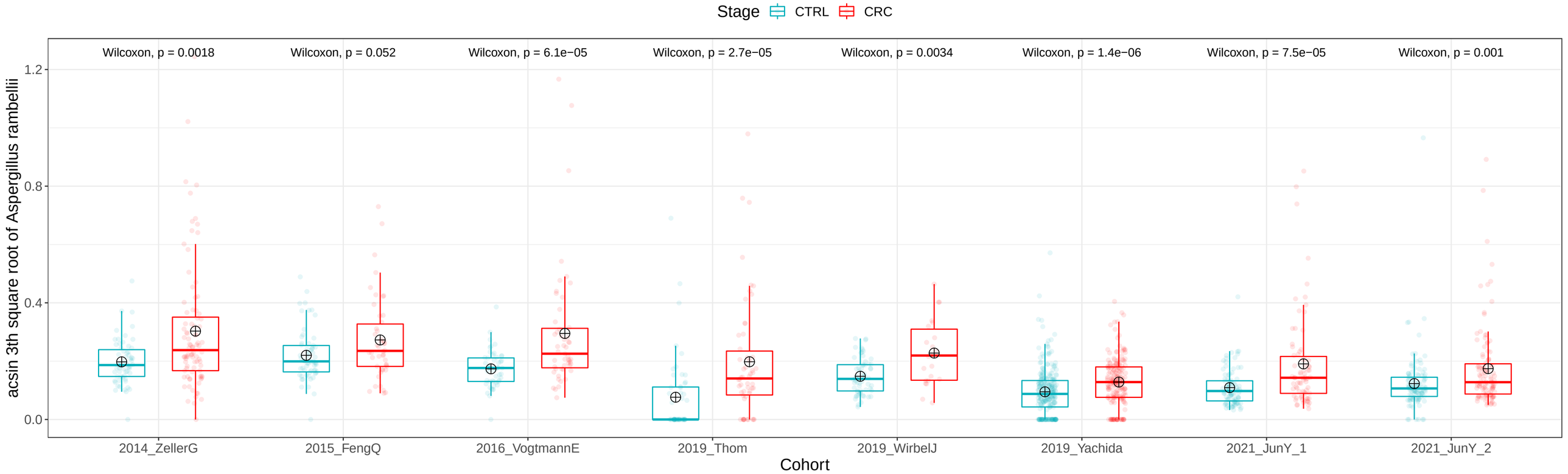
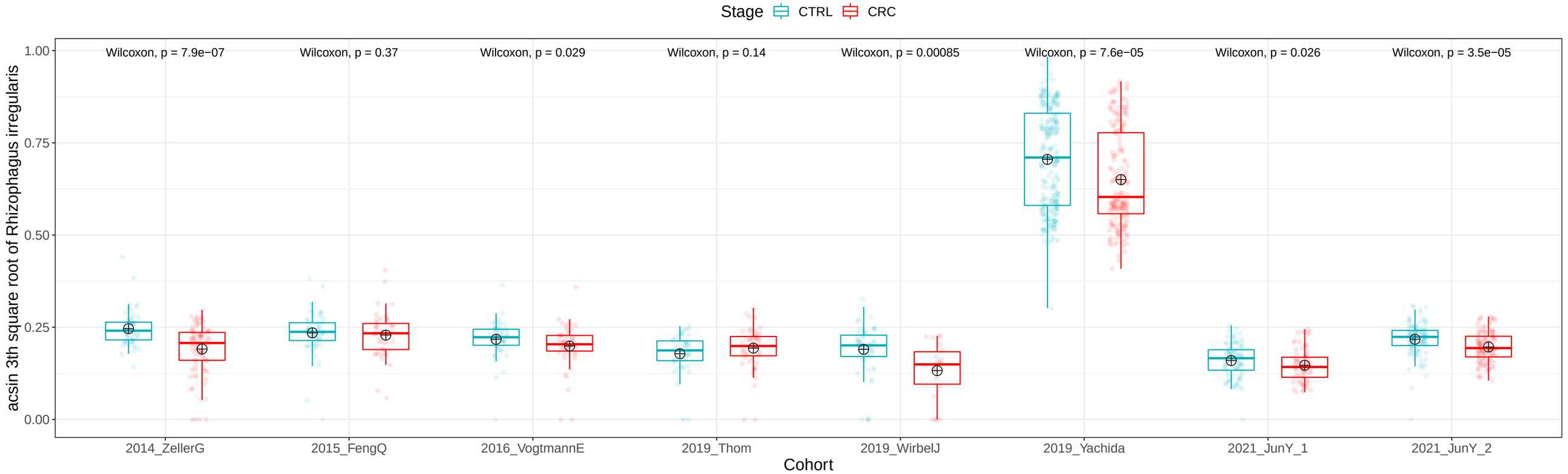
Aspergillus rambellii

8 cohorts has same trend in CRC entiched.
7 cohorts play significant different.



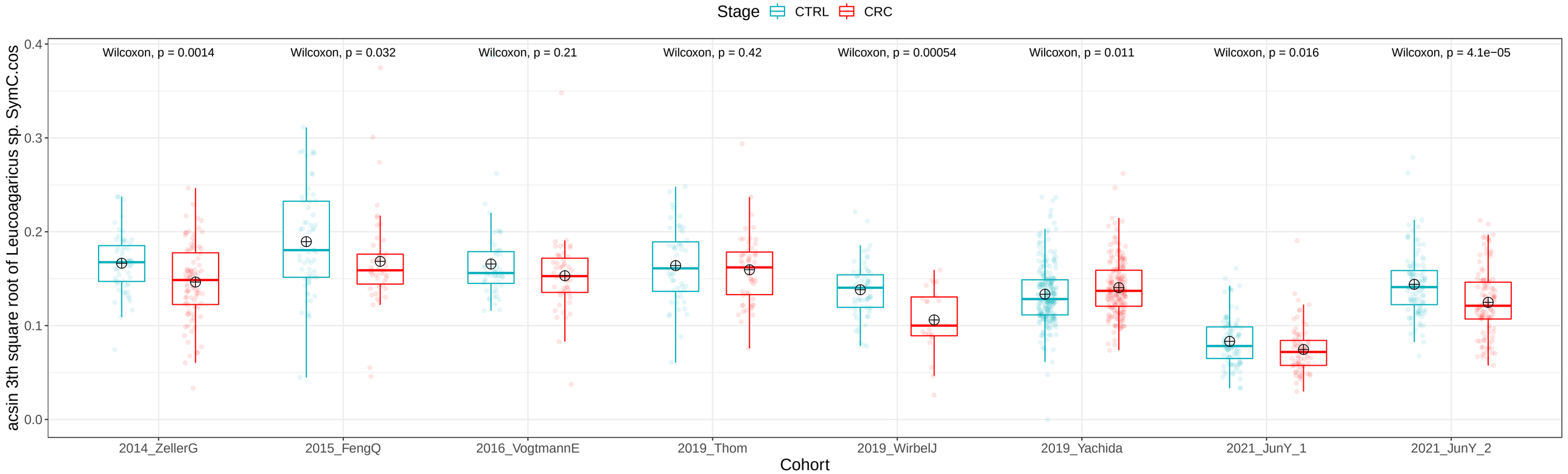
Rhizophagus irregularis

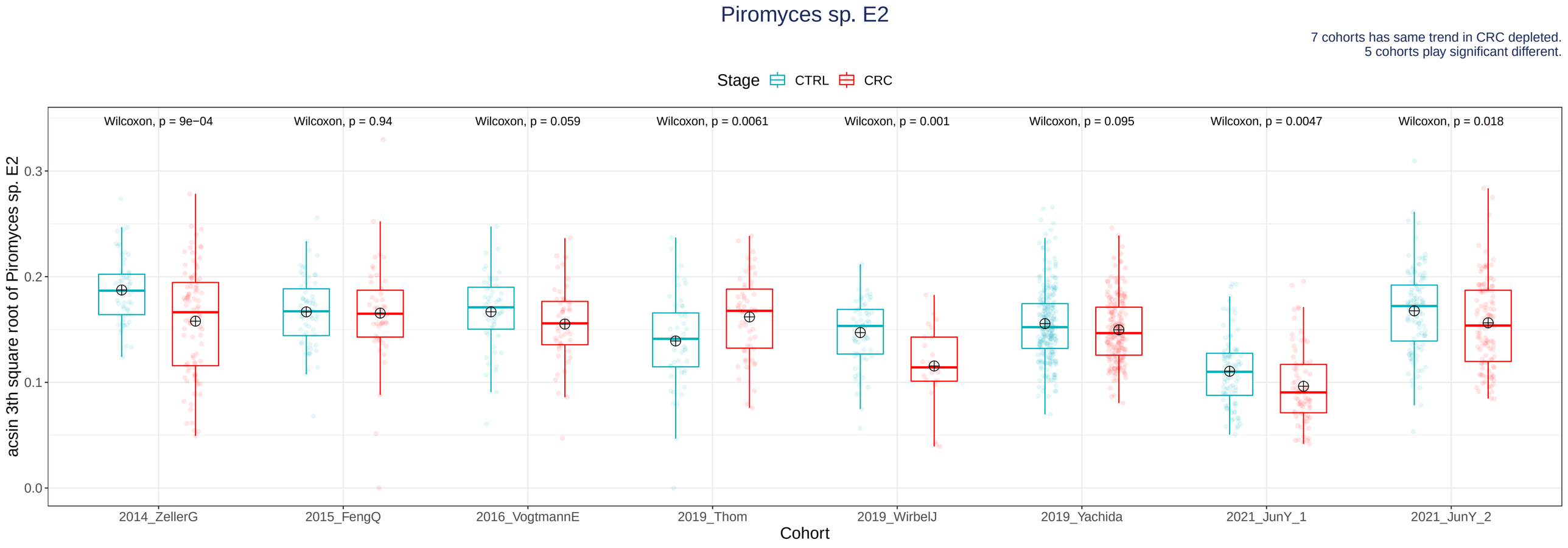
7 cohorts has same trend in CRC depleted.
6 cohorts play significant different.



Leucoagaricus sp. SymC.cos

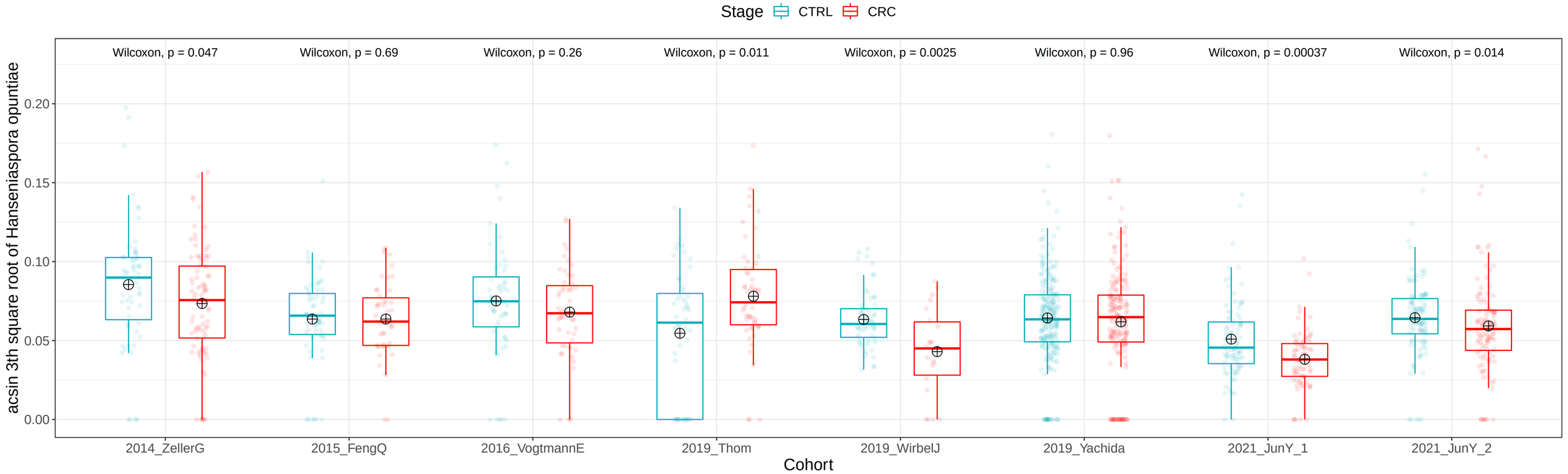
7 cohorts has same trend in CRC depleted.
6 cohorts play significant different.




Piromyces sp. E2
7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.
Stage
CTRL
CRC
Wilcoxon, p = 9e−04
Wilcoxon, p = 0.94
Wilcoxon, p = 0.059
Wilcoxon, p = 0.0061
Wilcoxon, p = 0.001
Wilcoxon, p = 0.095
Wilcoxon, p = 0.0047
Wilcoxon, p = 0.018
acsin 3th square root of Piromyces sp. E2
0.0
0.1
0.2
0.3
2014_ZellerG
2015_FengQ
2016_VogtmannE
2019_Thom
2019_WirbelJ
2019_Yachida
2021_JunY_1
2021_JunY_2
Cohort

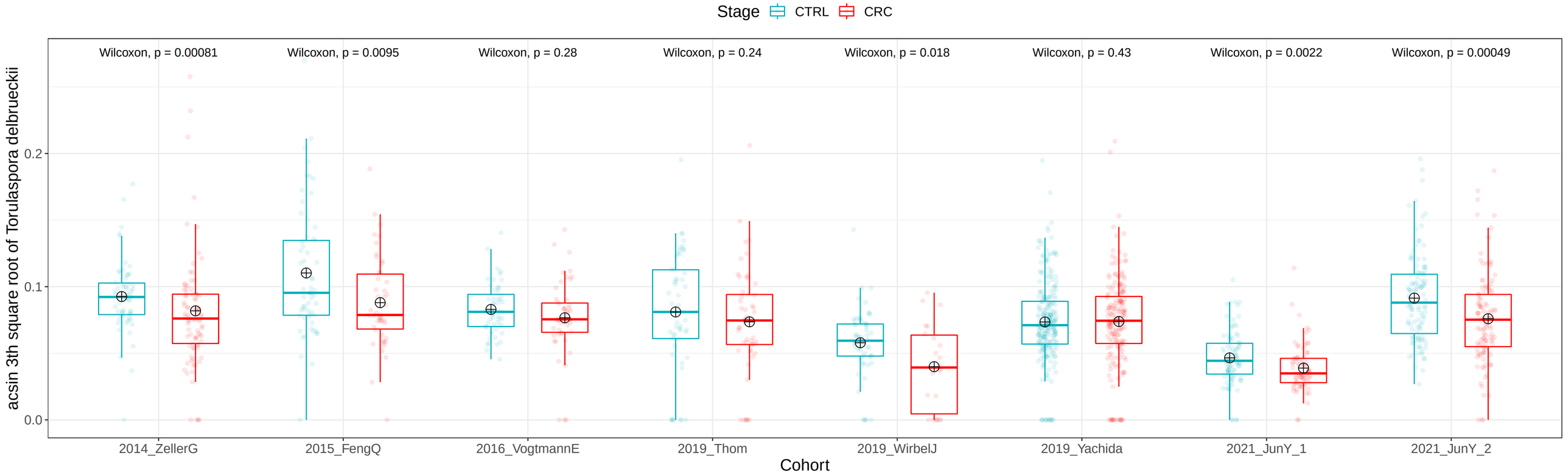
Hanseniaspora opuntiae

7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.



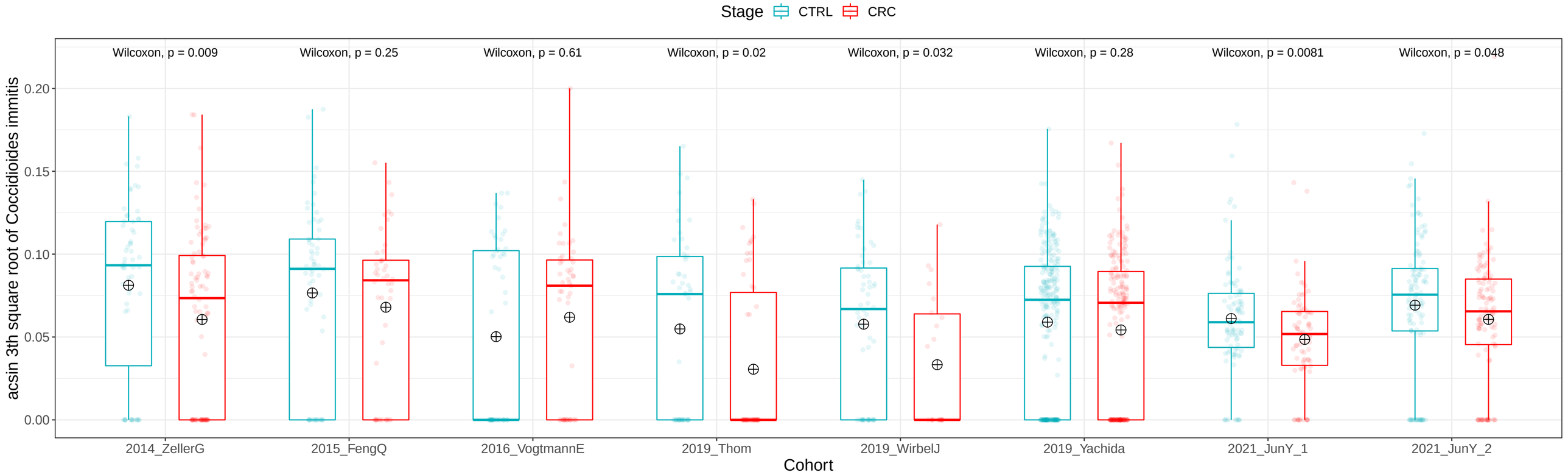
Torulaspora delbrueckii

7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.



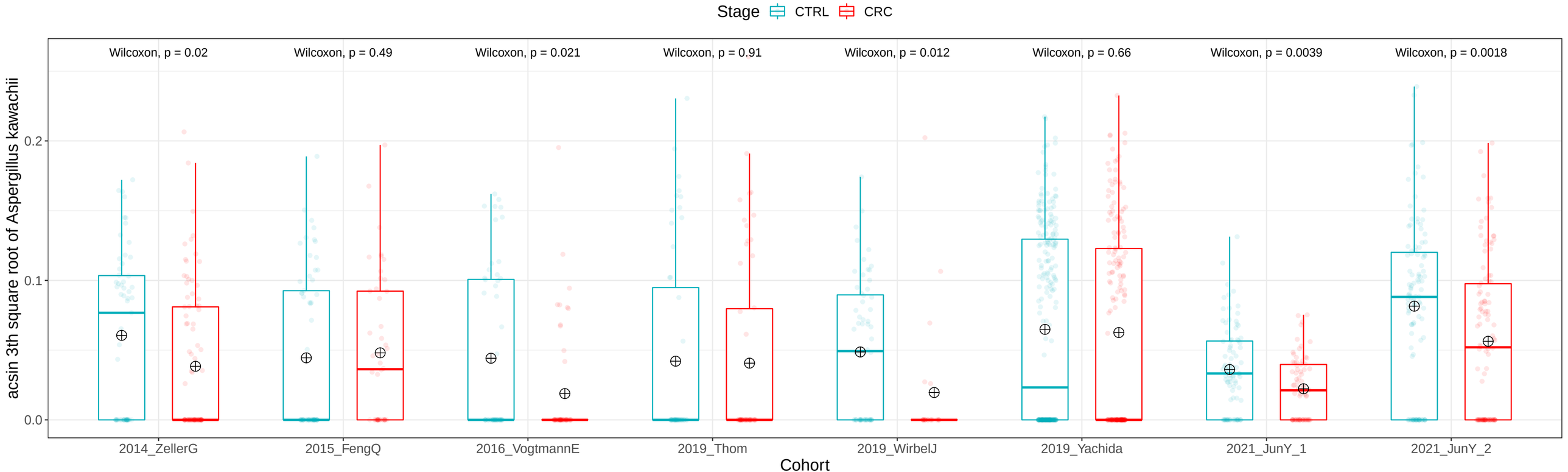
Coccidioides immitis

7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.



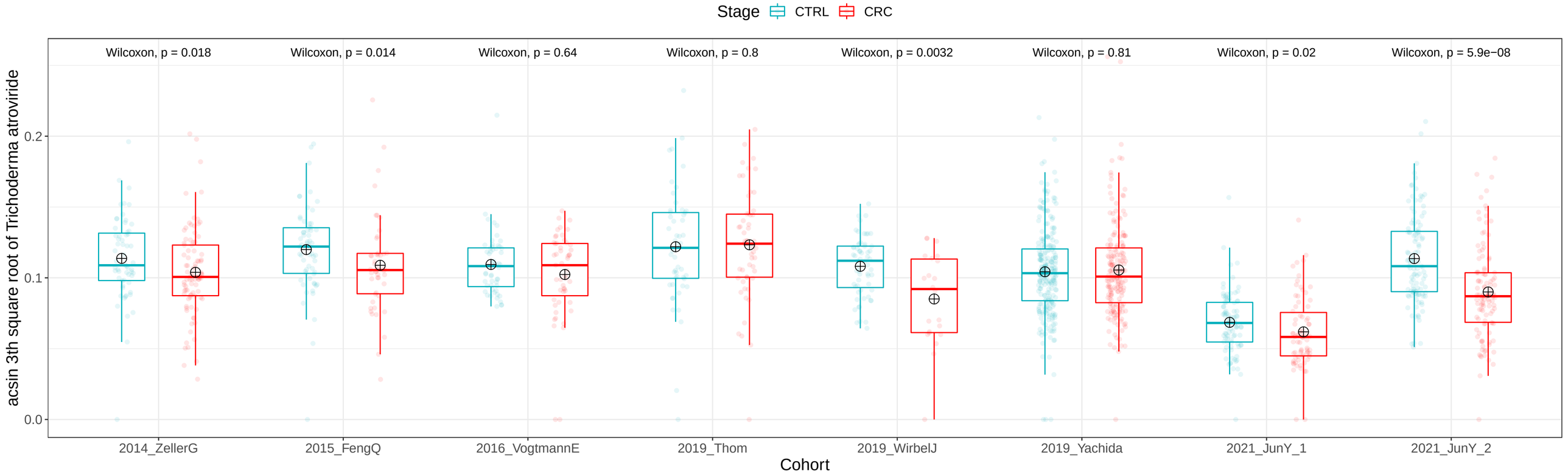
Aspergillus kawachii

7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.



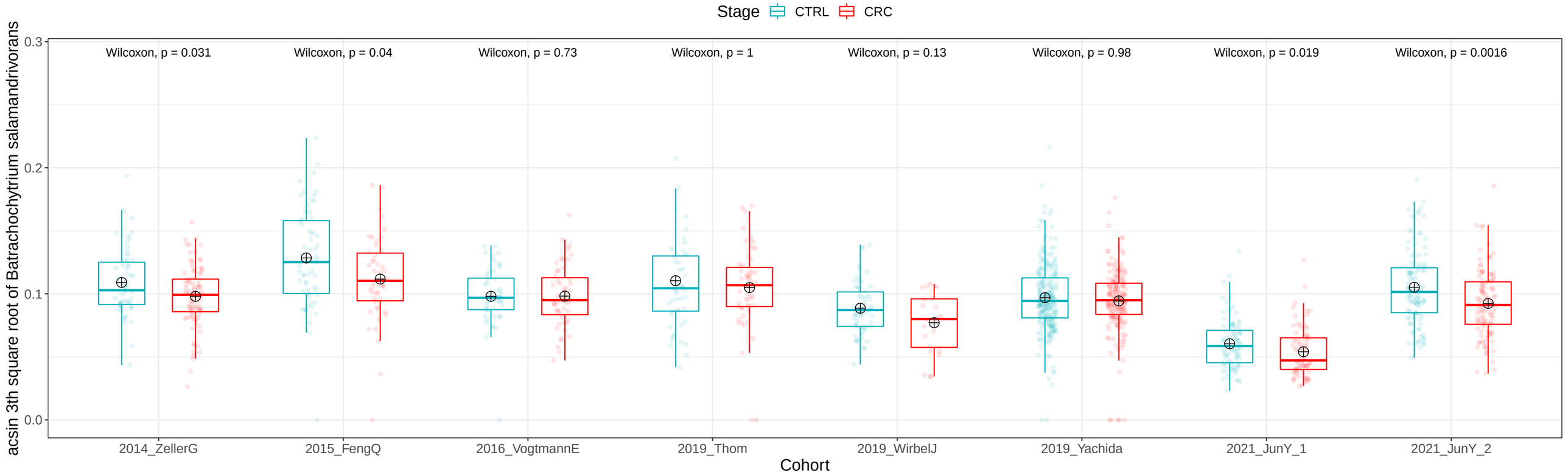
Trichoderma atroviride

7 cohorts has same trend in CRC depleted.
5 cohorts play significant different.



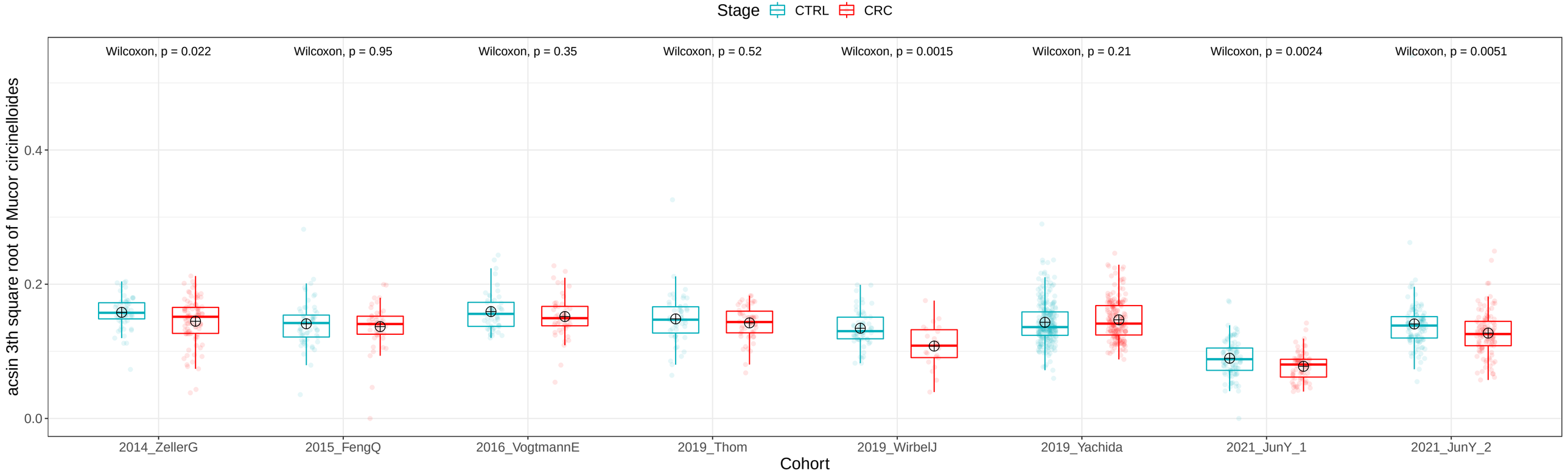
Batrachochytrium salamandrivorans

8 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



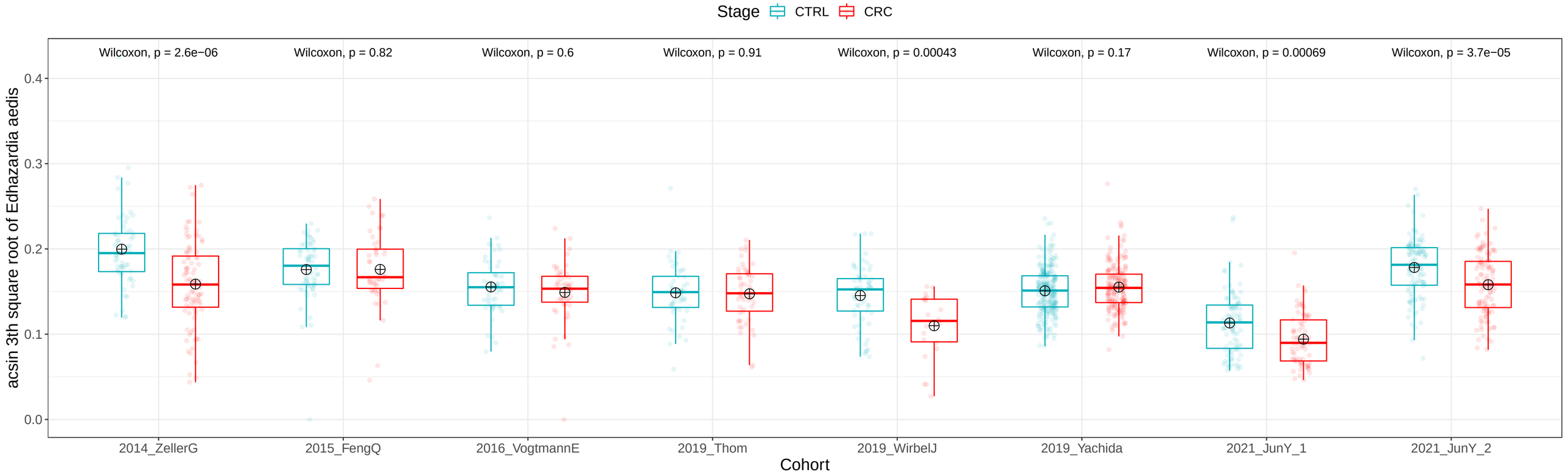
Mucor circinelloides

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



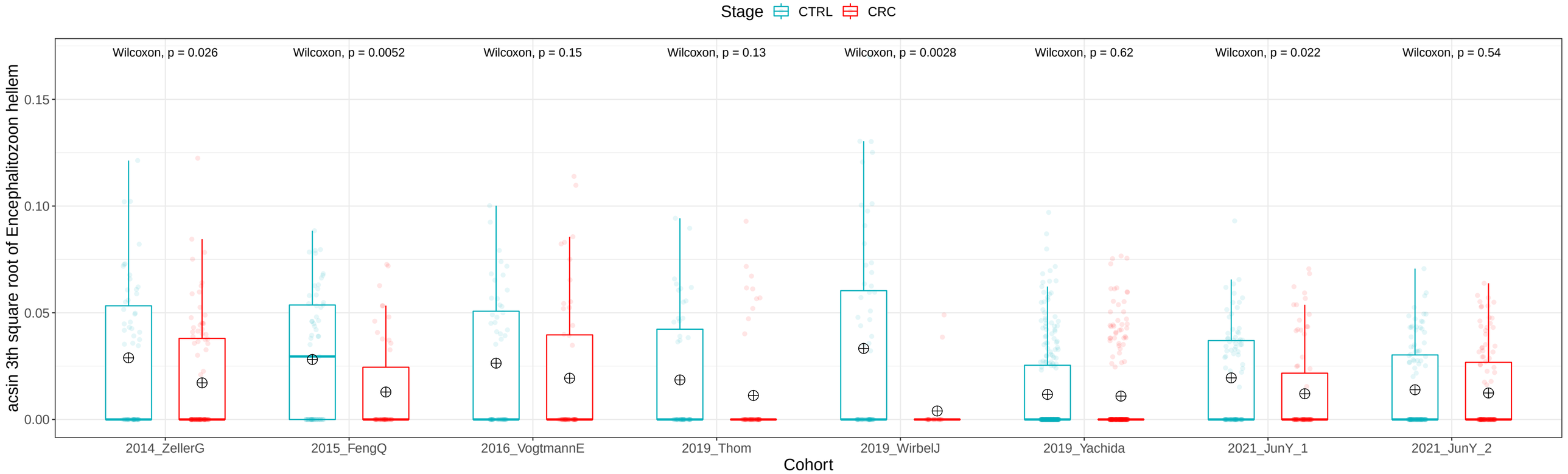
Edhazardia aedis

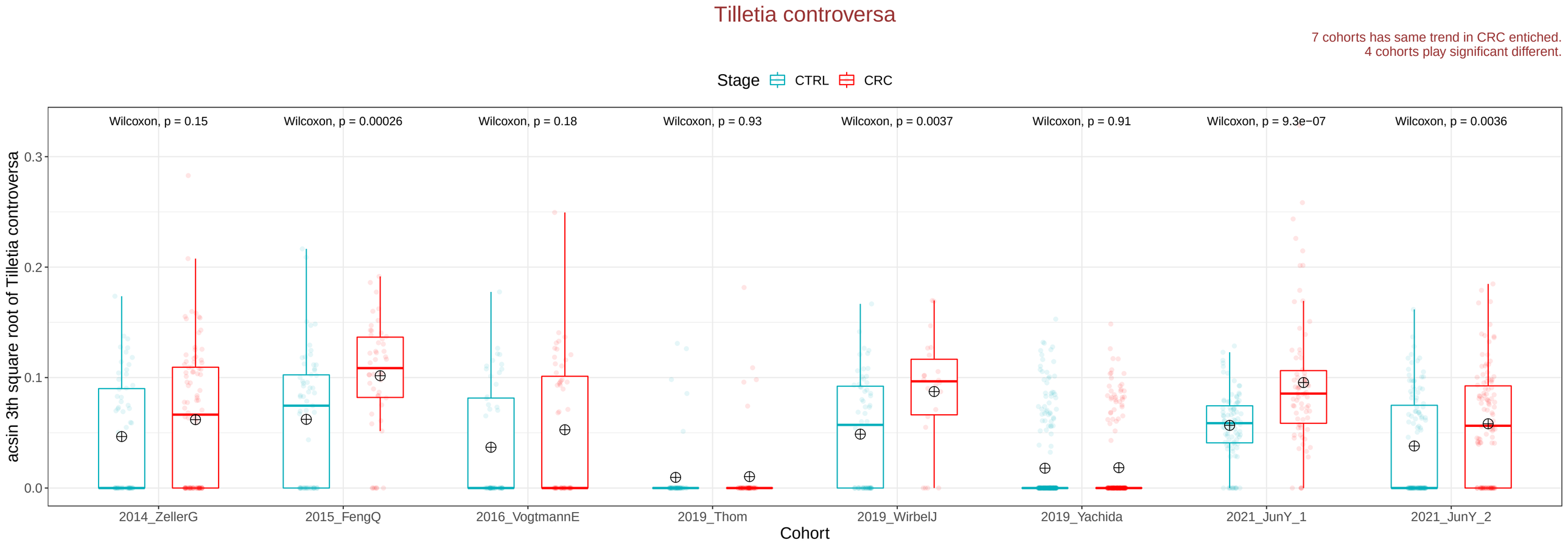
7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



Encephalitozoon hellem

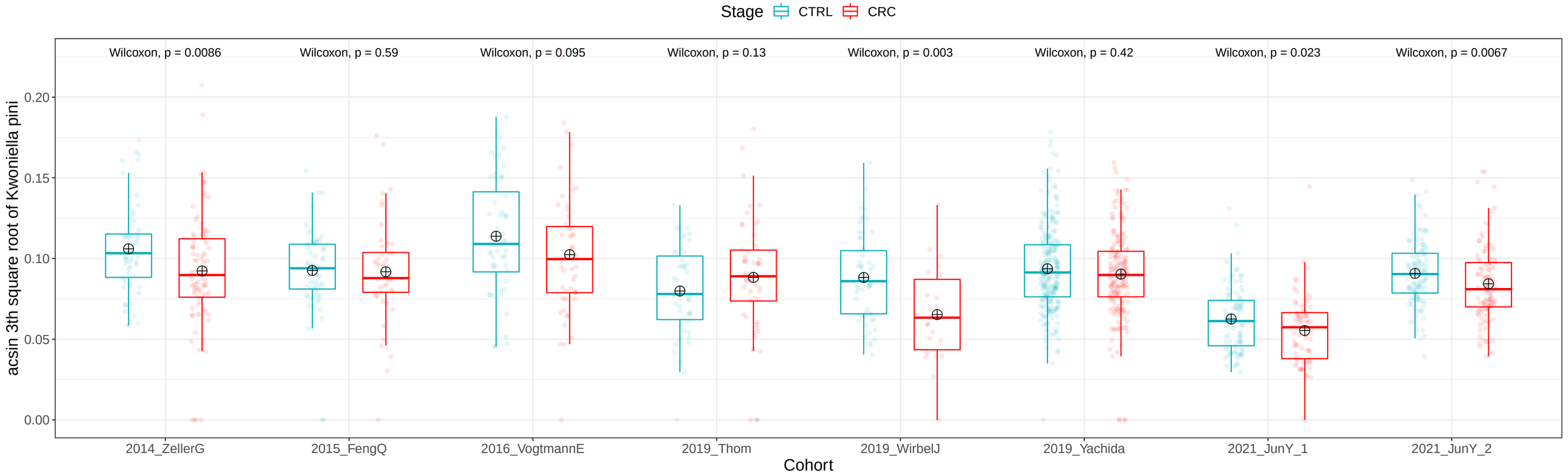
8 cohorts has same trend in CRC depleted.
4 cohorts play significant different.




Tilletia controversa
7 cohorts has same trend in CRC entiched.
4 cohorts play significant different.
Stage
CTRL
CRC
Wilcoxon, p = 0.15
Wilcoxon, p = 0.00026
Wilcoxon, p = 0.18
Wilcoxon, p = 0.93
Wilcoxon, p = 0.0037
Wilcoxon, p = 0.91
Wilcoxon, p = 9.3e−07
Wilcoxon, p = 0.0036
acsin 3th square root of Tilletia controversa
0.0
0.1
0.2
0.3
2014_ZellerG
2015_FengQ
2016_VogtmannE
2019_Thom
2019_WirbelJ
2019_Yachida
2021_JunY_1
2021_JunY_2
Cohort

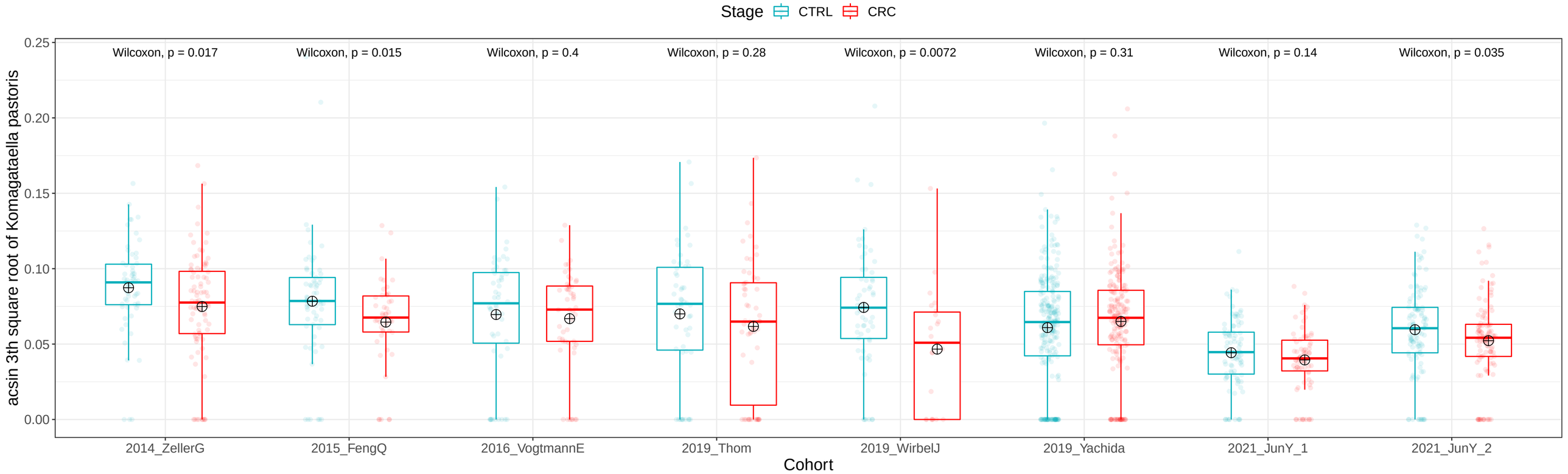
Kwoniella pini

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



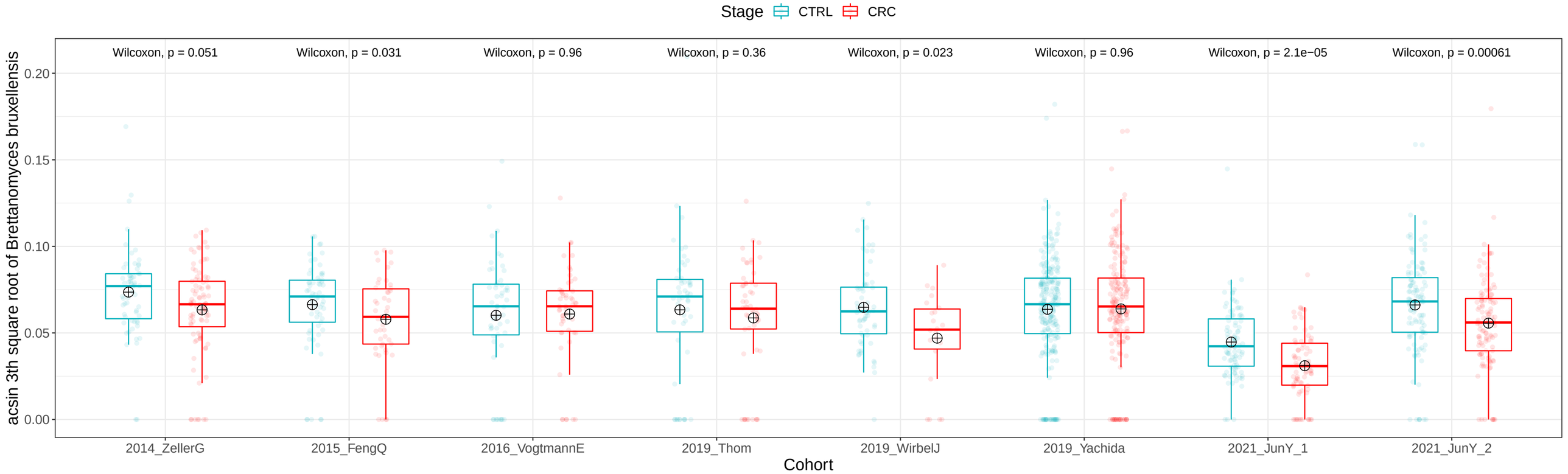
Komagataella pastoris

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



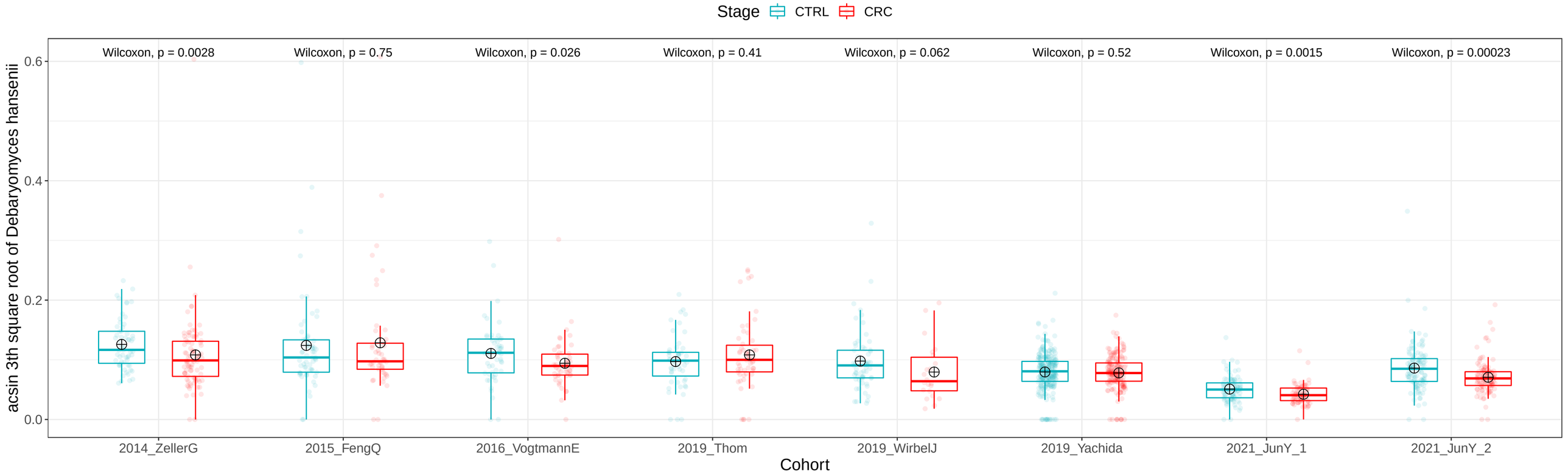
Brettanomyces bruxellensis

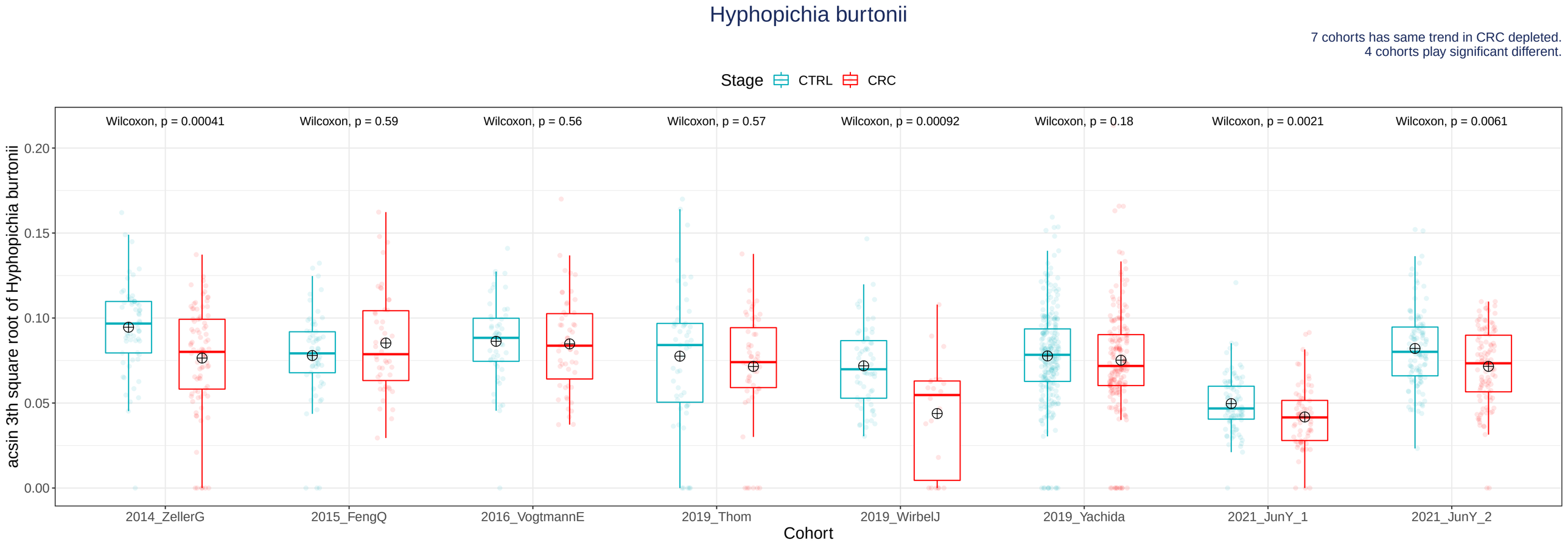
7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



Debaryomyces hansenii

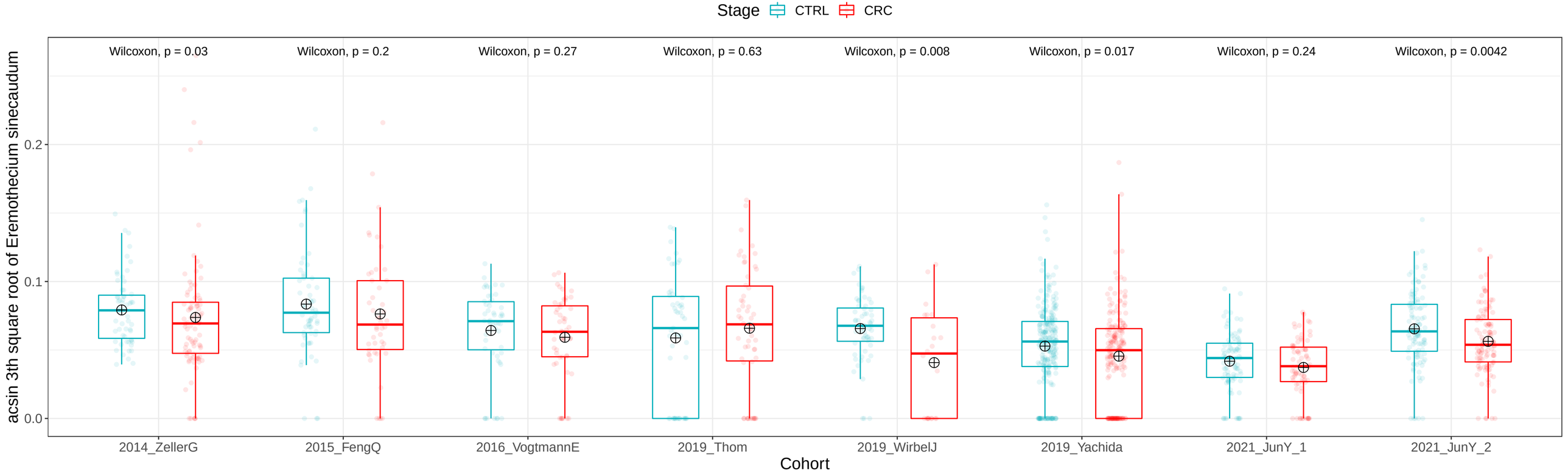
7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.




Hyphopichia burtonii
7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.
Stage
CTRL
CRC
Wilcoxon, p = 0.00041
Wilcoxon, p = 0.59
Wilcoxon, p = 0.56
Wilcoxon, p = 0.57
Wilcoxon, p = 0.00092
Wilcoxon, p = 0.18
Wilcoxon, p = 0.0021
Wilcoxon, p = 0.0061
acsin 3th square root of Hyphopichia burtonii
0.20
0.15
0.10
0.05
0.00
2014_ZellerG
2015_FengQ
2016_VogtmannE
2019_Thom
2019_WirbelJ
2019_Yachida
2021_JunY_1
2021_JunY_2
Cohort

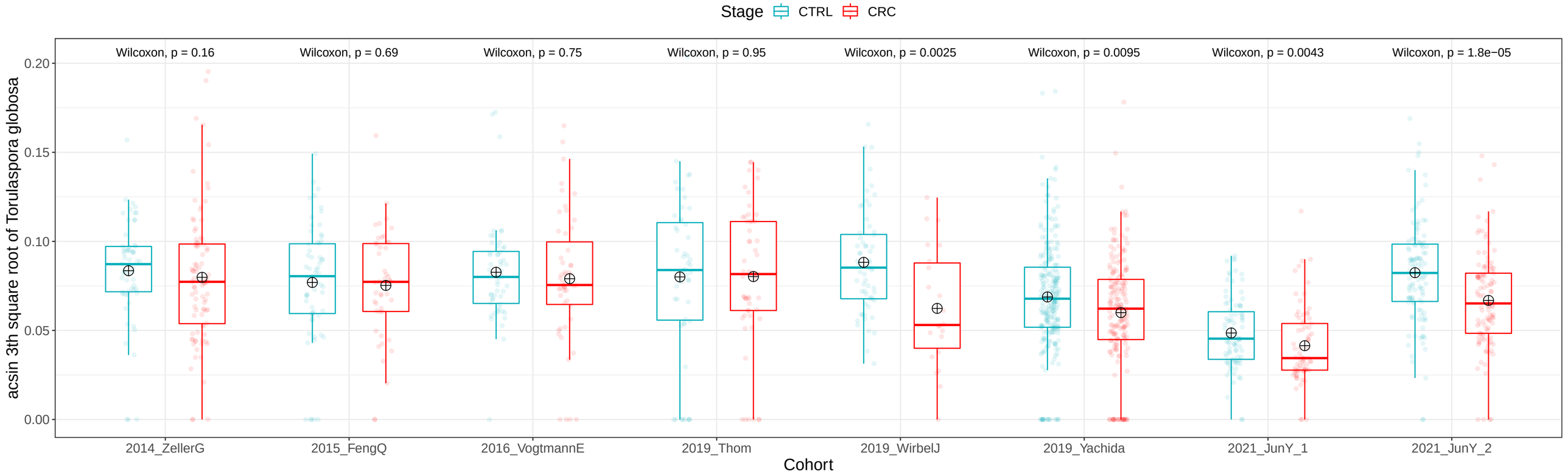
Eremothecium sinecaudum

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



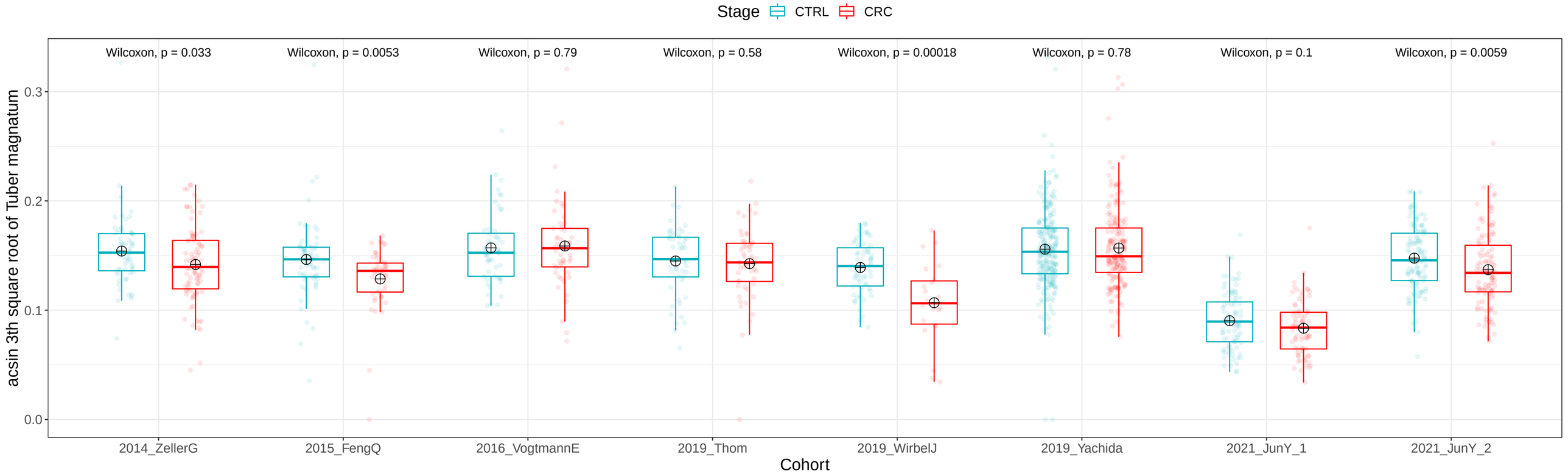
Torulaspora globosa

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



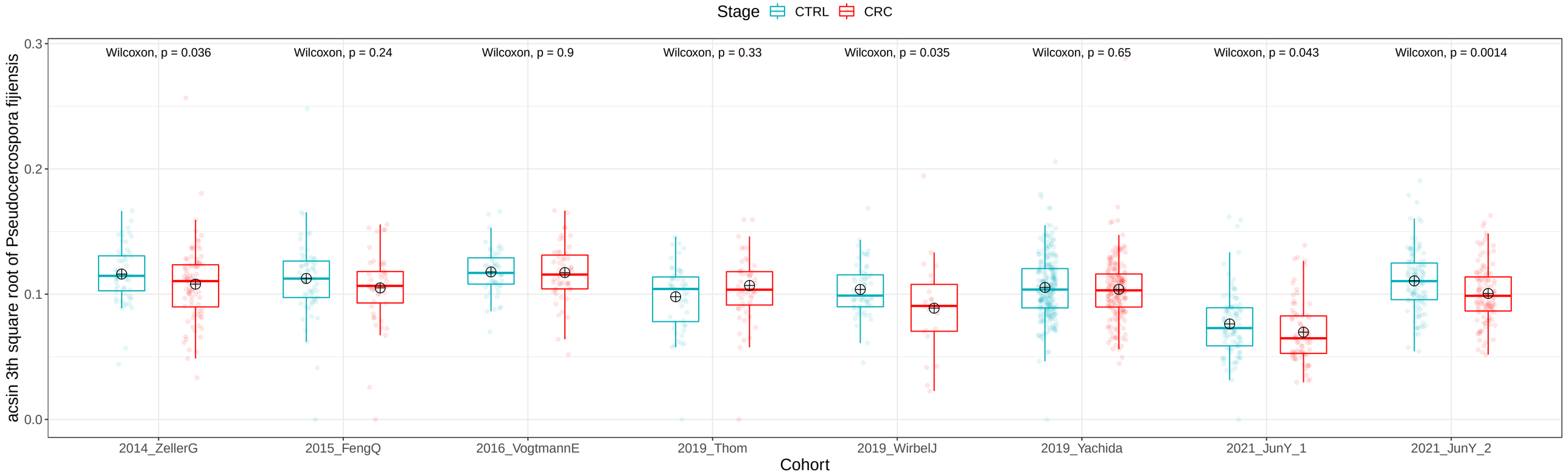
Tuber magnatum

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



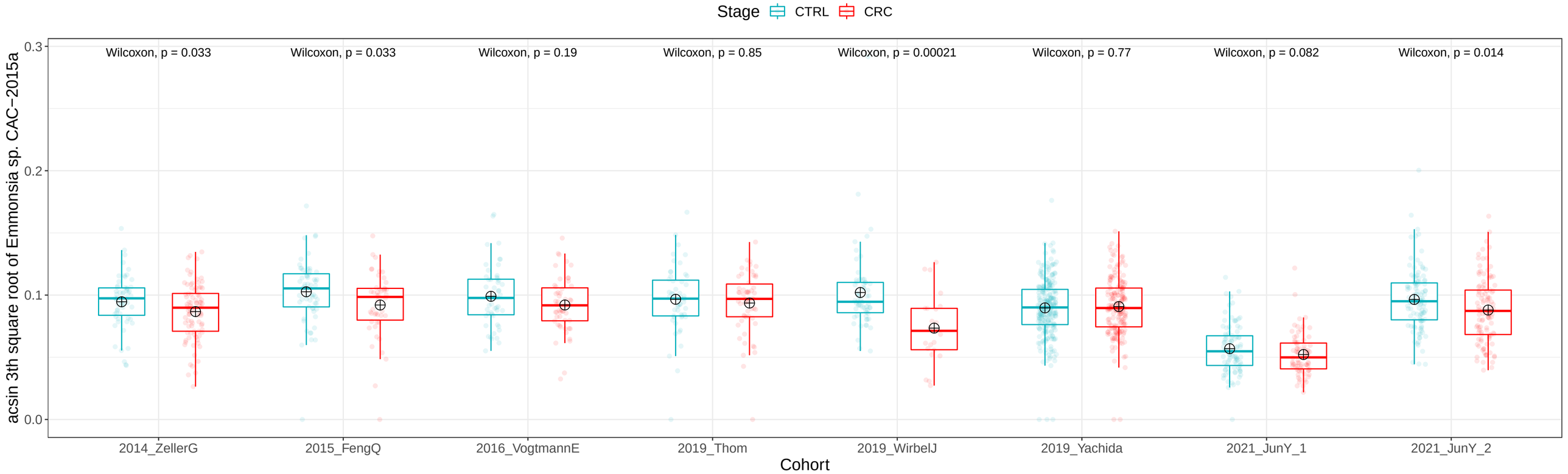
Pseudocercospora fijiensis

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



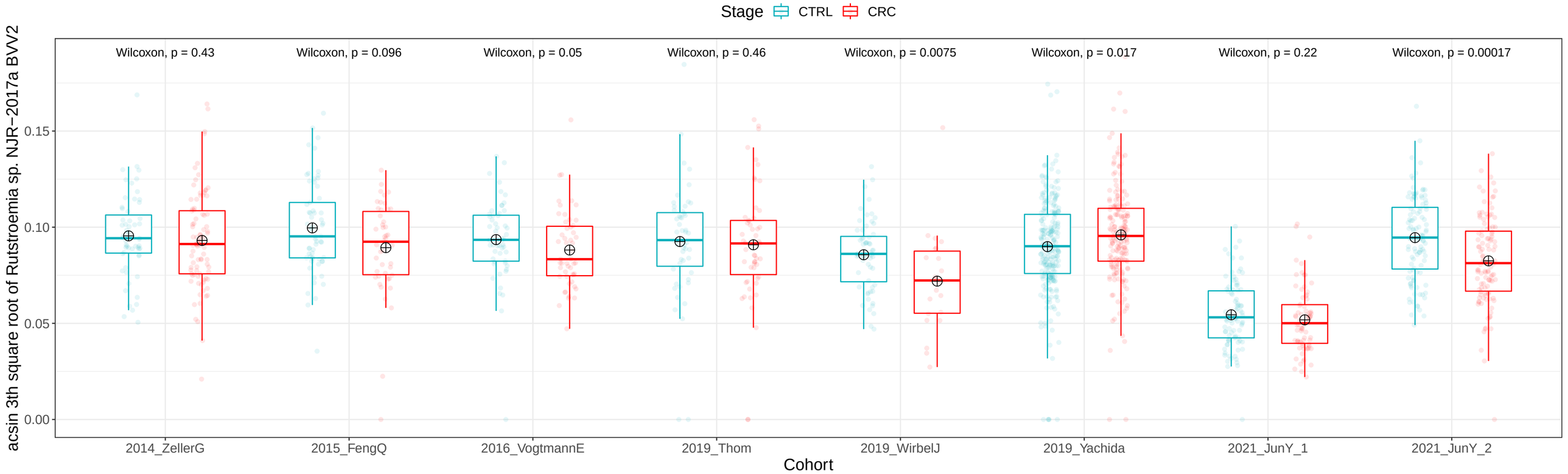
Emmonsia sp. CAC−2015a

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



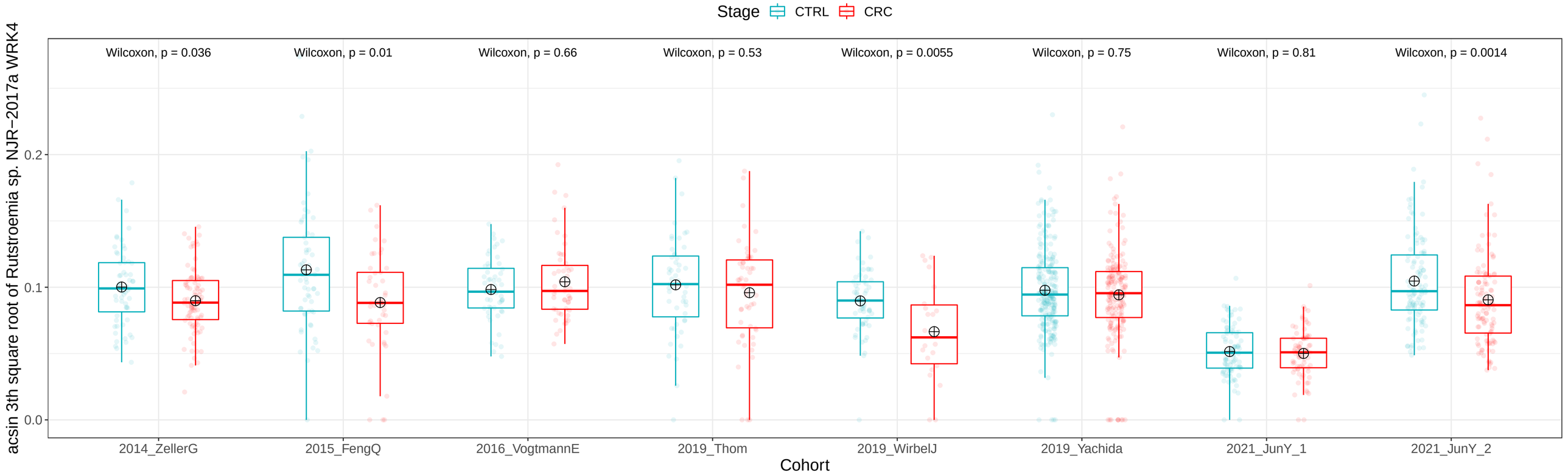
Rutstroemia sp. NJR−2017a BVV2

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



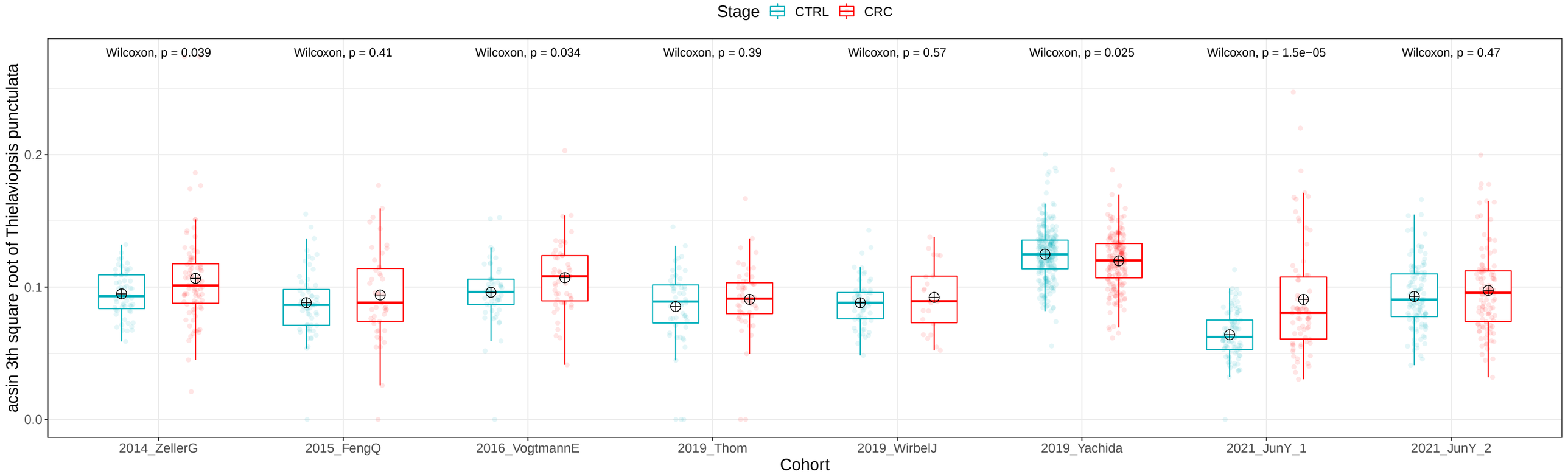
Rutstroemia sp. NJR−2017a WRK4

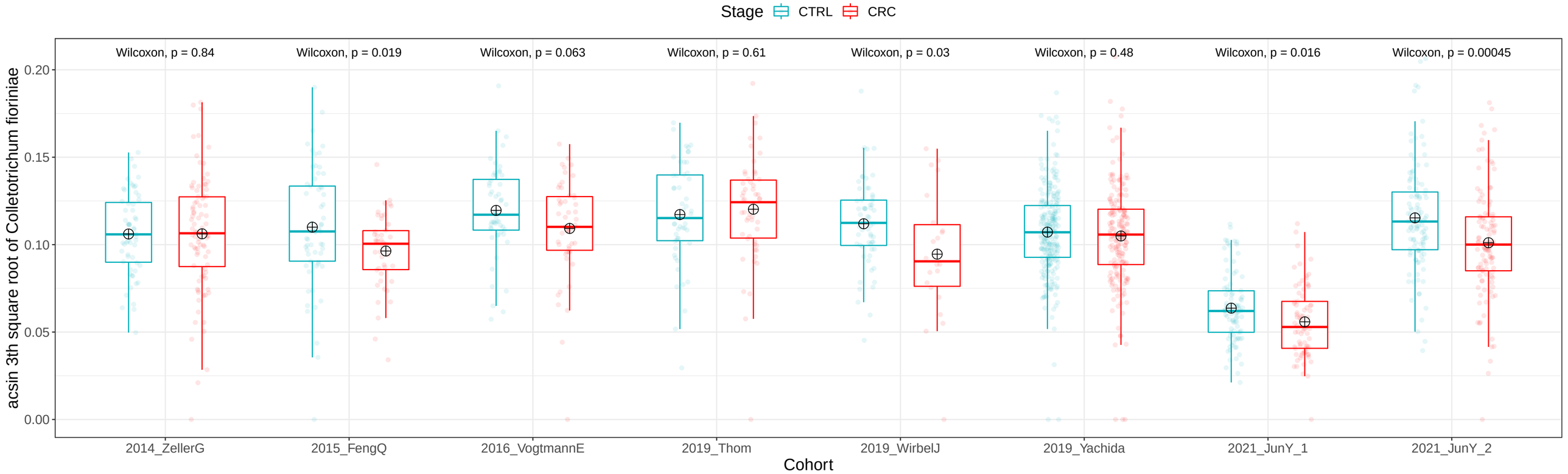
7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.



Thielaviopsis punctulata

7 cohorts has same trend in CRC entiched.
4 cohorts play significant different.



Colletotrichum fioriniae

7 cohorts has same trend in CRC depleted.
4 cohorts play significant different.