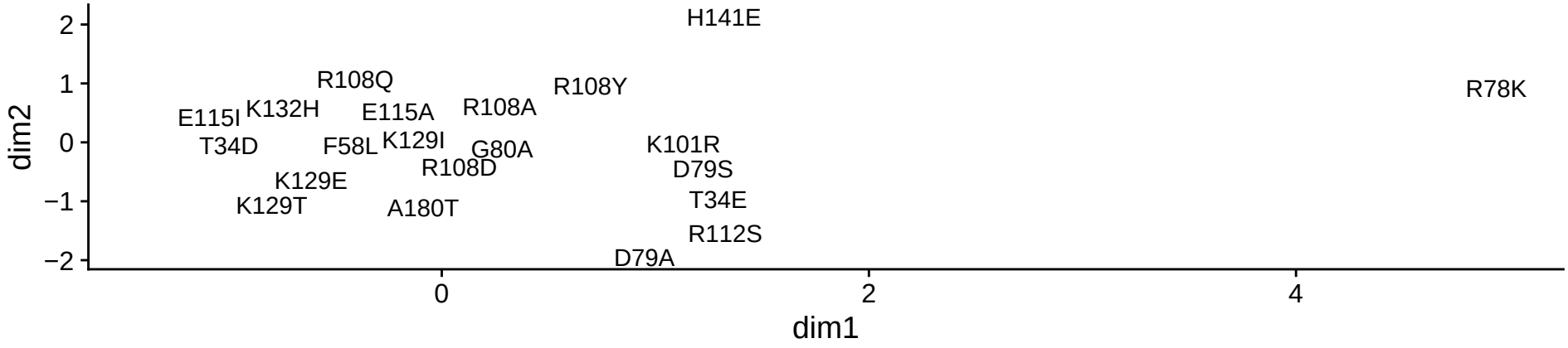
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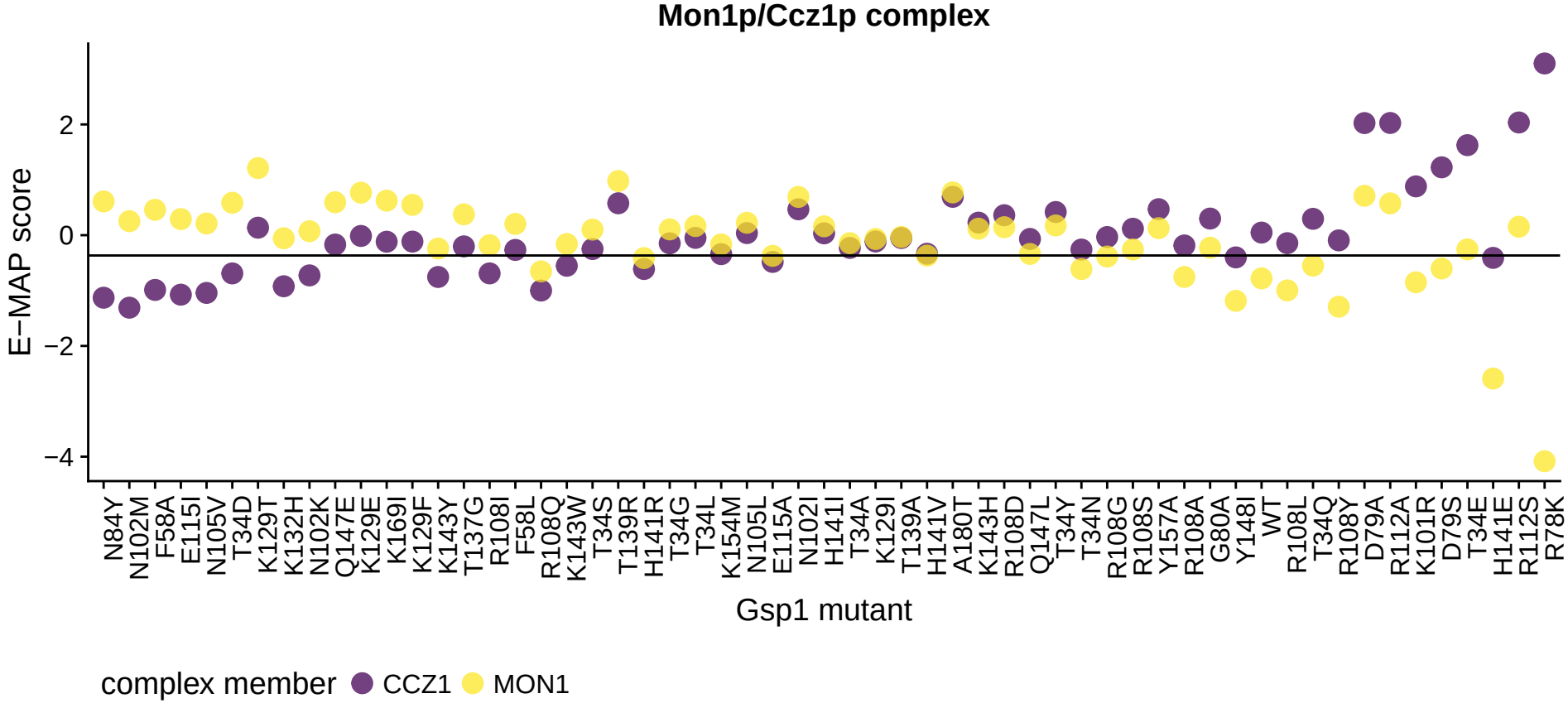
R108Q
R108Y
R78K
E115I
K132H
E115A
R108A
T34D
F58L
K129I
G80A
K101R
R108D
D79S
K129E
K129T
A180T
T34E
R112S
D79A
dim2
dim1
2
1
0
−1
−2
0
2
4


Mon1p/Ccz1p complex
E−MAP score
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0
−2
−4
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N102M
F58A
E115I
N105V
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K129T
K132H
N102K
Q147E
K129E
K169I
K129F
K143Y
T137G
R108I
F58L
R108Q
K143W
T34S
T139R
H141R
T34G
T34L
K154M
N105L
E115A
N102I
H141I
T34A
K129I
T139A
H141V
A180T
K143H
R108D
Q147L
T34Y
T34N
R108G
R108S
Y157A
R108A
G80A
Y148I
WT
R108L
T34Q
R108Y
D79A
R112A
K101R
D79S
T34E
H141E
R112S
R78K
Gsp1 mutant
complex member
CCZ1
MON1