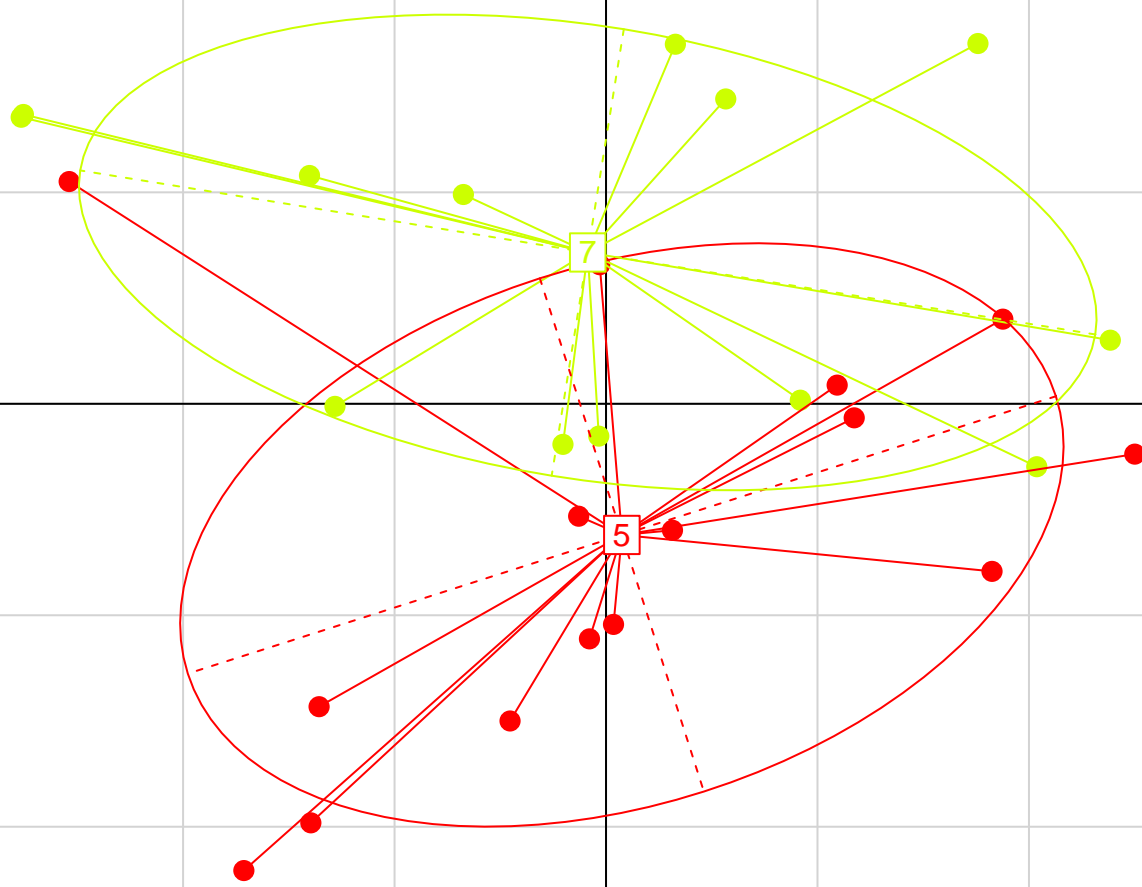
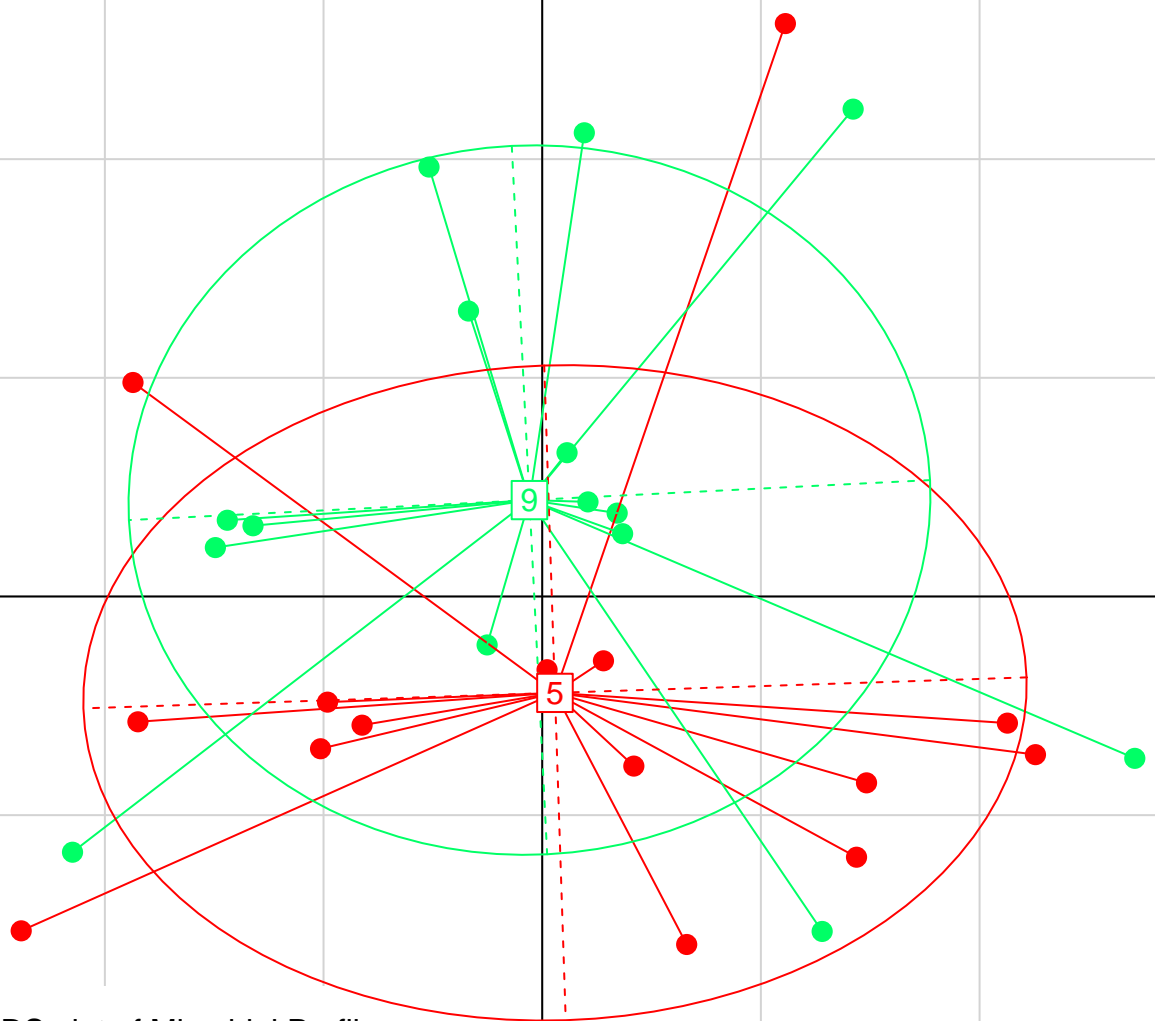


d = 0.1



MDS plot of Microbial Profiles  
5 – 7  
(p-value 0.019, corr. p-value 0.19)

d = 0.1

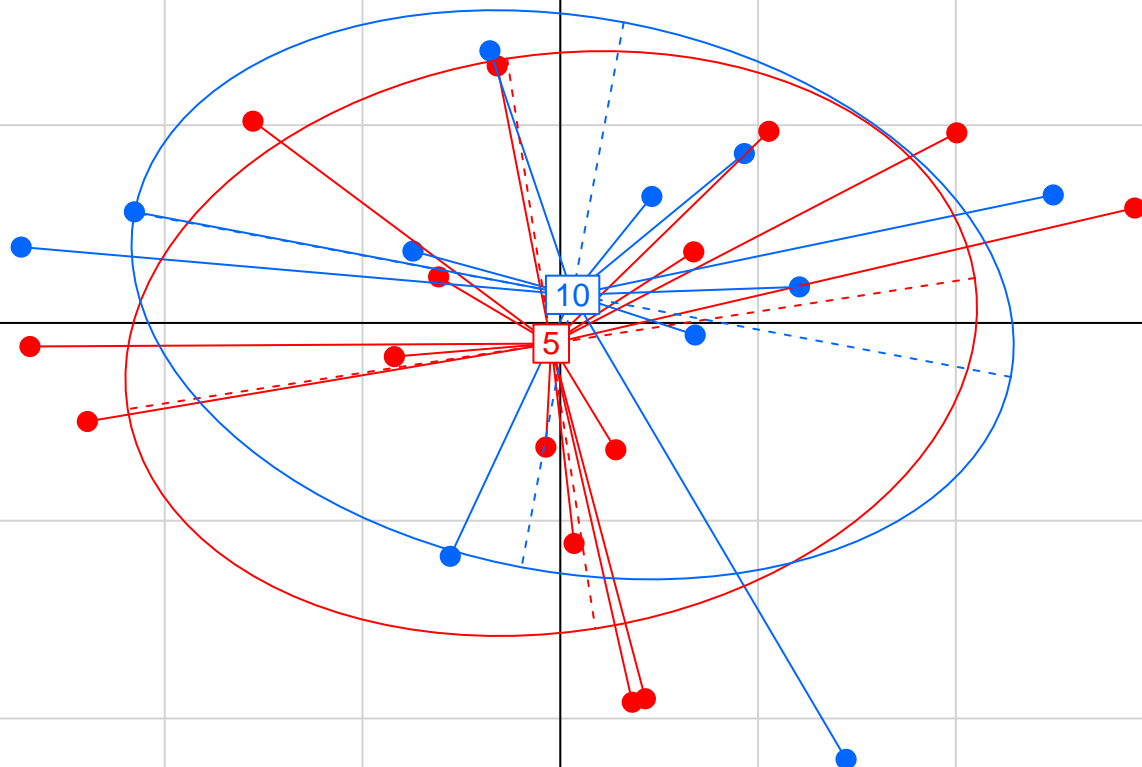


MDS plot of Microbial Profiles

5 – 9

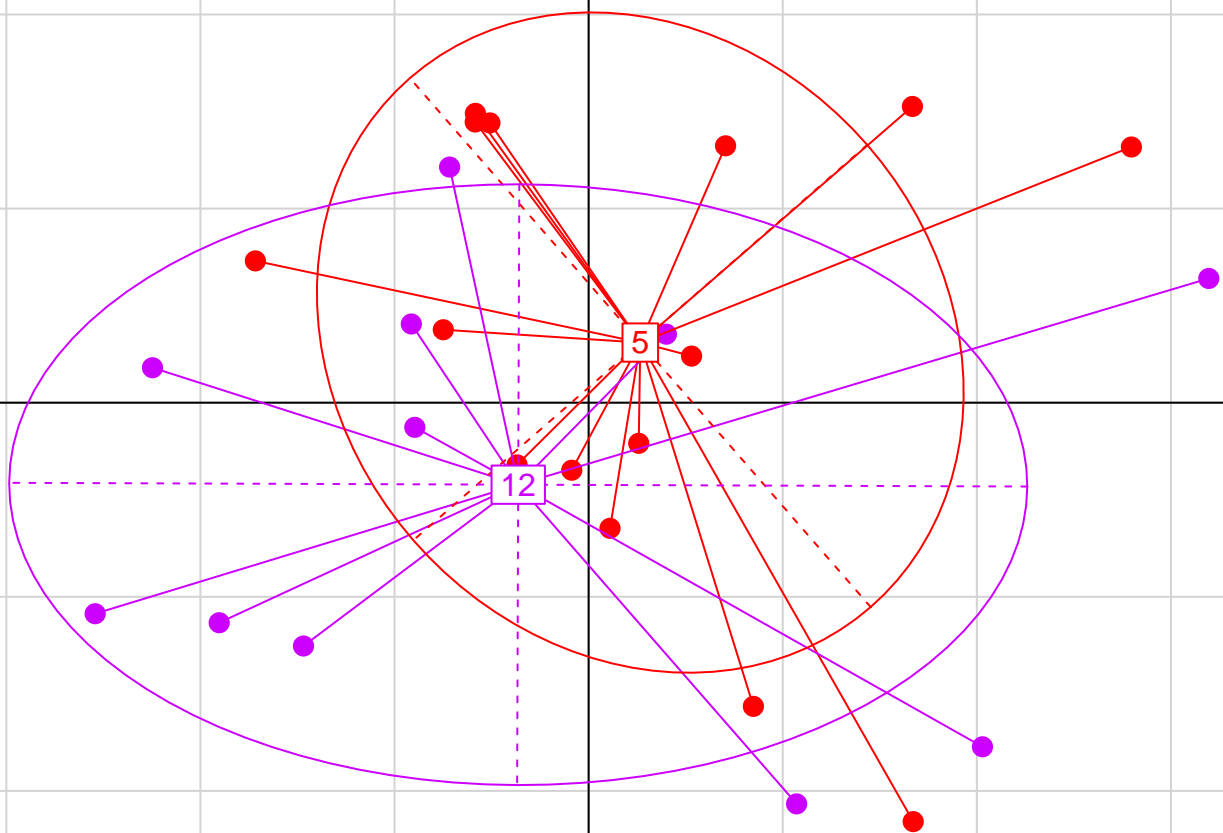
(p-value 0.154, corr. p-value 0.4875)

d = 0.1



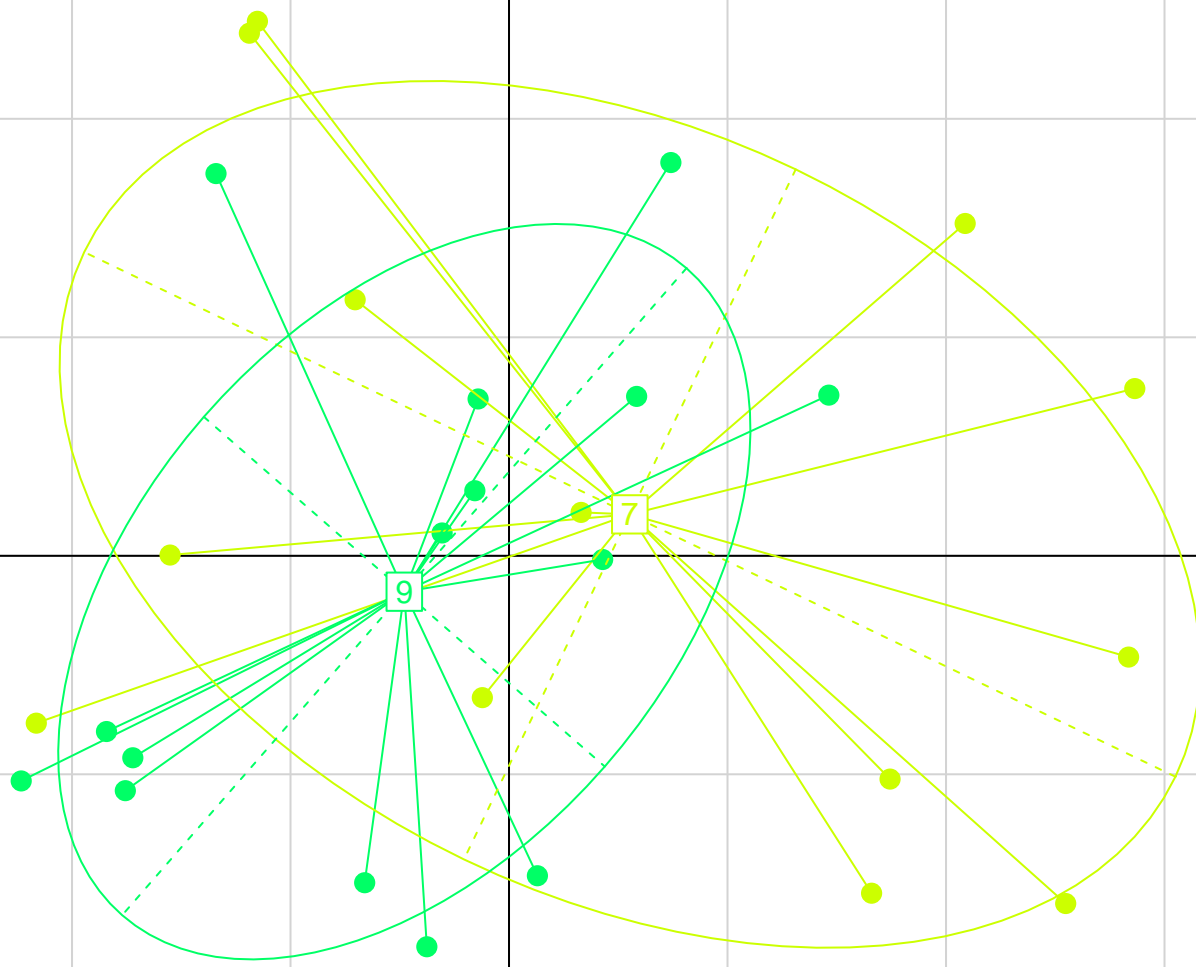
MDS plot of Microbial Profiles  
5 – 10  
(p-value 0.897, corr. p-value 0.897)

d = 0.1



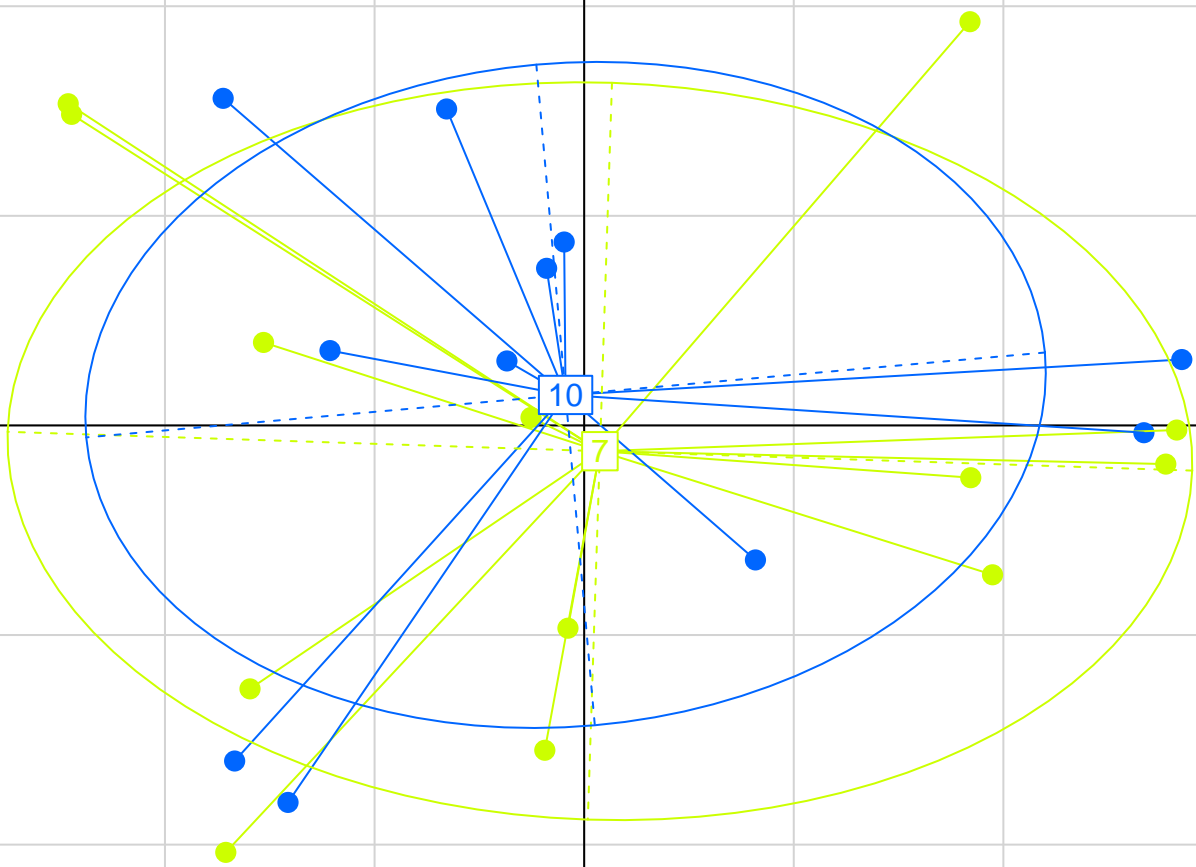
MDS plot of Microbial Profiles  
5 – 12  
(p-value 0.195, corr. p-value 0.4875)

d = 0.1



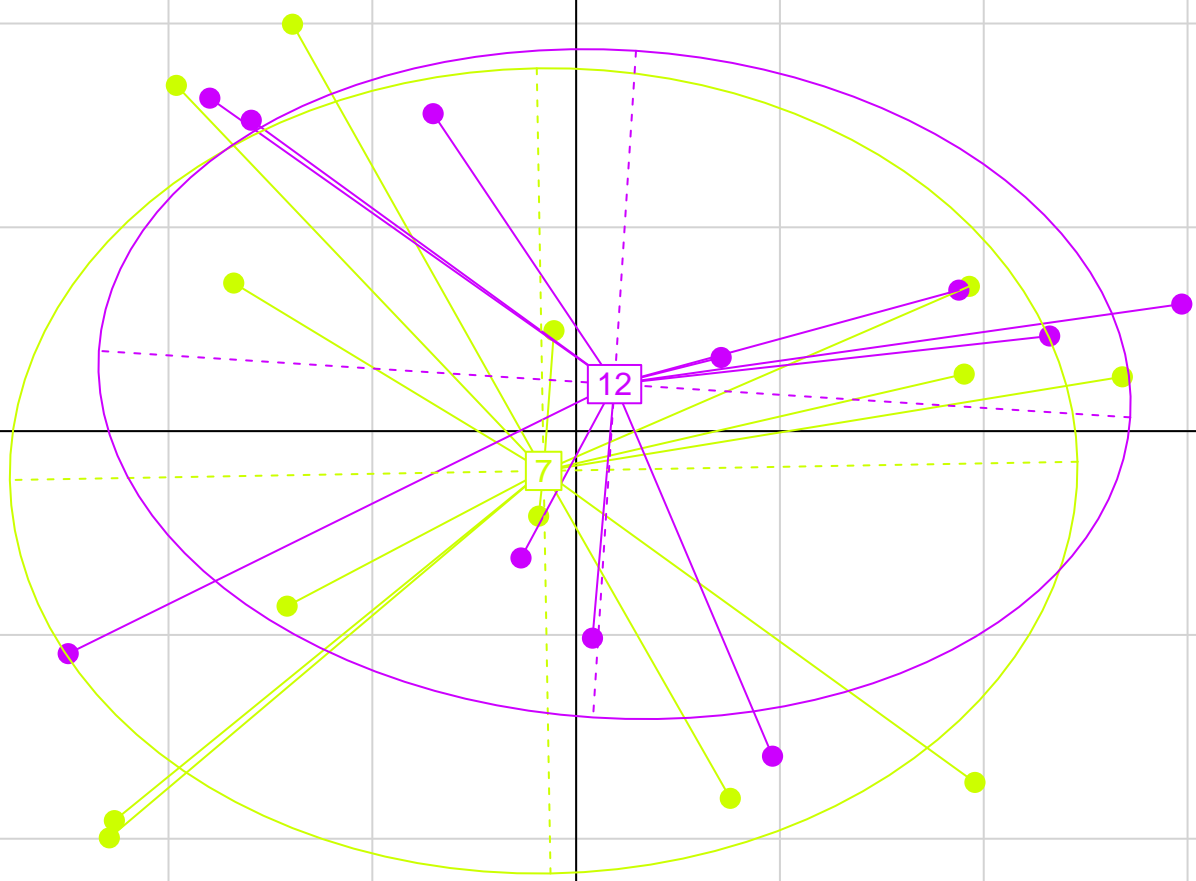
MDS plot of Microbial Profiles  
7 – 9  
(p-value 0.057, corr. p-value 0.285)

d = 0.1



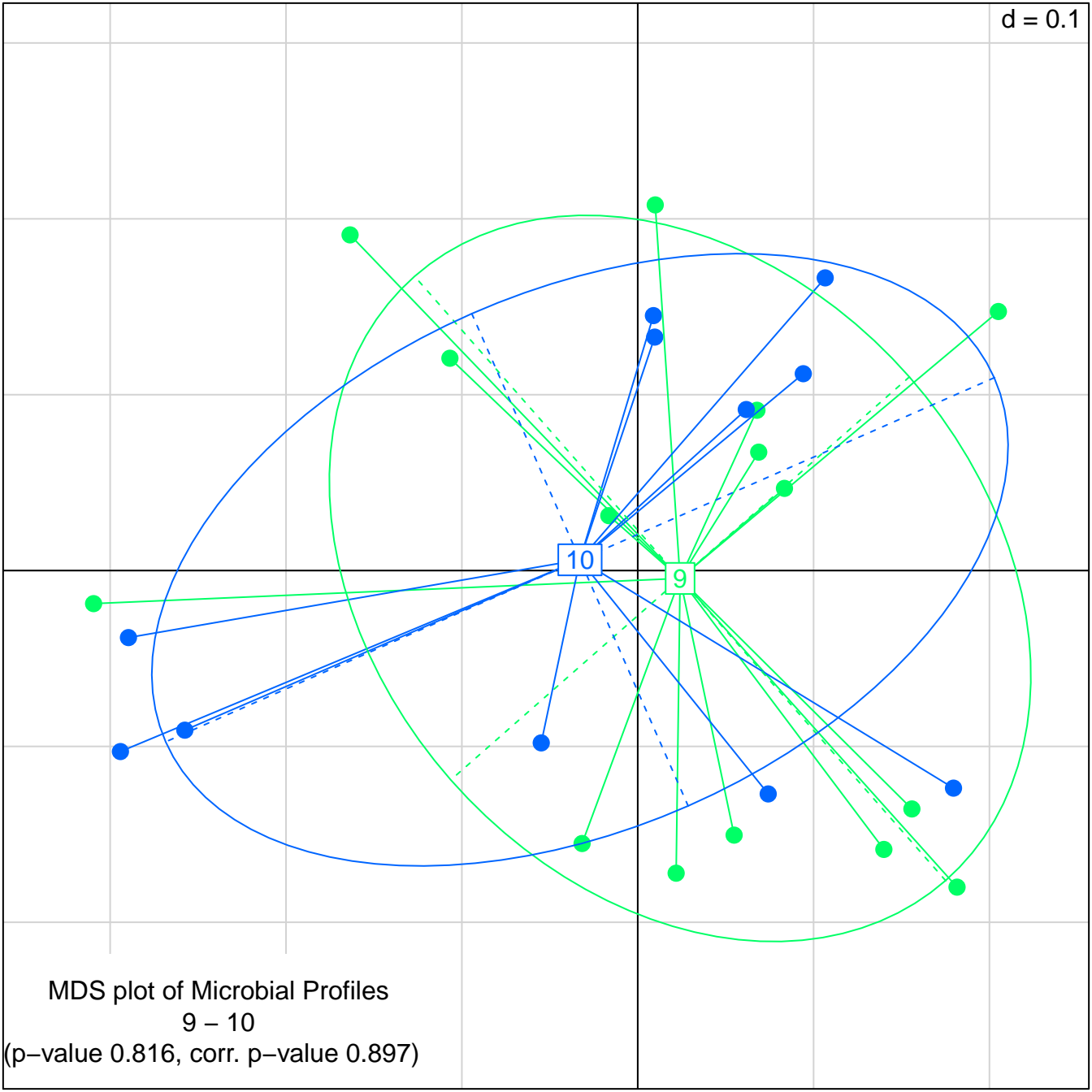
MDS plot of Microbial Profiles  
7 – 10  
(p-value 0.682, corr. p-value 0.897)

d = 0.1



MDS plot of Microbial Profiles  
7 – 12  
(p-value 0.494, corr. p-value 0.897)

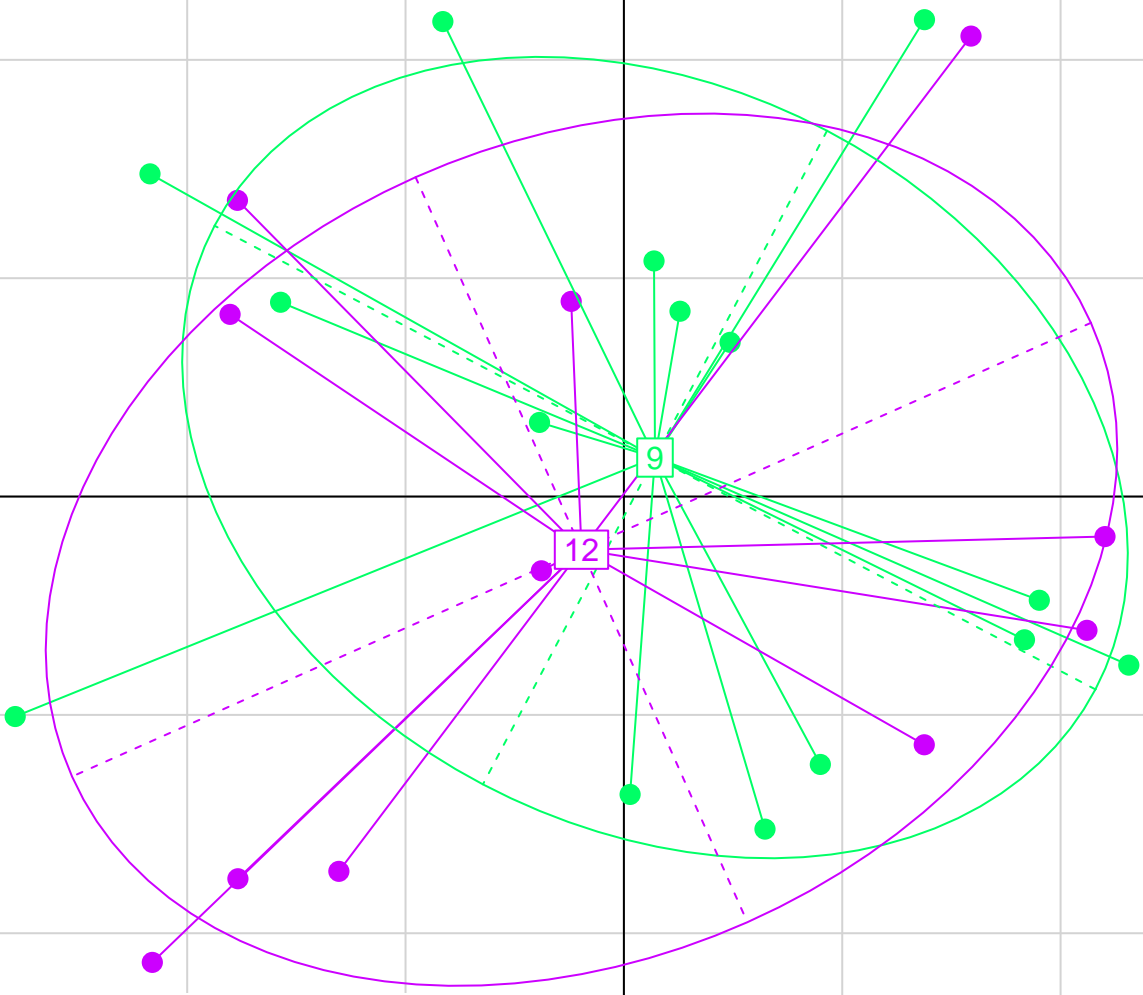
d = 0.1



MDS plot of Microbial Profiles  
9 - 10  
(p-value 0.816, corr. p-value 0.897)

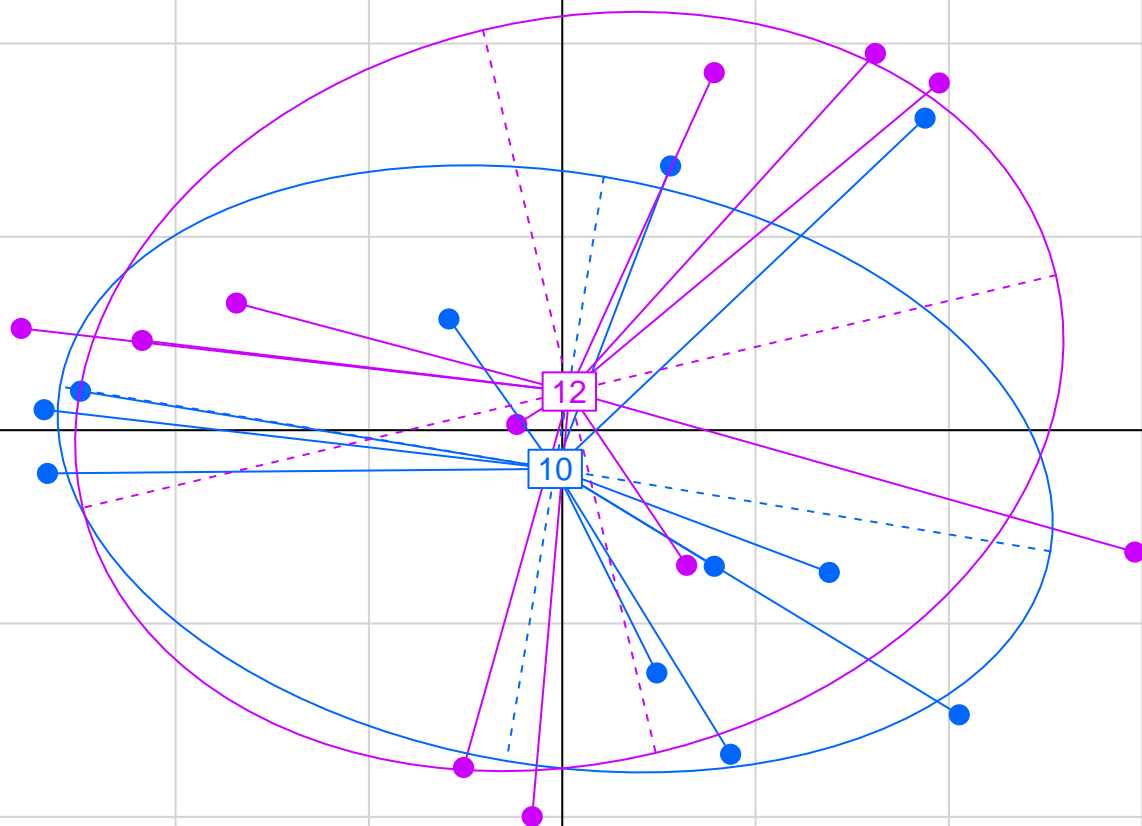


d = 0.1



MDS plot of Microbial Profiles  
9 - 12  
(p-value 0.738, corr. p-value 0.897)

d = 0.1



MDS plot of Microbial Profiles  
10 – 12  
(p-value 0.879, corr. p-value 0.897)