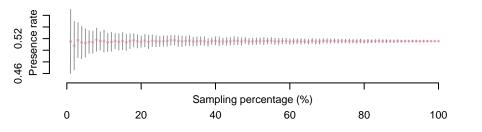
PYRIDOXSYN.PWY..pyridoxal.5..phosphate.biosynthesis.l

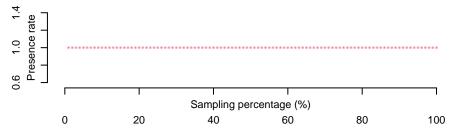




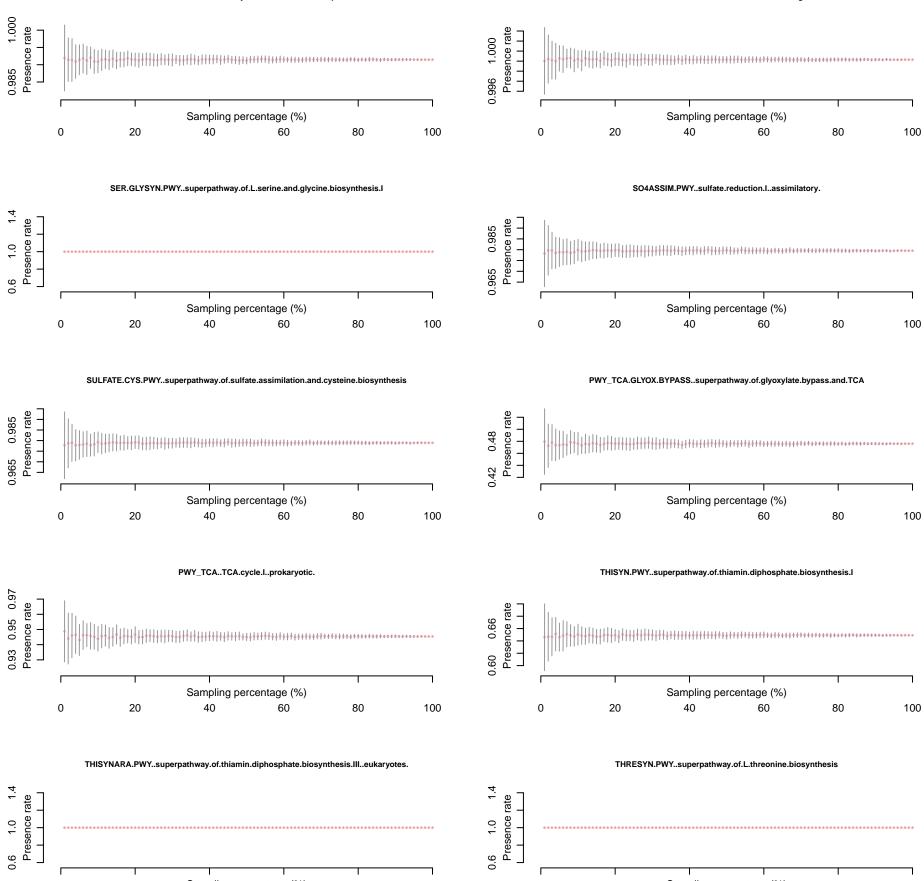
PWY_REDCITCYC..TCA.cycle.VIII..helicobacter.

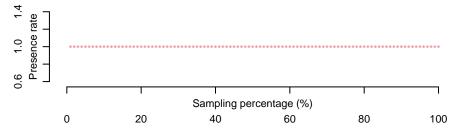
RHAMCAT.PWY..L.rhamnose.degradation.I

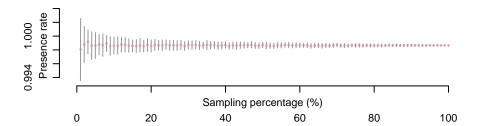




Sampling percentage (%)



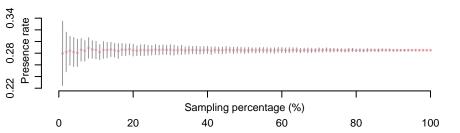


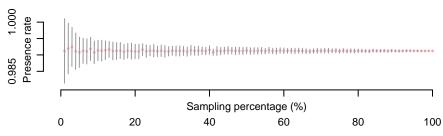


UBISYN.PWY..superpathway.of.ubiquinol.8.biosynthesis..prokaryotic.



UDPNAGSYN.PWY..UDP.N.acetyl.D.glucosamine.biosynthesis.I





URDEGR.PWY..superpathway.of.allantoin.degradation.in.plants



