####### 5S-Bacteria-aspralign #######

Method: single

Rand\_score 0.5529175050301811

Homogeneity\_score 0.18073456660159648

completeness\_score 0.4626212981445627

Method: complete

Rand\_score 0.7090543259557344

Homogeneity\_score 0.4551692141619585

completeness\_score 0.602272969566192

Method: average

Rand\_score 0.7090543259557344

Homogeneity\_score 0.4551692141619584

completeness\_score 0.602272969566192

####### 5S-Bacteria-pskorder #######

Method: single

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: complete

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: average

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

####### 5S-Bacteria-genus #######

Method: single

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: complete

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: average

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

####### 5S-Bacteria-psmalign #######

Method: single

Rand\_score 0.7476861167002012

Homogeneity\_score 0.5569767655811528

completeness\_score 0.7196911632891175

Method: complete

Rand\_score 0.929979879275654

Homogeneity\_score 0.8542496297556706

completeness\_score 0.8156210364501525

Method: average

Rand\_score 0.8297786720321931

Homogeneity\_score 0.6999999890933517

completeness\_score 0.783776866149487

####### 5S-Bacteria-eigevalues #######

Method: single

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: complete

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

Method: average

Rand\_score 0.42334004024144867

Homogeneity\_score 0.030934202035020876

completeness\_score 0.07918142595297606

####### 16S-Bacteria-aspralign #######

Method: single

Rand\_score 0.46145728643216083

Homogeneity\_score 0.2764155680473436

completeness\_score 0.5641430868416992

Method: complete

Rand\_score 0.8179396984924623

Homogeneity\_score 0.6840400443629563

completeness\_score 0.5824012865799807

Method: average

Rand\_score 0.82321608040201

Homogeneity\_score 0.6605102152073611

completeness\_score 0.6131245192673109

####### 16S-Bacteria-pskorder #######

Method: single

Rand\_score 0.3024120603015075

Homogeneity\_score 0.07355571504234057

completeness\_score 0.2792971795181419

Method: complete

Rand\_score 0.3024120603015075

Homogeneity\_score 0.07355571504234057

completeness\_score 0.2792971795181419

Method: average

Rand\_score 0.3024120603015075

Homogeneity\_score 0.07355571504234057

completeness\_score 0.2792971795181419

####### 16S-Bacteria-genus #######

Method: single

Rand\_score 0.40733668341708545

Homogeneity\_score 0.09983866588126618

completeness\_score 0.20620720537595327

Method: complete

Rand\_score 0.45256281407035176

Homogeneity\_score 0.14574744915515345

completeness\_score 0.27164905133901884

Method: average

Rand\_score 0.45256281407035176

Homogeneity\_score 0.14574744915515345

completeness\_score 0.27164905133901884

####### 16S-Bacteria-psmalign #######

Method: single

Rand\_score 0.4949246231155779

Homogeneity\_score 0.35739359443662366

completeness\_score 0.8432382514557765

Method: complete

Rand\_score 0.8466834170854272

Homogeneity\_score 0.8112186885235896

completeness\_score 0.6308516270110918

Method: average

Rand\_score 0.7587437185929649

Homogeneity\_score 0.6181210851619614

completeness\_score 0.713472058488121

####### 16S-Bacteria-eigevalues #######

Method: single

Rand\_score 0.6959296482412061

Homogeneity\_score 0.23968606779491505

completeness\_score 0.2146656835458438

Method: complete

Rand\_score 0.7071356783919598

Homogeneity\_score 0.2530677233226602

completeness\_score 0.21255168157892046

Method: average

Rand\_score 0.7042211055276382

Homogeneity\_score 0.24660149248078836

completeness\_score 0.2101892688874817

####### 23S-Bacteriaa-aspralign #######

Method: single

Rand\_score 0.4274640088593577

Homogeneity\_score 0.21003750295377194

completeness\_score 0.4710271908025647

Method: complete

Rand\_score 0.8349944629014396

Homogeneity\_score 0.6598700951203536

completeness\_score 0.6849792905216411

Method: average

Rand\_score 0.5016611295681063

Homogeneity\_score 0.30311087712860973

completeness\_score 0.553847399495723

####### 23S-Bacteria-pskorder #######

Method: single

Rand\_score 0.44518272425249167

Homogeneity\_score 0.19082713831329973

completeness\_score 0.42794629351687336

Method: complete

Rand\_score 0.44739756367663347

Homogeneity\_score 0.16238125298649841

completeness\_score 0.30127111475474533

Method: average

Rand\_score 0.44739756367663347

Homogeneity\_score 0.16238125298649841

completeness\_score 0.30127111475474533

####### 23S-Bacteria-genus #######

Method: single

Rand\_score 0.4318936877076412

Homogeneity\_score 0.18416826200382552

completeness\_score 0.4130131898670756

Method: complete

Rand\_score 0.44629014396456257

Homogeneity\_score 0.18050545157965572

completeness\_score 0.3633229175084606

Method: average

Rand\_score 0.44629014396456257

Homogeneity\_score 0.18050545157965572

completeness\_score 0.3633229175084606

####### 23S-Bacteria-psmalign #######

Method: single

Rand\_score 0.8084163898117387

Homogeneity\_score 0.6749791744423378

completeness\_score 0.7464723008145635

Method: complete

Rand\_score 0.8970099667774086

Homogeneity\_score 0.8709633725236959

completeness\_score 0.8140322805751424

Method: average

Rand\_score 0.8504983388704319

Homogeneity\_score 0.7780514248683934

completeness\_score 0.7963473497294369

####### 23S-Bacteria-eigevalues #######

Method: single

Rand\_score 0.583610188261351

Homogeneity\_score 0.2161933823099246

completeness\_score 0.22963103460946435

Method: complete

Rand\_score 0.5913621262458472

Homogeneity\_score 0.22339298193595716

completeness\_score 0.22007585490161516

Method: average

Rand\_score 0.5913621262458472

Homogeneity\_score 0.2233929819359571

completeness\_score 0.22007585490161516