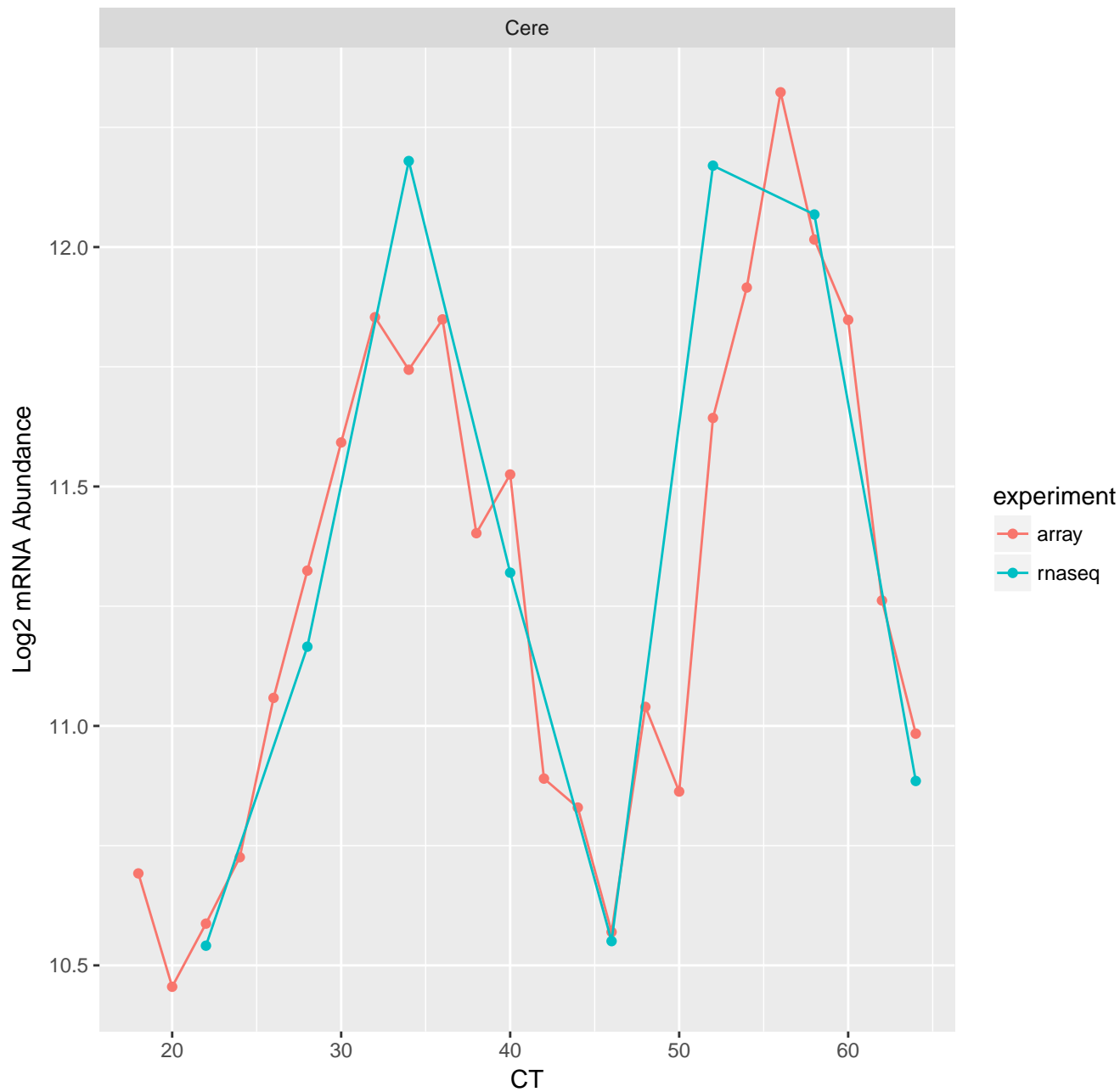
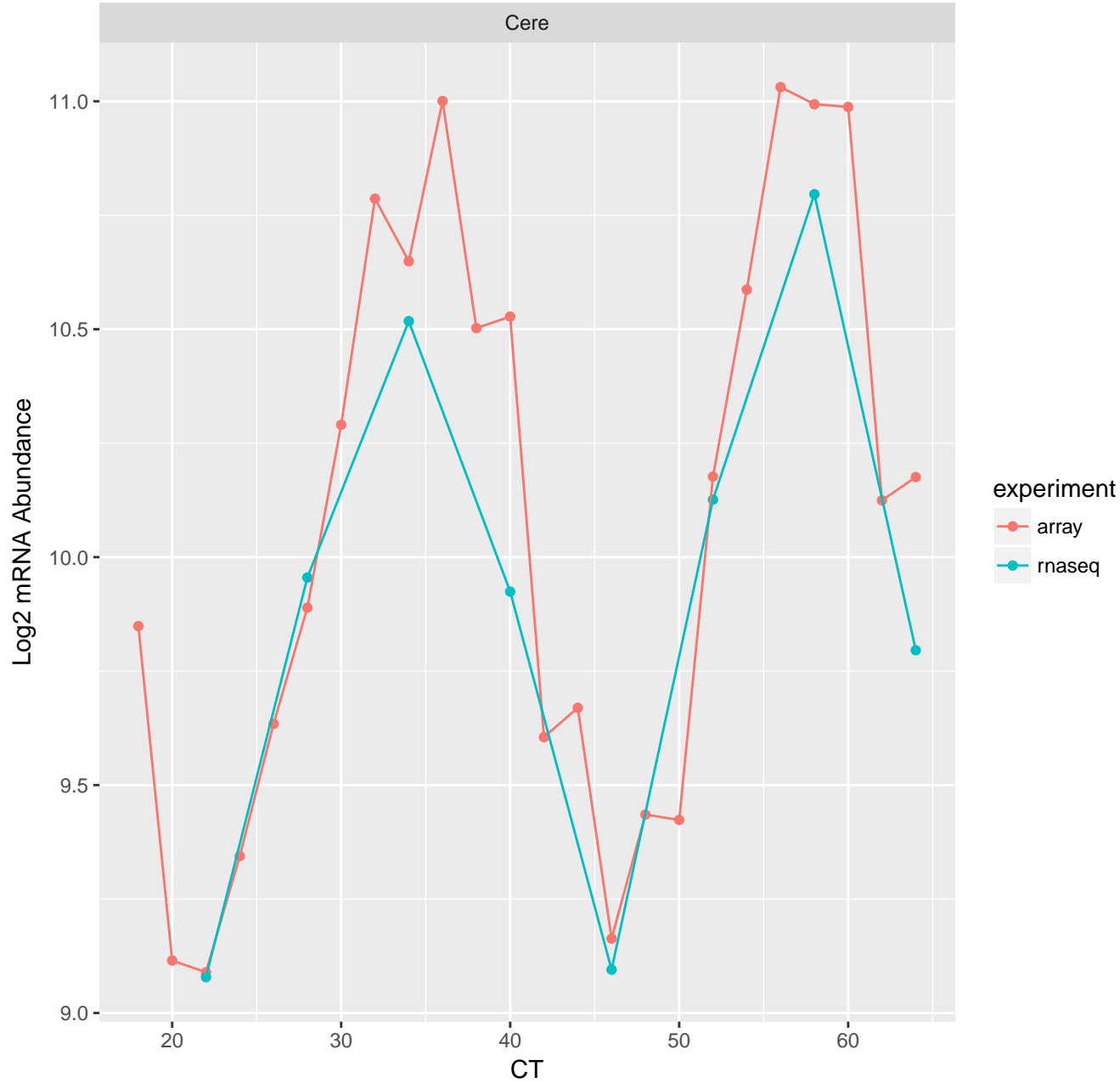


Nr1d1



Dbp



Per2

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

CT

11.0
10.5
10.0
9.5
9.0

20

30

40

50

60

11.0
10.5
10.0
9.5
9.0

Per3

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

20

CT

50

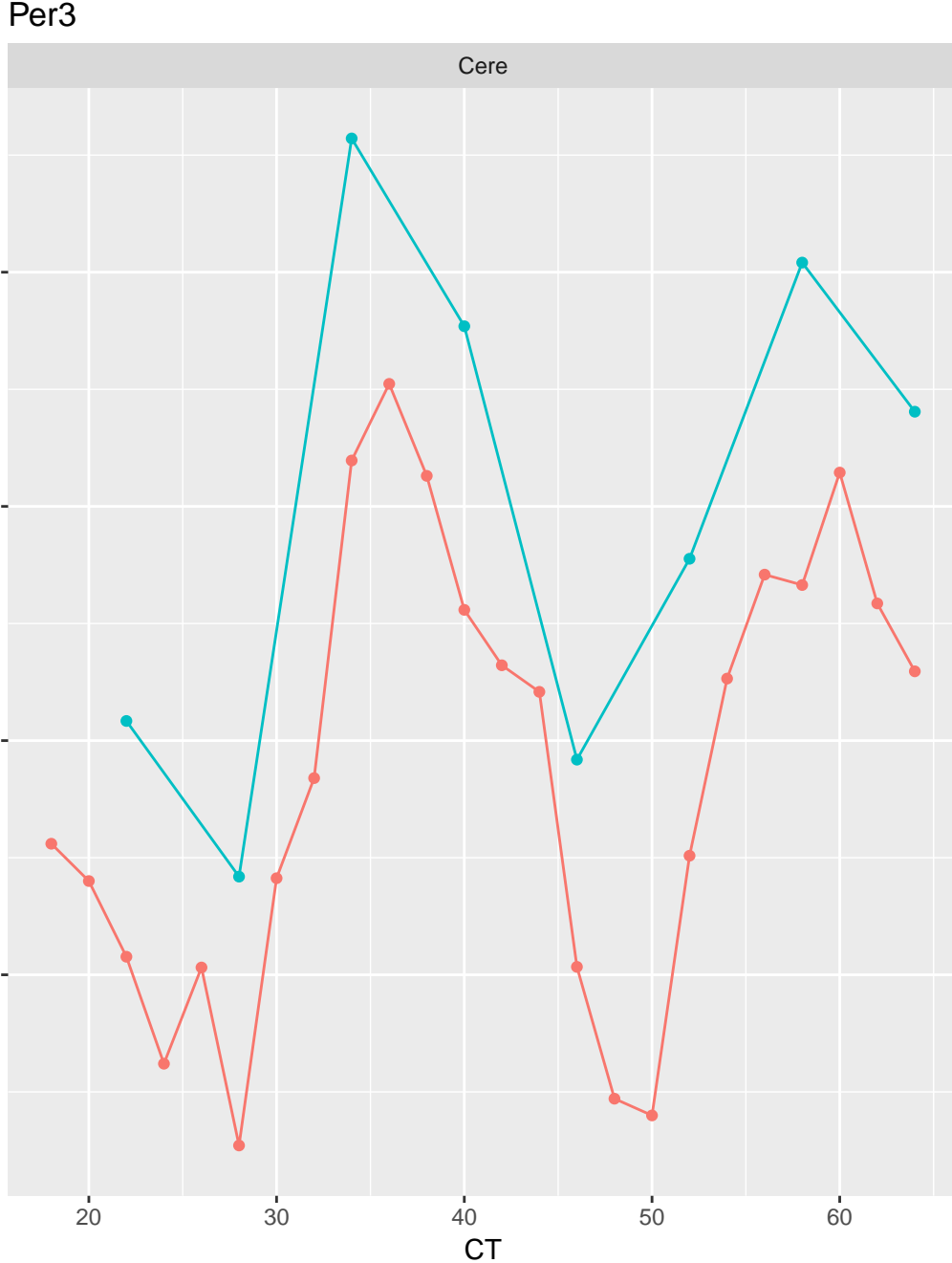
60

11.00

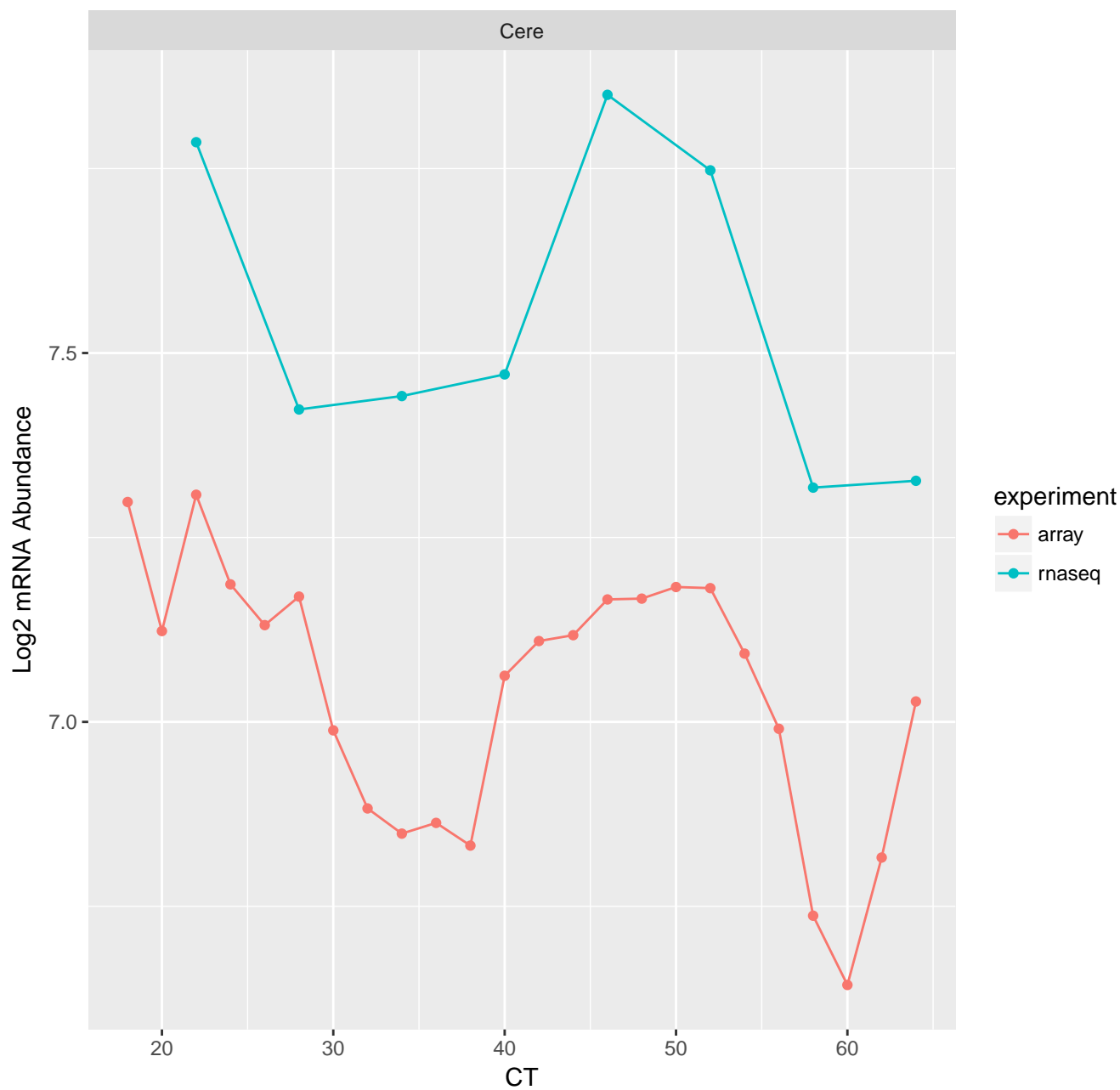
10.75

10.50

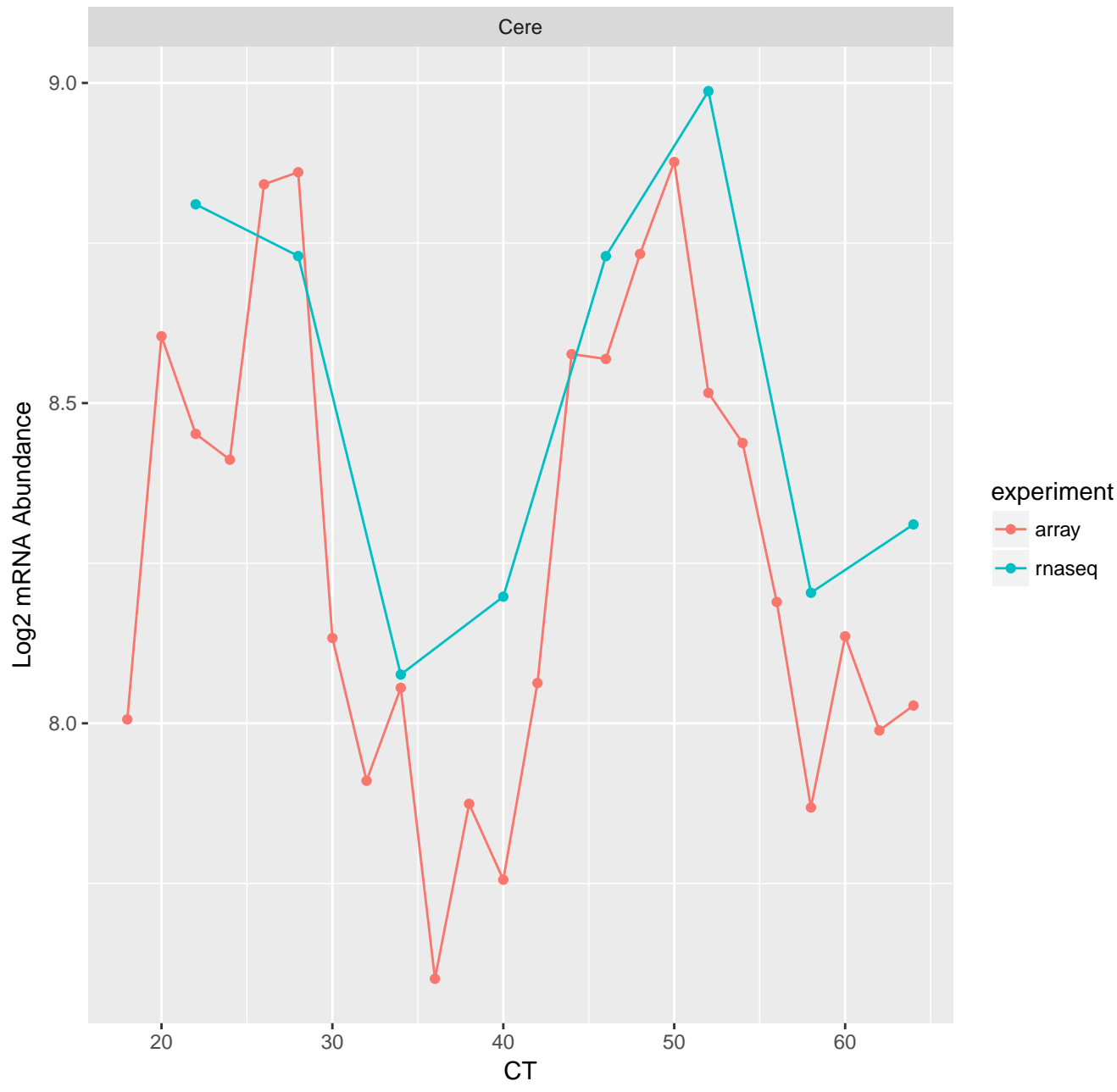
10.25



Rhoj



Mmp14



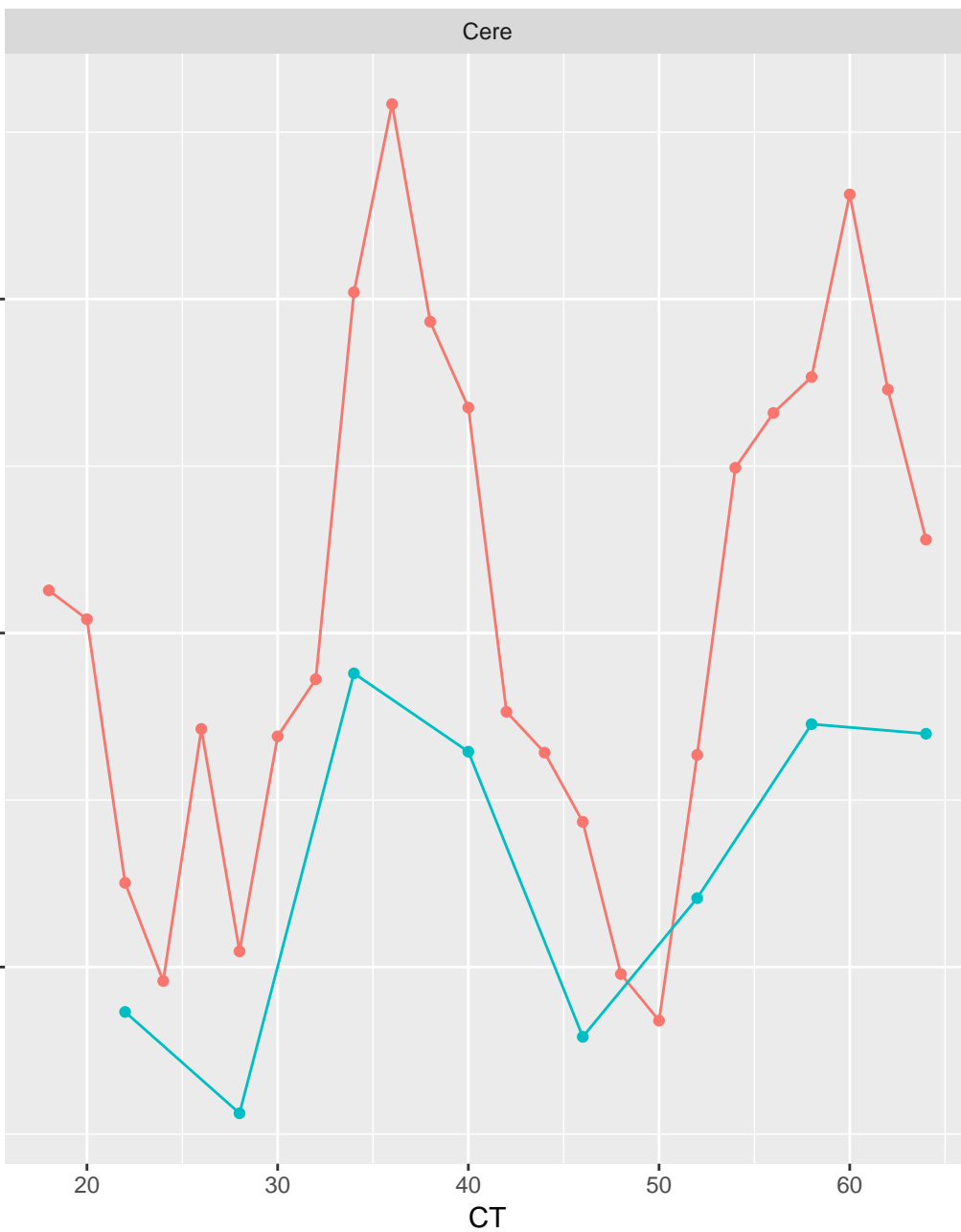
Tef

Cere

Log2 mRNA Abundance

experiment

array
rnaseq



Arntl

Cere

Log2 mRNA Abundance

CT

experiment

array
rnaseq

10.8

10.4

10.0

9.6

20

30

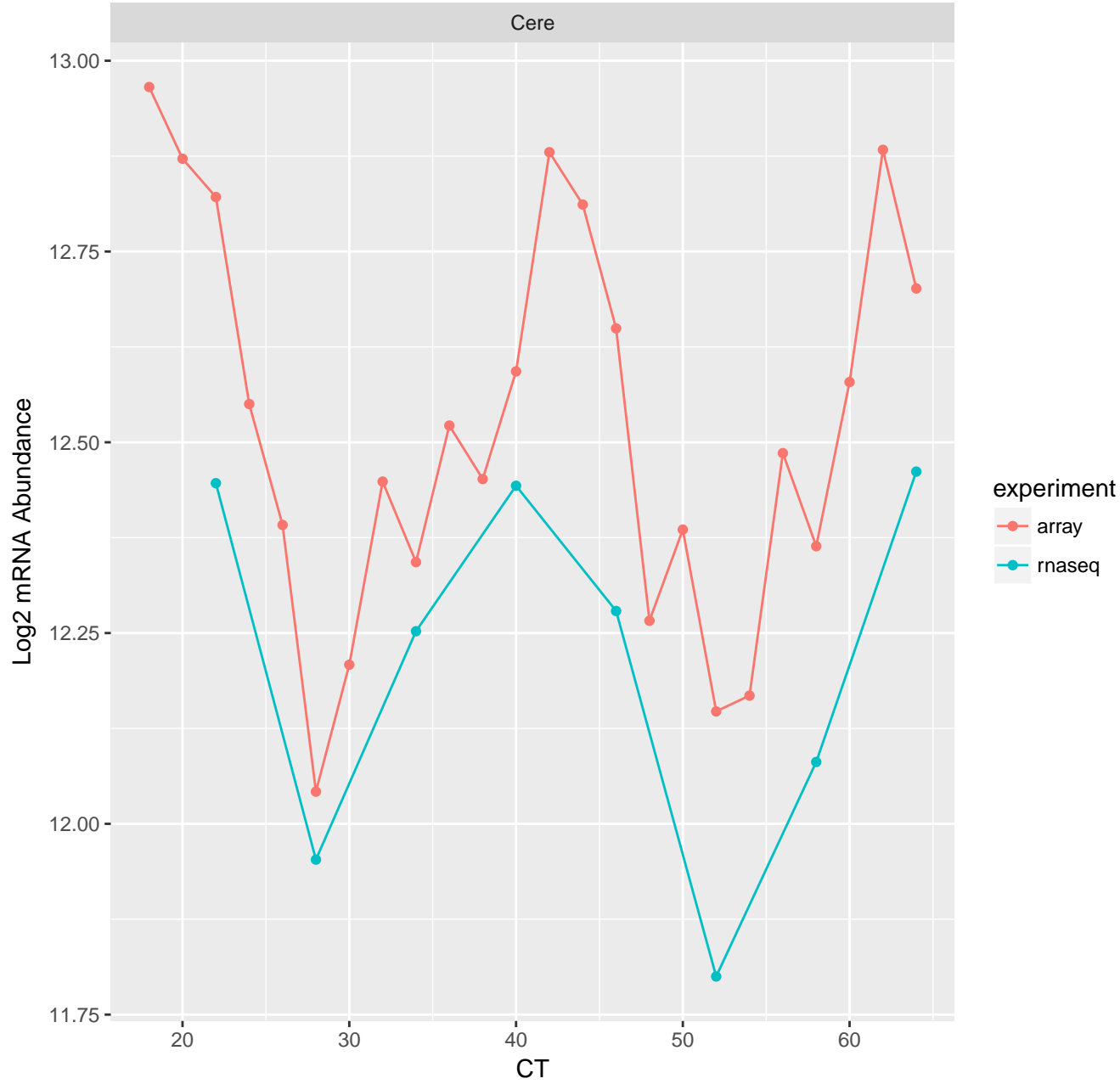
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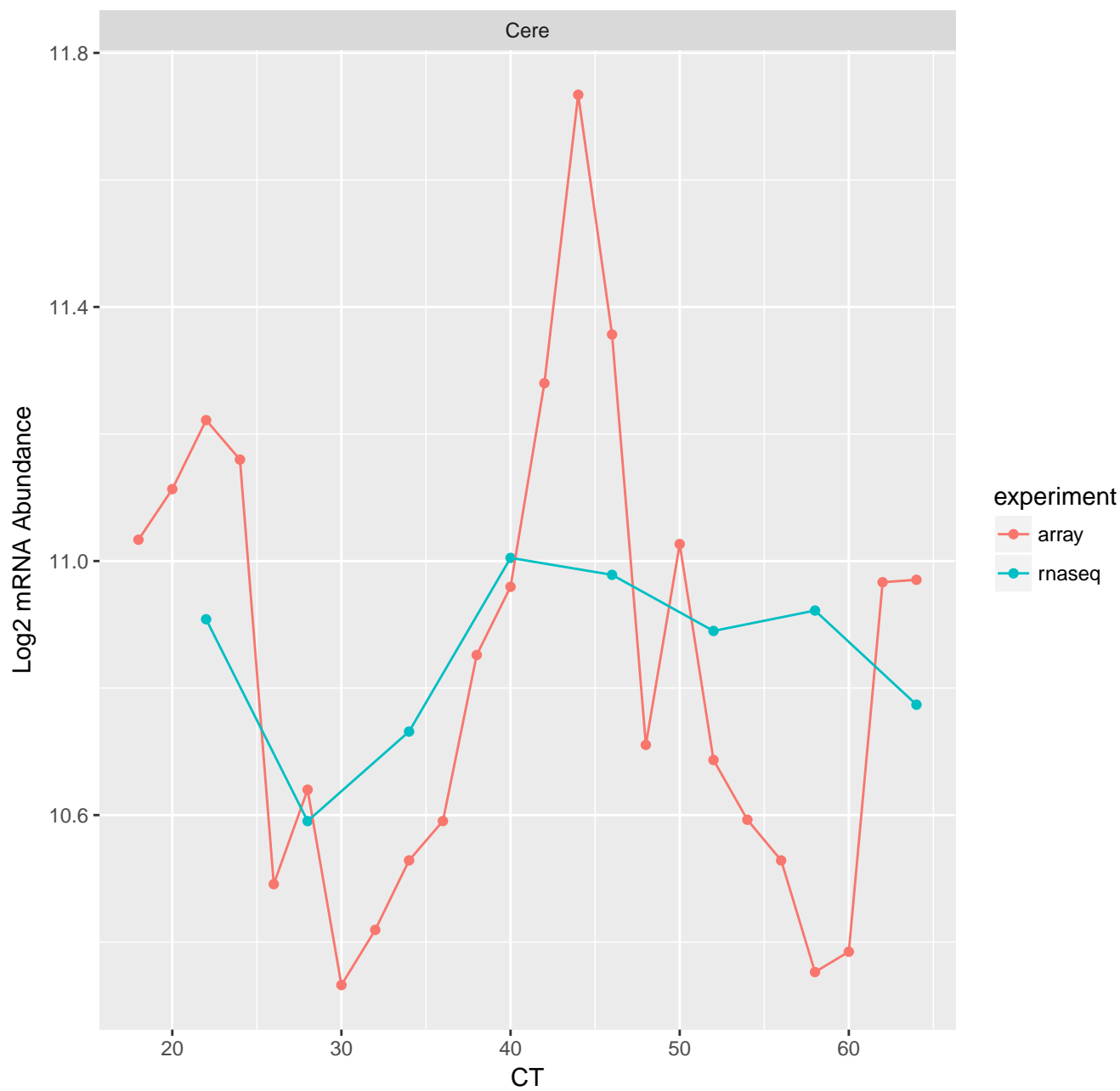
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CT

Paqr8



Ptpn22



Mt2

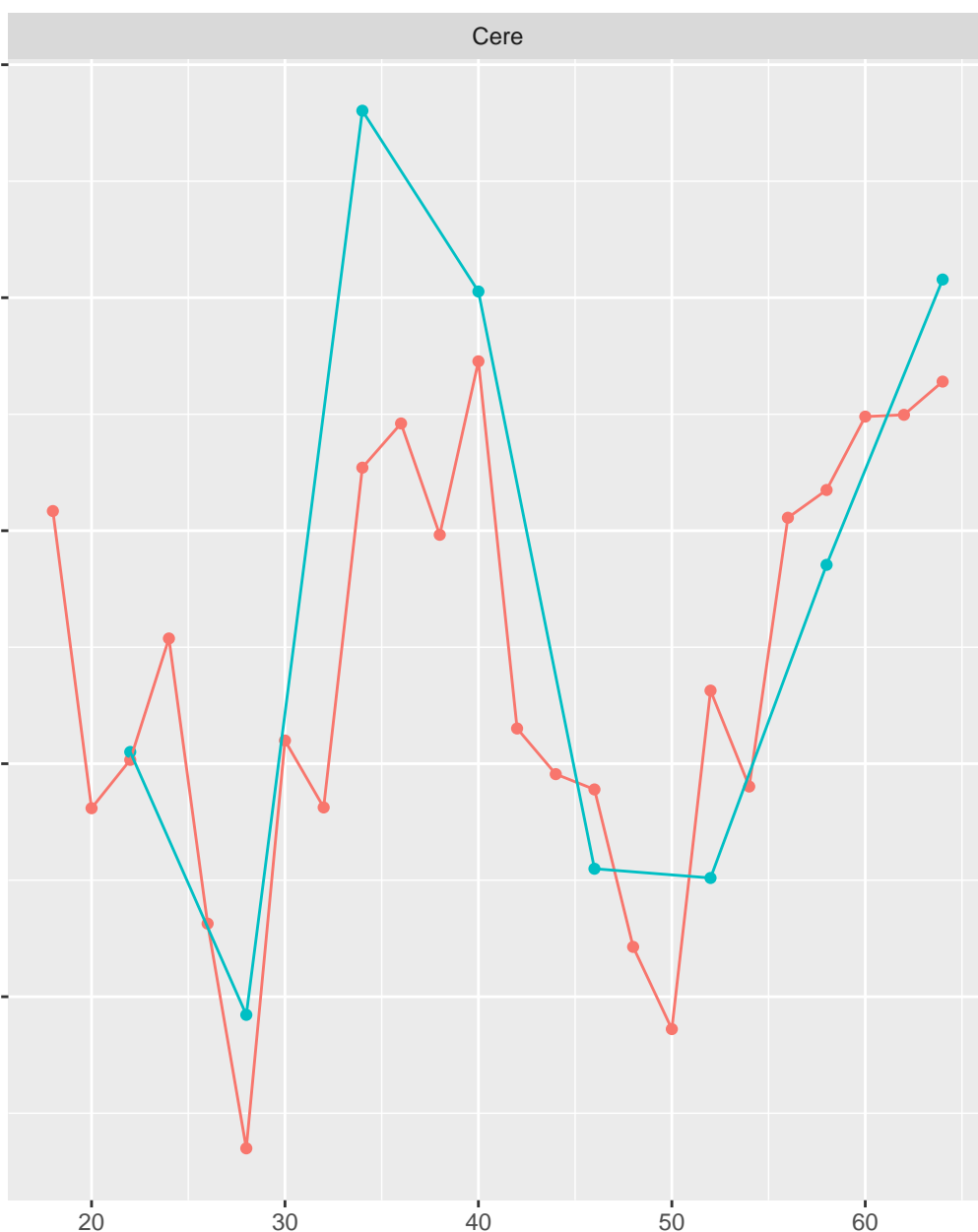
Cere

Log2 mRNA Abundance

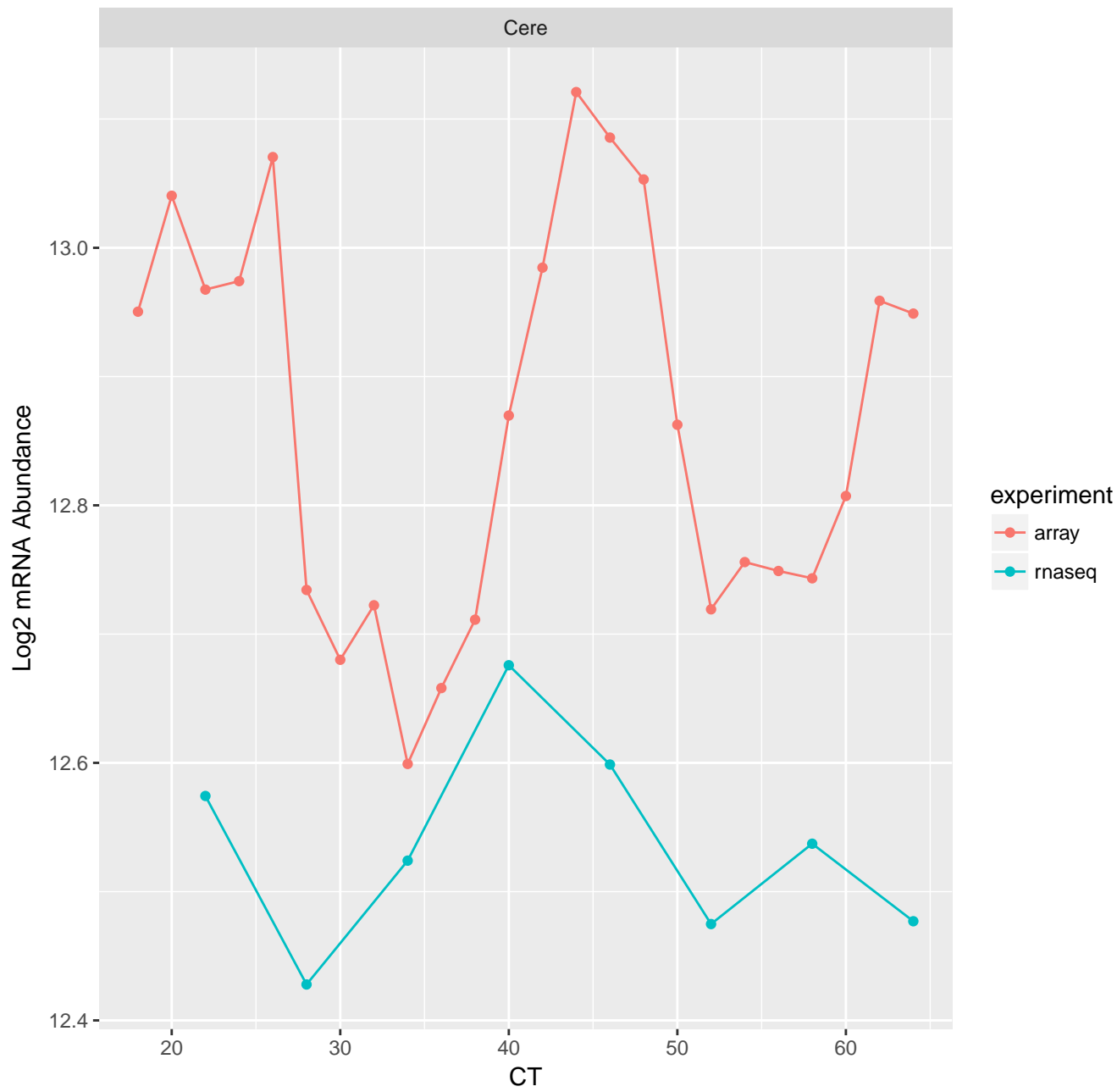
experiment

array
rnaseq

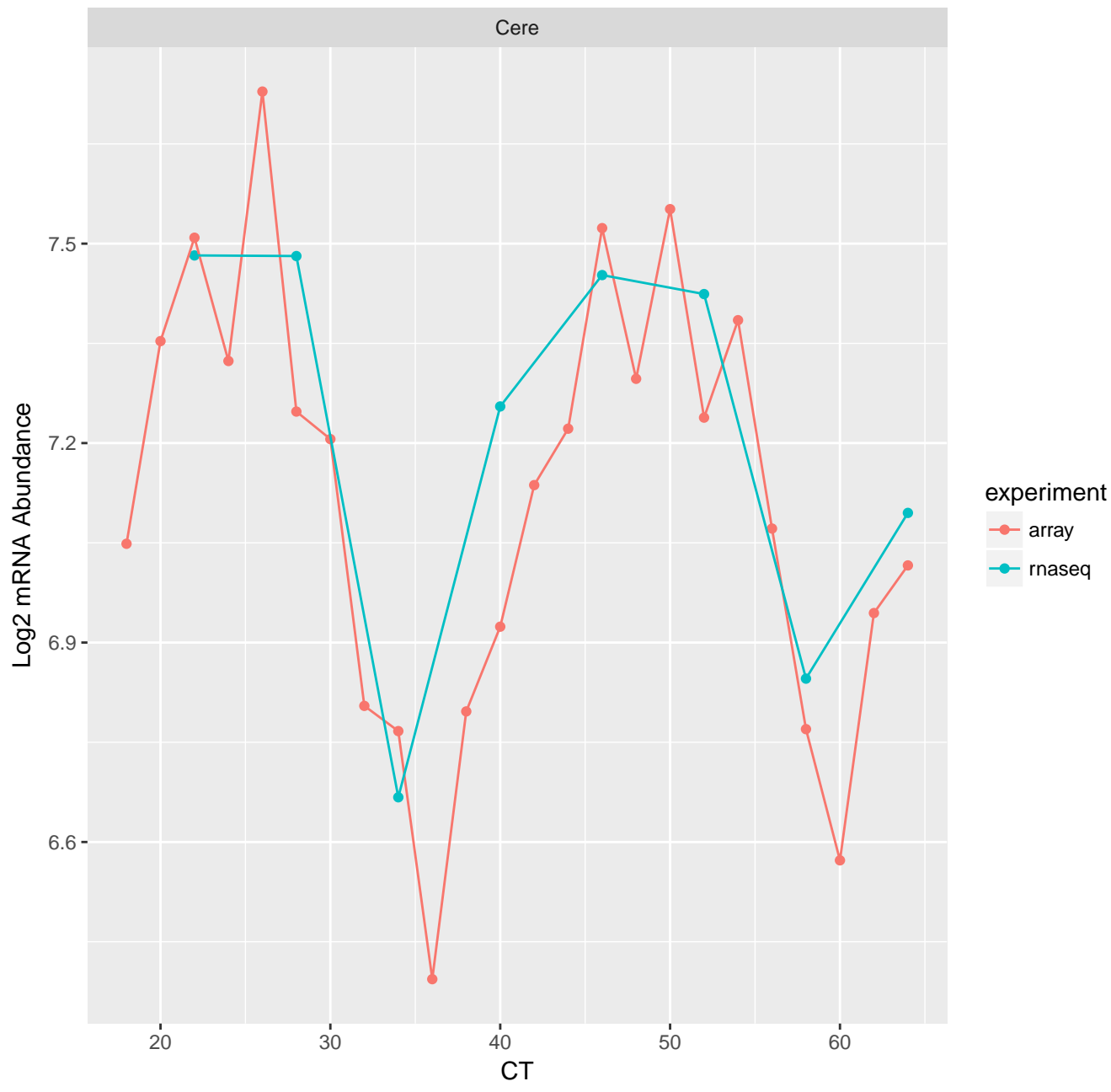
CT



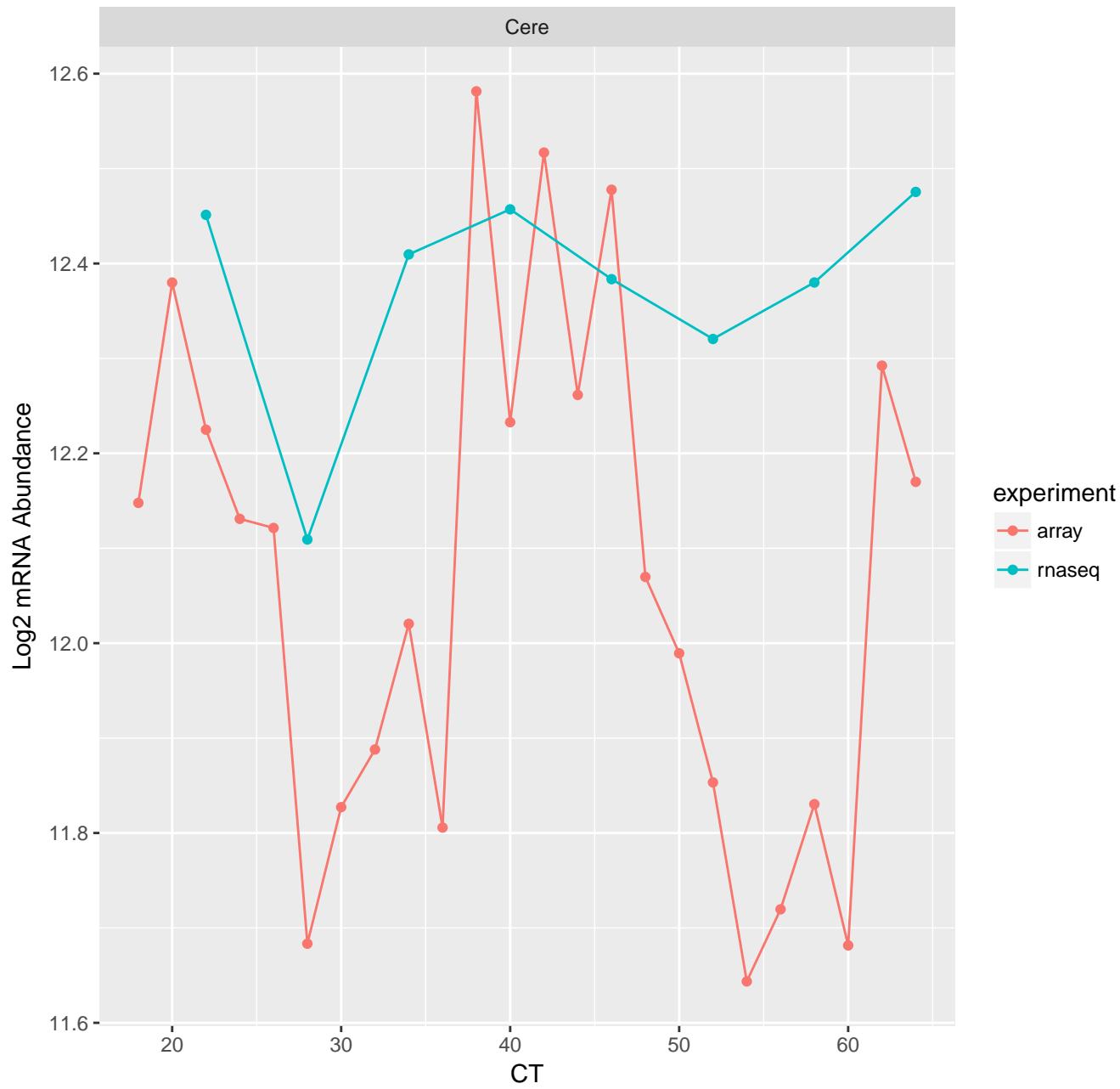
Tcf4



Vstm4



Dusp11



Hlf

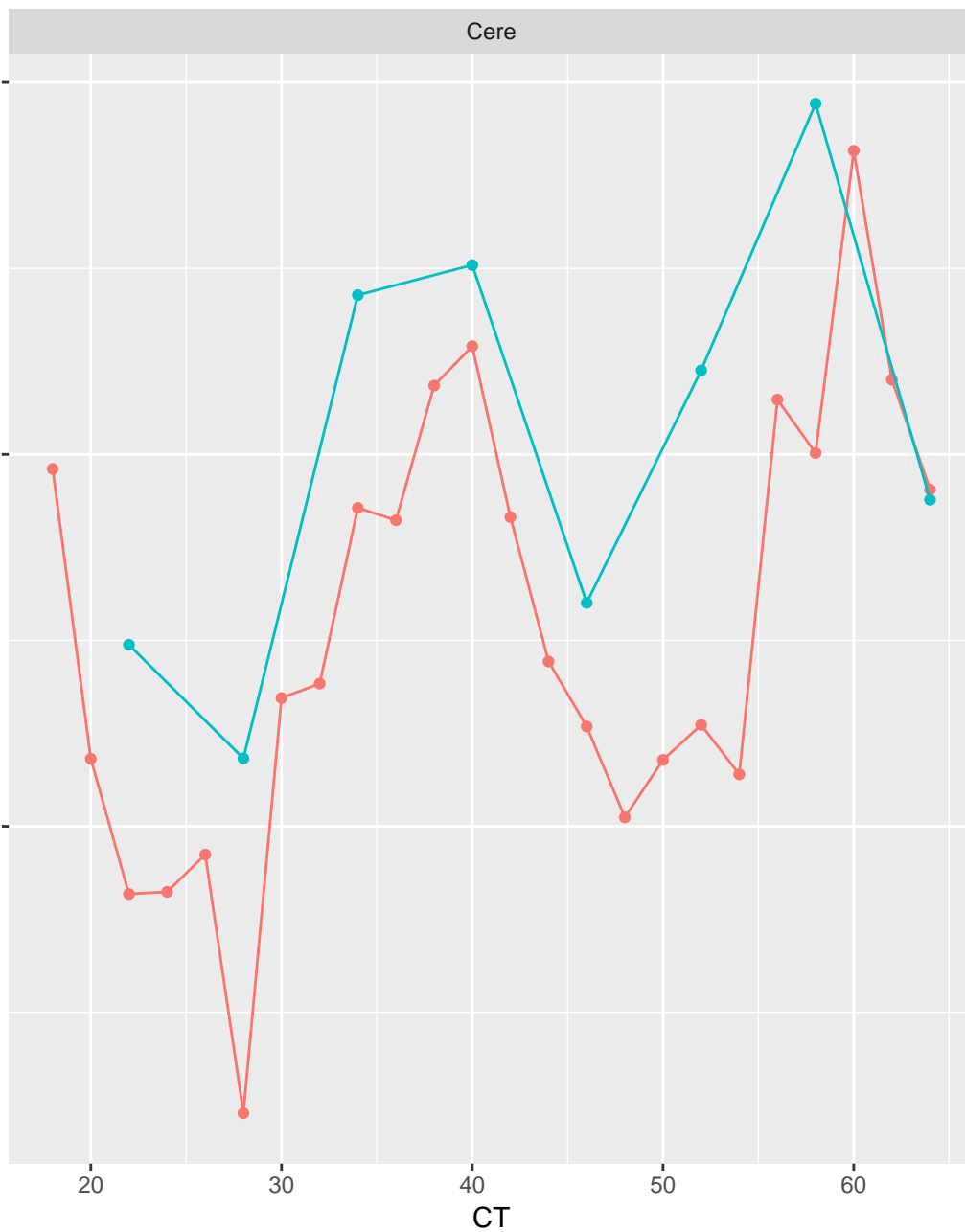
Cere

Log2 mRNA Abundance

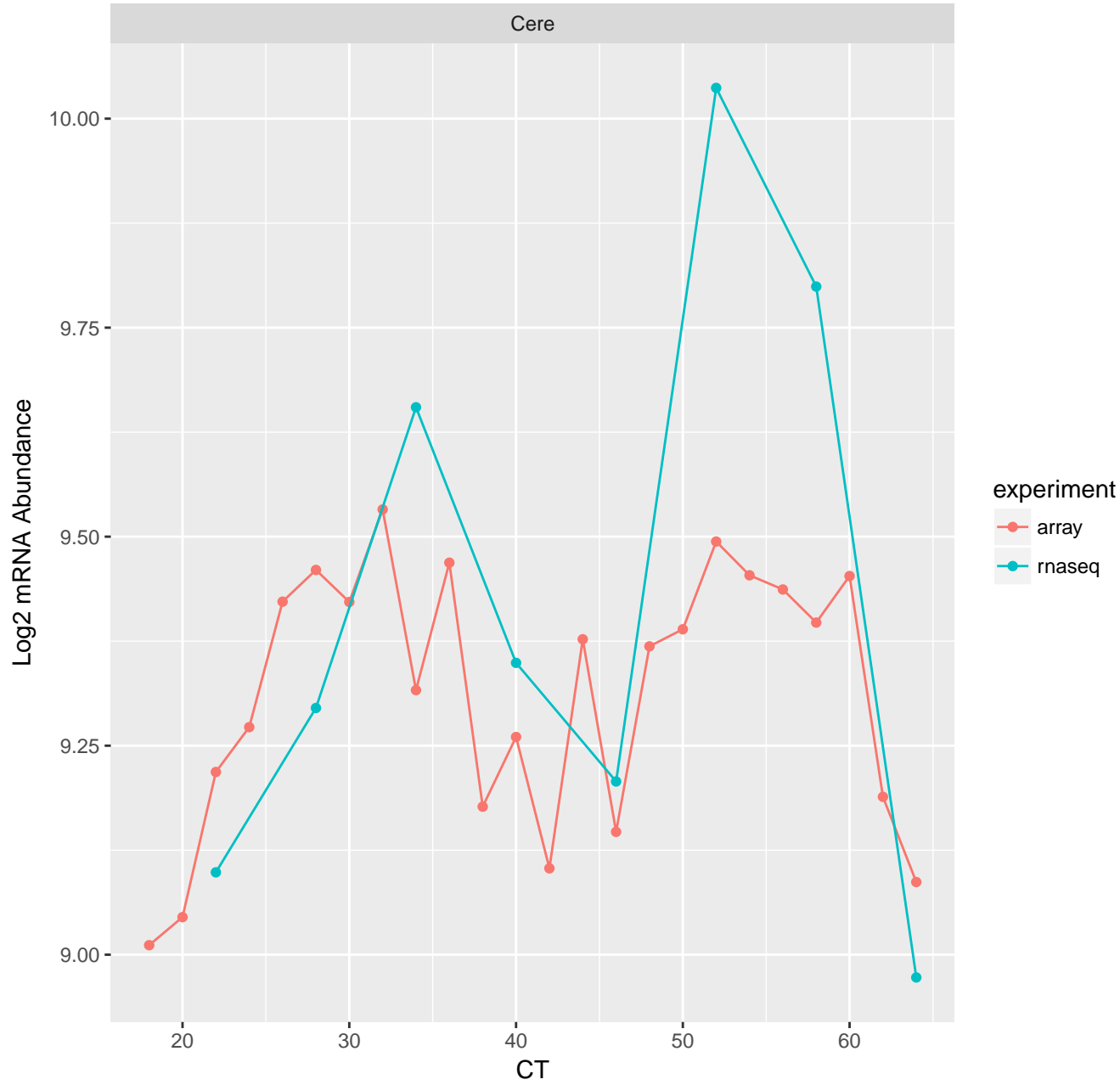
CT

experiment

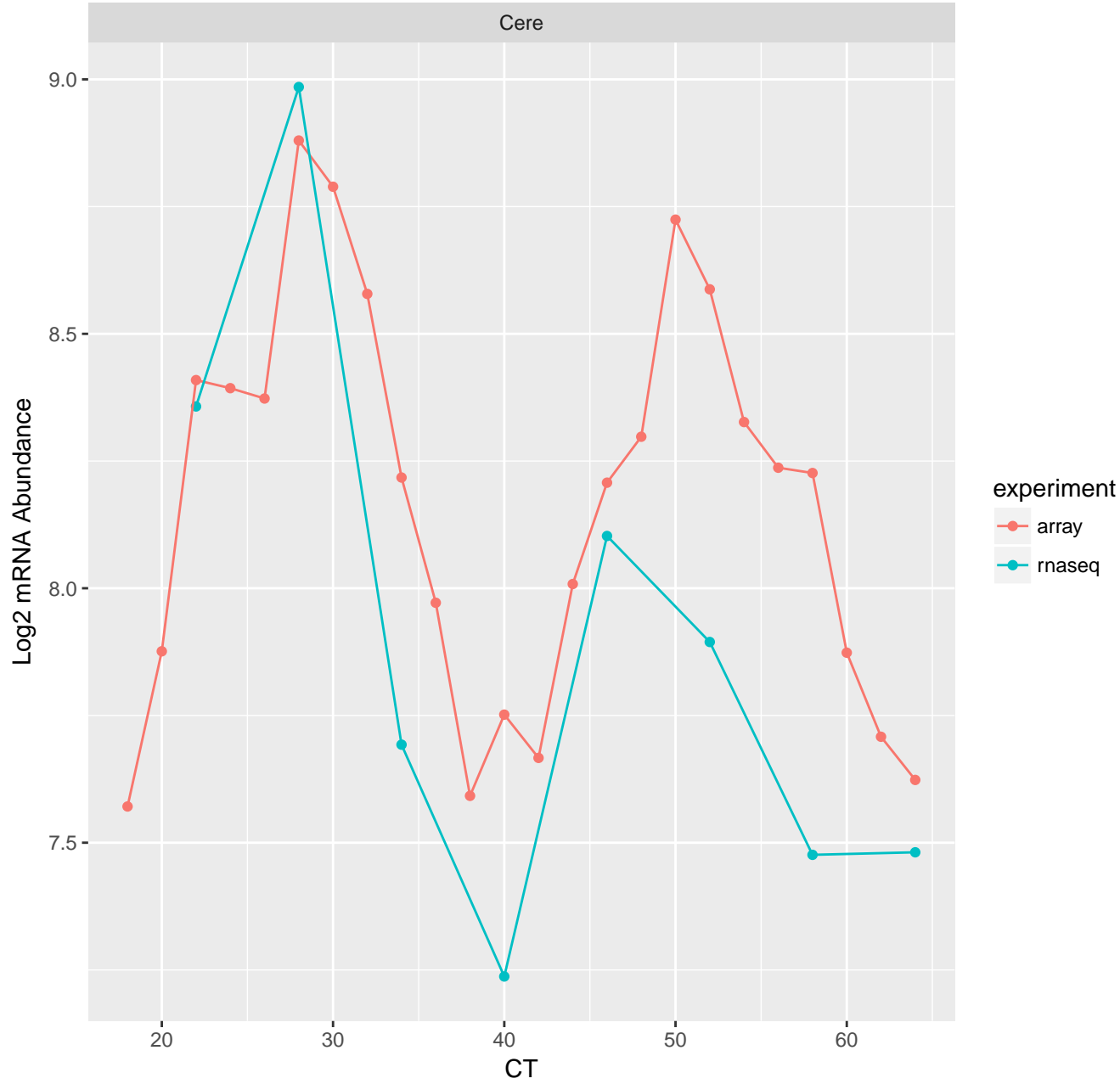
array
rnaseq



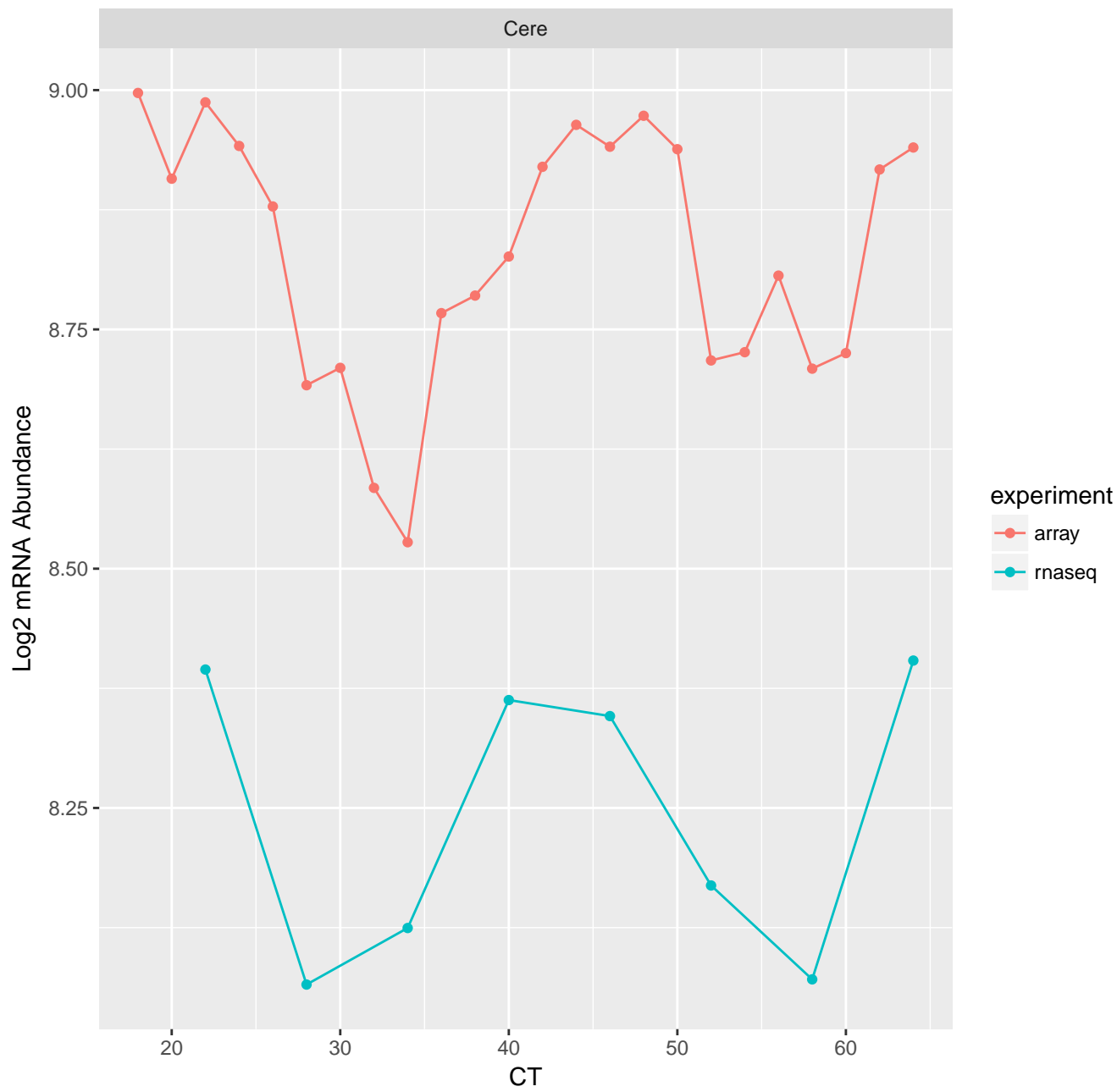
Cpne2



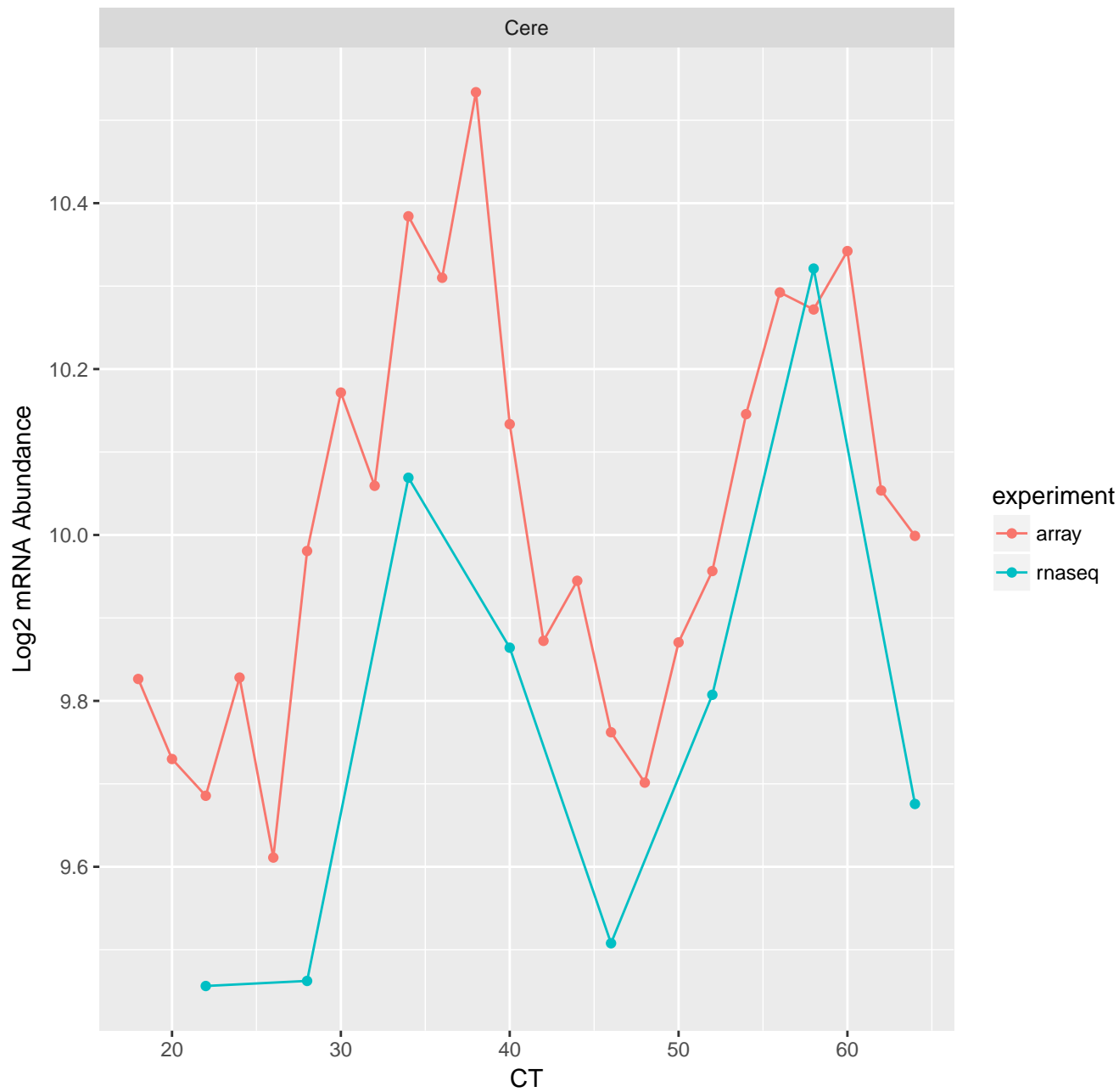
Fabp7



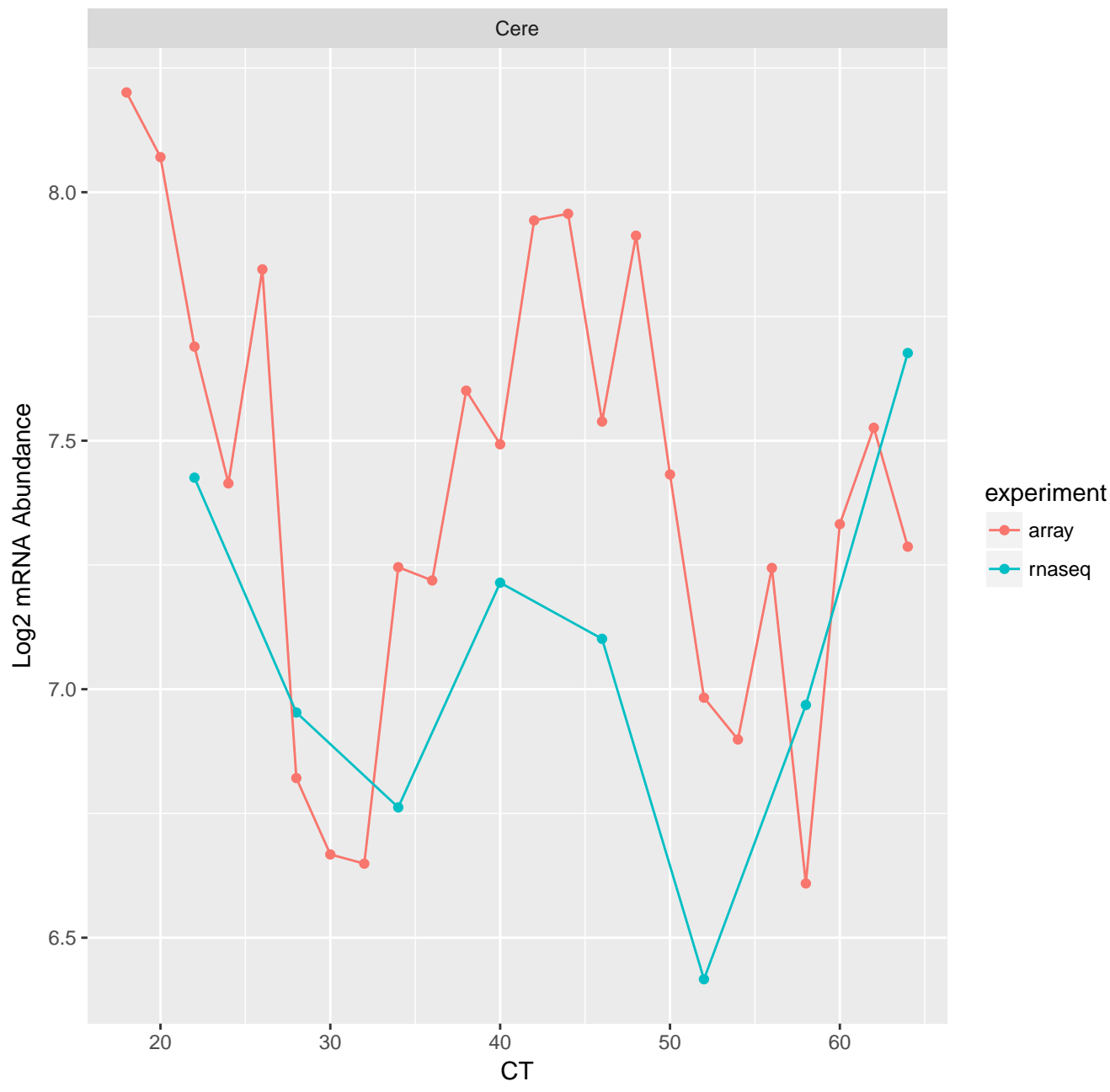
Bbs9



Bhlhe41



Lonrf3



Atxn1

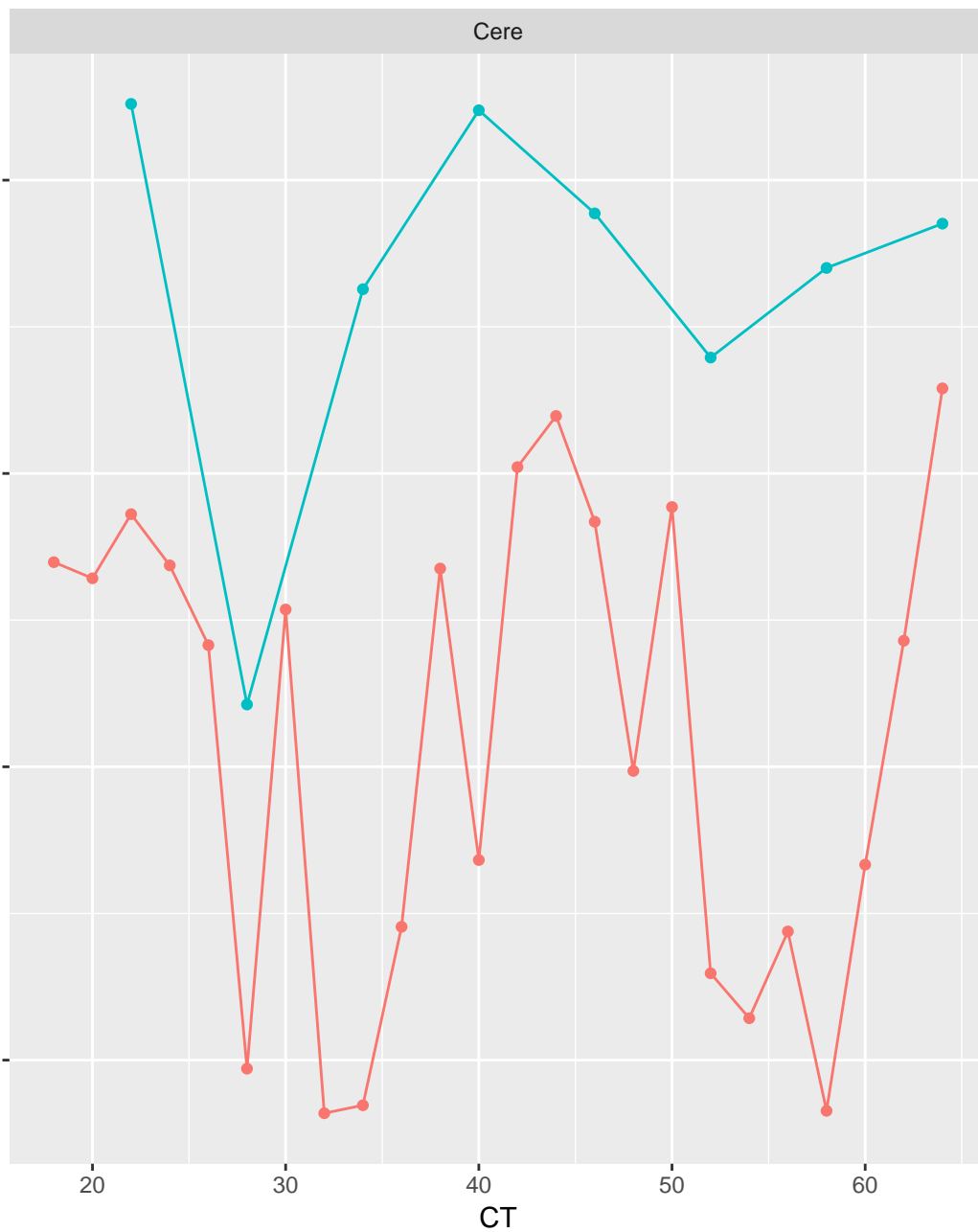
Cere

Log2 mRNA Abundance

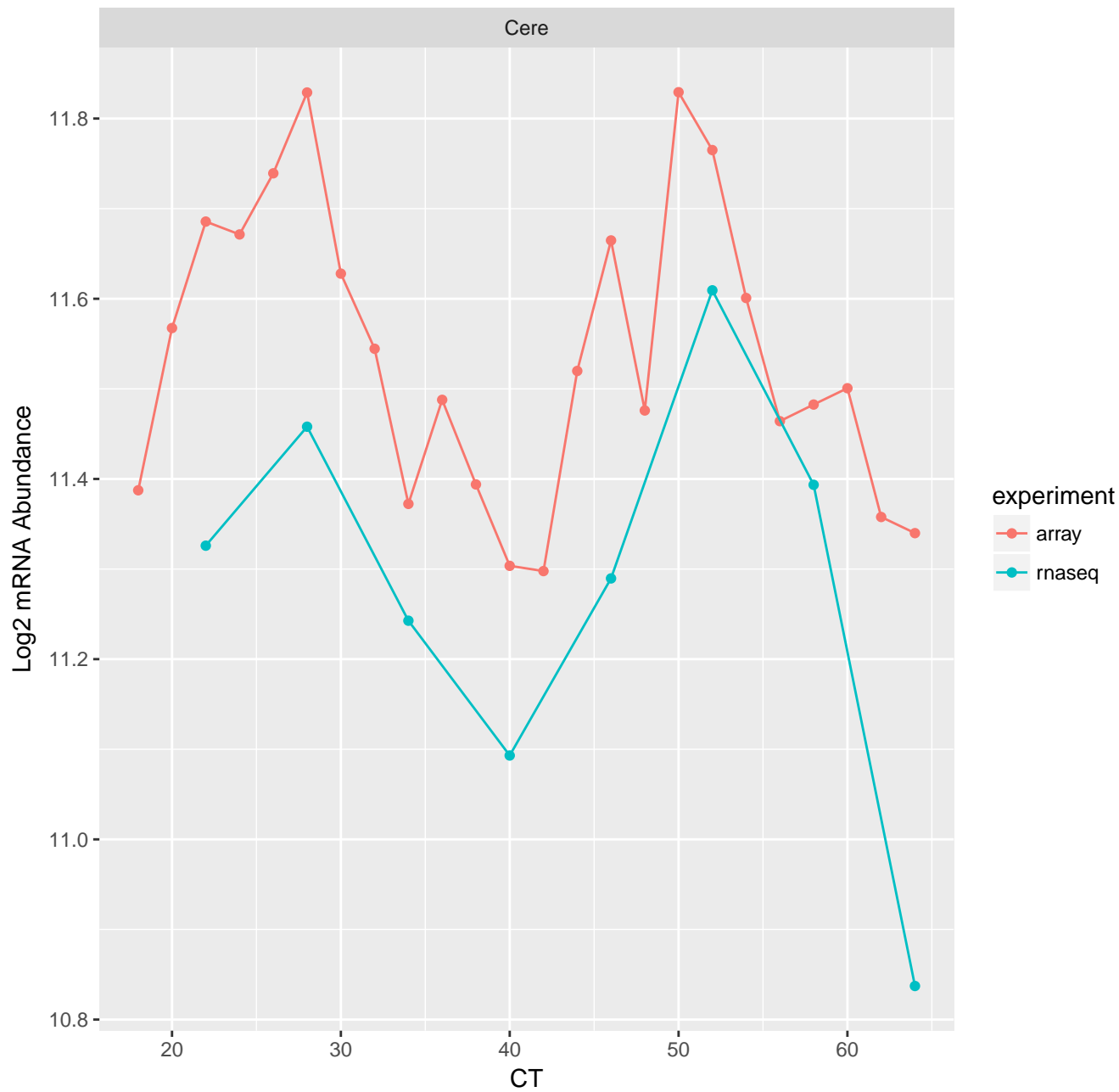
experiment

array
rnaseq

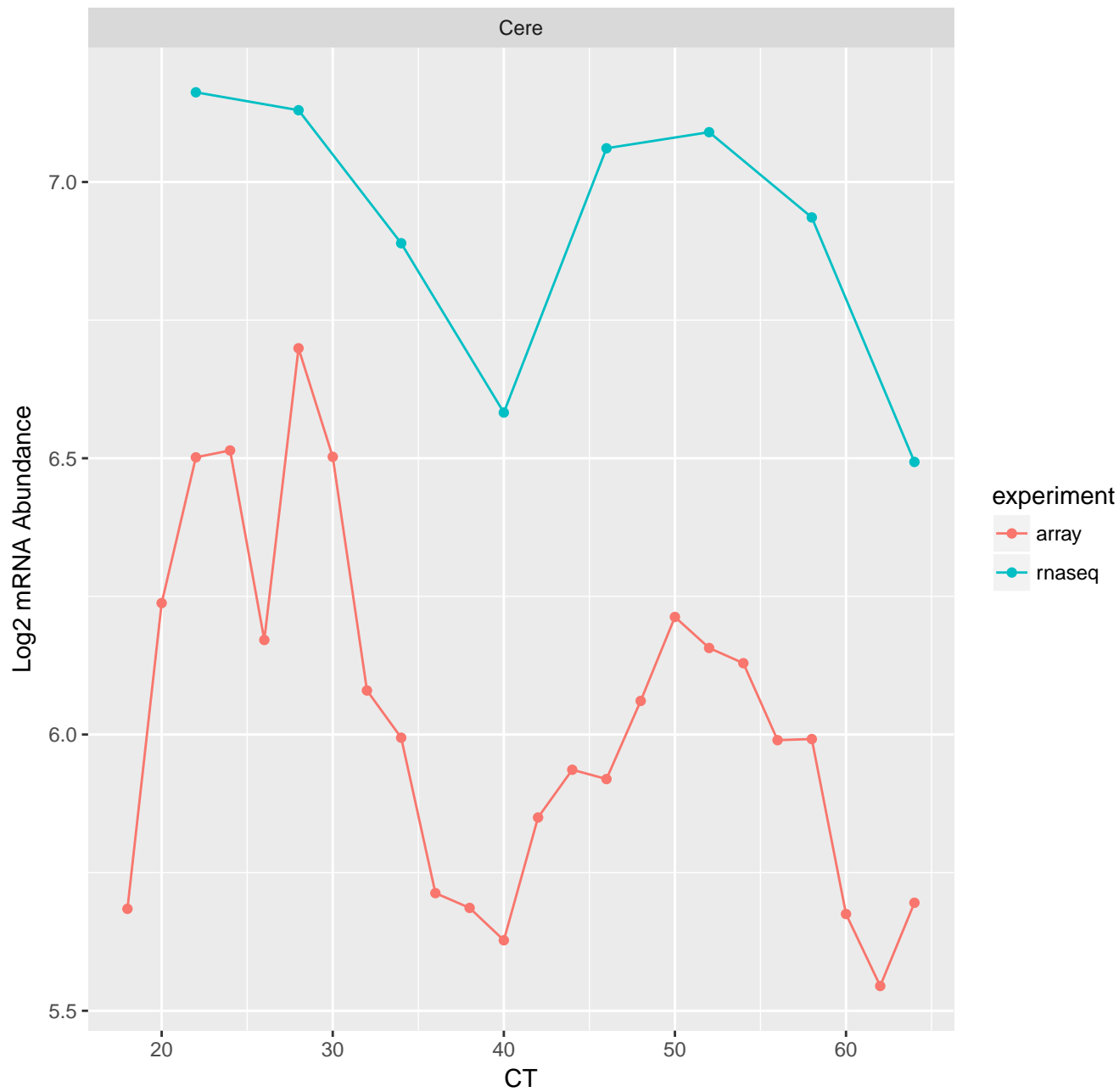
CT



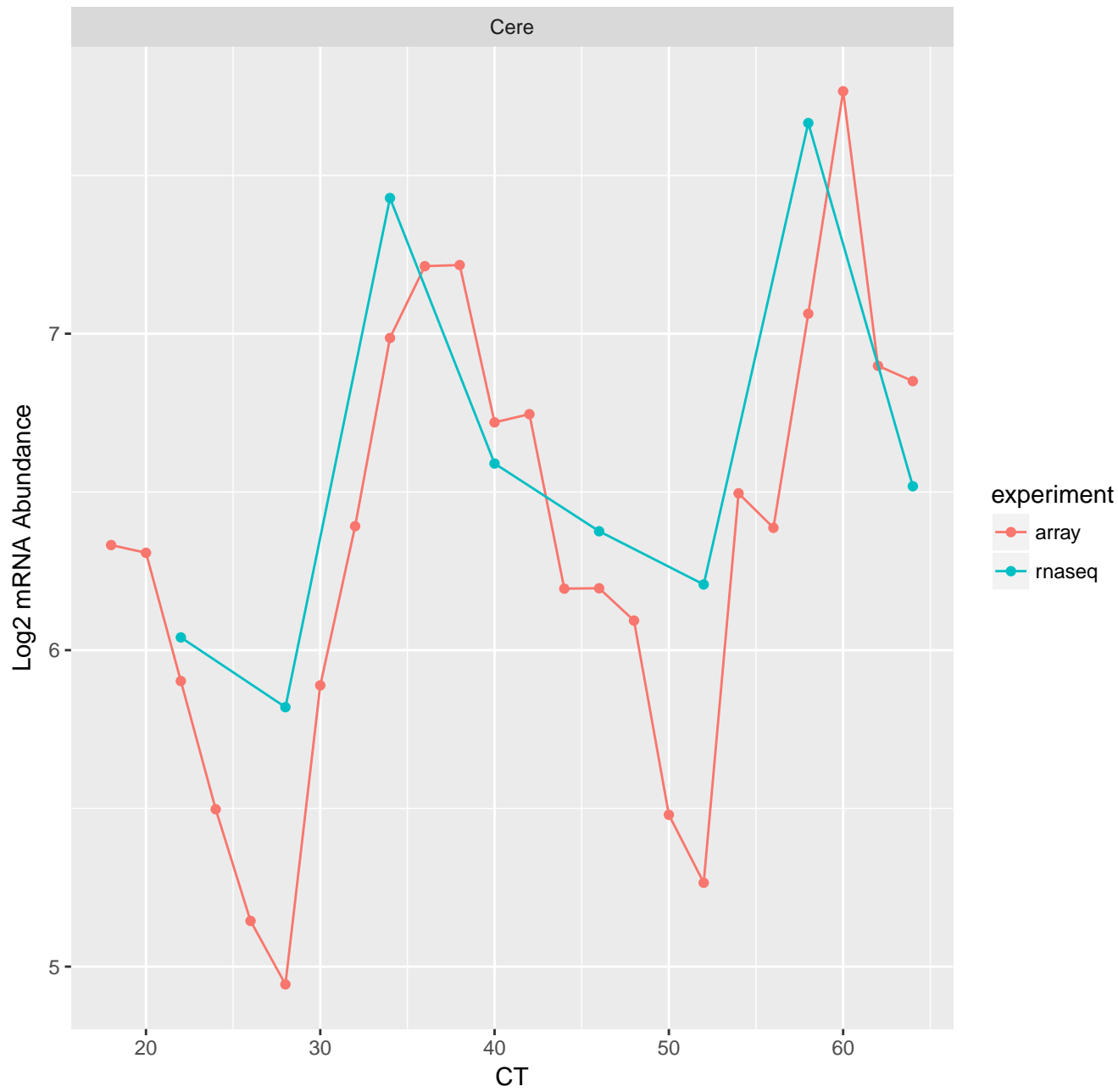
Vat1l



Asxl3



Fmo2



Ift80

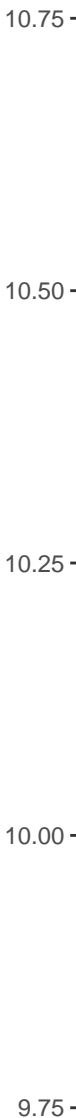
Cere

Log2 mRNA Abundance

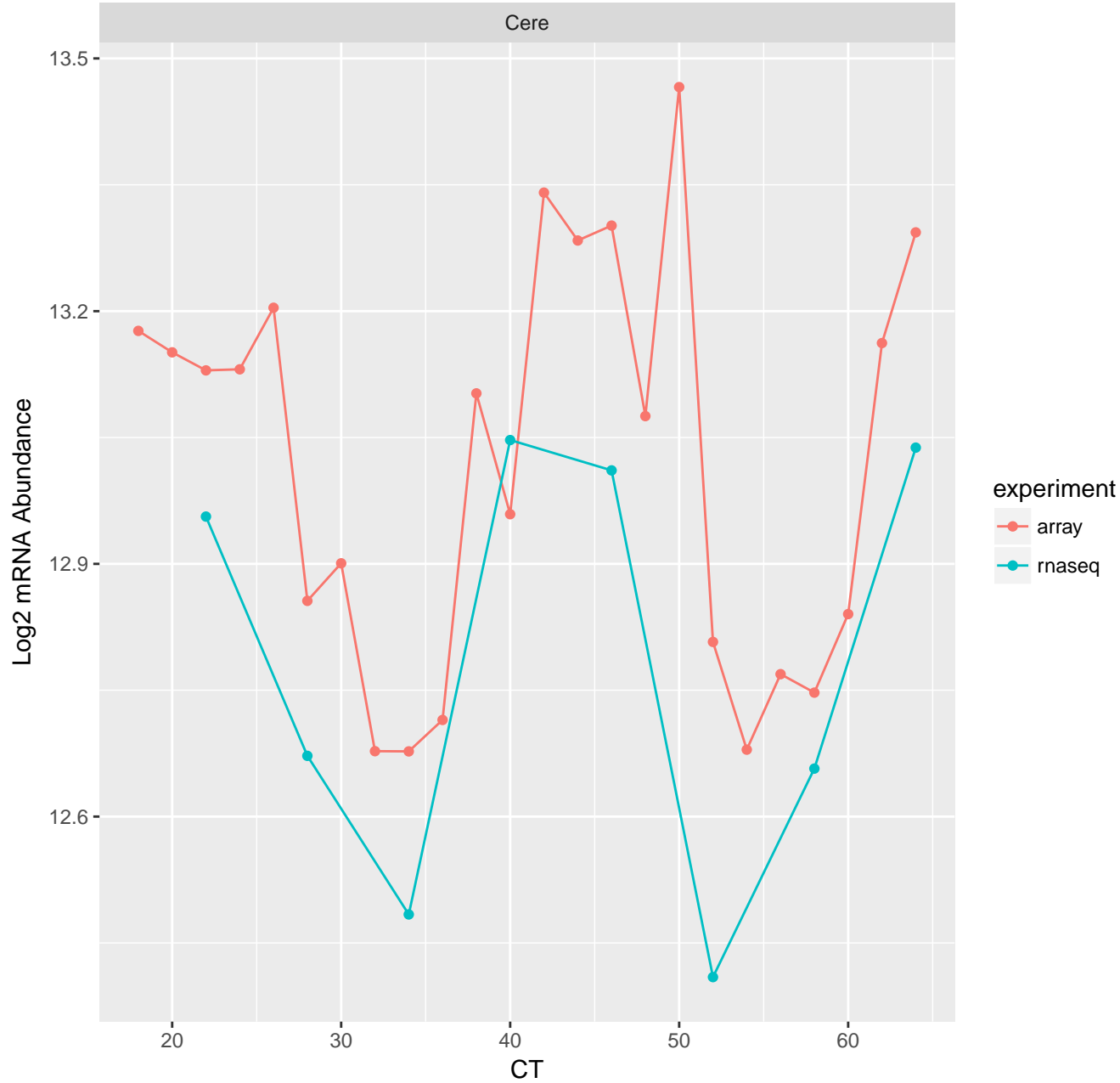
experiment

array
rnaseq

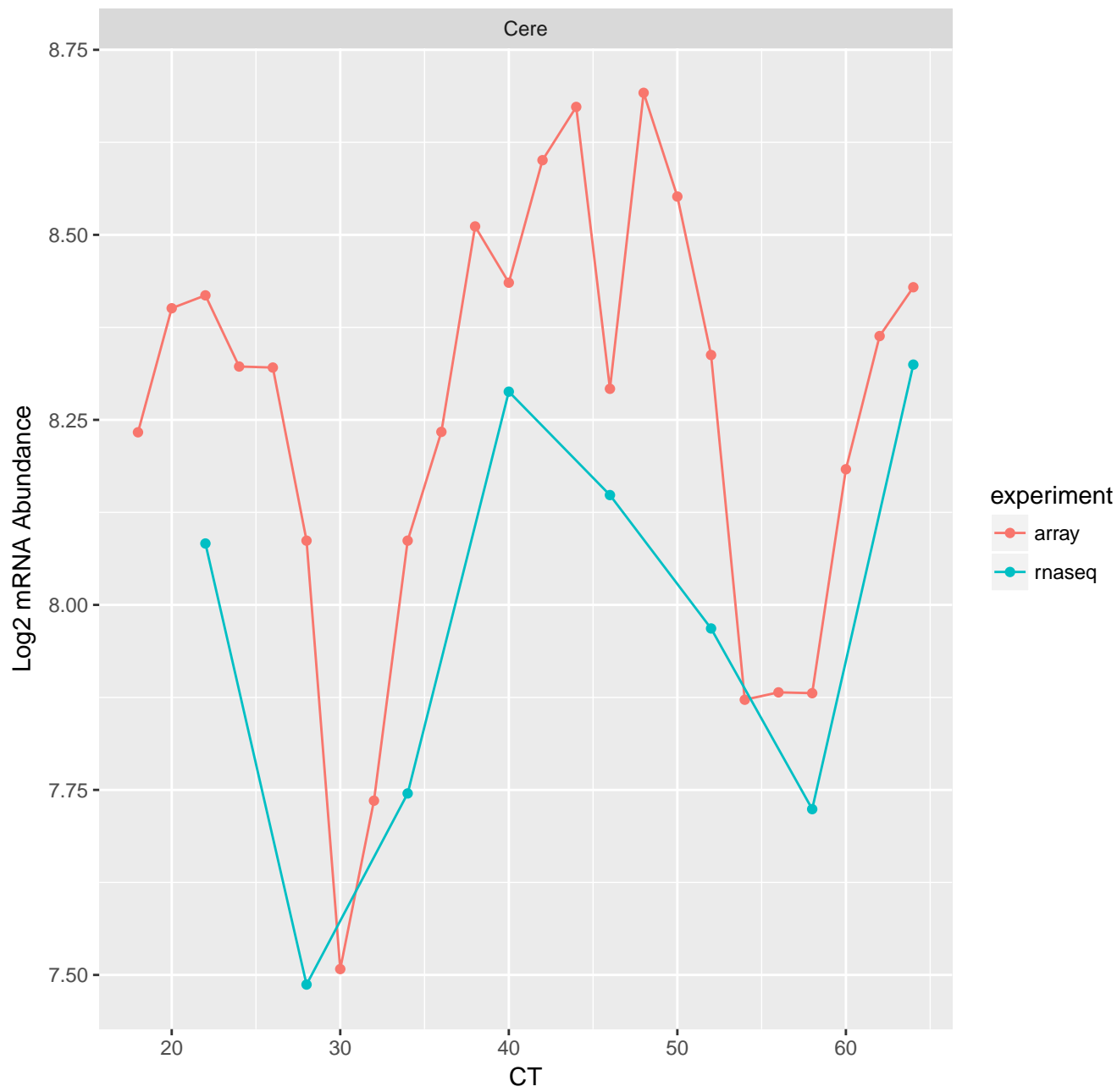
CT



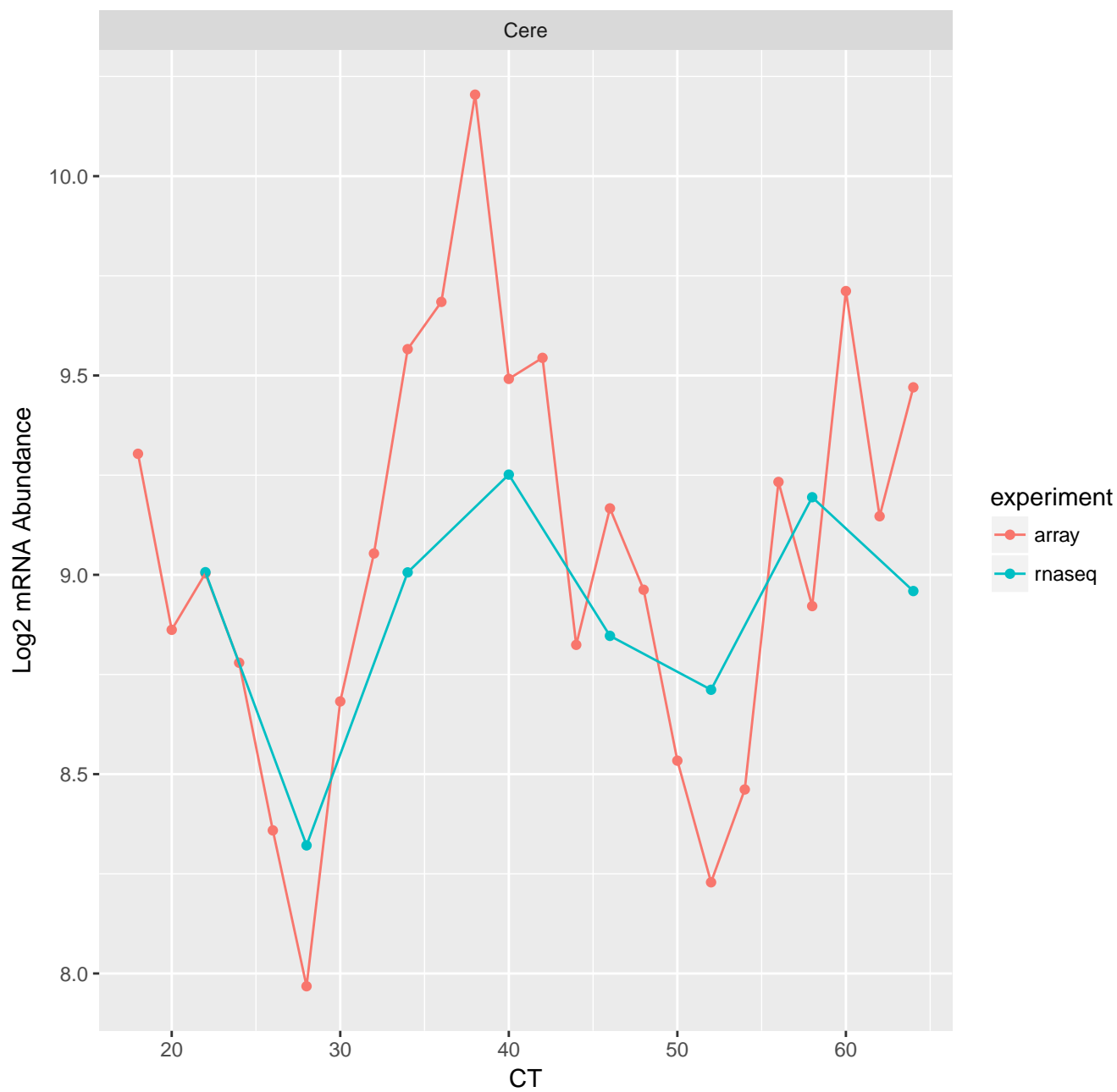
Hsp90b1



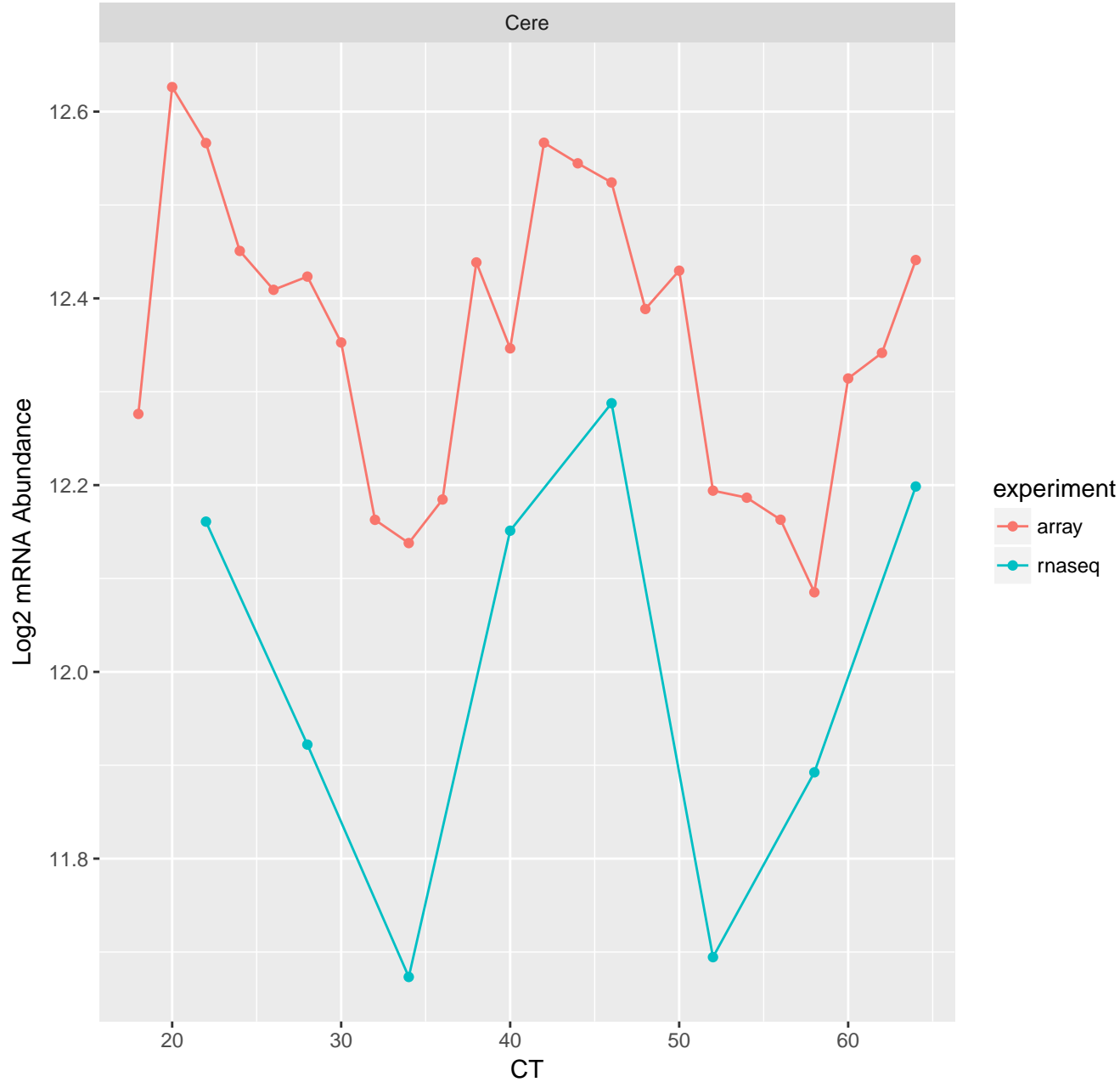
Thbd



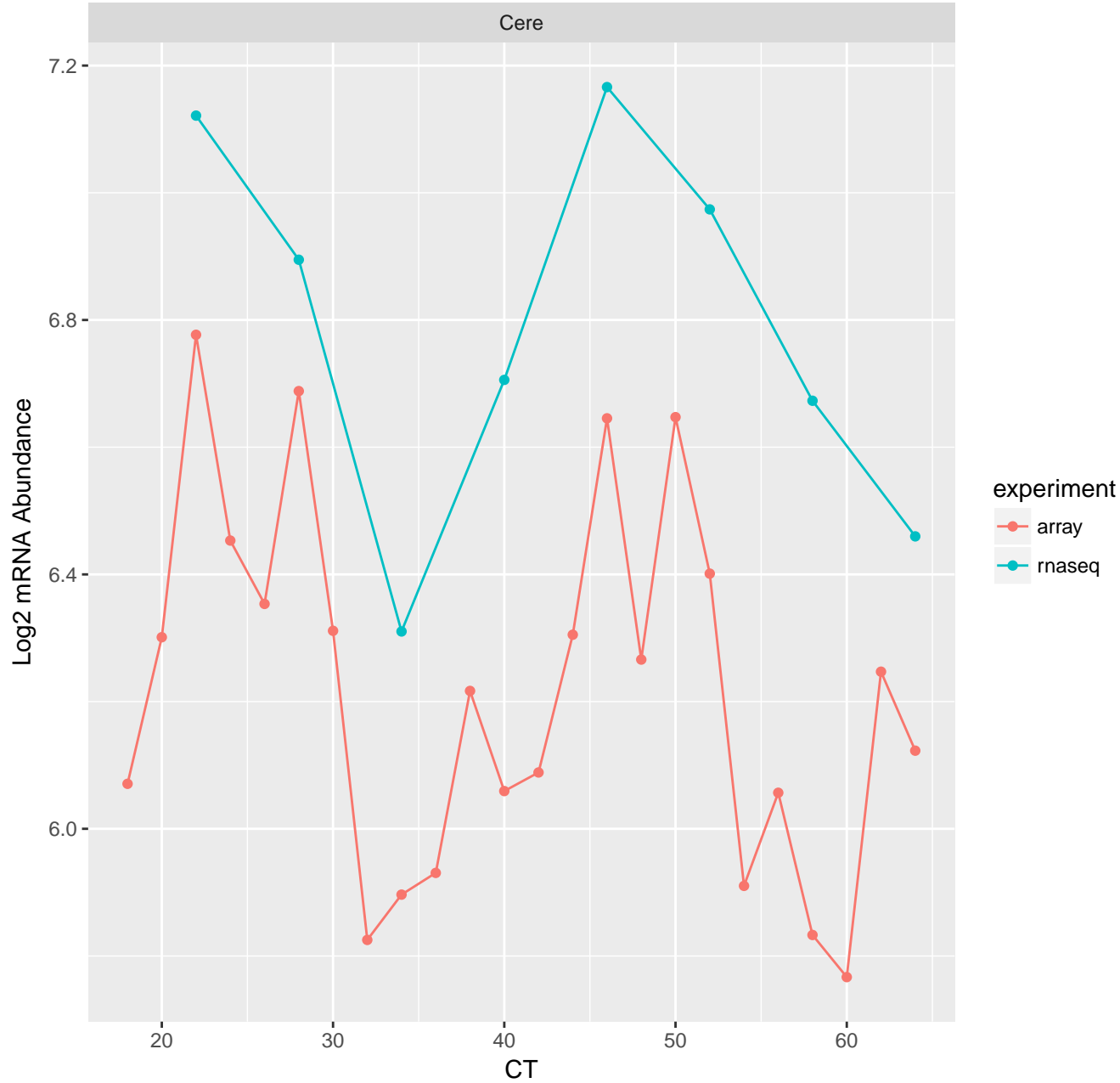
Stk32a



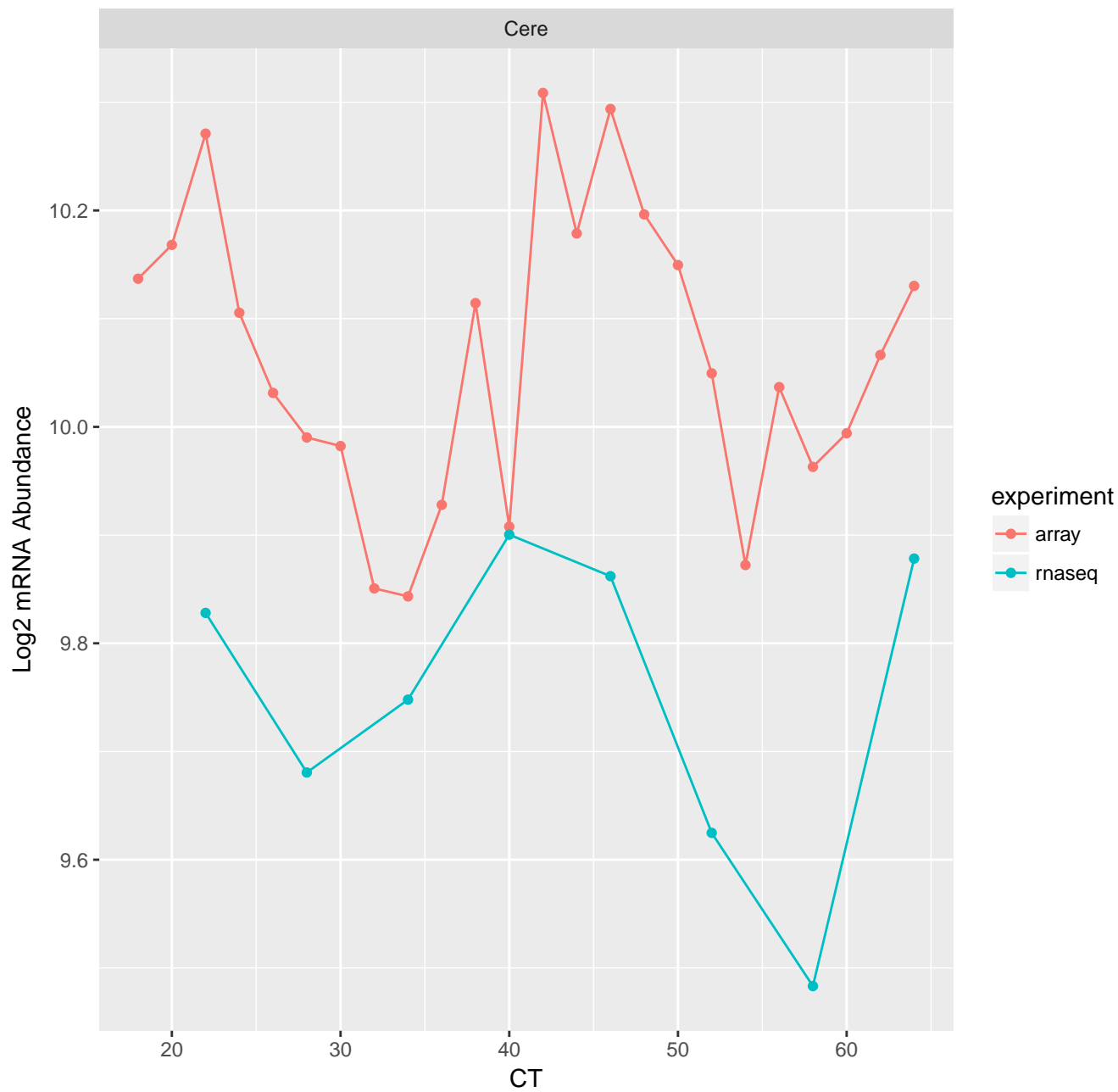
Hnrrnpm



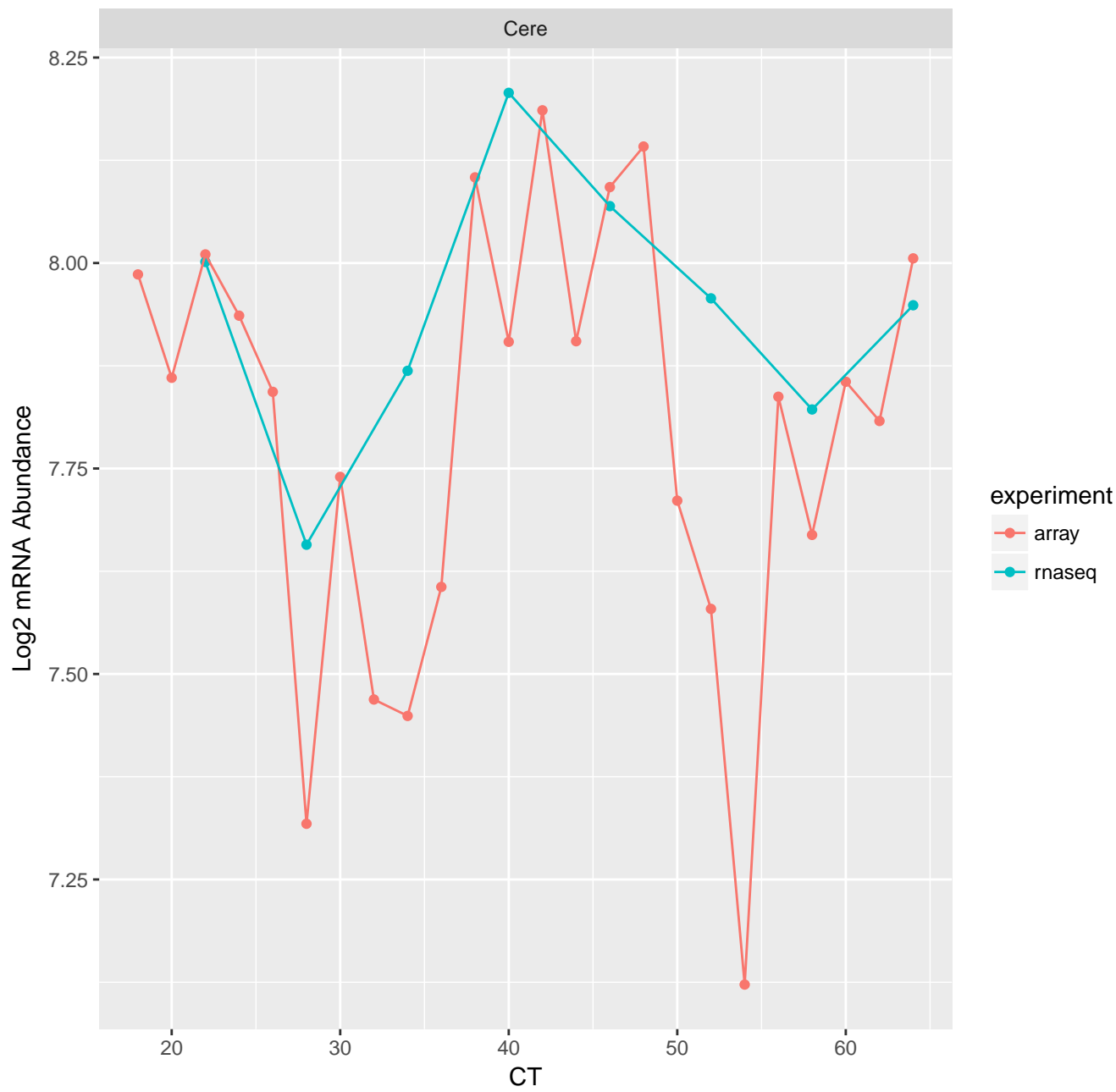
Npas2



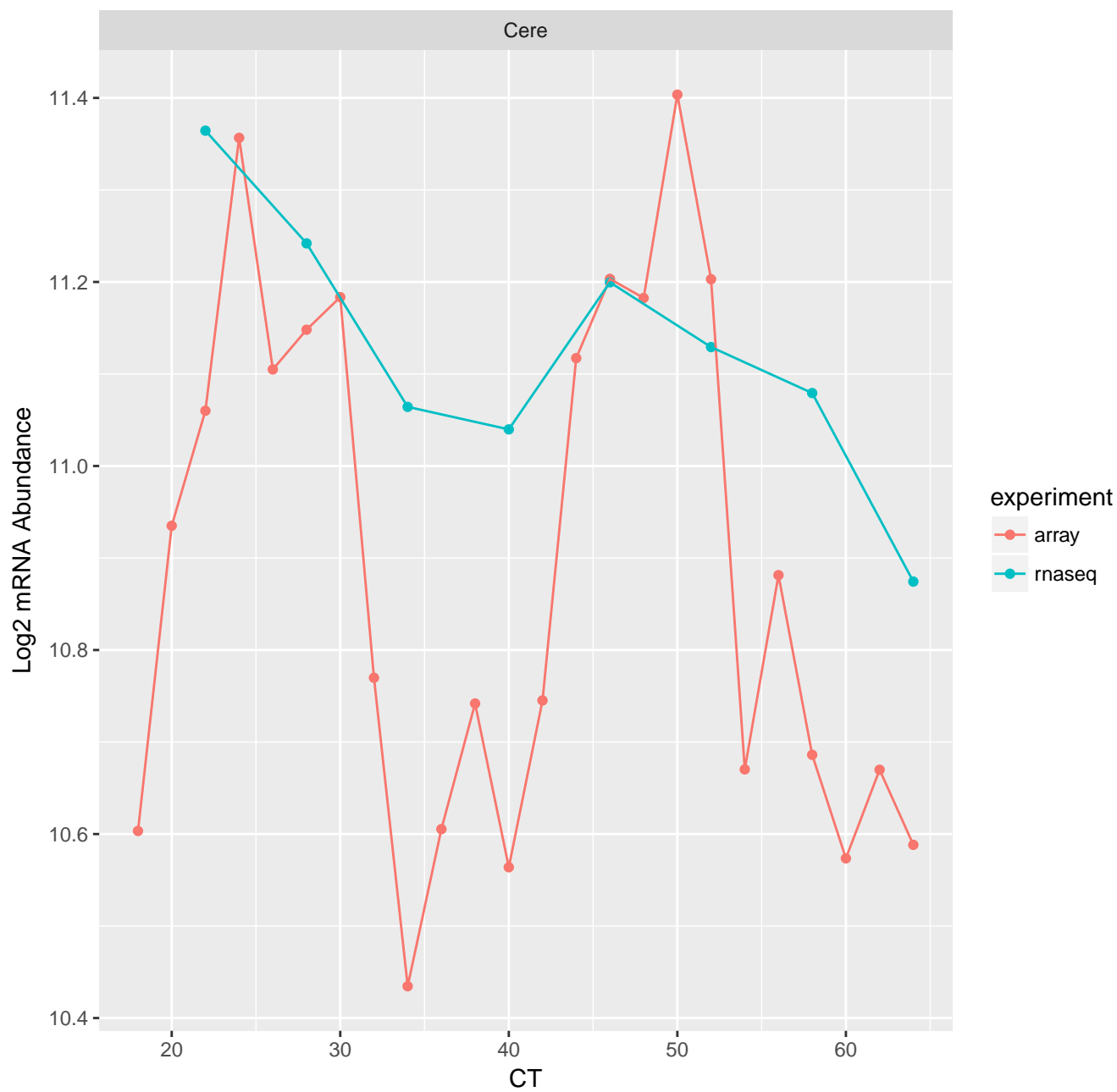
Lman2l



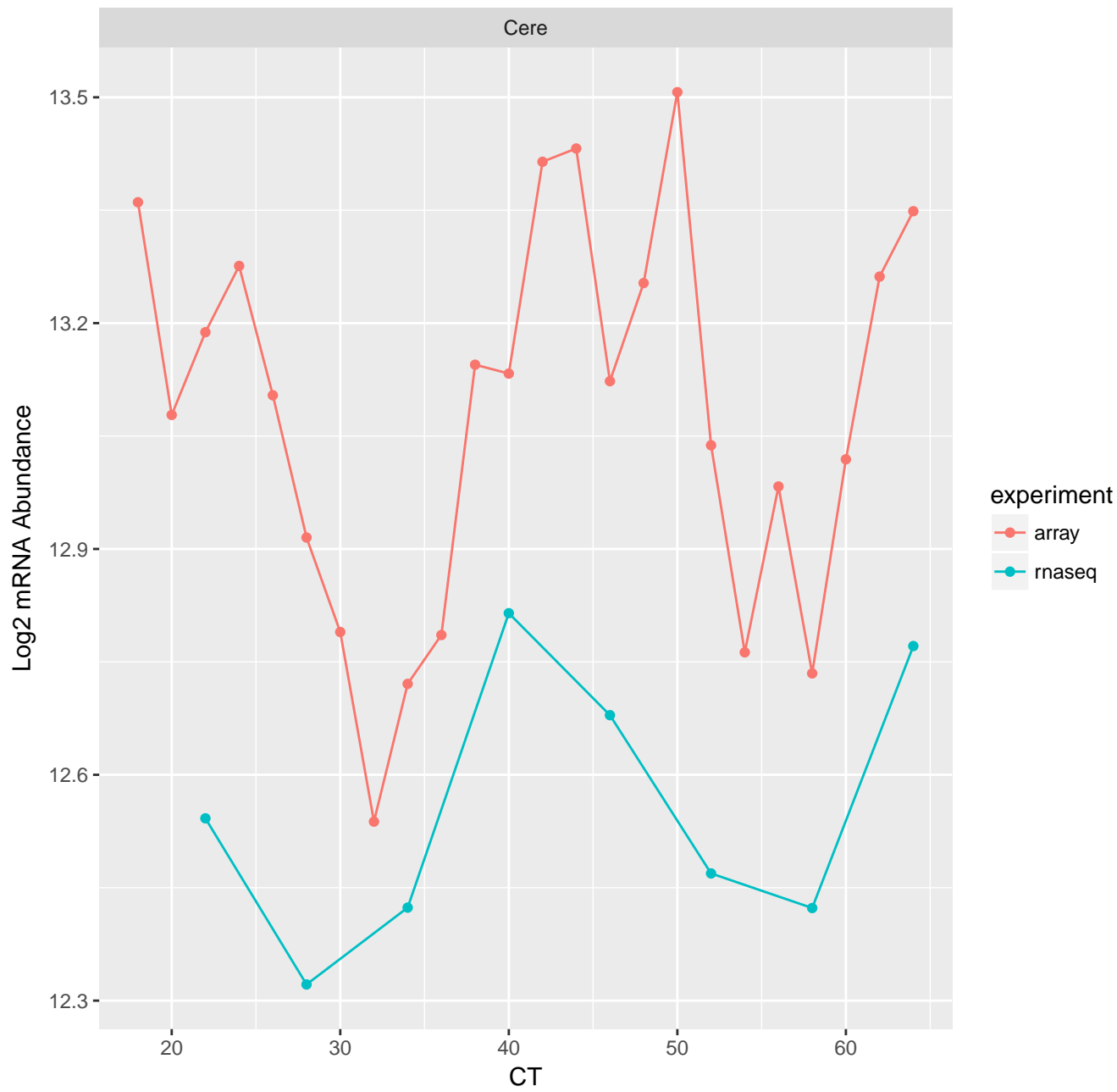
Fam227a



Clock



Calr



Per1

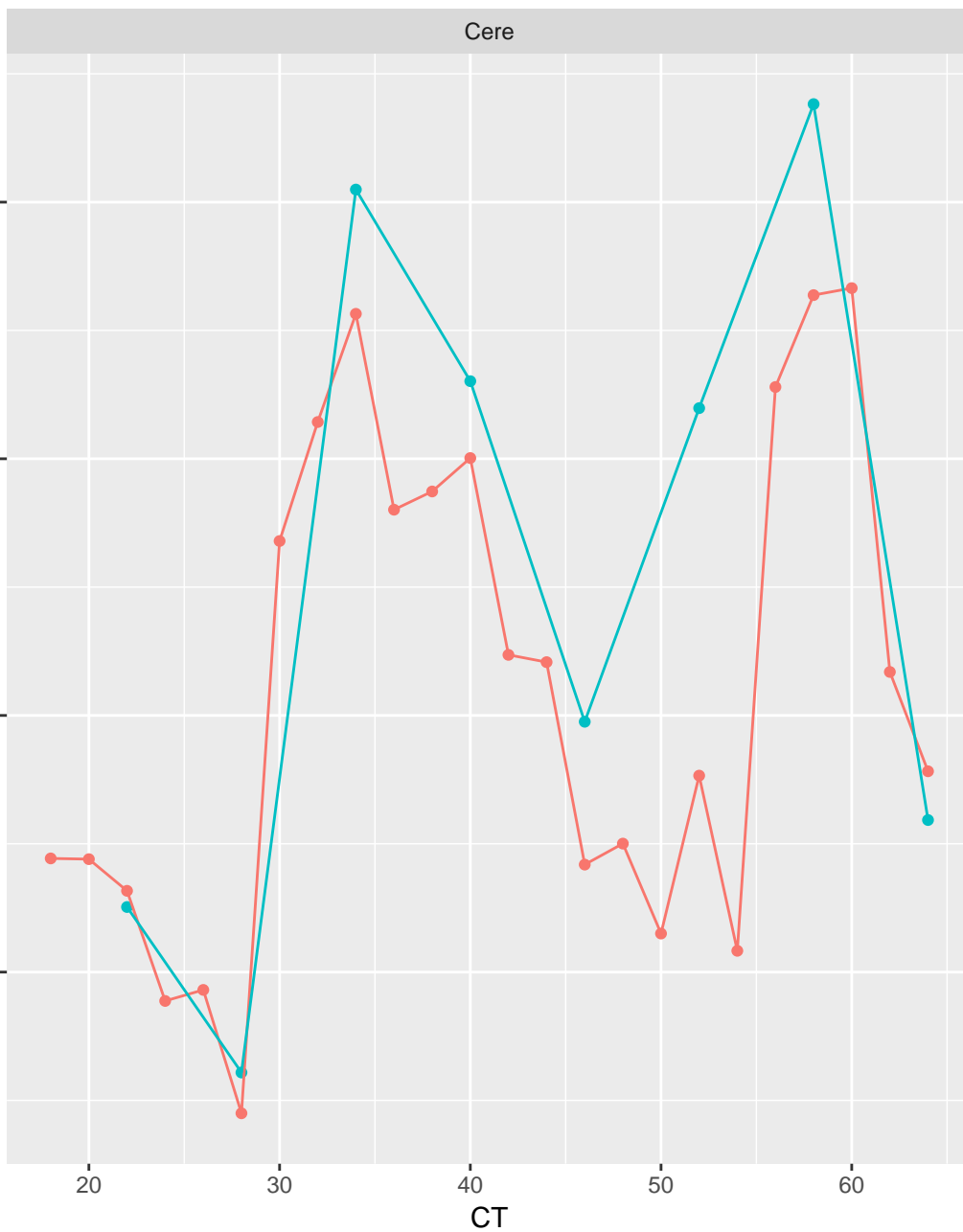
Cere

Log2 mRNA Abundance

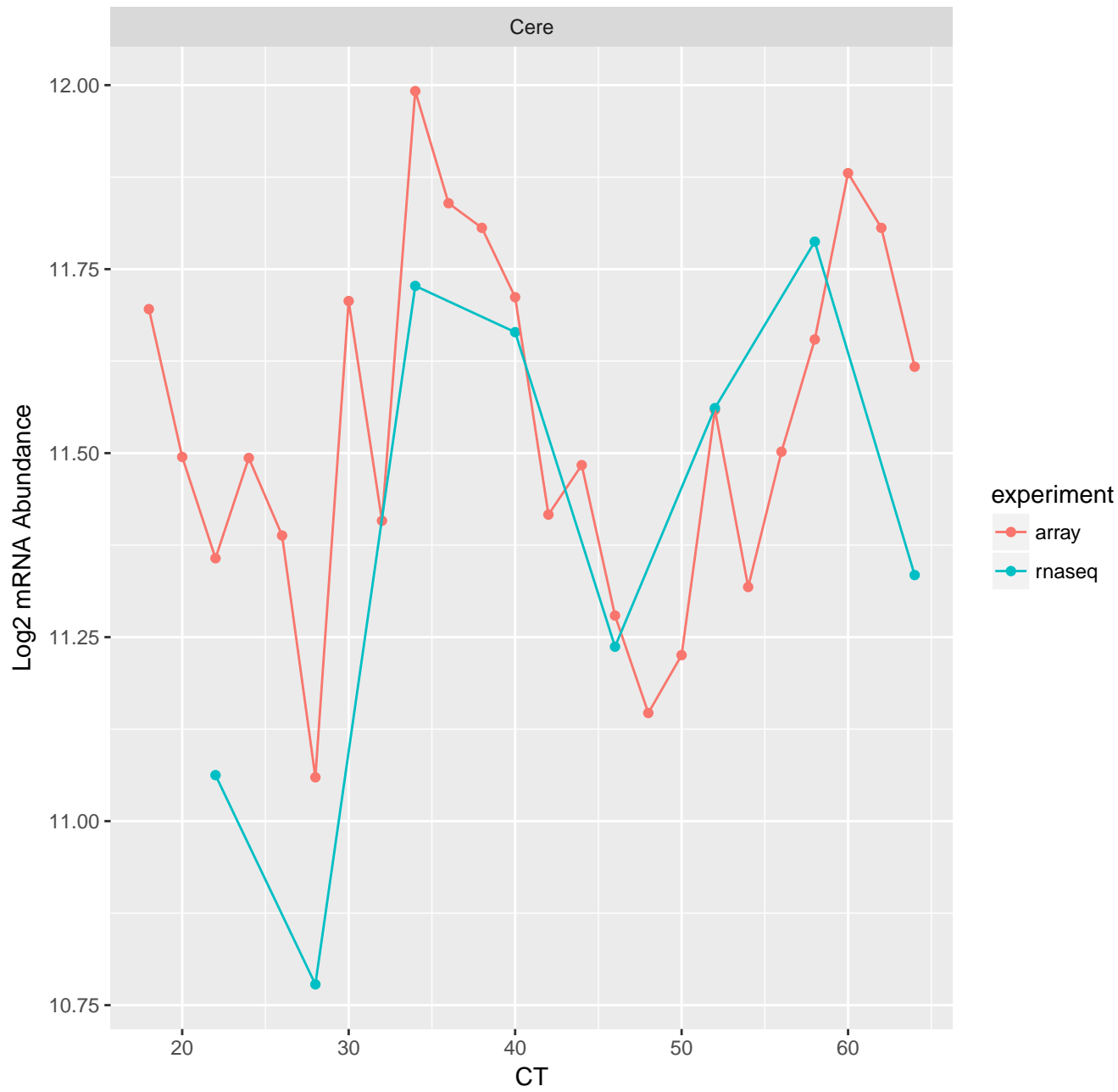
experiment

array
rnaseq

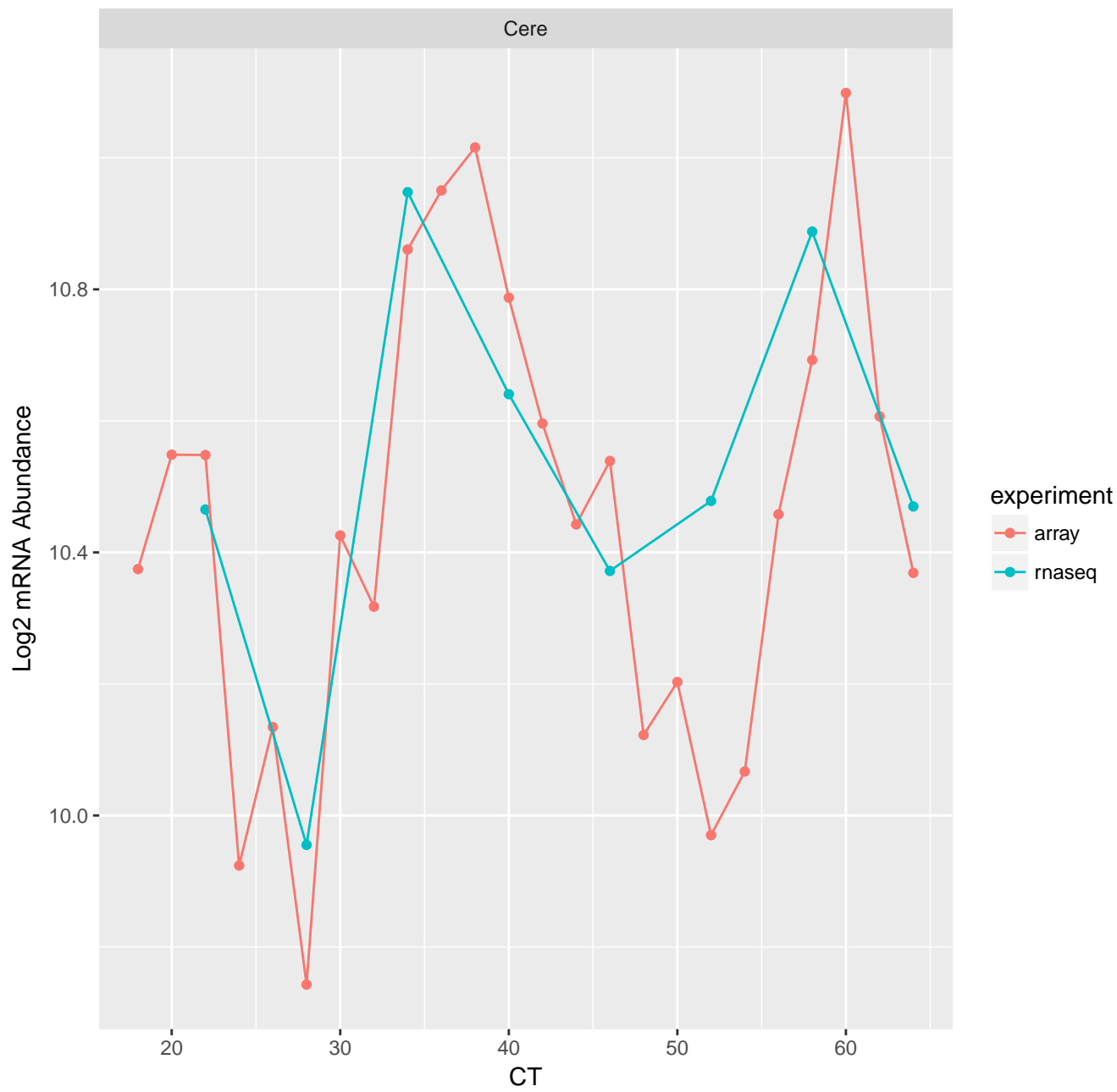
CT



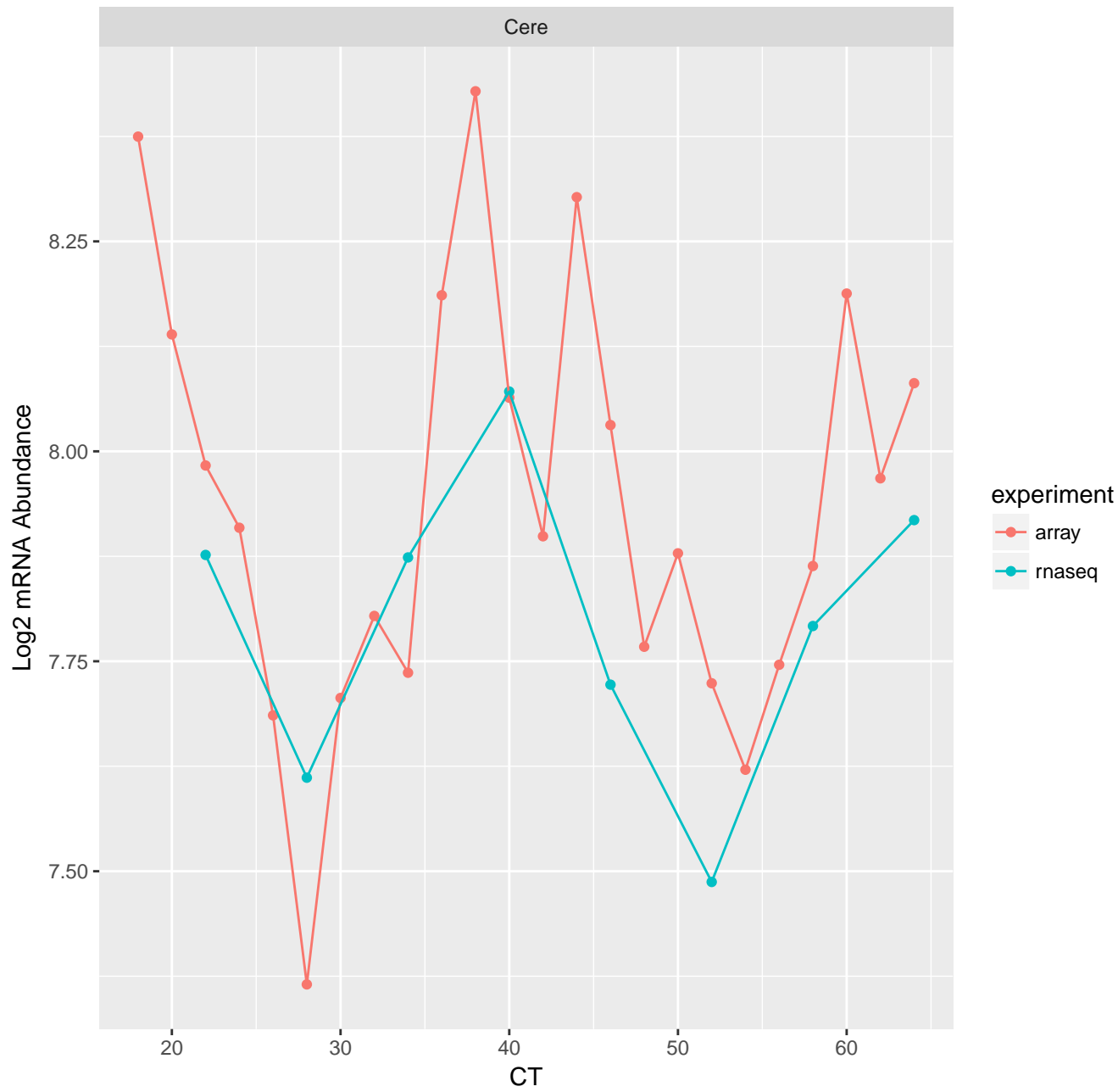
Mt1



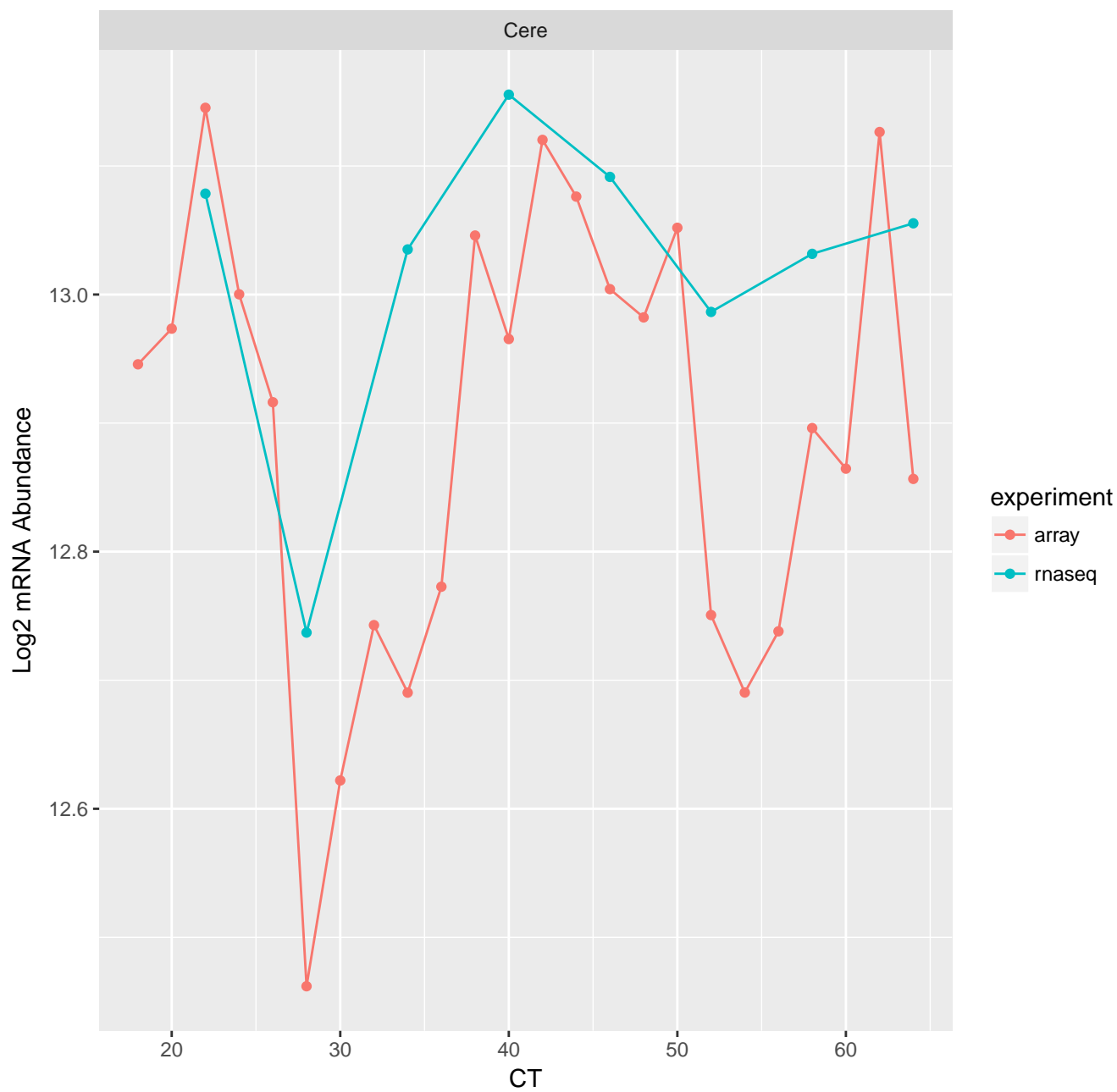
Tsc22d3



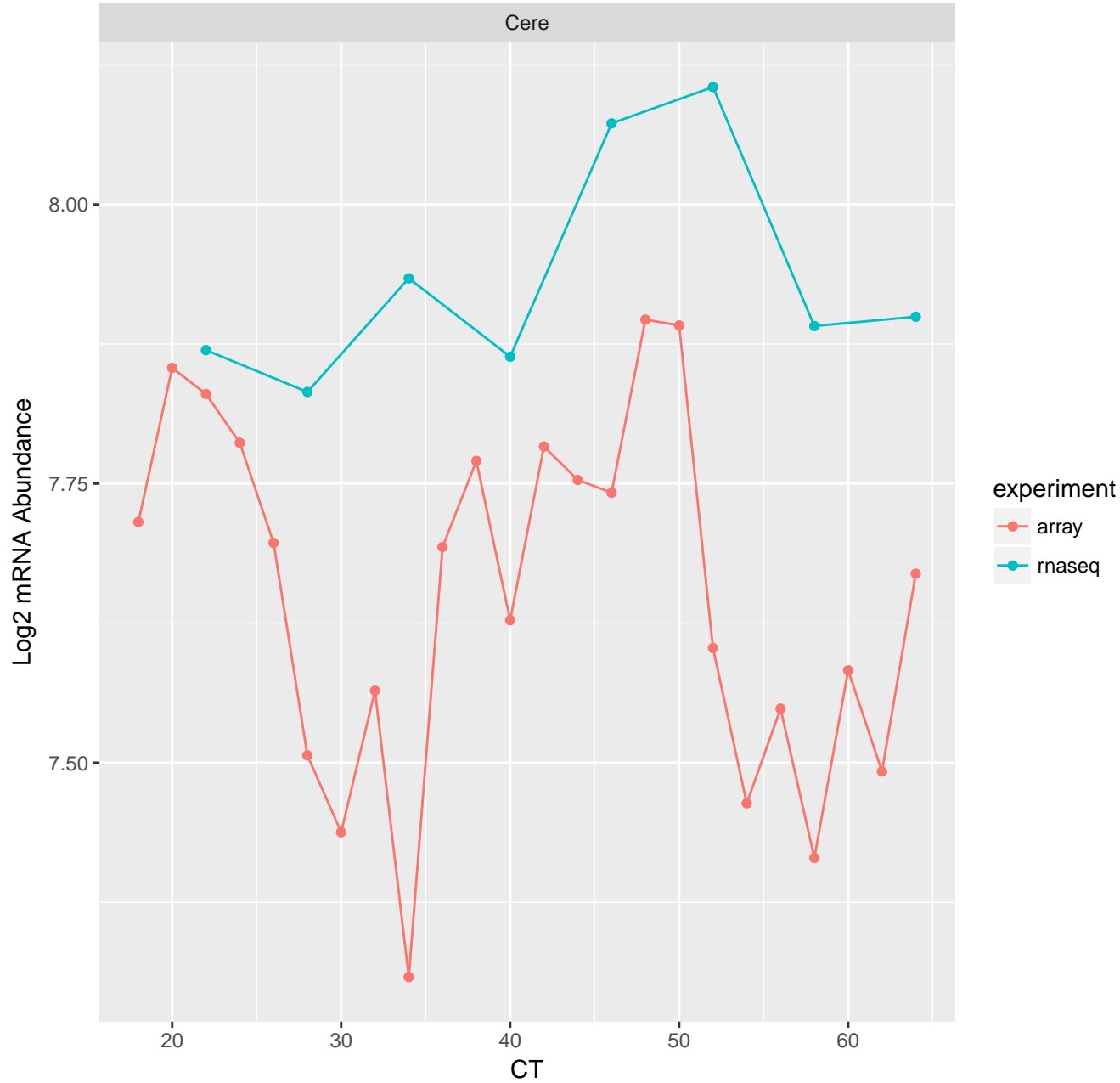
Wee1



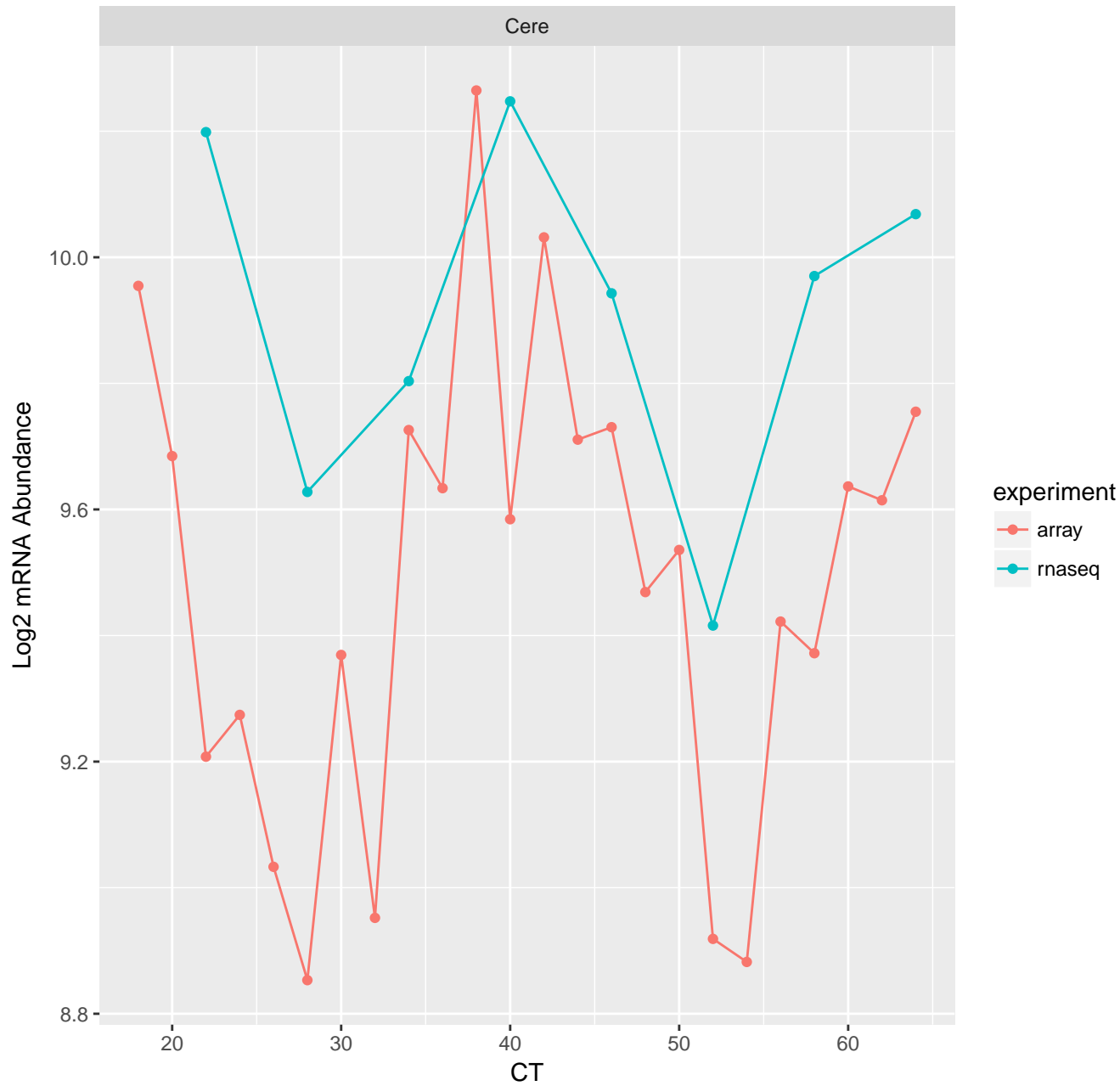
Setd7



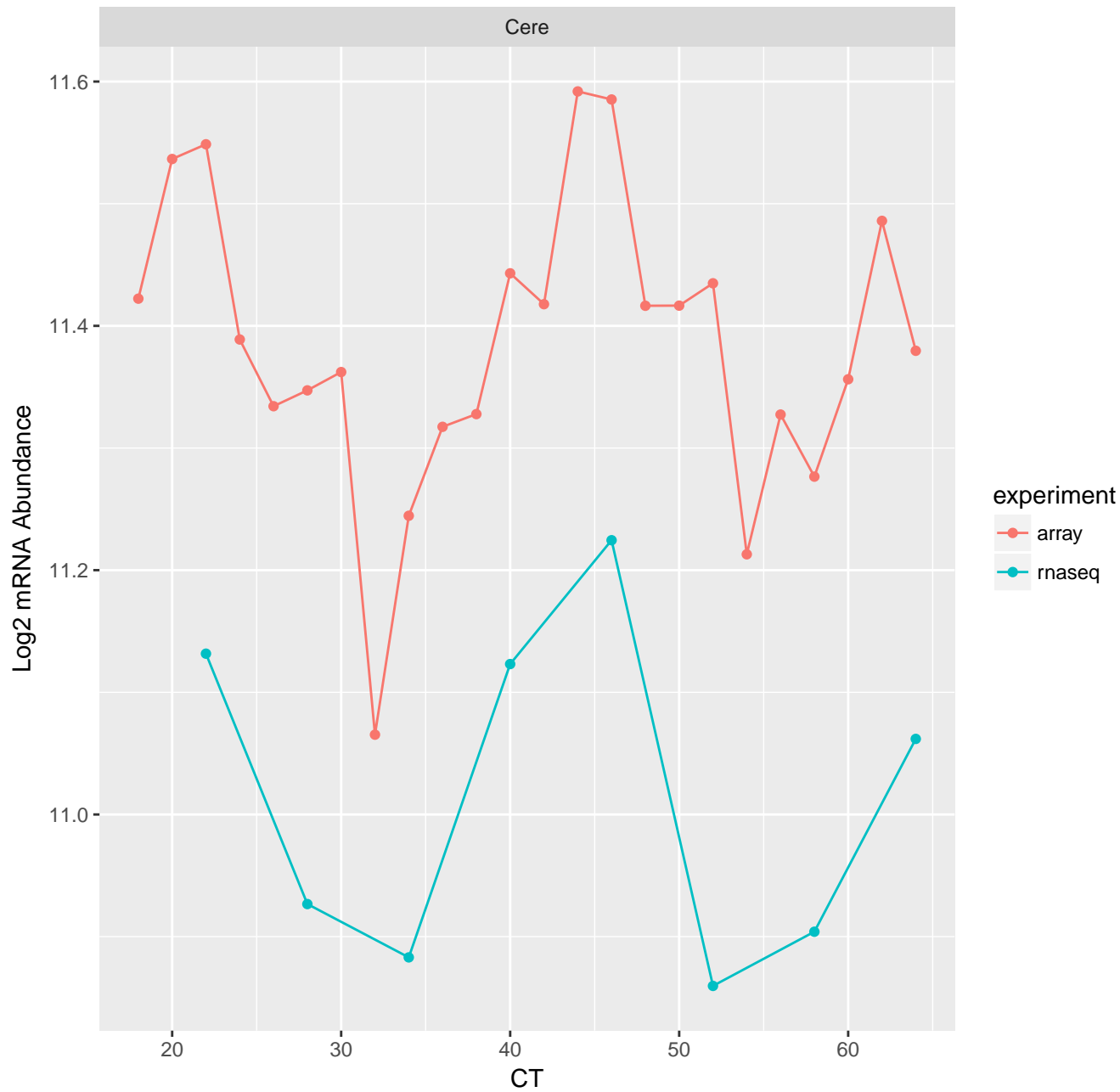
Napepld



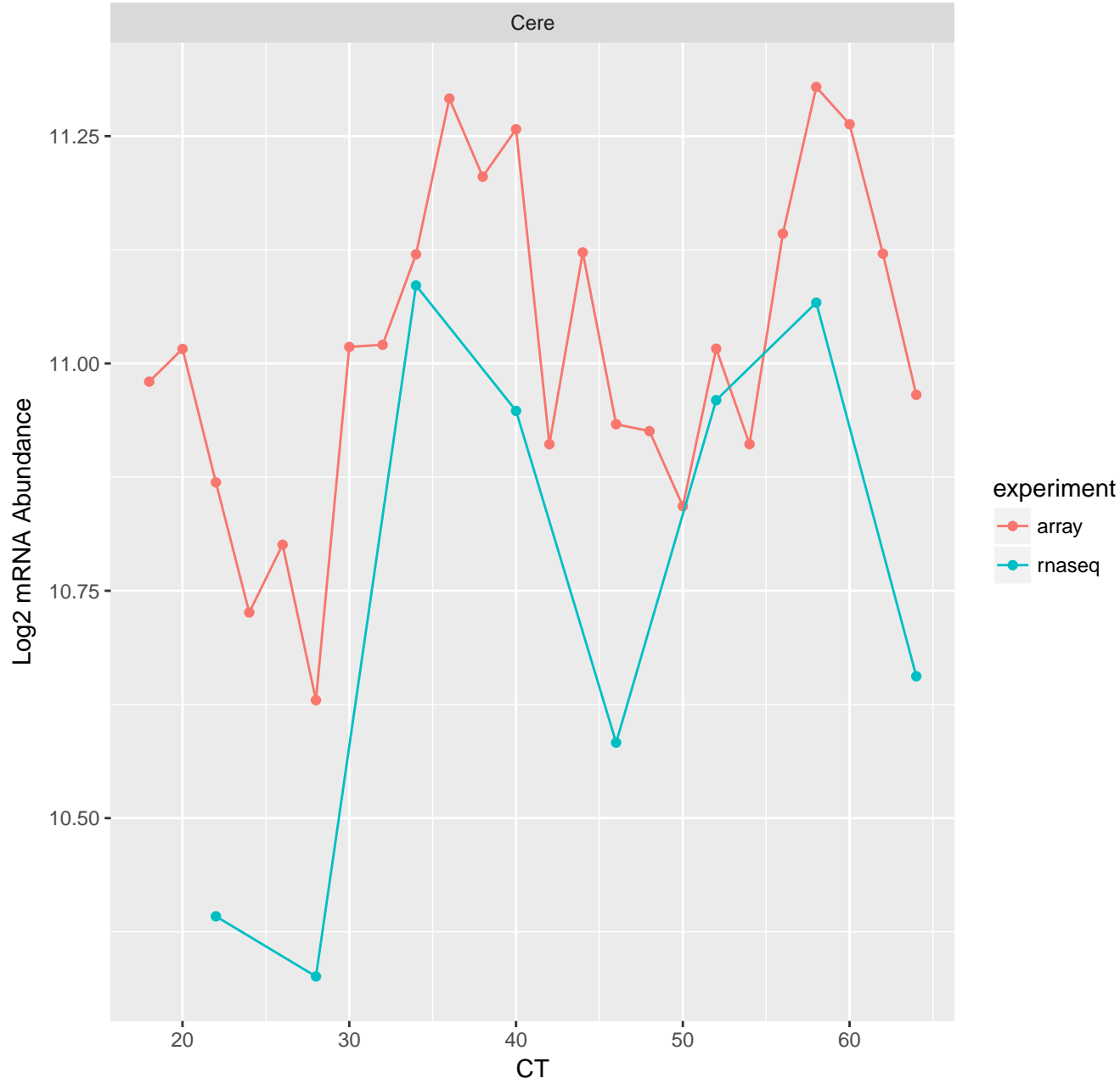
Vps13a



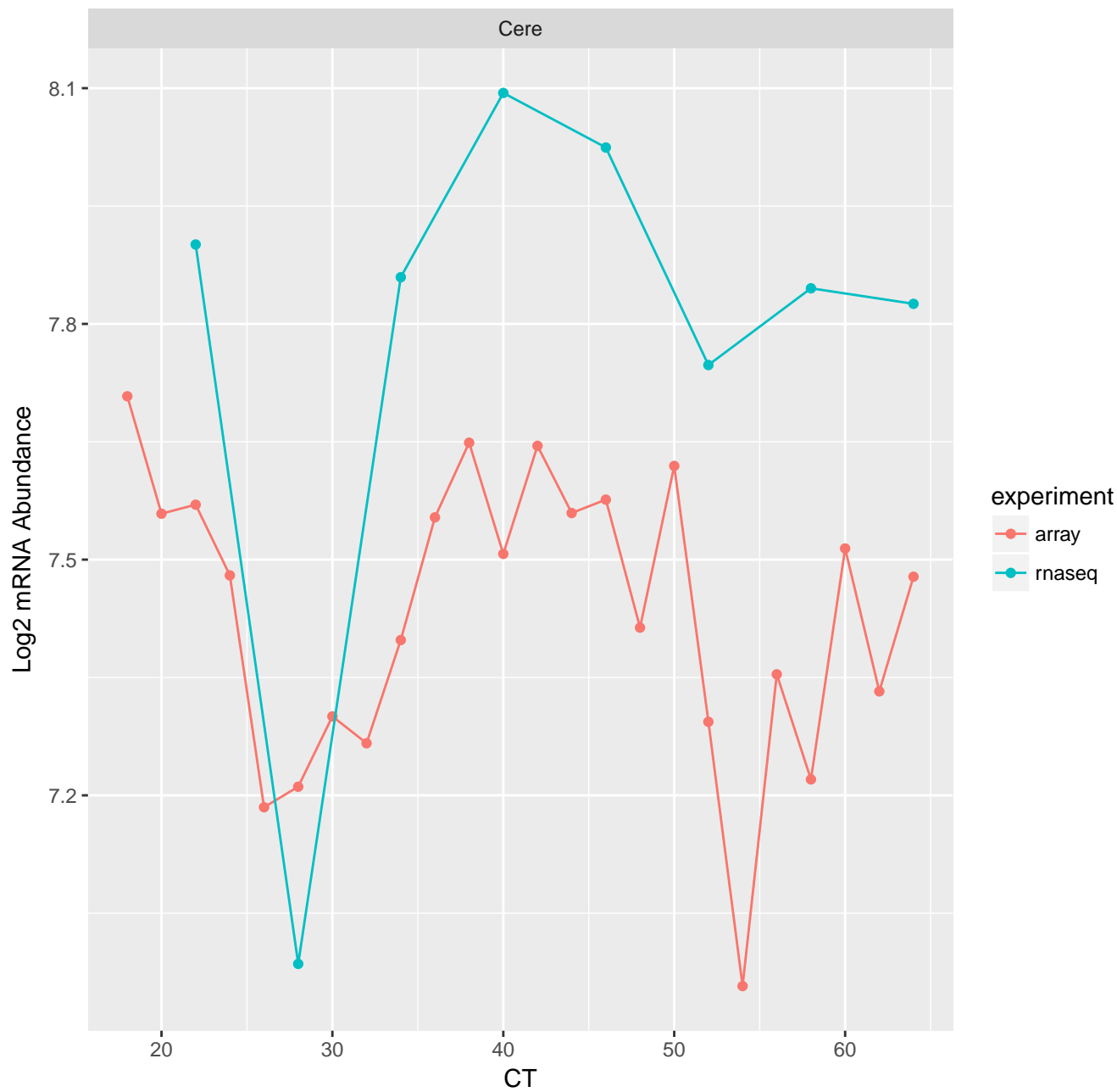
Htra1



Usp2



Folh1



Opalin

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

6.0

6.5

7.0

7.5

8.0

20

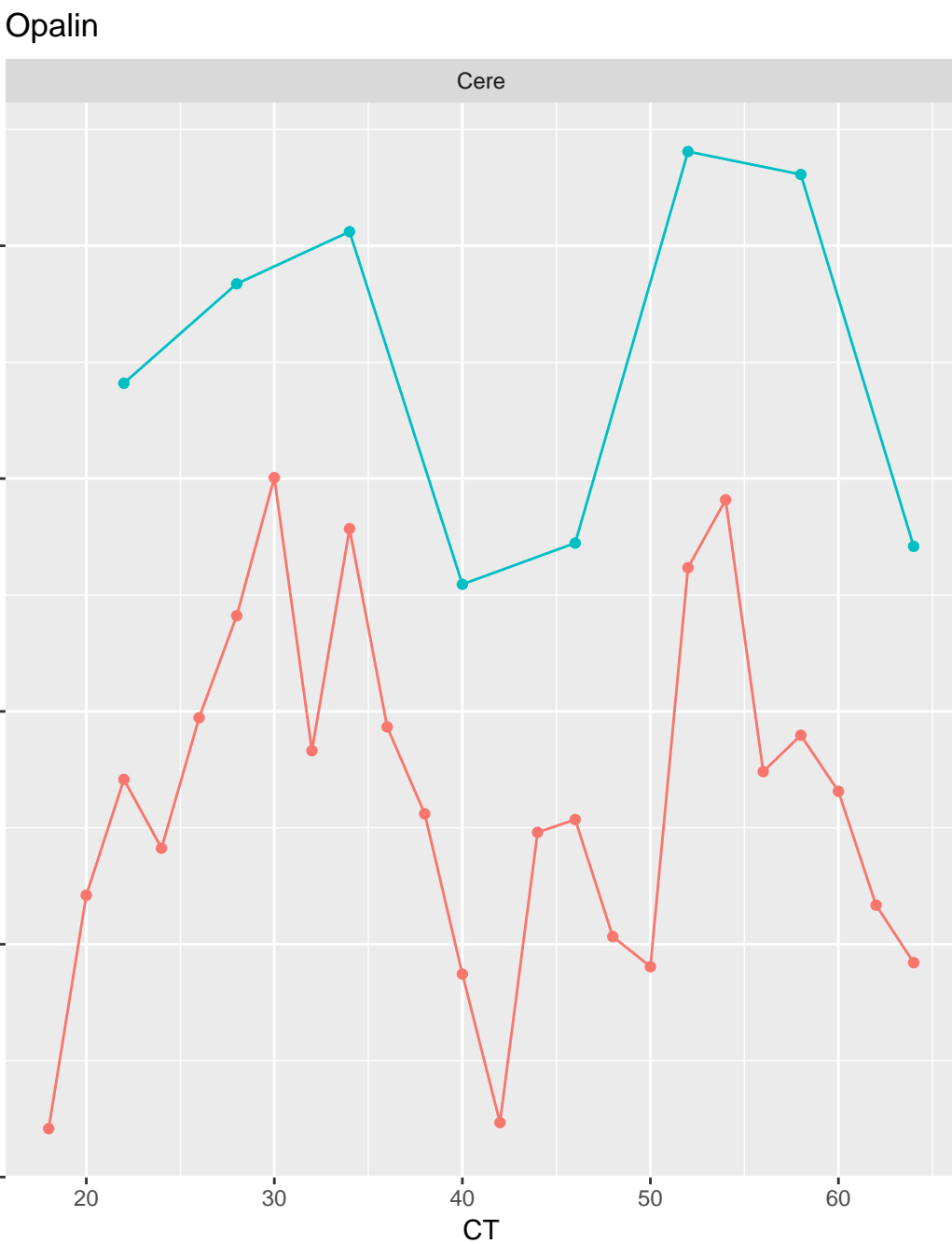
30

CT

40

50

60



Por

Cere

Log2 mRNA Abundance

CT

experiment

array
rnaseq

10.2
10.0
9.8
9.6

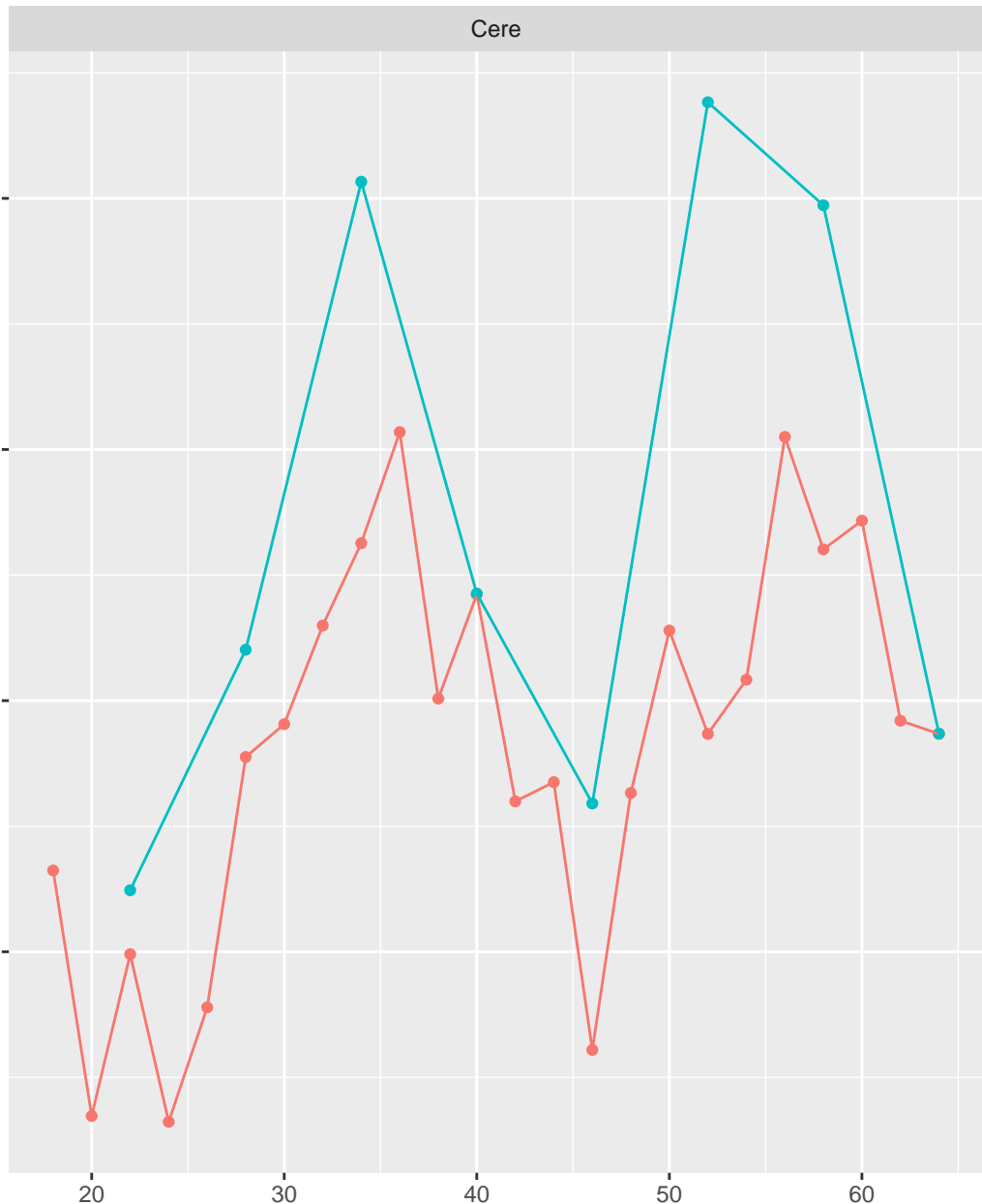
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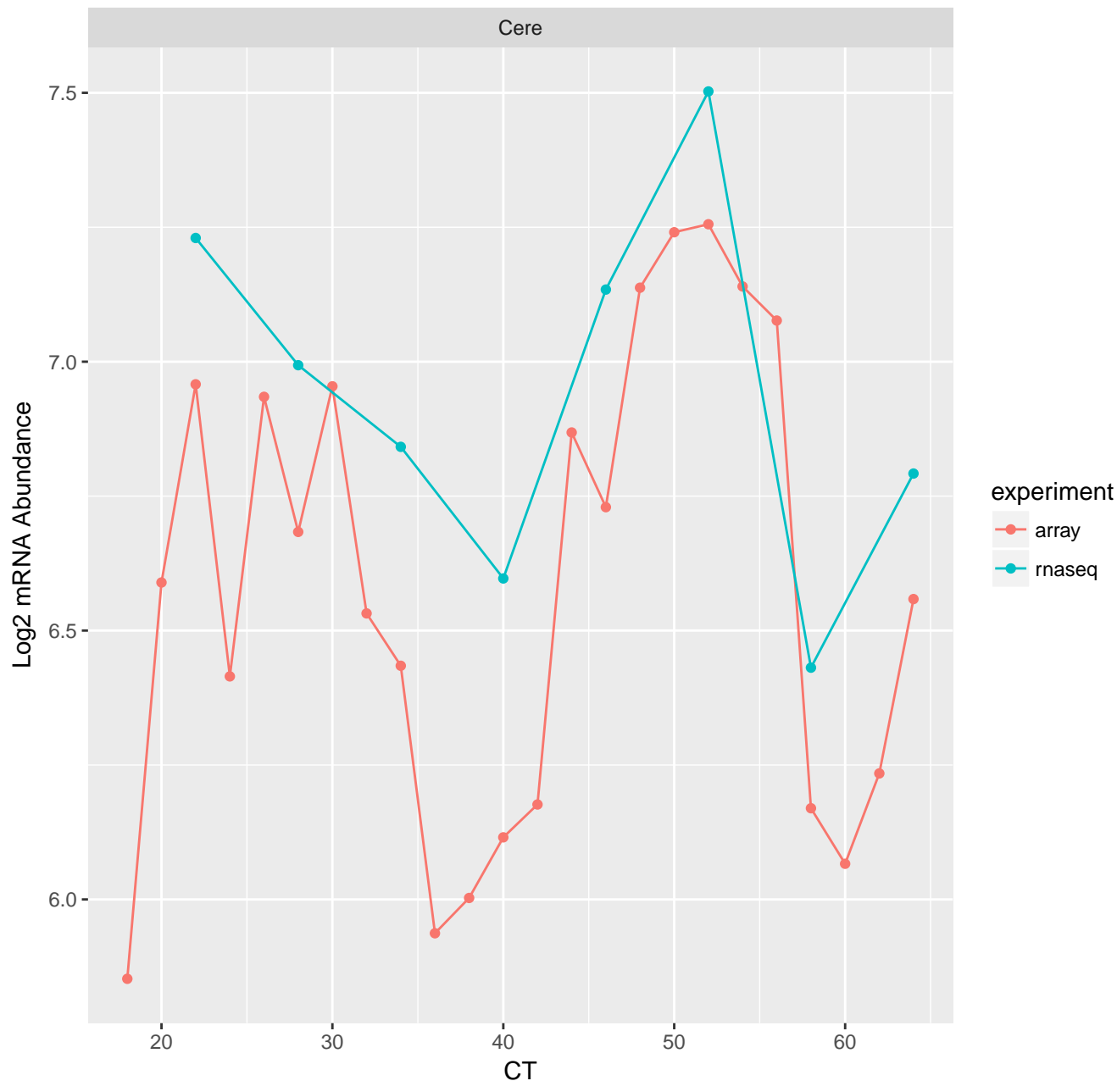
40

50

60



Hmgcs2



Mfsd2a

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

7.2

7.4

7.6

7.8

8.0

CT

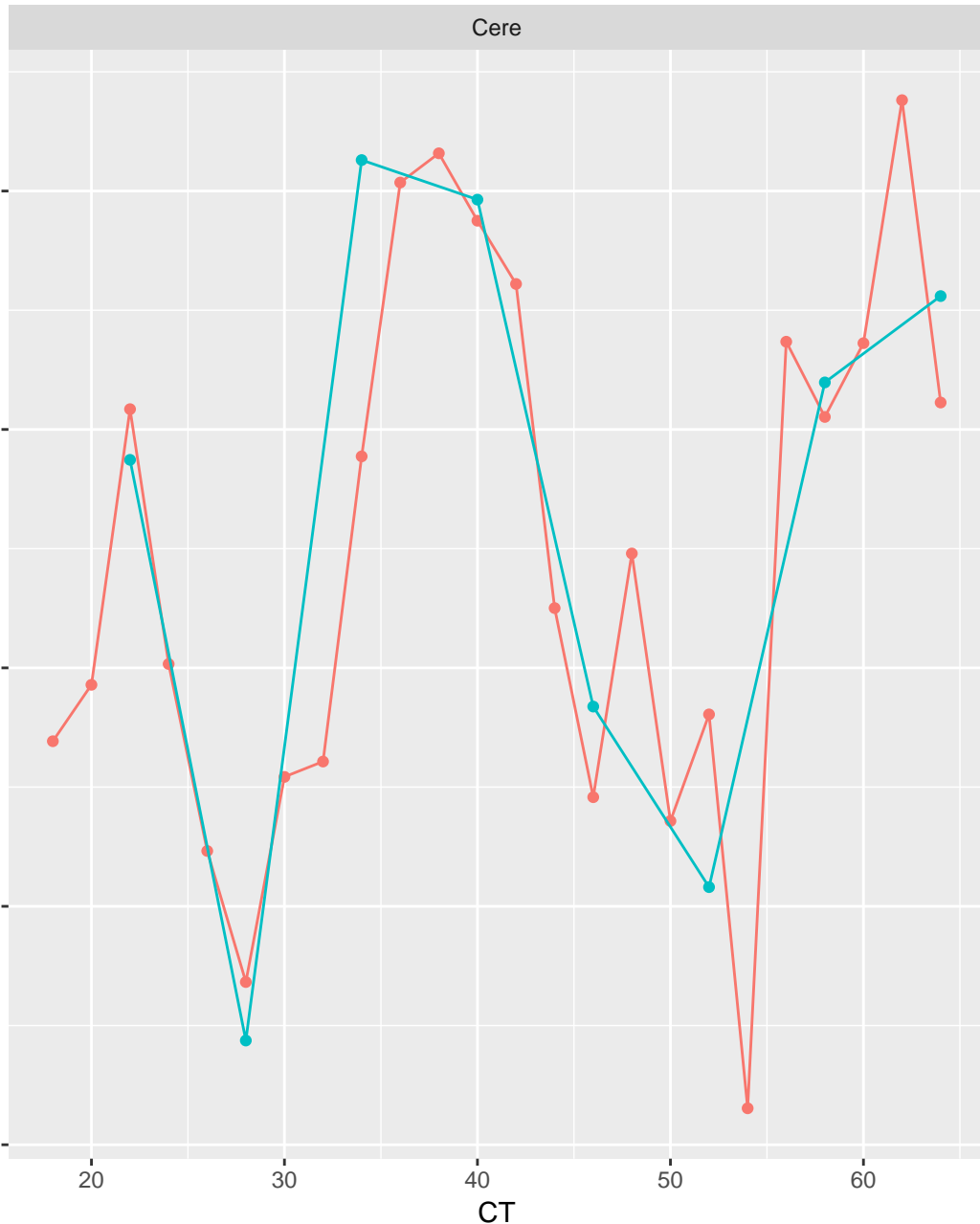
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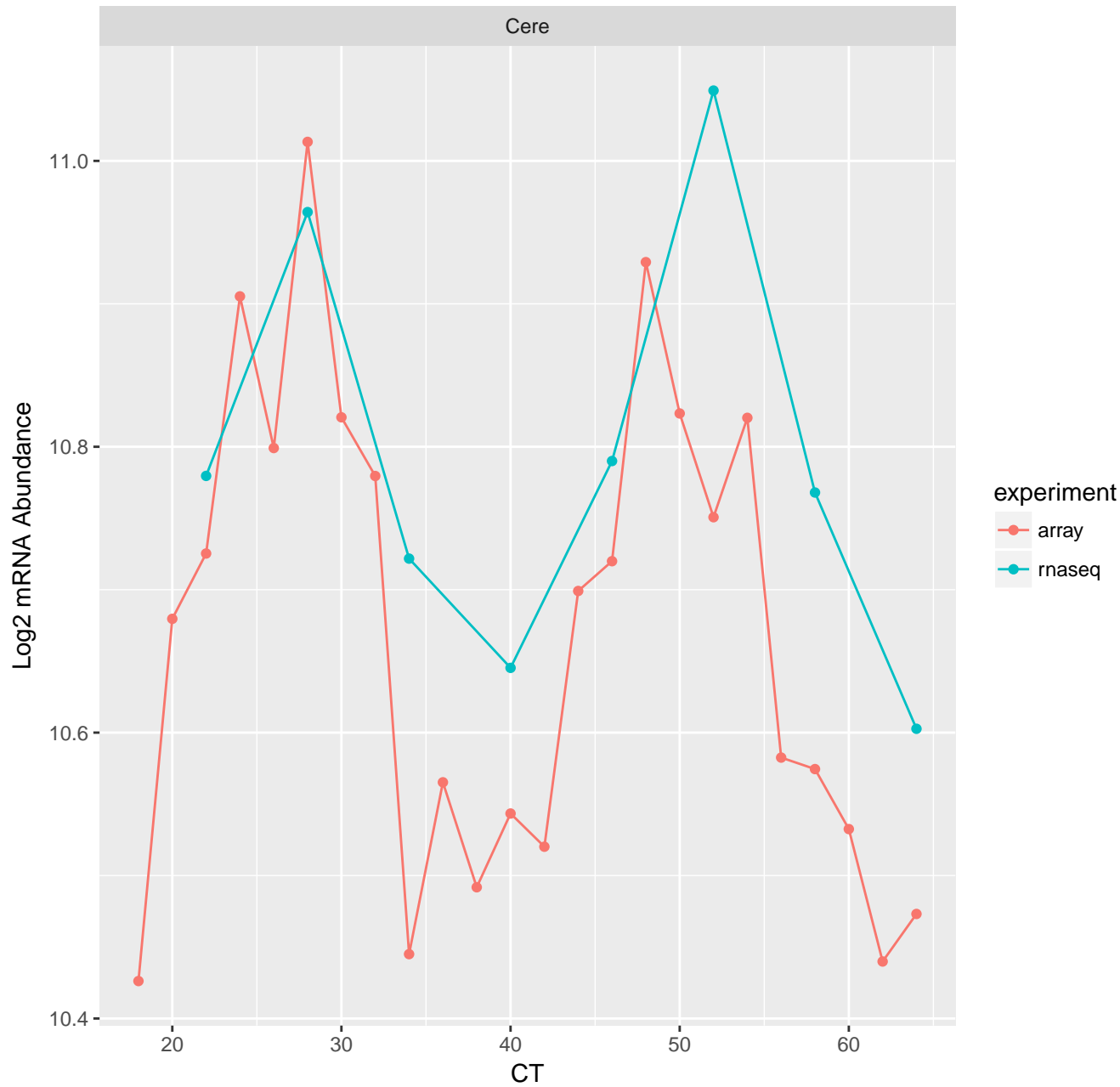
40

50

60



Eps8



Aass

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

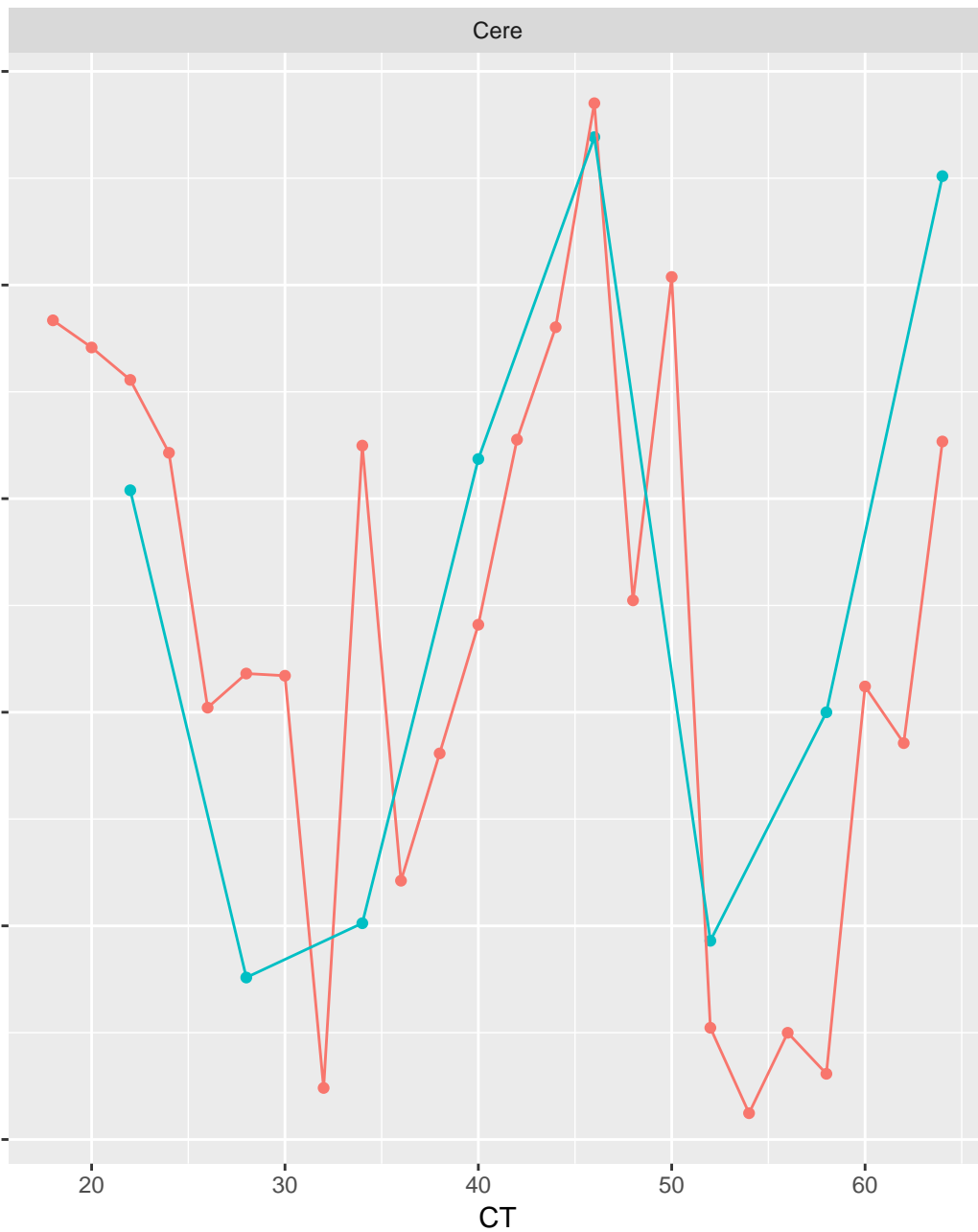
7.6
7.4
7.2
7.0
6.8
6.6

20

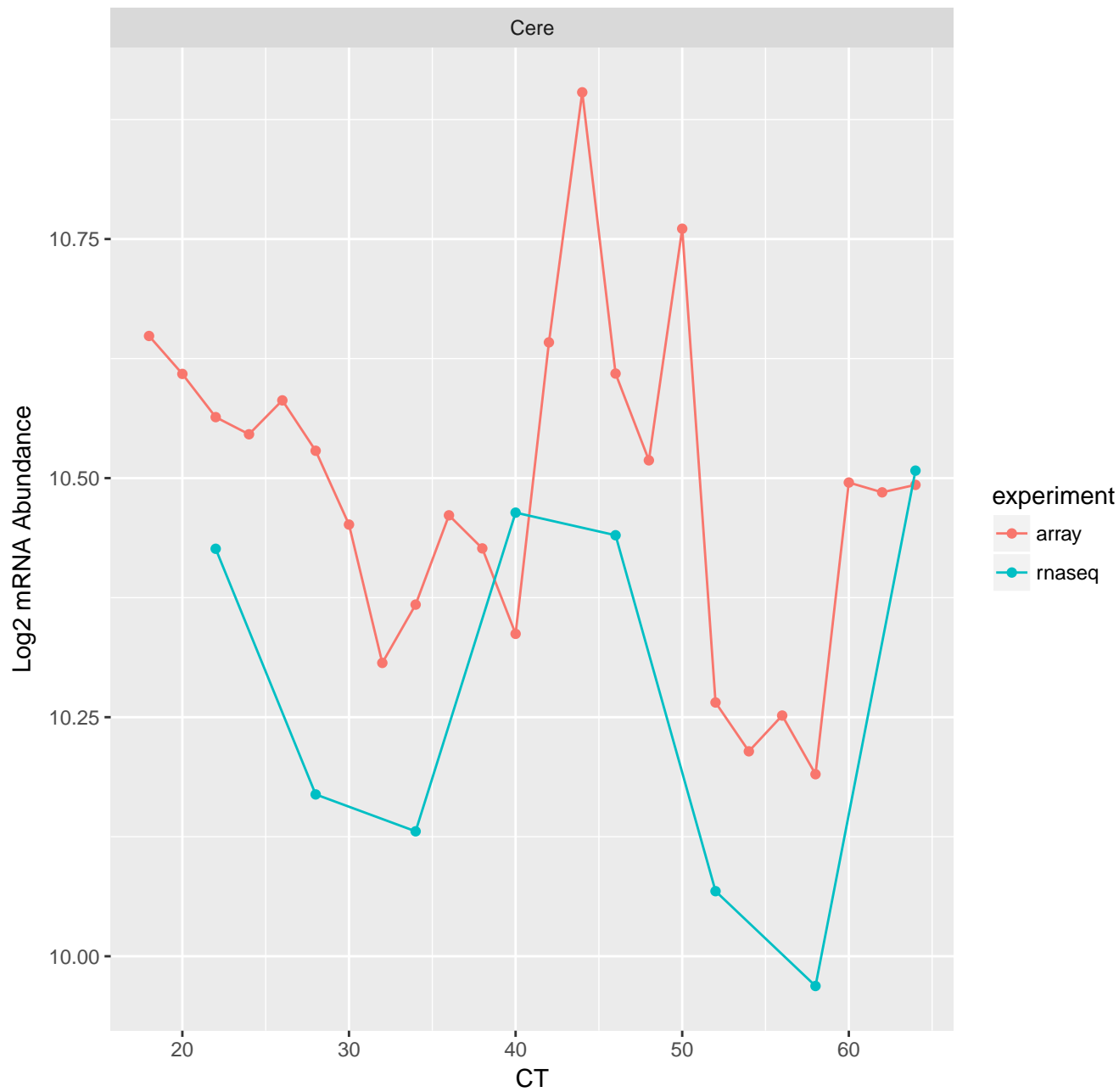
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CT

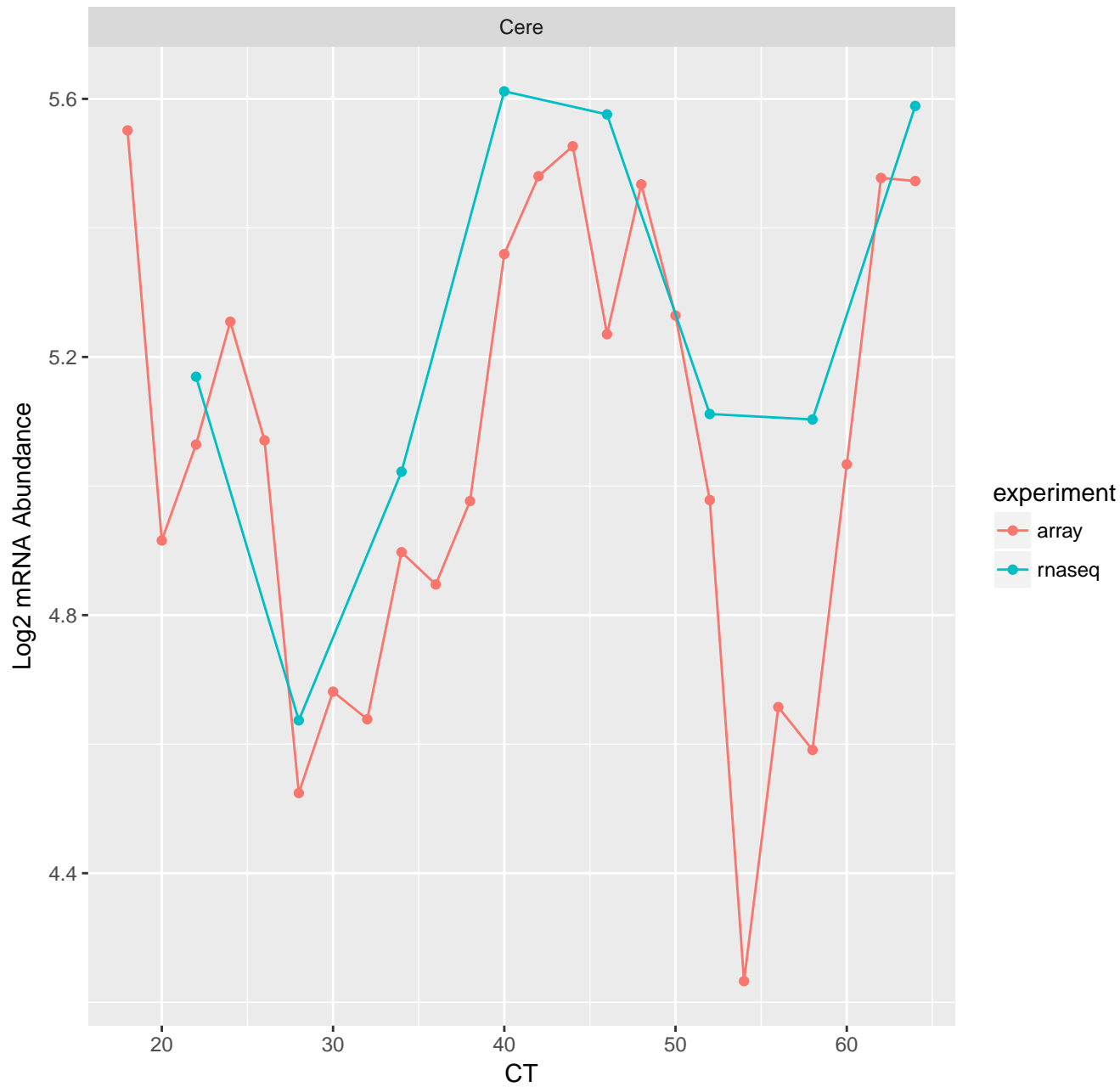
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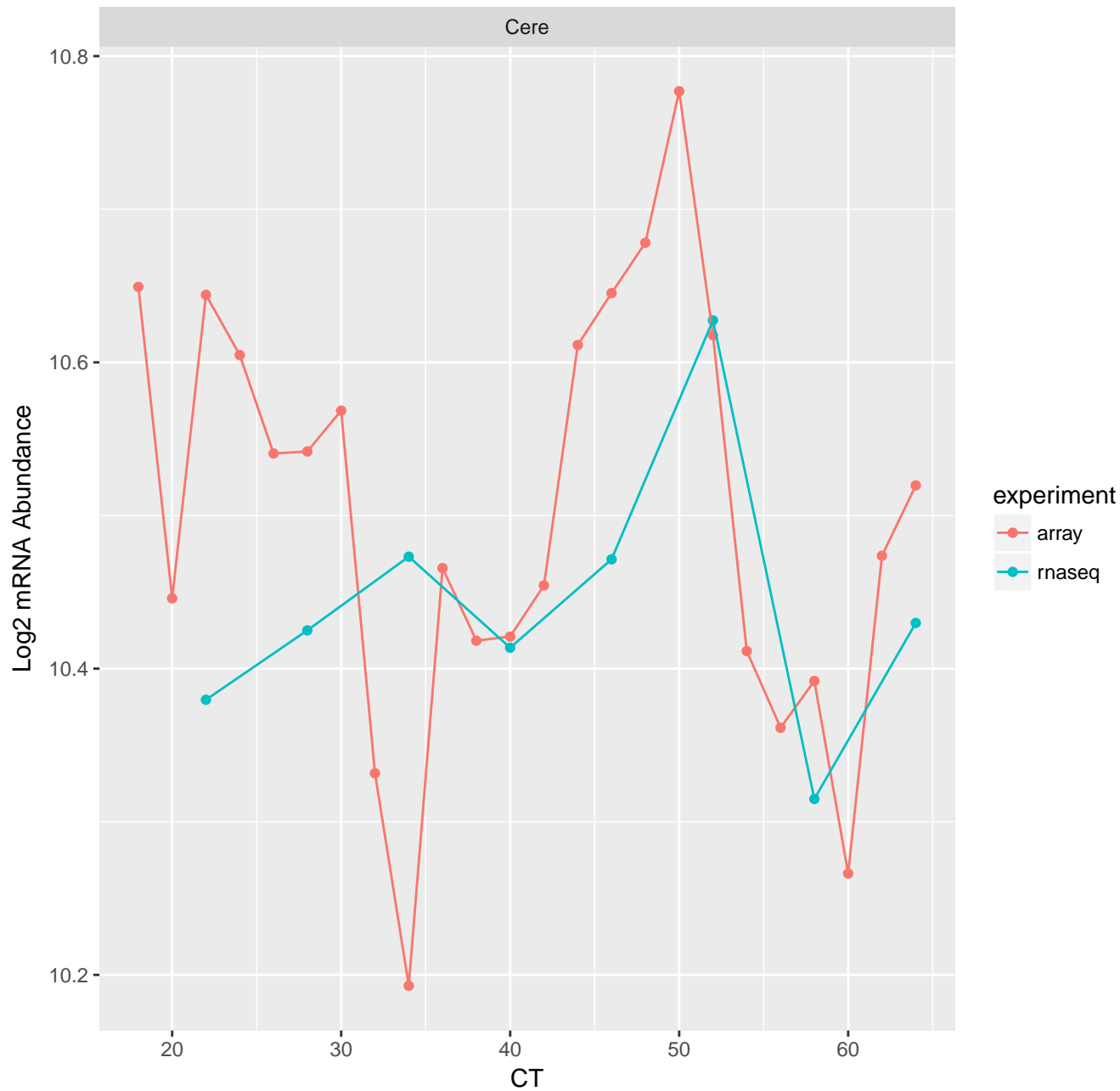
Rbm28



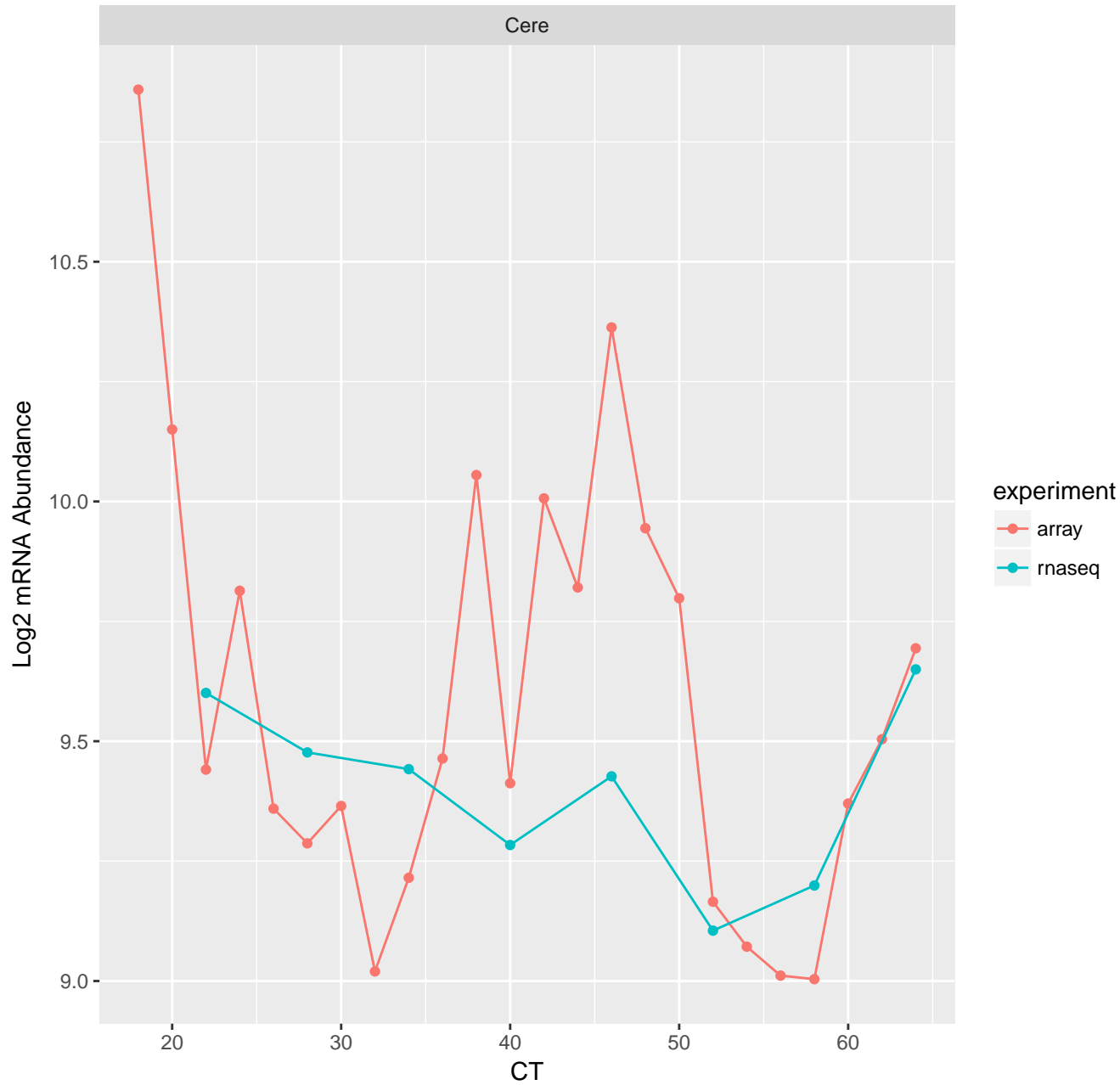
Hspb1



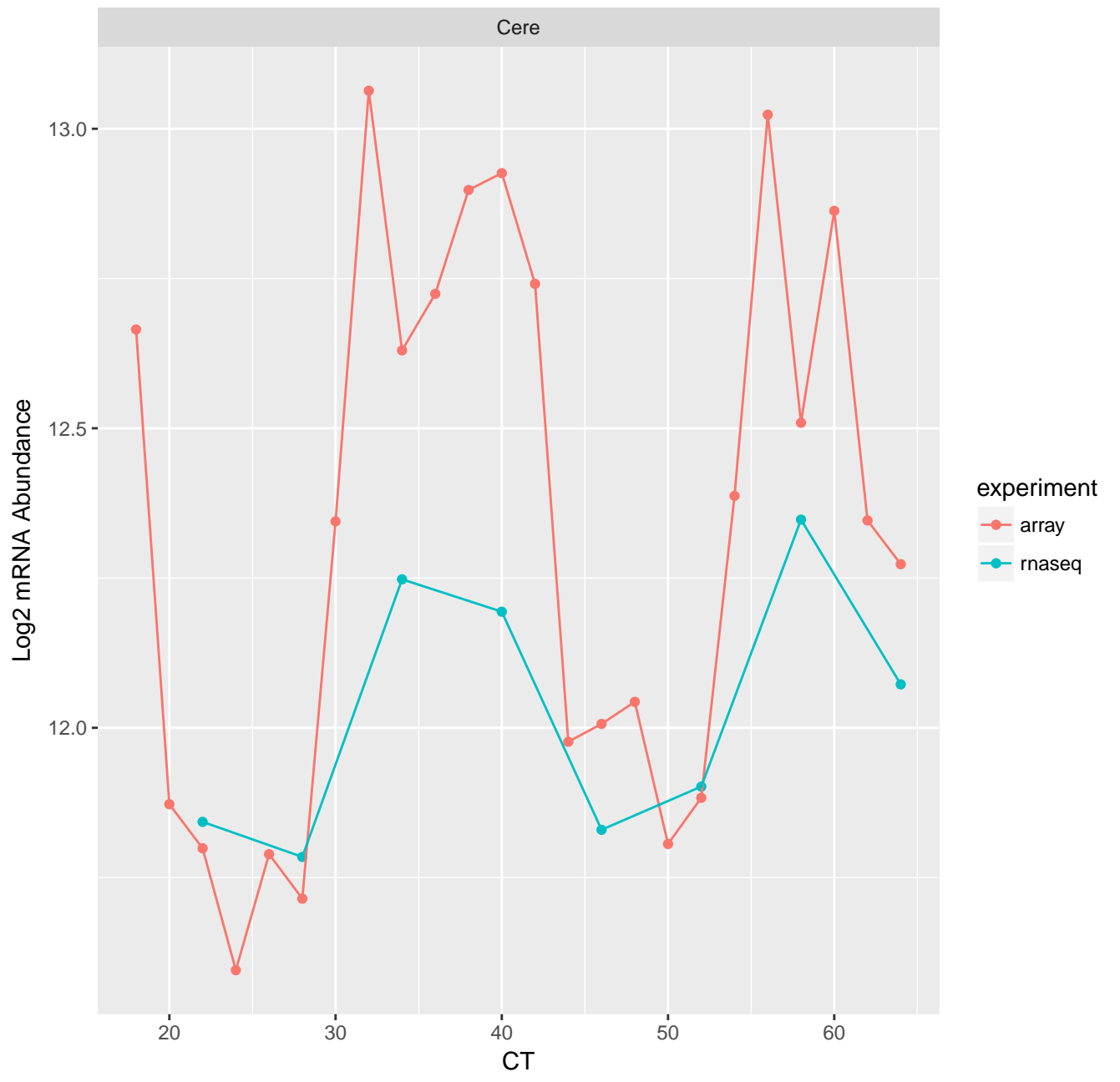
Akt2



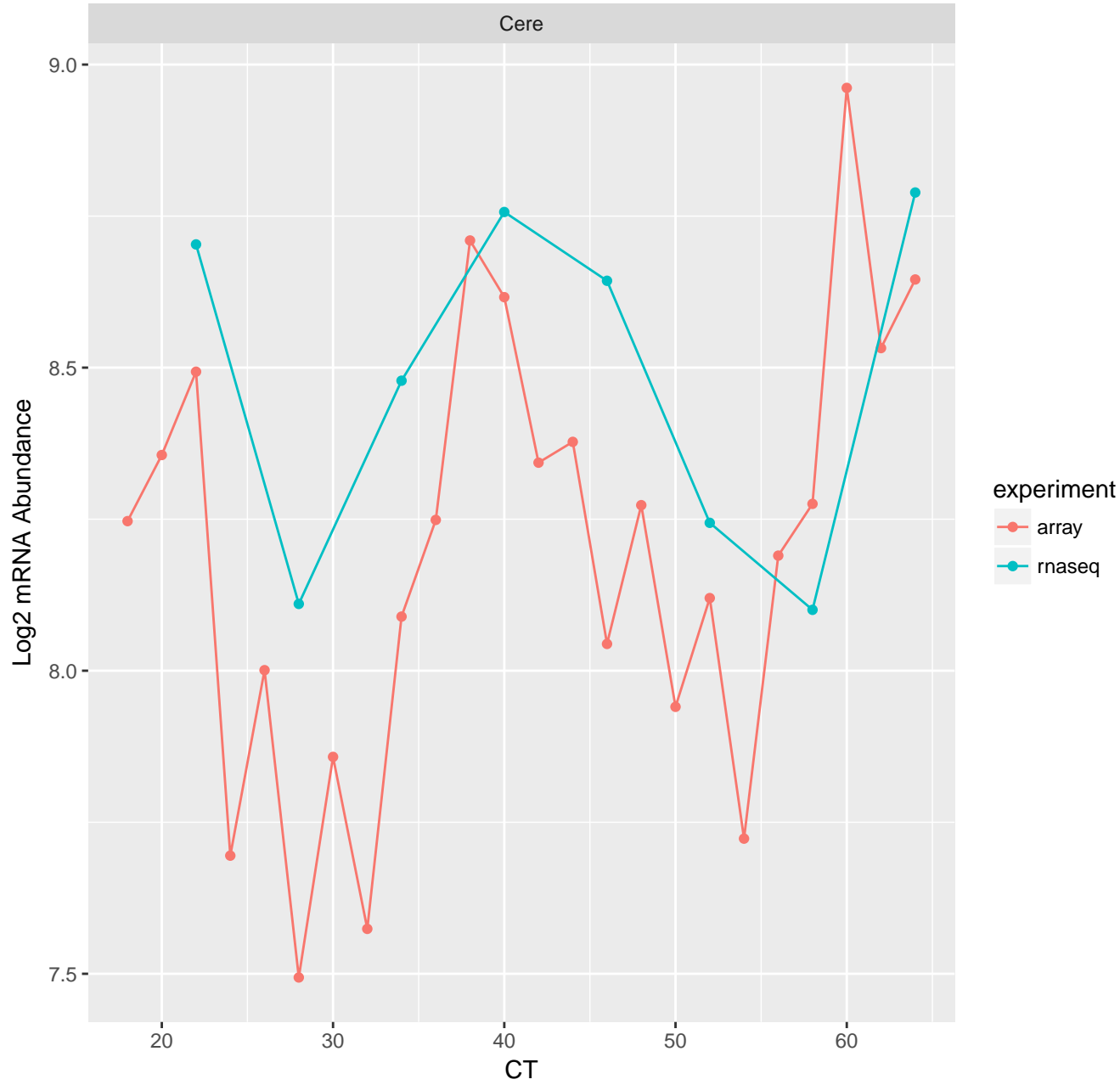
Zfp760



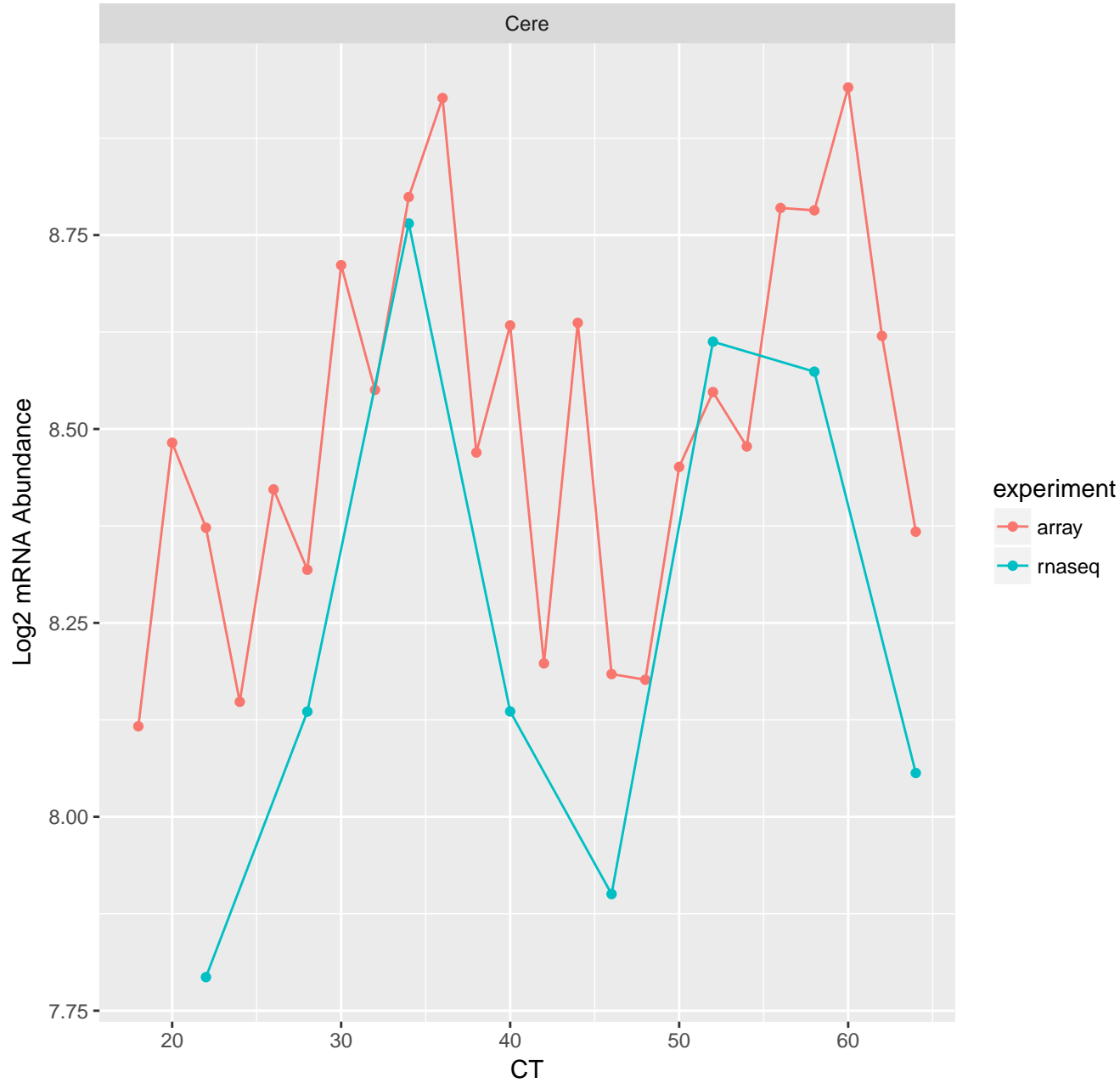
Nr1d2



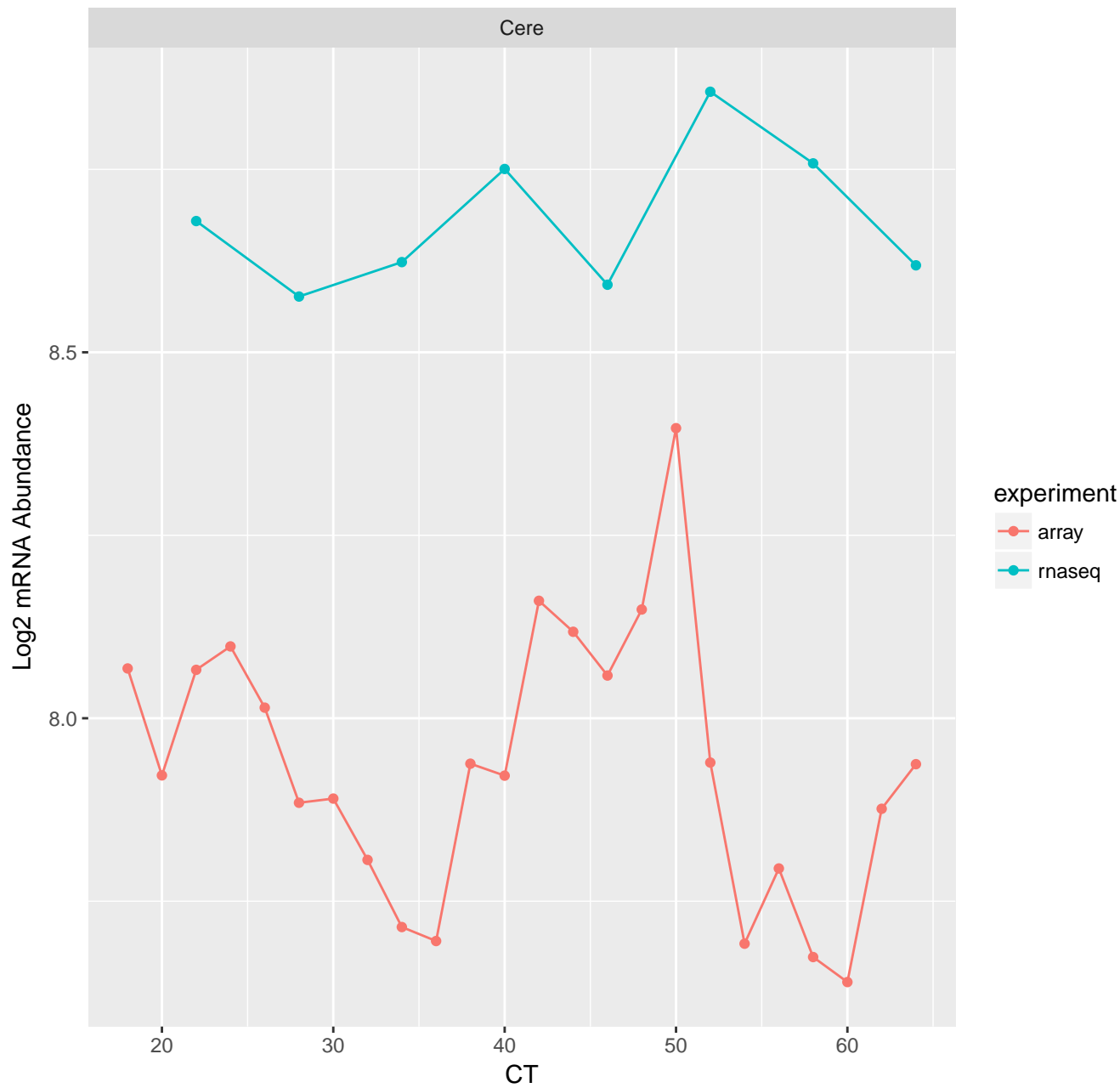
Pnpla7



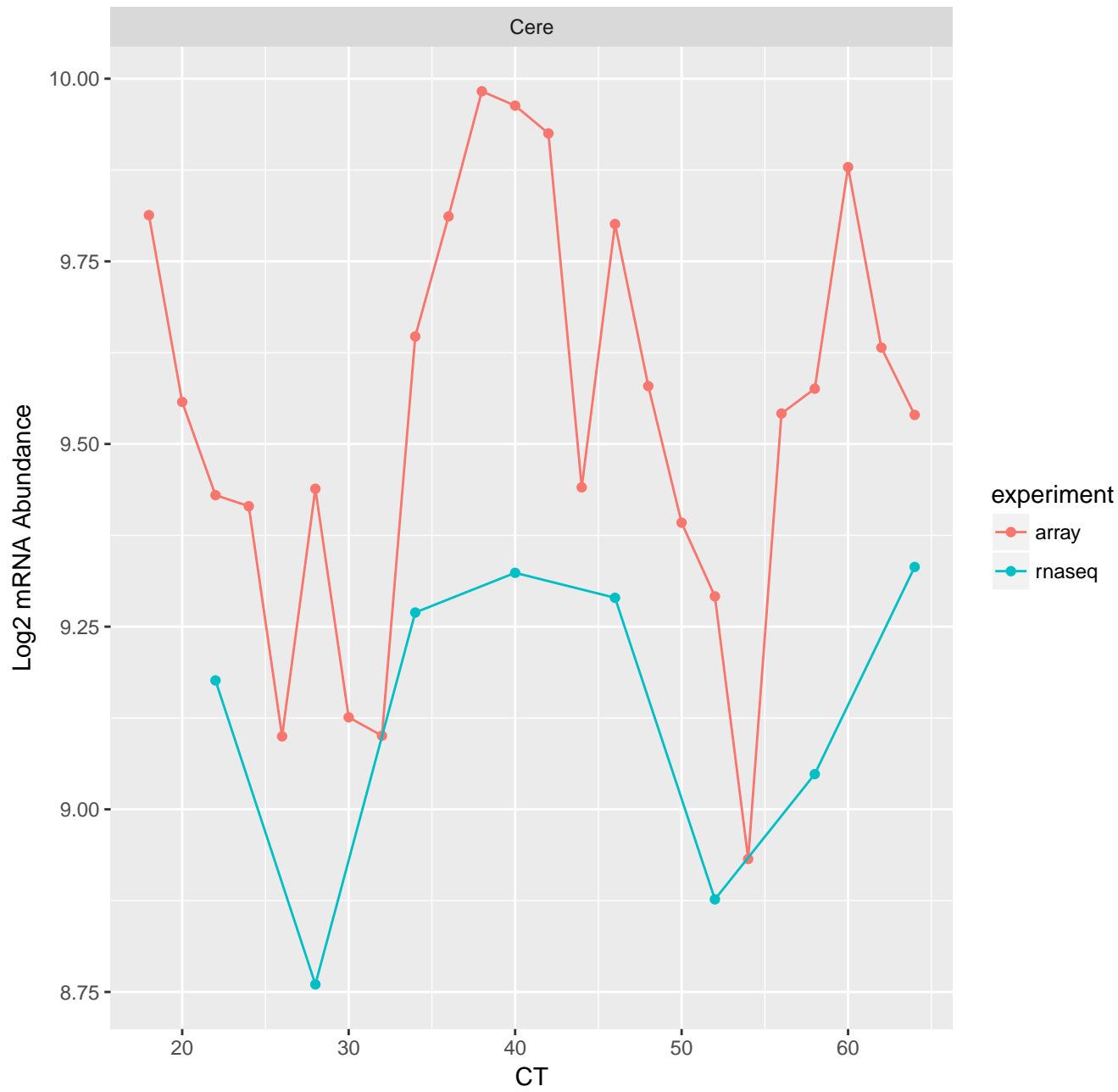
Lfng



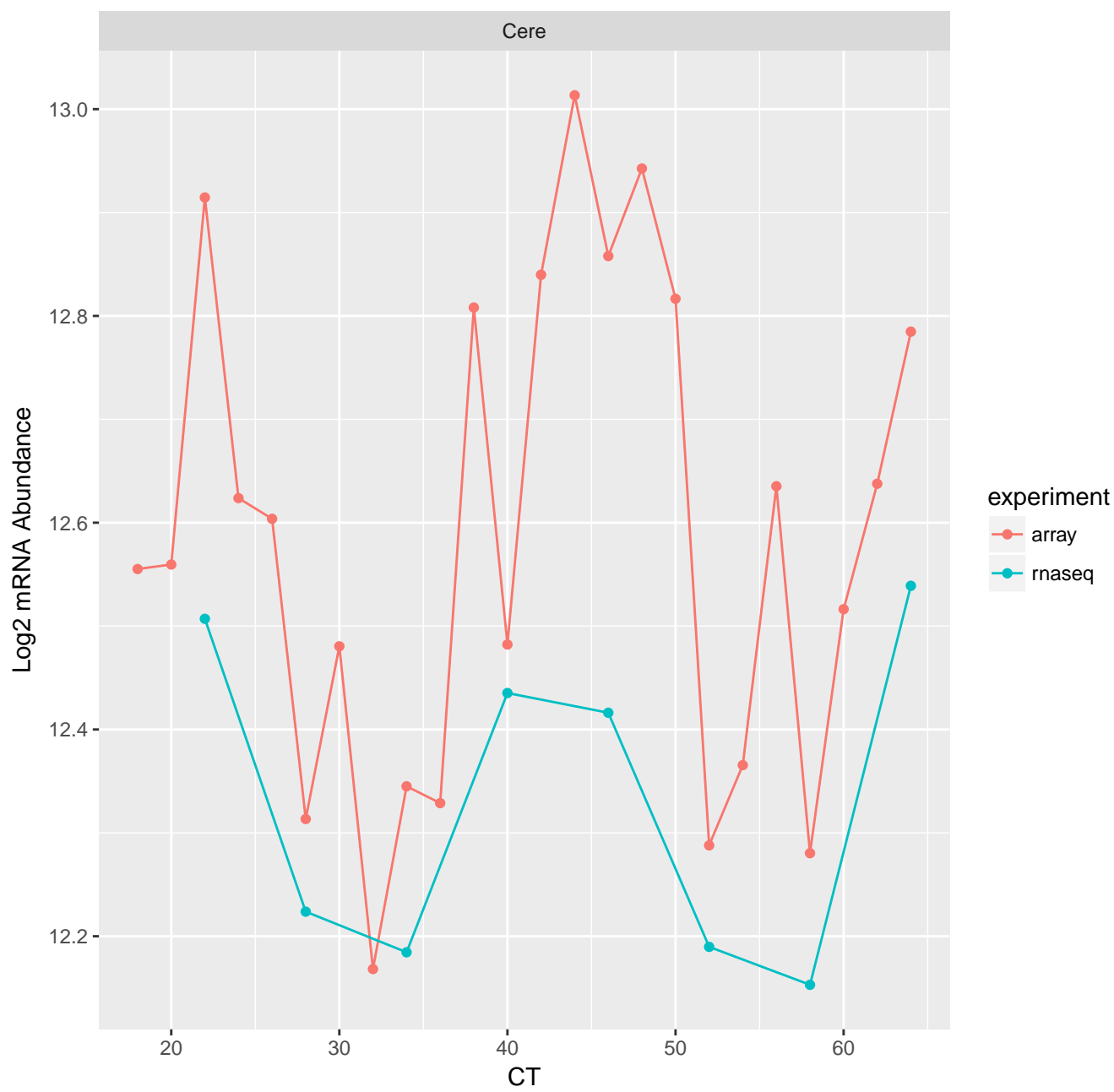
Rasgrp3



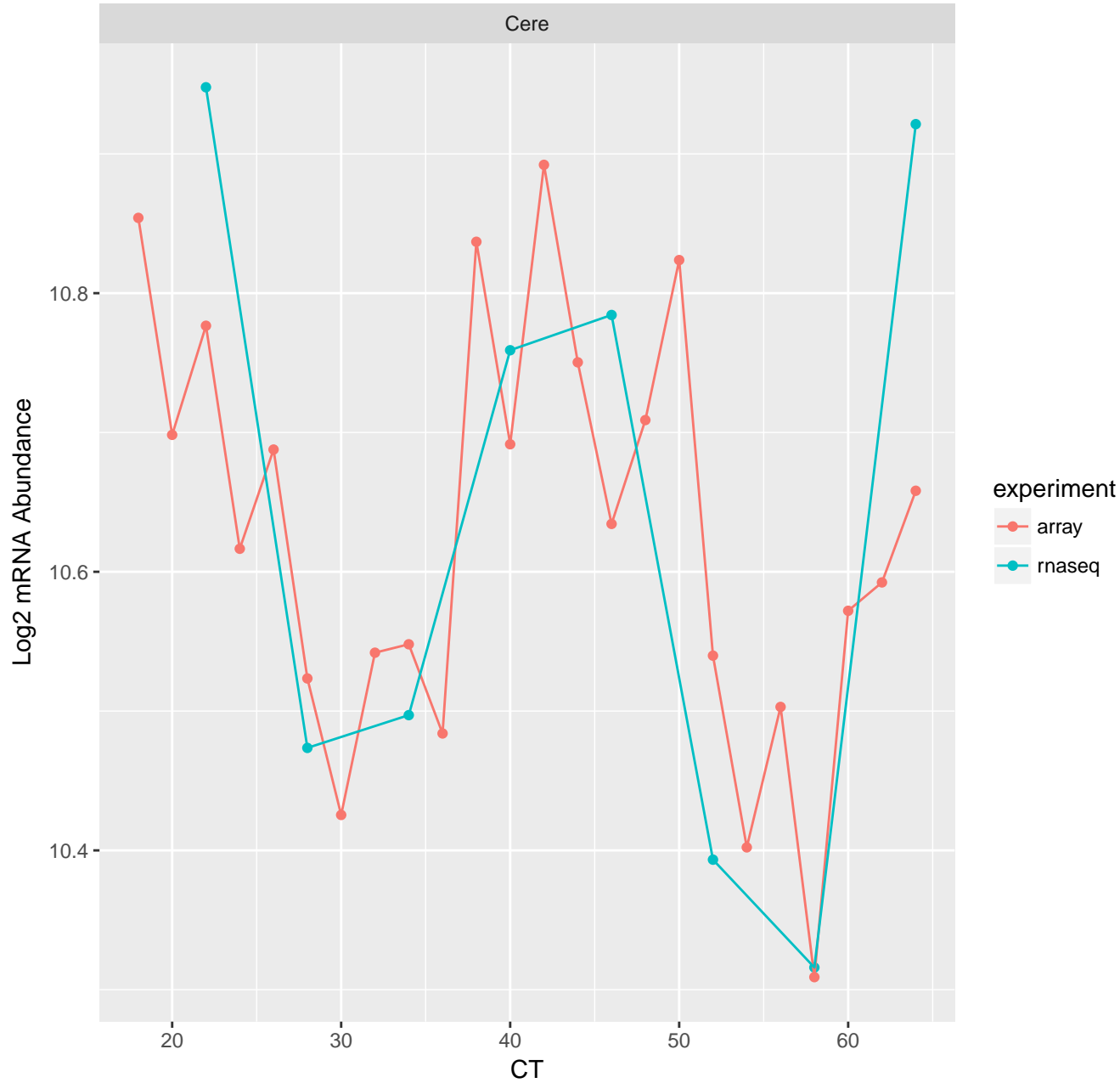
Mettl7a1



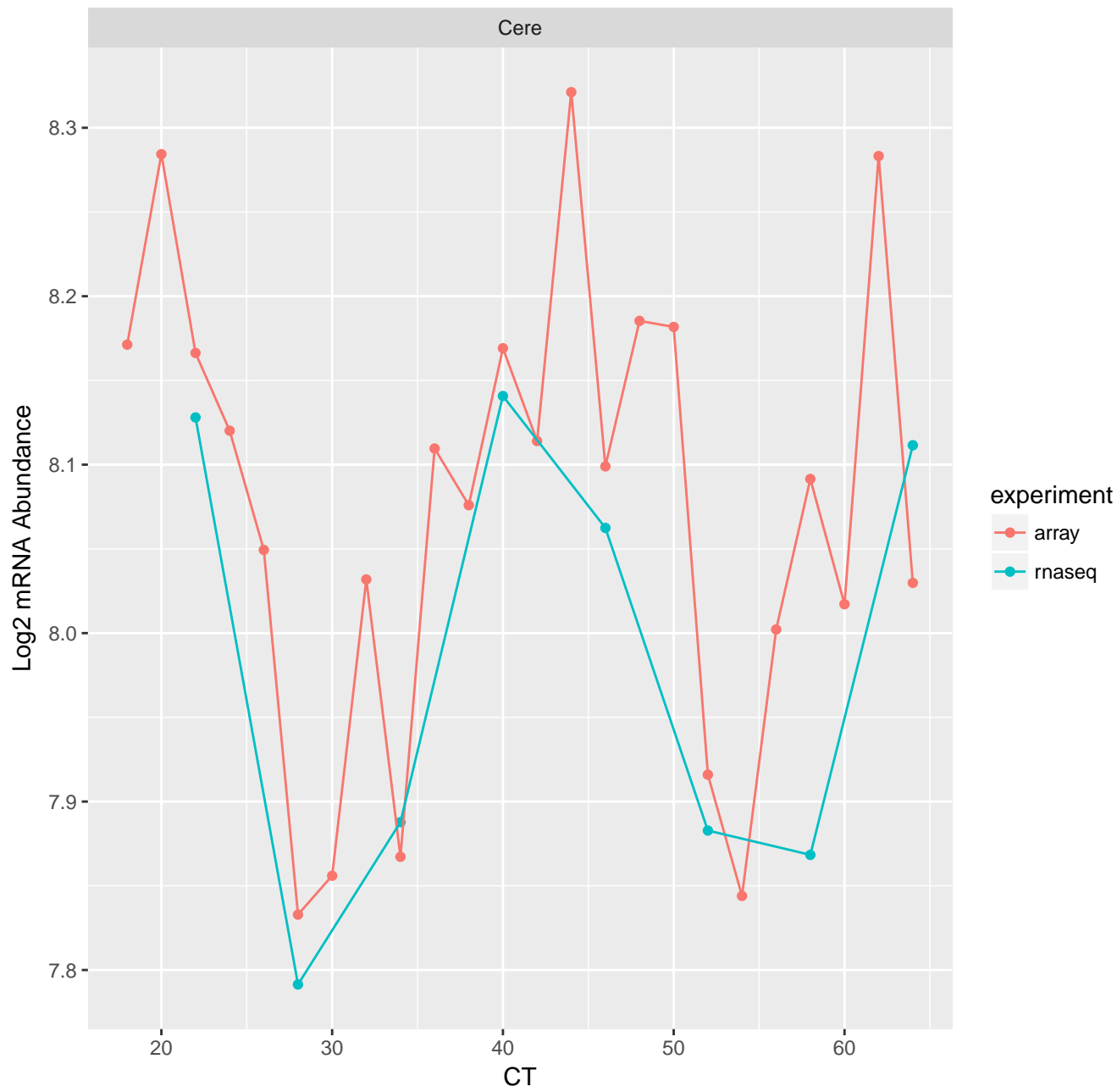
Abcc5



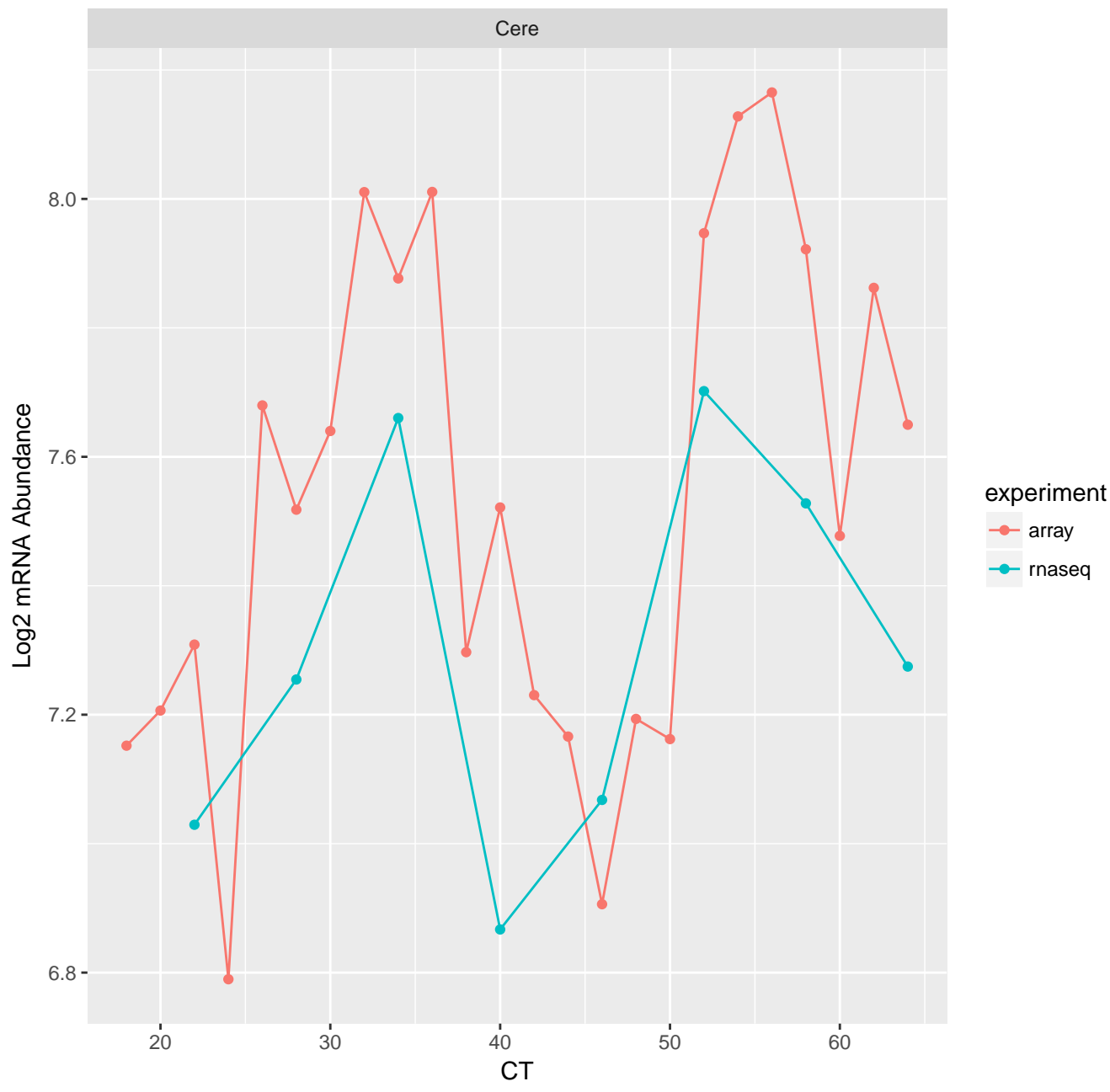
Syne2



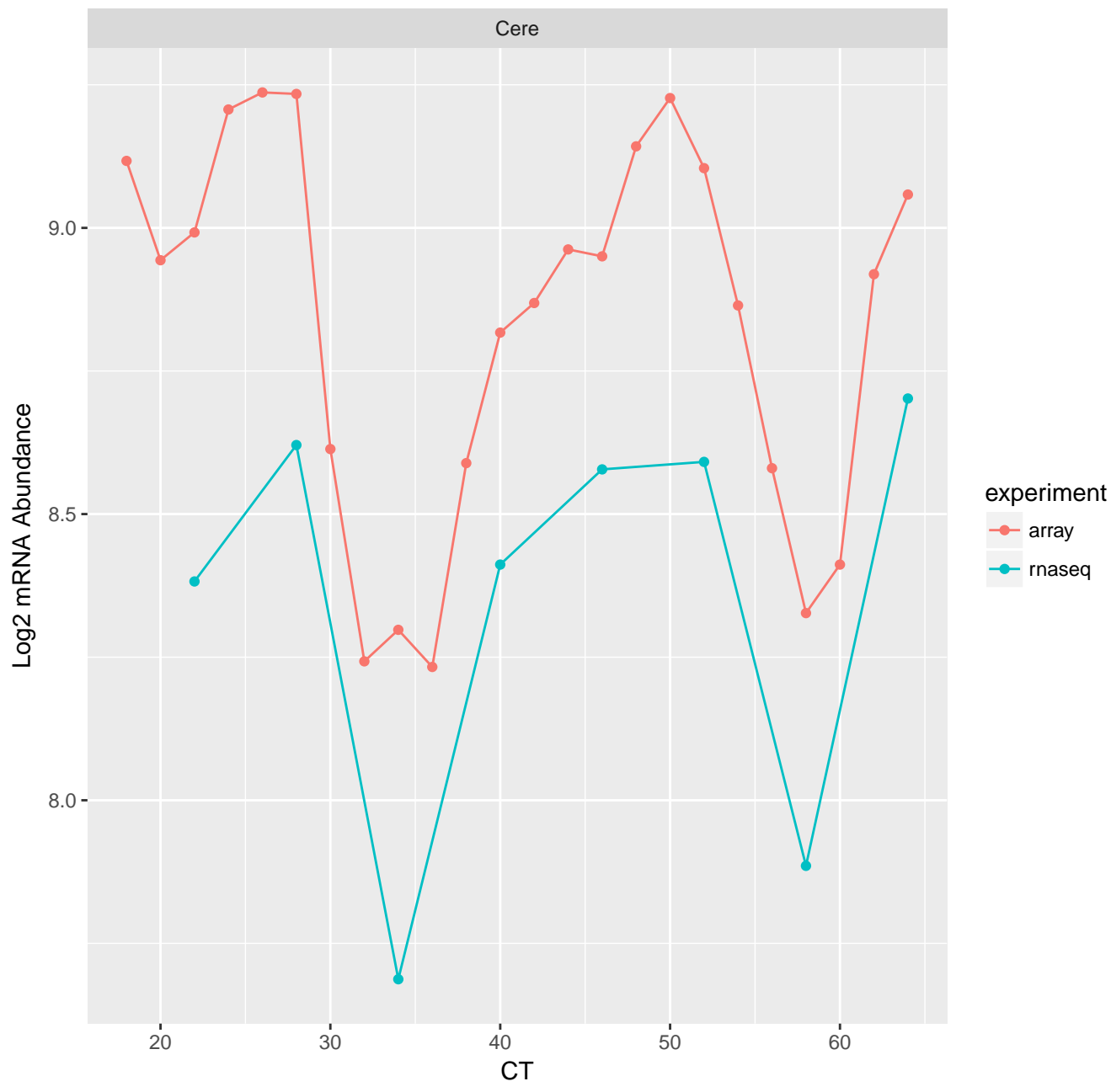
ErbB3



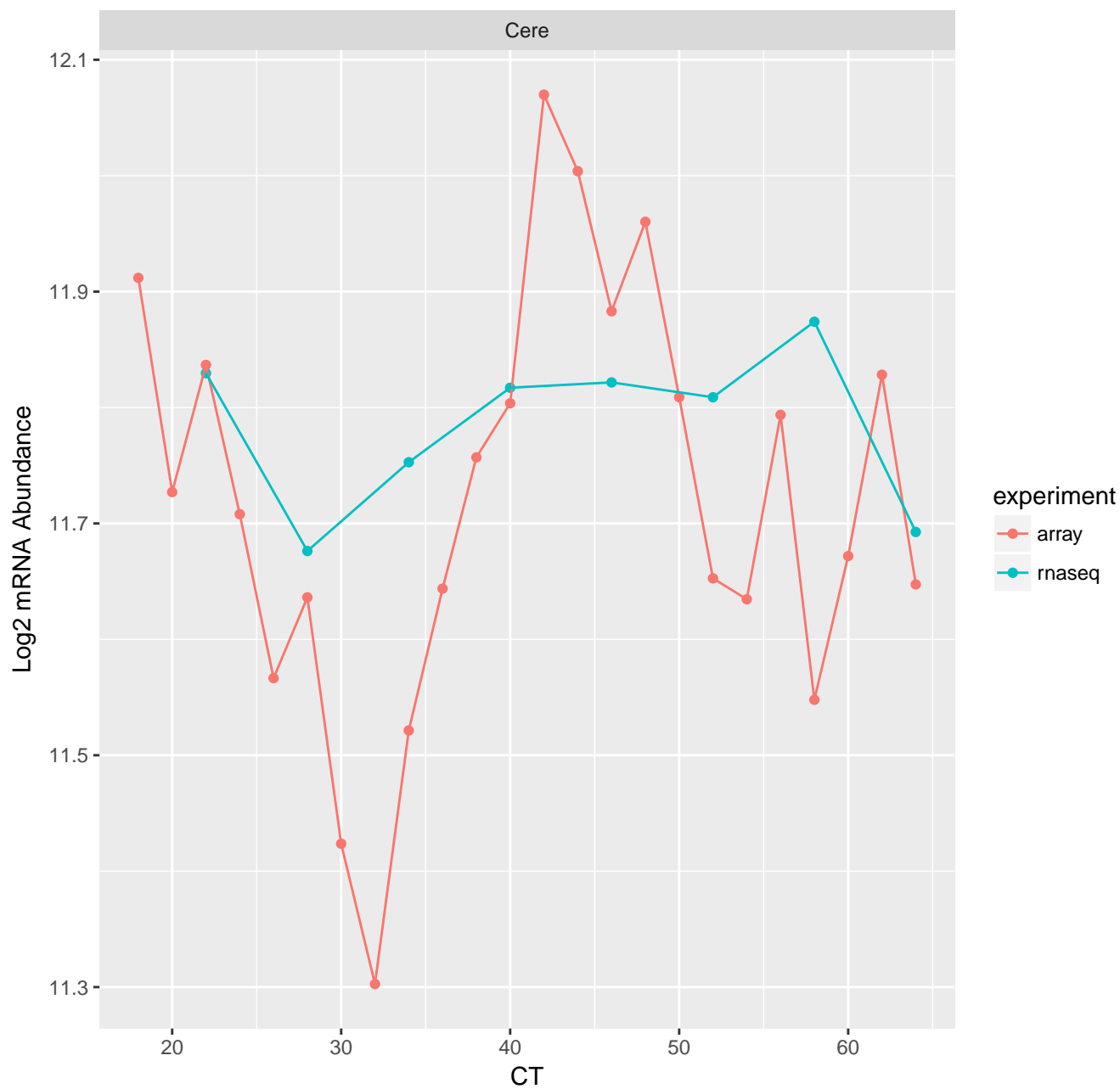
Tmem86a



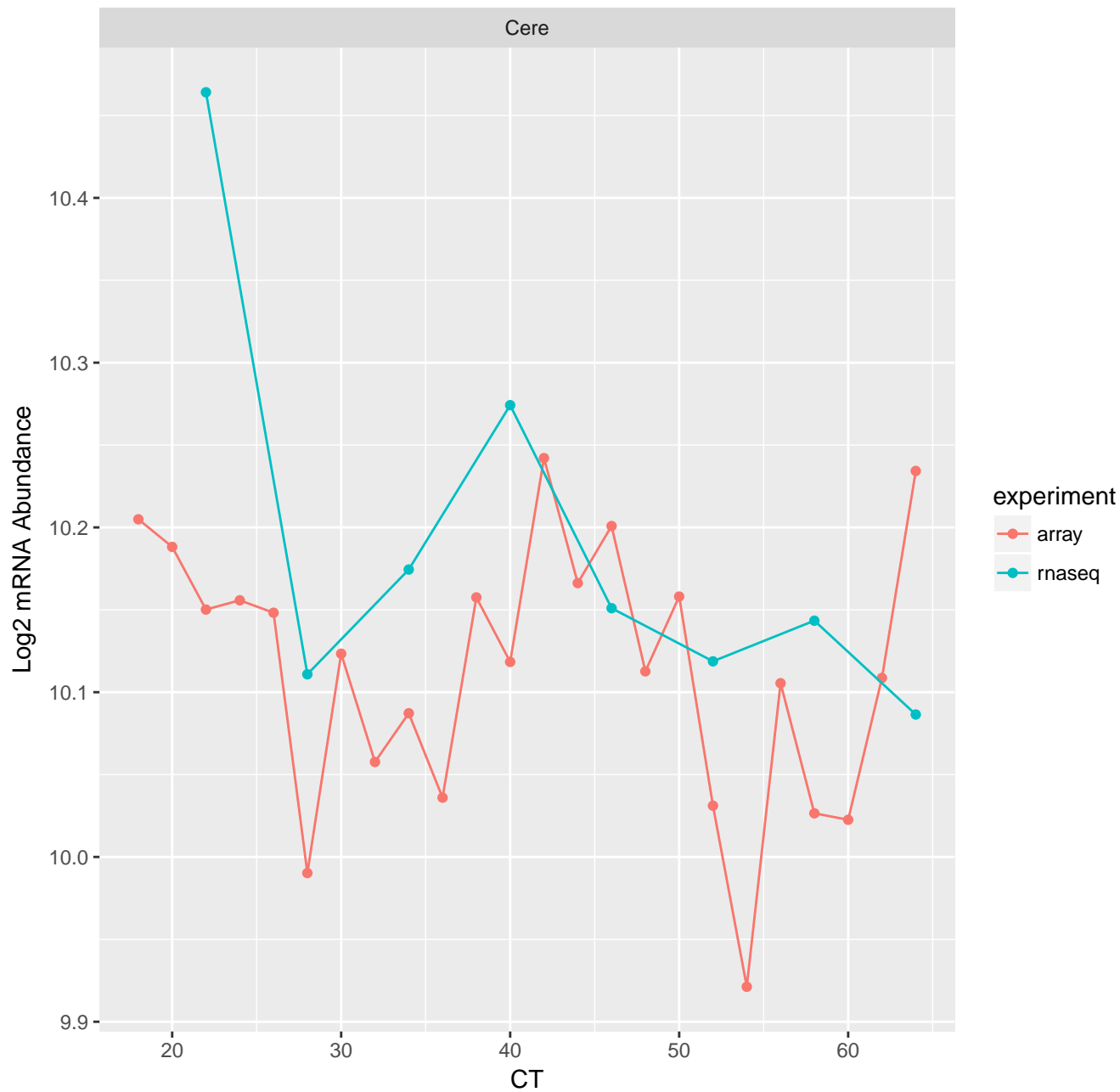
Cldn5



Osbp16



Alms1



Mcc

Cere

Log2 mRNA Abundance

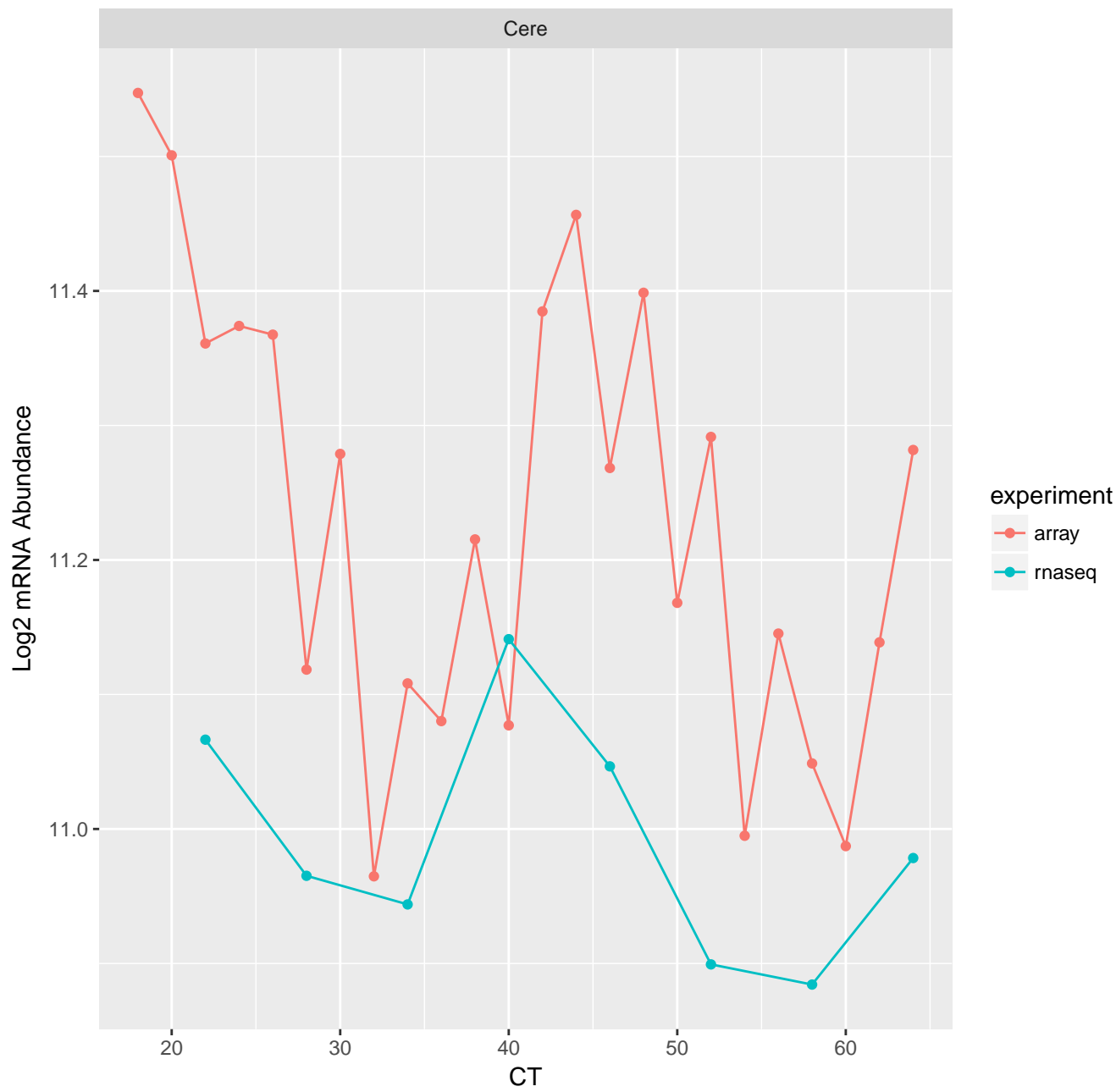
CT

experiment

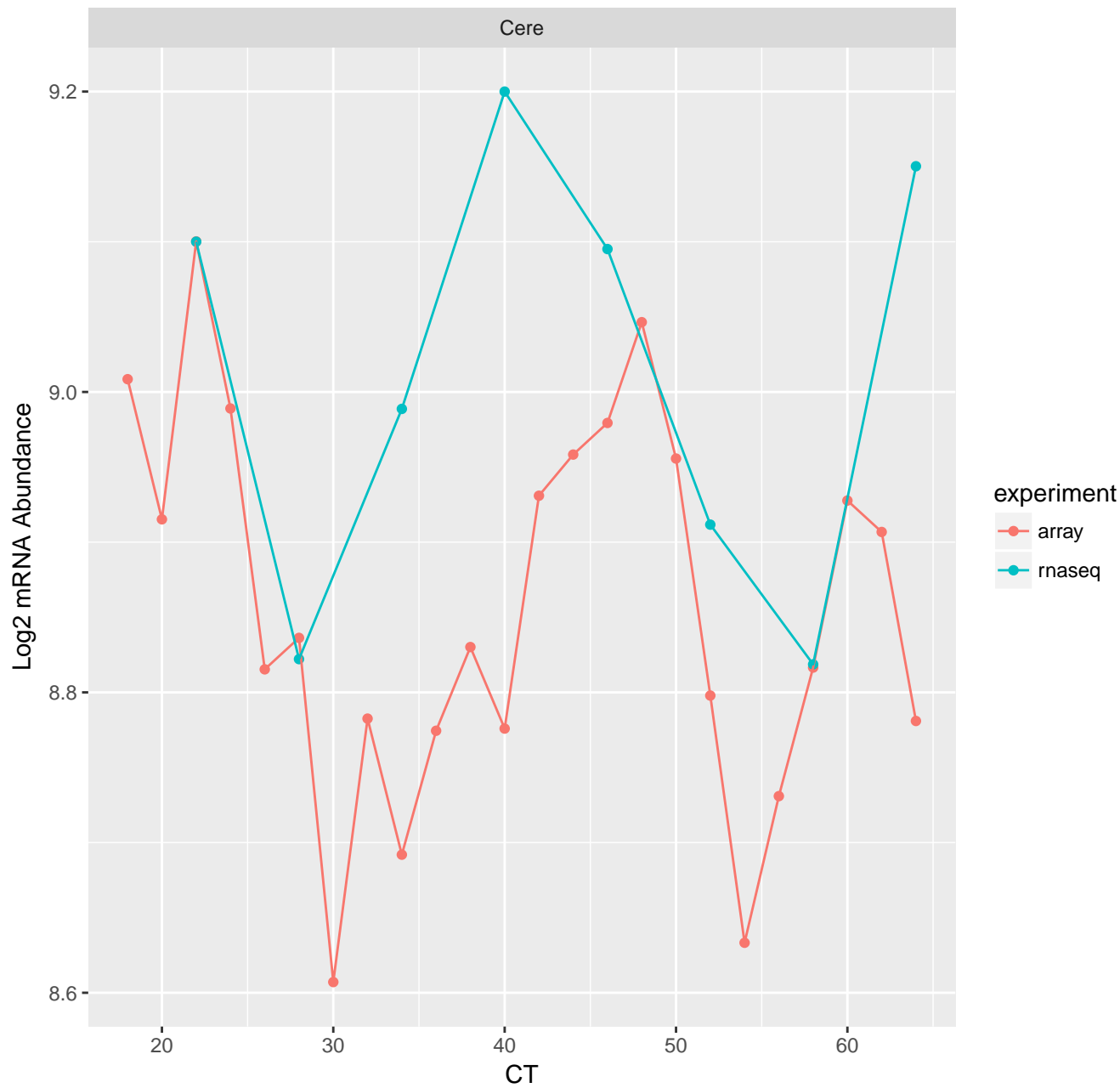
array
rnaseq

10.2
10.0
9.8

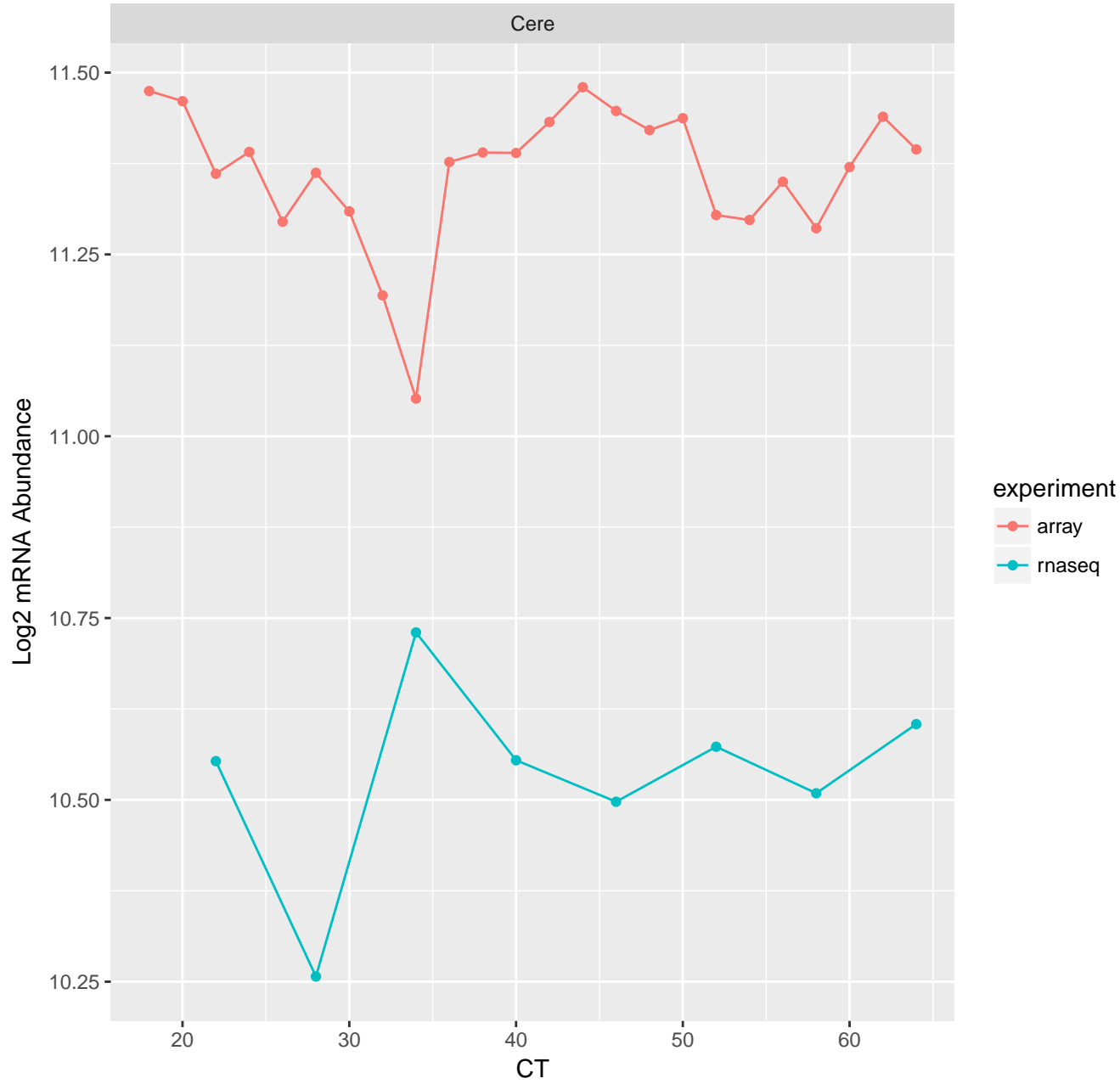
Cntn6



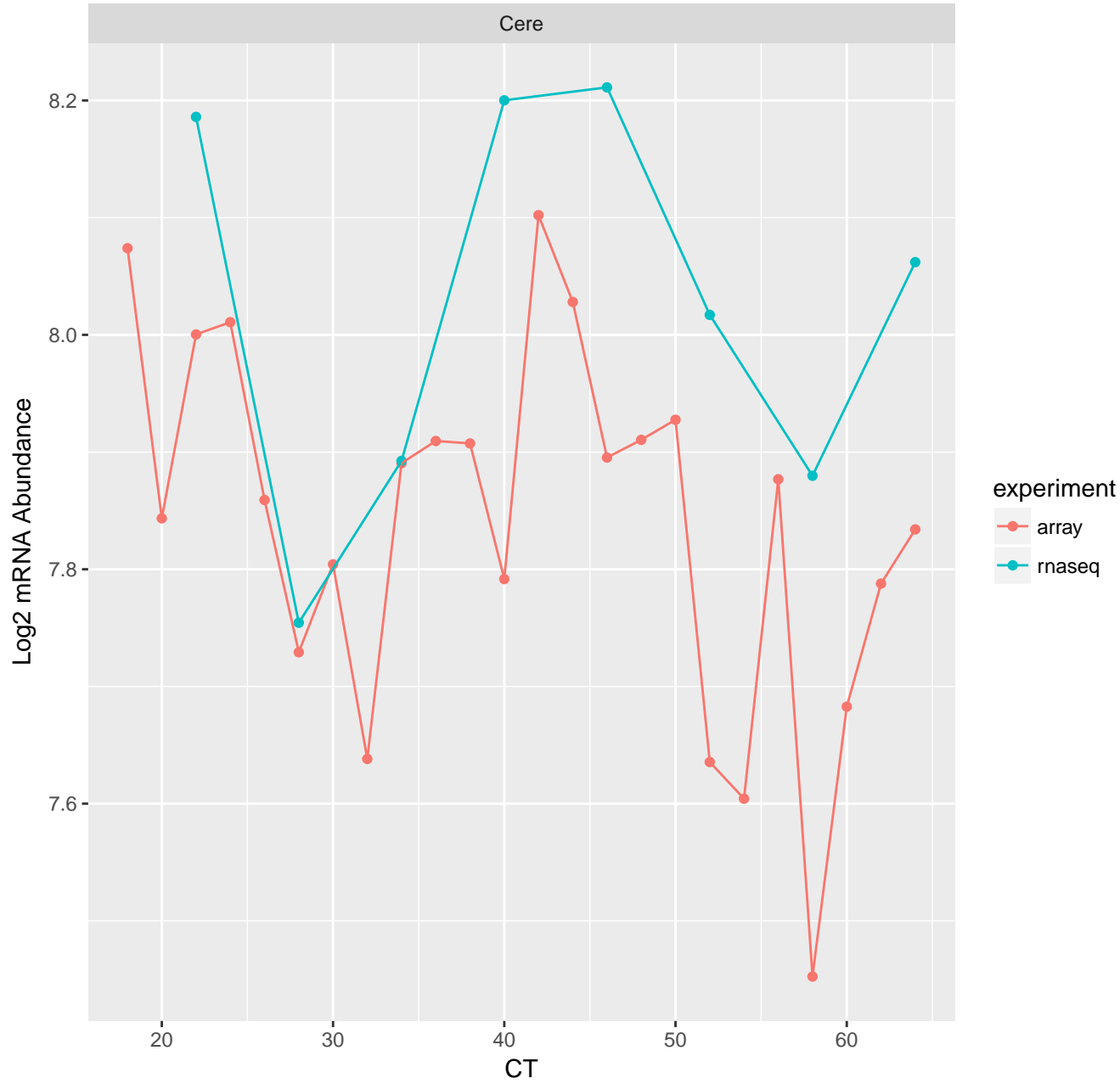
Nfya



Dpysl2



Pdgfd



Clic5

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

7.5

7.0

6.5

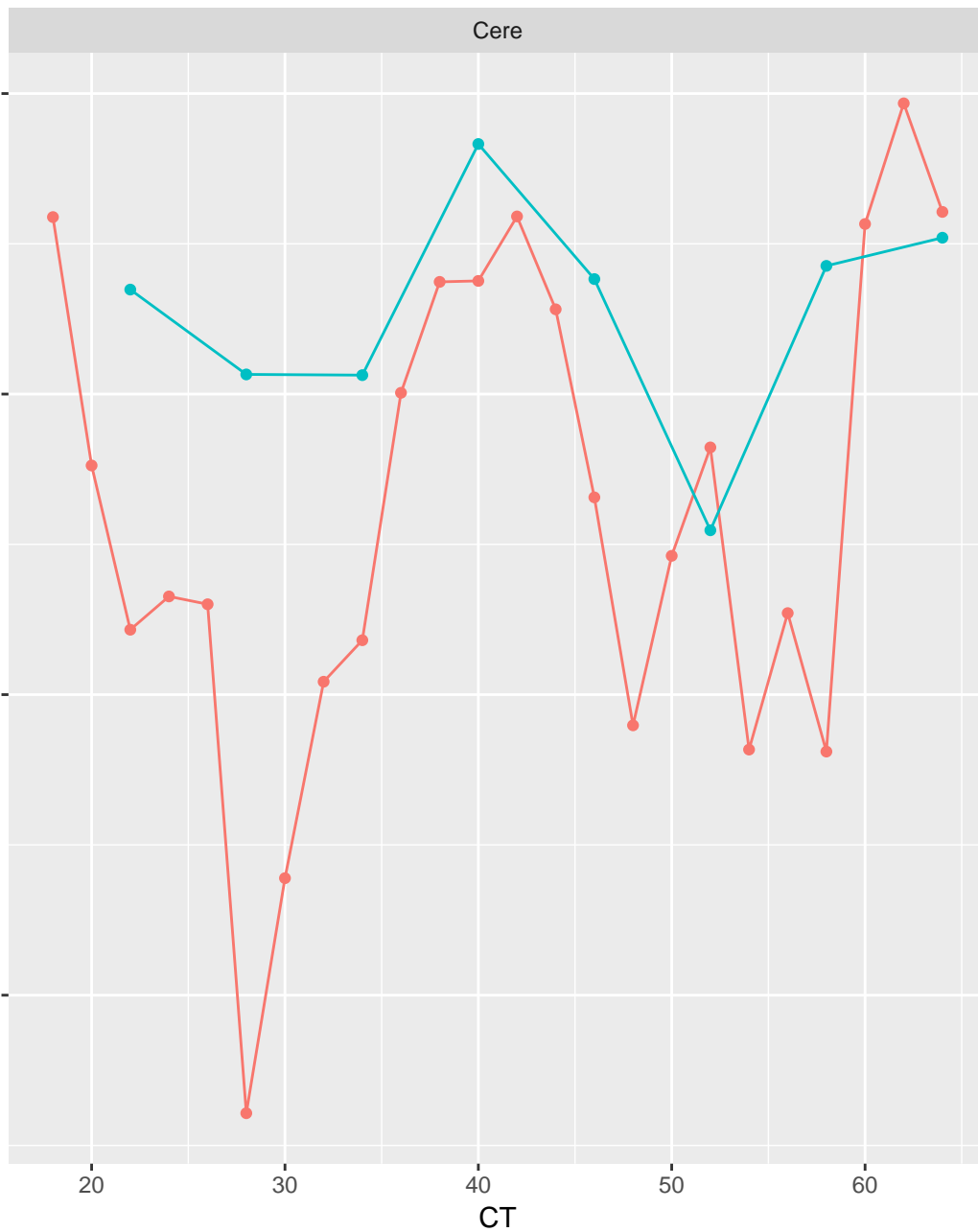
6.0

20

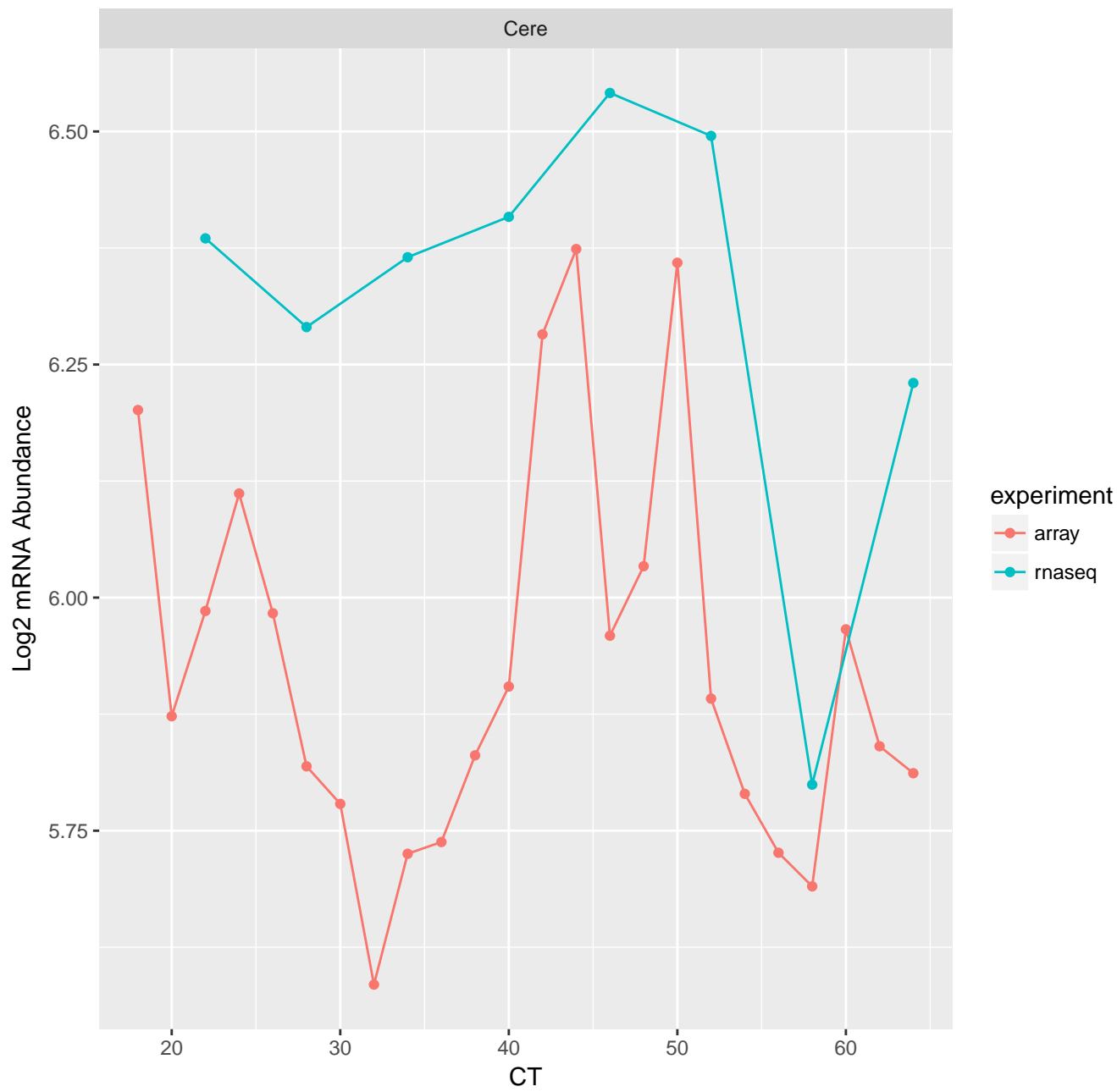
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CT

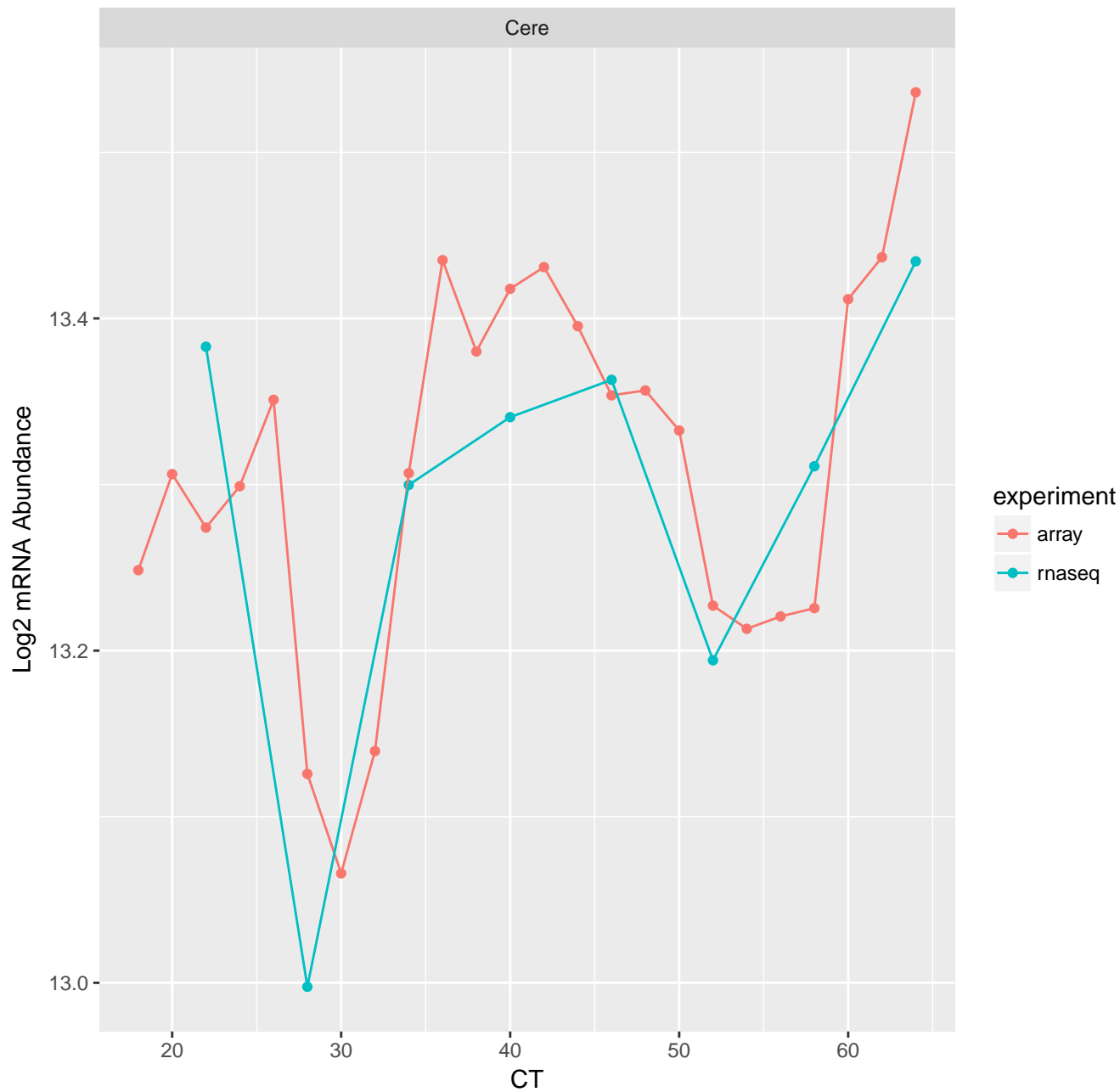
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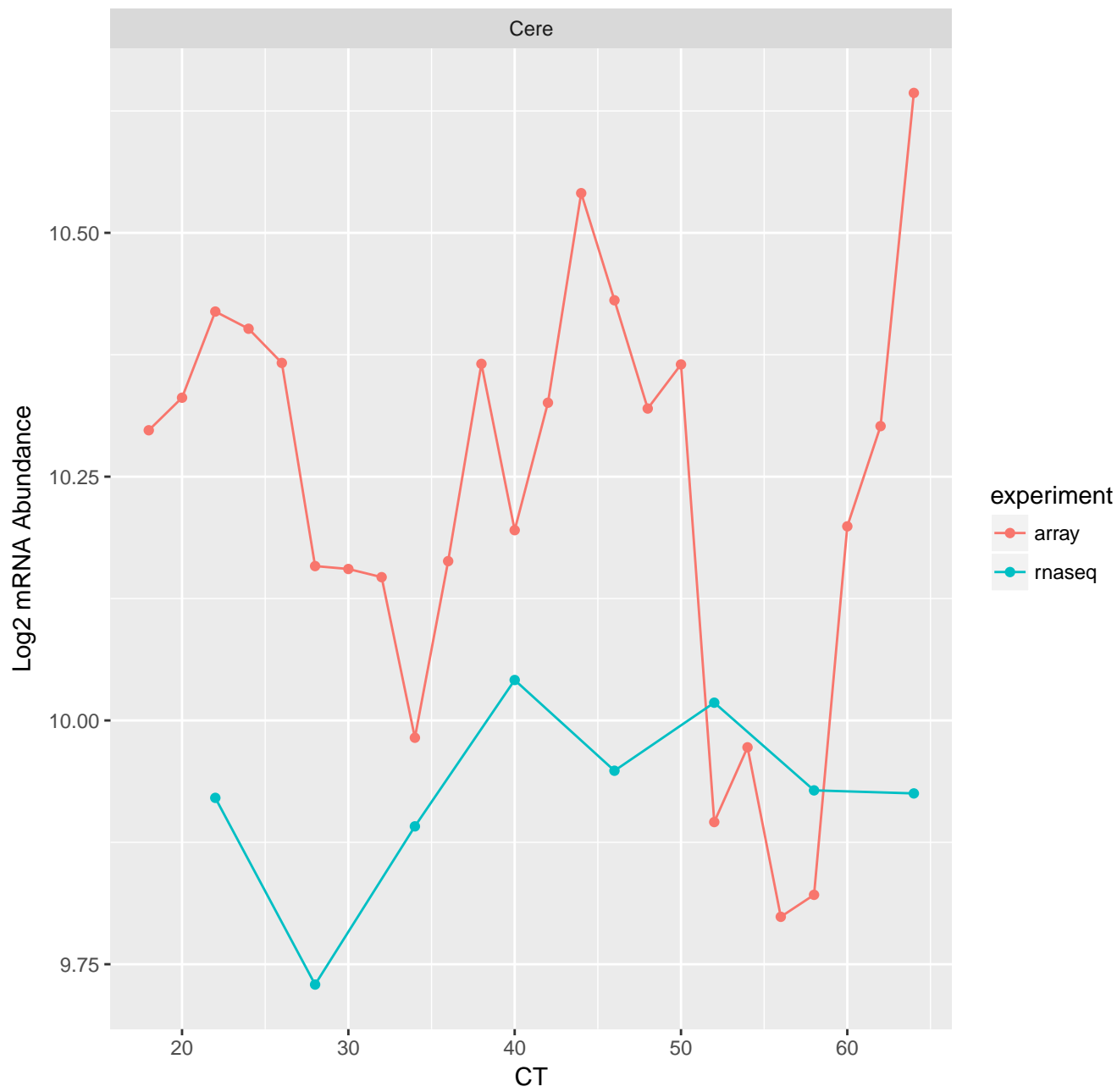
Ceacam1



Ablim1



0610037L13Rik



Pclo

Cere

Log2 mRNA Abundance

experiment

array
rnaseq

14.0
13.5
13.0

CT

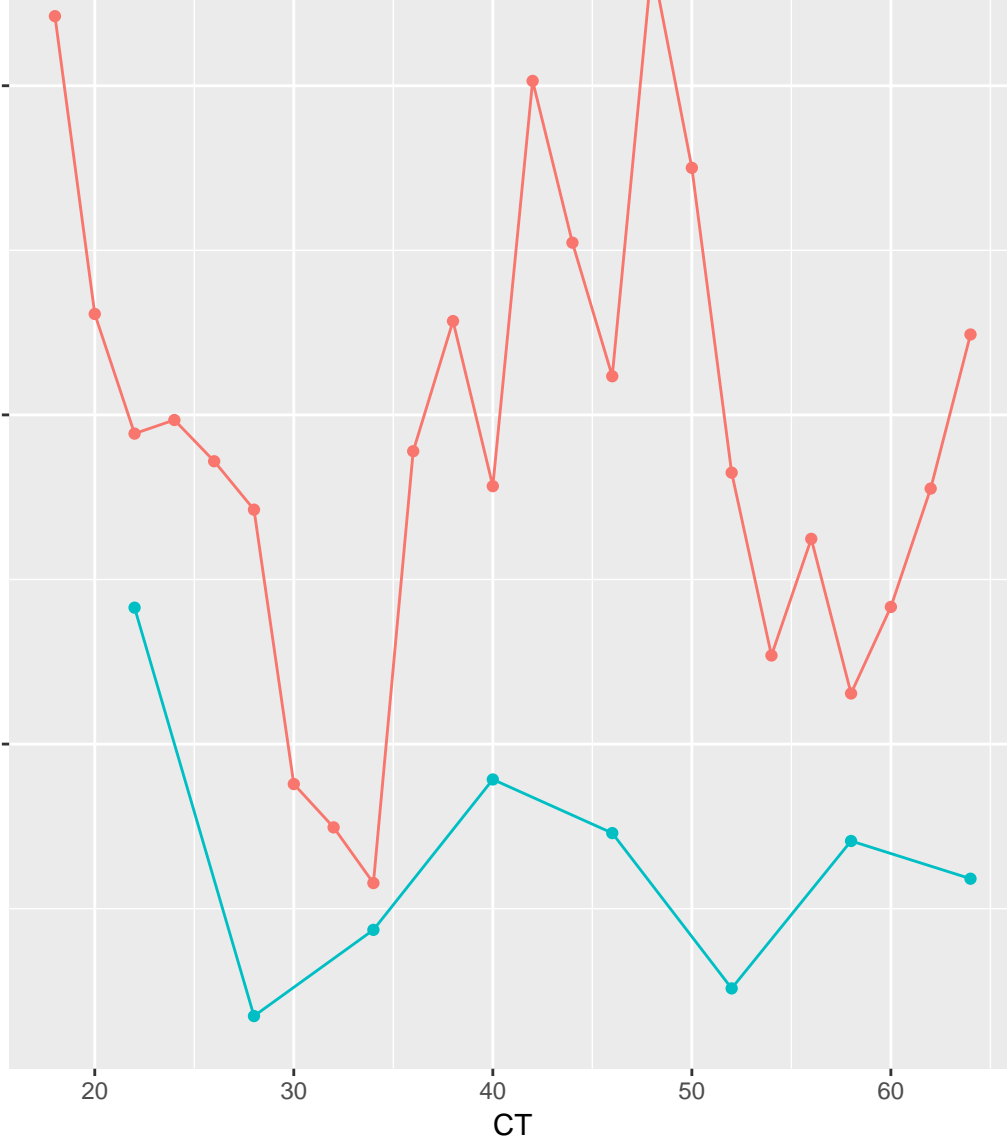
20

30

40

50

60



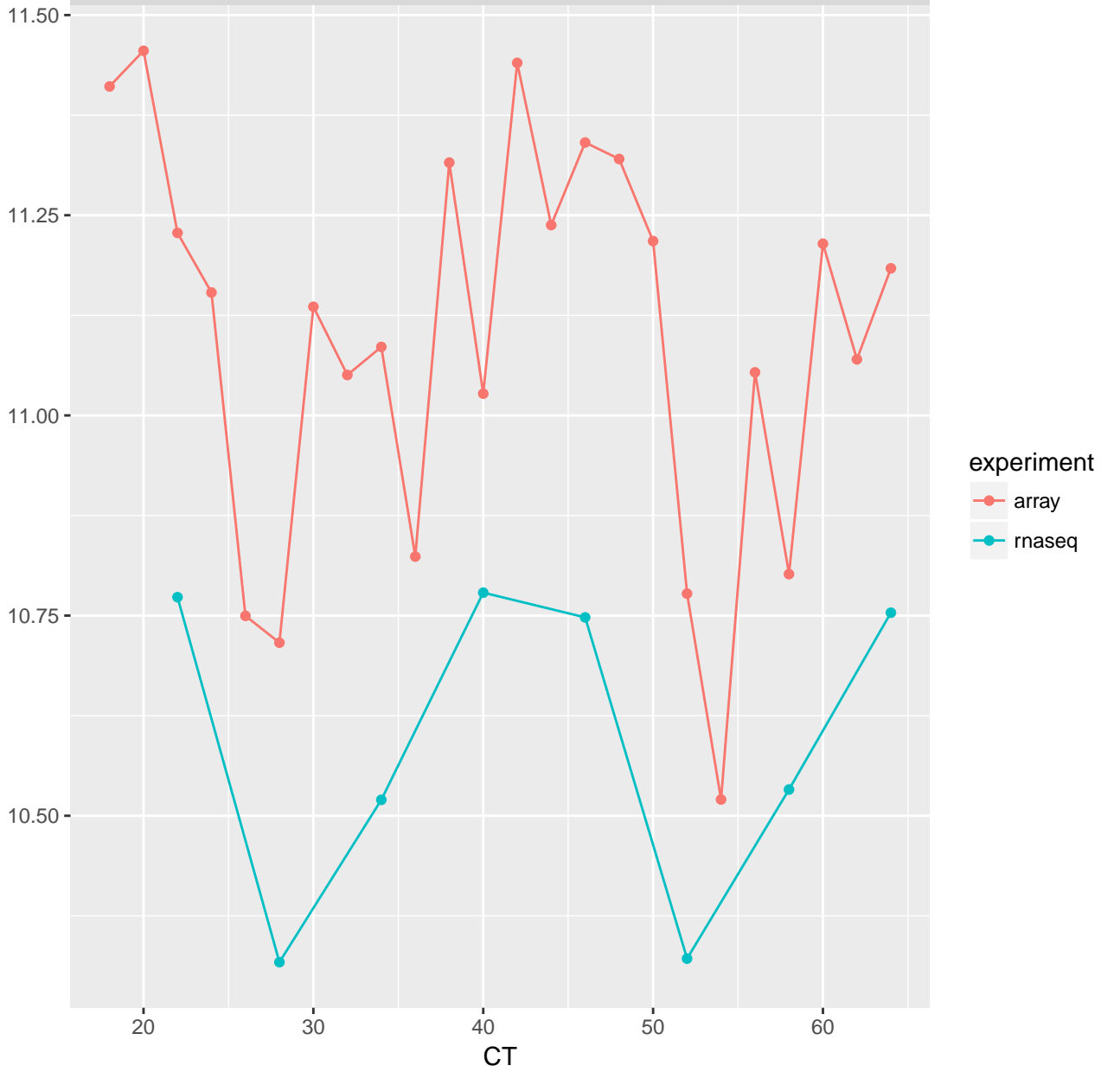
Tra2a

Cere

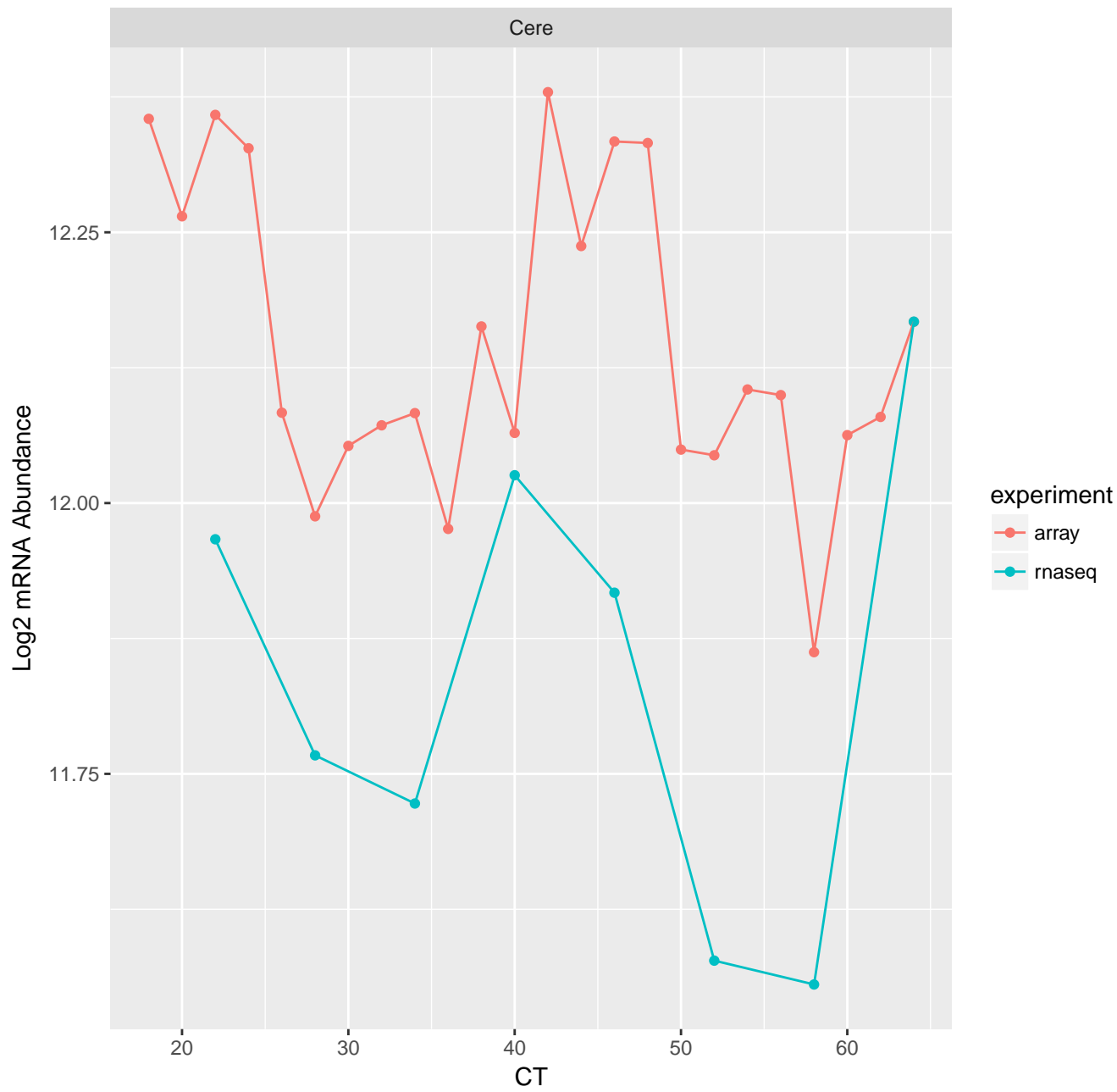
Log2 mRNA Abundance

experiment

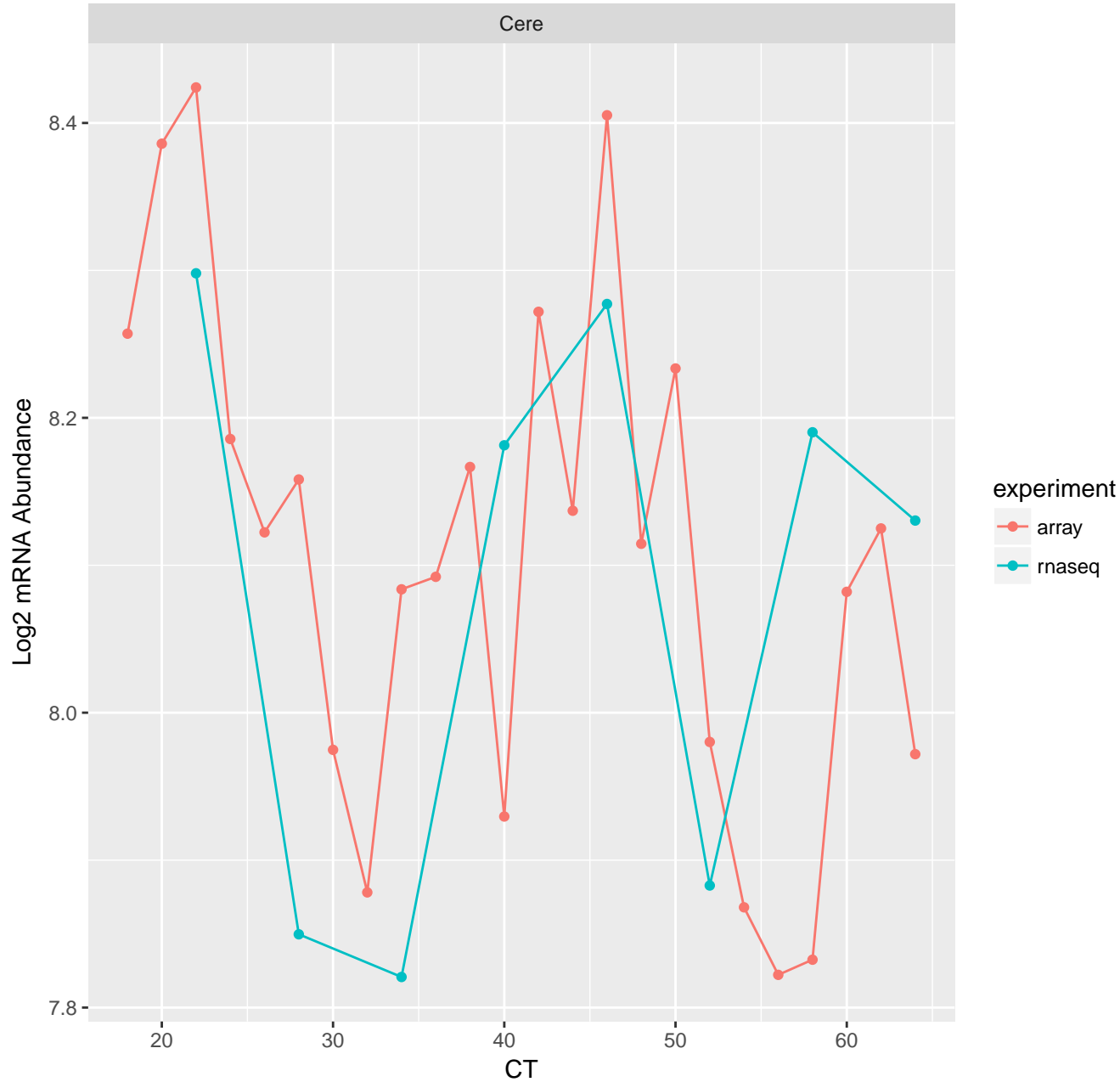
array
rnaseq



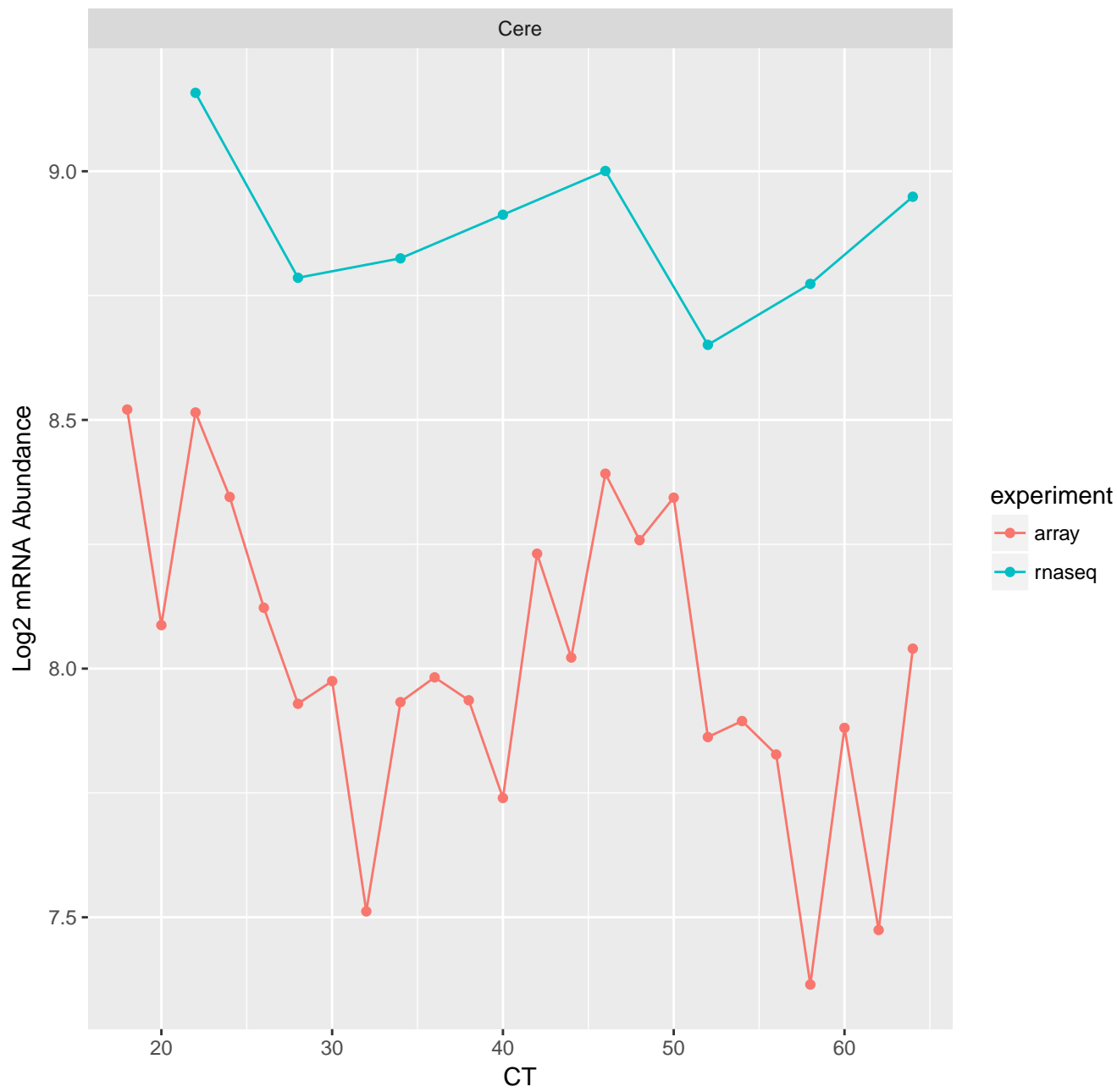
Ccn12



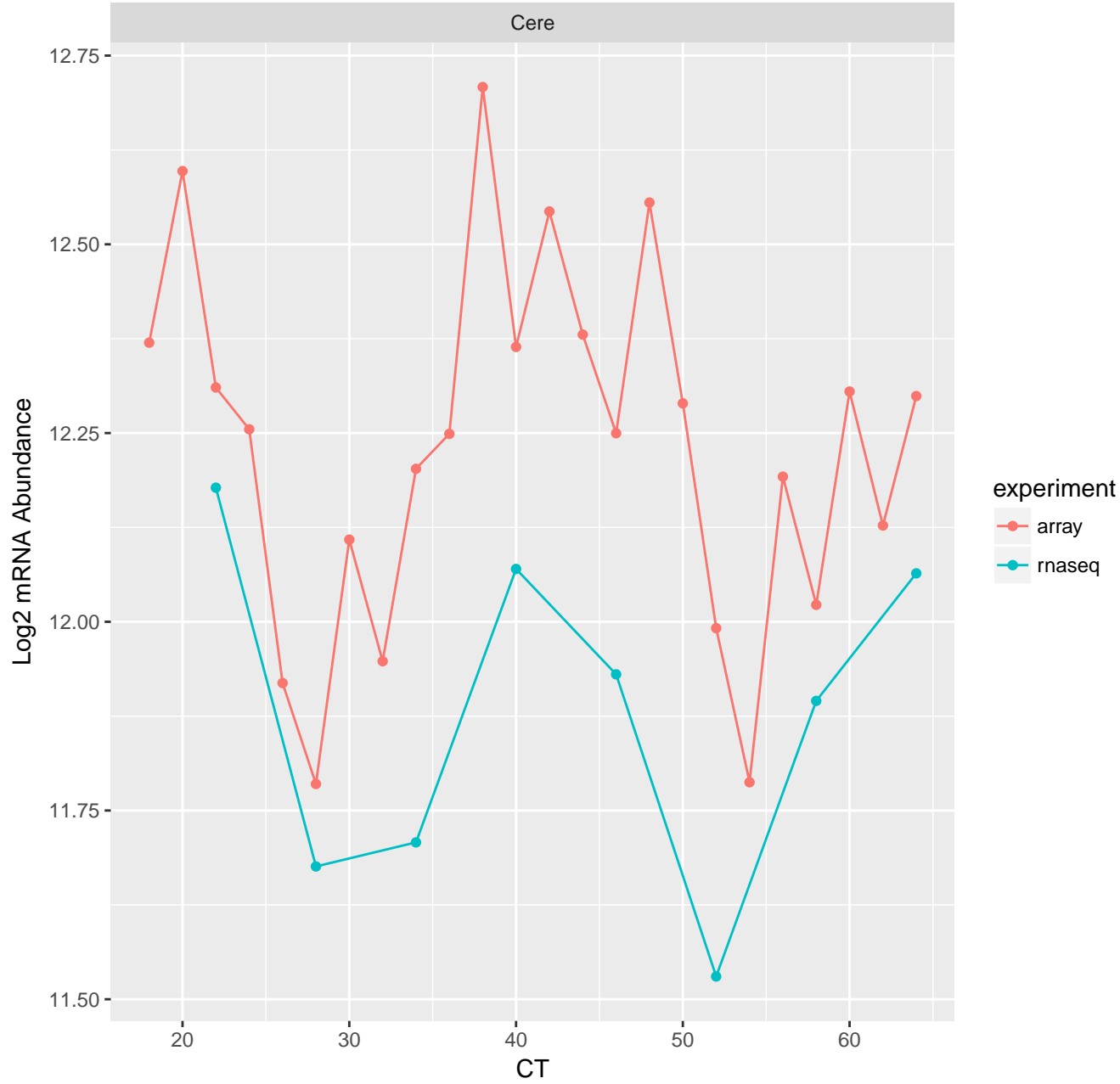
Igfbp3



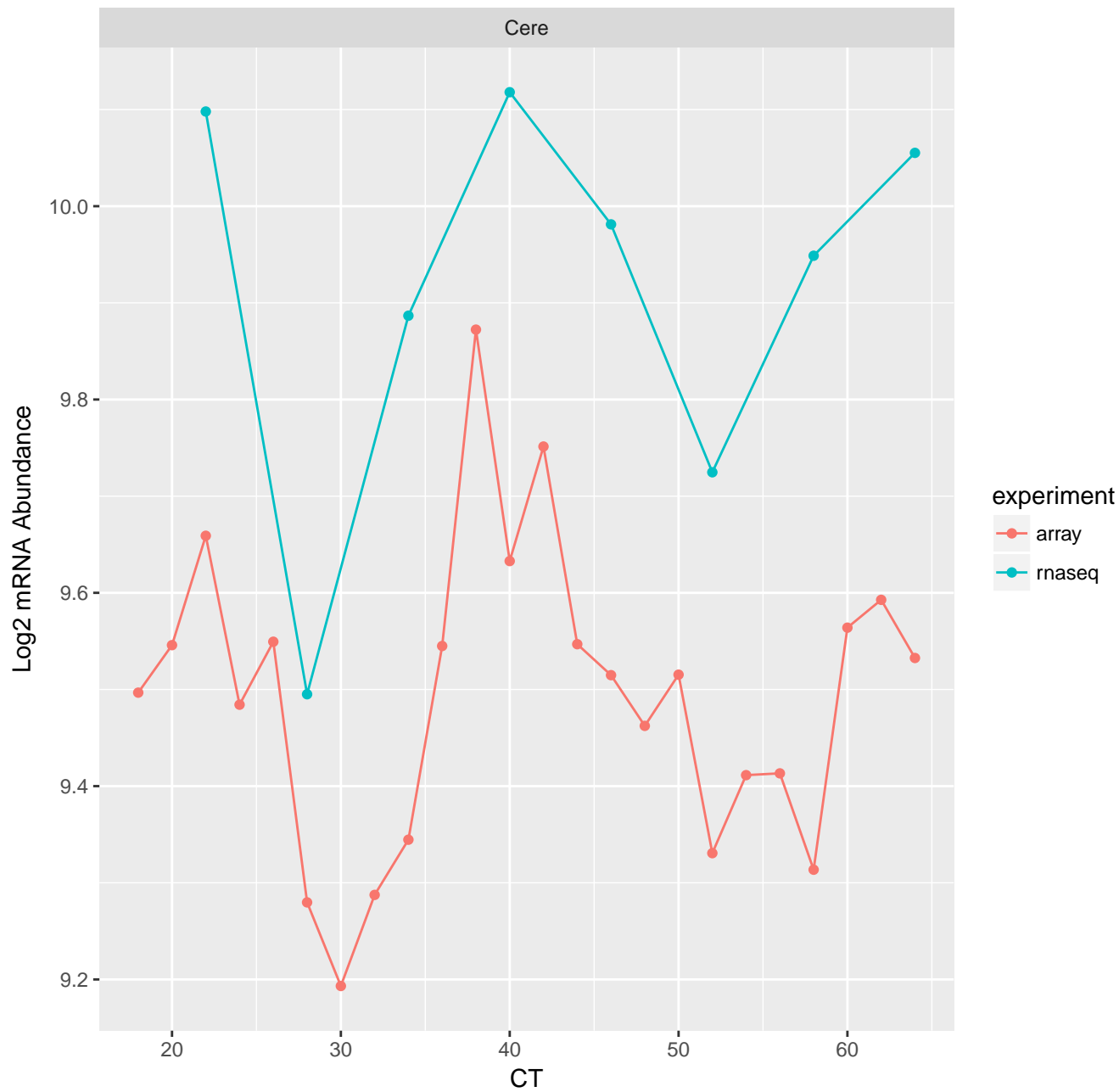
Elmod2



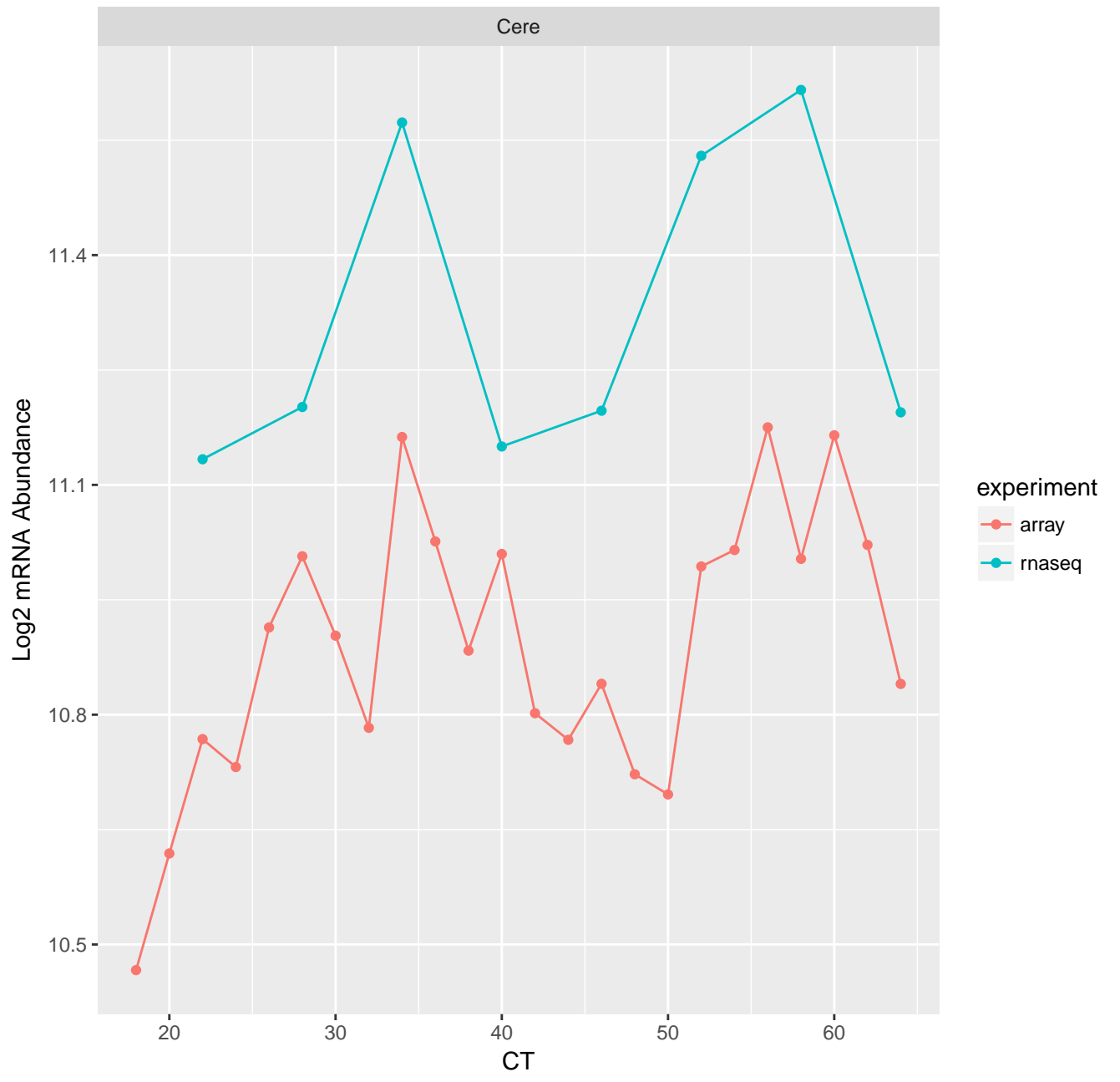
Sfpq

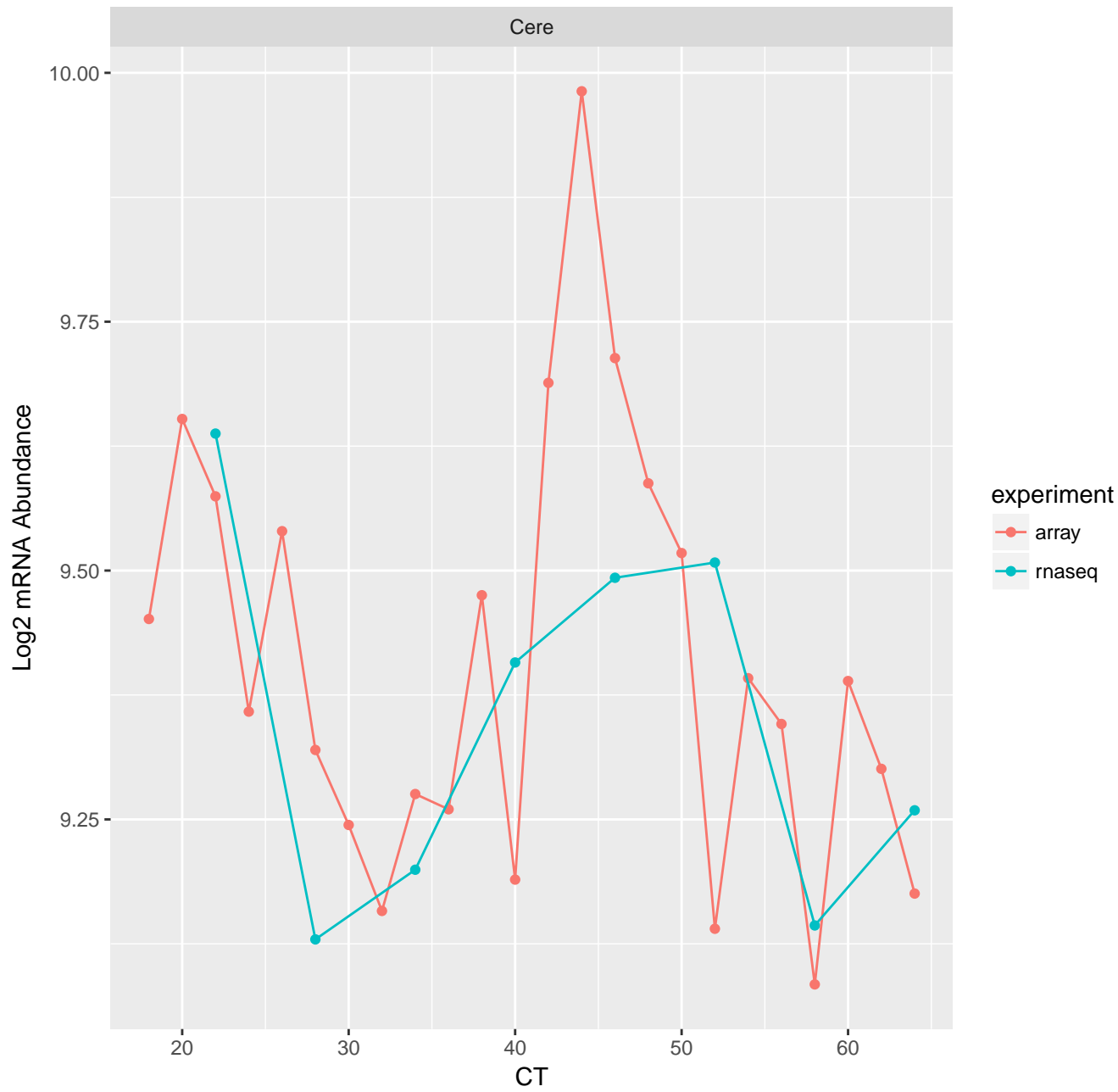


Plice1

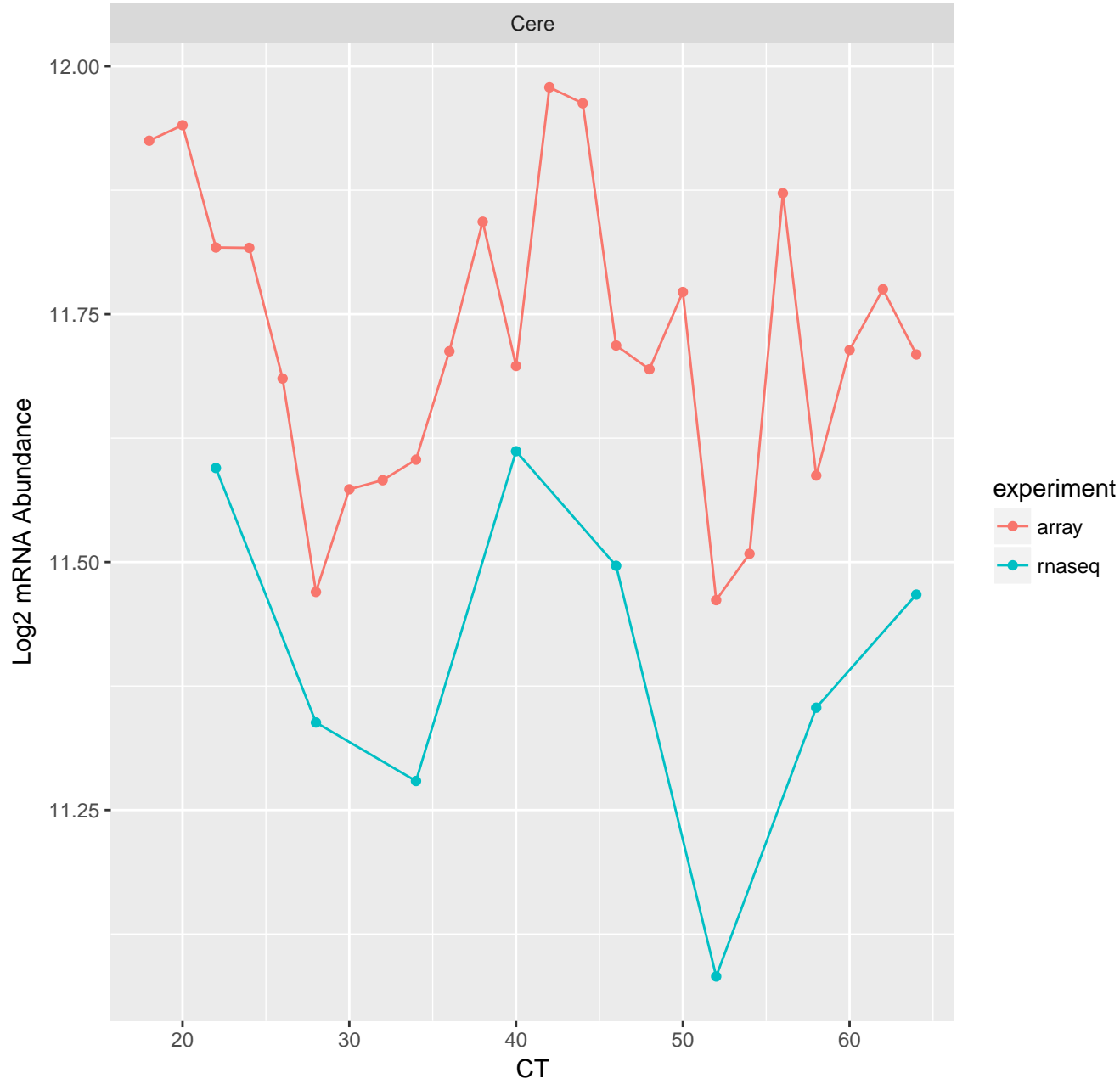


Med12

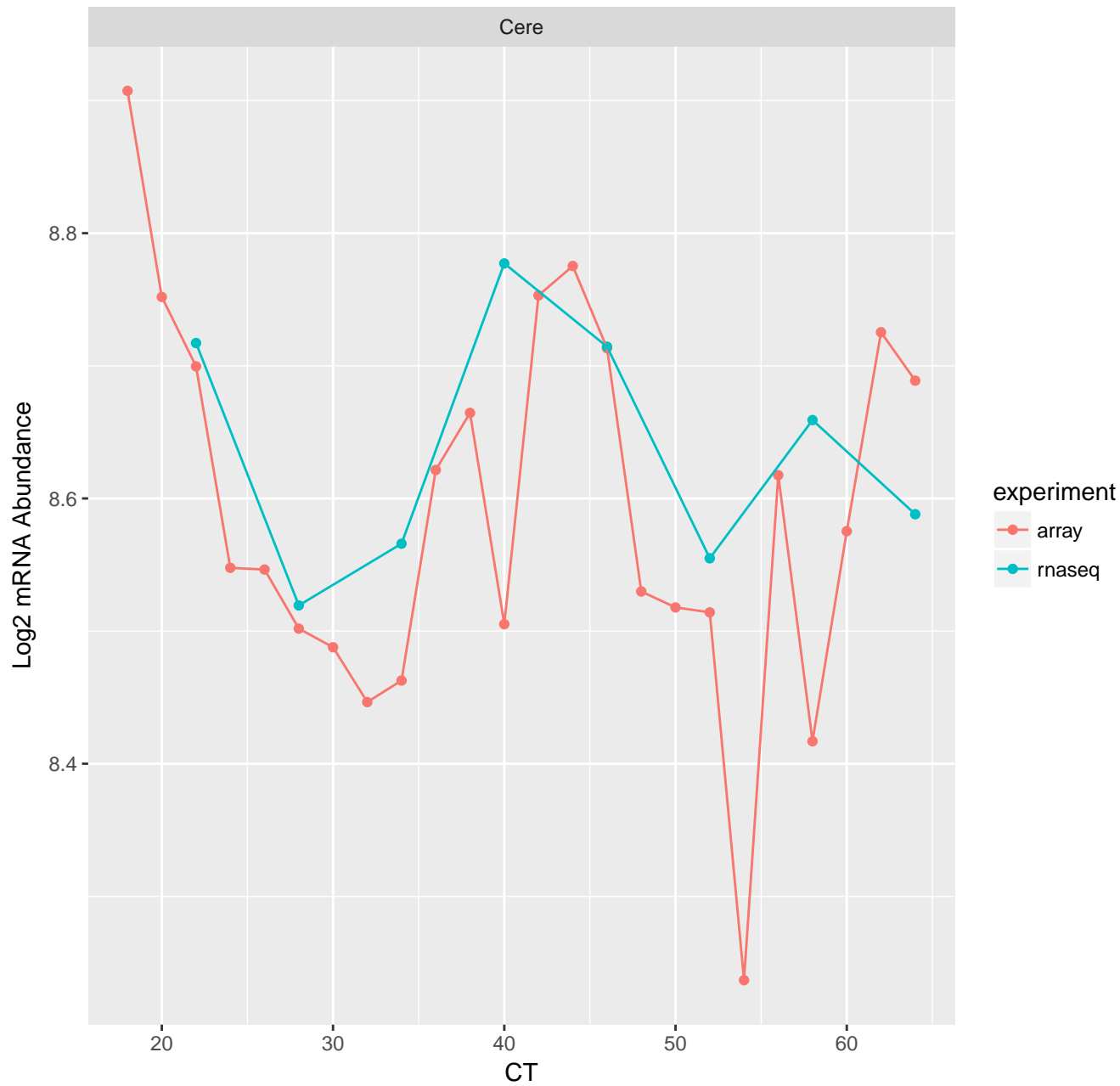




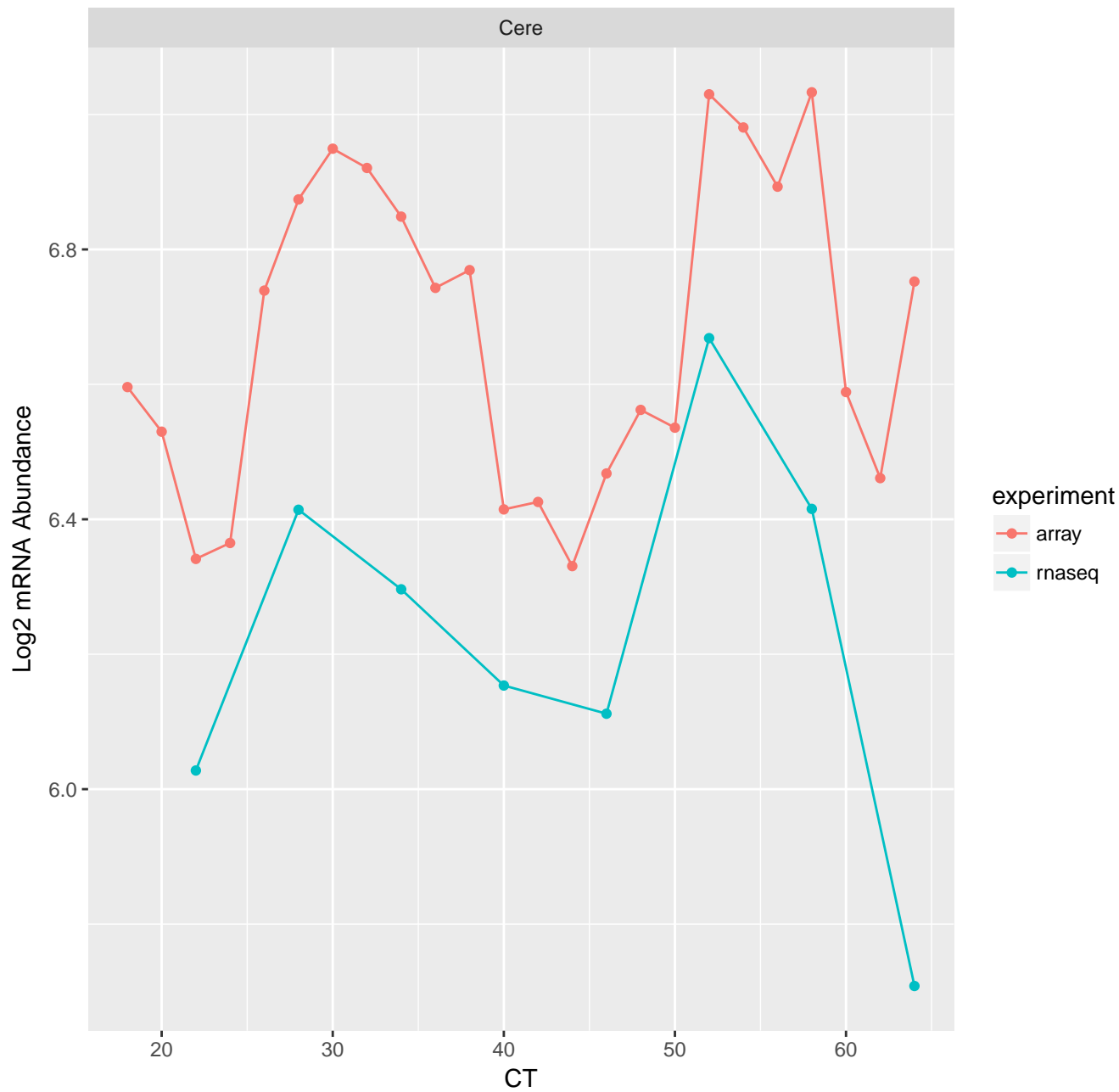
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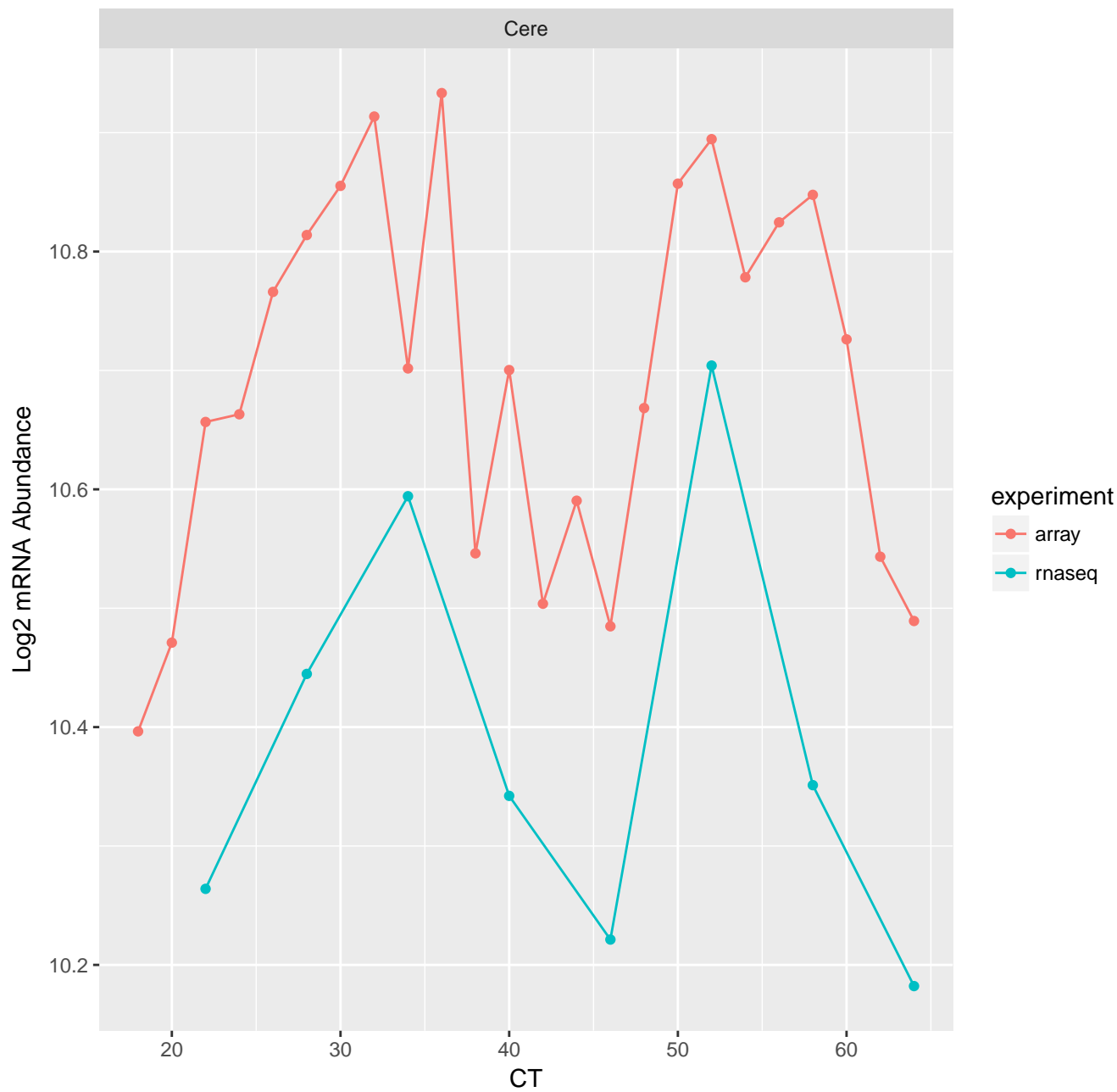
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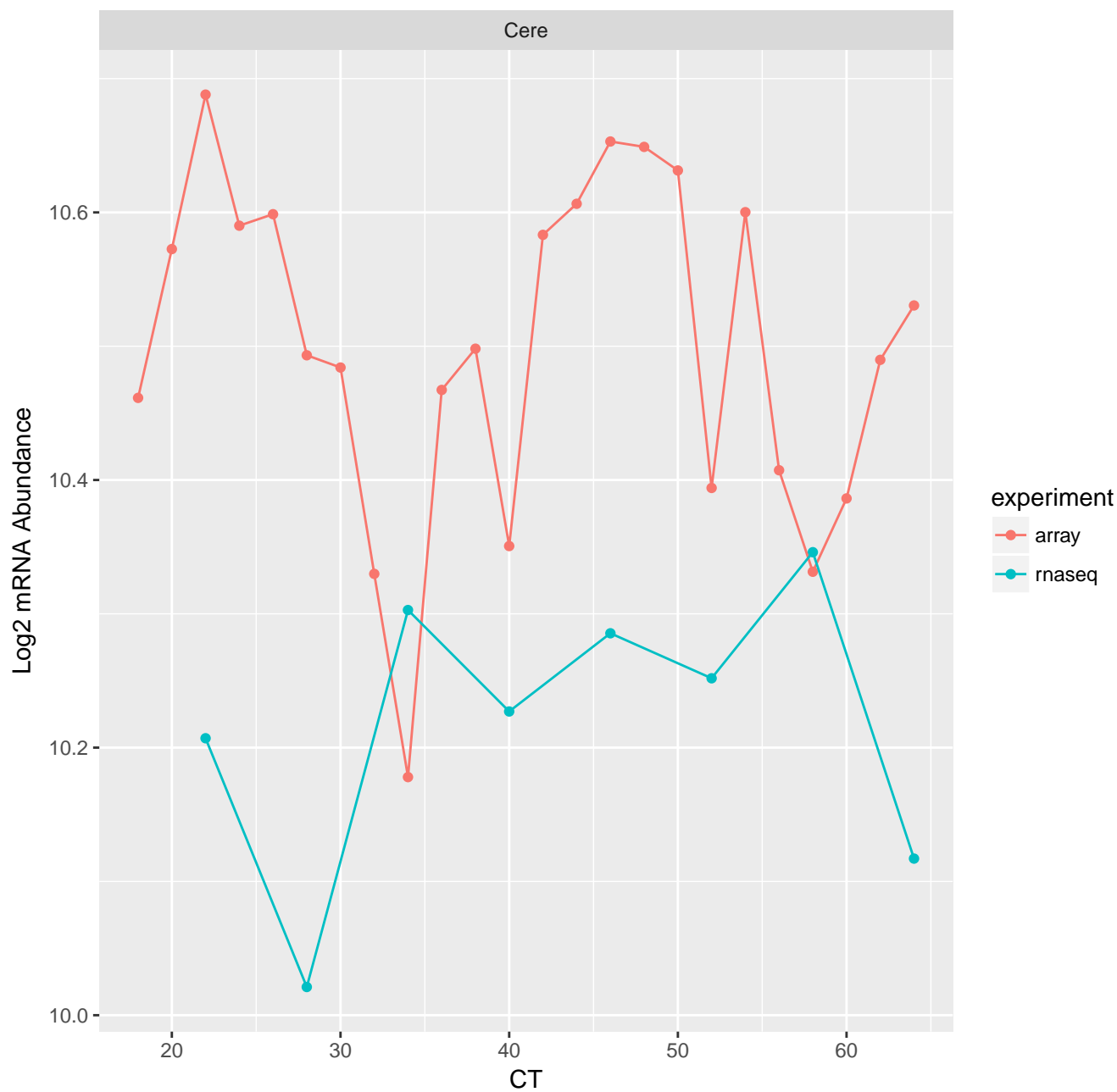
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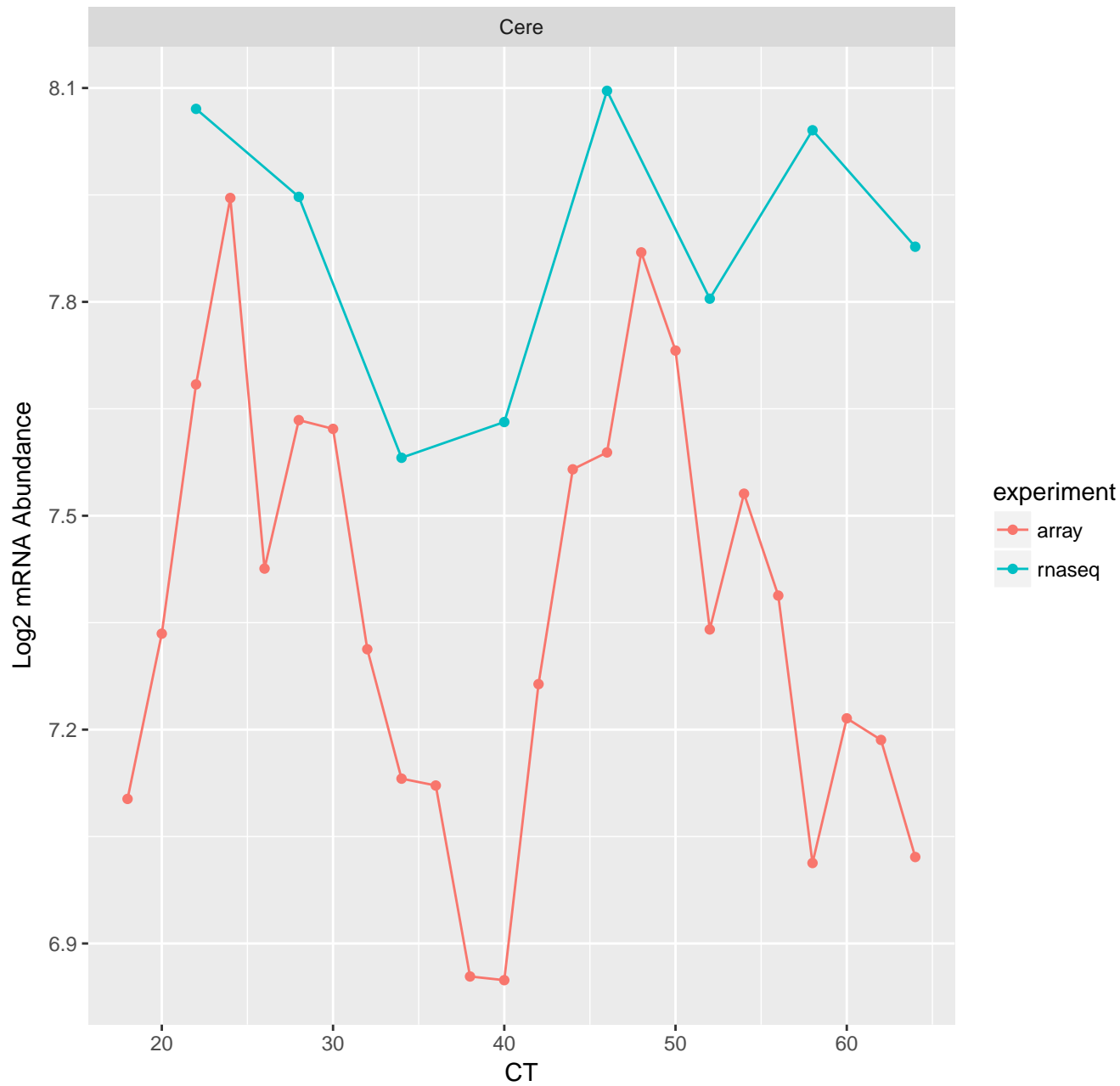
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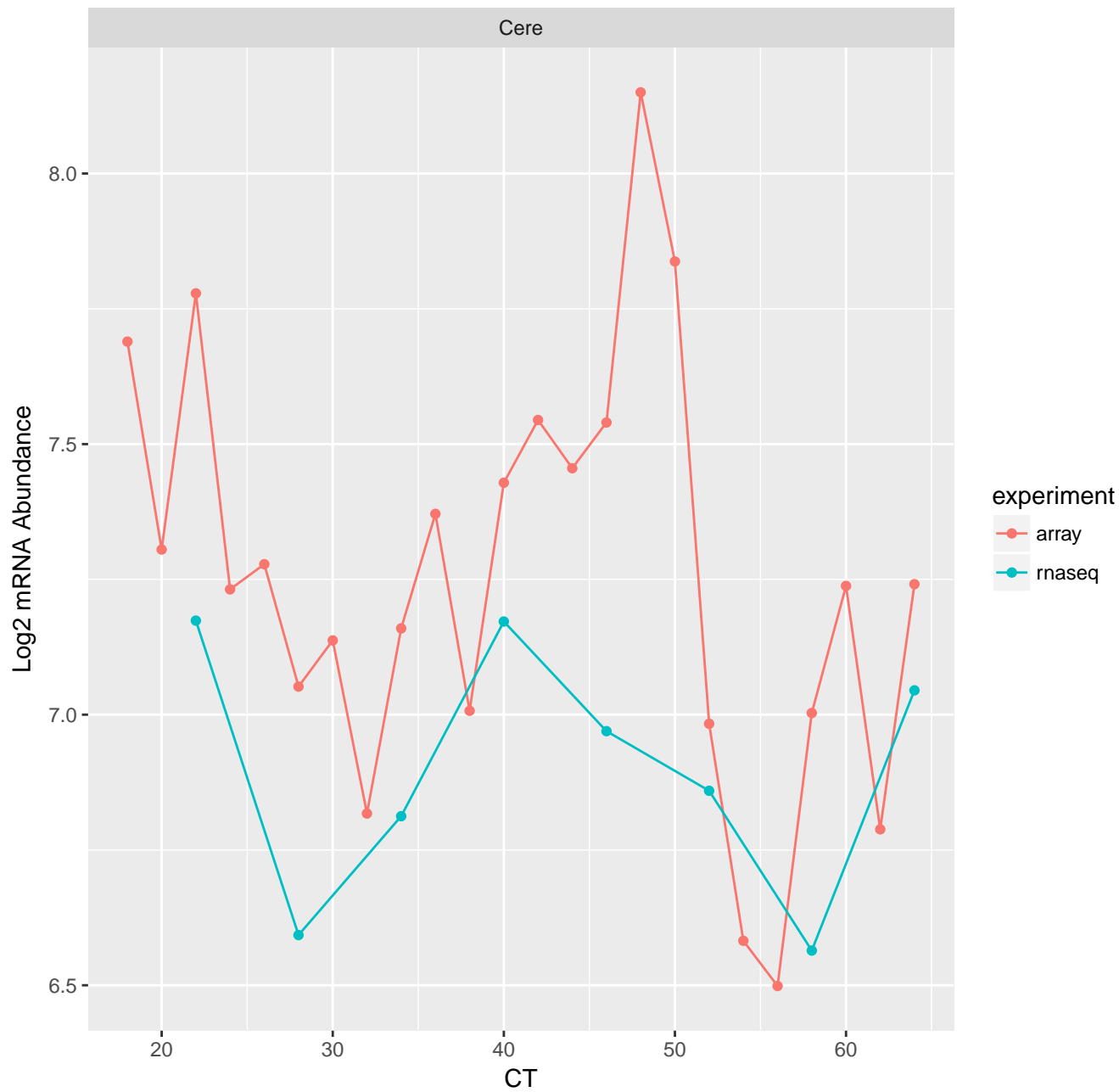
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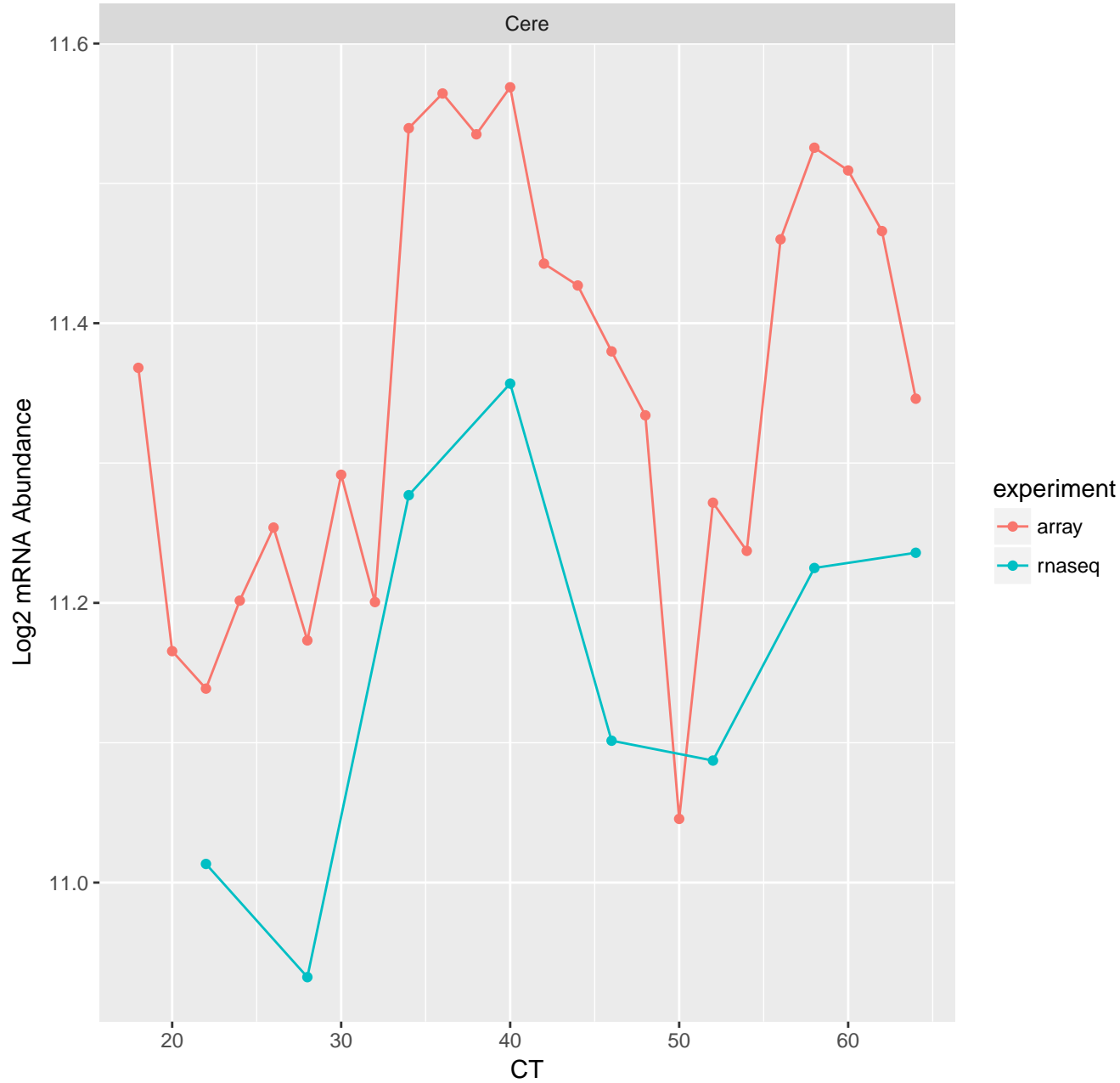
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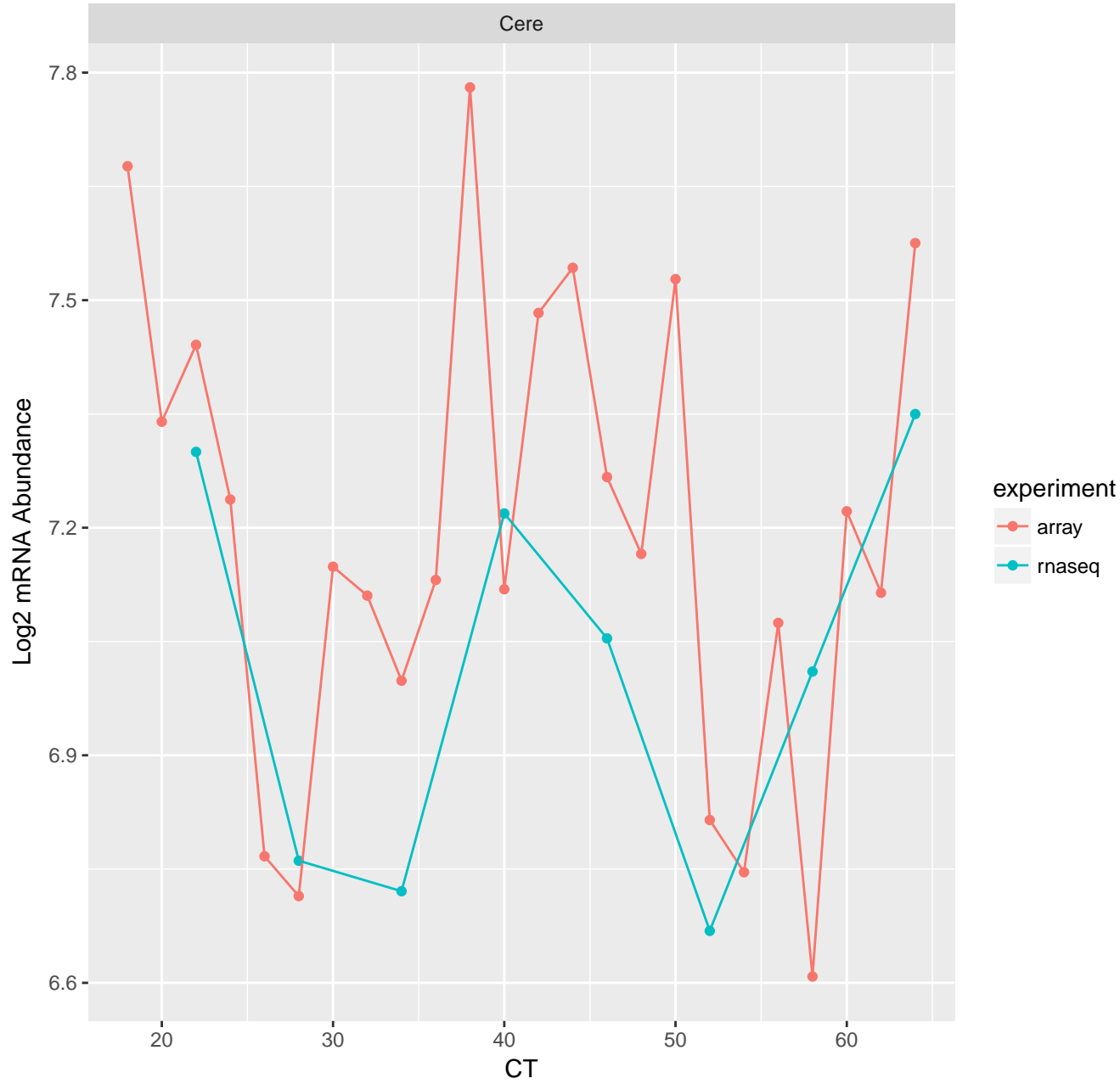
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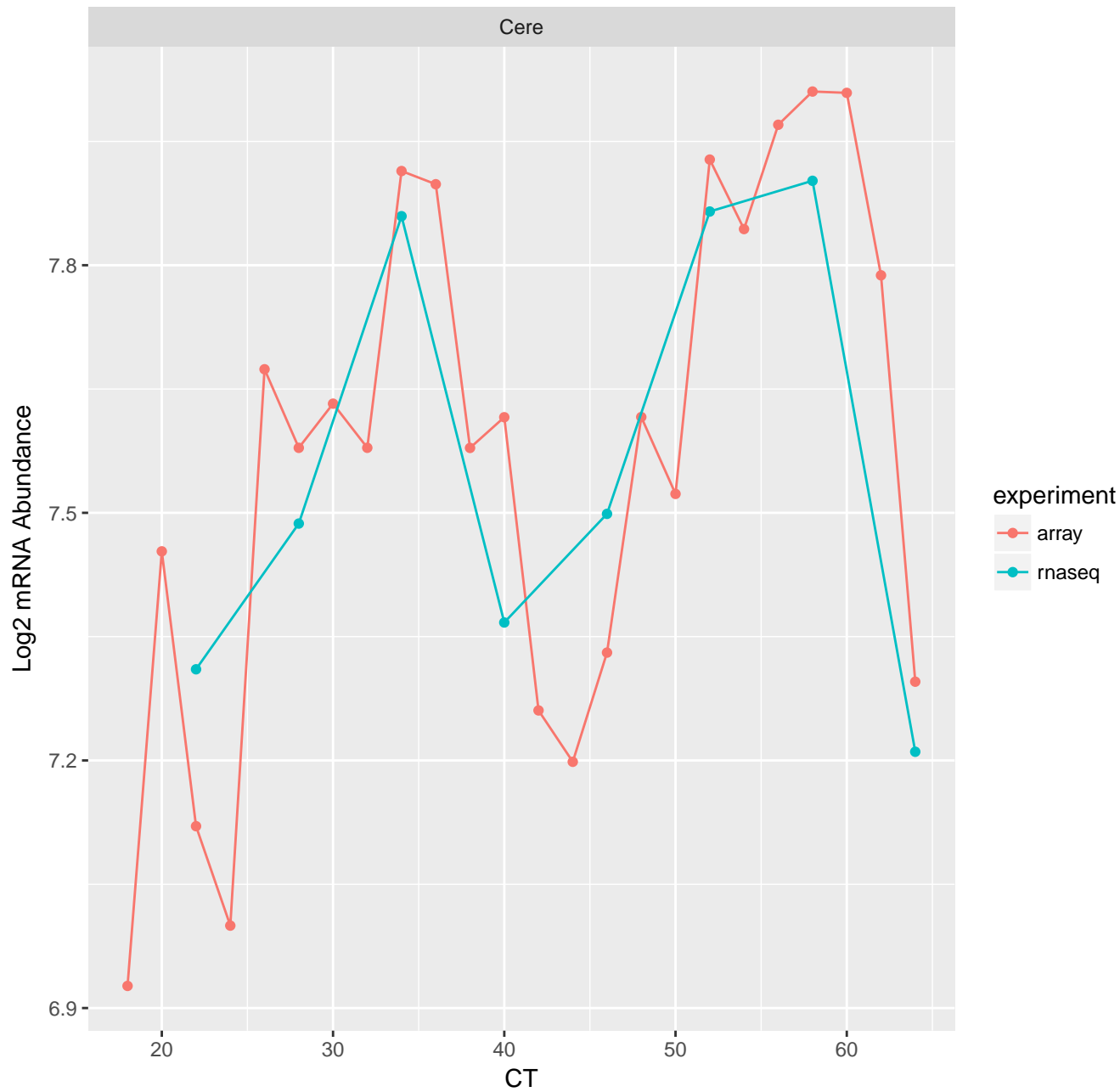
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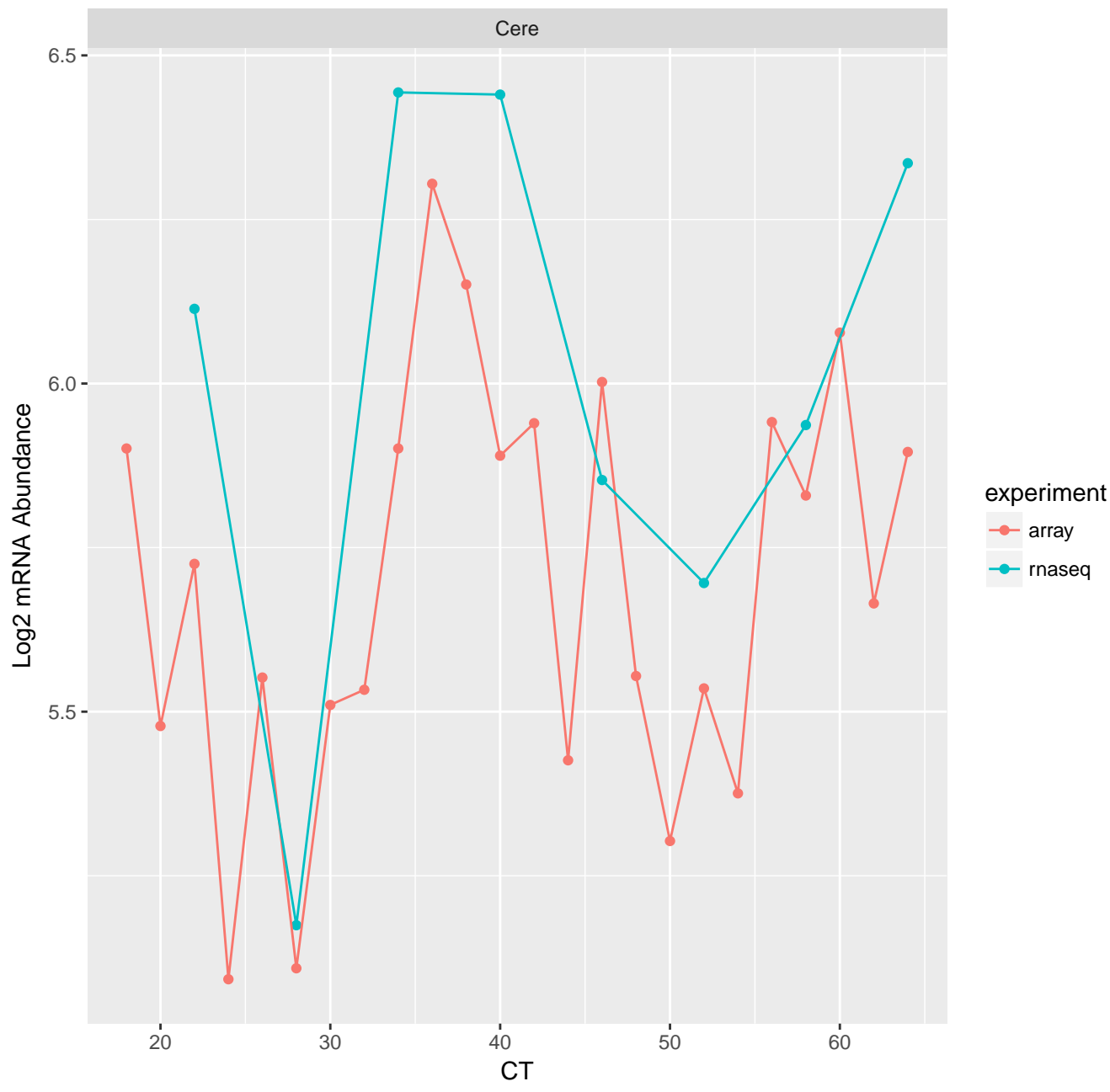
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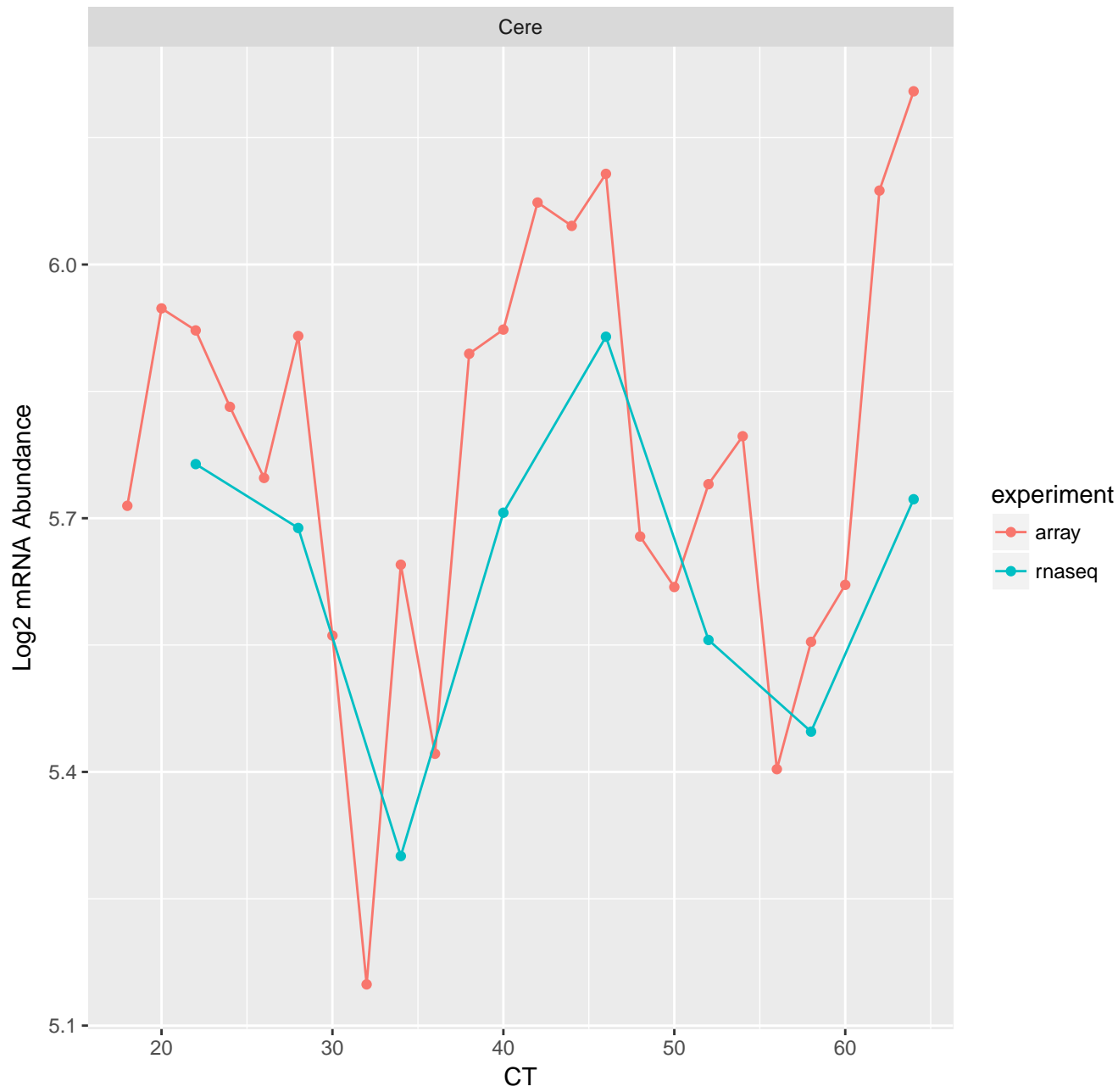
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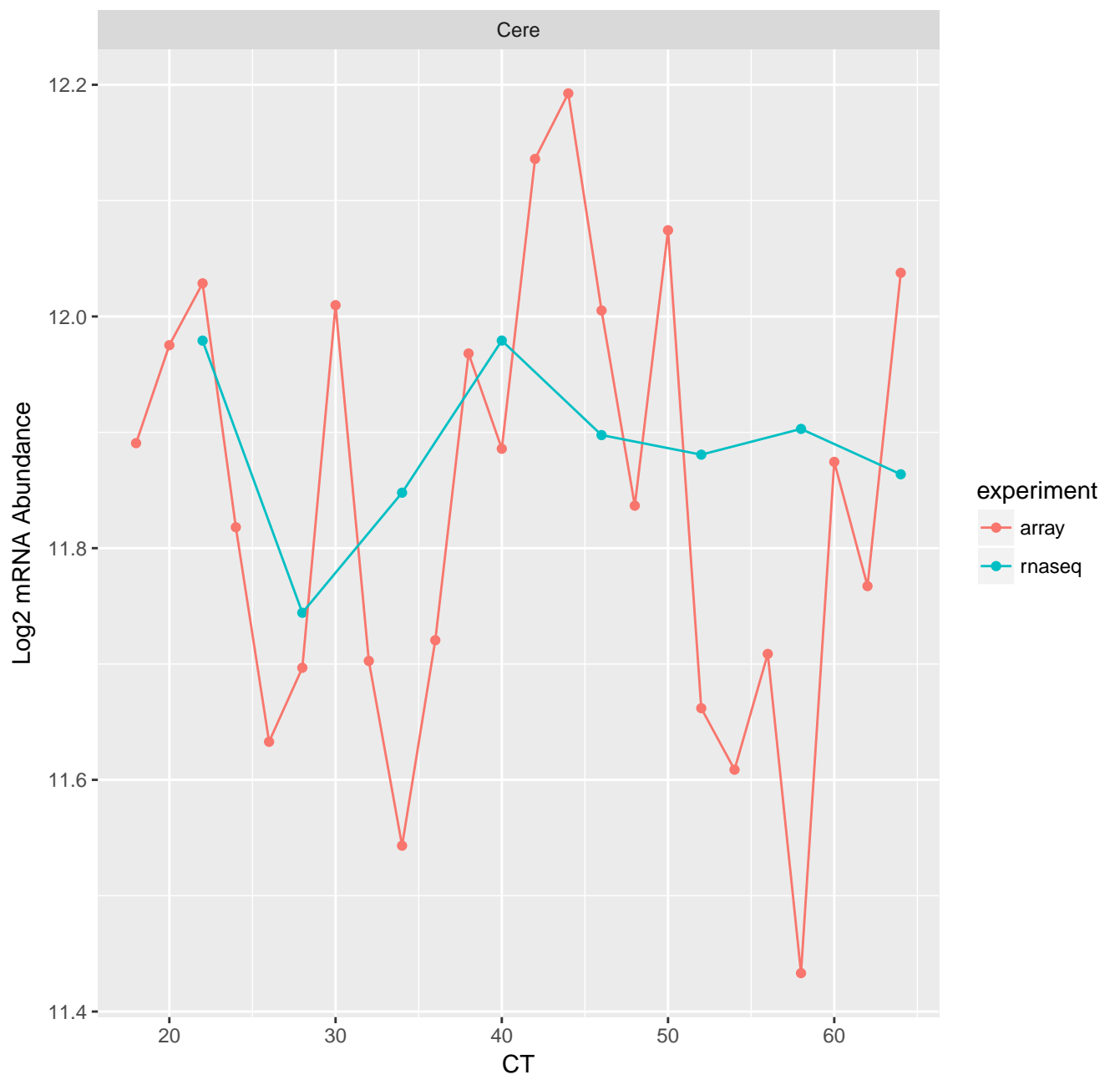
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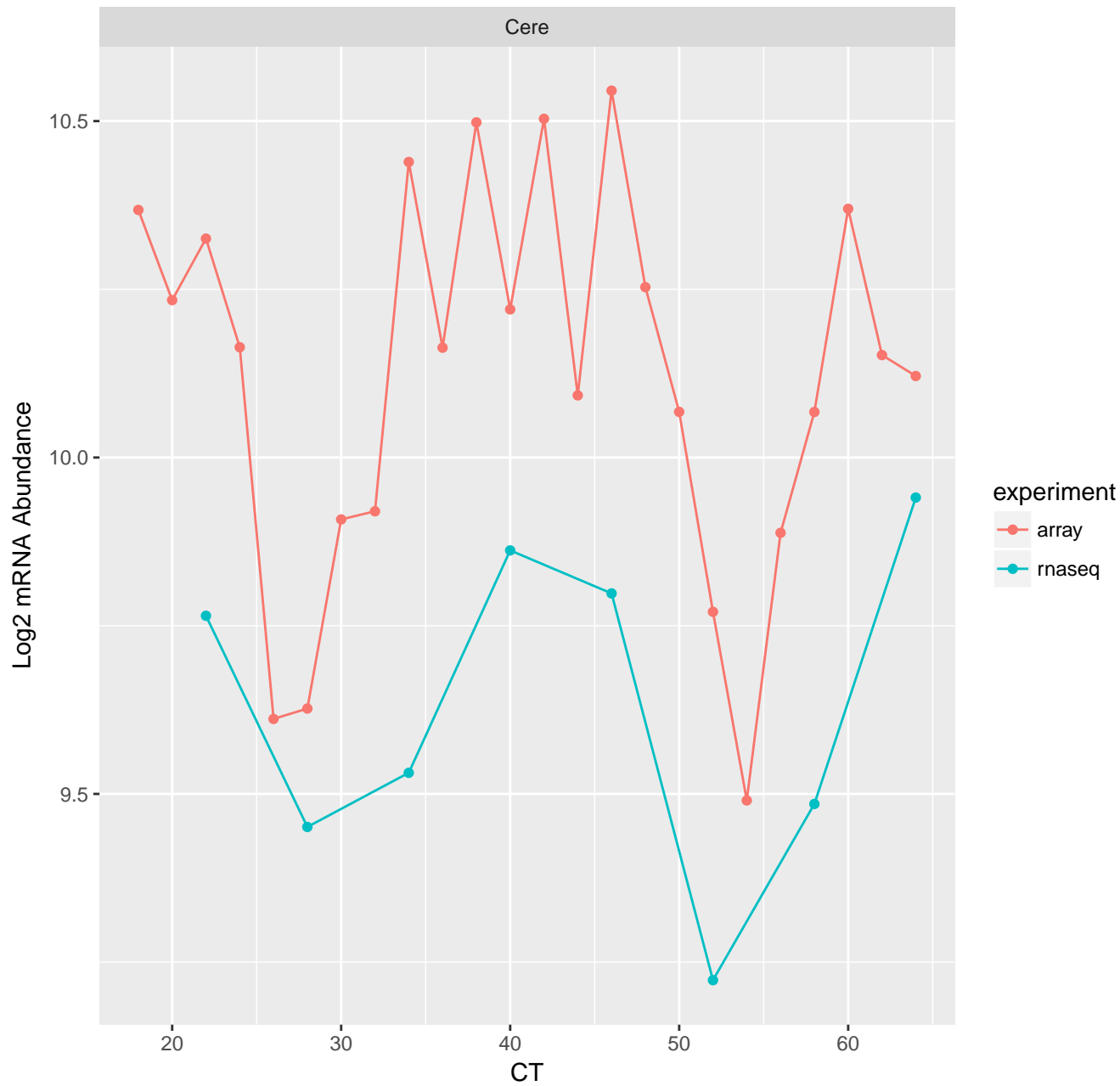
Edn1



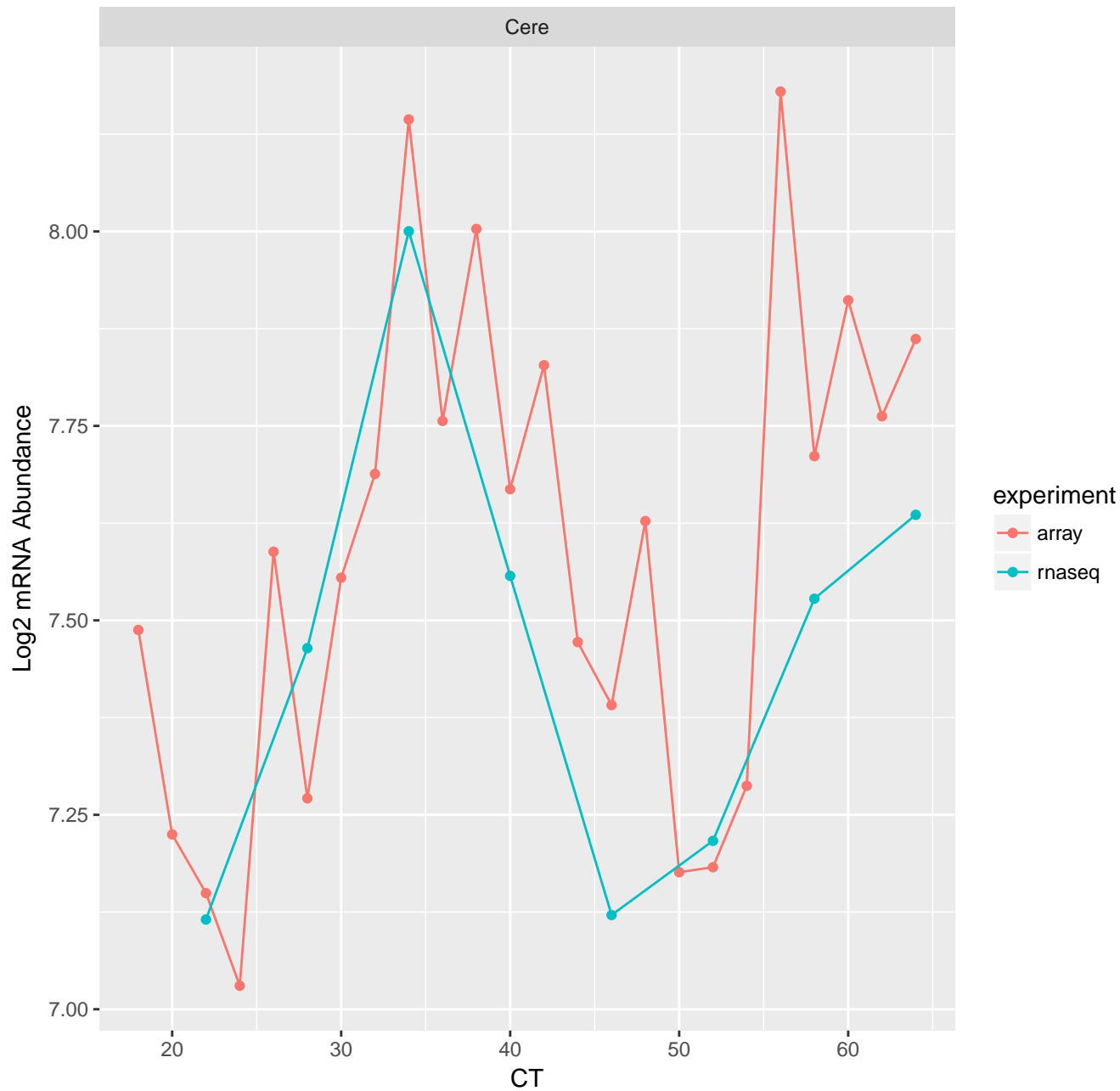
Zscan26



Cry1



Ifit3



Sptssb

