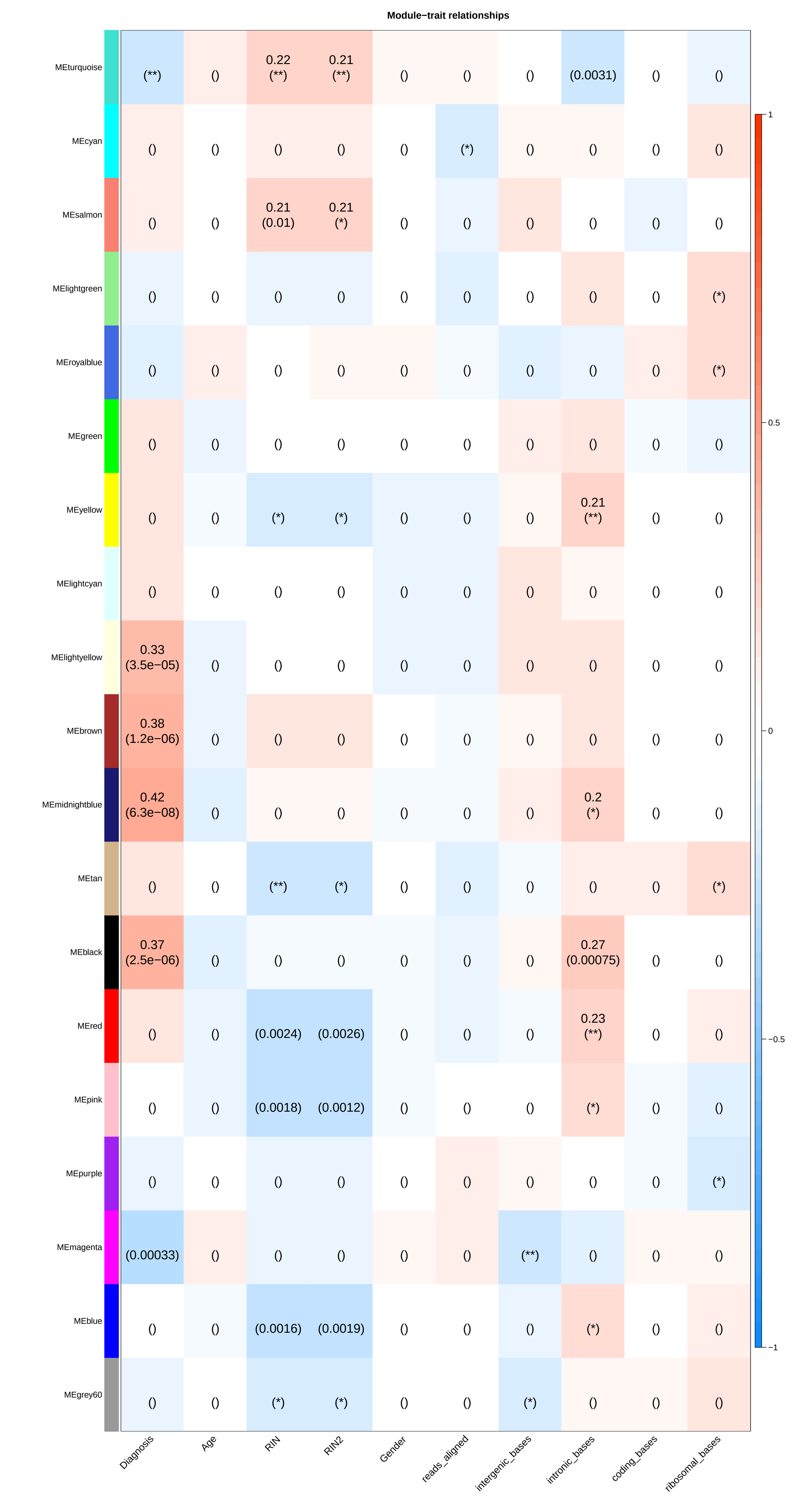
Module−trait relationships
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MEcyan
MEsalmon
MElightgreen
MEroyalblue
MEgreen
MEyellow
MElightcyan
MElightyellow
MEbrown
MEmidnightblue
MEtan
MEblack
MEred
MEpink
MEpurple
MEmagenta
MEblue
MEgrey60
Diagnosis
Age
RIN
RIN2
Gender
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intergenic_bases
intronic_bases
coding_bases
ribosomal_bases
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0
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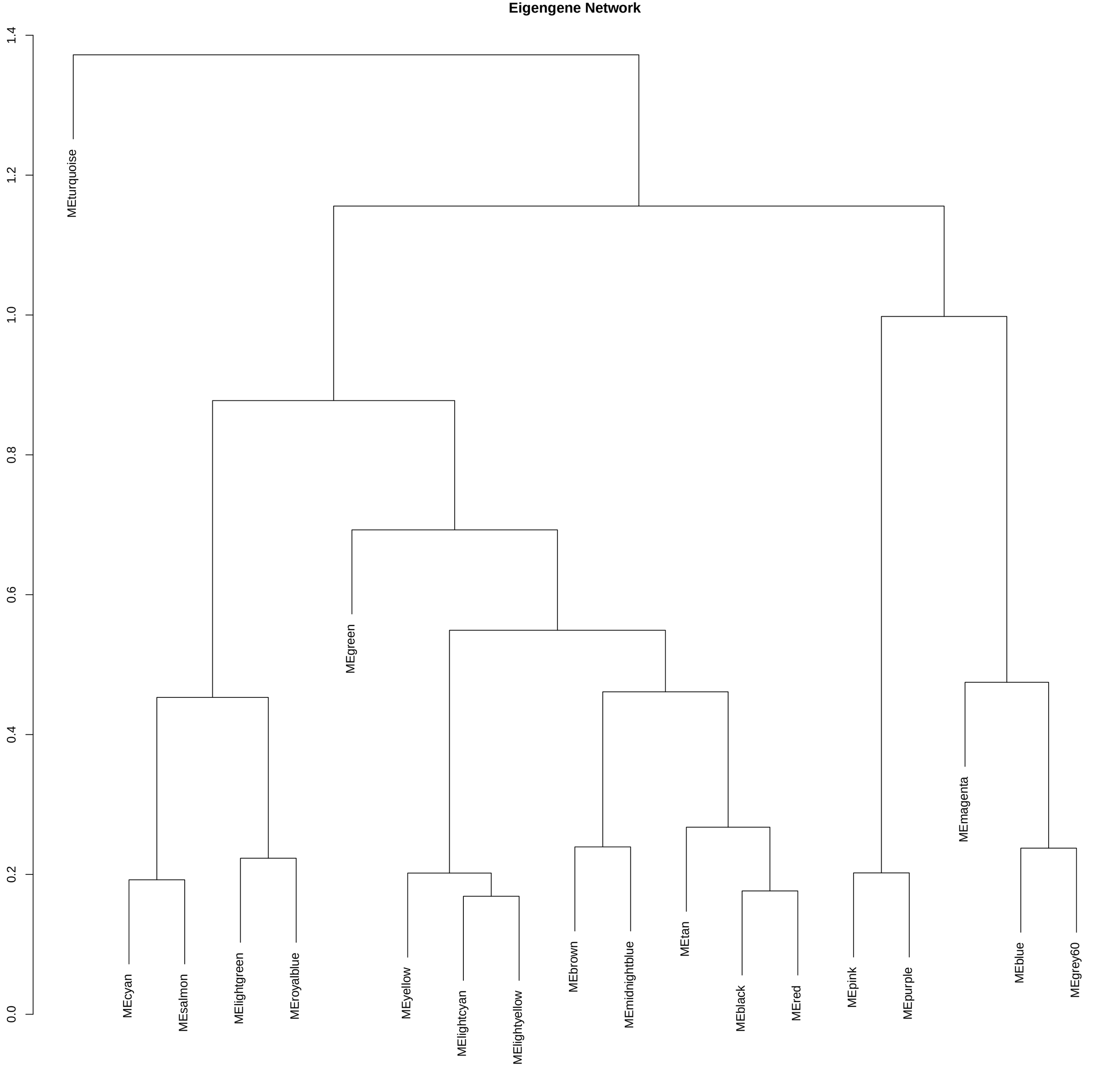
−1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0.2
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MEcyan
MEsalmon
MElightgreen
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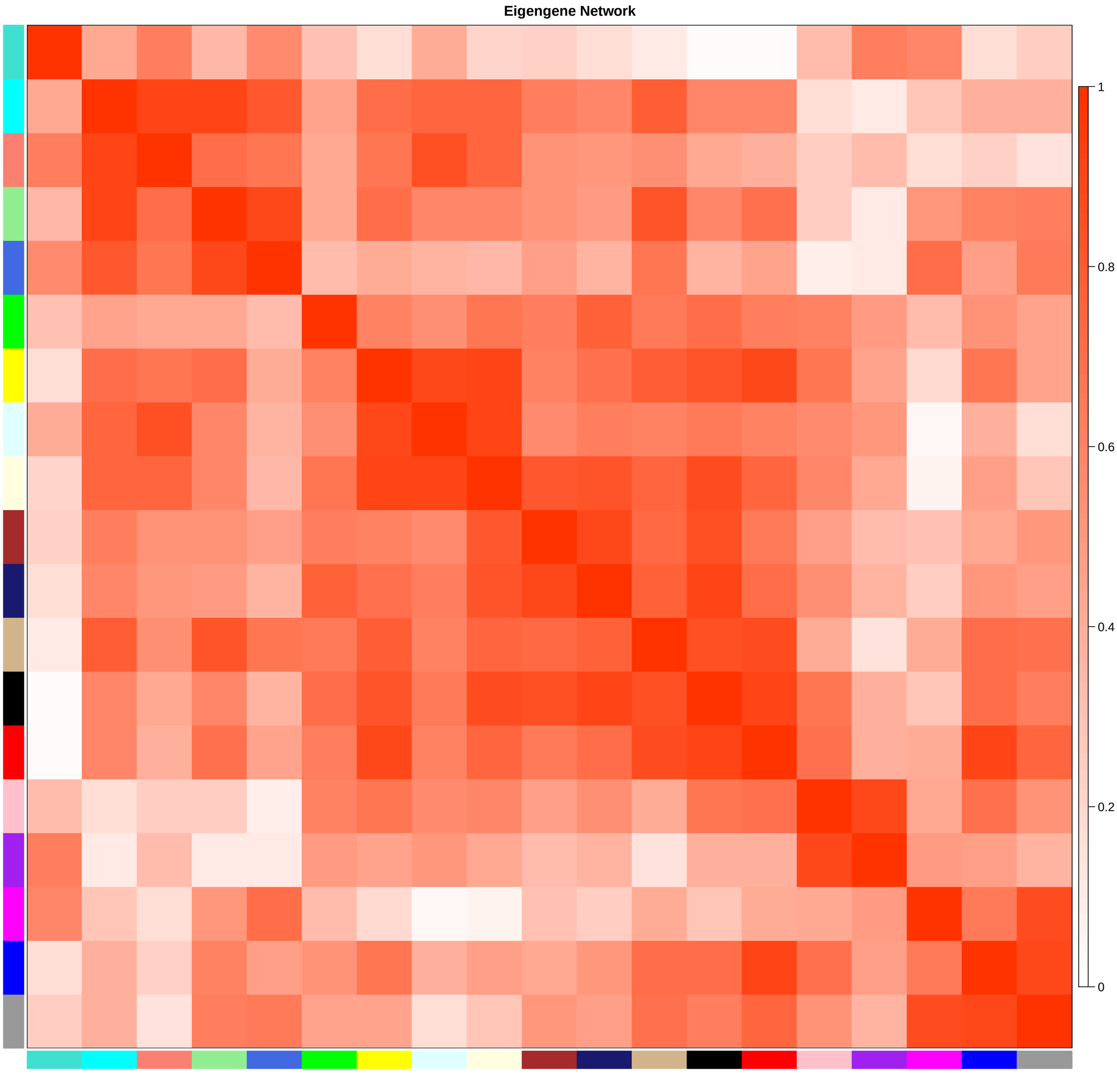
Eigengene Network
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0.2
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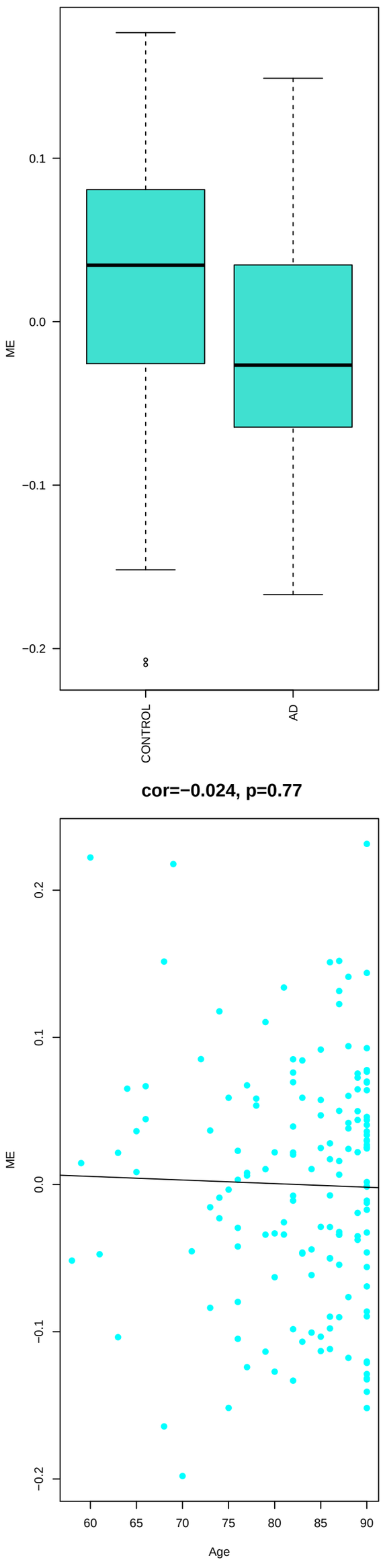
MEturquoise



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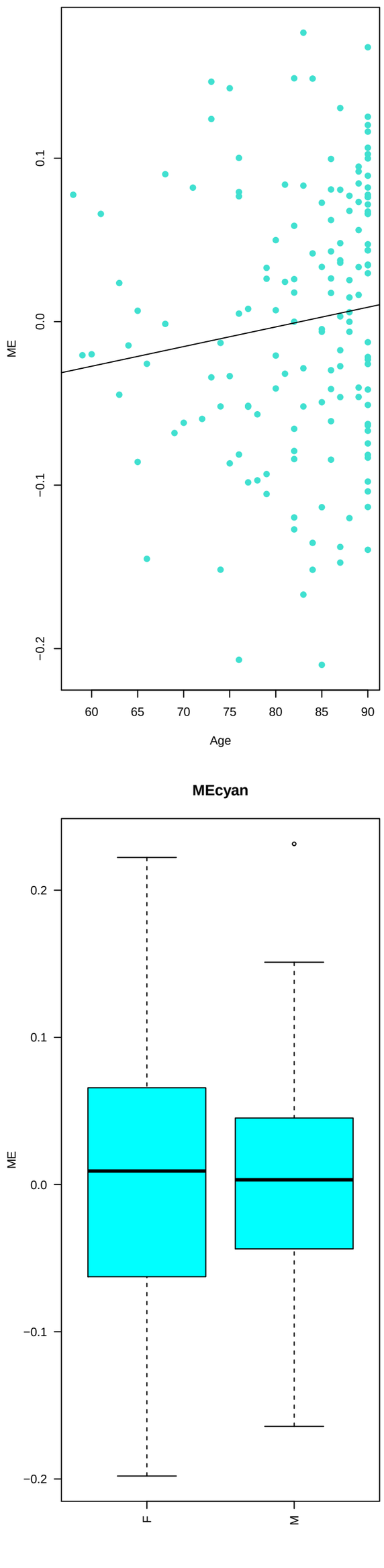

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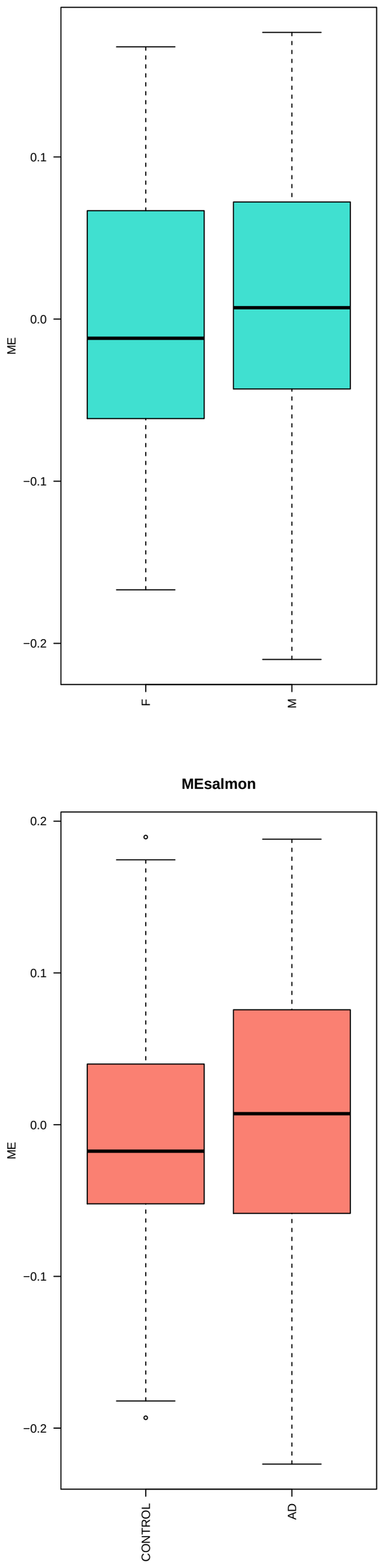
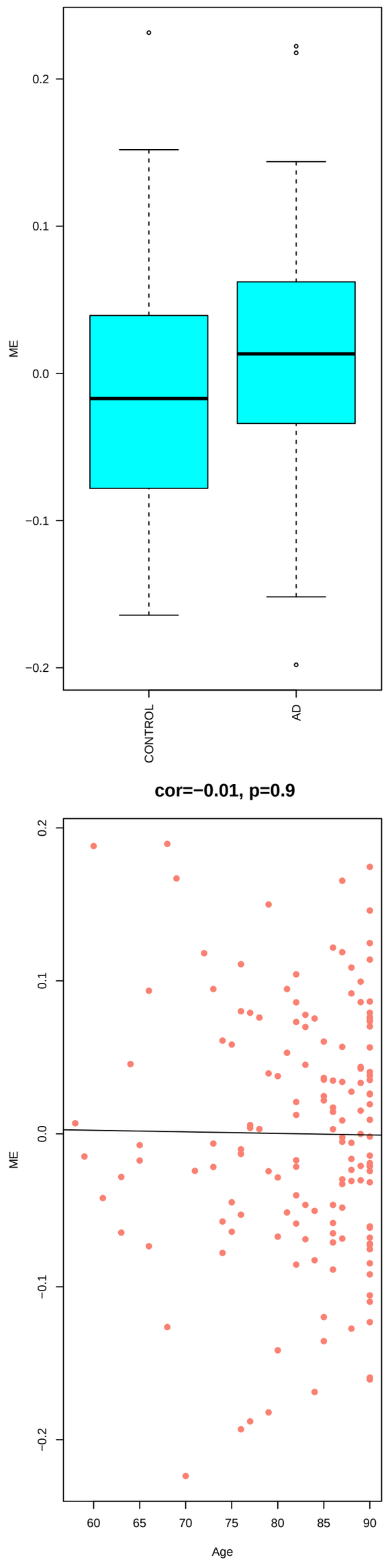


MEcyan



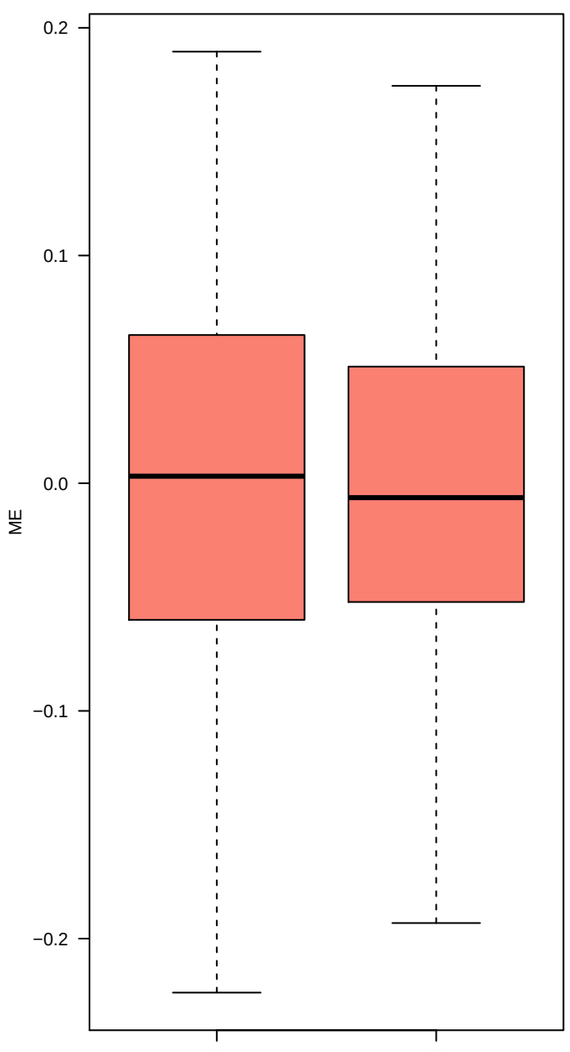
cor=−0.024, p=0.77

MEcyan

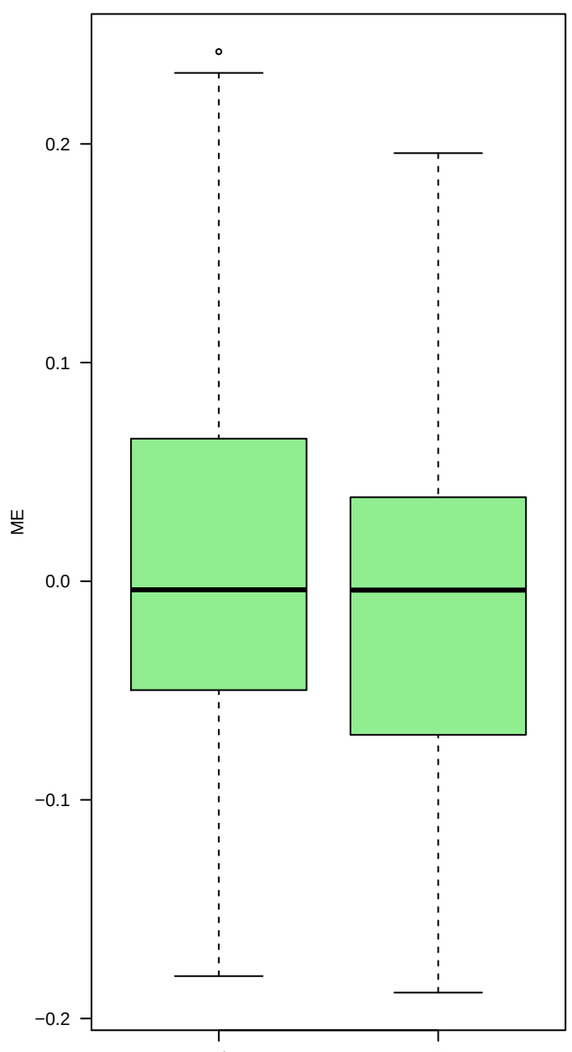
MEsalmon

cor=−0.01, p=0.9

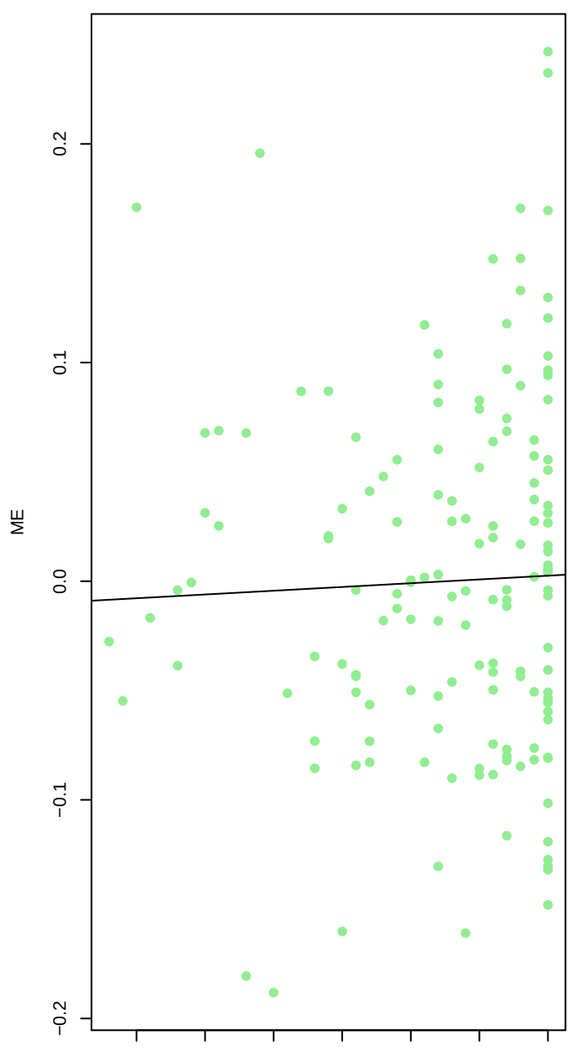
MEsalmon



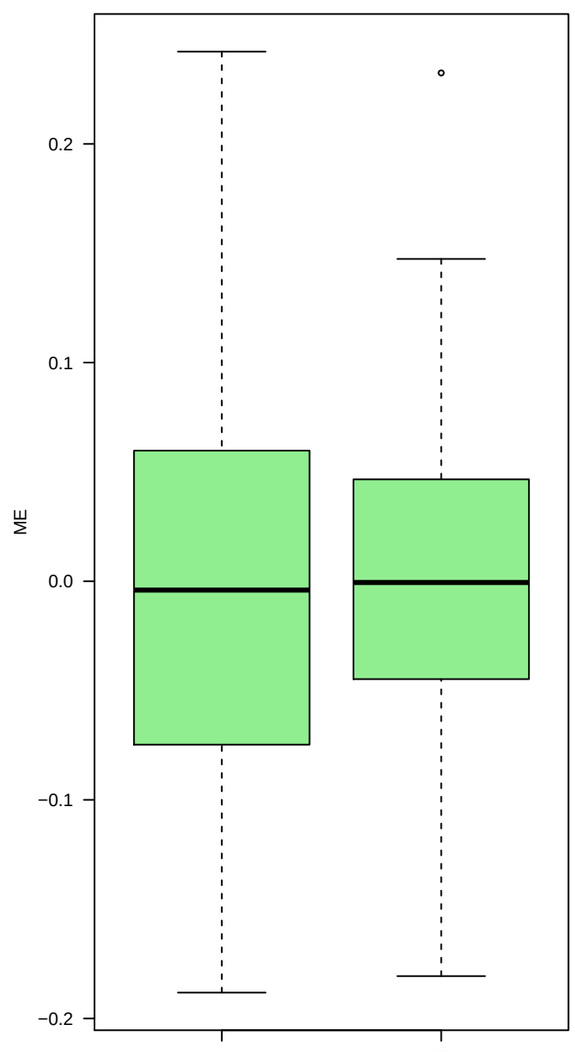
MElightgreen



cor=0.034, p=0.68



MElightgreen



MEroyalblue

cor=0.11, p=0.18

MEroyalblue

MEgreen

cor=−0.12, p=0.14

MEgreen

MEyellow

cor=−0.066, p=0.42

MEyellow

MElightcyan

cor=−0.021, p=0.8

MElightcyan

MElightyellow

cor=−0.11, p=0.18

MElightyellow

MEbrown

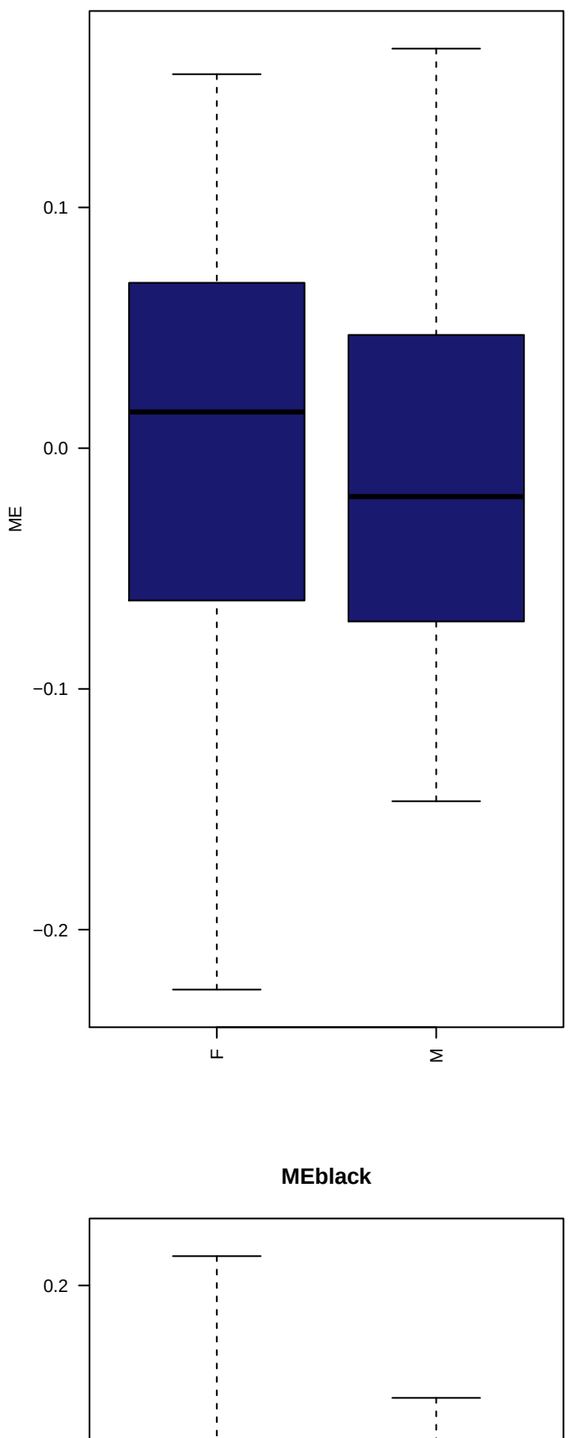
cor=−0.12, p=0.14

MEbrown

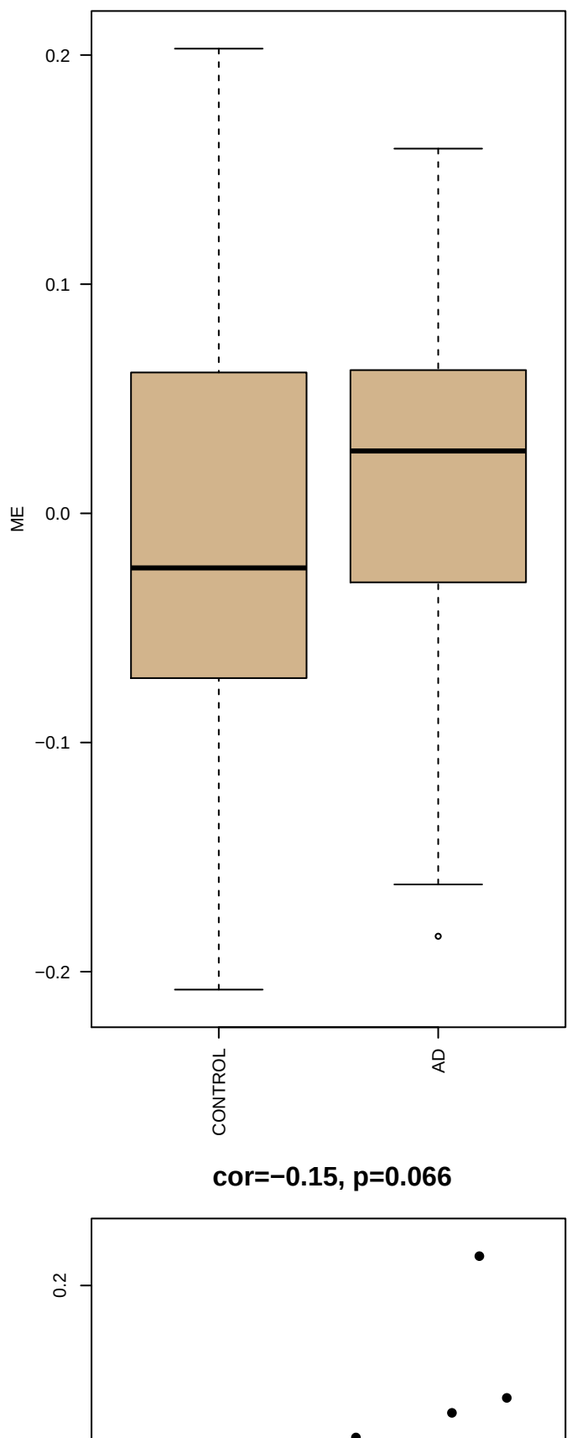
MEmidnightblue

cor=−0.15, p=0.066

MEmidnightblue



MEtan



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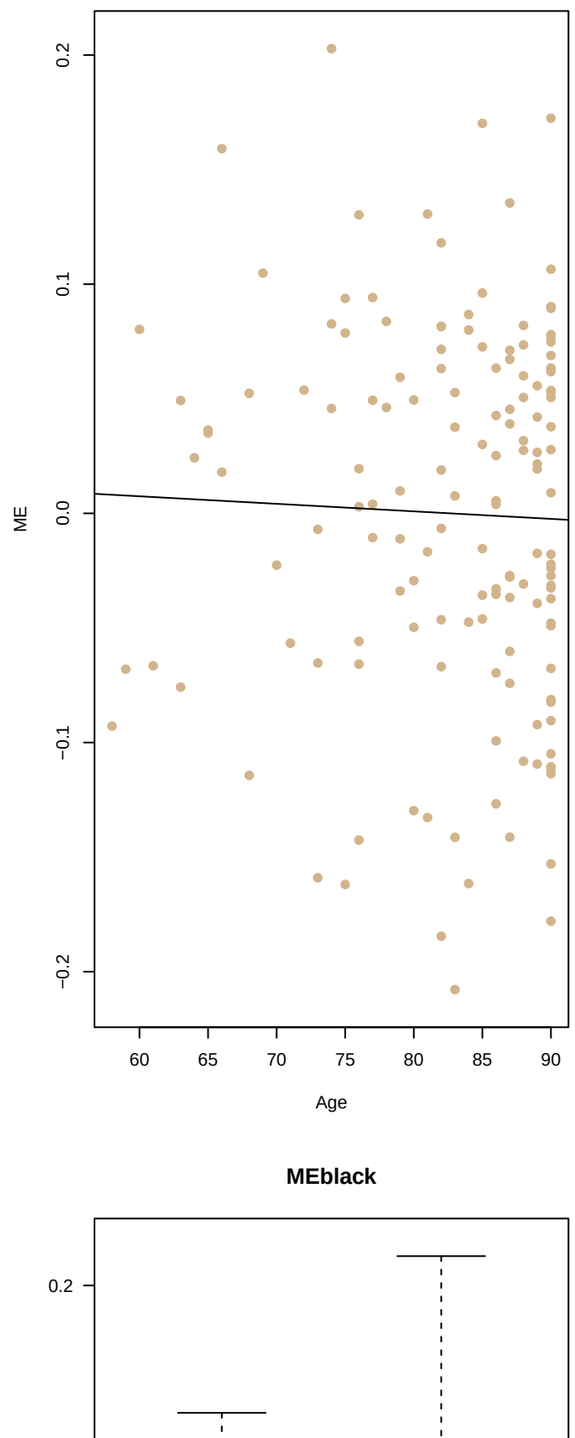
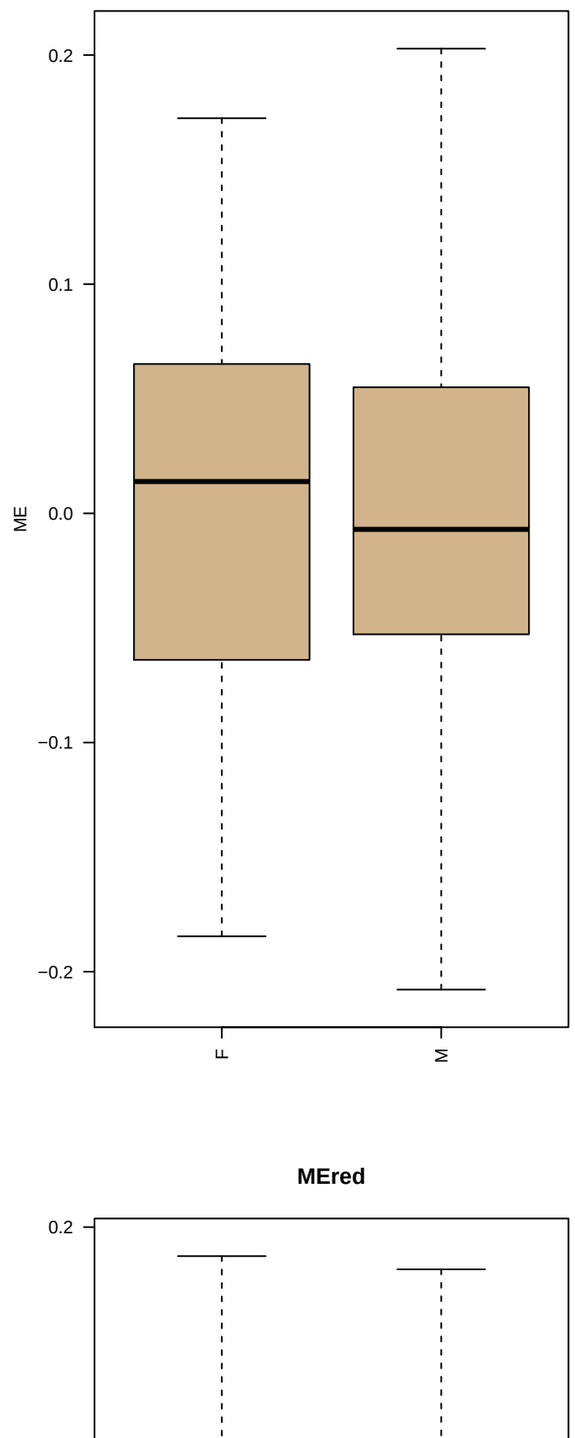


MEtan



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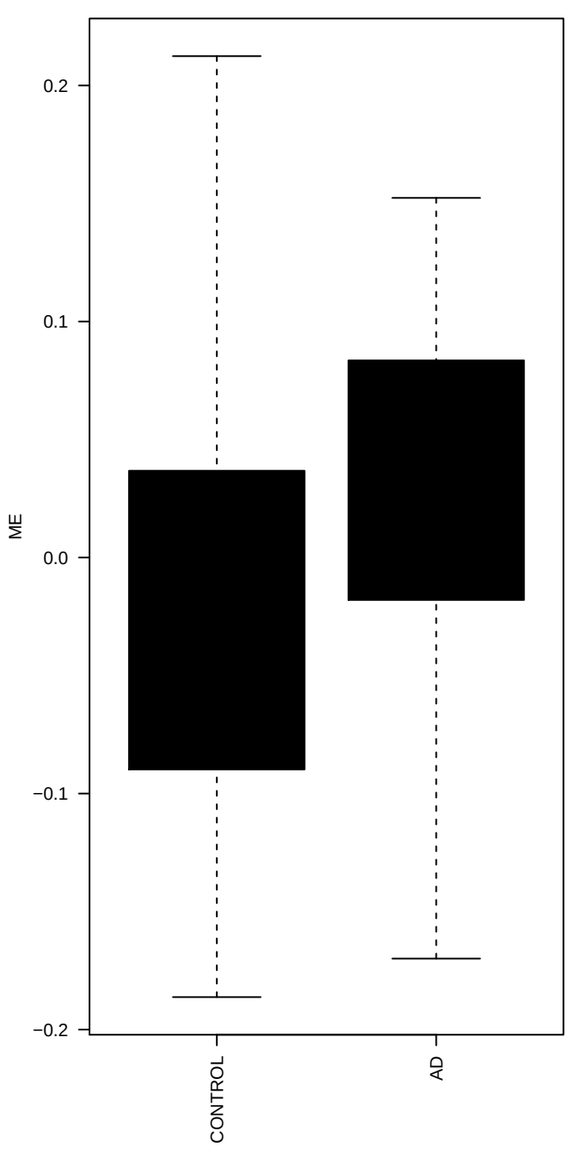

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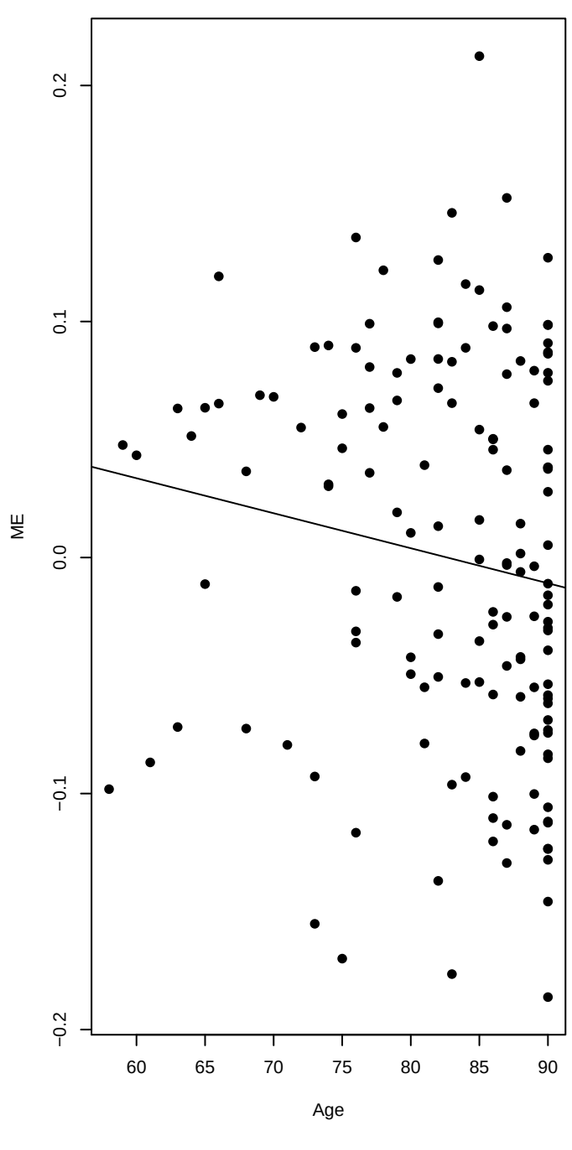

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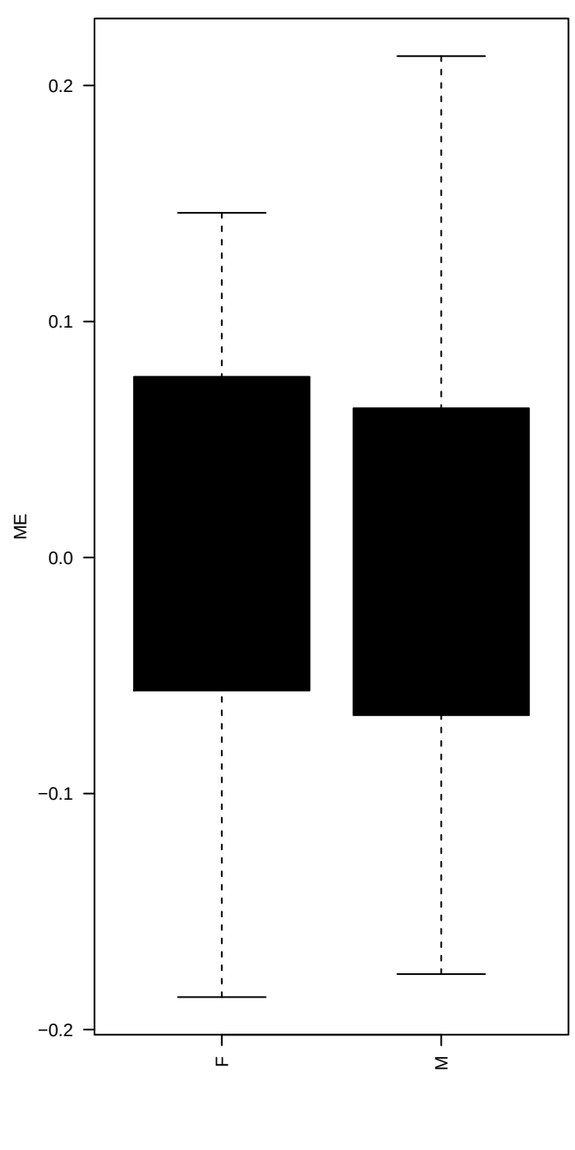
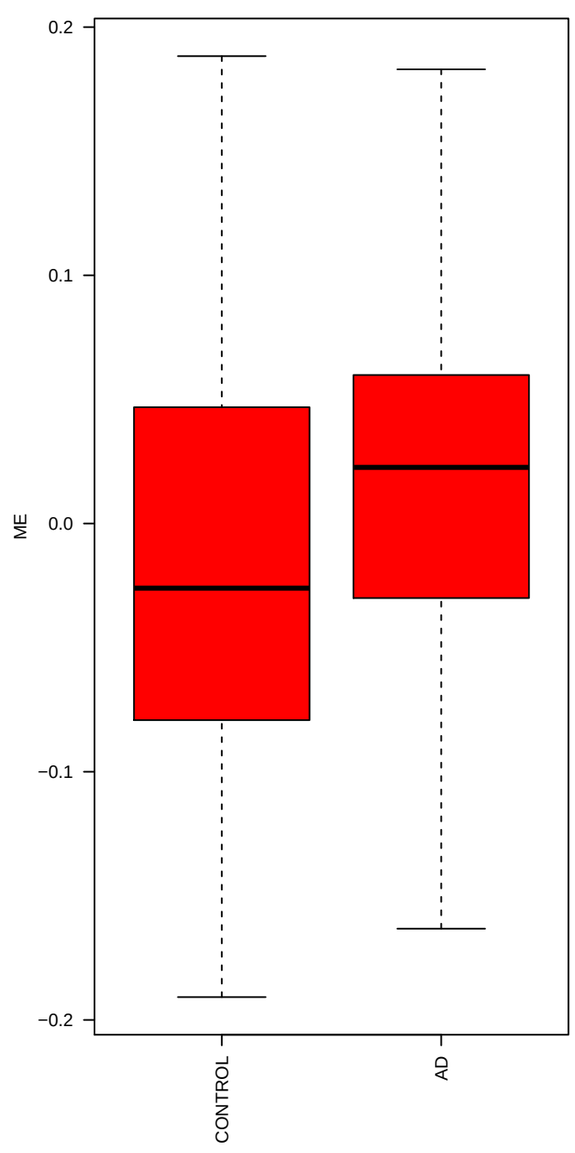
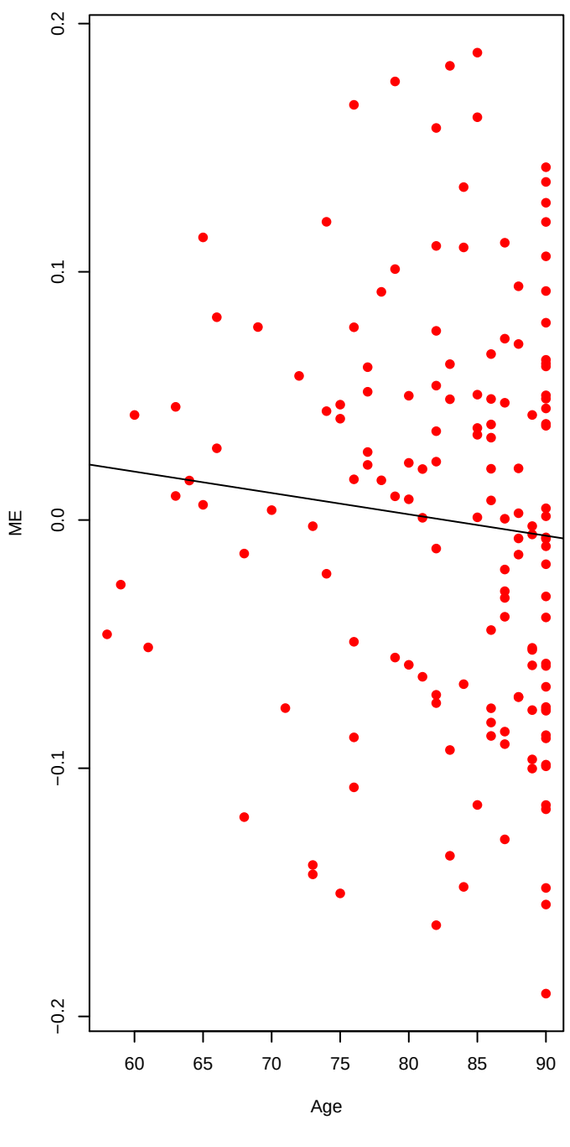


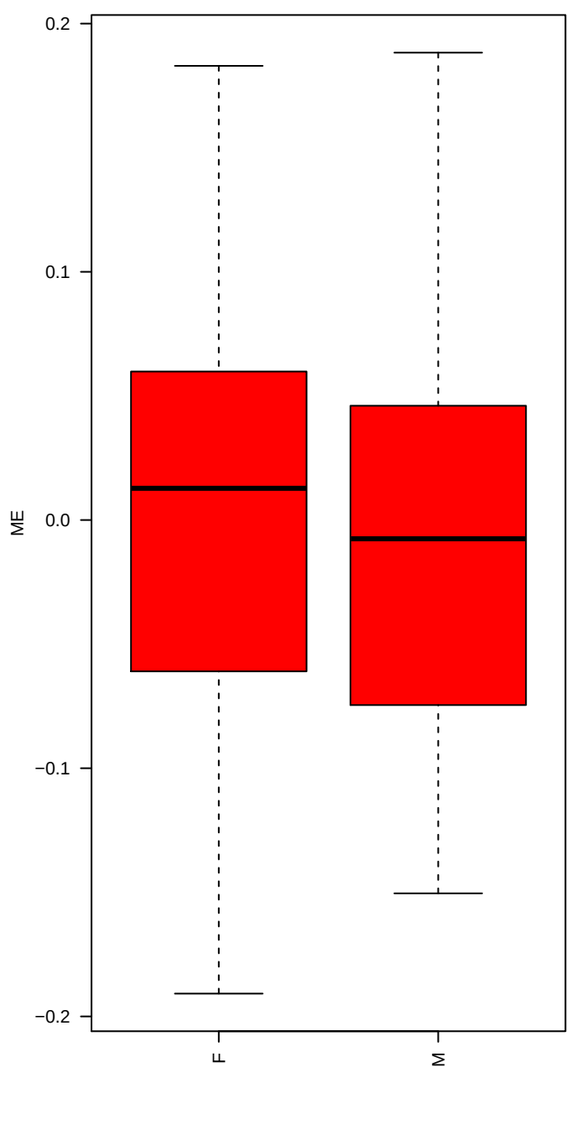
MEred



cor=−0.085, p=0.3



MEred



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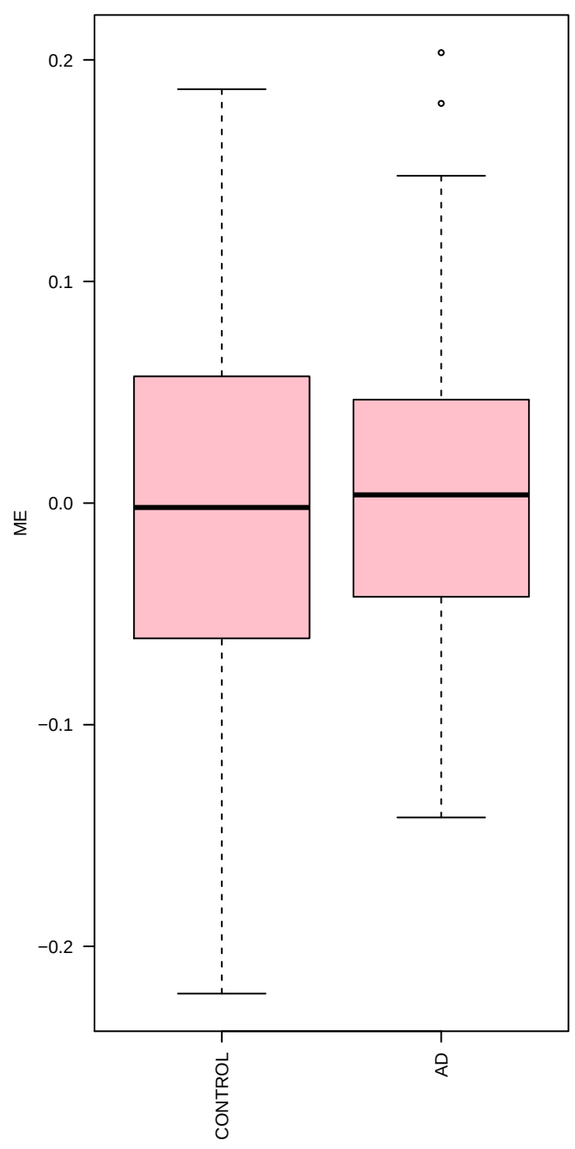

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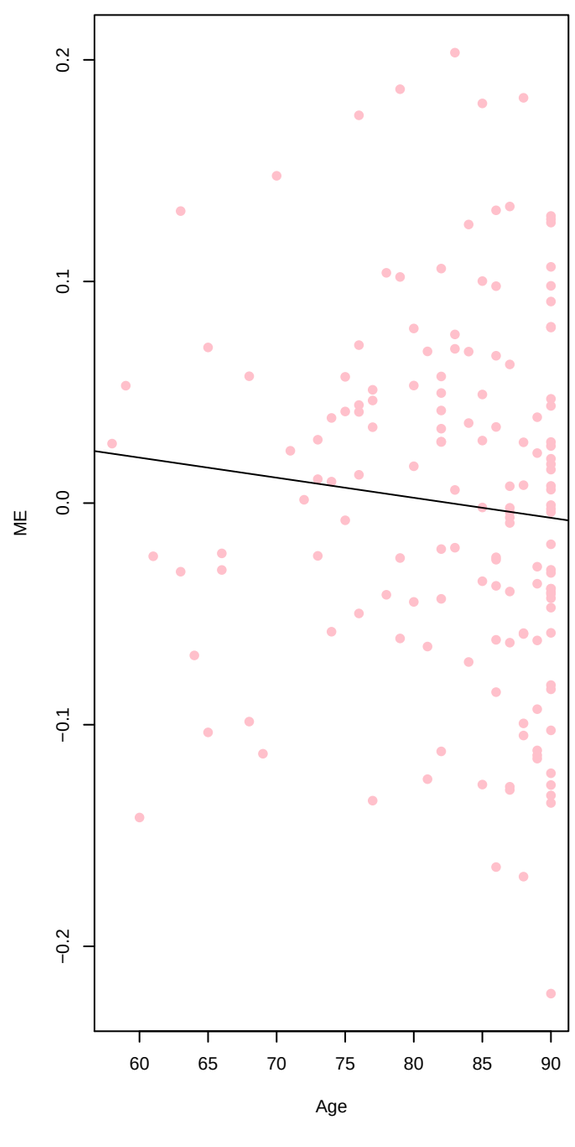
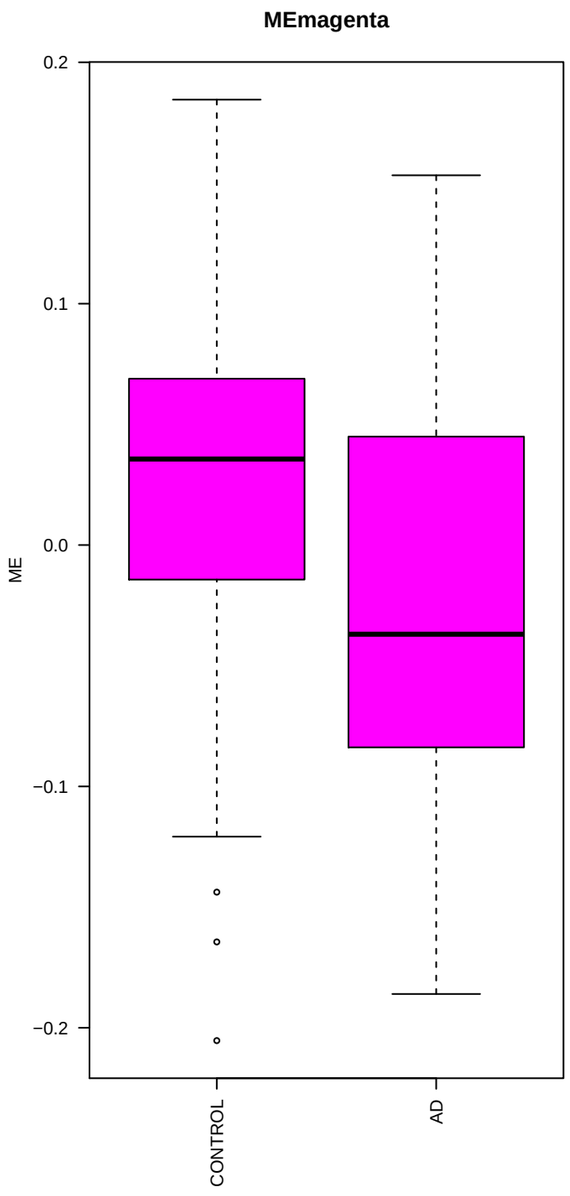


MEpurple

MEpurple

cor=−0.014, p=0.86

MEpurple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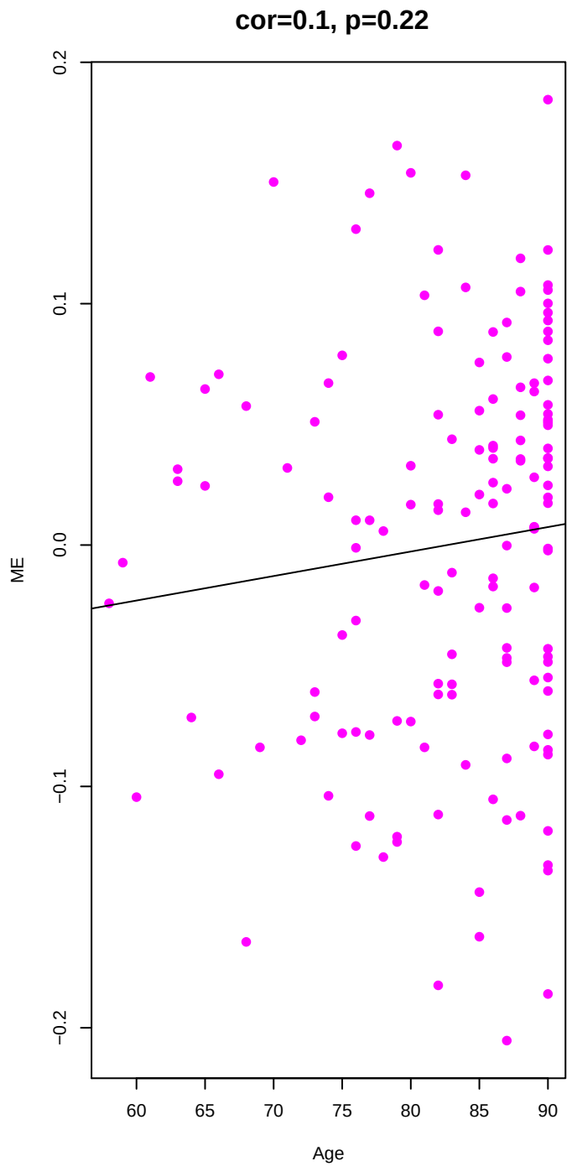
0.0
−0.1
−0.2
CONTROL
AD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−0.2
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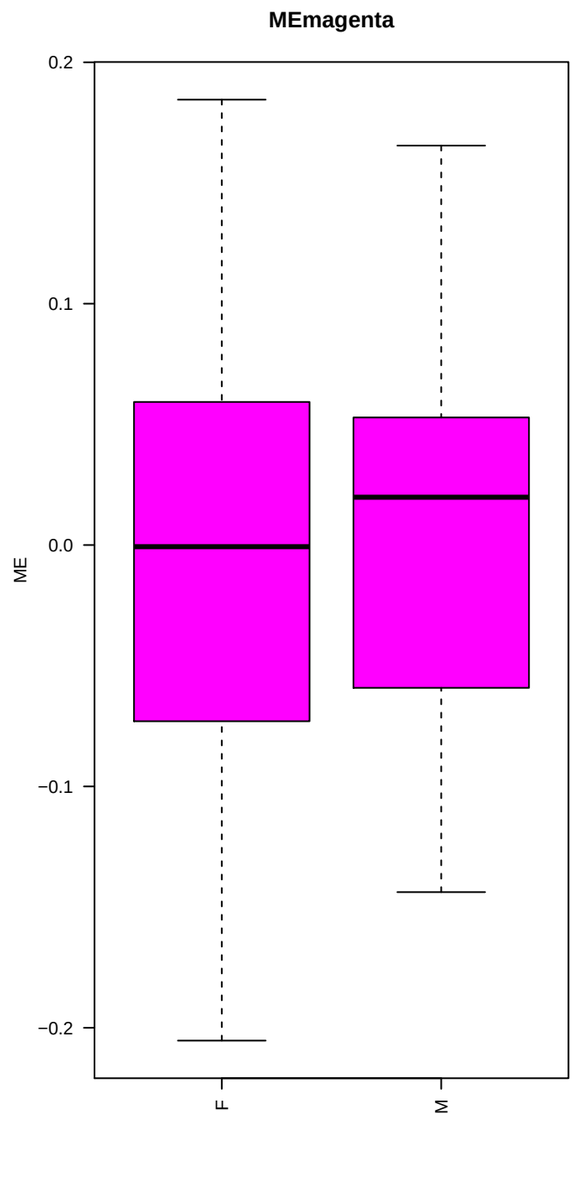
75
80
85
90
Age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0.1
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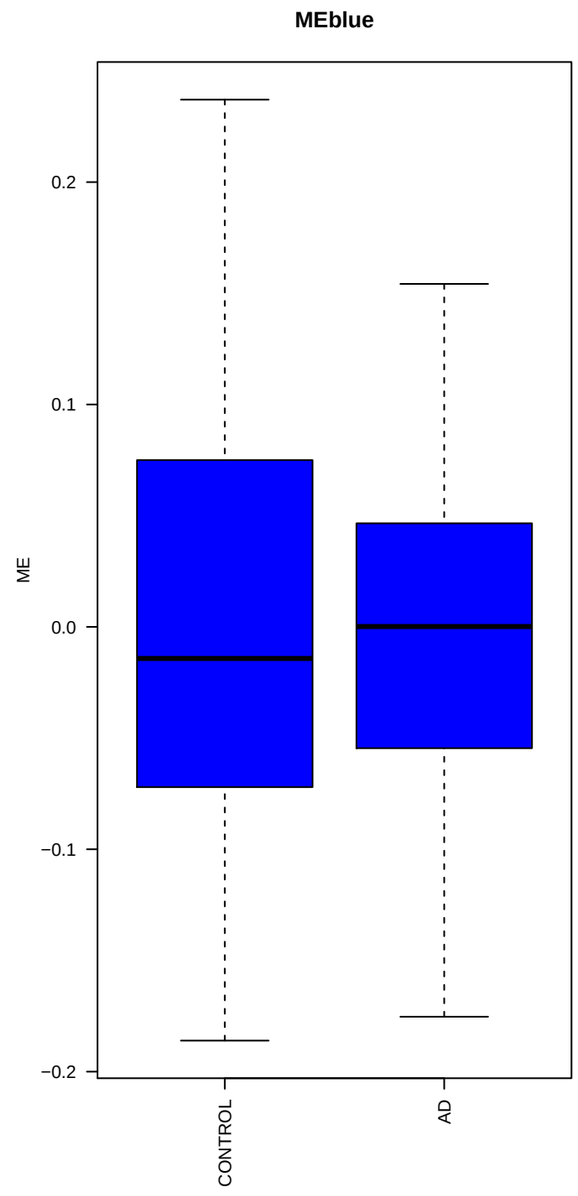
−0.1
−0.2
F
M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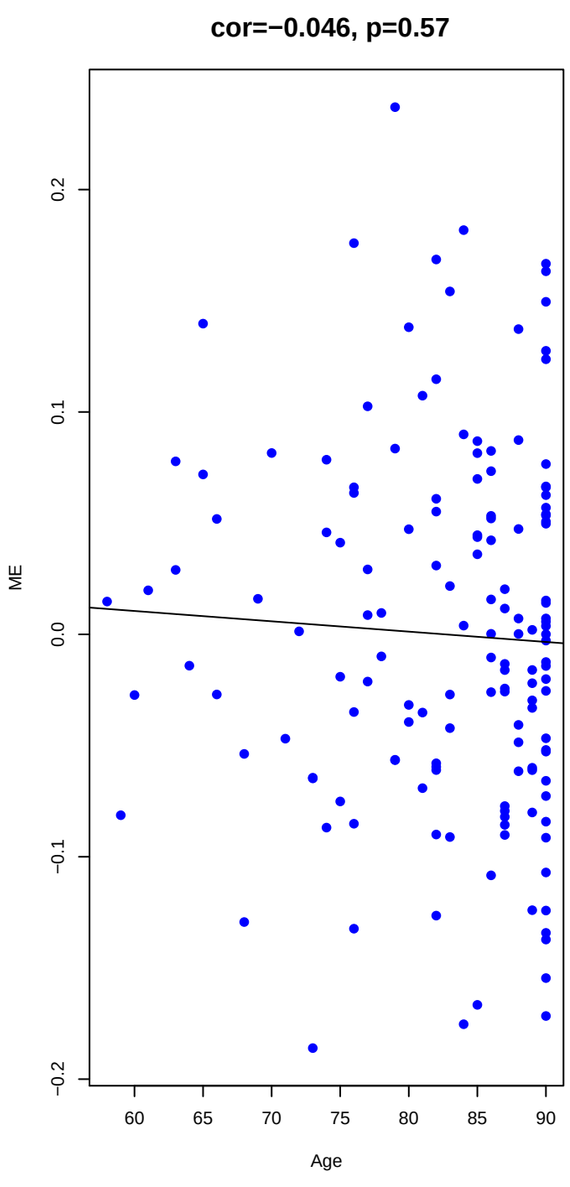
−0.2
CONTROL
AD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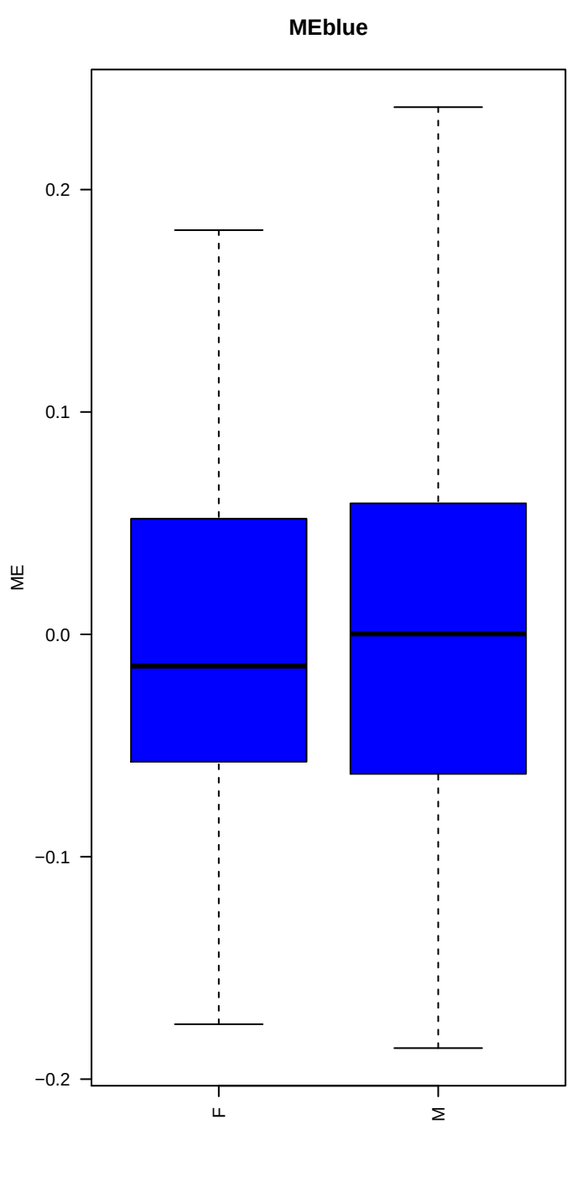
Age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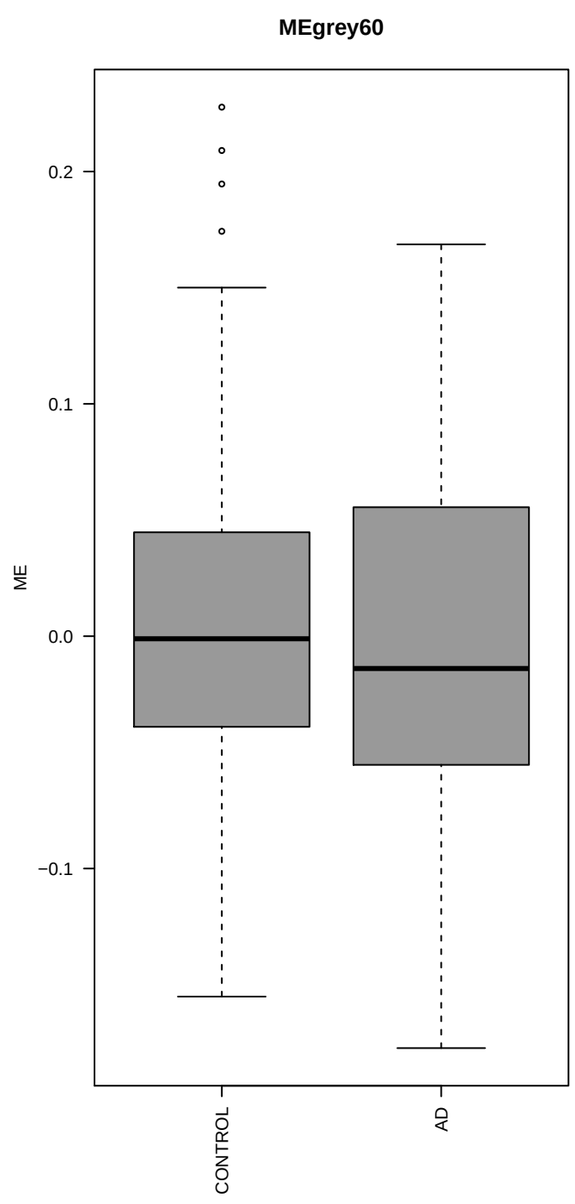
M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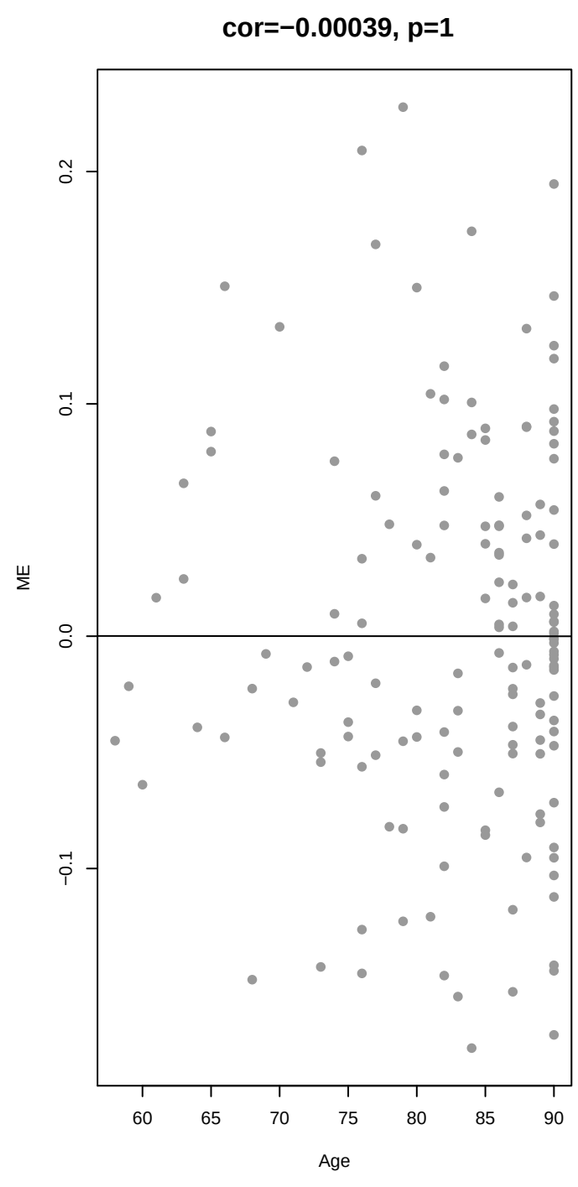
−0.1
CONTROL
AD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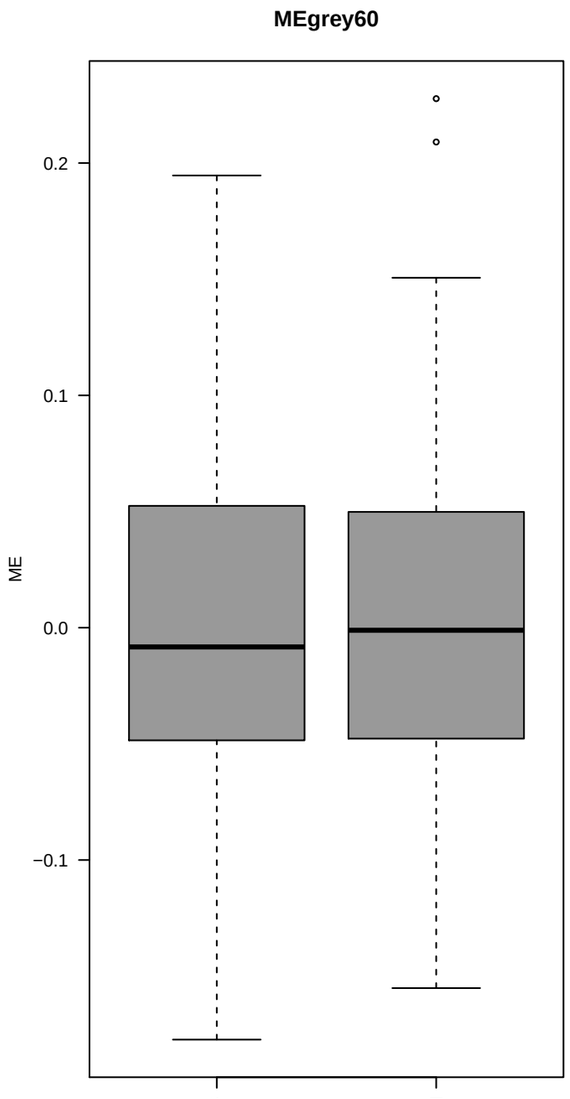
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0.1
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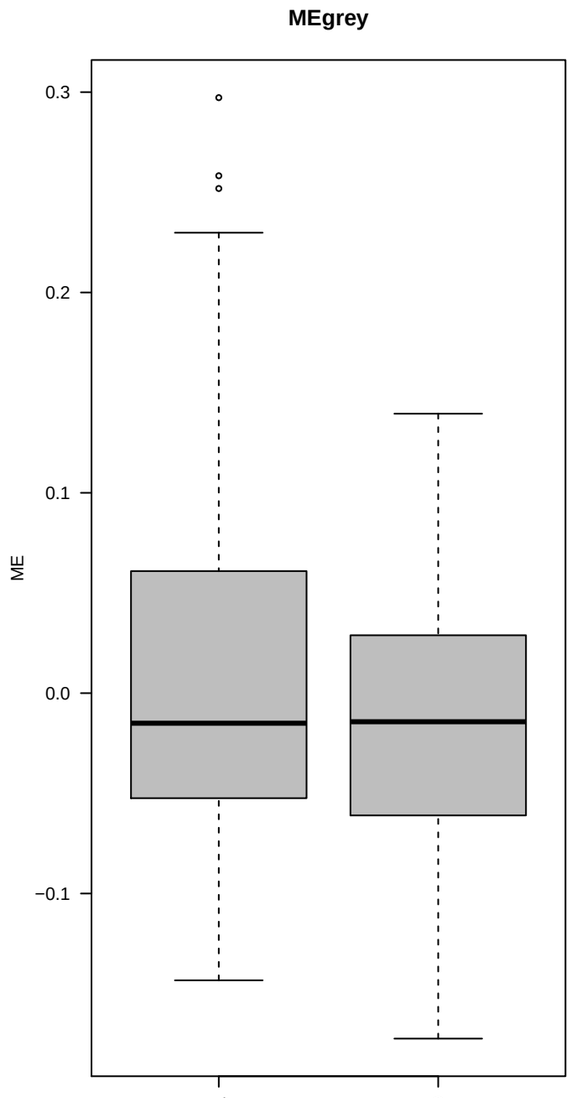
F
M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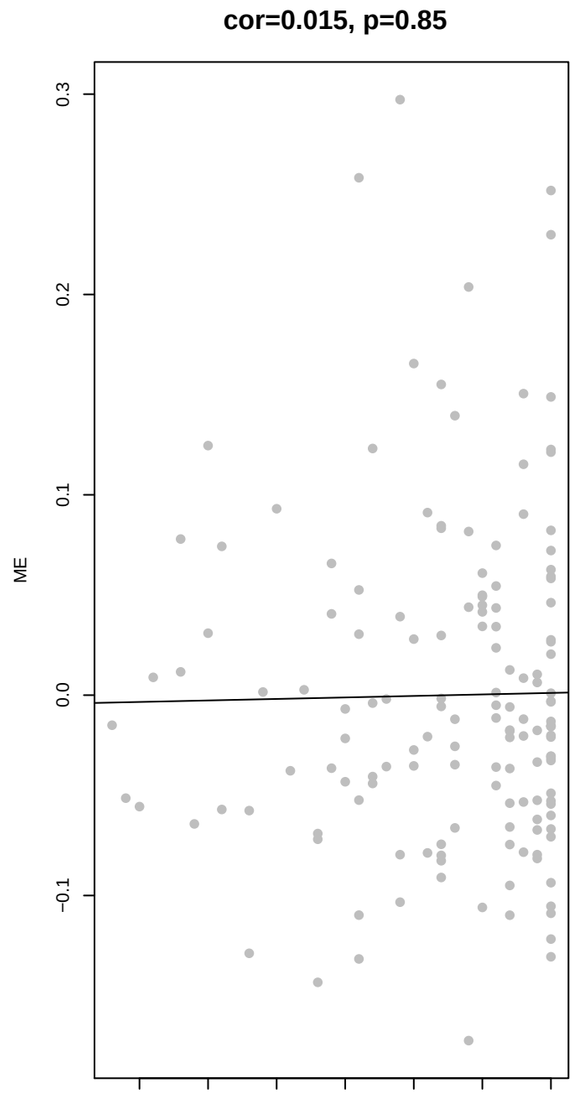
0.2
0.1
0.0
−0.1
CONTROL
AD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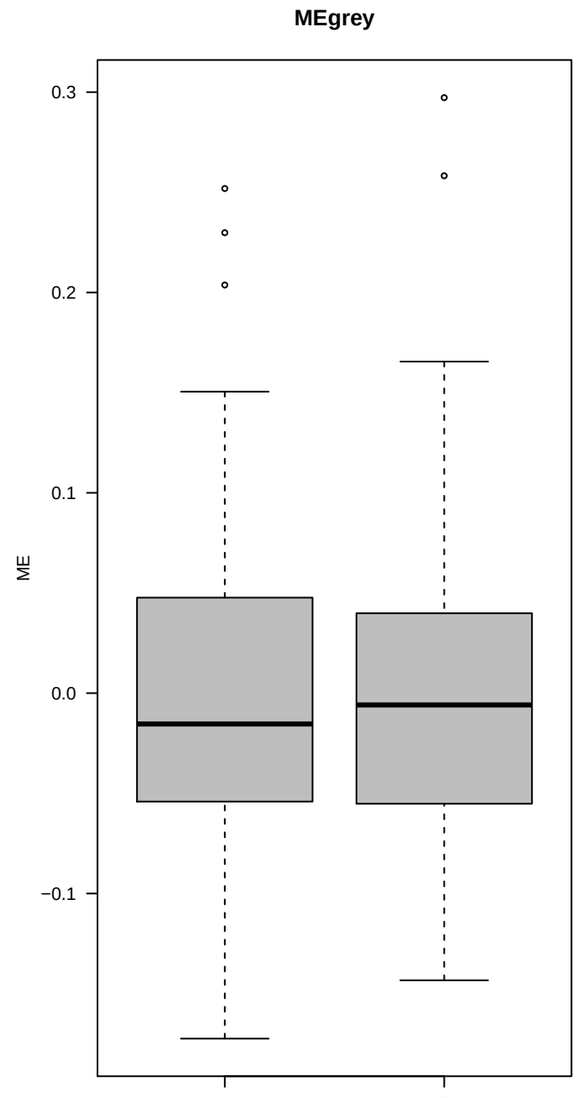
MEgrey
ME
0.3
0.2
0.1
0.0
−0.1
F
M