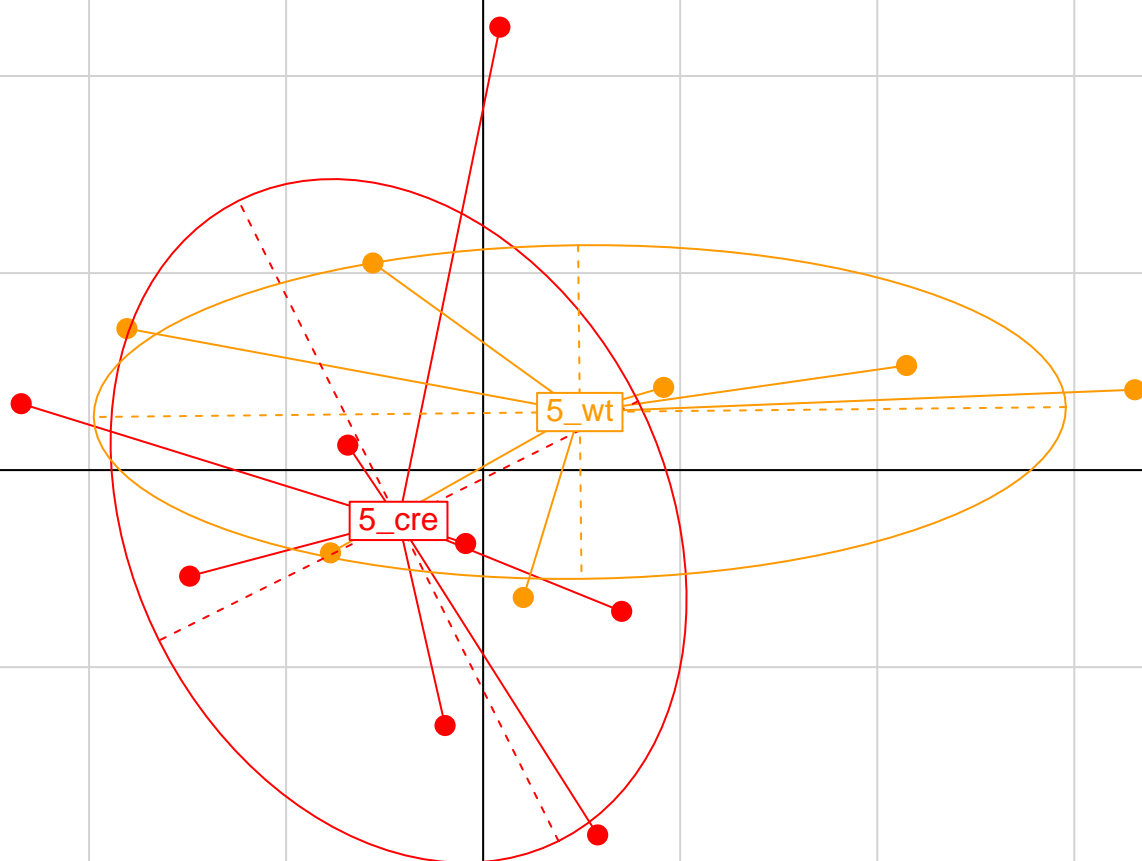
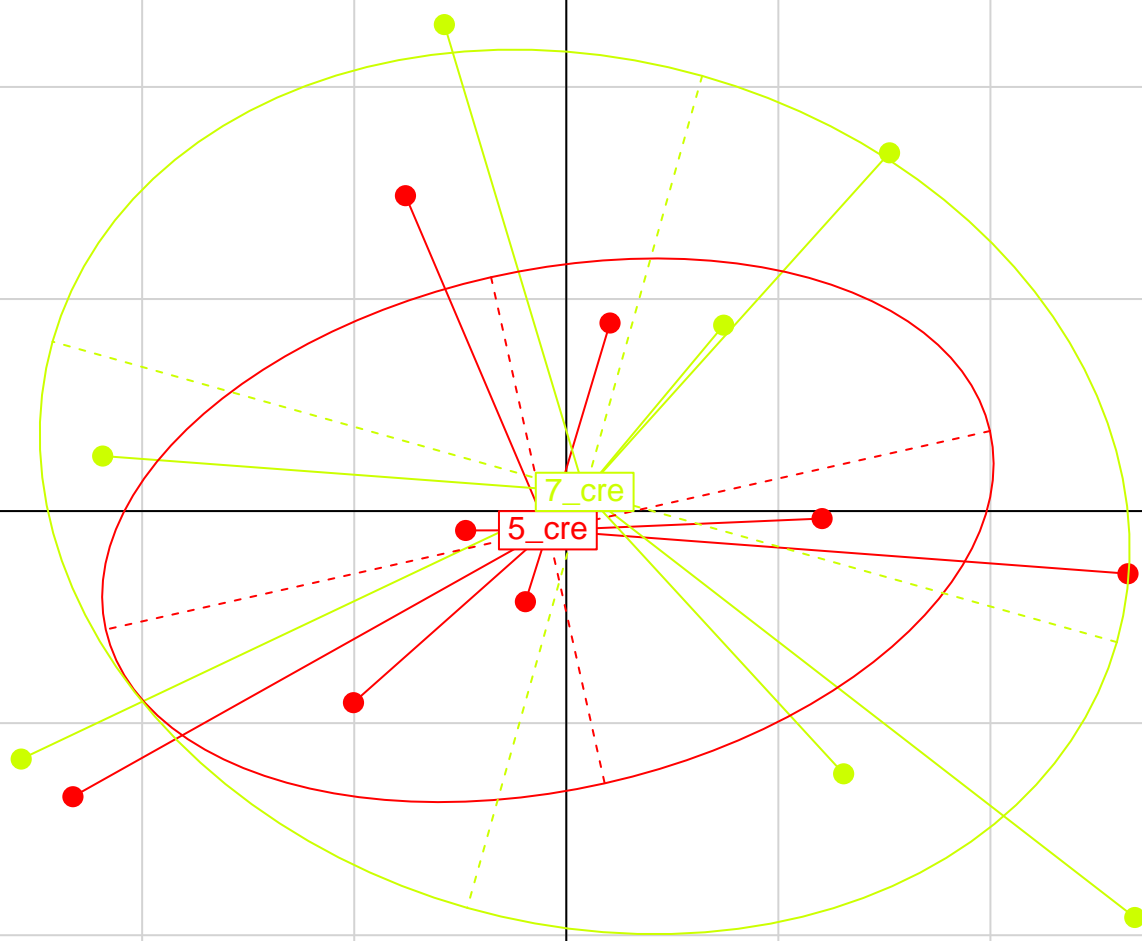


d = 0.1



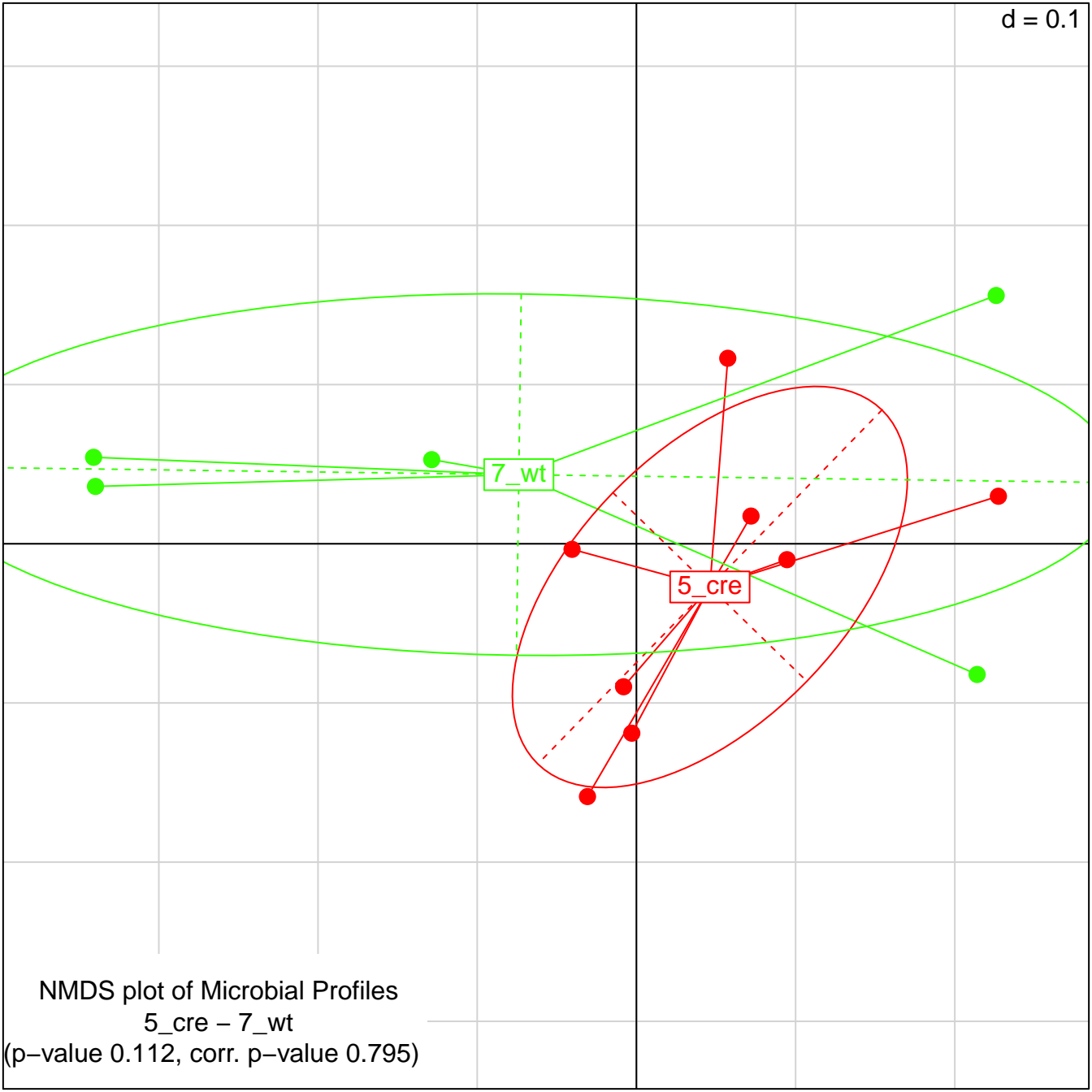
NMDS plot of Microbial Profiles  
5\_cre – 5\_wt  
(p-value 0.191, corr. p-value 0.795)

d = 0.1

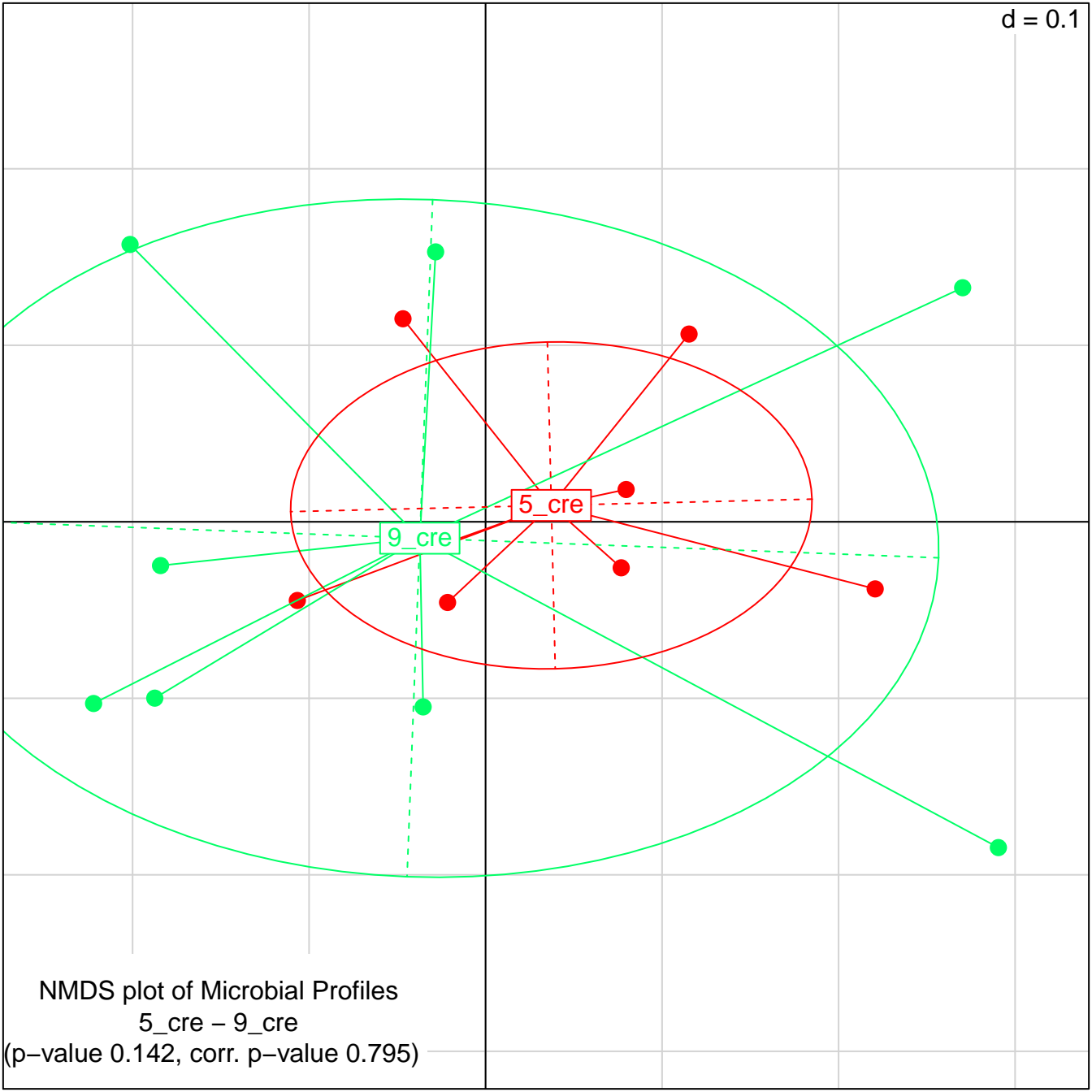


NMDS plot of Microbial Profiles  
5\_cre – 7\_cre  
(p-value 0.95, corr. p-value 0.95)

d = 0.1

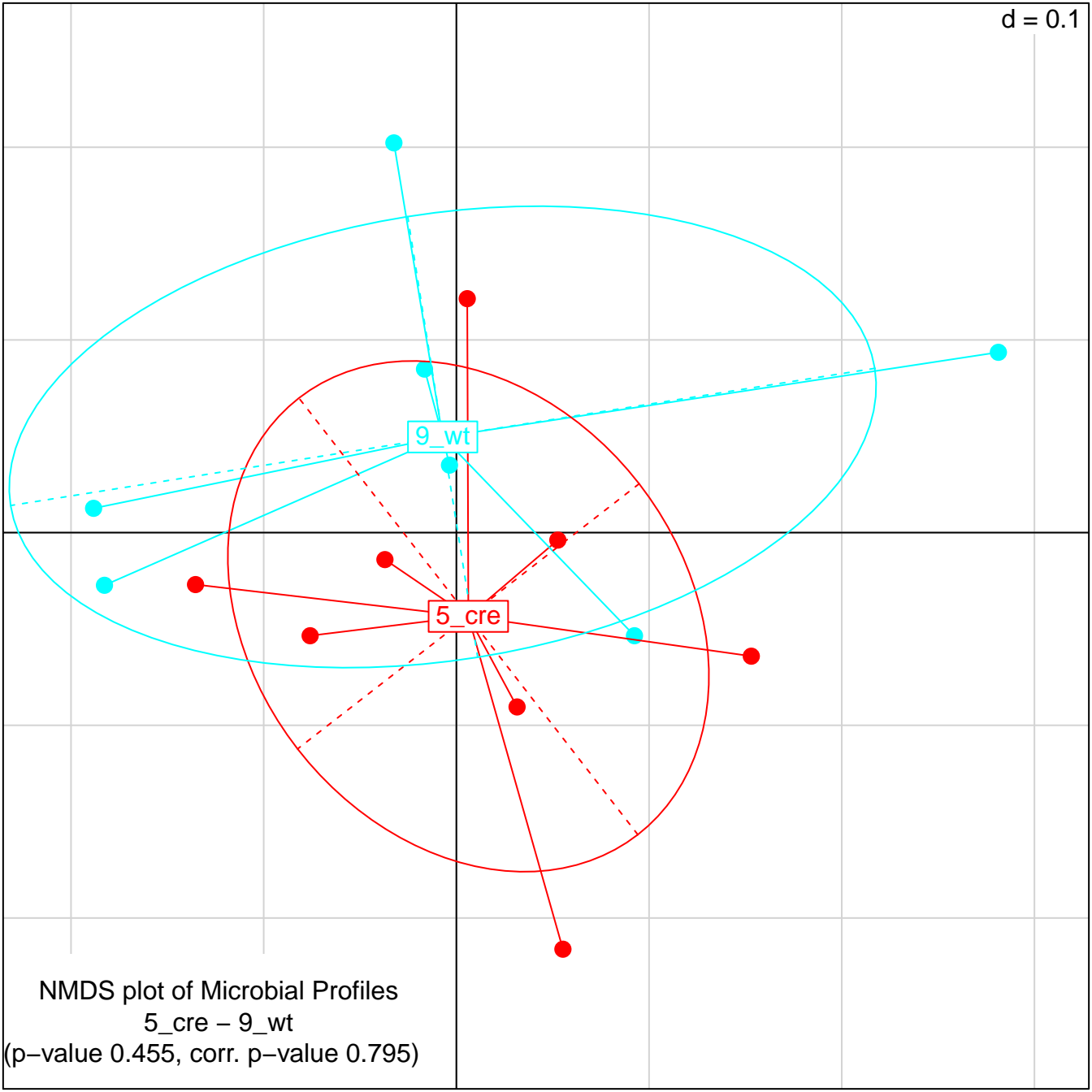


d = 0.1

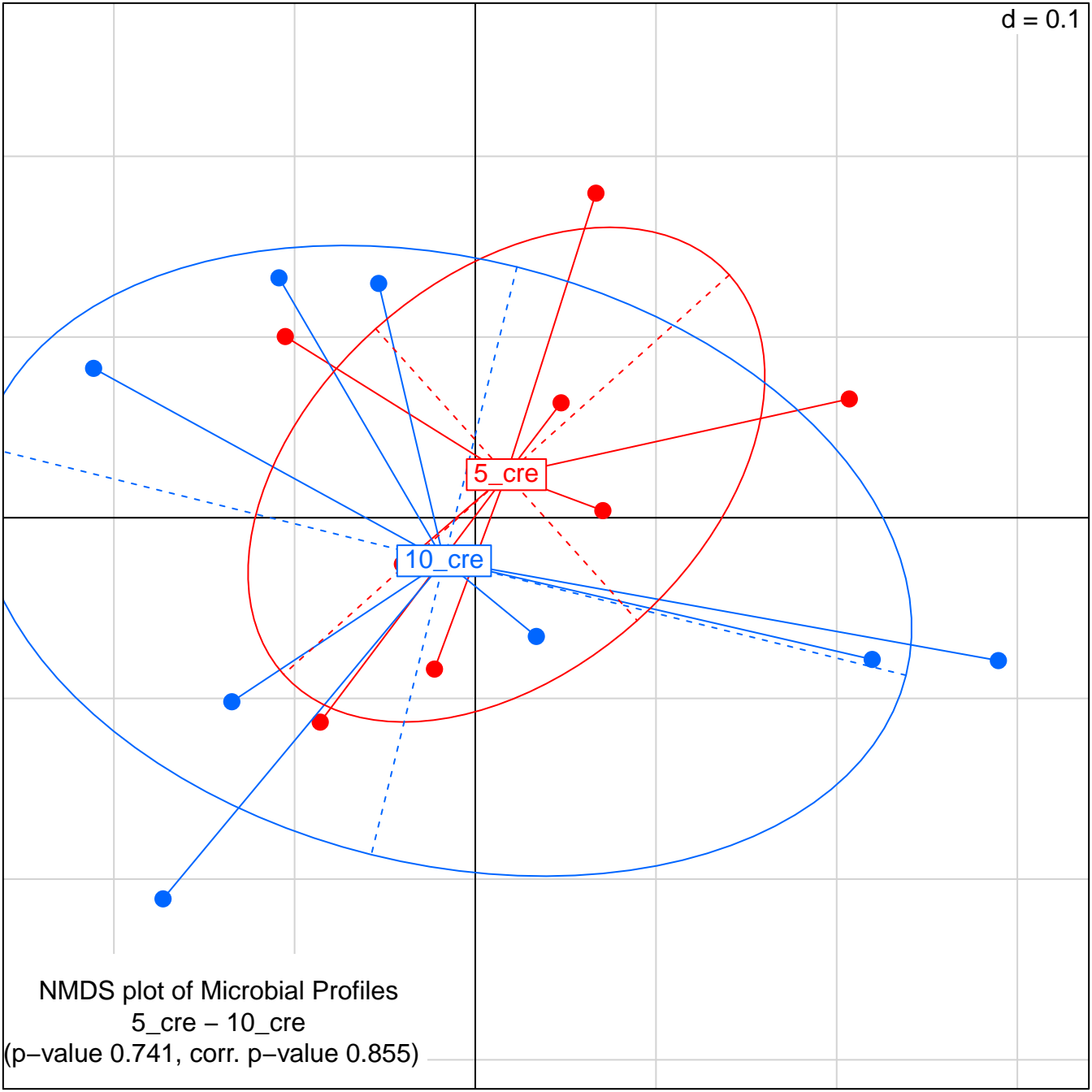


NMDS plot of Microbial Profiles  
5\_cre – 9\_cre  
(p-value 0.142, corr. p-value 0.795)

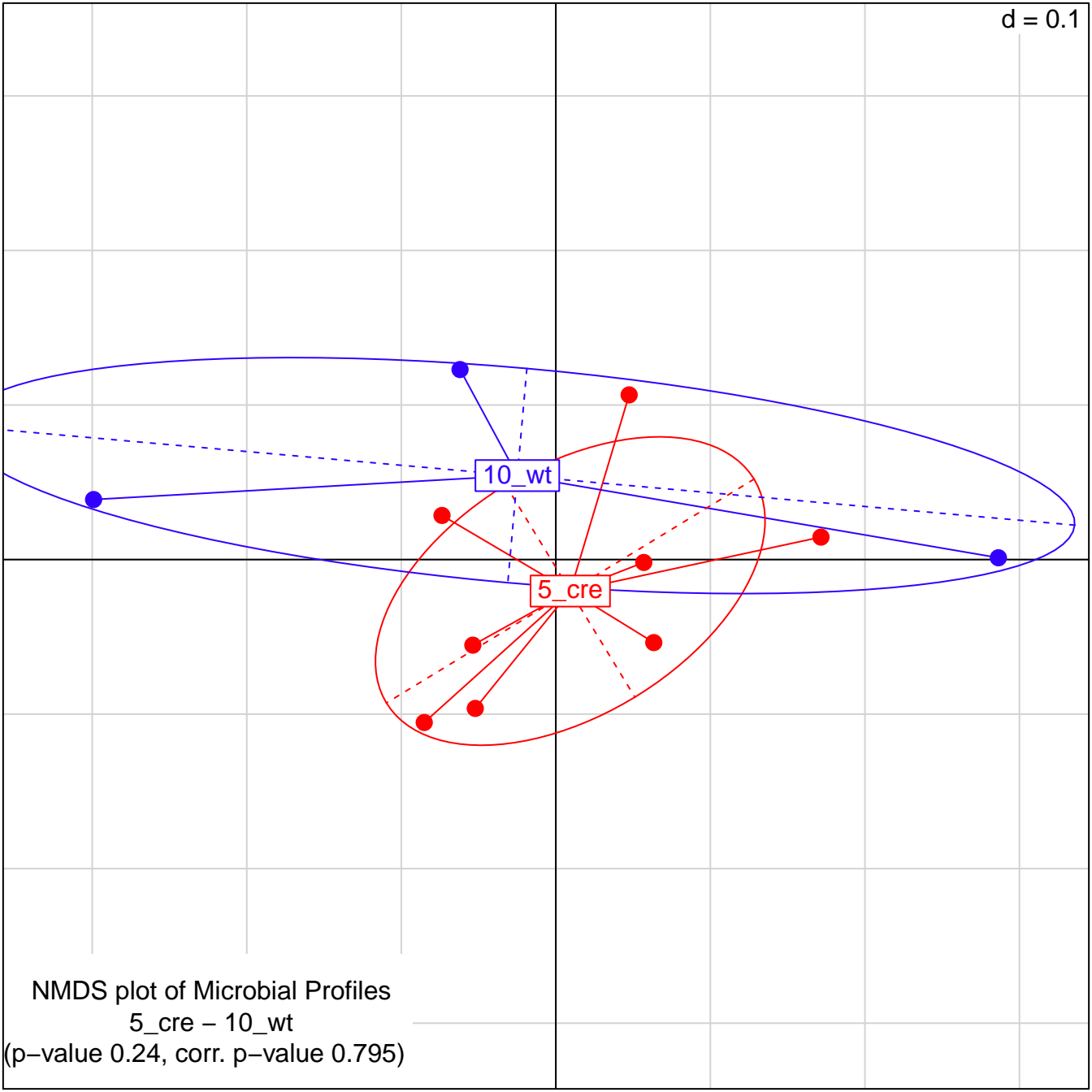
d = 0.1



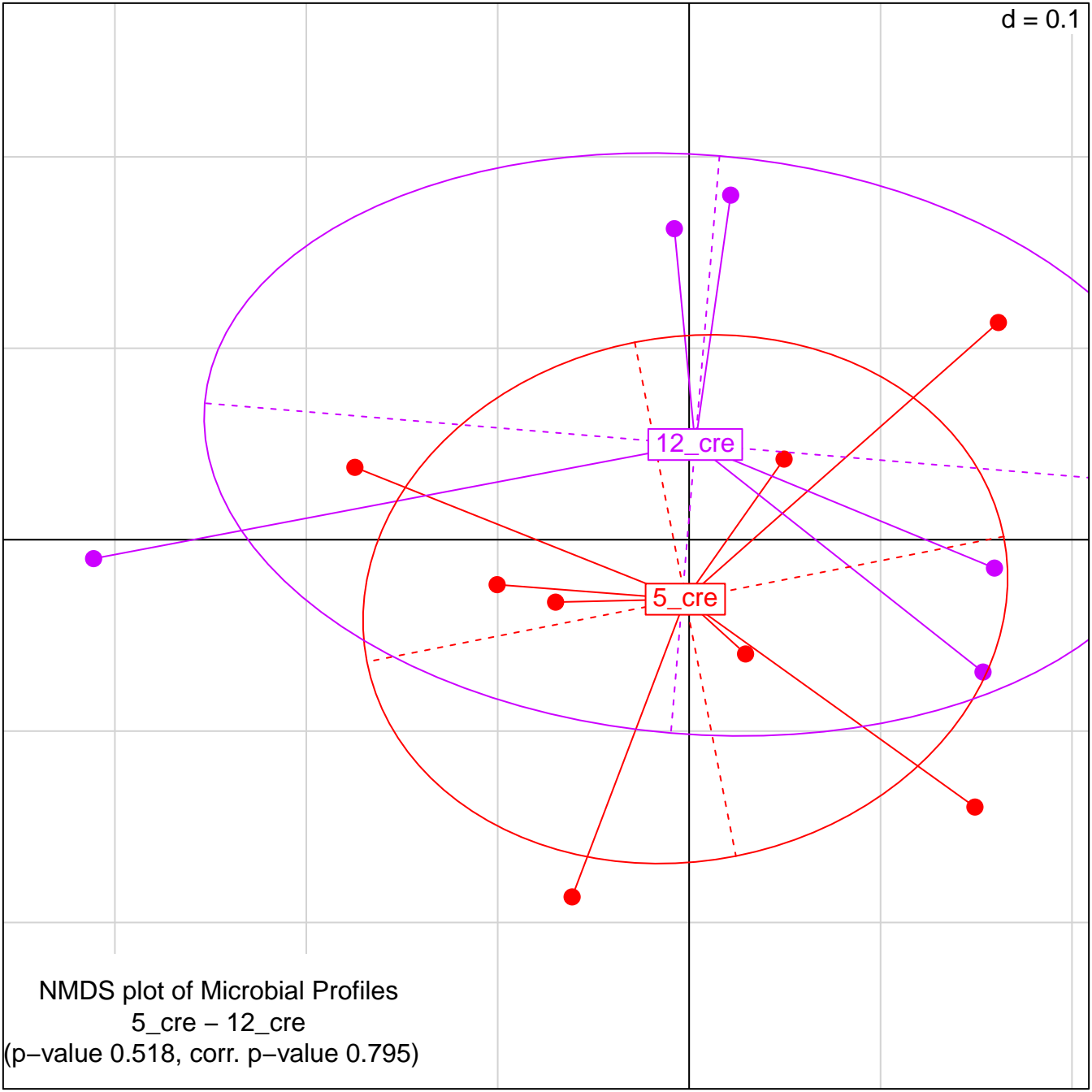
d = 0.1



d = 0.1



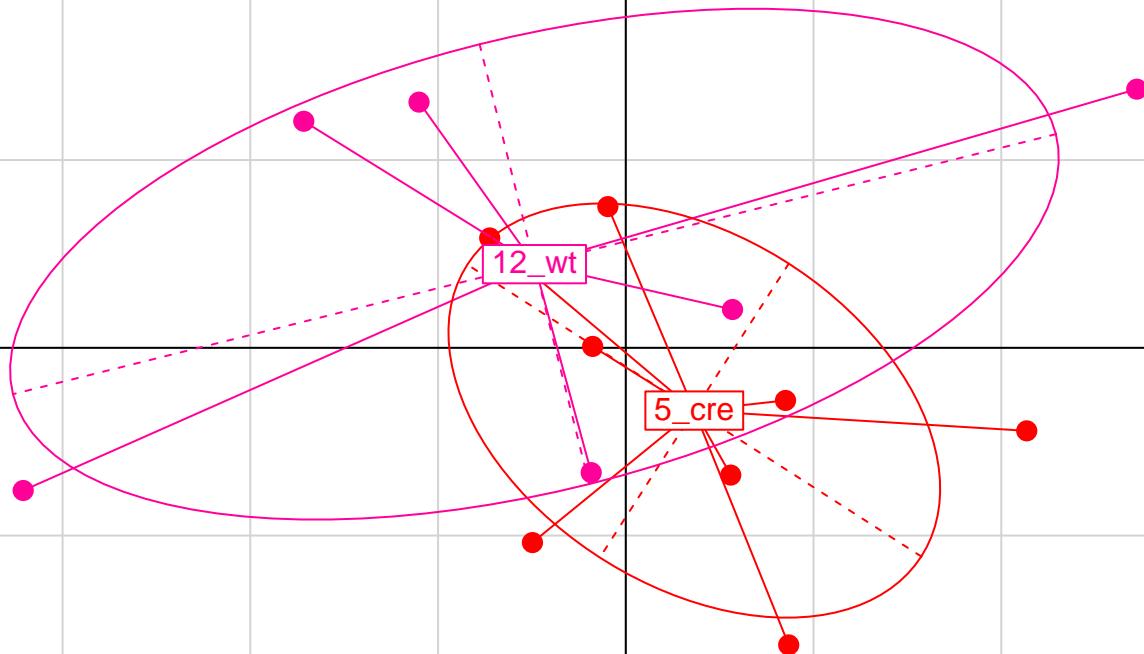
d = 0.1



NMDS plot of Microbial Profiles  
5\_cre – 12\_cre  
(p-value 0.518, corr. p-value 0.795)

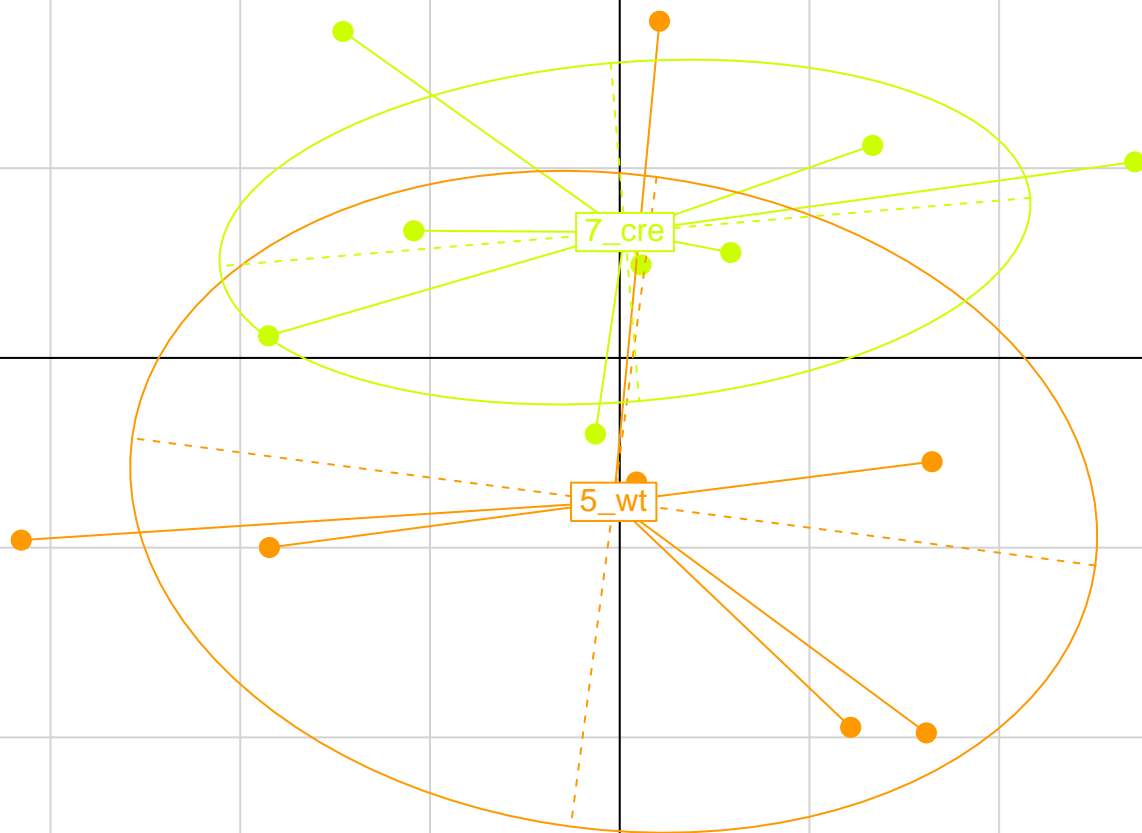


d = 0.1



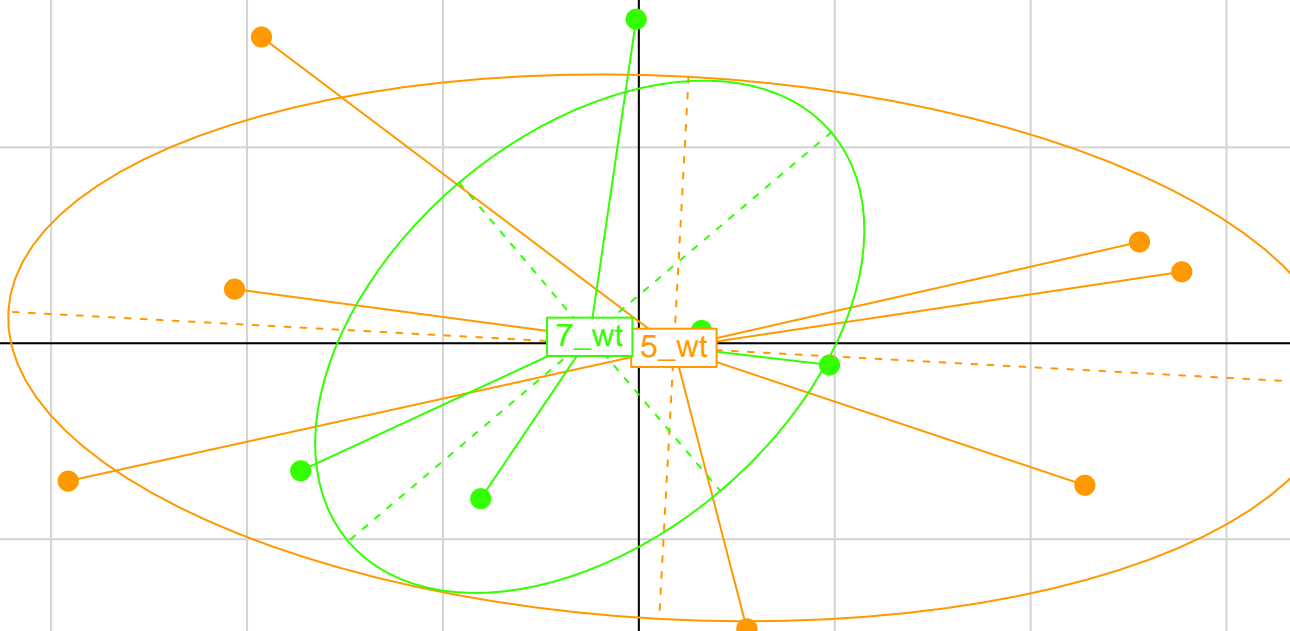
NMDS plot of Microbial Profiles  
5\_cre – 12\_wt  
(p-value 0.202, corr. p-value 0.795)

d = 0.1



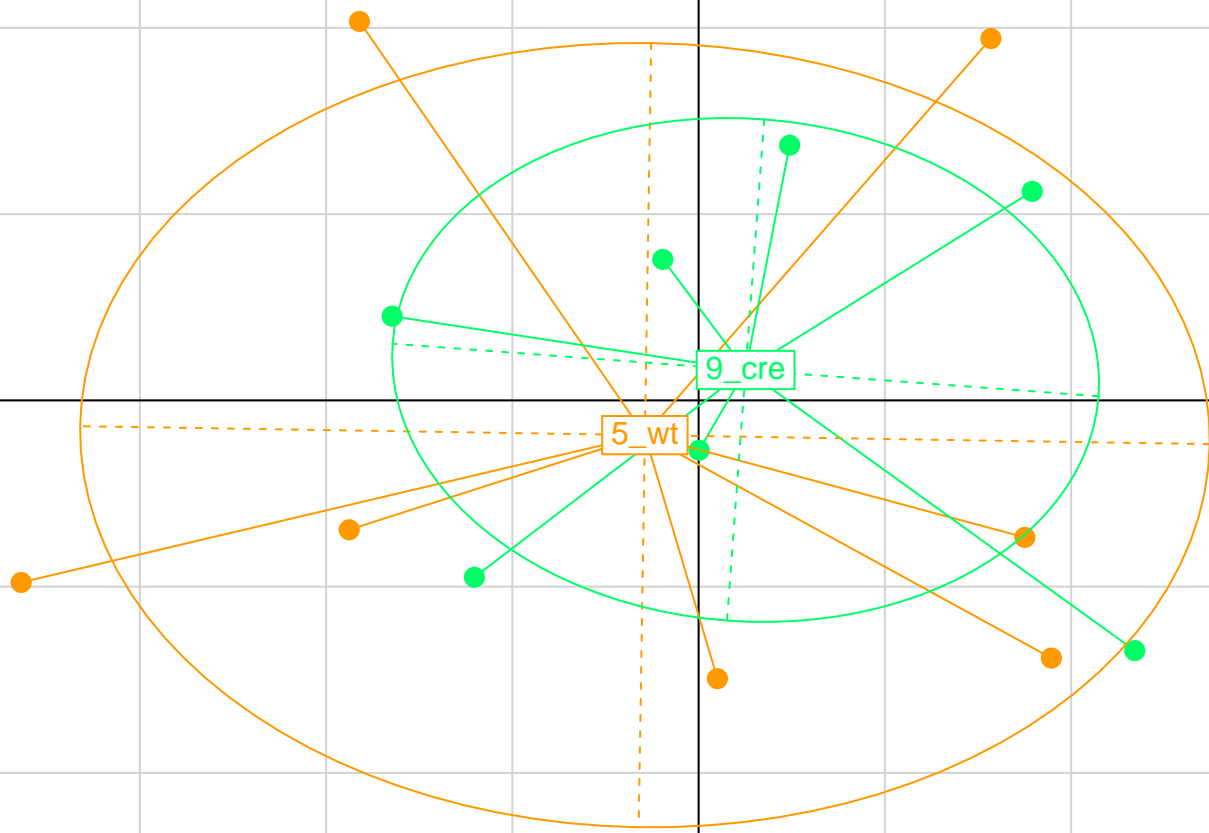
NMDS plot of Microbial Profiles  
5\_wt – 7\_cre  
(p-value 0.019, corr. p-value 0.795)

d = 0.1



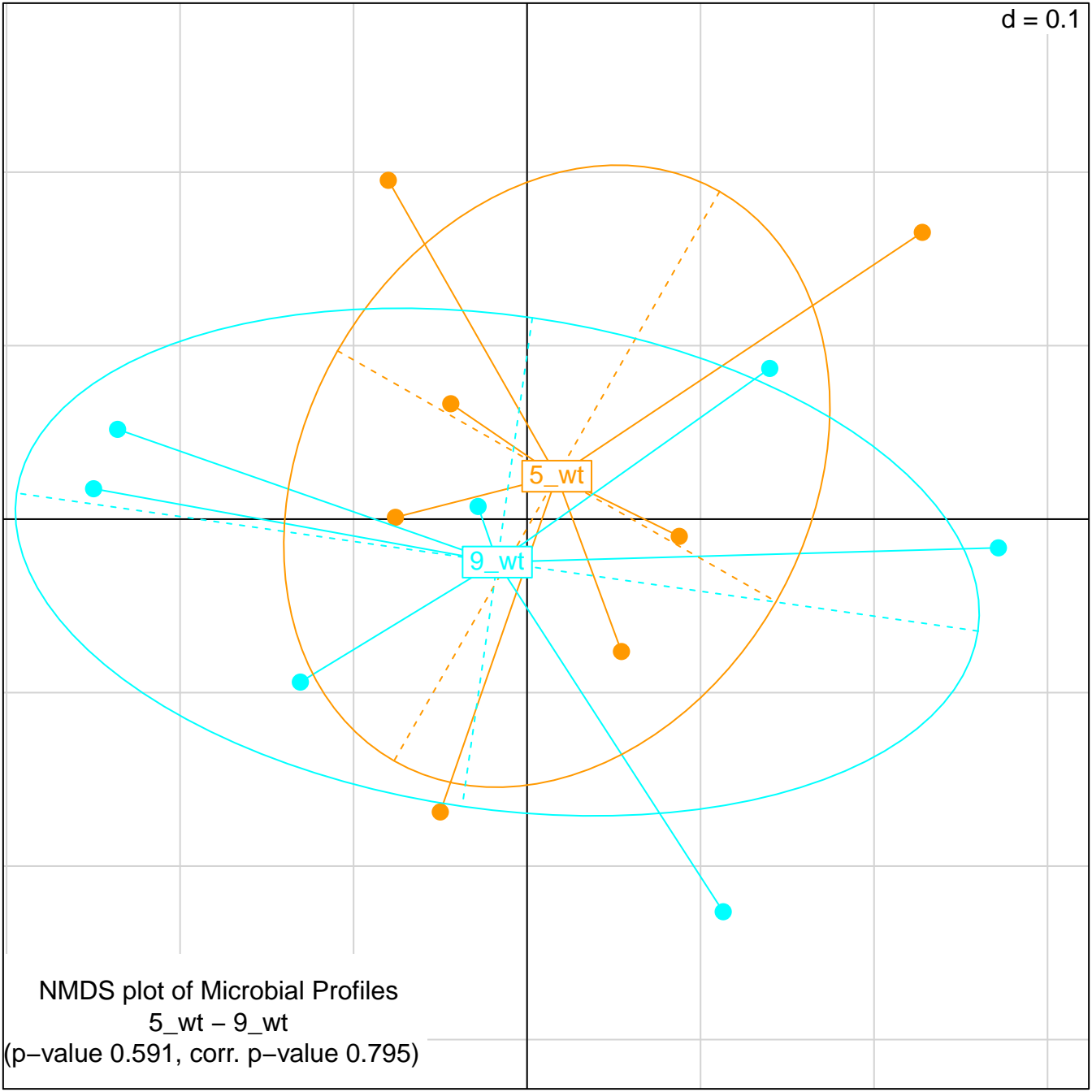
NMDS plot of Microbial Profiles  
5\_wt – 7\_wt  
(p-value 0.632, corr. p-value 0.795)

d = 0.1

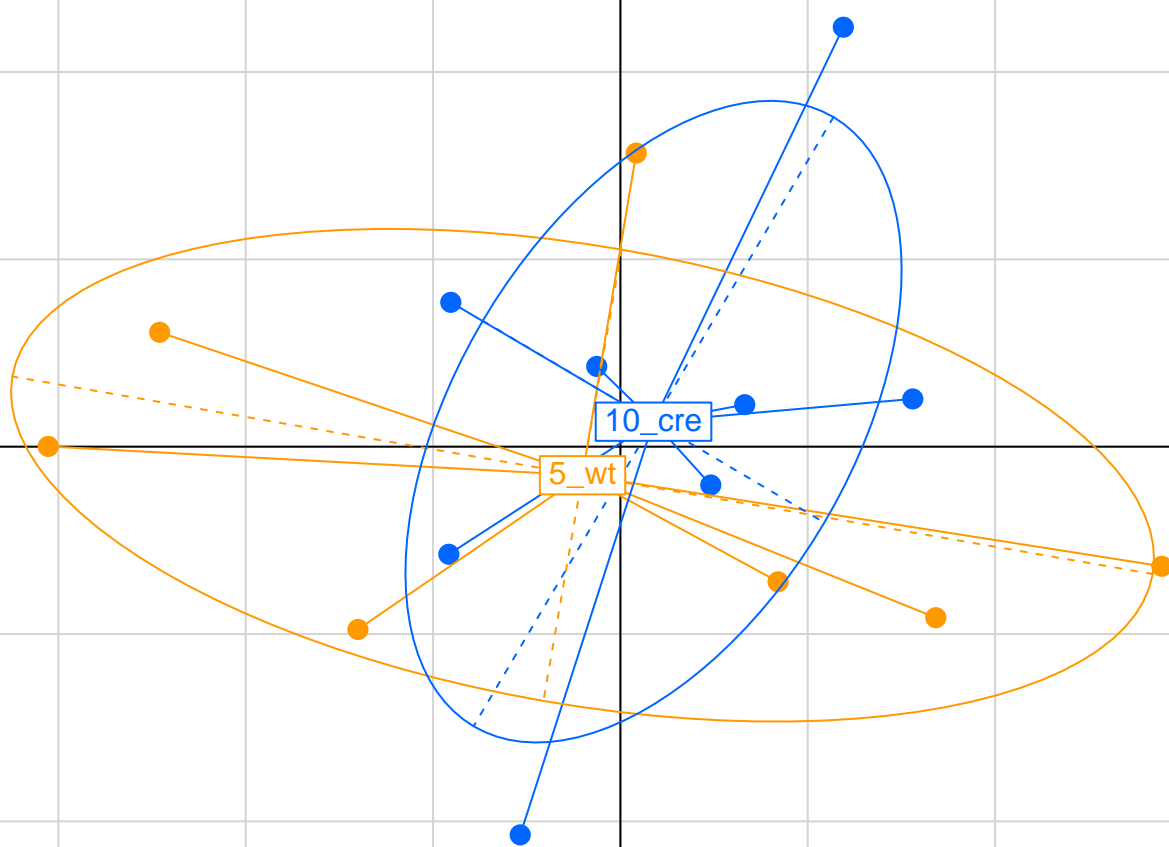


NMDS plot of Microbial Profiles  
5\_wt – 9\_cre  
(p-value 0.485, corr. p-value 0.795)

d = 0.1

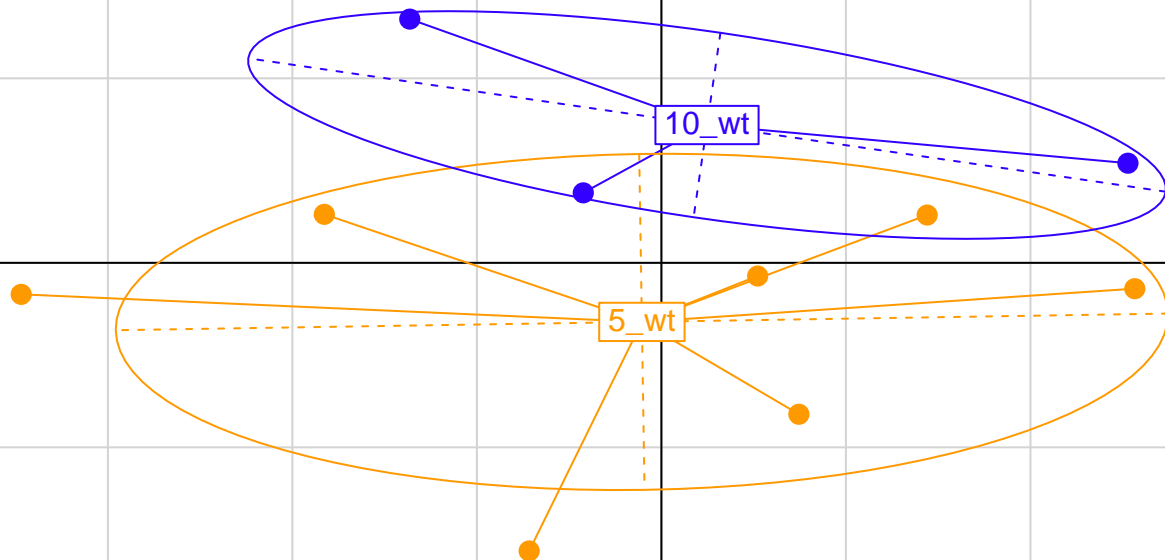


d = 0.1



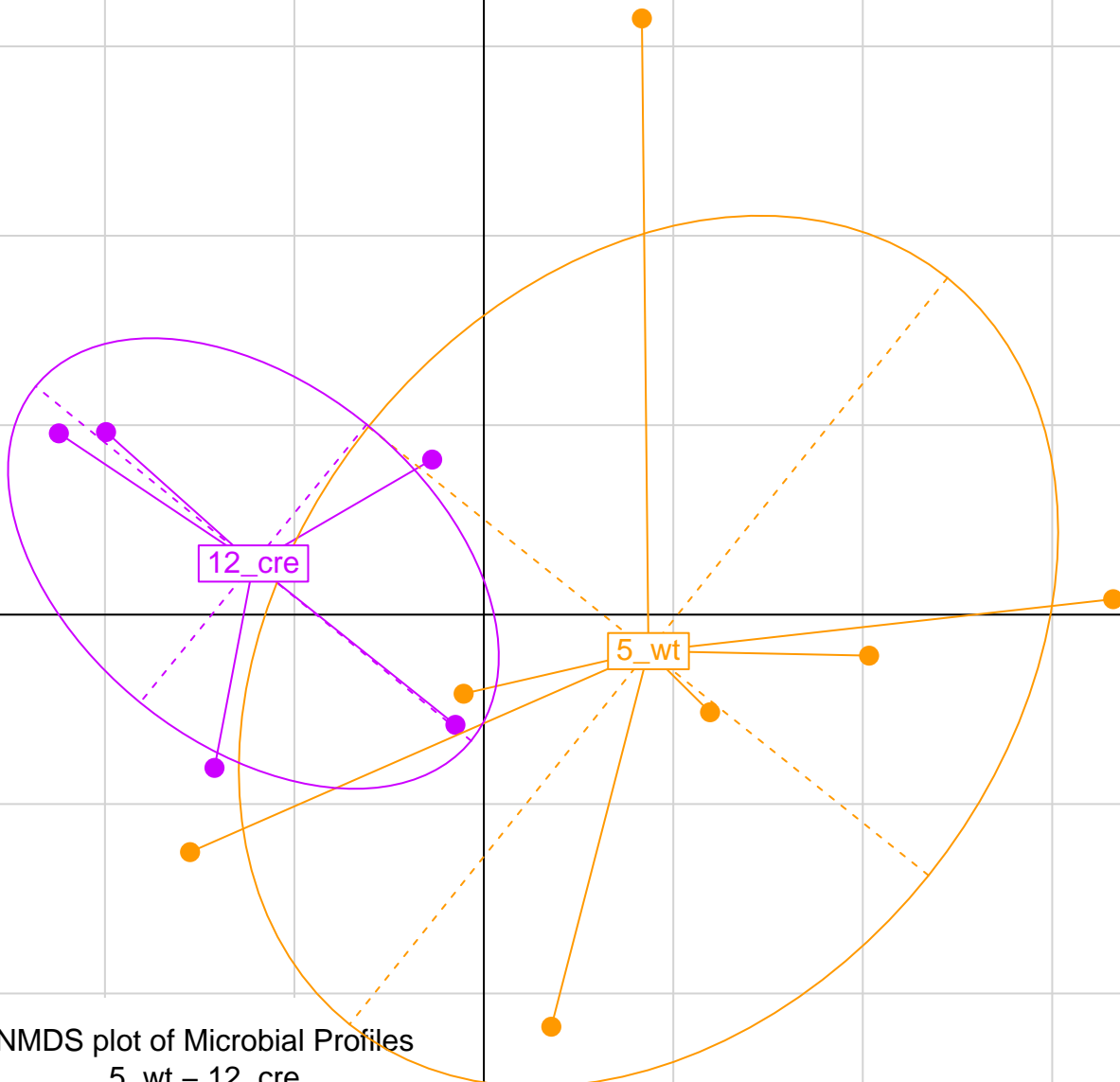
NMDS plot of Microbial Profiles  
5\_wt – 10\_cre  
(p-value 0.564, corr. p-value 0.795)

d = 0.1



NMDS plot of Microbial Profiles  
5\_wt - 10\_wt  
(p-value 0.726, corr. p-value 0.855)

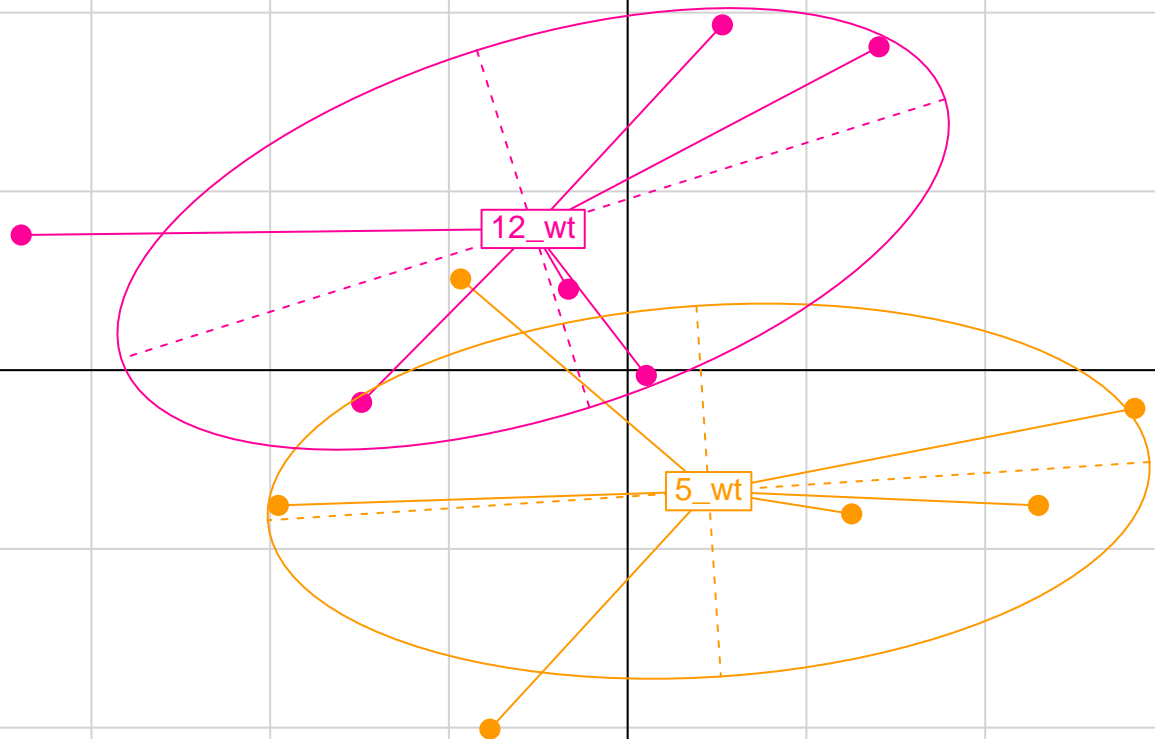
d = 0.1



NMDS plot of Microbial Profiles  
5\_wt – 12\_cre  
(p-value 0.066, corr. p-value 0.795)

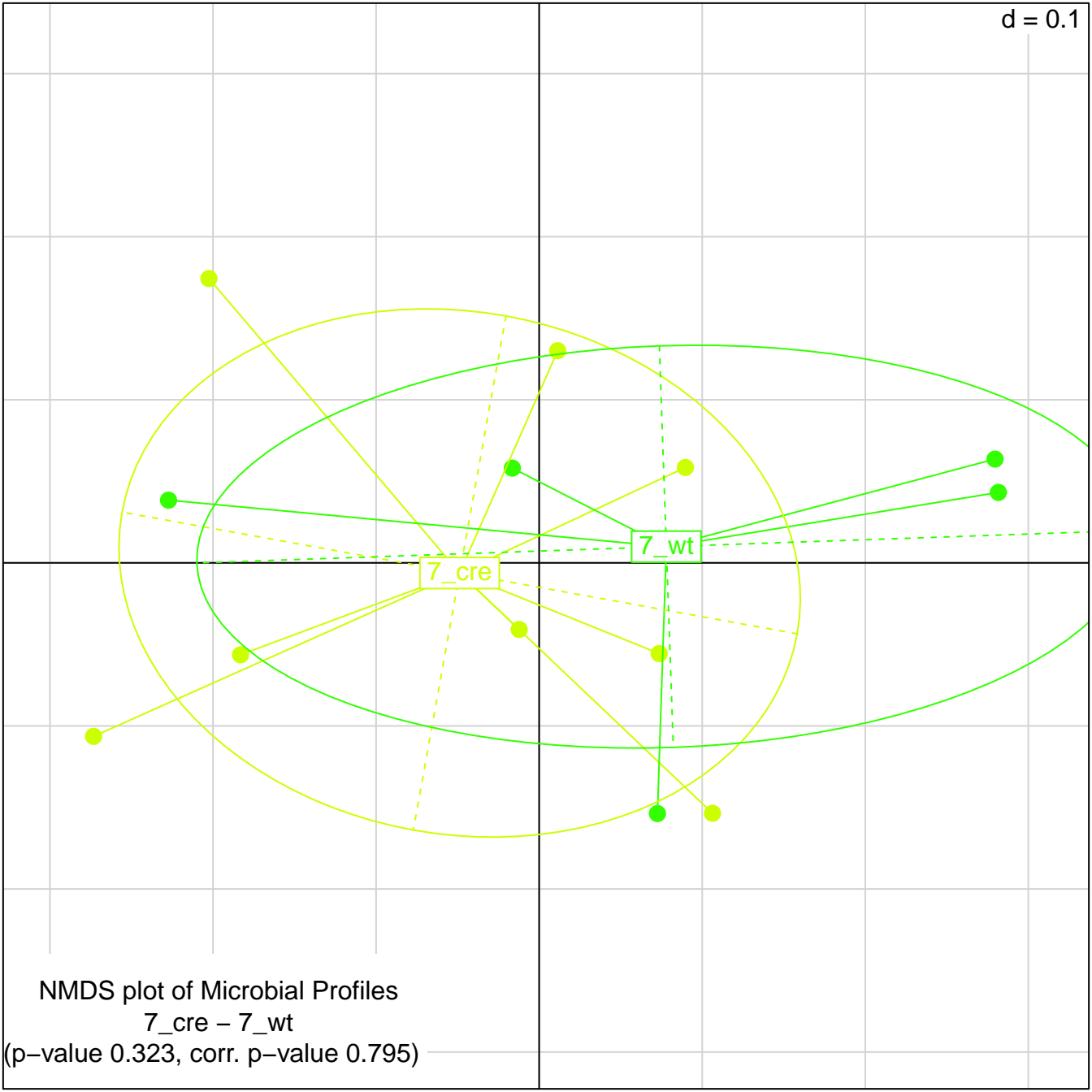


d = 0.1



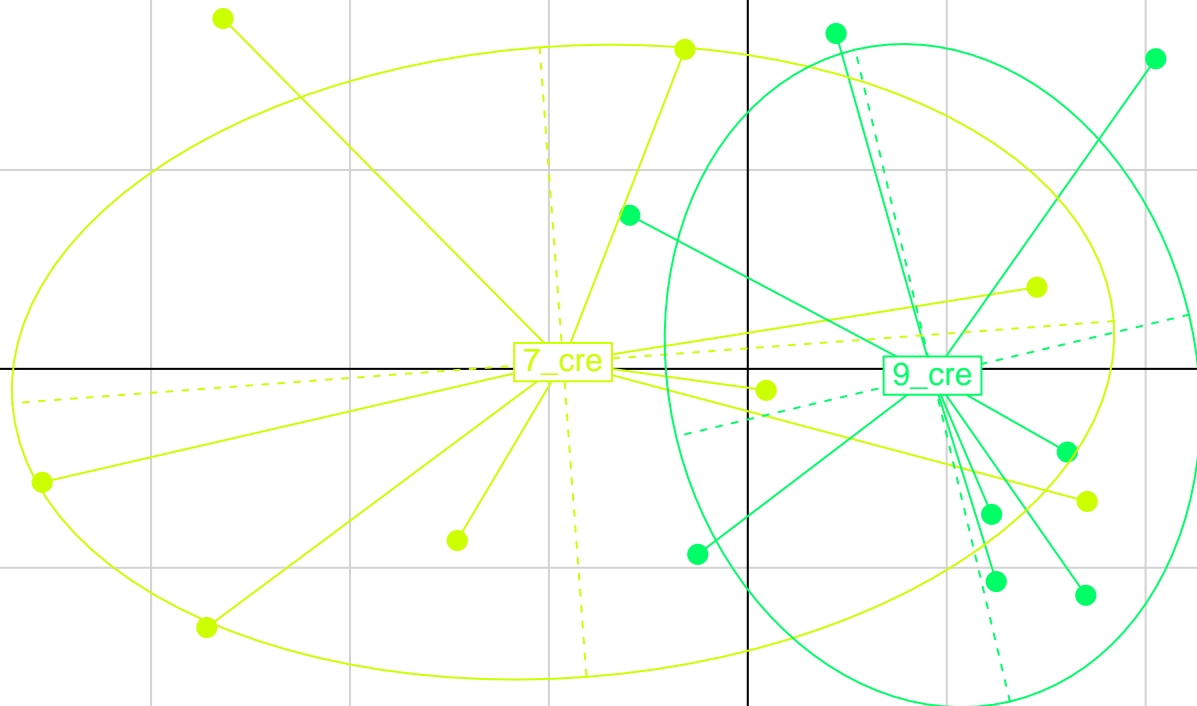
NMDS plot of Microbial Profiles  
5\_wt – 12\_wt  
(p-value 0.072, corr. p-value 0.795)

d = 0.1



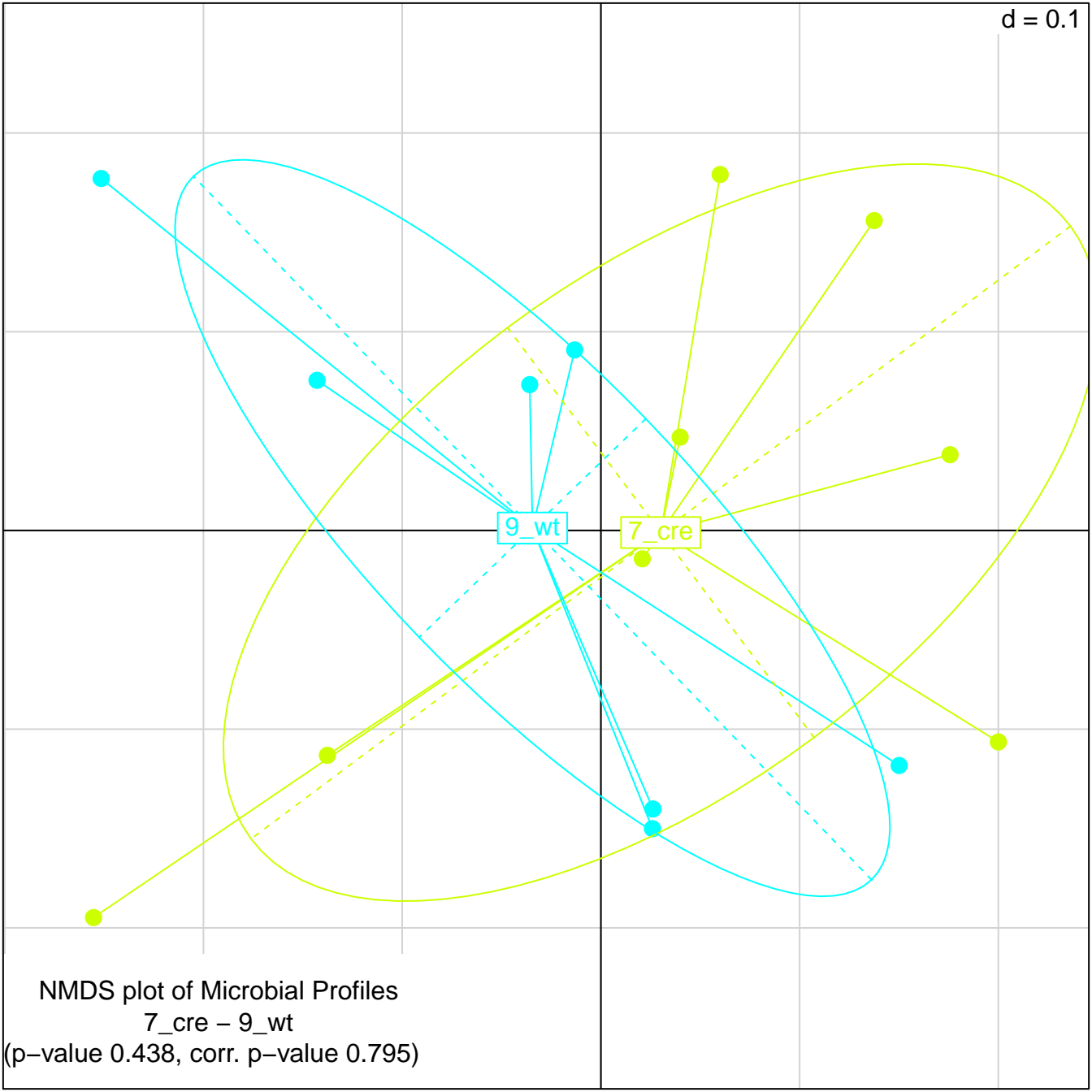
NMDS plot of Microbial Profiles  
7\_cre – 7\_wt  
(p-value 0.323, corr. p-value 0.795)

d = 0.1



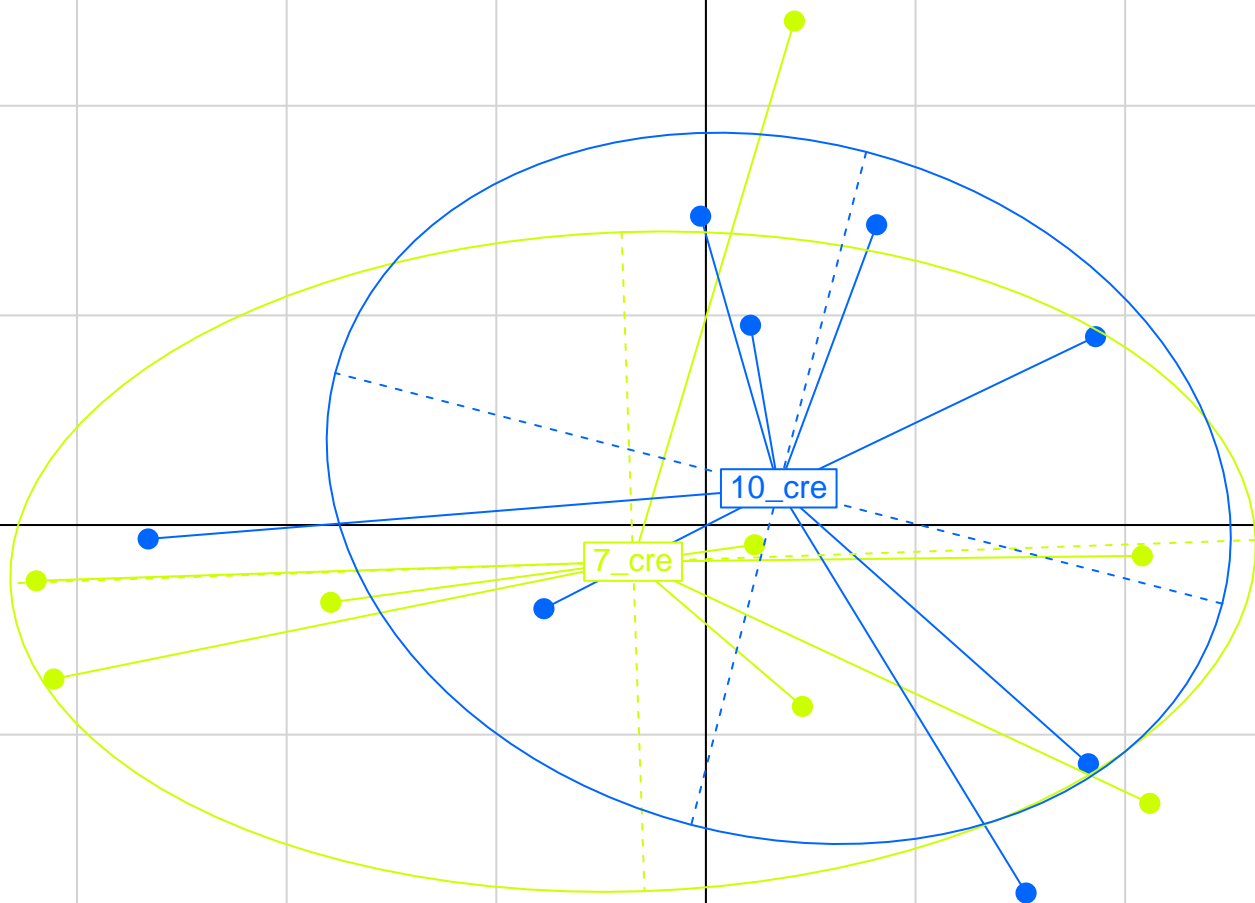
NMDS plot of Microbial Profiles  
7\_cre – 9\_cre  
(p-value 0.055, corr. p-value 0.795)

d = 0.1



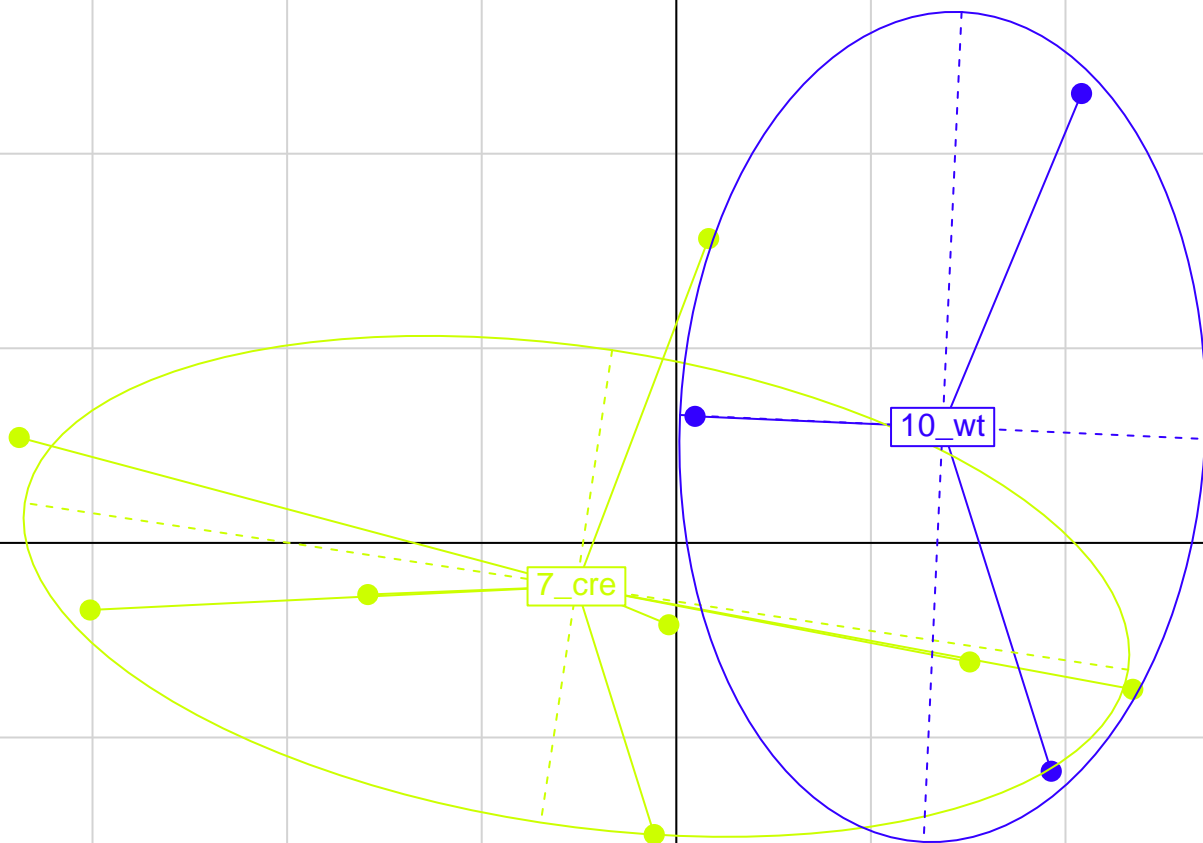
NMDS plot of Microbial Profiles  
7\_cre – 9\_wt  
(p-value 0.438, corr. p-value 0.795)

d = 0.1



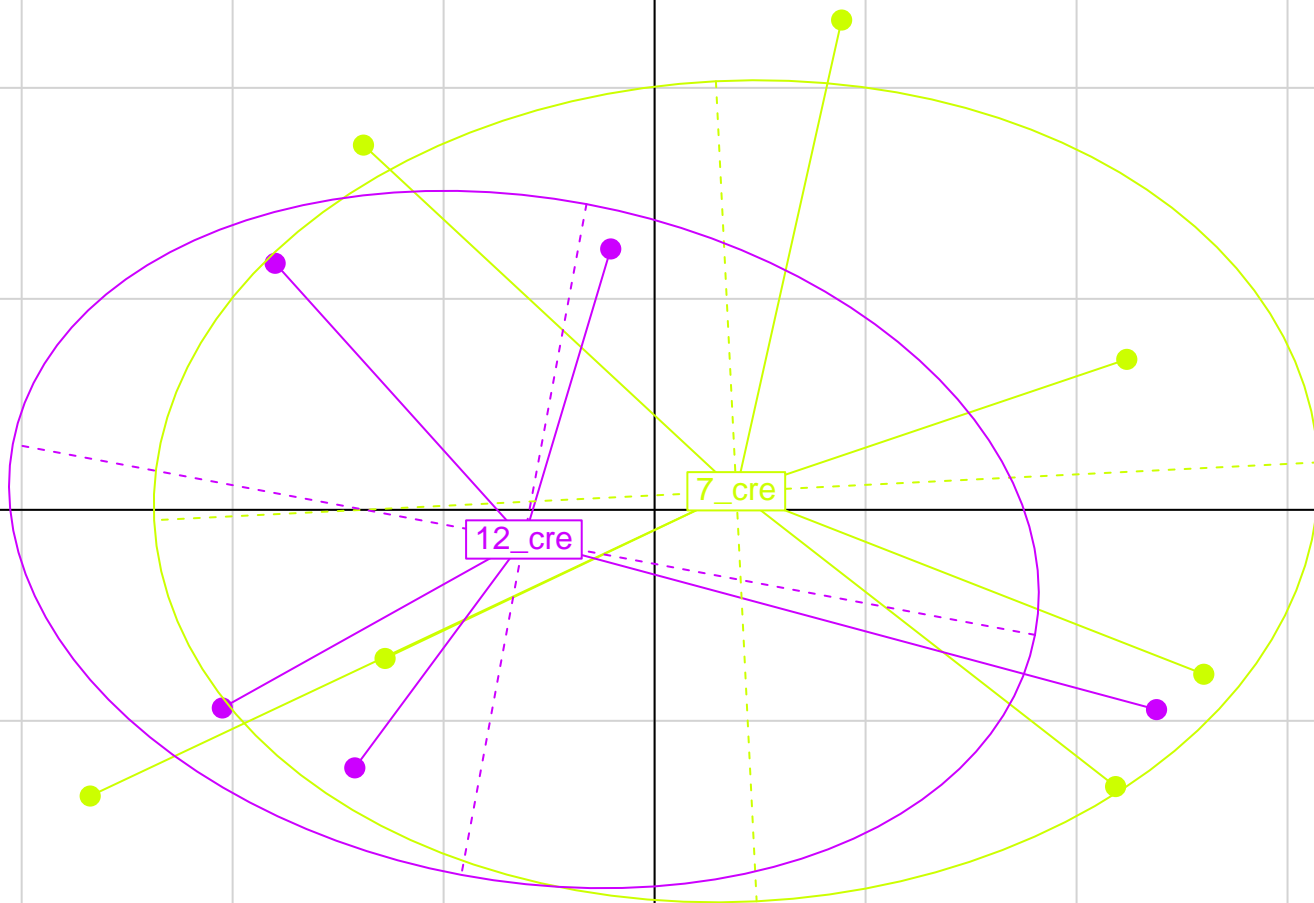
NMDS plot of Microbial Profiles  
7\_cre – 10\_cre  
(p-value 0.332, corr. p-value 0.795)

d = 0.1



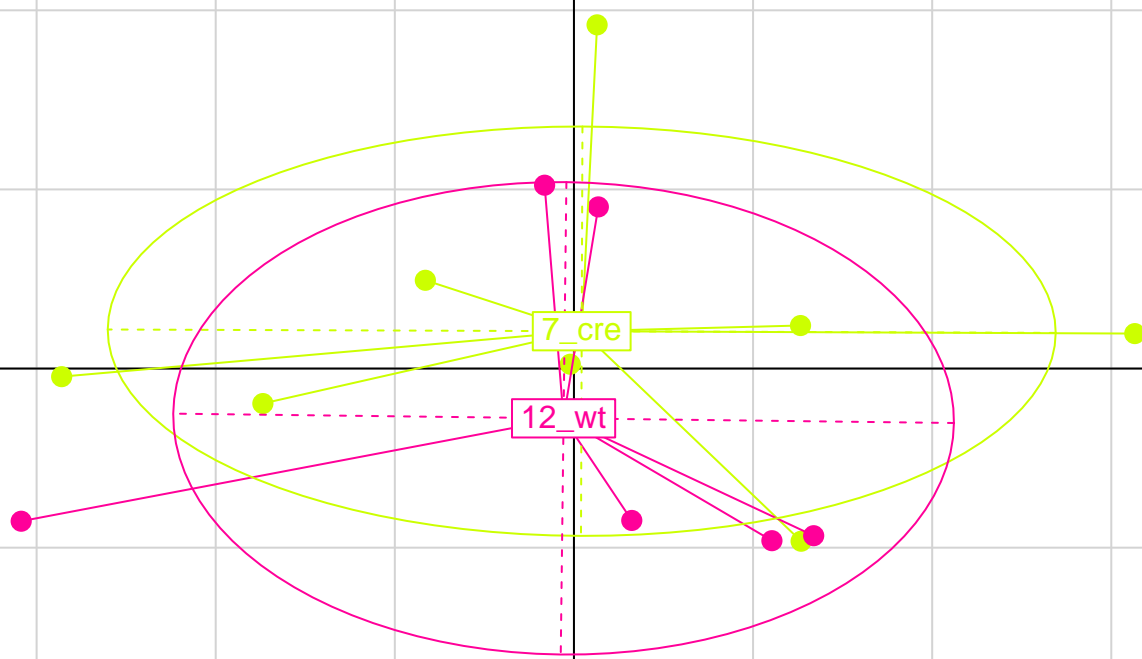
NMDS plot of Microbial Profiles  
7\_cre – 10\_wt  
(p-value 0.358, corr. p-value 0.795)

d = 0.1



NMDS plot of Microbial Profiles  
7\_cre – 12\_cre  
(p-value 0.521, corr. p-value 0.795)

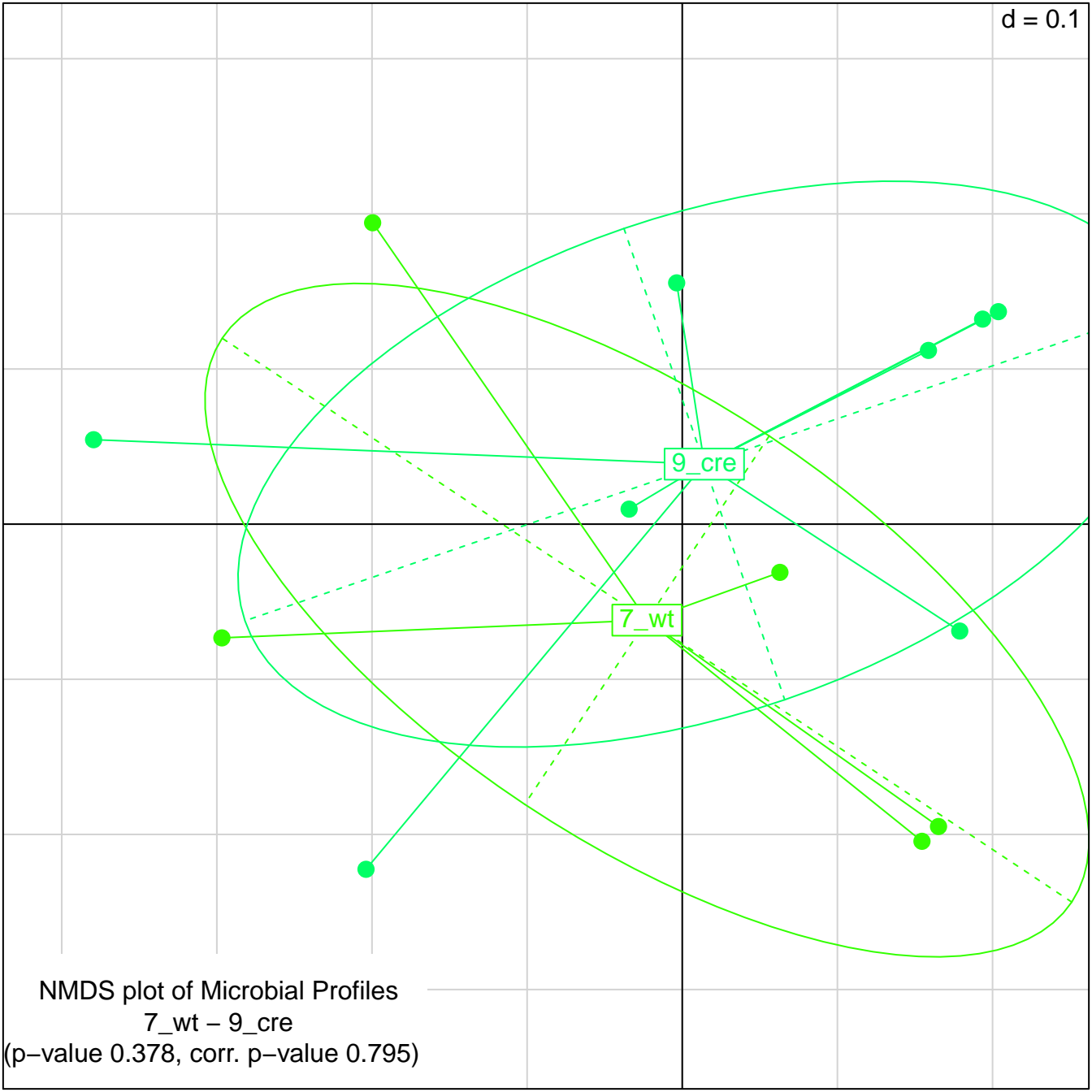
d = 0.1



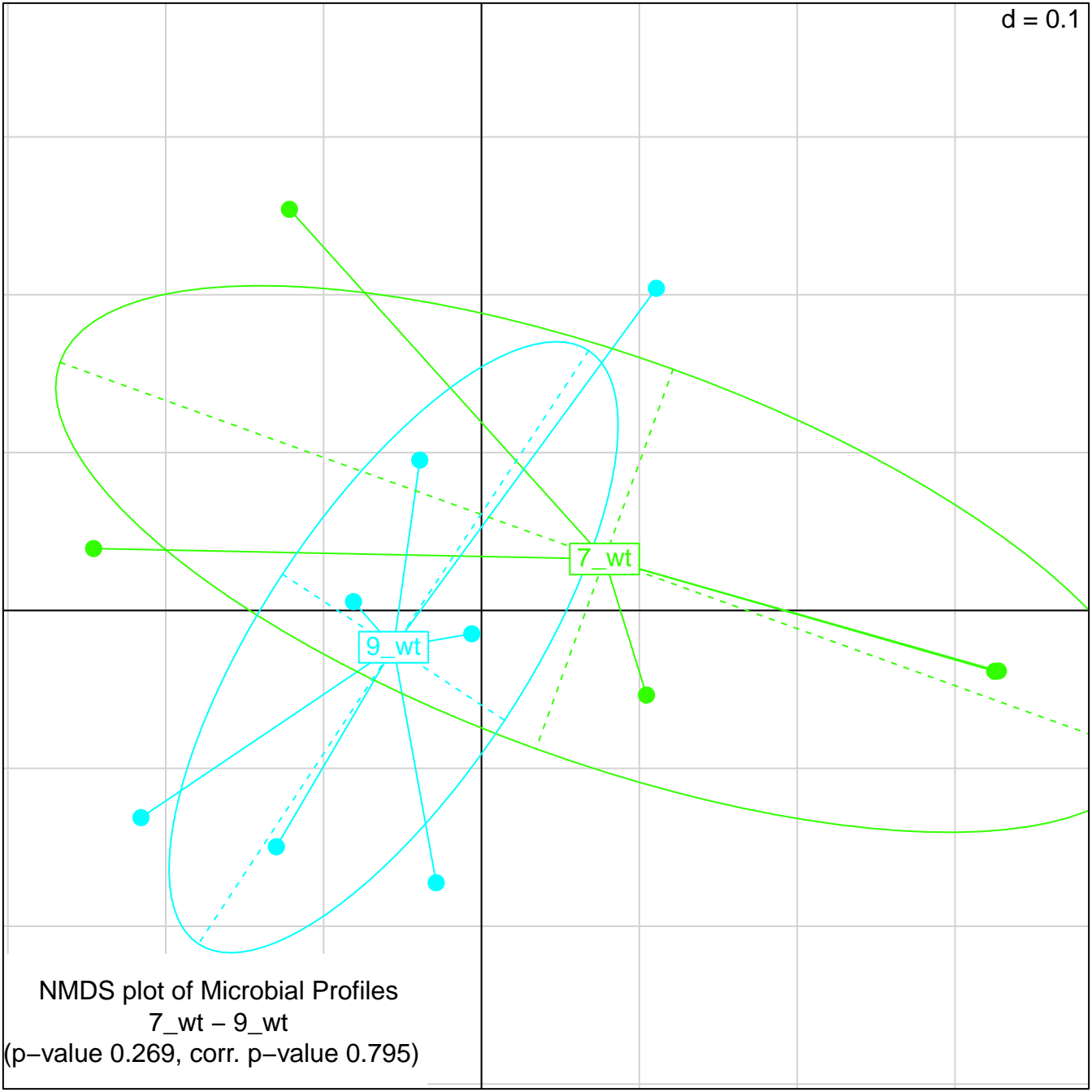
NMDS plot of Microbial Profiles  
7\_cre – 12\_wt  
(p-value 0.829, corr. p-value 0.892674418604651)



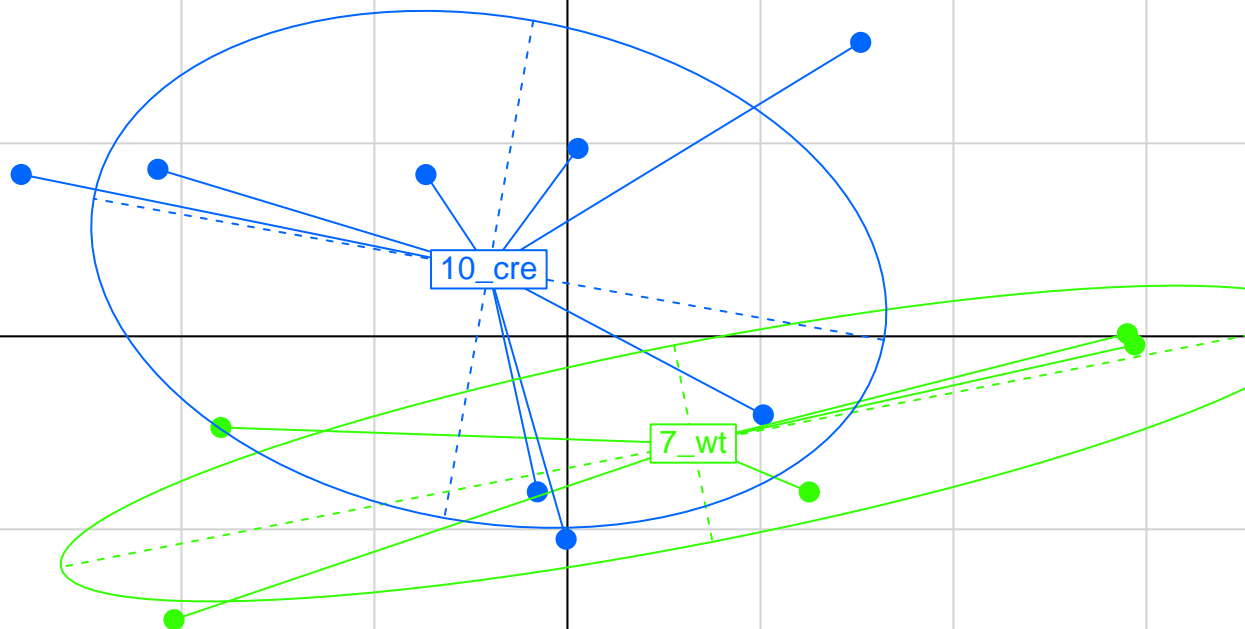
d = 0.1



d = 0.1

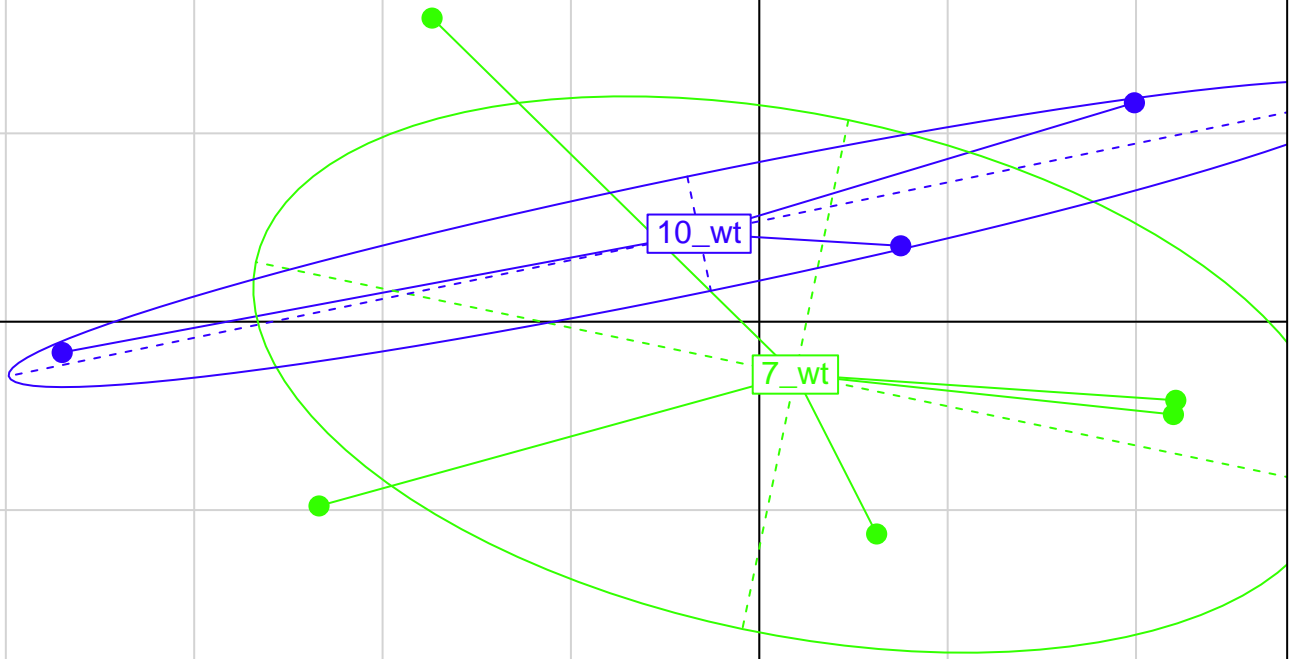


d = 0.1



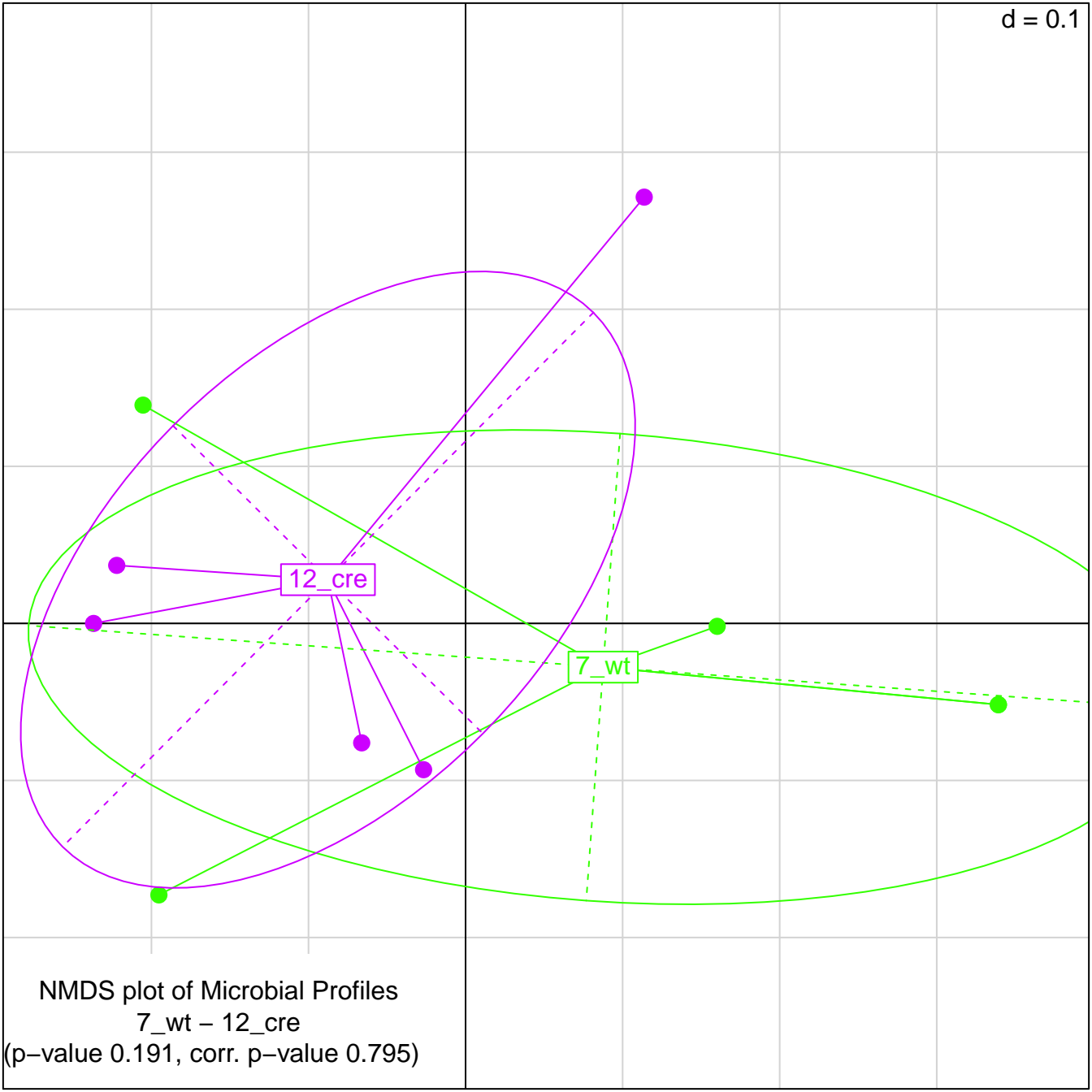
NMDS plot of Microbial Profiles  
7\_wt - 10\_cre  
(p-value 0.222, corr. p-value 0.795)

d = 0.1

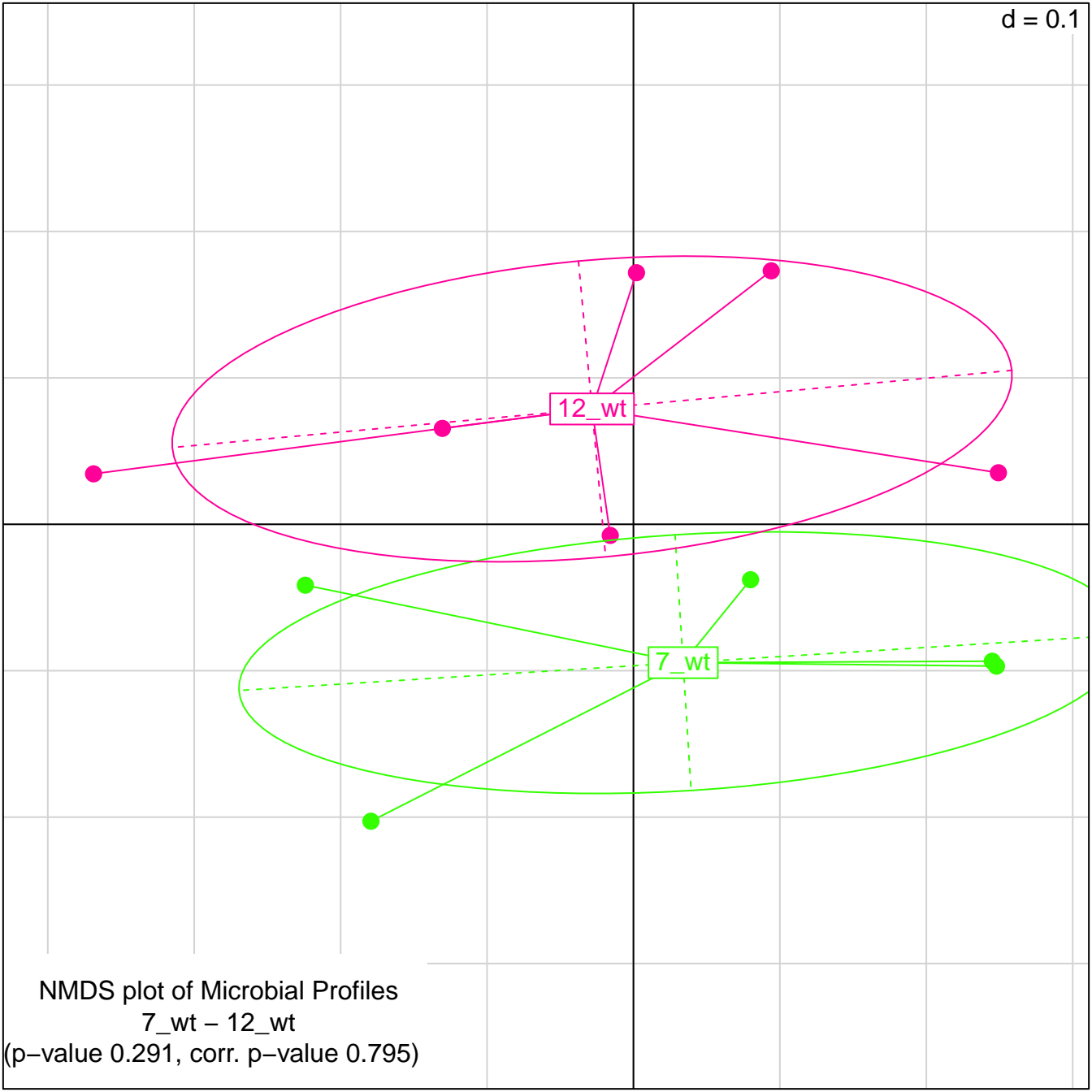


NMDS plot of Microbial Profiles  
7\_wt – 10\_wt  
(p-value 0.785, corr. p-value 0.883125)

d = 0.1

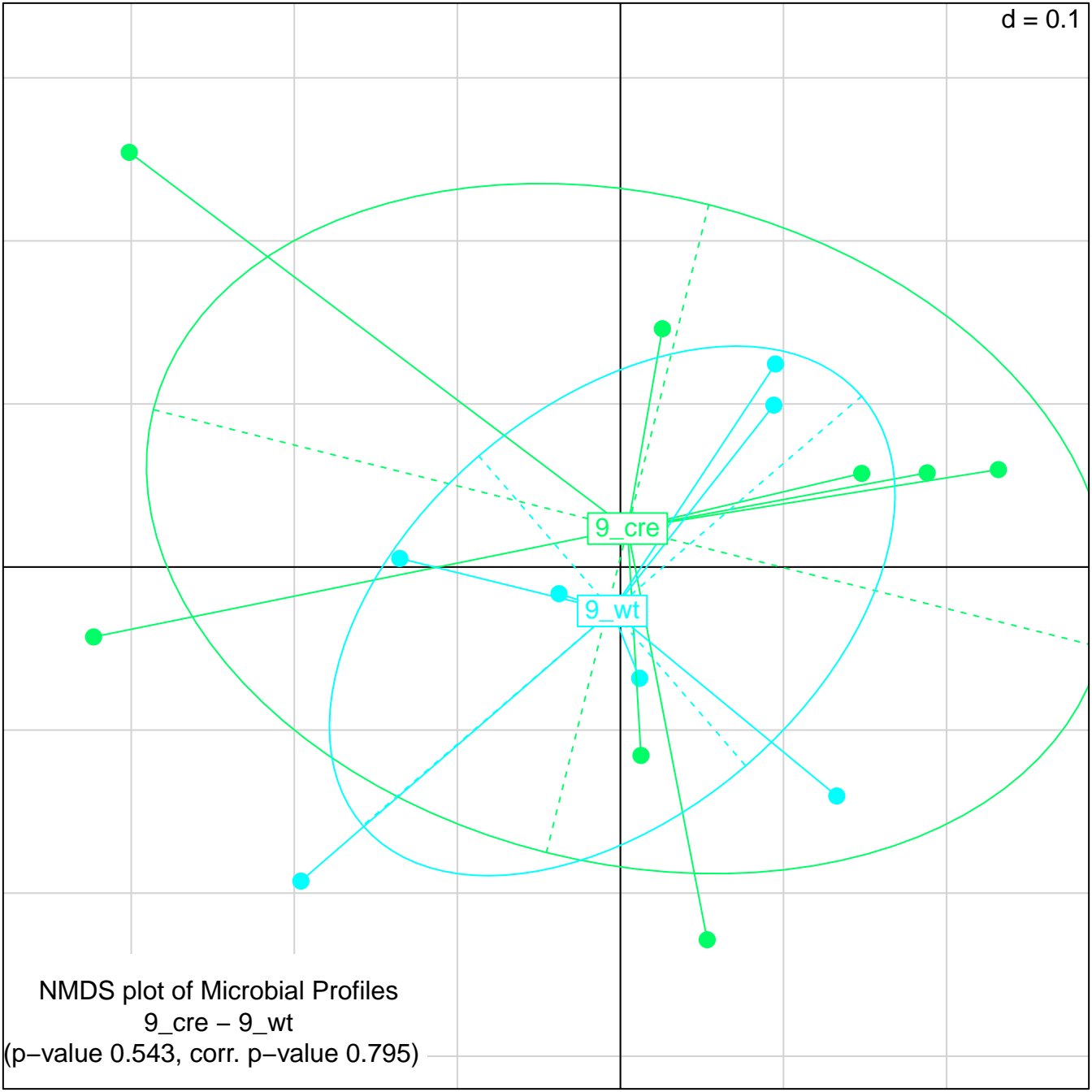


d = 0.1



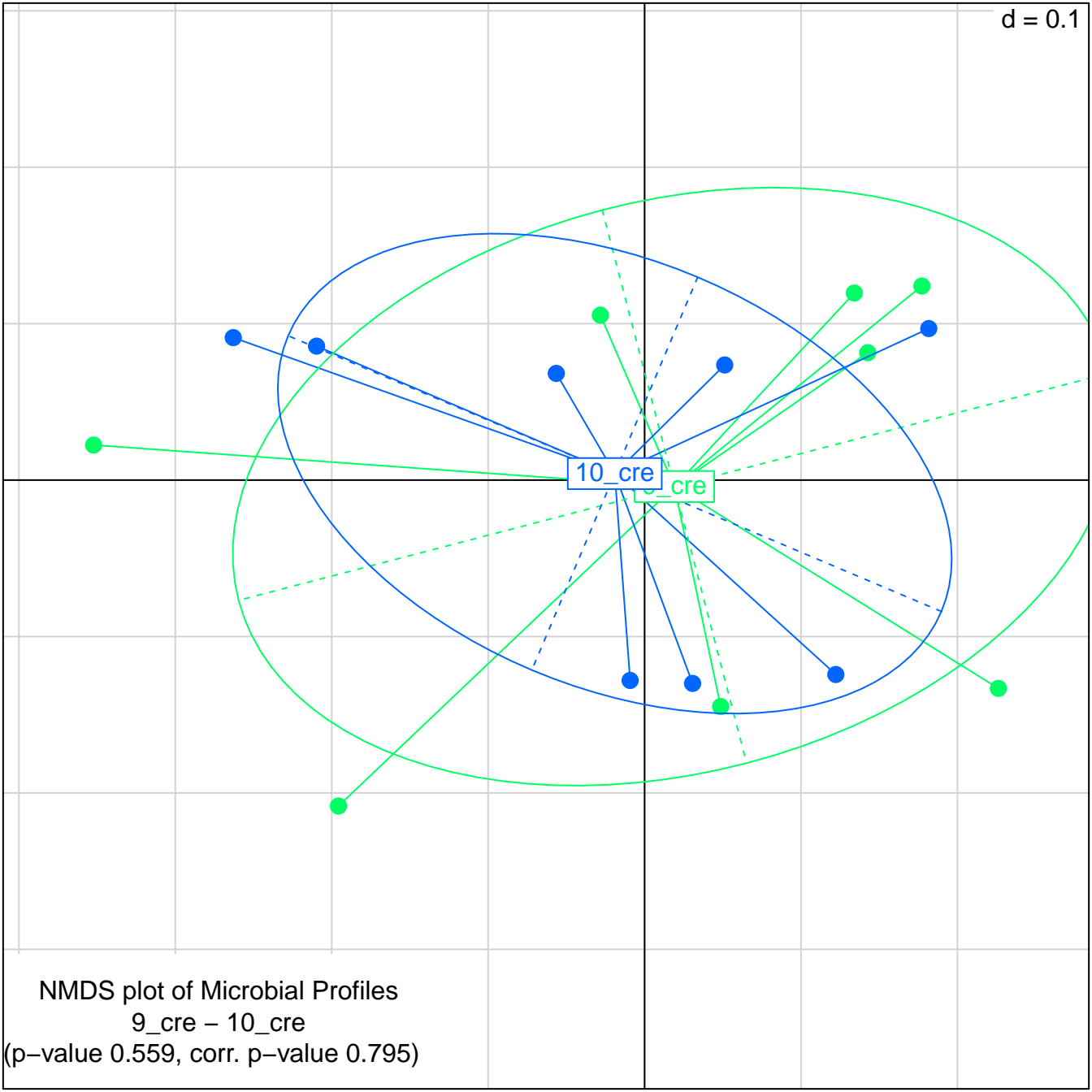
NMDS plot of Microbial Profiles  
7\_wt – 12\_wt  
(p-value 0.291, corr. p-value 0.795)

d = 0.1



NMDS plot of Microbial Profiles  
9\_cre – 9\_wt  
(p-value 0.543, corr. p-value 0.795)

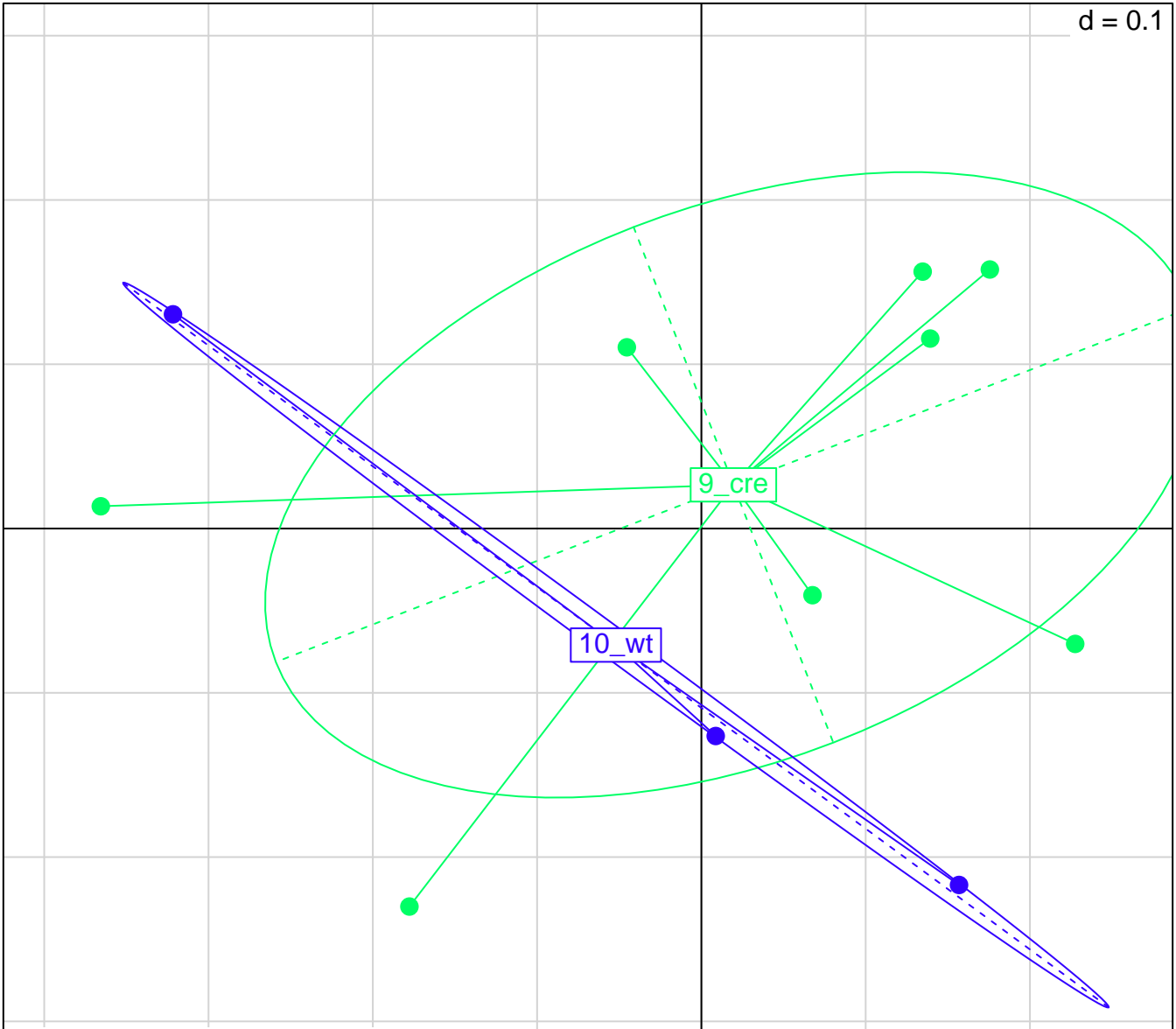
d = 0.1



NMDS plot of Microbial Profiles  
9\_cre – 10\_cre  
(p-value 0.559, corr. p-value 0.795)

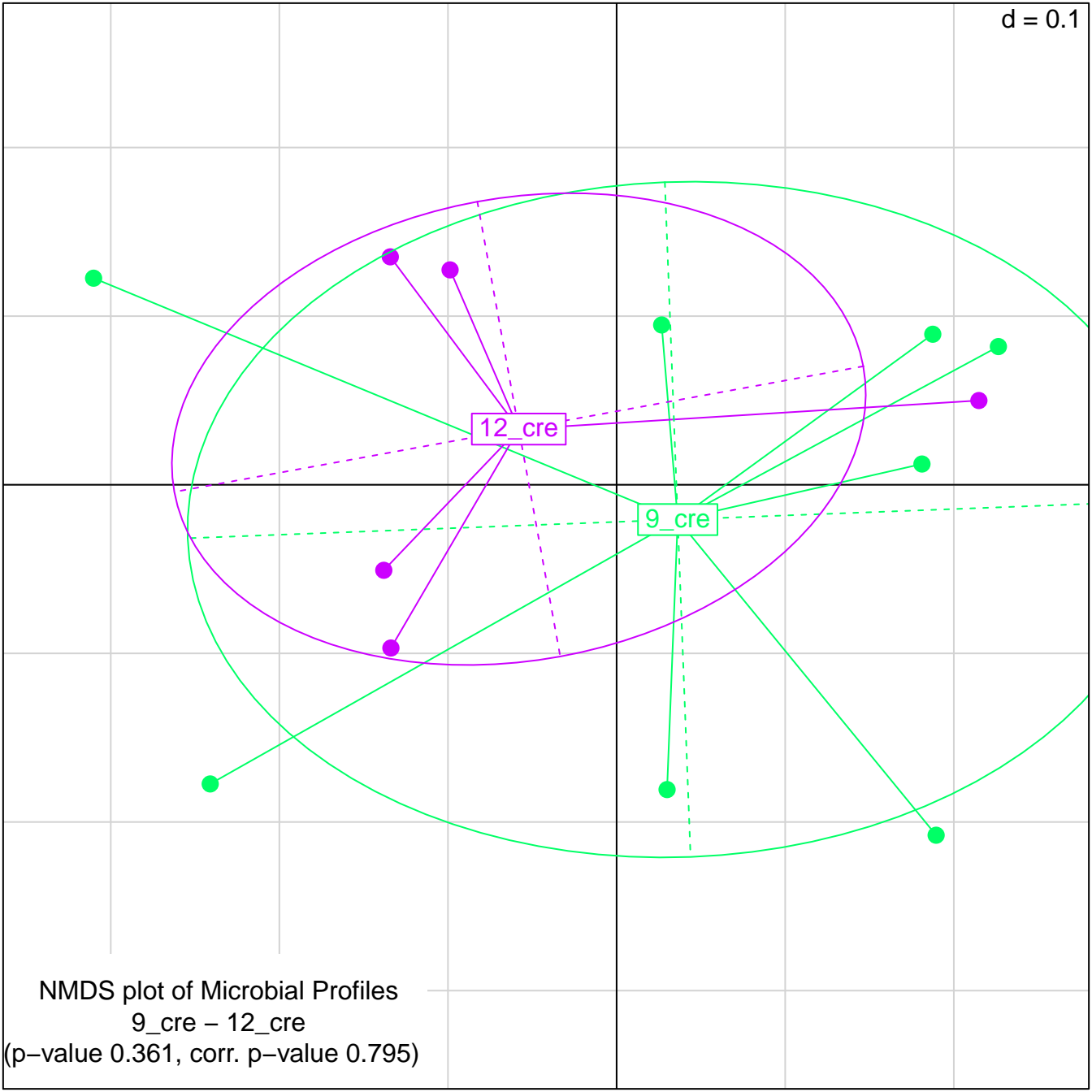


d = 0.1



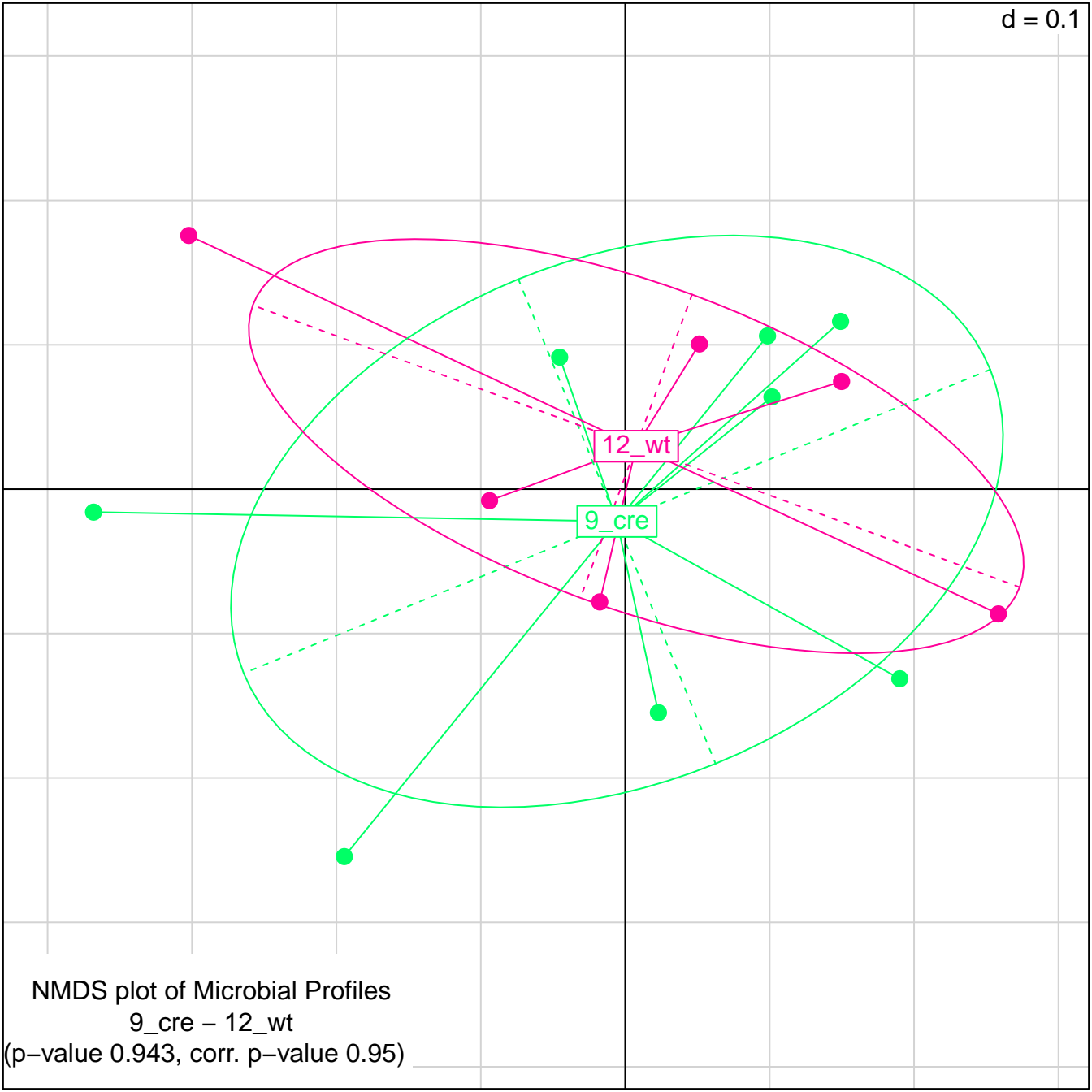
NMDS plot of Microbial Profiles  
9\_cre – 10\_wt  
(p-value 0.608, corr. p-value 0.795)

d = 0.1



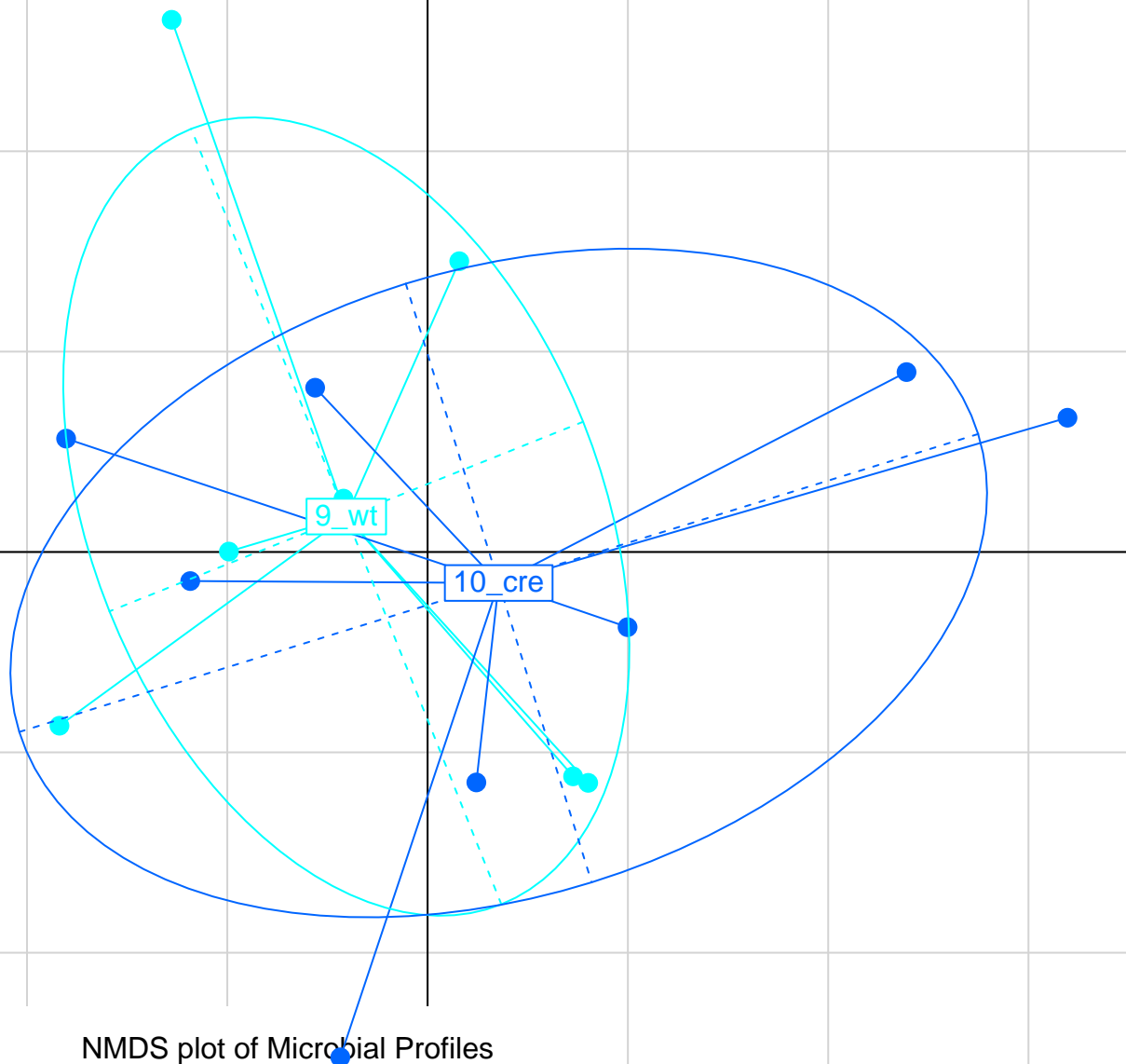
NMDS plot of Microbial Profiles  
9\_cre – 12\_cre  
(p-value 0.361, corr. p-value 0.795)

d = 0.1



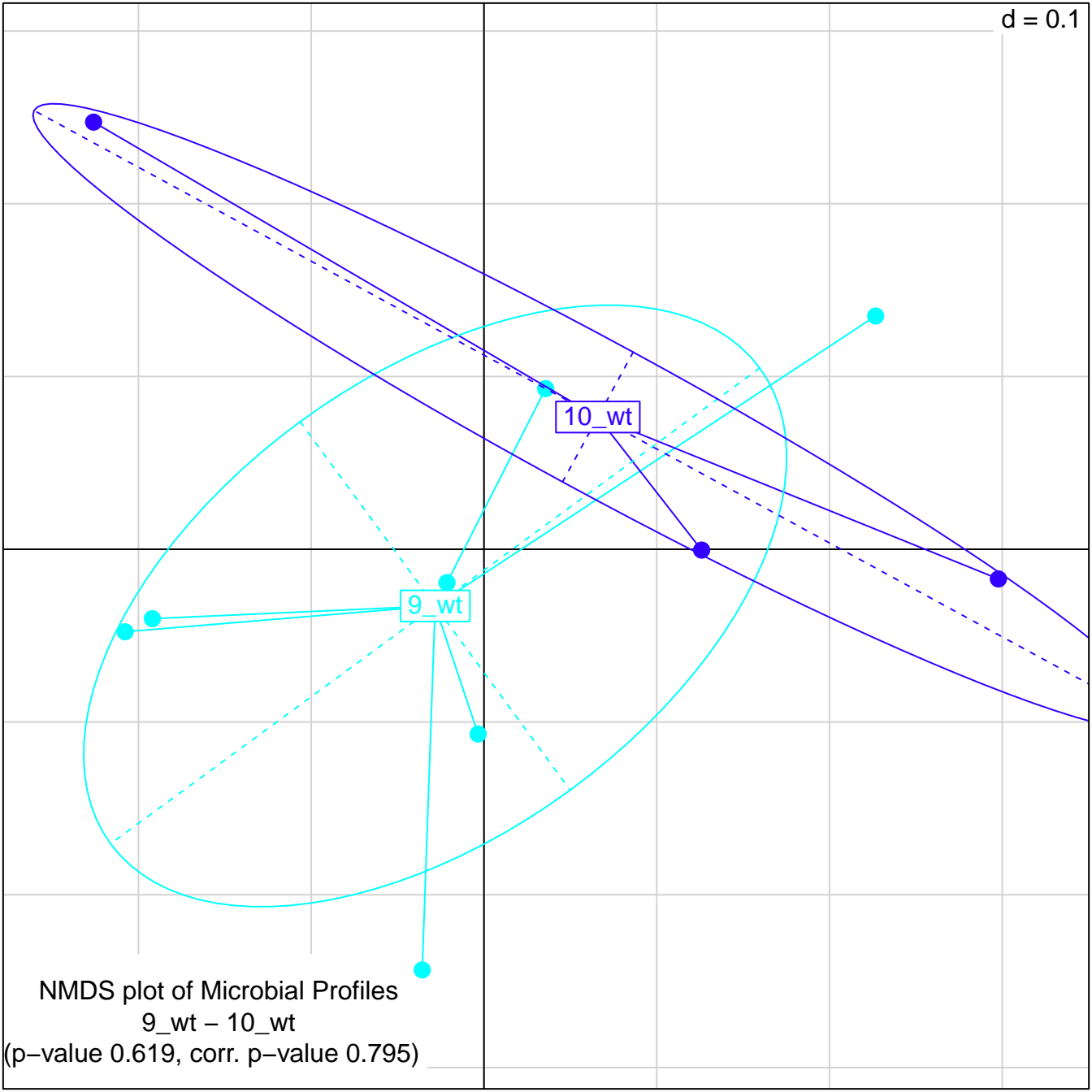
NMDS plot of Microbial Profiles  
9\_cre – 12\_wt  
(p-value 0.943, corr. p-value 0.95)

d = 0.1

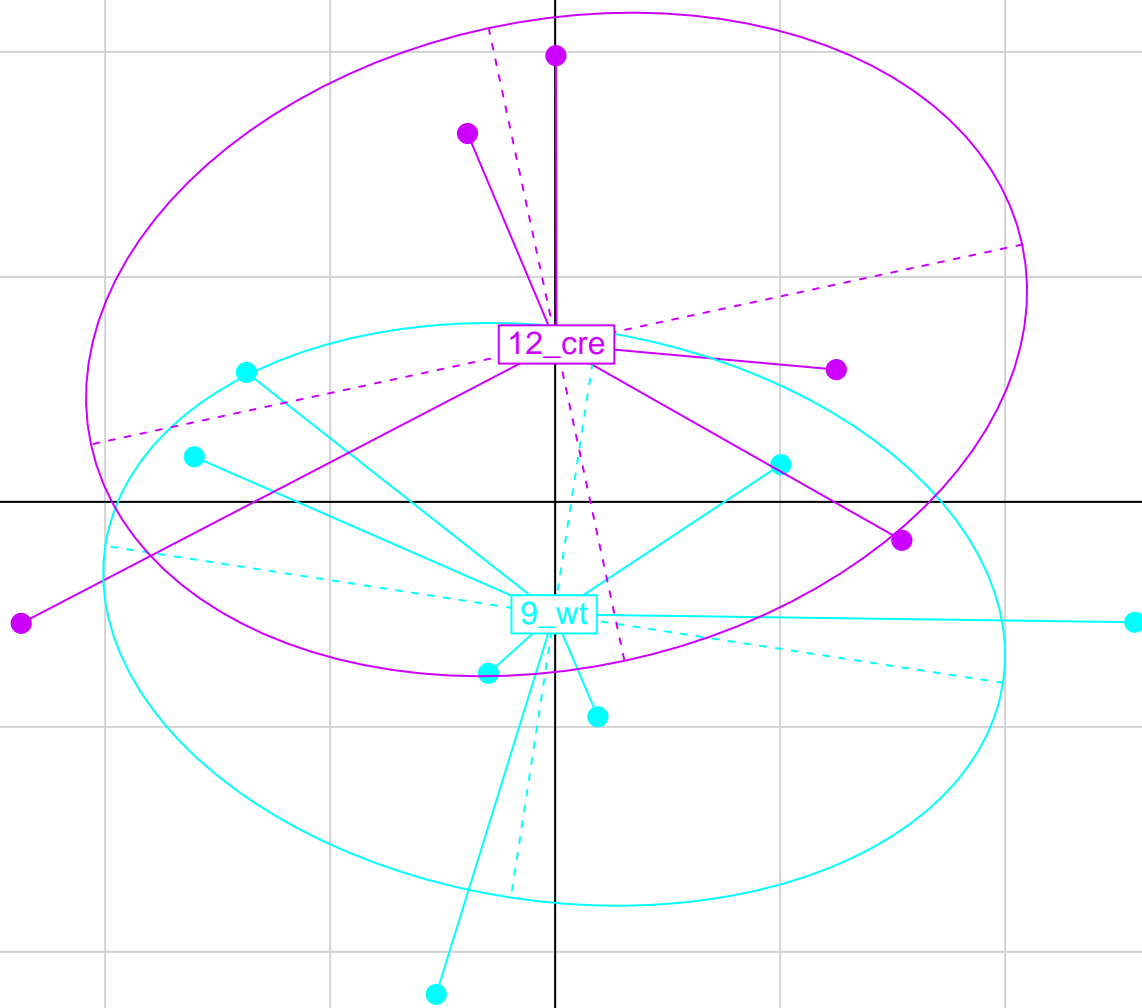


NMDS plot of Microbial Profiles  
9\_wt – 10\_cre  
(p-value 0.853, corr. p-value 0.892674418604651)

d = 0.1

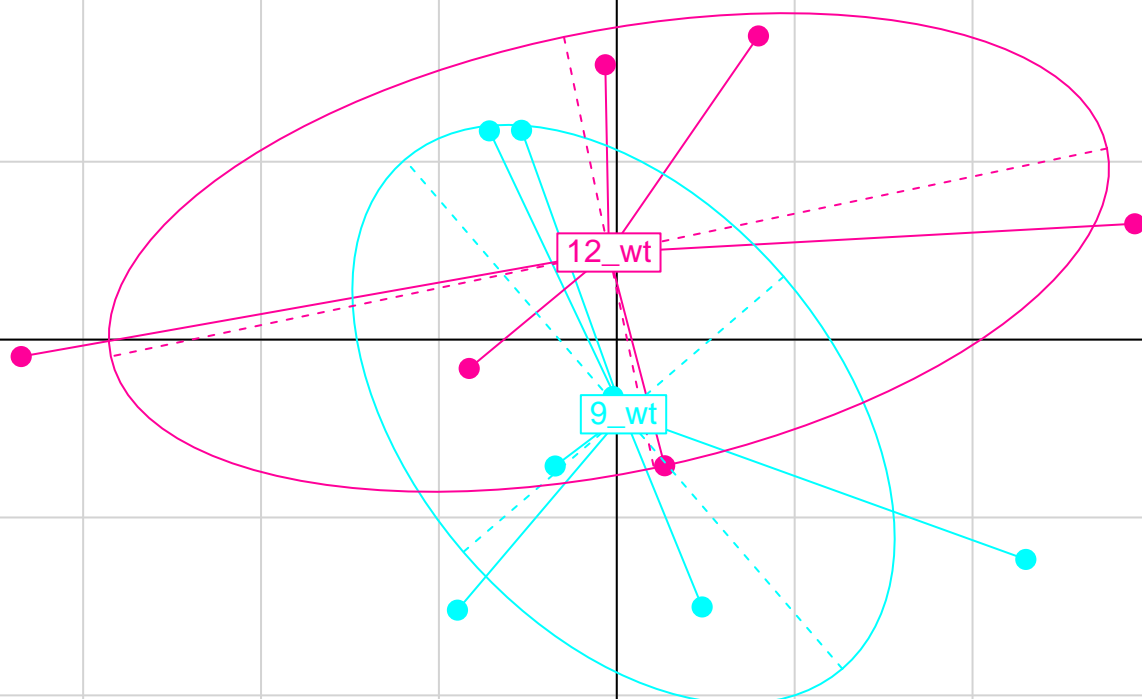


d = 0.1



NMDS plot of Microbial Profiles  
9\_wt – 12\_cre  
(p-value 0.328, corr. p-value 0.795)

d = 0.1

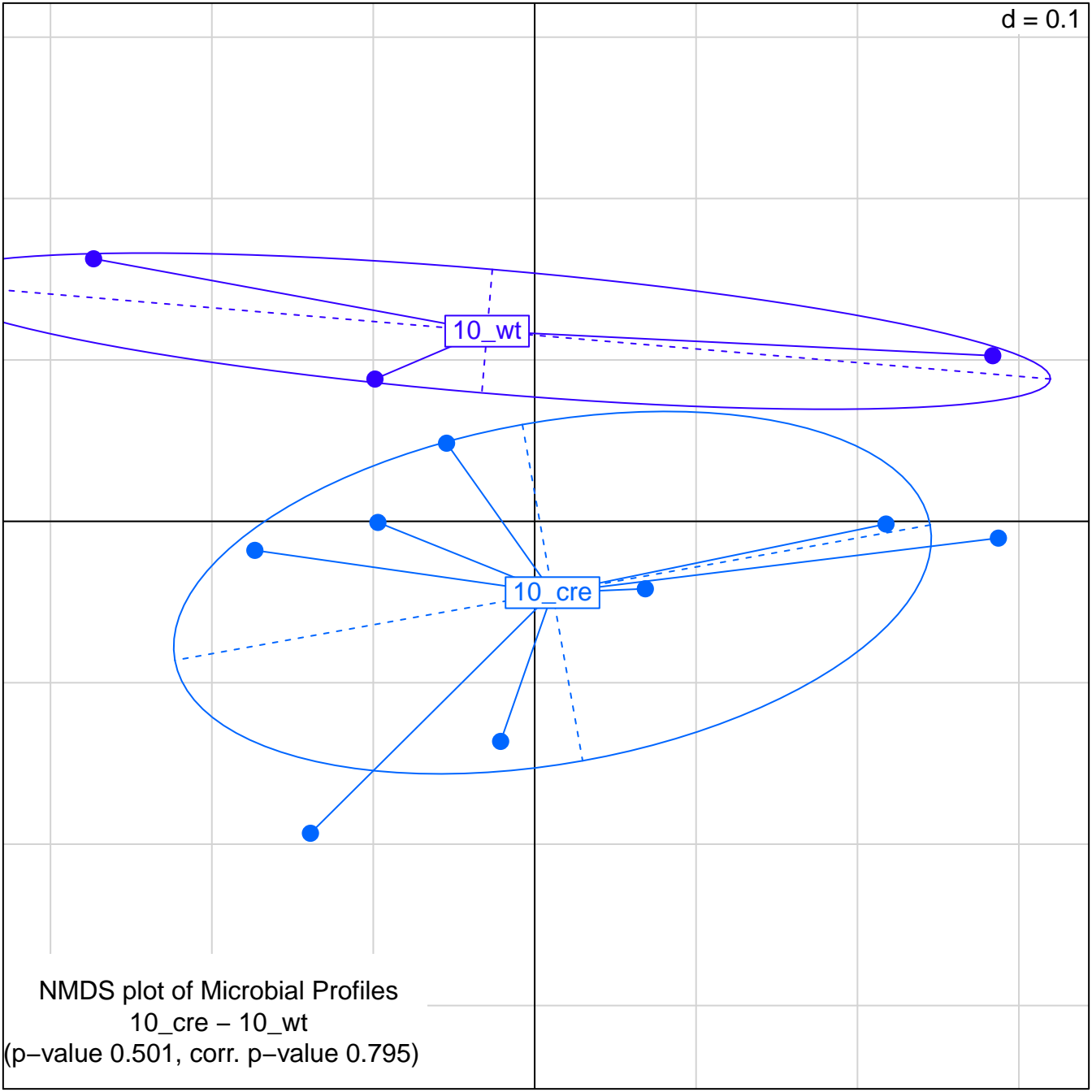


NMDS plot of Microbial Profiles

9\_wt – 12\_wt

(p-value 0.841, corr. p-value 0.892674418604651)

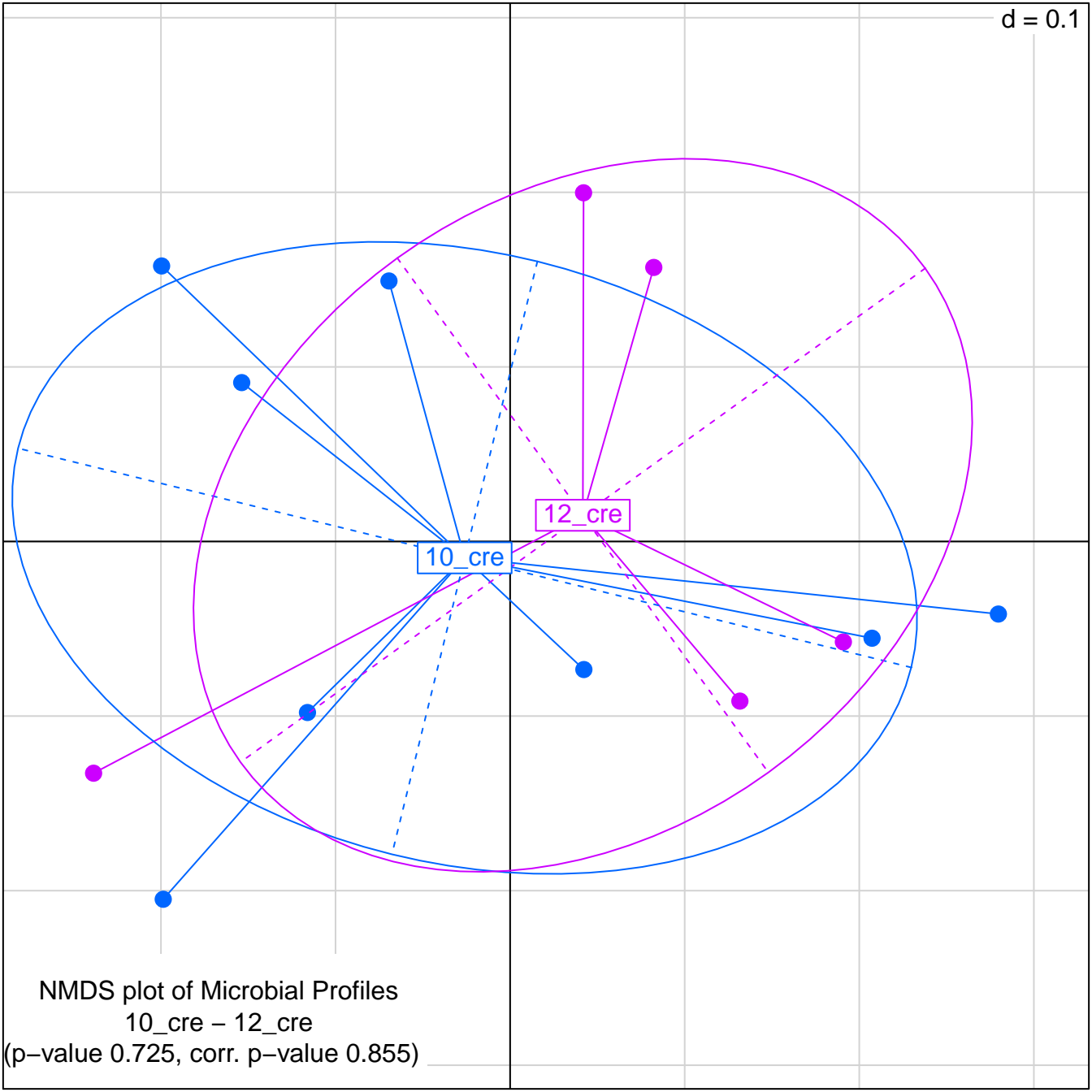
d = 0.1



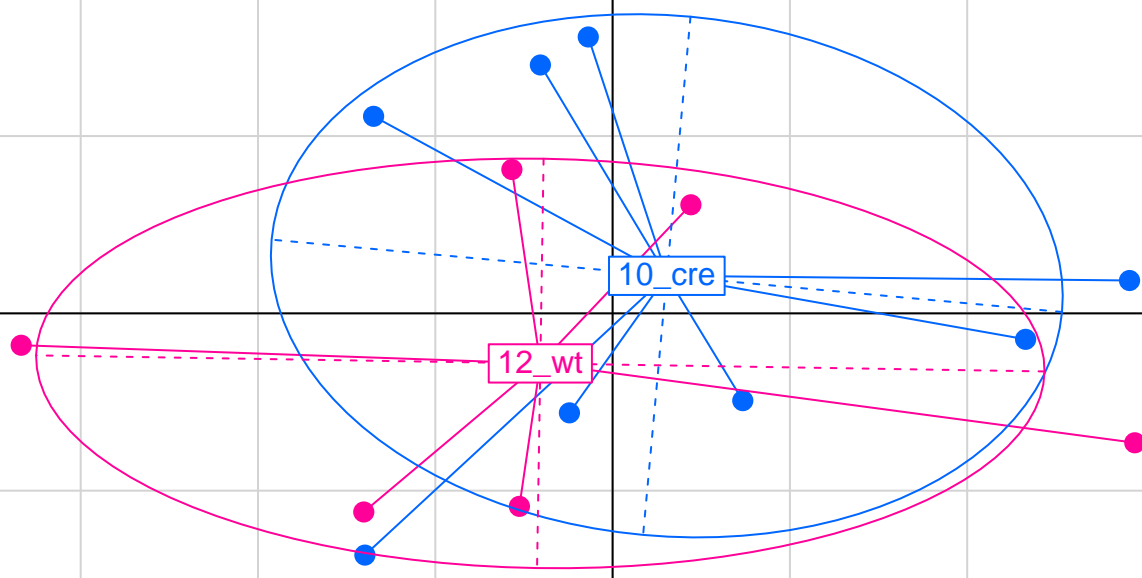
NMDS plot of Microbial Profiles  
10\_cre - 10\_wt  
(p-value 0.501, corr. p-value 0.795)



d = 0.1

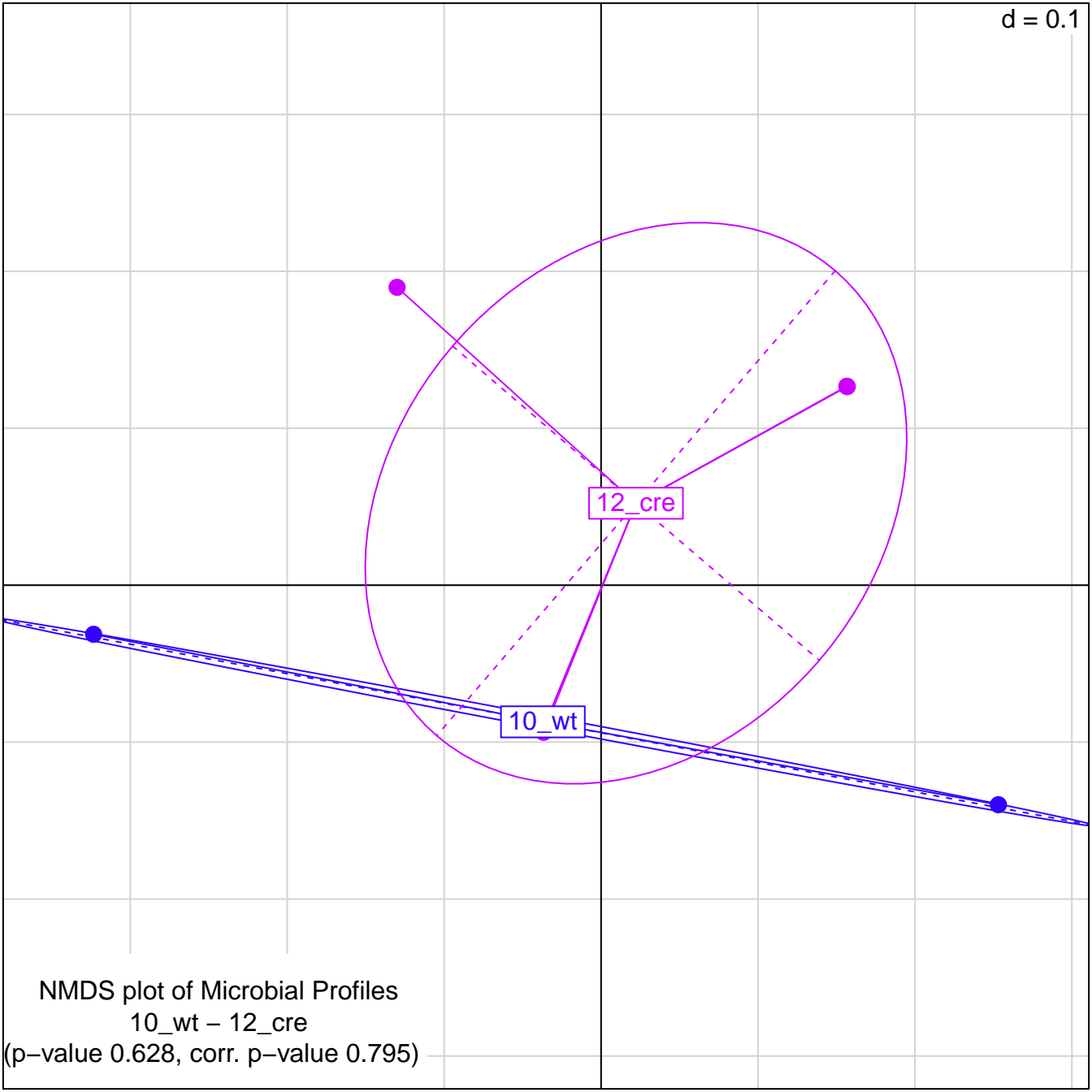


d = 0.1

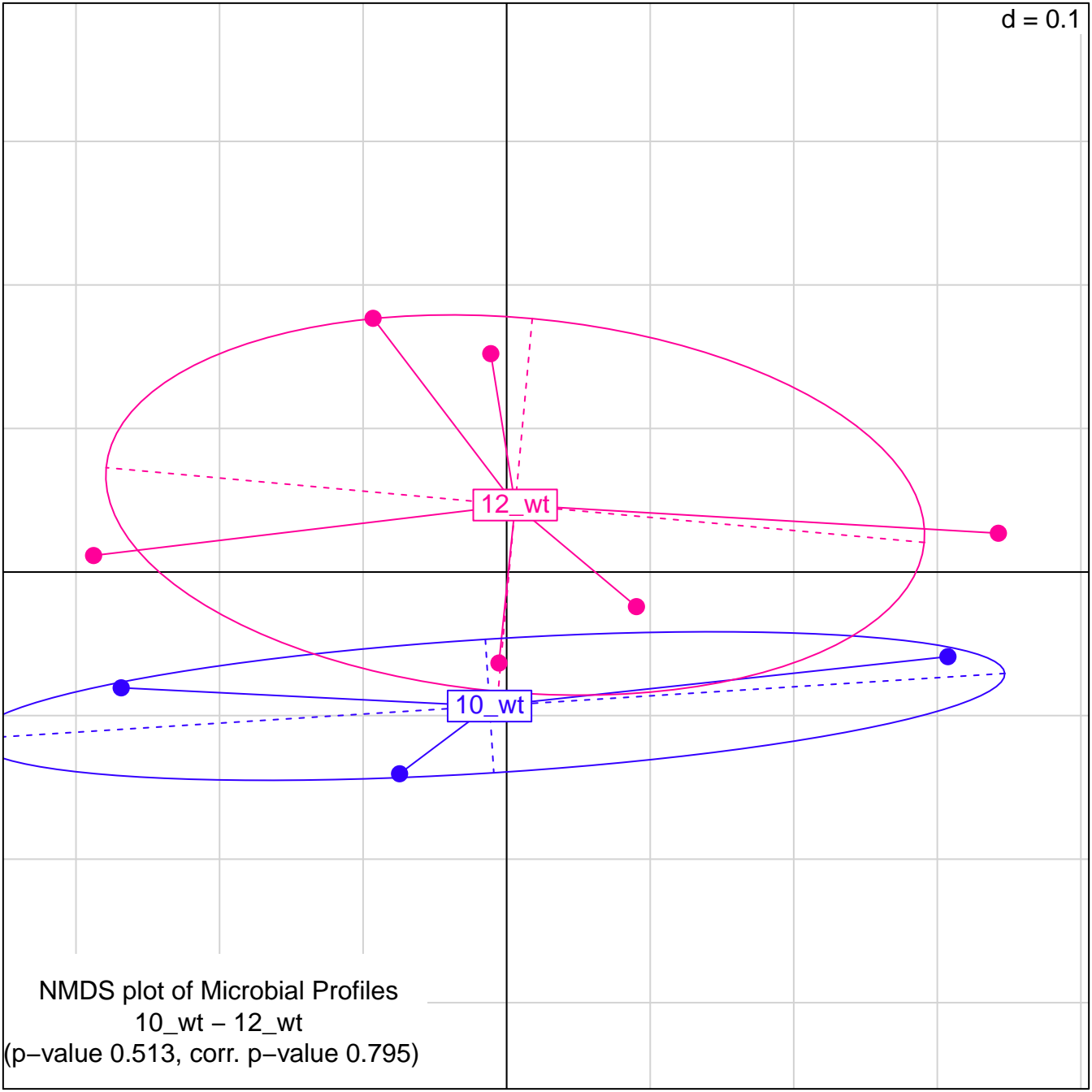


NMDS plot of Microbial Profiles  
10\_cre – 12\_wt  
(p-value 0.636, corr. p-value 0.795)

d = 0.1

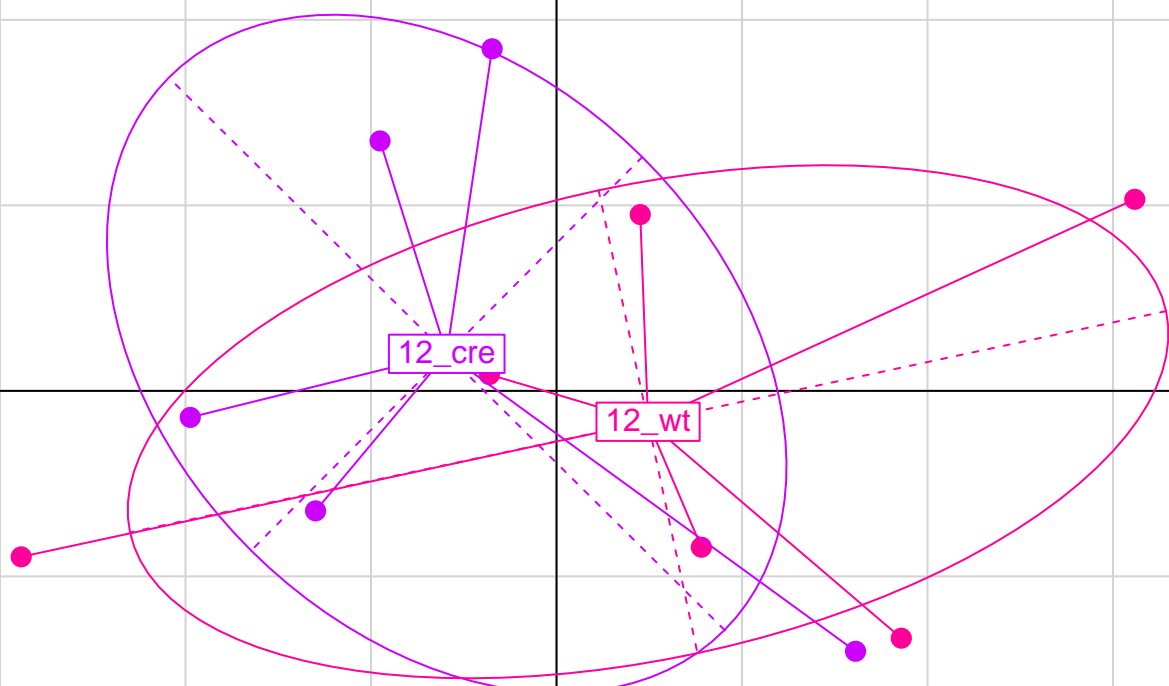


d = 0.1



NMDS plot of Microbial Profiles  
10\_wt – 12\_wt  
(p-value 0.513, corr. p-value 0.795)

d = 0.1



NMDS plot of Microbial Profiles  
12\_cre – 12\_wt  
(p-value 0.397, corr. p-value 0.795)