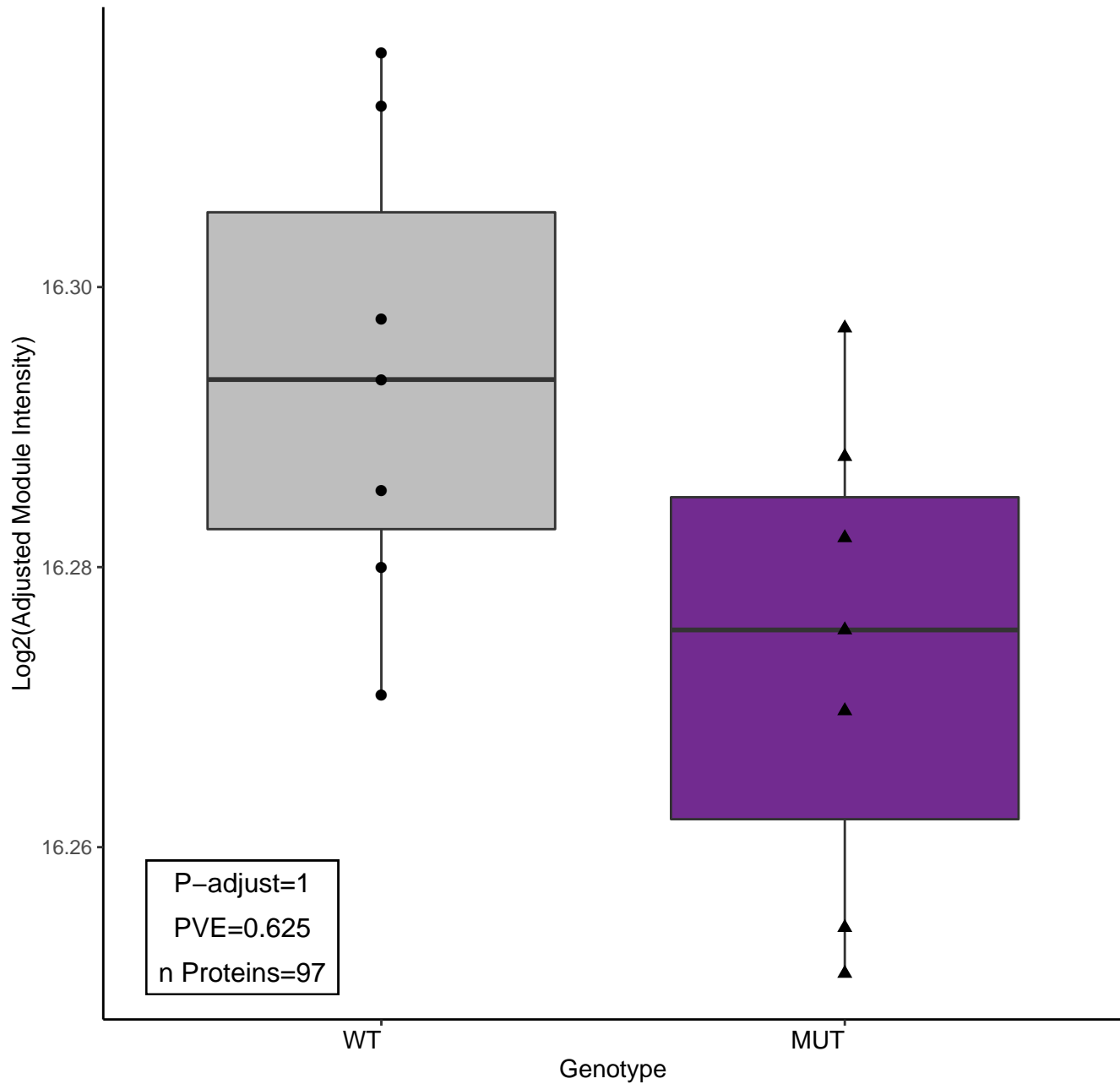
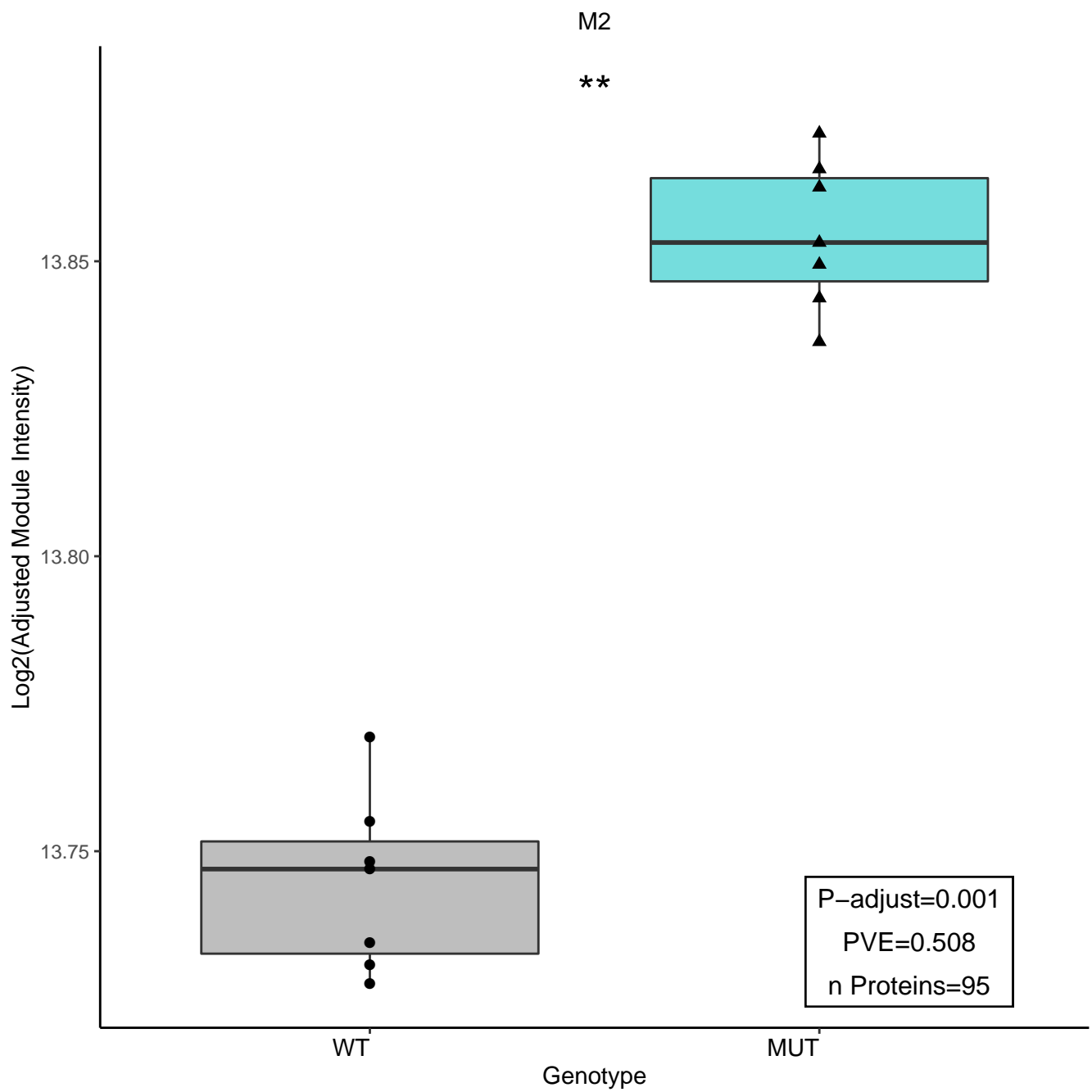
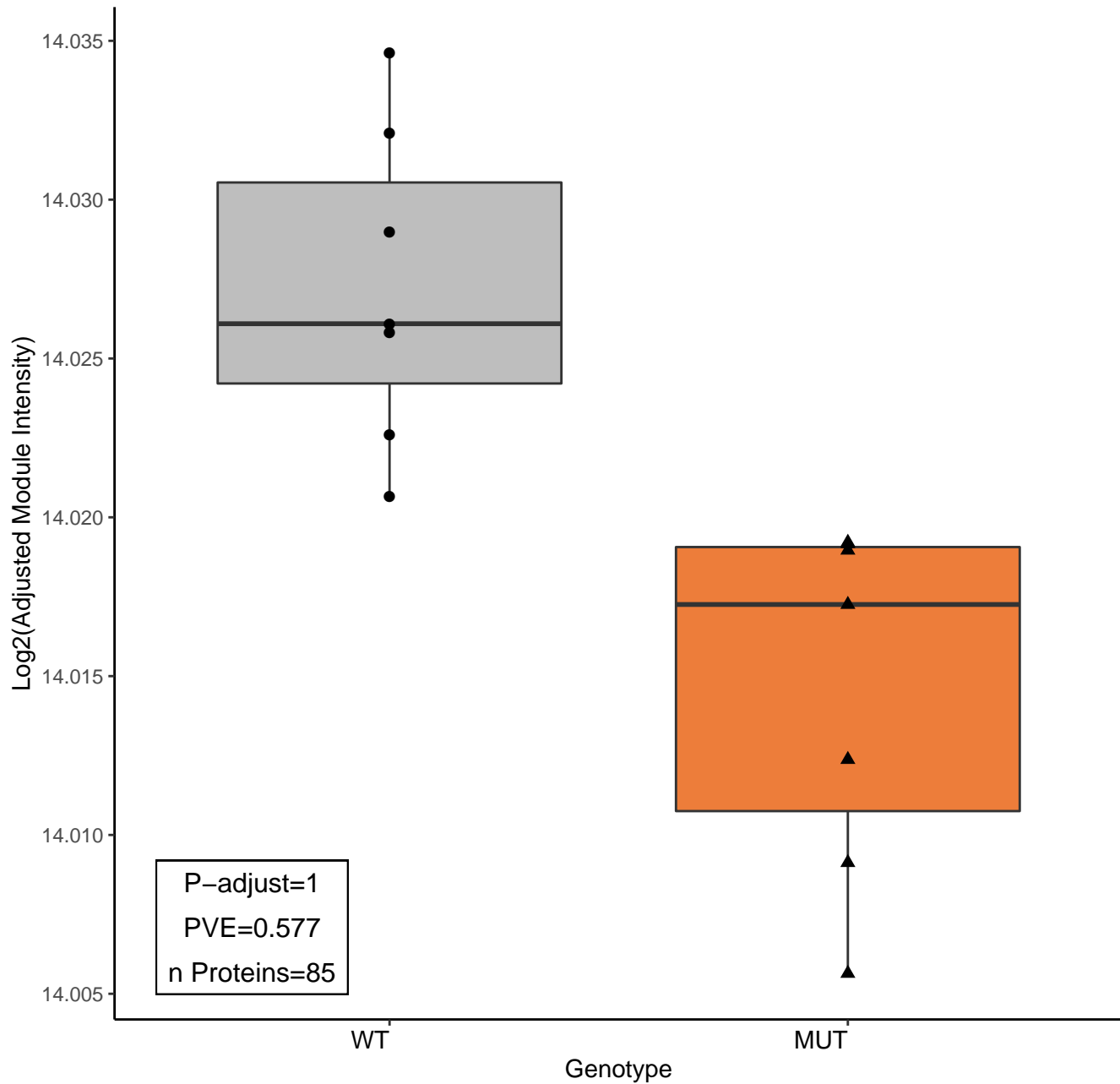


M1

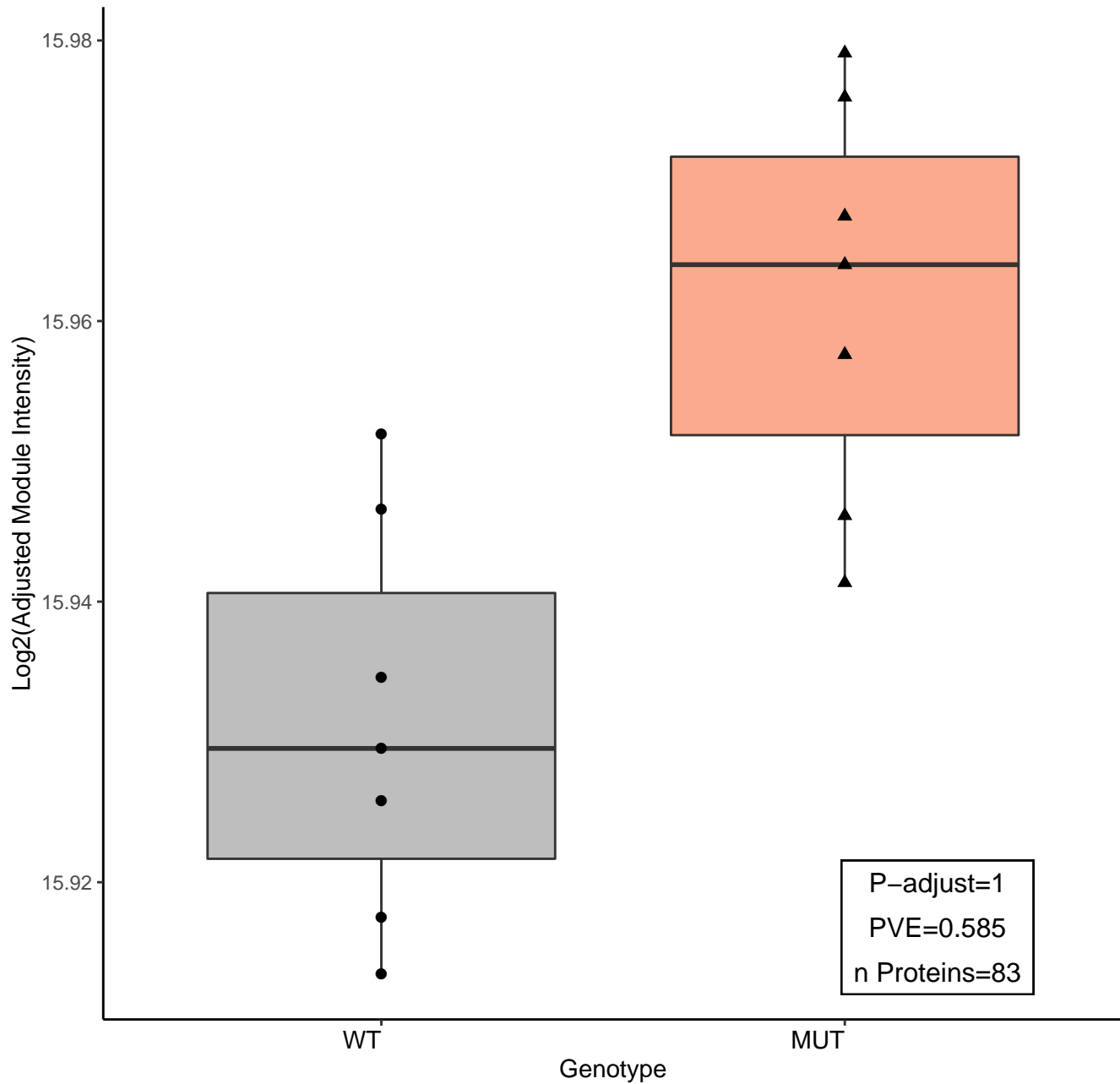




M3



M4



M5

Log2(Adjusted Module Intensity)

13.72

13.71

13.70

13.69

WT

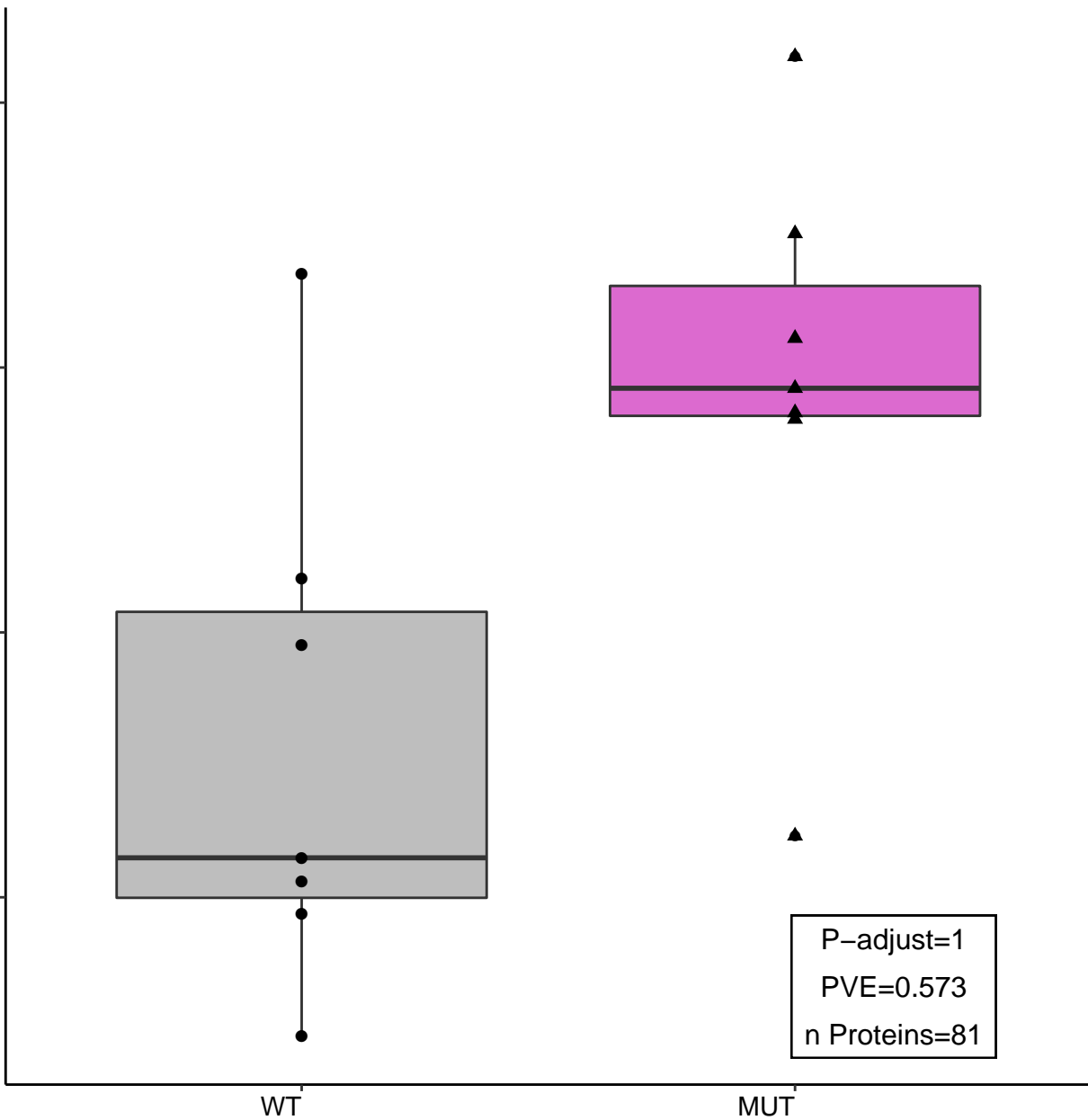
Genotype

MUT

P-adjust=1

PVE=0.573

n Proteins=81



M6

Log2(Adjusted Module Intensity)

14.86

14.84

14.82

14.80

WT

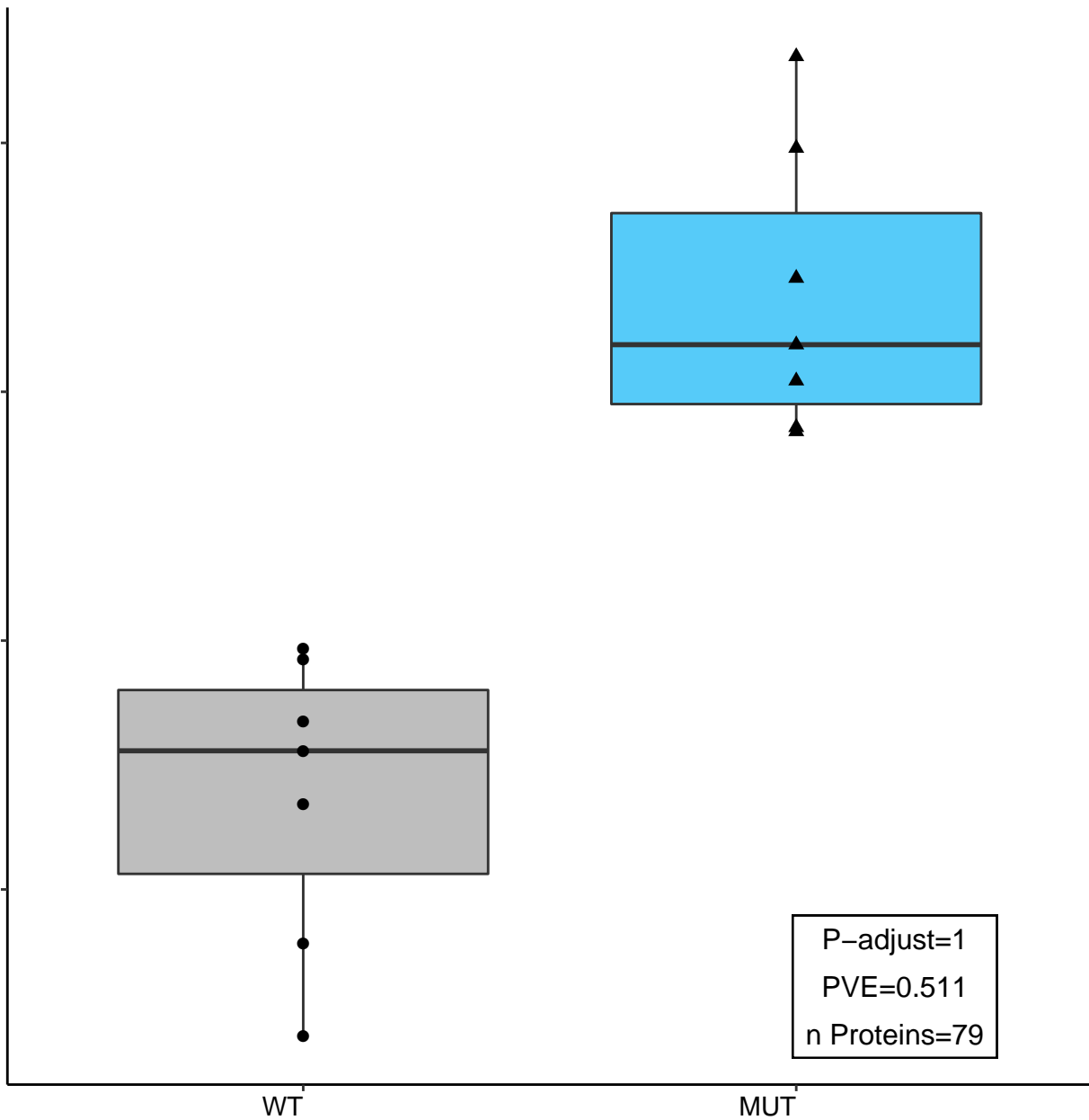
MUT

Genotype

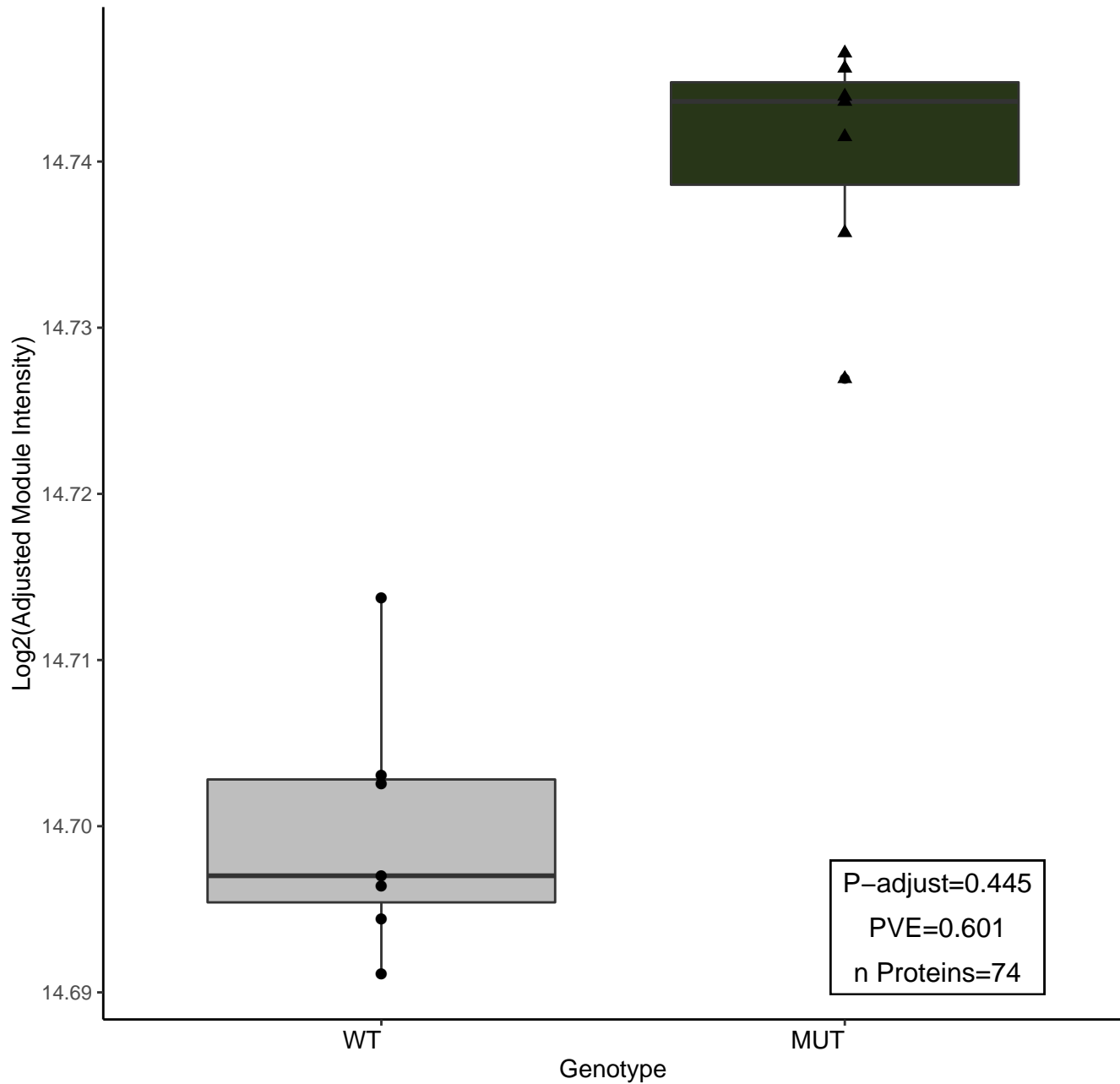
P-adjust=1

PVE=0.511

n Proteins=79



M7



M8

Log2(Adjusted Module Intensity)

13.22

13.20

13.18

WT

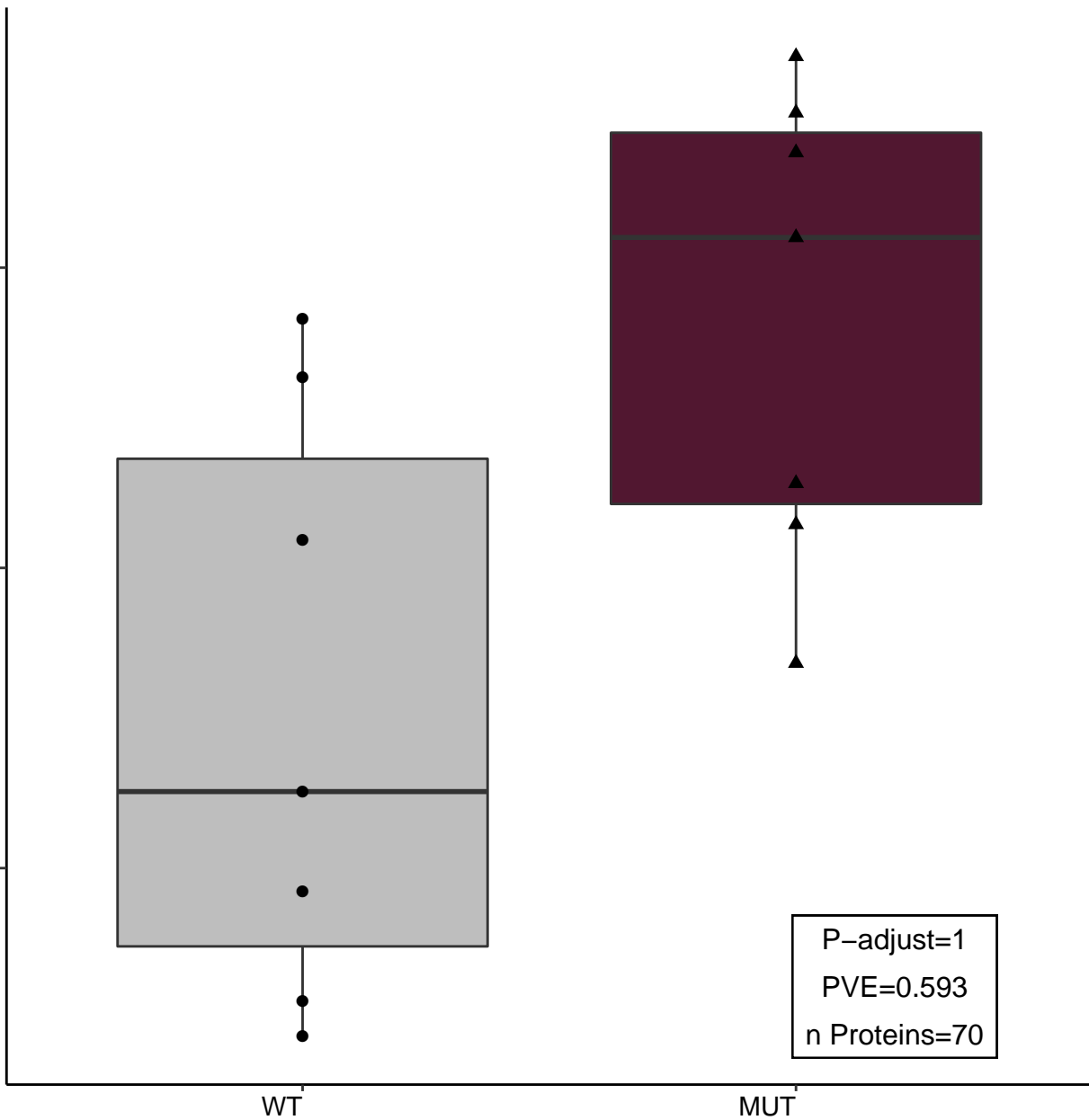
Genotype

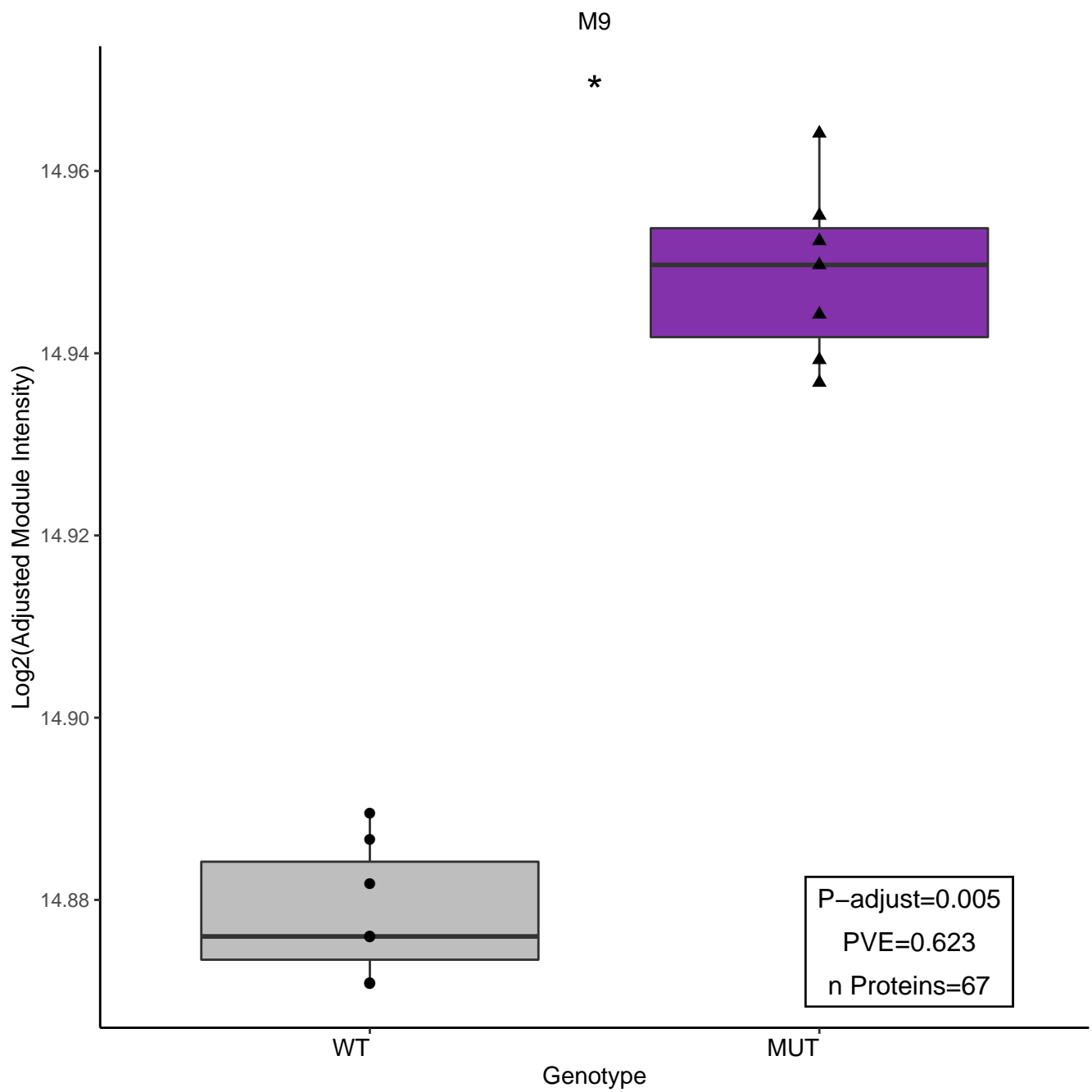
MUT

P-adjust=1

PVE=0.593

n Proteins=70





M10

Log2(Adjusted Module Intensity)

12.25

12.24

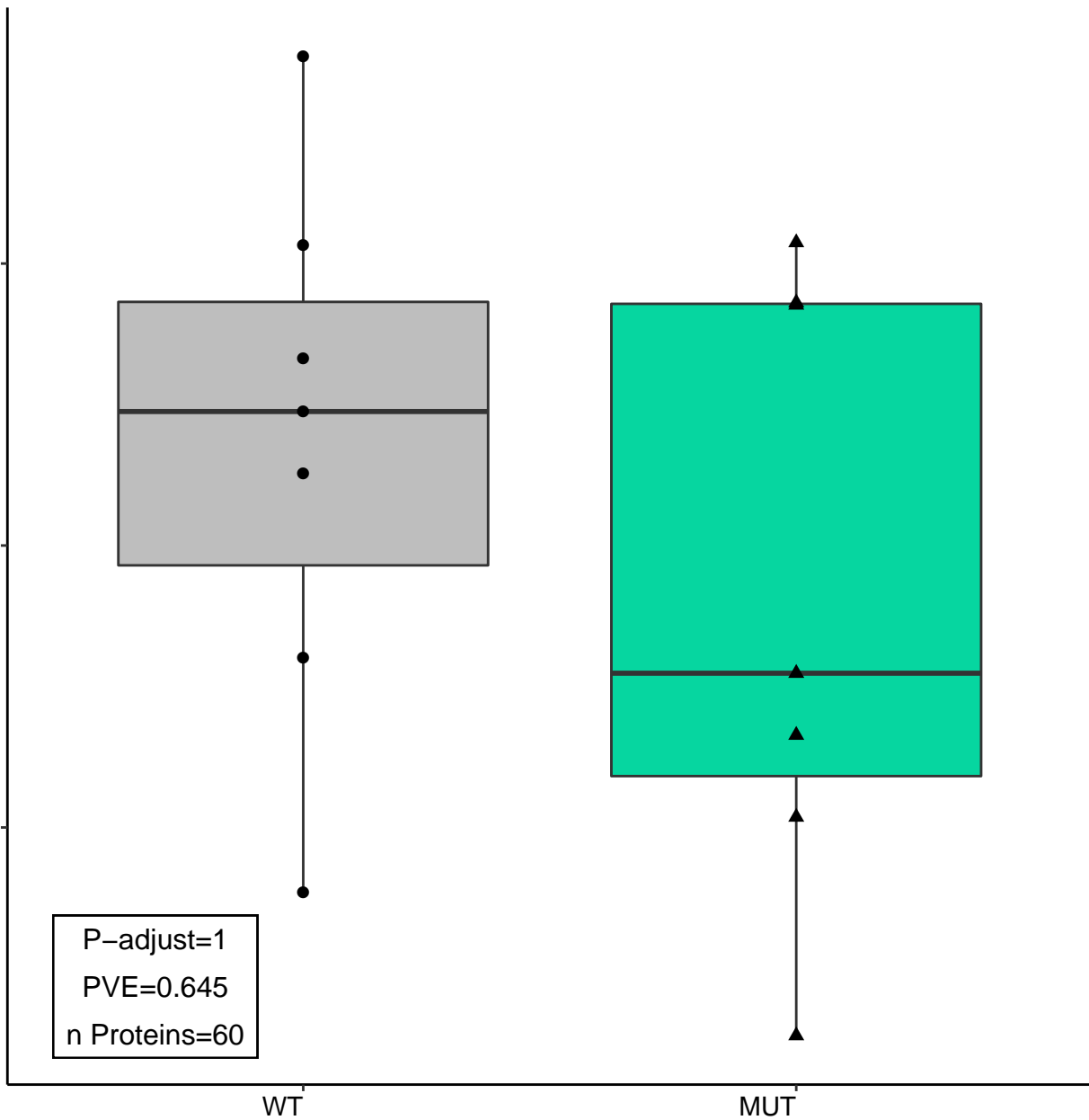
12.23

P-adjust=1
PVE=0.645
n Proteins=60

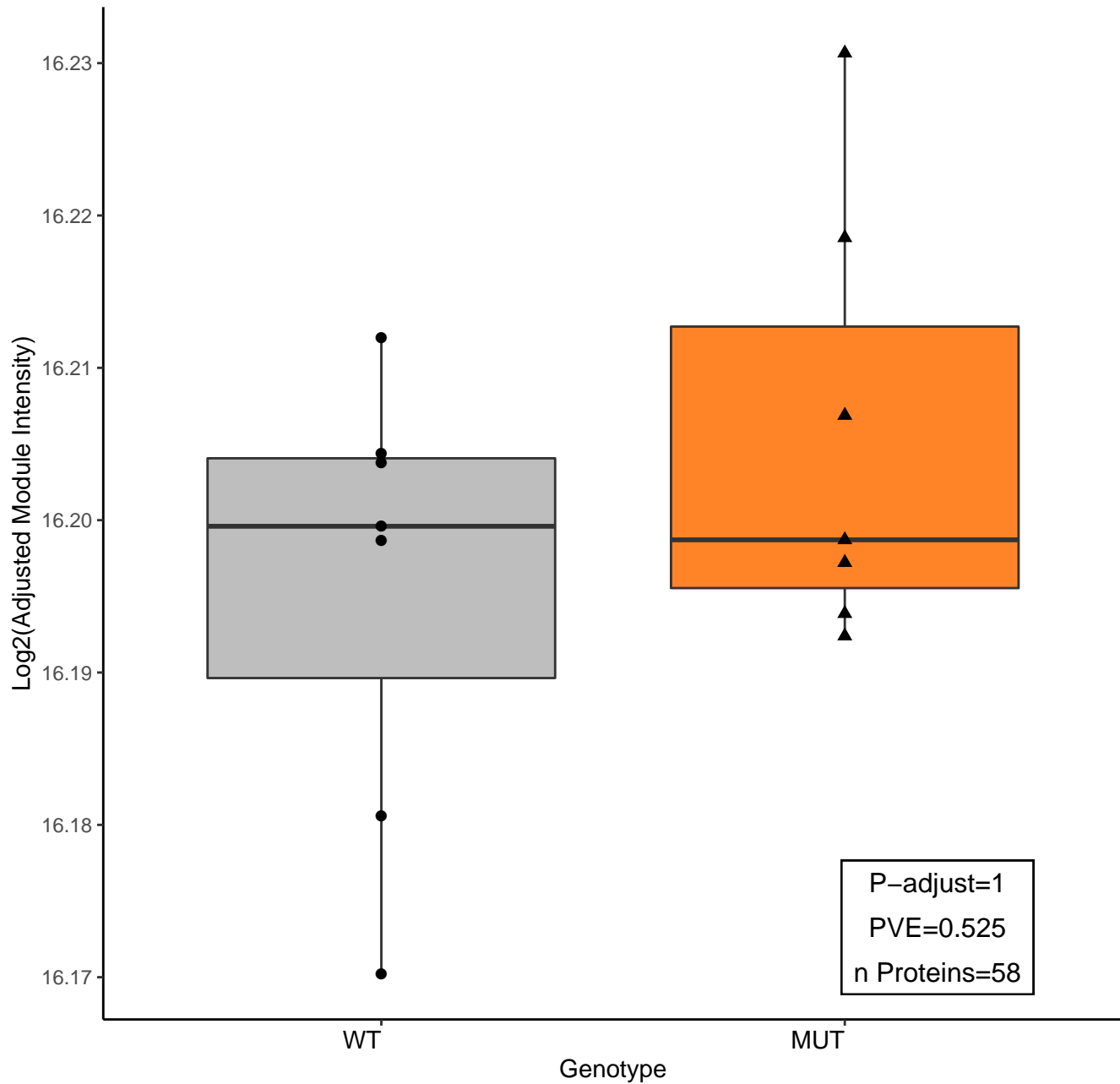
WT

Genotype

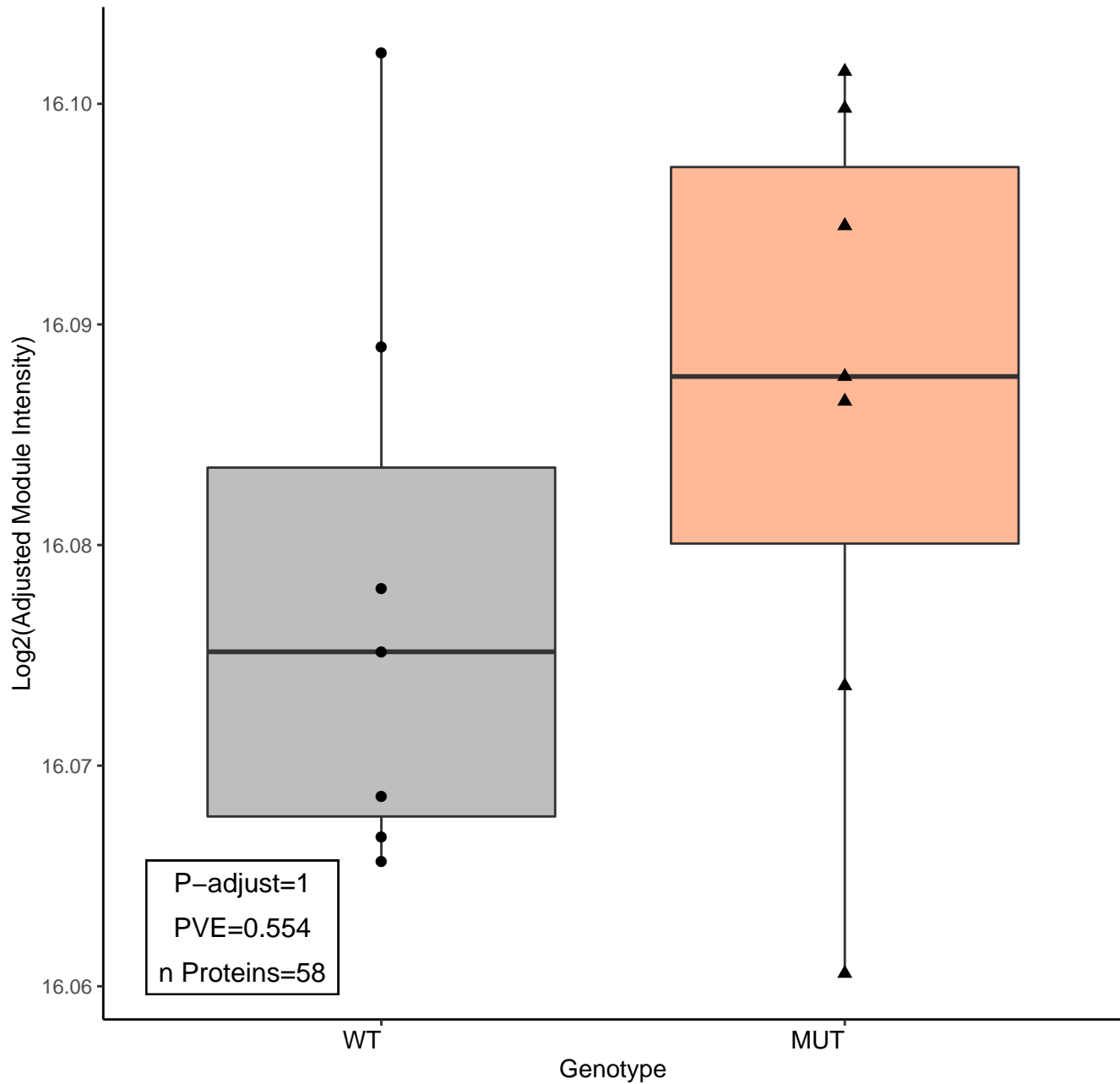
MUT



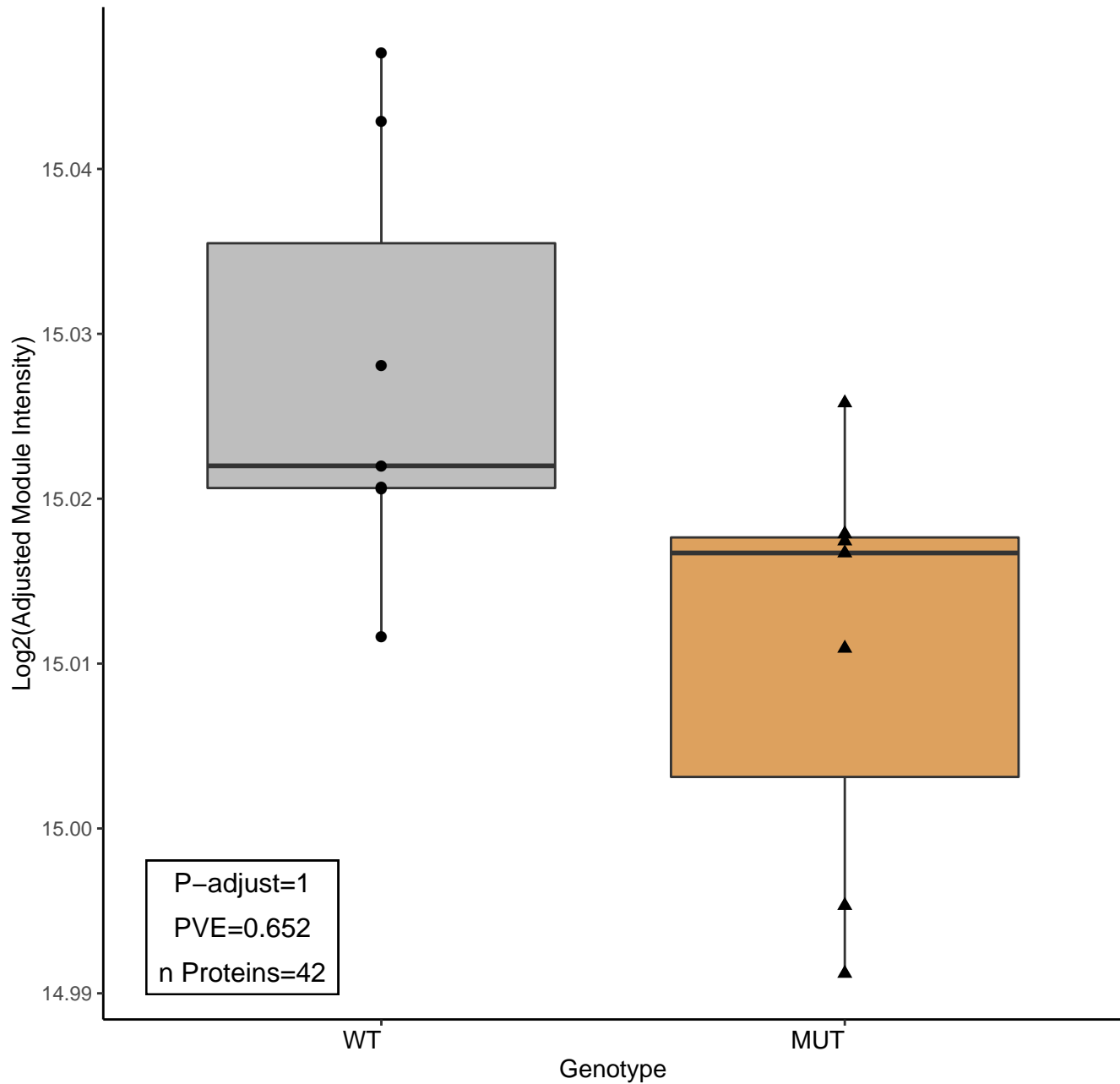
M11

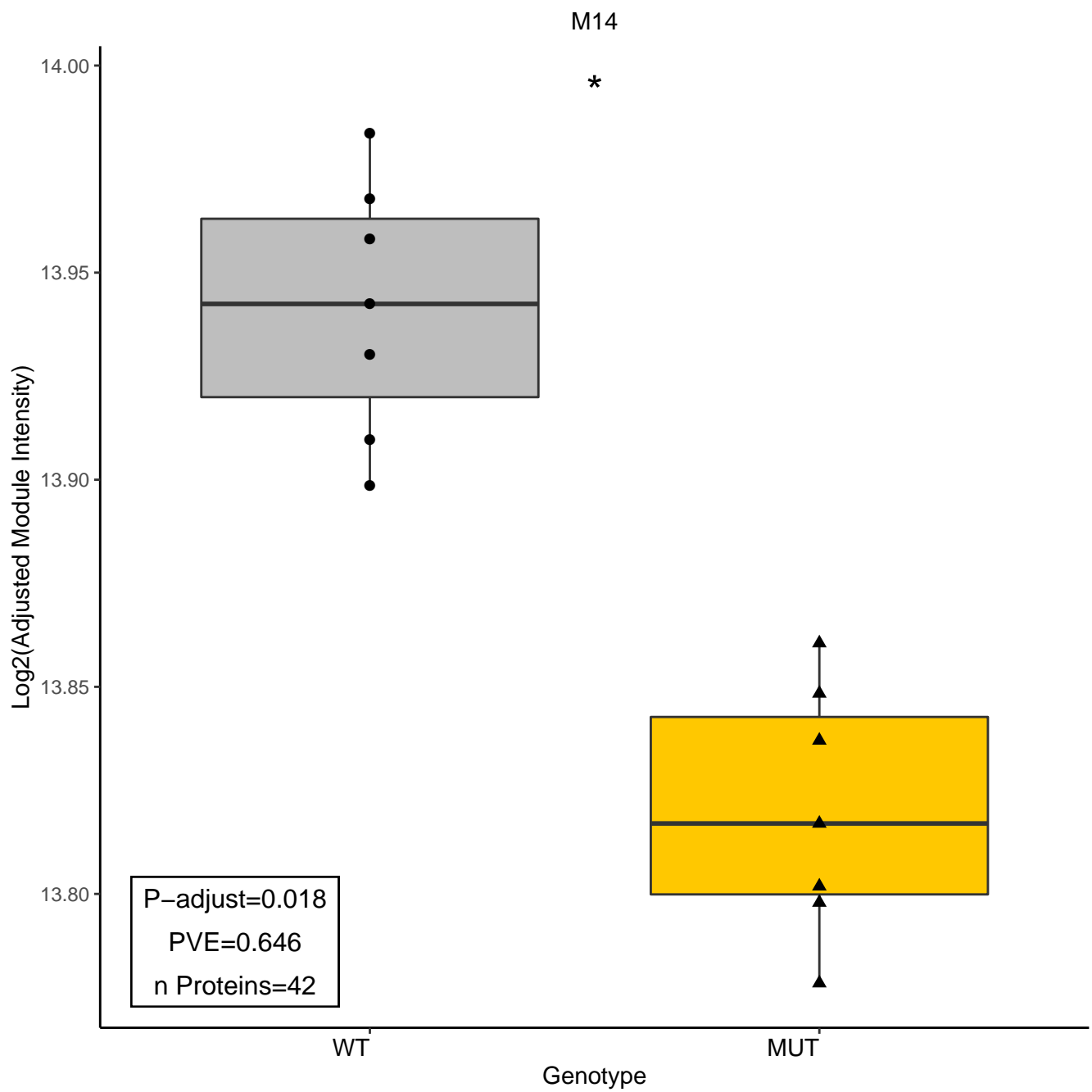


M12



M13





M15

Log2(Adjusted Module Intensity)

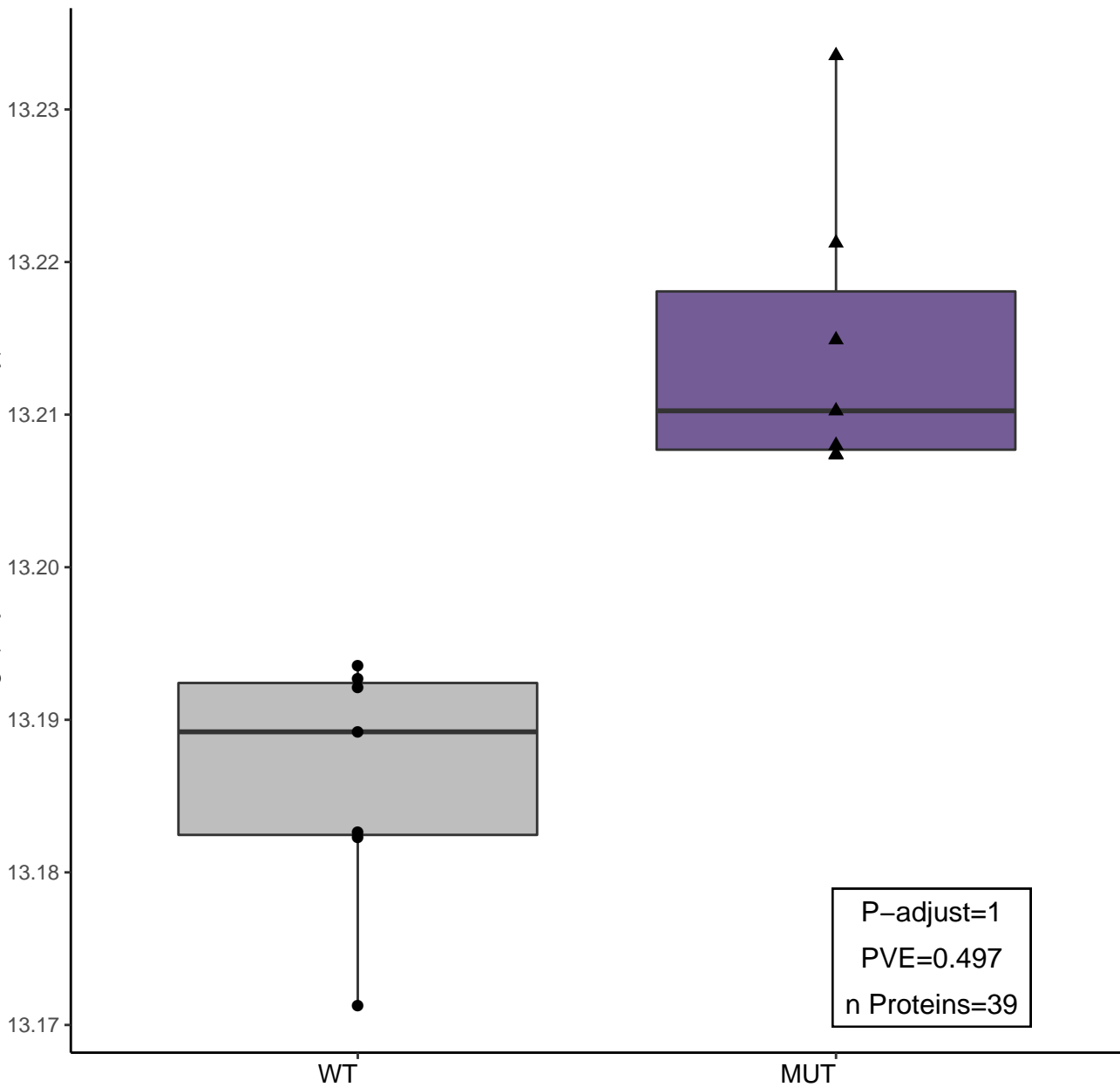
13.23
13.22
13.21
13.20
13.19
13.18
13.17

WT

MUT

Genotype

P-adjust=1
PVE=0.497
n Proteins=39



M16

Log2(Adjusted Module Intensity)

12.26

12.28

12.30

12.32

12.34

WT

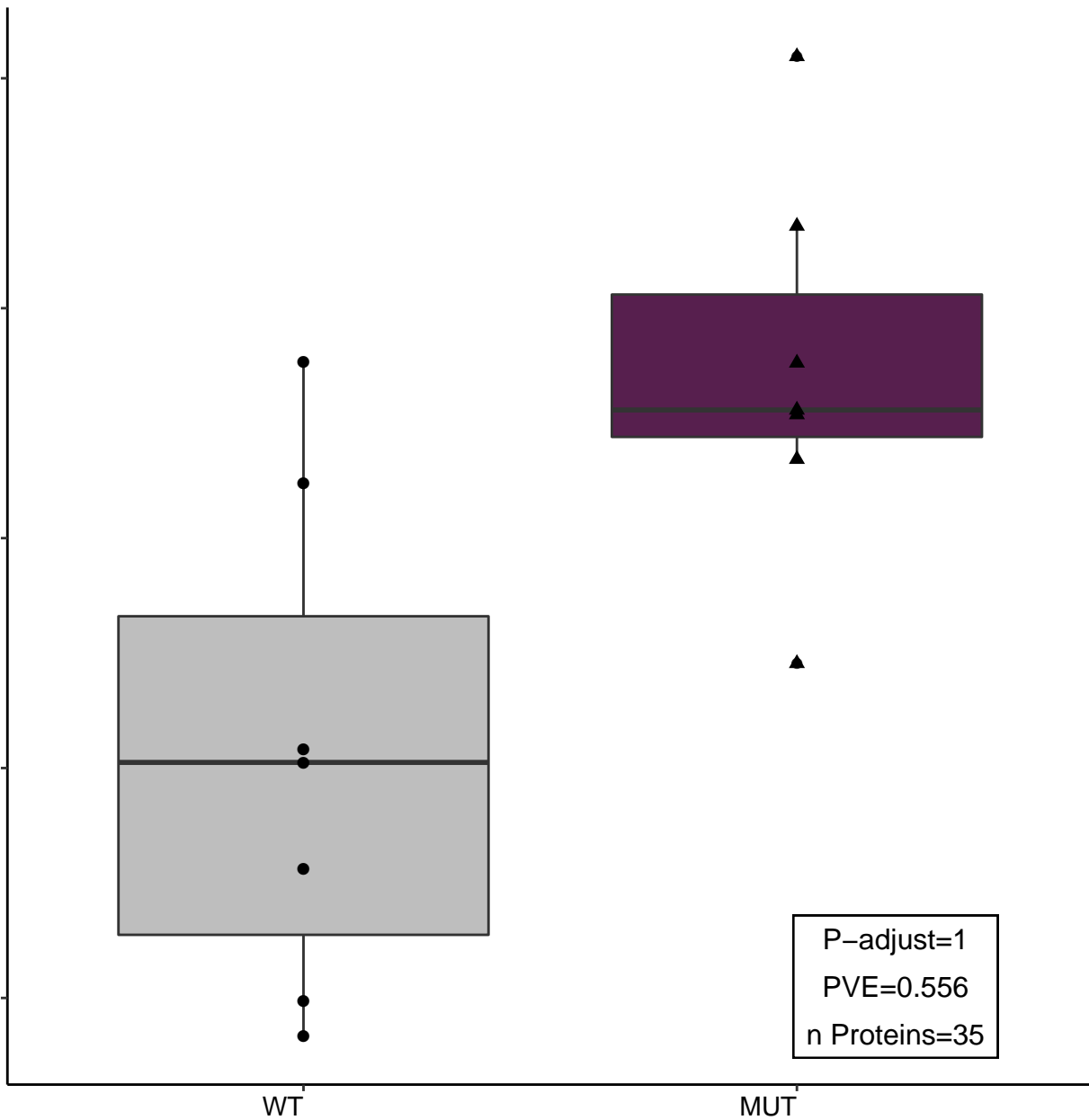
Genotype

MUT

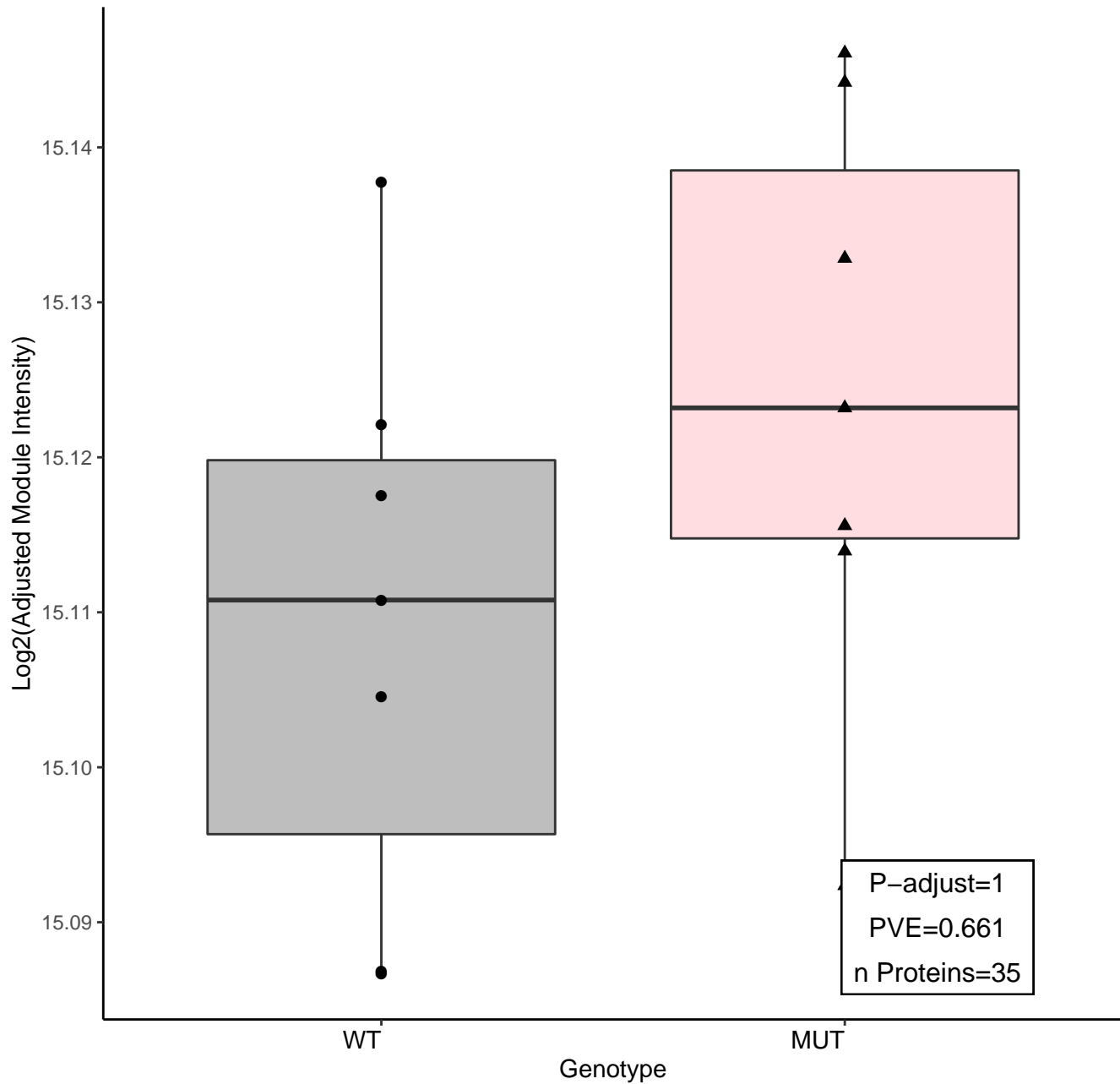
P-adjust=1

PVE=0.556

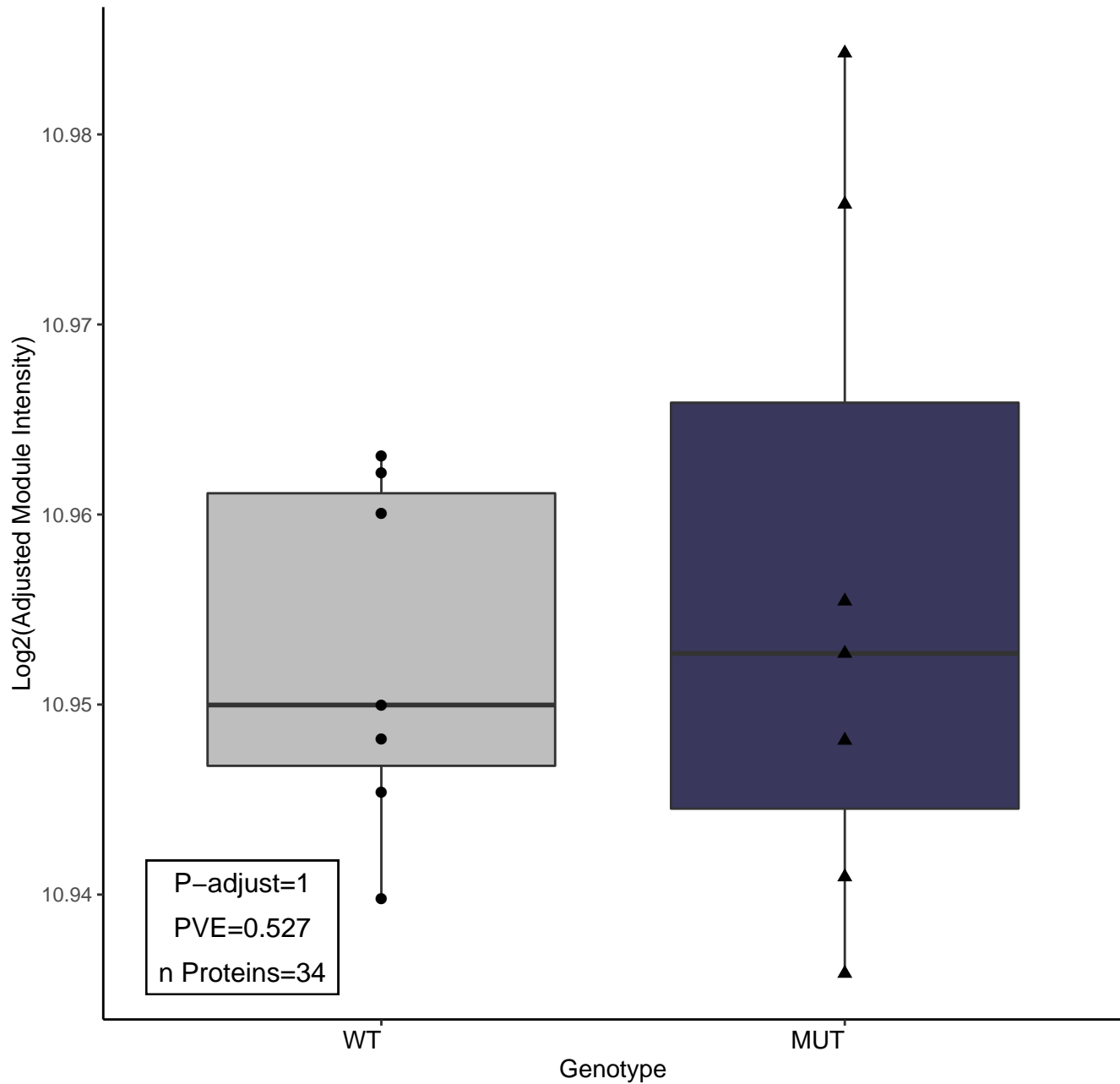
n Proteins=35



M17



M18



M19

**

Log2(Adjusted Module Intensity)

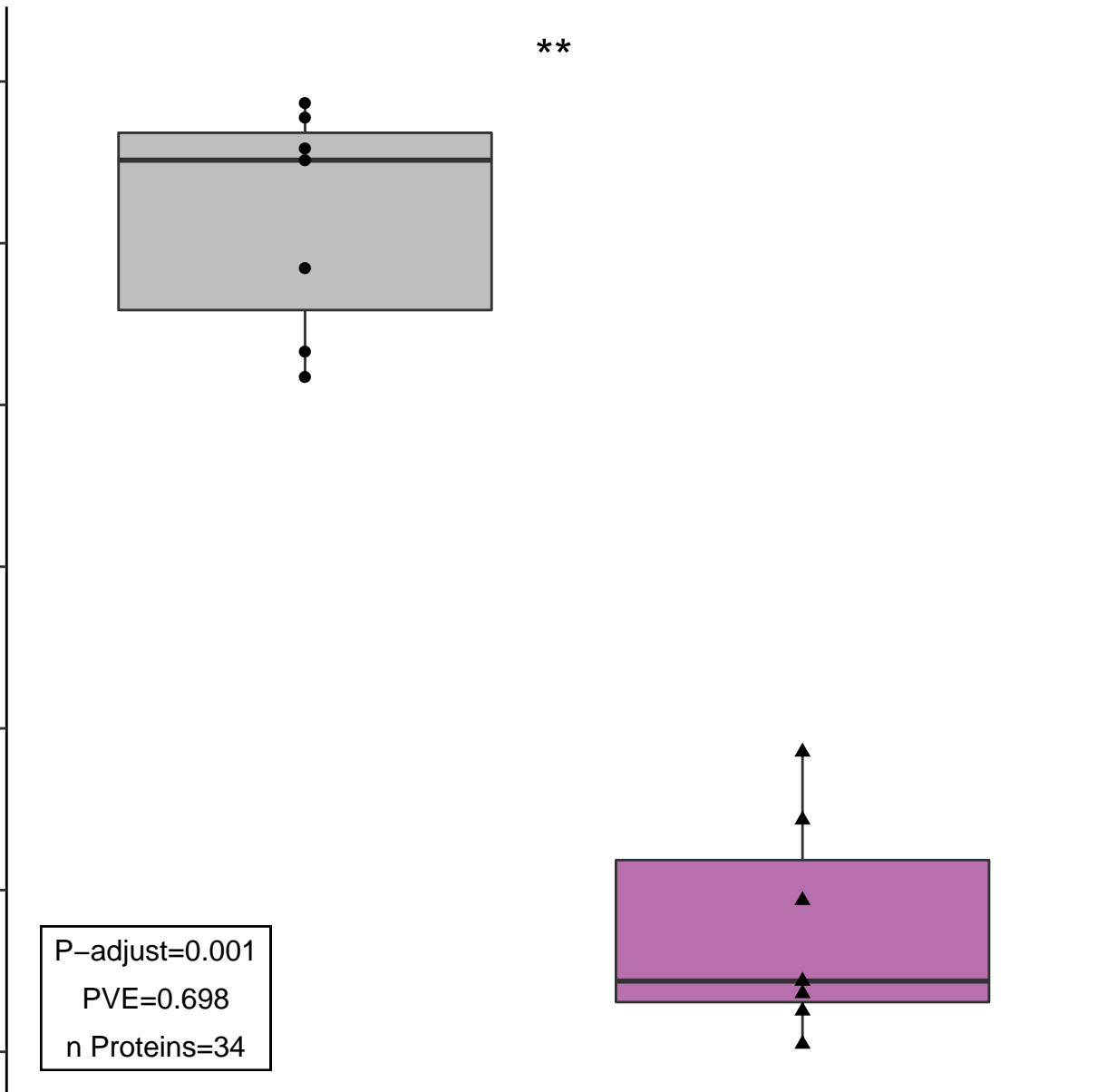
13.25
13.20
13.15
13.10
13.05
13.00
12.95

P-adjust=0.001
PVE=0.698
n Proteins=34

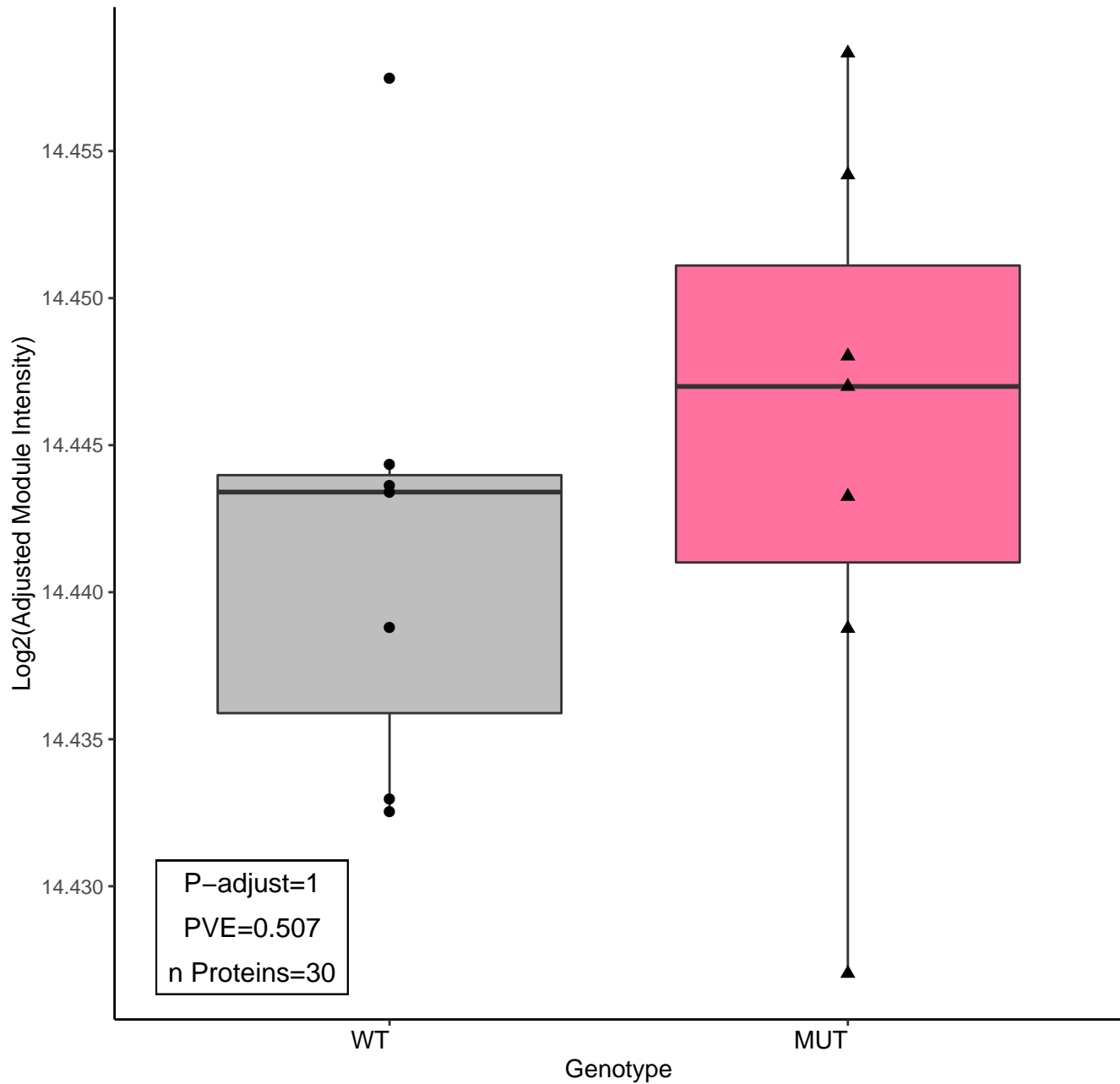
WT

Genotype

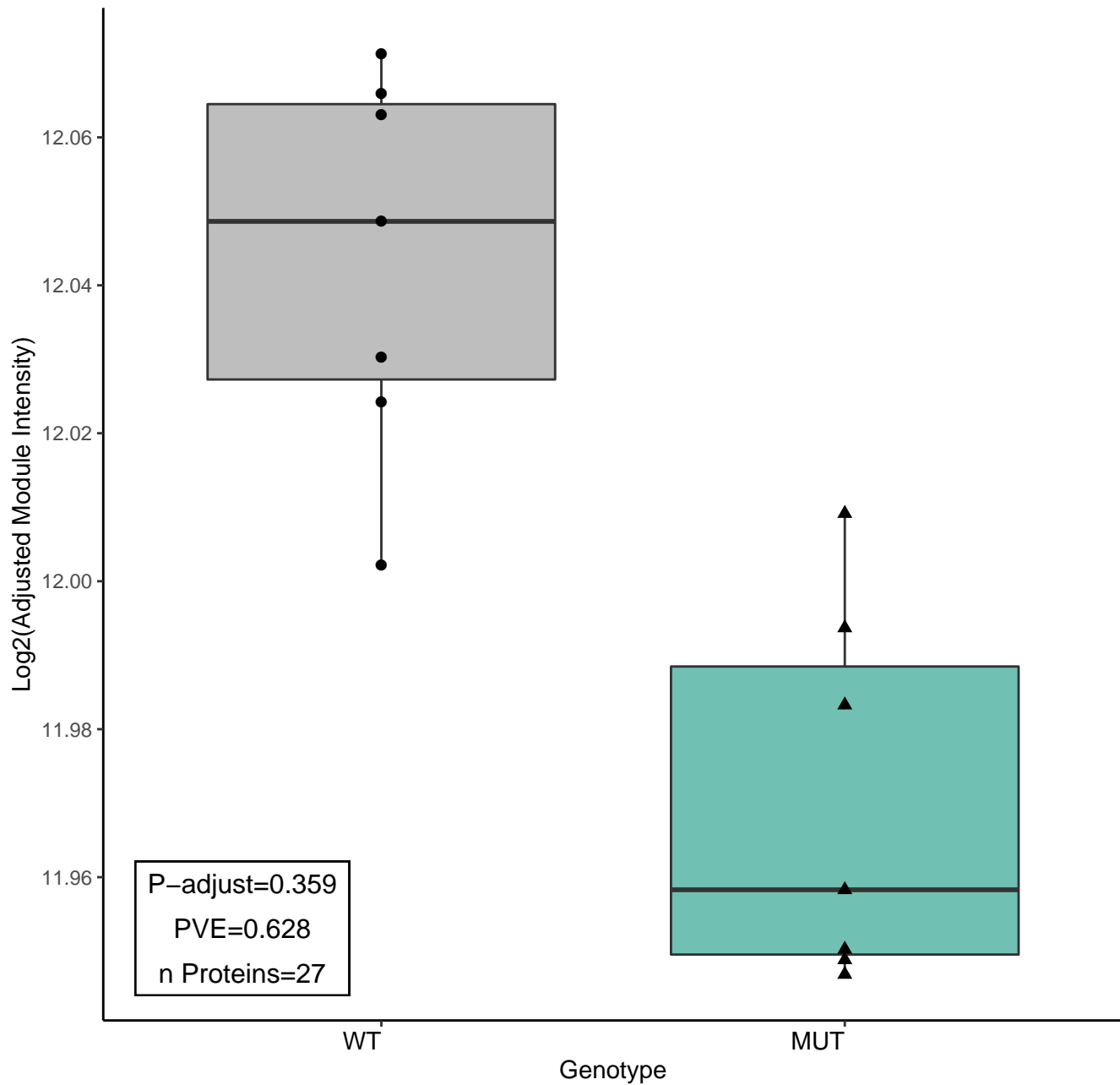
MUT



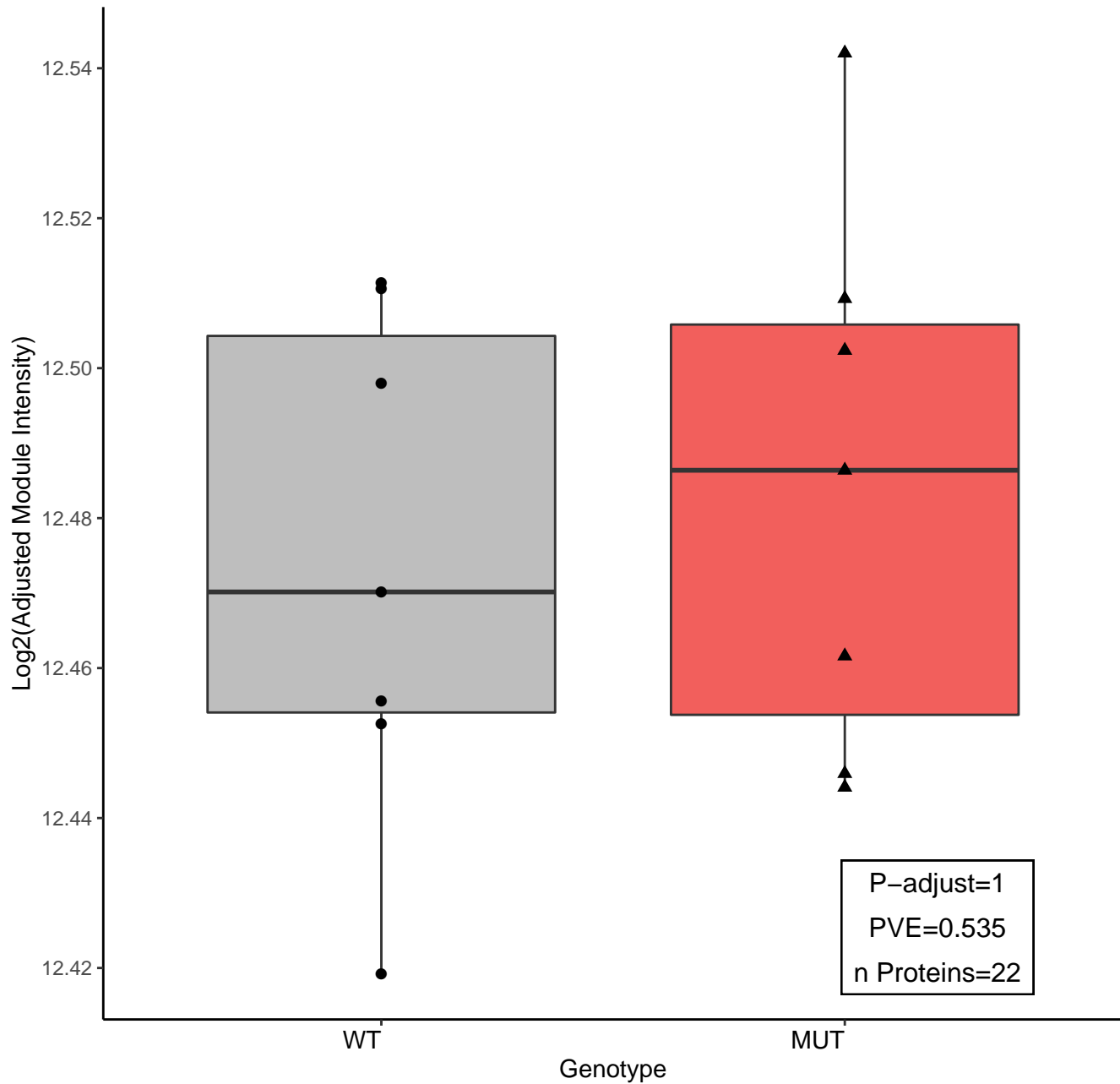
M20

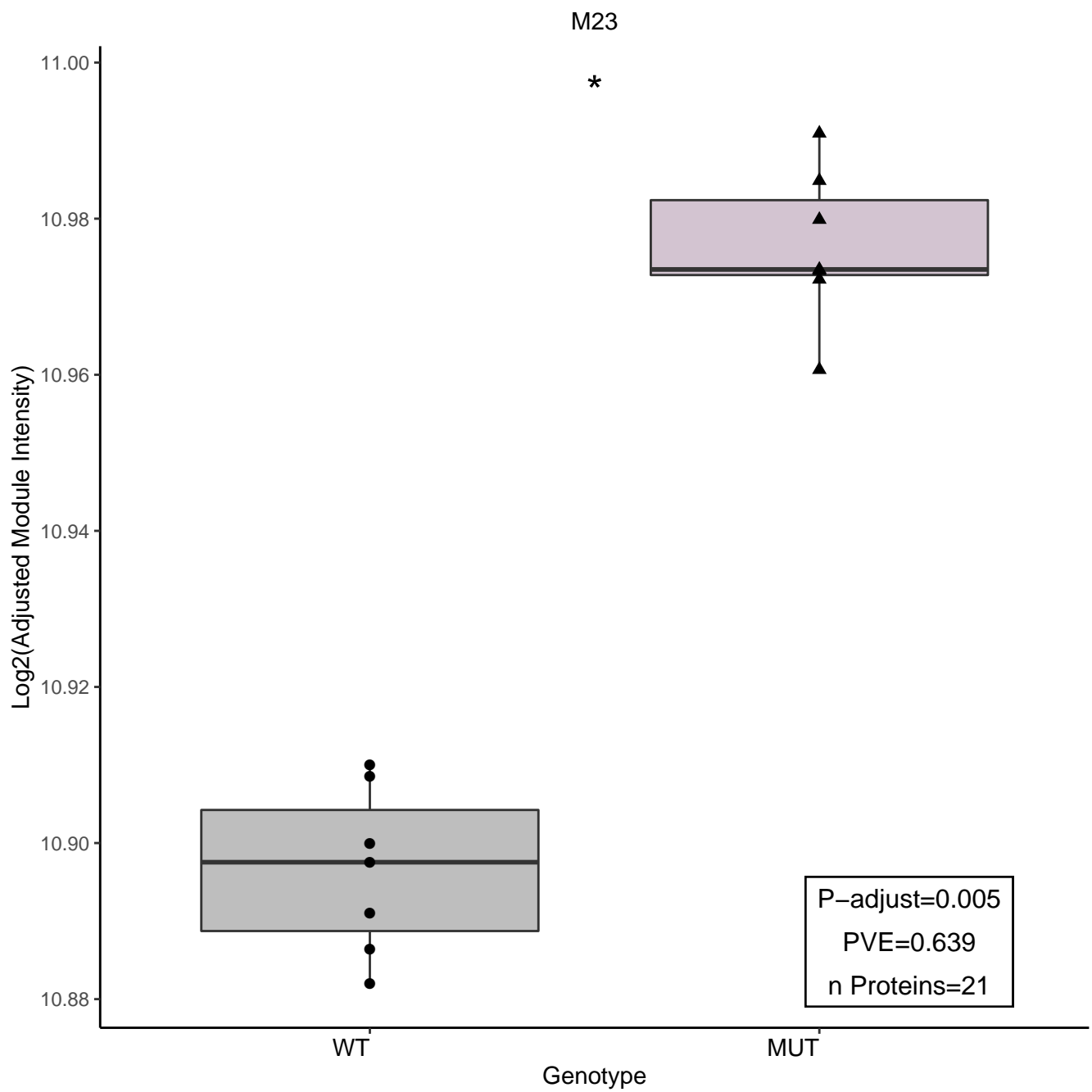


M21

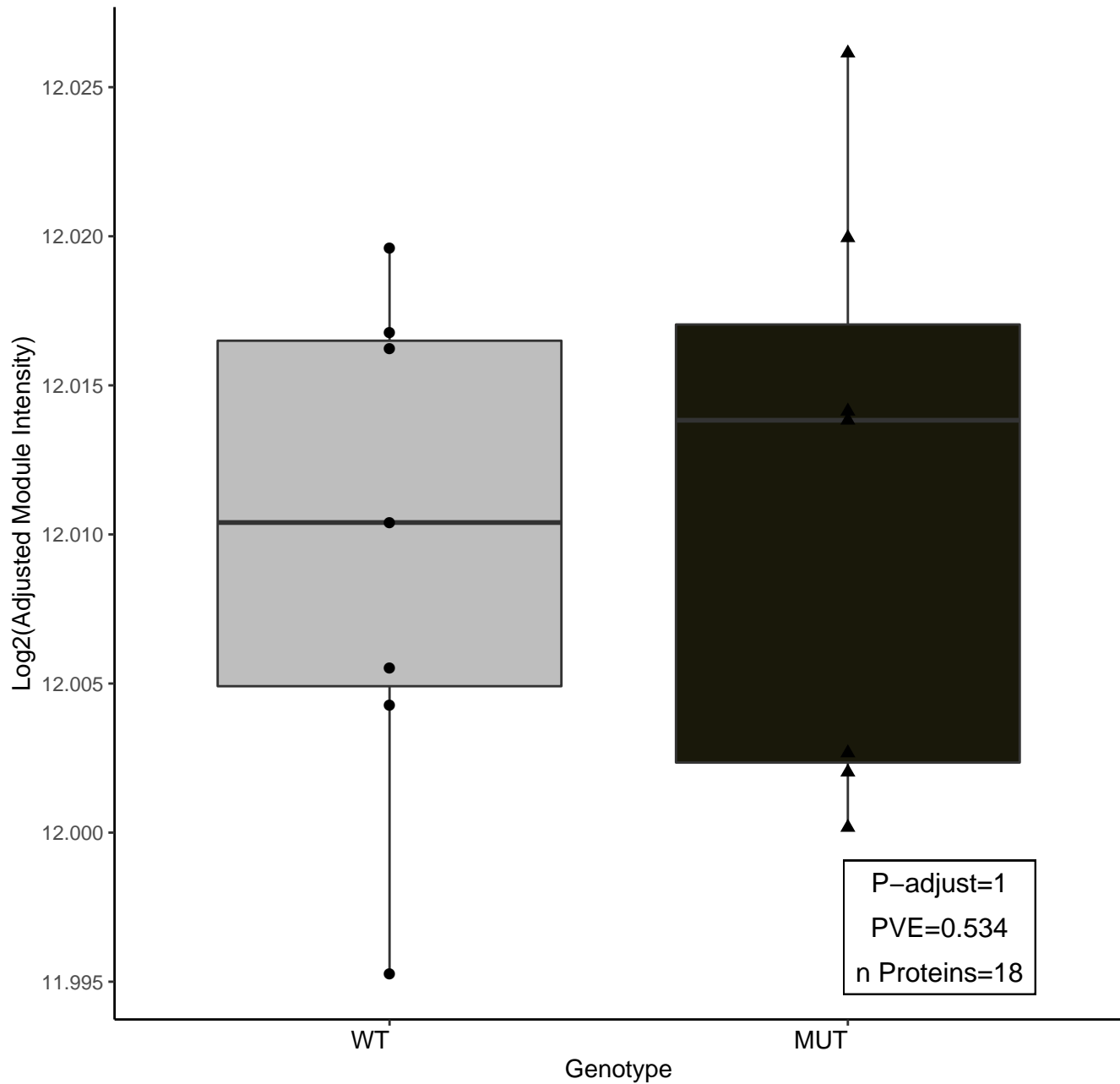


M22

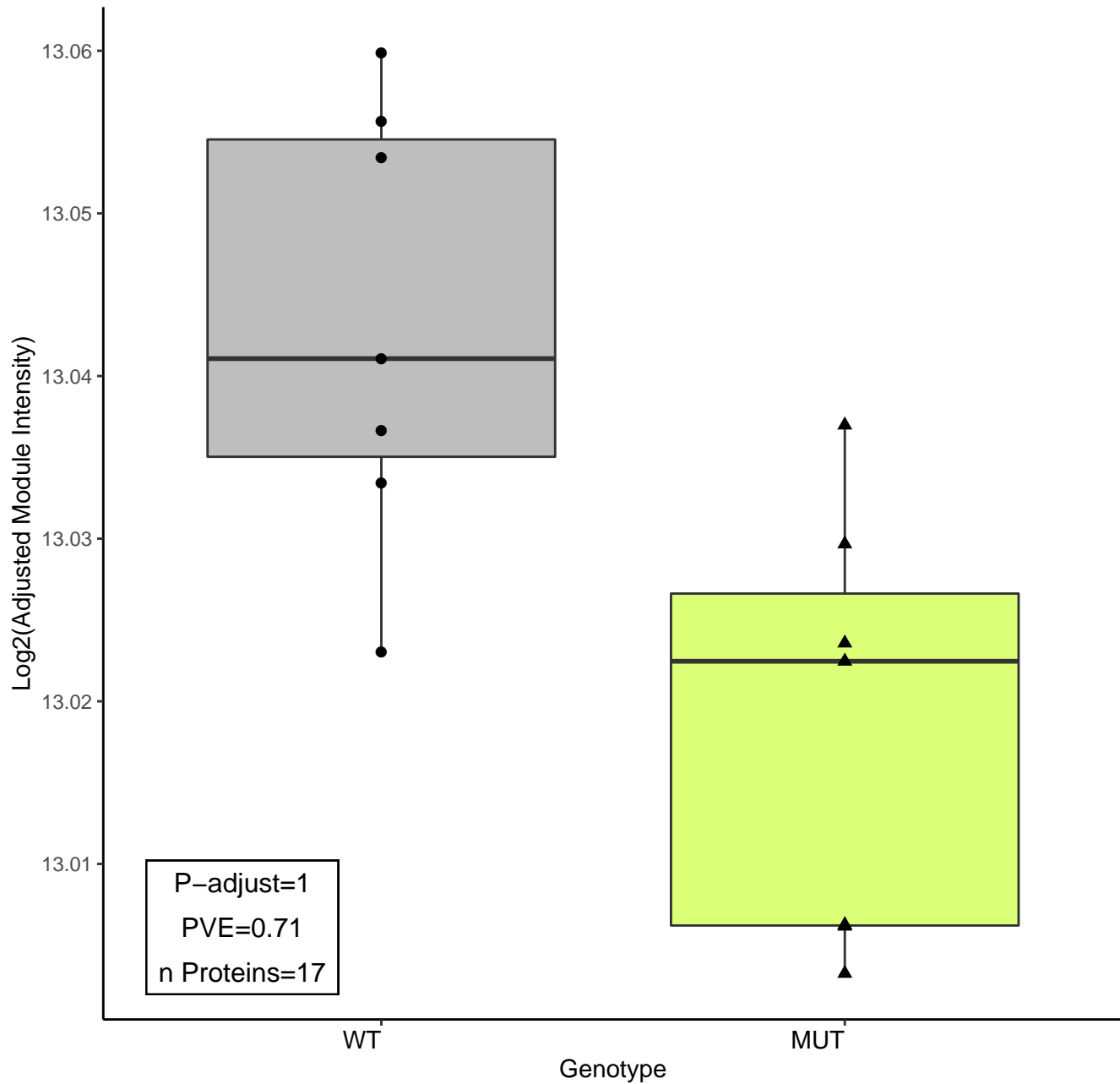




M24



M25



M26

Log2(Adjusted Module Intensity)

11.51

11.50

11.49

11.48

11.47

WT

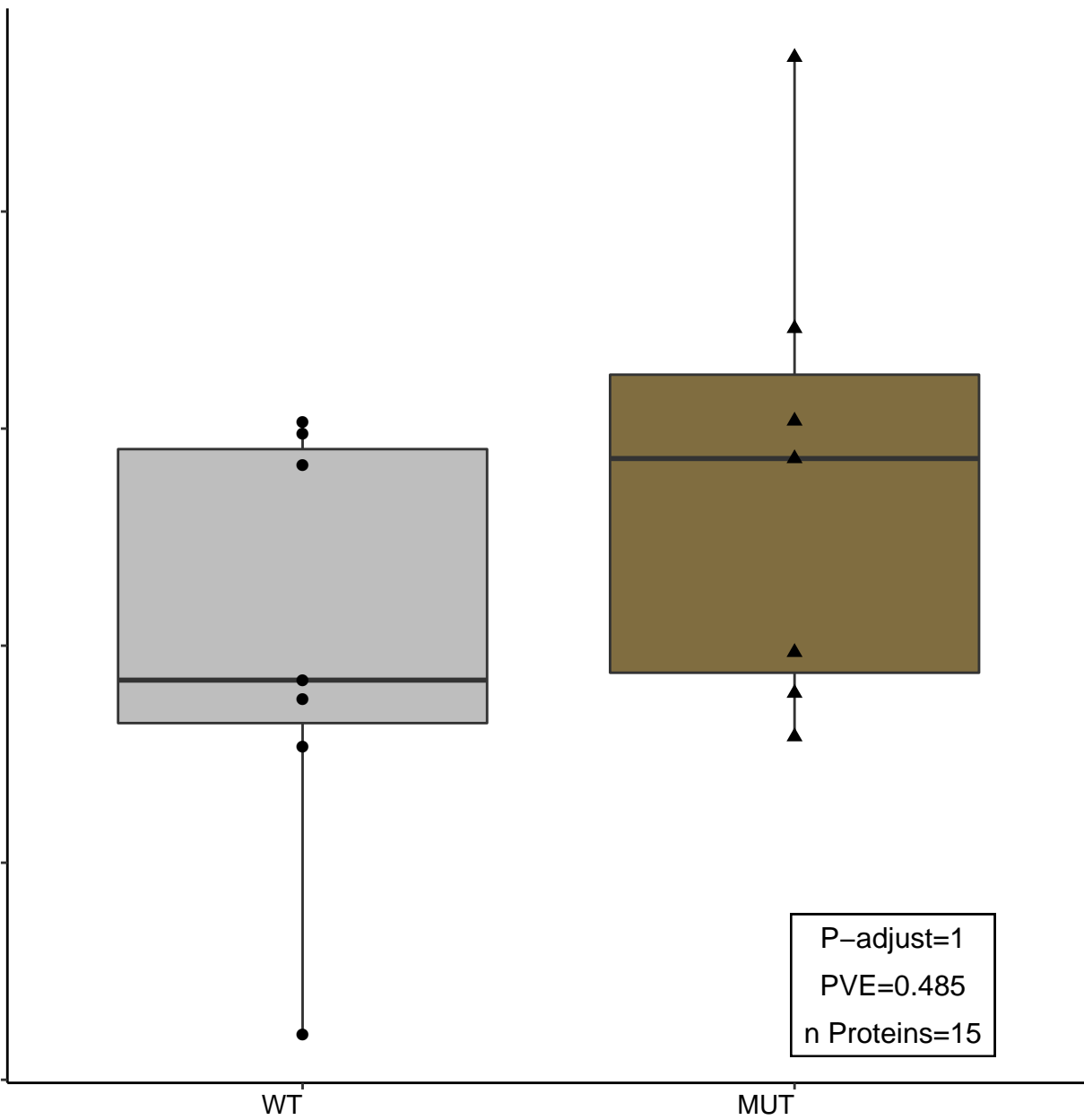
Genotype

MUT

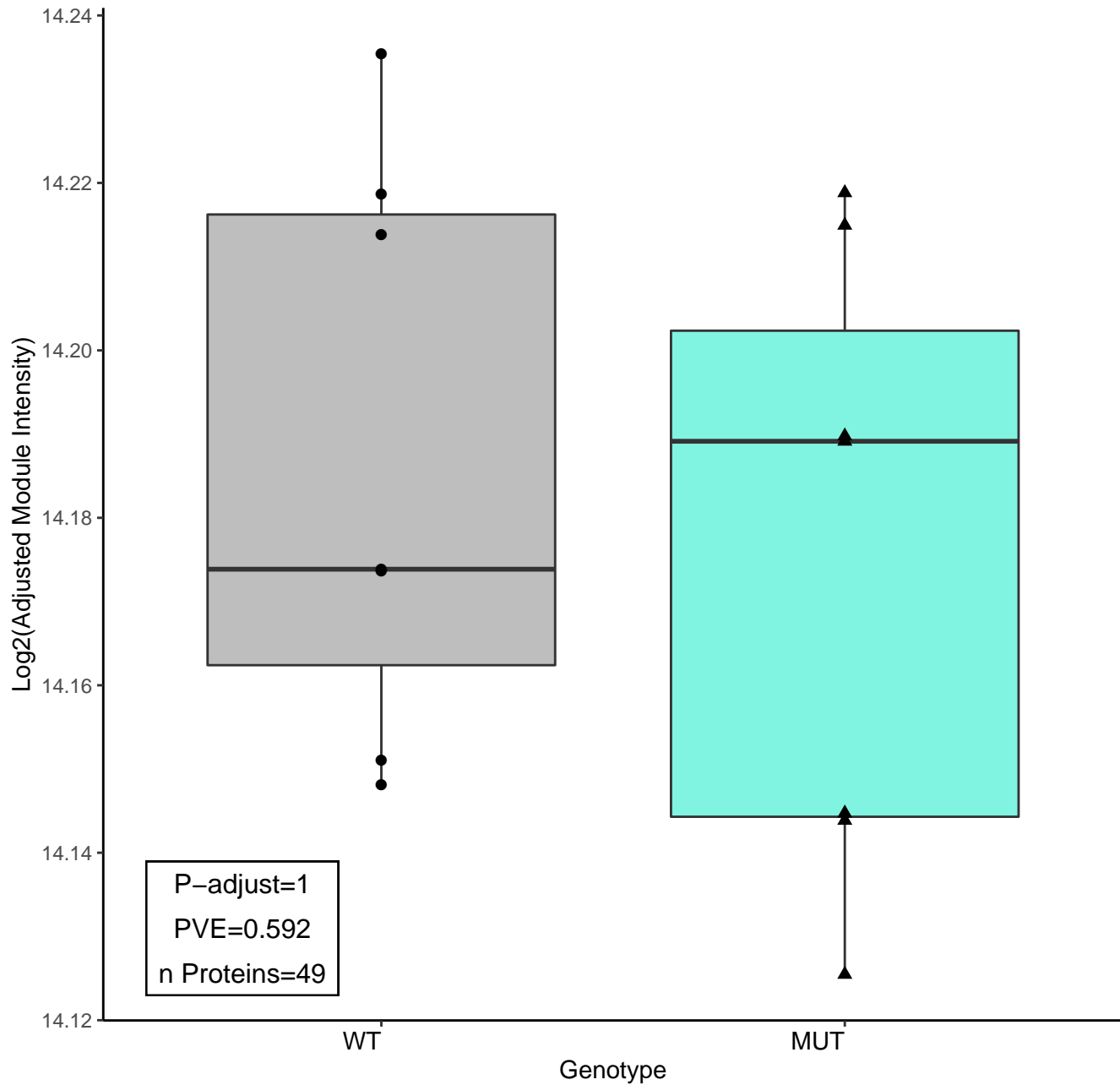
P-adjust=1

PVE=0.485

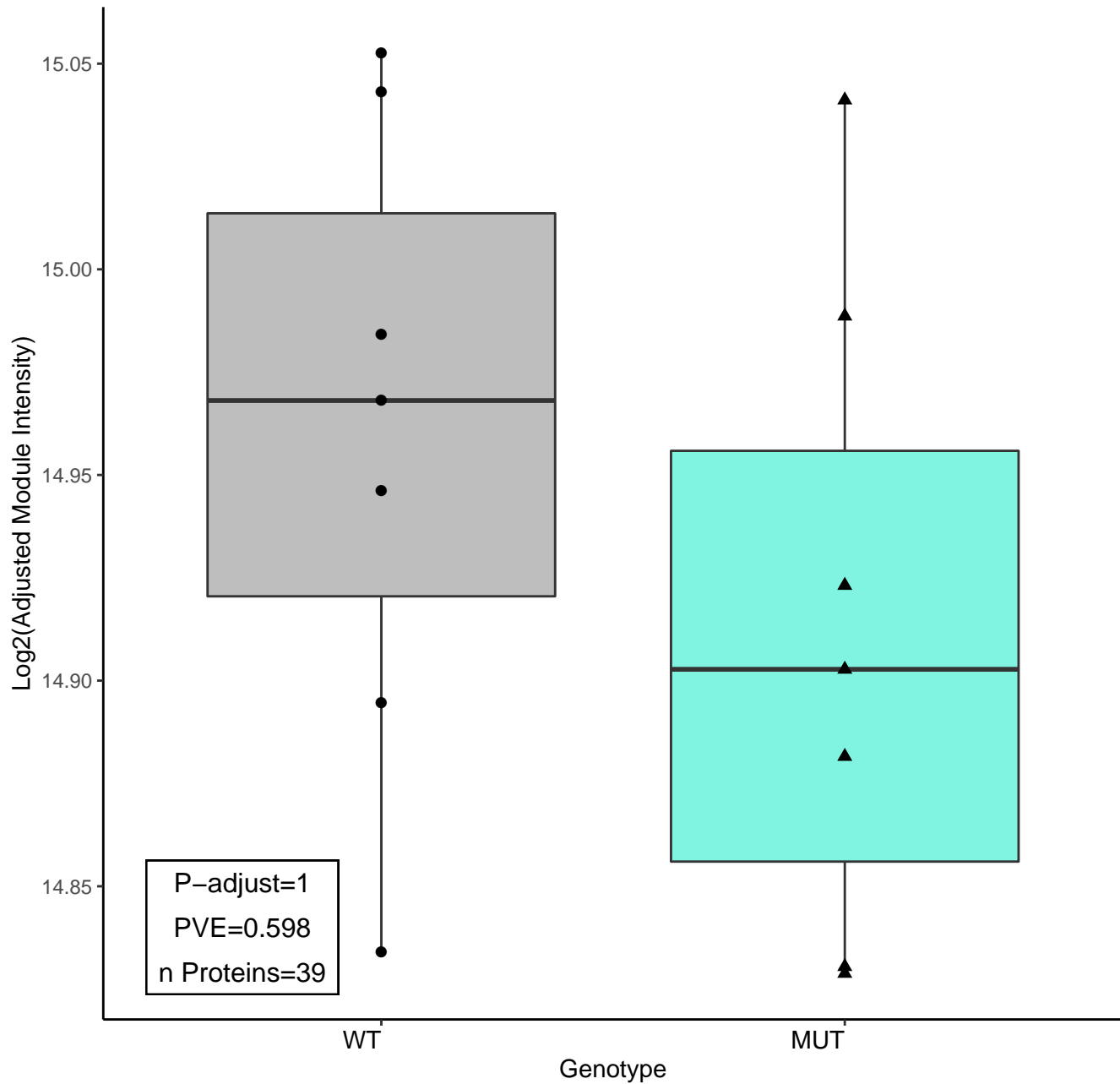
n Proteins=15

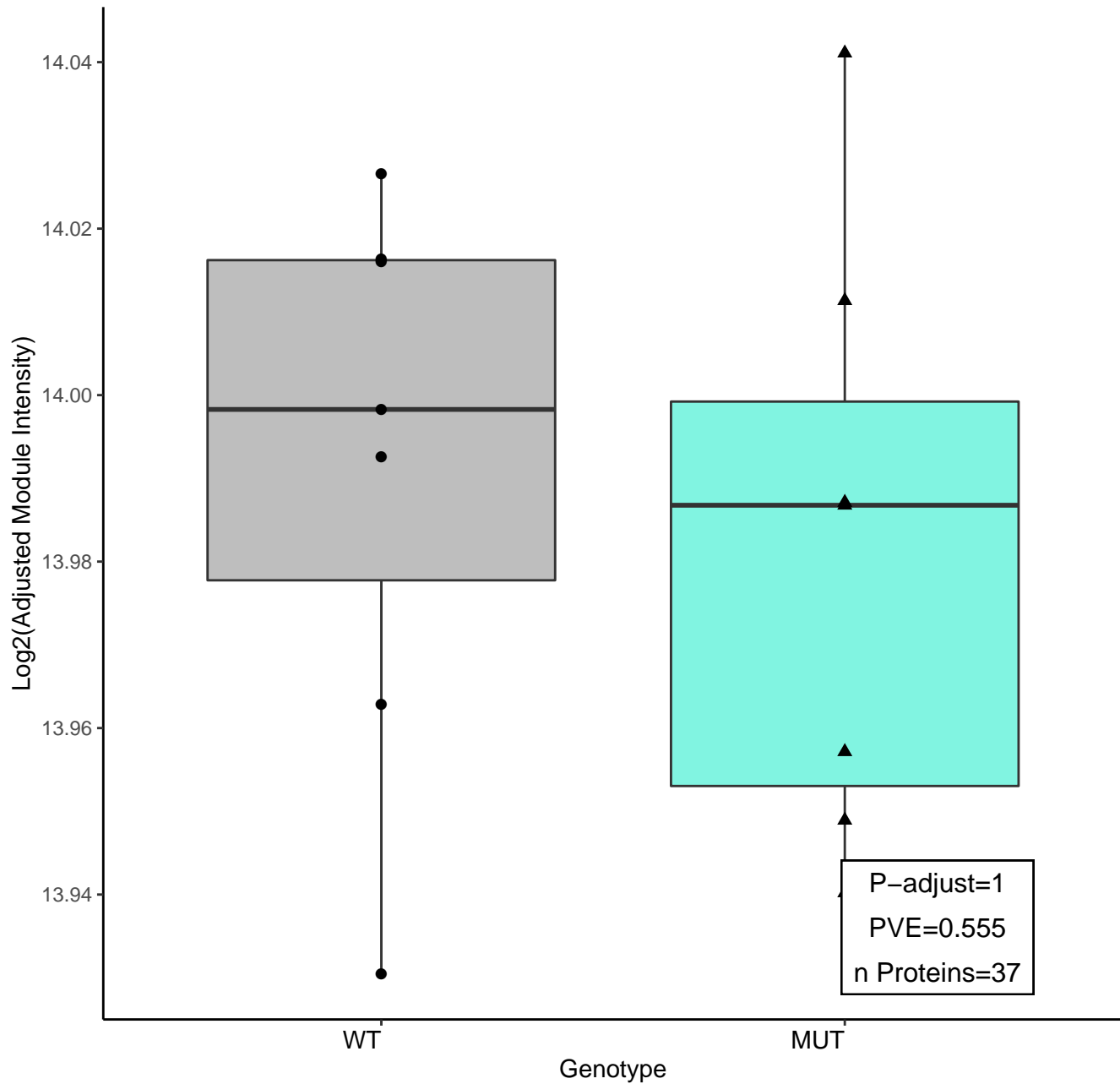


M27

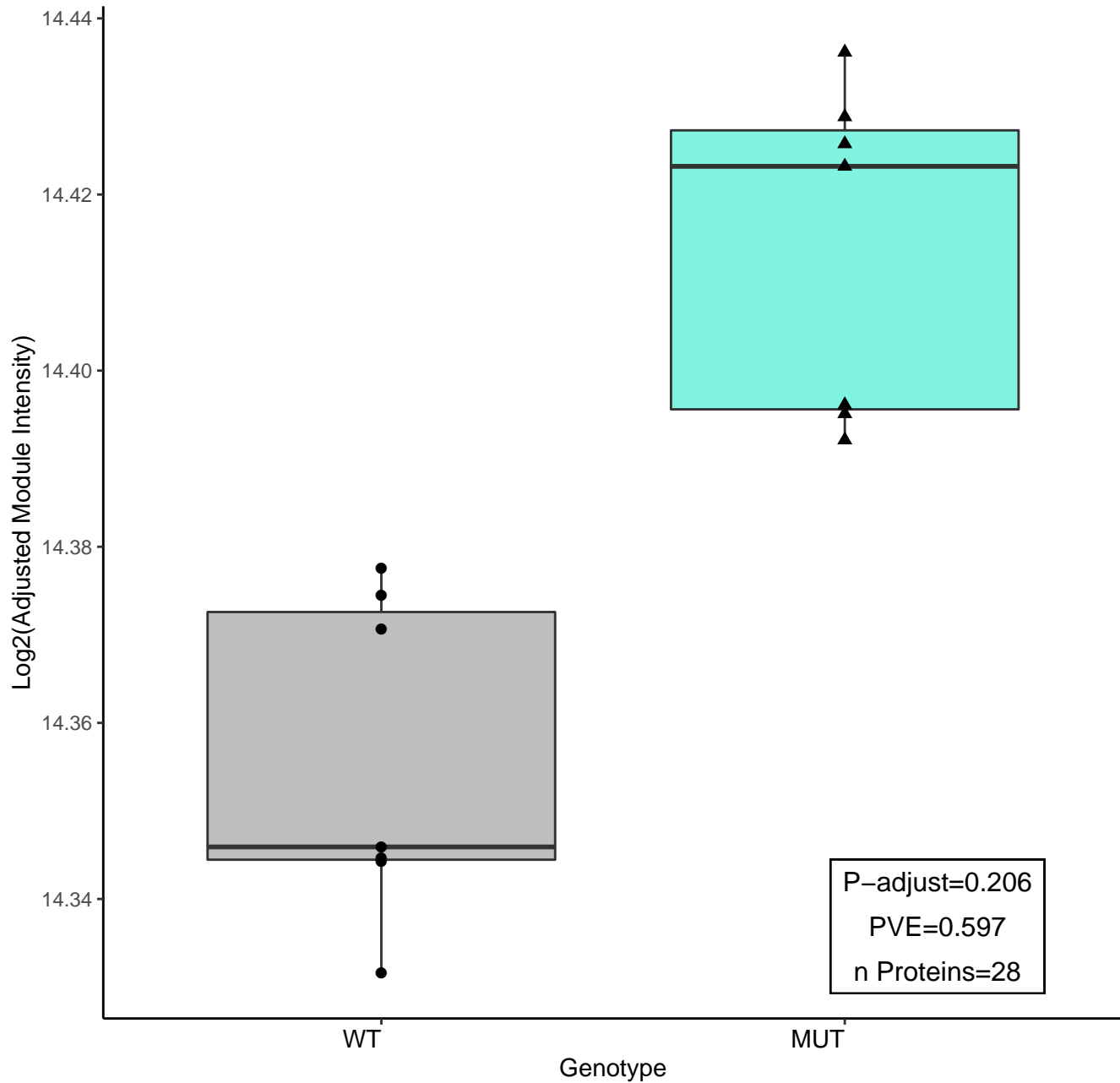


M28





M30



M31

Log2(Adjusted Module Intensity)

12.45

12.40

12.35

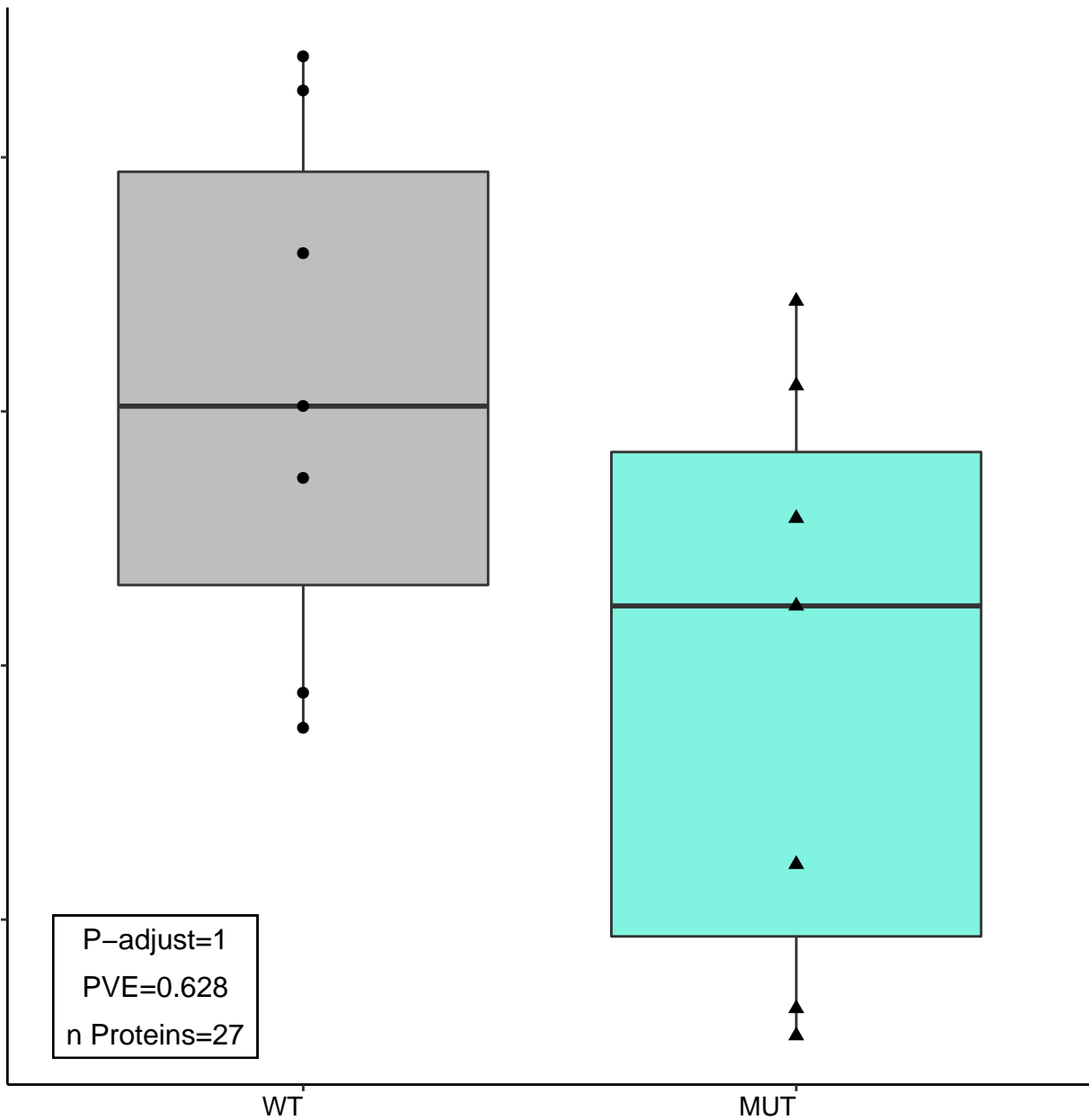
12.30

P-adjust=1
PVE=0.628
n Proteins=27

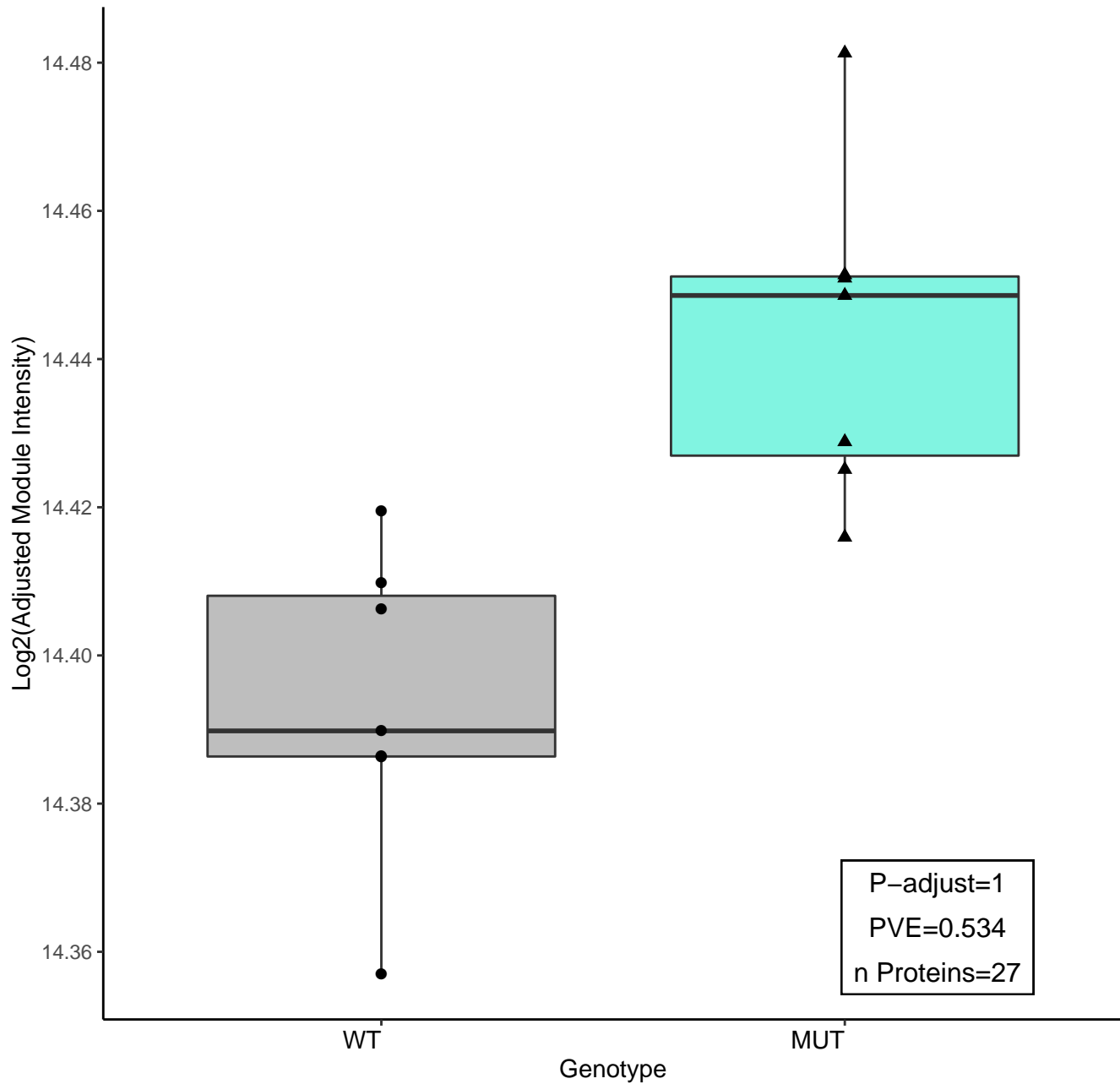
WT

Genotype

MUT



M32



M33

Log2(Adjusted Module Intensity)

12.00

11.95

11.90

WT

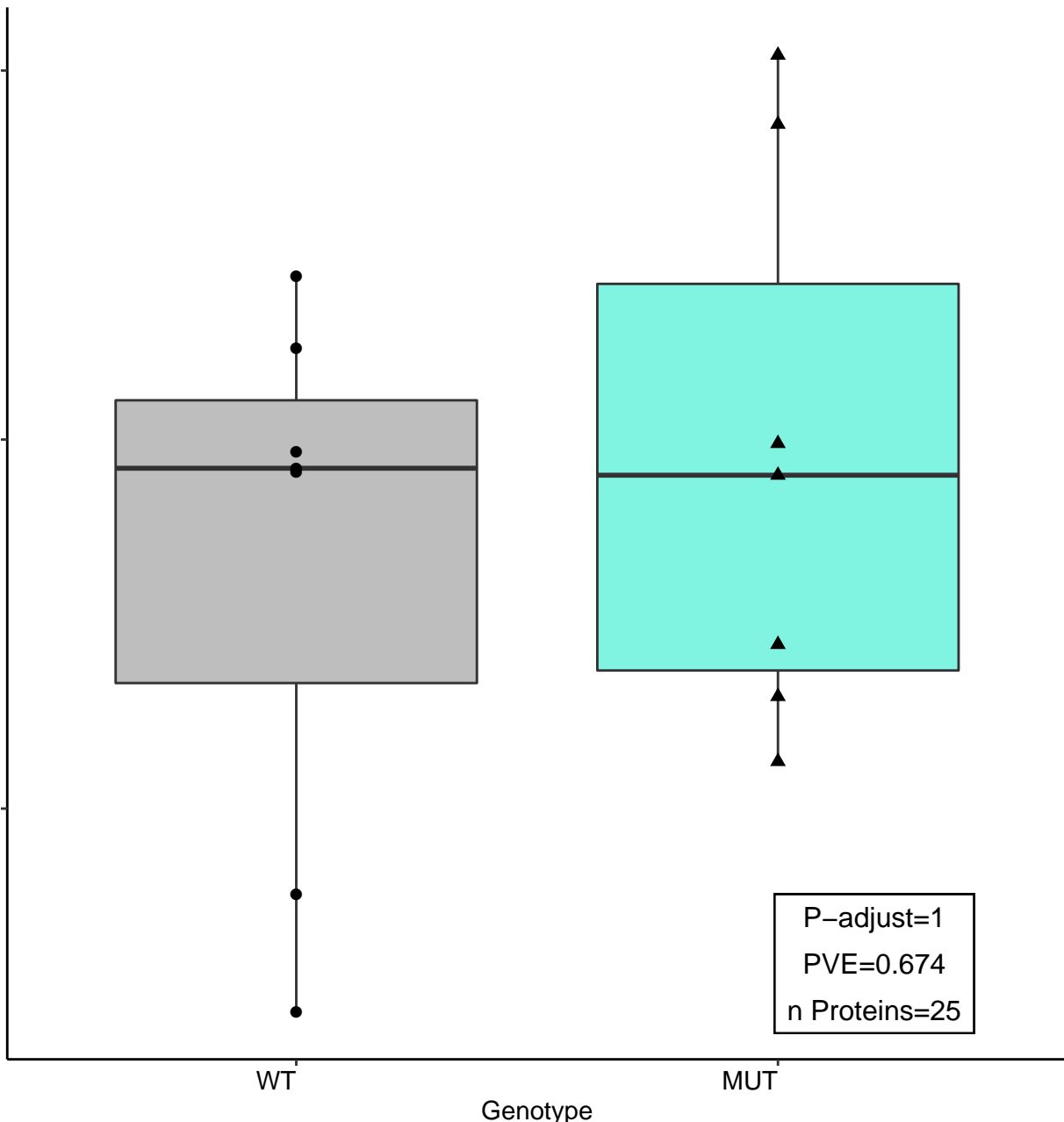
Genotype

MUT

P-adjust=1

PVE=0.674

n Proteins=25



M34

Log2(Adjusted Module Intensity)

12.35

12.30

12.25

P-adjust=1

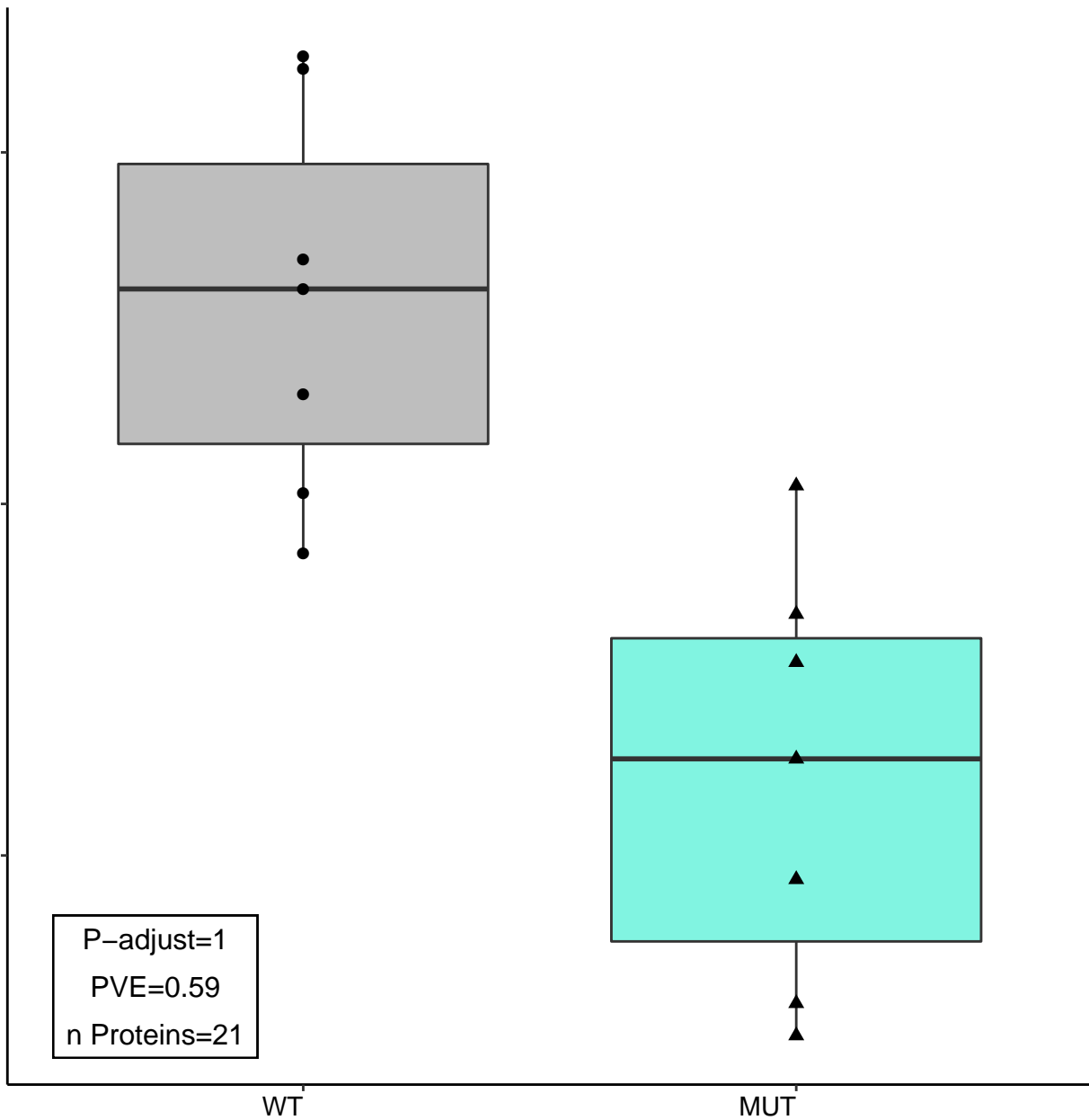
PVE=0.59

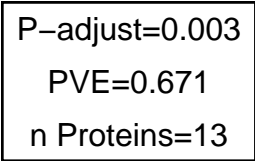
n Proteins=21

WT

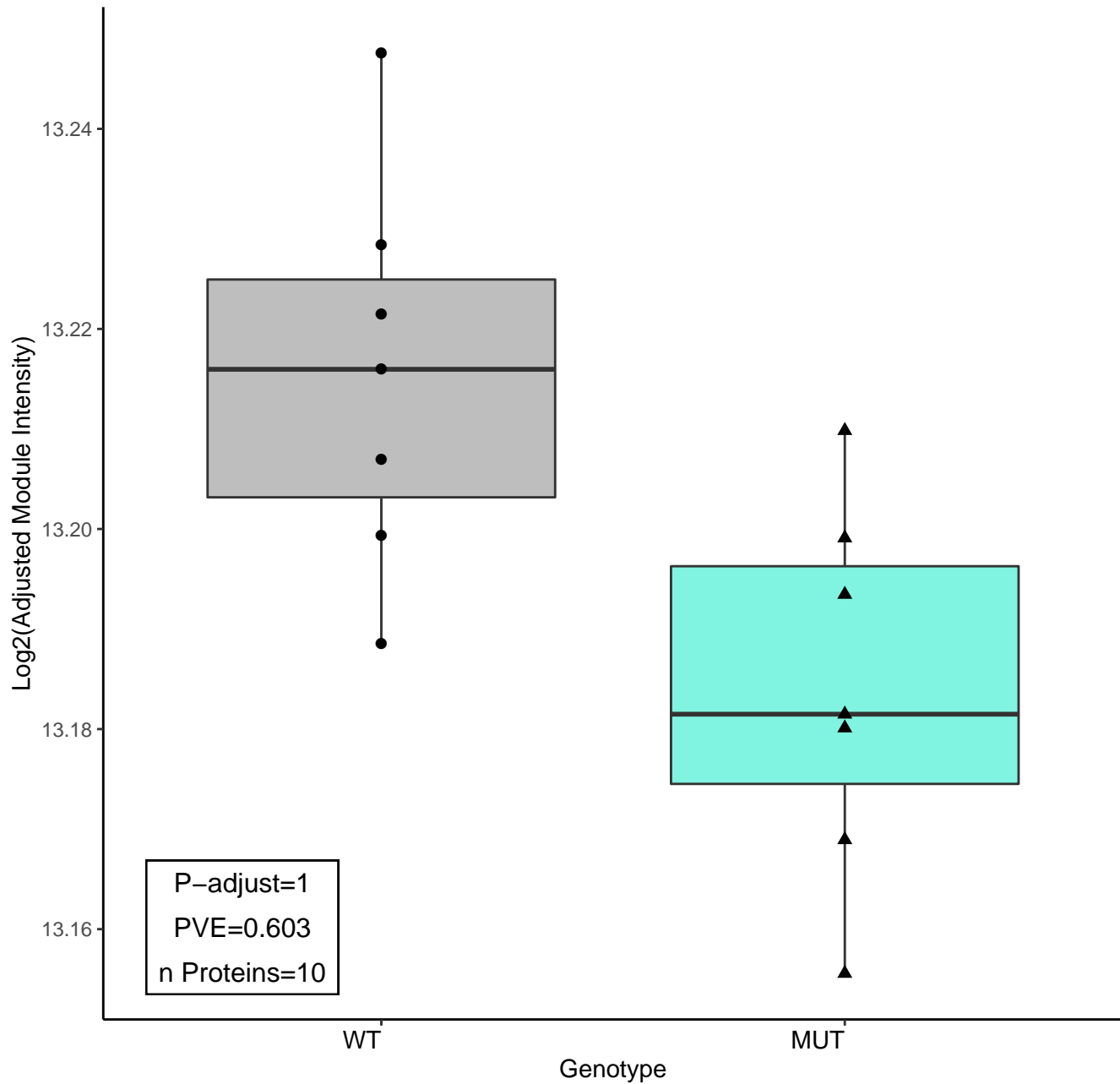
Genotype

MUT

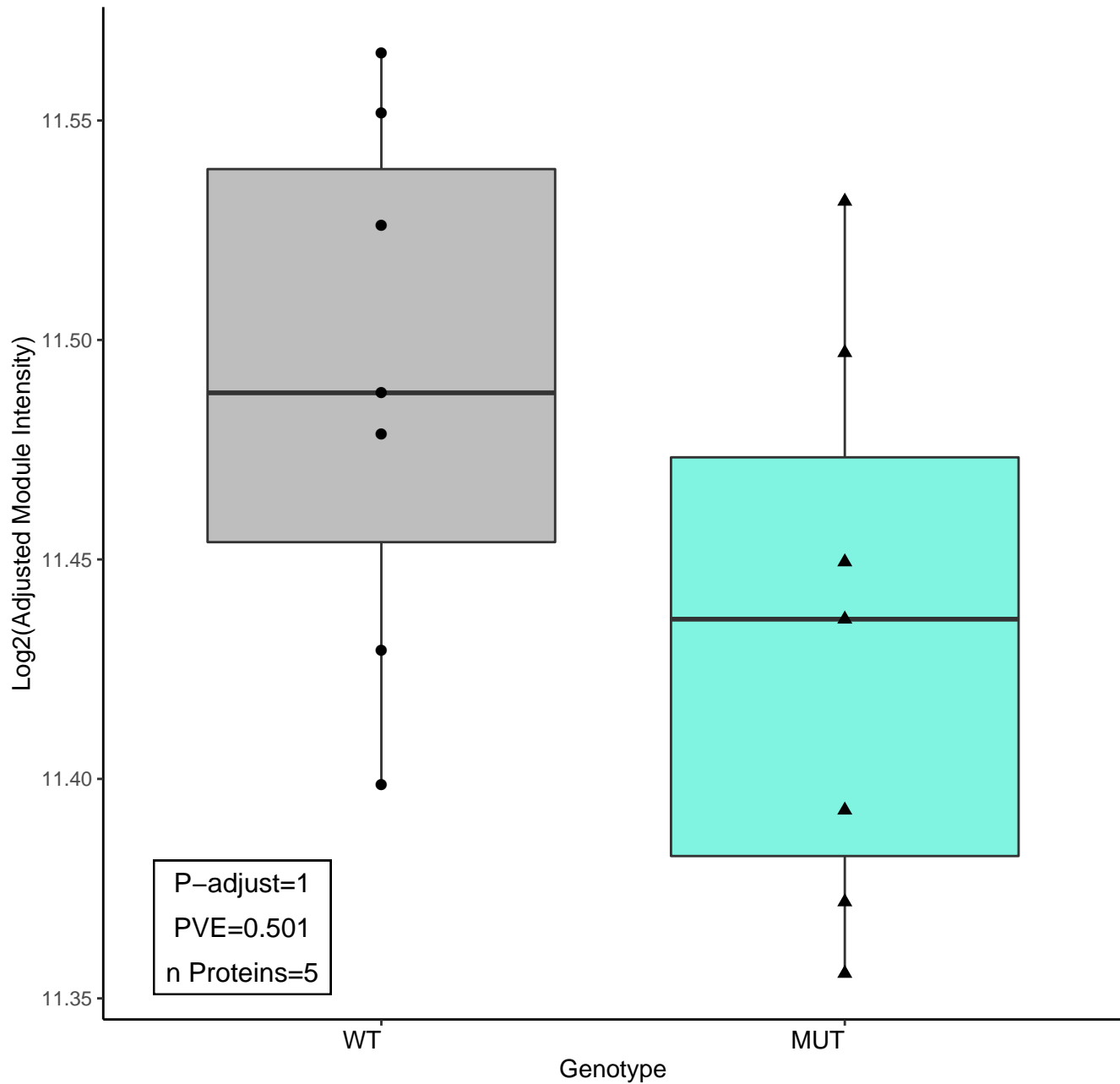




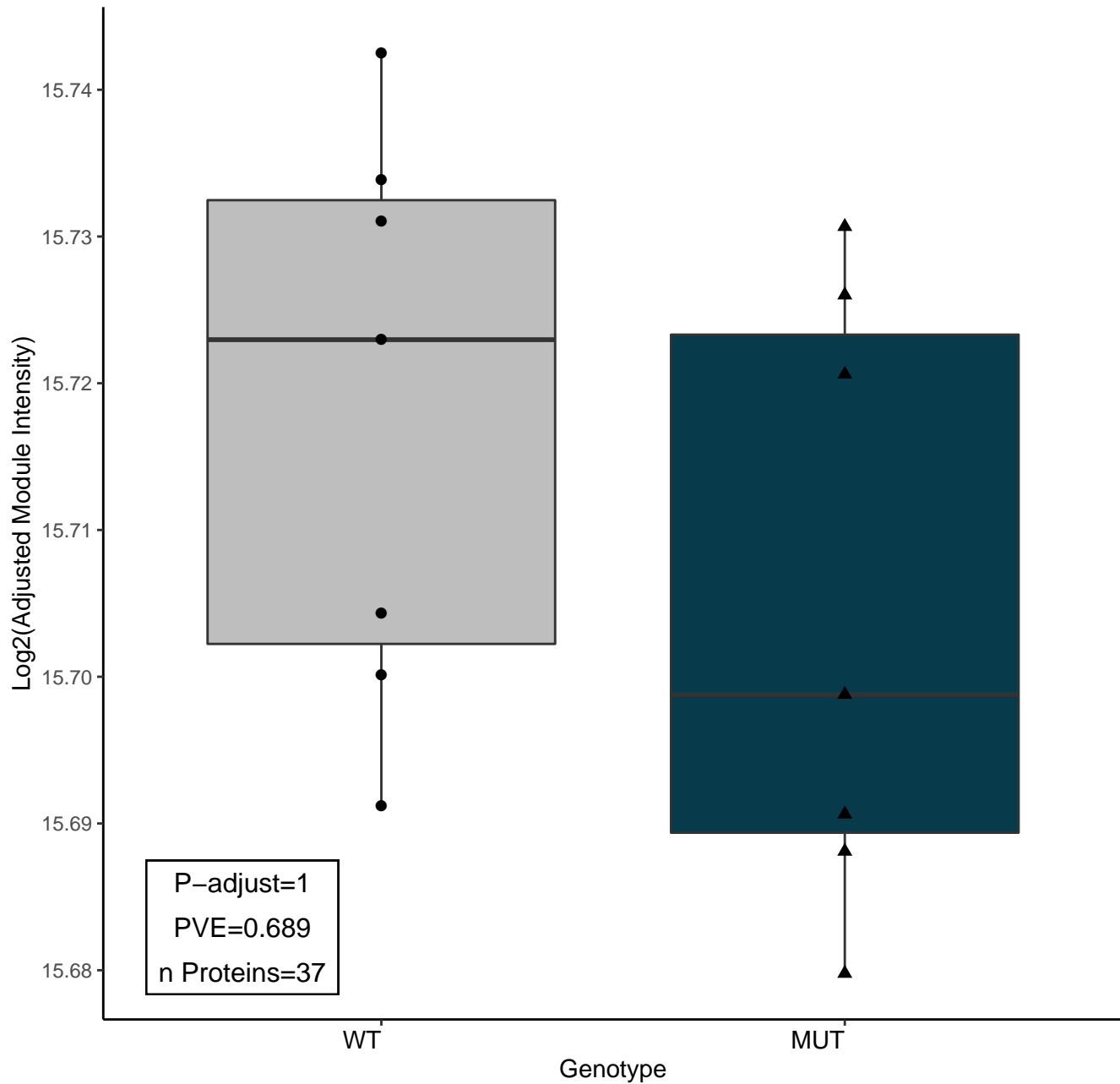
M36



M37



M38



M39

Log2(Adjusted Module Intensity)

14.50

14.48

14.46

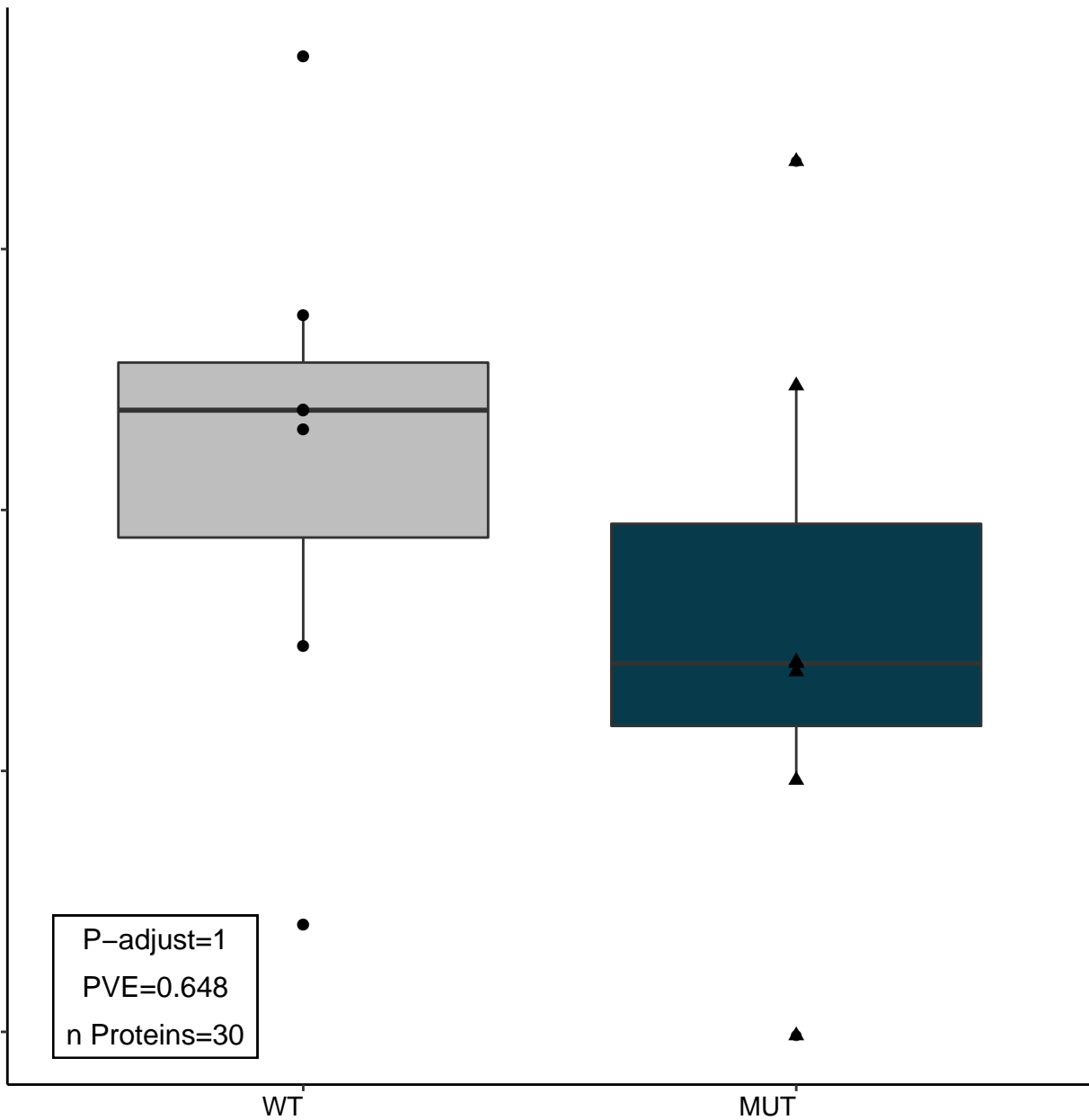
14.44

P-adjust=1
PVE=0.648
n Proteins=30

WT

Genotype

MUT



M40

Log2(Adjusted Module Intensity)

13.42

13.40

13.38

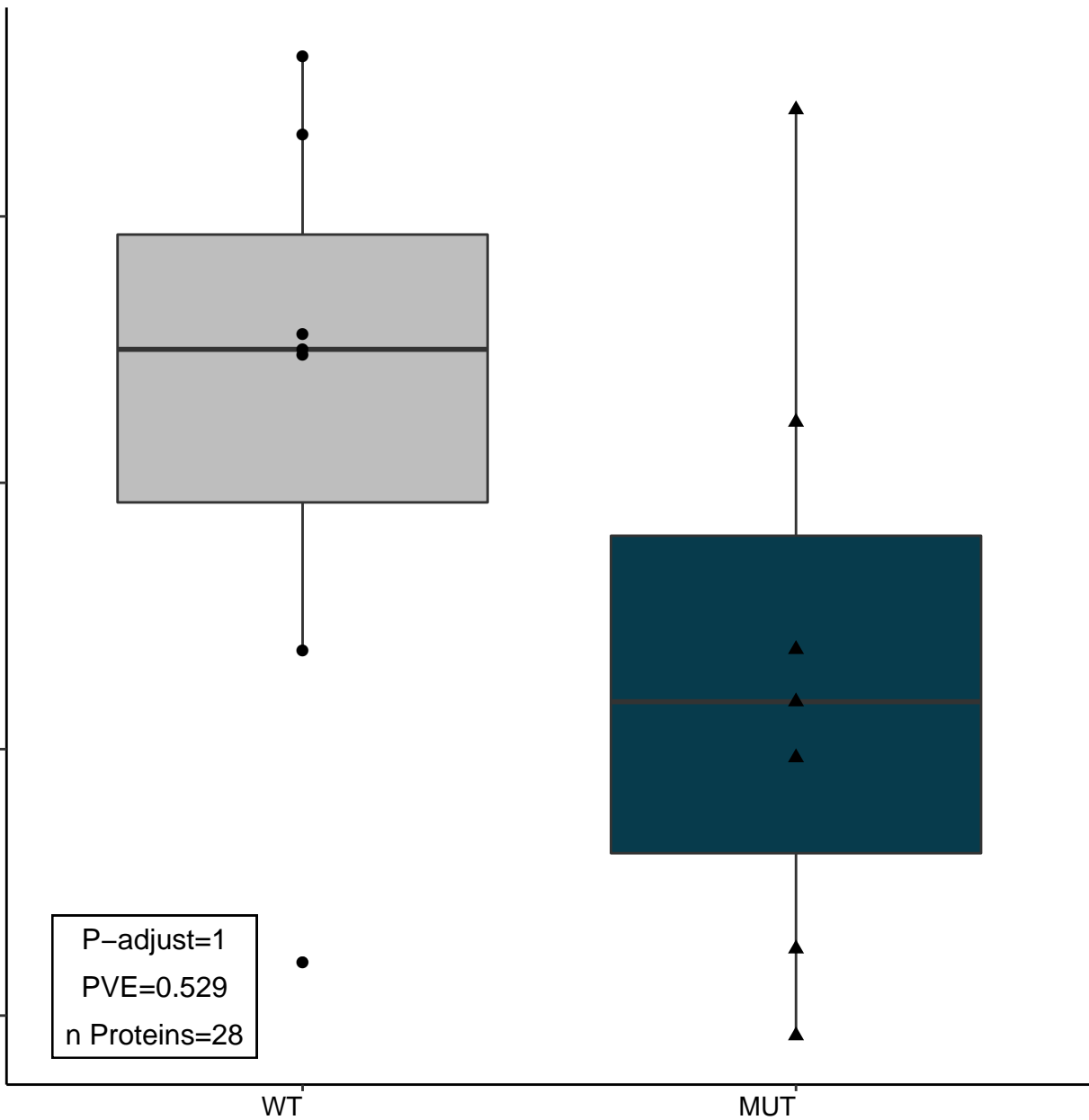
13.36

P-adjust=1
PVE=0.529
n Proteins=28

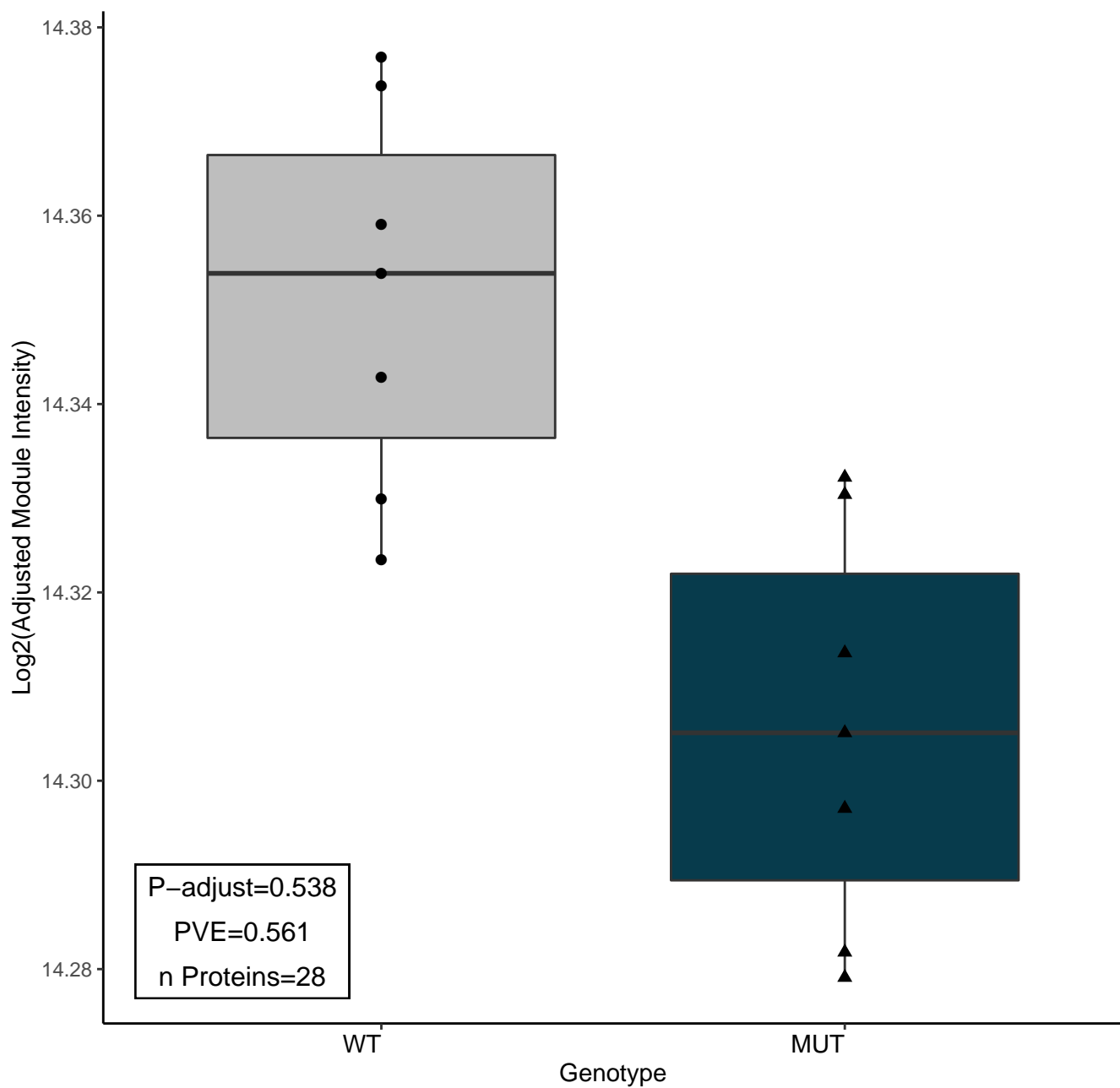
WT

Genotype

MUT



M41



M42

Log2(Adjusted Module Intensity)

14.45

14.40

14.35

WT

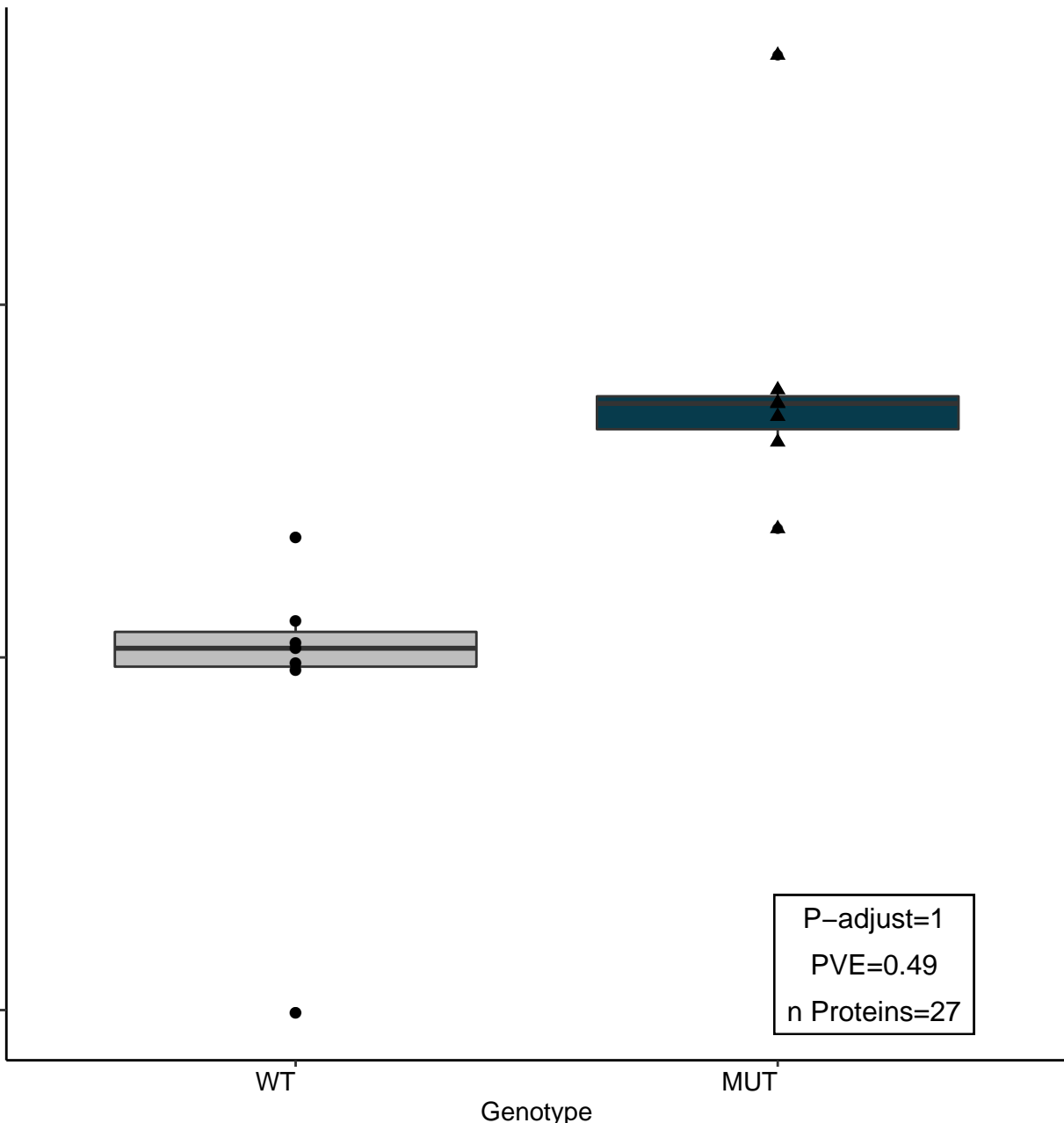
Genotype

MUT

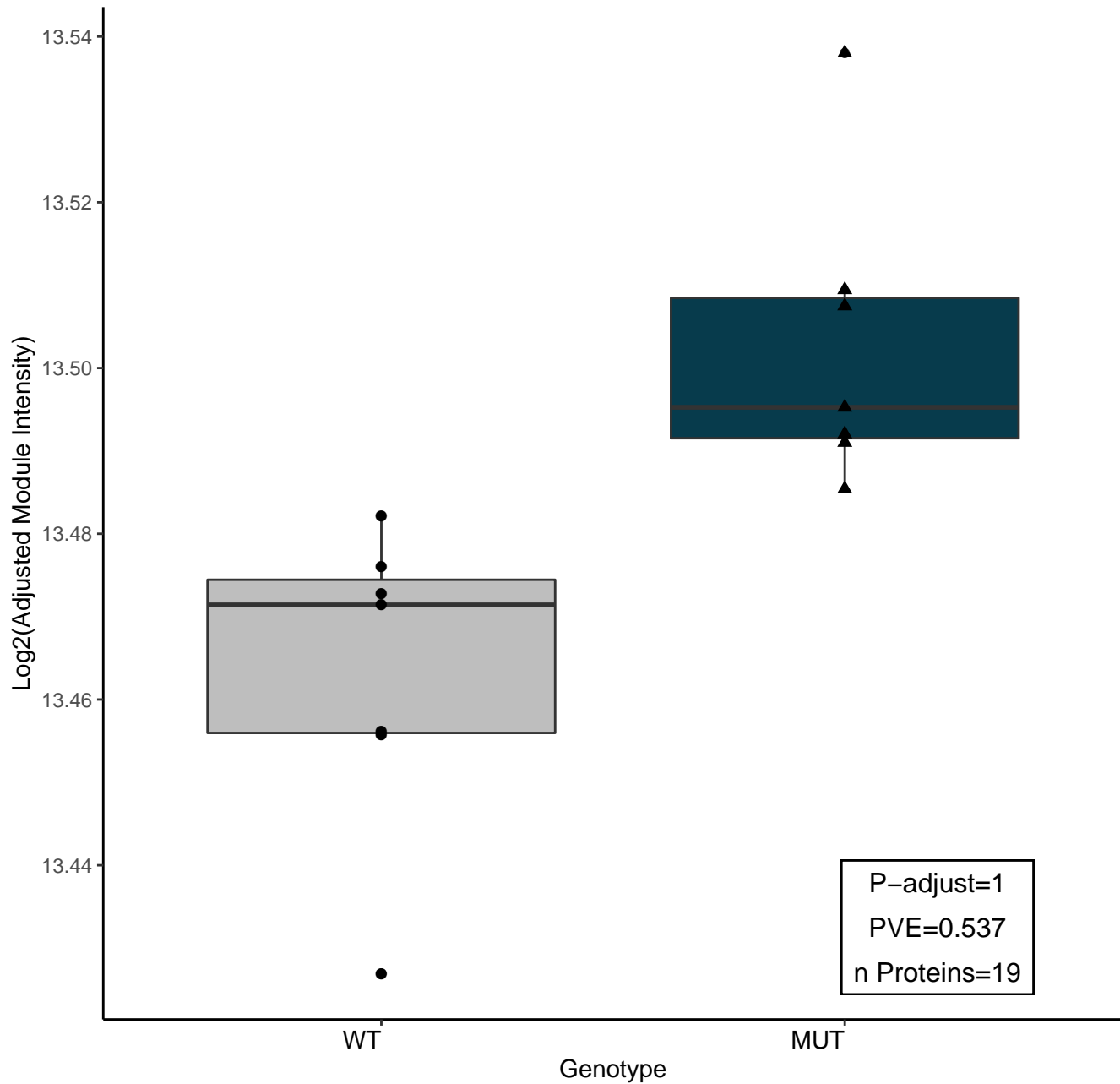
P-adjust=1

PVE=0.49

n Proteins=27



M43



M44

Log2(Adjusted Module Intensity)

13.94

13.92

13.90

WT

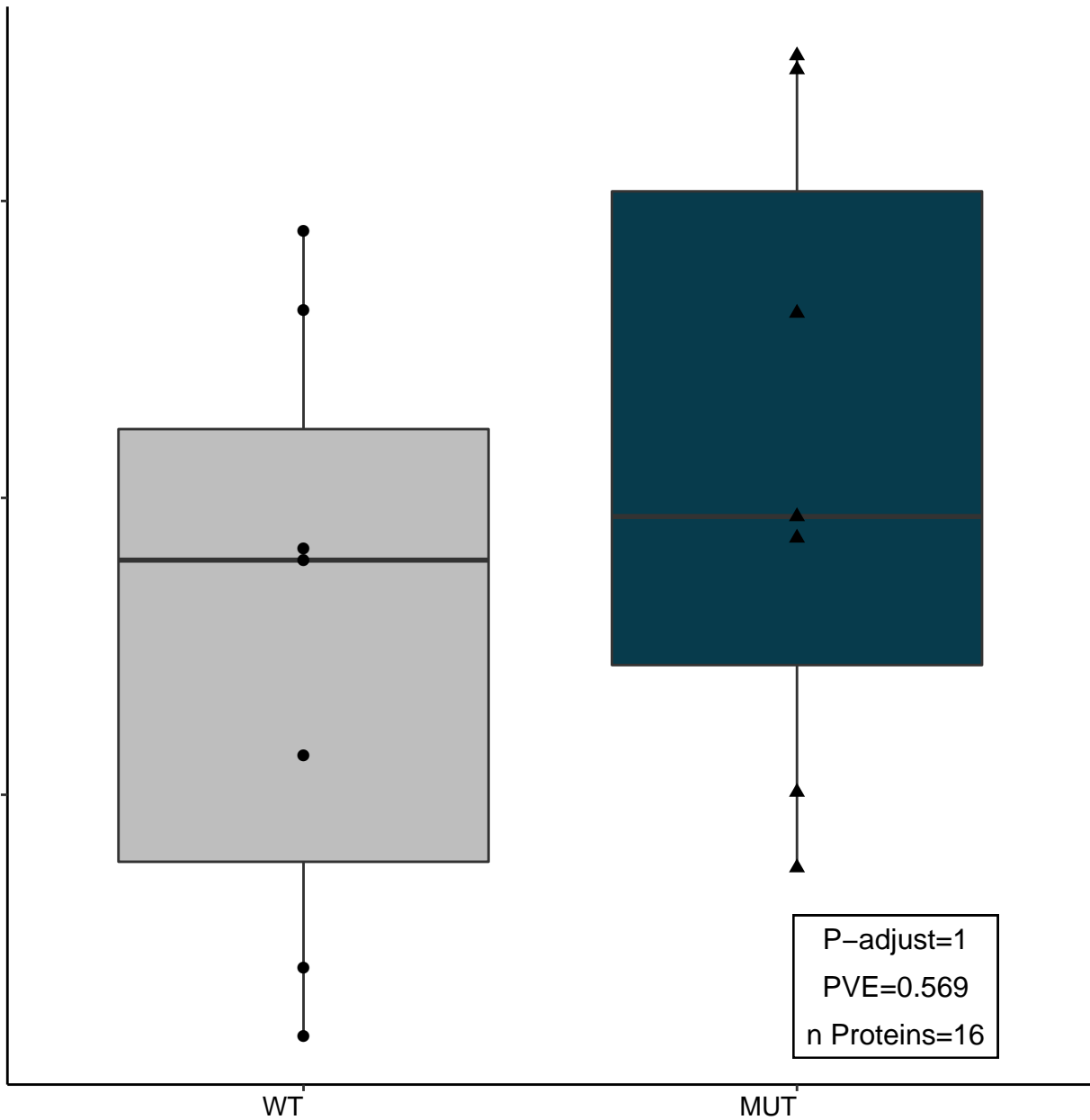
Genotype

MUT

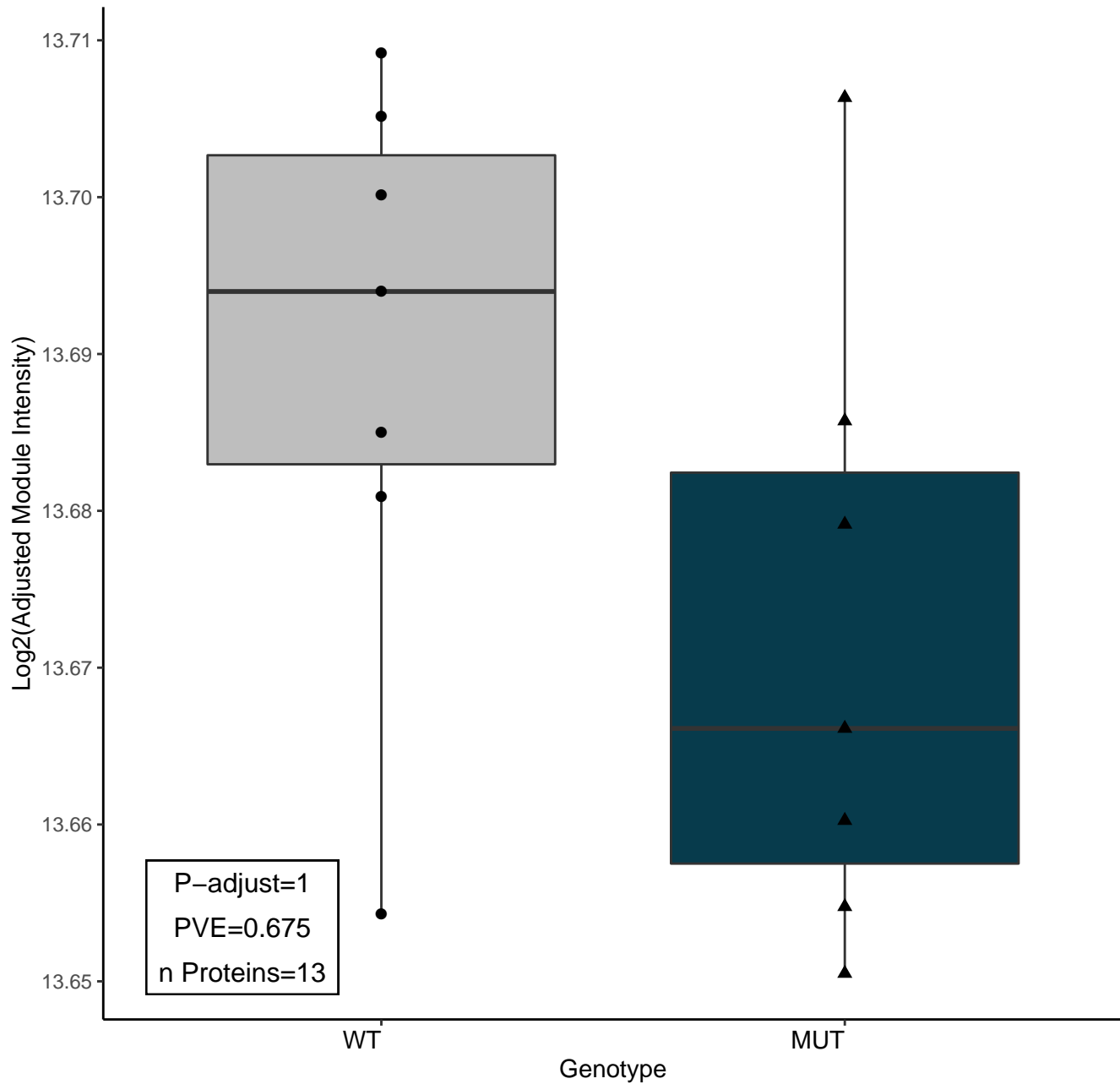
P-adjust=1

PVE=0.569

n Proteins=16



M45



M46

Log2(Adjusted Module Intensity)

12.58

12.56

12.54

12.52

12.50

12.48

WT

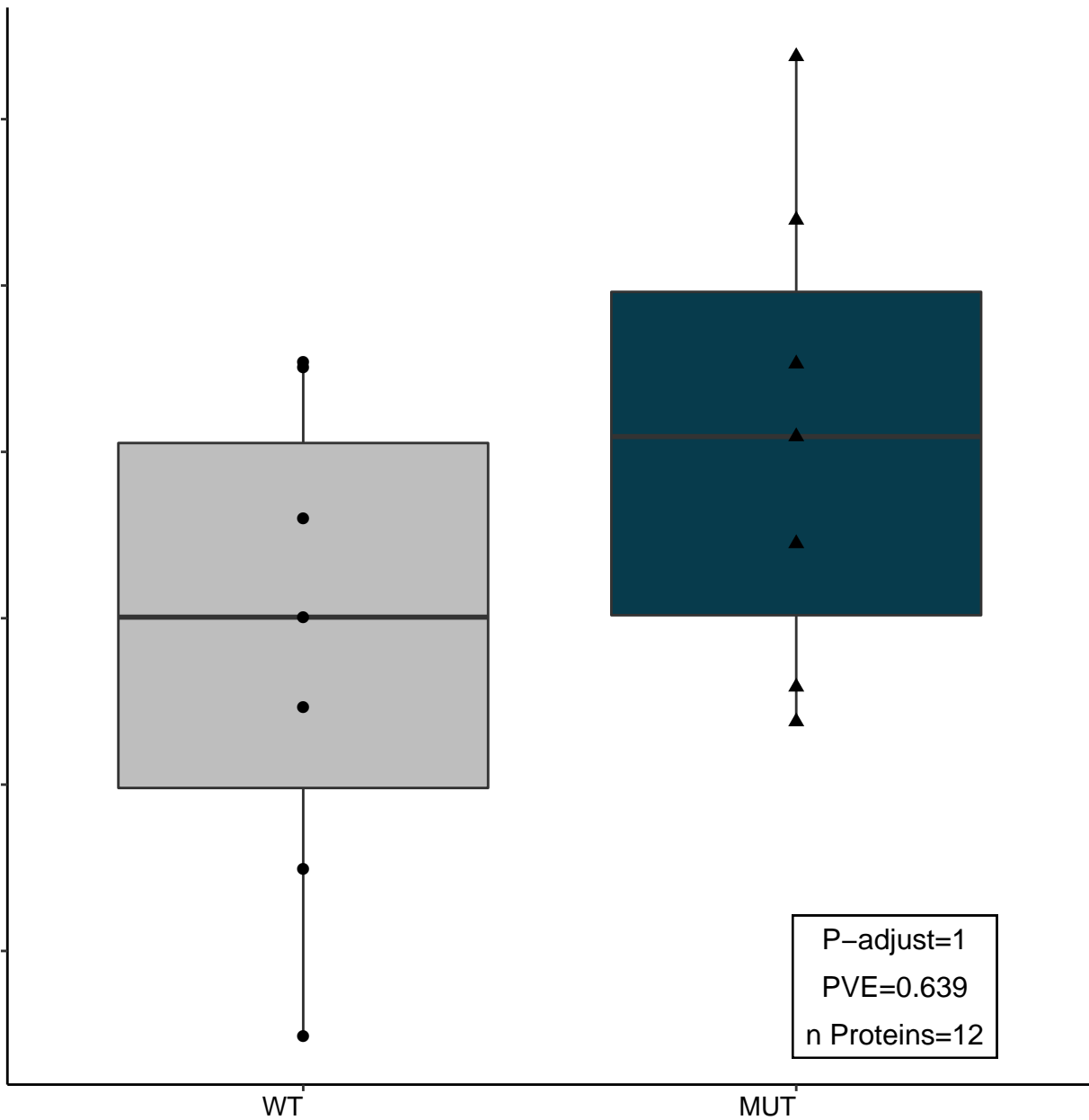
Genotype

MUT

P-adjust=1

PVE=0.639

n Proteins=12



M47

Log2(Adjusted Module Intensity)

12.55

12.50

12.45

P-adjust=0.138

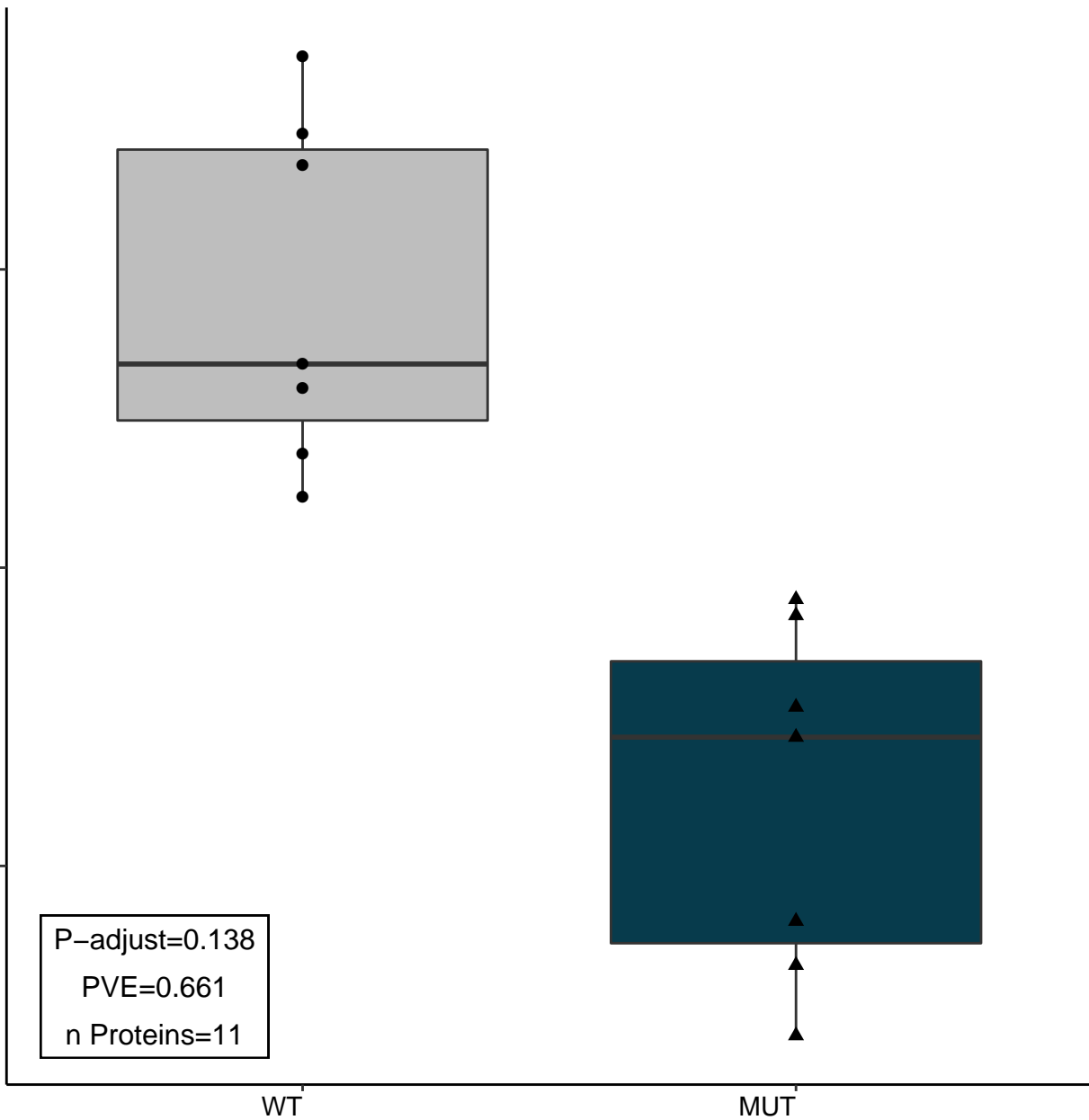
PVE=0.661

n Proteins=11

WT

Genotype

MUT



M48

Log2(Adjusted Module Intensity)

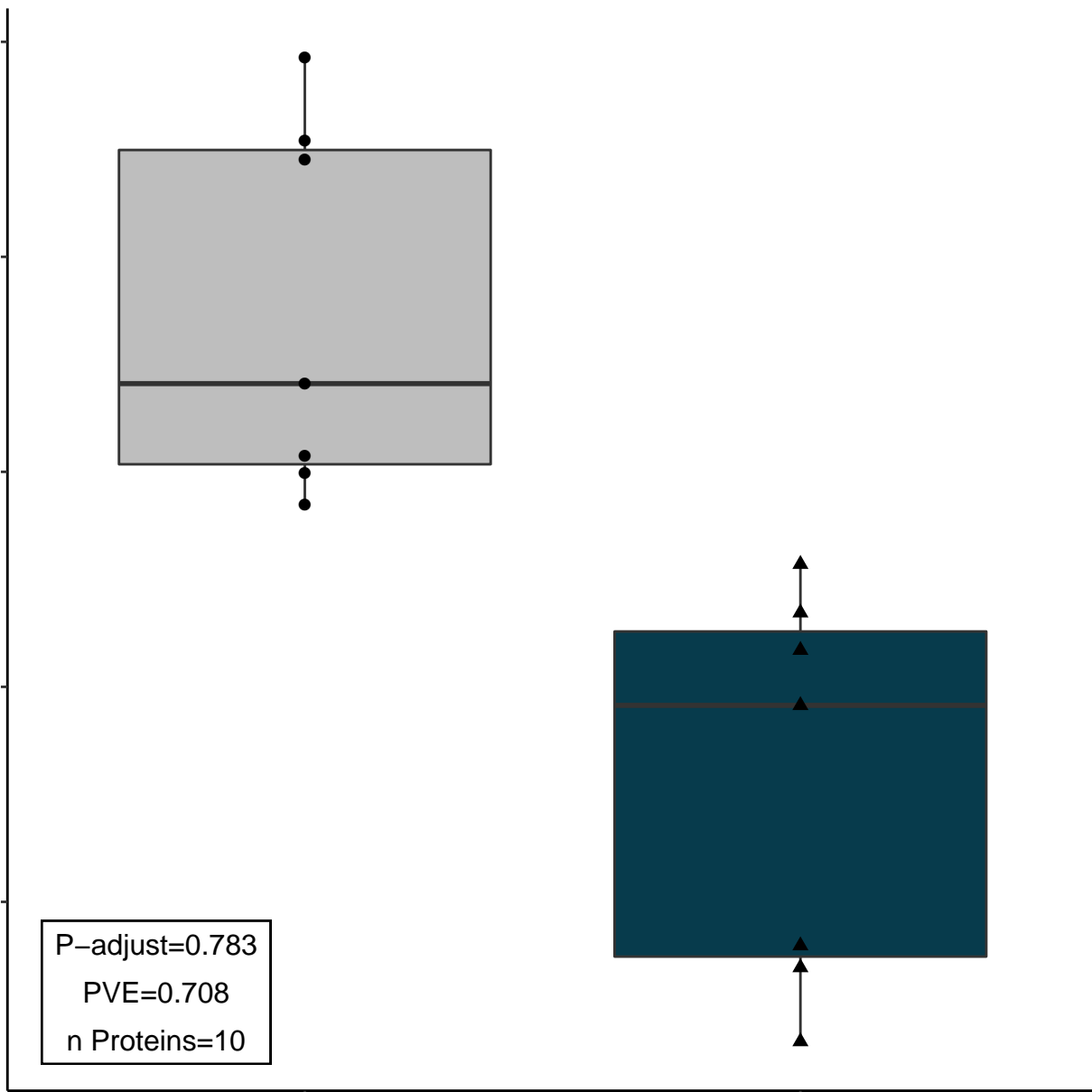
12.40
12.38
12.36
12.34
12.32

P-adjust=0.783
PVE=0.708
n Proteins=10

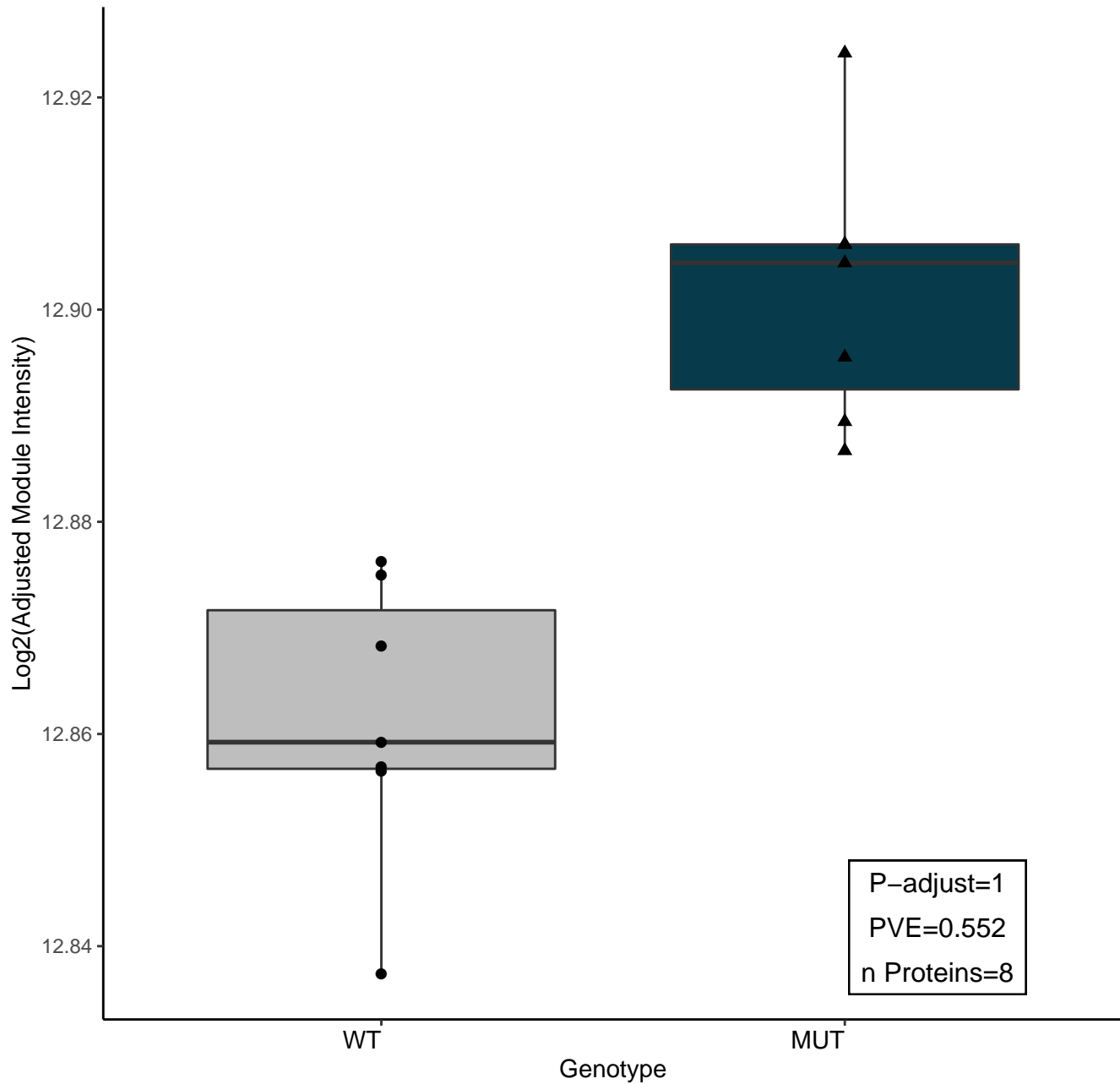
WT

Genotype

MUT



M49



M50

Log2(Adjusted Module Intensity)

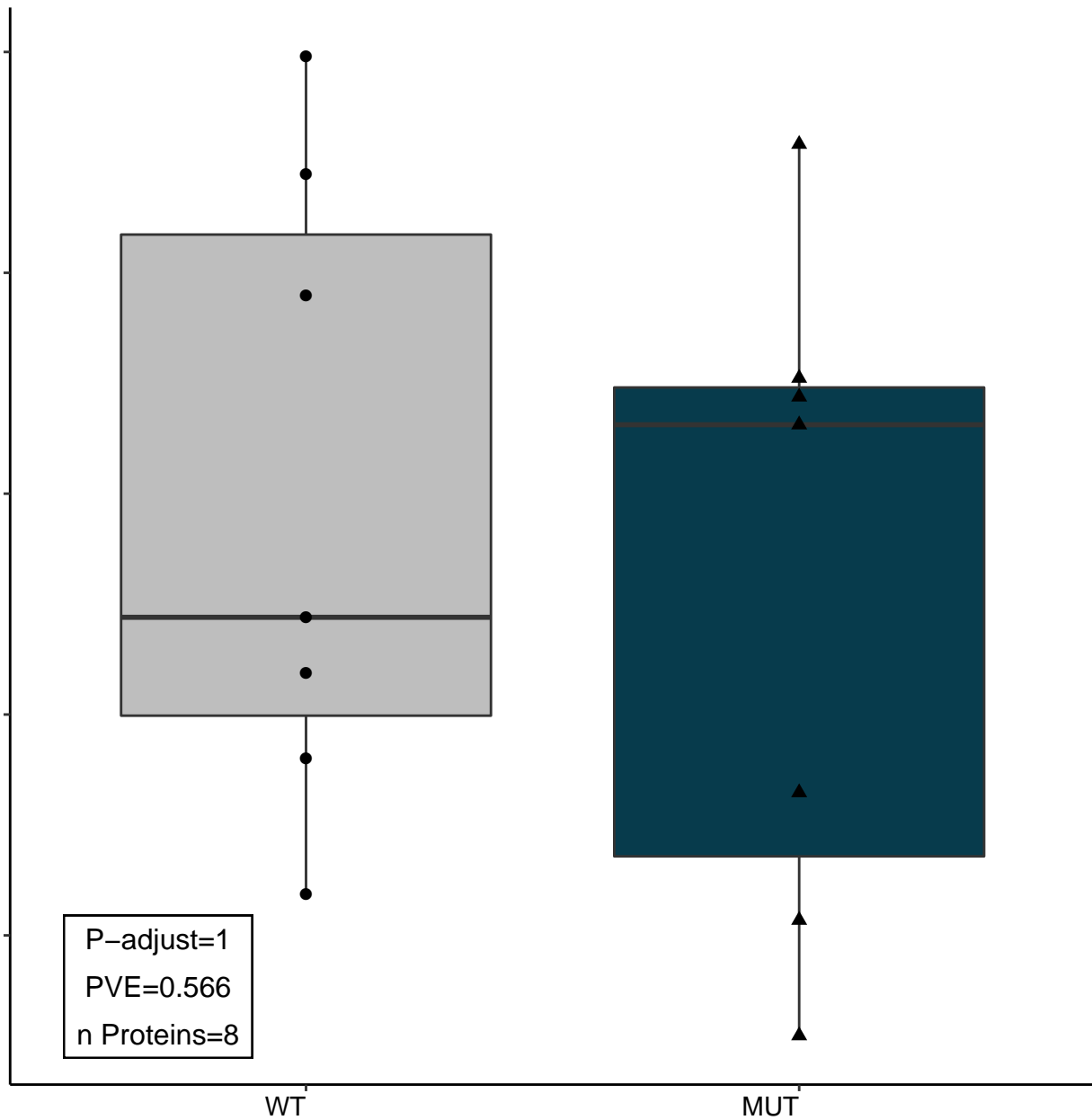
11.41
11.40
11.39
11.38
11.37

P-adjust=1
PVE=0.566
n Proteins=8

WT

Genotype

MUT



M51

Log2(Adjusted Module Intensity)

11.26

11.25

11.24

11.23

11.22

P-adjust=1

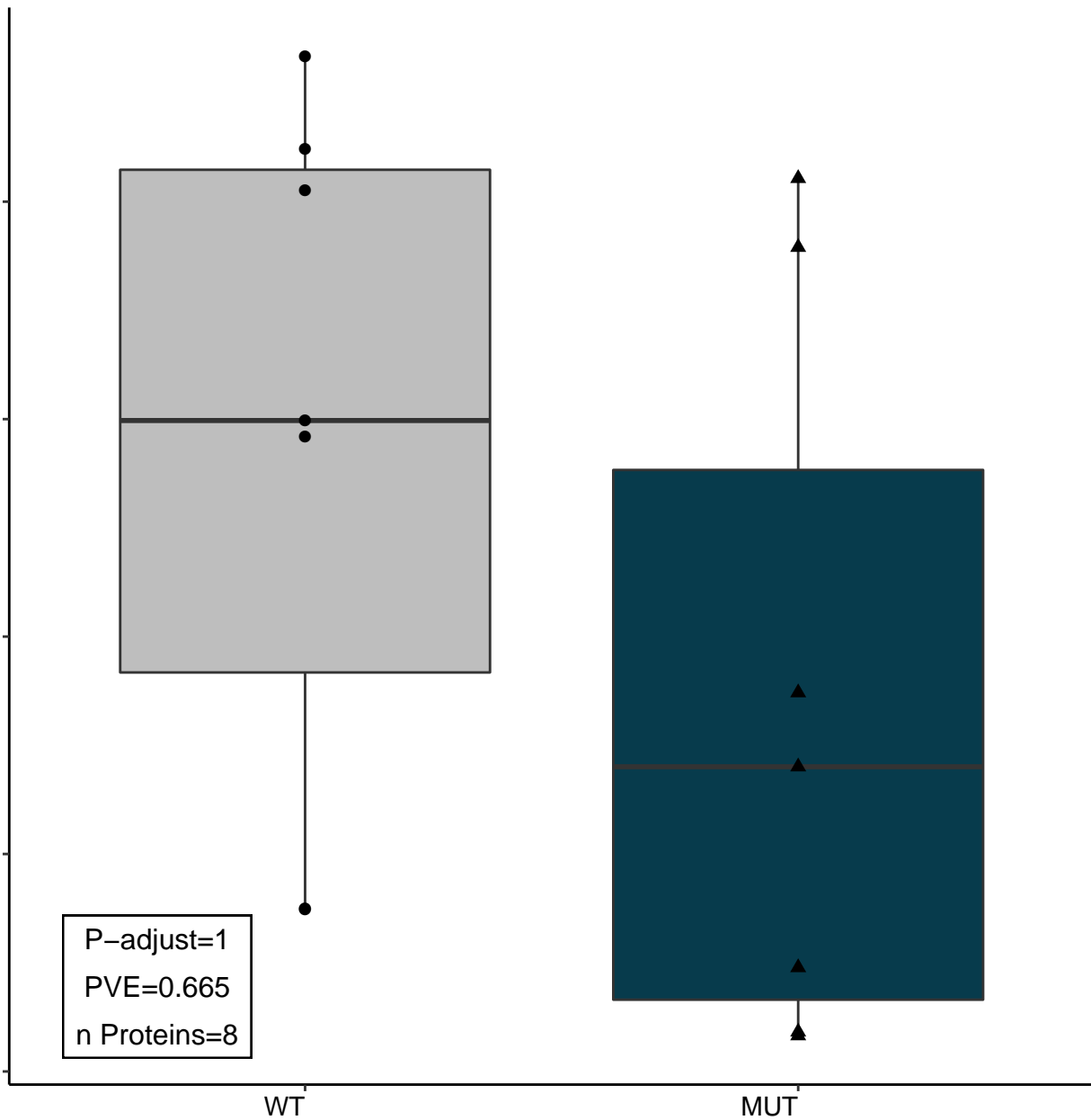
PVE=0.665

n Proteins=8

WT

Genotype

MUT



M52

Log2(Adjusted Module Intensity)

11.25

11.24

11.23

11.22

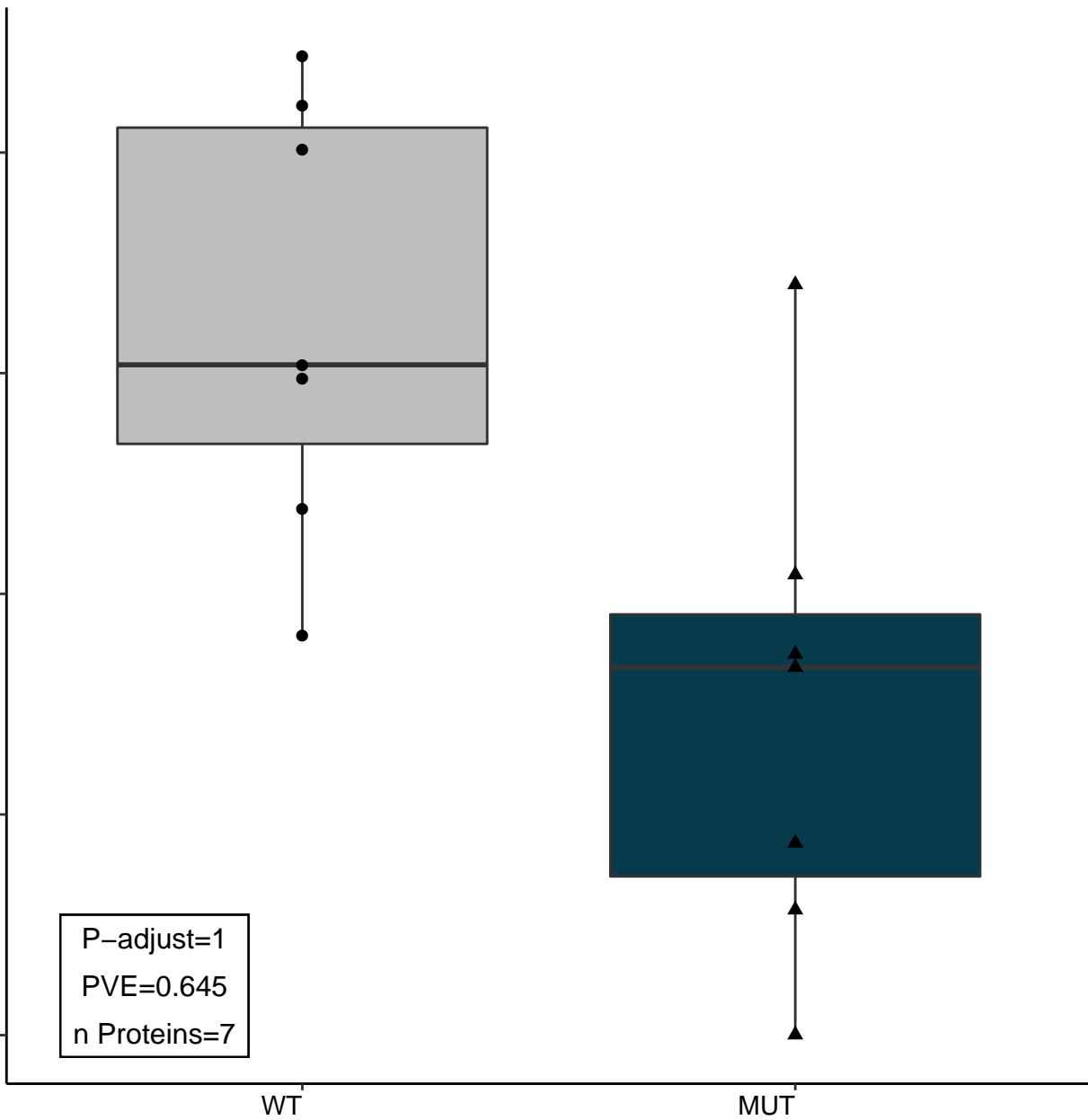
11.21

P-adjust=1
PVE=0.645
n Proteins=7

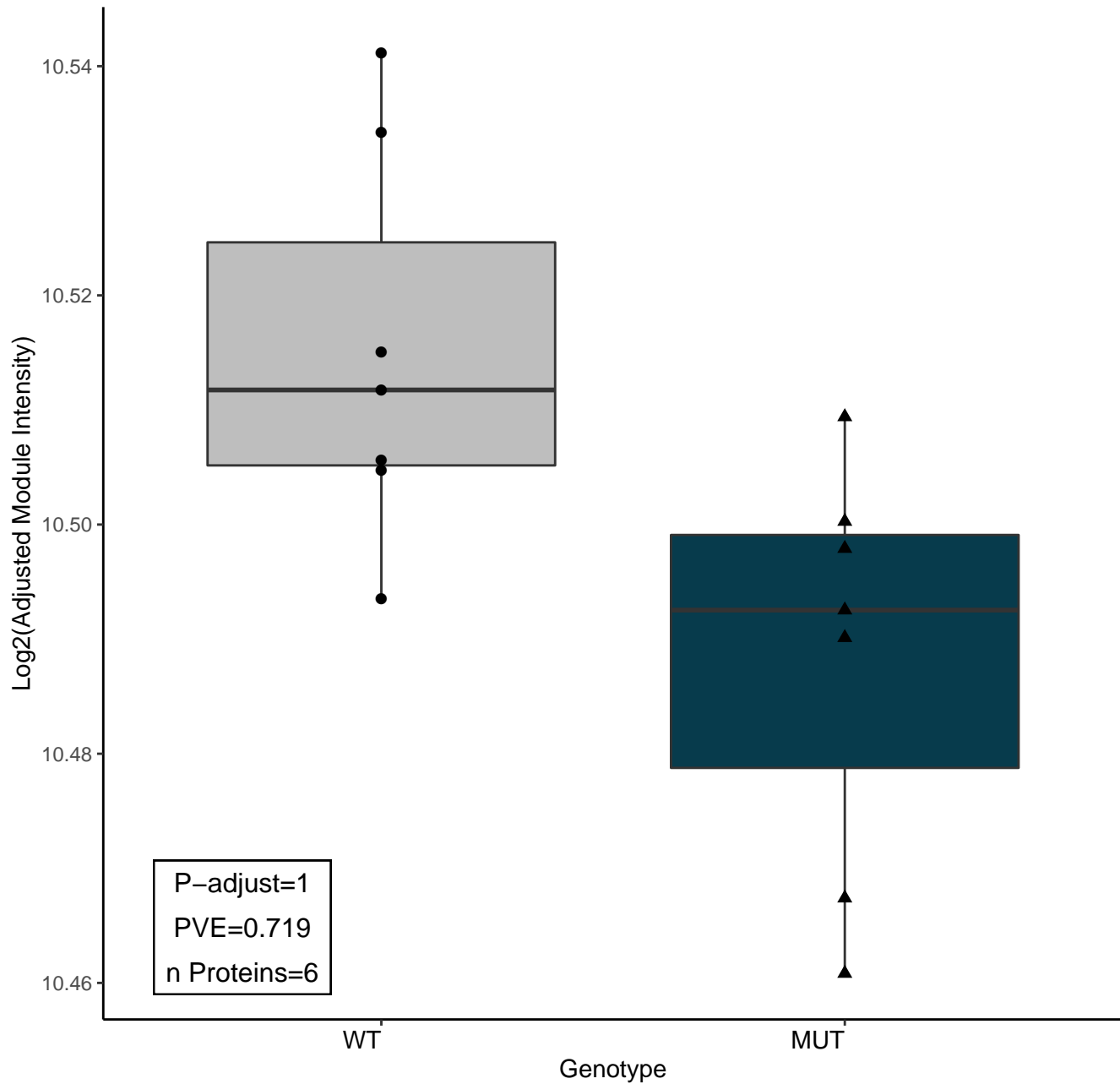
WT

Genotype

MUT



M53



M54

Log2(Adjusted Module Intensity)

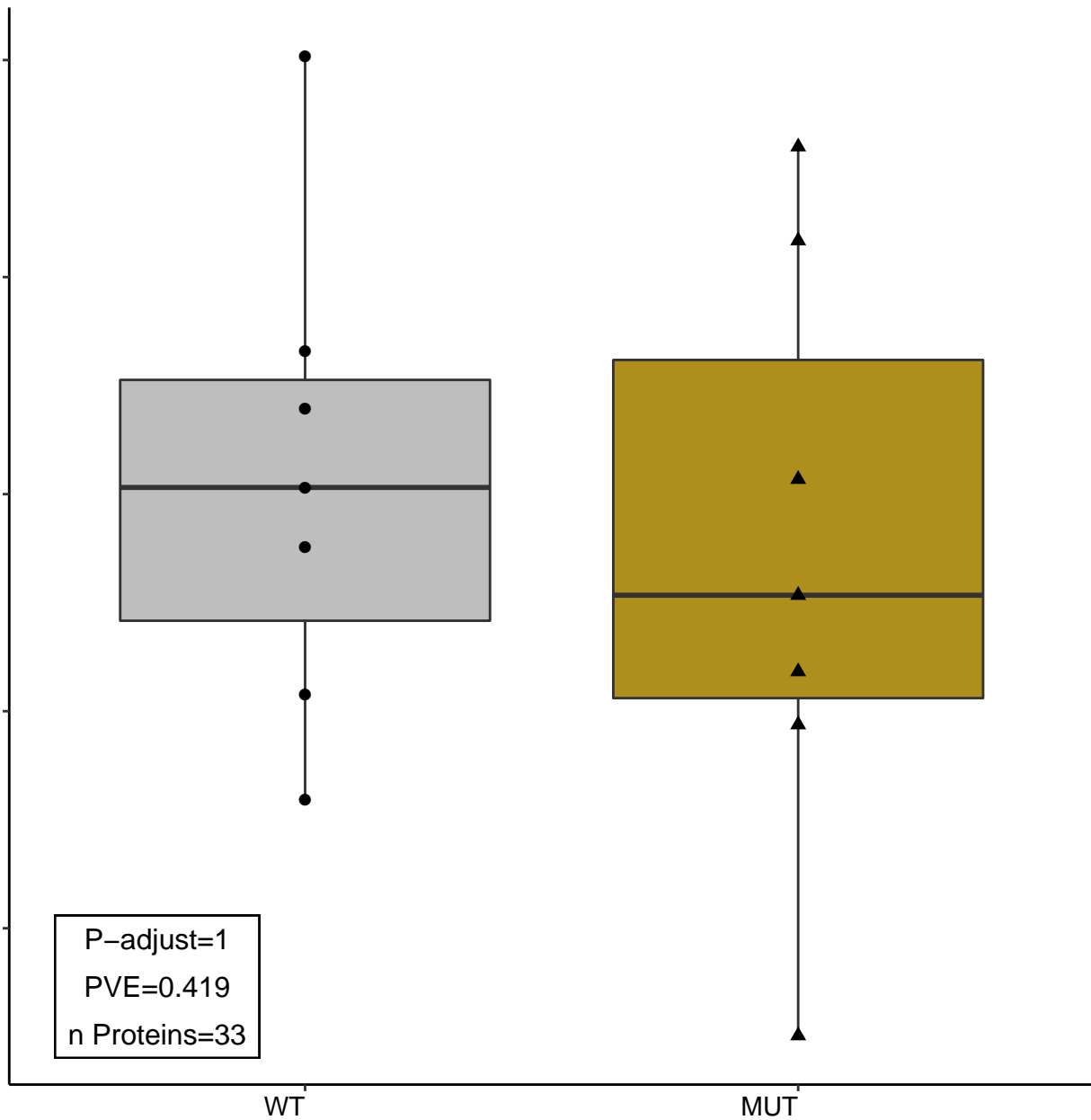
13.31
13.30
13.29
13.28
13.27

P-adjust=1
PVE=0.419
n Proteins=33

WT

Genotype

MUT



M55

Log2(Adjusted Module Intensity)

13.78

13.77

13.76

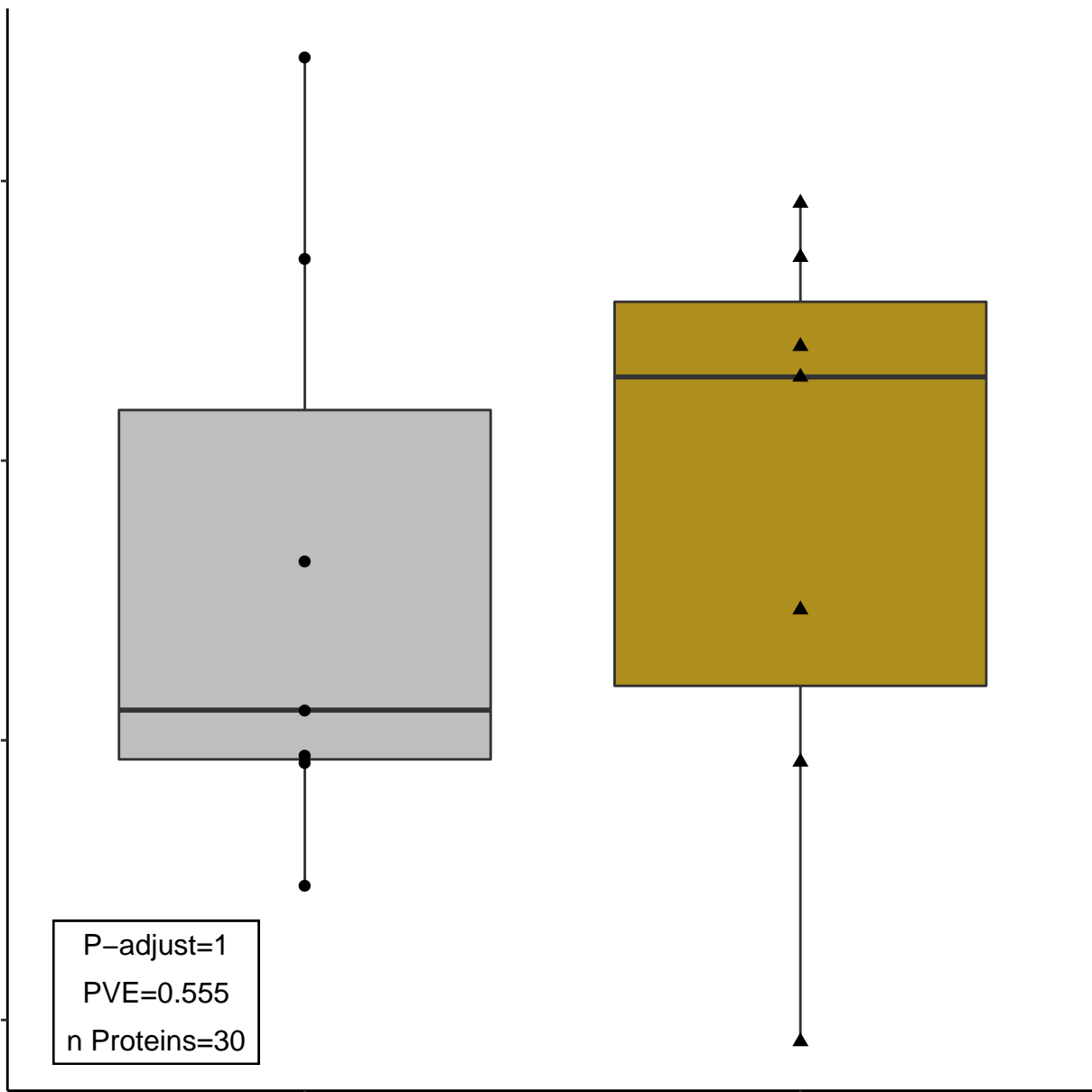
13.75

P-adjust=1
PVE=0.555
n Proteins=30

WT

Genotype

MUT



M56

Log2(Adjusted Module Intensity)

12.970

12.965

12.960

WT

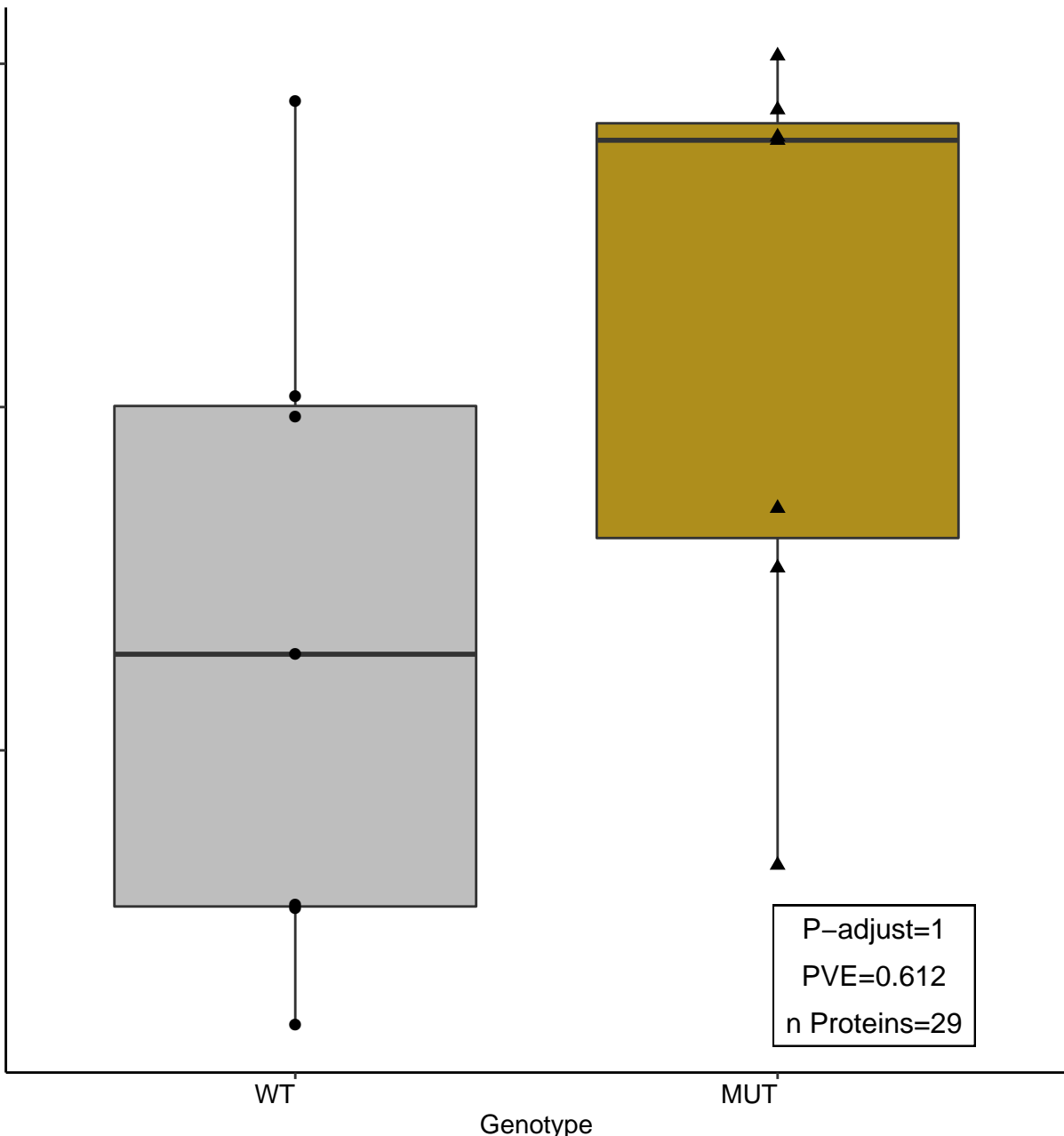
Genotype

MUT

P-adjust=1

PVE=0.612

n Proteins=29



M57

Log2(Adjusted Module Intensity)

14.47

14.46

14.45

14.44

14.43

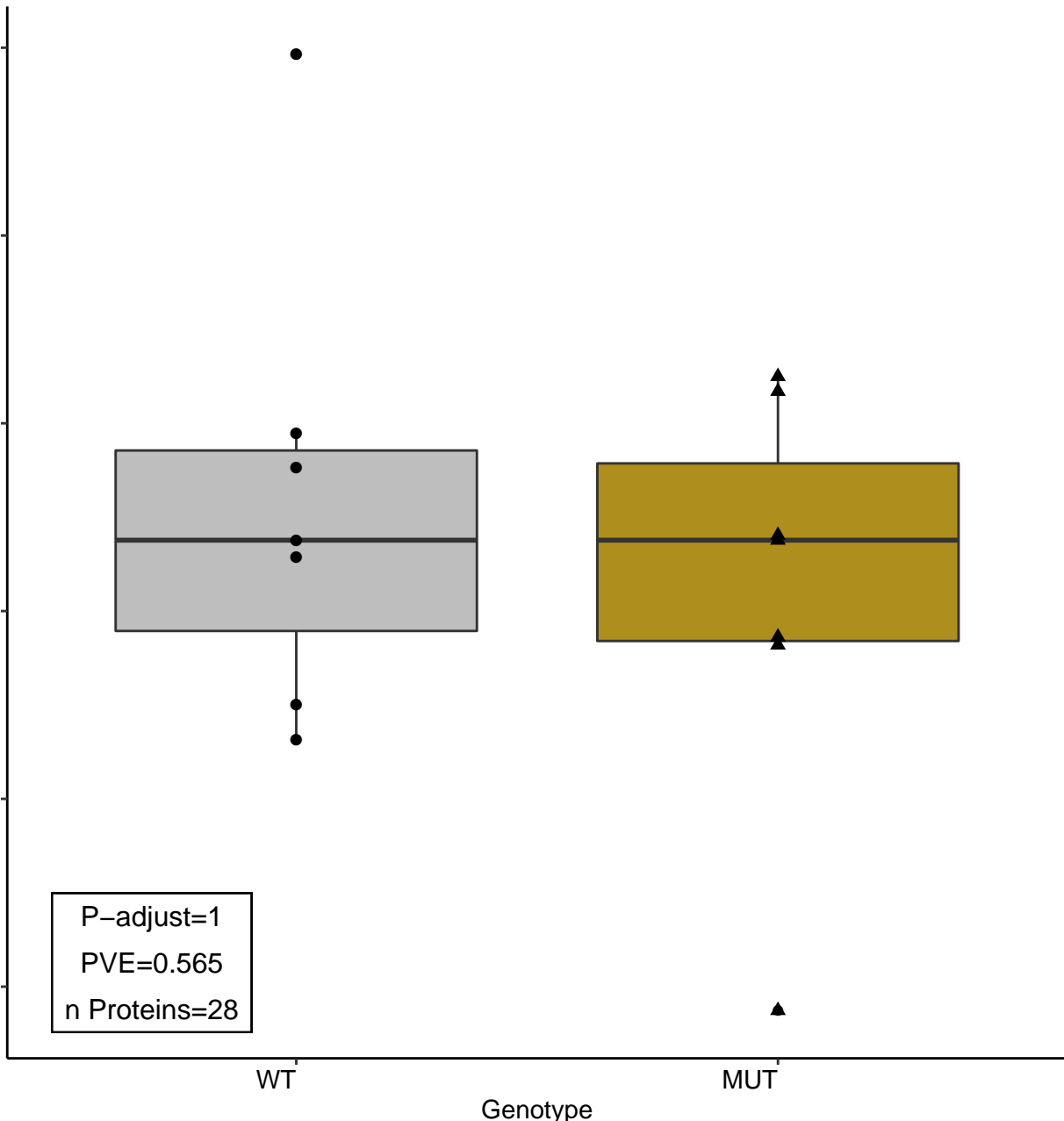
14.42

WT

MUT

Genotype

P-adjust=1
PVE=0.565
n Proteins=28



M58

Log2(Adjusted Module Intensity)

12.98

12.97

12.96

12.95

WT

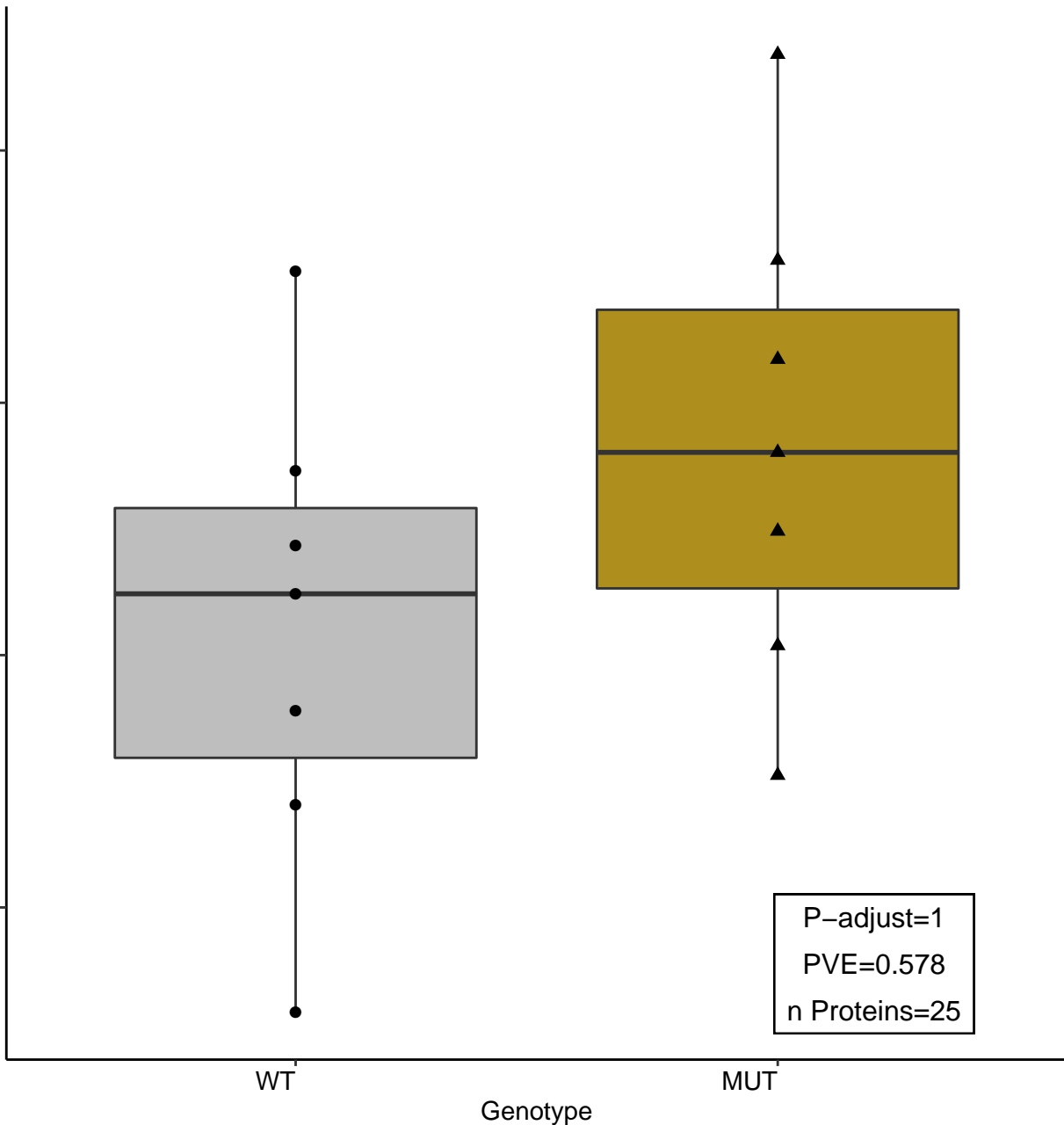
Genotype

MUT

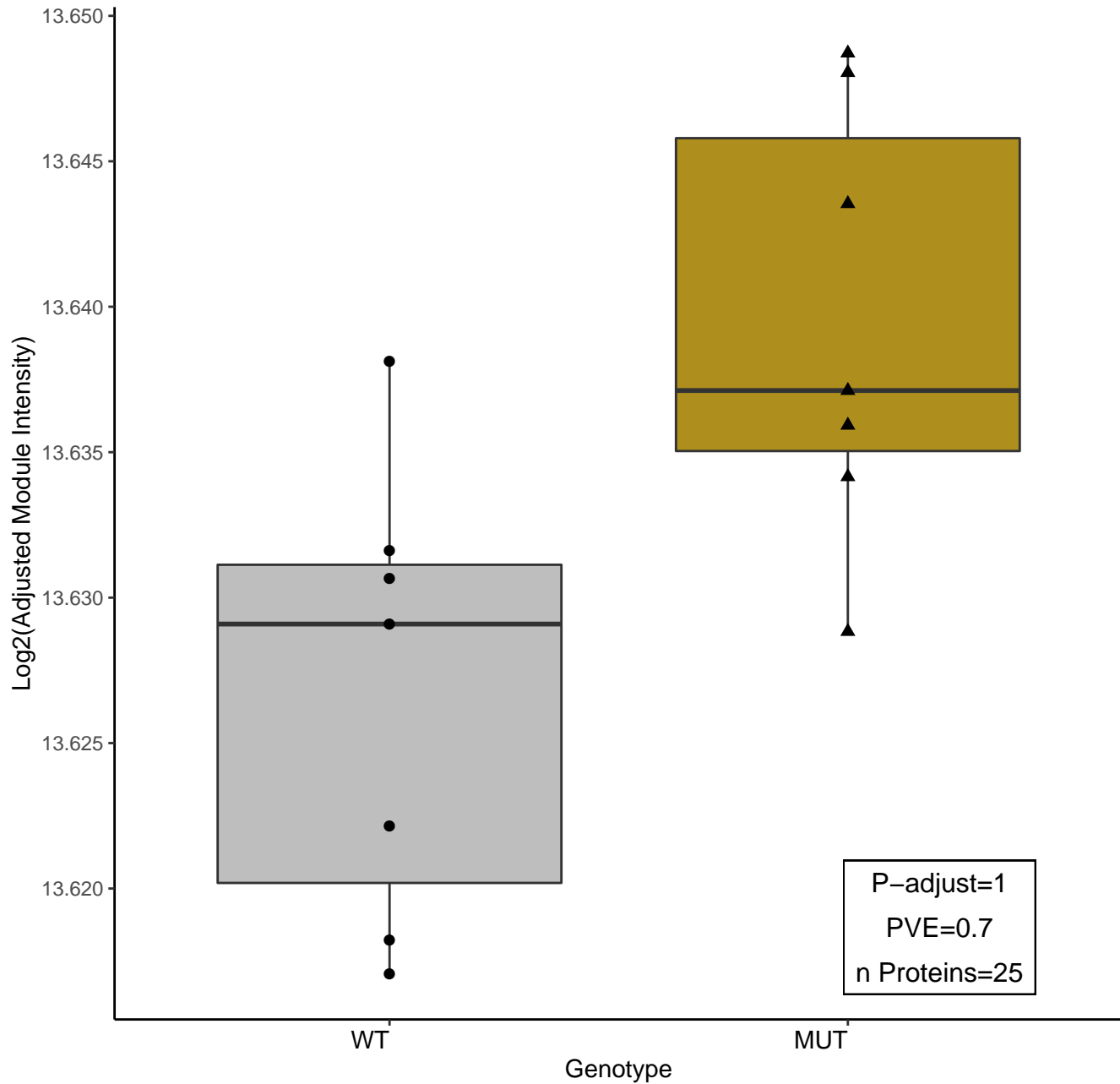
P-adjust=1

PVE=0.578

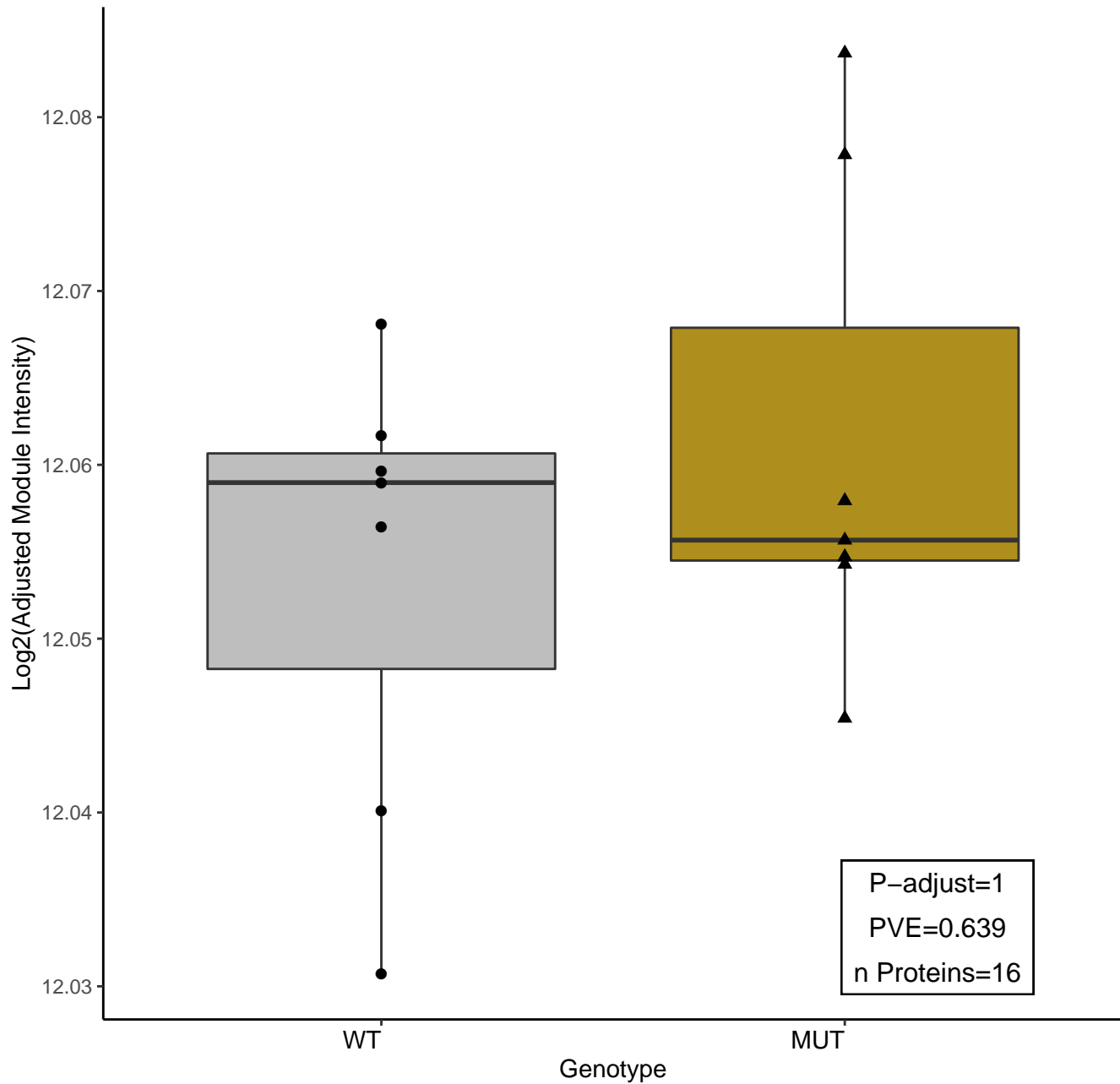
n Proteins=25



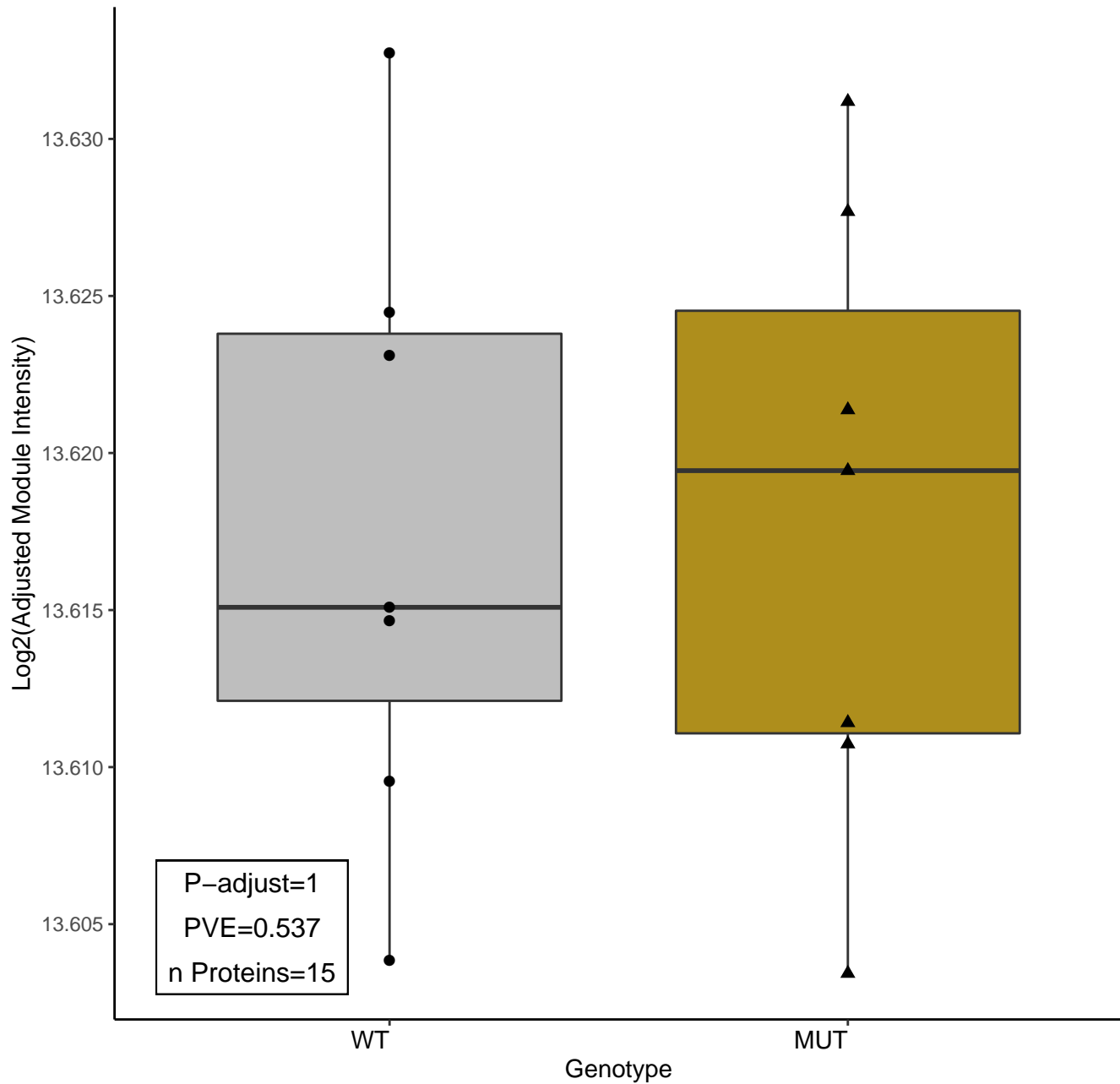
M59



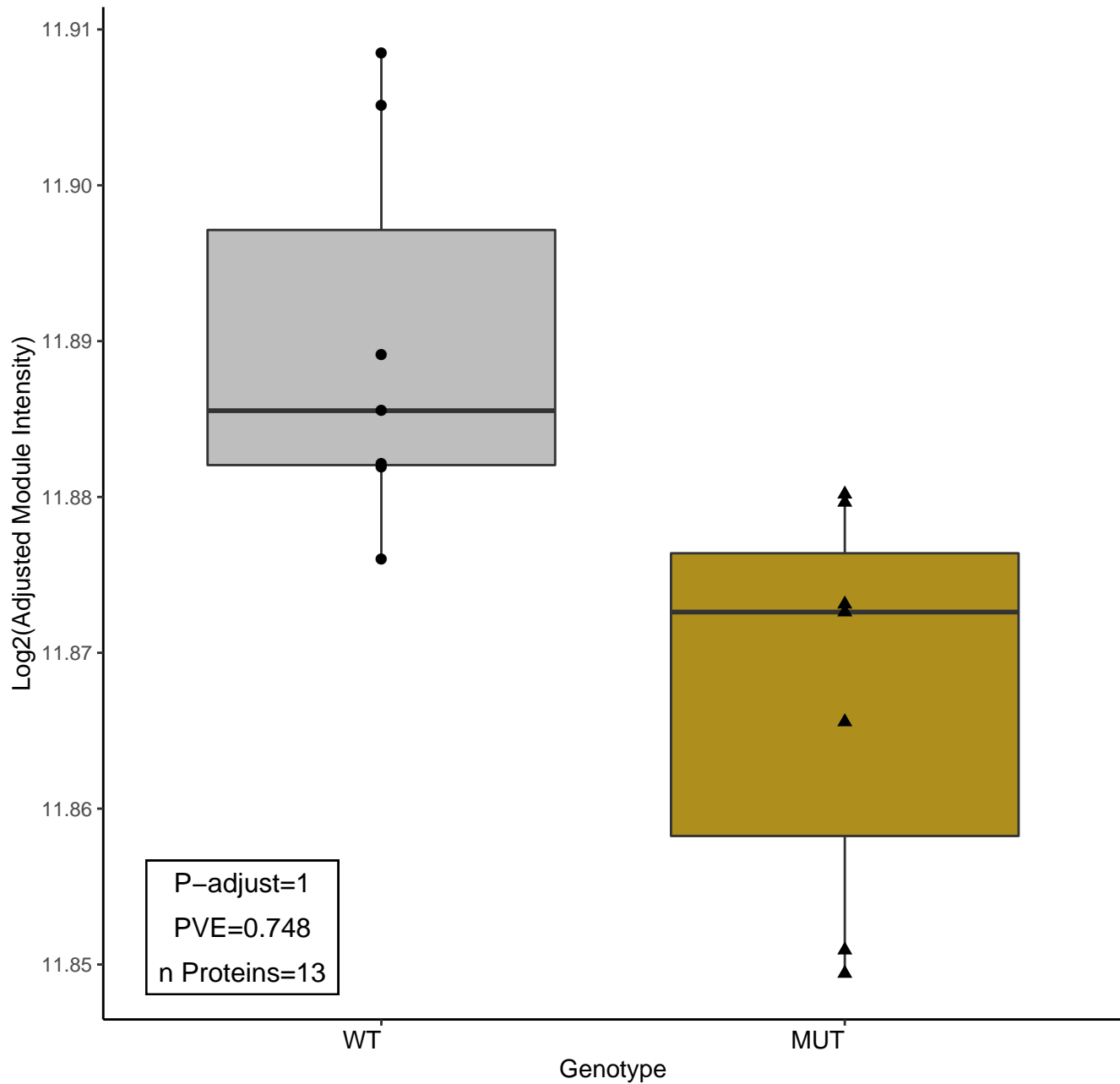
M60



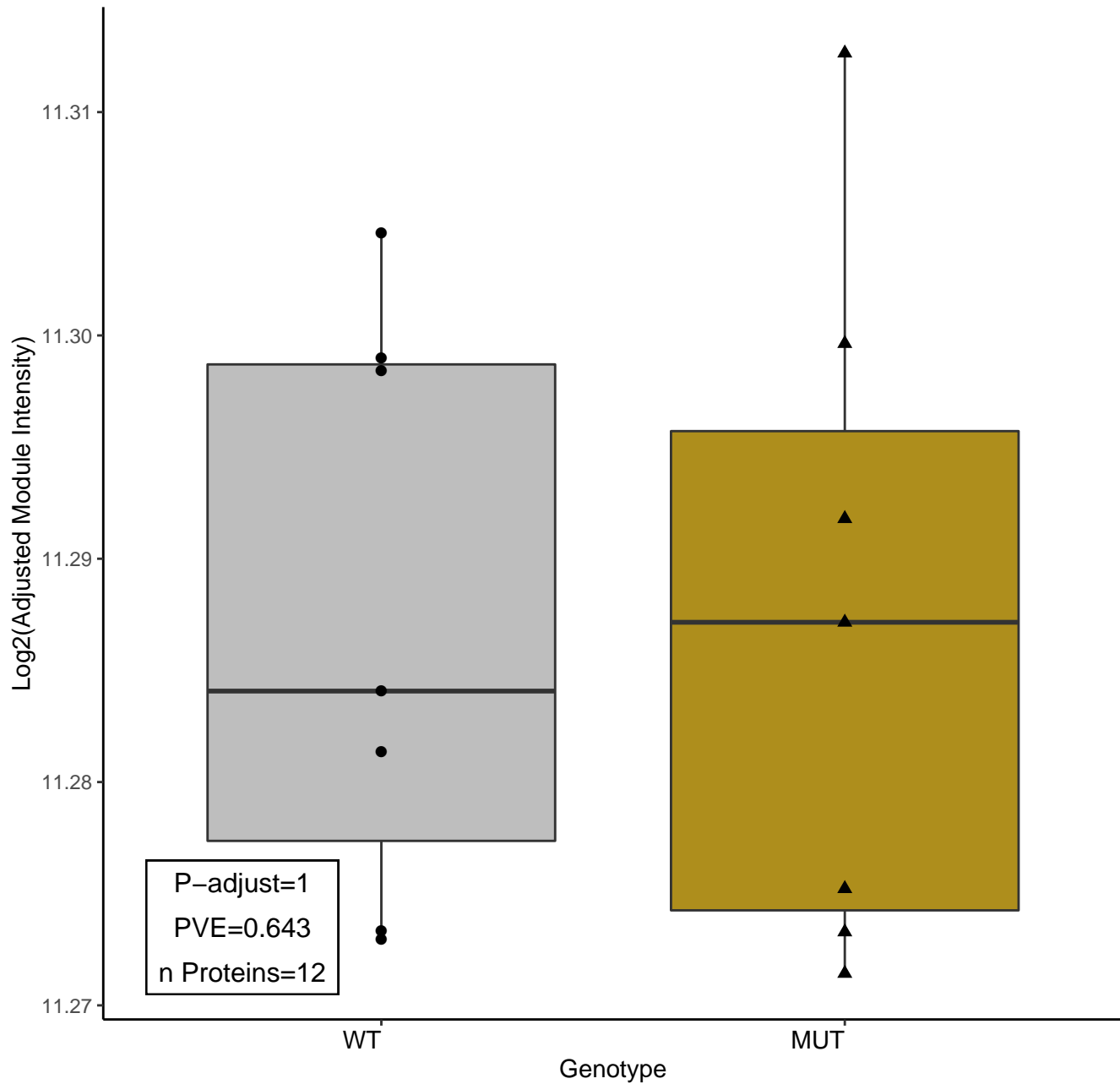
M61



M62



M63



M64

Log2(Adjusted Module Intensity)

13.17

13.16

13.15

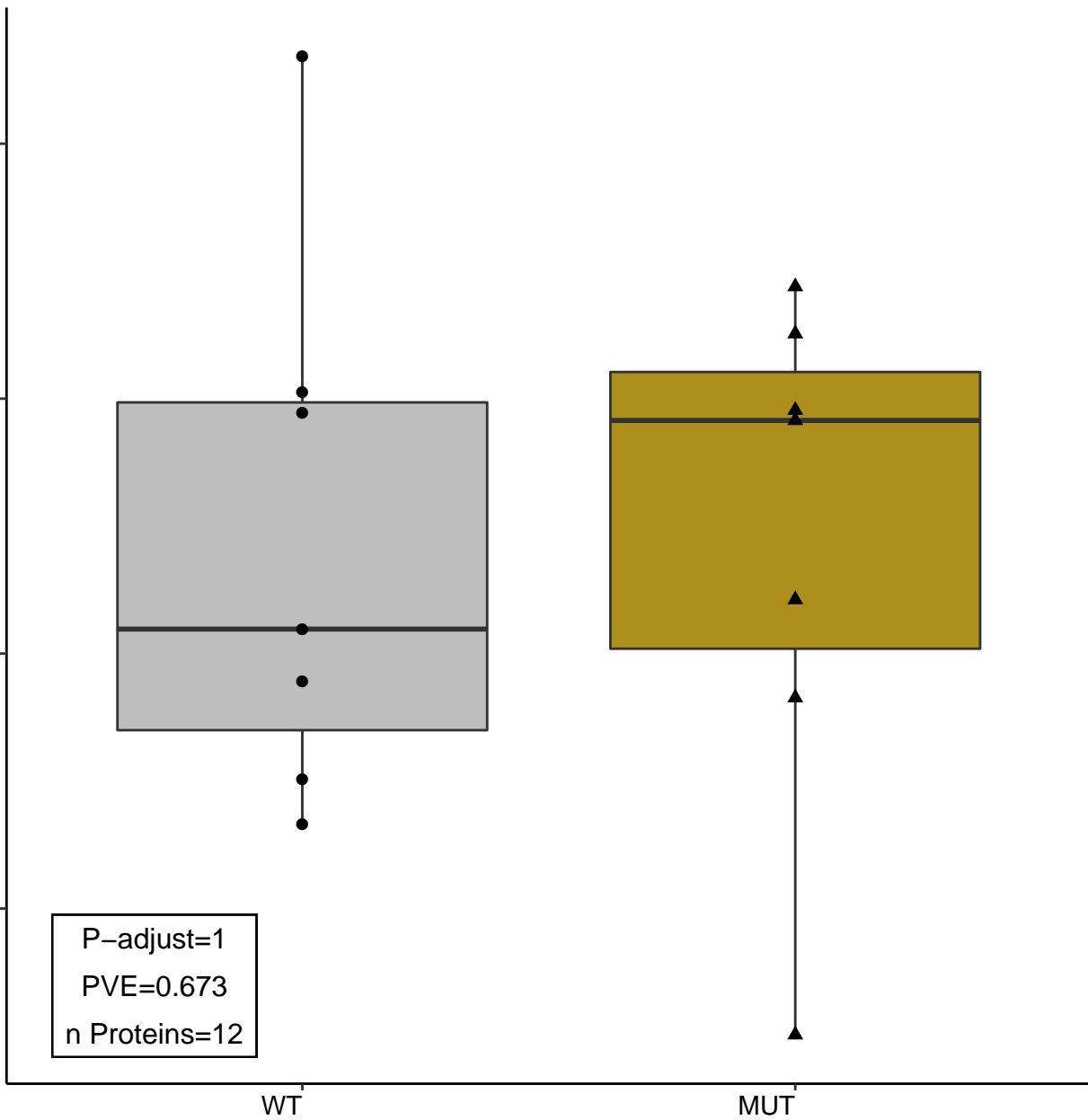
13.14

P-adjust=1
PVE=0.673
n Proteins=12

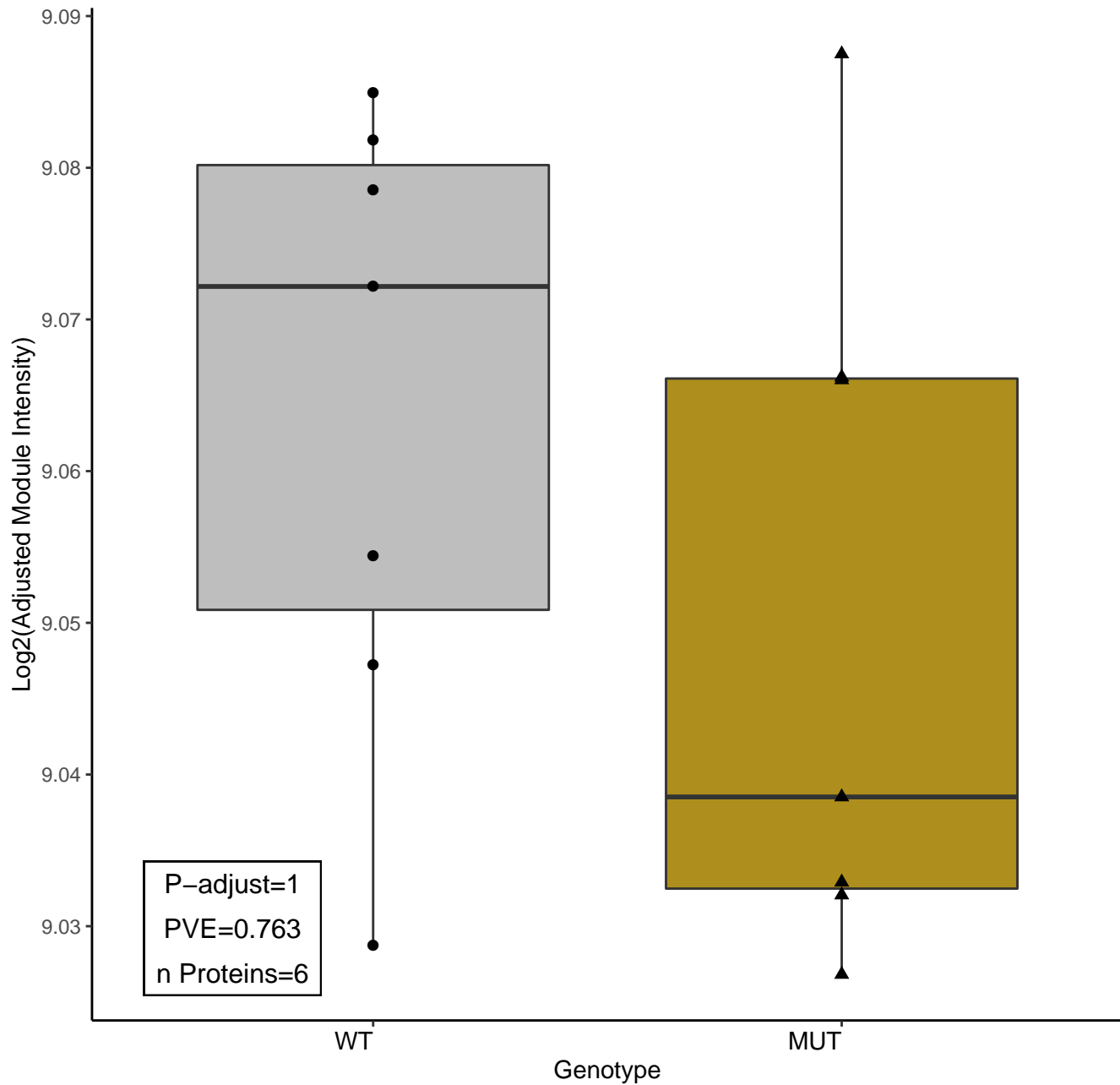
WT

Genotype

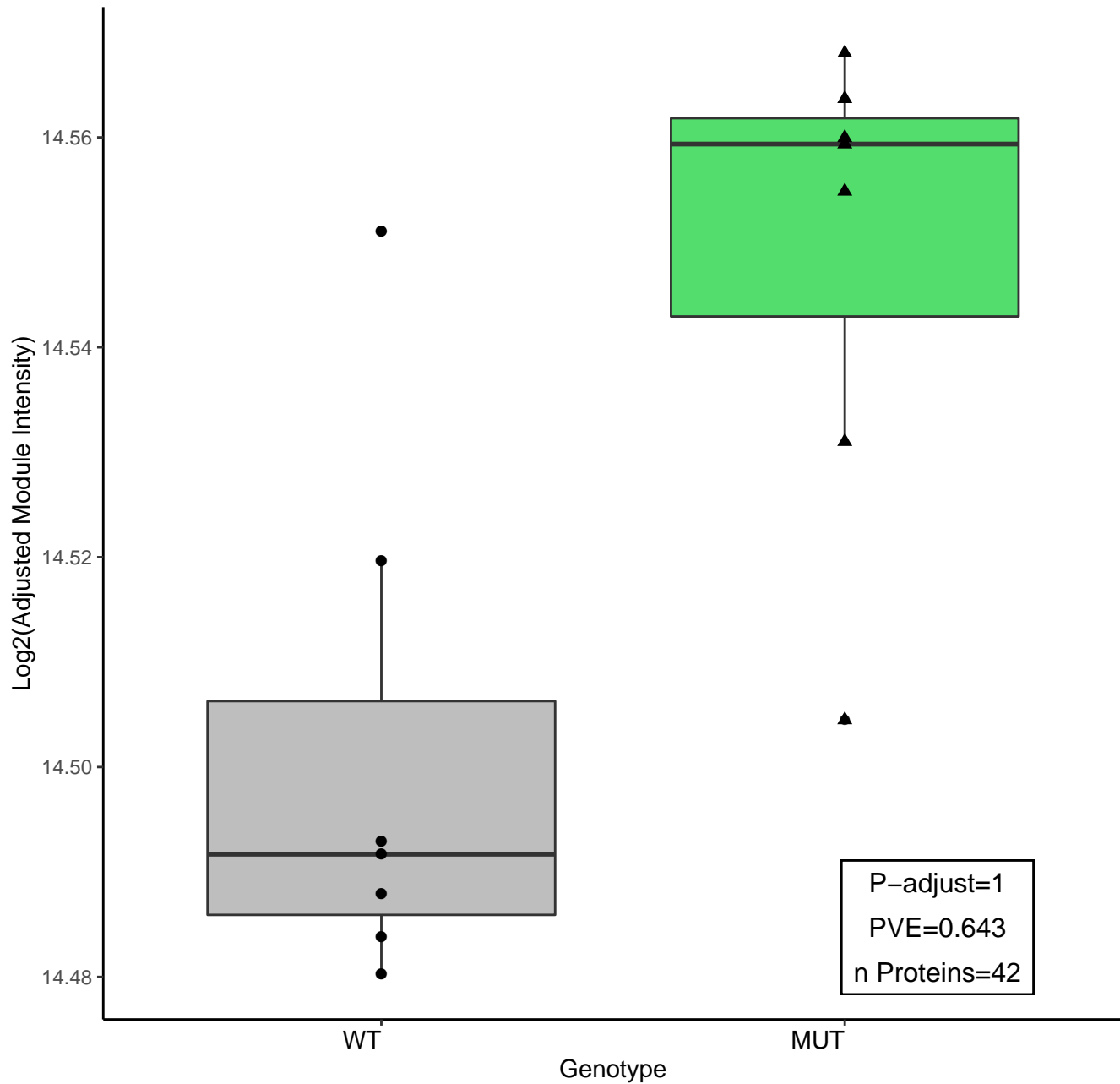
MUT



M65



M66



M67

Log2(Adjusted Module Intensity)

13.76

13.74

13.72

13.70

WT

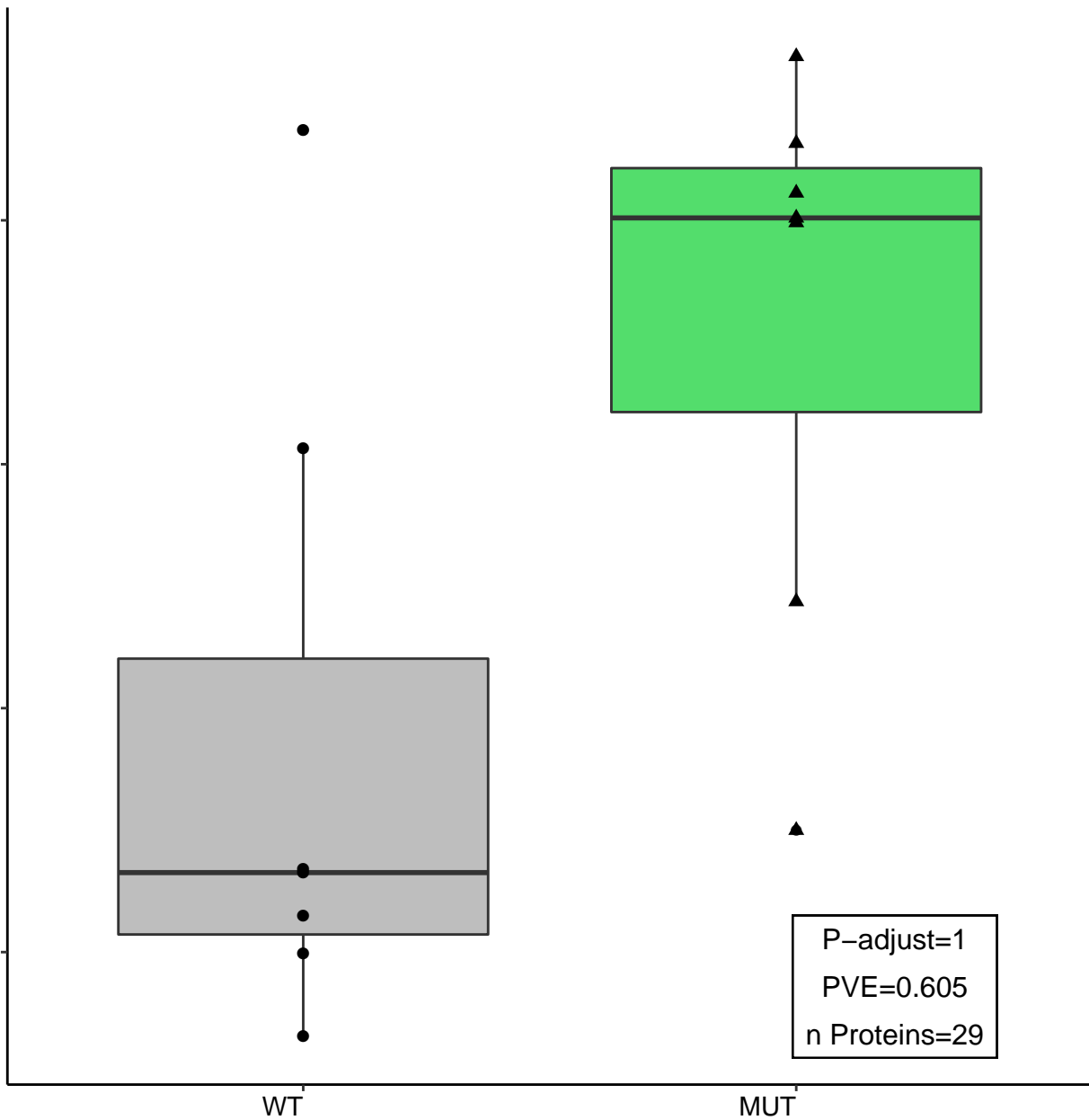
Genotype

MUT

P-adjust=1

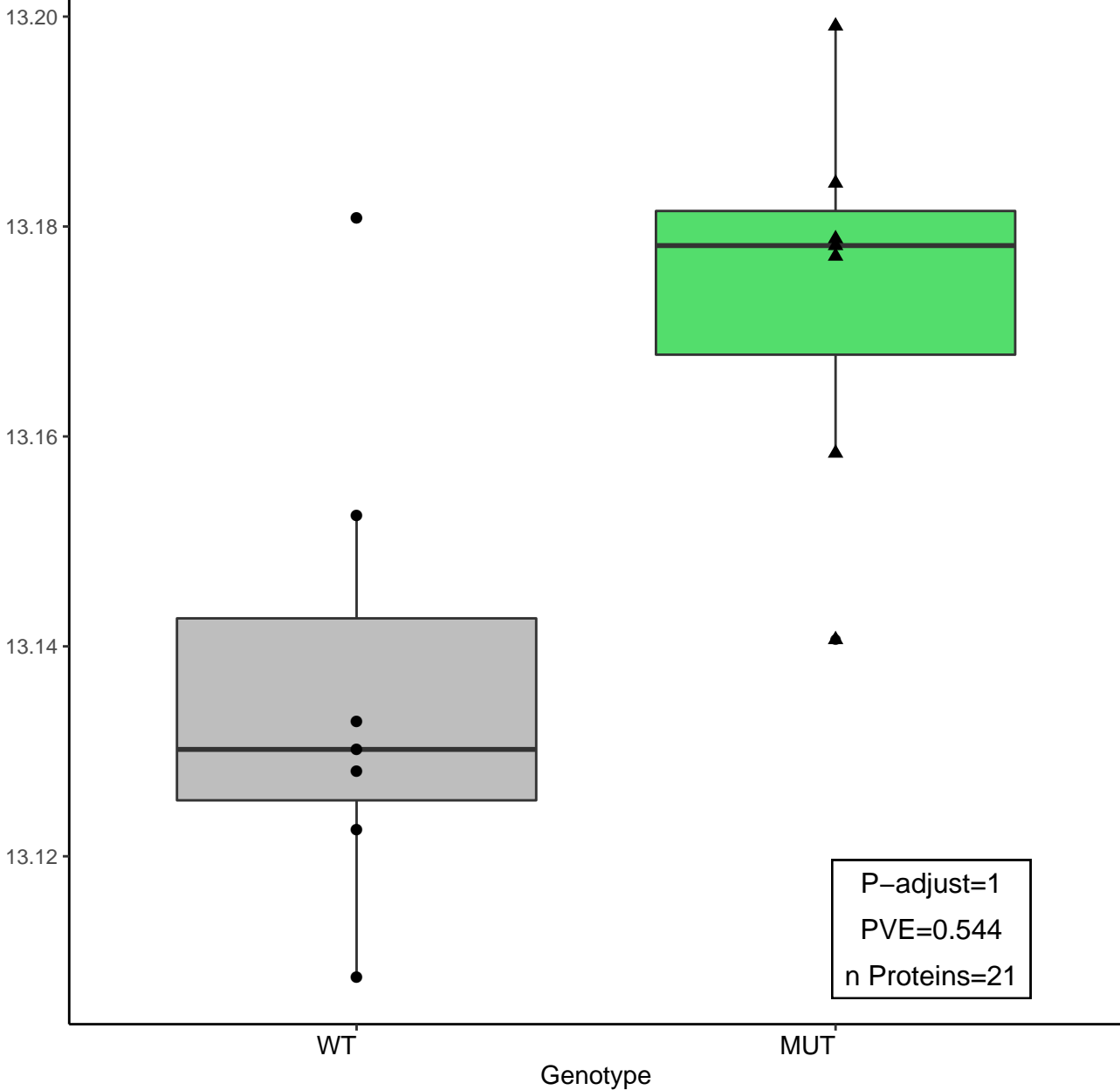
PVE=0.605

n Proteins=29



M68

Log2(Adjusted Module Intensity)



M69

Log2(Adjusted Module Intensity)

13.10

13.05

13.00

WT

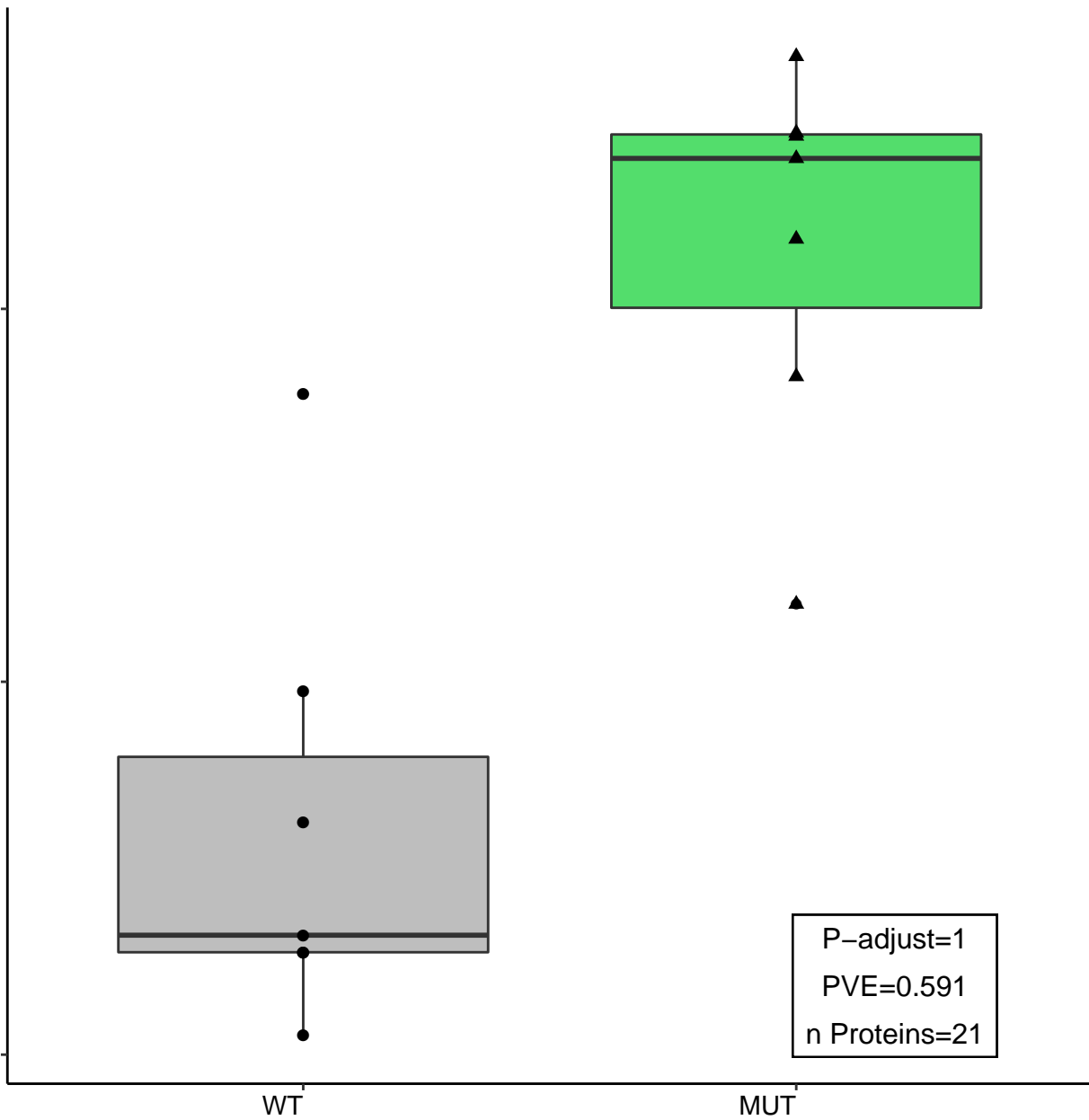
MUT

Genotype

P-adjust=1

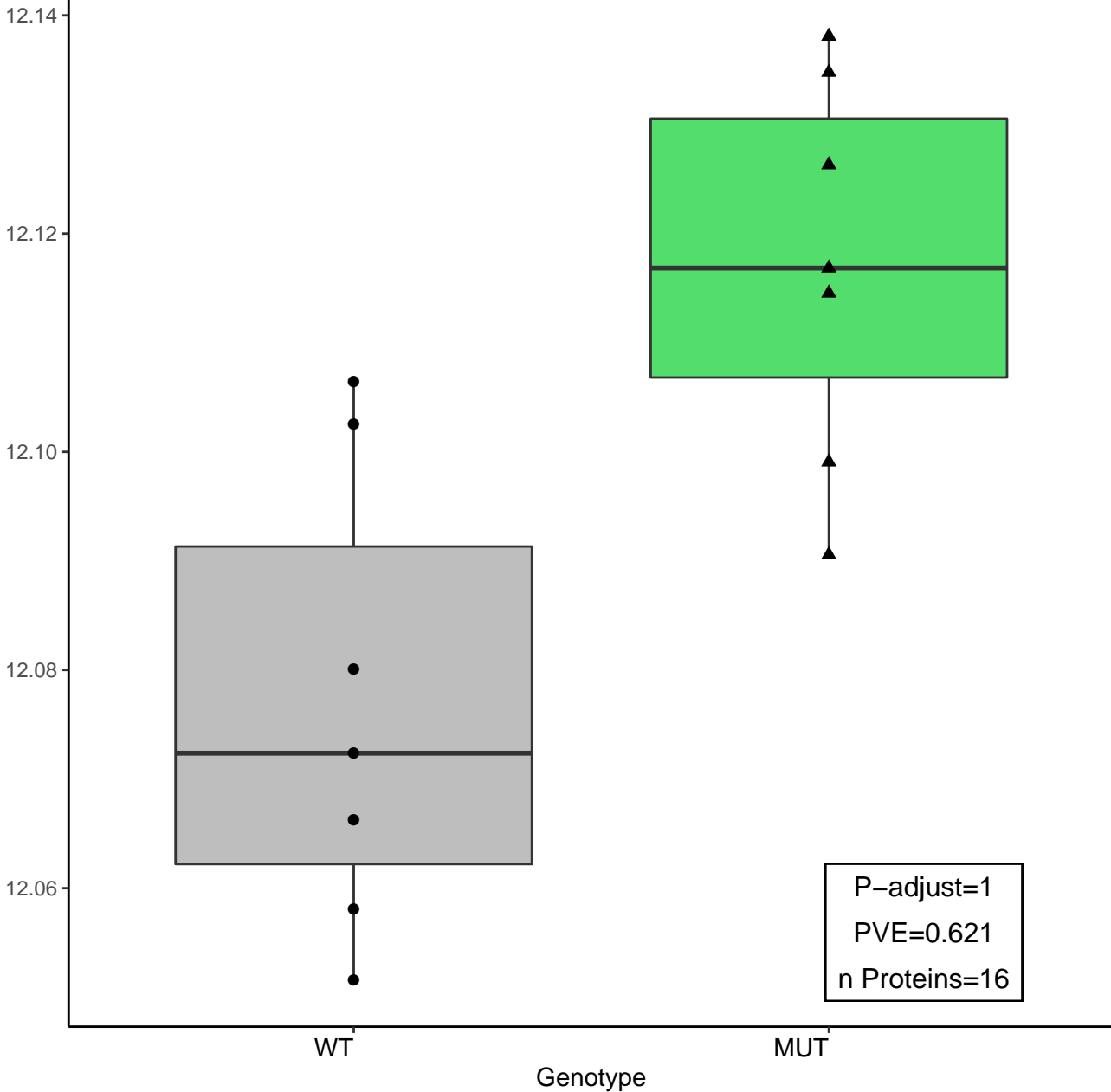
PVE=0.591

n Proteins=21

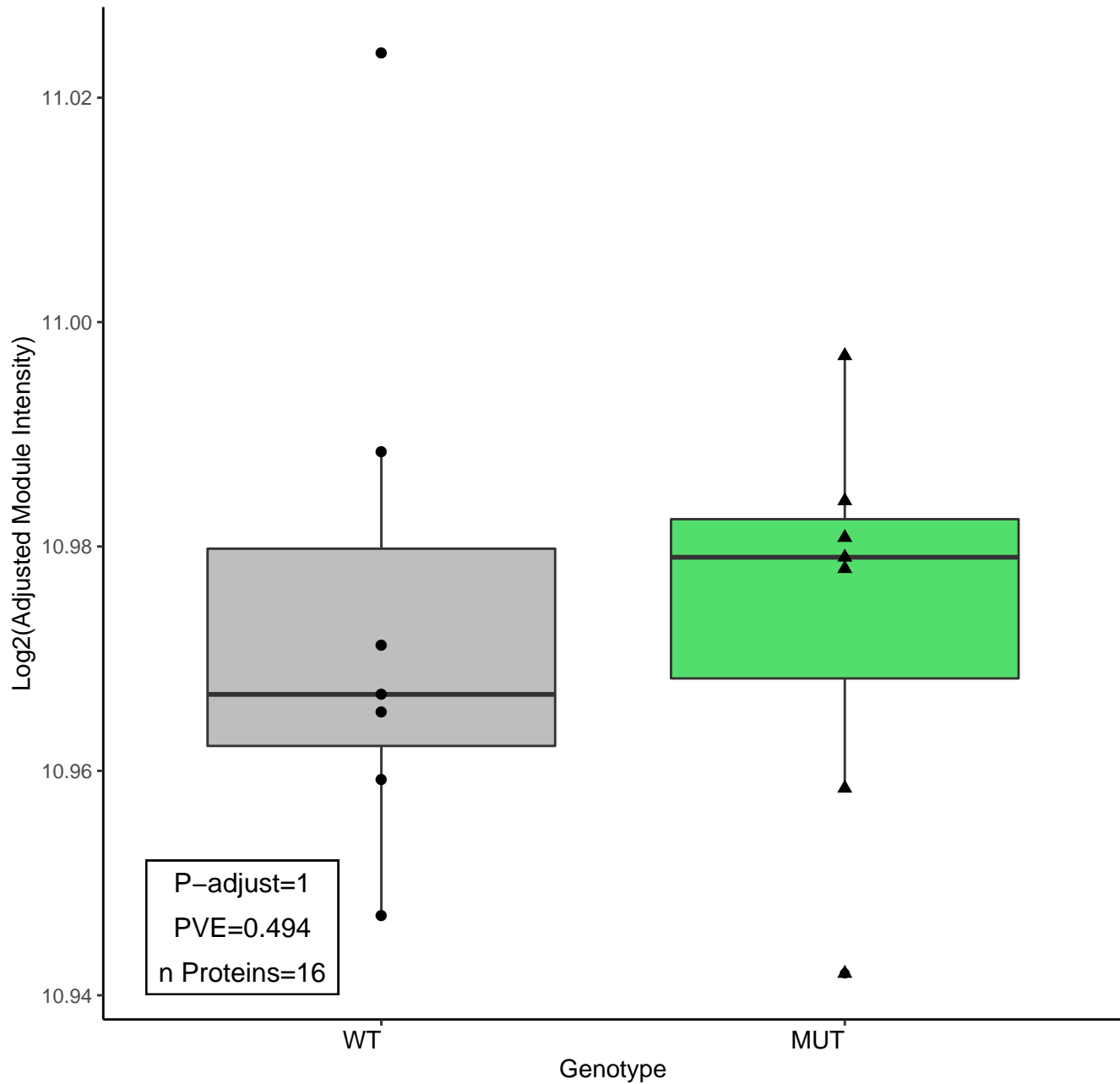


M70

Log2(Adjusted Module Intensity)



M71



M72

Log2(Adjusted Module Intensity)

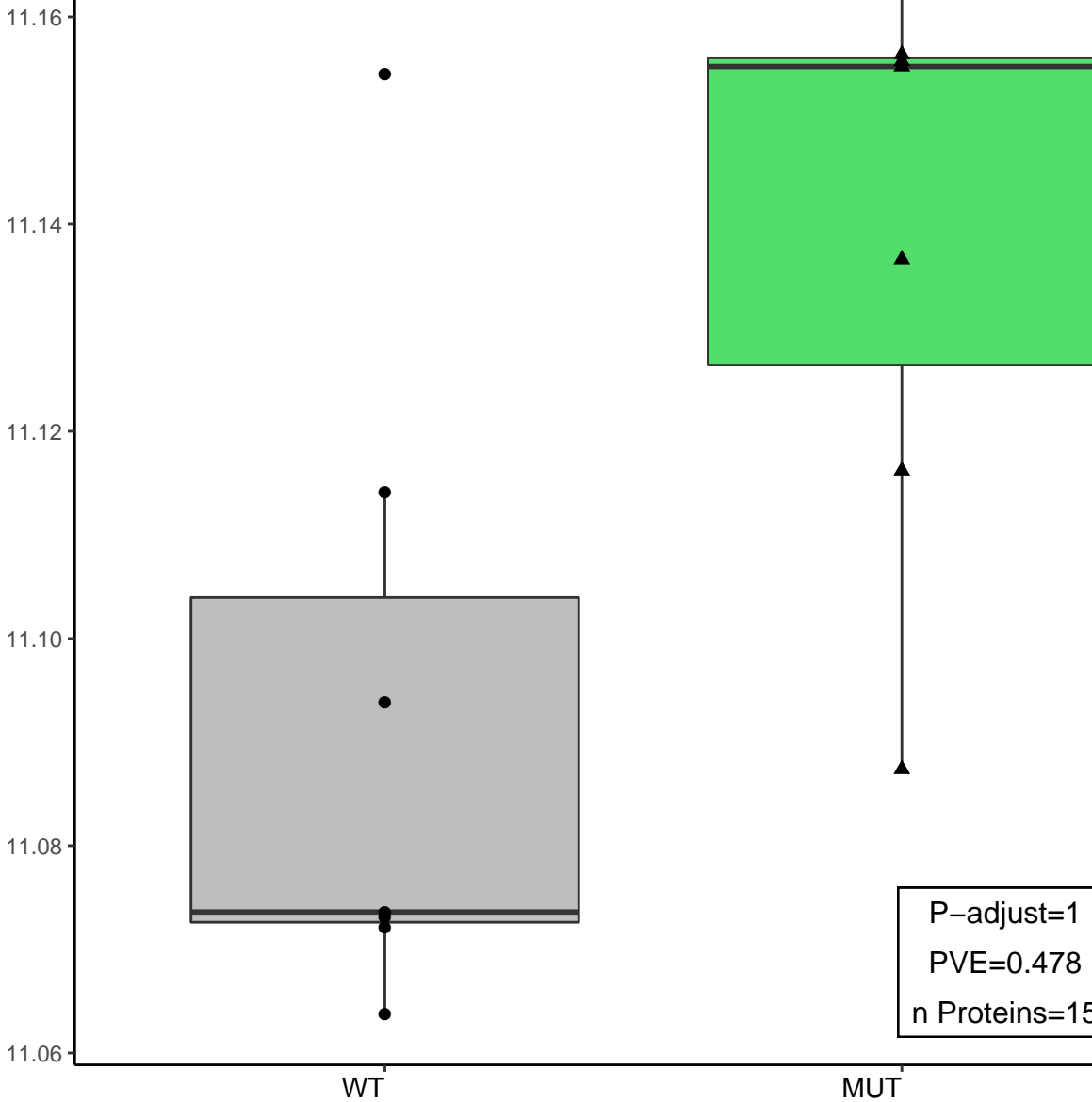
11.16
11.14
11.12
11.10
11.08
11.06

WT

MUT

Genotype

P-adjust=1
PVE=0.478
n Proteins=15



M73

Log2(Adjusted Module Intensity)

11.04

11.02

11.00

10.98

10.96

10.94

WT

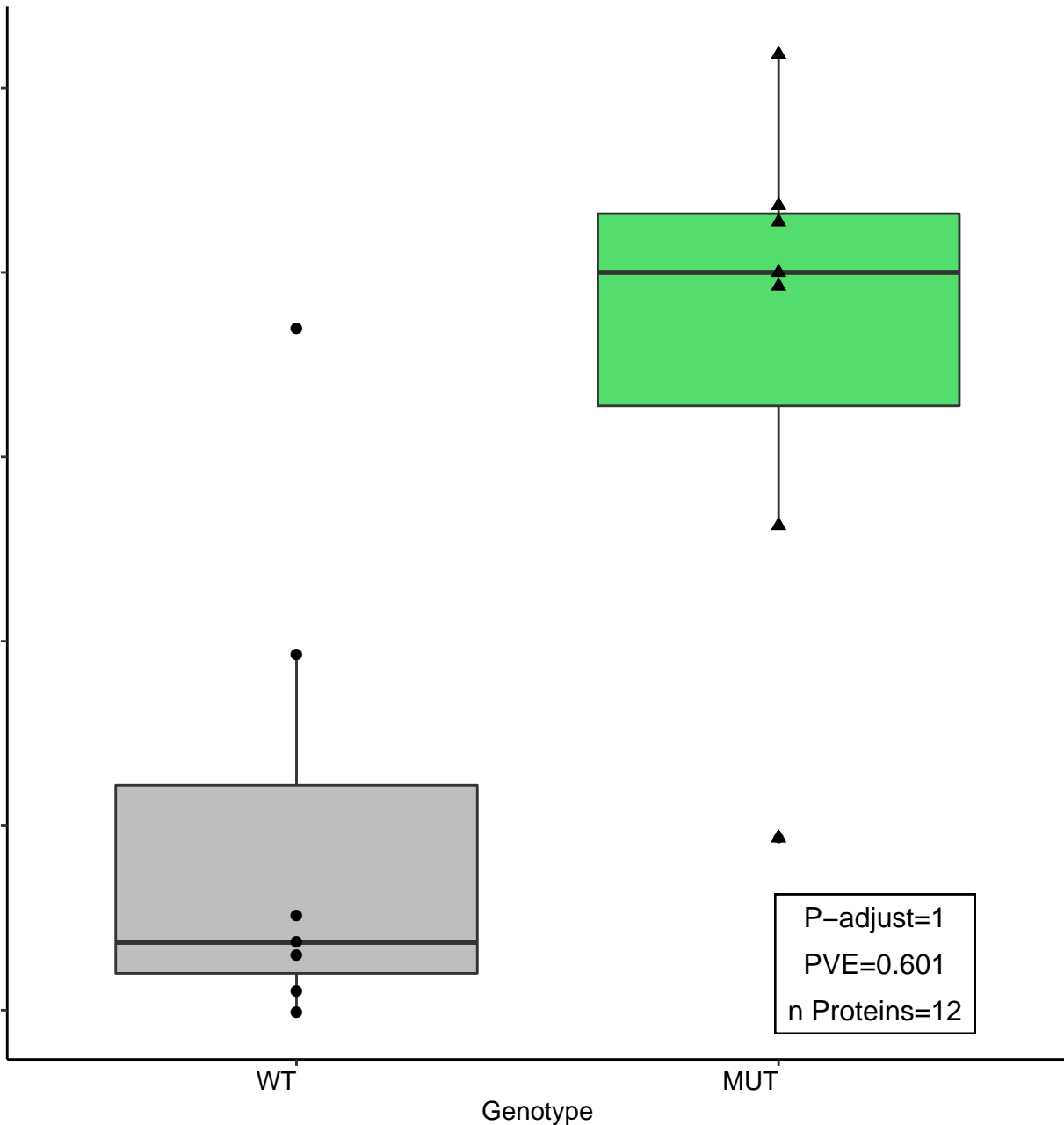
MUT

Genotype

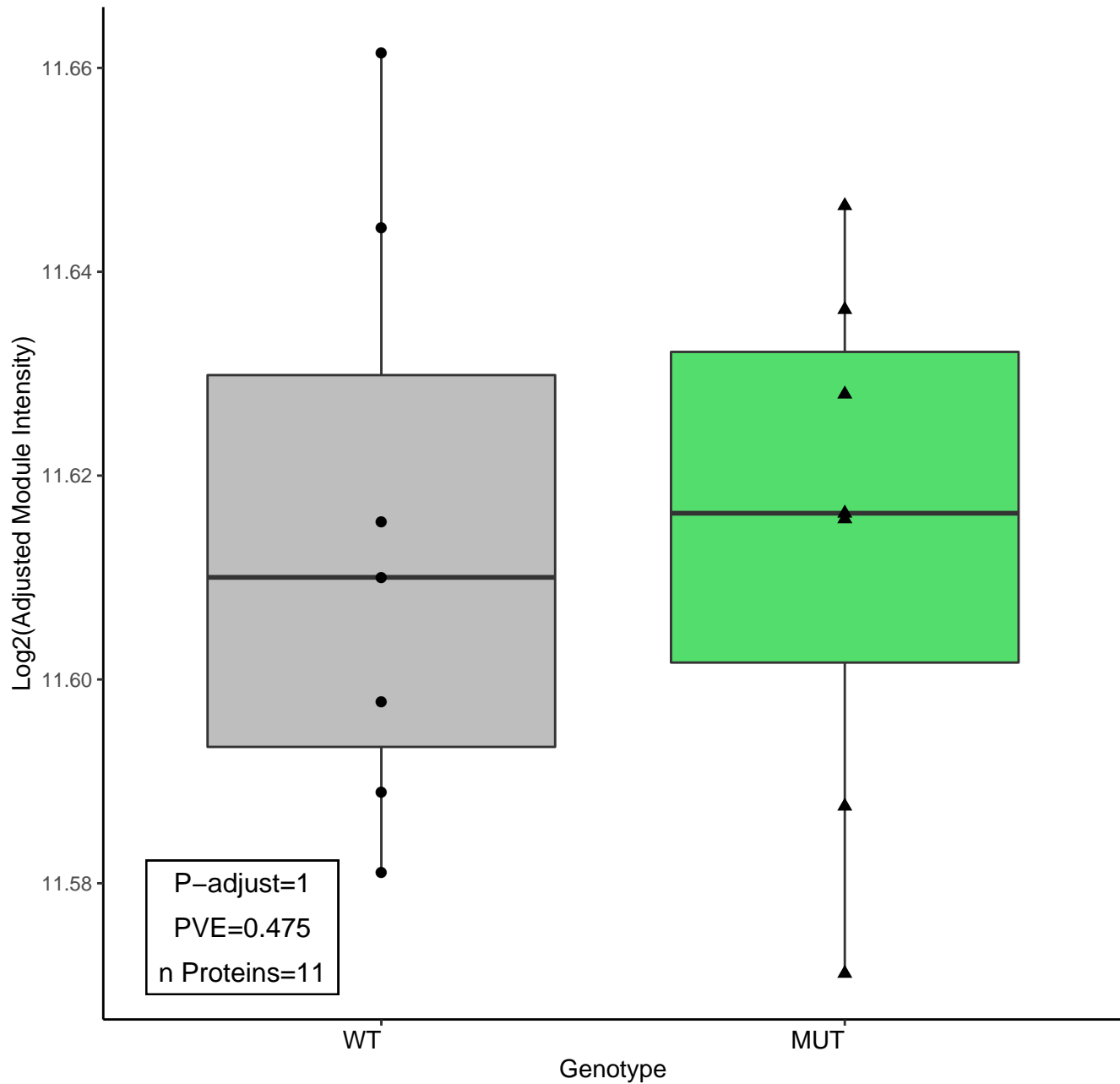
P-adjust=1

PVE=0.601

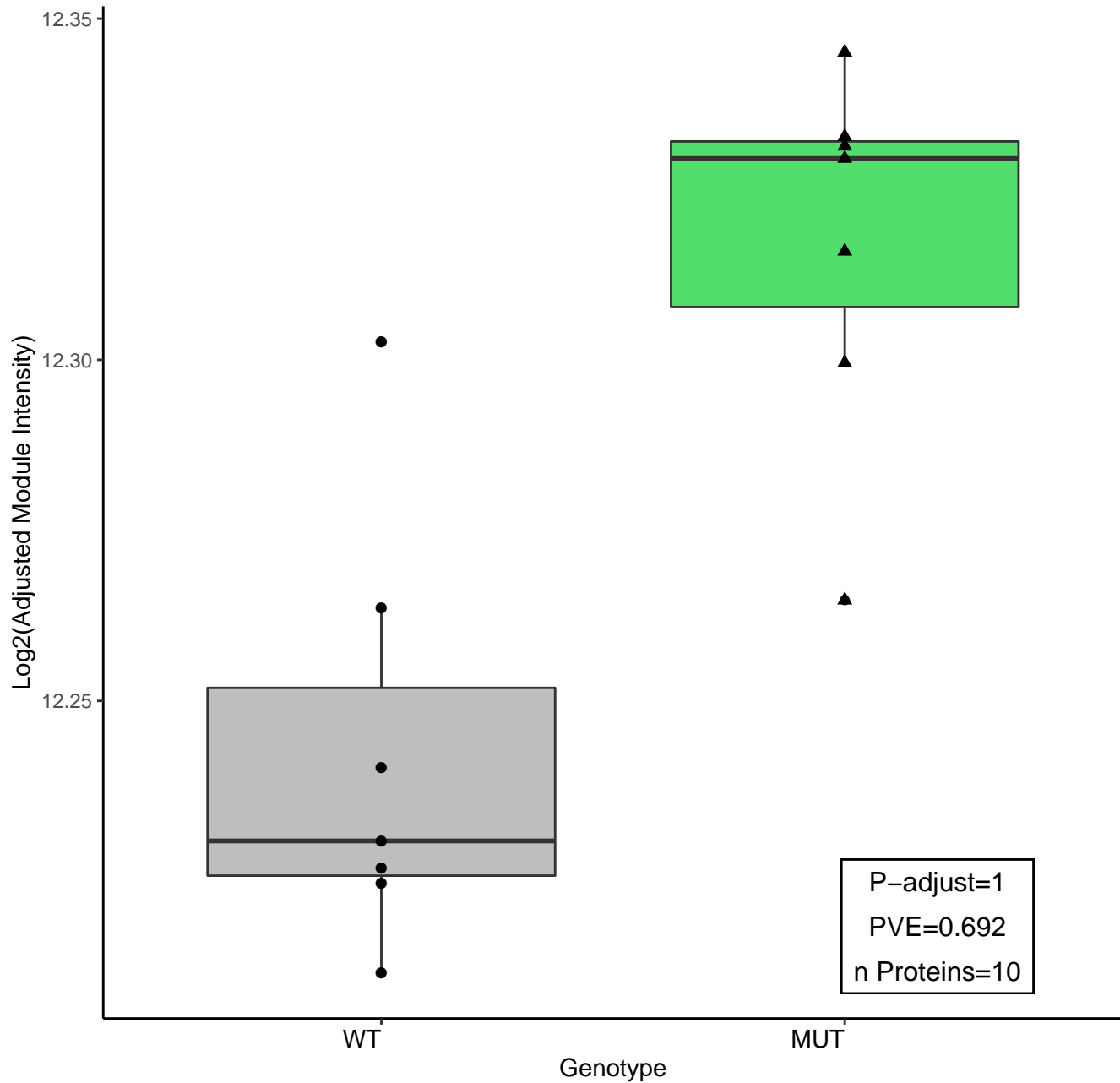
n Proteins=12

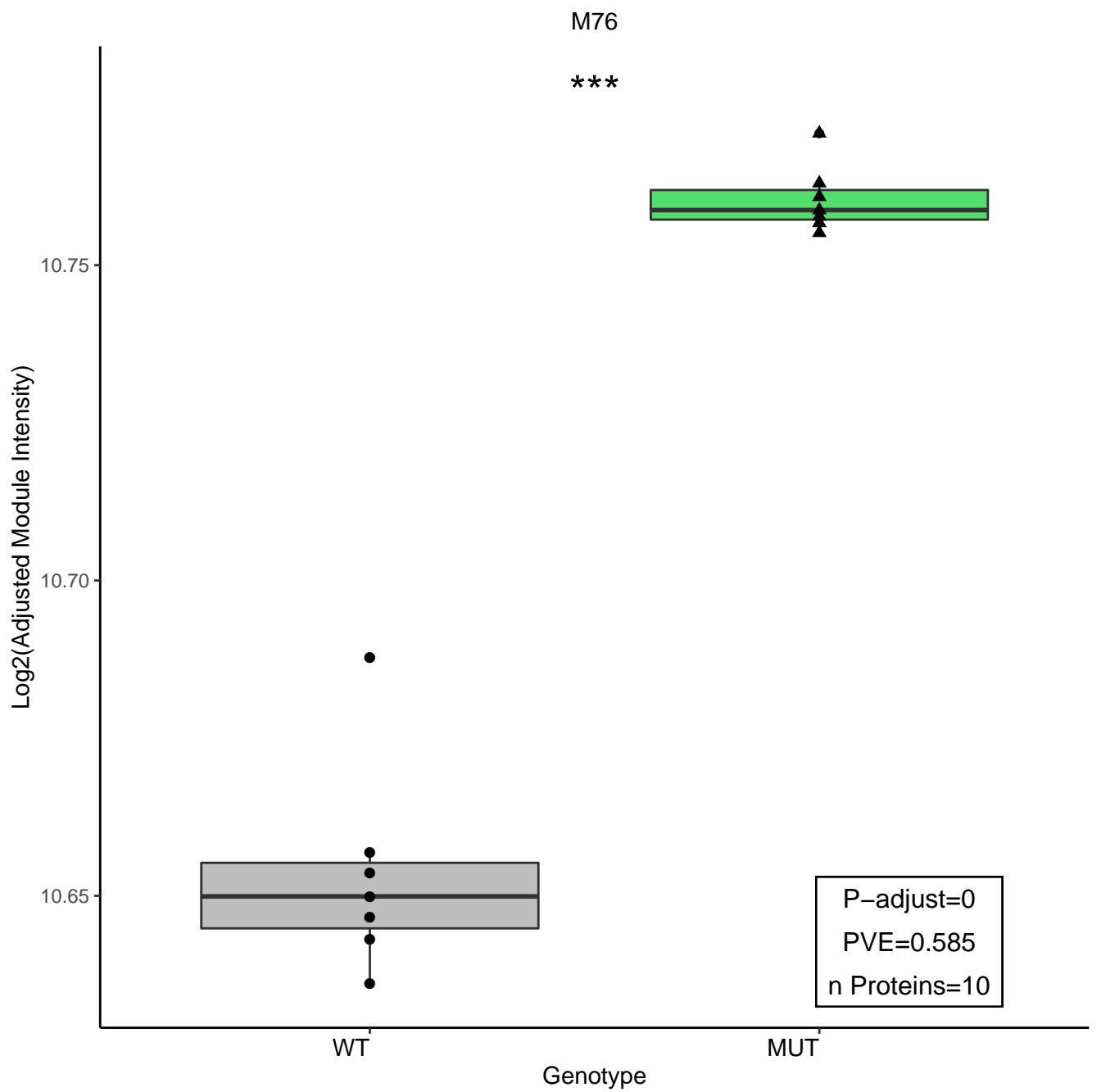


M74

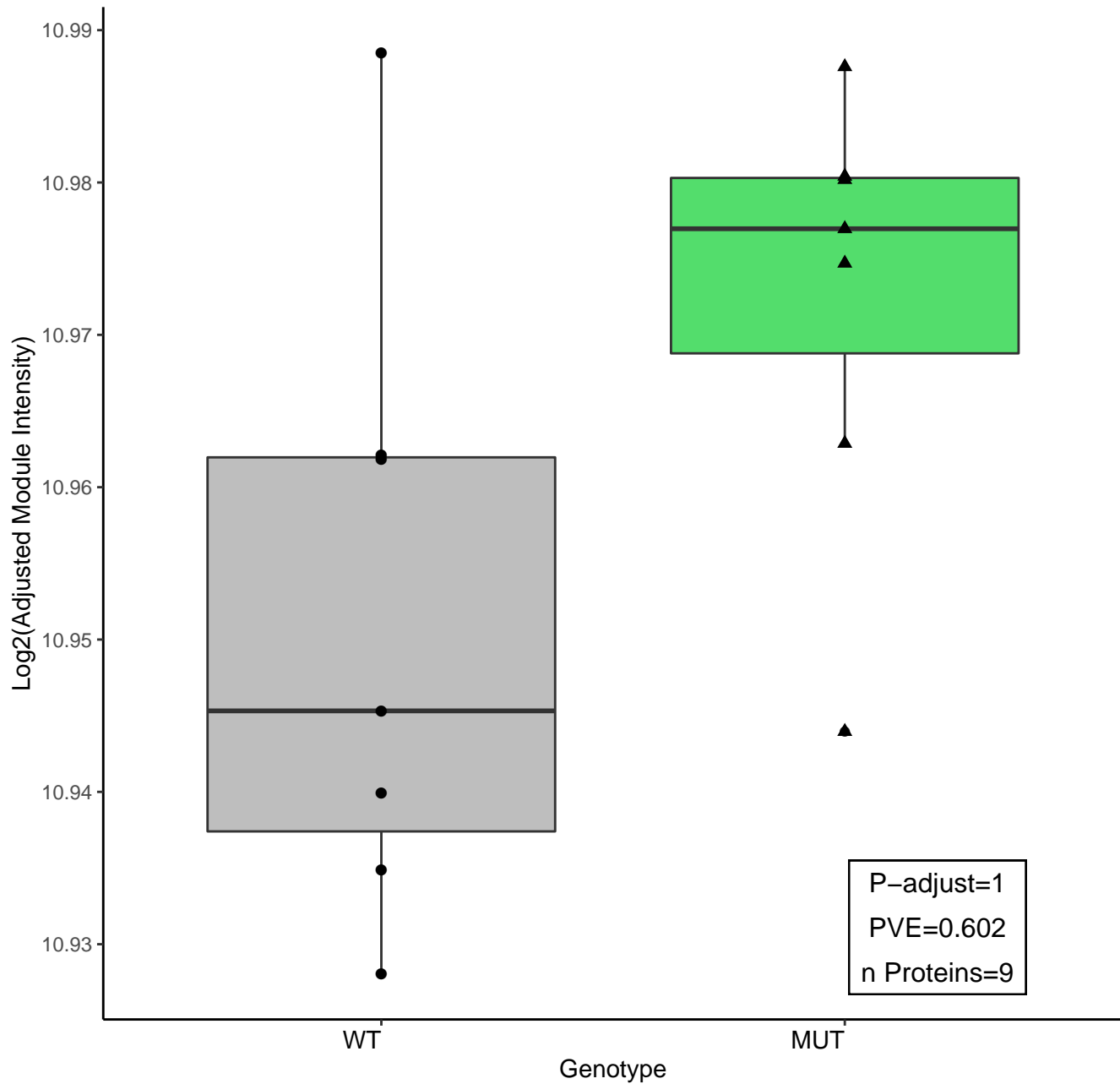


M75

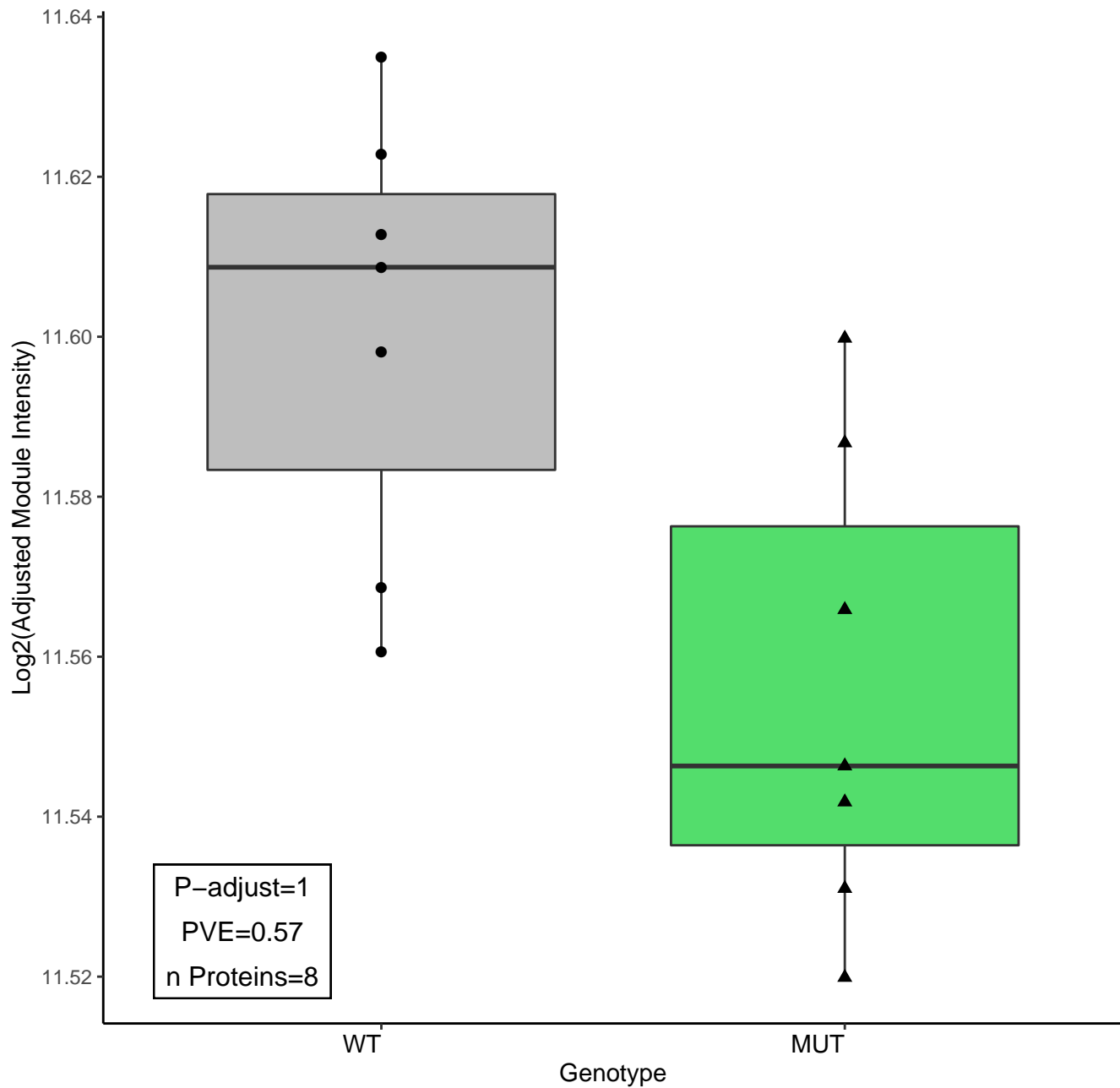




M77



M78



M79

Log2(Adjusted Module Intensity)

9.55

9.50

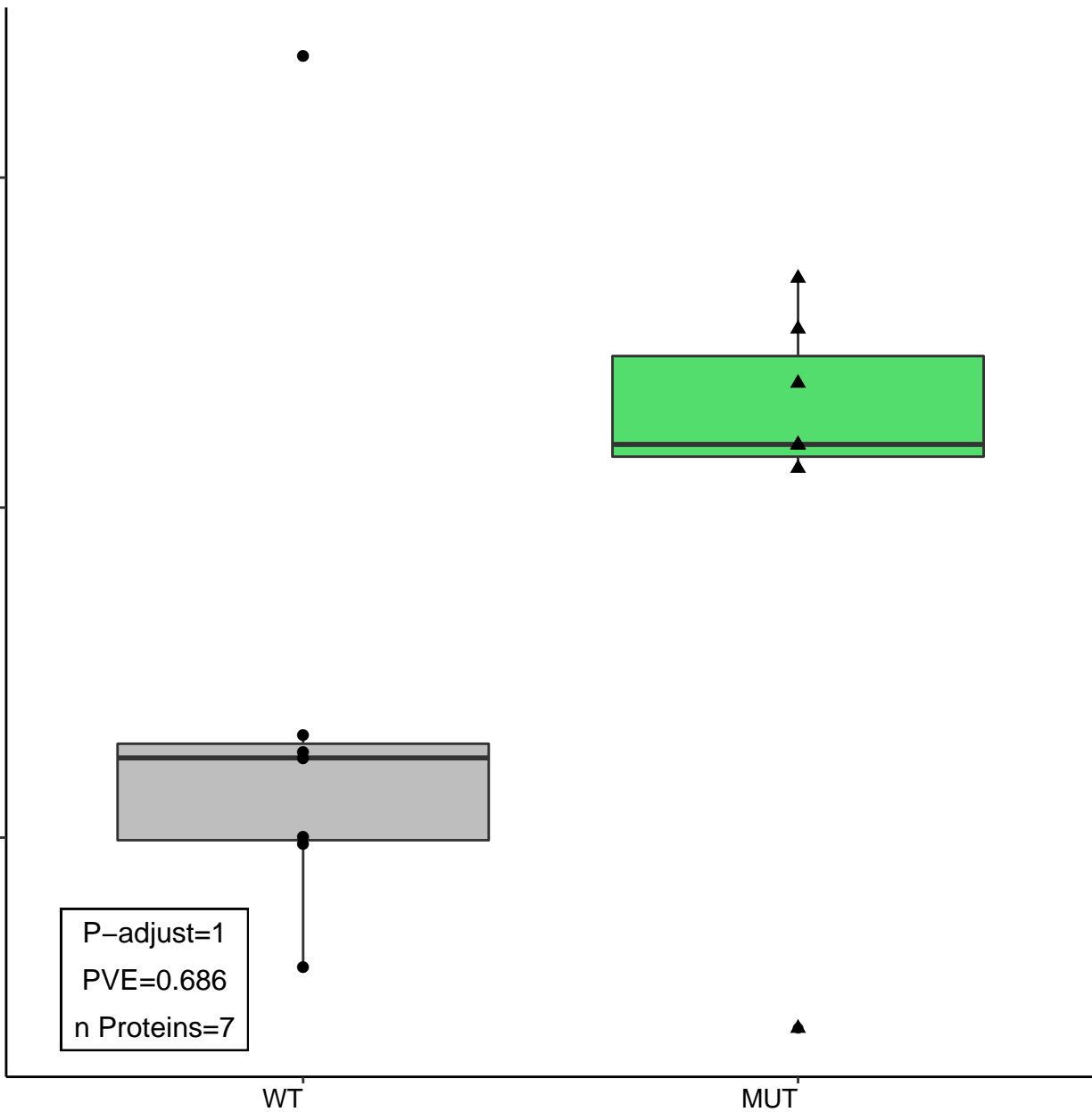
9.45

P-adjust=1
PVE=0.686
n Proteins=7

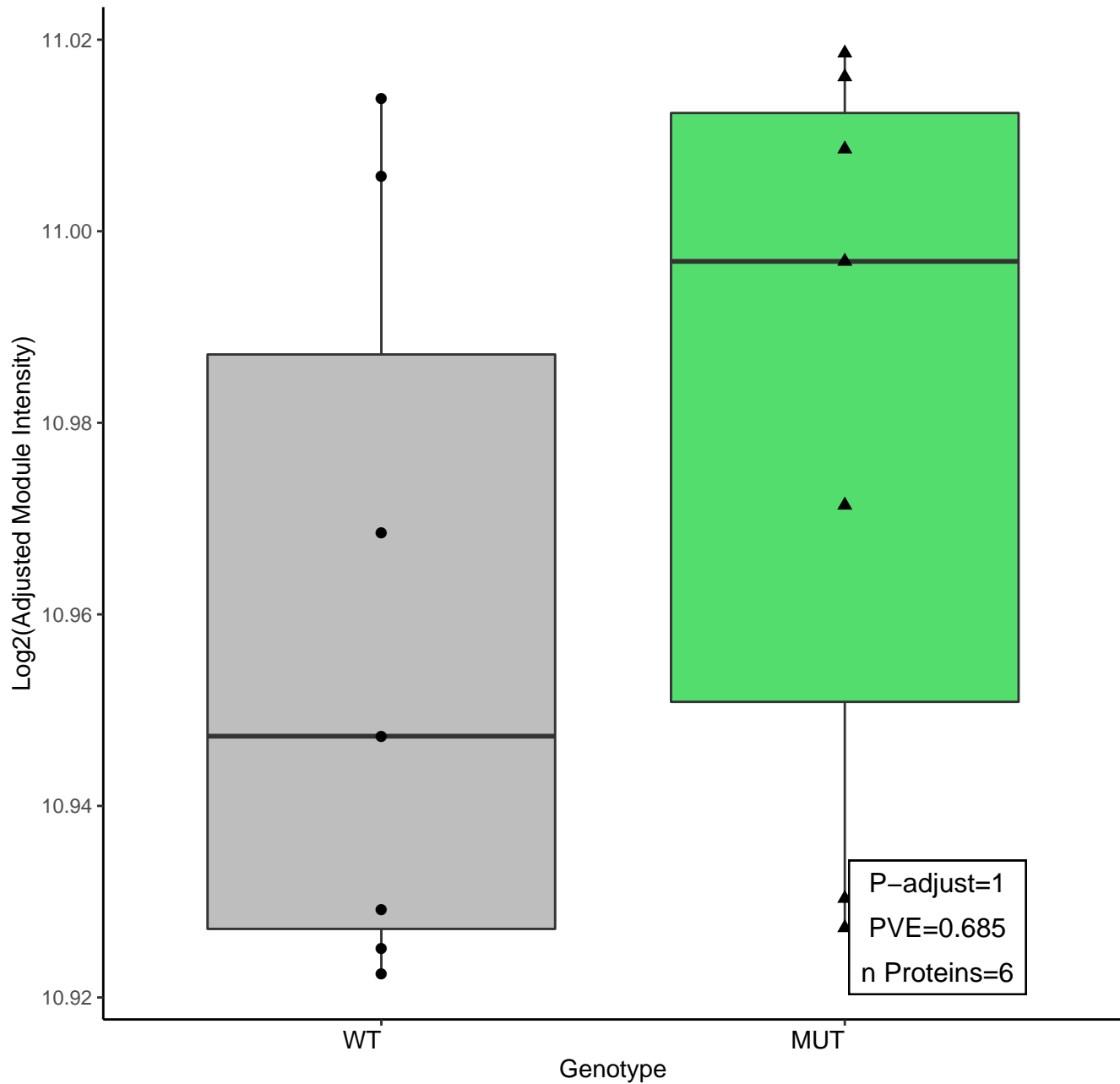
WT

Genotype

MUT



M80



M81

Log2(Adjusted Module Intensity)

15.20

15.15

15.10

15.05

WT

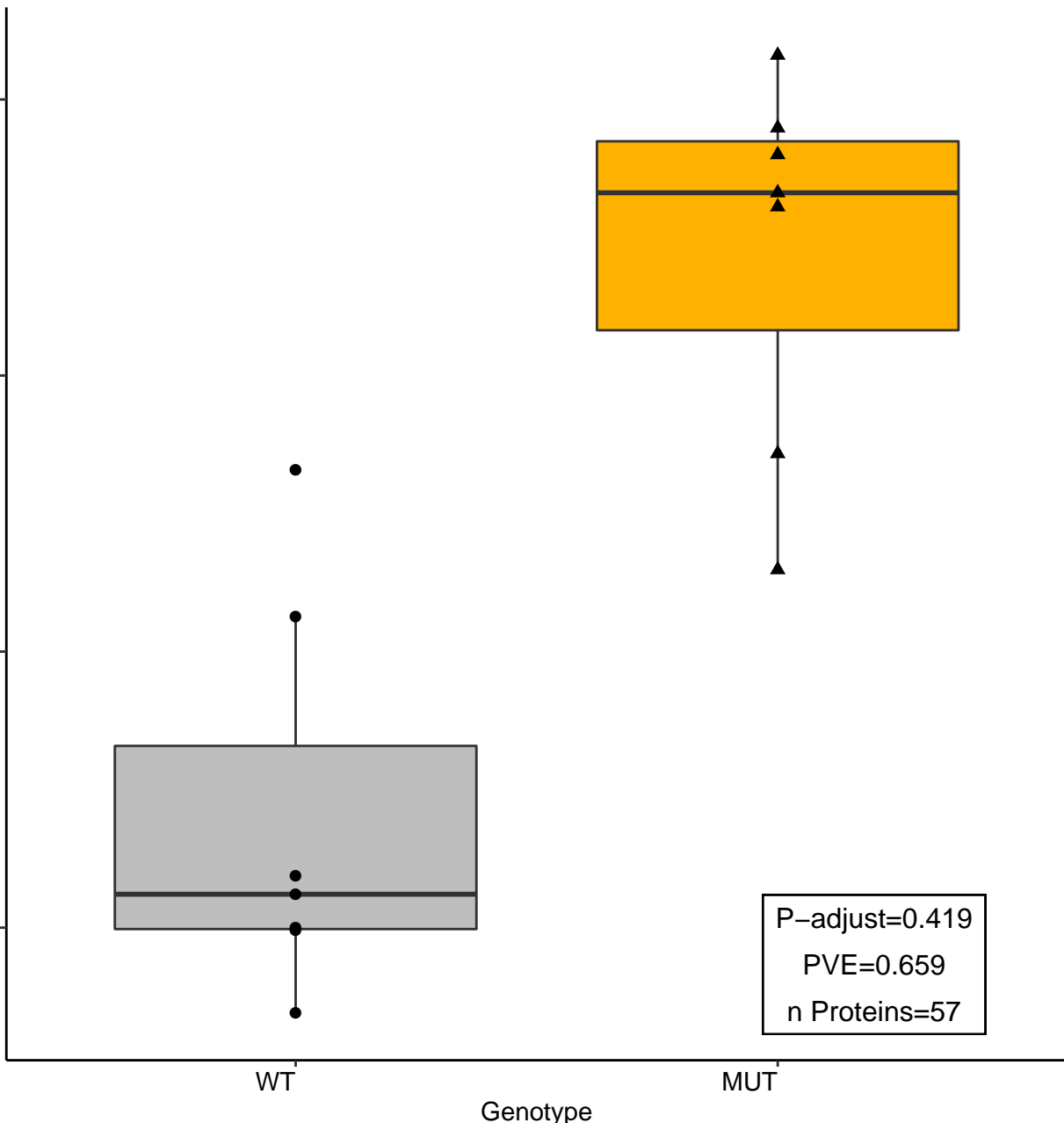
Genotype

MUT

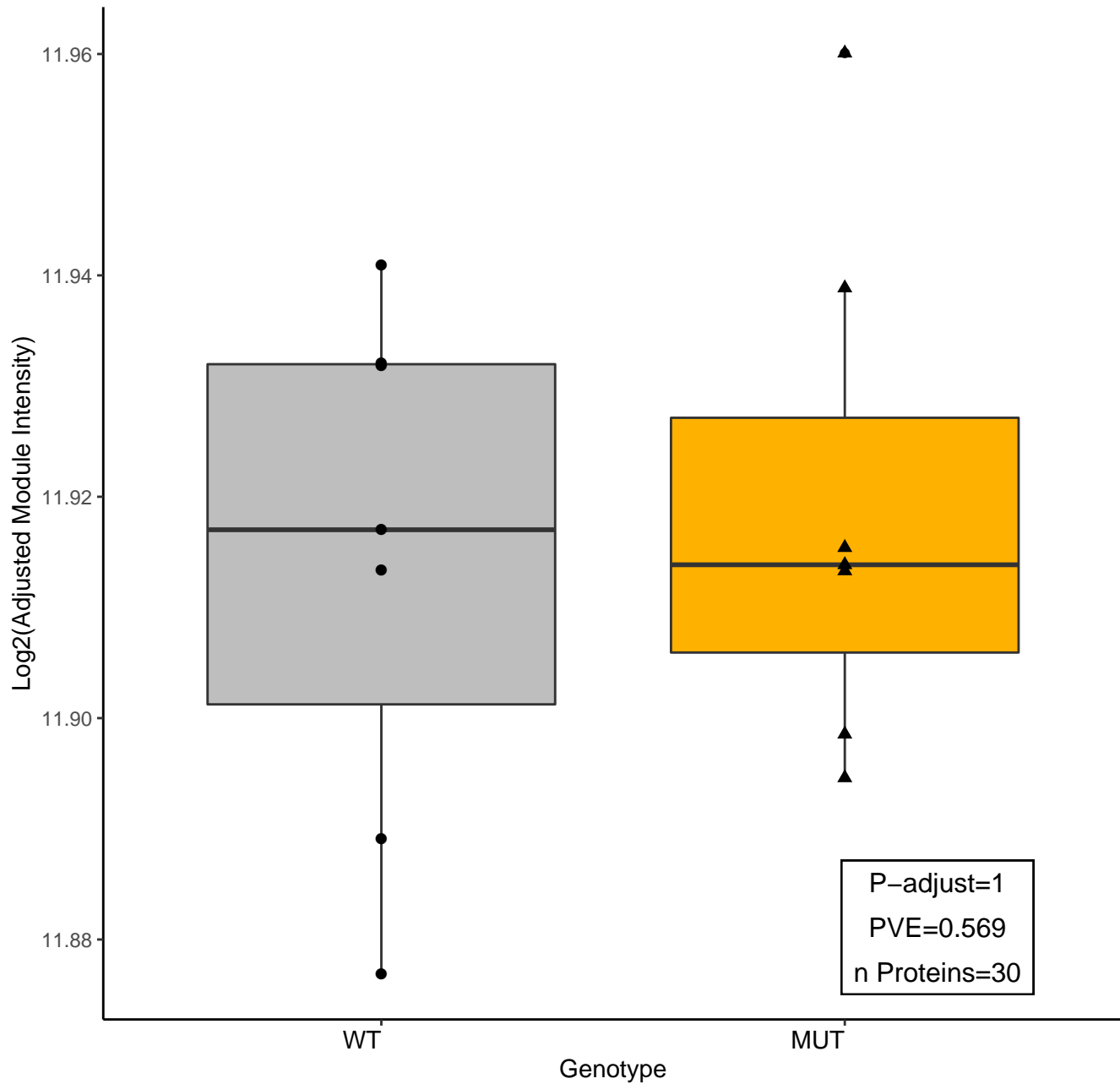
P-adjust=0.419

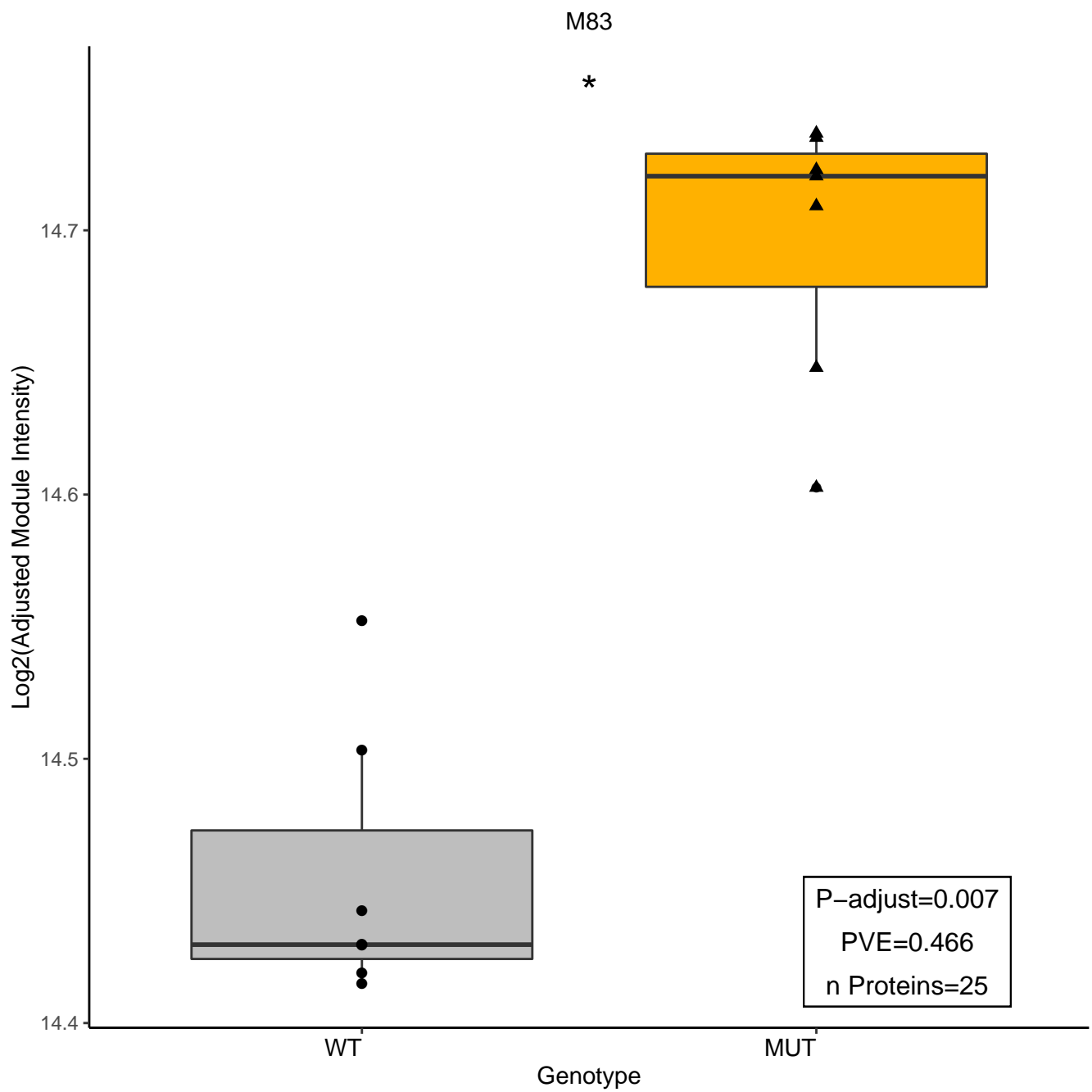
PVE=0.659

n Proteins=57



M82





M84

Log2(Adjusted Module Intensity)

11.58

11.56

11.54

11.52

11.50

11.48

WT

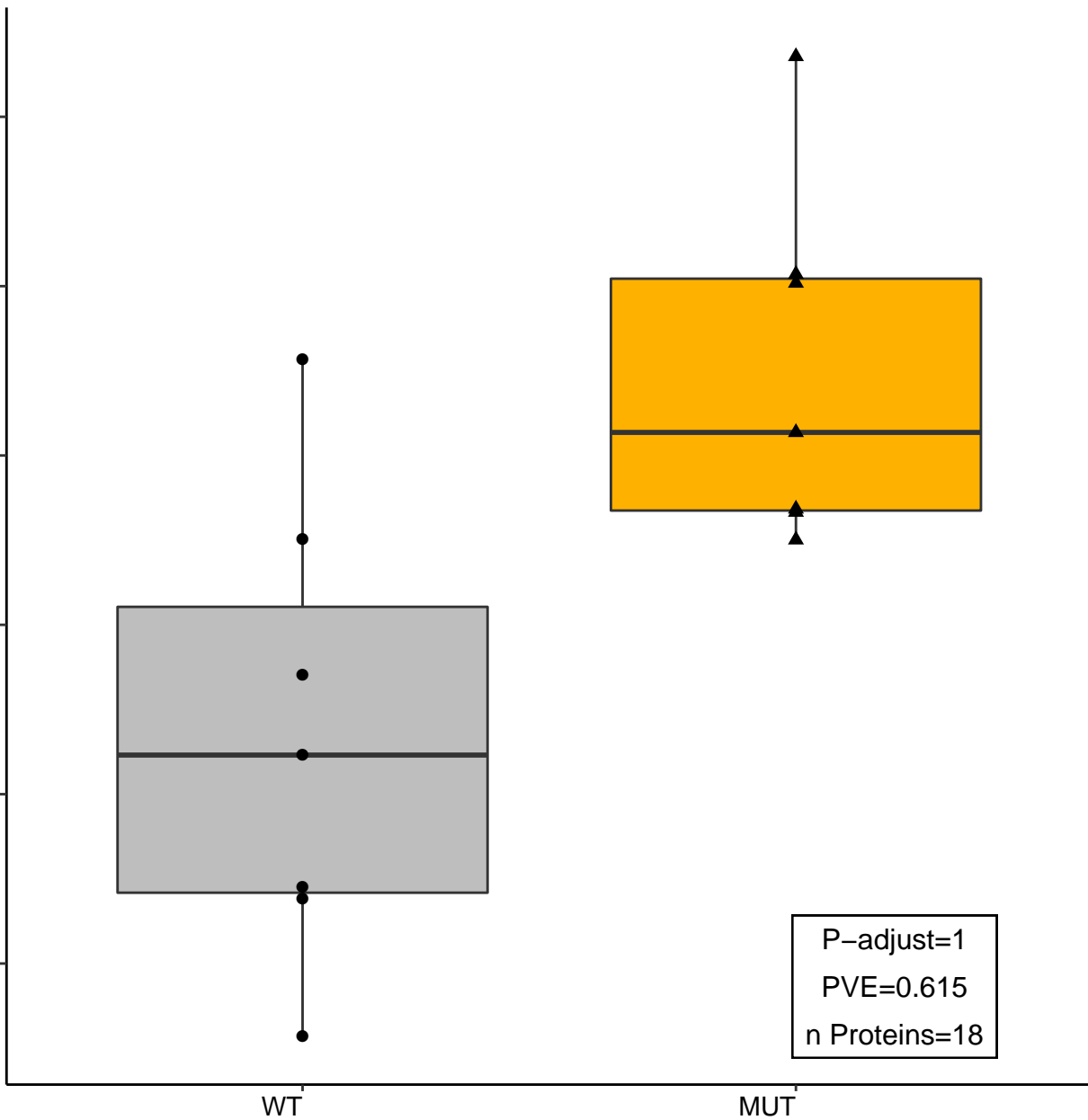
MUT

Genotype

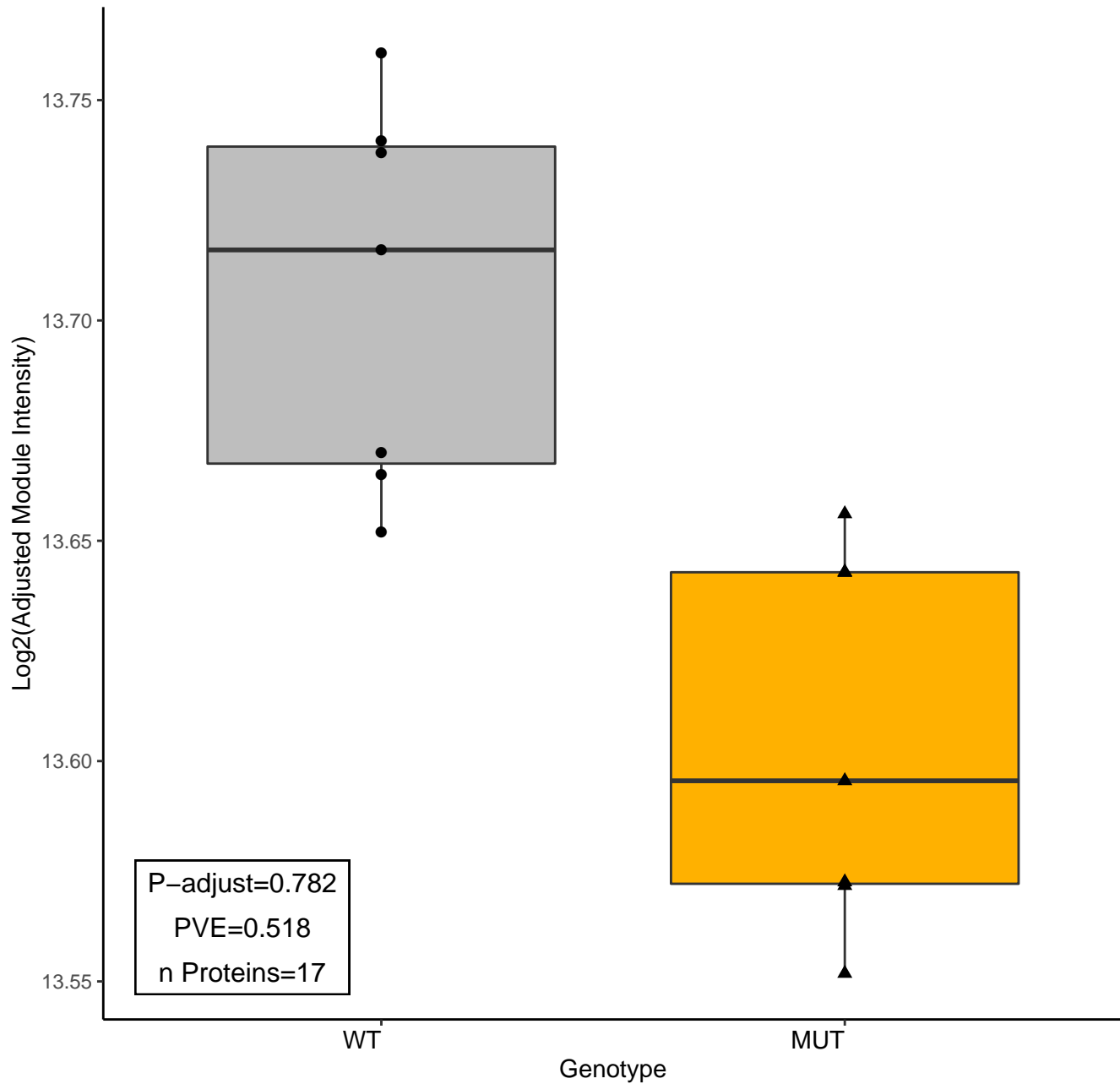
P-adjust=1

PVE=0.615

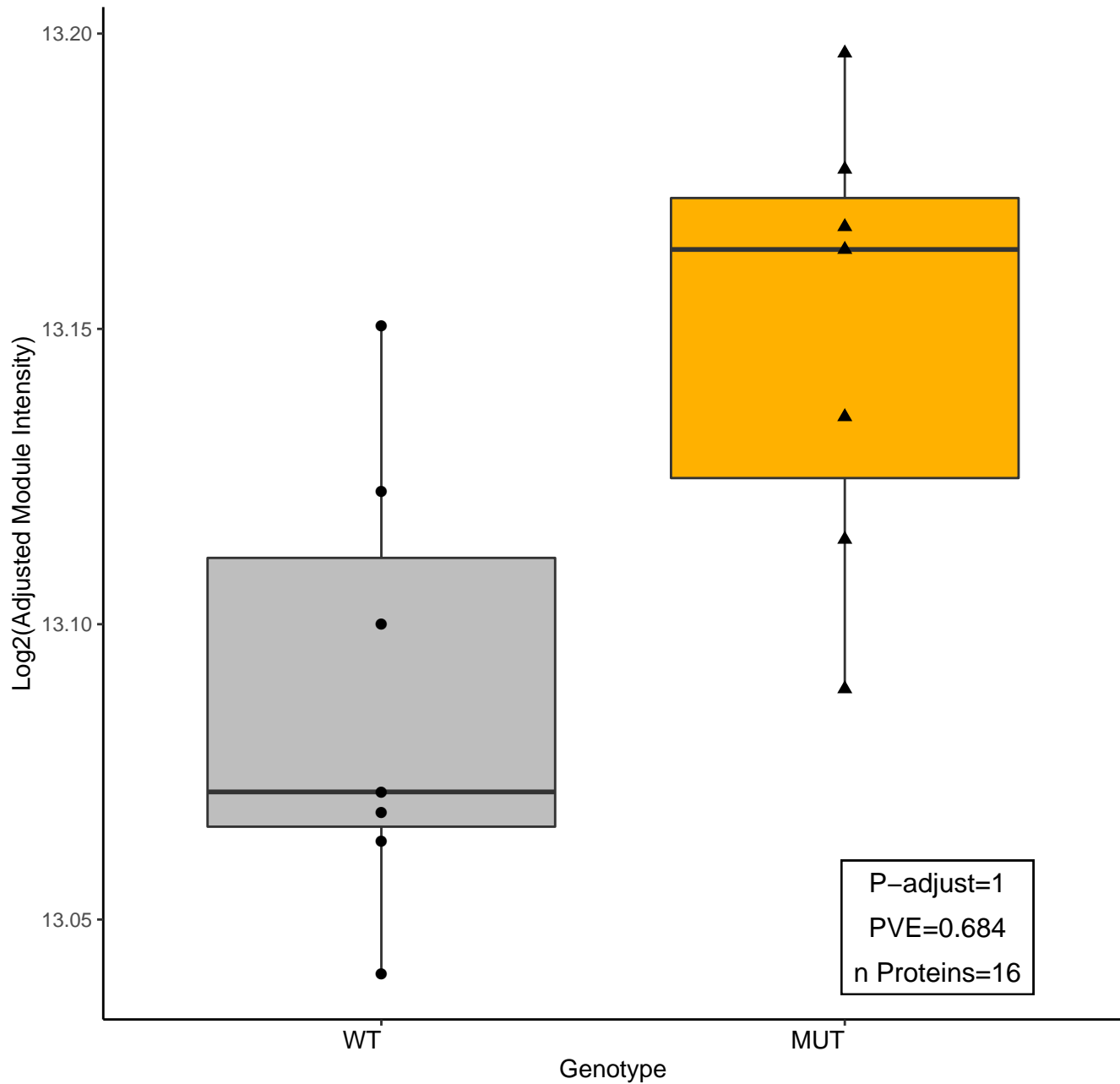
n Proteins=18



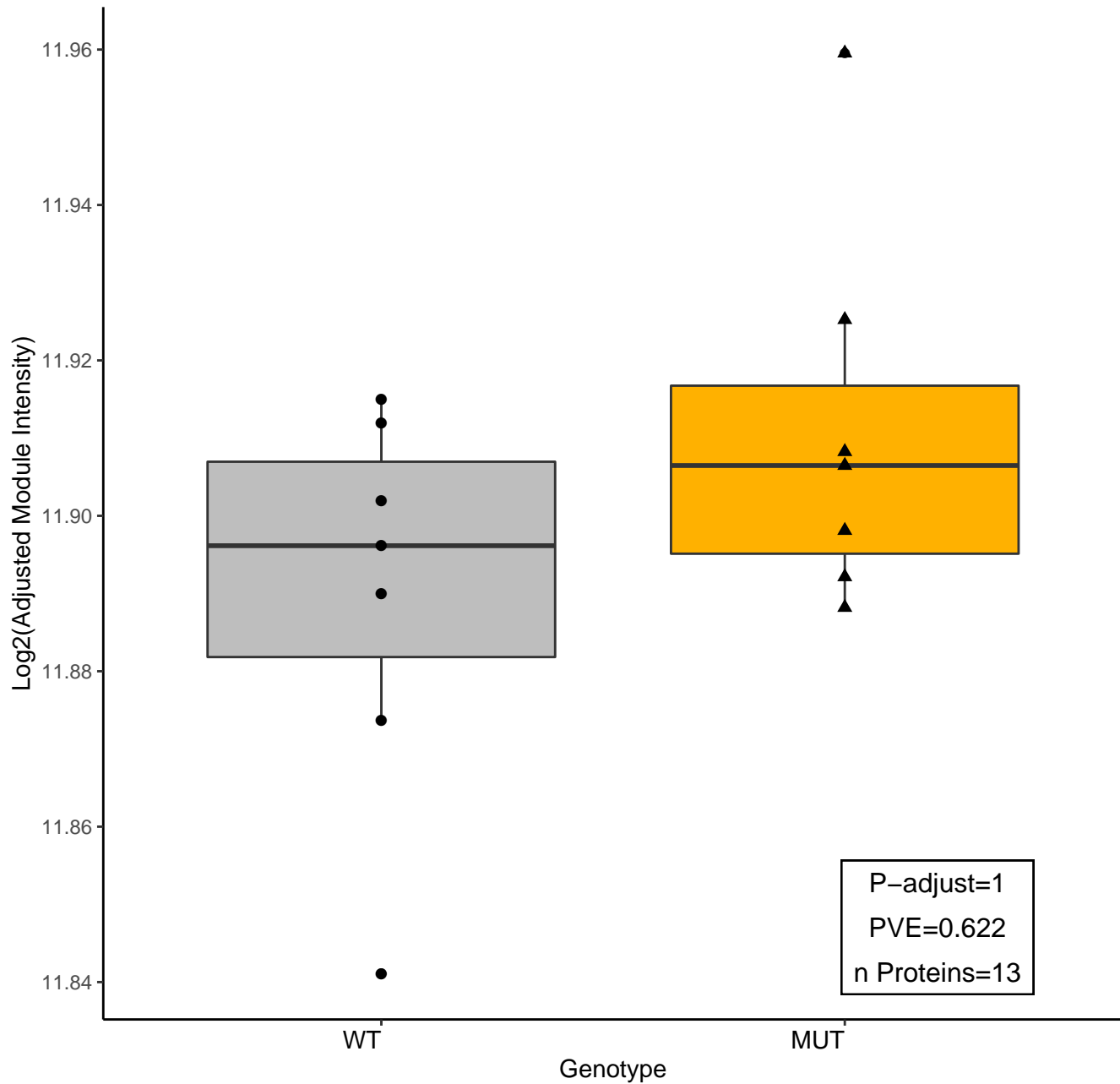
M85



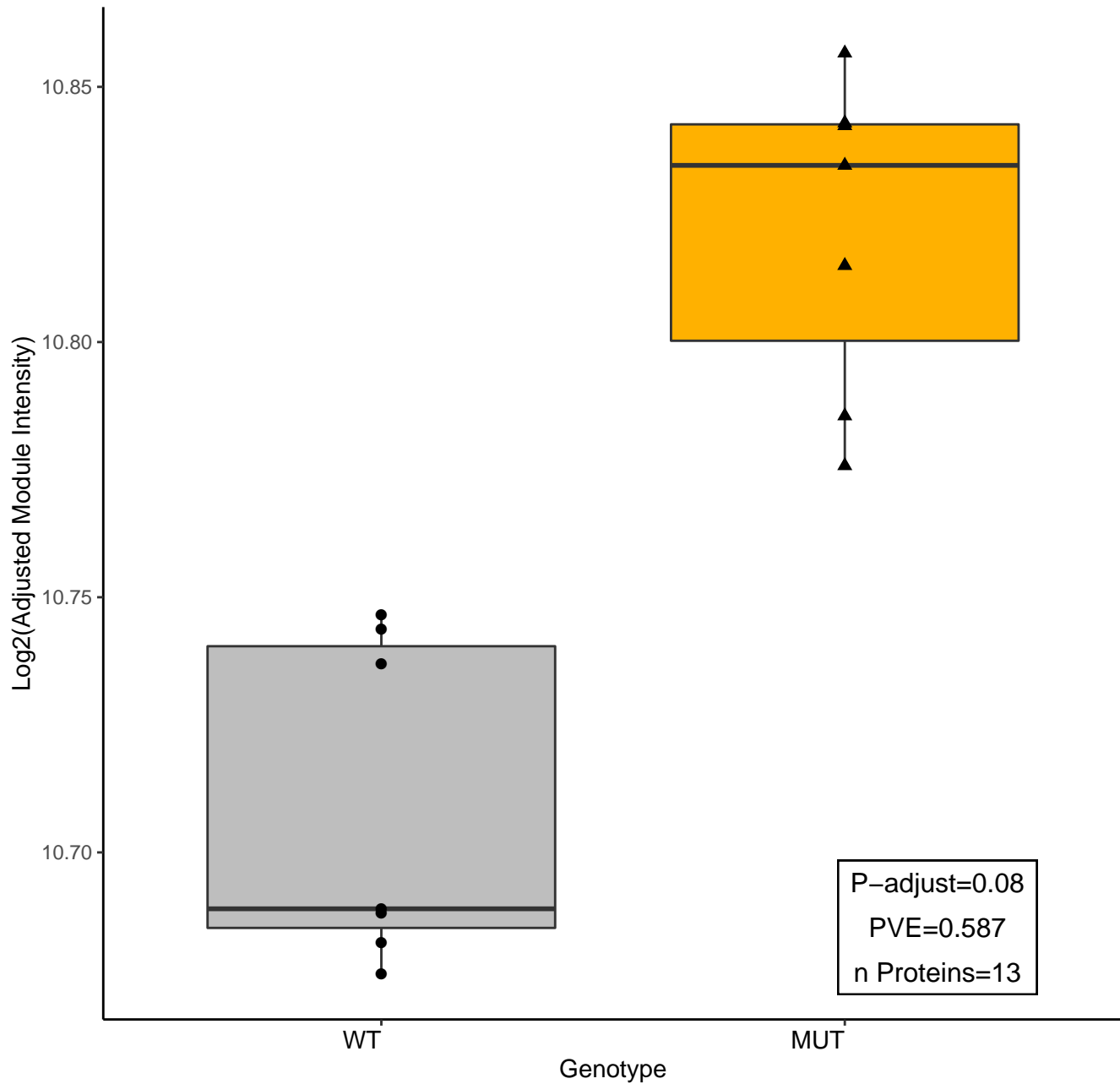
M86



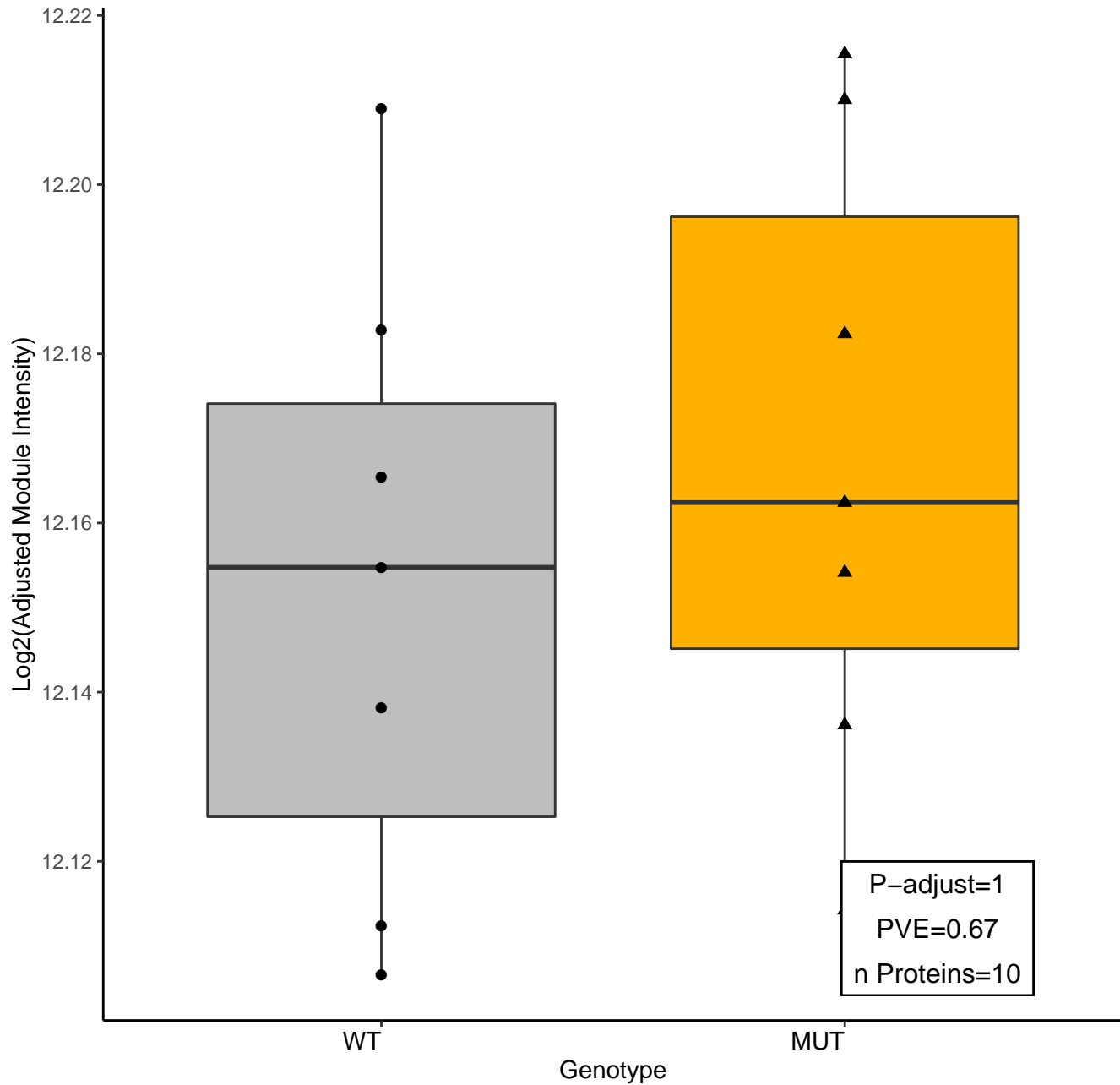
M87



M88



M89



M90

Log2(Adjusted Module Intensity)

9.20

9.15

9.10

WT

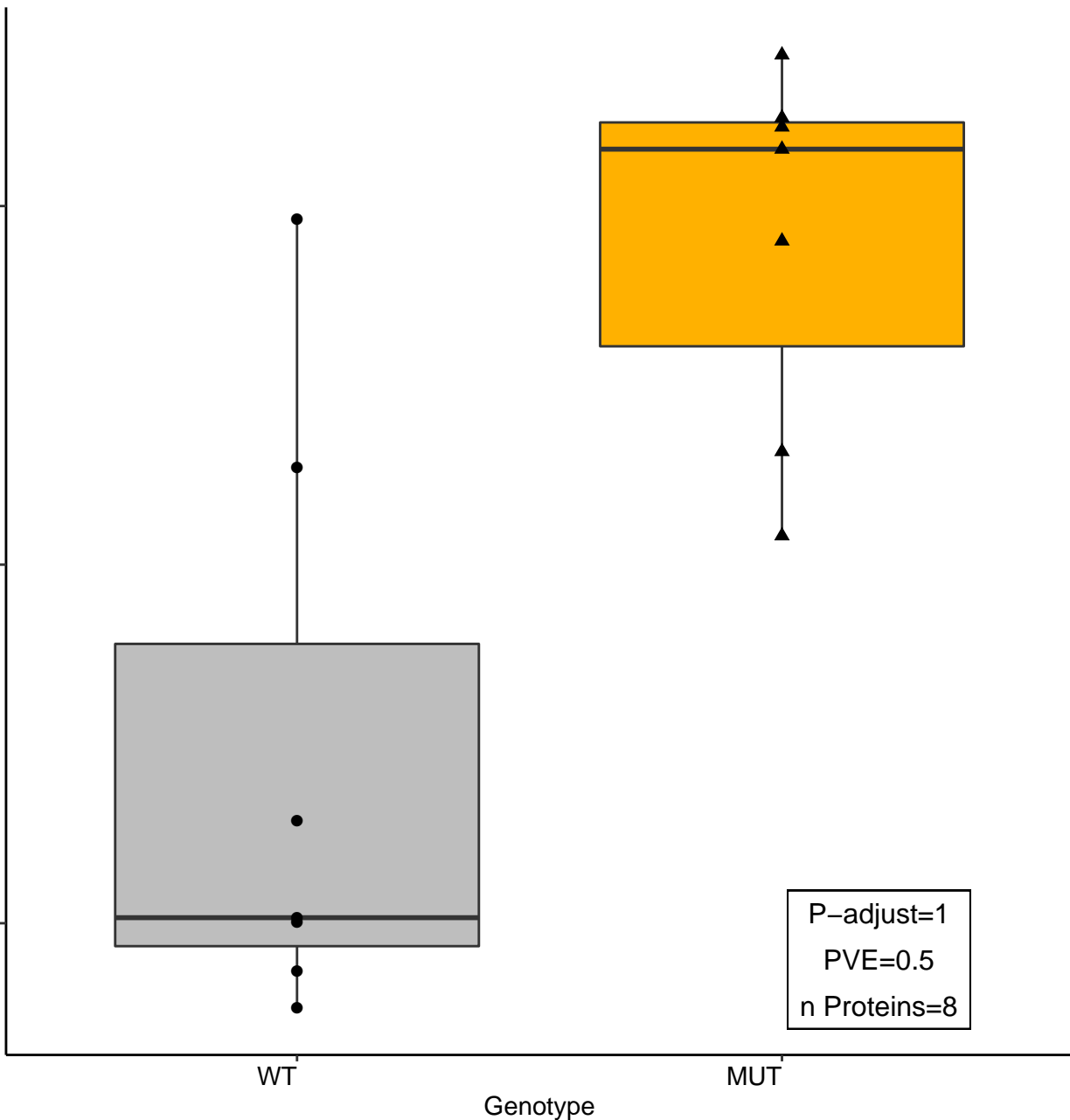
MUT

Genotype

P-adjust=1

PVE=0.5

n Proteins=8



M91

Log2(Adjusted Module Intensity)

10.04

10.02

10.00

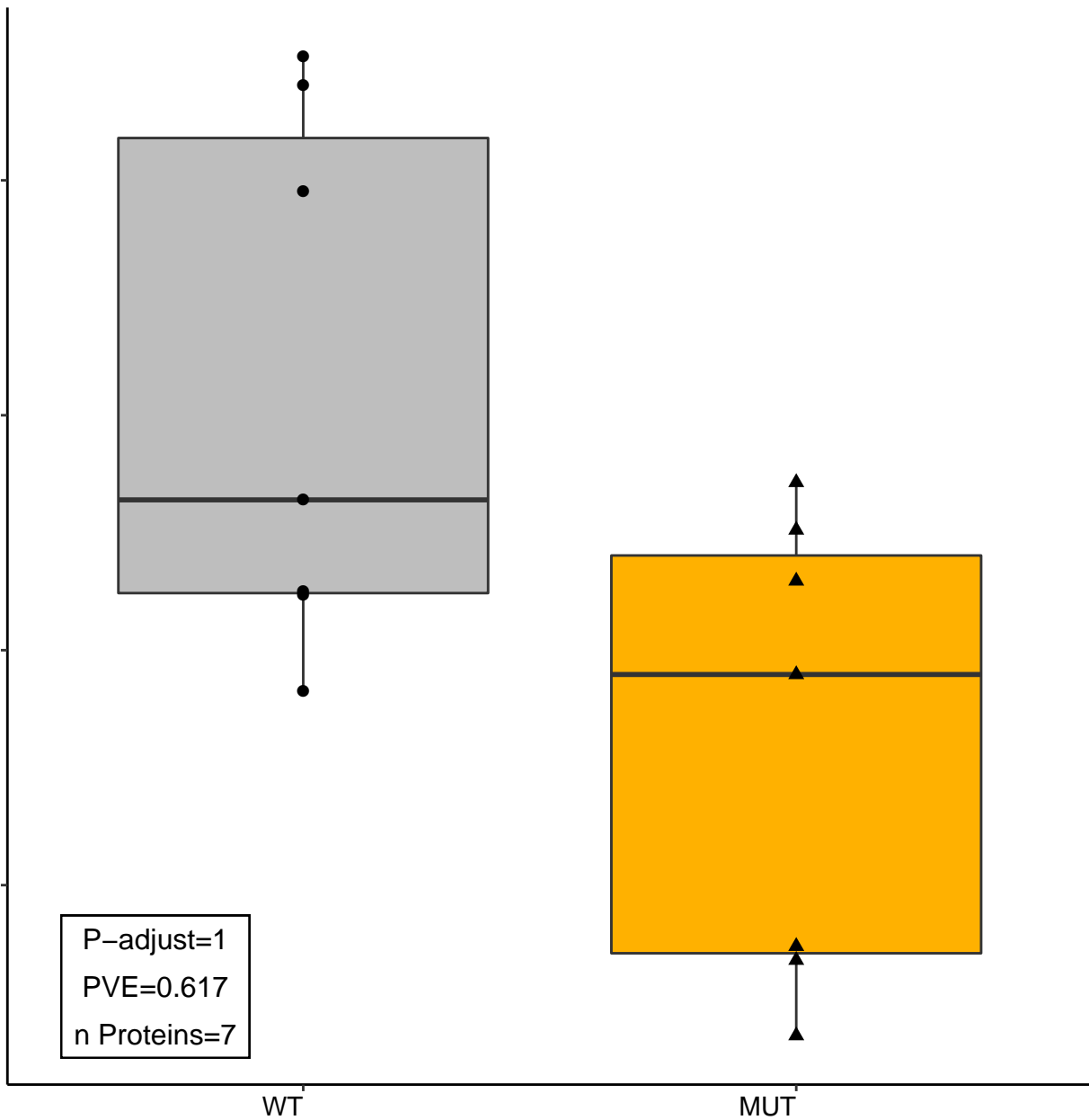
9.98

P-adjust=1
PVE=0.617
n Proteins=7

WT

Genotype

MUT



M92

Log2(Adjusted Module Intensity)

14.08

14.06

14.04

14.02

WT

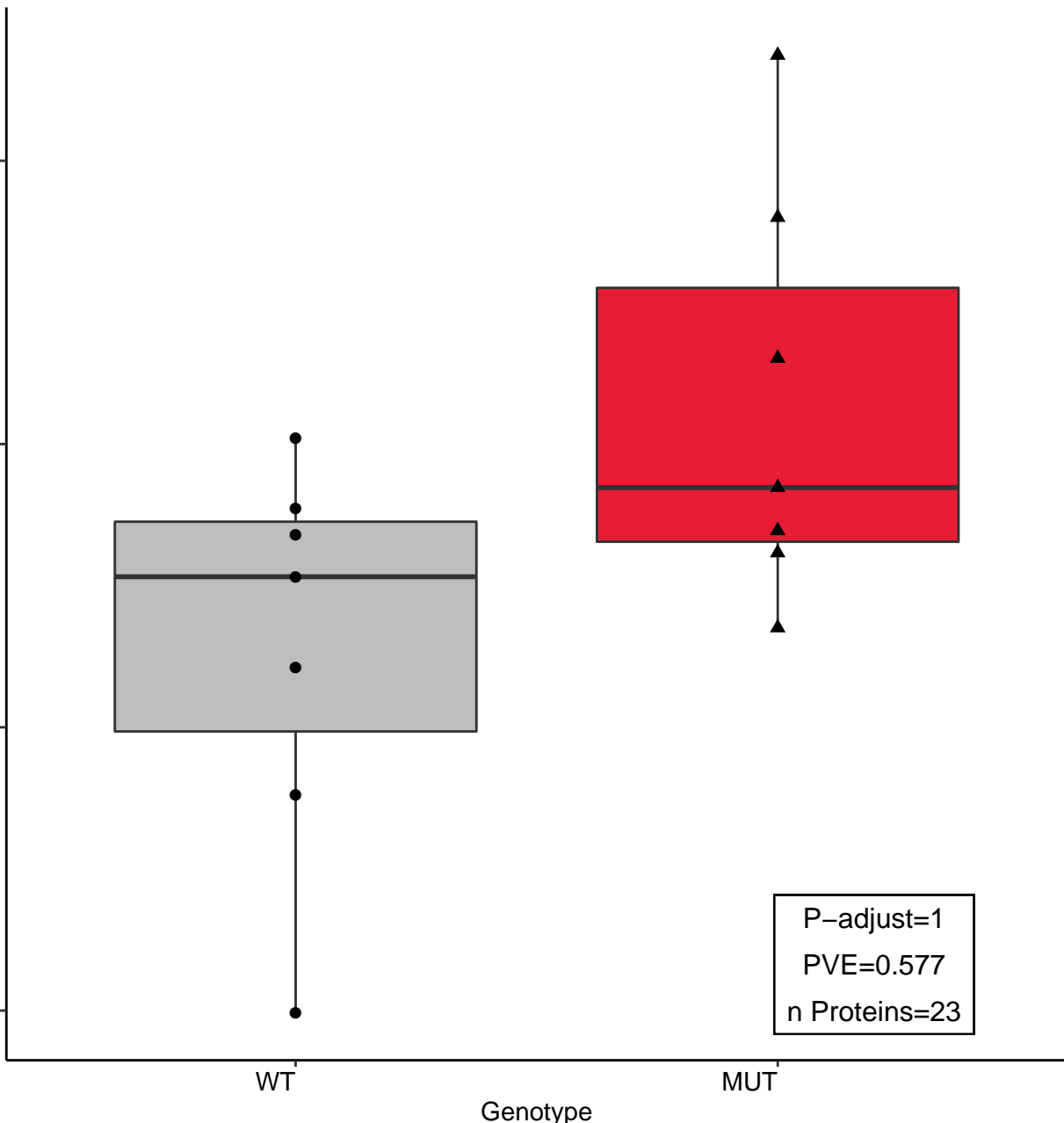
Genotype

MUT

P-adjust=1

PVE=0.577

n Proteins=23



M93

Log2(Adjusted Module Intensity)

13.66

13.64

13.62

13.60

WT

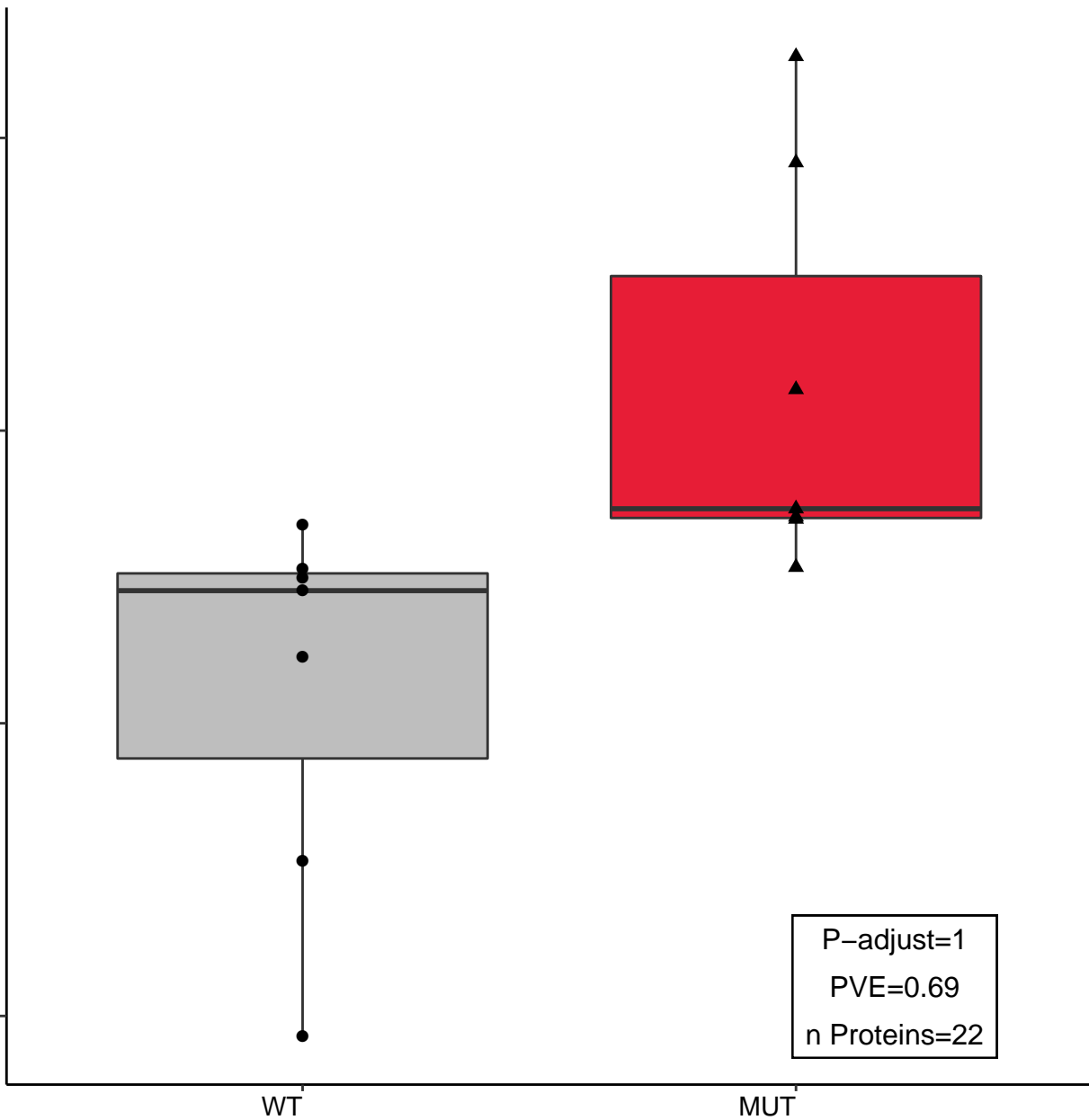
Genotype

MUT

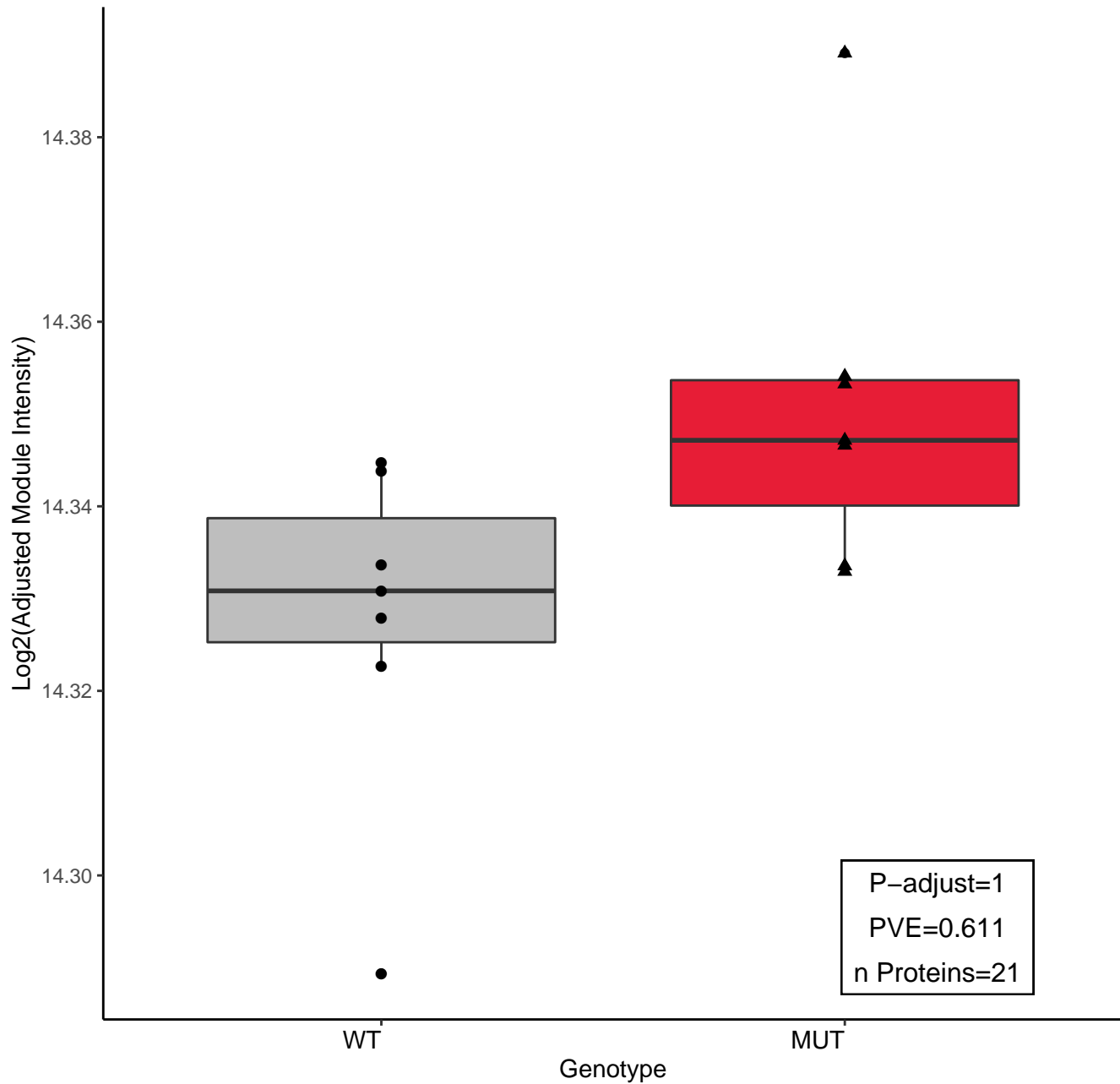
P-adjust=1

PVE=0.69

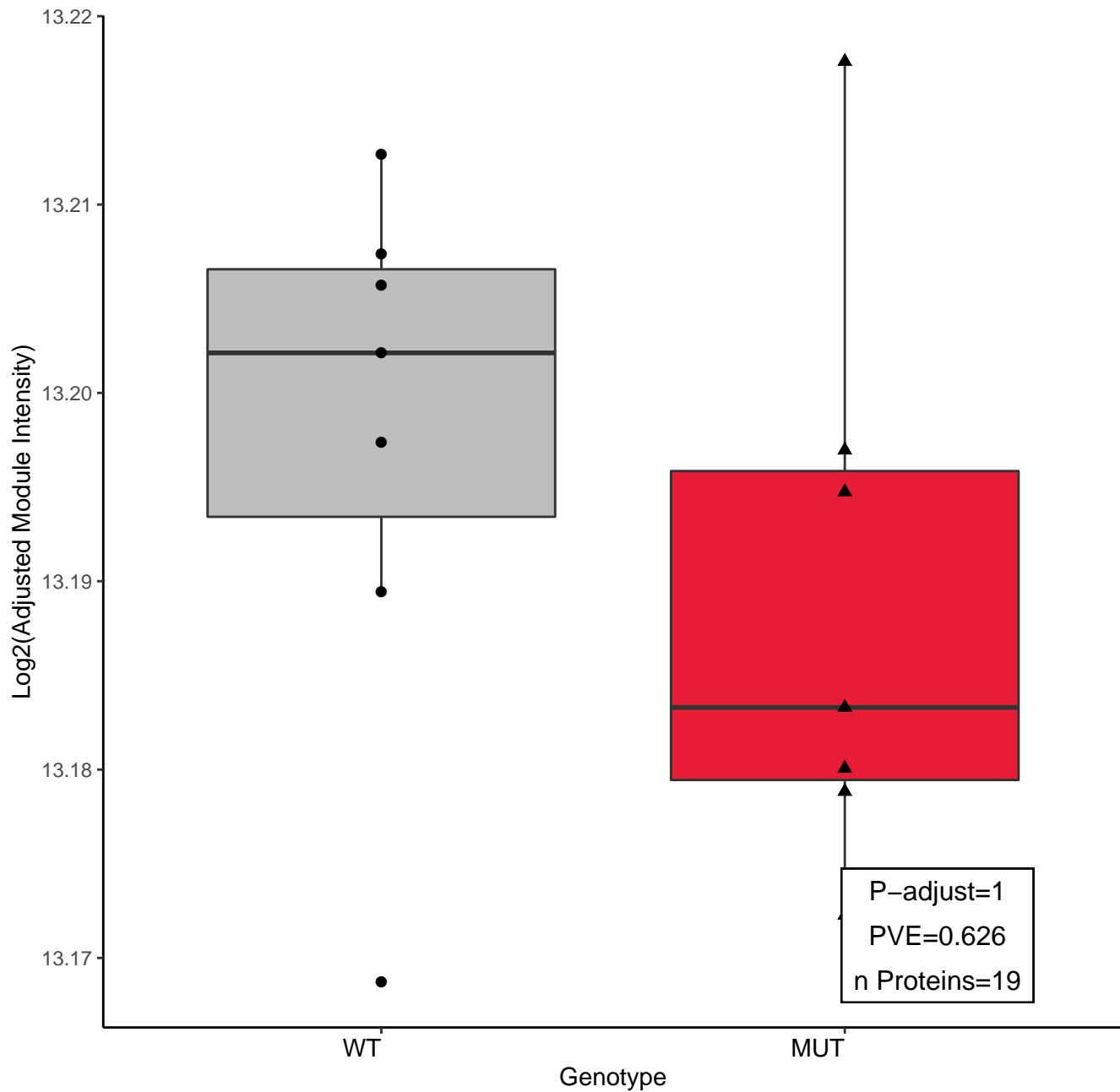
n Proteins=22



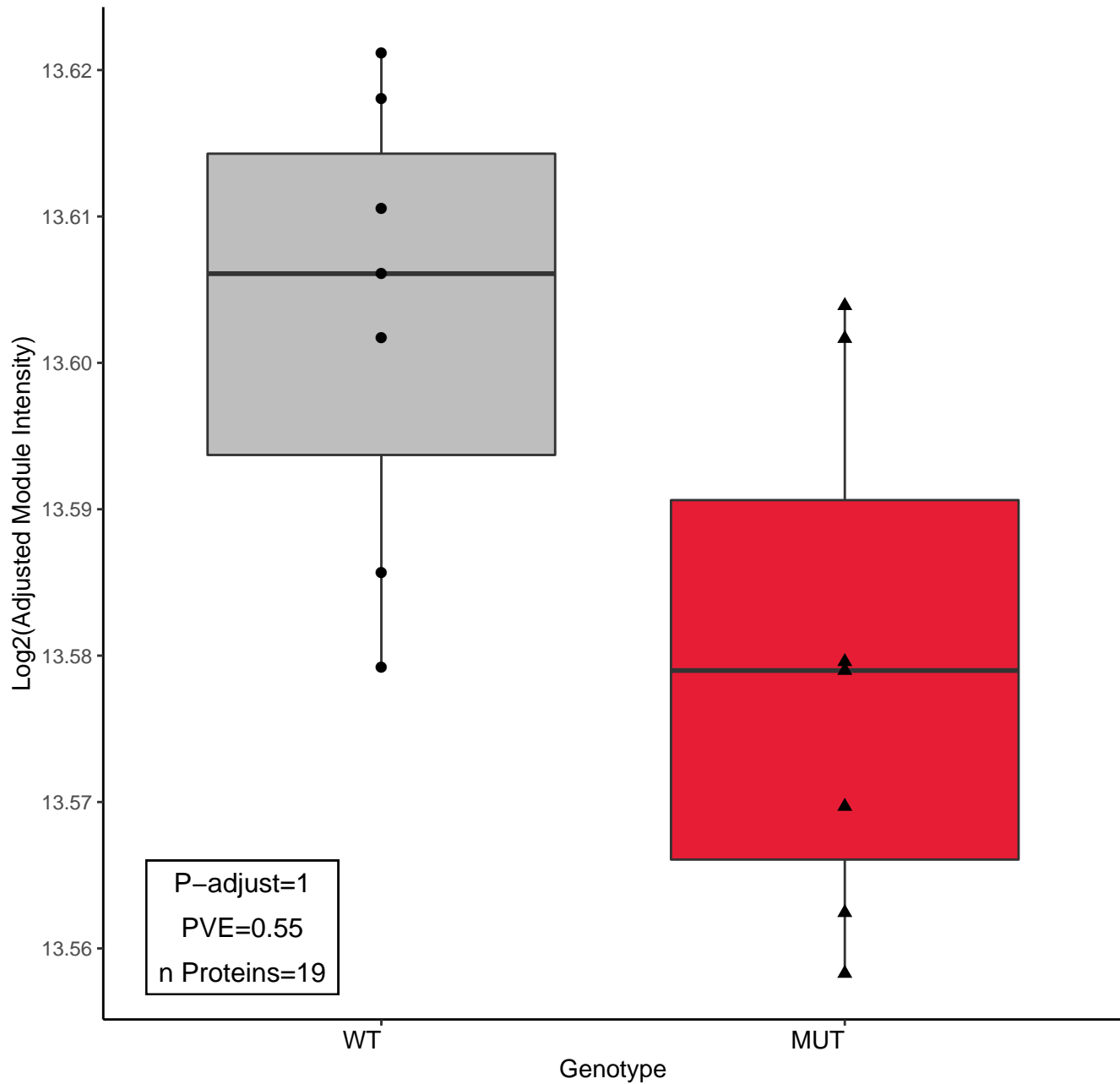
M94

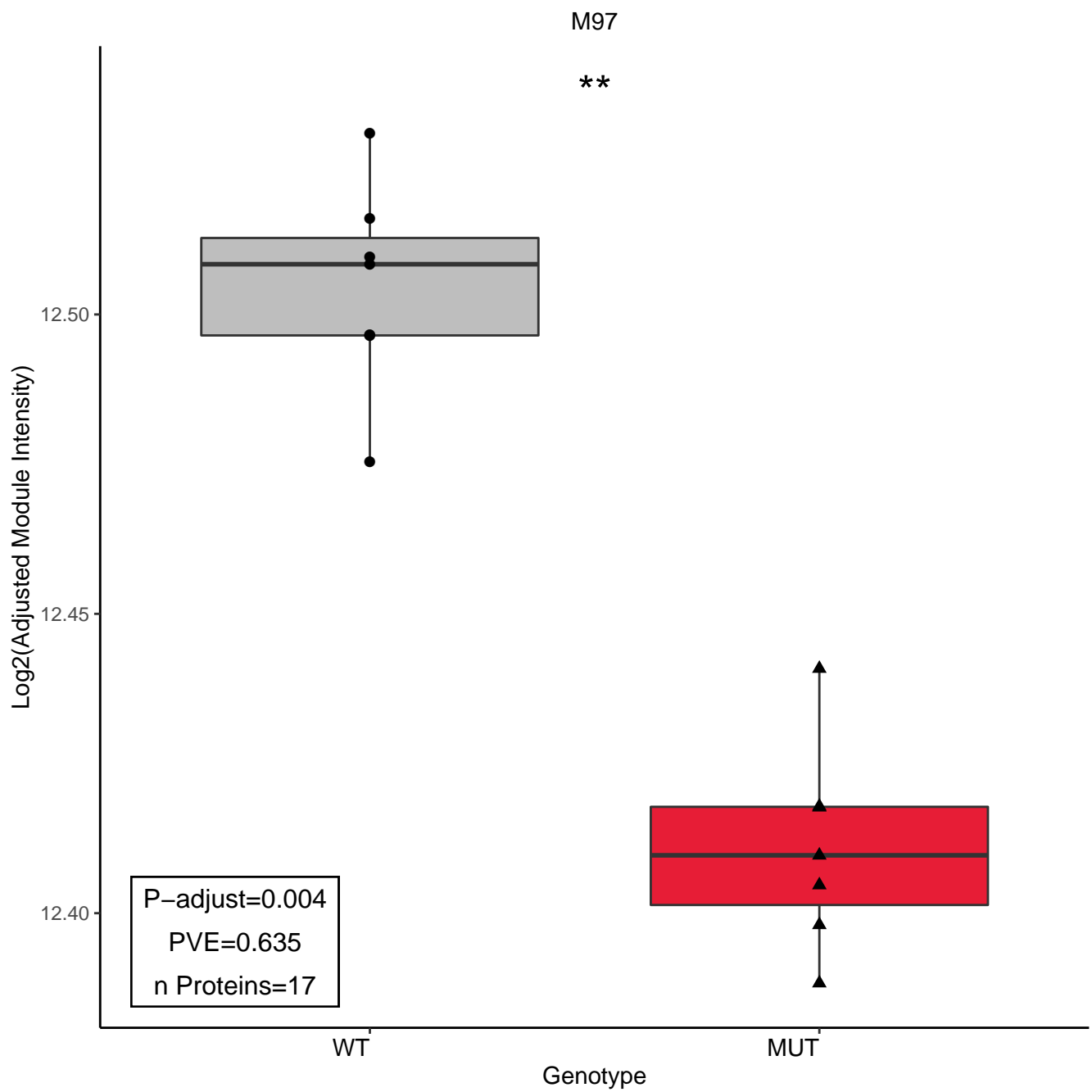


M95

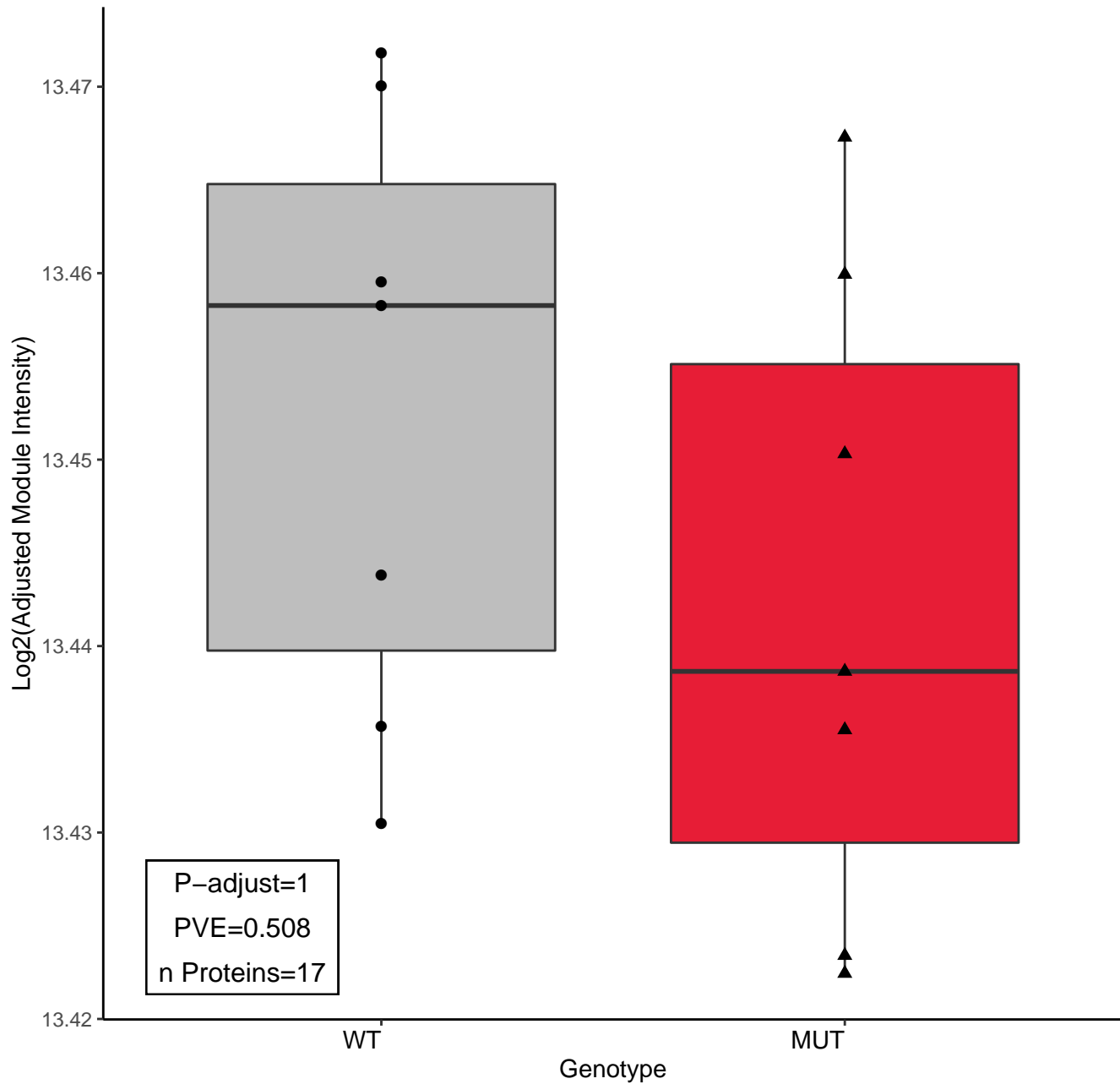


M96

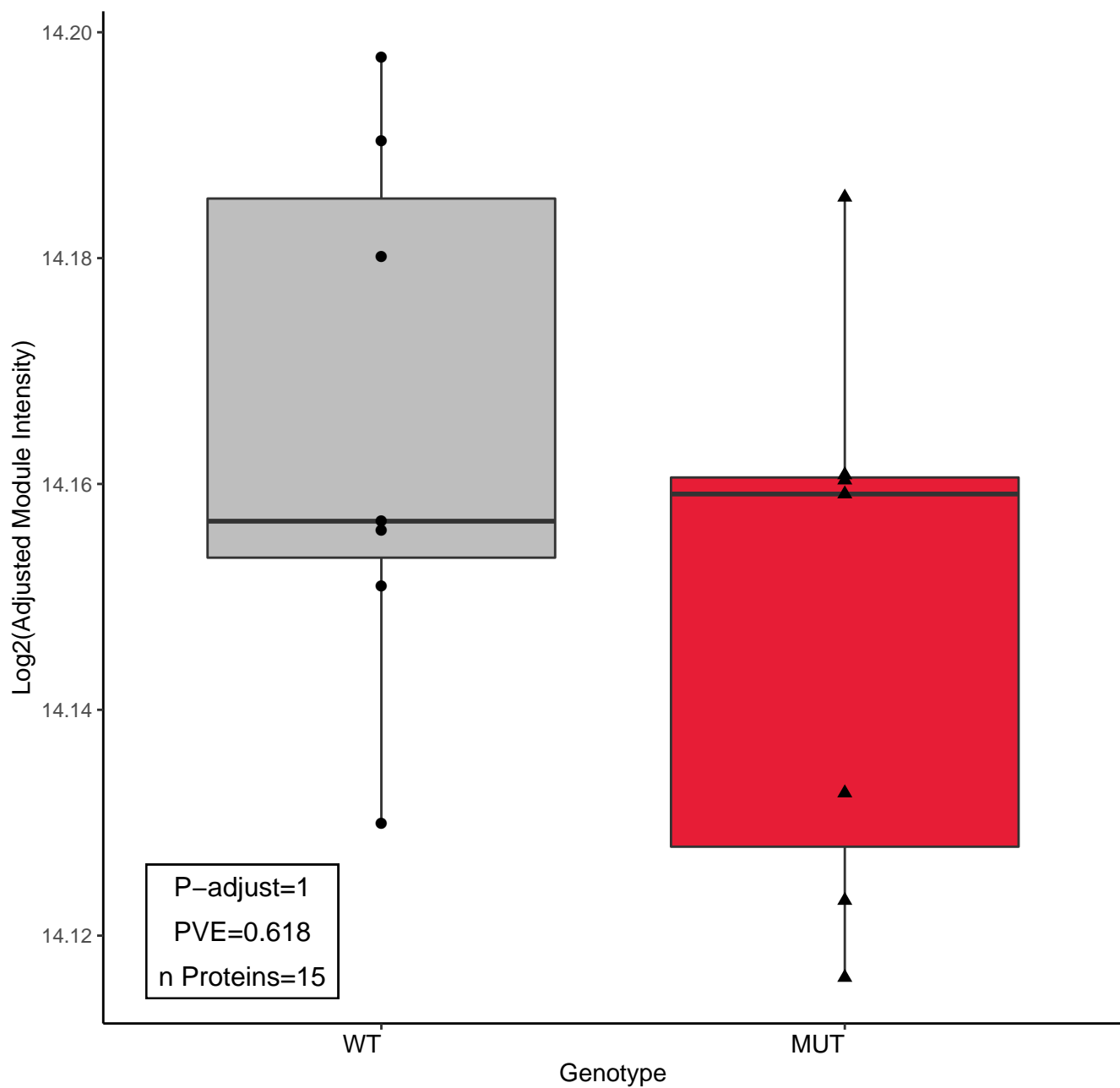




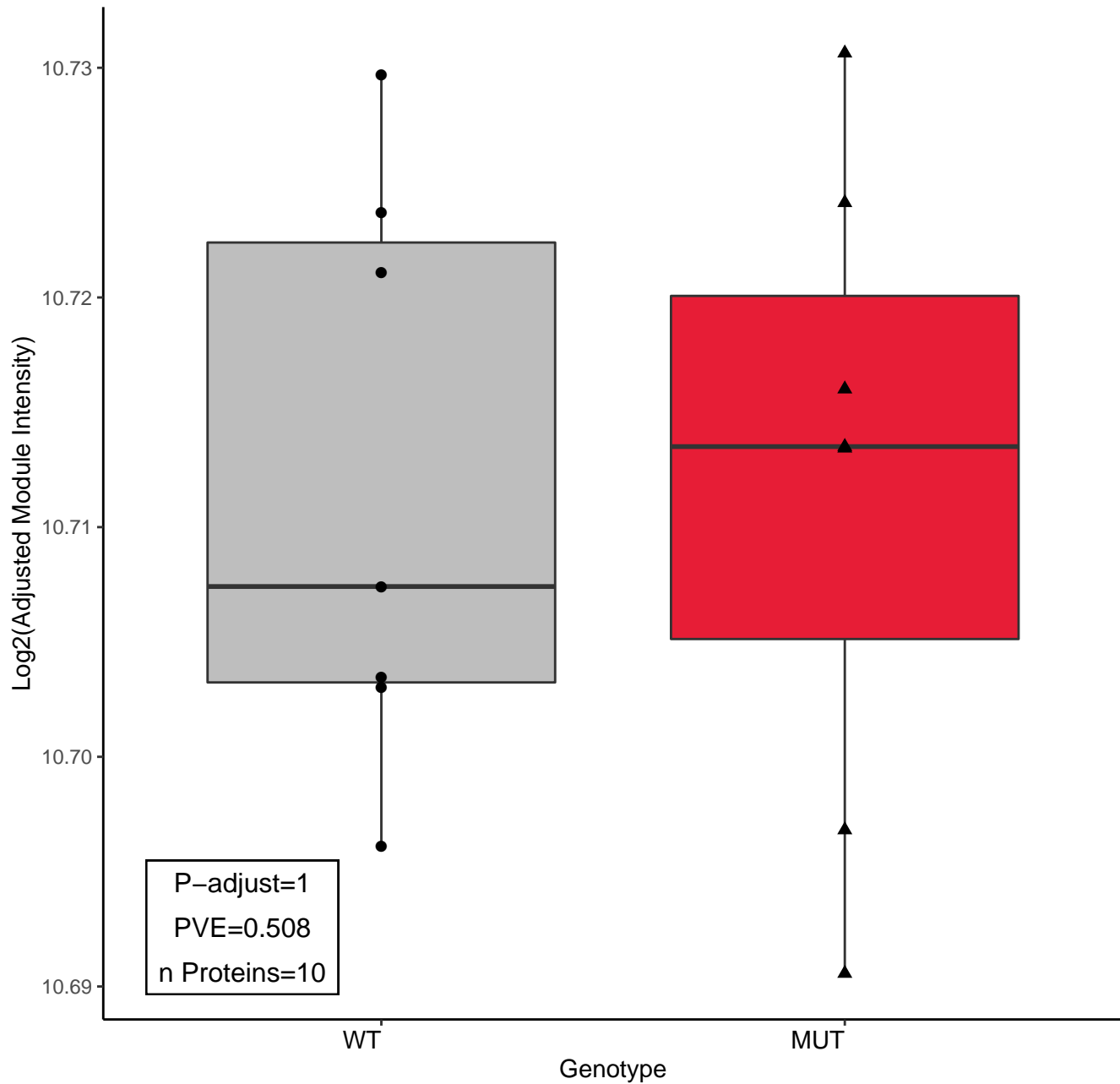
M98



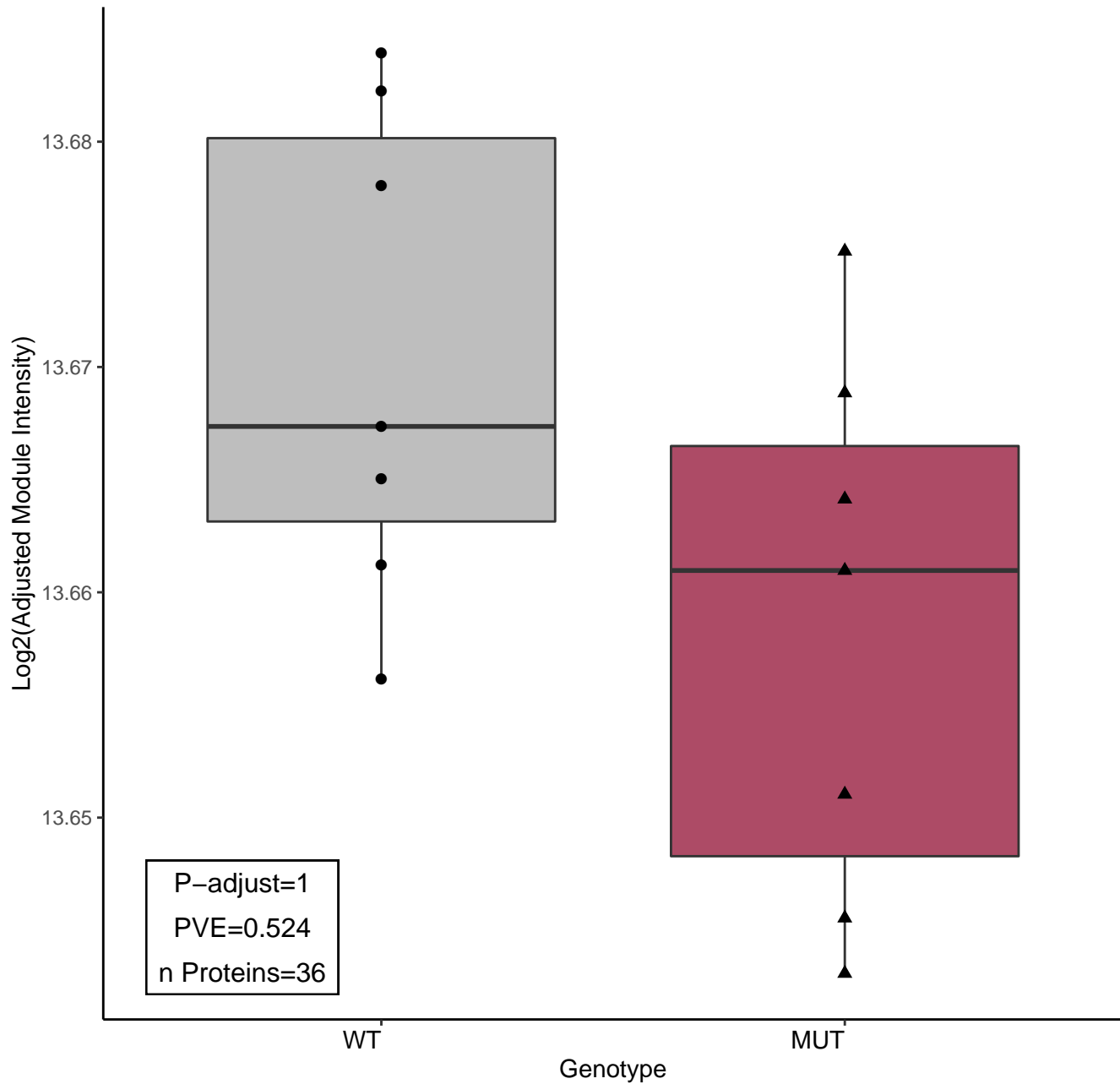
M99



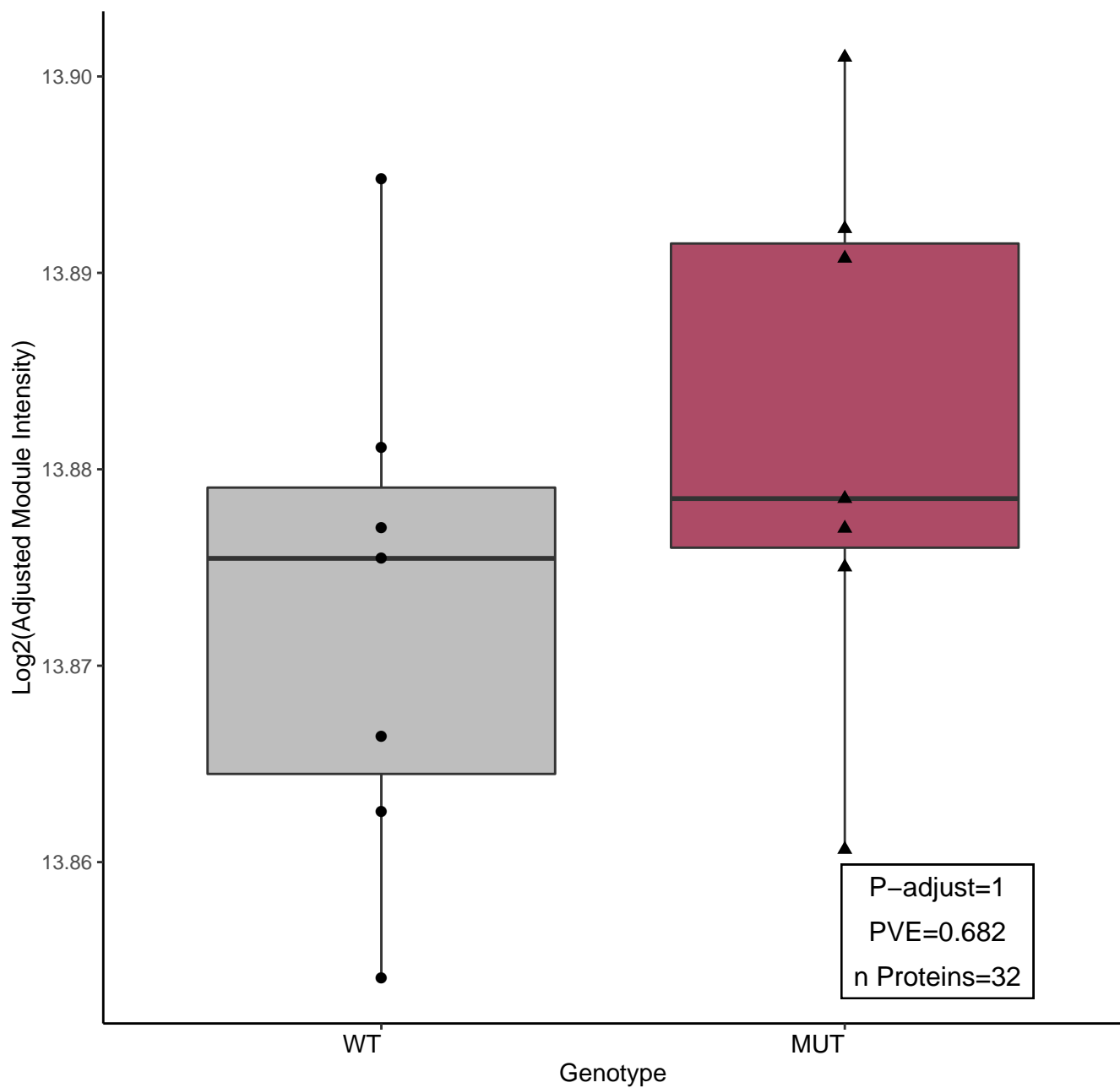
M100



M101

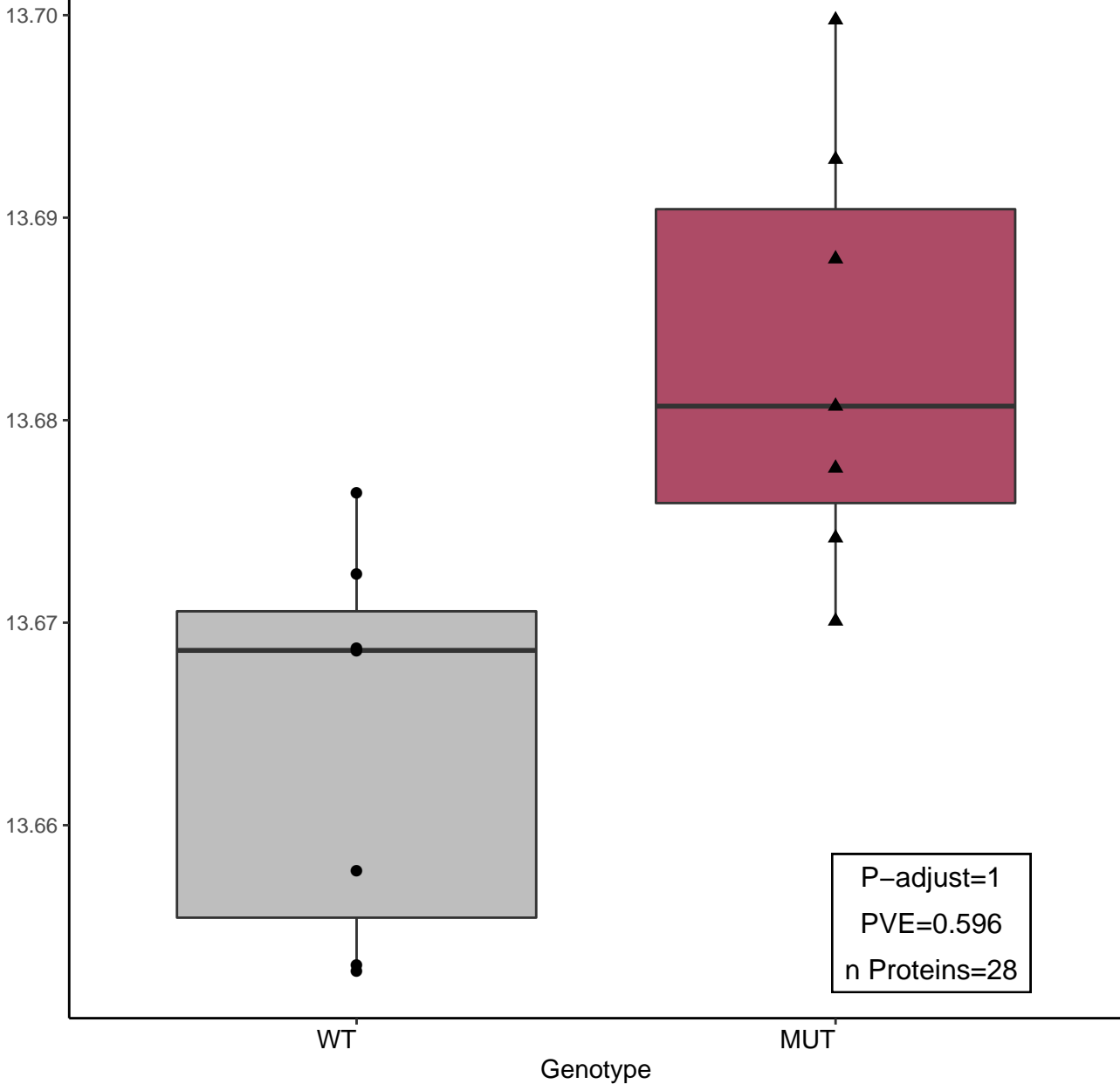


M102

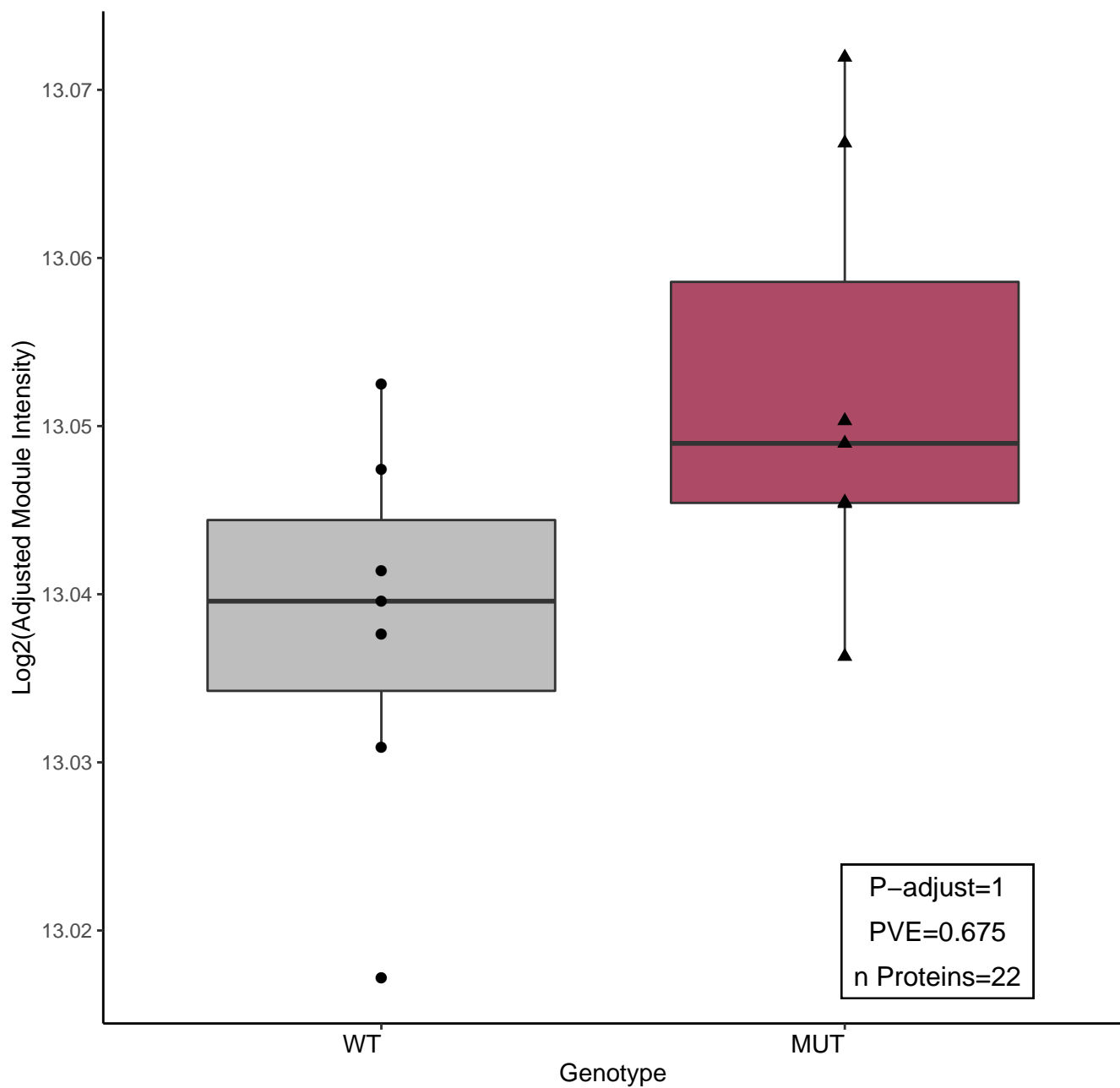


M103

Log2(Adjusted Module Intensity)

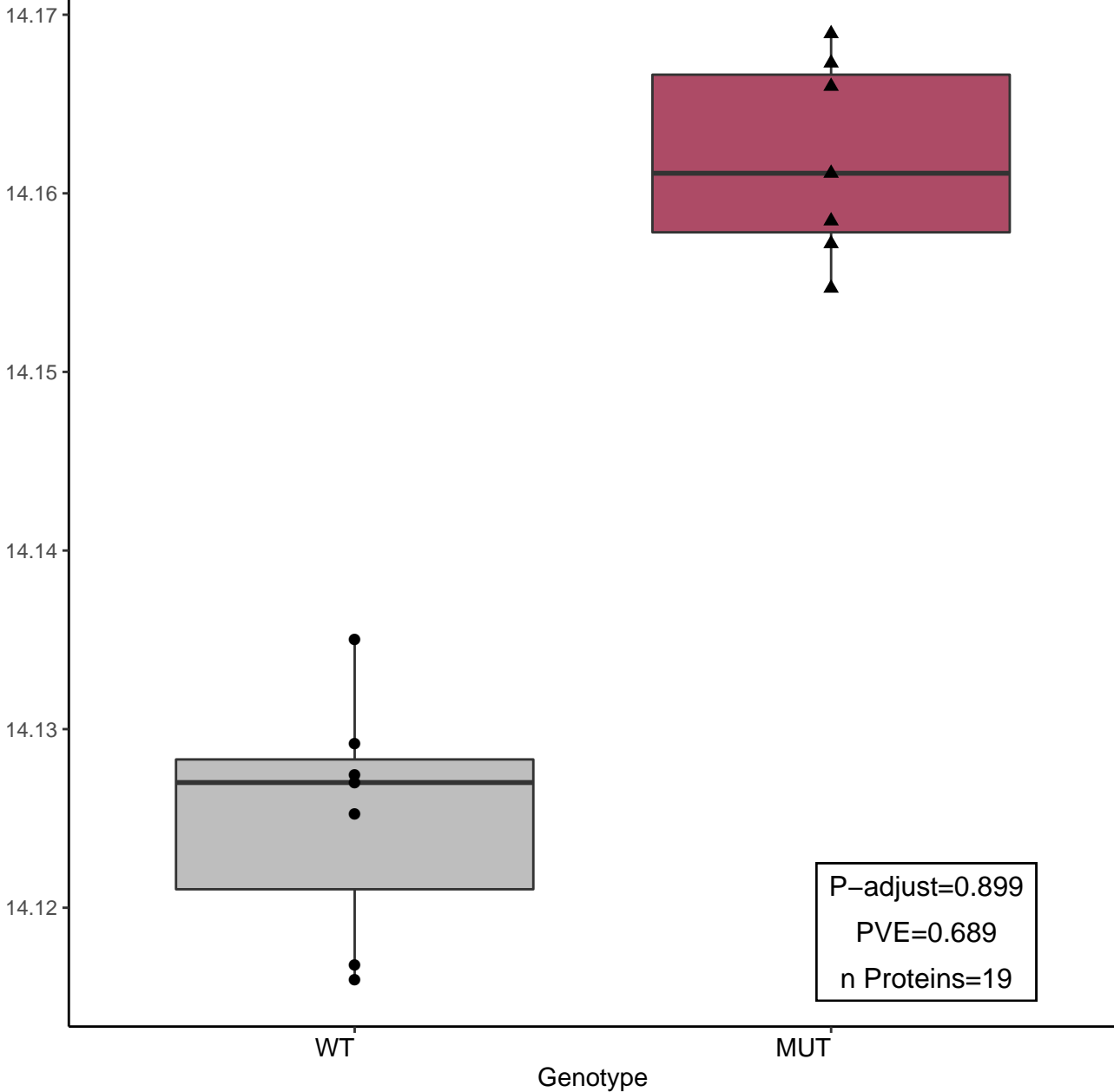


M104

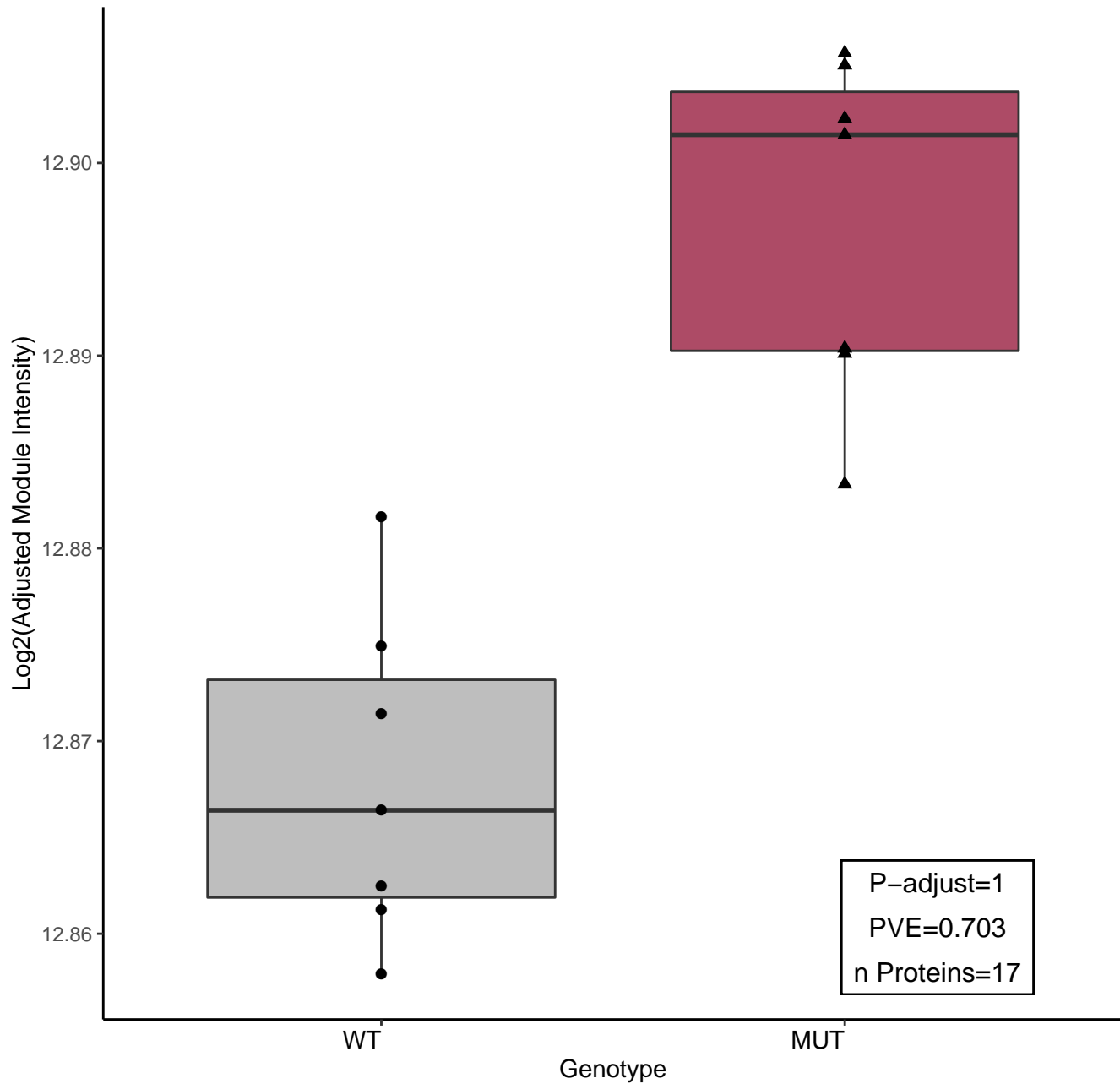


M105

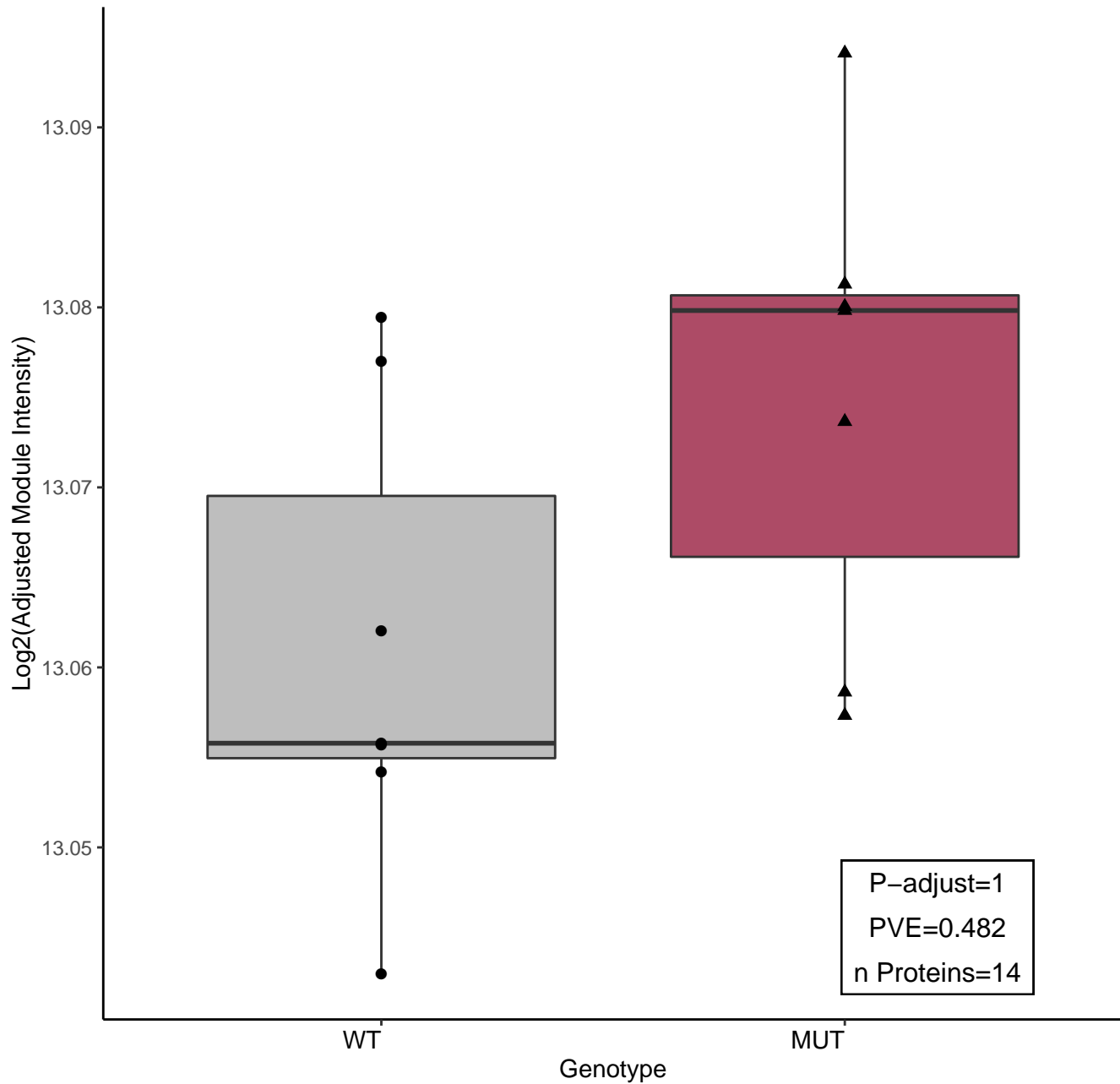
Log2(Adjusted Module Intensity)



M106



M107



M108

Log2(Adjusted Module Intensity)

10.20

10.18

10.16

10.14

WT

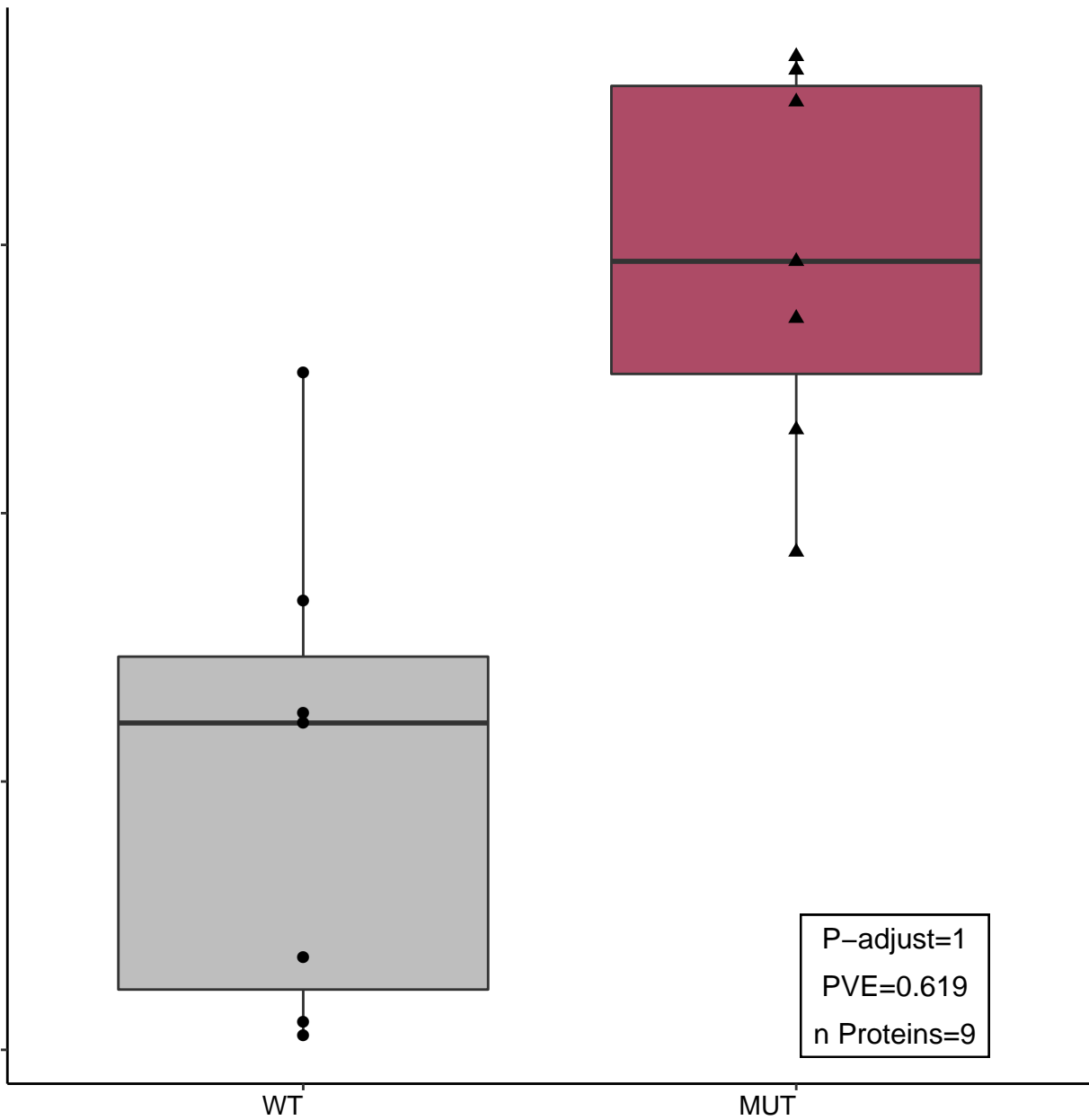
Genotype

MUT

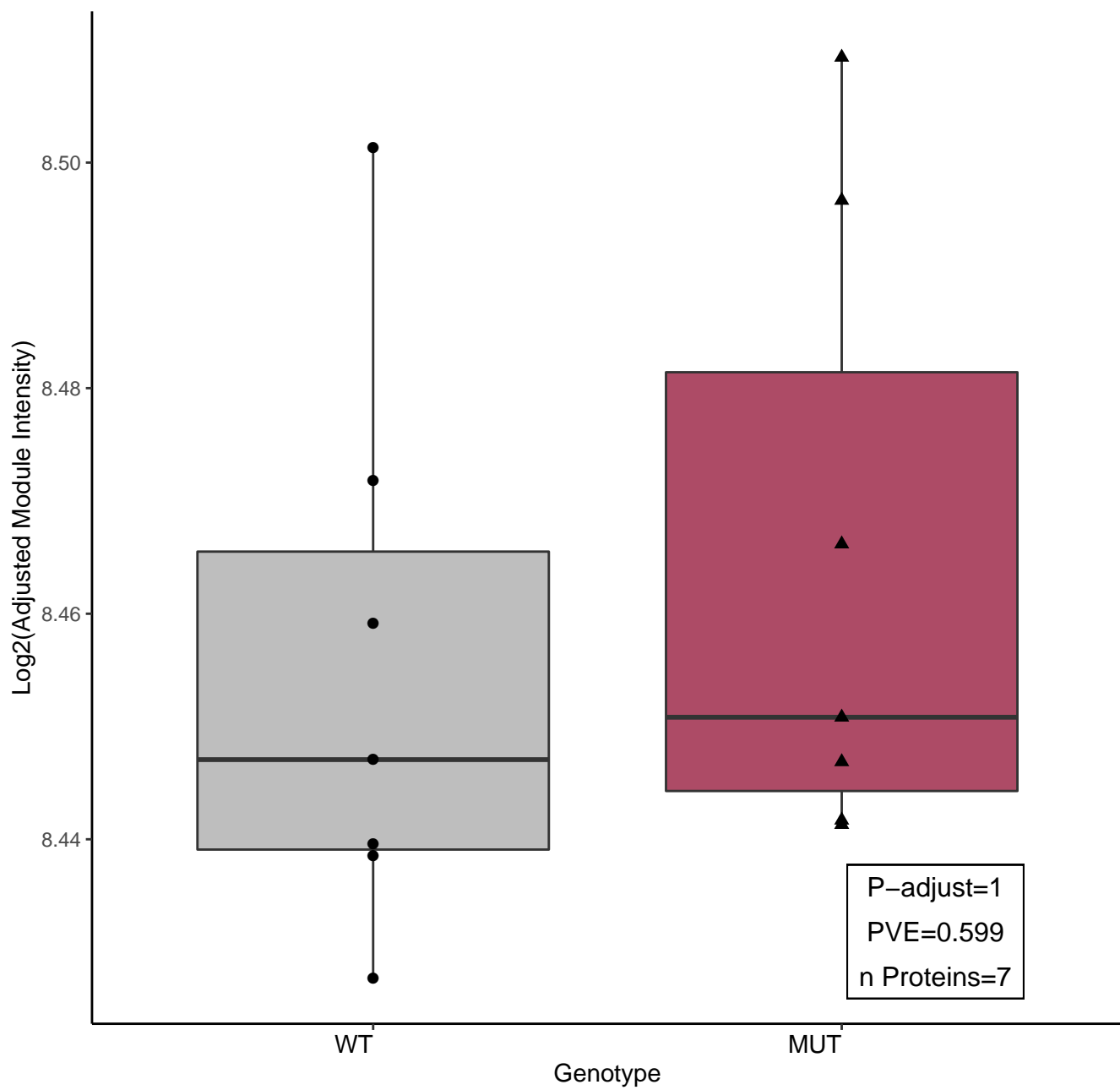
P-adjust=1

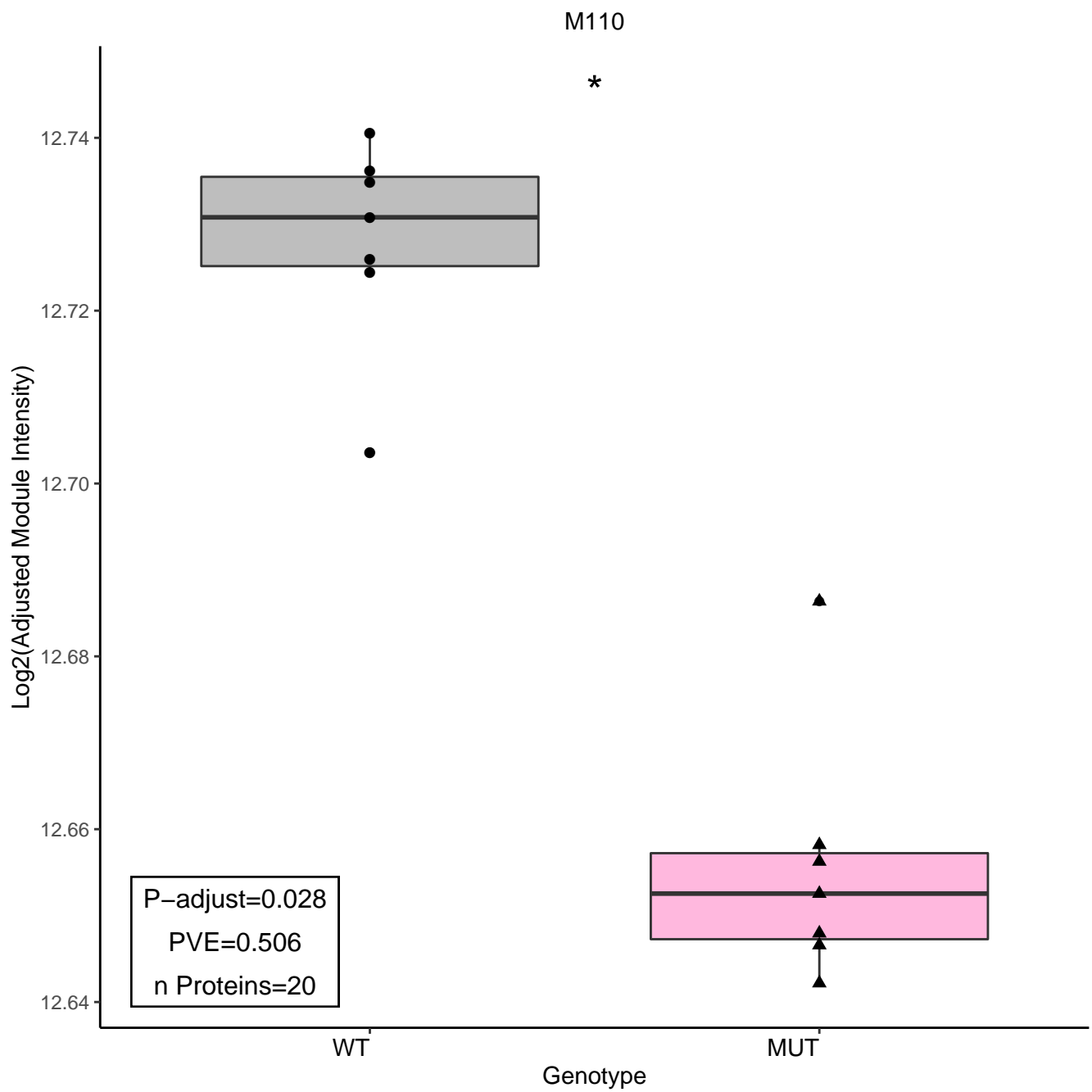
PVE=0.619

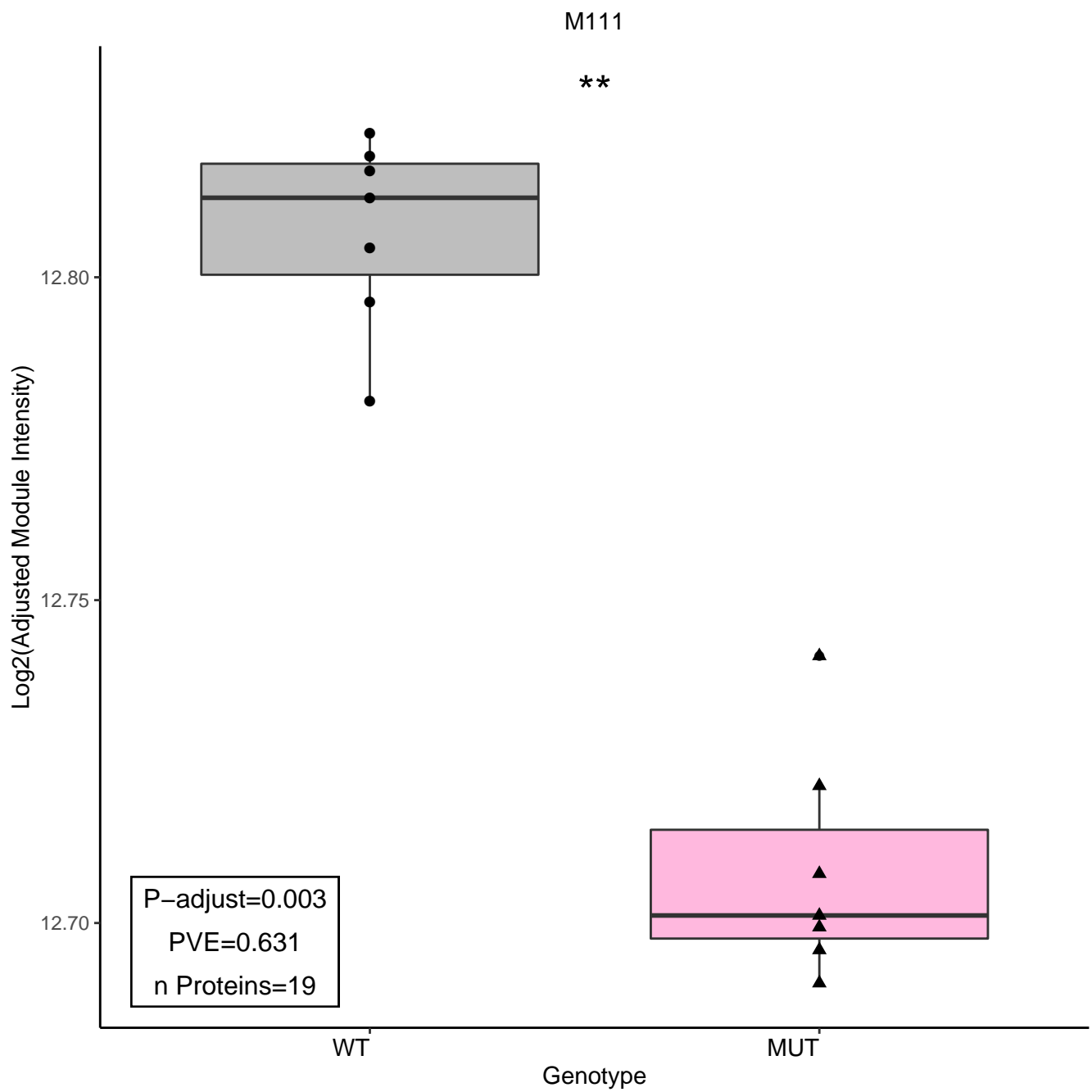
n Proteins=9



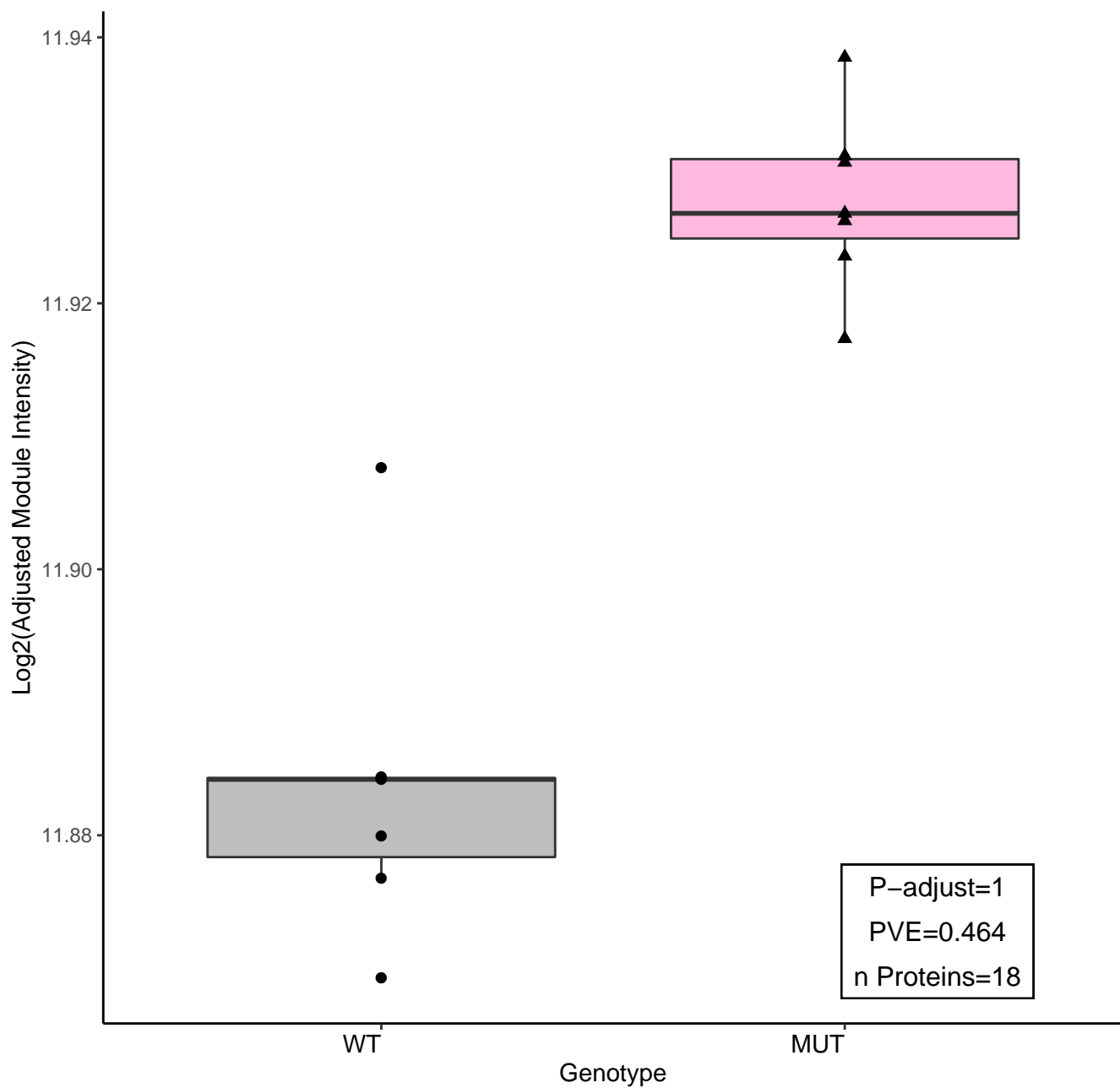
M109







M112



M113

Log2(Adjusted Module Intensity)

13.25

13.20

13.15

P-adjust=0.198

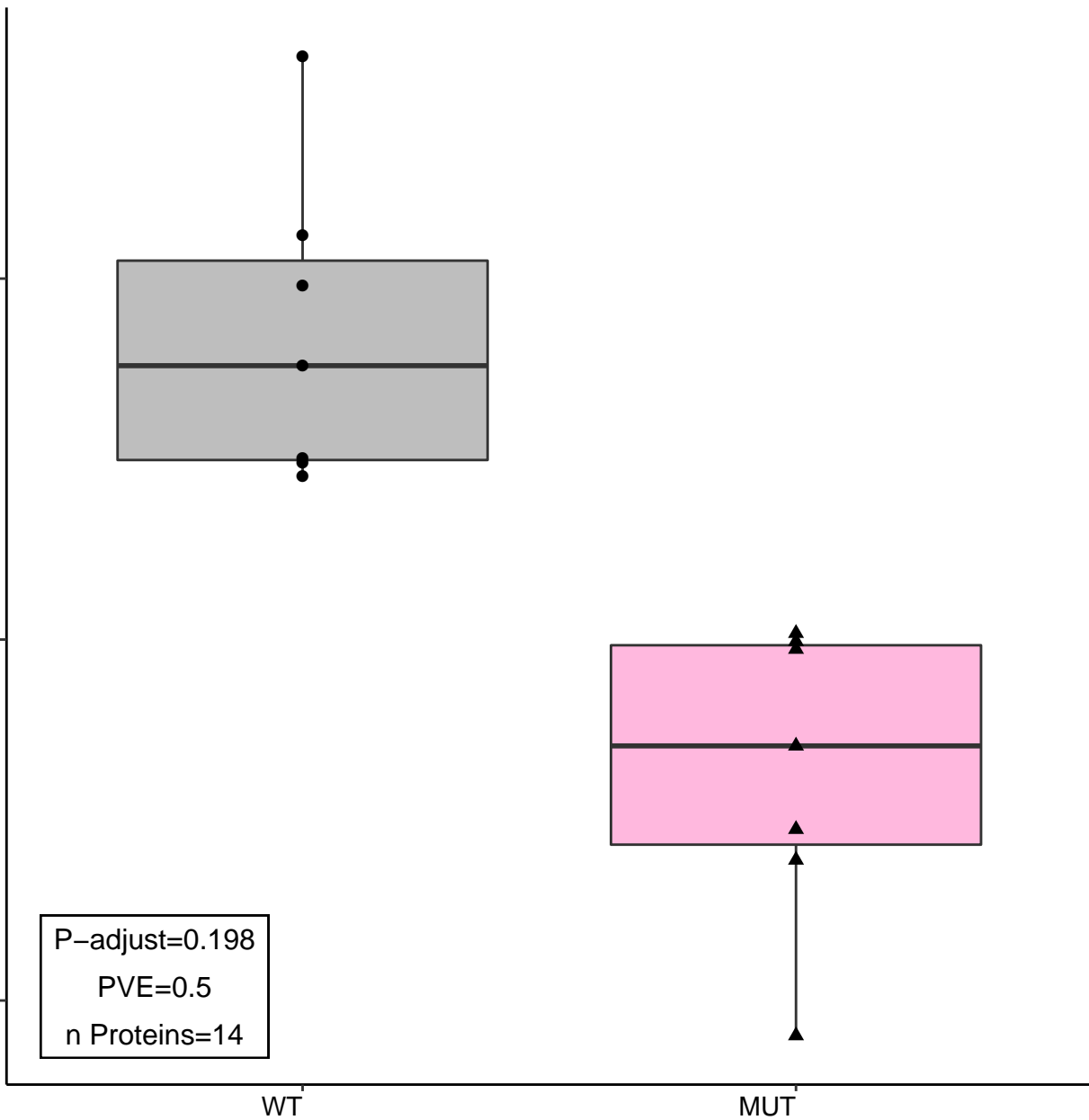
PVE=0.5

n Proteins=14

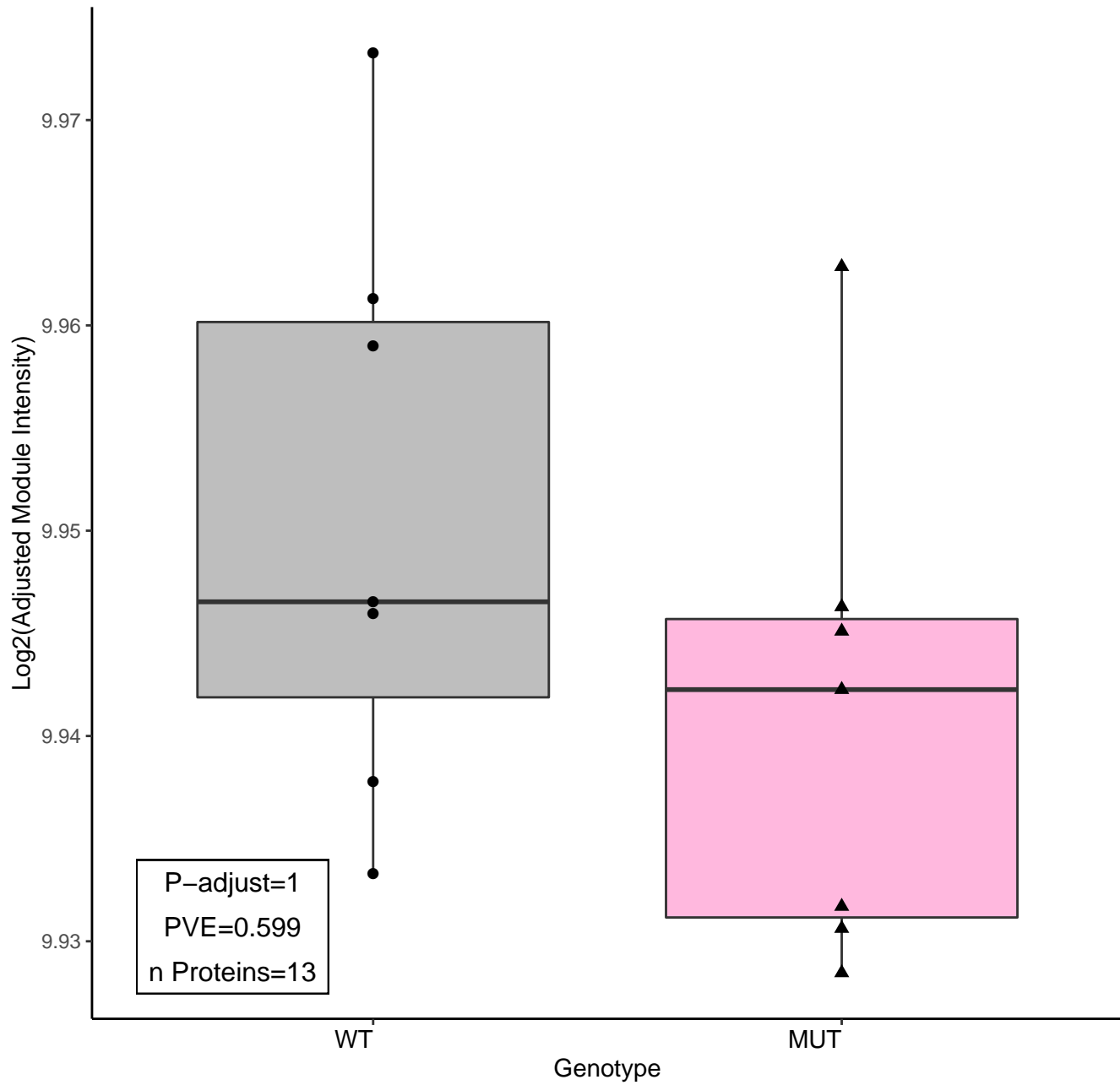
WT

Genotype

MUT



M114



M115

*

Log2(Adjusted Module Intensity)

13.85

13.80

P-adjust=0.018

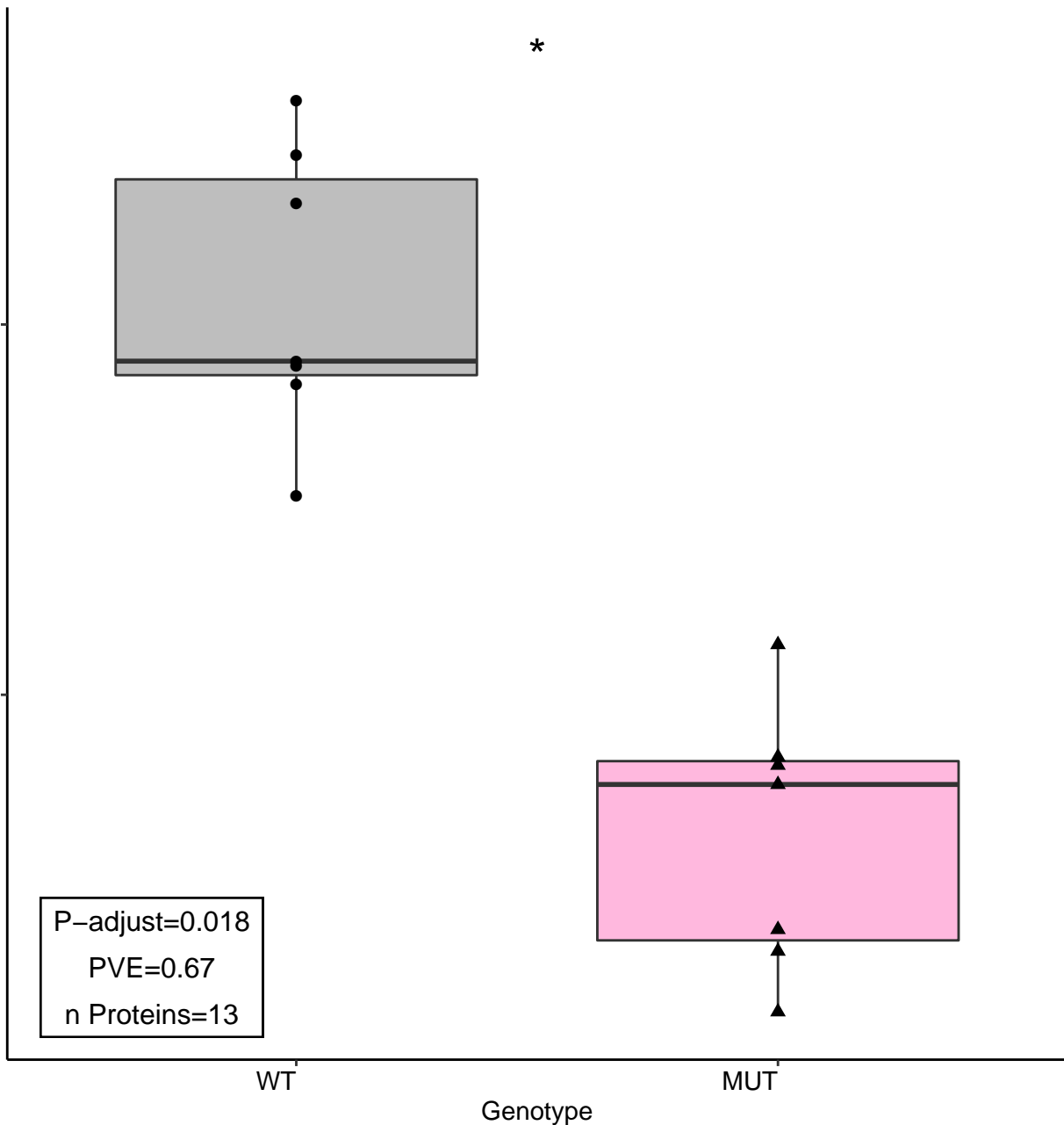
PVE=0.67

n Proteins=13

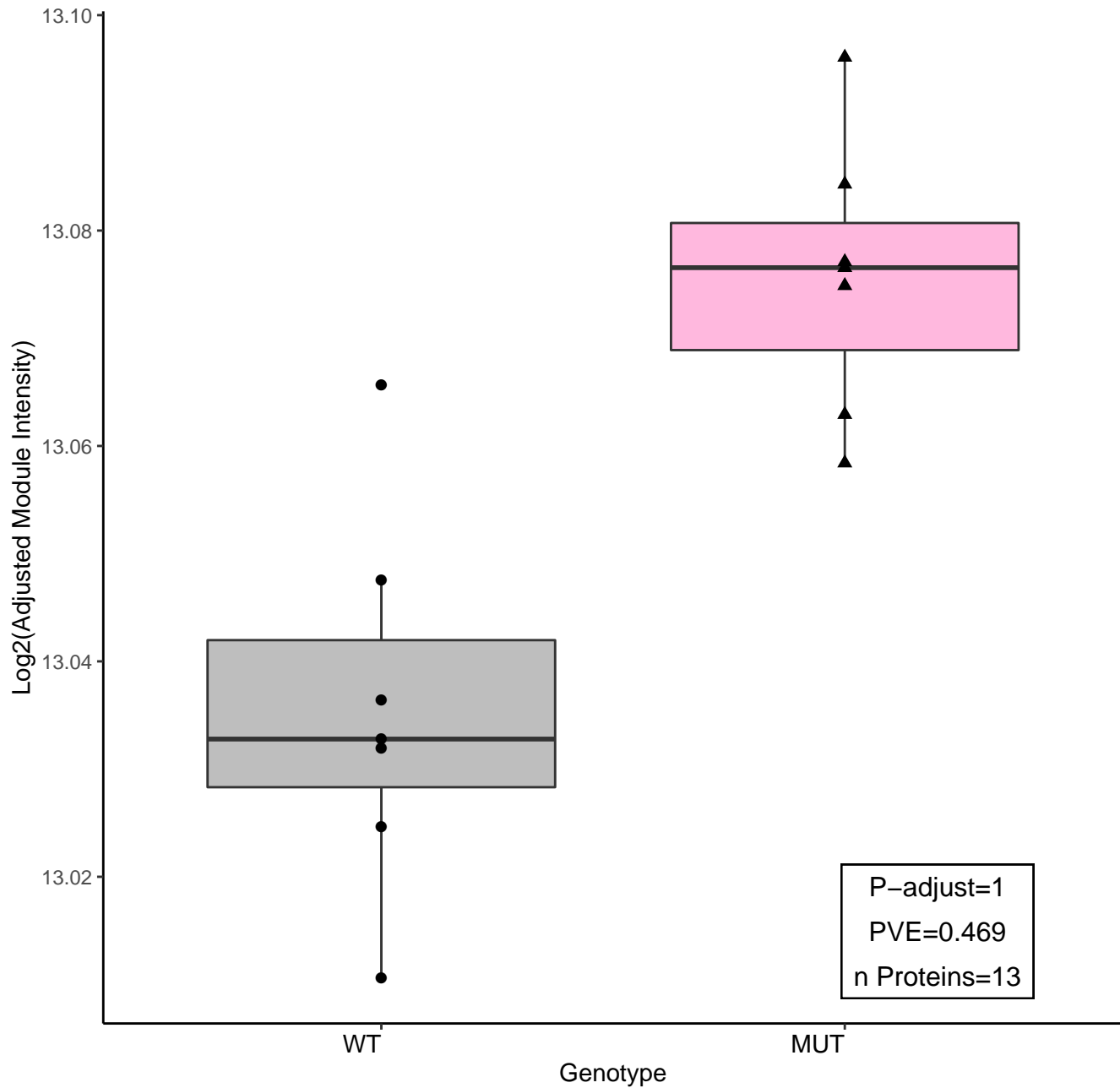
WT

Genotype

MUT



M116



M117

Log2(Adjusted Module Intensity)

11.92

11.90

11.88

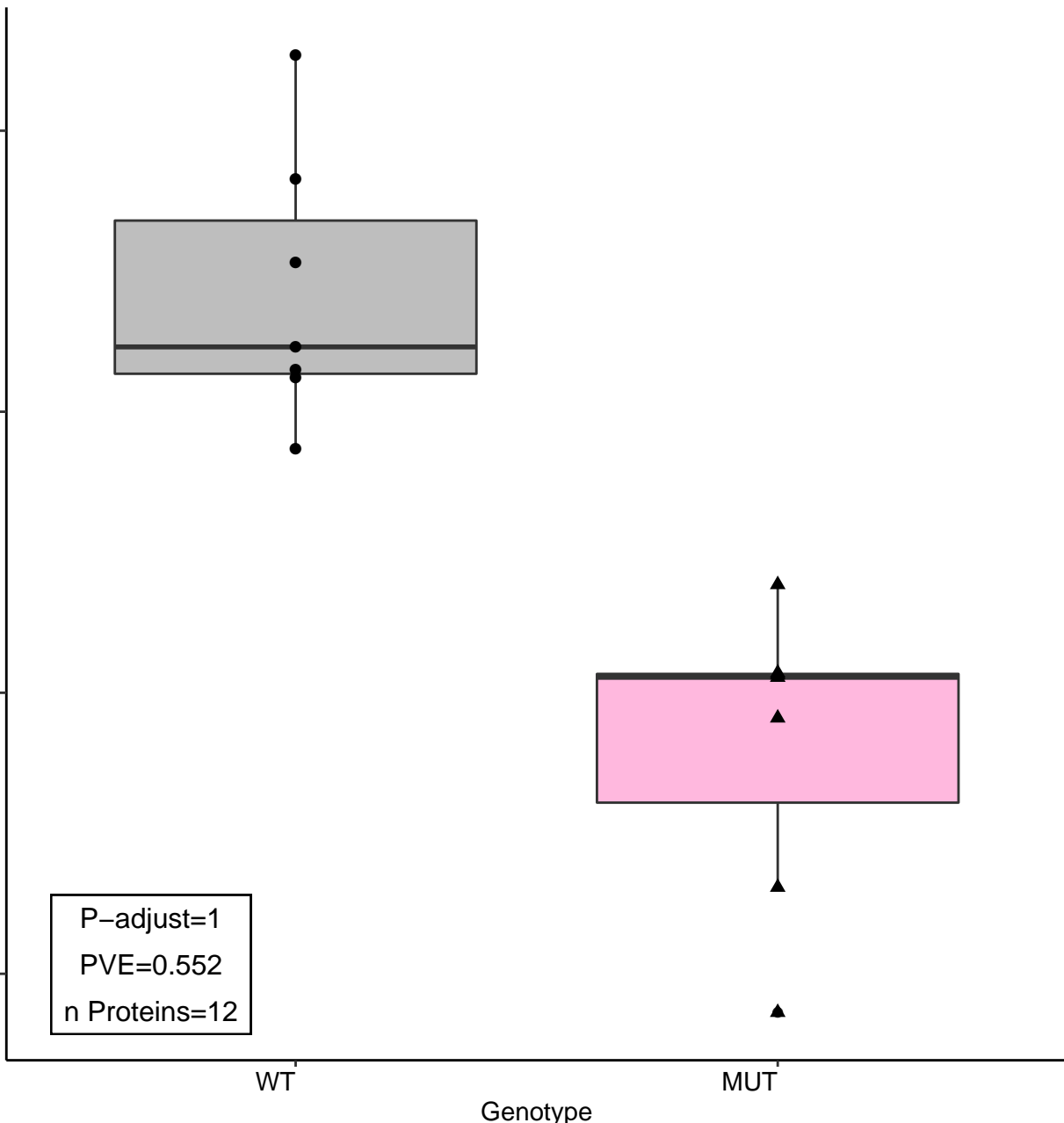
11.86

P-adjust=1
PVE=0.552
n Proteins=12

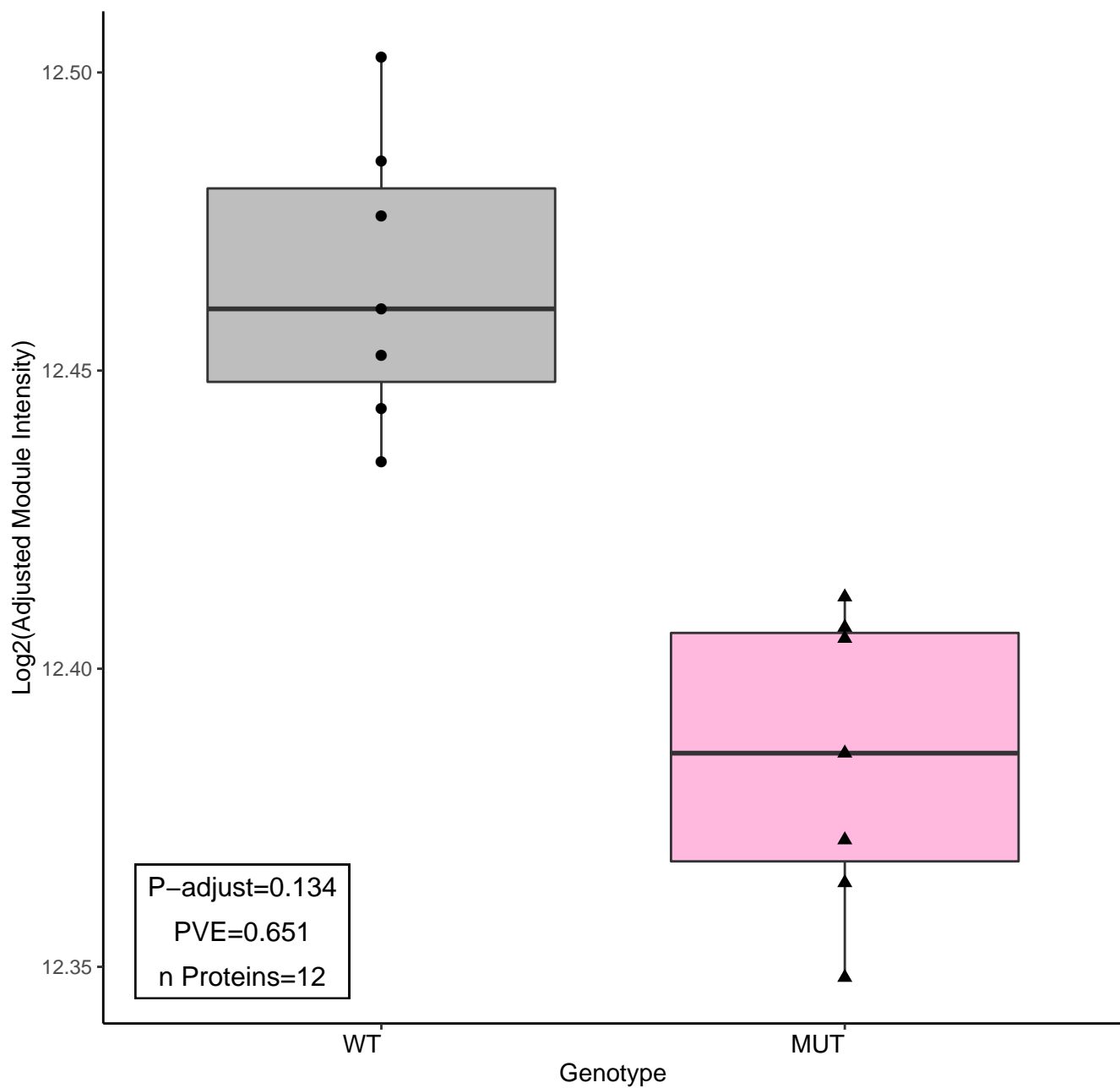
WT

Genotype

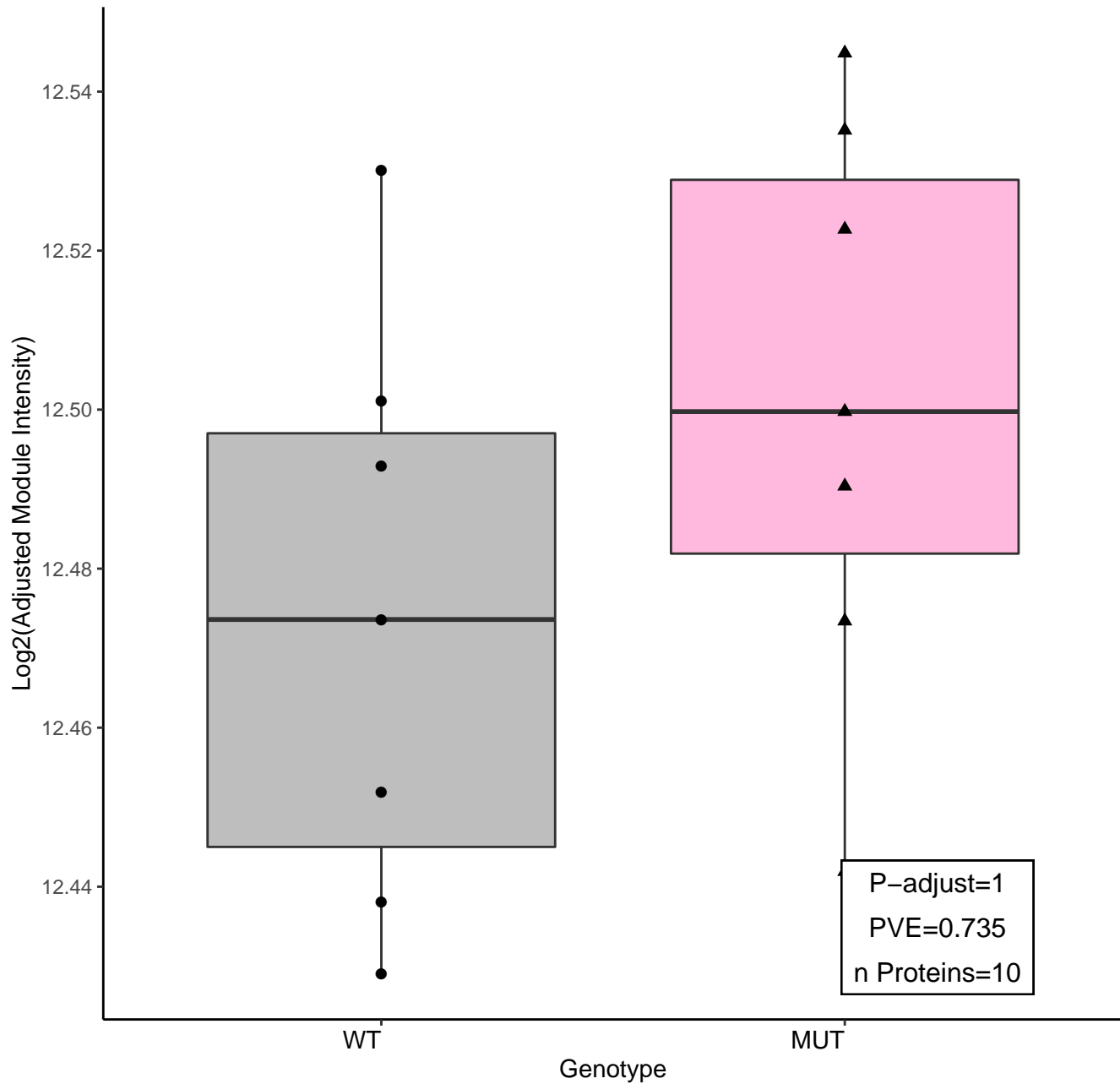
MUT



M118



M119



M120

Log2(Adjusted Module Intensity)

11.08

11.06

11.04

WT

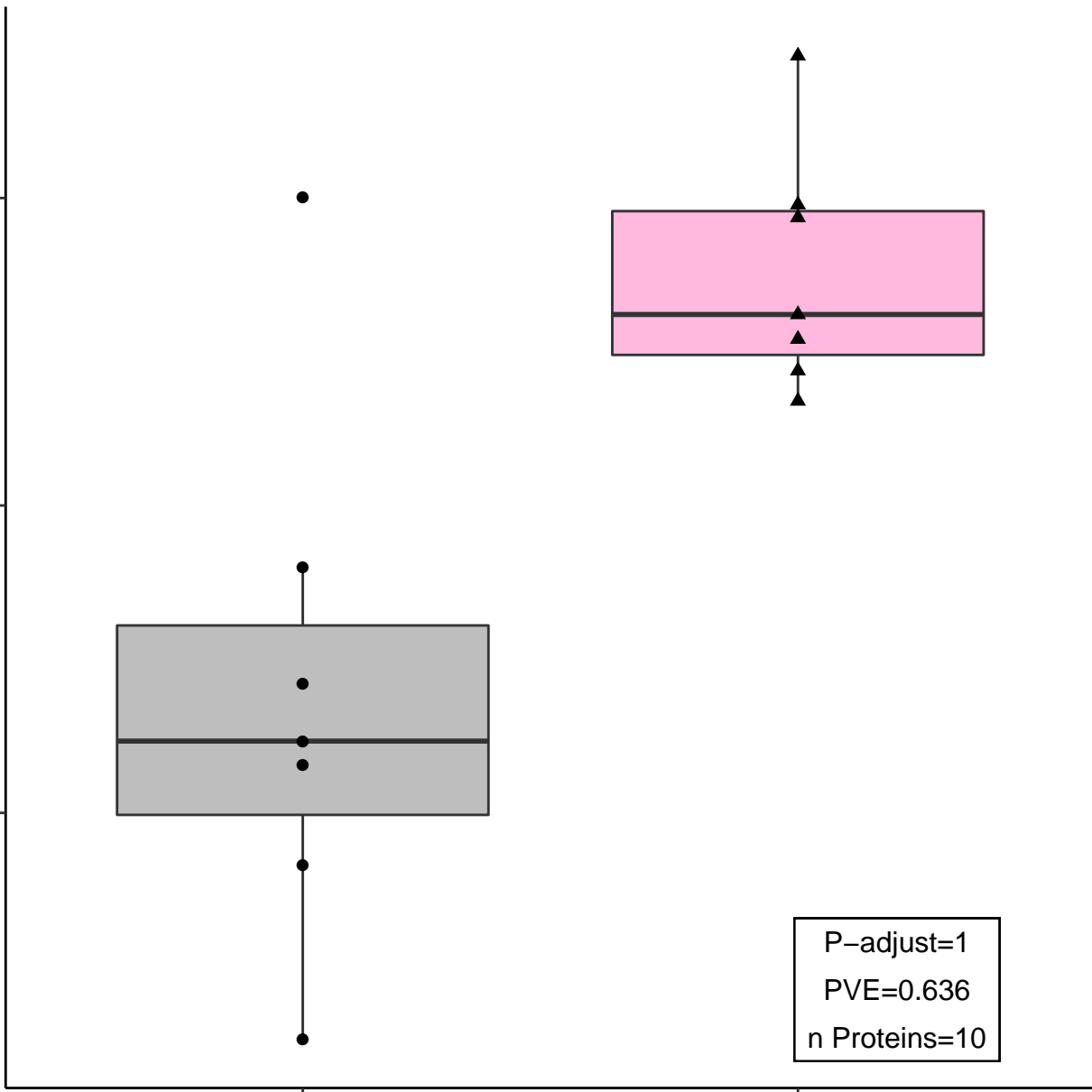
MUT

Genotype

P-adjust=1

PVE=0.636

n Proteins=10



M121

Log2(Adjusted Module Intensity)

10.44

10.42

10.40

10.38

P-adjust=1

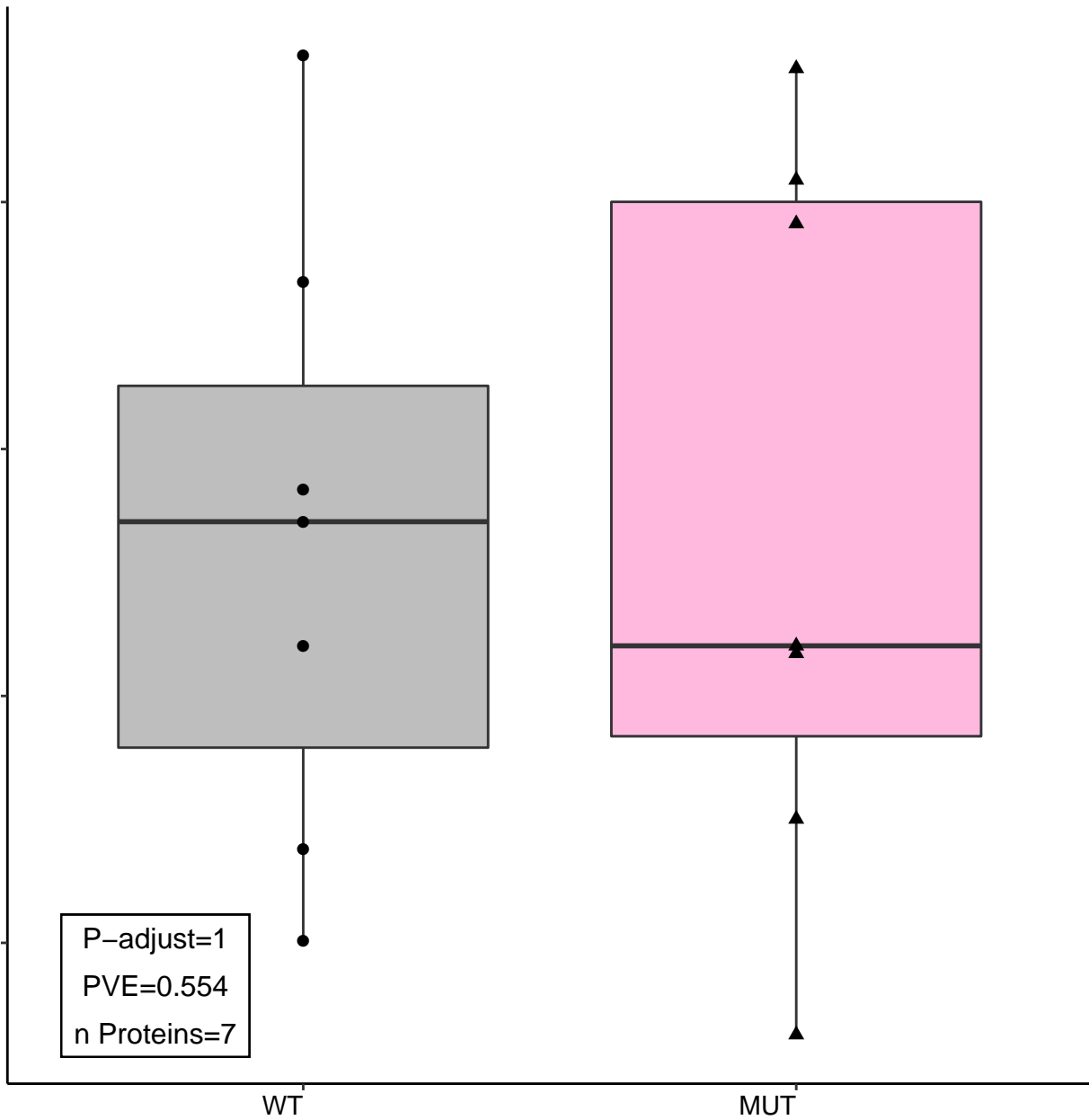
PVE=0.554

n Proteins=7

WT

Genotype

MUT



M122

Log2(Adjusted Module Intensity)

12.15

12.10

12.05

P-adjust=0.078

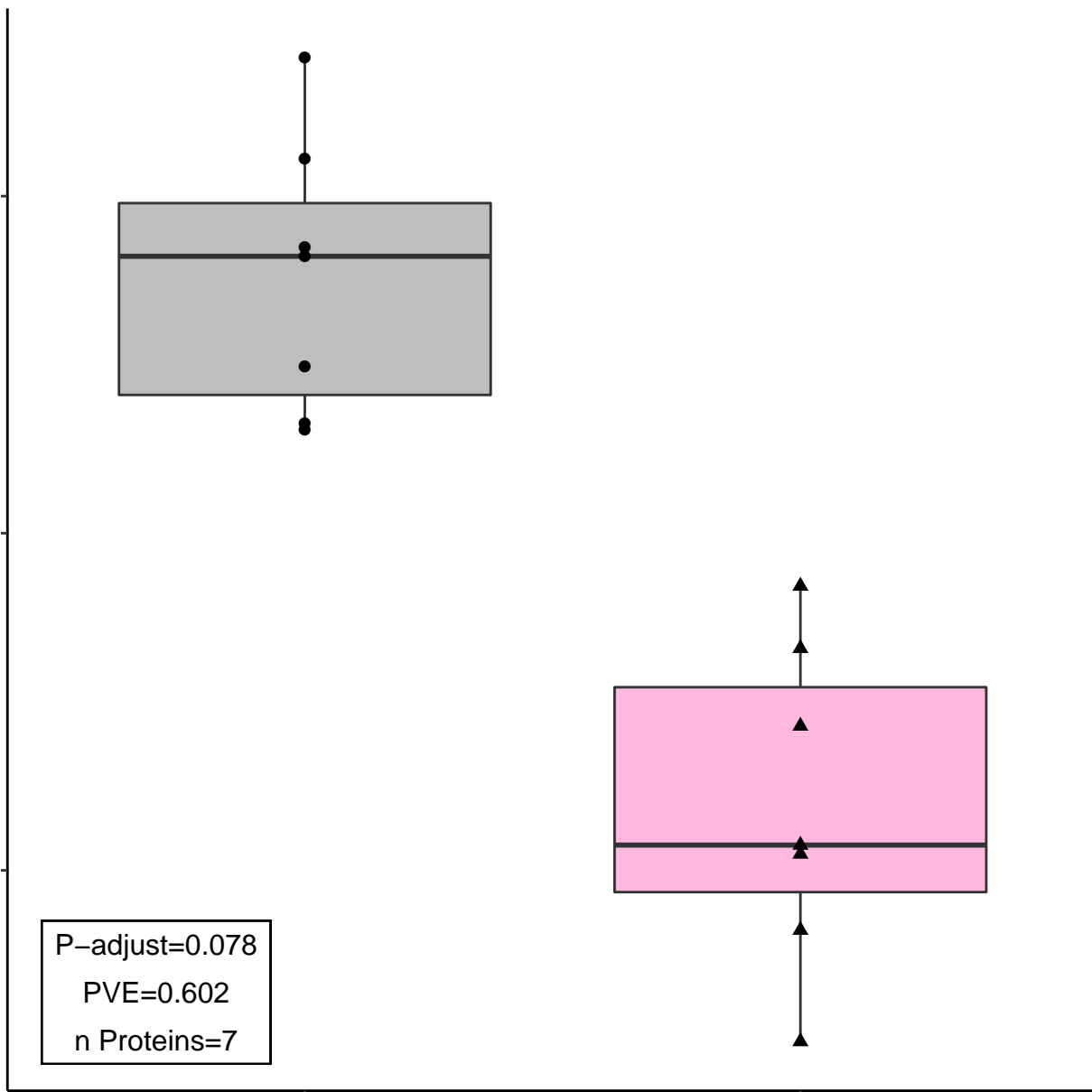
PVE=0.602

n Proteins=7

WT

Genotype

MUT



M123

Log2(Adjusted Module Intensity)

15.00

14.98

14.96

WT

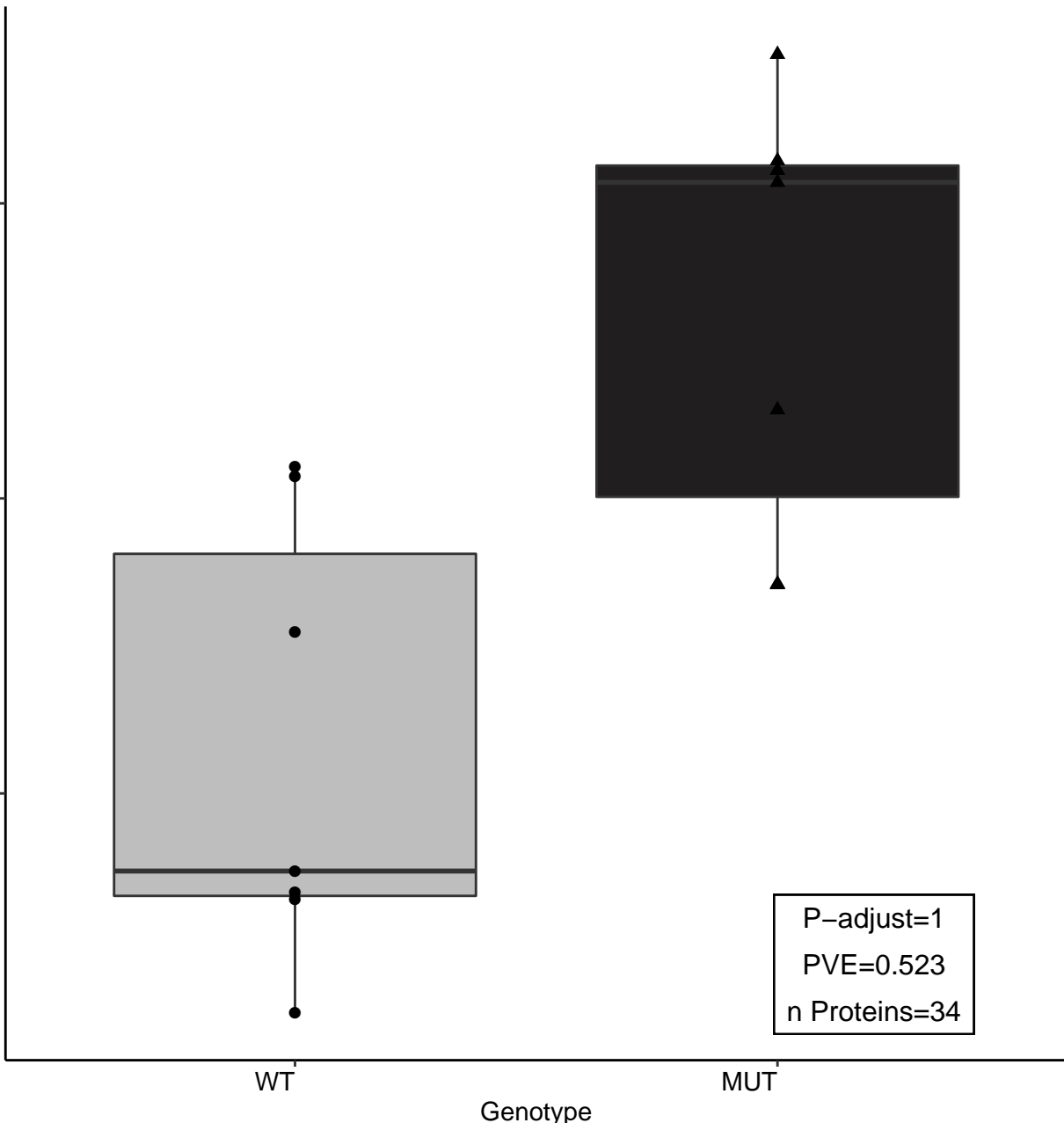
MUT

Genotype

P-adjust=1

PVE=0.523

n Proteins=34



M124

Log2(Adjusted Module Intensity)

12.76

12.75

12.74

WT

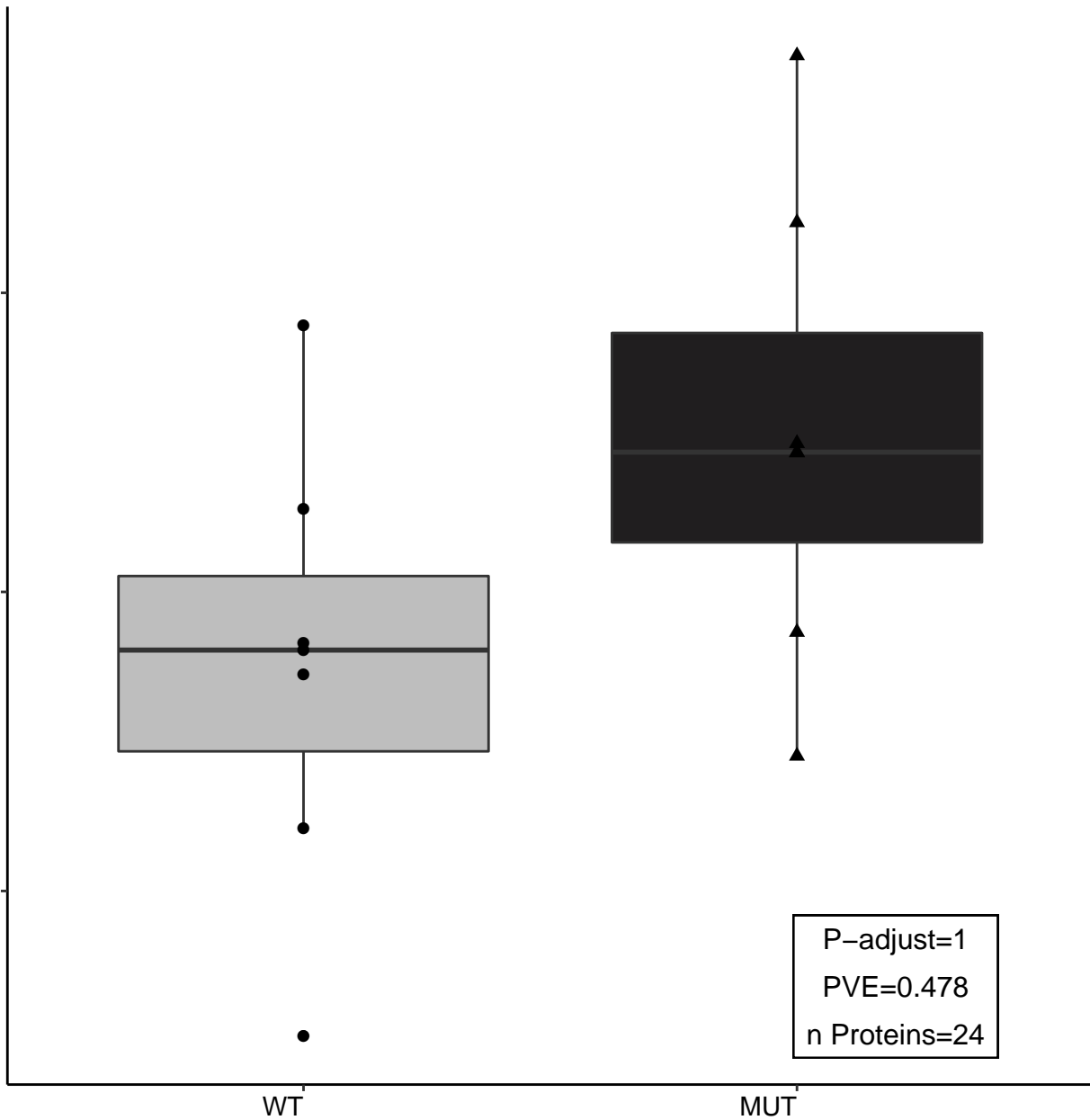
Genotype

MUT

P-adjust=1

PVE=0.478

n Proteins=24



M125

Log2(Adjusted Module Intensity)

11.48

11.46

11.44

11.42

WT

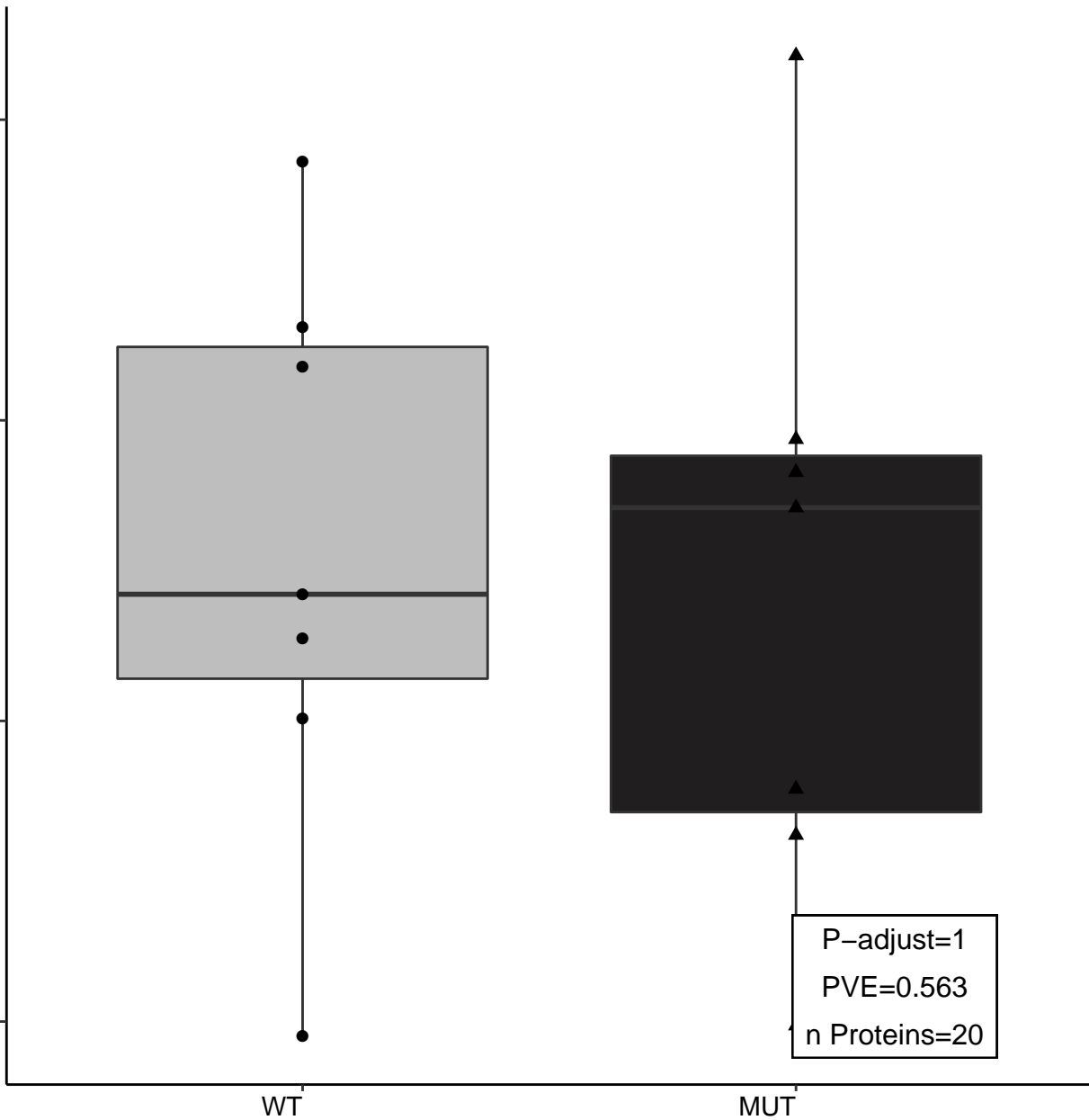
Genotype

MUT

P-adjust=1

PVE=0.563

n Proteins=20



M126

Log2(Adjusted Module Intensity)

11.23

11.22

11.21

11.20

11.19

WT

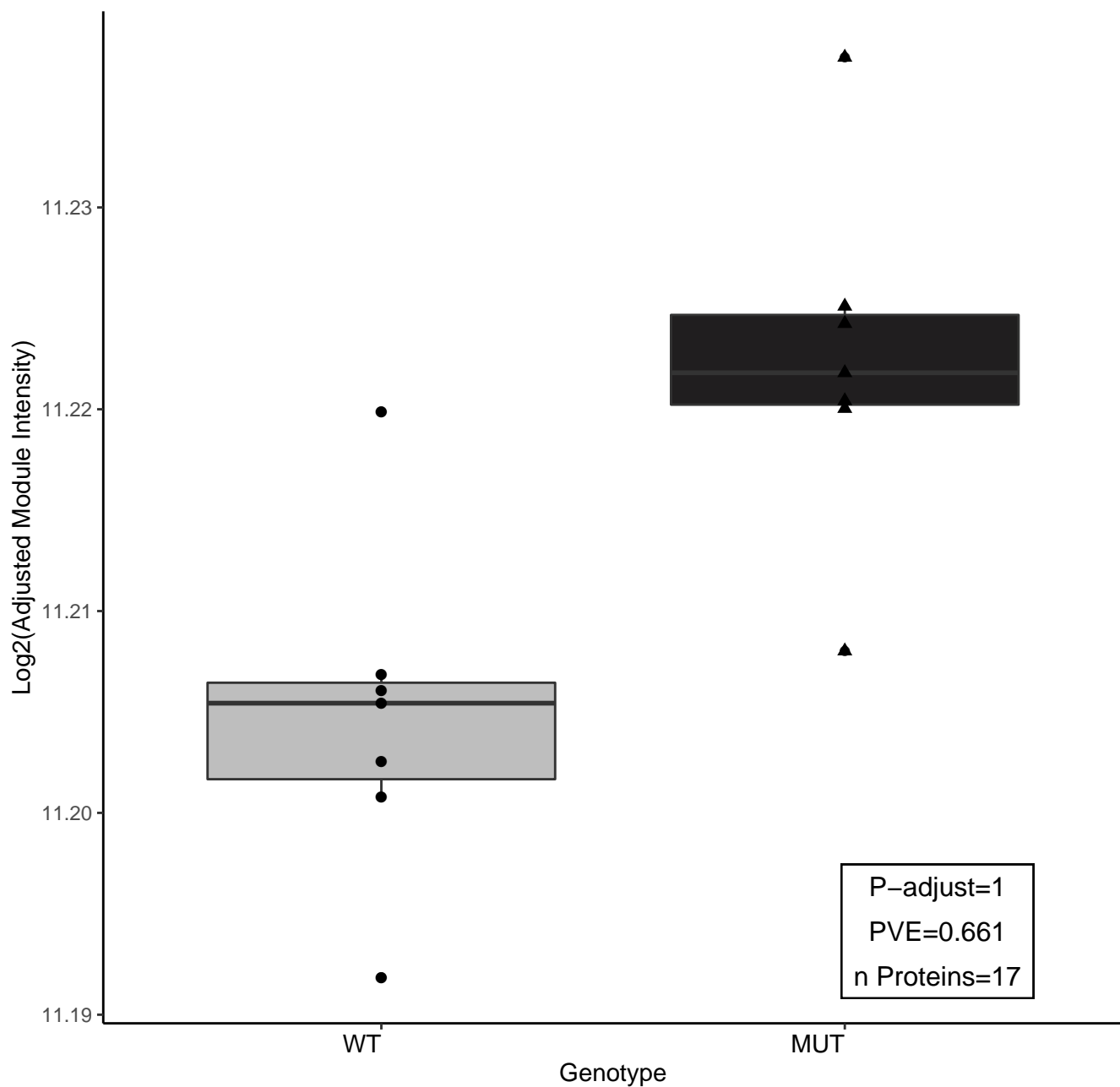
MUT

Genotype

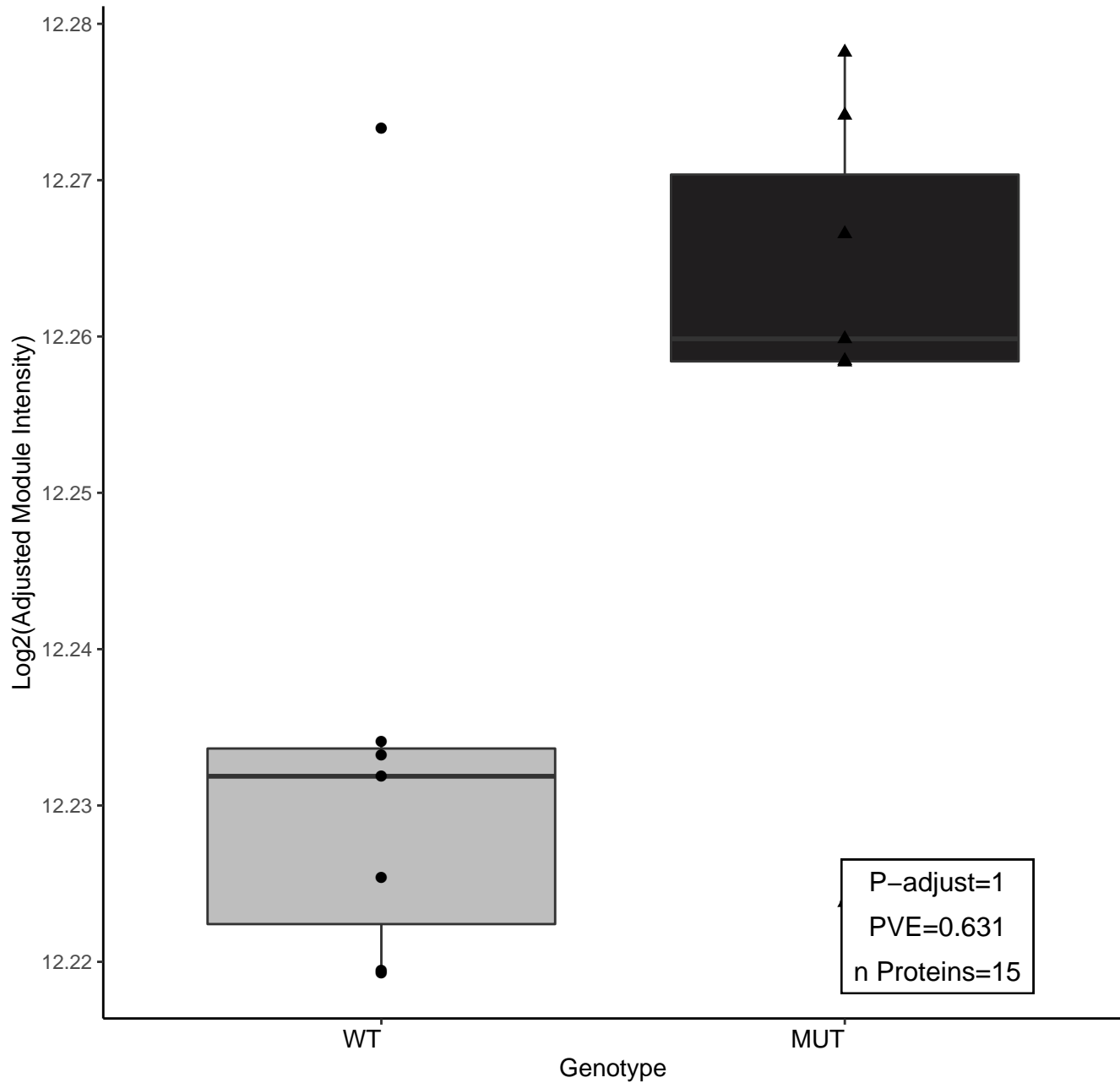
P-adjust=1

PVE=0.661

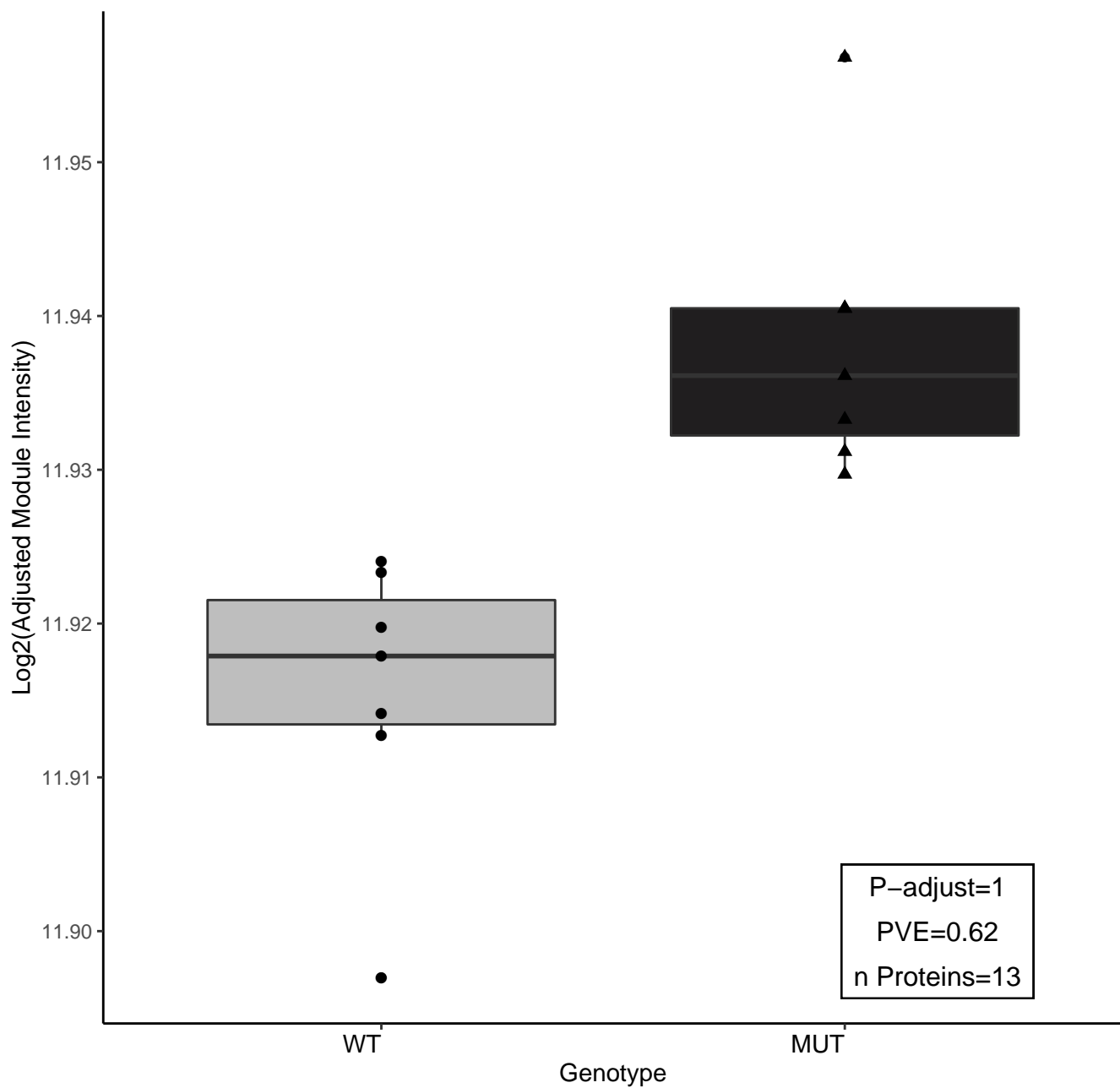
n Proteins=17



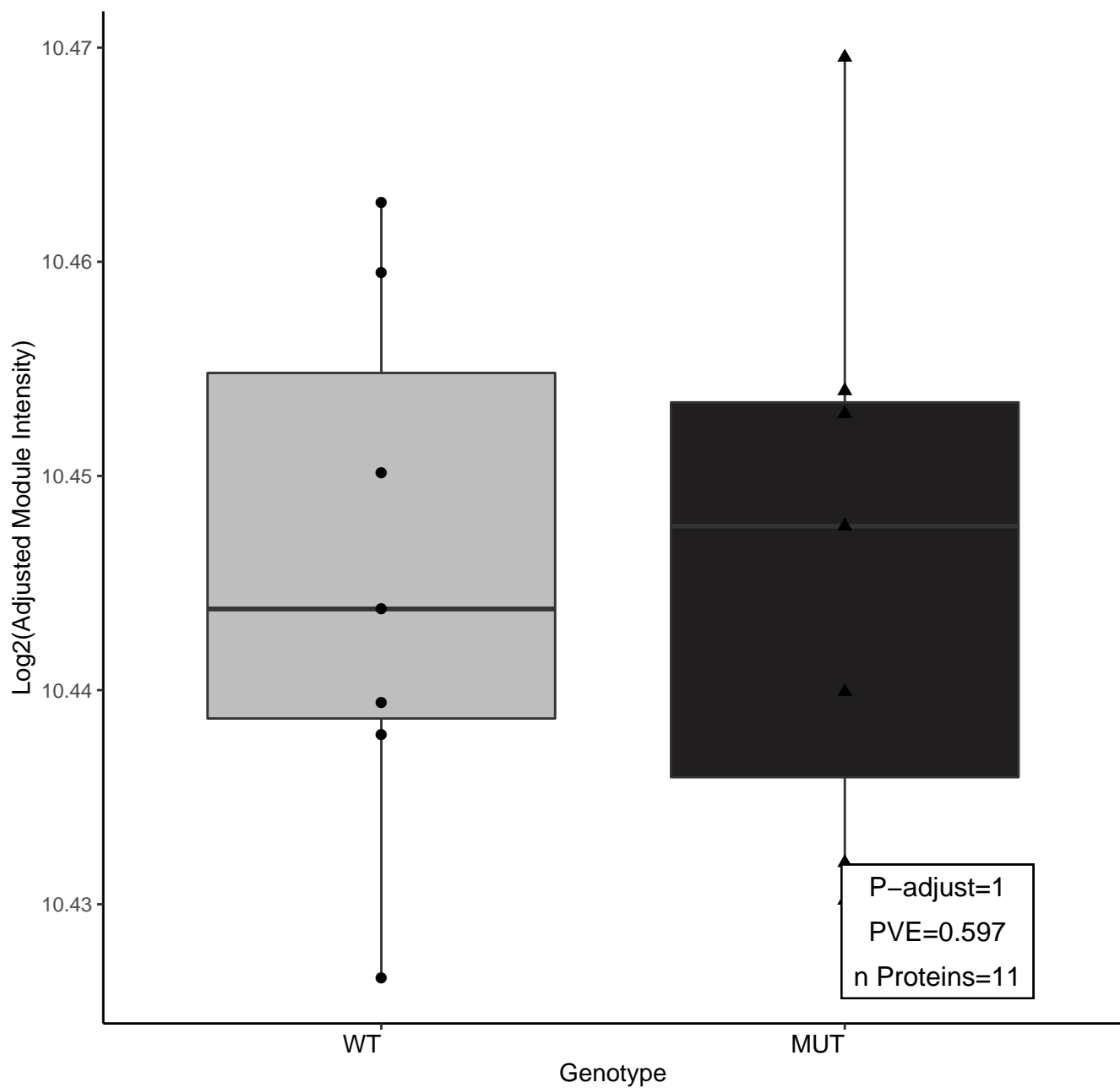
M127



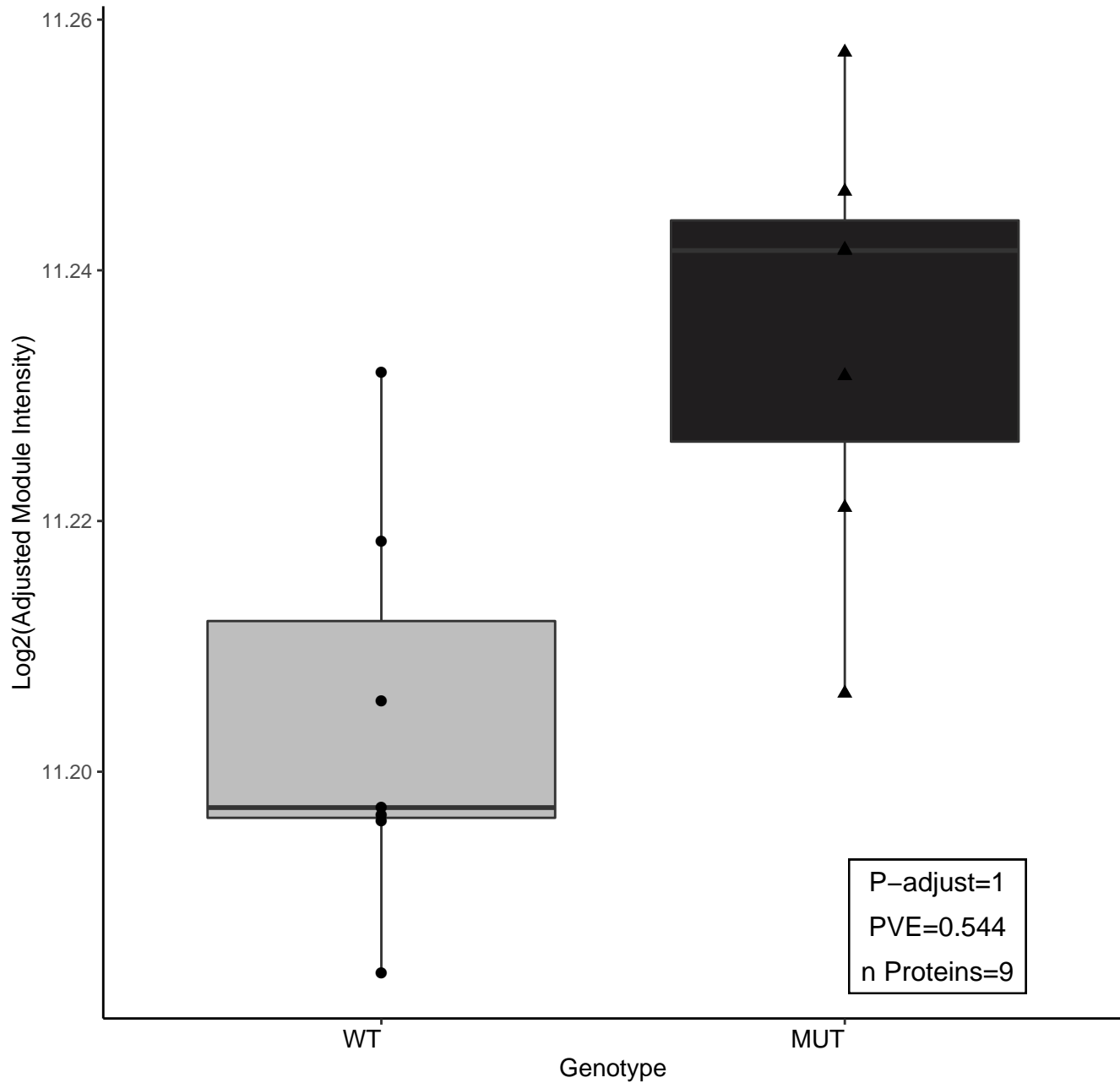
M128



M129



M130



M131

Log2(Adjusted Module Intensity)

12.70

12.68

12.66

12.64

12.62

12.60

WT

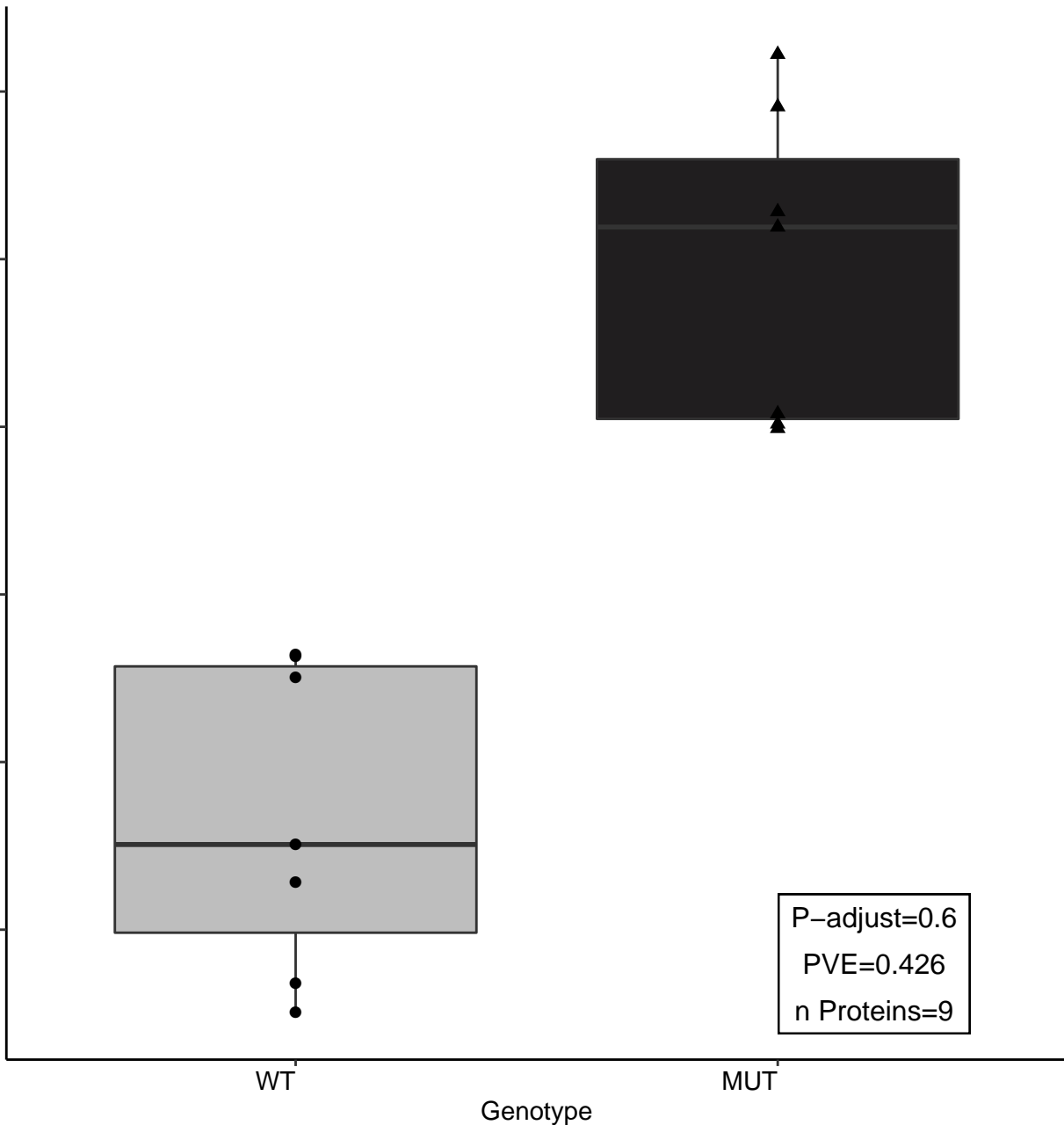
MUT

Genotype

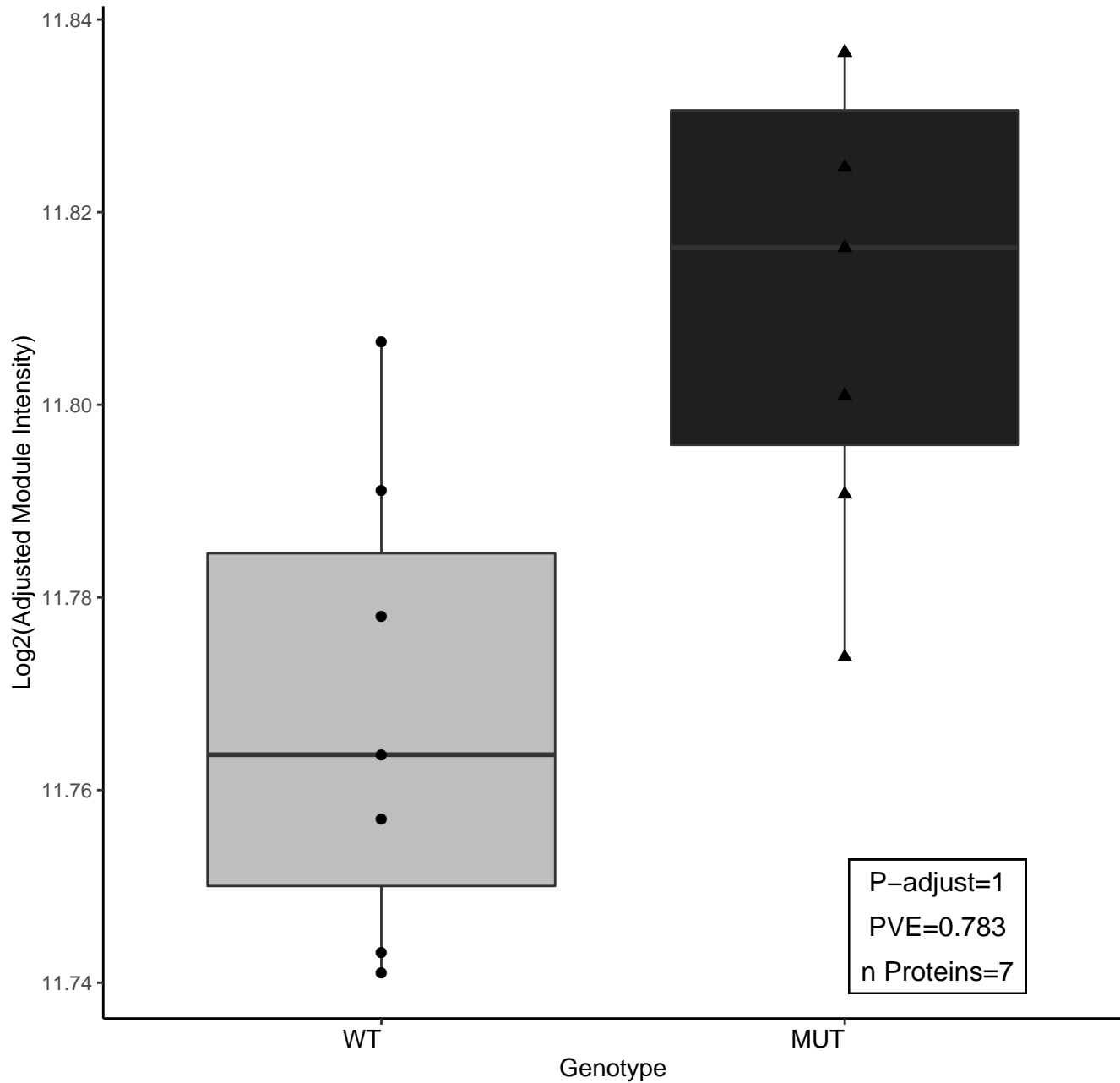
P-adjust=0.6

PVE=0.426

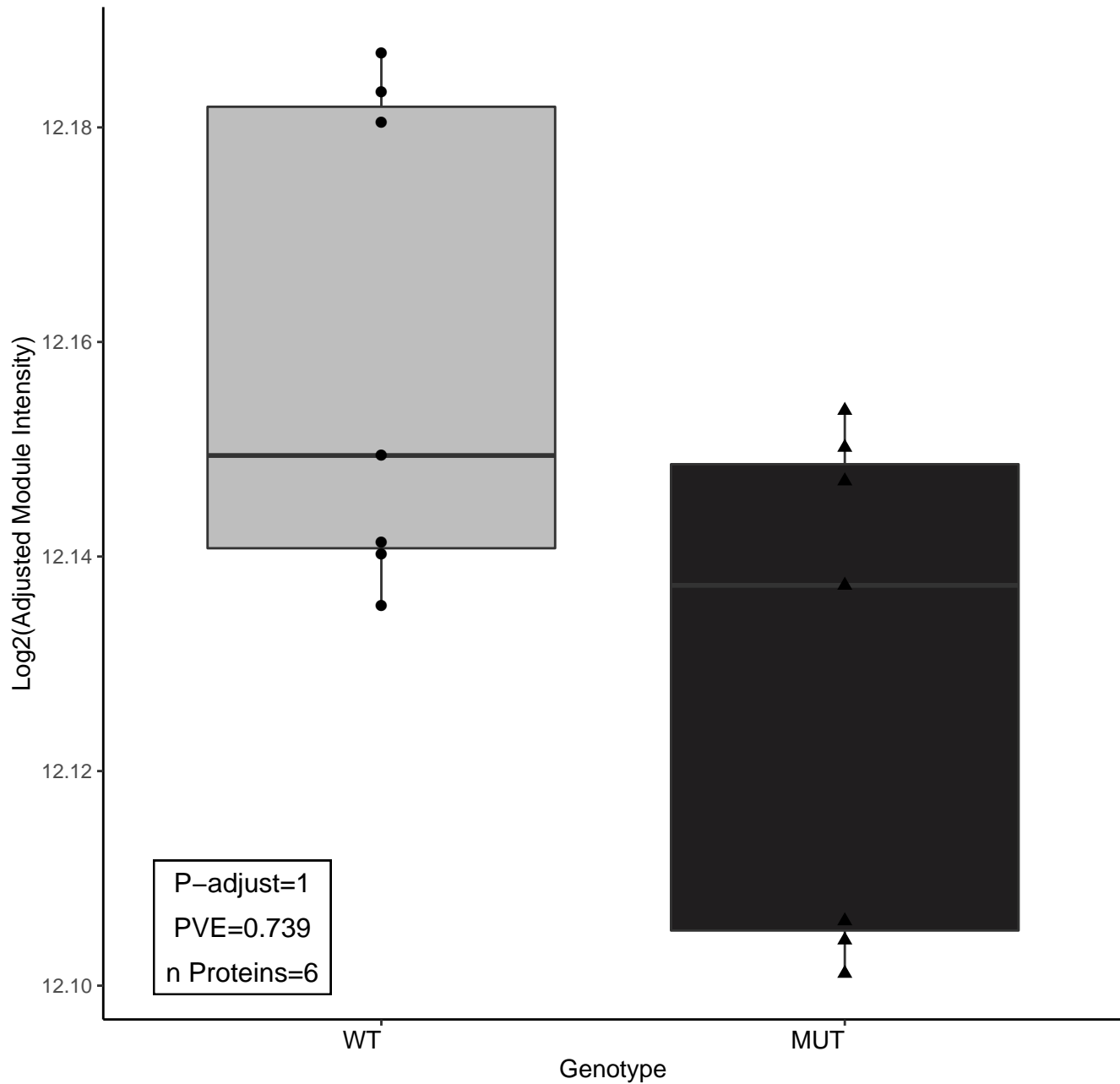
n Proteins=9

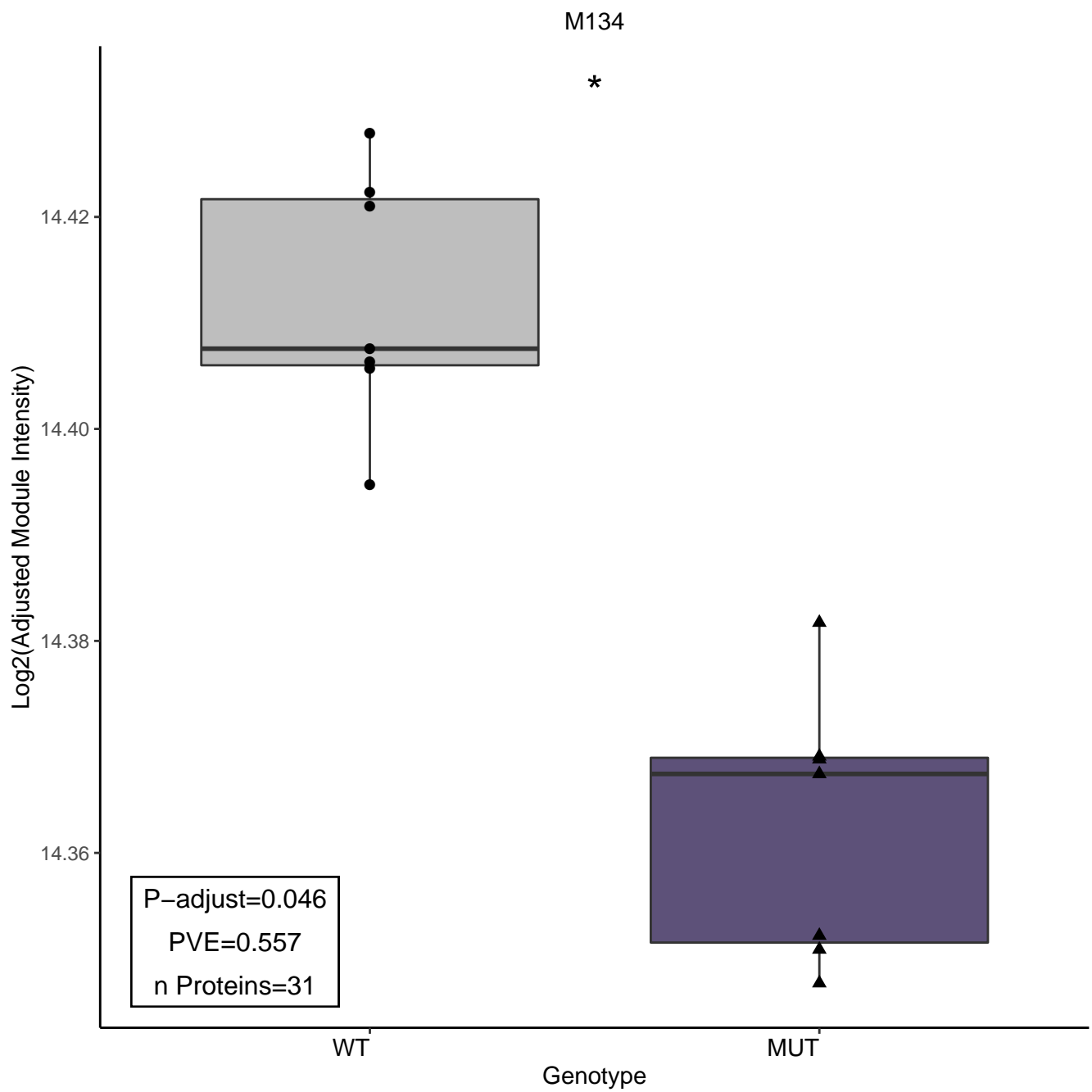


M132

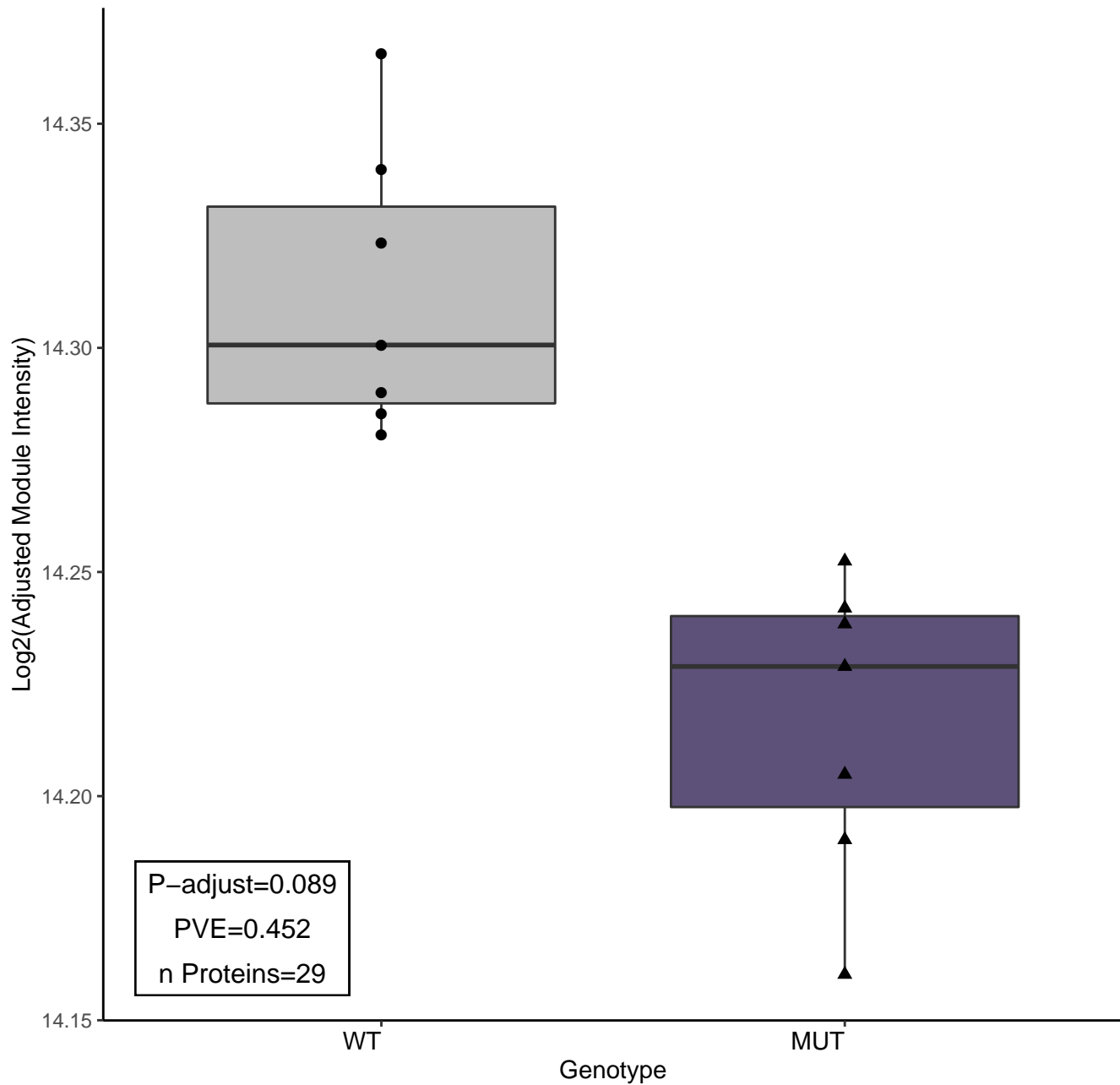


M133





M135



M136

Log2(Adjusted Module Intensity)

13.70

13.65

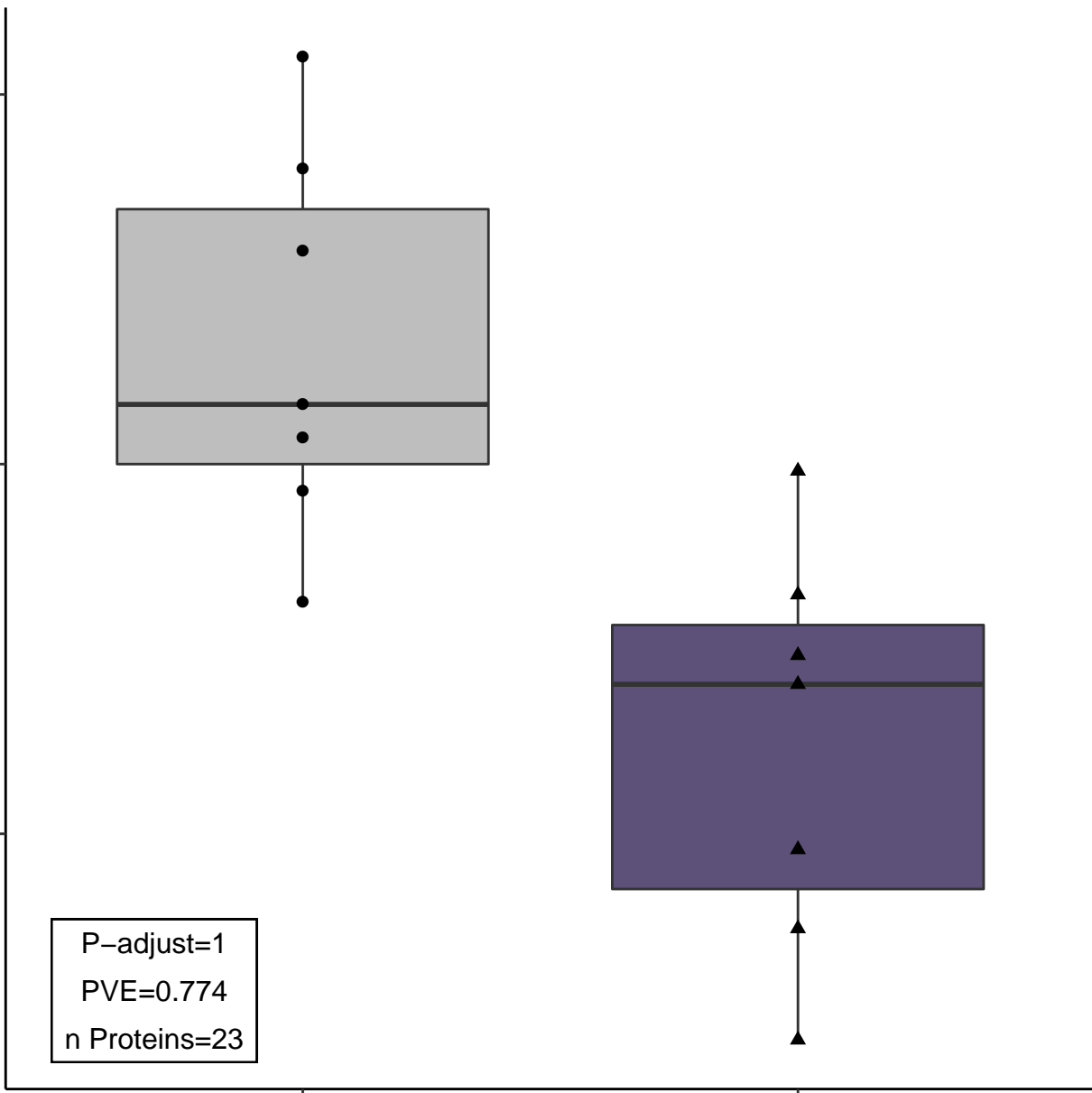
13.60

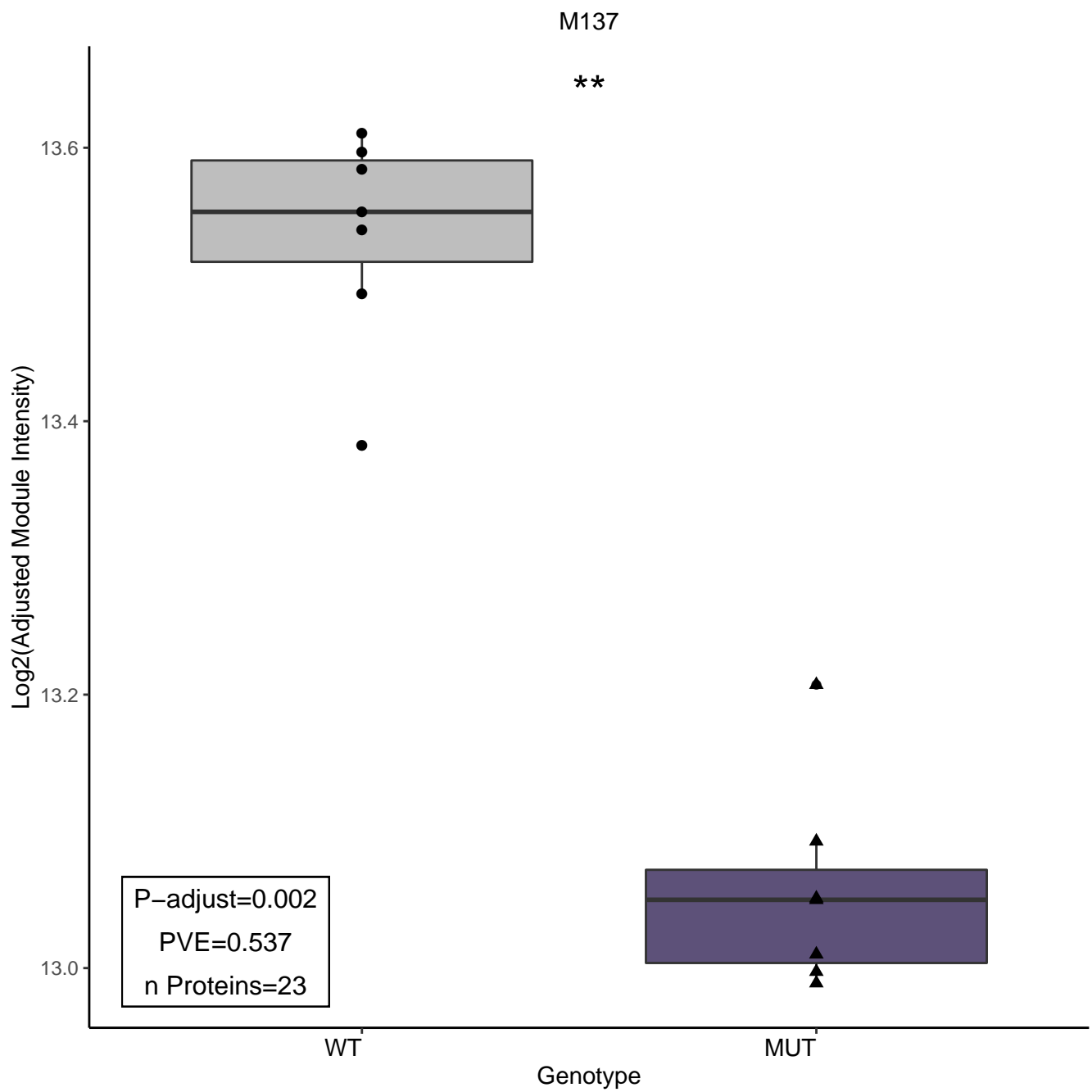
P-adjust=1
PVE=0.774
n Proteins=23

WT

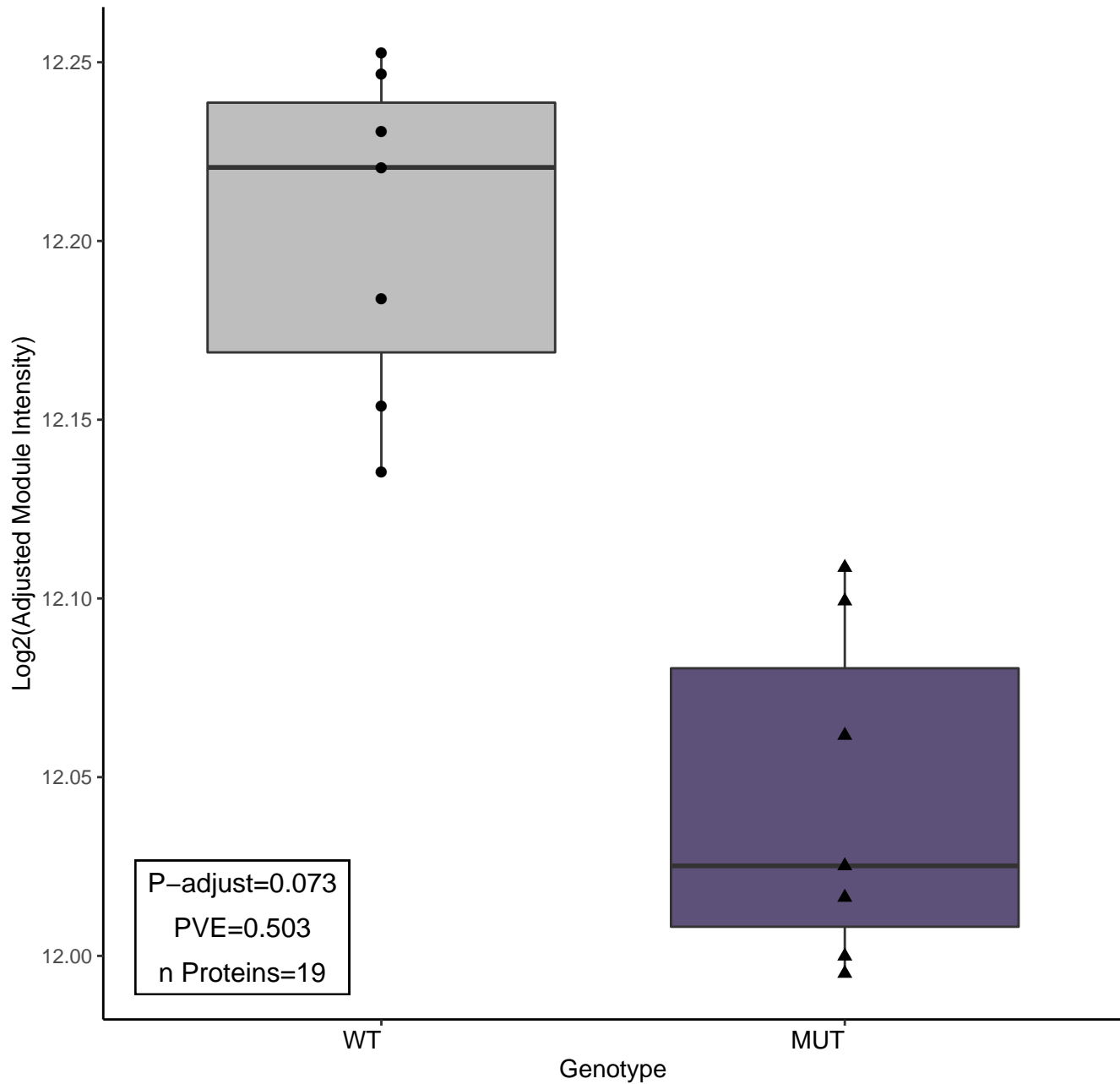
Genotype

MUT

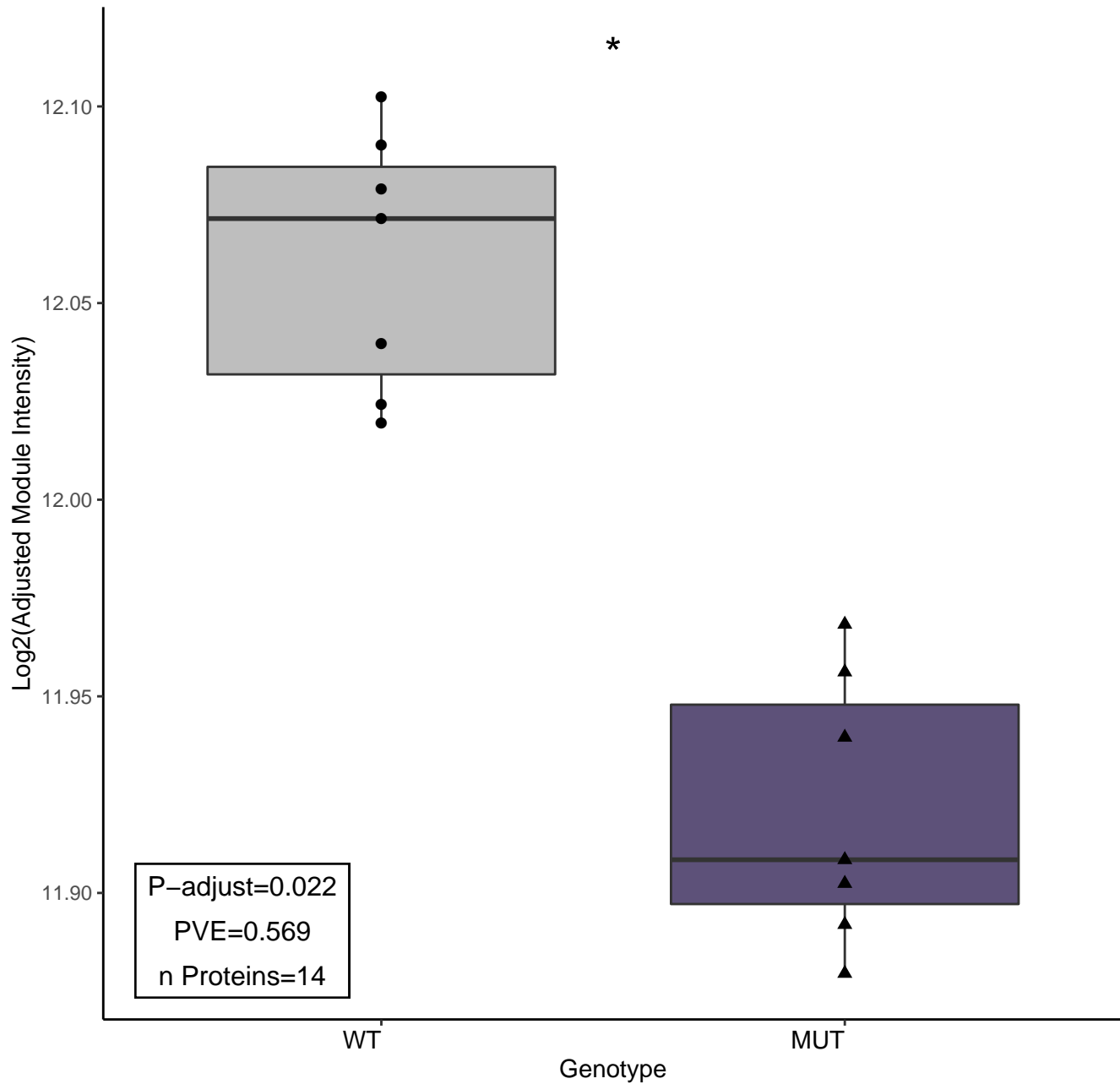




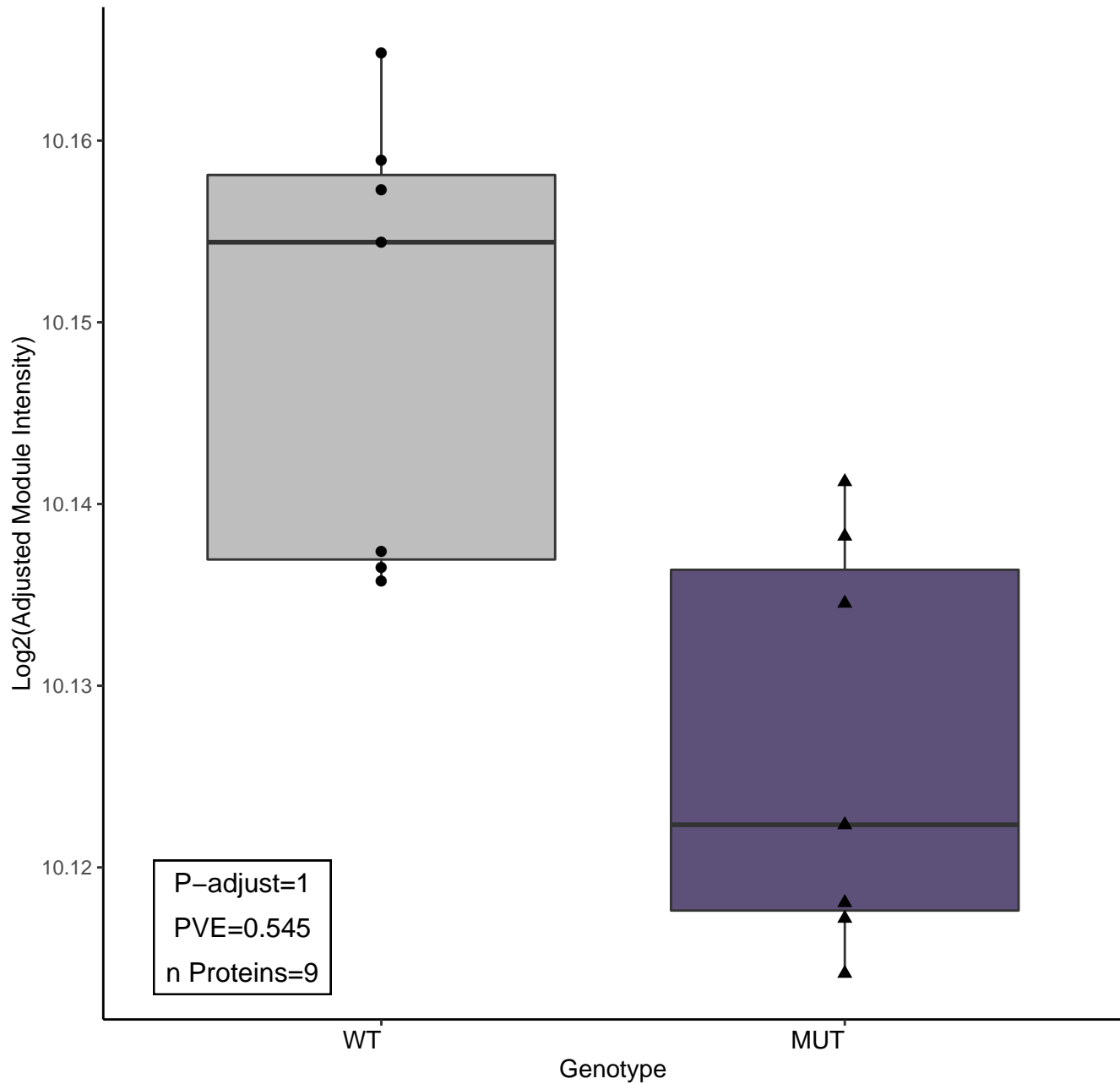
M138



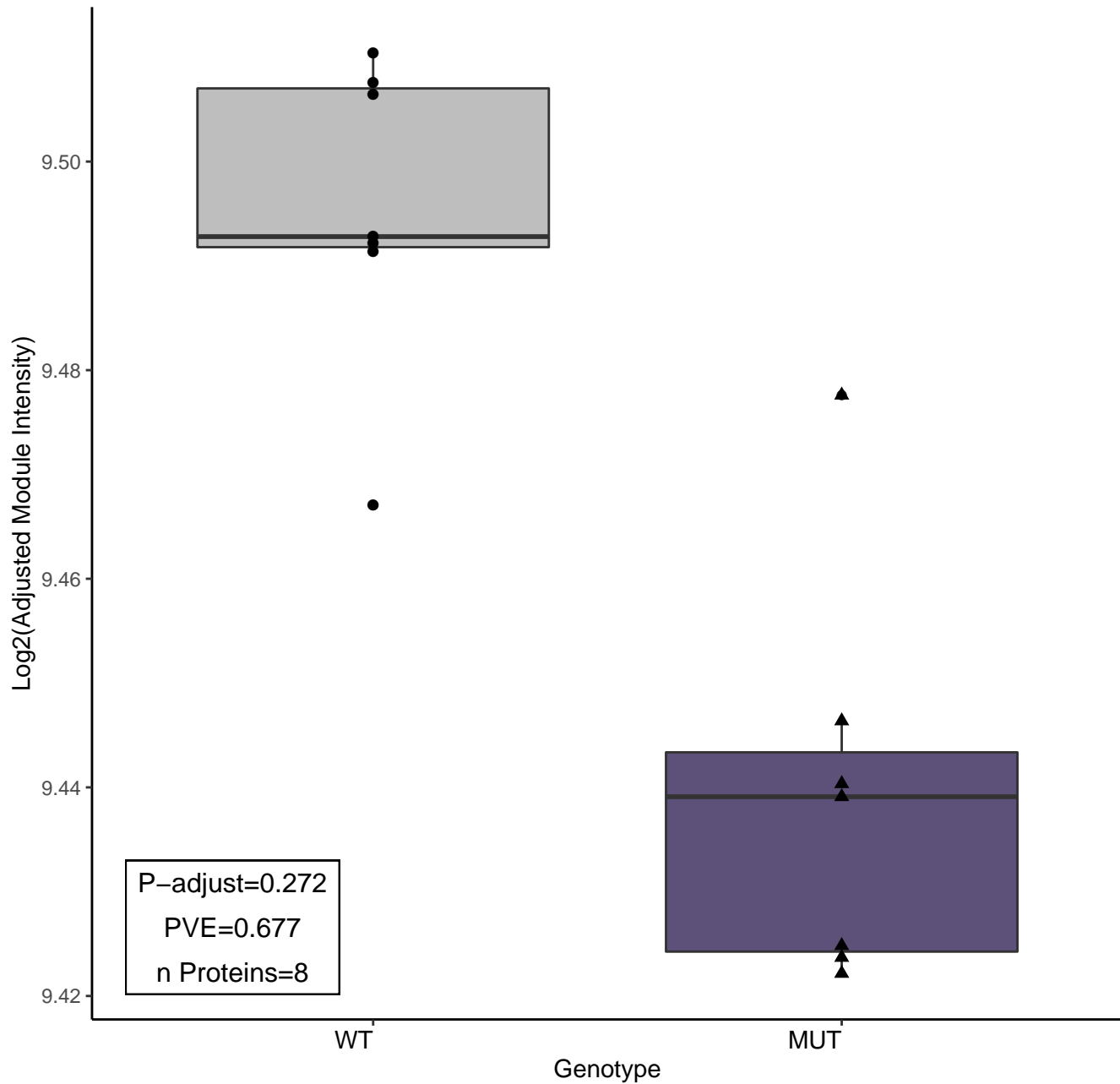
M139



M140



M141



M142

Log2(Adjusted Module Intensity)

14.55

14.50

14.45

P-adjust=0.16

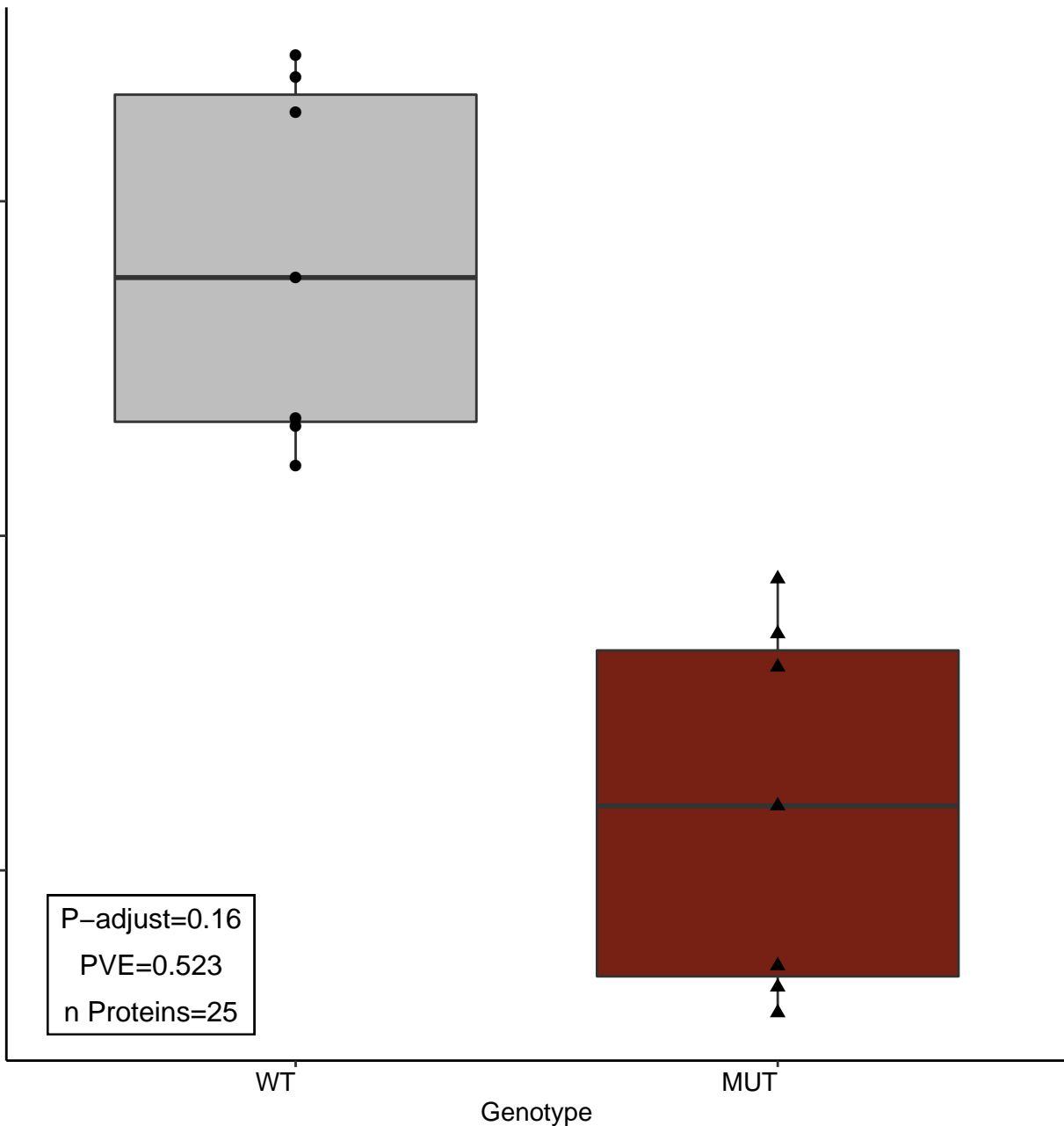
PVE=0.523

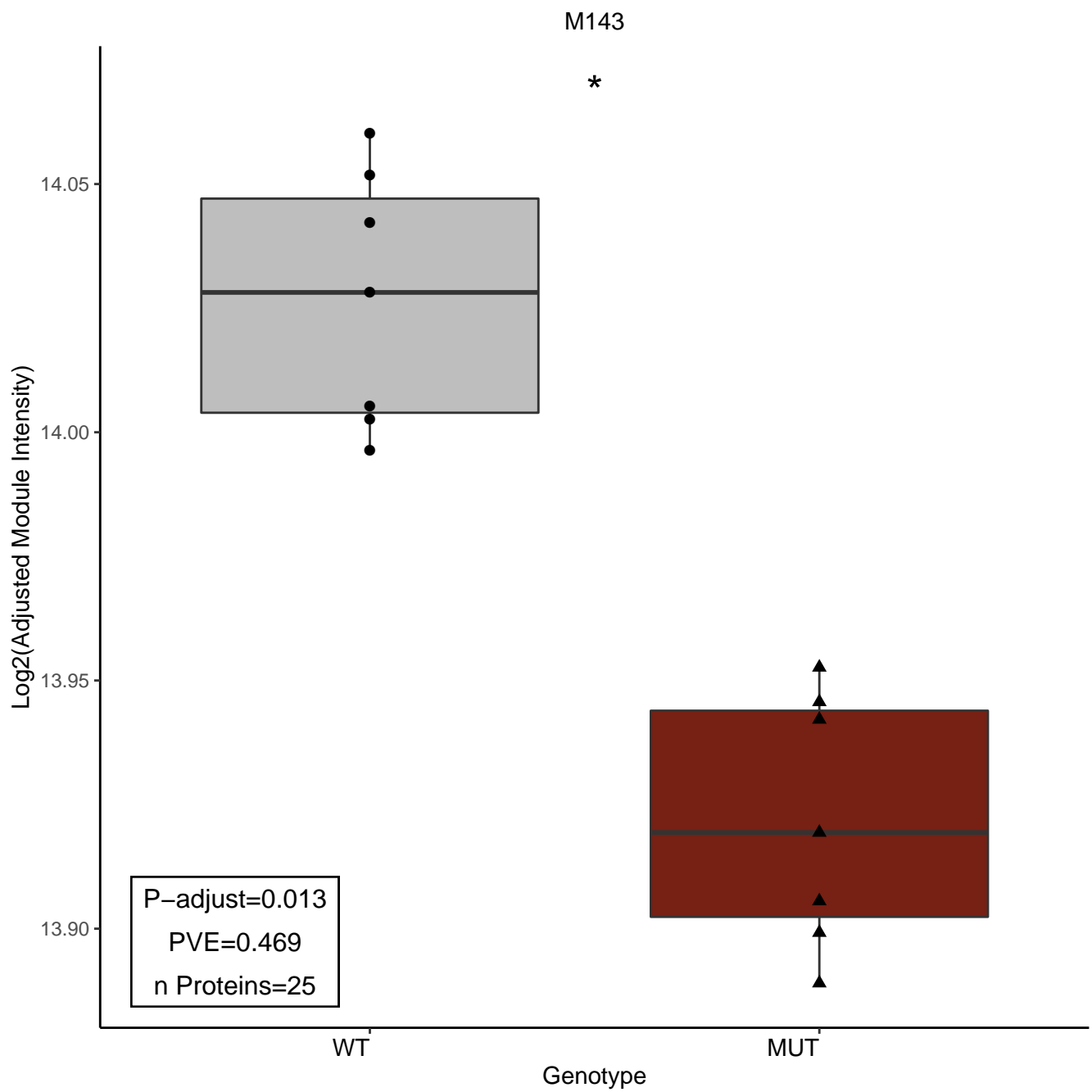
n Proteins=25

WT

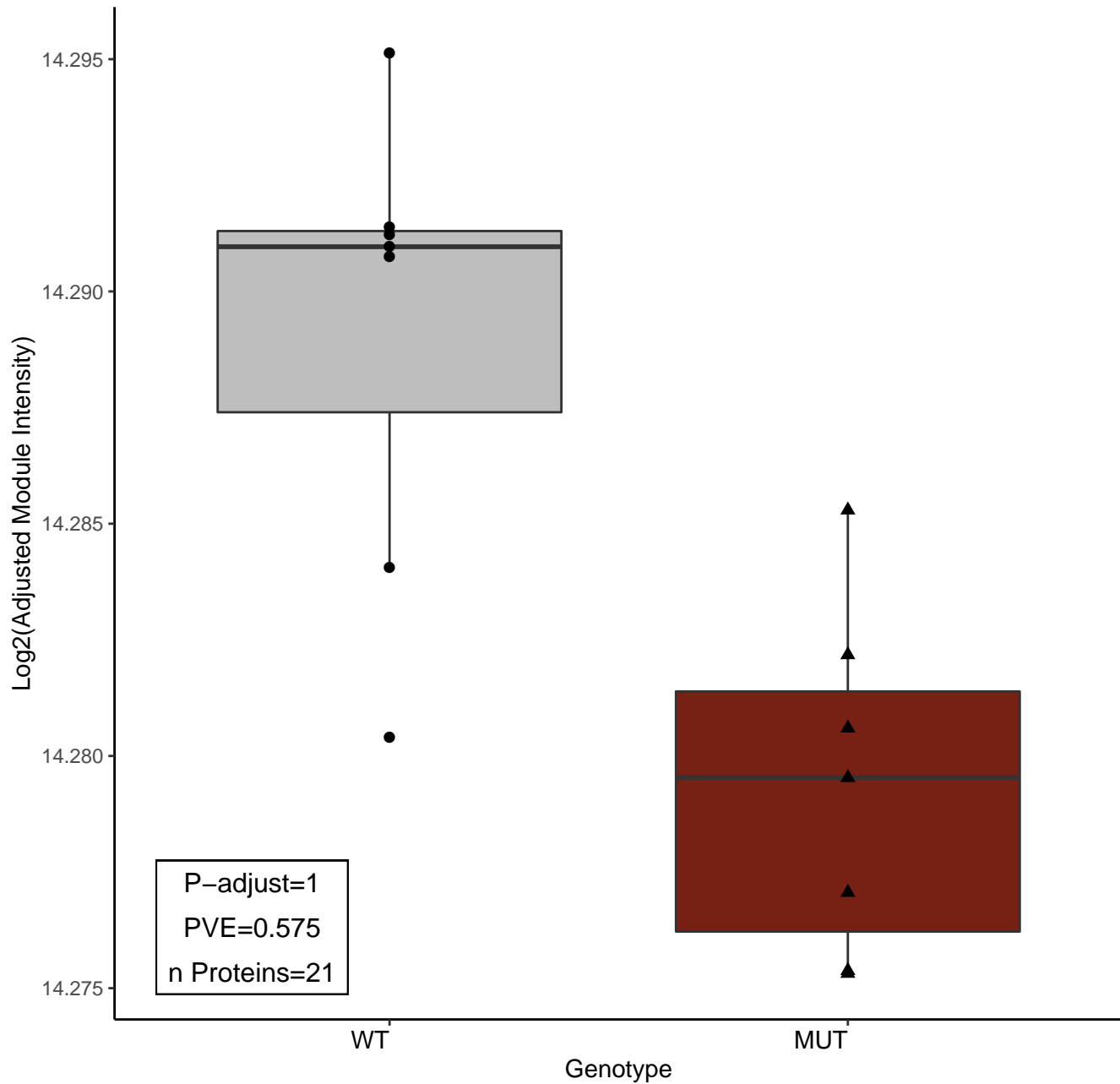
Genotype

MUT

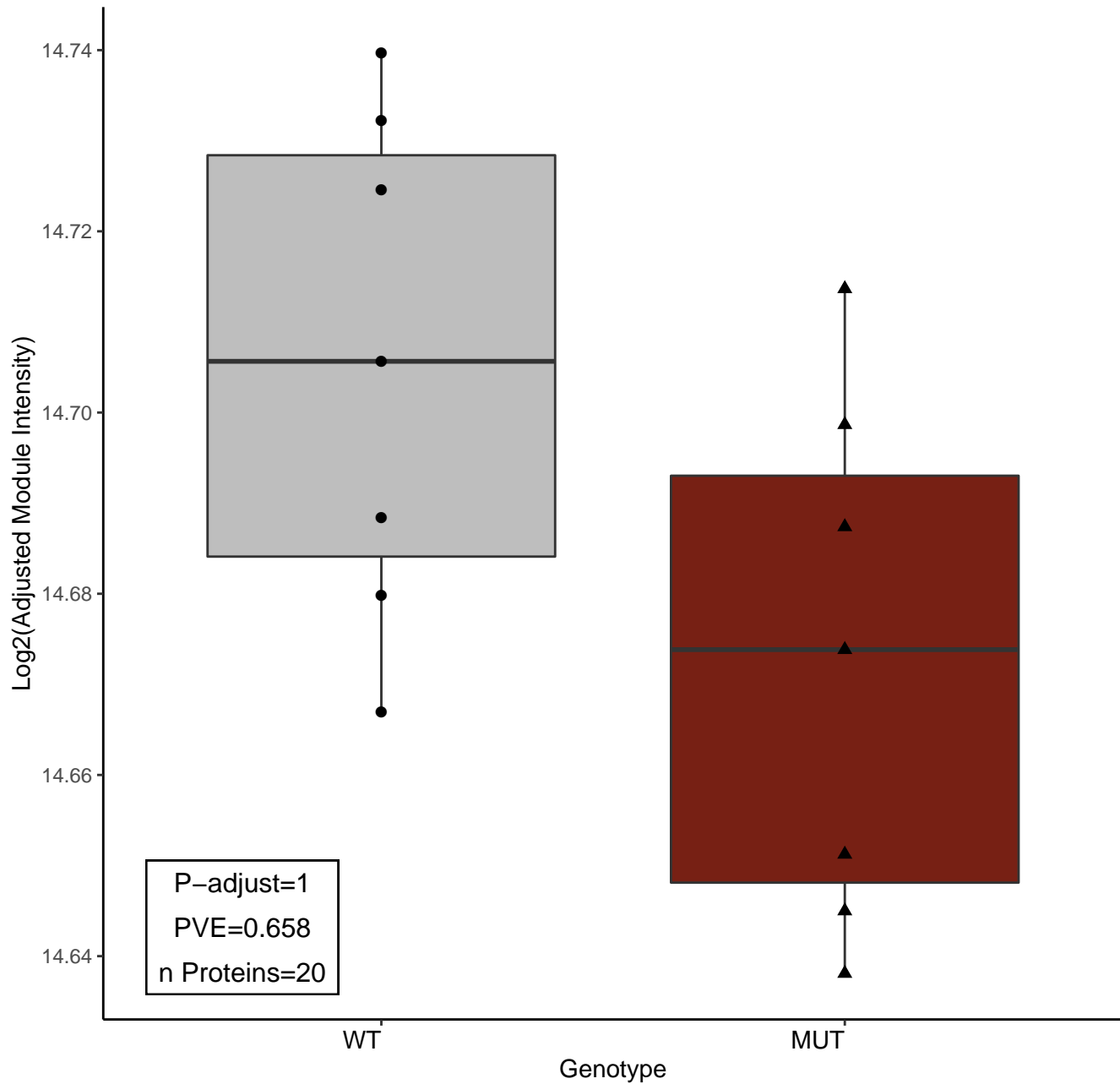




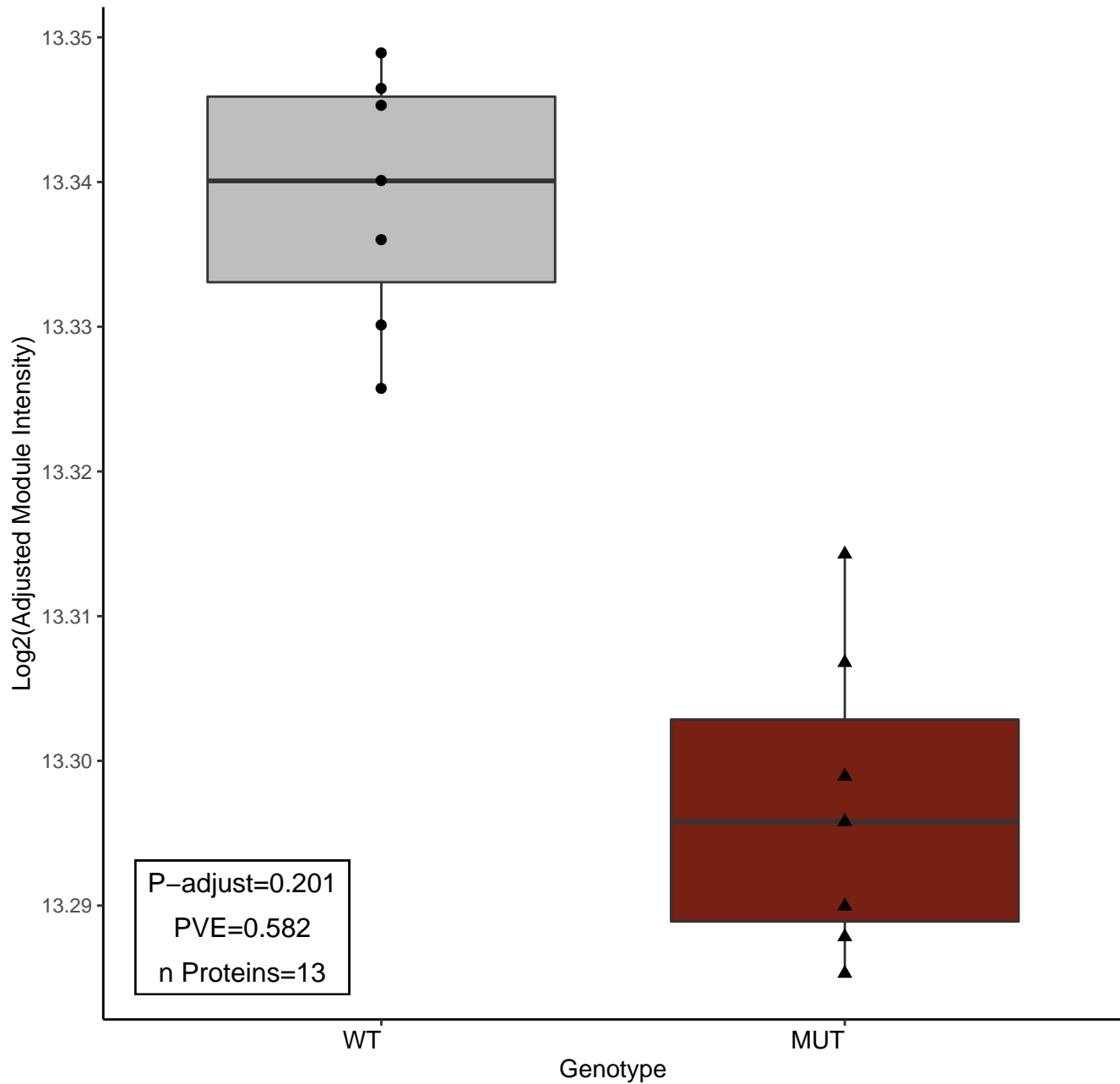
M144



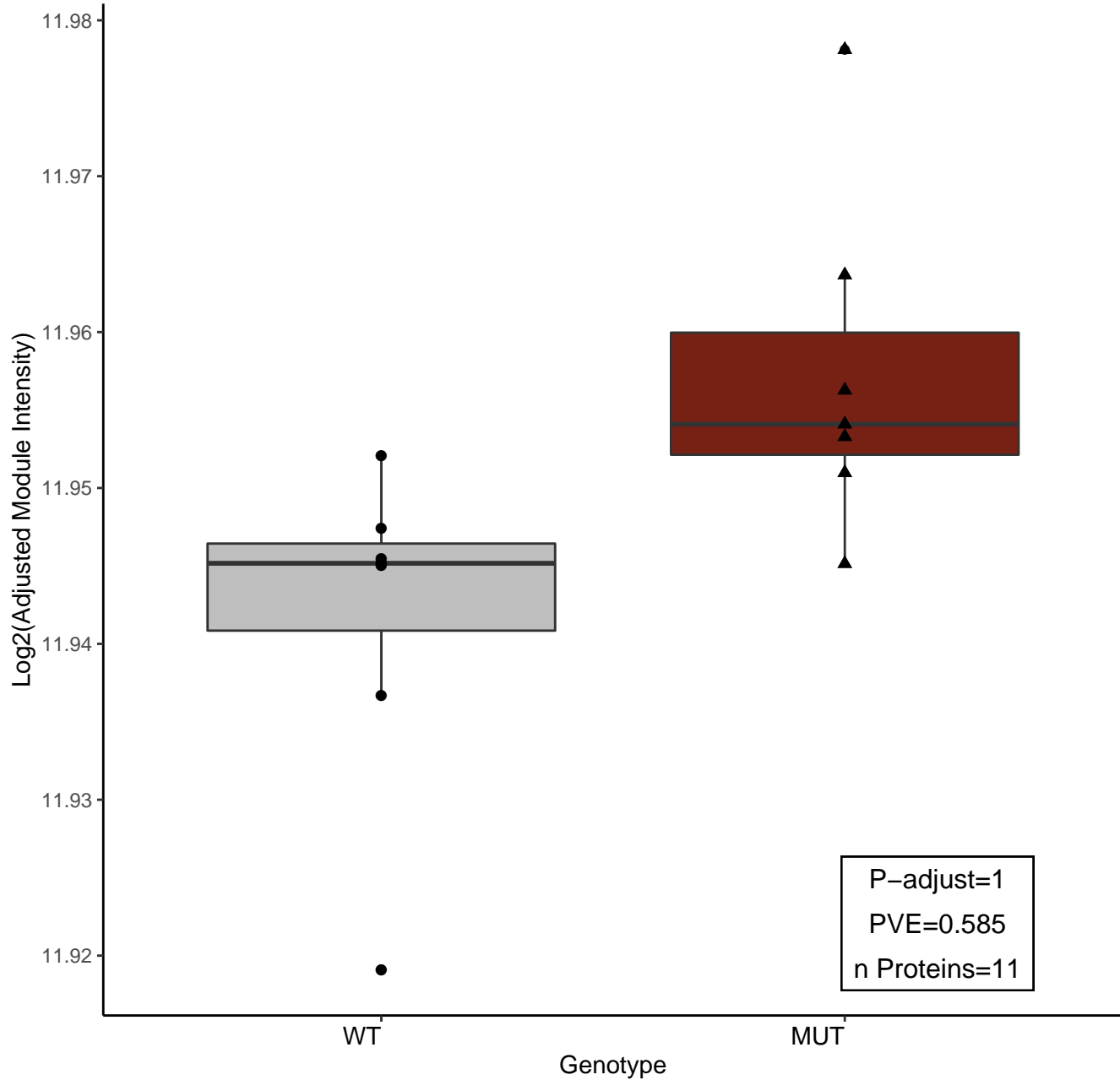
M145



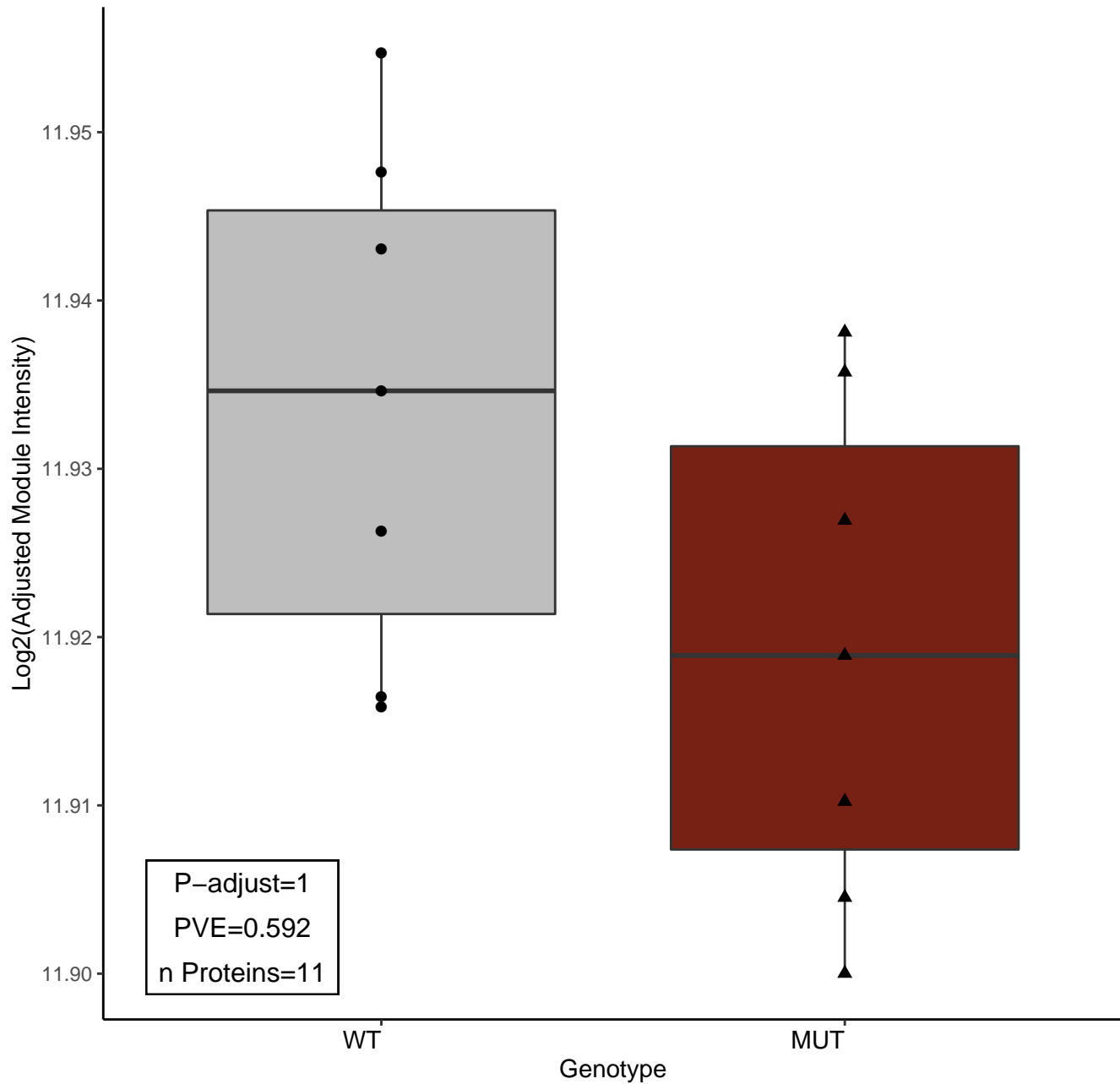
M146



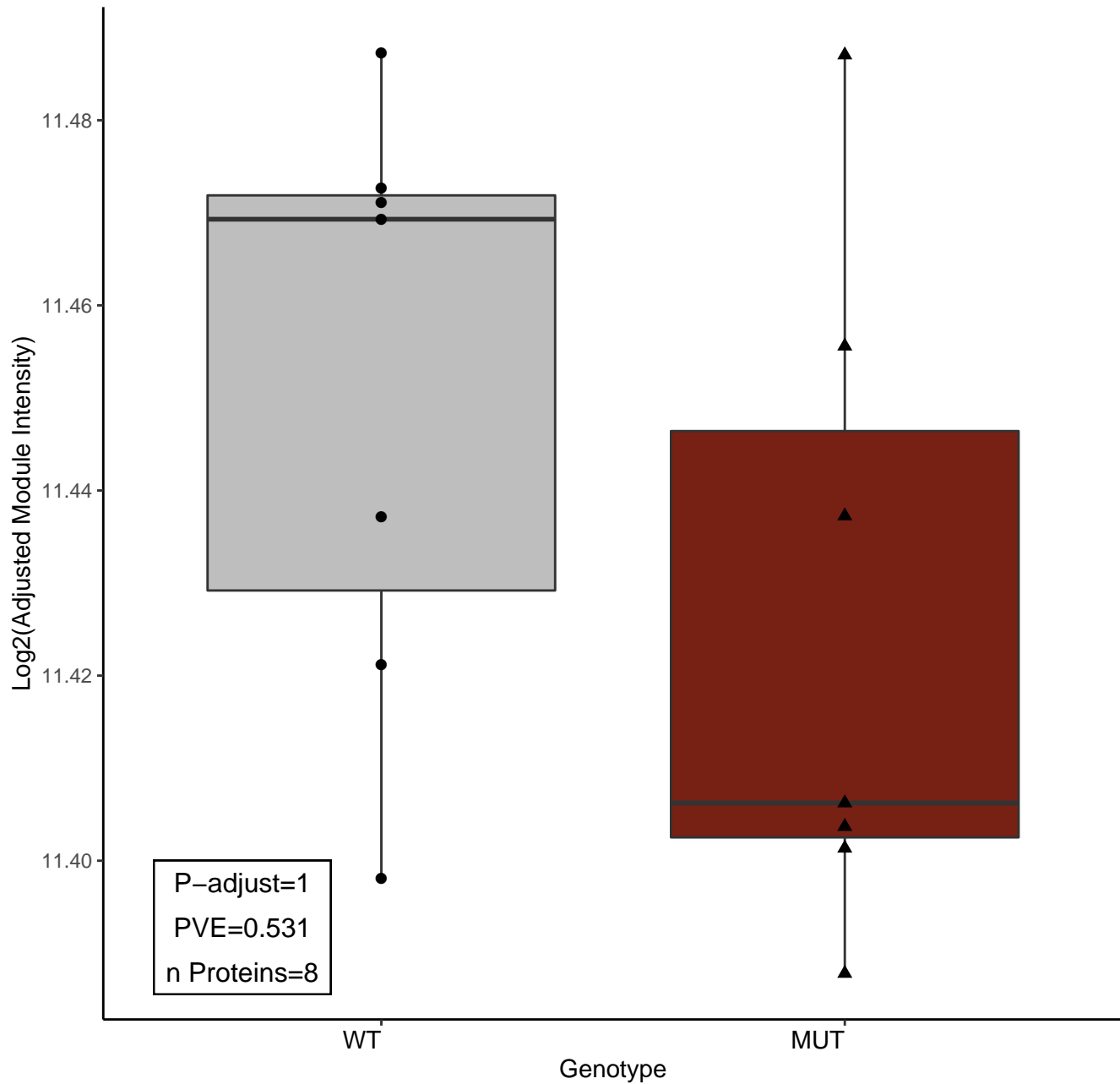
M147



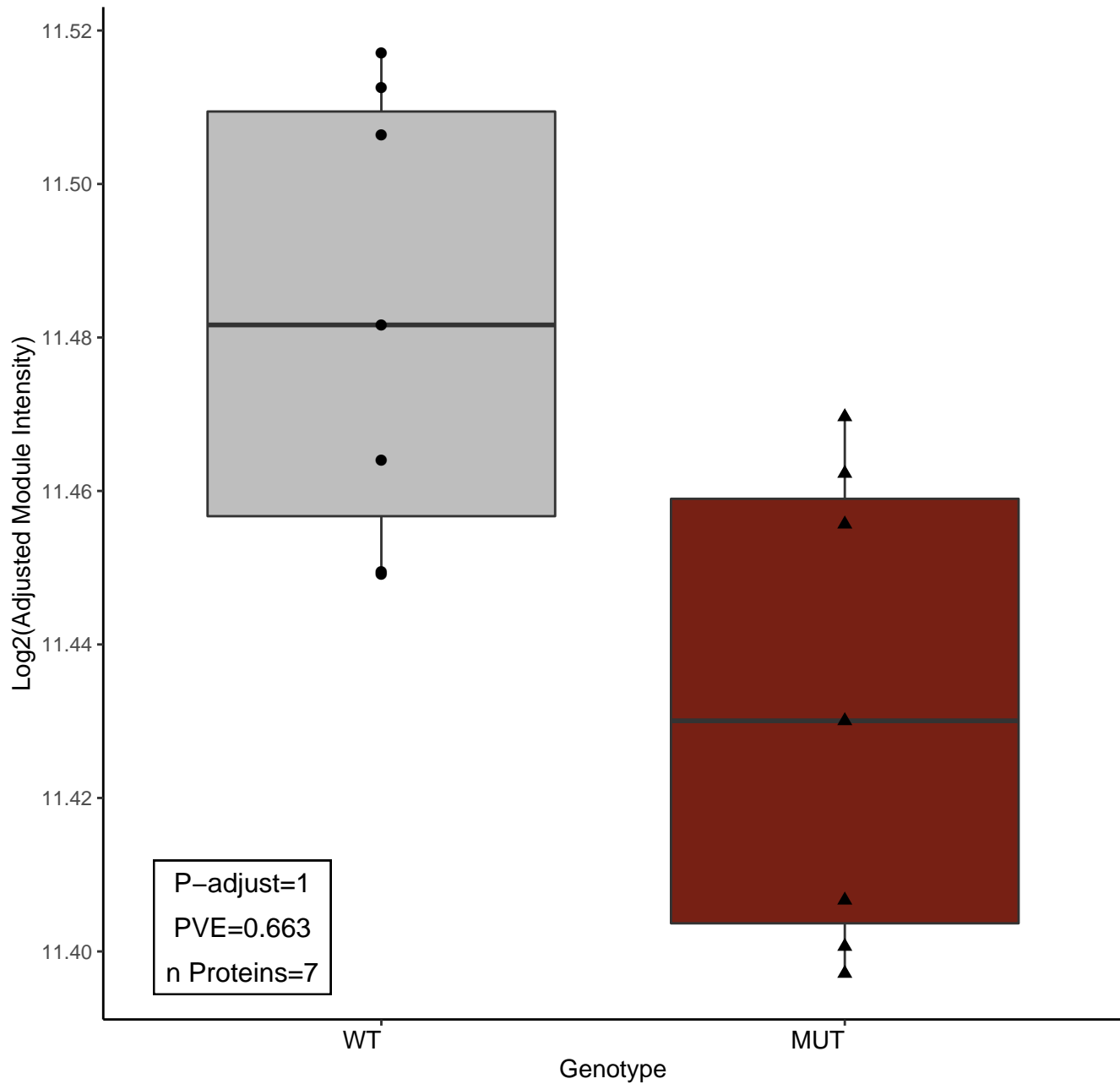
M148

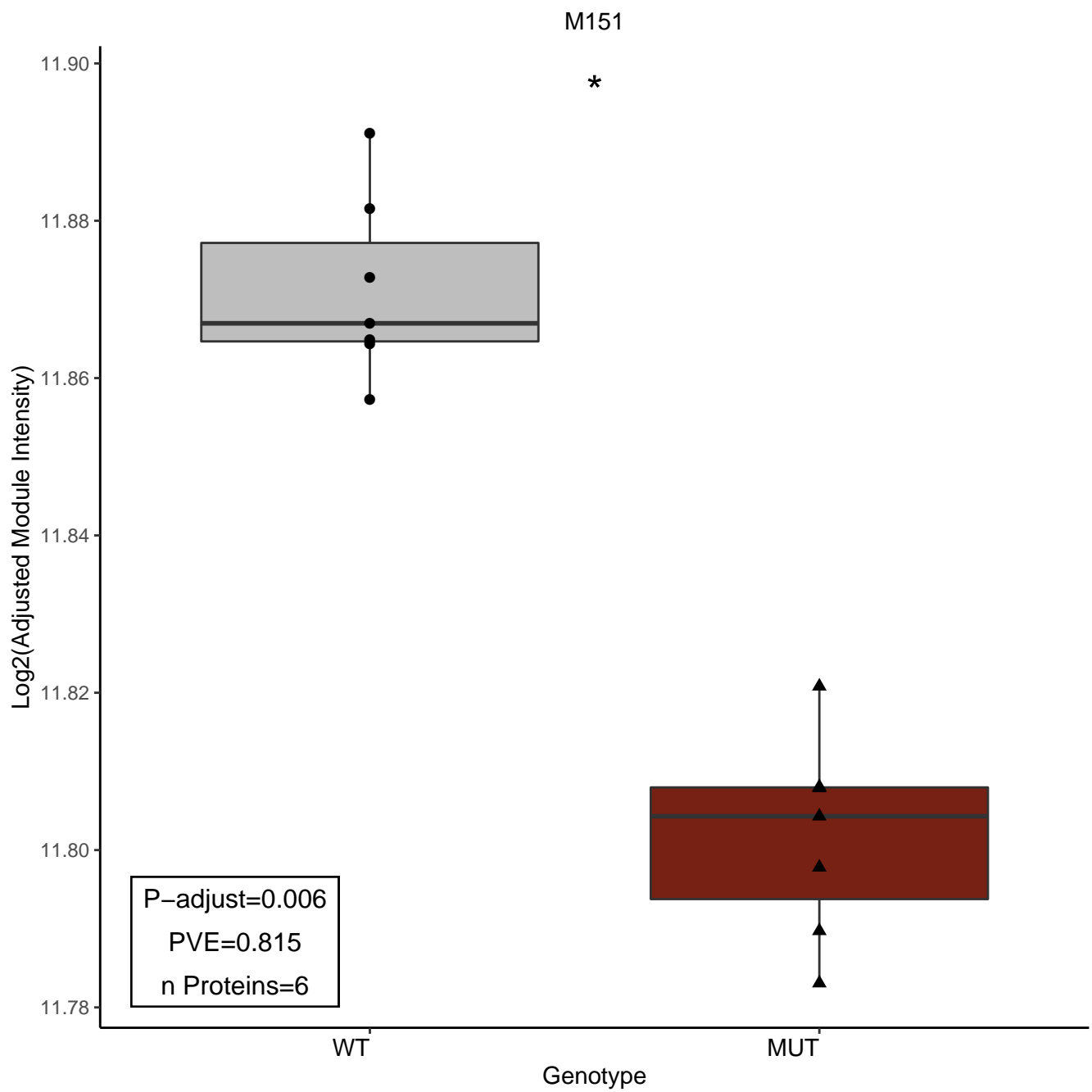


M149



M150





M152

Log2(Adjusted Module Intensity)

12.82

12.80

12.78

12.76

WT

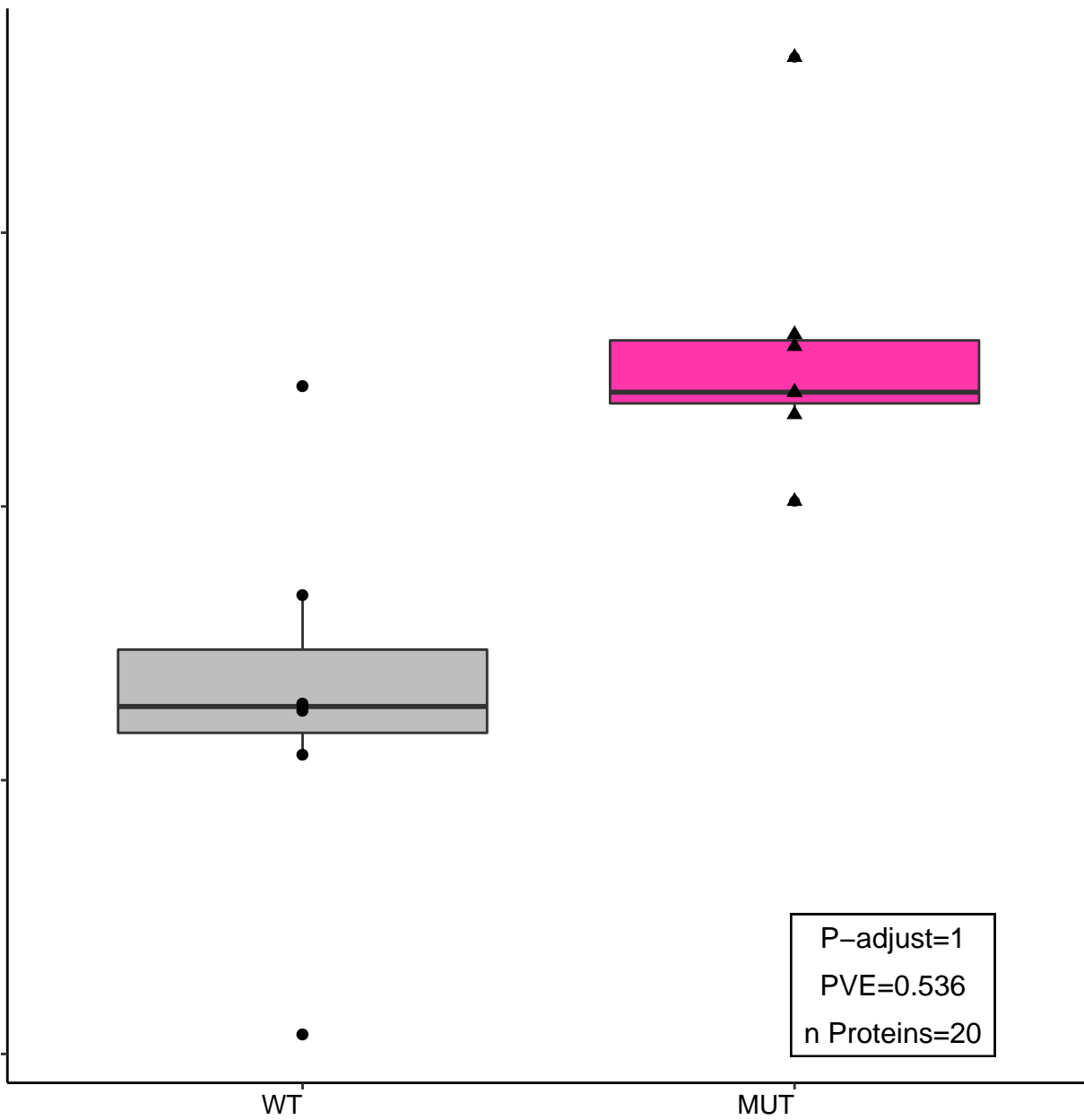
Genotype

MUT

P-adjust=1

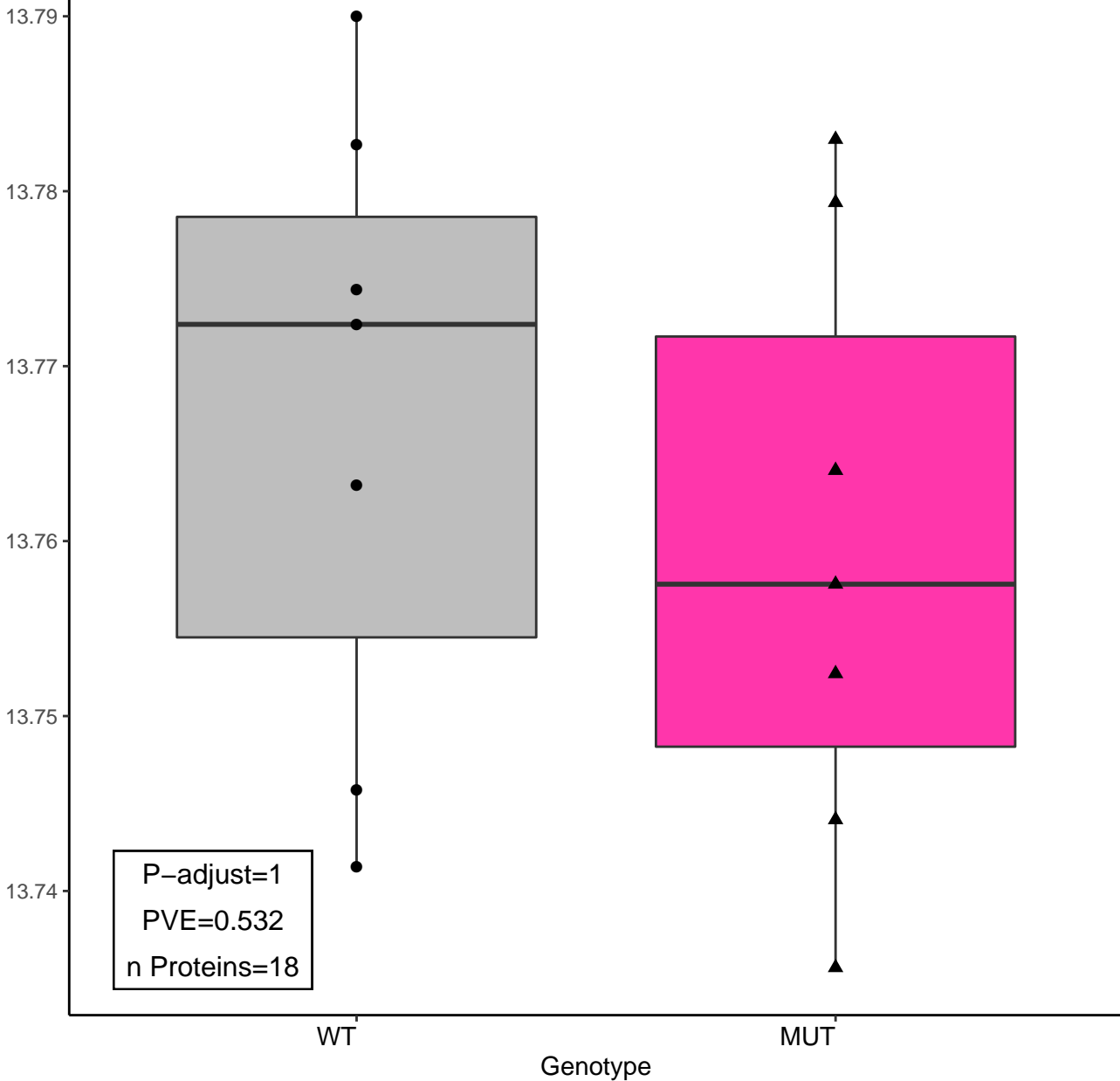
PVE=0.536

n Proteins=20



M153

Log2(Adjusted Module Intensity)



M154

Log2(Adjusted Module Intensity)

14.64

14.63

14.62

14.61

WT

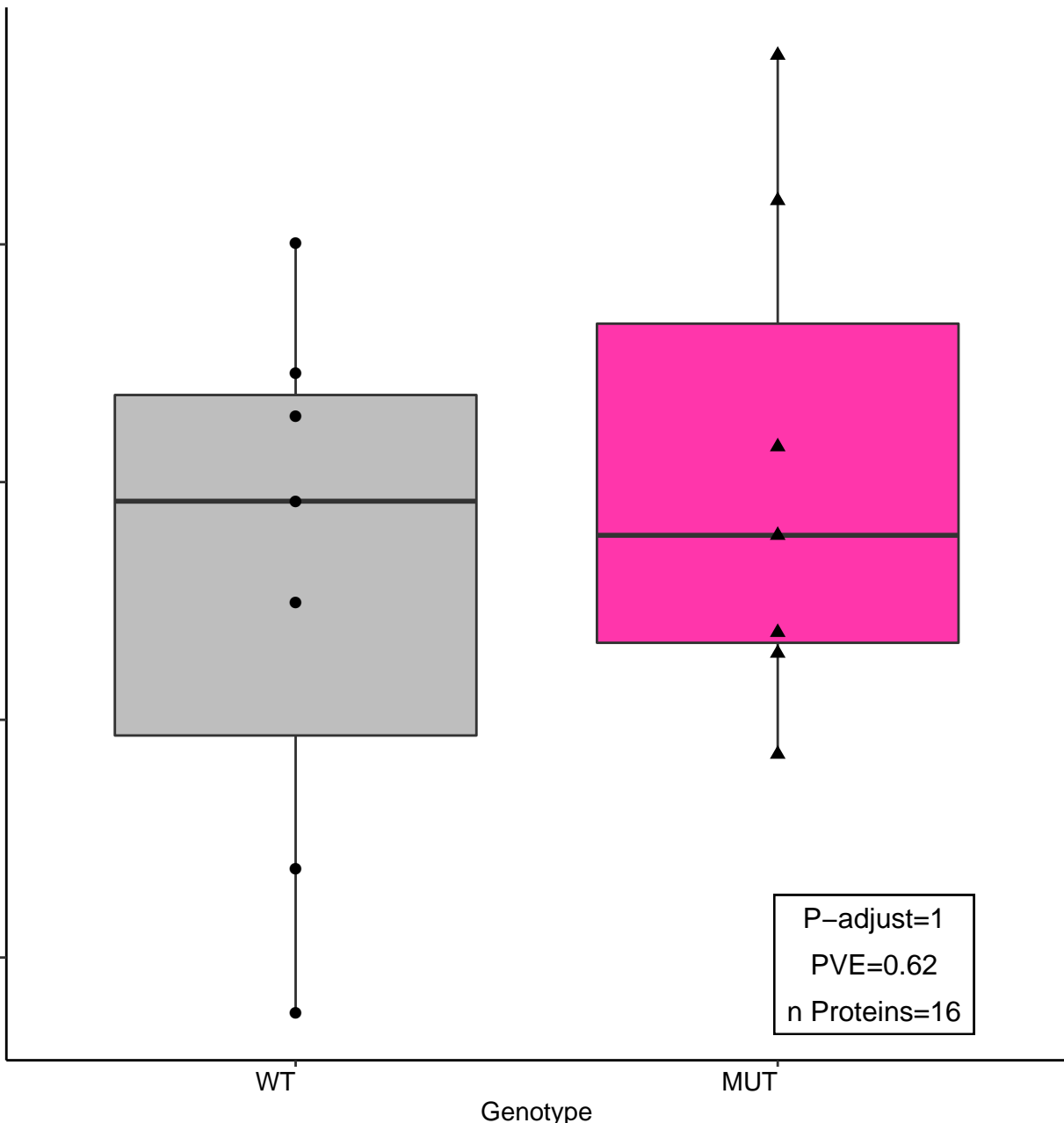
Genotype

MUT

P-adjust=1

PVE=0.62

n Proteins=16



M155

Log2(Adjusted Module Intensity)

13.35

13.30

13.25

WT

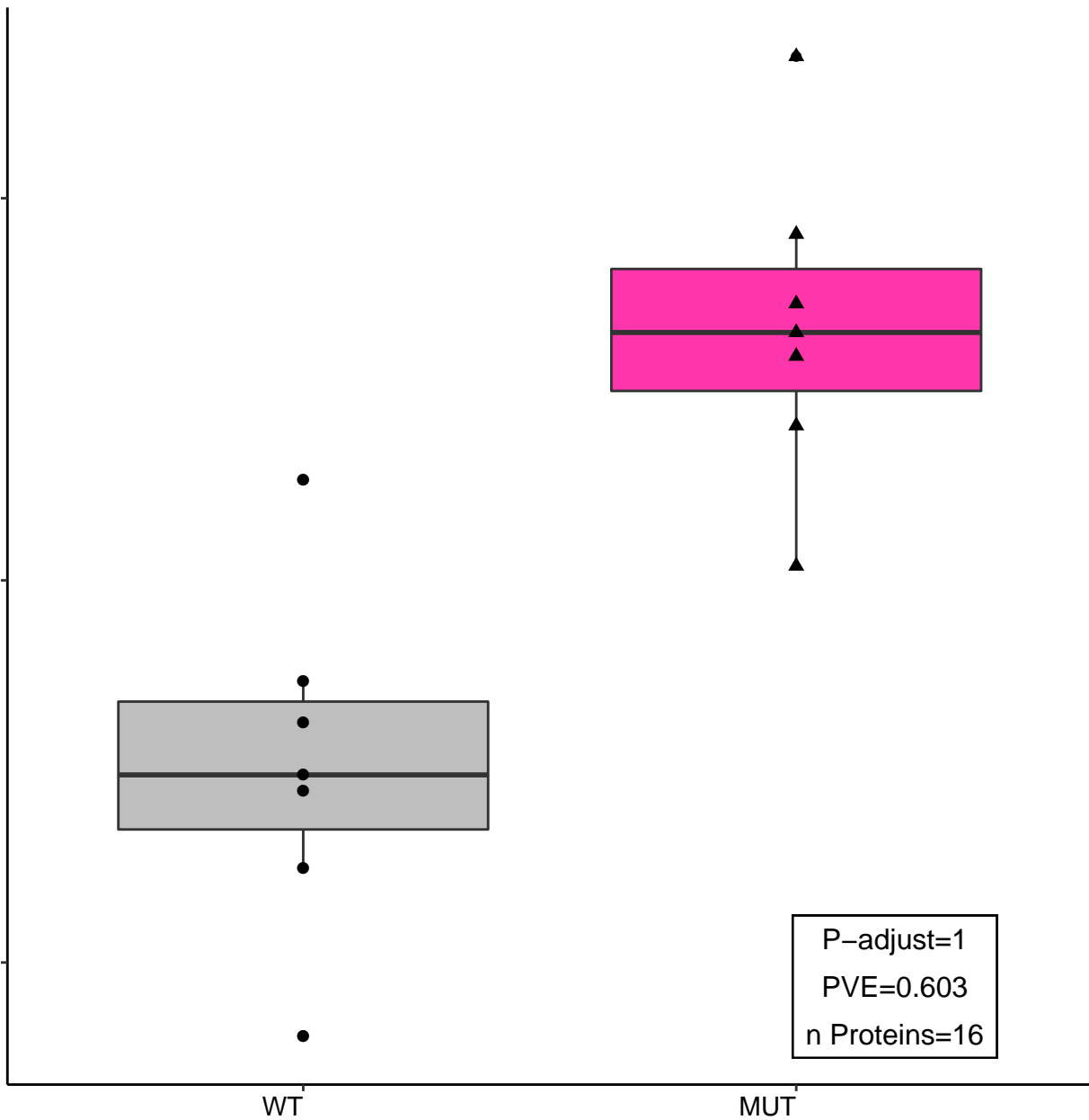
MUT

Genotype

P-adjust=1

PVE=0.603

n Proteins=16



M156

**

Log2(Adjusted Module Intensity)

12.65

12.60

12.55

P-adjust=0.004

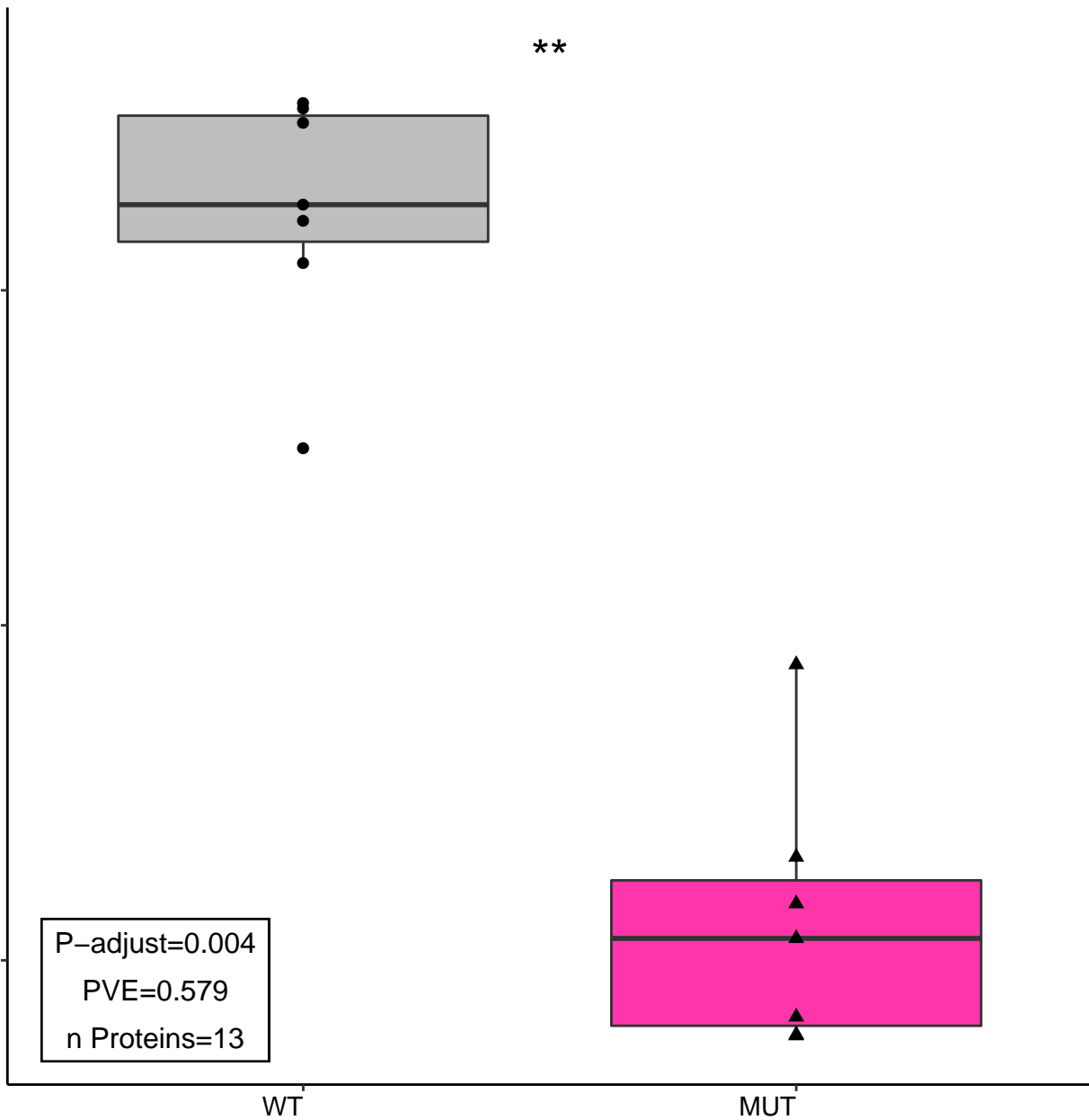
PVE=0.579

n Proteins=13

WT

Genotype

MUT



M157

Log2(Adjusted Module Intensity)

10.95

10.90

10.85

10.80

WT

Genotype

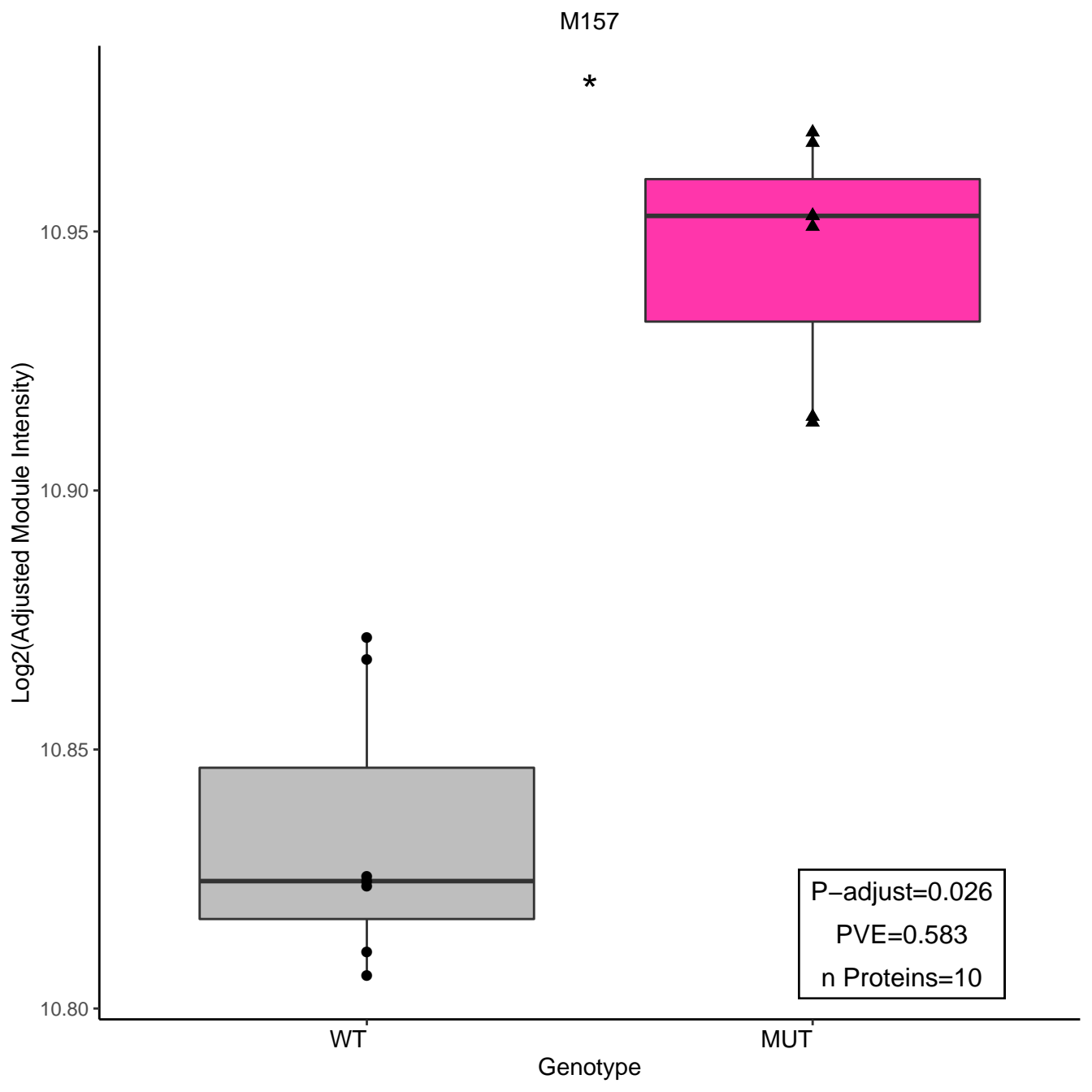
MUT

P-adjust=0.026

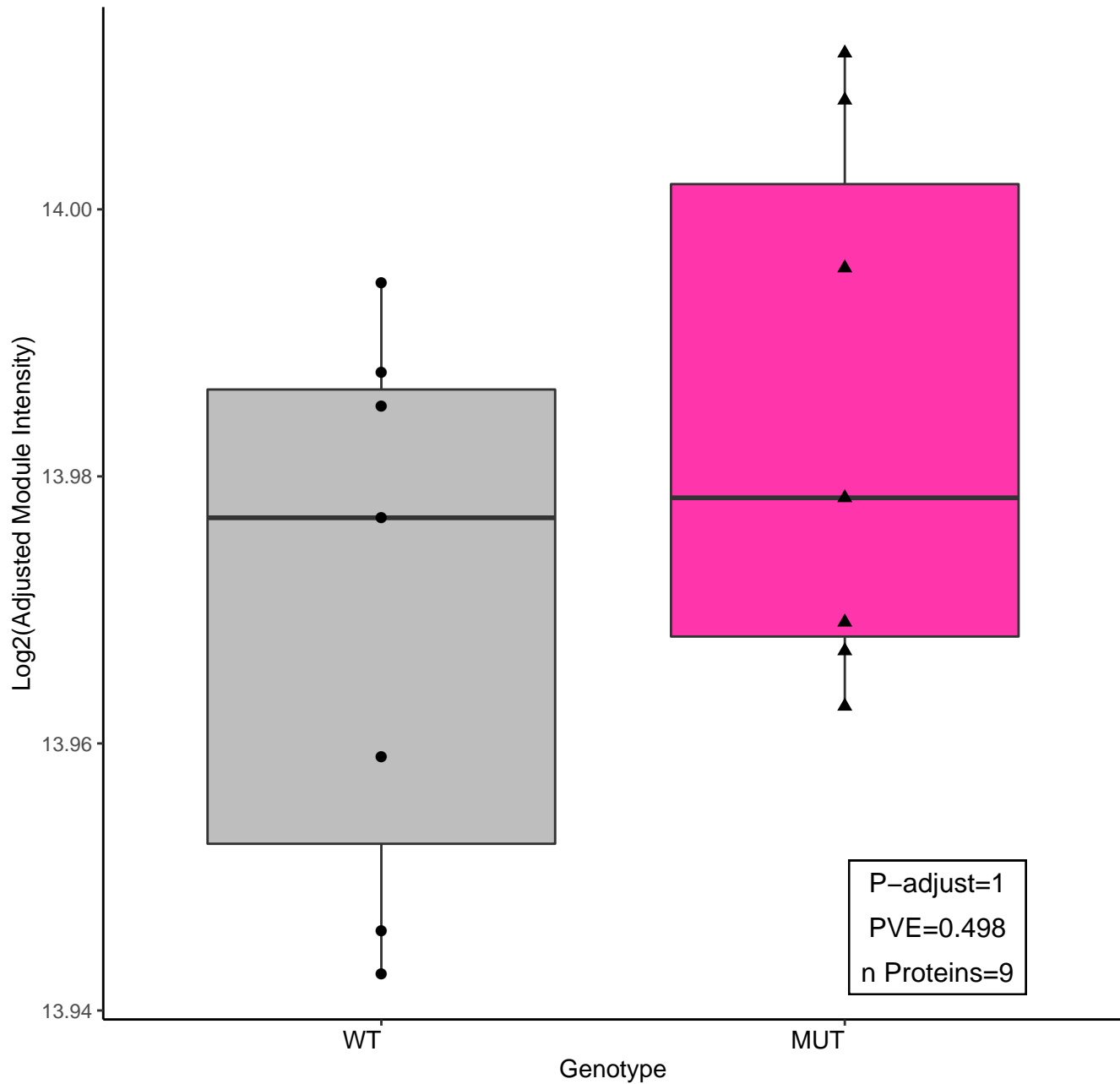
PVE=0.583

n Proteins=10

*



M158



M159

Log2(Adjusted Module Intensity)

11.30

11.28

11.26

11.24

11.22

11.20

P-adjust=0.679

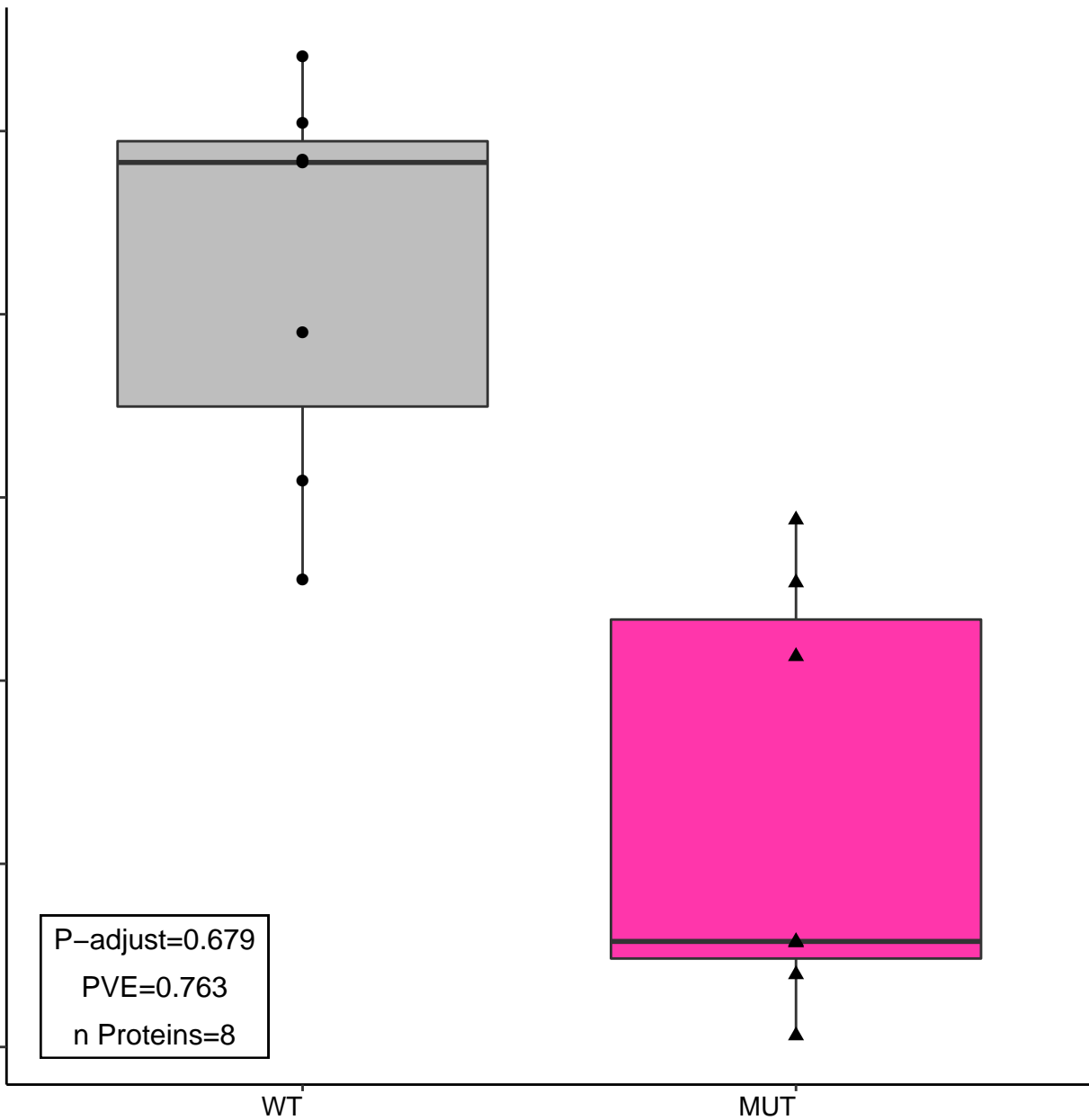
PVE=0.763

n Proteins=8

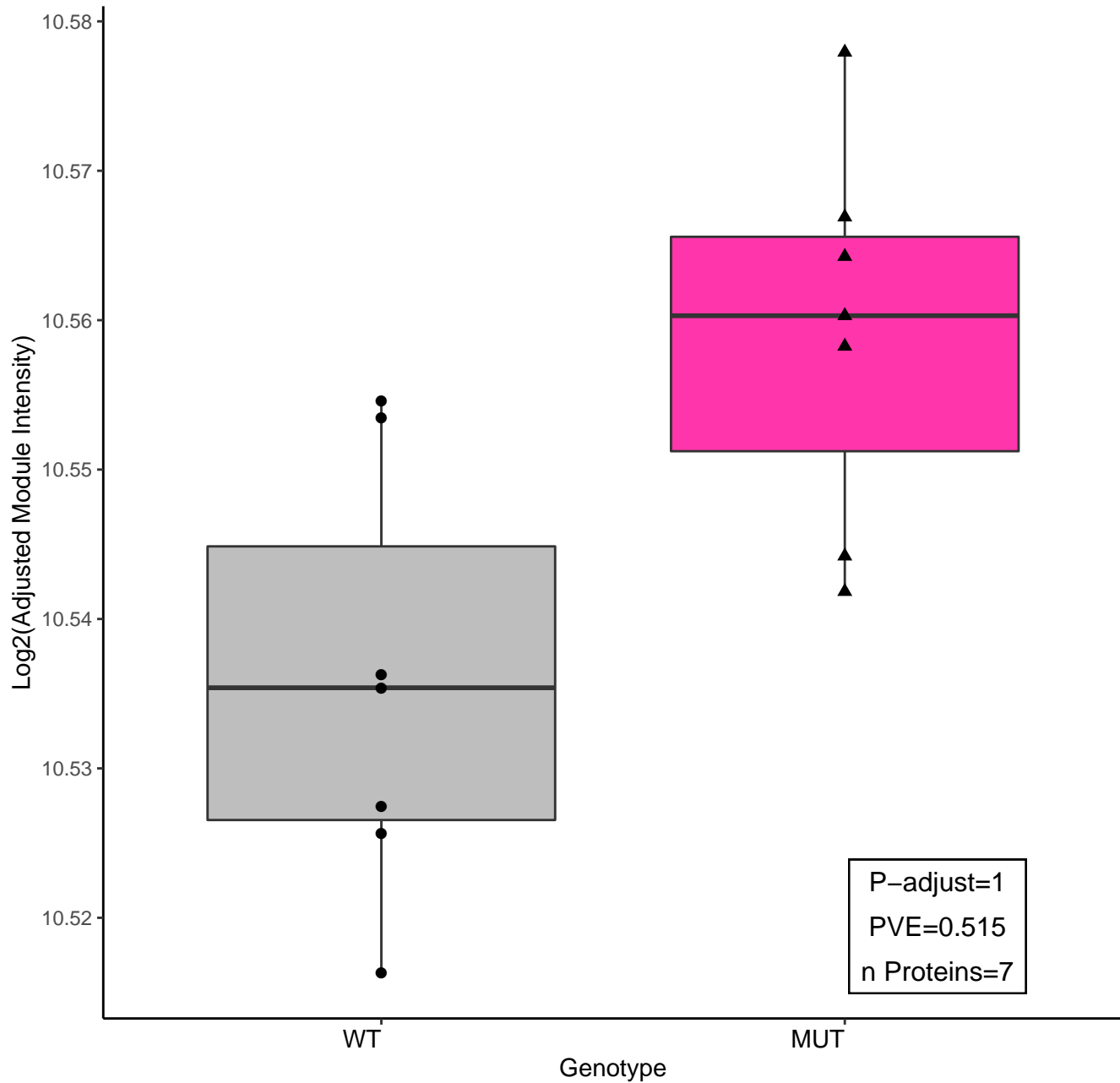
WT

Genotype

MUT



M160



M161

Log2(Adjusted Module Intensity)

14.46

14.44

14.42

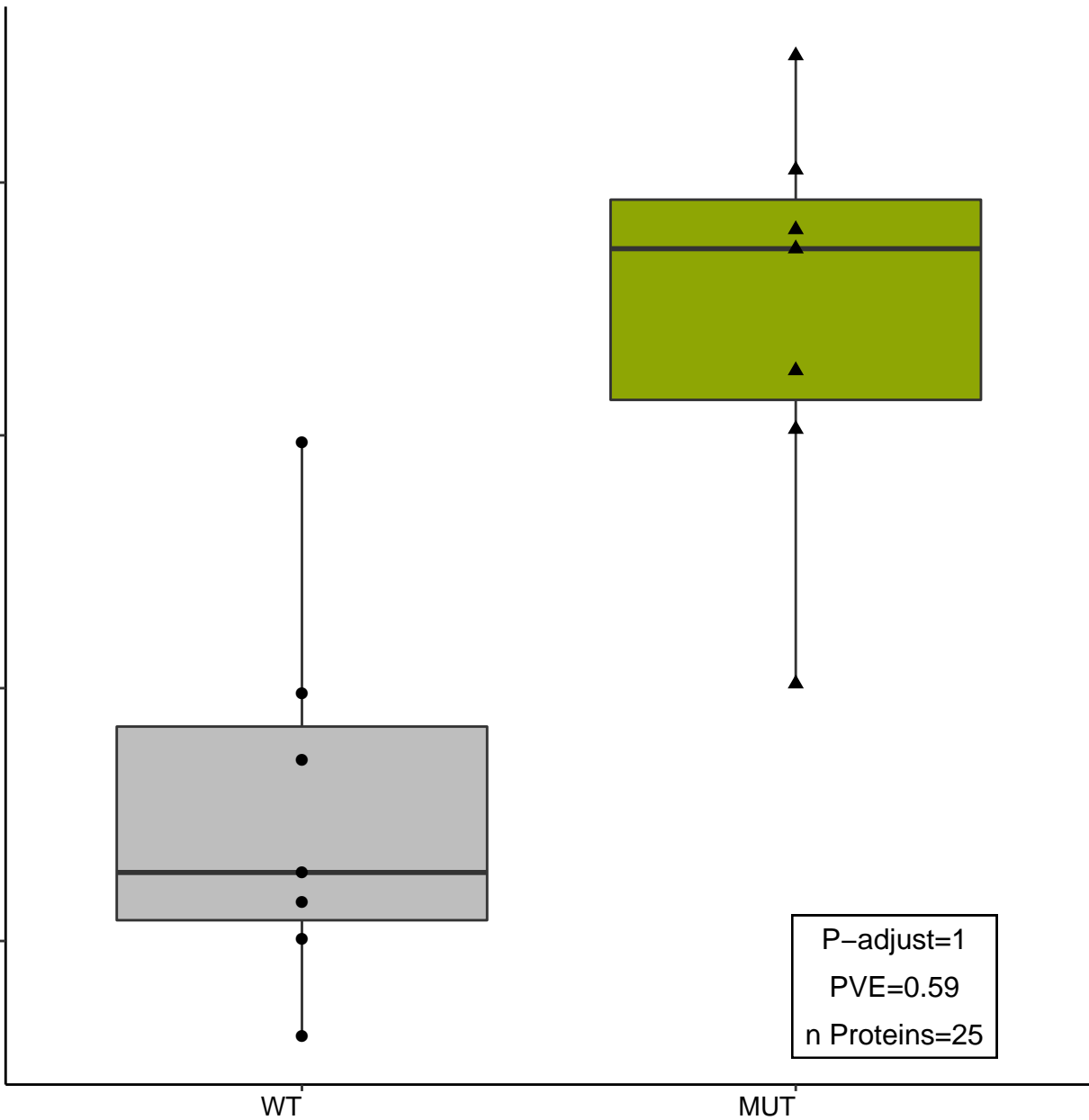
14.40

WT

Genotype

MUT

P-adjust=1
PVE=0.59
n Proteins=25



M162

Log2(Adjusted Module Intensity)

13.20

13.18

13.16

WT

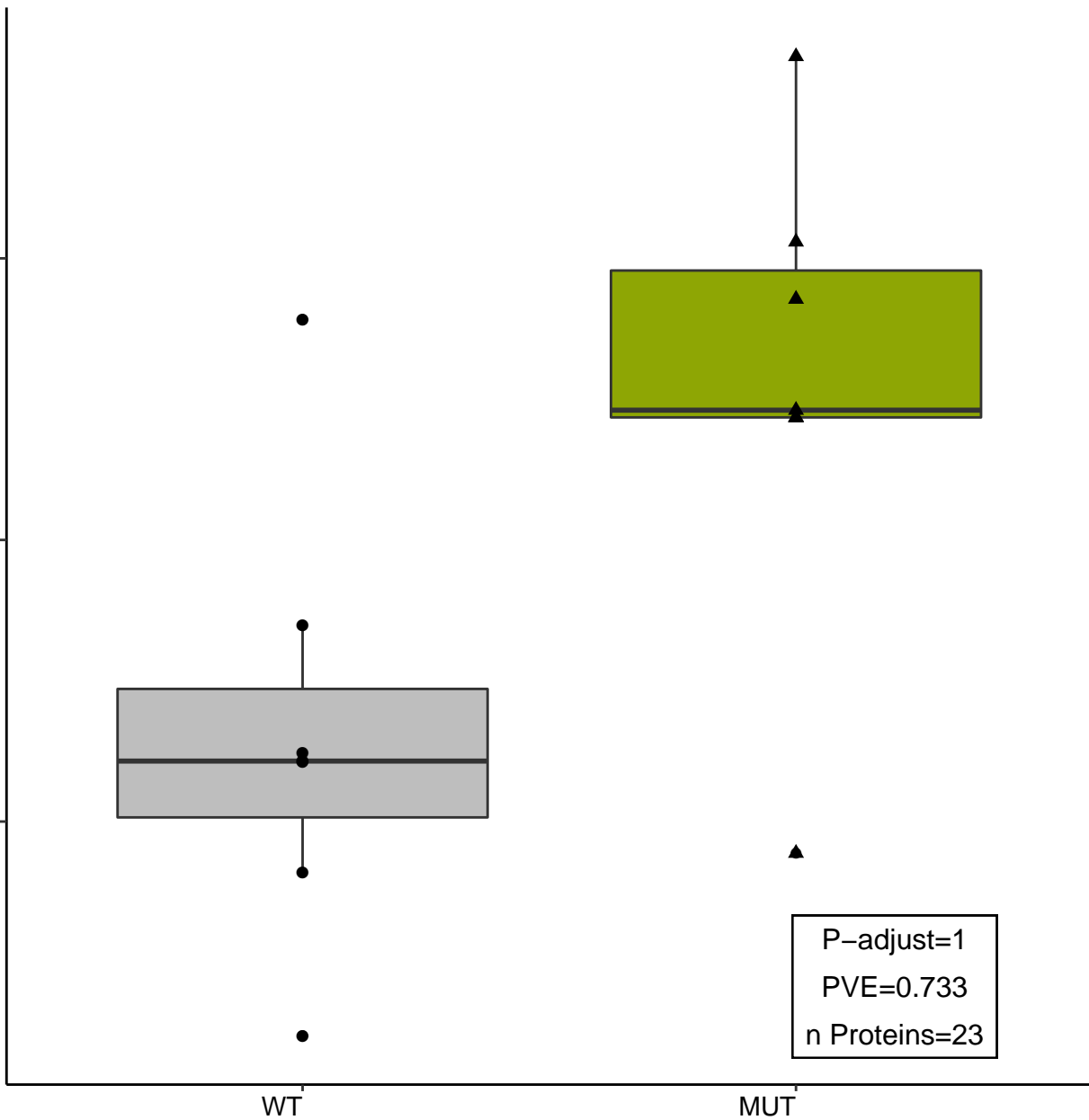
MUT

Genotype

P-adjust=1

PVE=0.733

n Proteins=23



M163

Log2(Adjusted Module Intensity)

13.74

13.72

13.70

WT

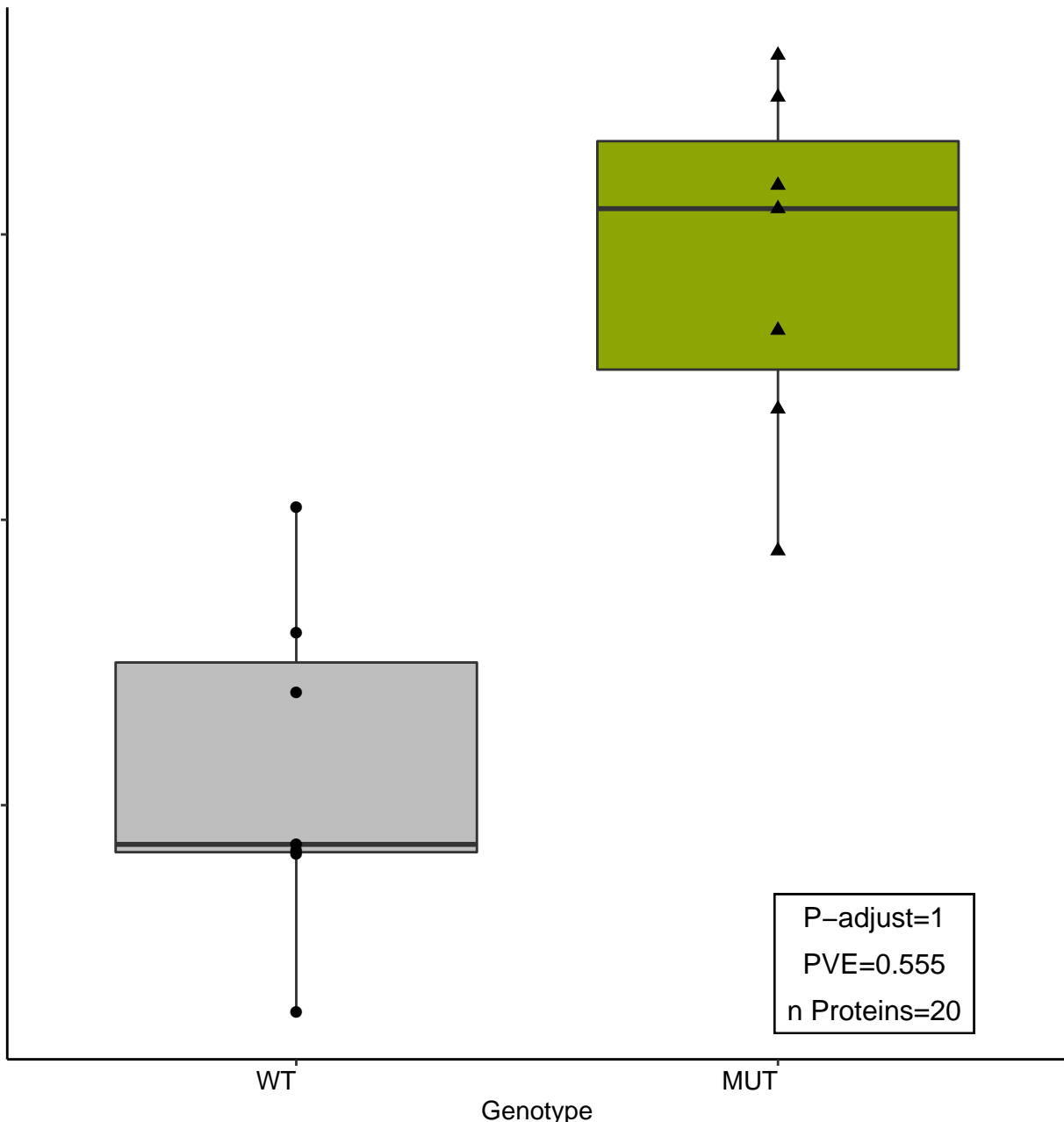
Genotype

MUT

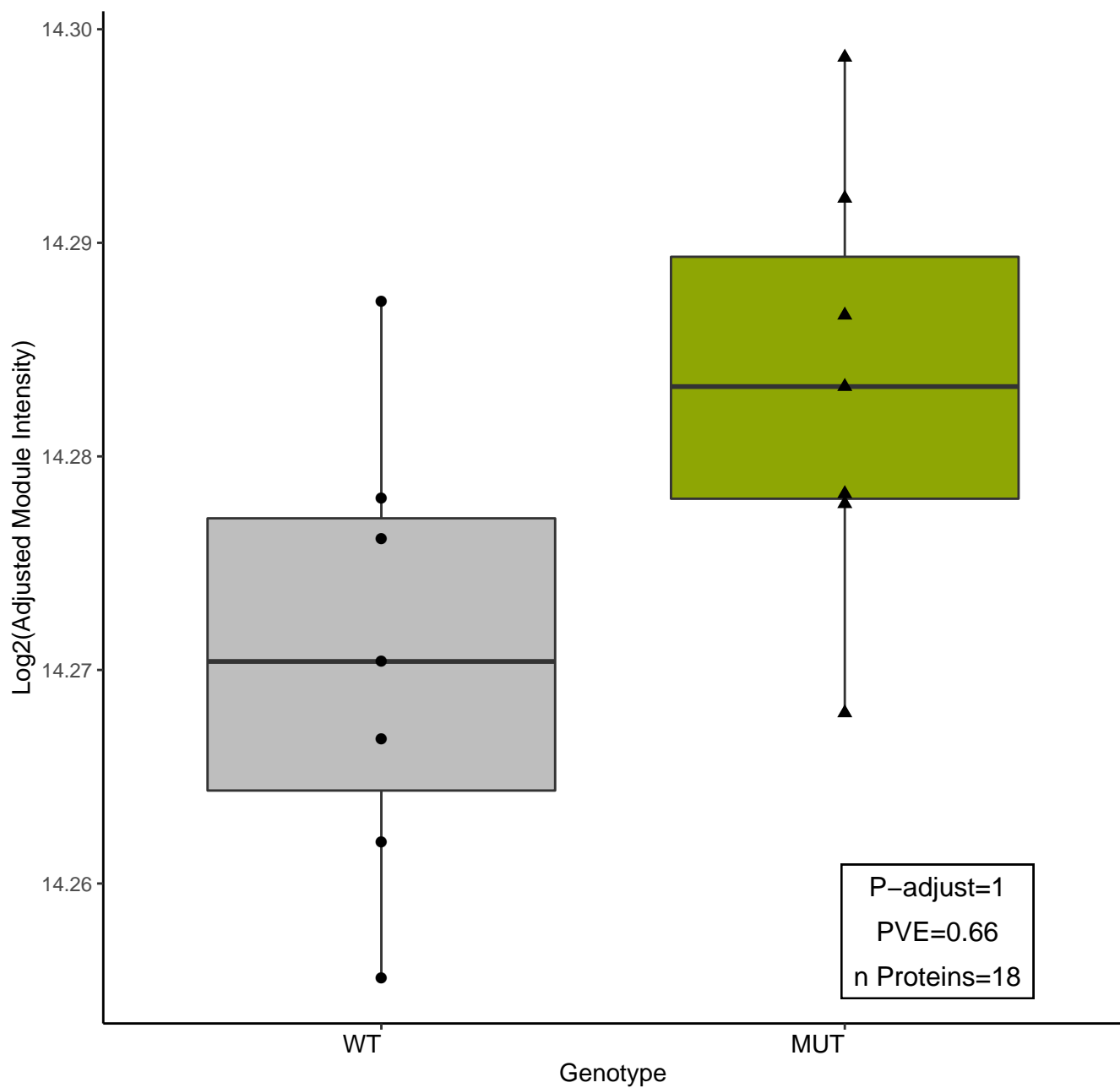
P-adjust=1

PVE=0.555

n Proteins=20



M164



M165

Log2(Adjusted Module Intensity)

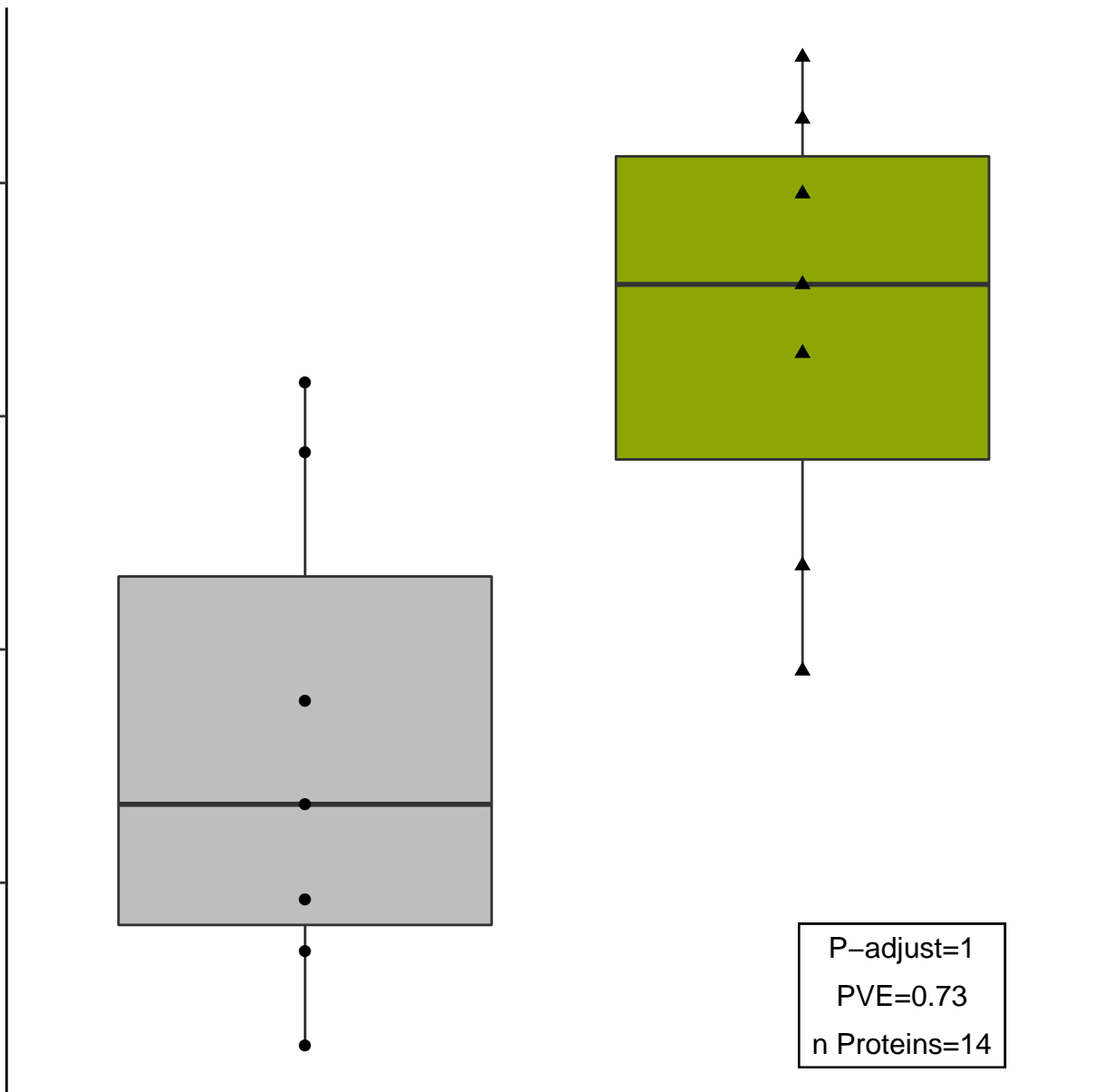
13.93
13.92
13.91
13.90

WT

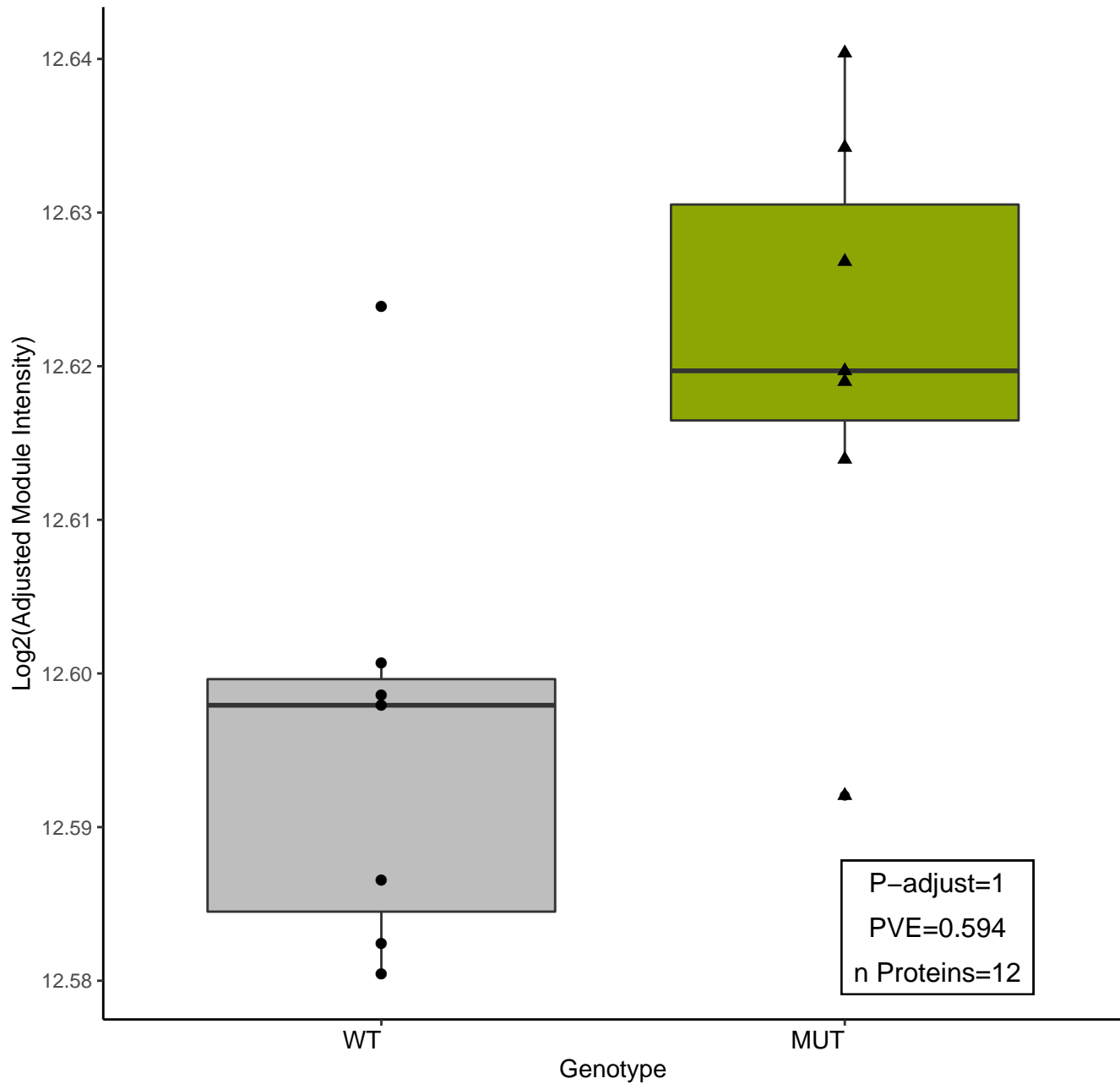
Genotype

MUT

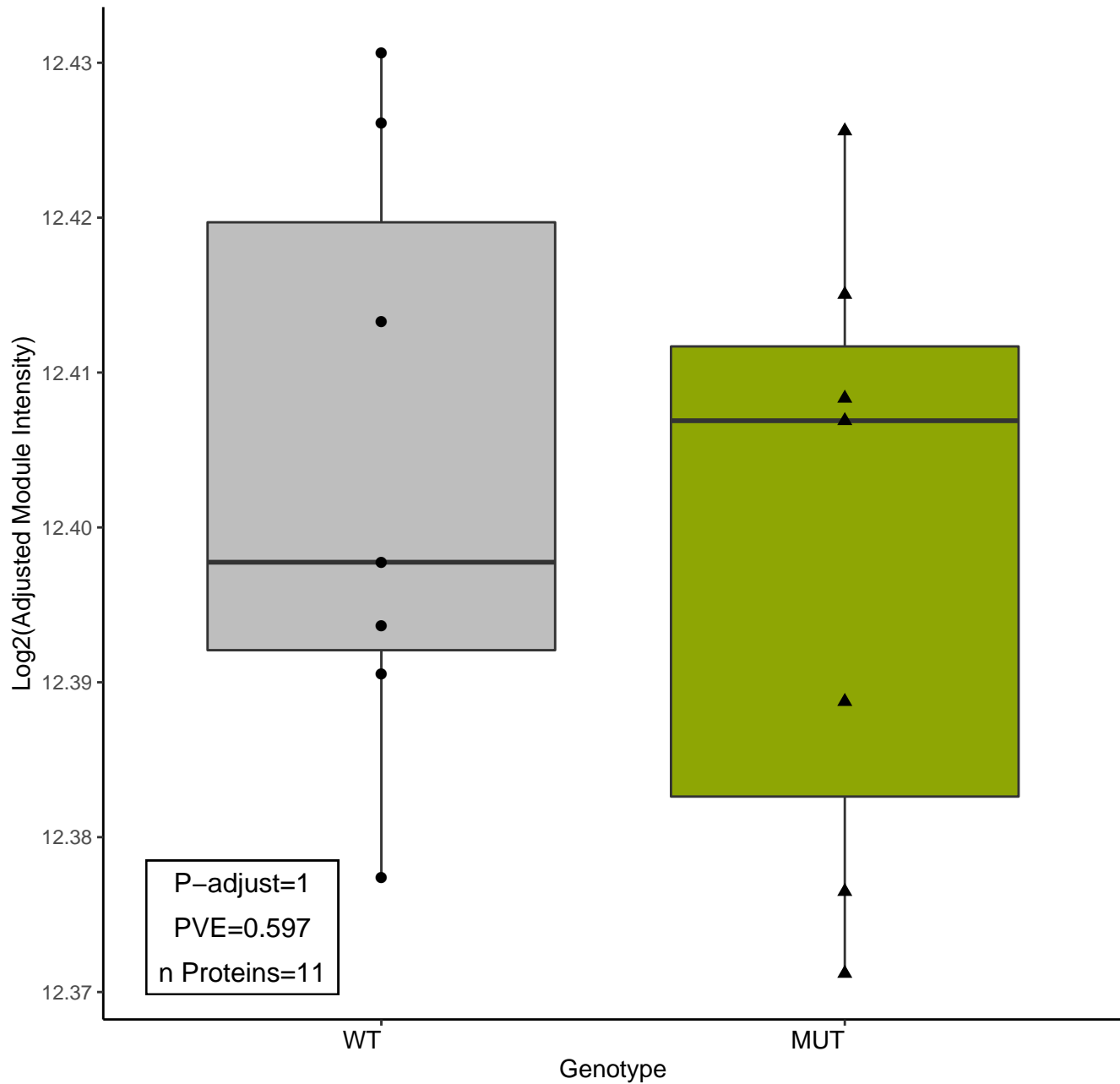
P-adjust=1
PVE=0.73
n Proteins=14



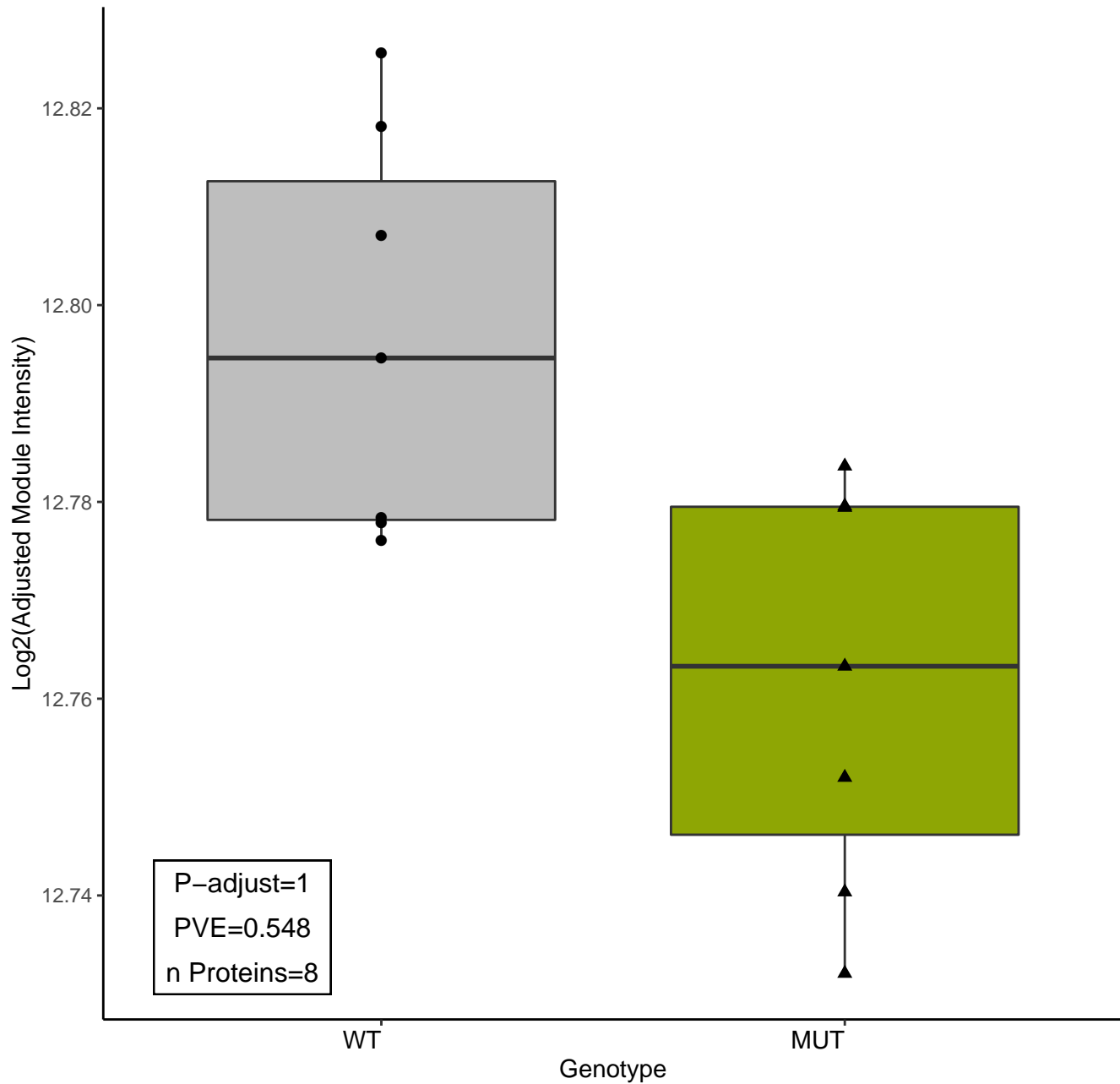
M166



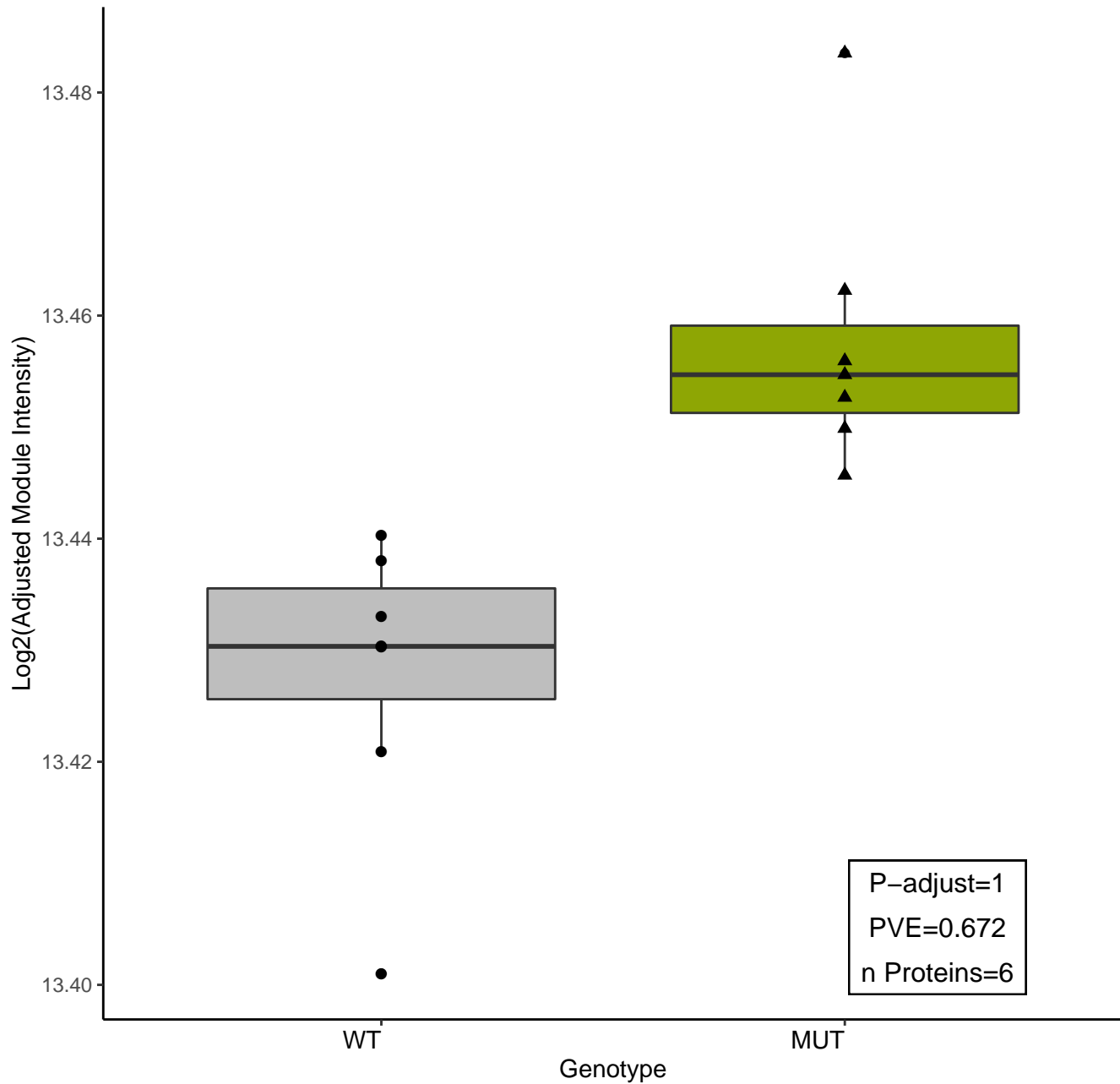
M167



M168

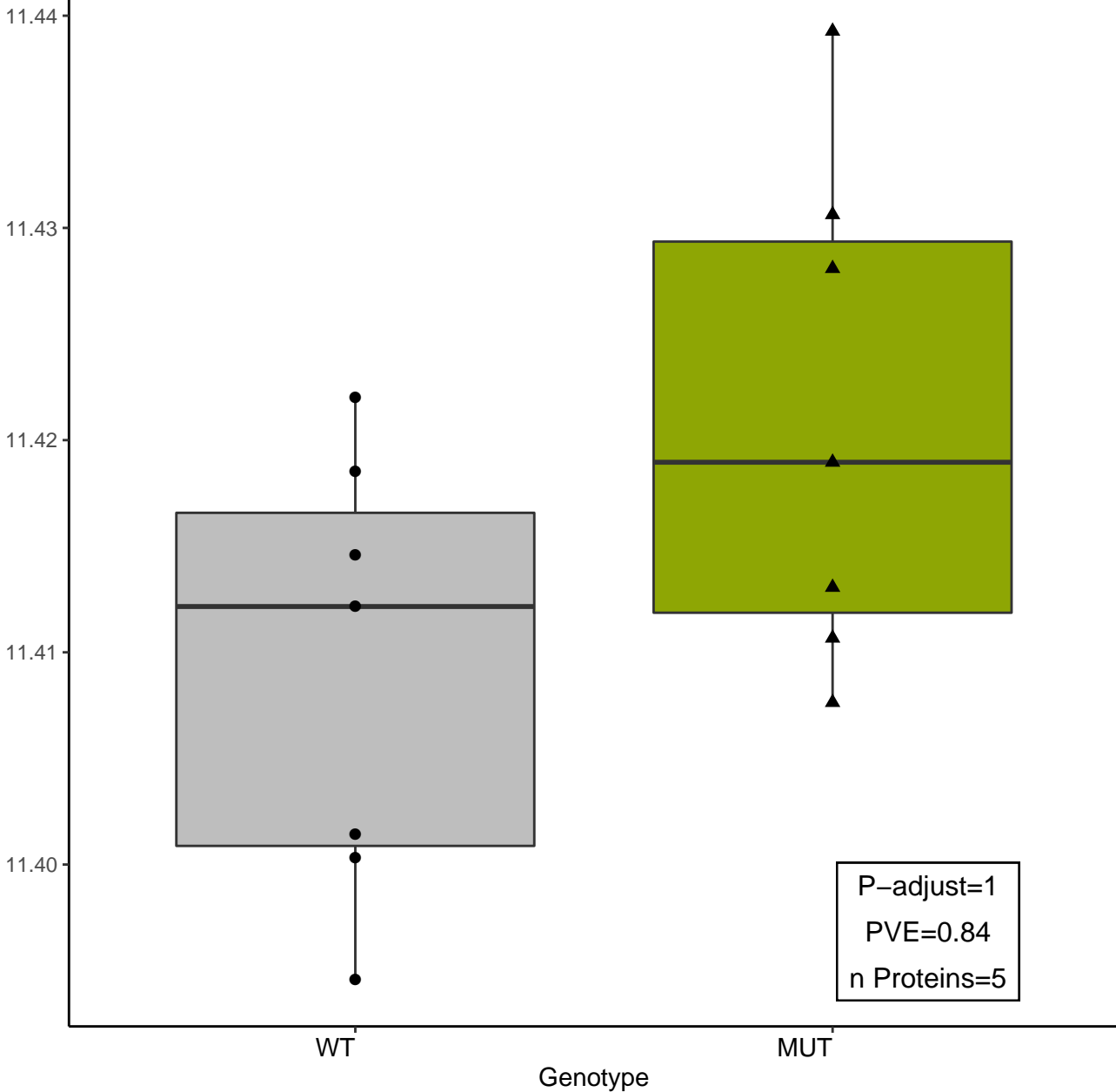


M169

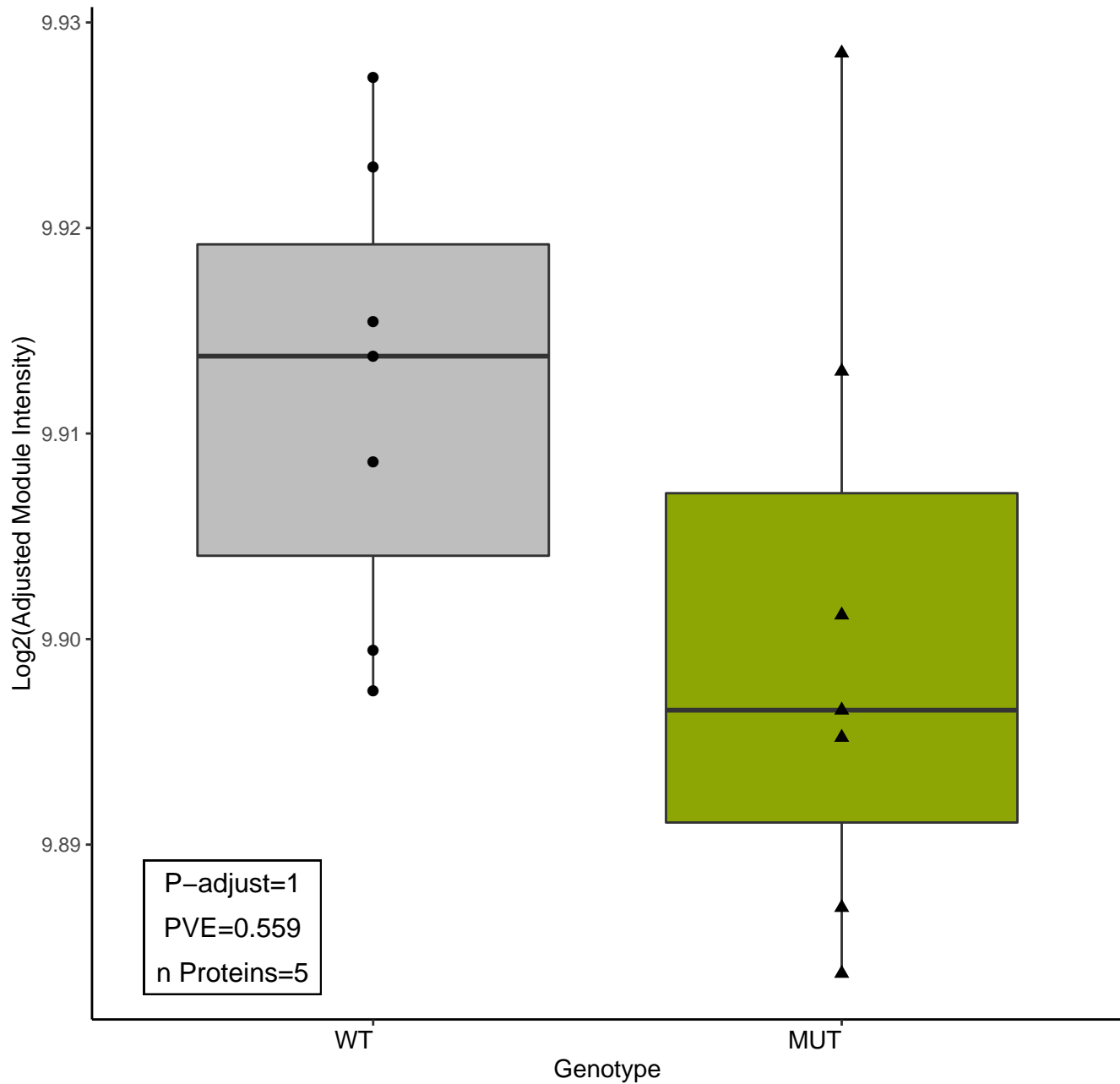


M170

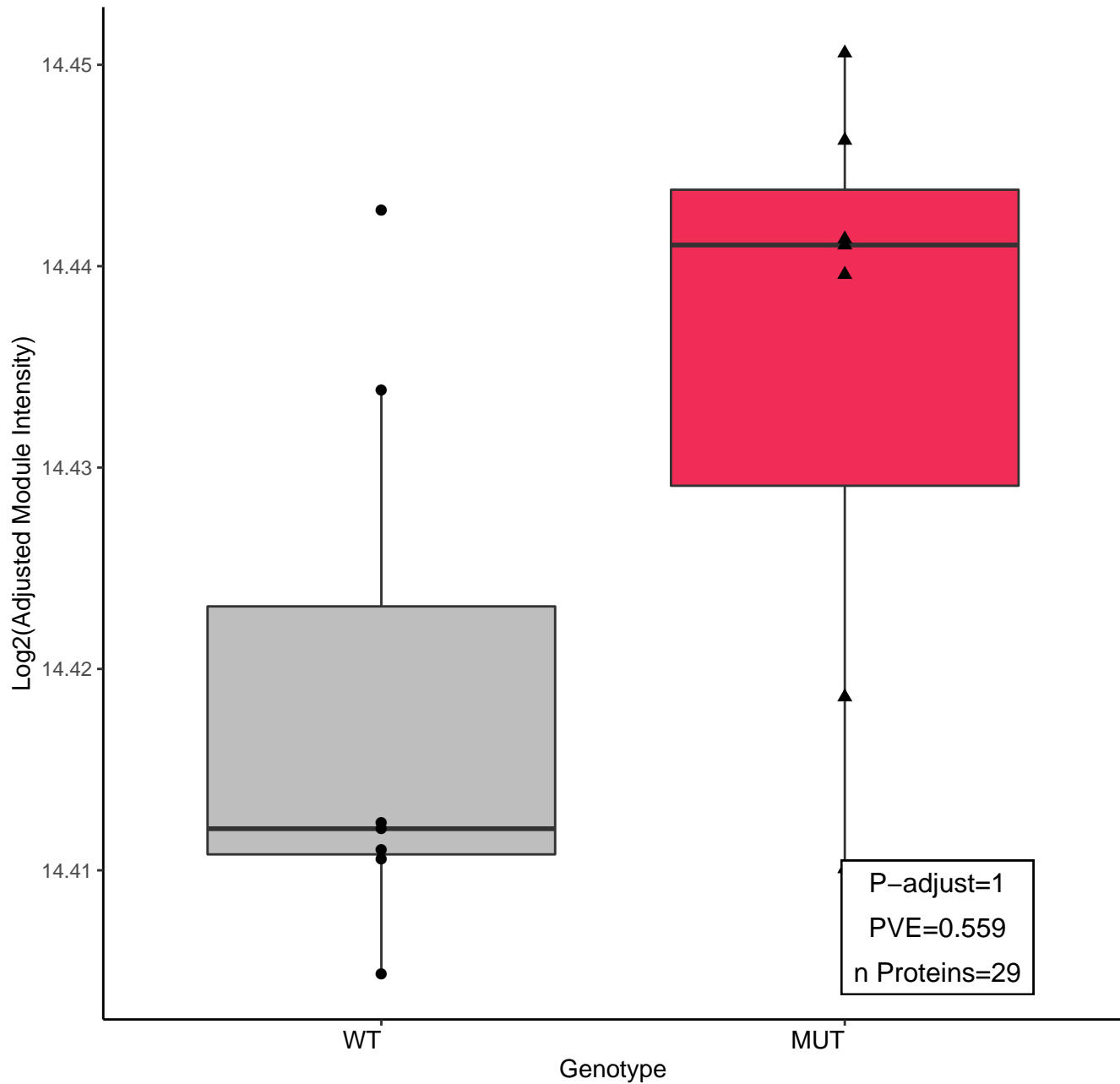
Log2(Adjusted Module Intensity)



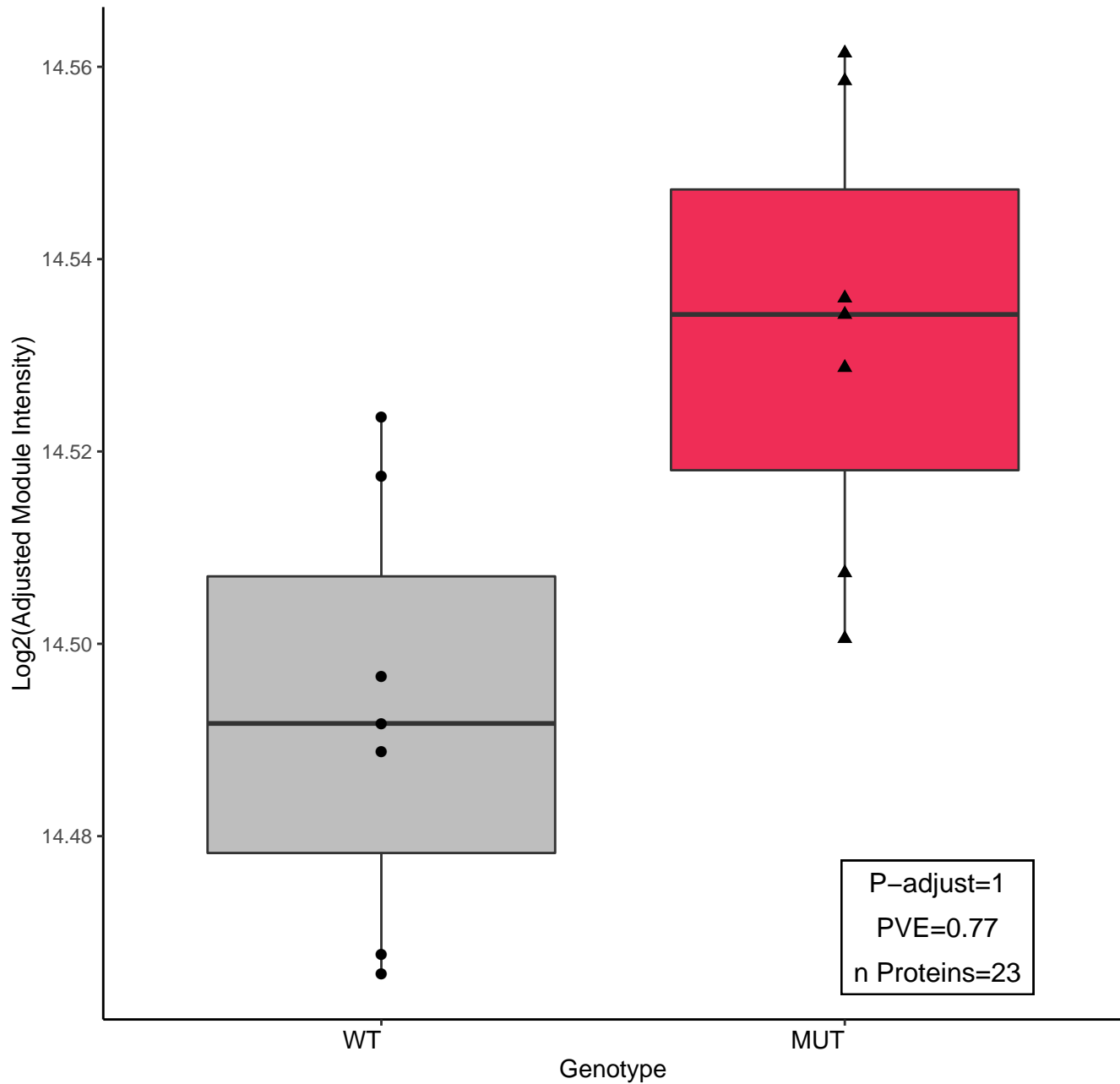
M171



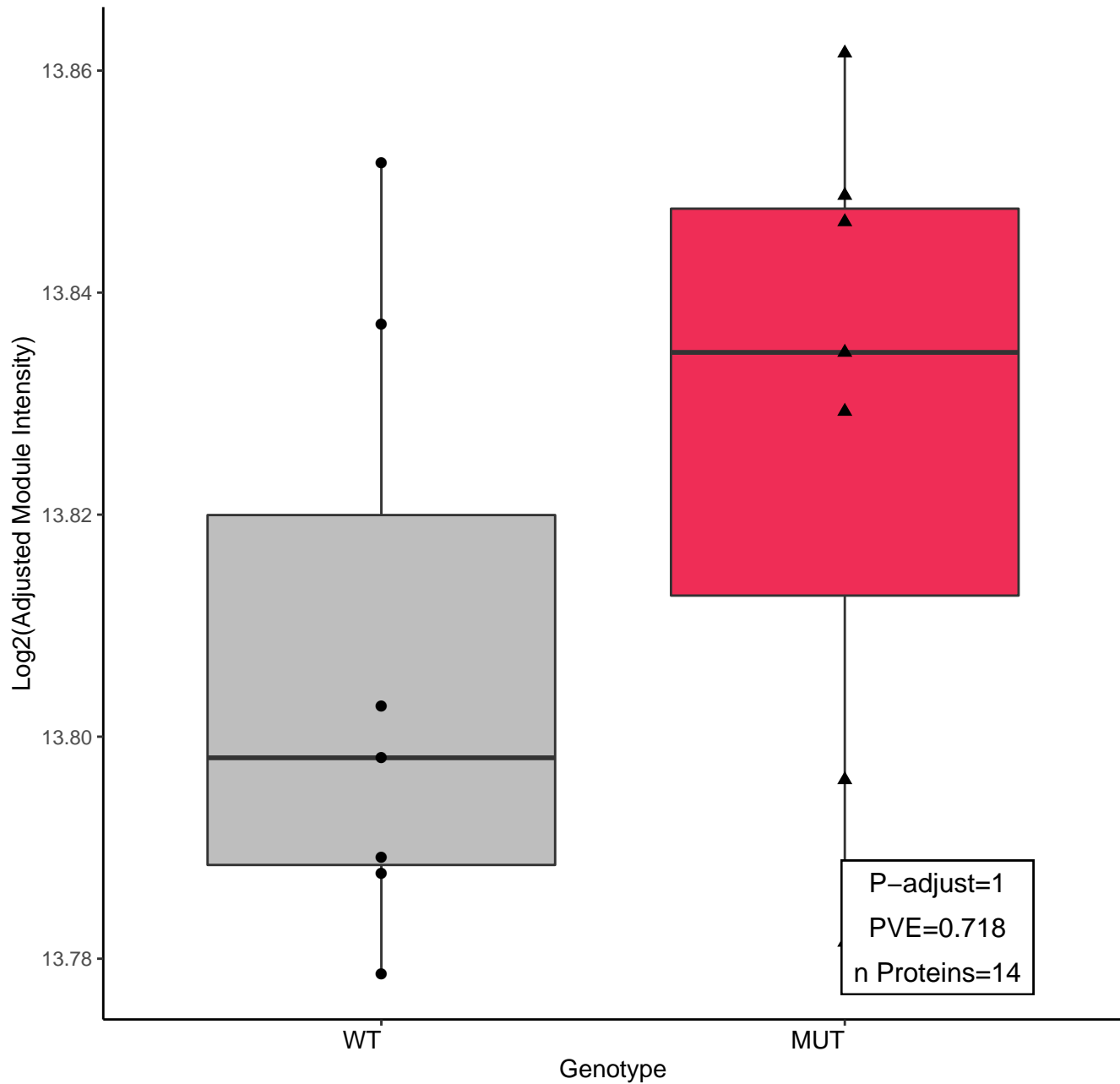
M172



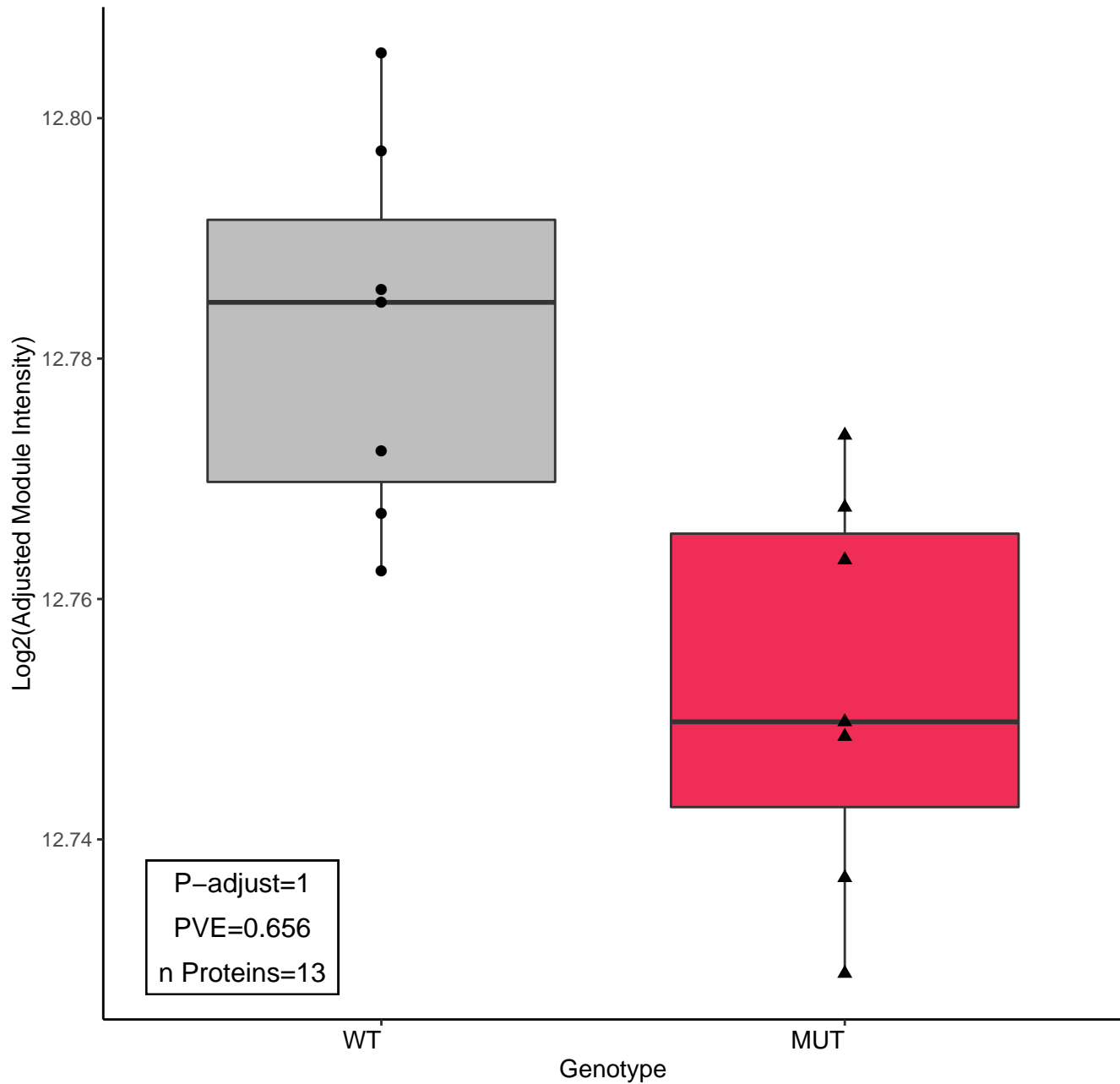
M173



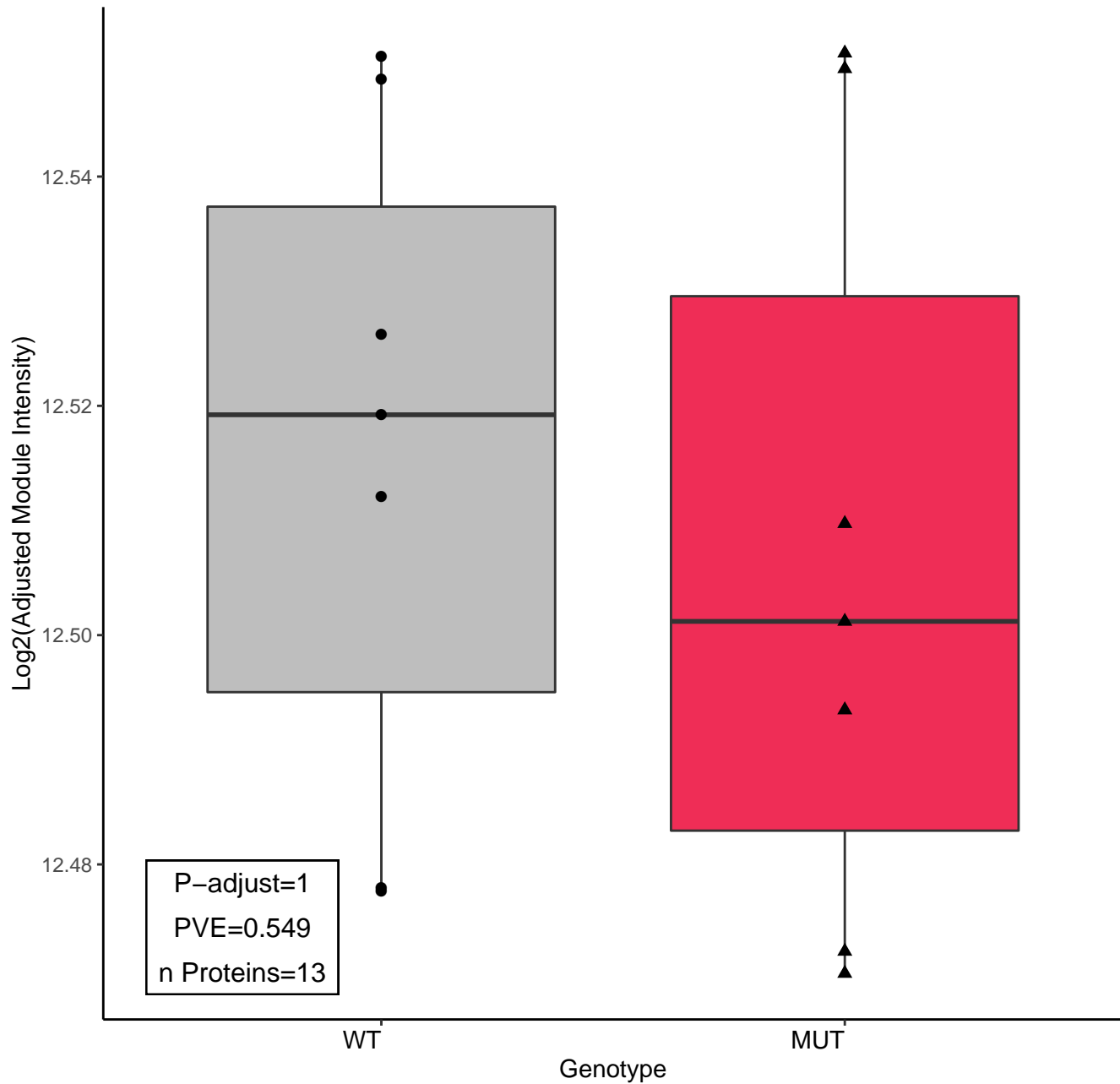
M174



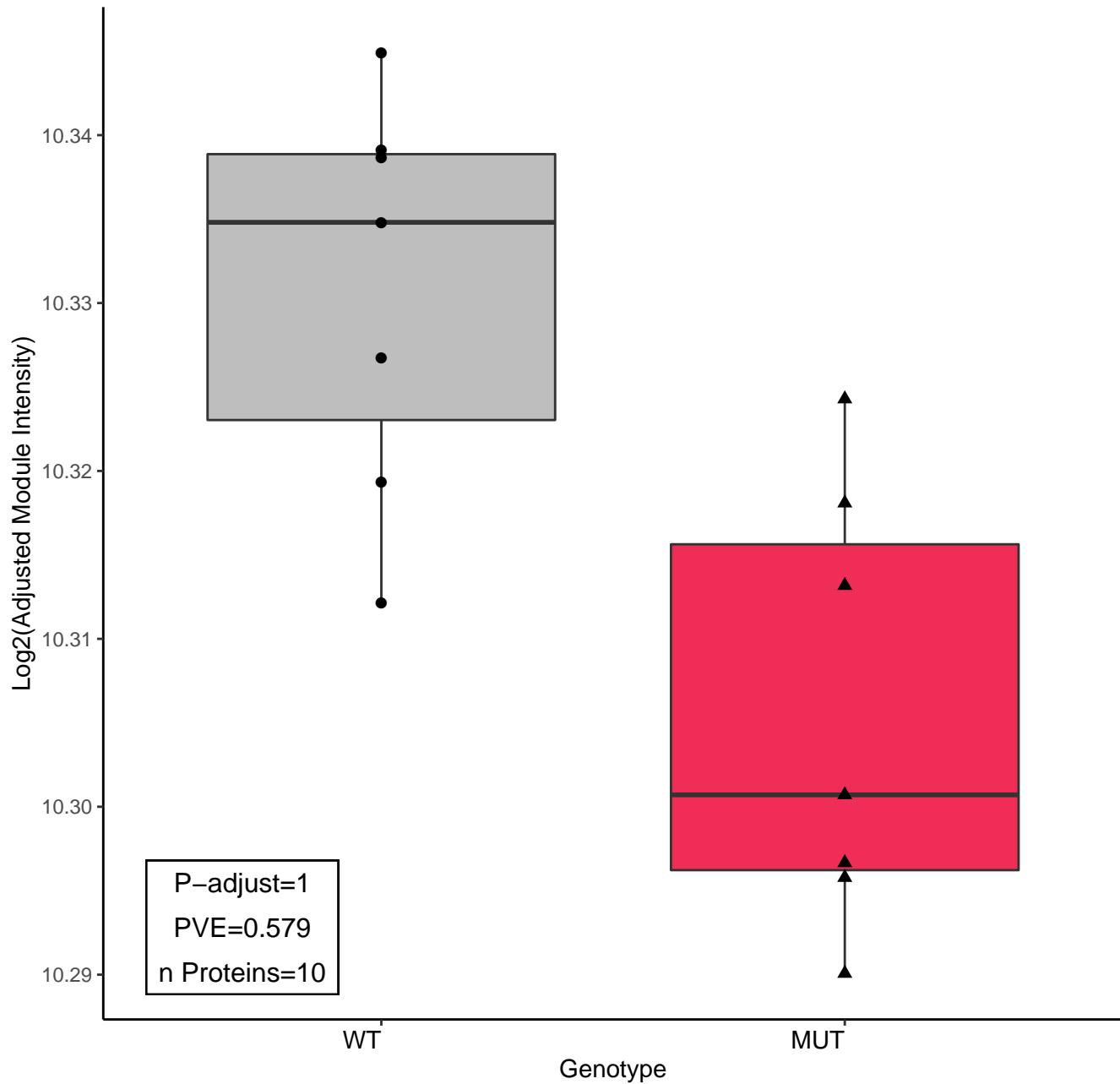
M175



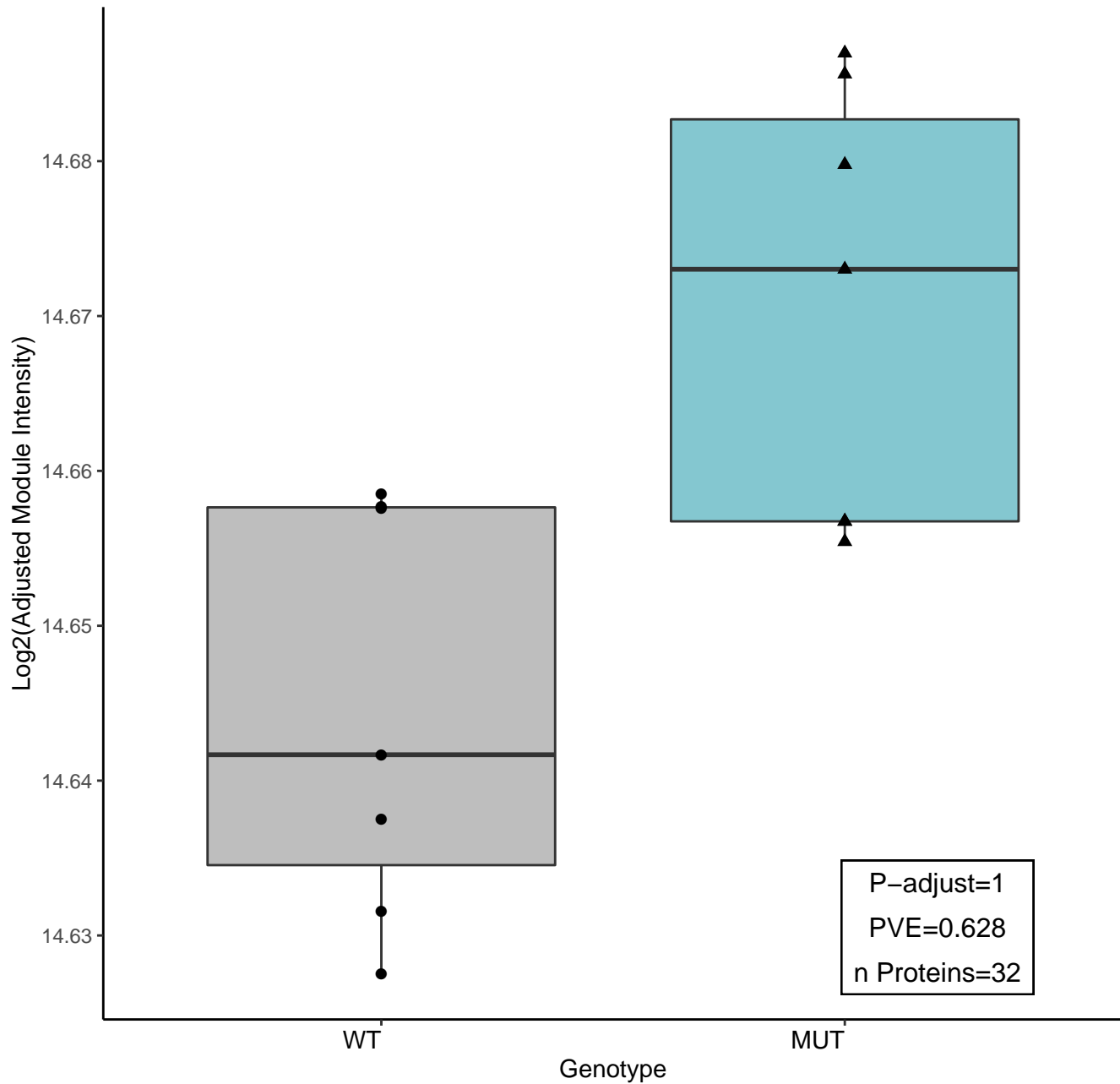
M176



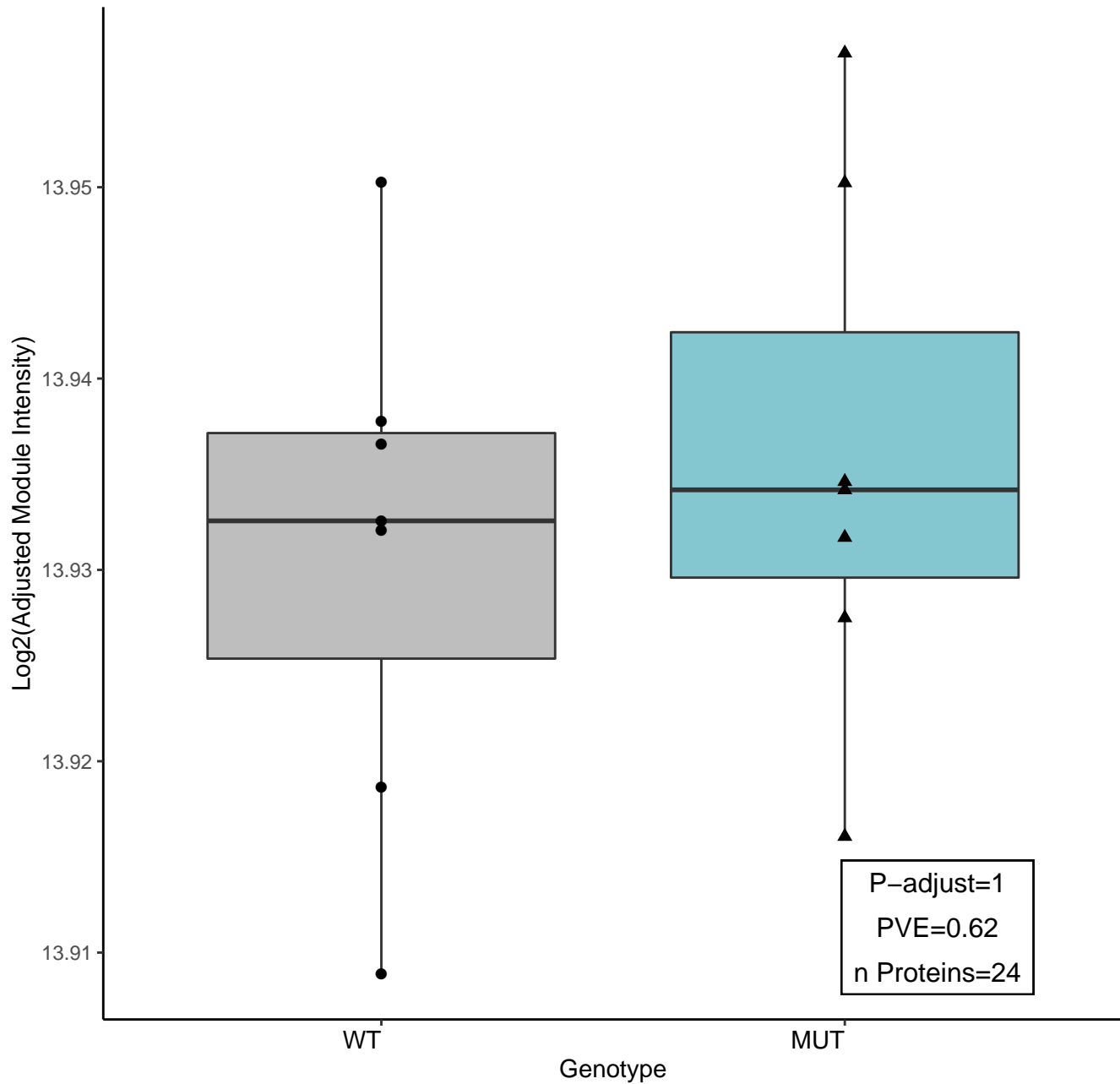
M177



M178



M179



M180

Log2(Adjusted Module Intensity)

13.94

13.92

13.90

13.88

WT

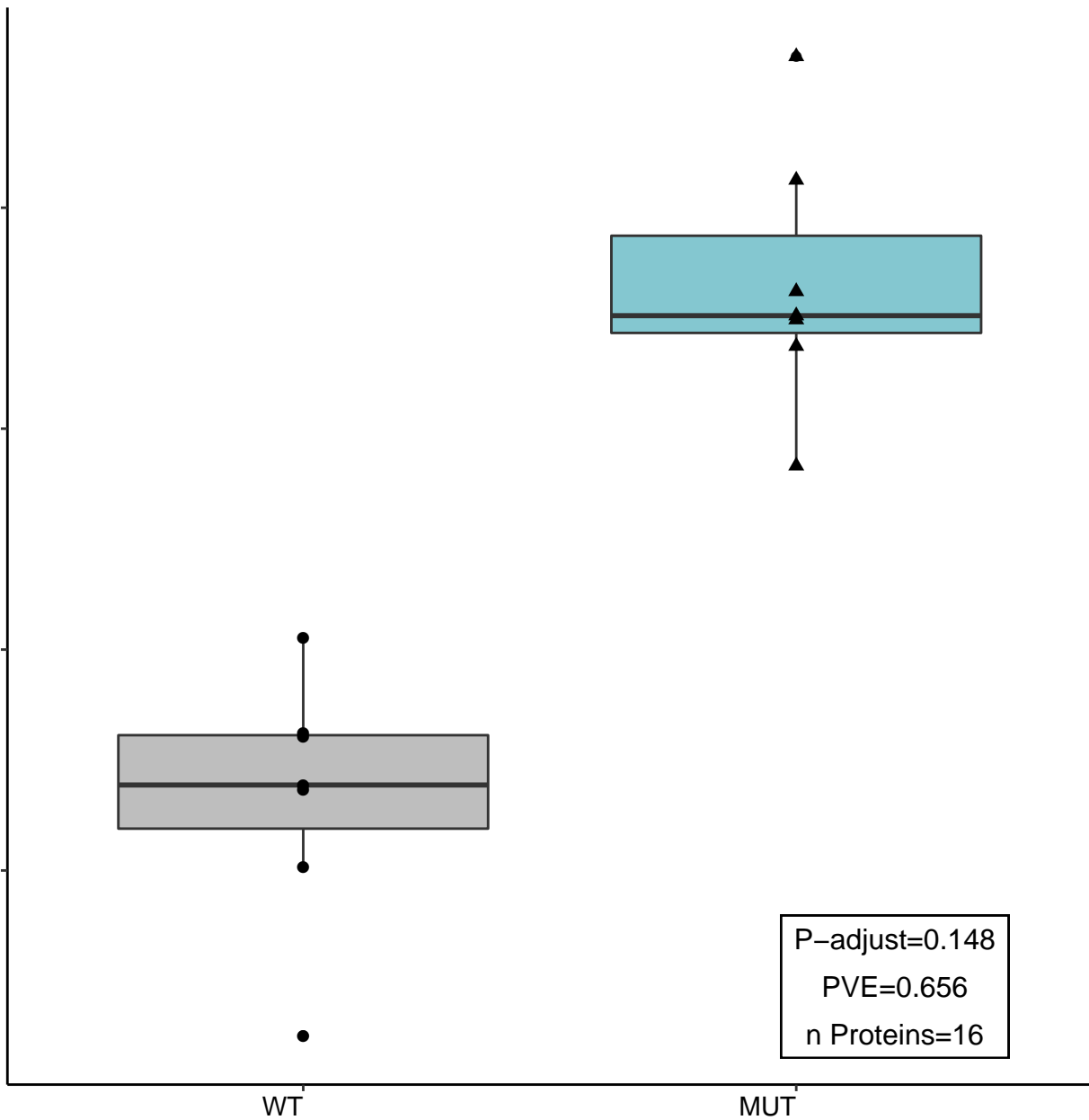
Genotype

MUT

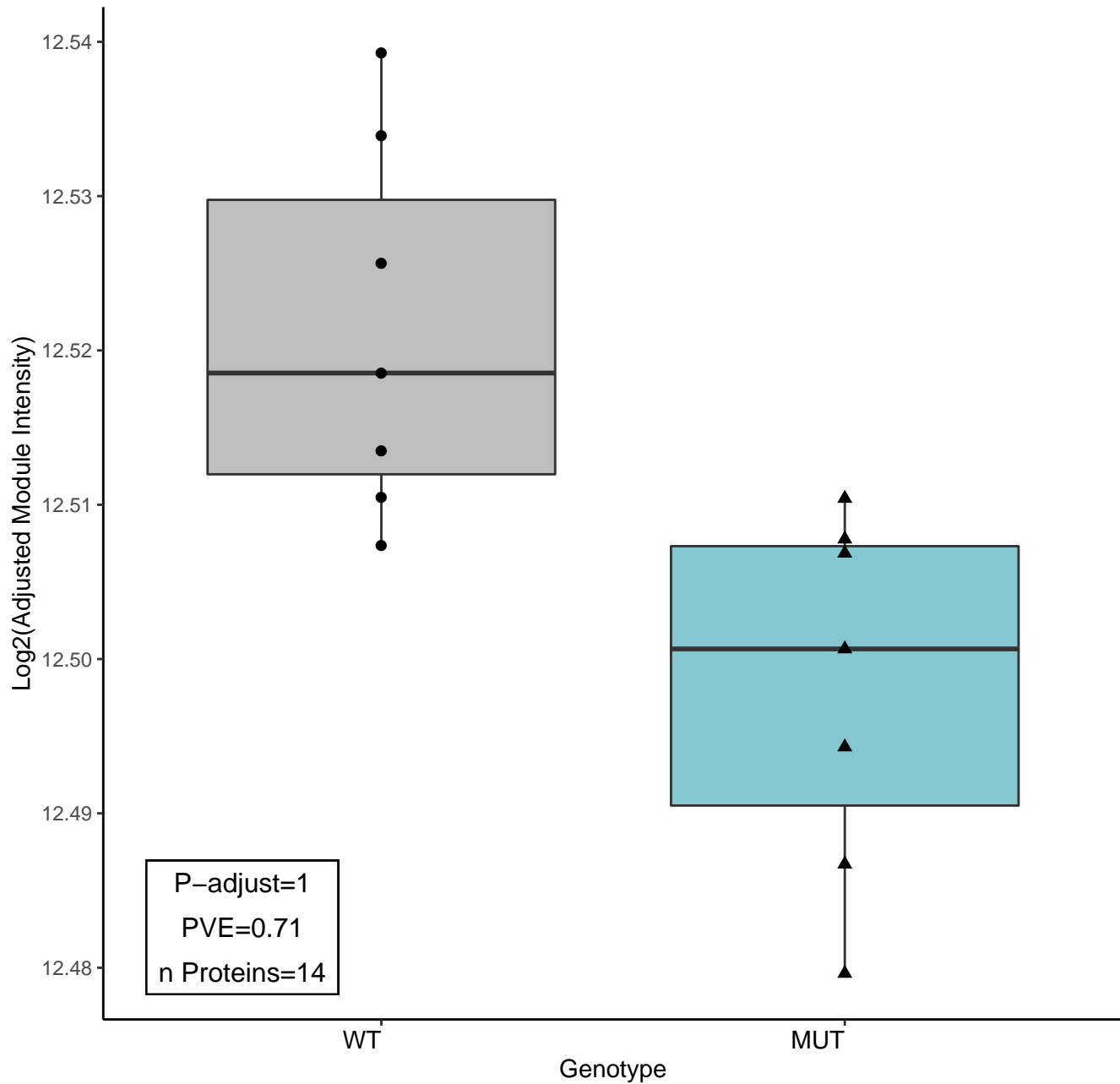
P-adjust=0.148

PVE=0.656

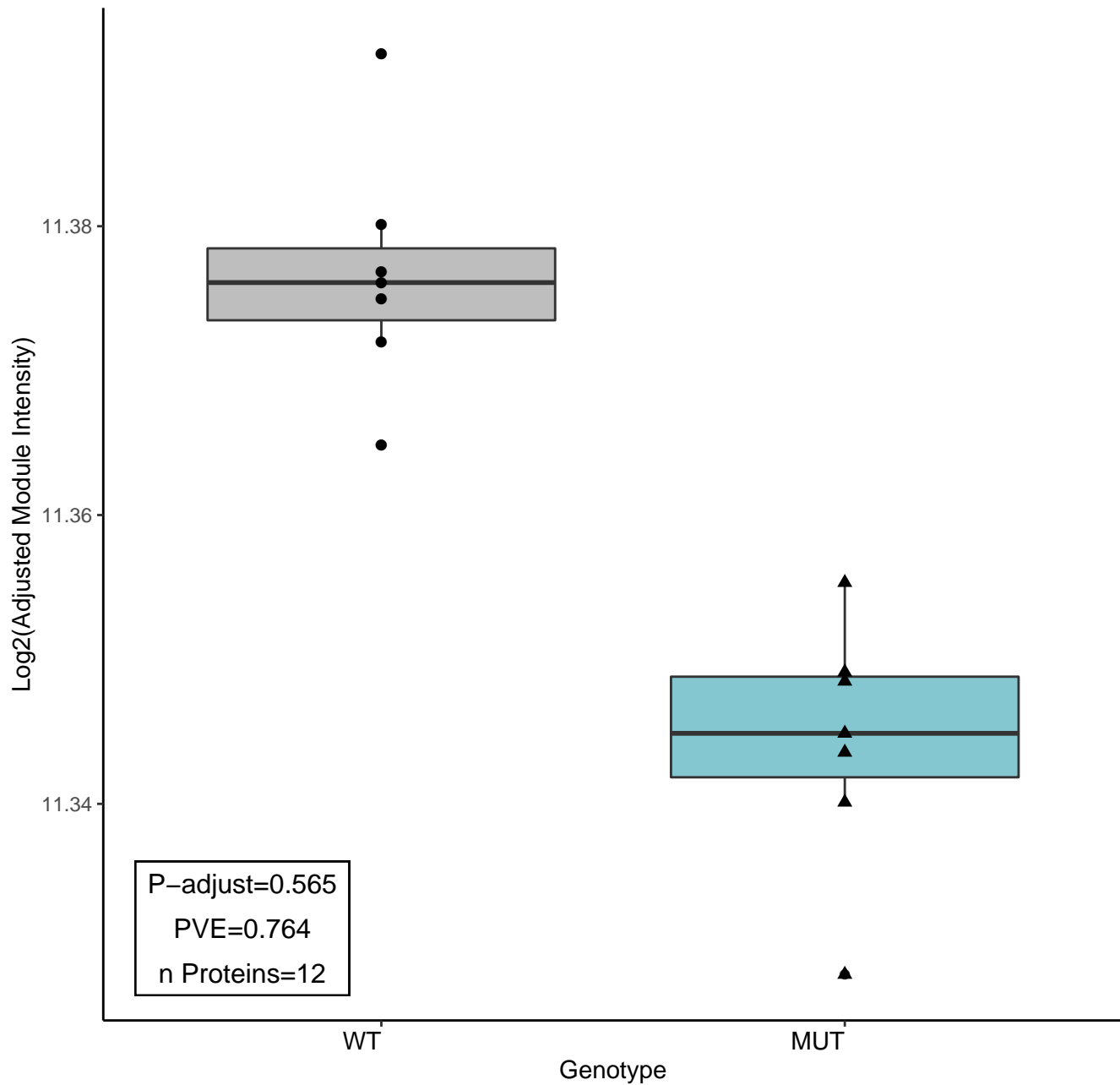
n Proteins=16



M181



M182



M183

Log2(Adjusted Module Intensity)

12.04

12.02

12.00

11.98

WT

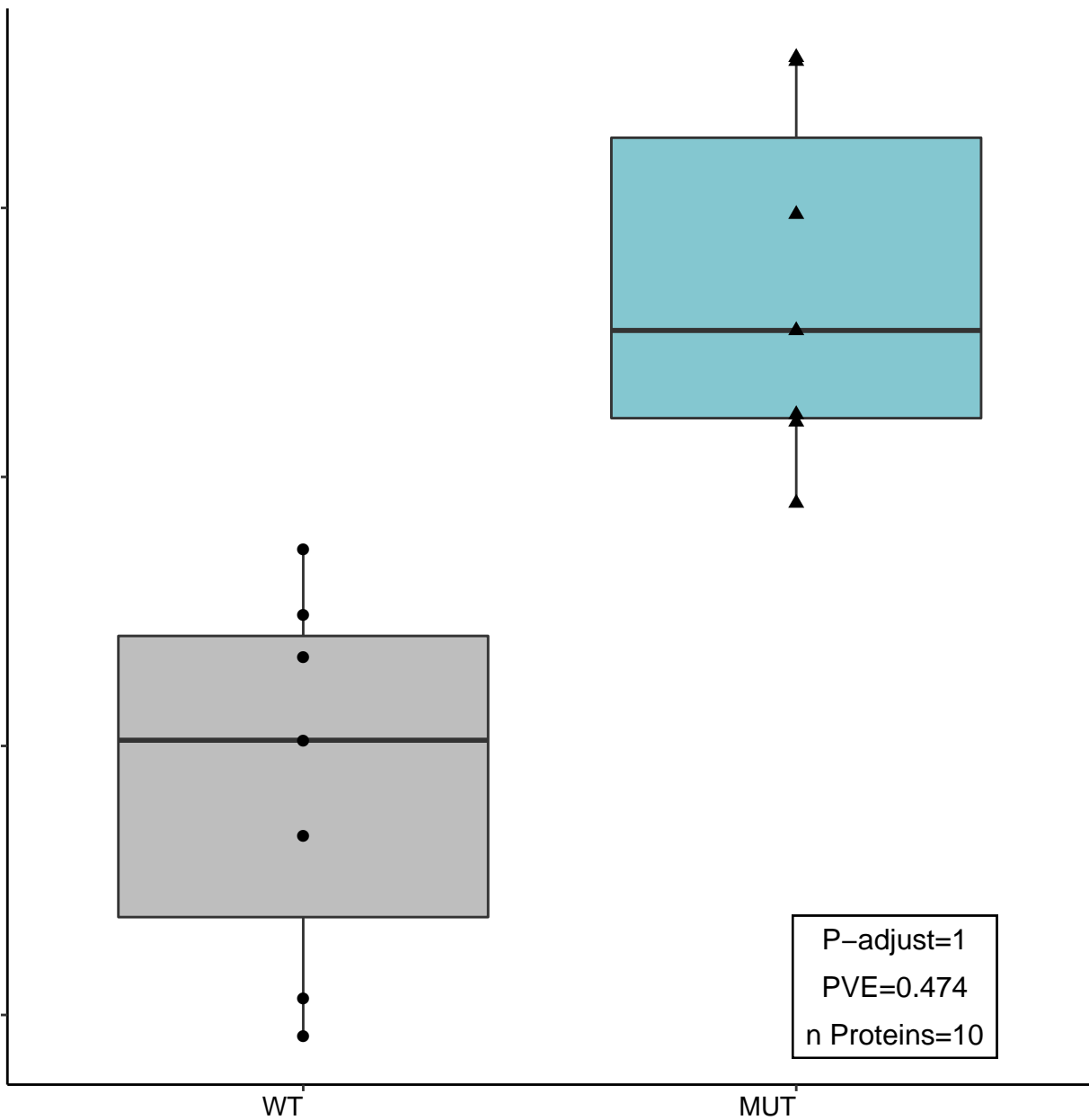
MUT

Genotype

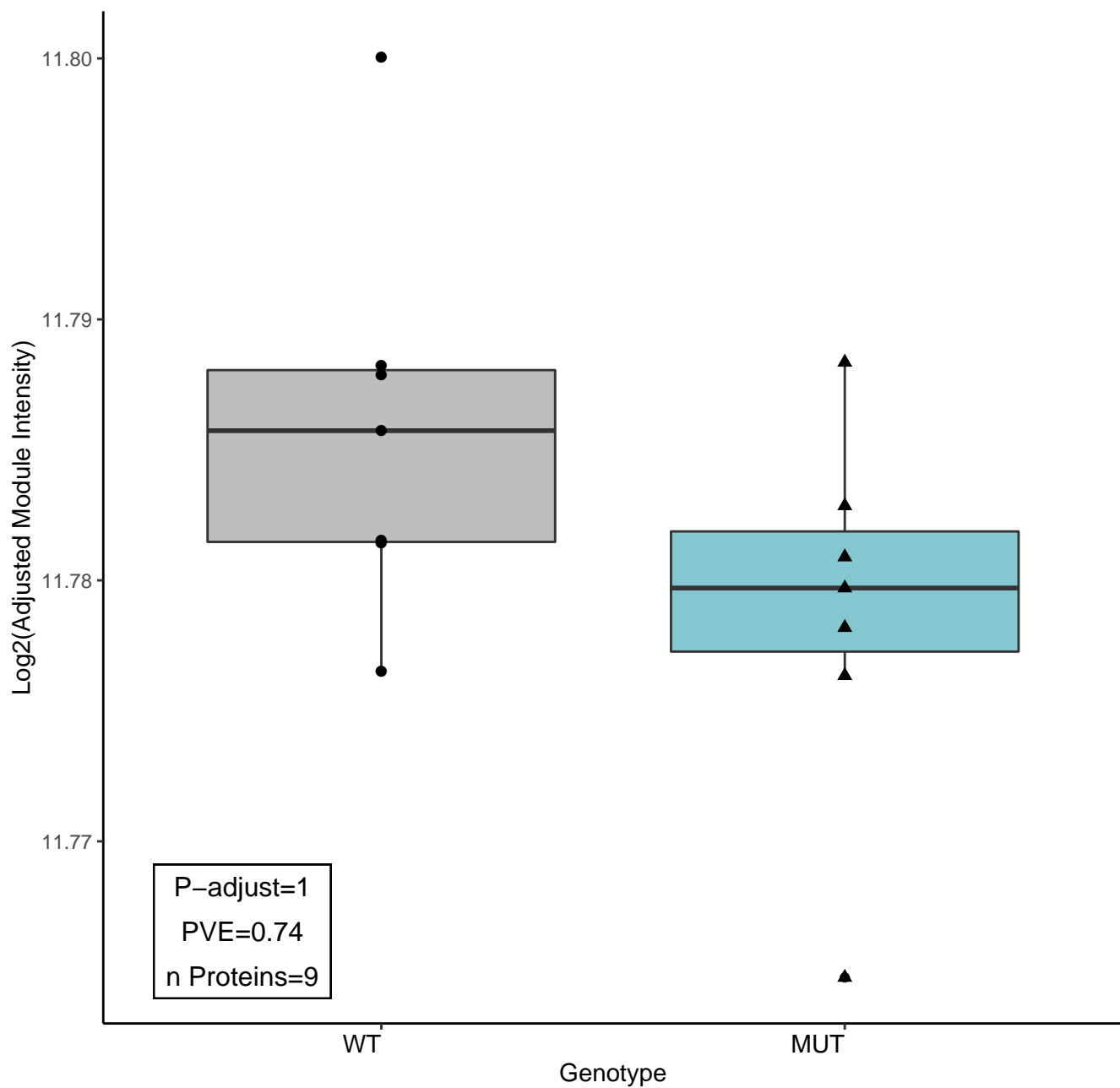
P-adjust=1

PVE=0.474

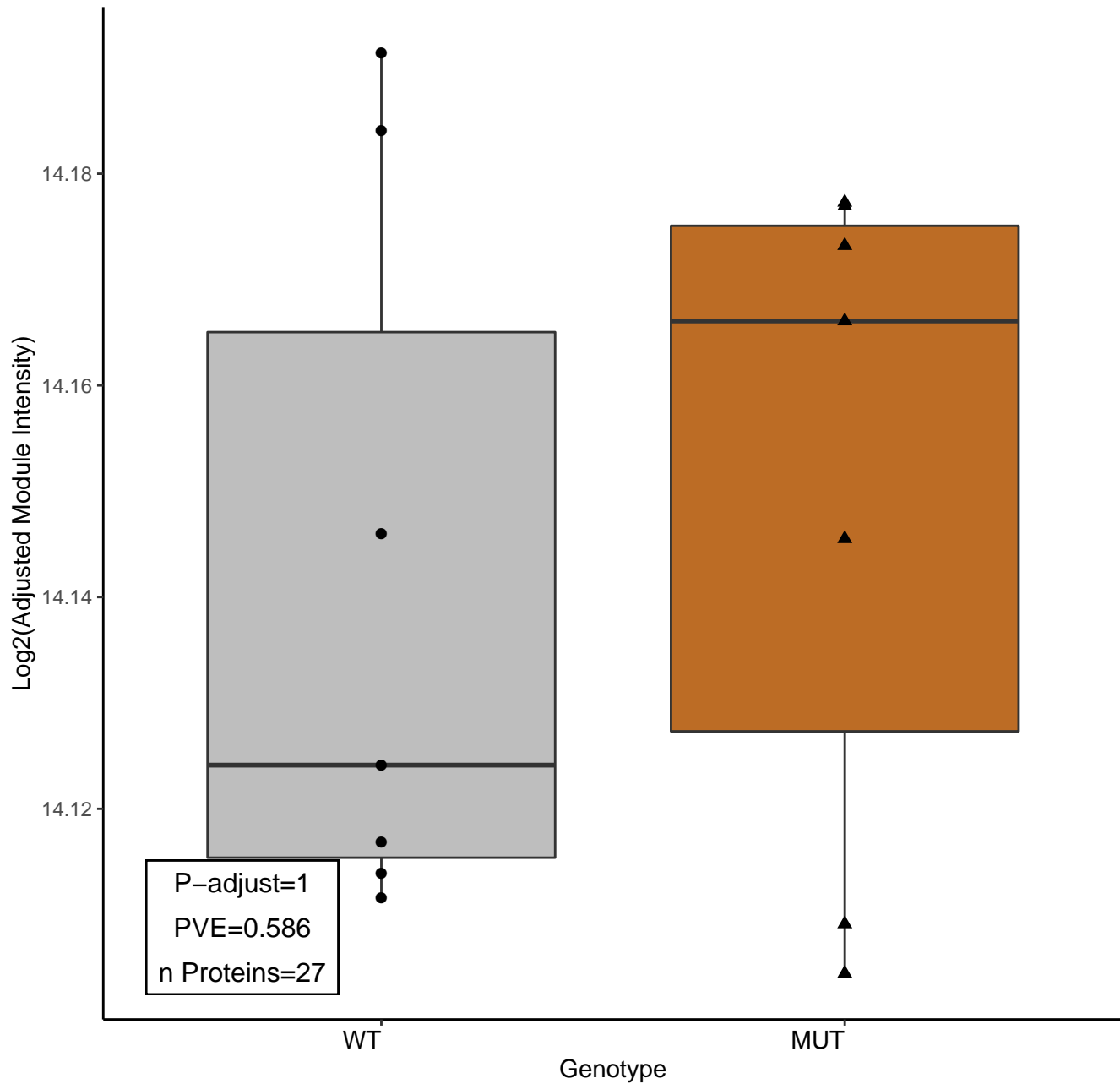
n Proteins=10



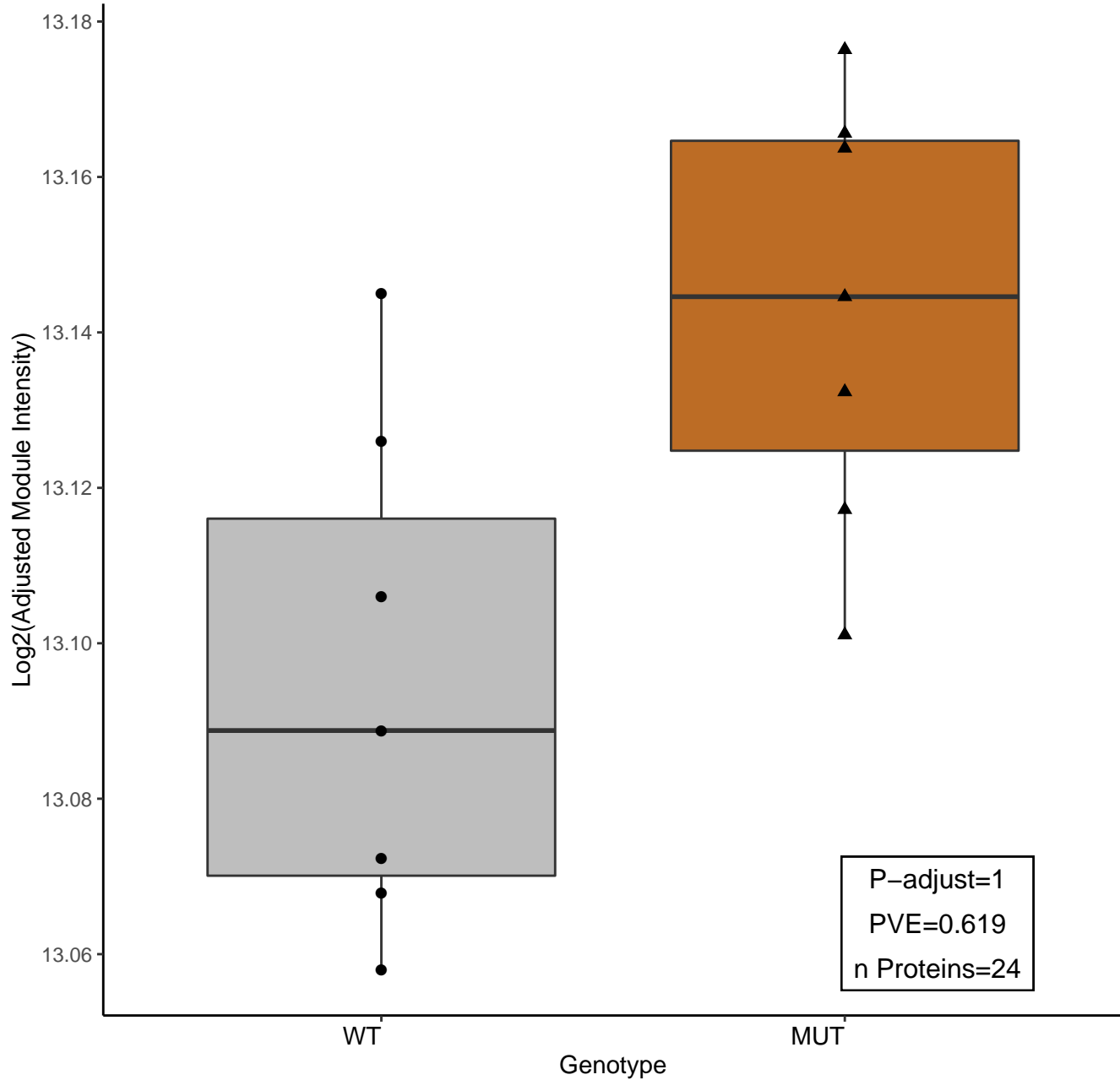
M184



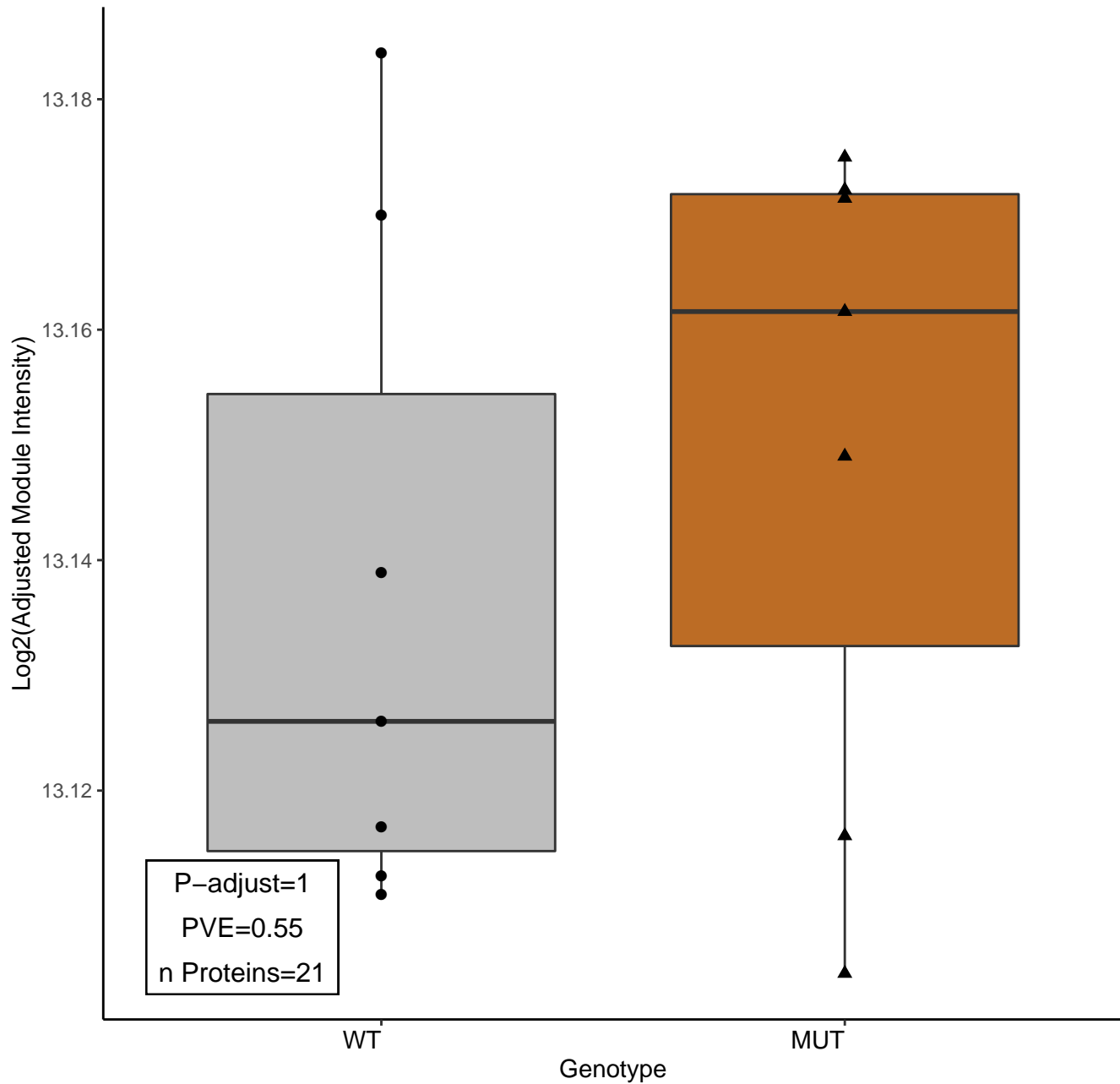
M185



M186

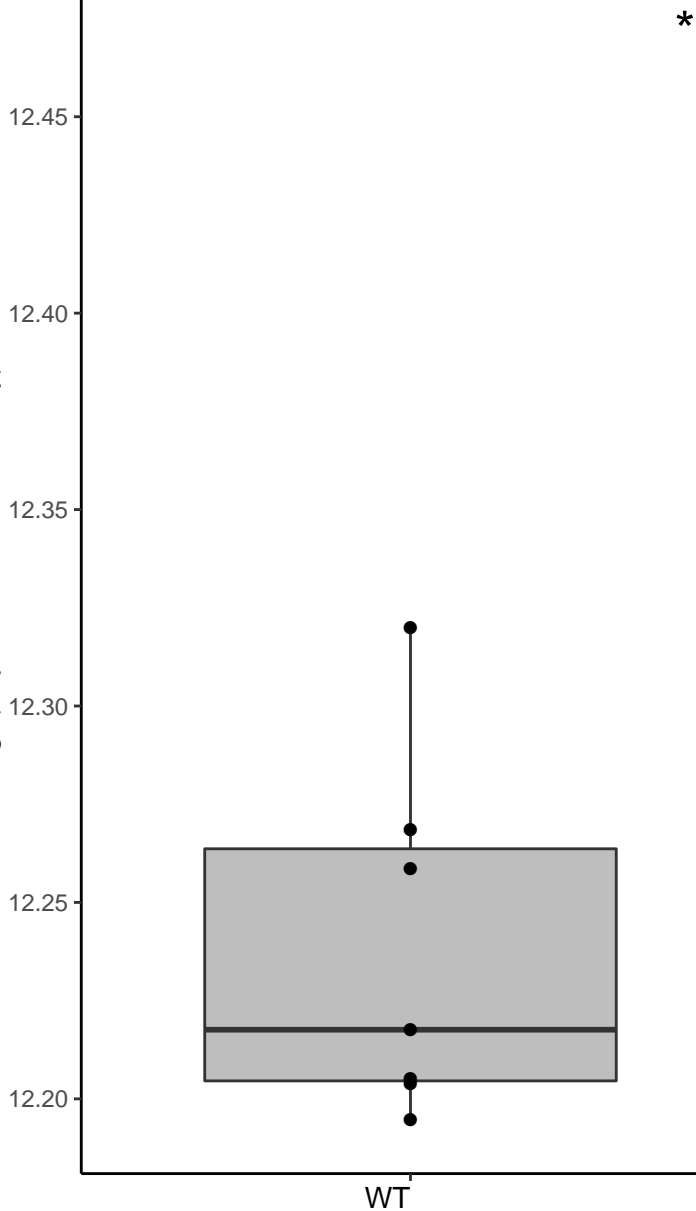


M187



M188

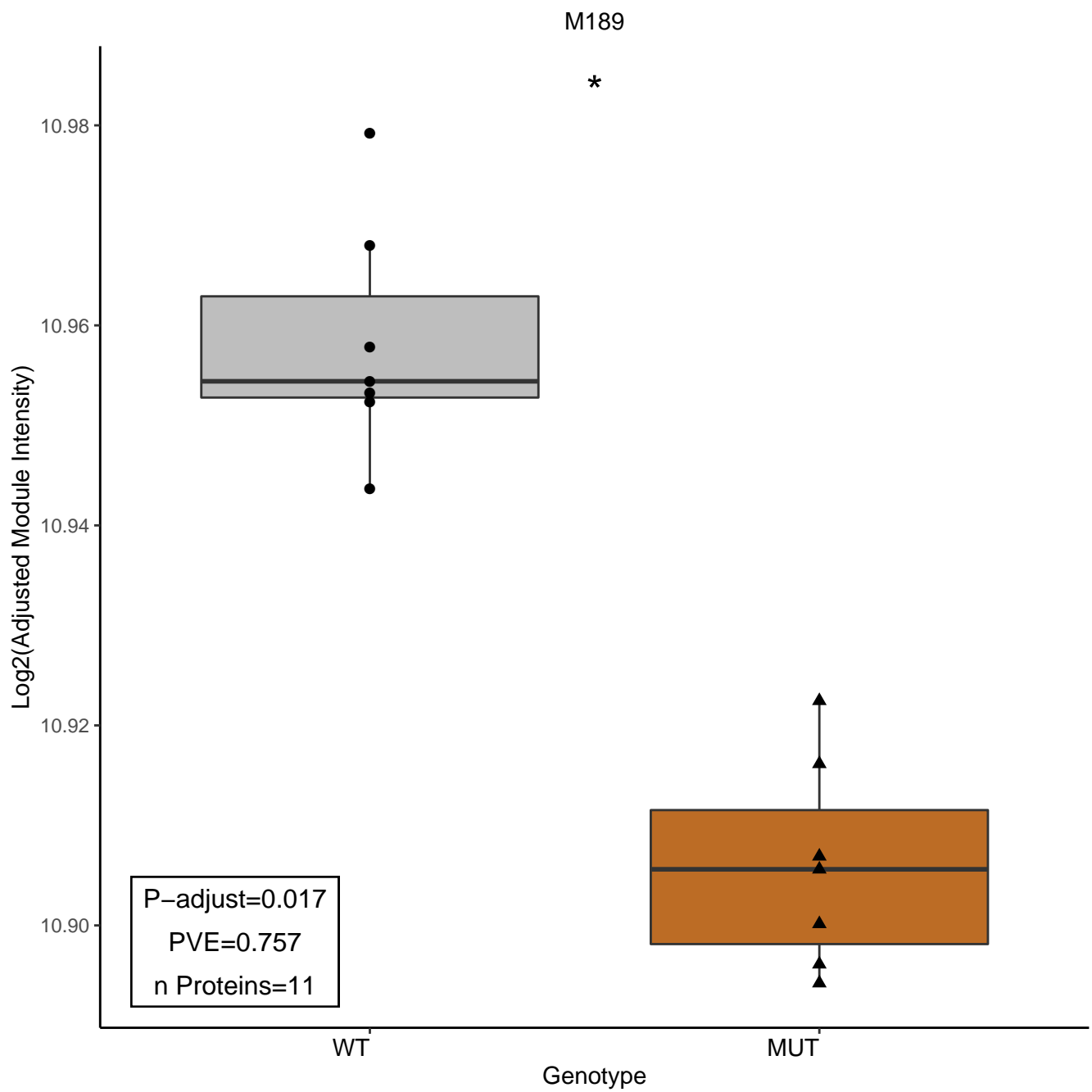
Log2(Adjusted Module Intensity)



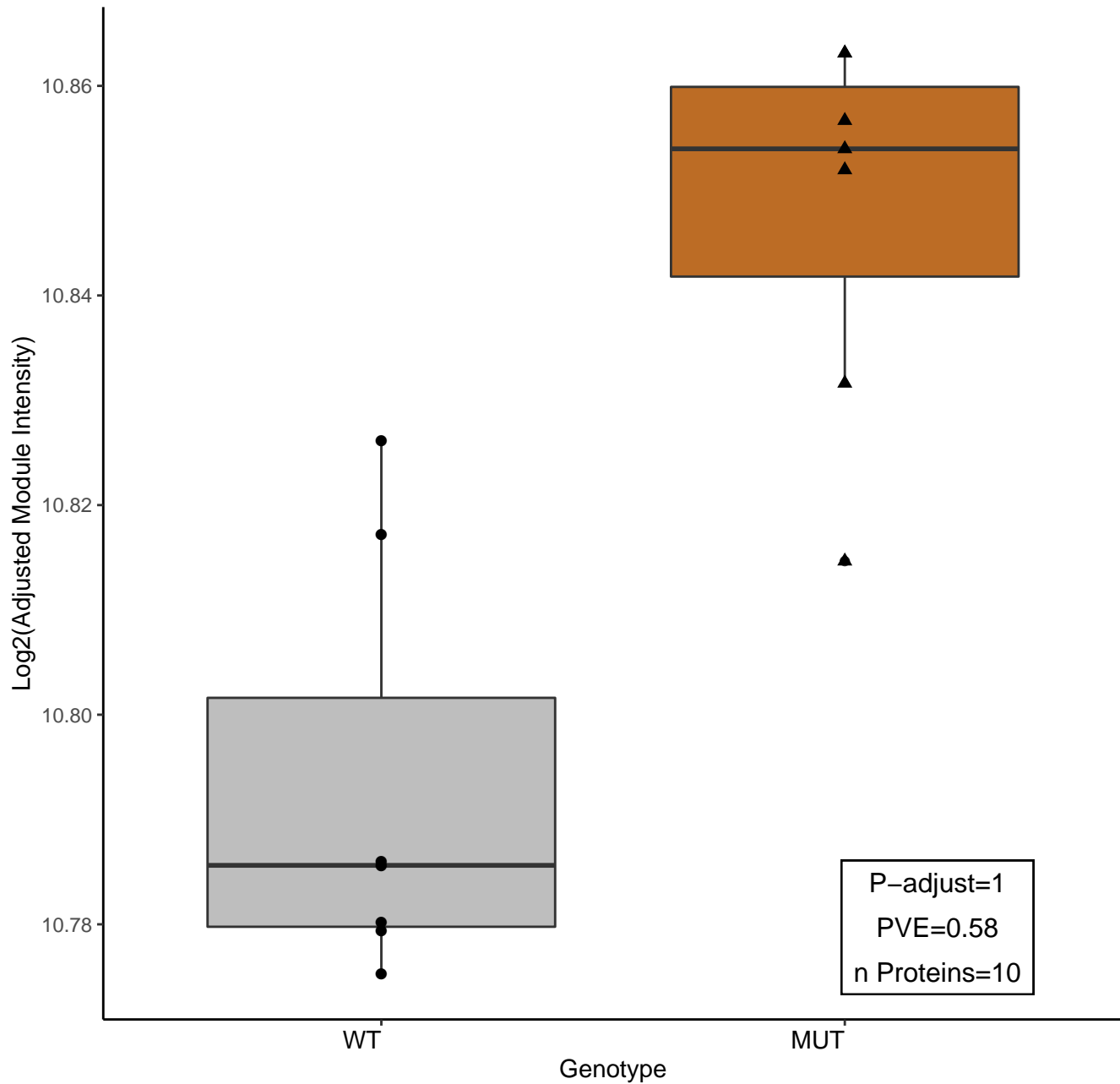
MUT

Genotype

P-adjust=0.026
PVE=0.61
n Proteins=17



M190



M191

Log2(Adjusted Module Intensity)

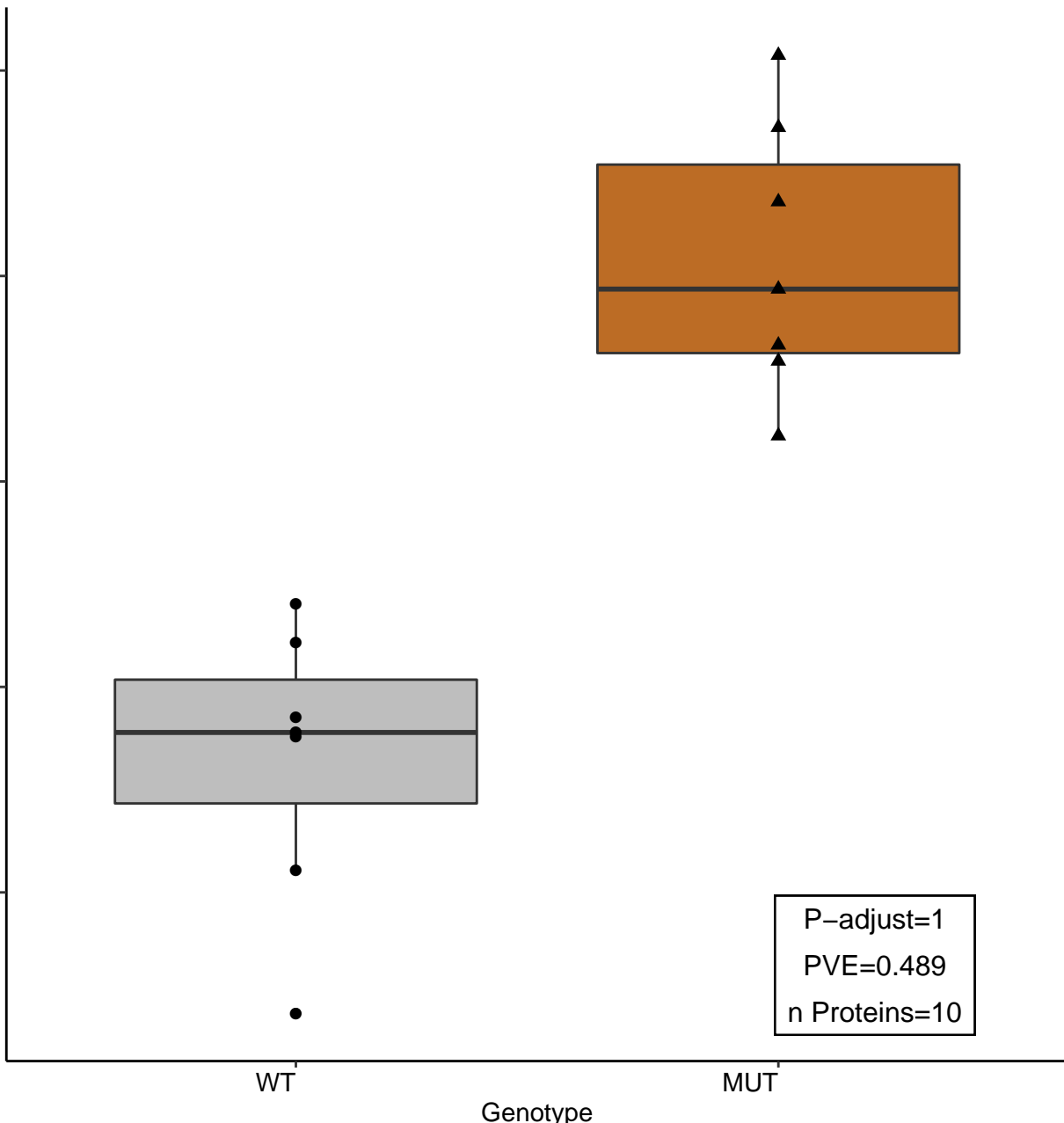
11.36
11.35
11.34
11.33
11.32

WT

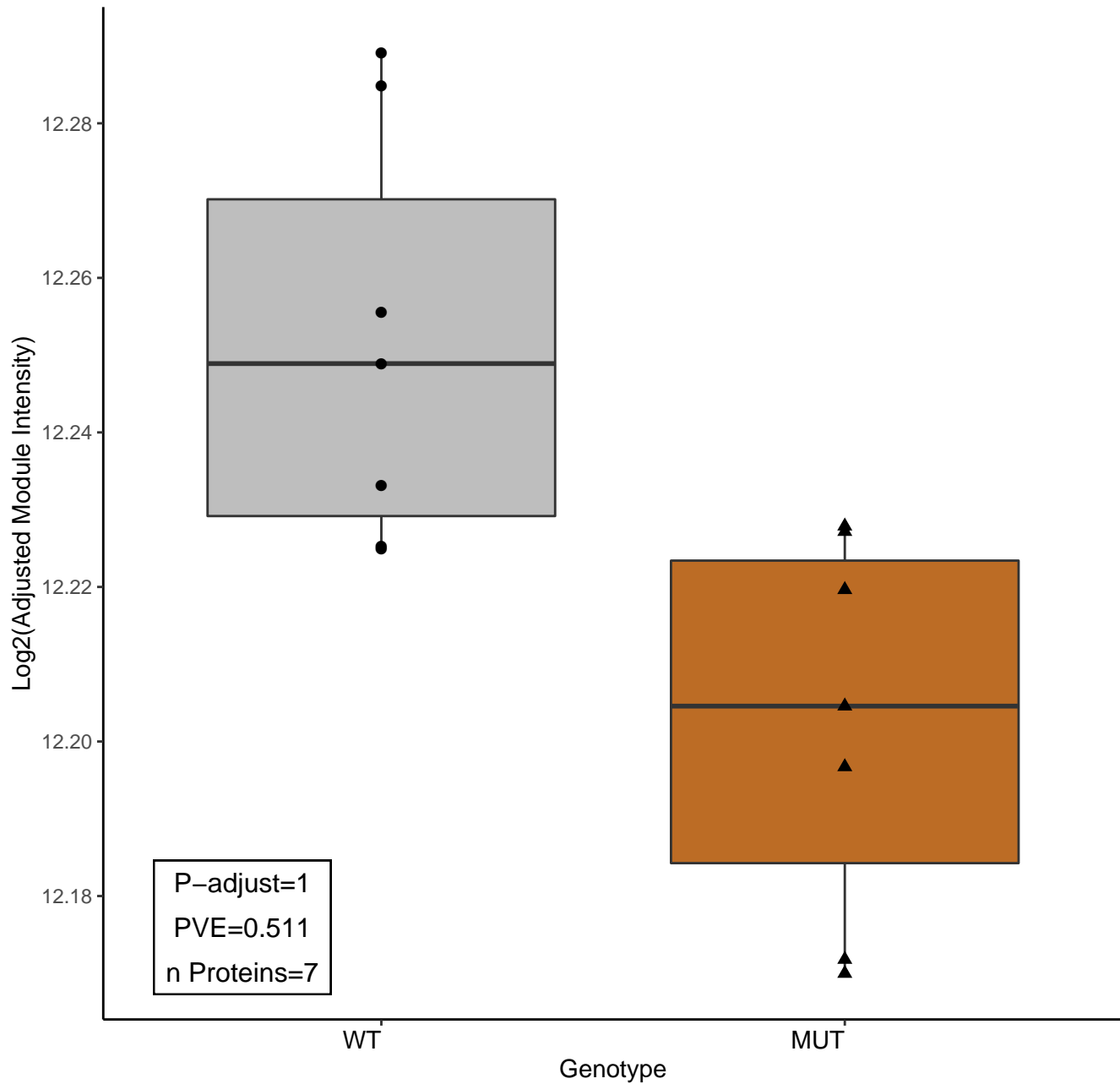
MUT

Genotype

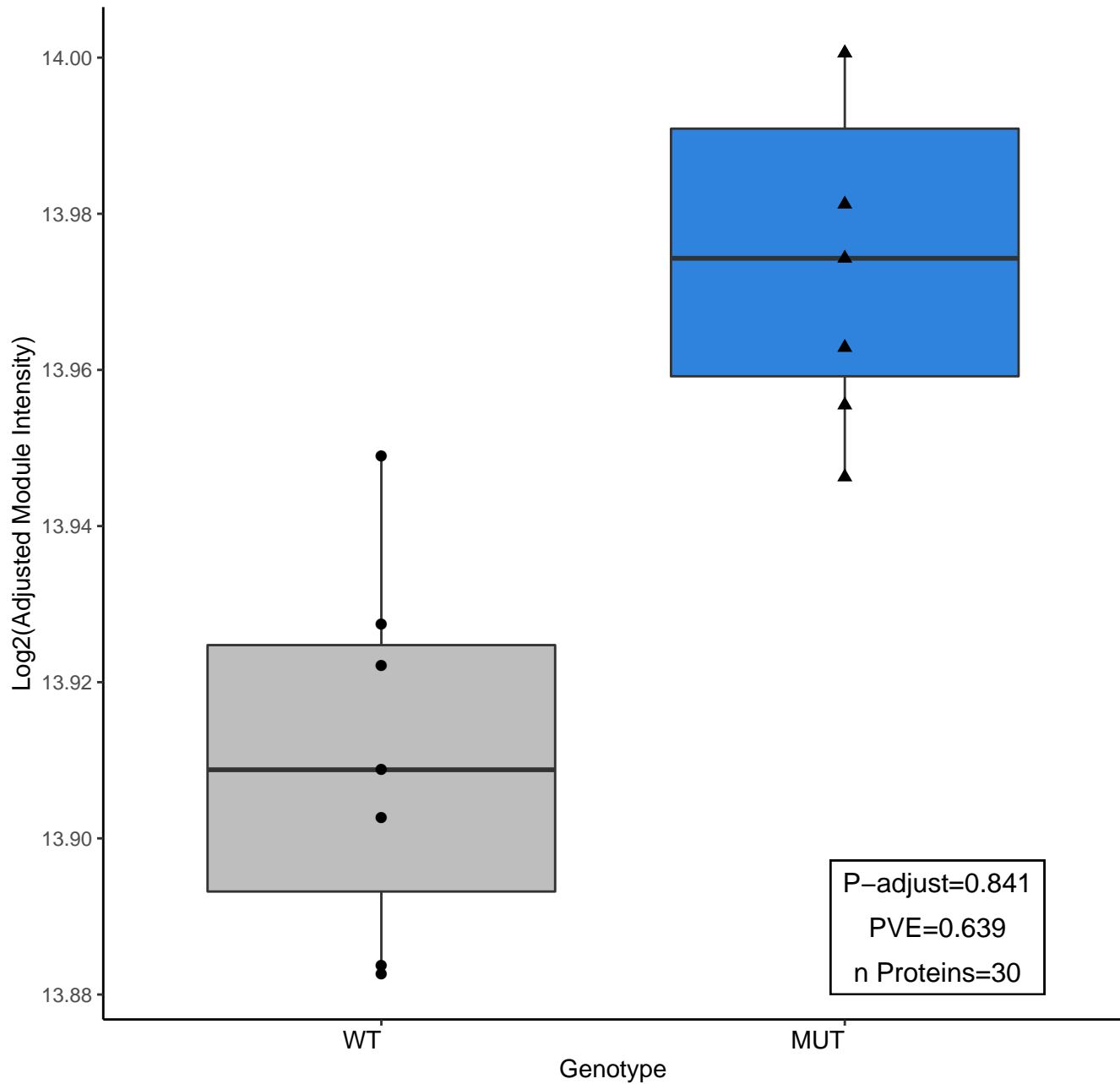
P-adjust=1
PVE=0.489
n Proteins=10



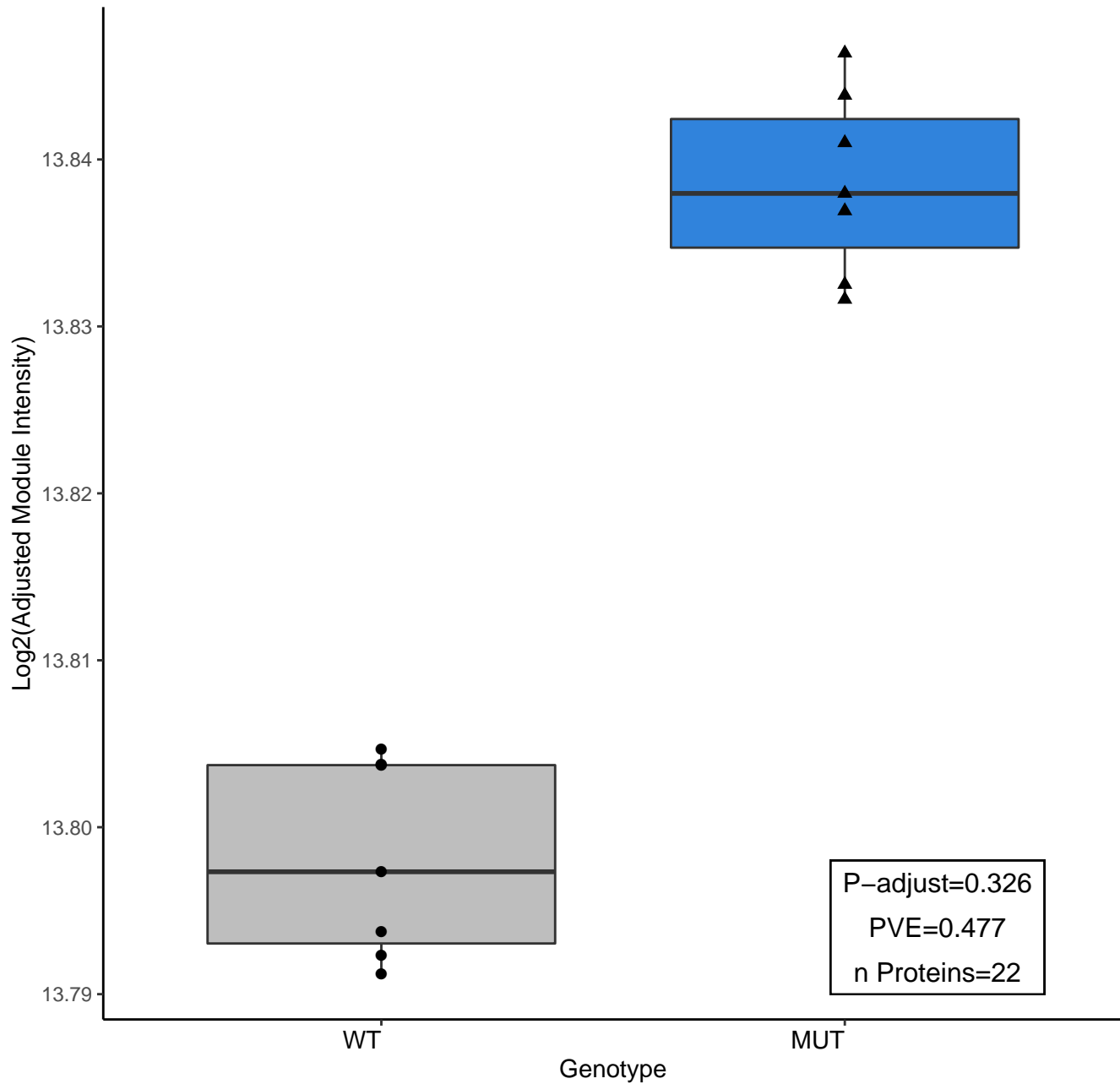
M192



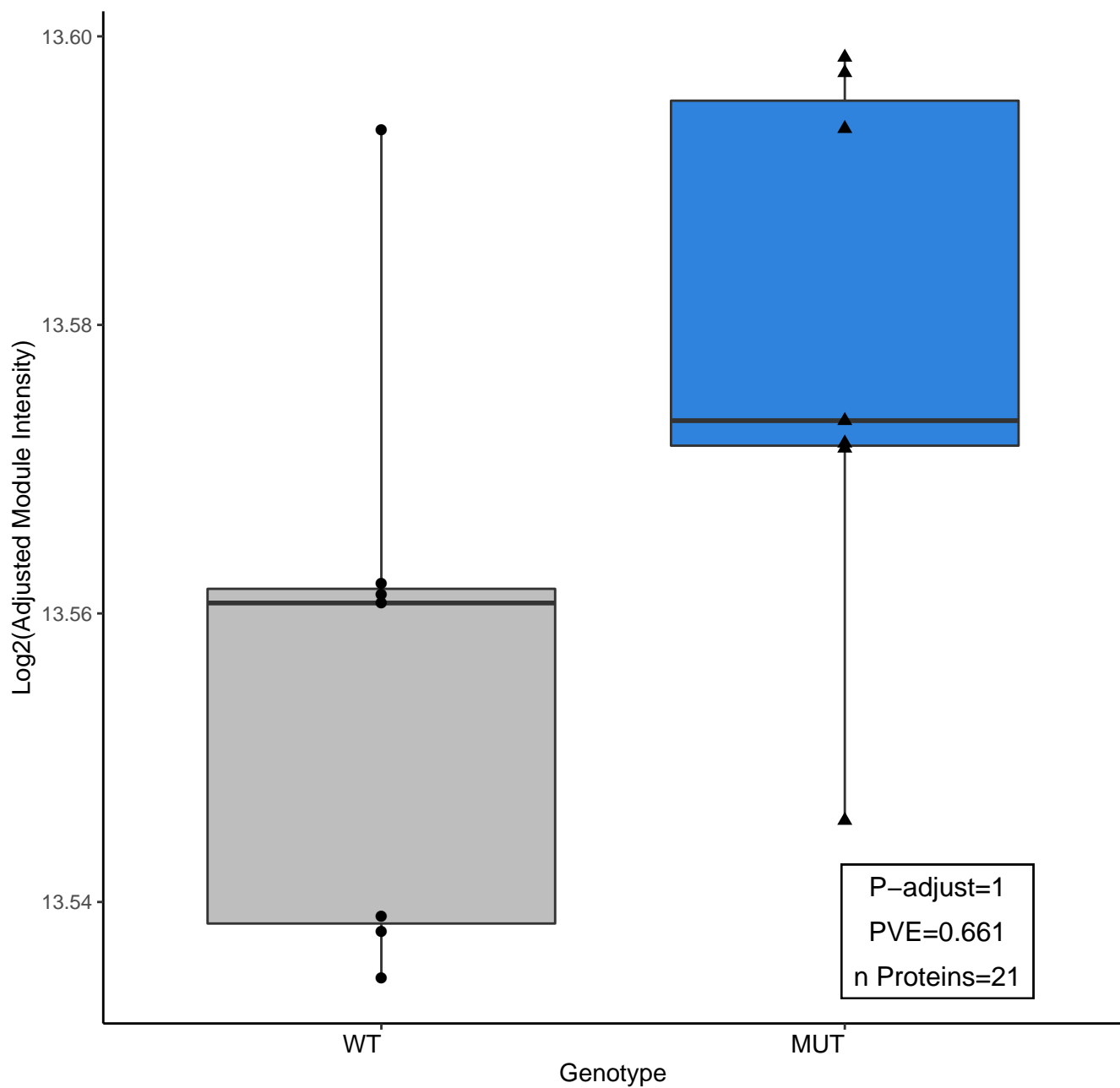
M193



M194



M195



M196

**

Log2(Adjusted Module Intensity)

12.40

12.35

12.30

P-adjust=0.003

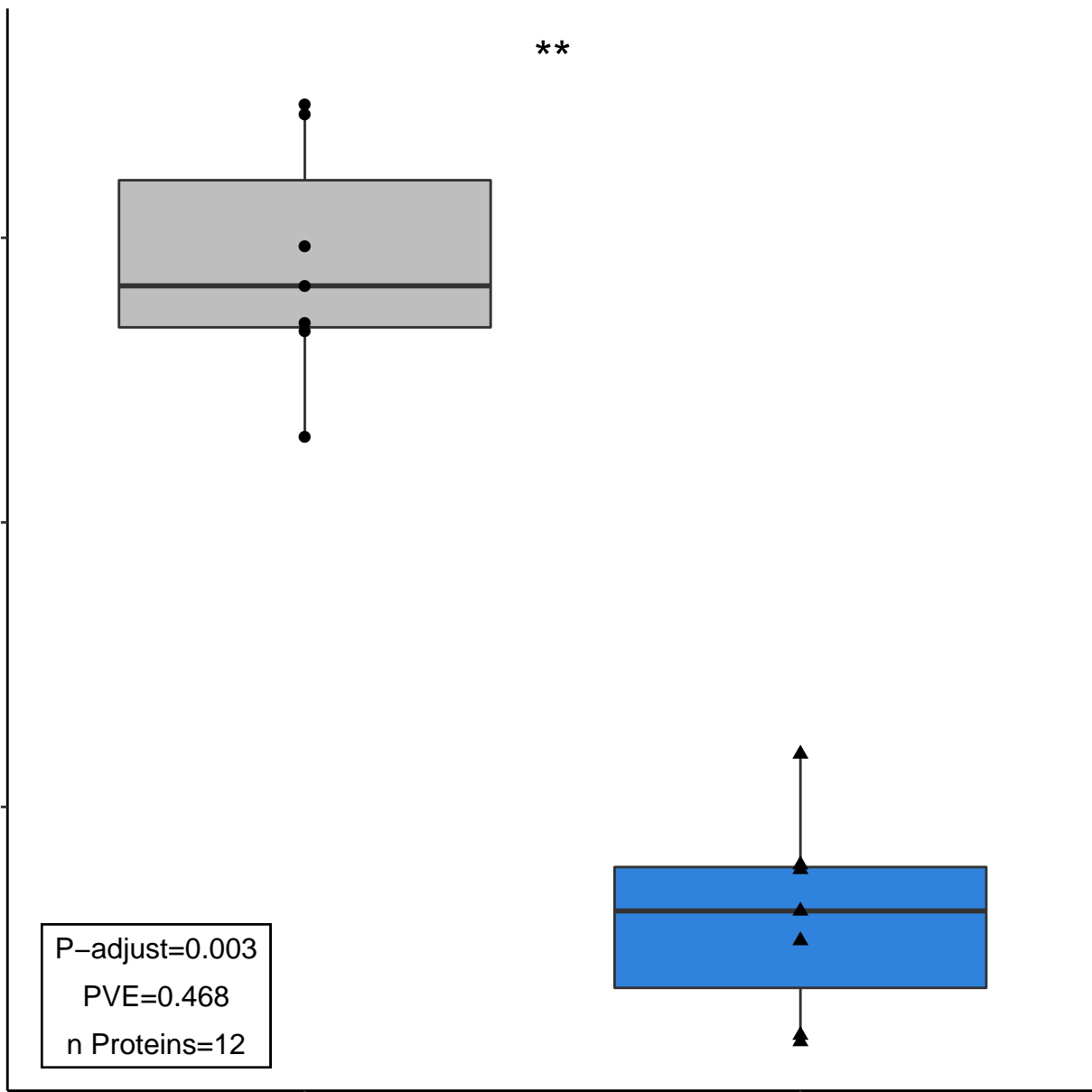
PVE=0.468

n Proteins=12

WT

Genotype

MUT



M197

Log2(Adjusted Module Intensity)

13.80

13.75

13.70

13.65

WT

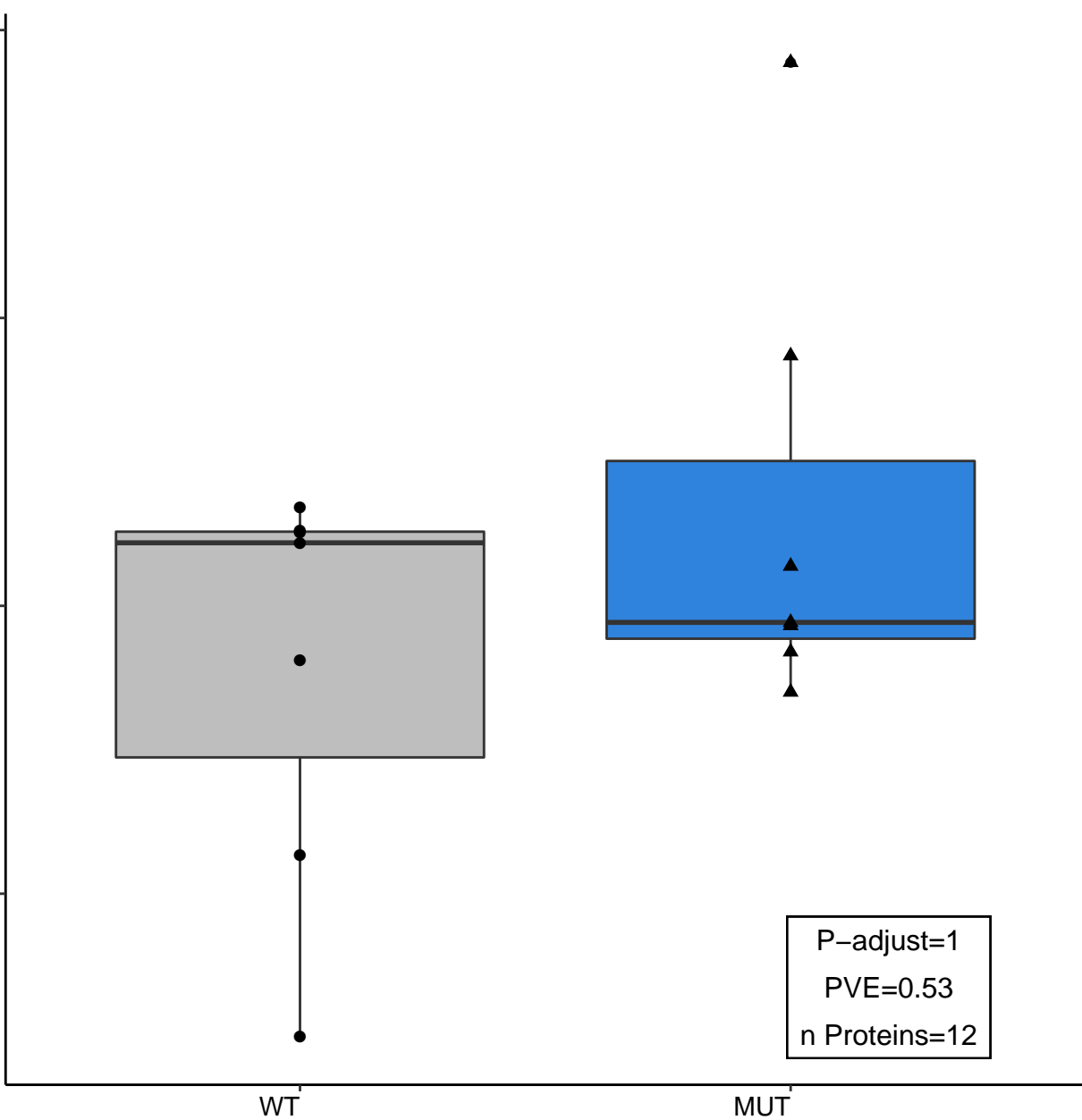
Genotype

MUT

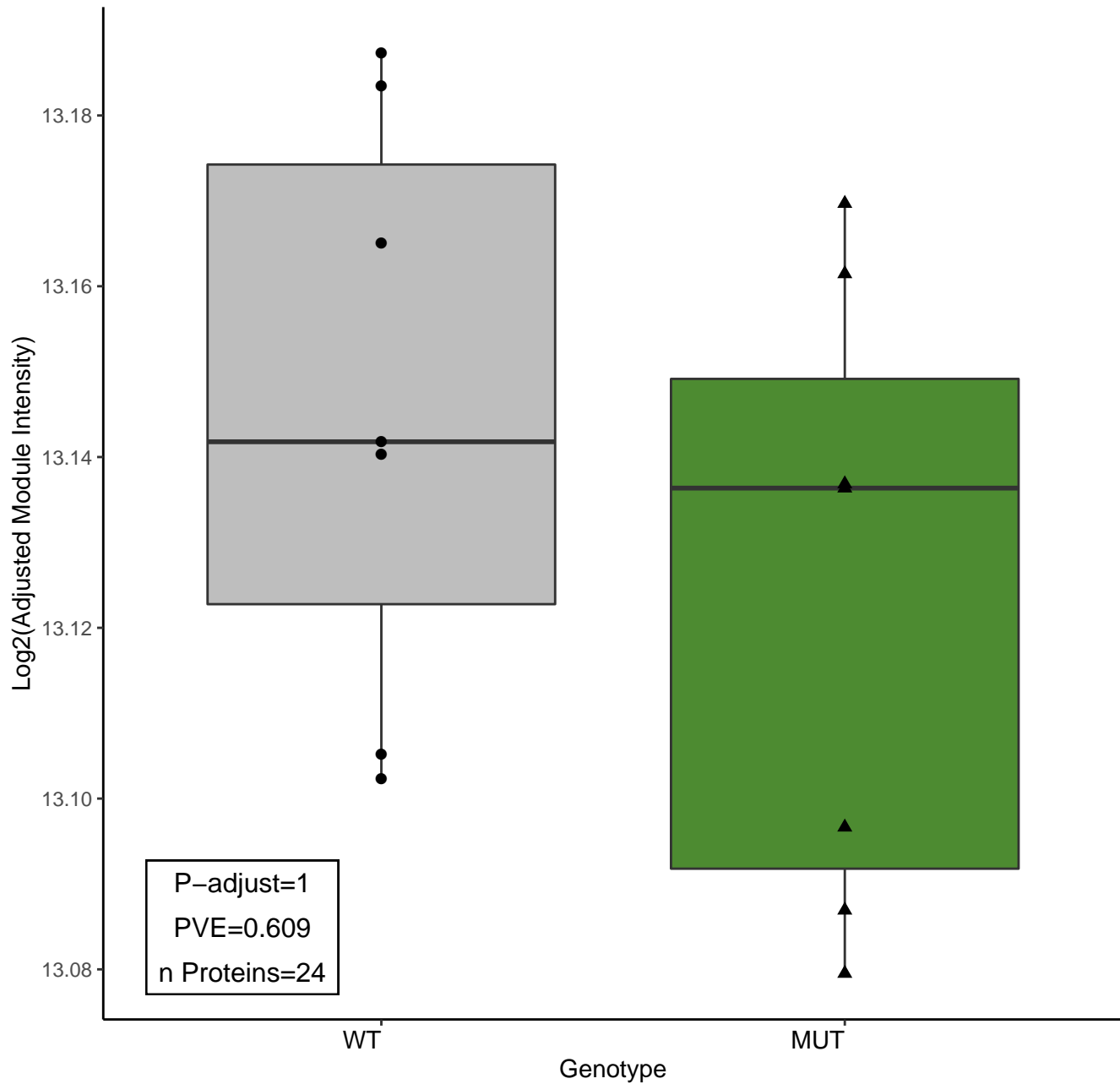
P-adjust=1

PVE=0.53

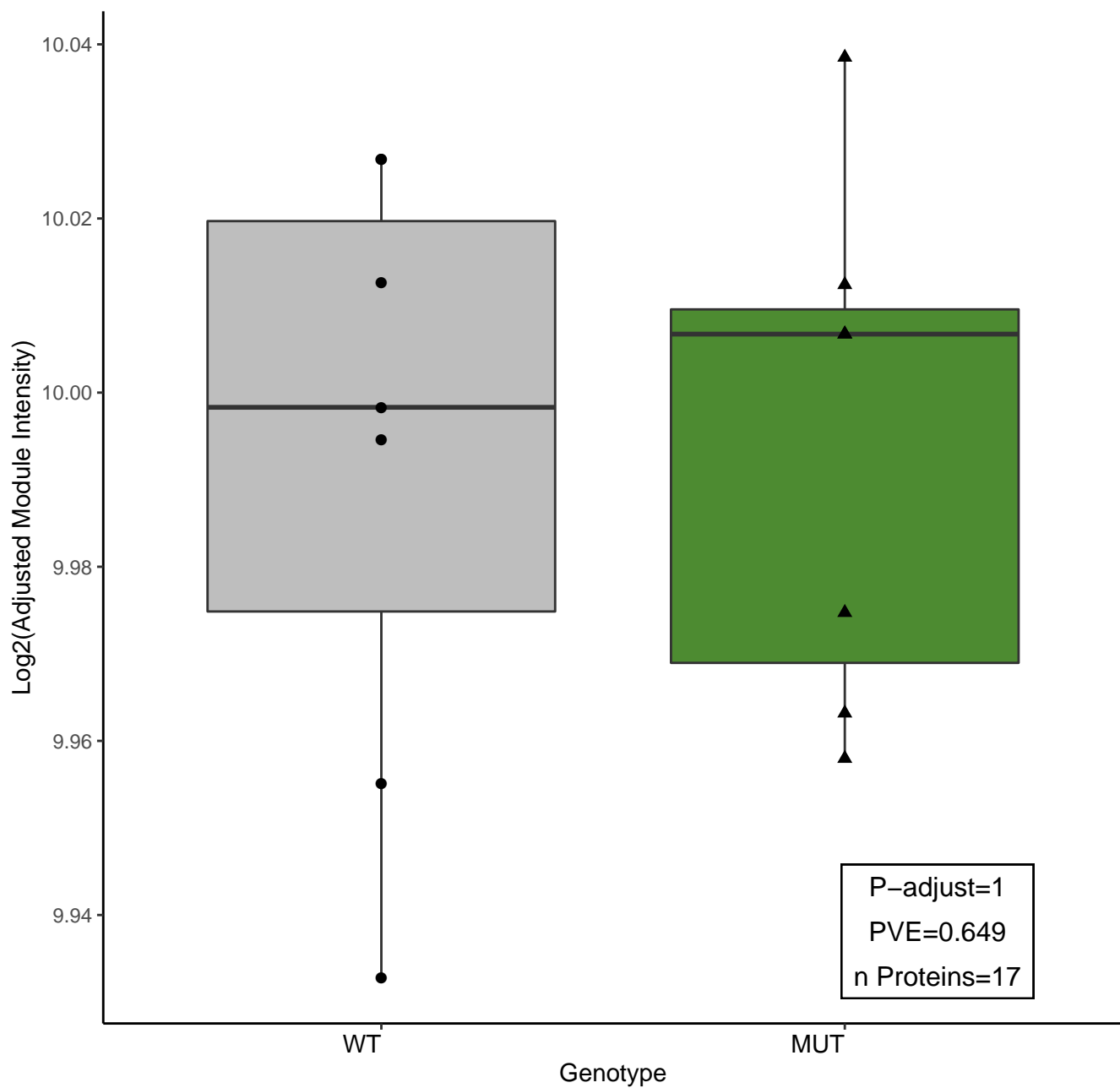
n Proteins=12

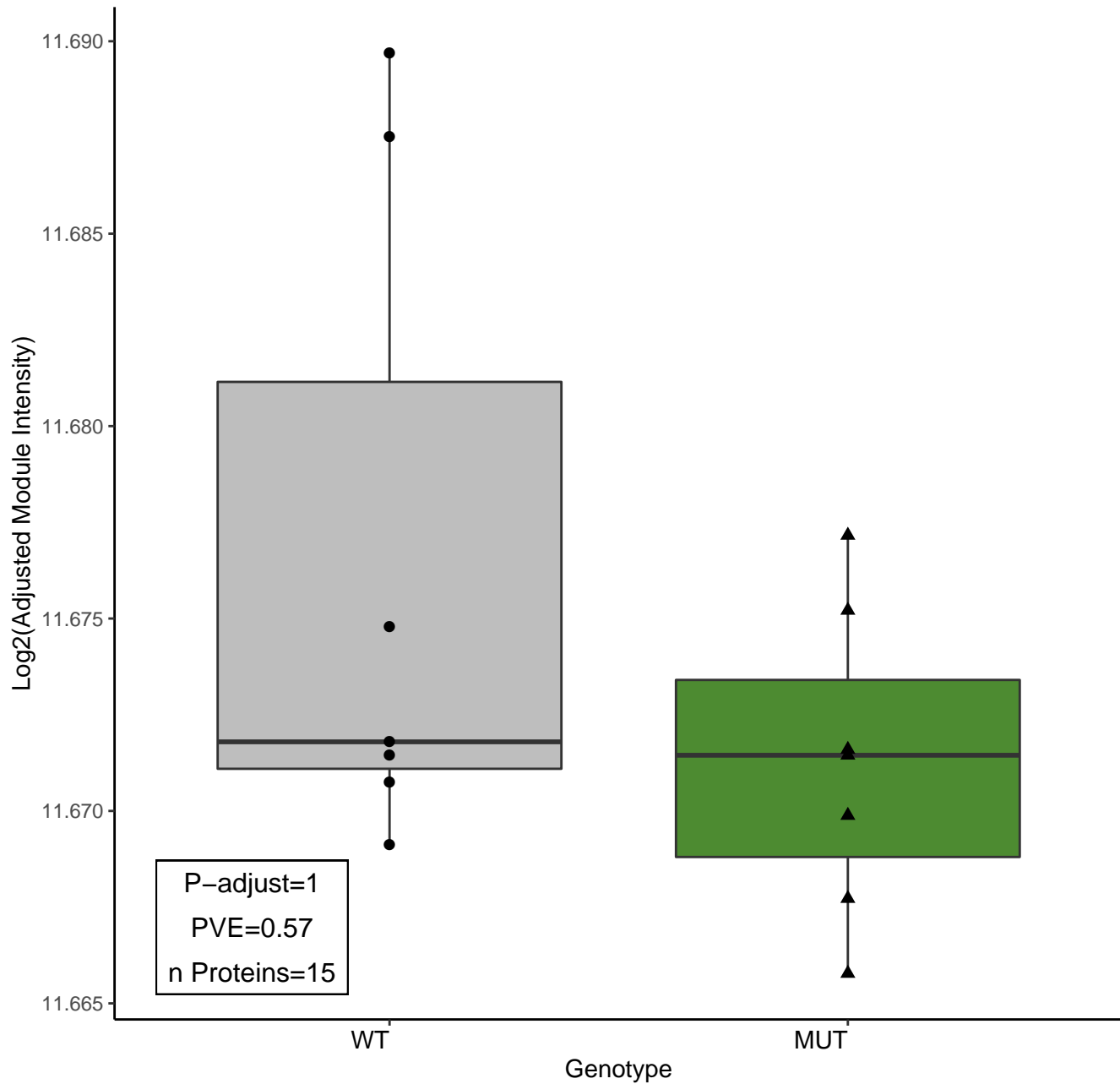


M198



M199





M201

Log2(Adjusted Module Intensity)

13.65

13.60

13.55

13.50

P-adjust=0.478

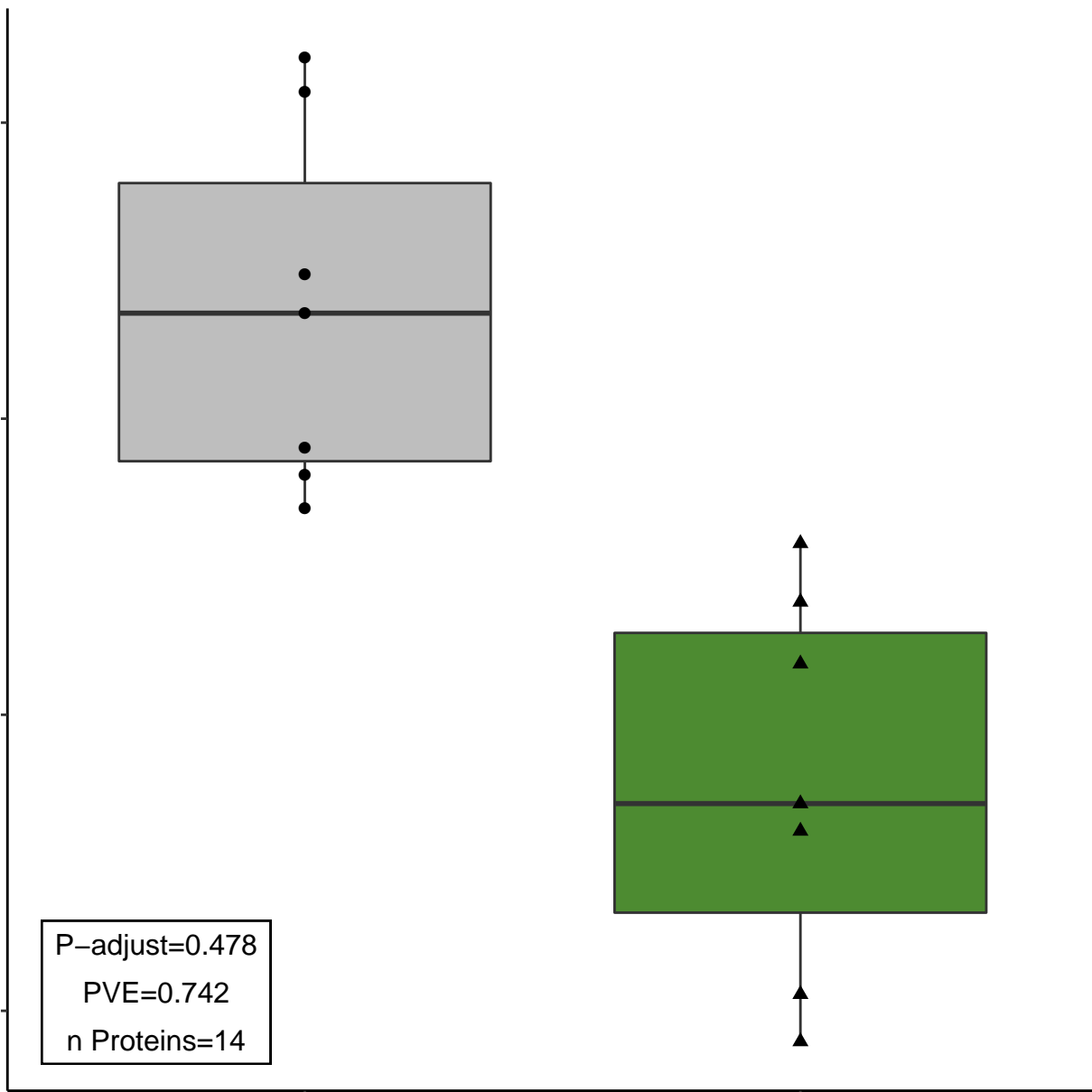
PVE=0.742

n Proteins=14

WT

Genotype

MUT



M202

Log2(Adjusted Module Intensity)

10.20

10.18

10.16

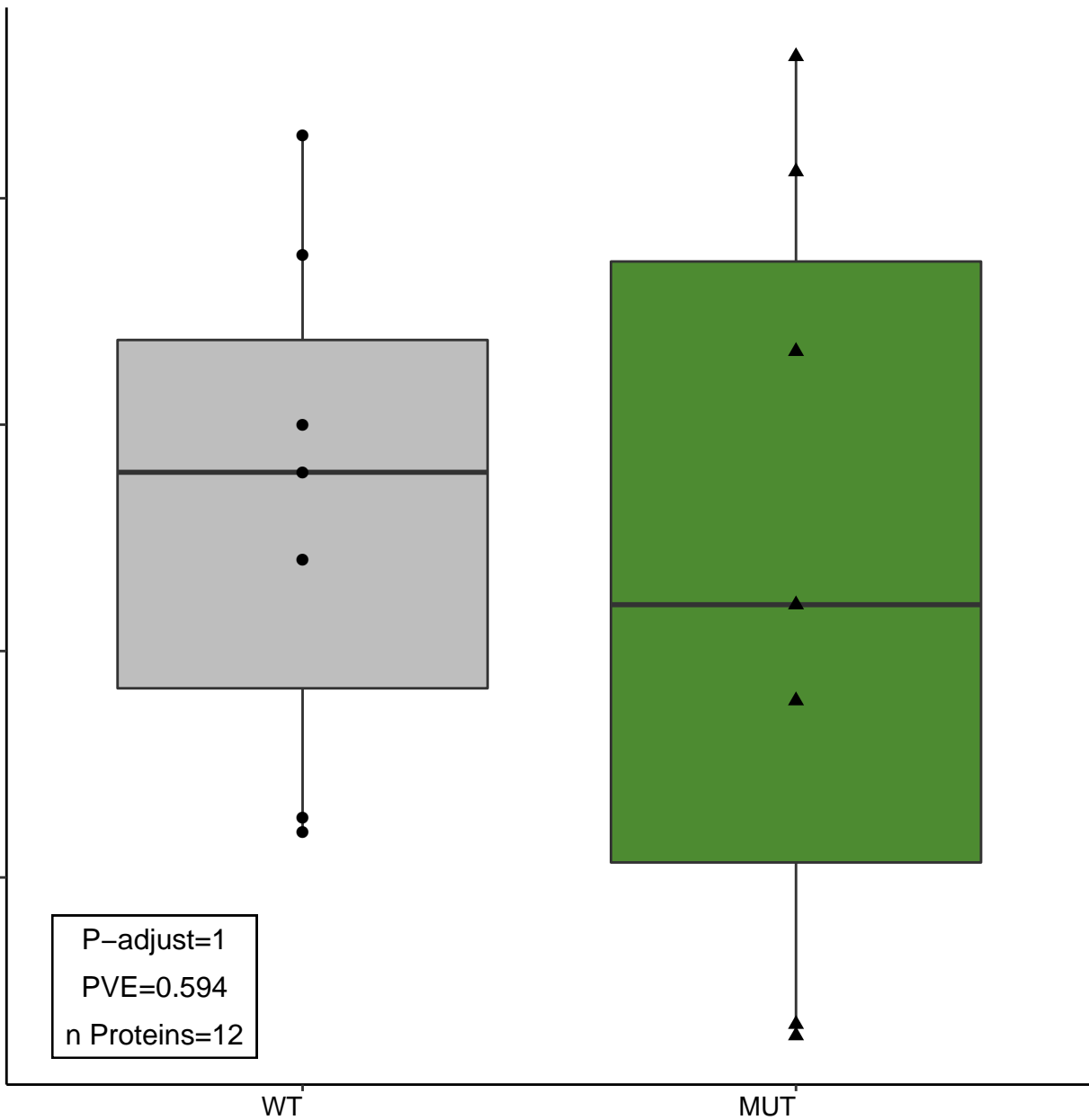
10.14

P-adjust=1
PVE=0.594
n Proteins=12

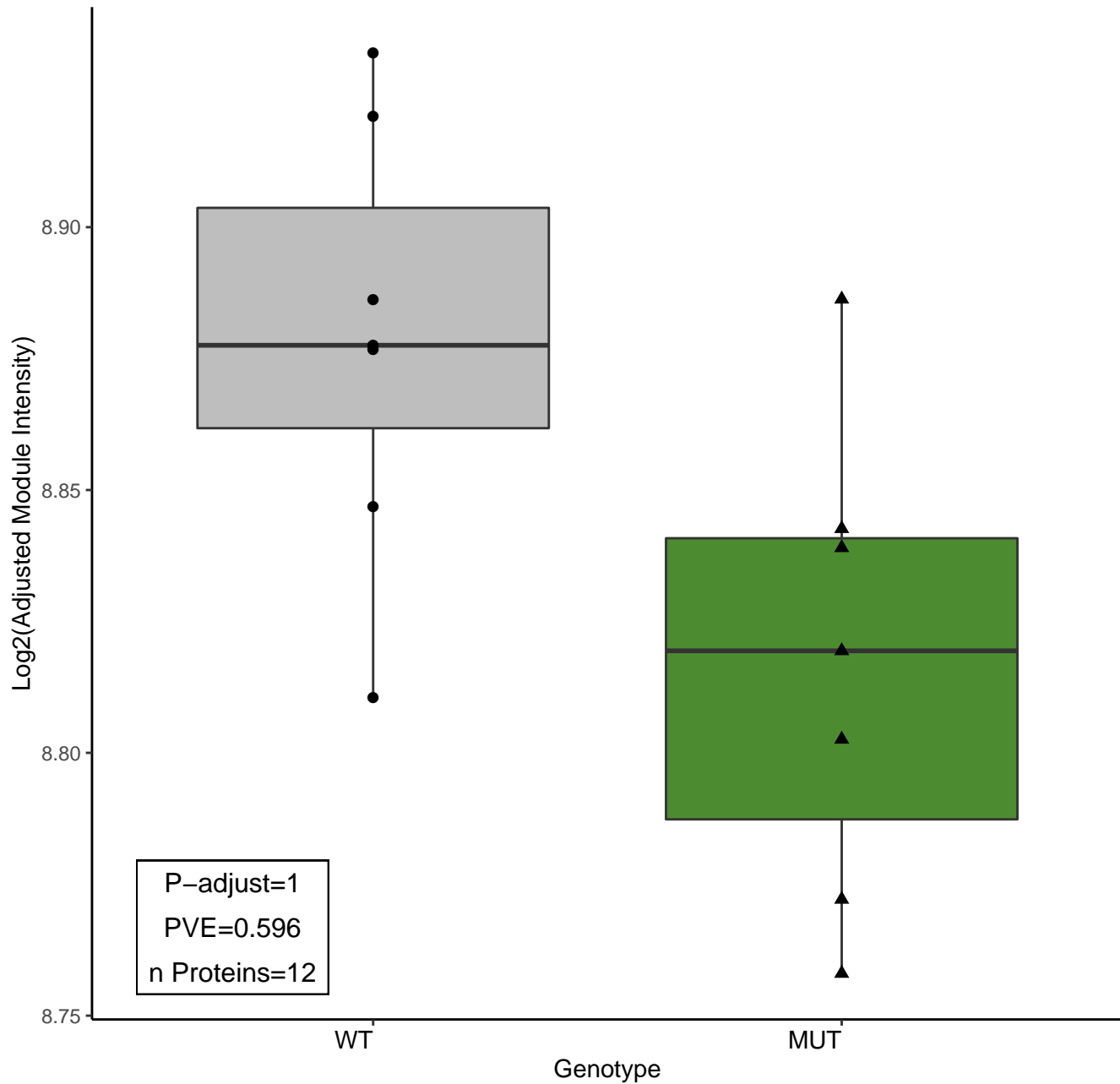
WT

Genotype

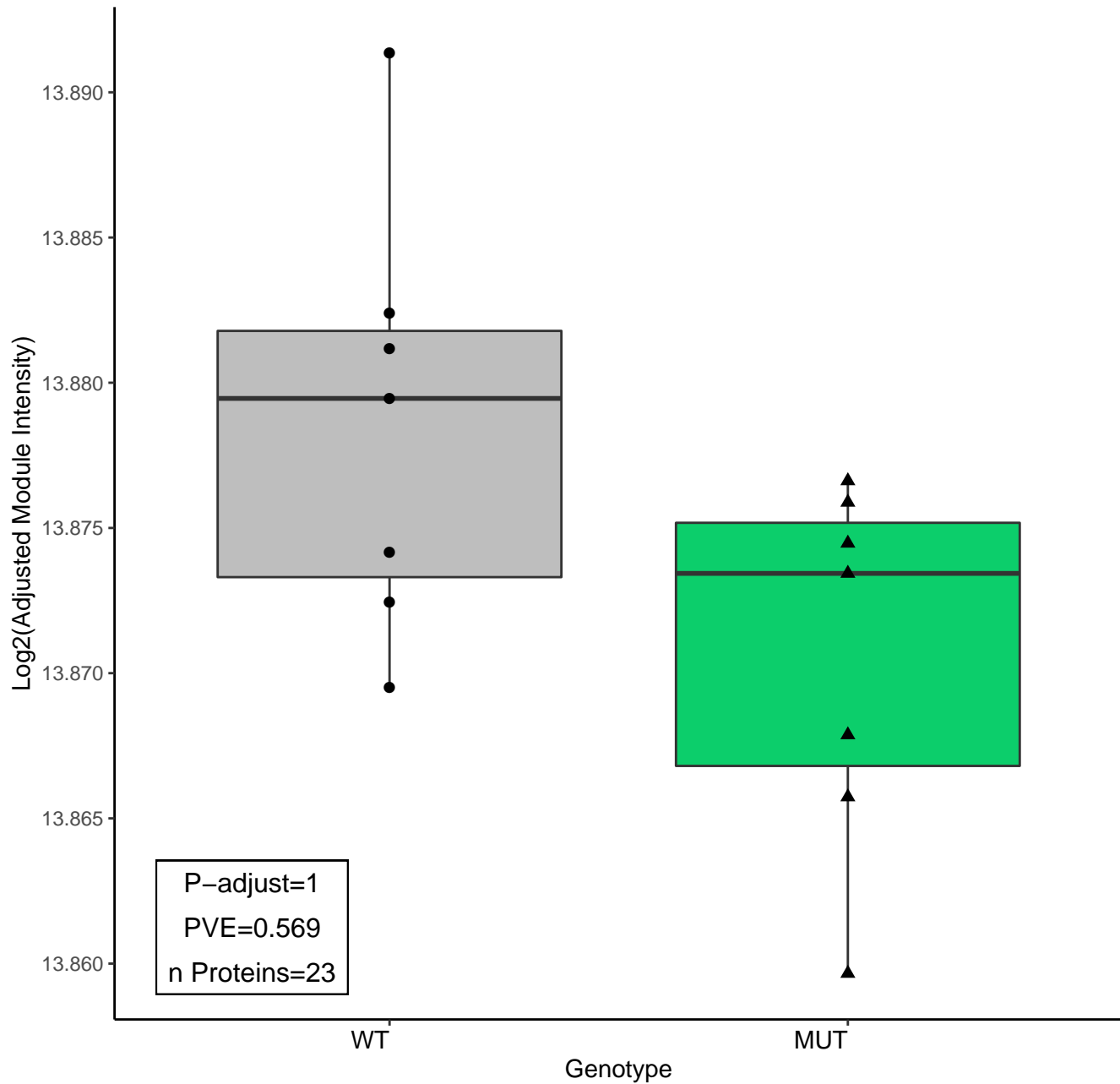
MUT



M203



M204



M205

**

Log2(Adjusted Module Intensity)

12.25

12.20

12.15

12.10

WT

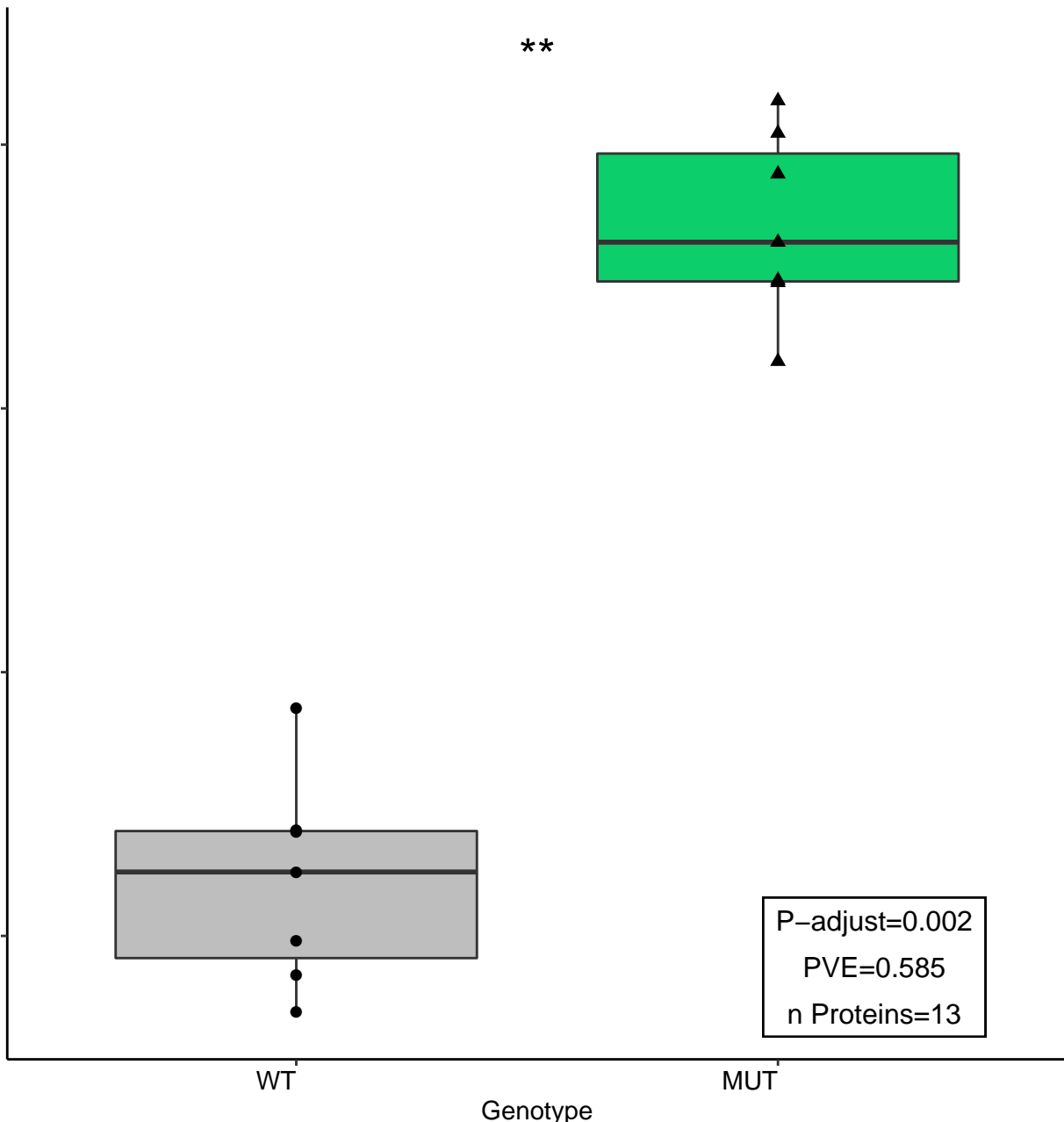
MUT

Genotype

P-adjust=0.002

PVE=0.585

n Proteins=13



M206

Log2(Adjusted Module Intensity)

13.05

13.00

12.95

WT

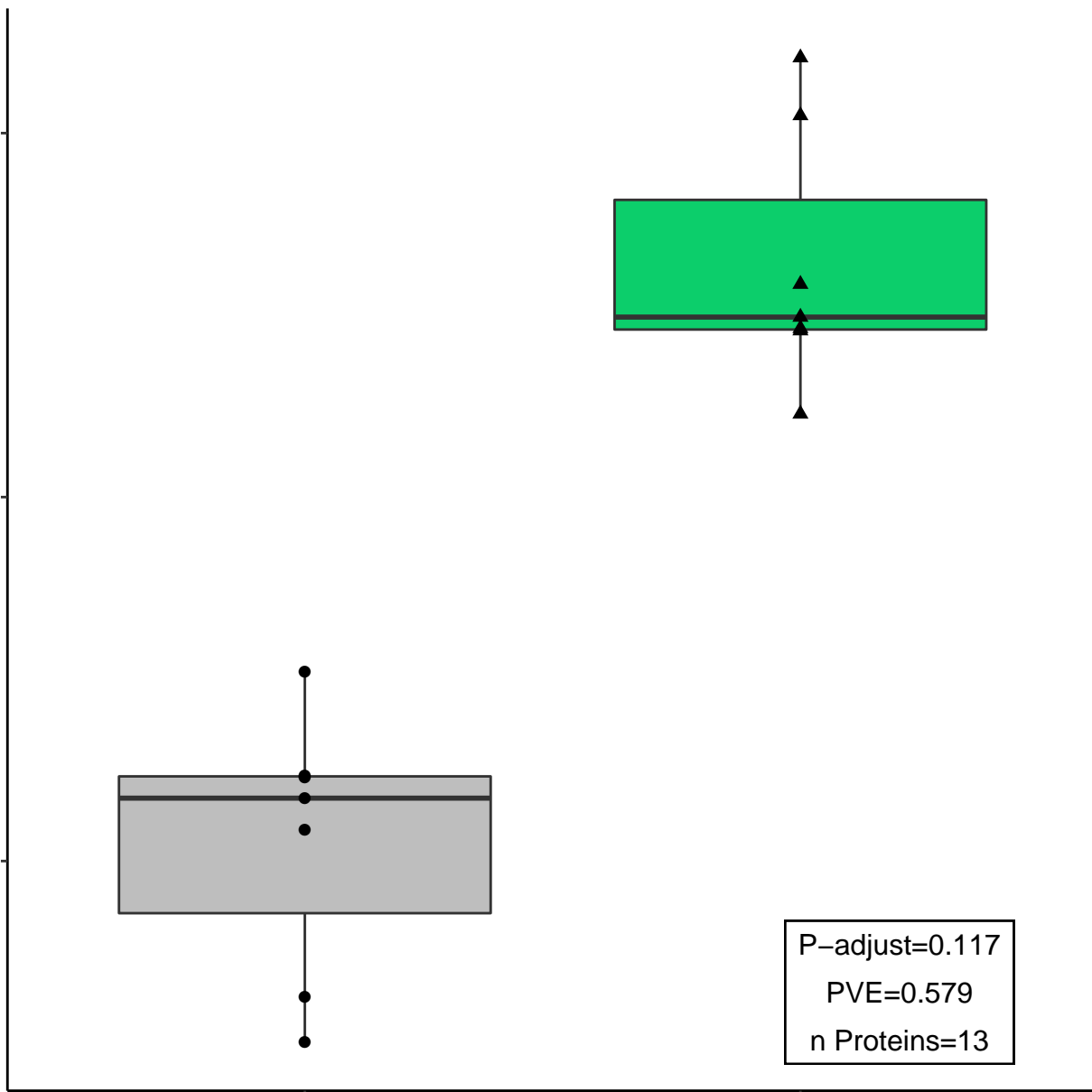
MUT

Genotype

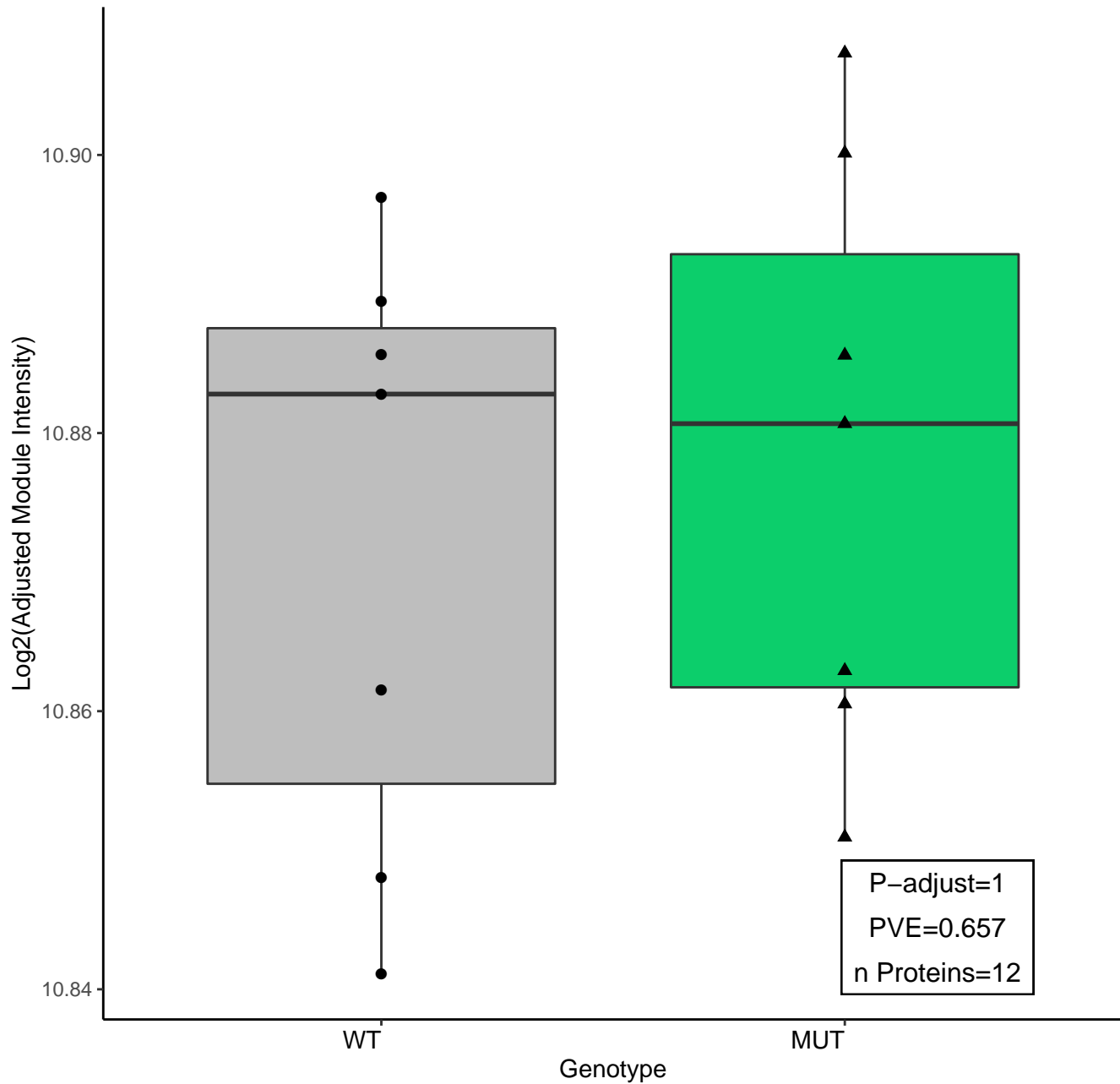
P-adjust=0.117

PVE=0.579

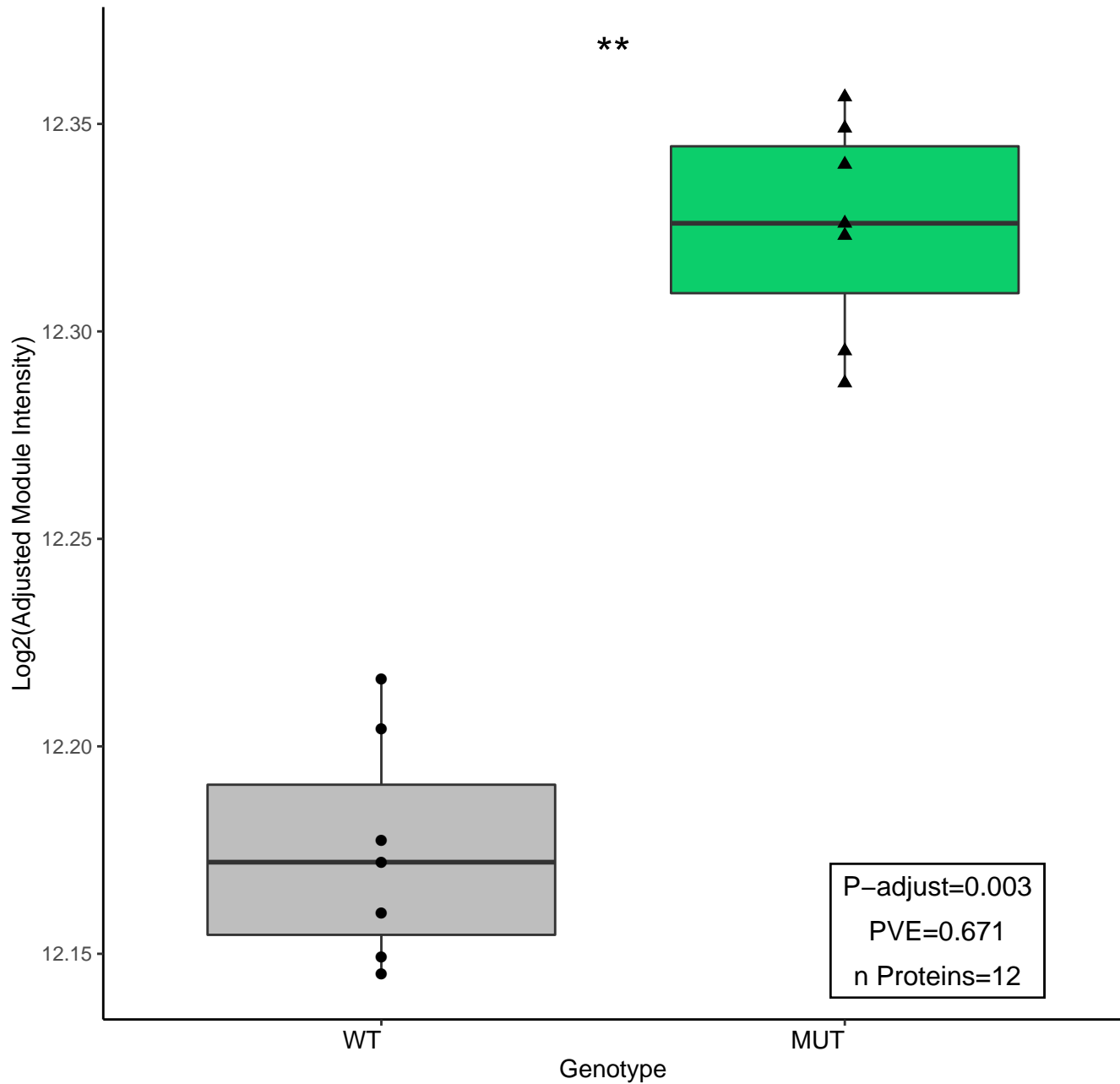
n Proteins=13



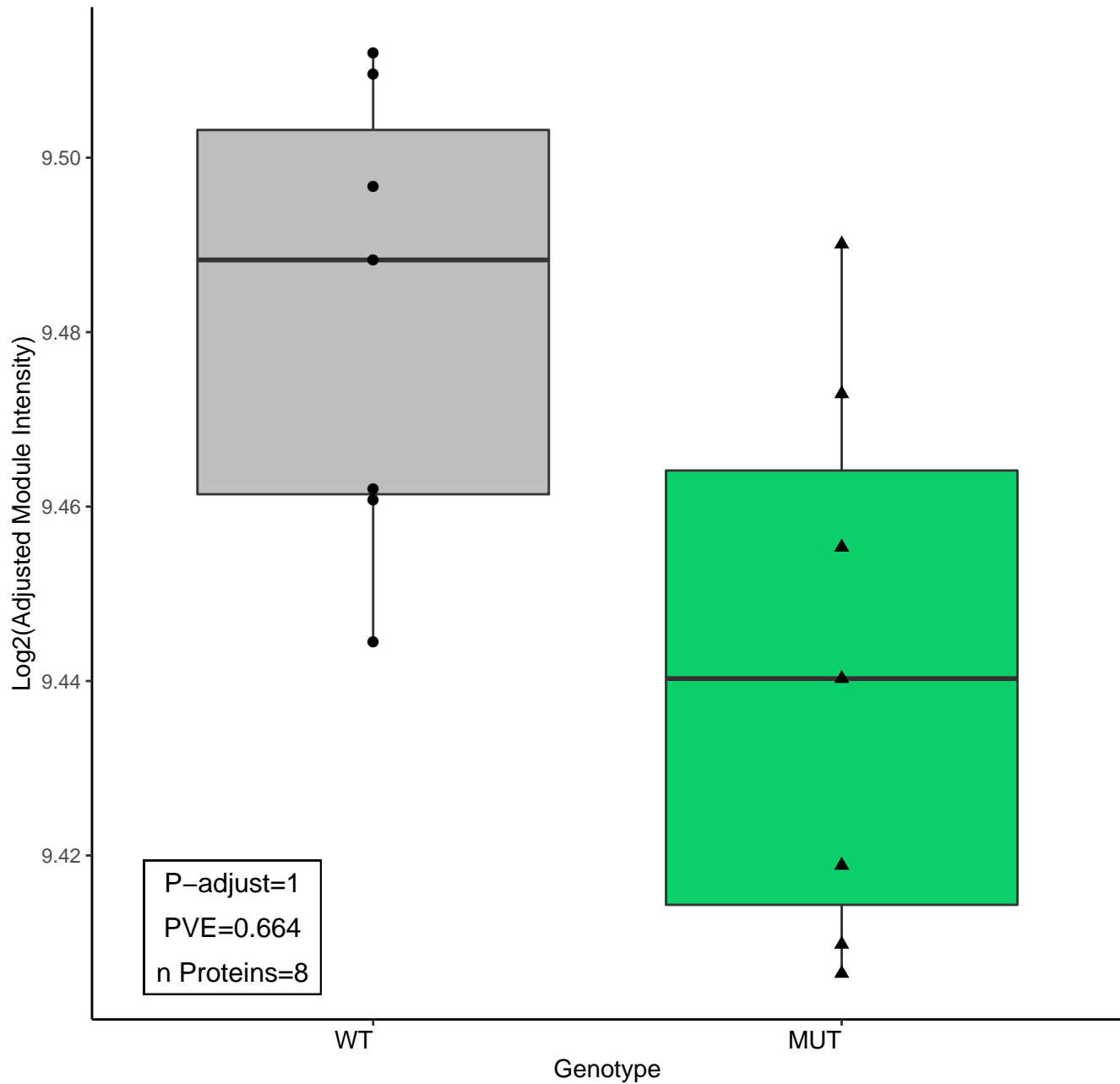
M207



M208



M209



M210

Log2(Adjusted Module Intensity)

10.64

10.63

10.62

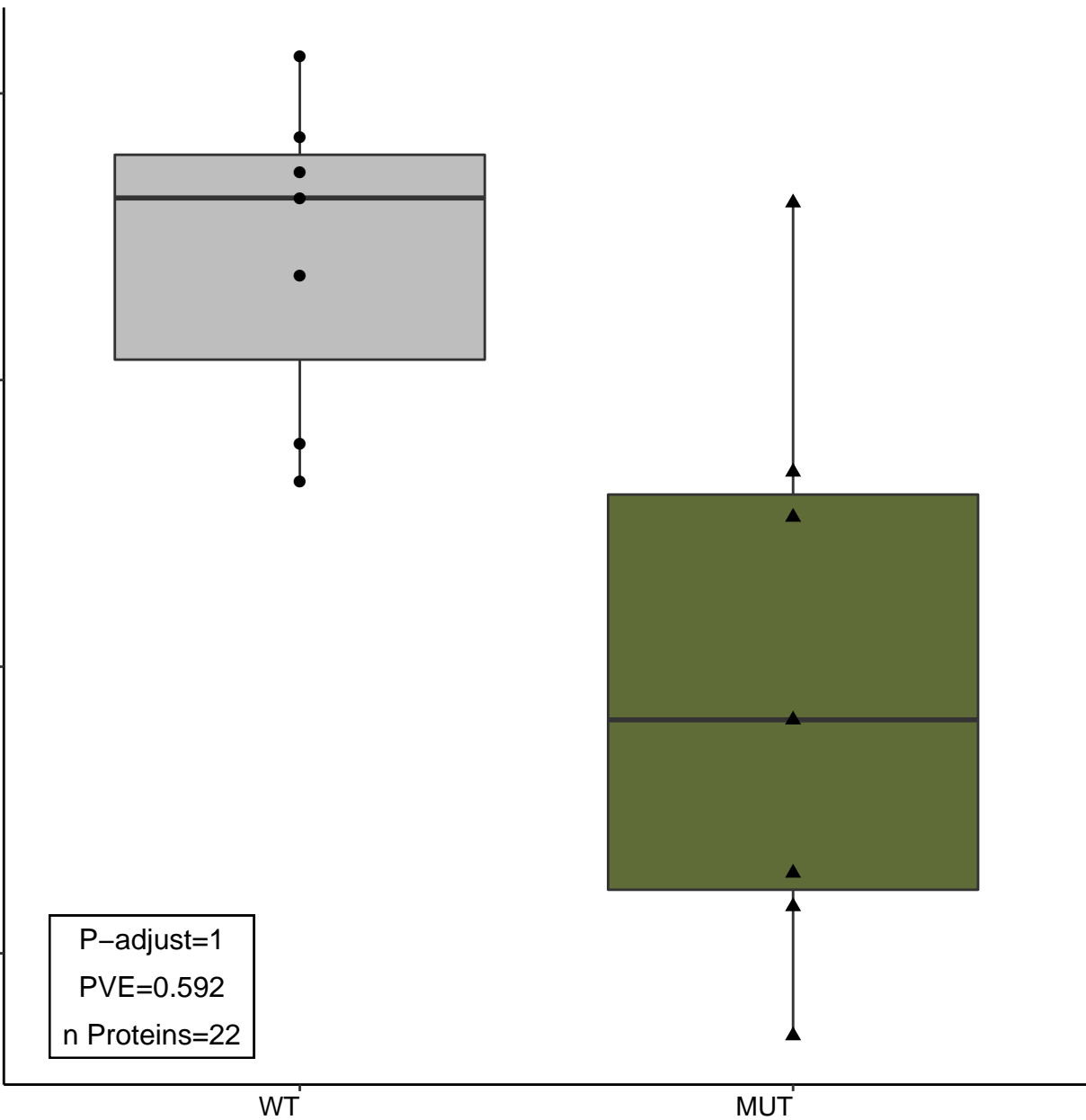
10.61

P-adjust=1
PVE=0.592
n Proteins=22

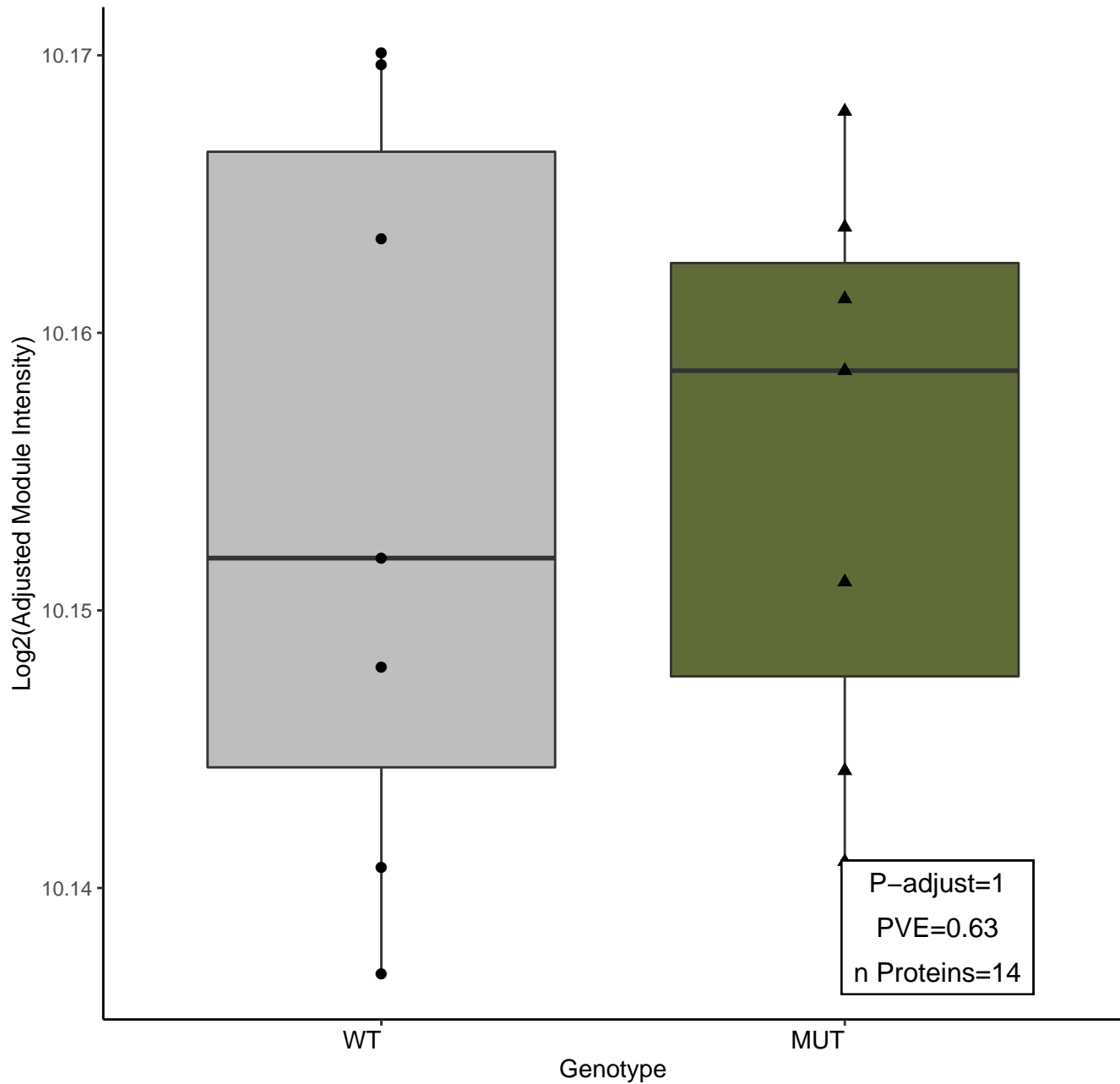
WT

Genotype

MUT



M211



M212

Log2(Adjusted Module Intensity)

10.31

10.30

10.29

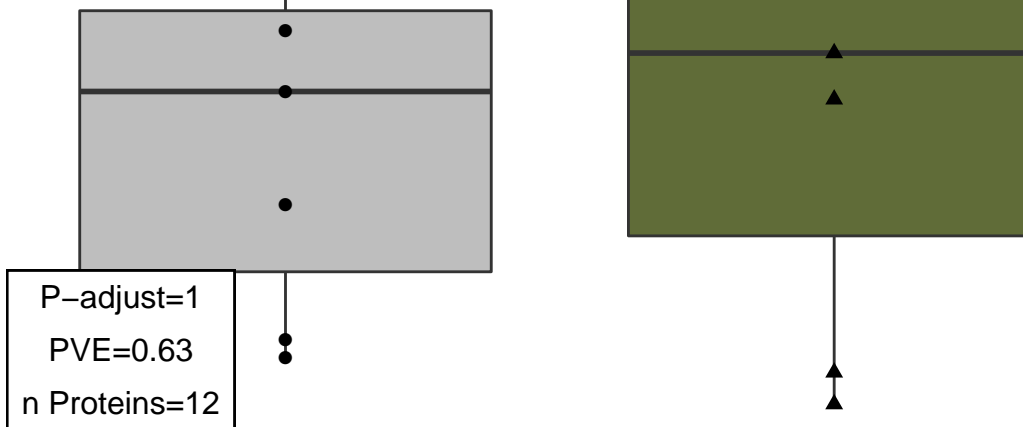
10.28

WT

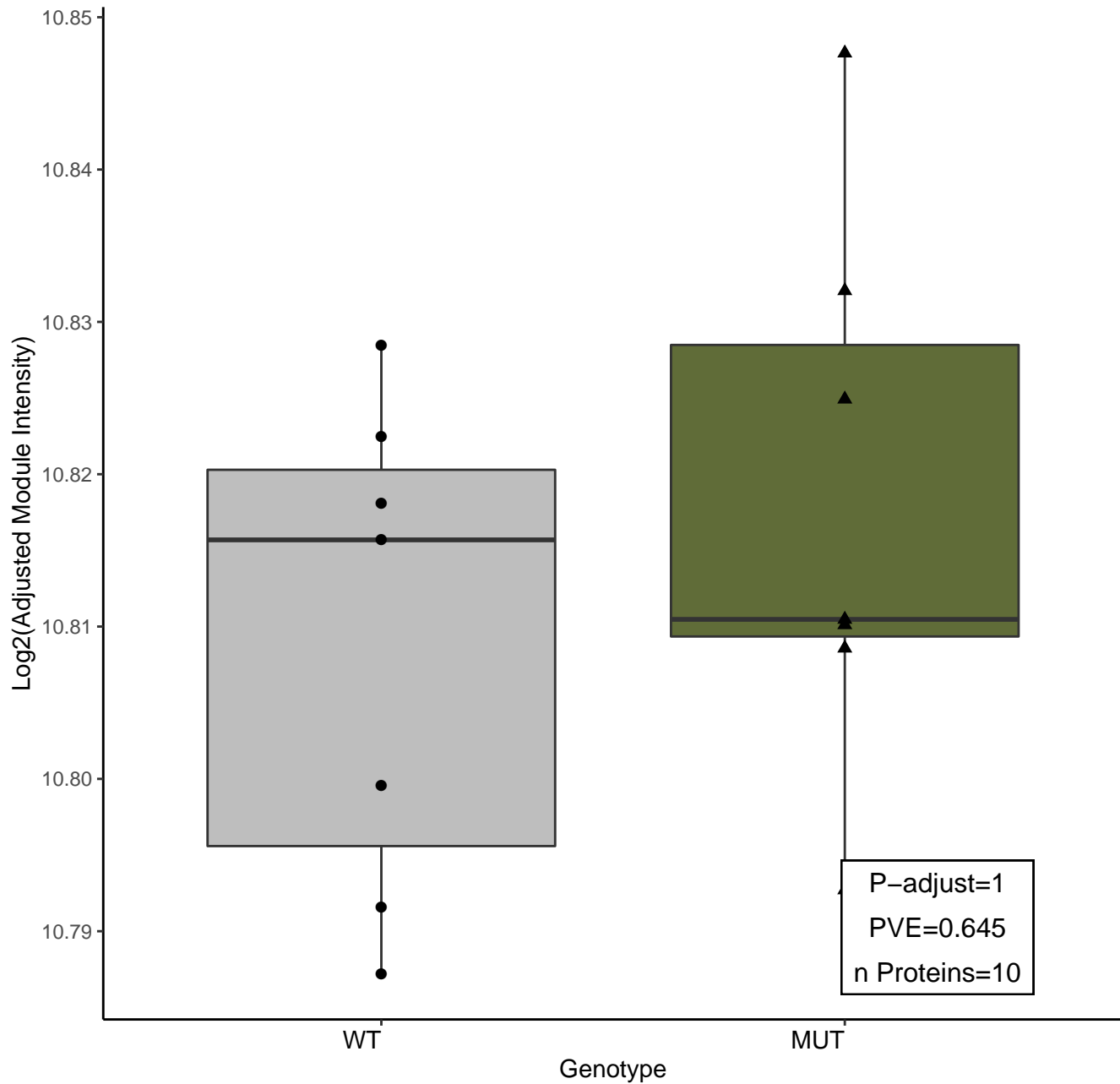
MUT

Genotype

P-adjust=1
PVE=0.63
n Proteins=12



M213



M214

Log2(Adjusted Module Intensity)

9.02

9.00

8.98

8.96

WT

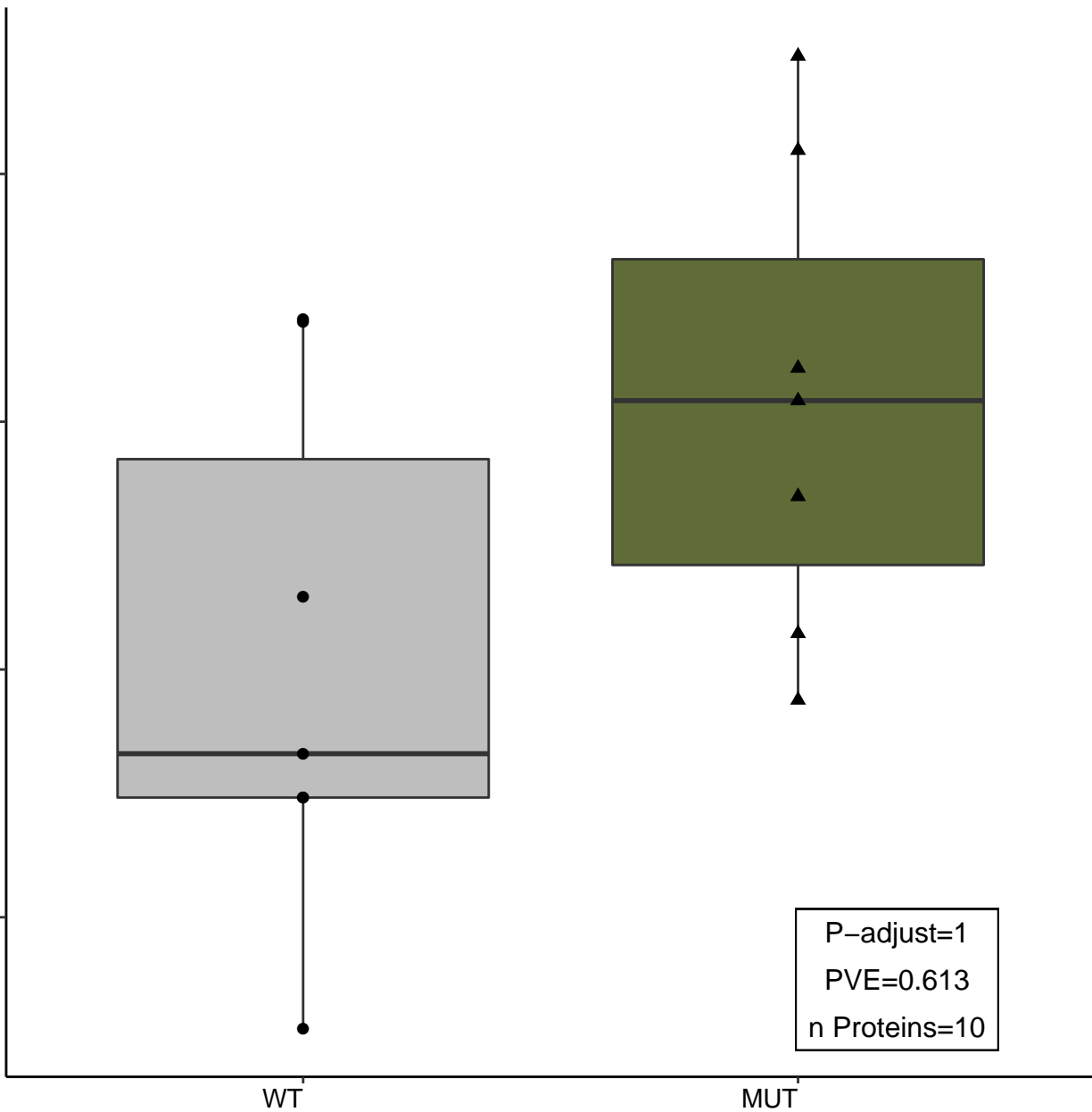
Genotype

MUT

P-adjust=1

PVE=0.613

n Proteins=10



M215

Log2(Adjusted Module Intensity)

11.05

11.00

10.95

10.90

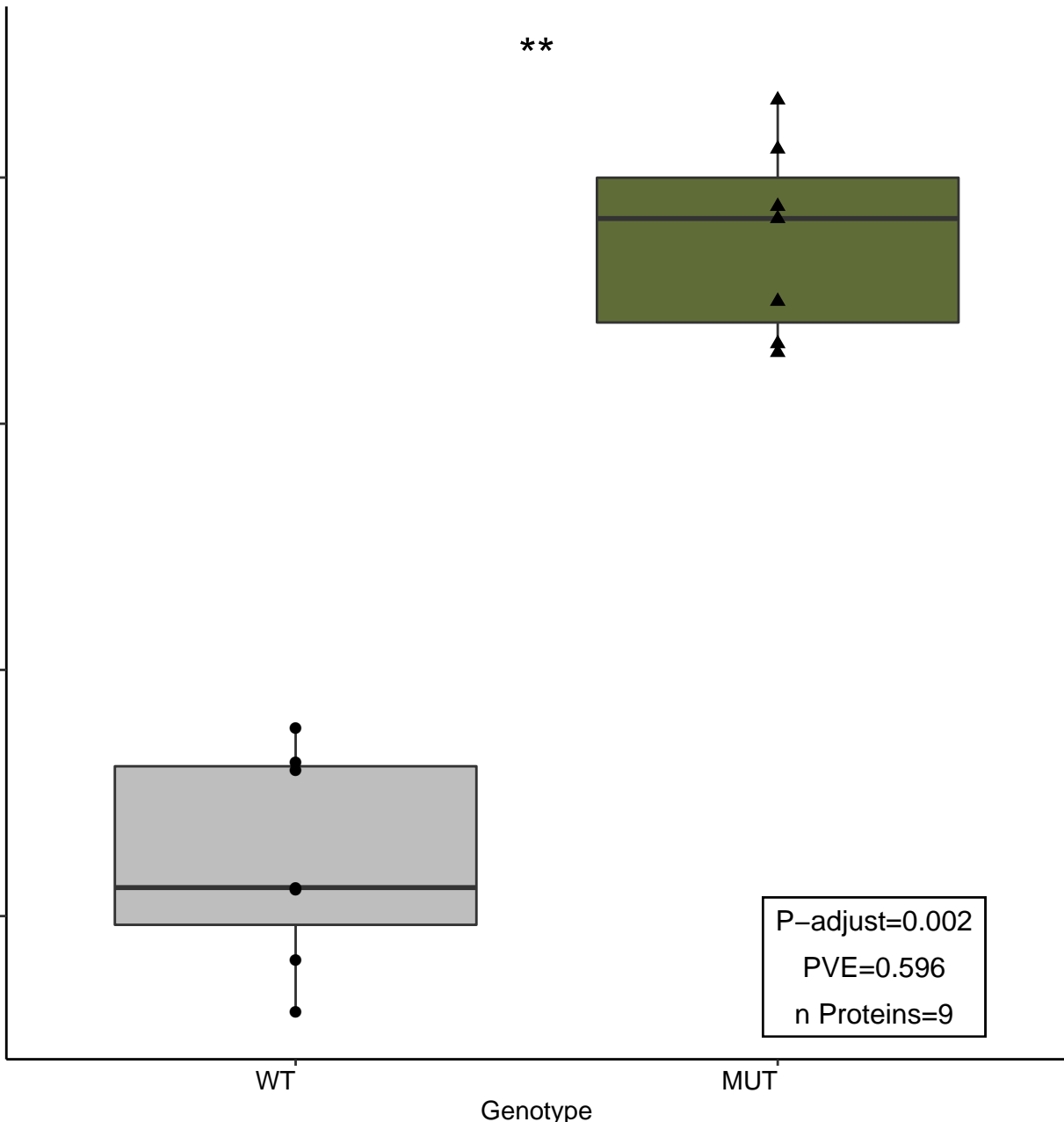
WT

MUT

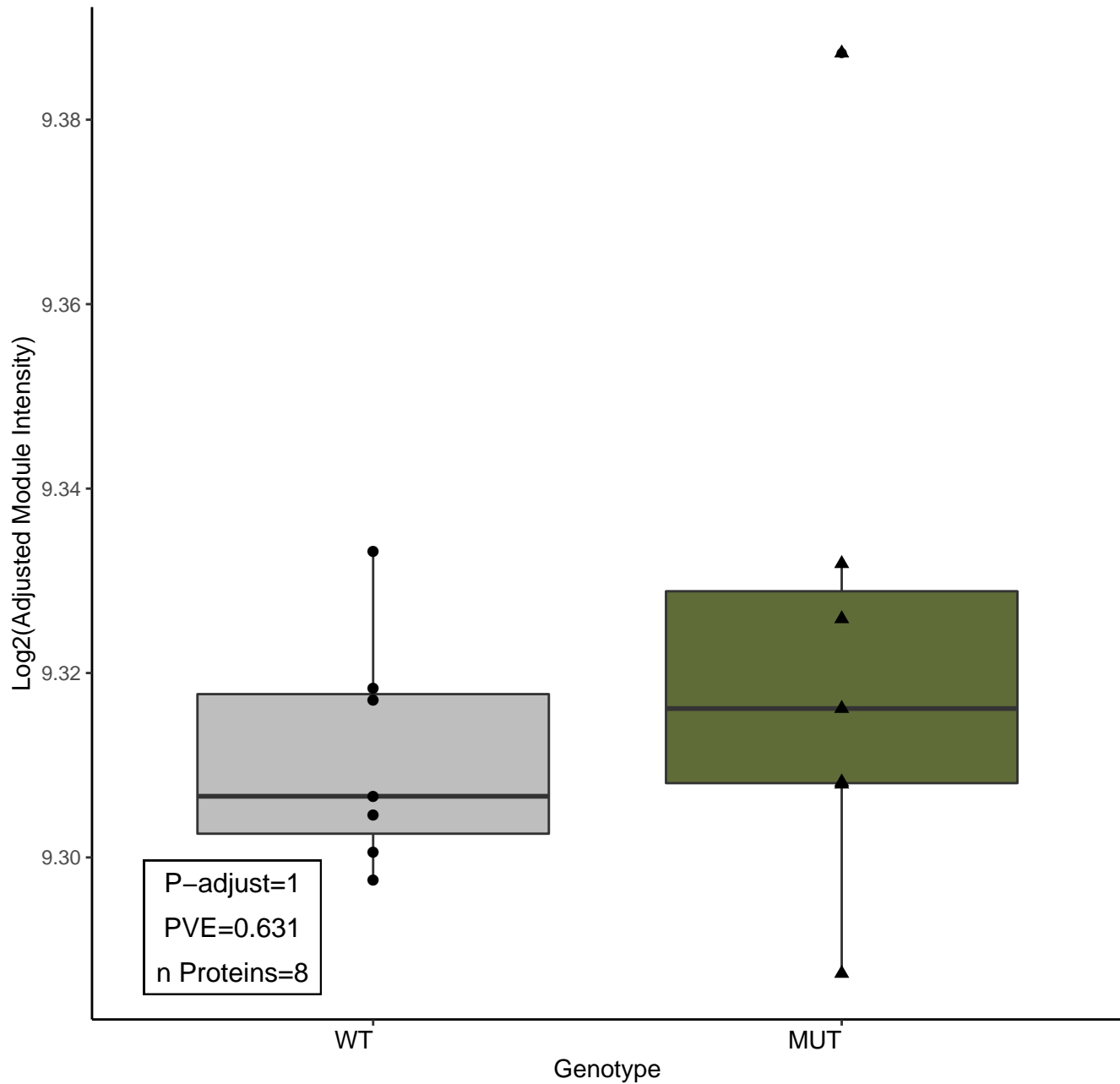
Genotype

**

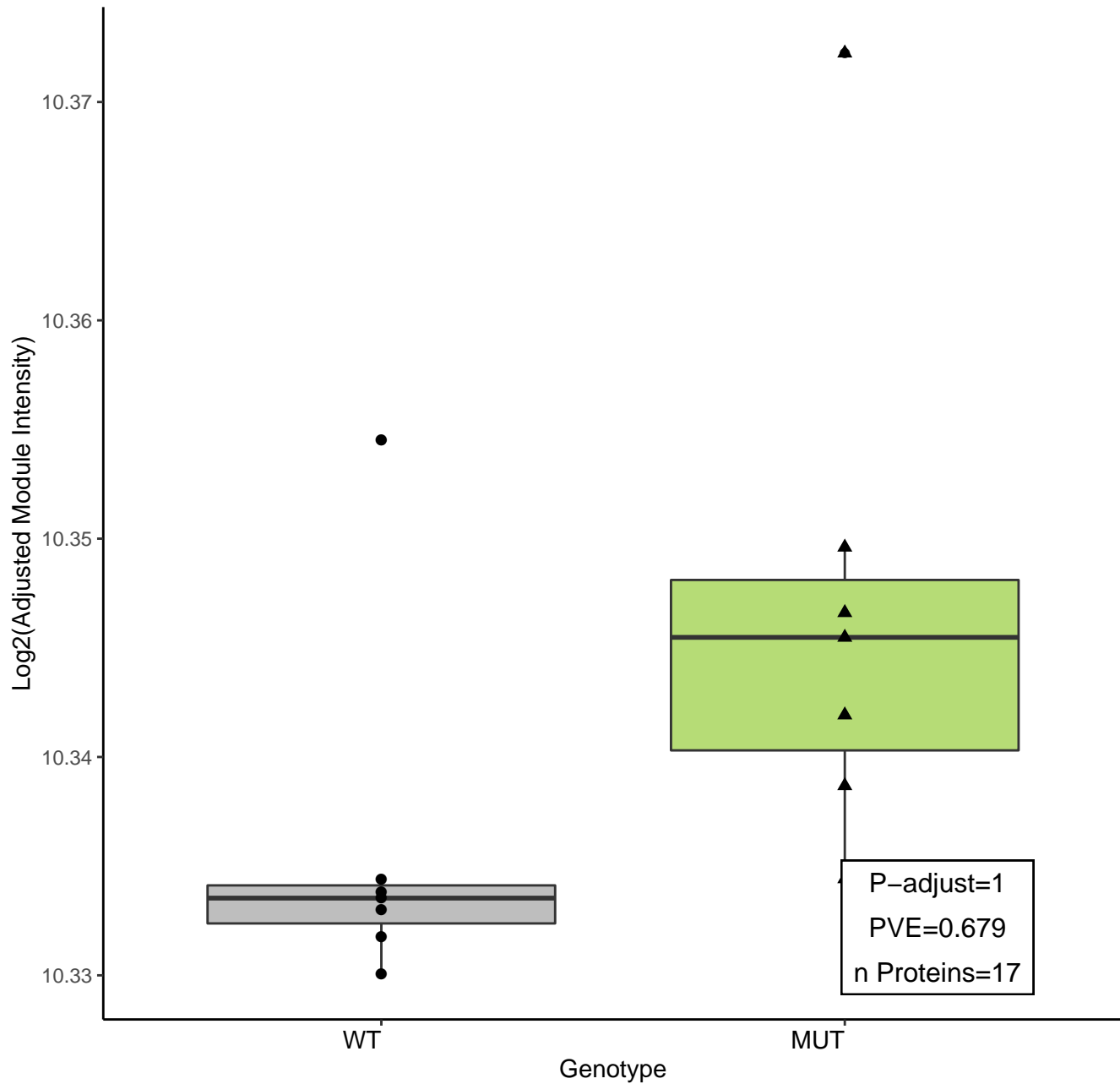
P-adjust=0.002
PVE=0.596
n Proteins=9



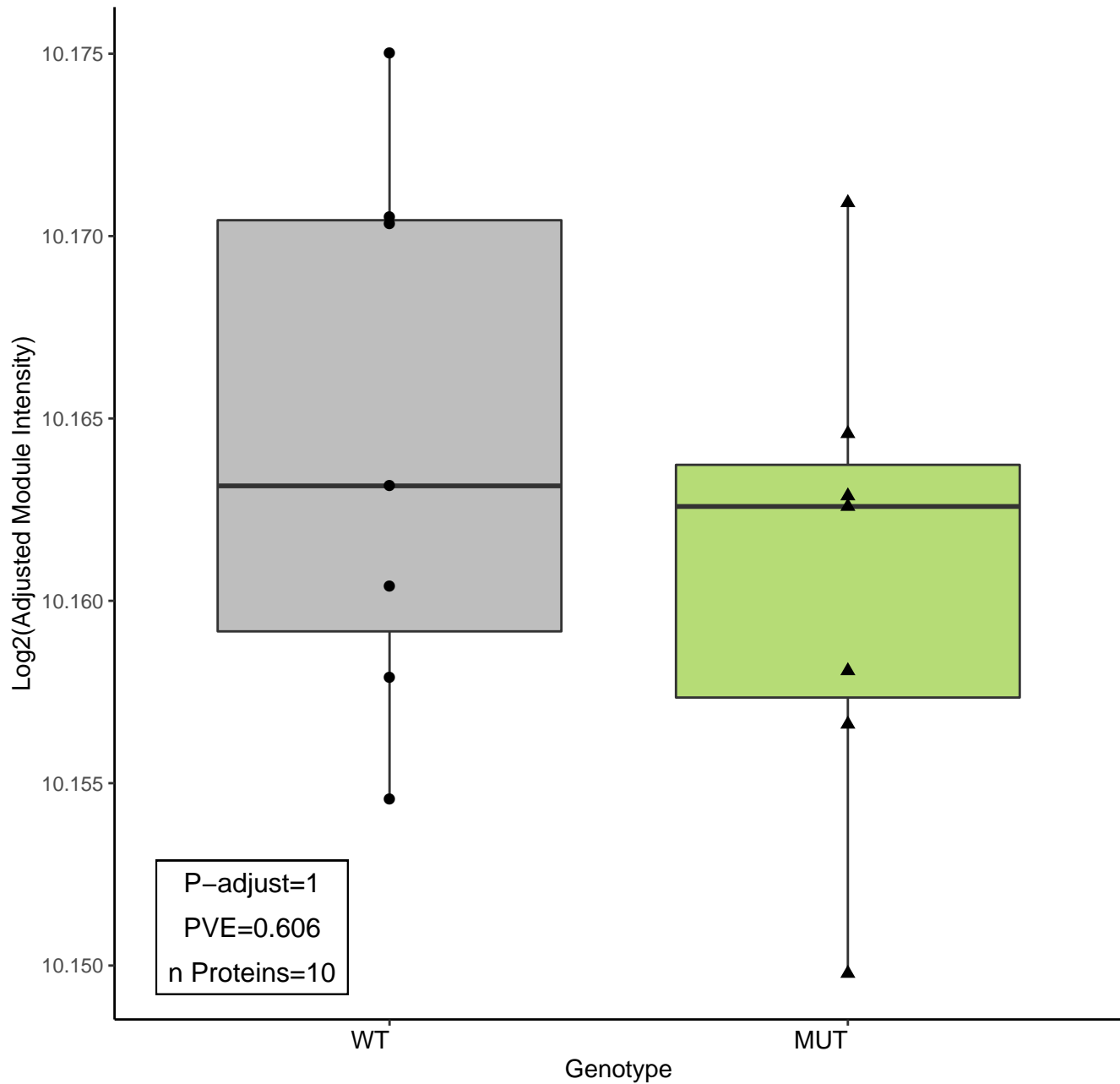
M216



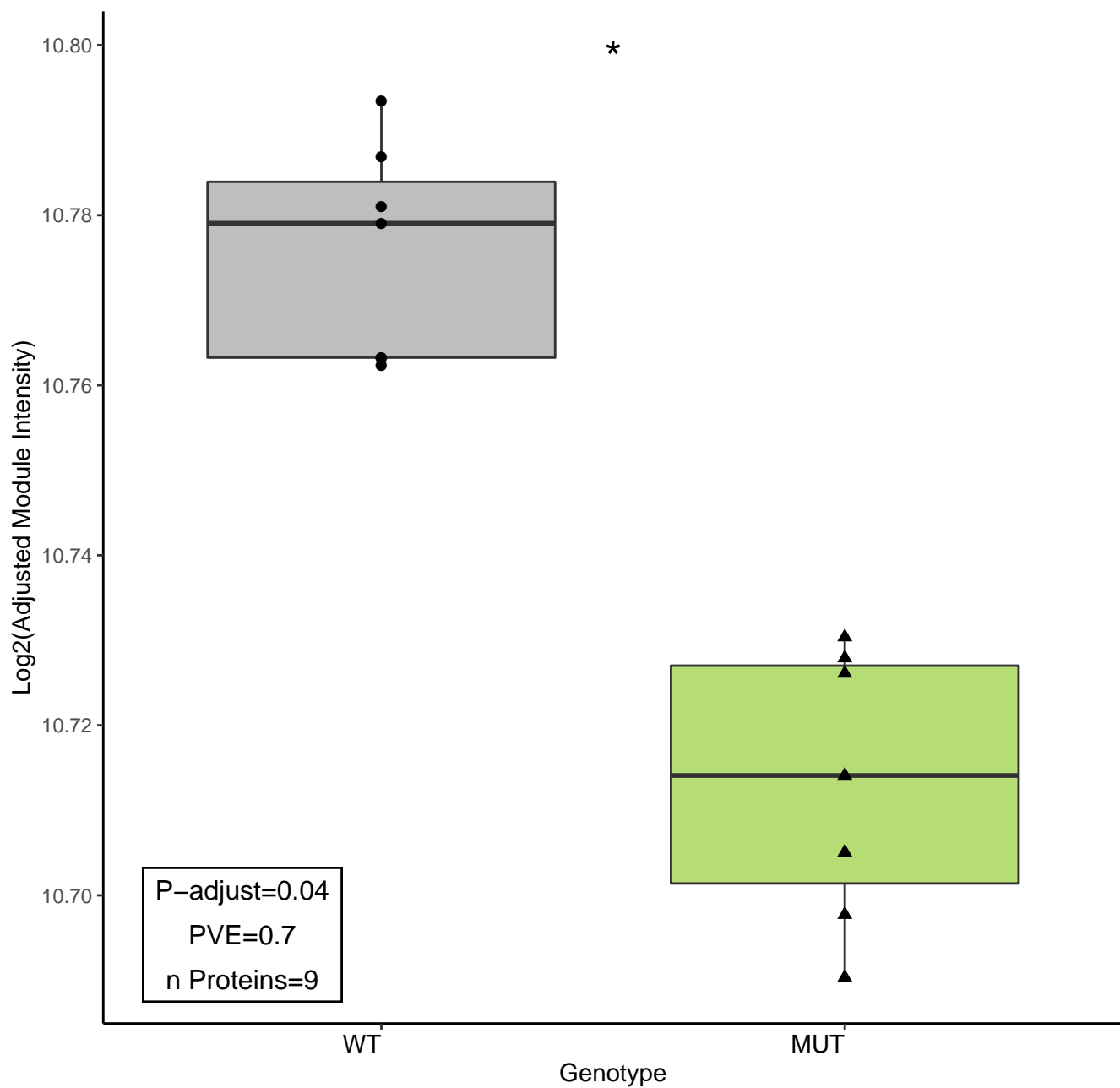
M217



M218



M219



M220

Log2(Adjusted Module Intensity)

10.66

10.64

10.62

P-adjust=0.654

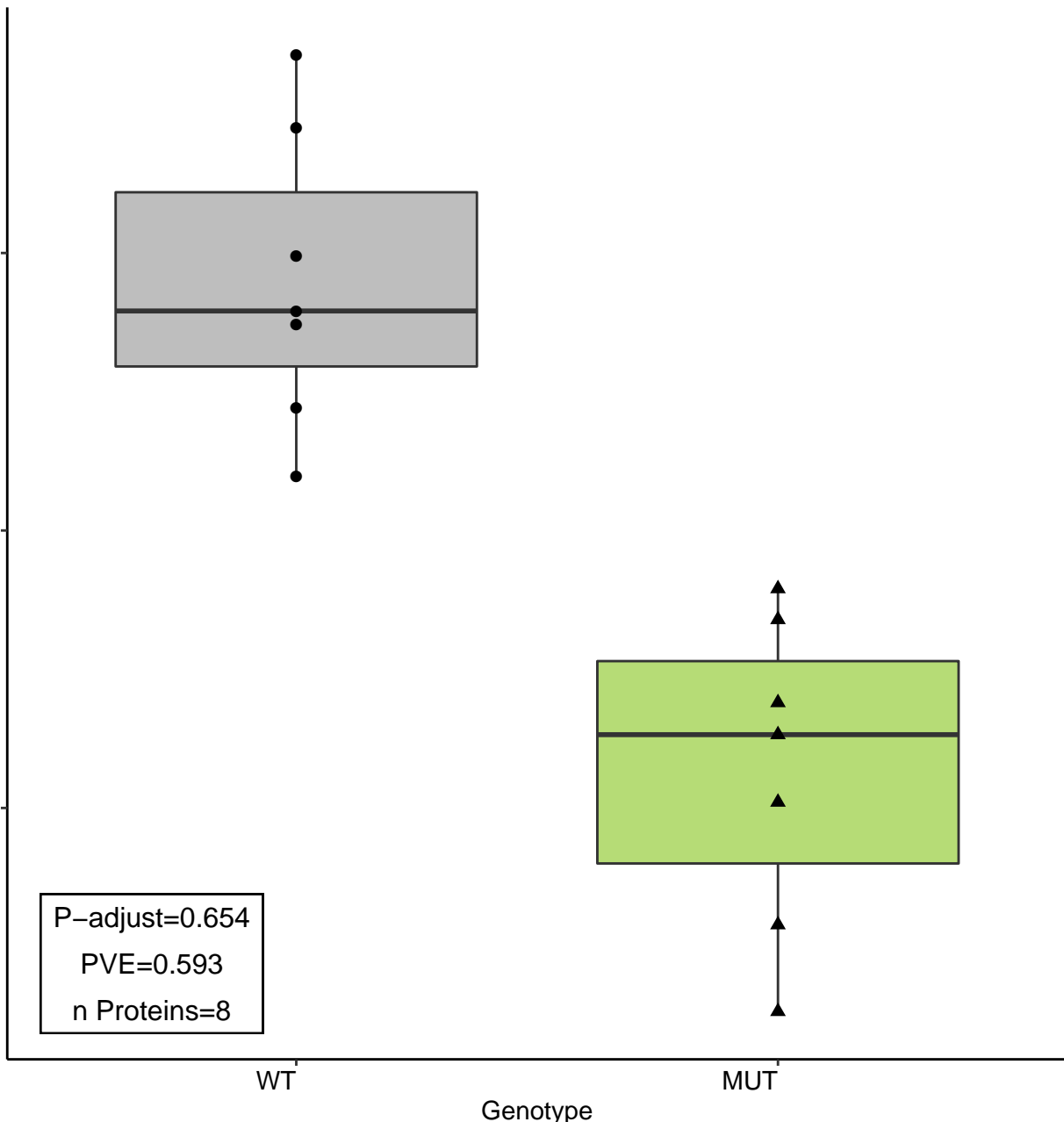
PVE=0.593

n Proteins=8

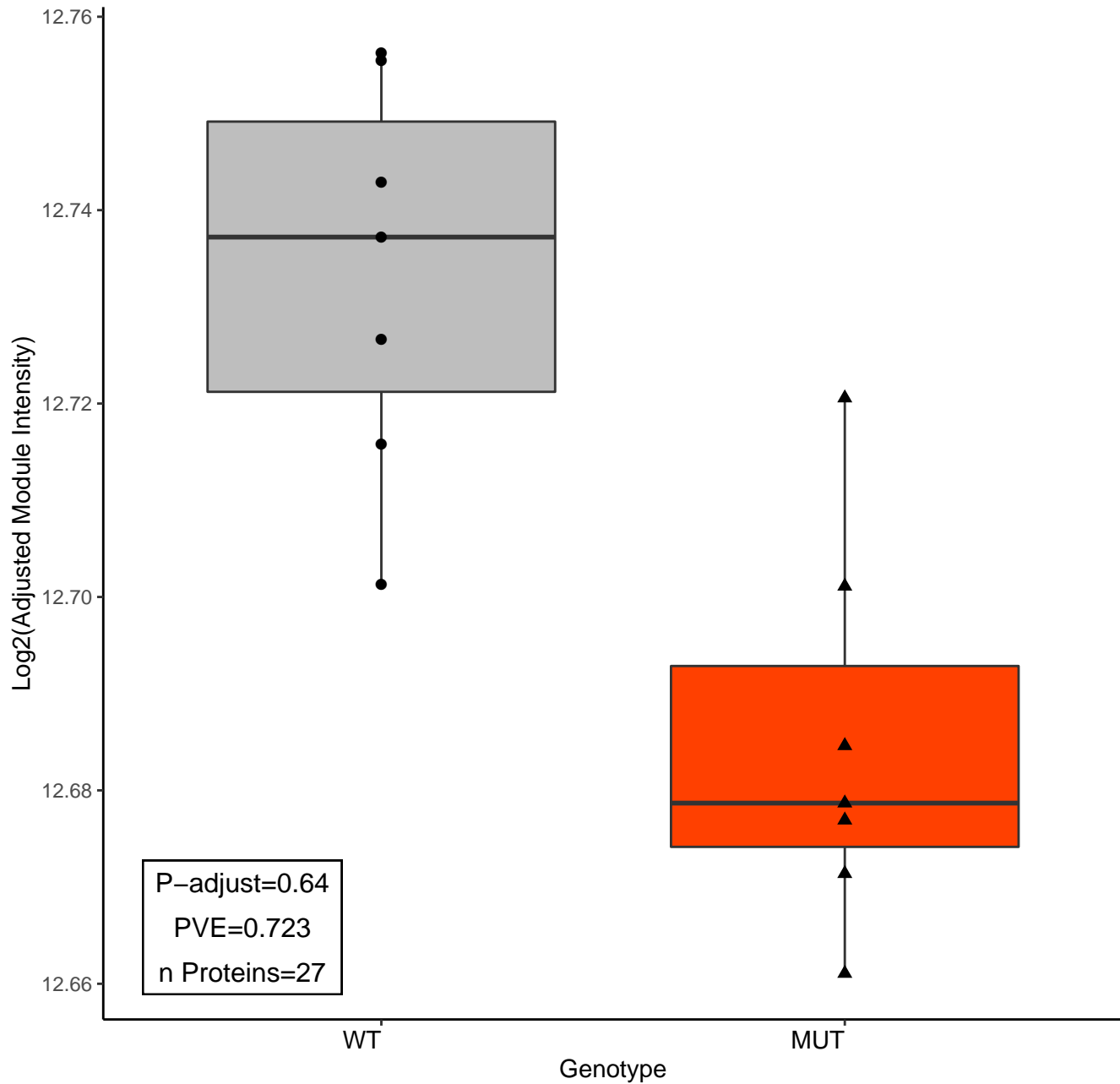
WT

Genotype

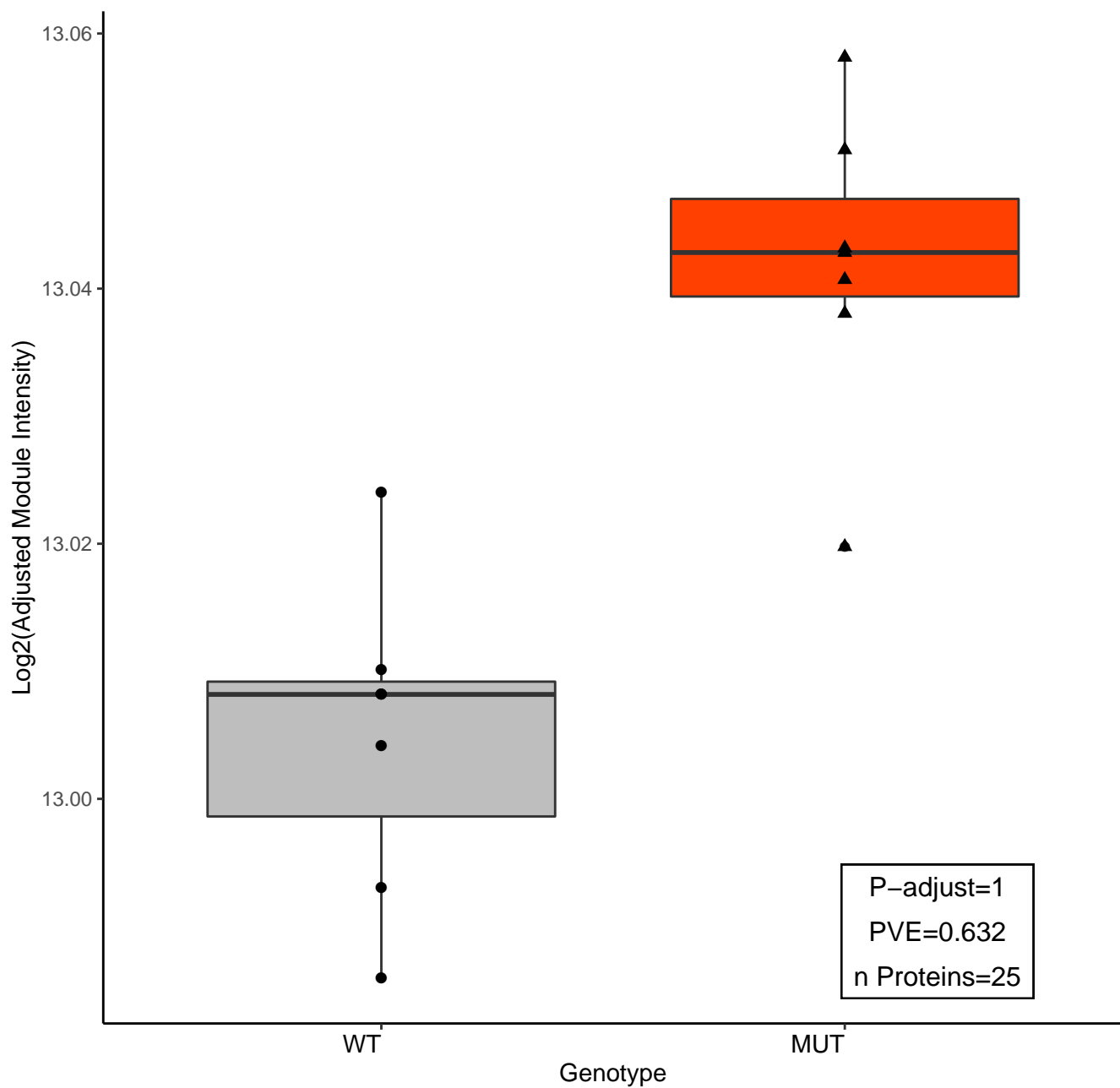
MUT

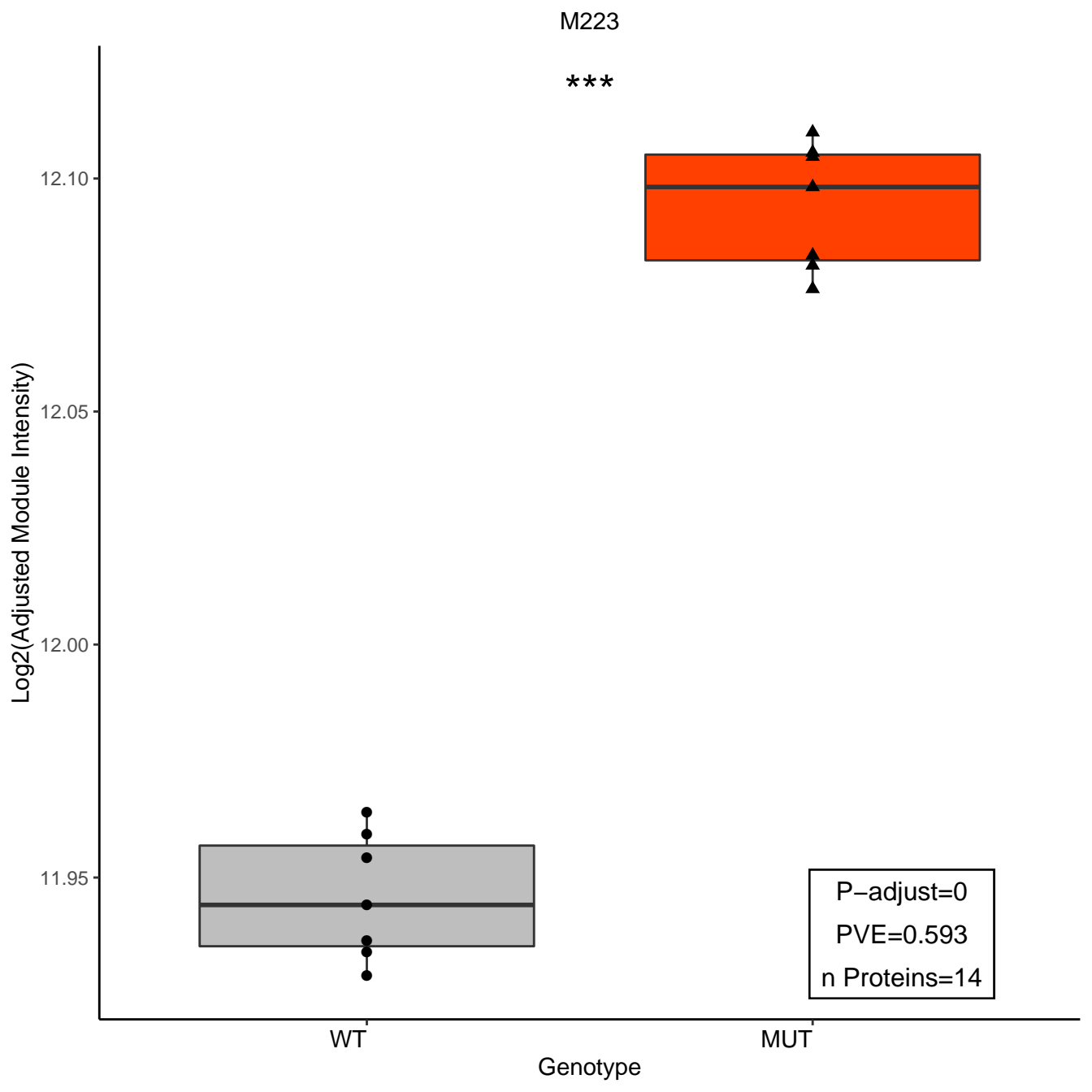


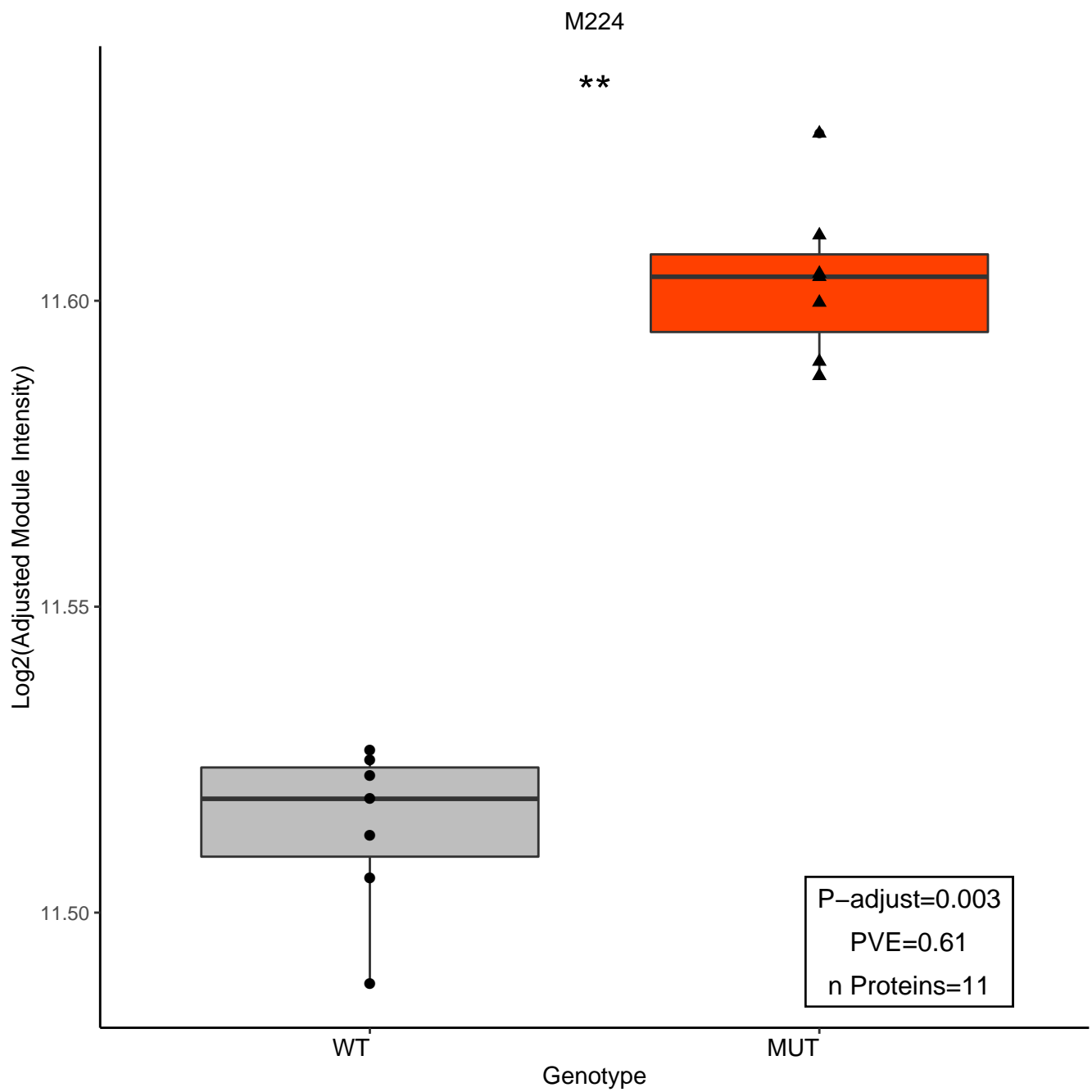
M221



M222







M225

**

Log2(Adjusted Module Intensity)

10.62

10.60

10.58

10.56

10.54

10.52

10.50

P-adjust=0.001

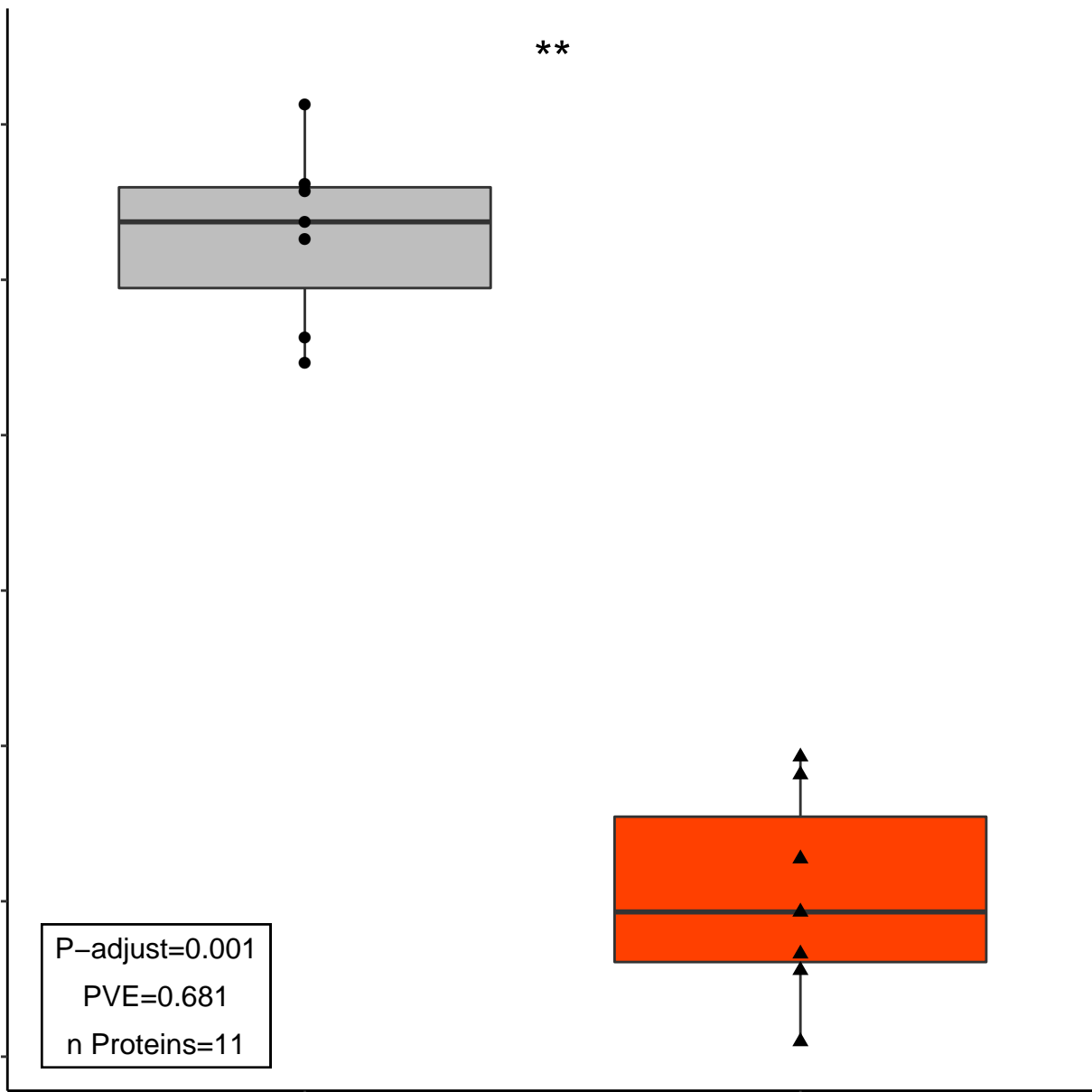
PVE=0.681

n Proteins=11

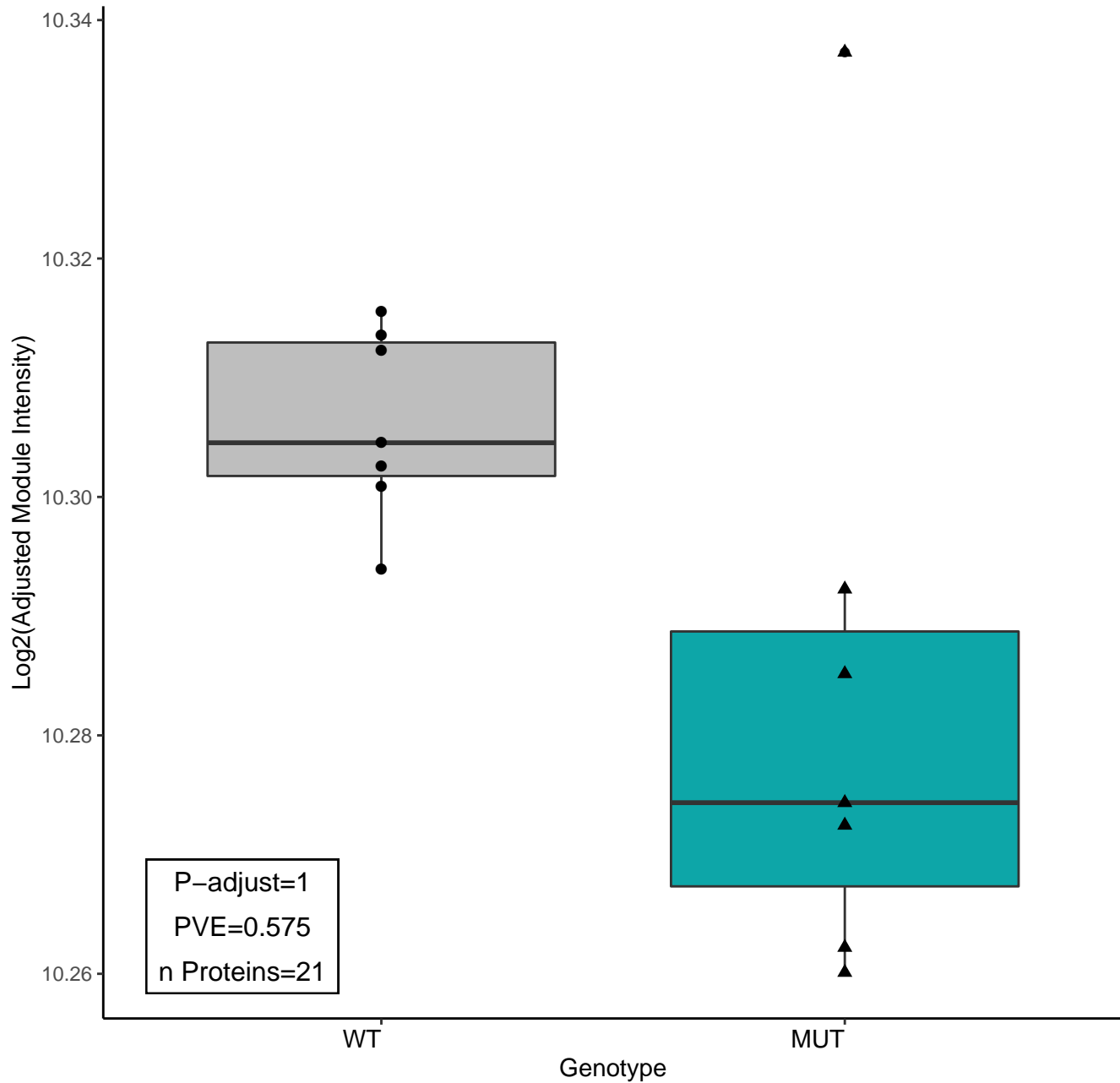
WT

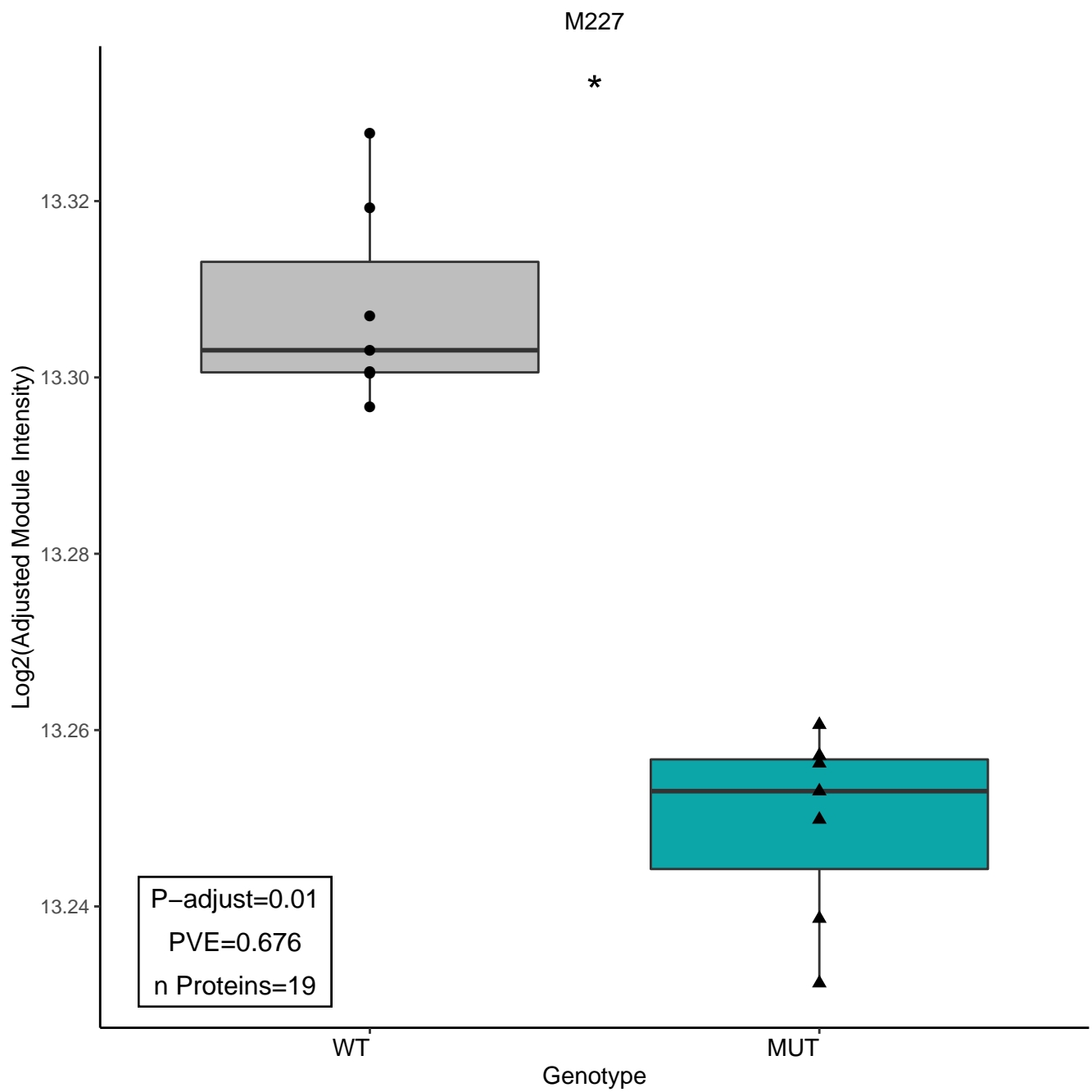
Genotype

MUT

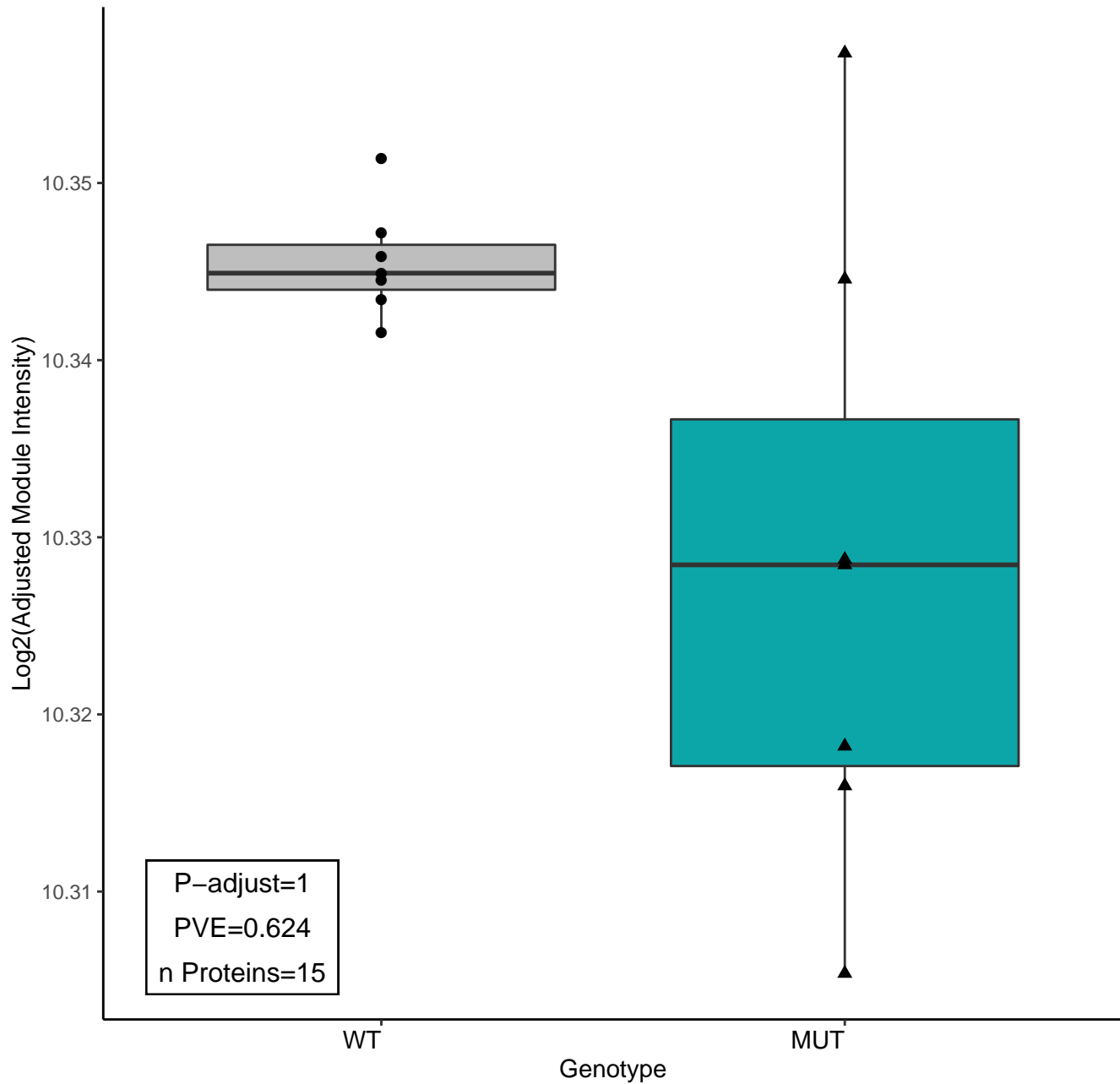


M226

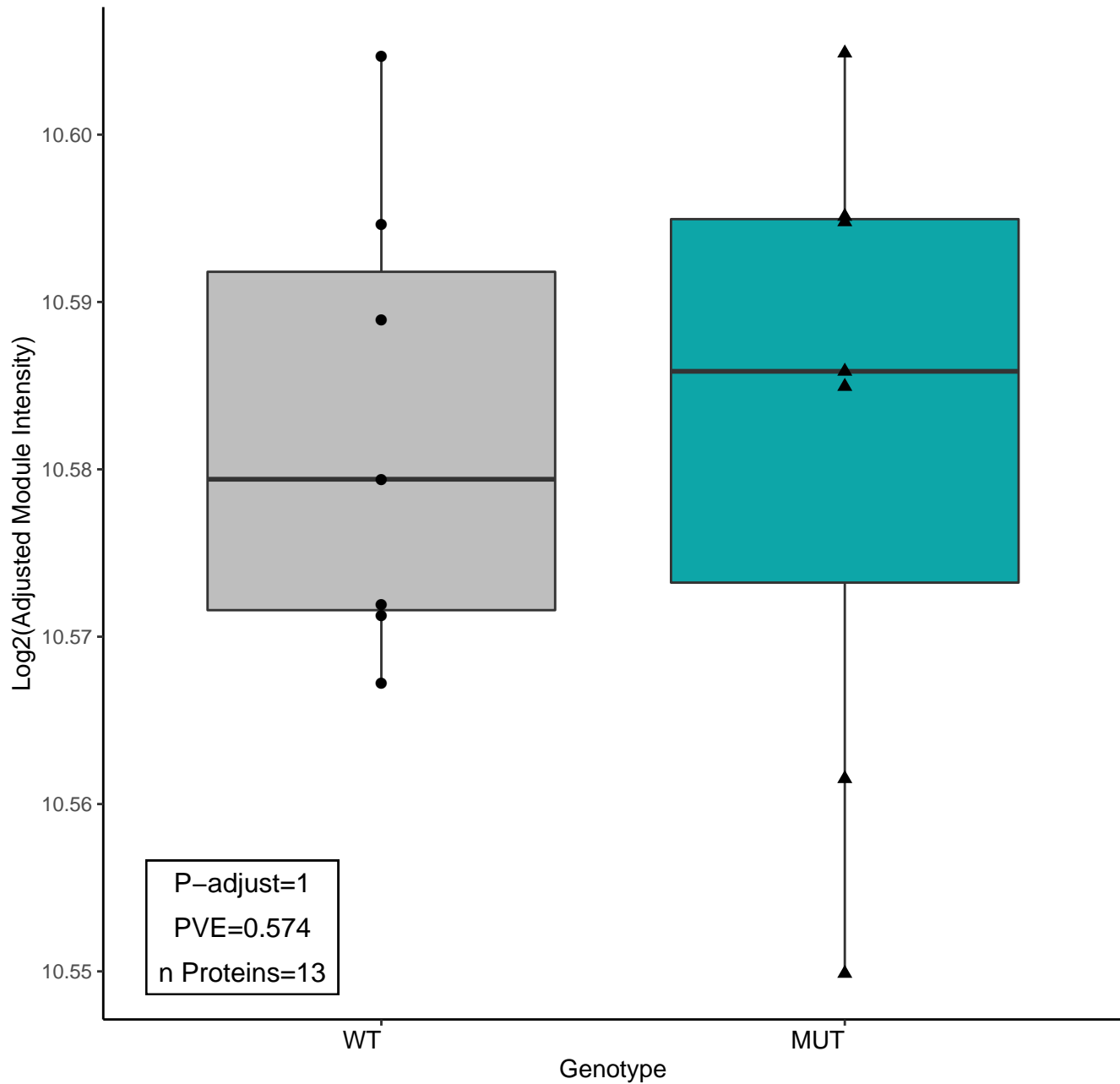




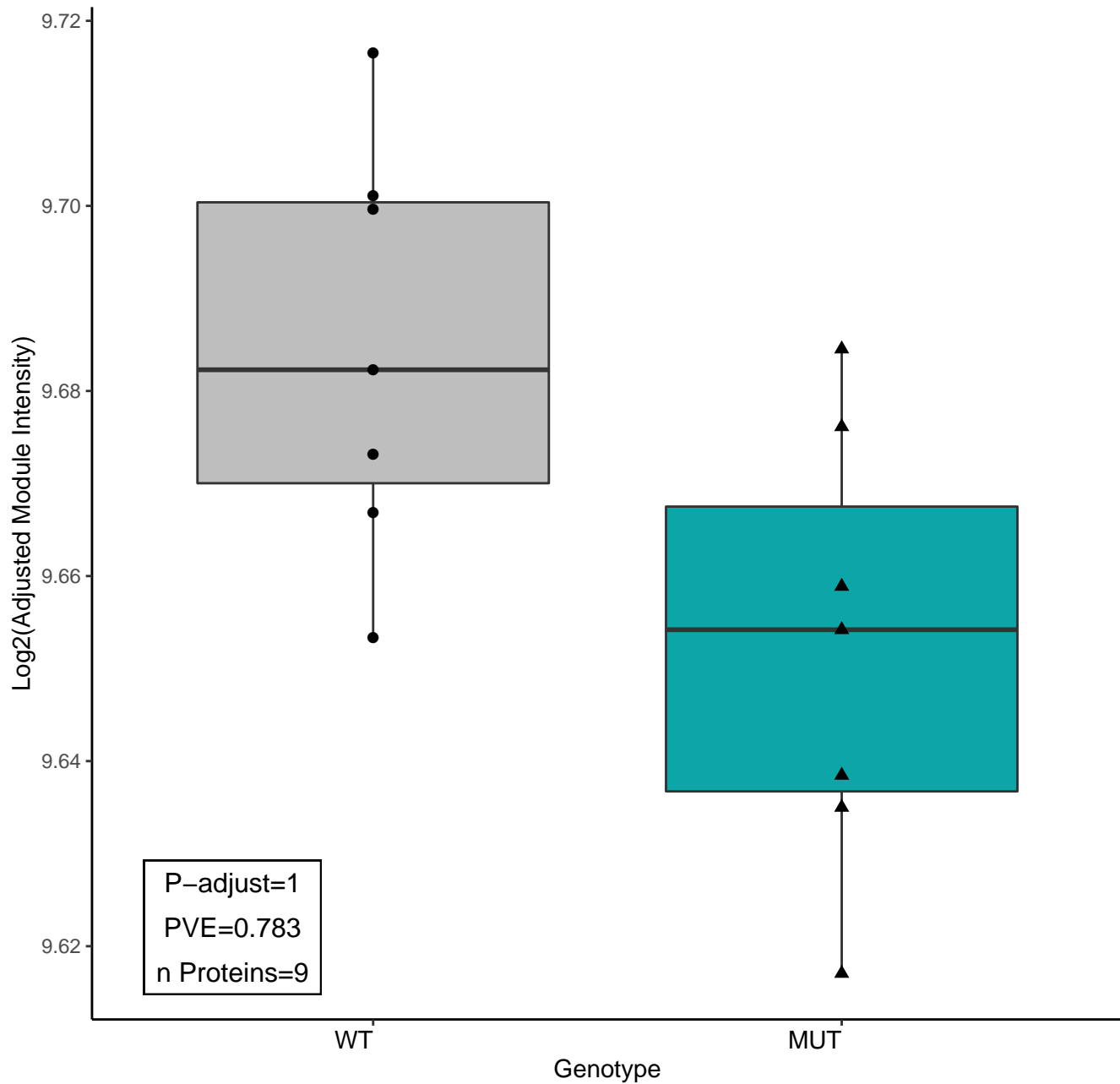
M228

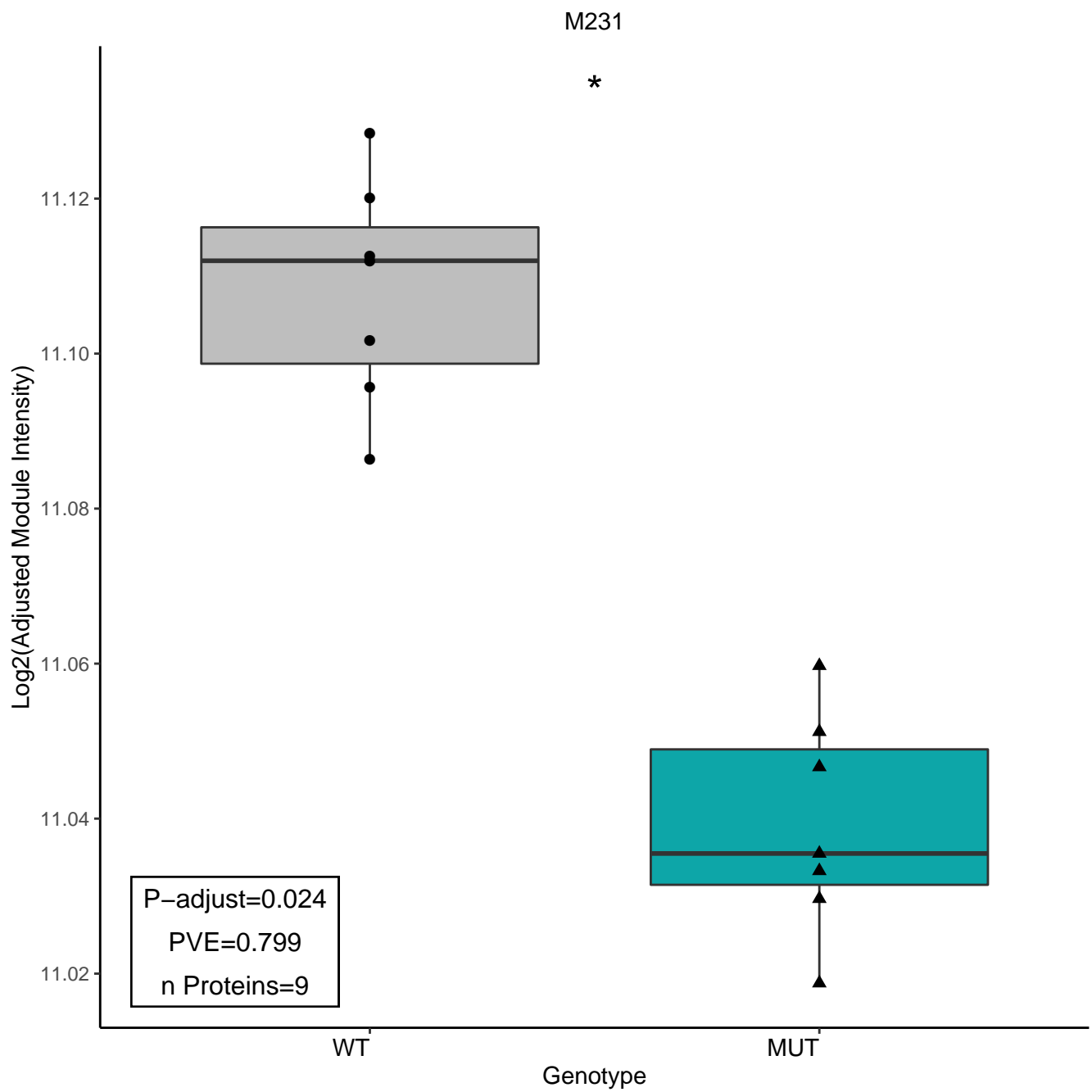


M229

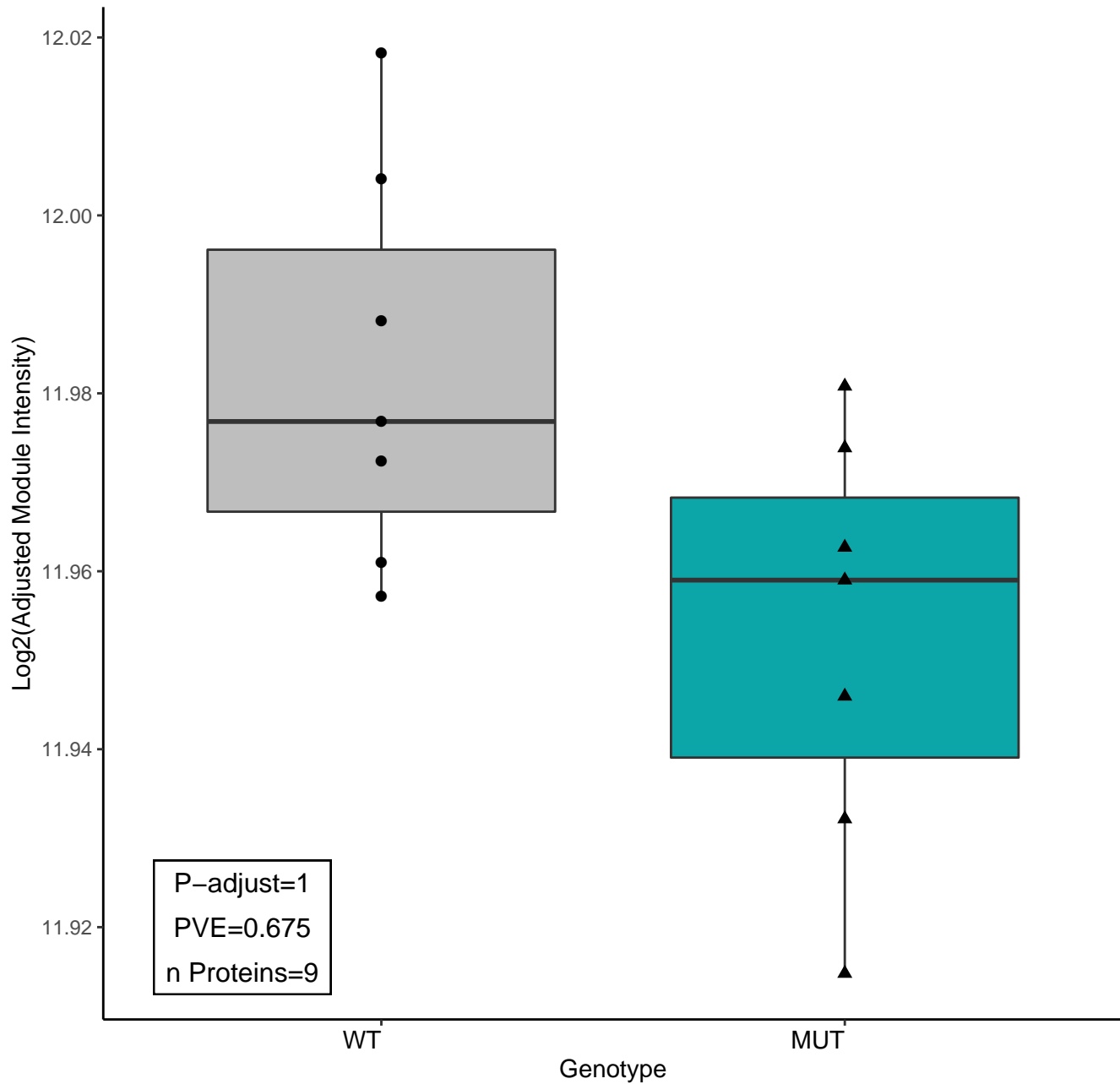


M230





M232



M233

Log2(Adjusted Module Intensity)

11.50

11.48

11.46

11.44

P-adjust=0.078

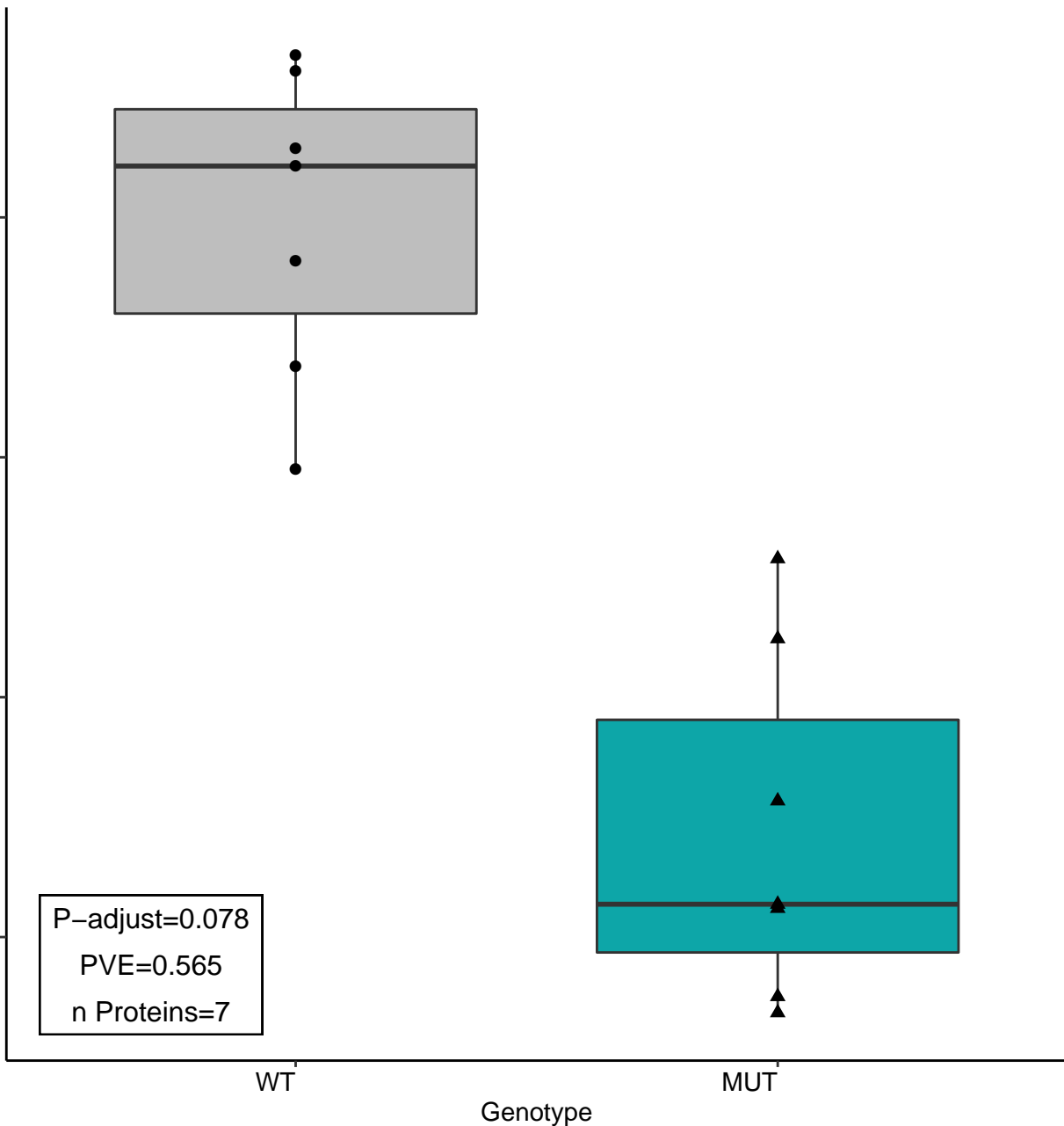
PVE=0.565

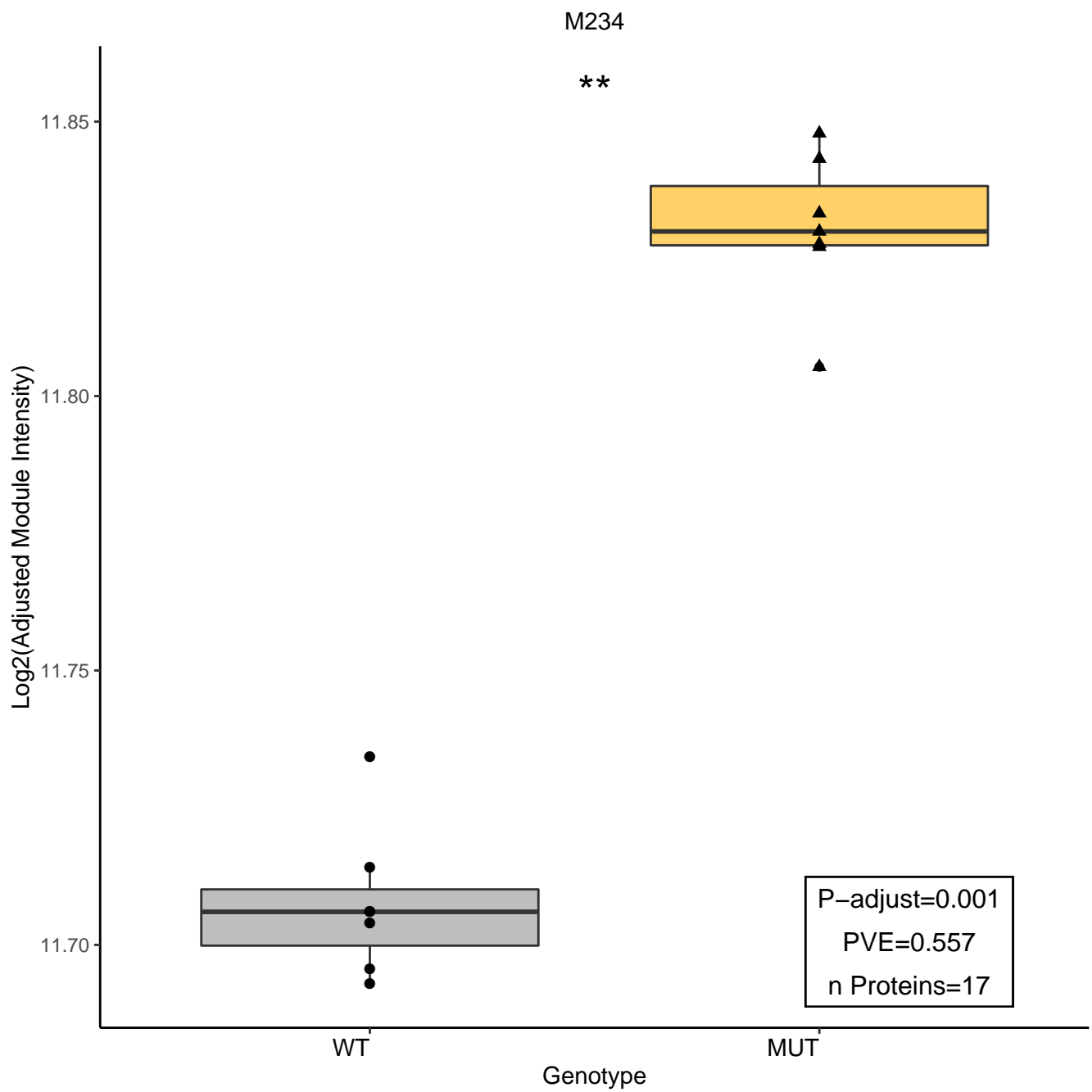
n Proteins=7

WT

Genotype

MUT





M235

Log2(Adjusted Module Intensity)

13.48

13.46

13.44

13.42

13.40

WT

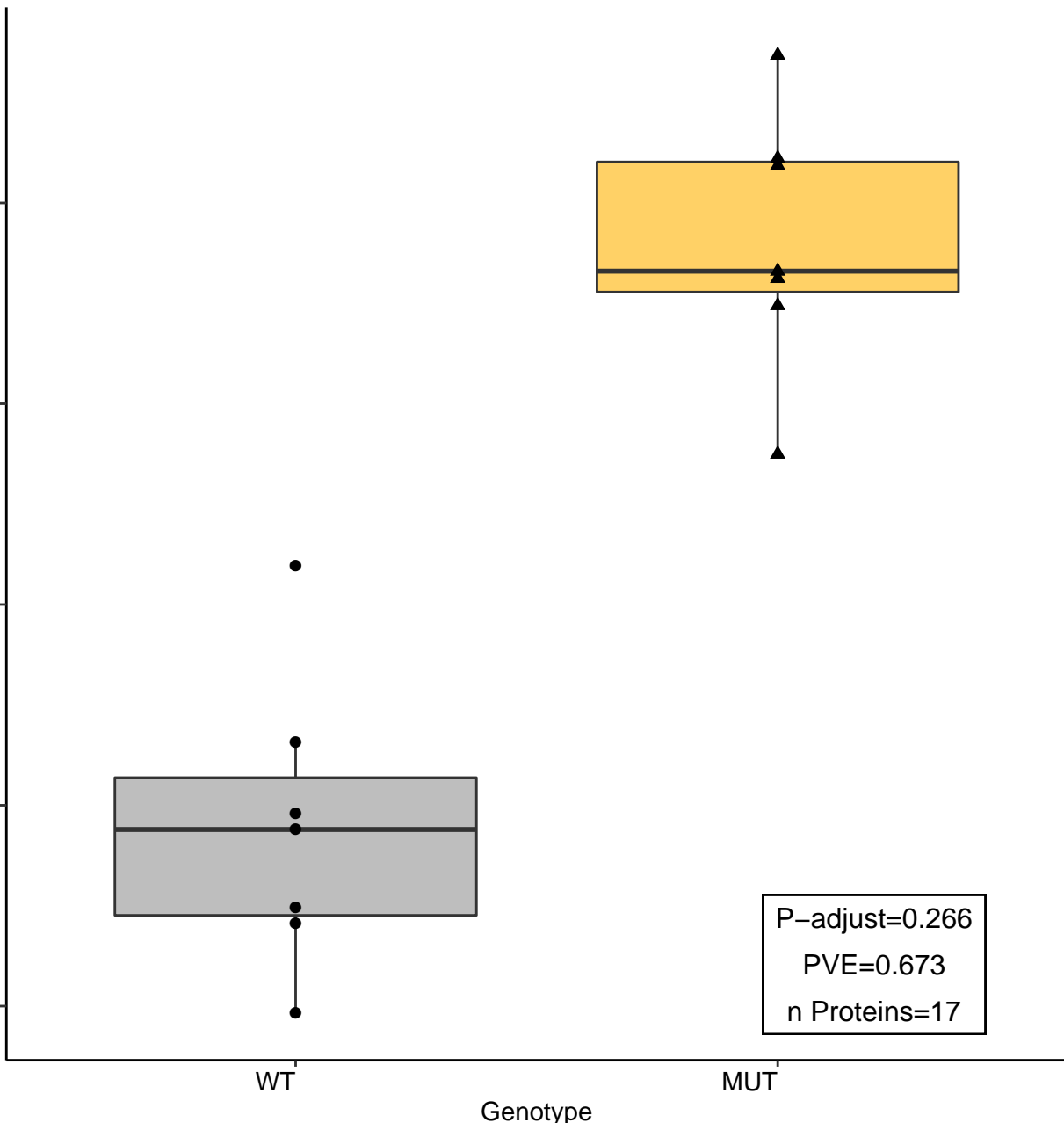
MUT

Genotype

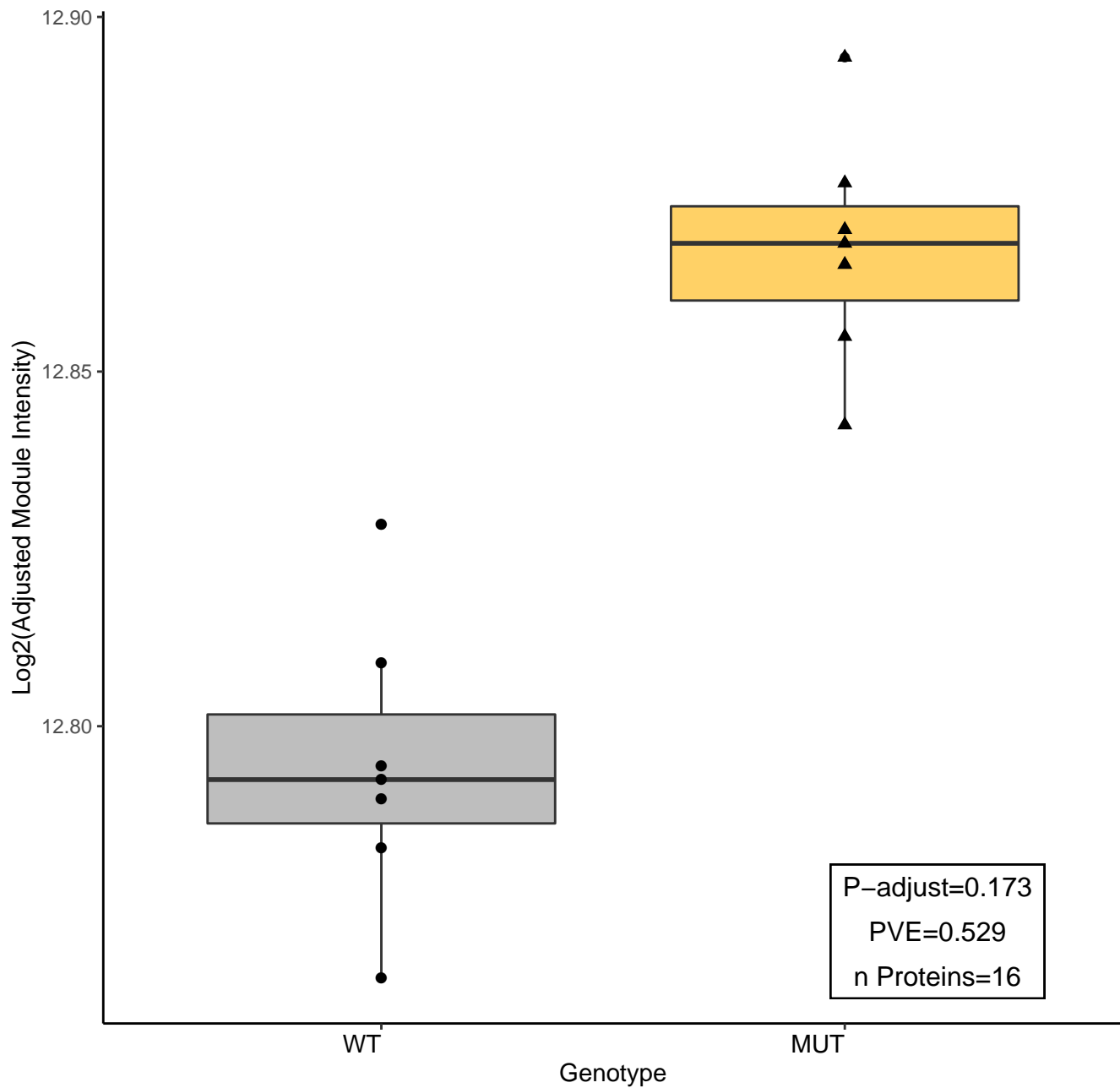
P-adjust=0.266

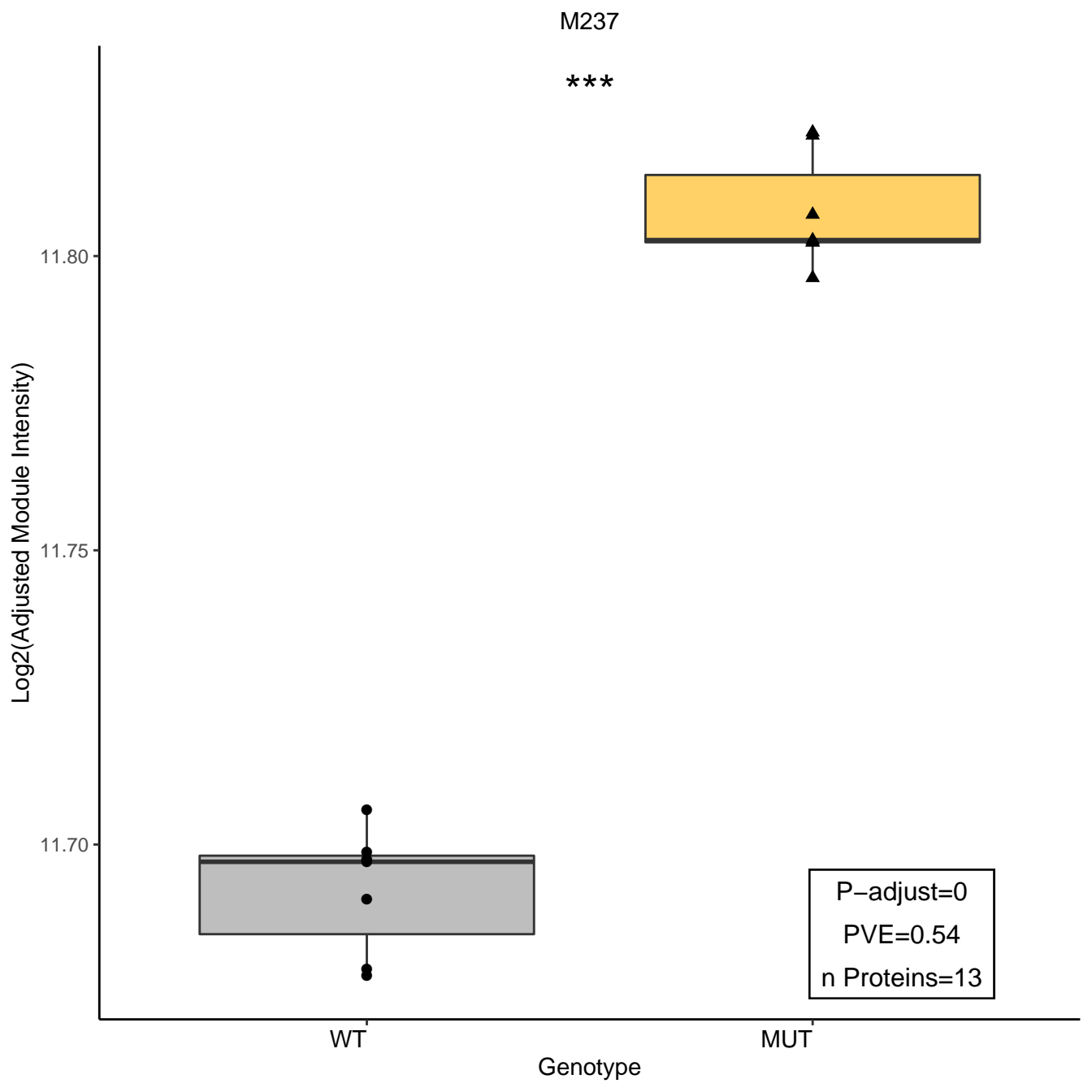
PVE=0.673

n Proteins=17

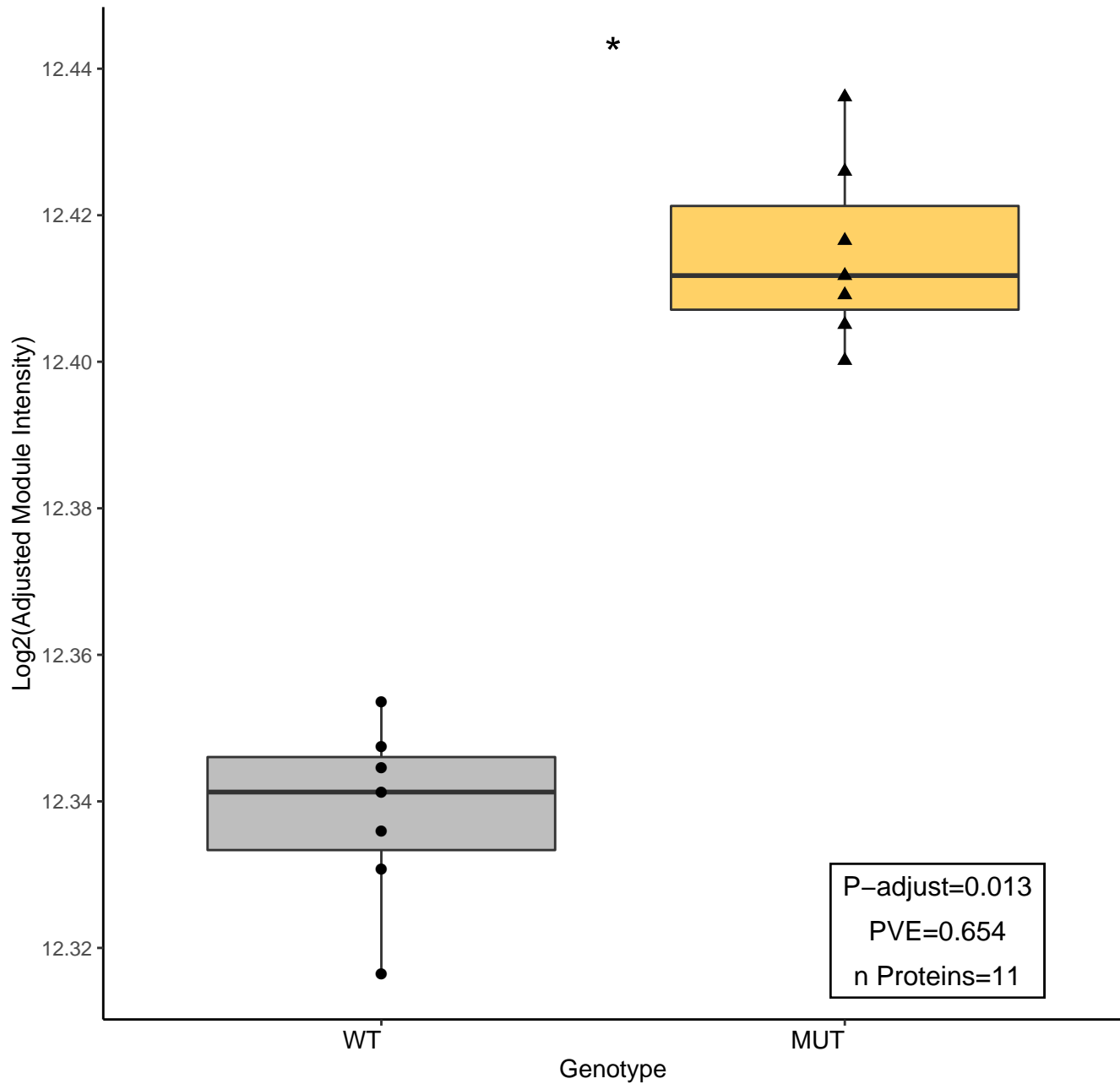


M236





M238



M239

Log2(Adjusted Module Intensity)

11.50

11.45

11.40

11.35

WT

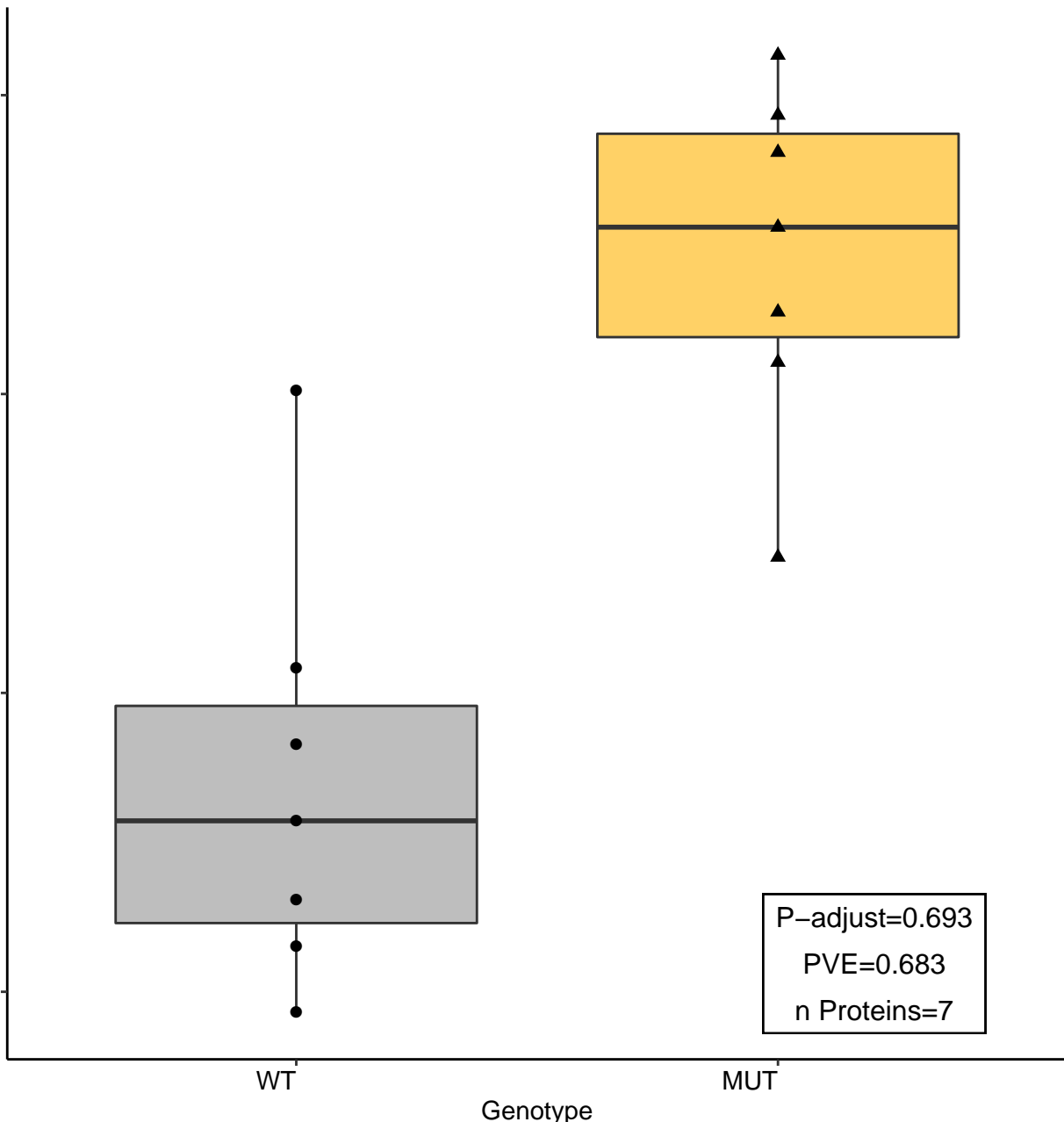
Genotype

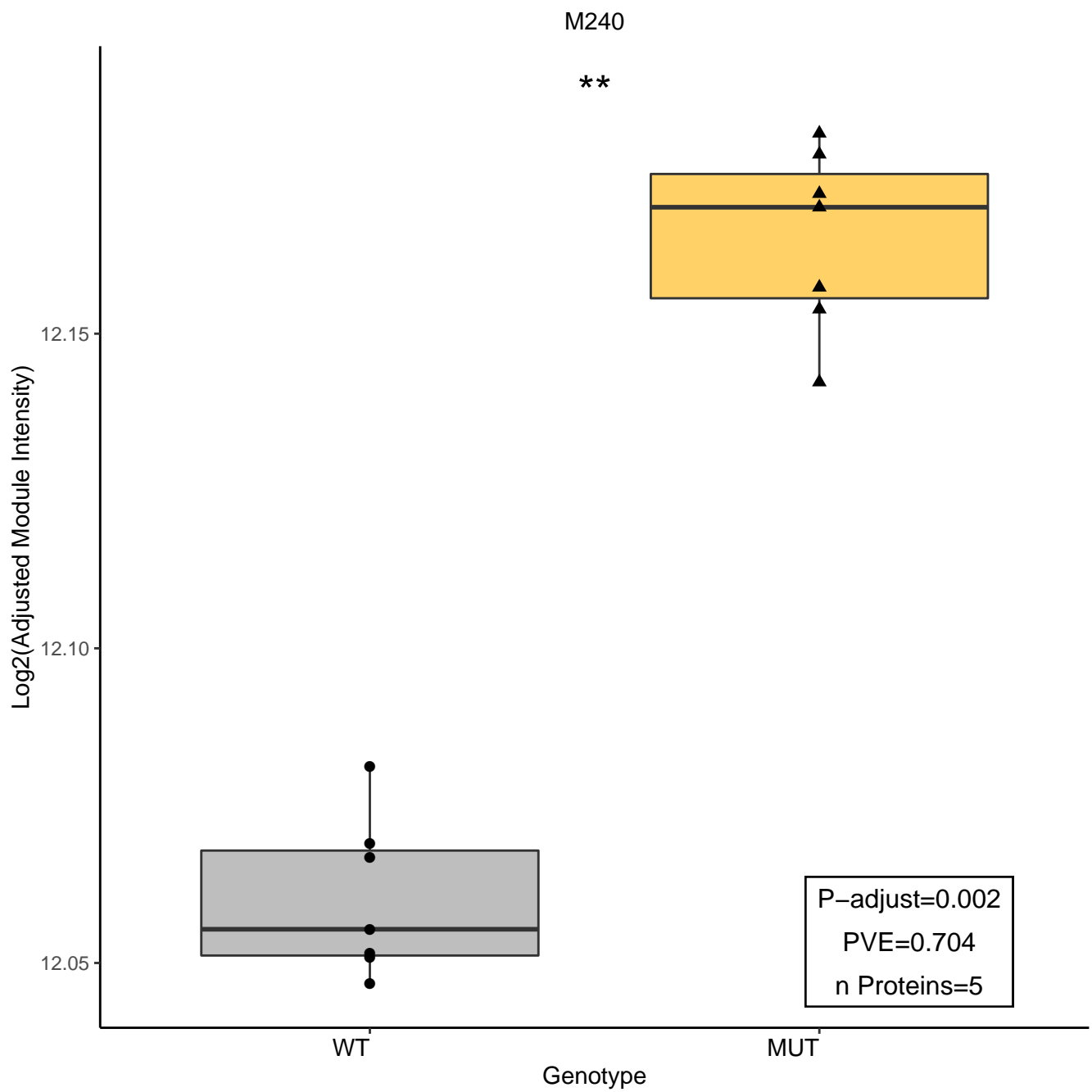
MUT

P-adjust=0.693

PVE=0.683

n Proteins=7





M241

Log2(Adjusted Module Intensity)

12.60

12.55

12.50

WT

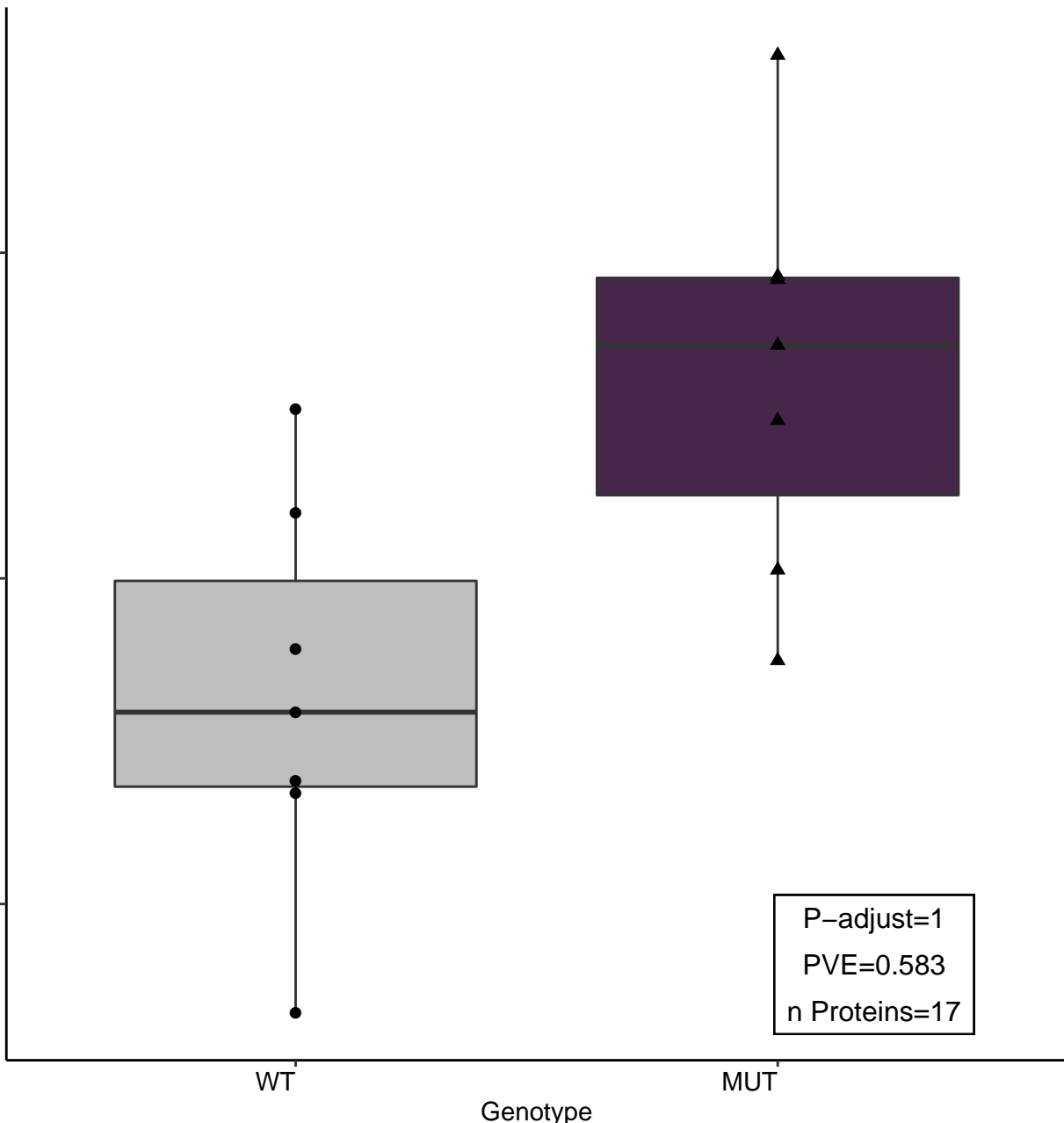
Genotype

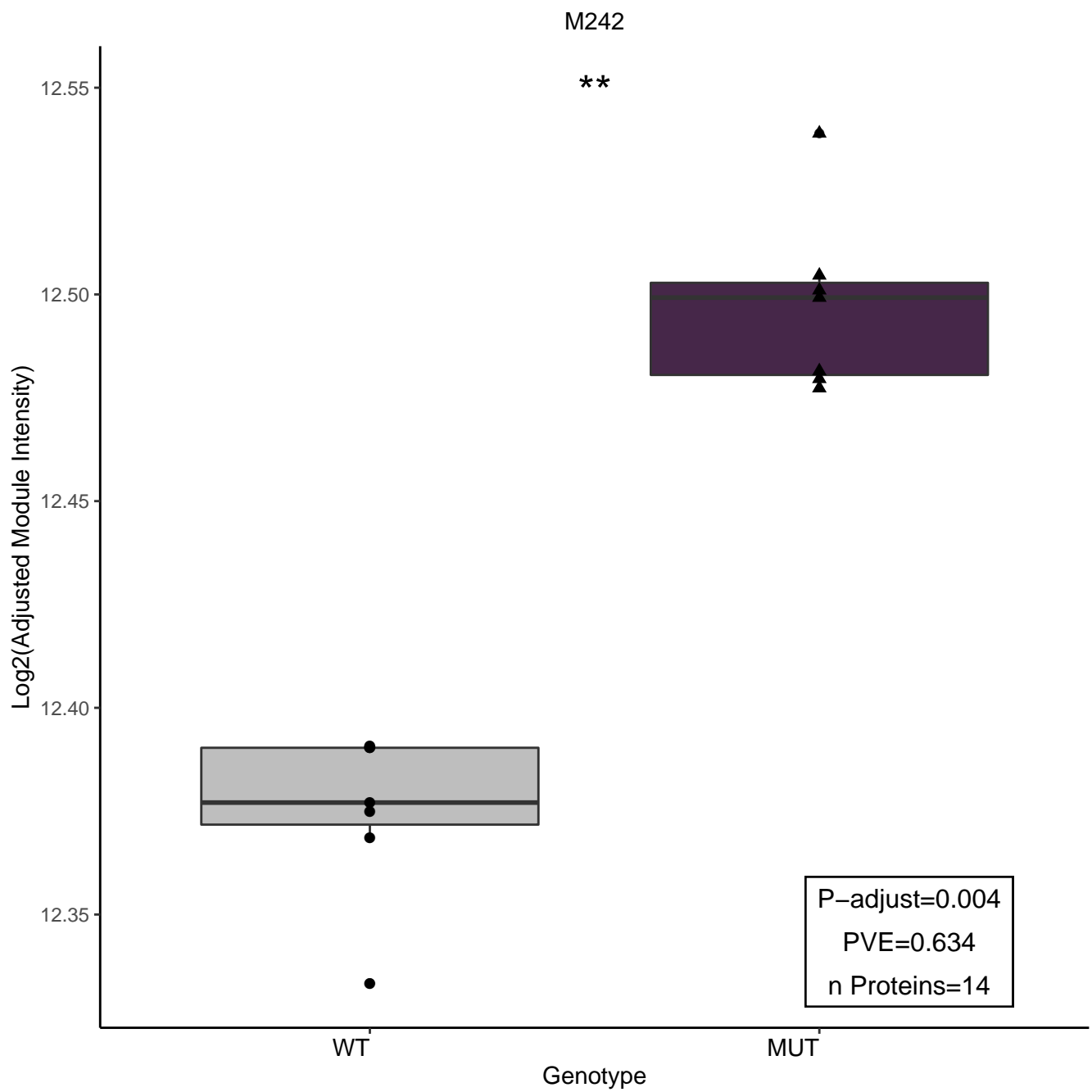
MUT

P-adjust=1

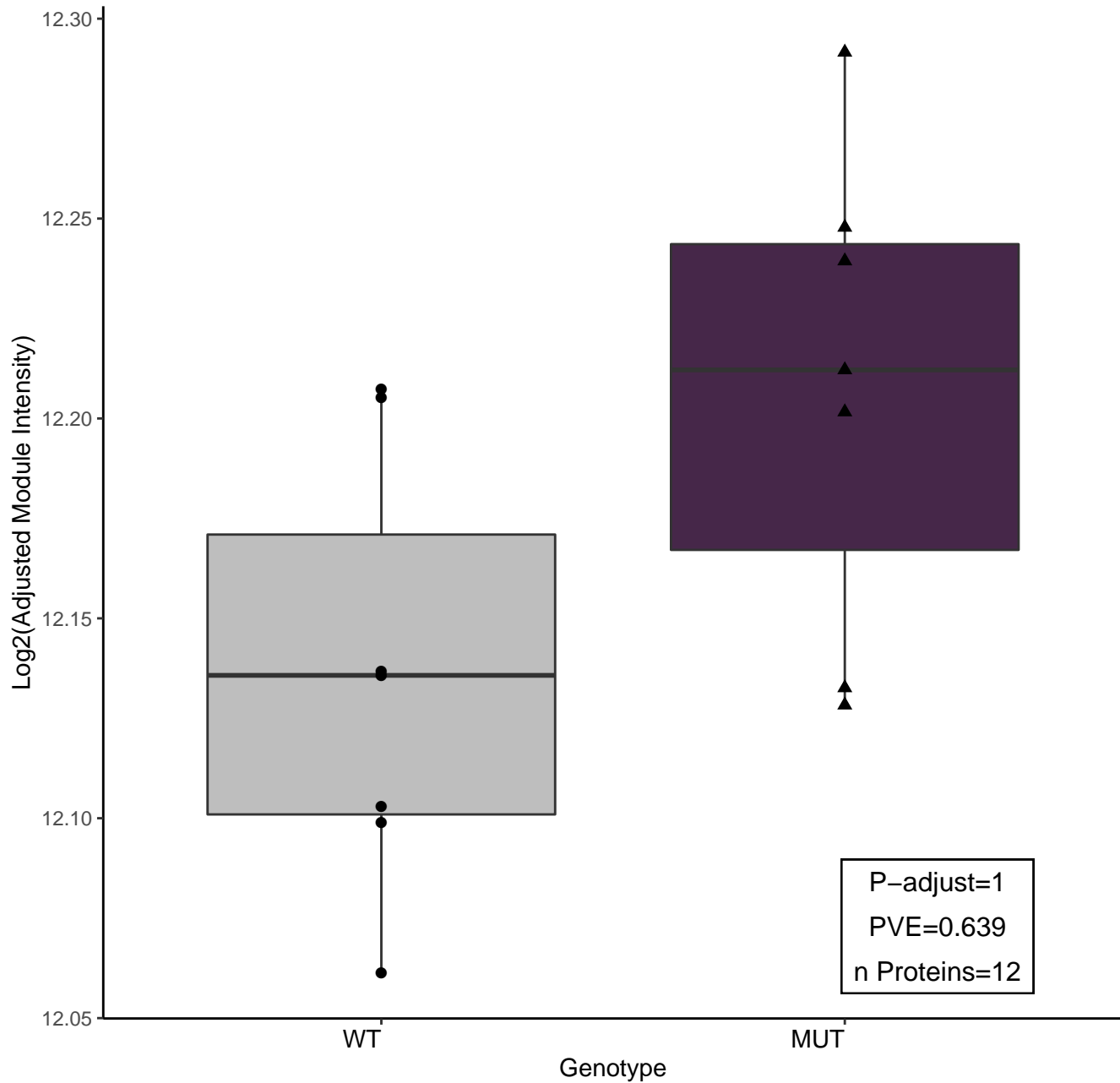
PVE=0.583

n Proteins=17

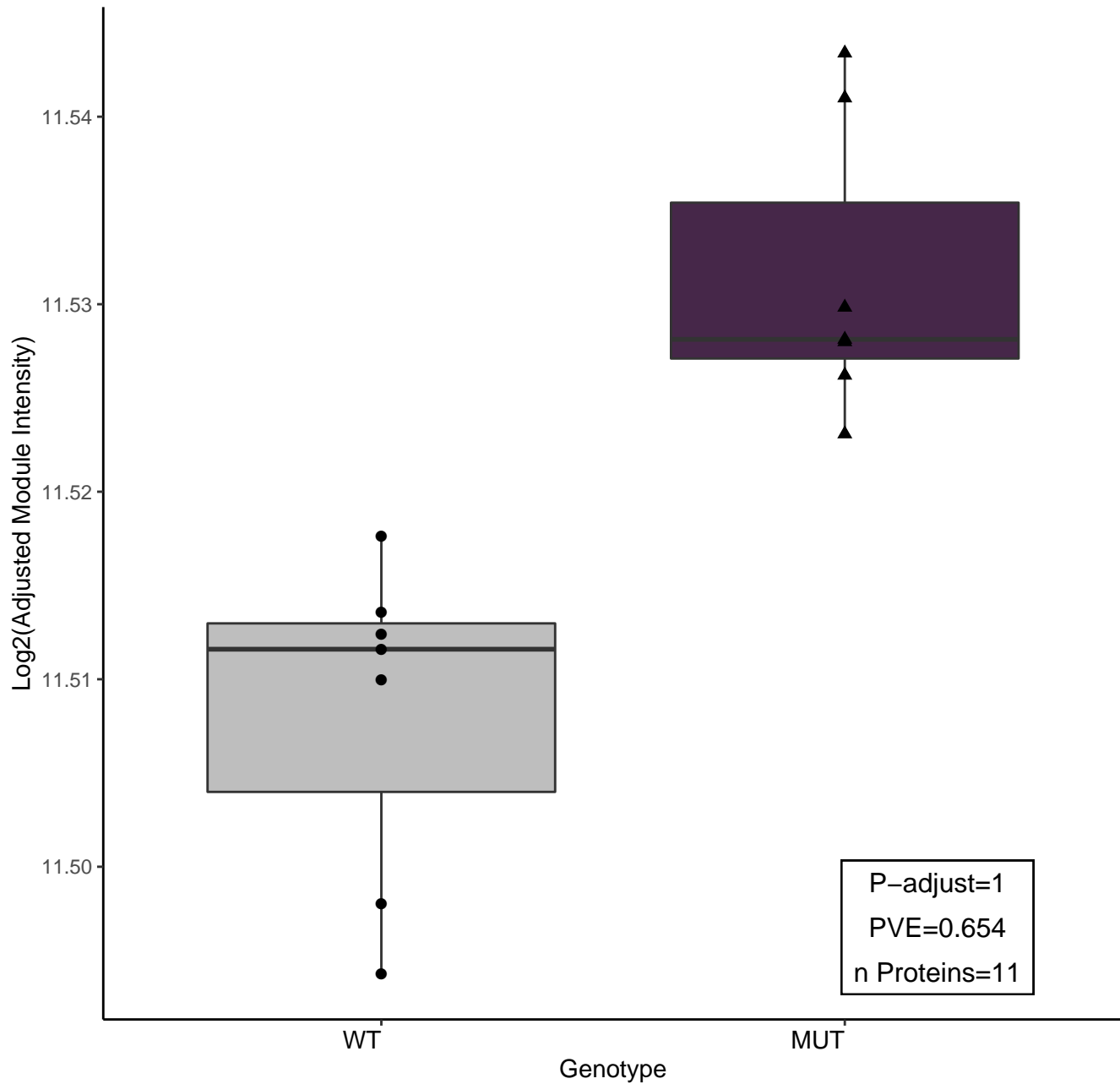




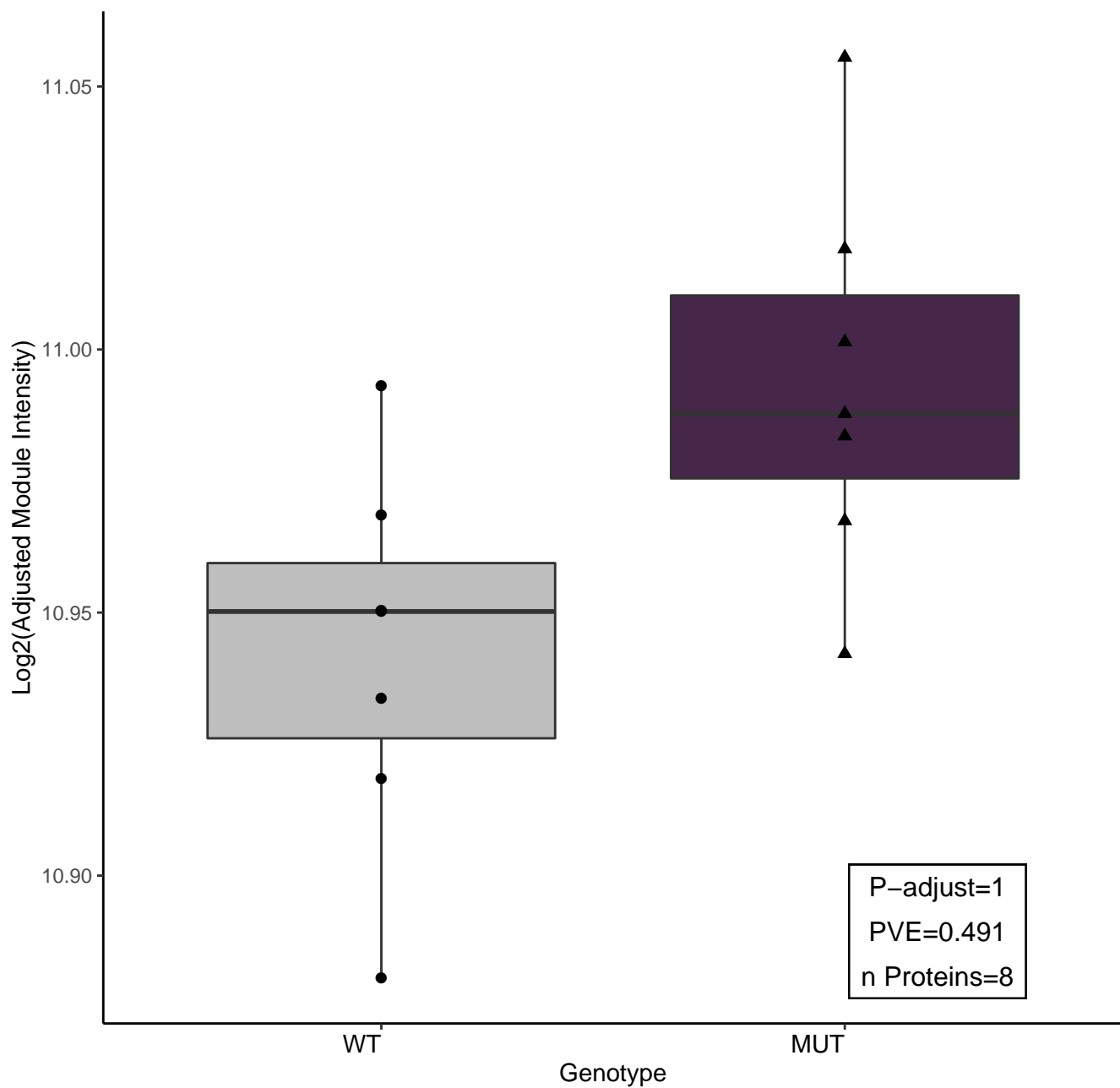
M243



M244



M245



M246

Log2(Adjusted Module Intensity)

13.86
13.84
13.82
13.80
13.78
13.76

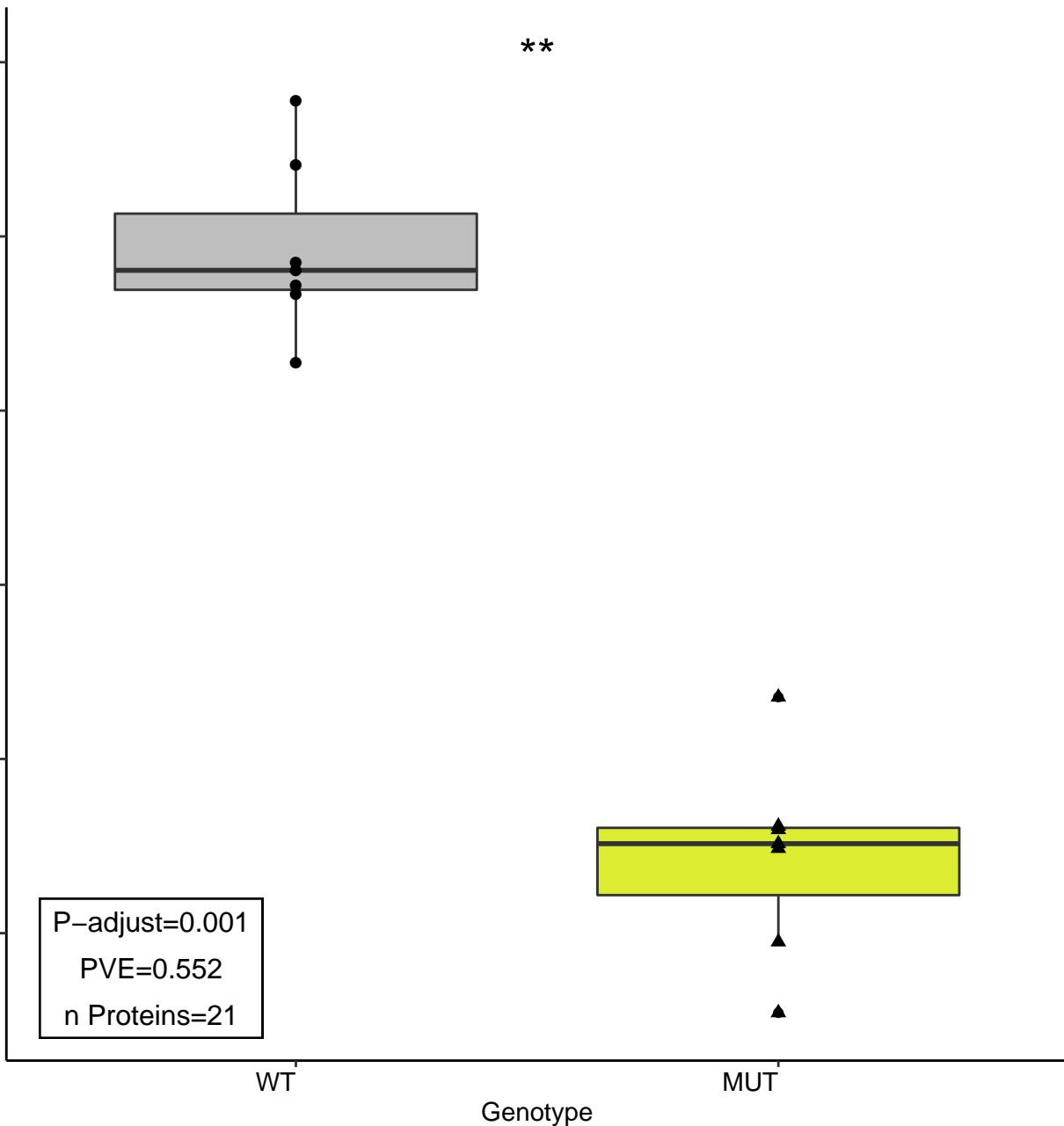
**

P-adjust=0.001
PVE=0.552
n Proteins=21

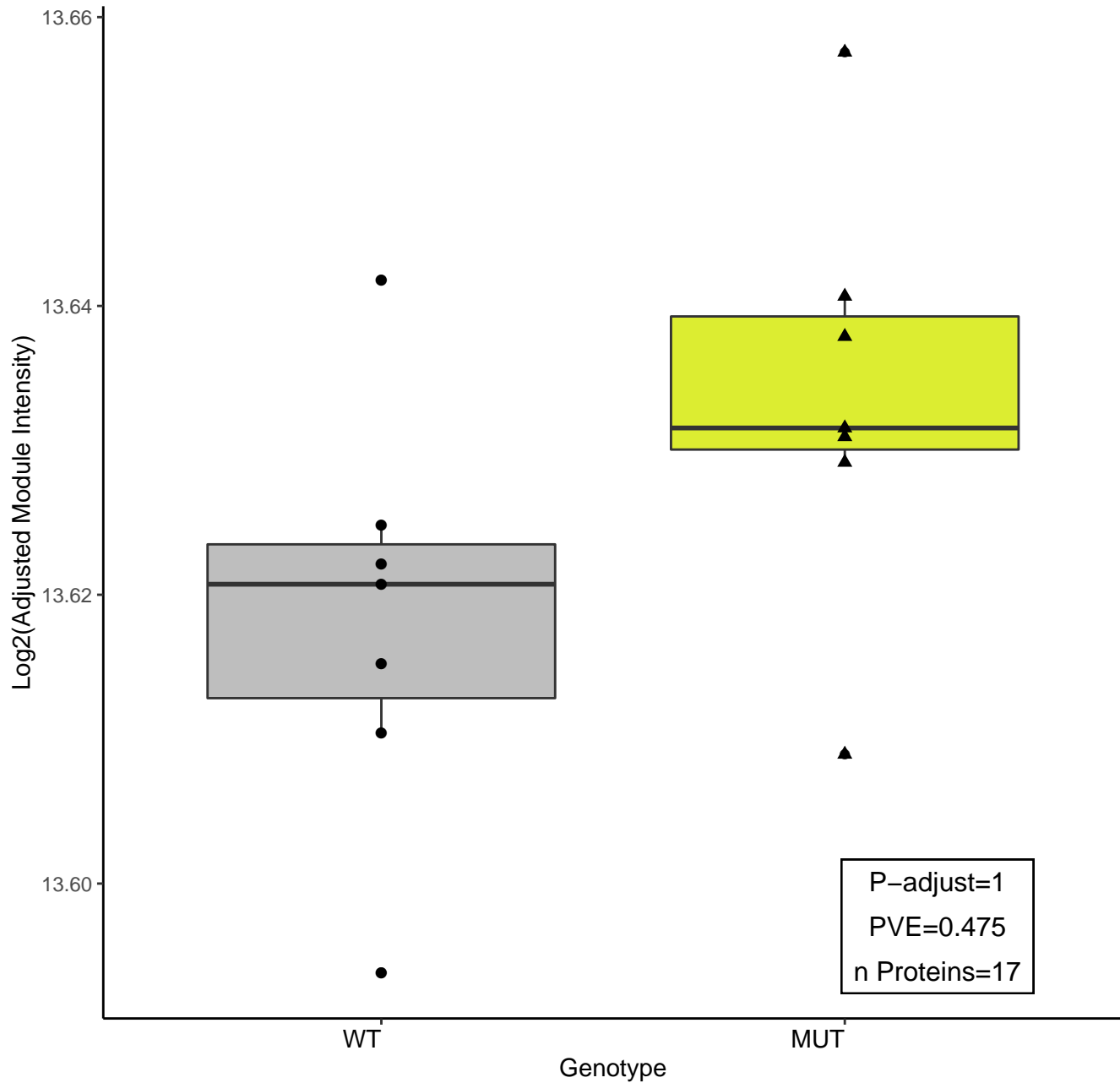
WT

Genotype

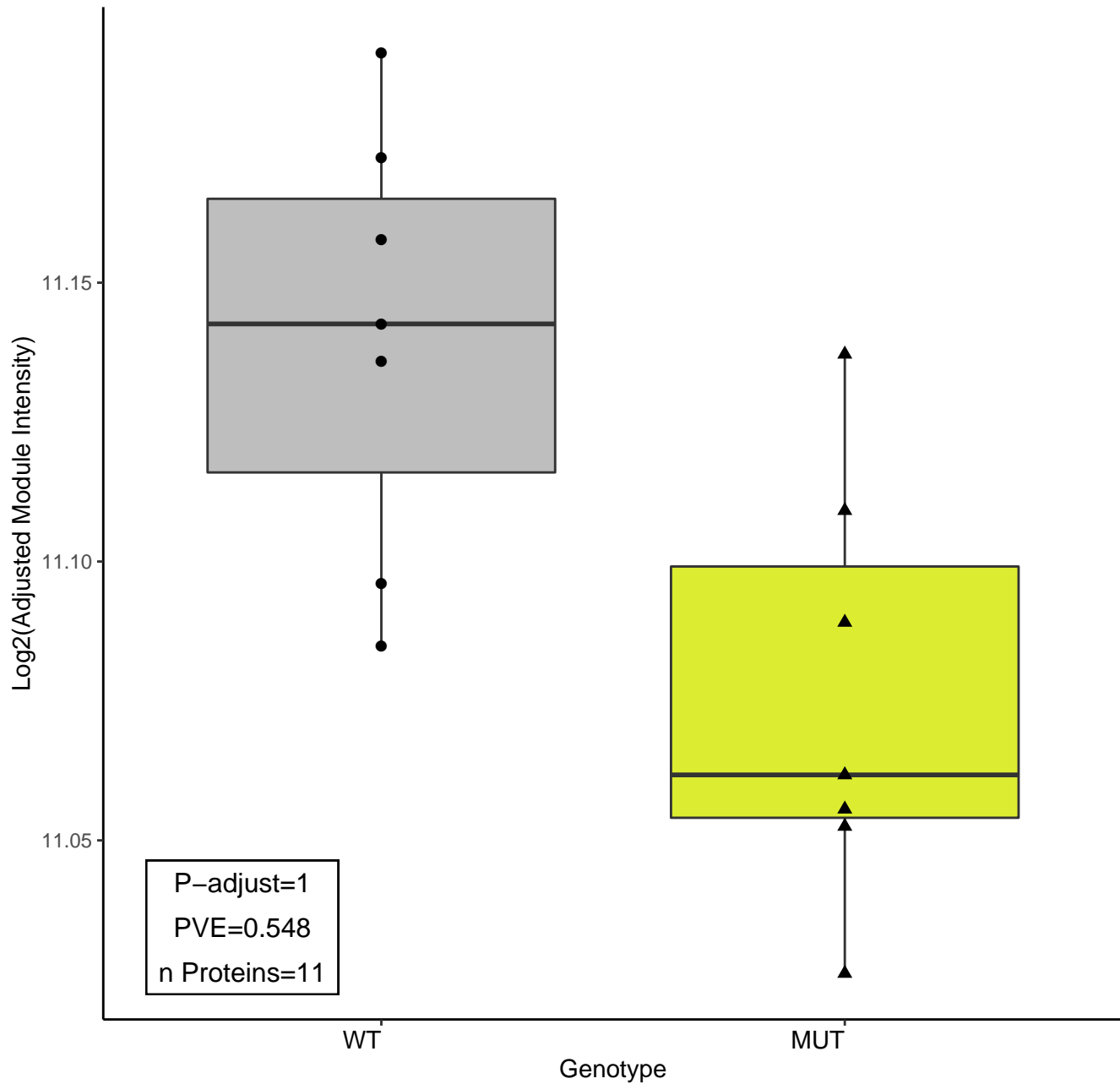
MUT



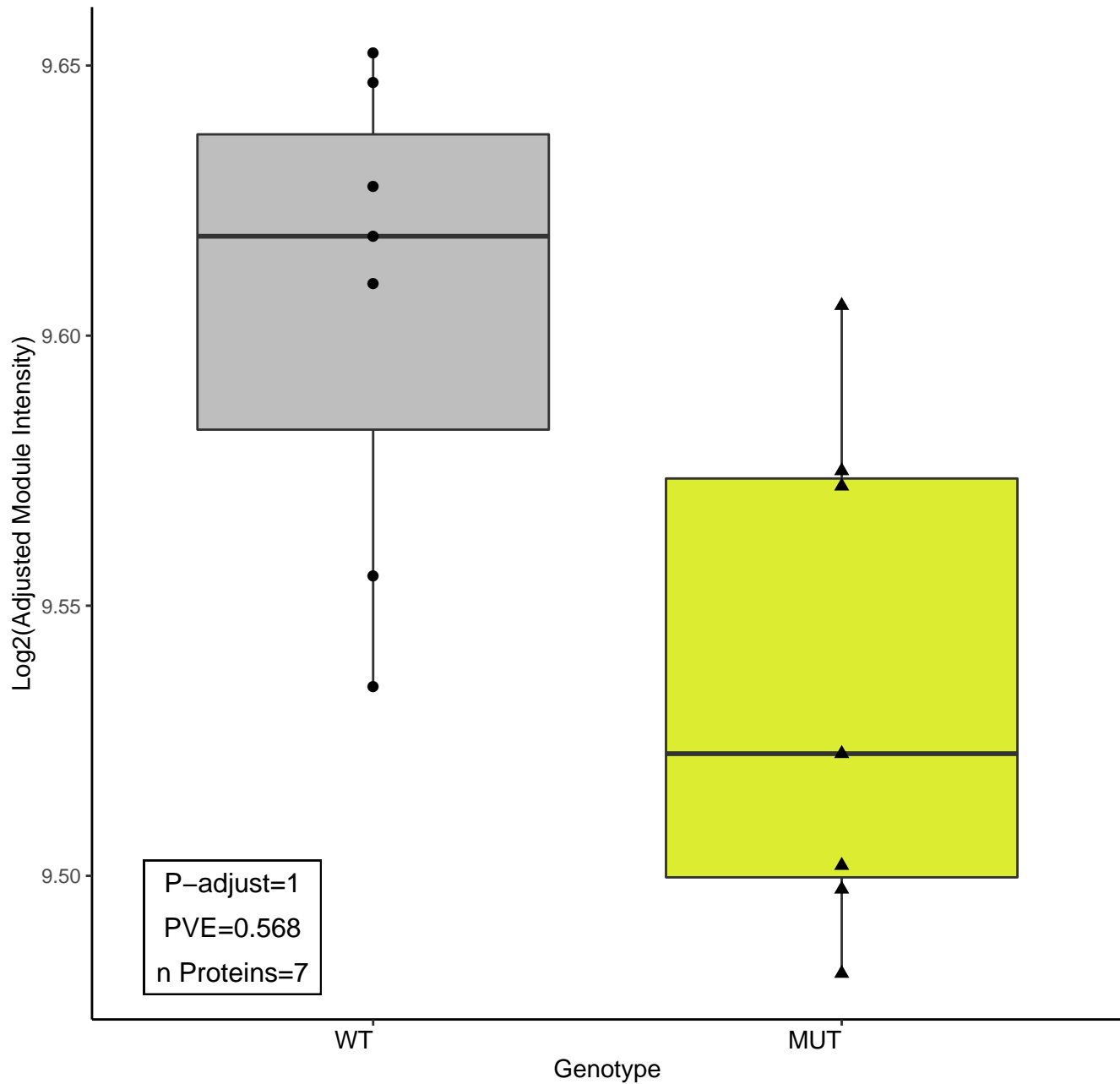
M247



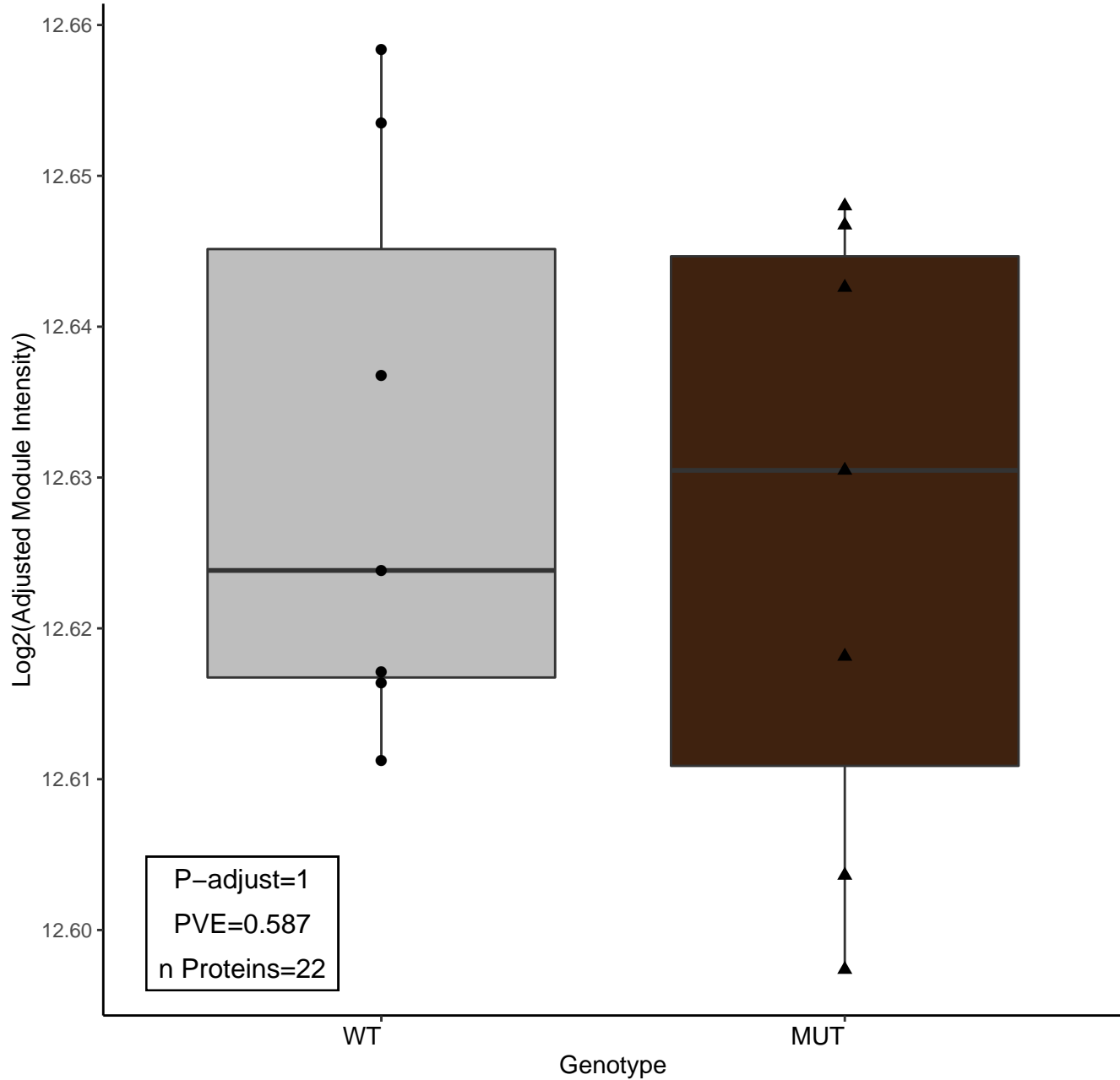
M248



M249



M250



M251

