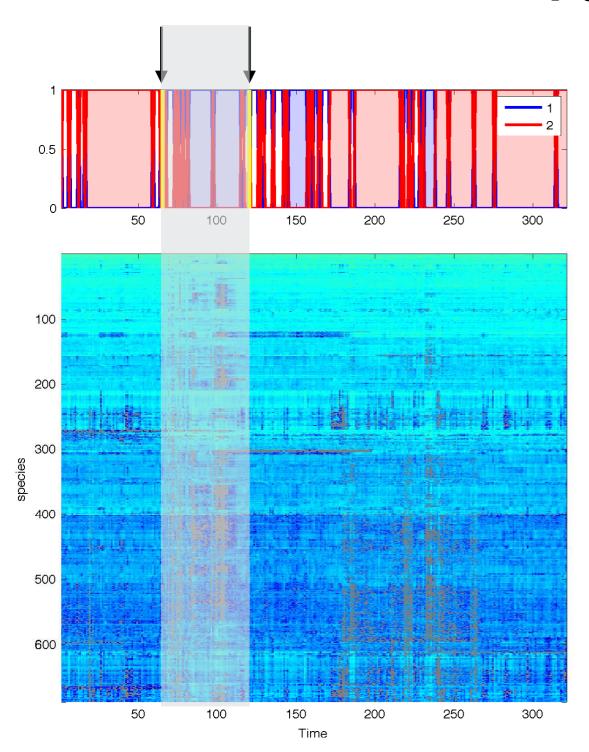
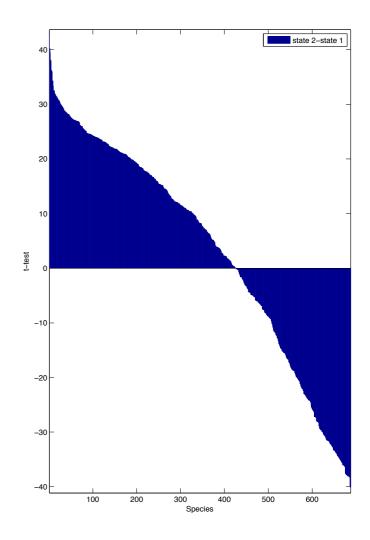
Patient A



Which are the species that vary the most between the 2 states?



t-test>35:

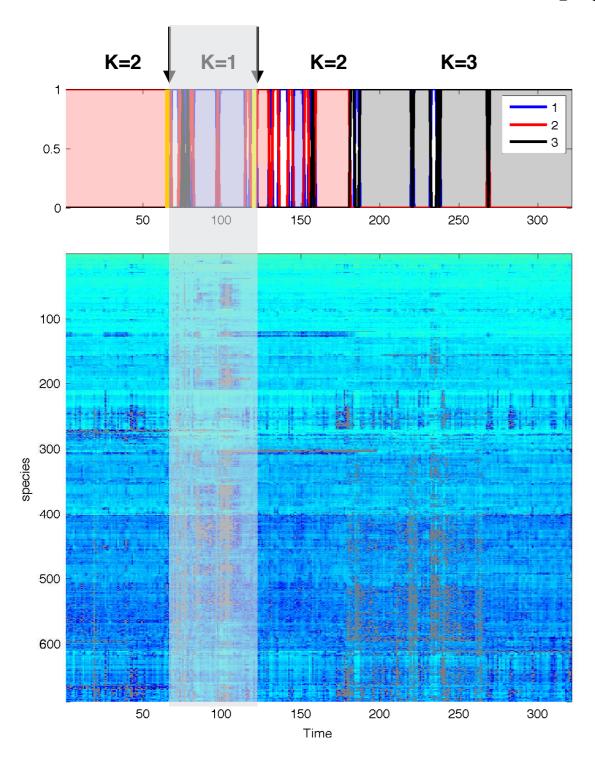
gnavus_CR_OTU_188333 Blautia_CR_OTU_184984 Blautia_CR_OTU_195186 Lachnospiraceae_CR_OTU_292399 Acinetobacter_CR_OTU_4425634 Bifidobacterium_CR_OTU_553611 Lachnospiraceae_CR_OTU_195759

t-test<-38:

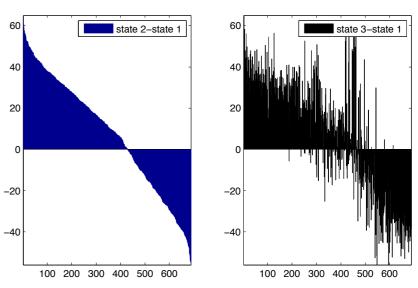
Lachnospiraceae_CR_OTU_300716 Ruminococcaceae_CR_OTU_197750 Bacteroides_CR_OTU_4449054 Dorea_CR_OTU_305329 Roseburia_CR_OTU_536831 Bacteroides_CR_OTU_197367 Lachnospiraceae_CR_OTU_176954 gnavus_CR_OTU_193853

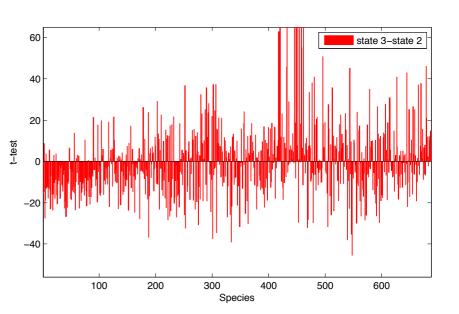
Select hidden state K=3

Patient A



Which are the species that vary the most between the 3 states?

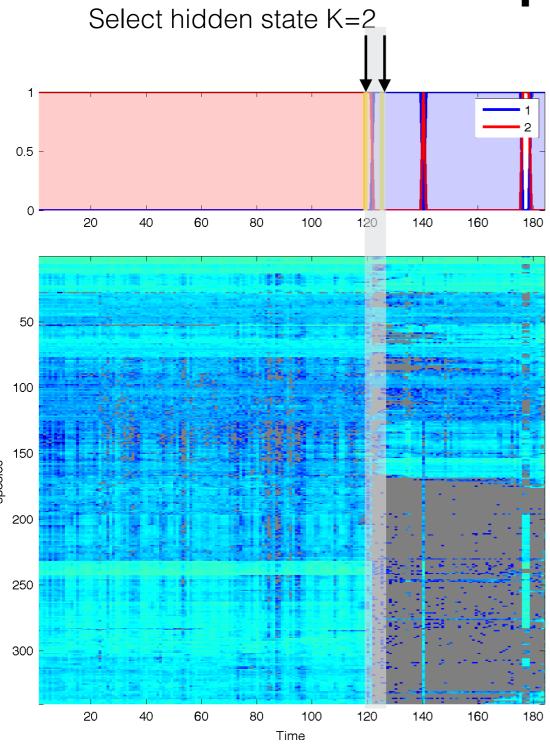




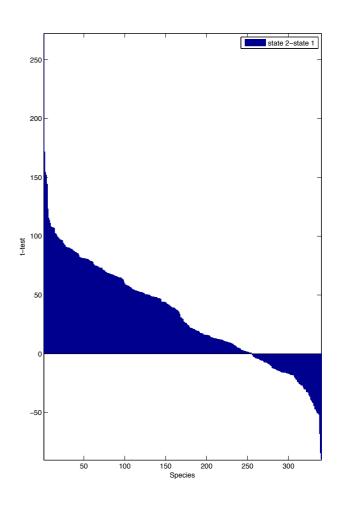
t-test>50:

Blautia_CR_OTU_184984
Blautia_CR_OTU_4468384
Lachnospiraceae_CR_OTU_357963
Lachnospiraceae_CR_OTU_185864
Ruminococcaceae_CR_OTU_312882
Ruminococcus_CR_OTU_4472091
Lachnospiraceae_CR_OTU_195465
Bacteroides_CR_OTU_198788
Lachnospiraceae_CR_OTU_176297
Ruminococcaceae_CR_OTU_176757

Patient B



Which are the species that vary the most between the 3 states?



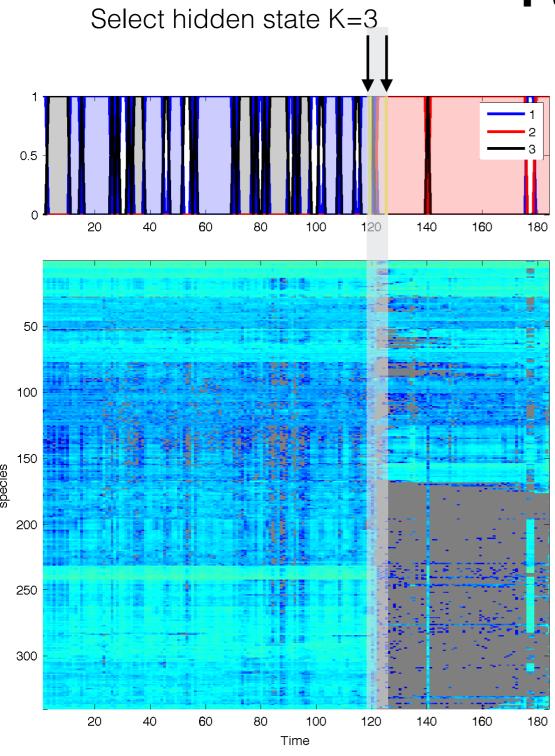
t-test>100:

Lachnospiraceae_CR_OTU_4456027
Ruminococcaceae_CR_OTU_360995
Dorea_CR_OTU_305329
Bacteroides_CR_OTU_1944498
Clostridiaceae_CR_OTU_4387453
Erysipelotrichaceae_CR_OTU_191718
Blautia_CR_OTU_358798
Lachnobacterium_CR_OTU_2220496
Bacteroides_CR_OTU_533373
Clostridiales_CR_OTU_177032
Roseburia_CR_OTU_518613
Ruminococcaceae_CR_OTU_173863
Ruminococcaceae_CR_OTU_1599042
Ruminococcaceae_CR_OTU_182089
Ruminococcaceae_CR_OTU_4431545

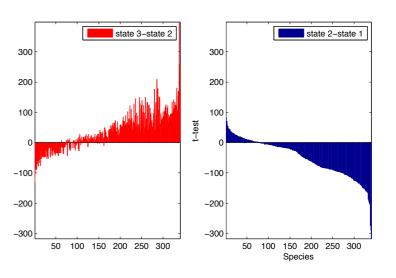
t-test<-50:

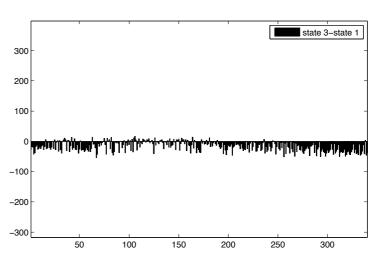
Christensenellaceae_CR_OTU_2442706 Lachnospiraceae_CR_OTU_361972 Ruminococcaceae_CR_OTU_327218 Phascolarctobacterium_CR_OTU_174049 Lachnospiraceae_CR_OTU_194112 Lachnospiraceae_CR_OTU_2420187

Patient B



Which are the species that vary the most between the 3 states?





t-test<-45:

Dorea_CR_OTU_1667433
Lachnospiraceae_CR_OTU_163494
Lachnospiraceae_CR_OTU_196526
Ruminococcaceae_CR_OTU_179864
Ruminococcaceae_CR_OTU_339566
Clostridiales_CR_OTU_4347520
prausnitzii_CR_OTU_319275
Bacteroides_CR_OTU_1698008
Lachnospiraceae_CR_OTU_189292
Leuconostocaceae_CR_OTU_189292
Lachnospiraceae_CR_OTU_175698
Clostridiales_CR_OTU_4462599