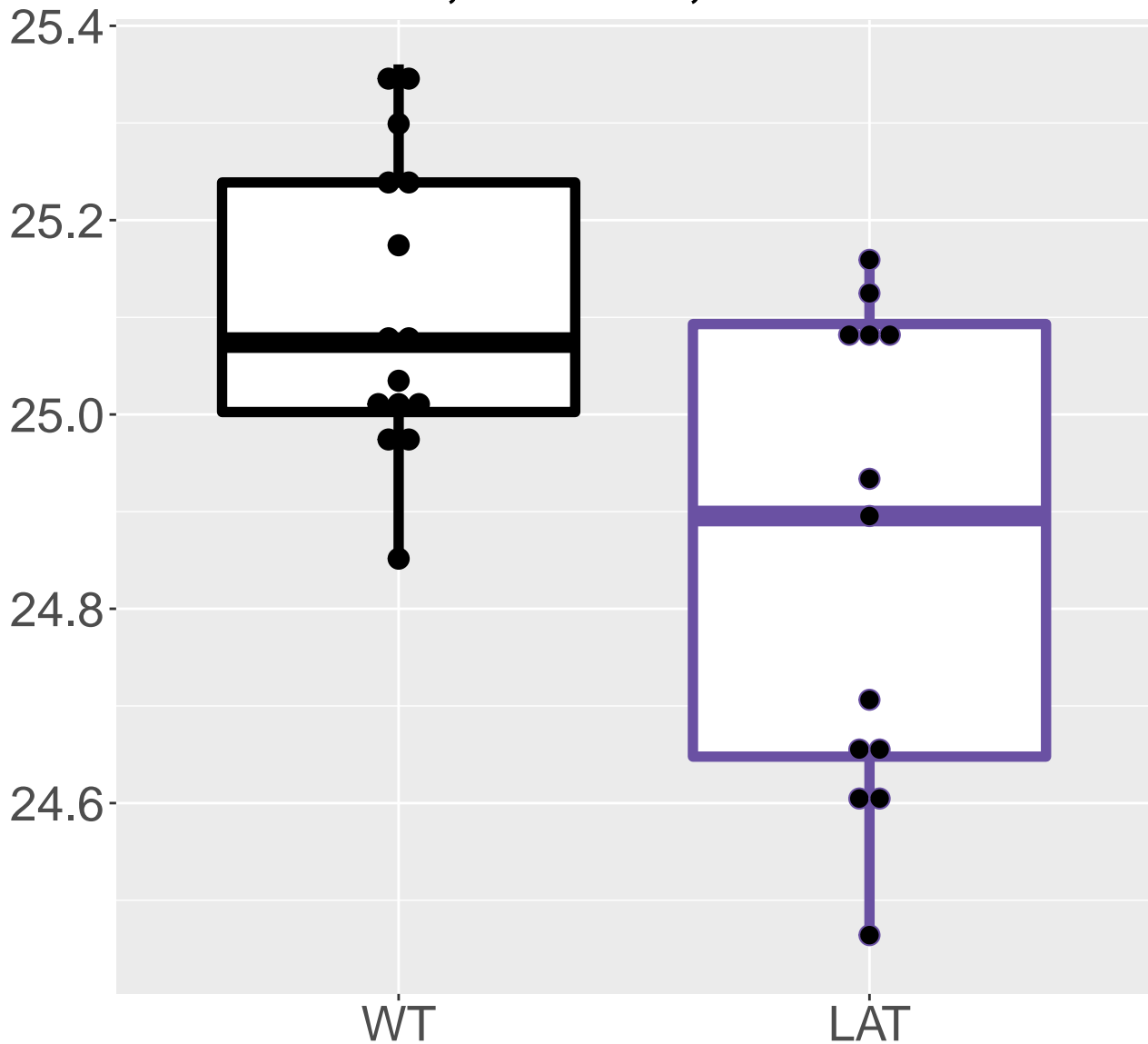
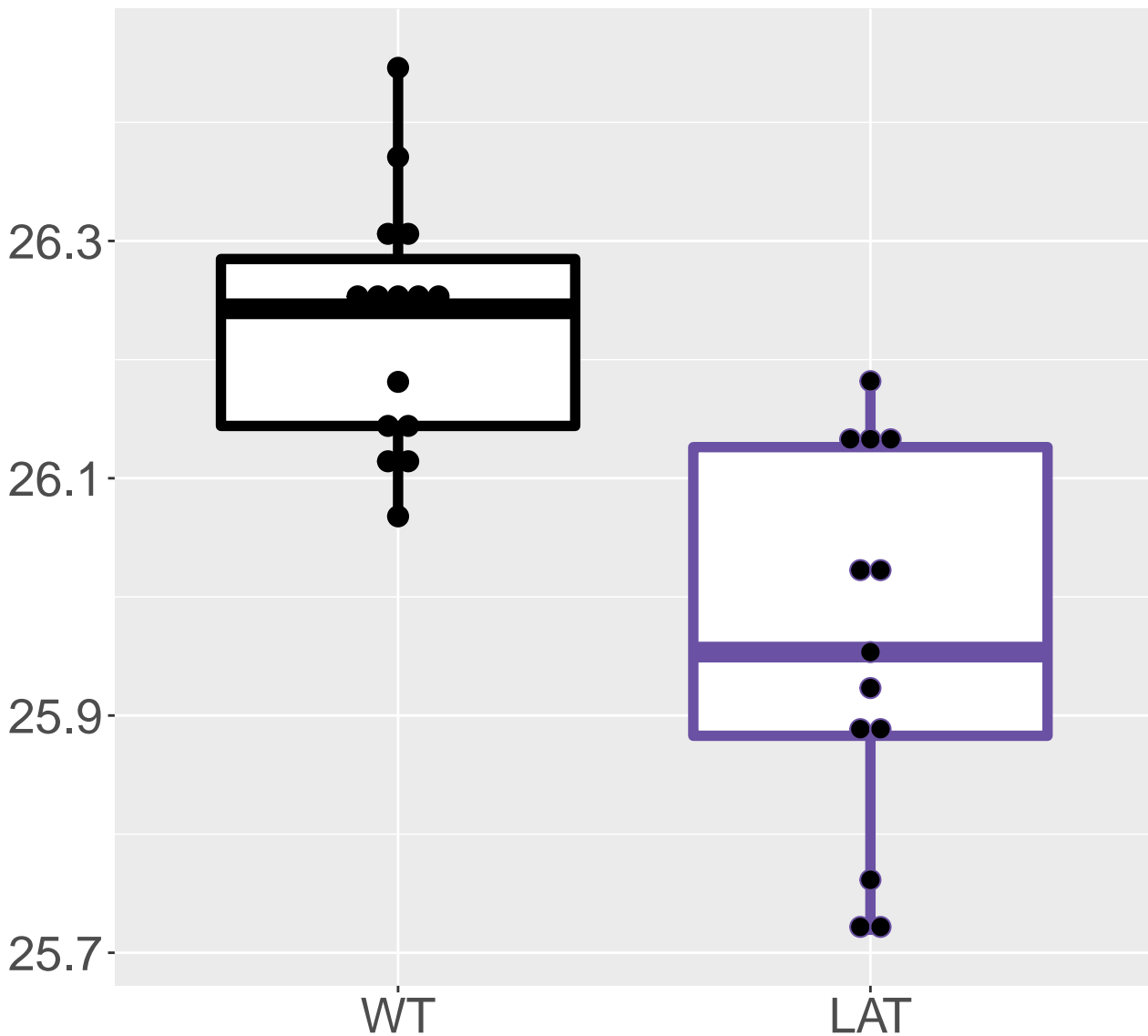


Q8R1I1_Cytochrome b-c1 complex .
FDR = 0.00044, FC = -0.6, sex***

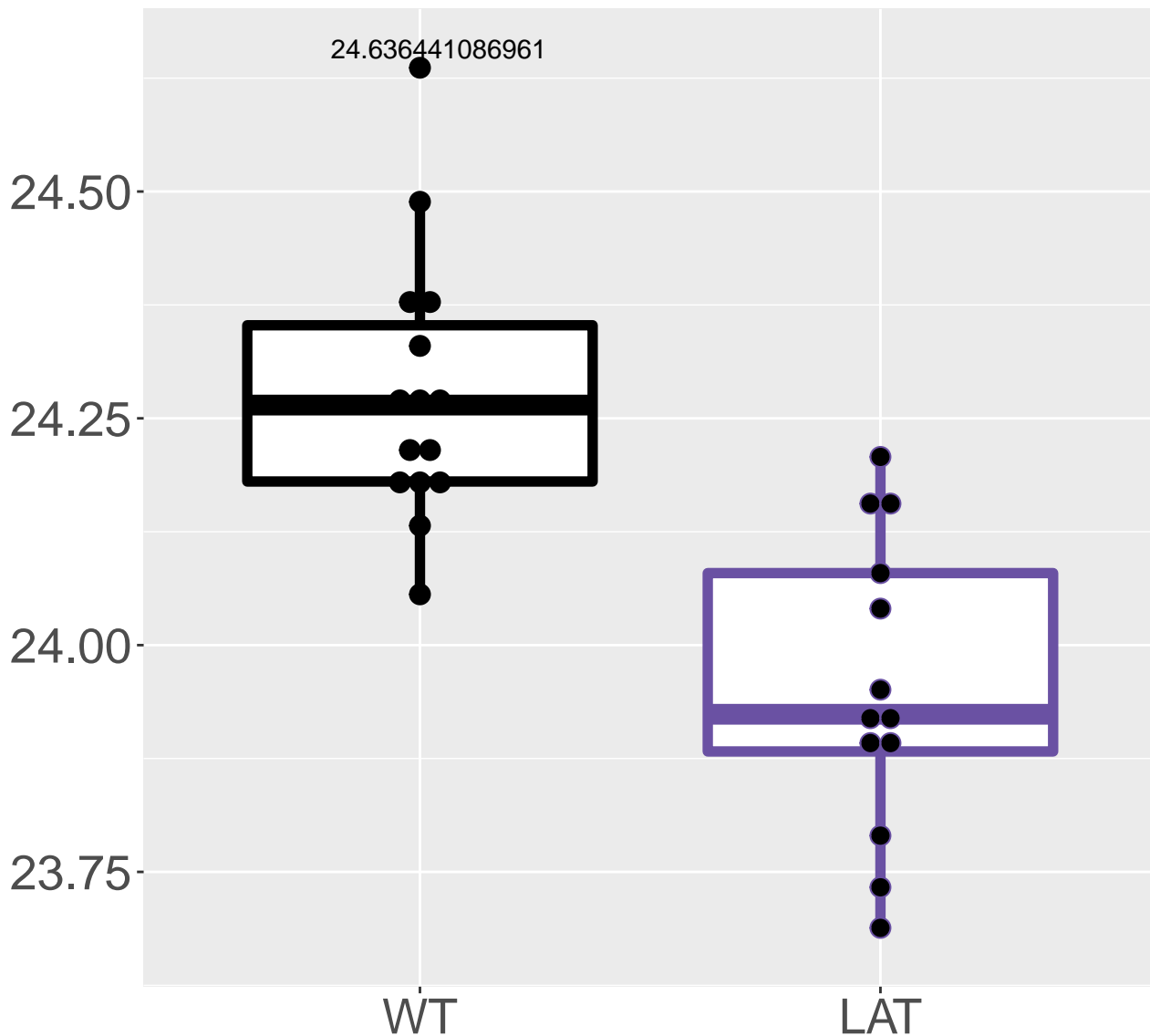


P62852_40S ribosomal protein S25

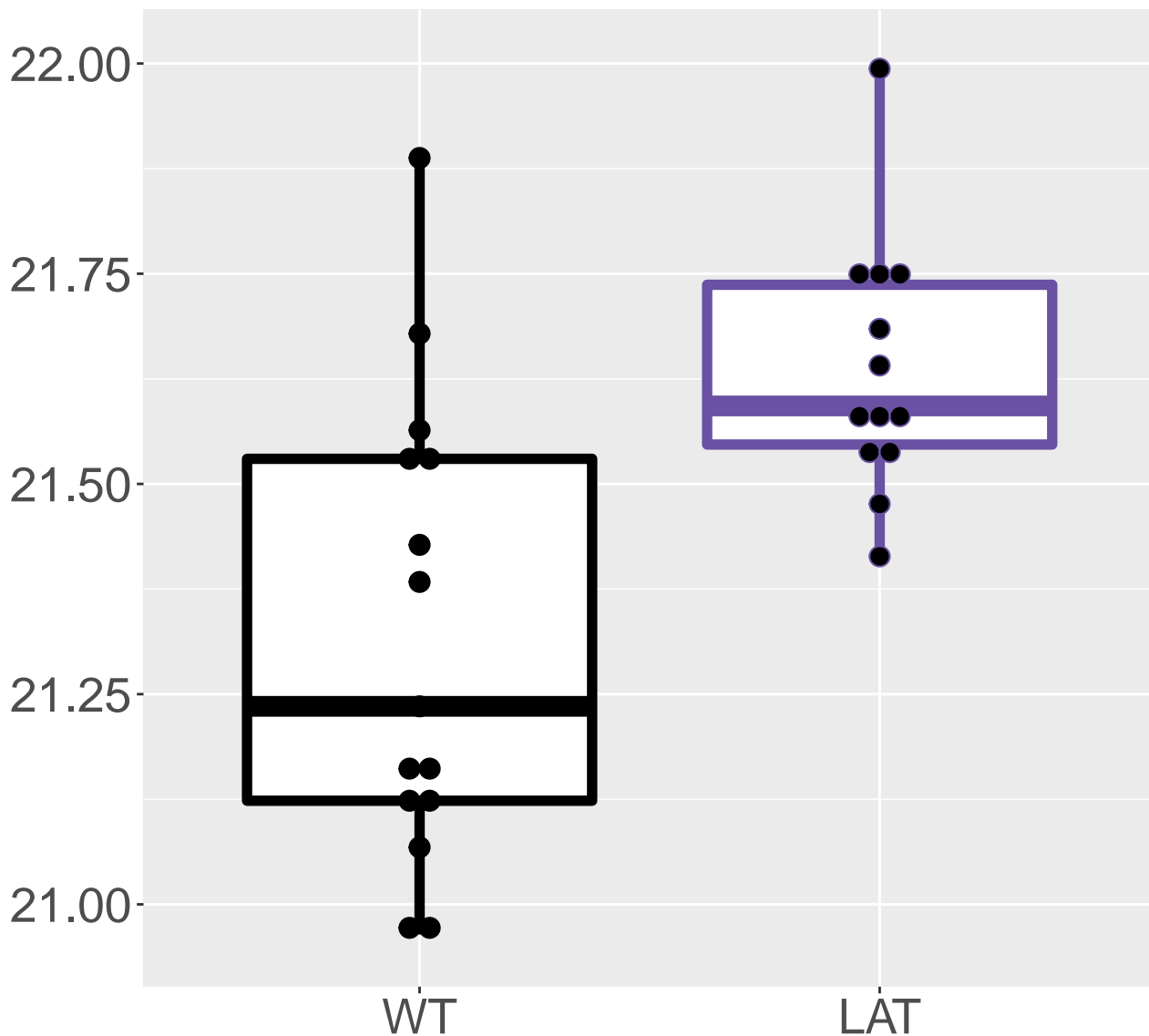
FDR = 0.00044, FC = -0.48, sex**



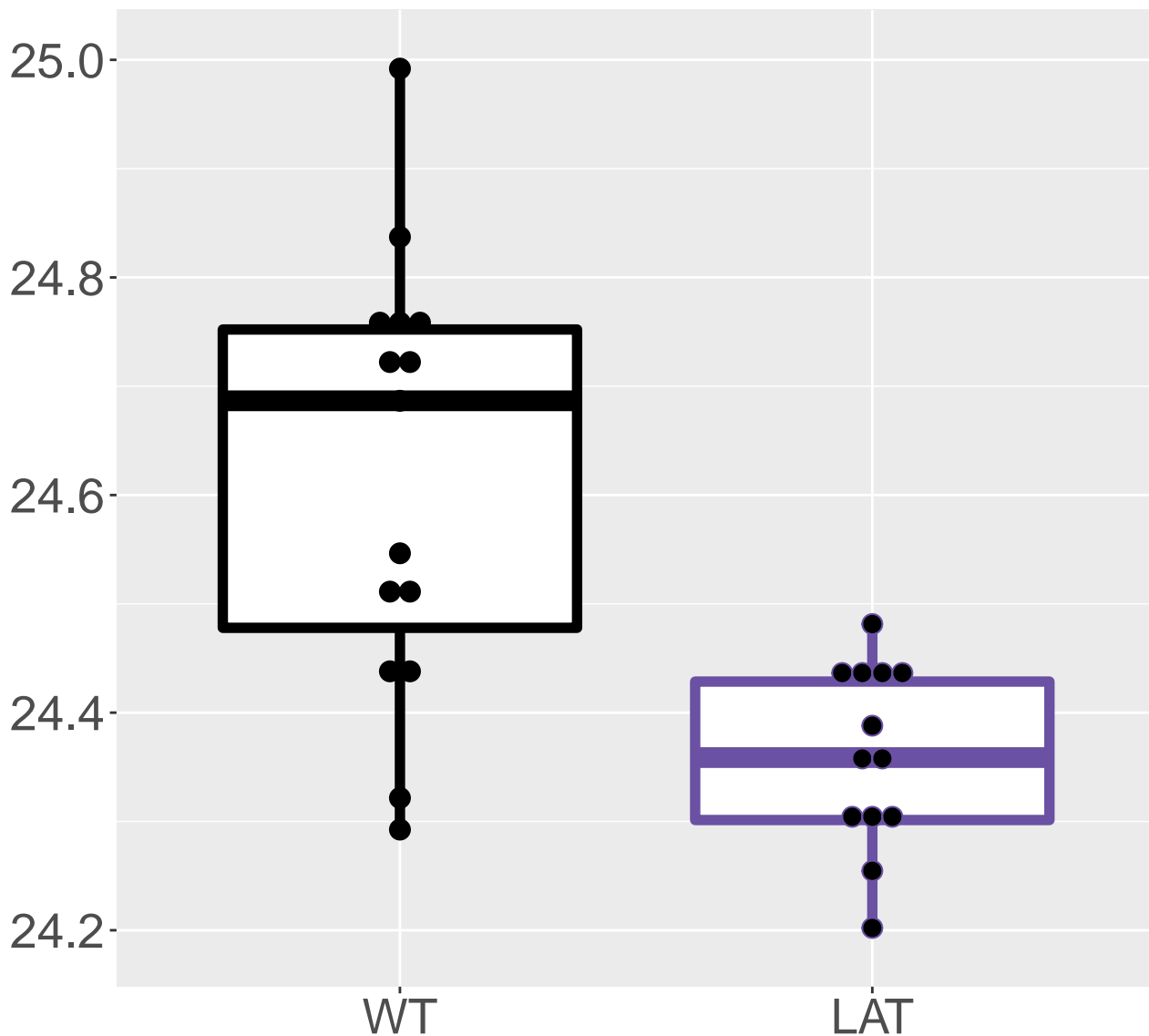
O08997_Copper transport protein.
FDR = 0.00065, FC = -0.55, sex*



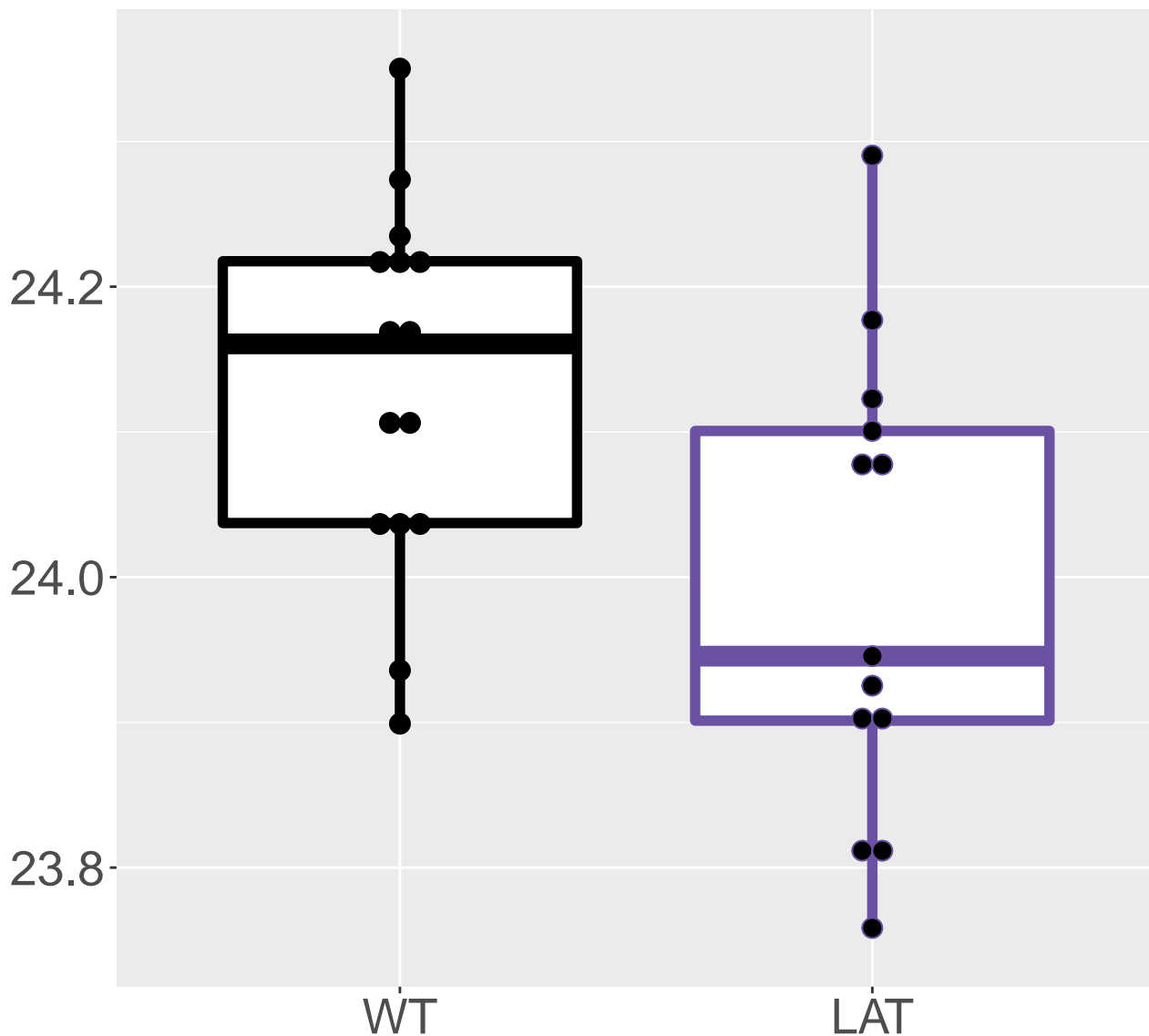
P03911_NADH-ubiquinone oxidored.
FDR = 0.00065, FC = 0.52, sex***



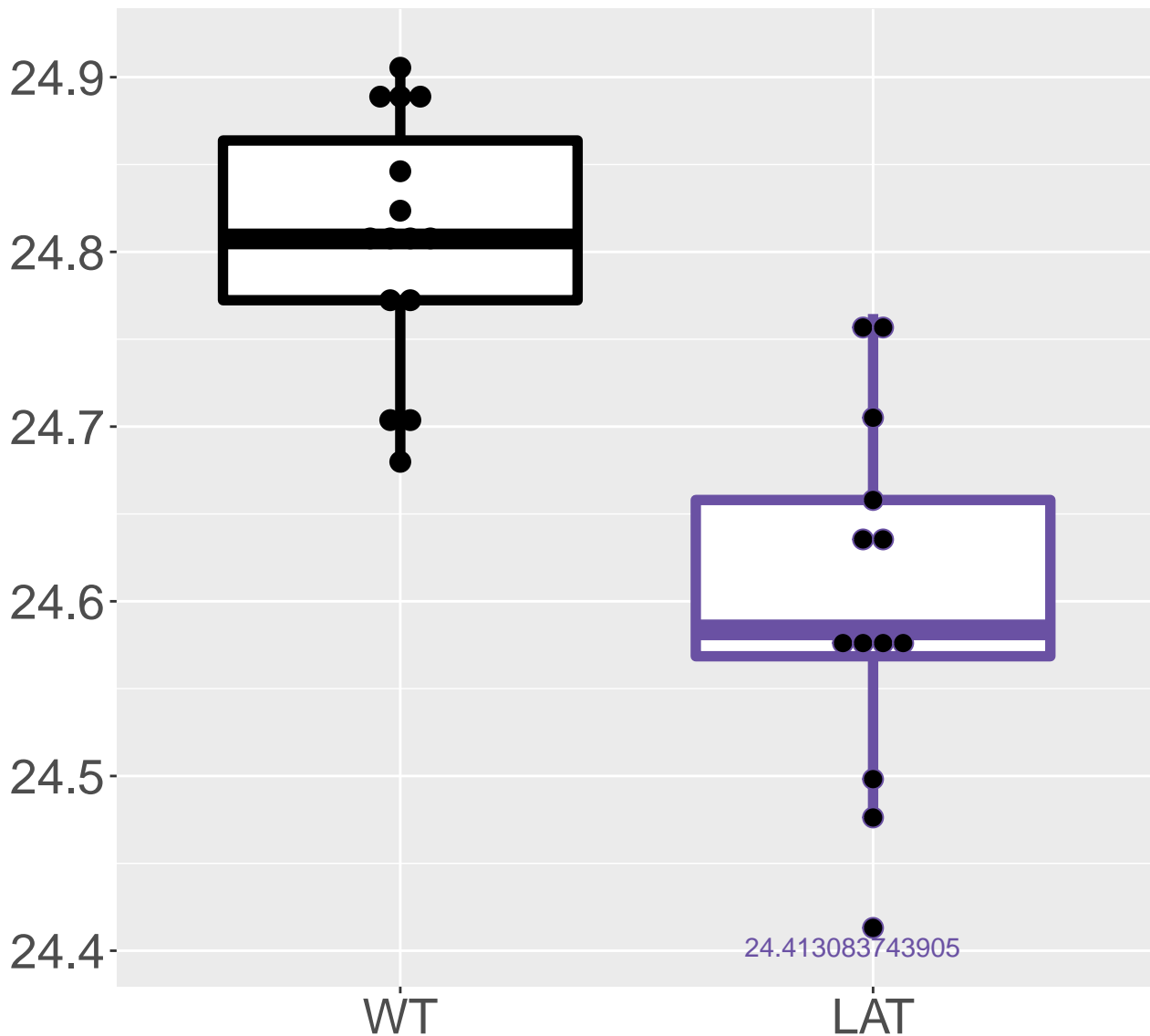
P61804_Dolichyl-diphosphooligos.
FDR = 0.00065, FC = -0.45, sex***



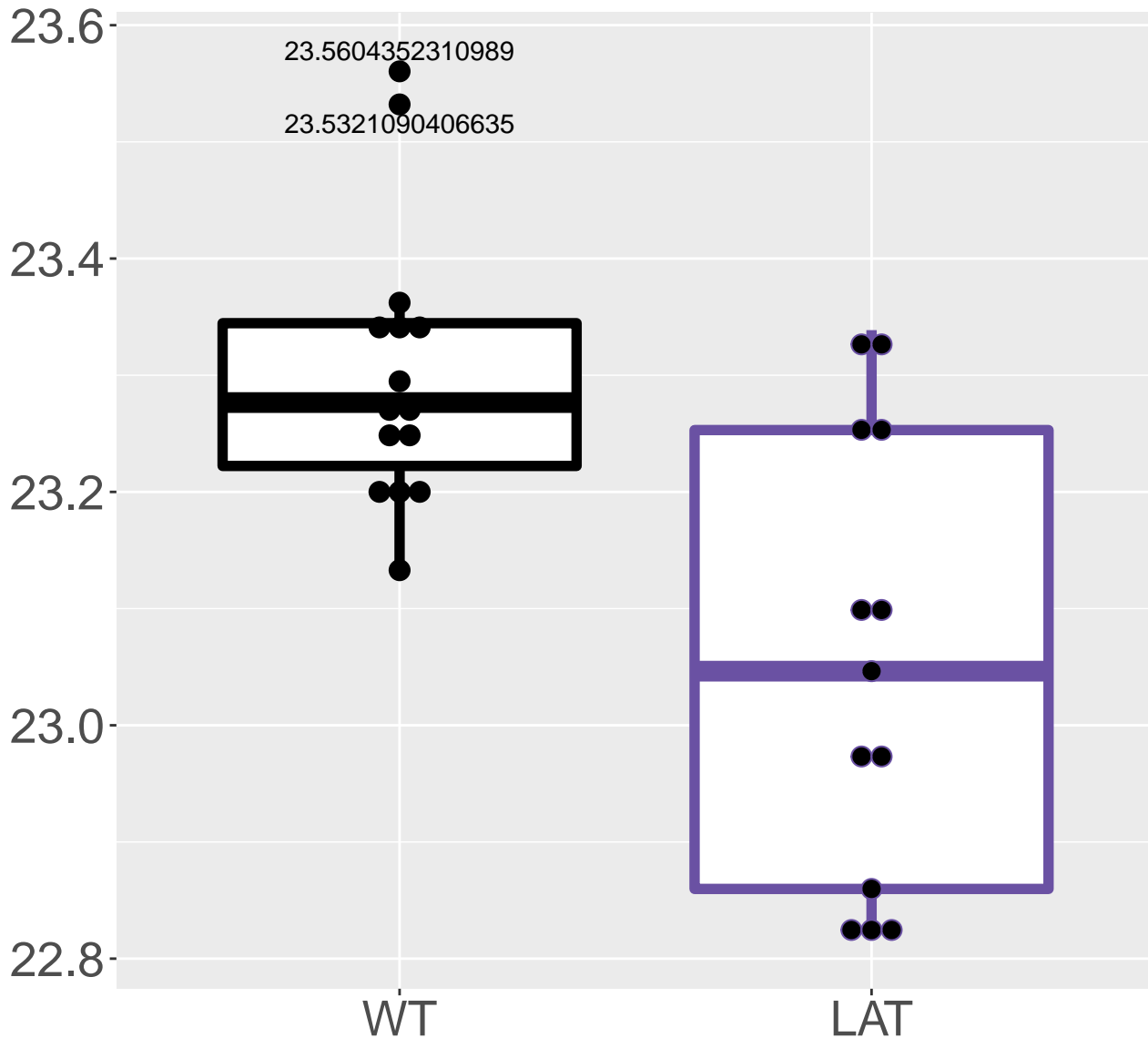
Q9CQC7_NADH dehydrogenase [ubiq.
FDR = 0.00065, FC = -0.36, sex*



Q9CQR2_40S ribosomal protein S21
FDR = 0.00065, FC = -0.32, sex*

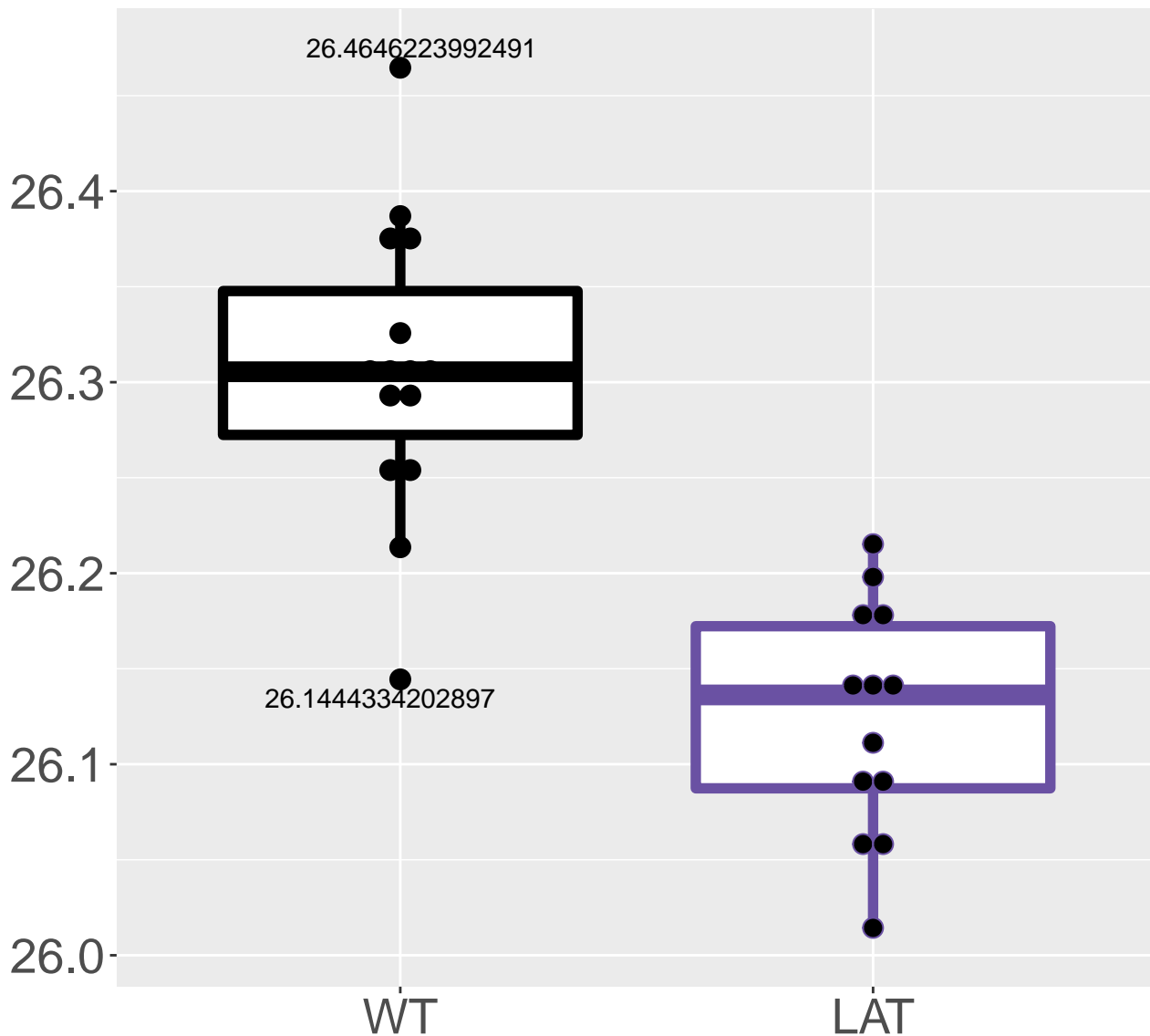


Q9CQZ6_NADH dehydrogenase [ubiq.
FDR = 0.0018, FC = -0.37, sex*

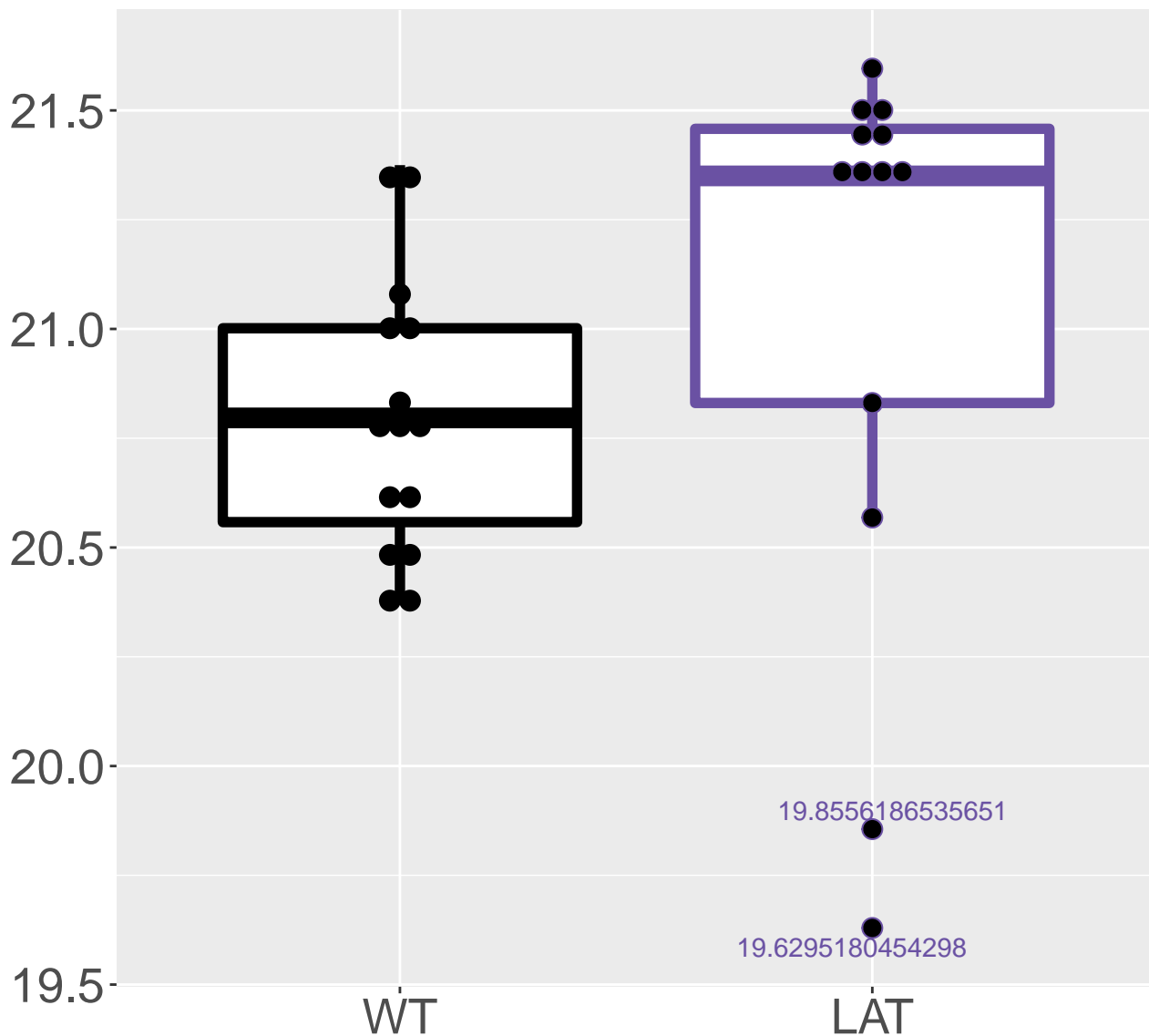


P62830_60S ribosomal protein L23

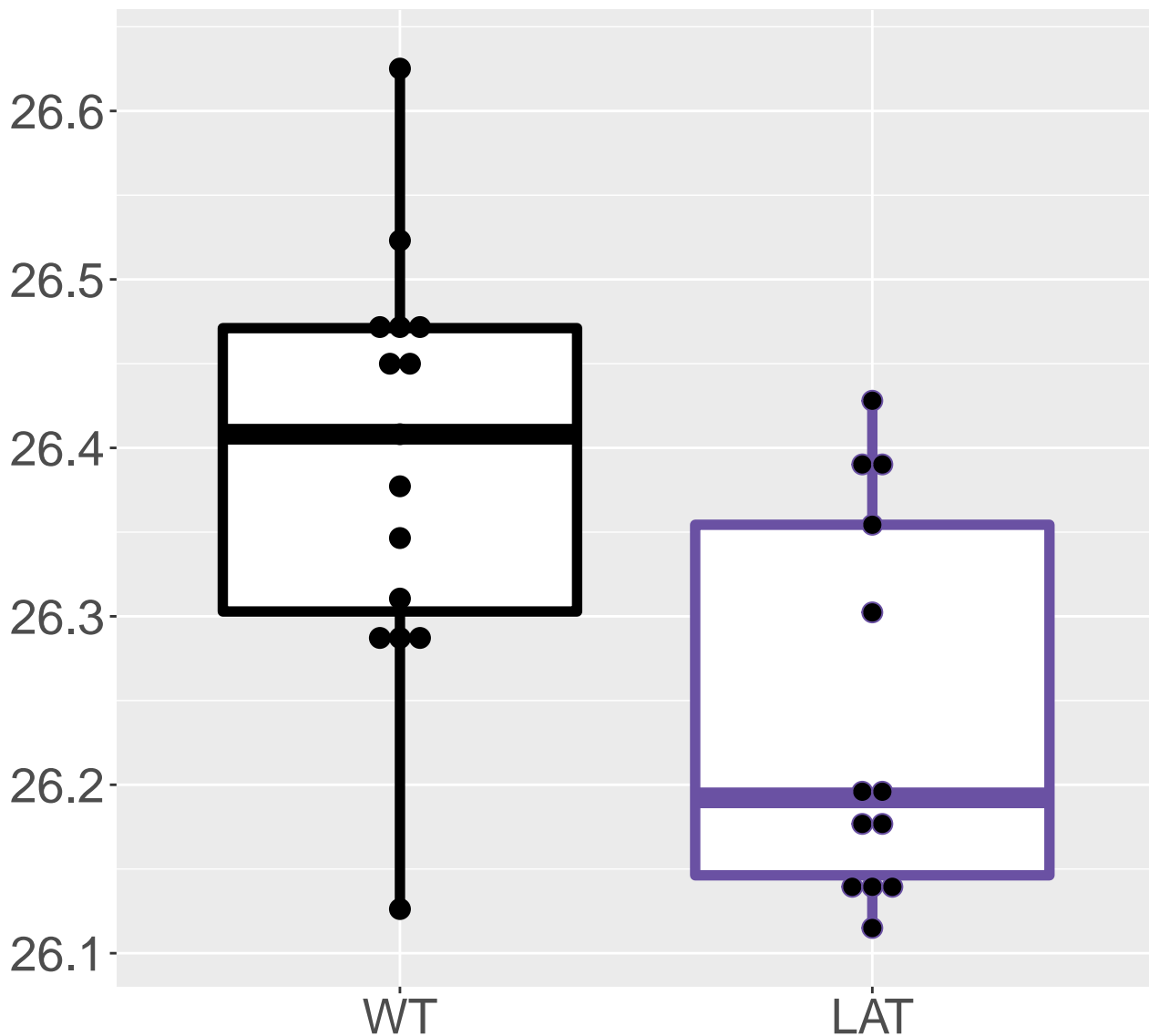
FDR = 0.0018, FC = -0.25



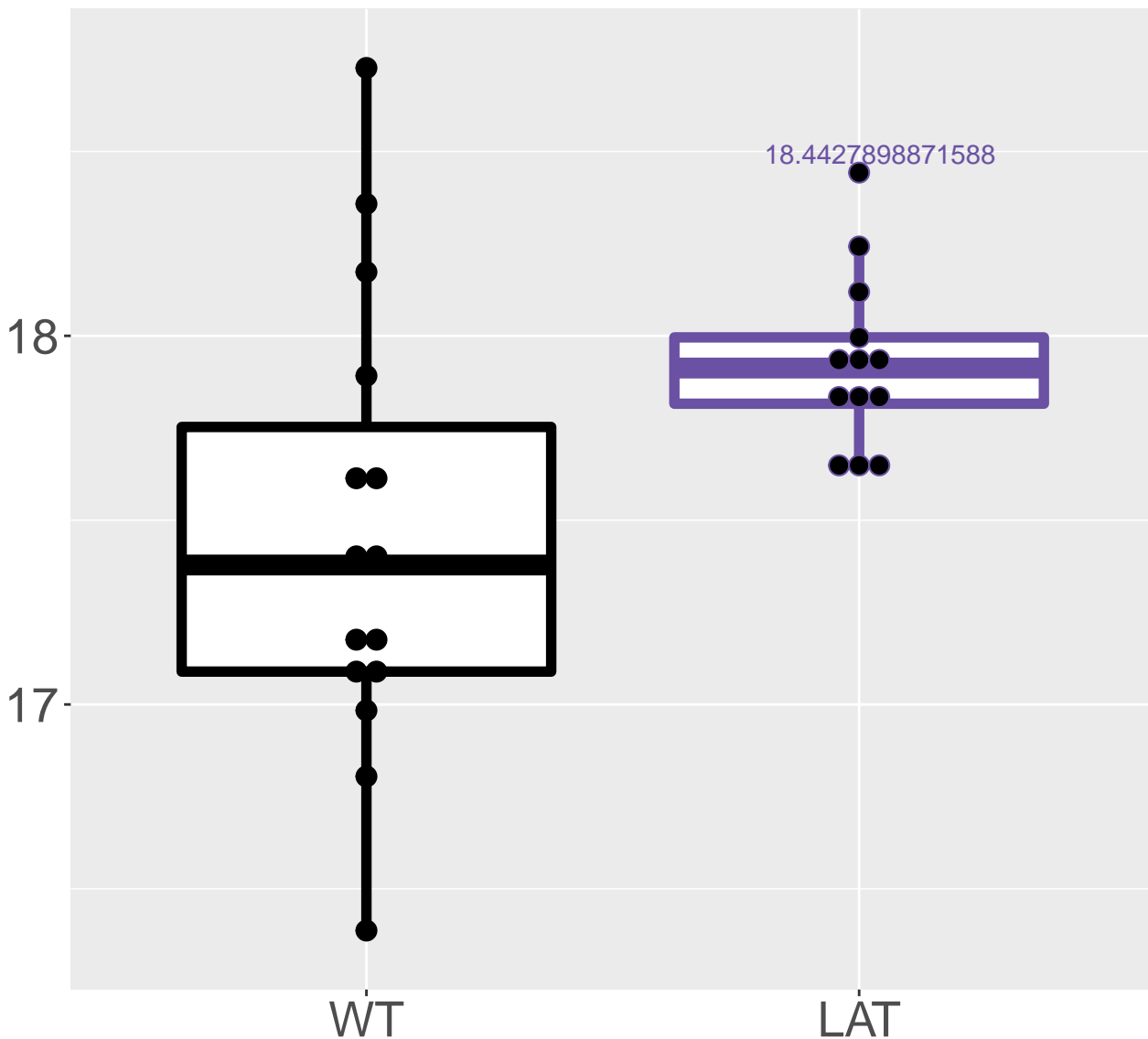
O70133_ATP-dependent RNA helica.
FDR = 0.0024, FC = 0.76, sex**



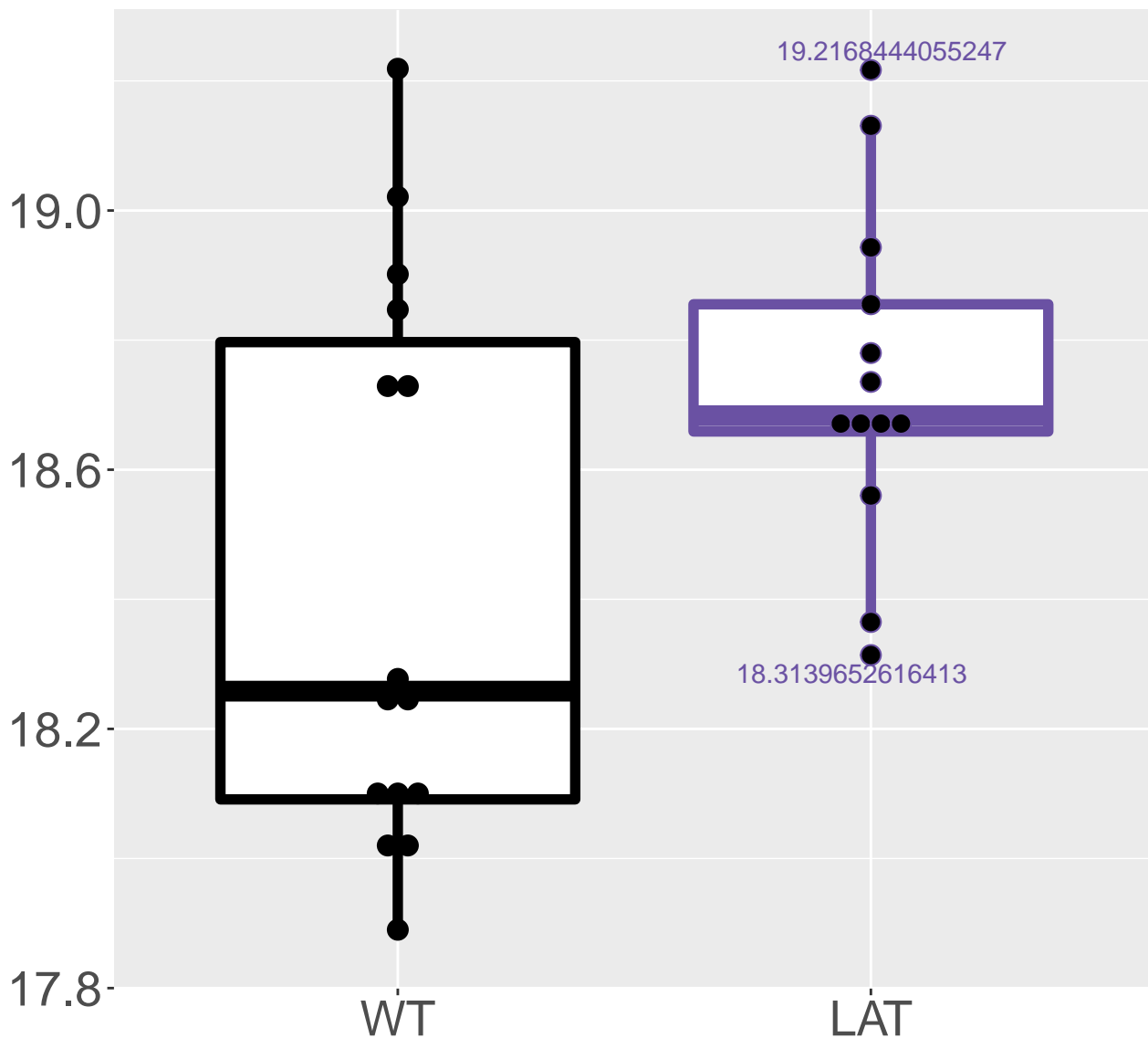
Q06185_ATP synthase subunit e, .
FDR = 0.0025, FC = -0.31, sex*



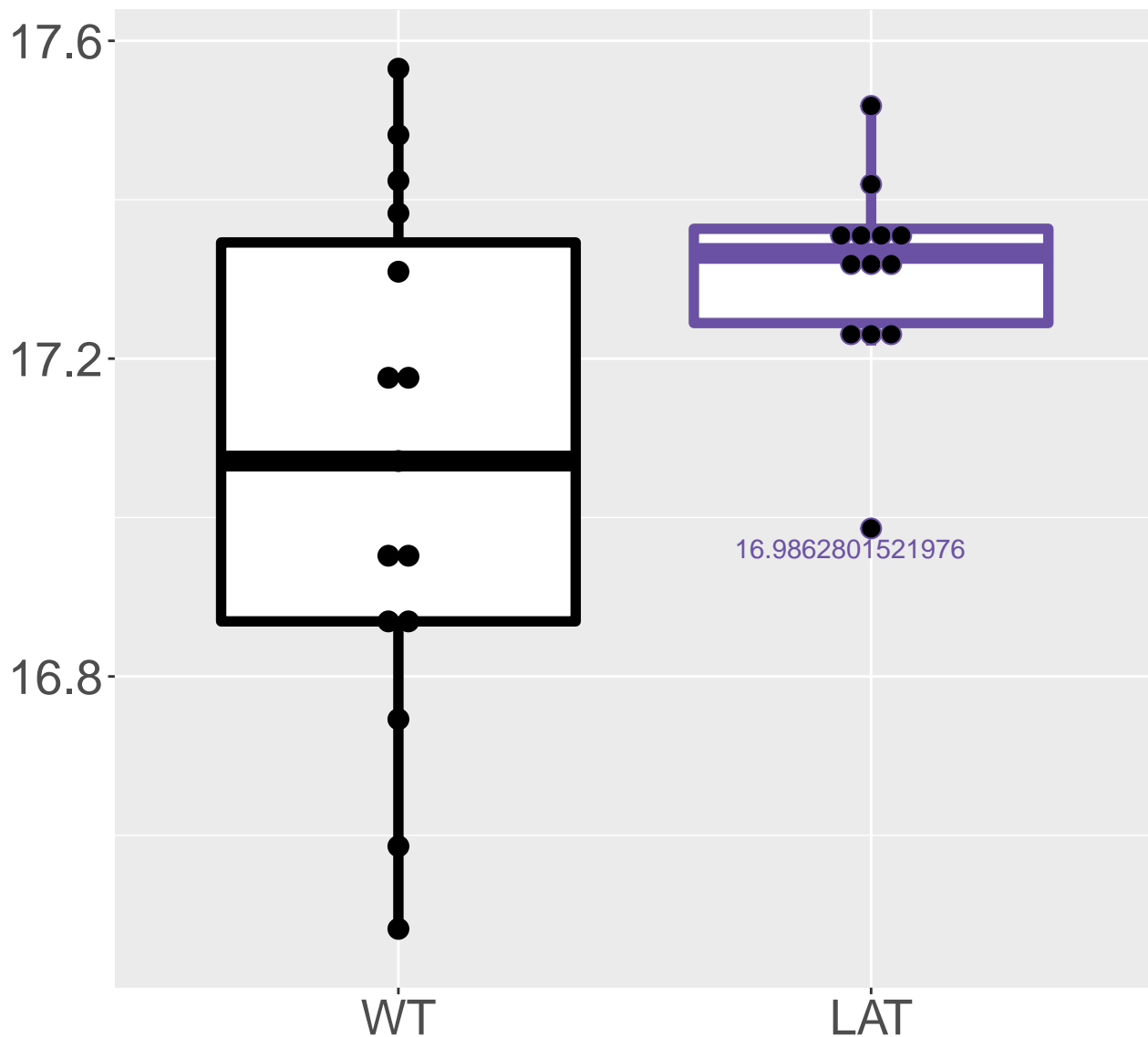
Q99JH8_ER lumen protein-retaini.
FDR = 0.0027, FC = 0.98, sex**



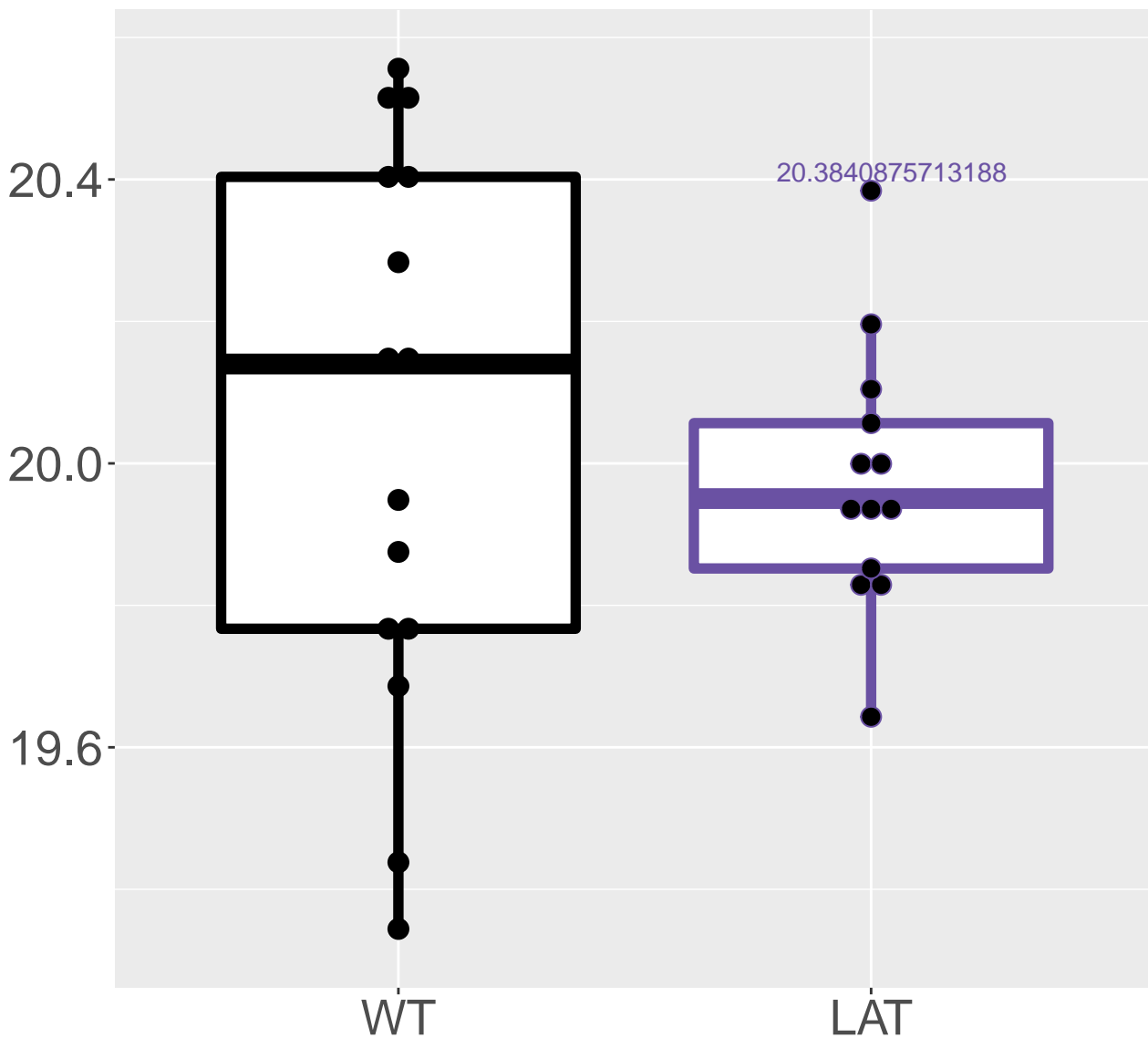
Q9CQM2_ER lumen protein-retaini.
FDR = 0.0027, FC = 0.69, sex***



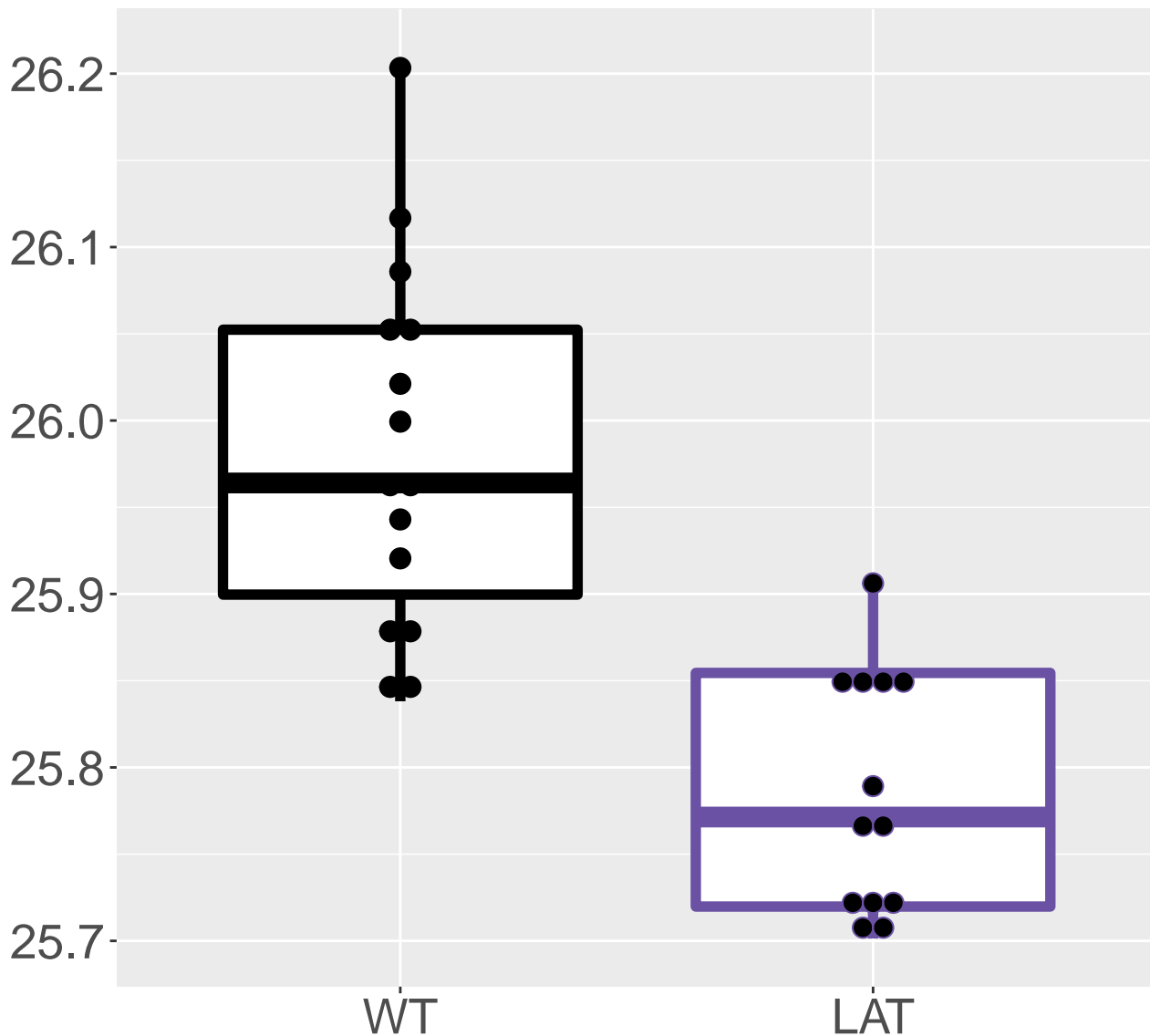
Q8BZS9_Putative pre-mRNA-splici.
FDR = 0.0027, FC = 0.55, sex***



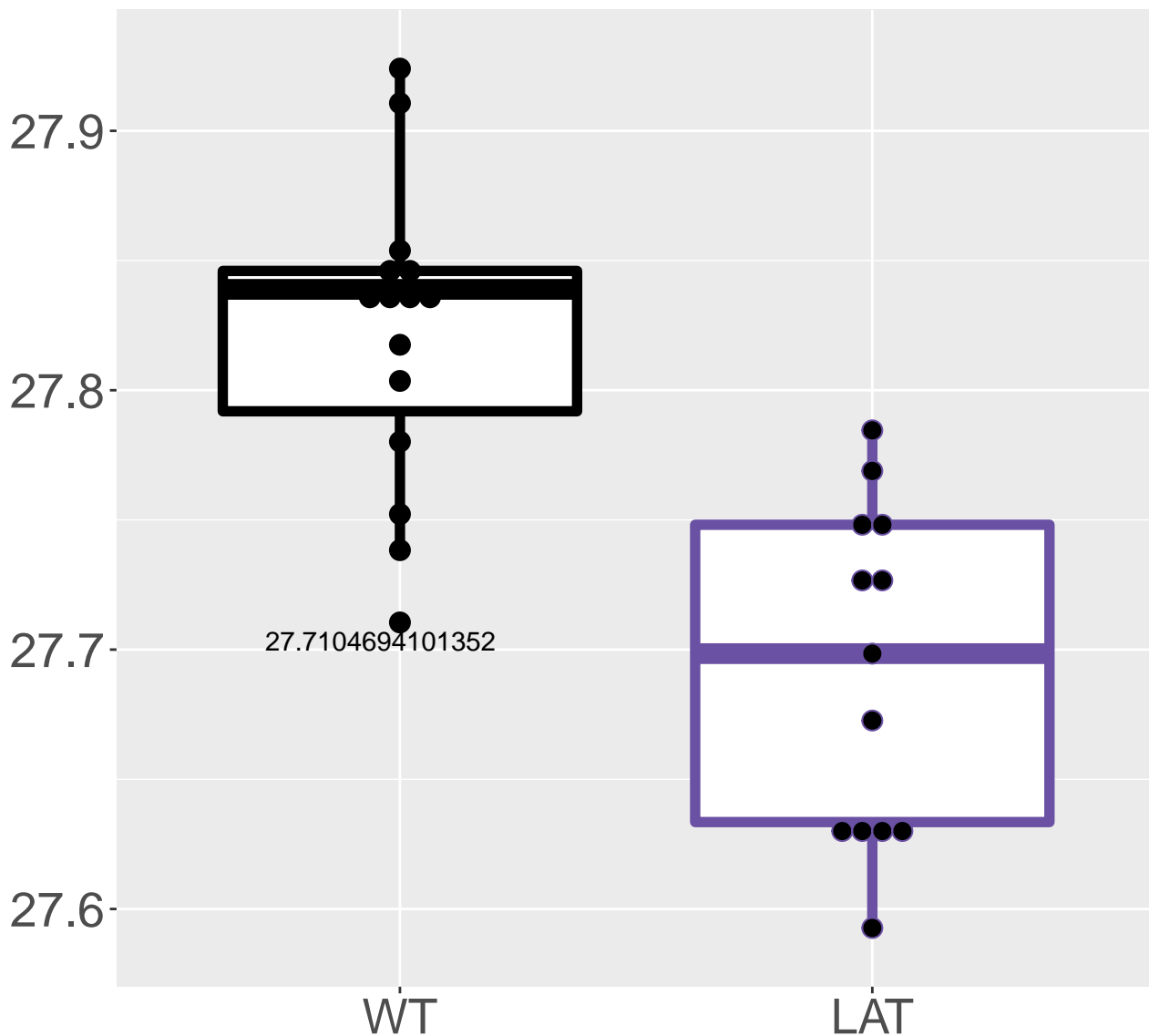
Q9CZY3_Ubiquitin-conjugating en.
FDR = 0.0027, FC = -0.46, sex***



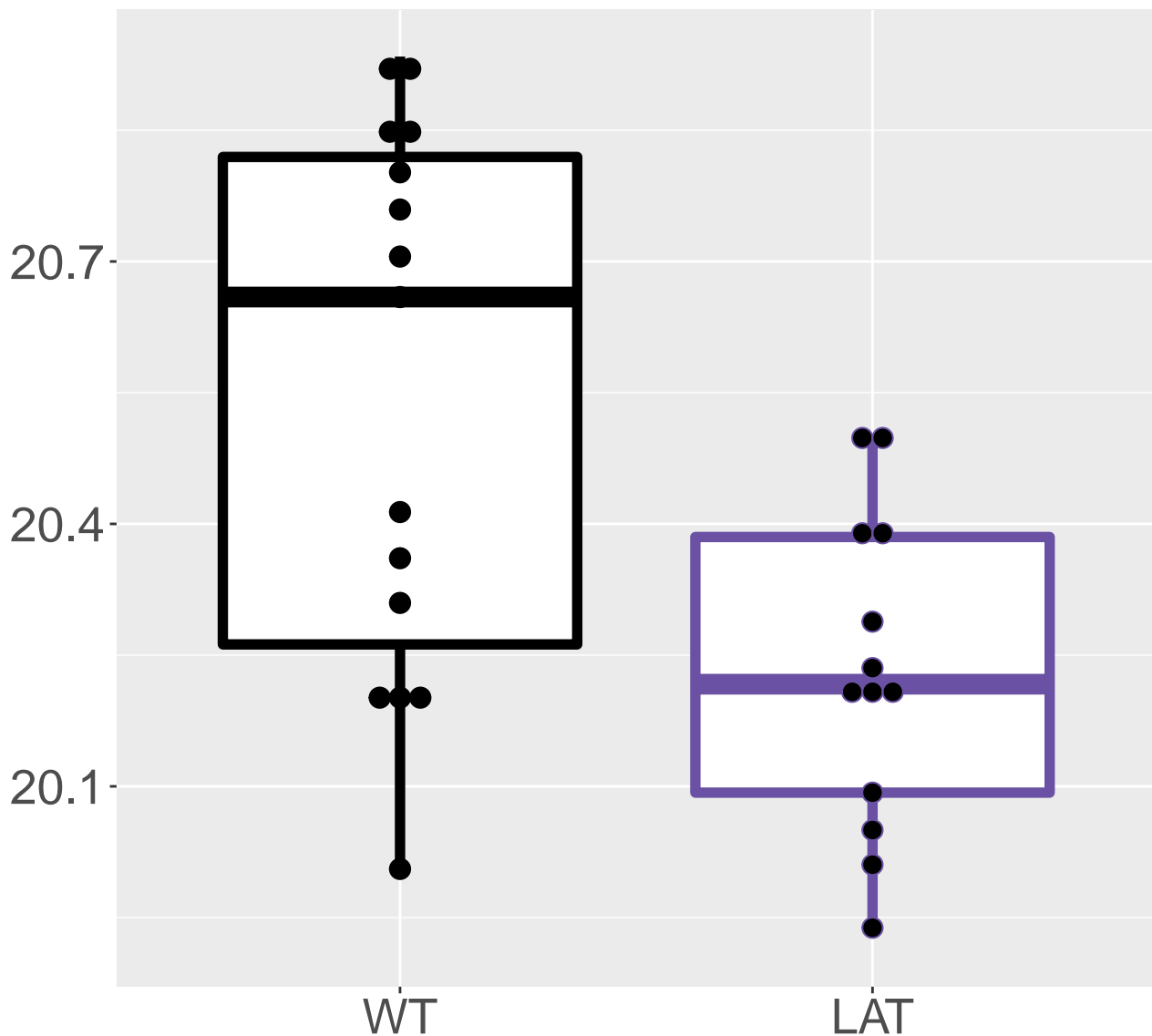
P62889_60S ribosomal protein L30
FDR = 0.0027, FC = -0.27



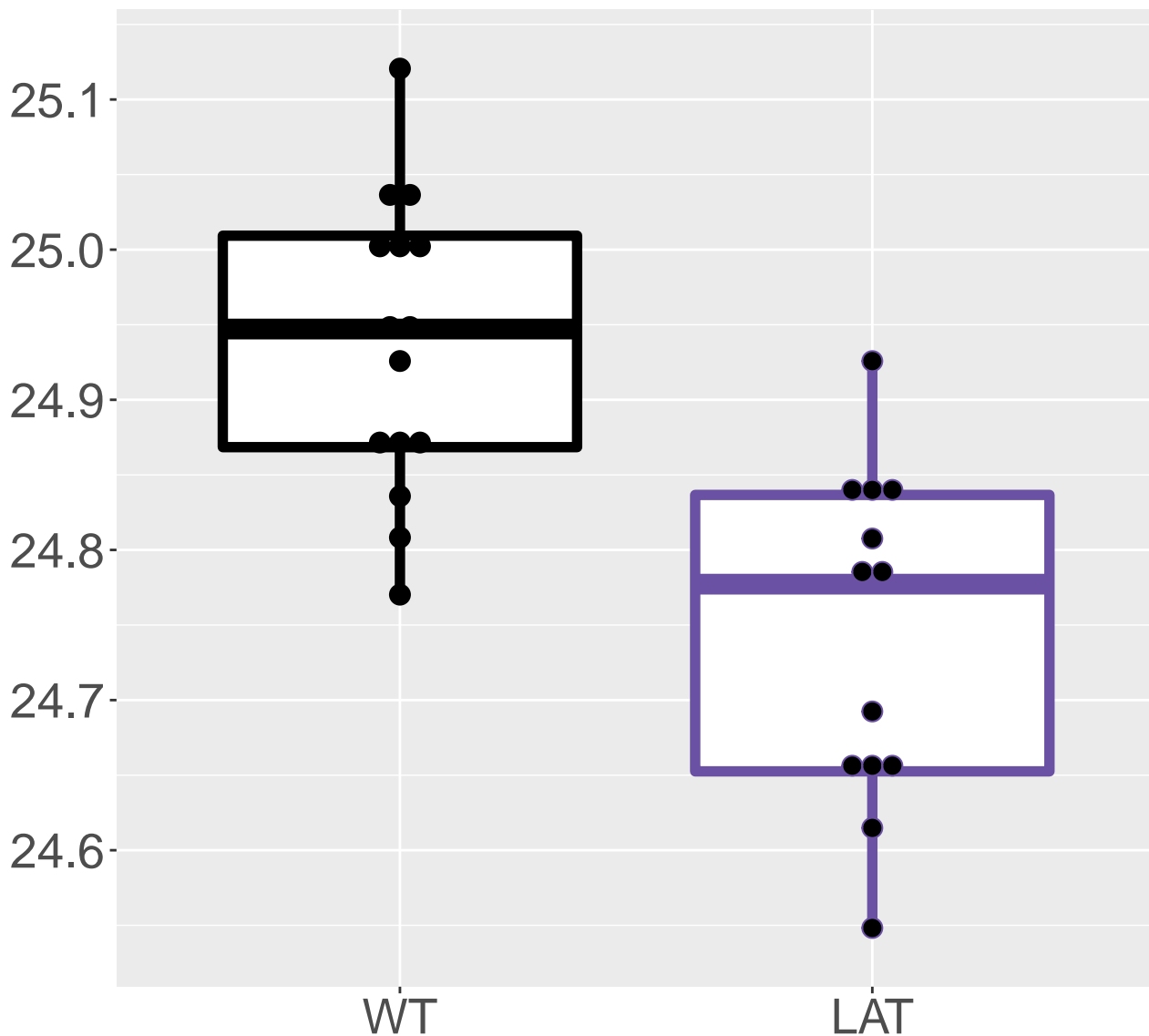
Q6ZWN5_40S ribosomal protein S9
FDR = 0.0027, FC = -0.21, sex*



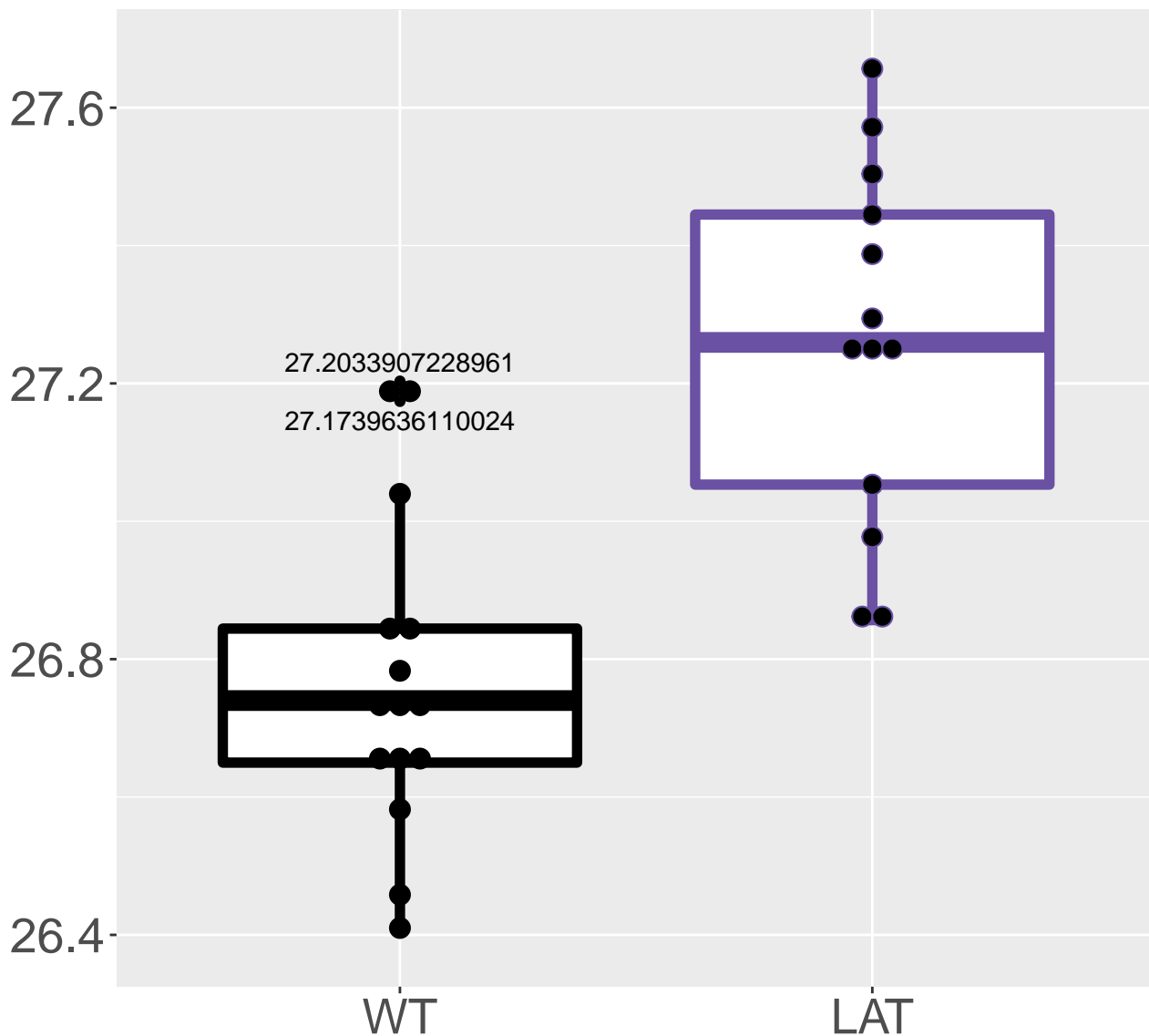
Q6P8J2_Diamine acetyltransferas.
FDR = 0.0027, FC = -0.49, sex***



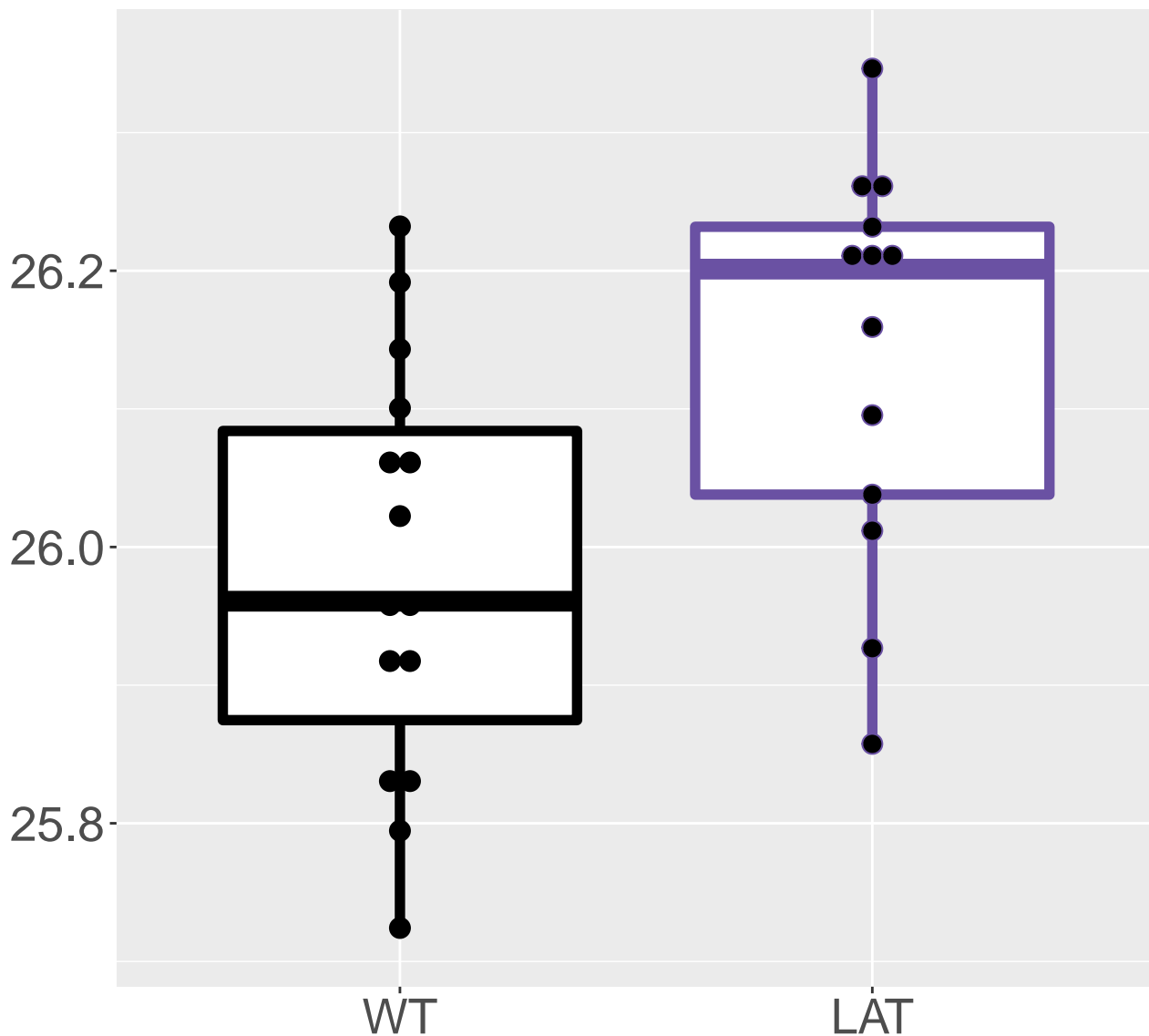
O55142_60S ribosomal protein L3.
FDR = 0.0031, FC = -0.33, sex**



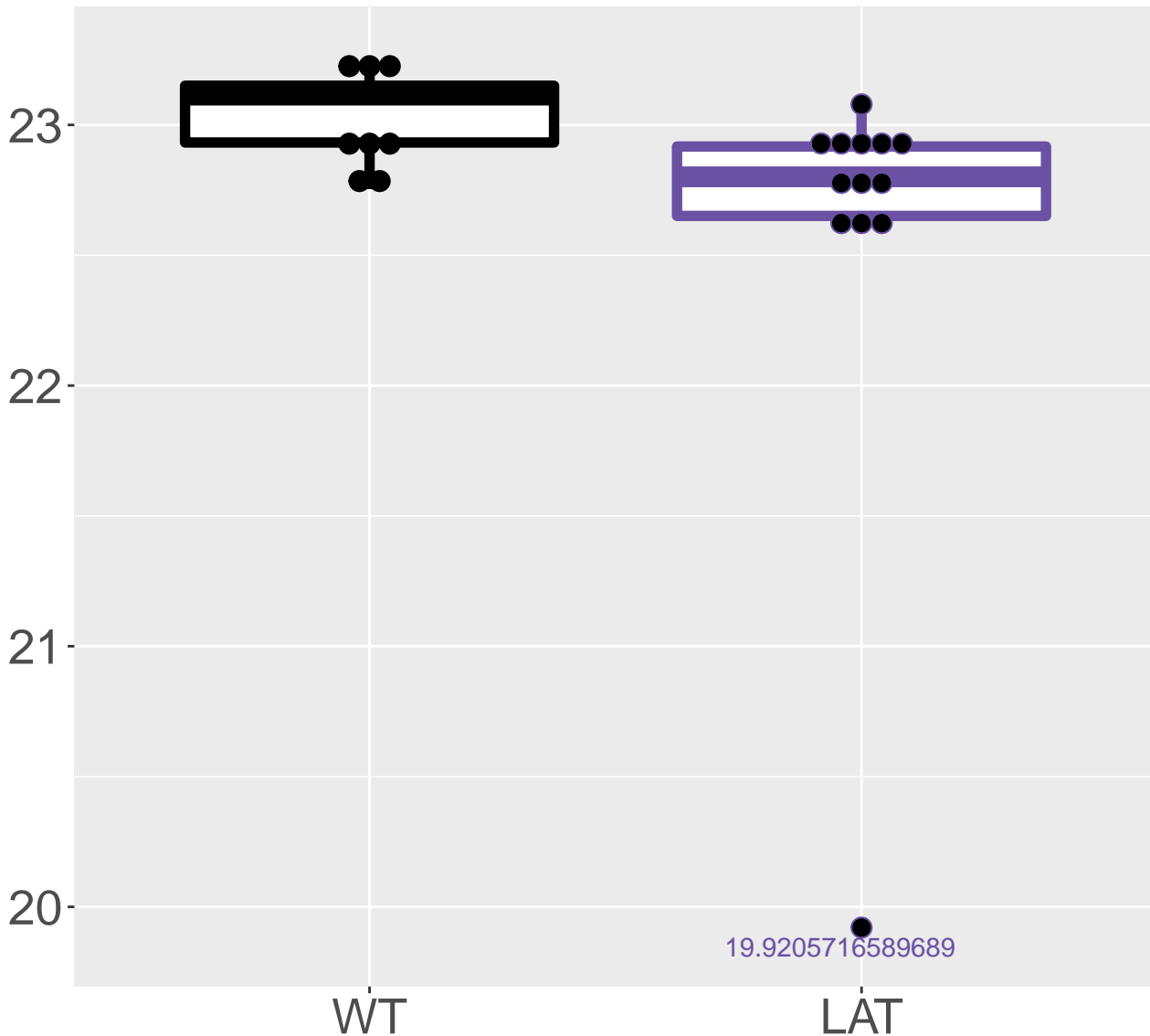
Q99K67_Alpha-aminoadipic semial.
FDR = 0.0032, FC = 0.5, sex*



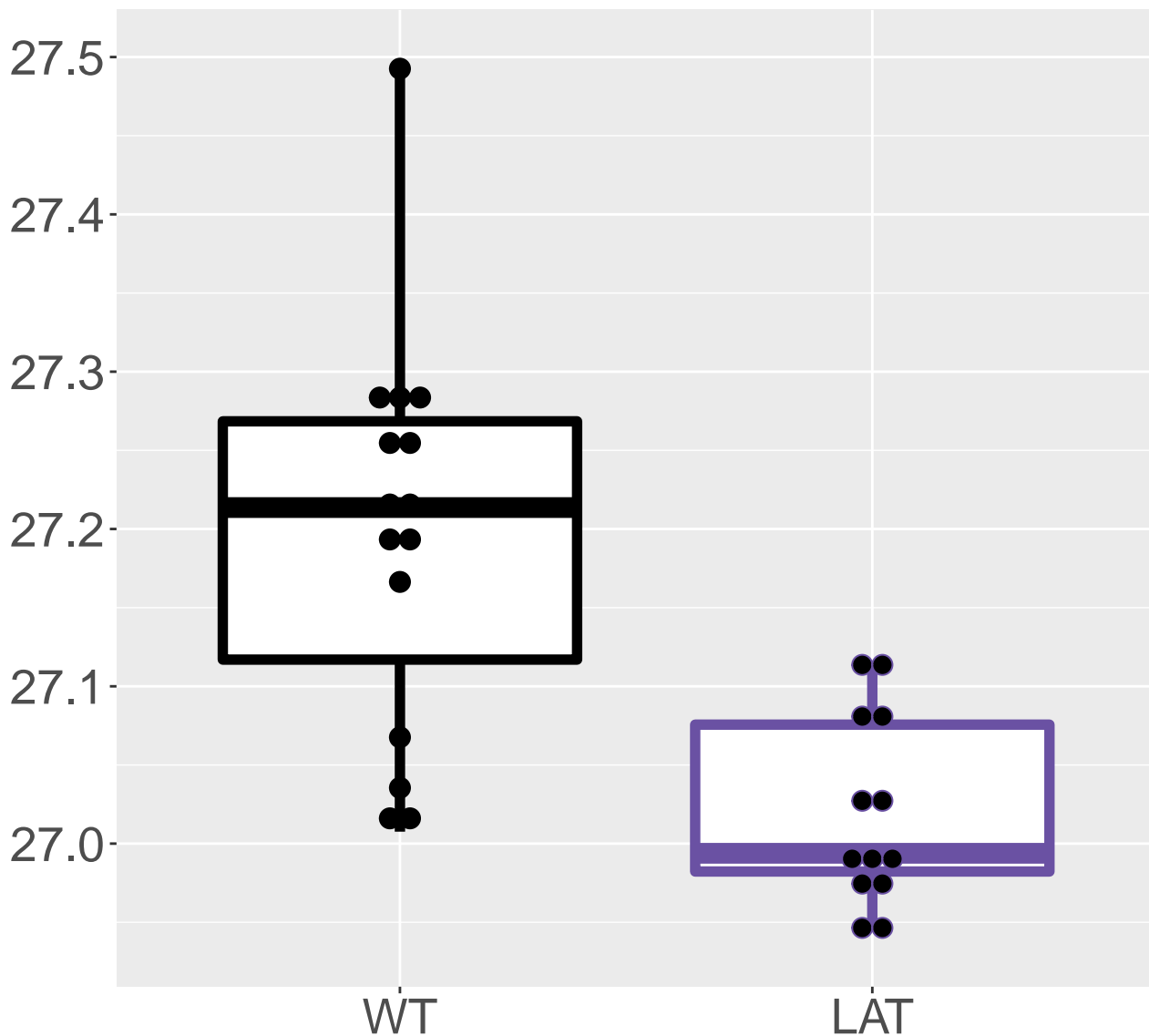
P97494_Glutamate--cysteine liga.
FDR = 0.0033, FC = 0.34, sex**



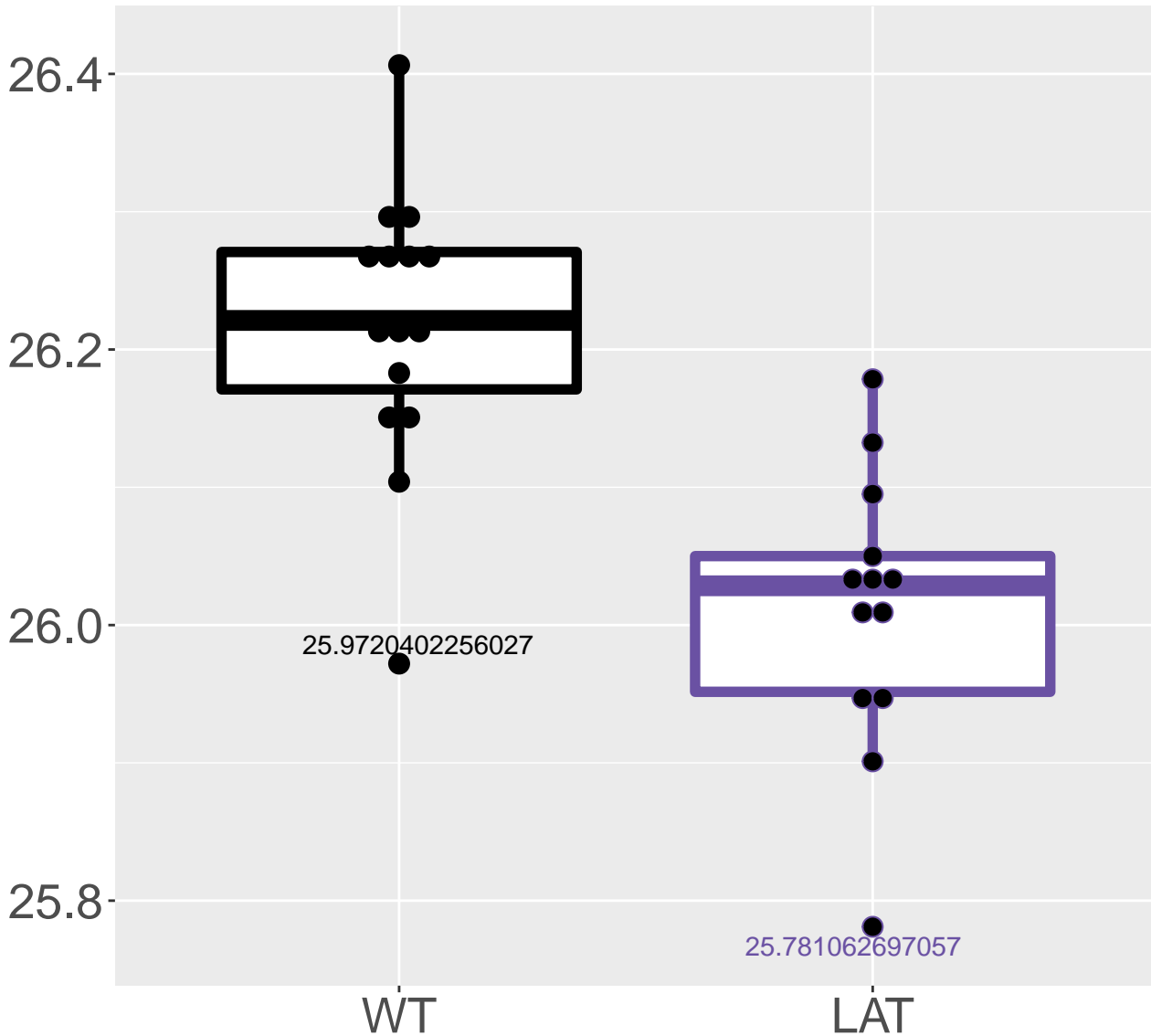
Q9Z1J3_Cysteine desulfurase, mi.
FDR = 0.0033, FC = -0.34, sex*



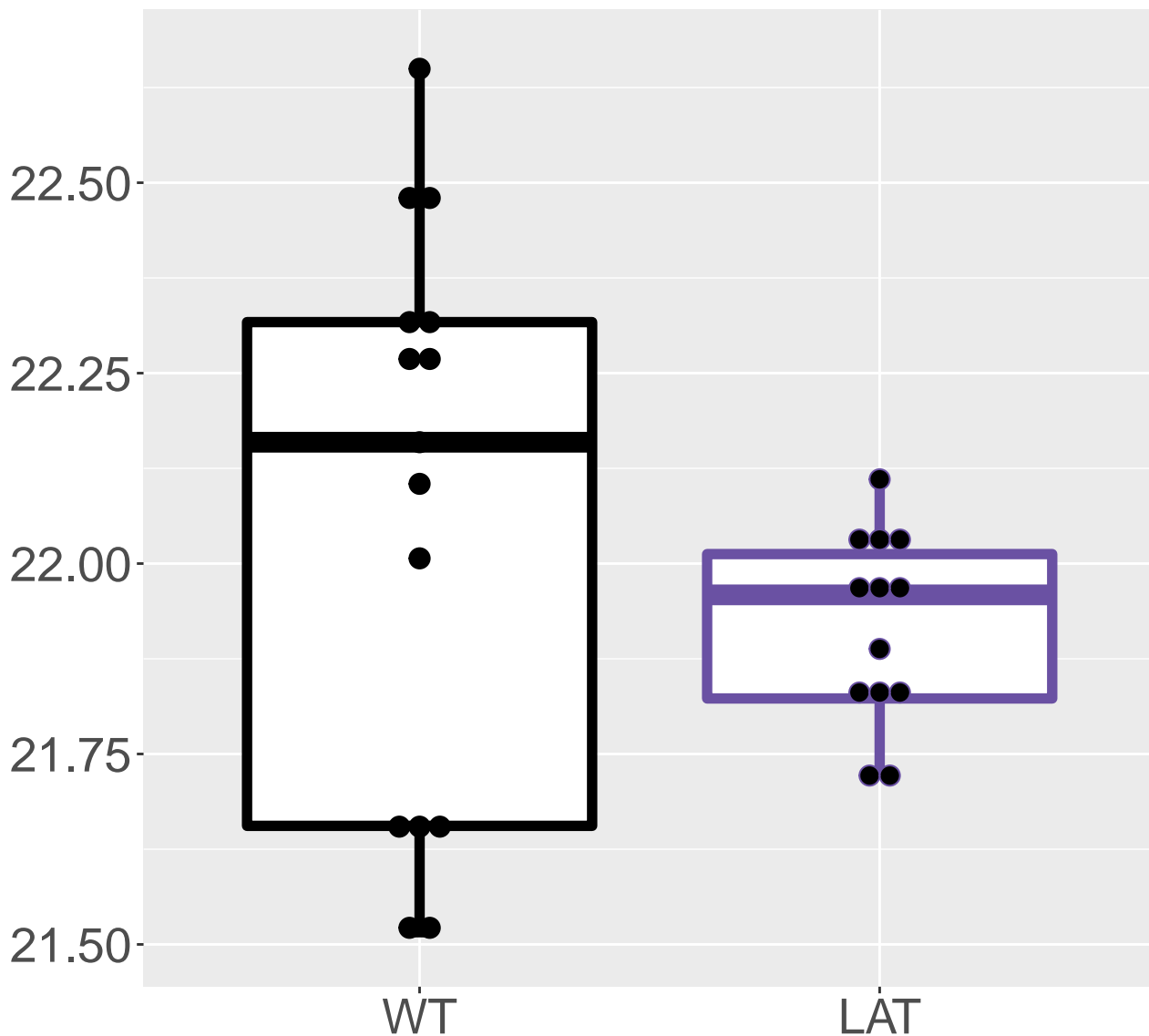
P35980_60S ribosomal protein L18
FDR = 0.0033, FC = -0.29, sex*



P14115_60S ribosomal protein L2.
FDR = 0.0034, FC = -0.29

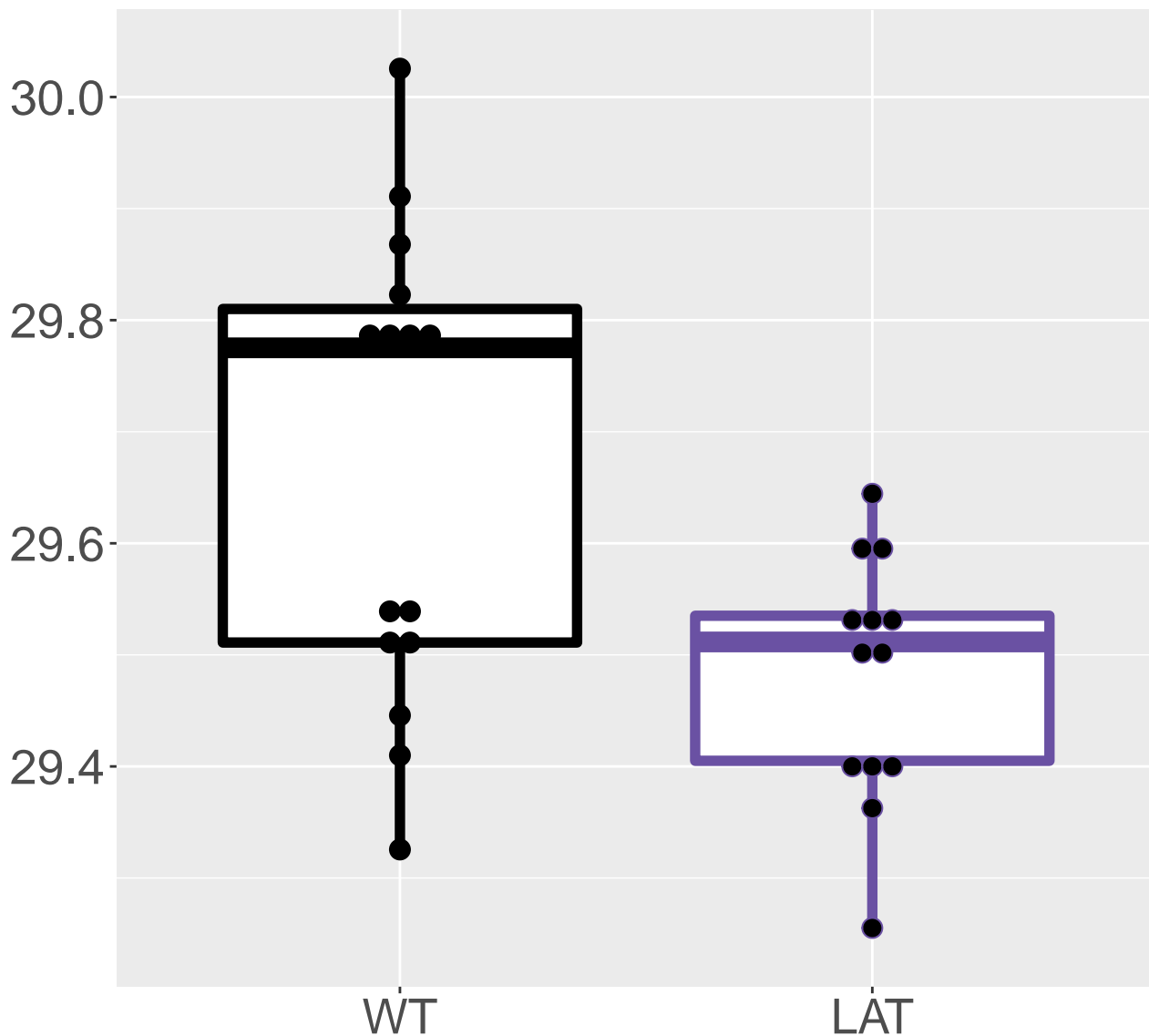


Q9D6K5_Synaptojanin-2-binding p.
FDR = 0.0035, FC = -0.47, sex***

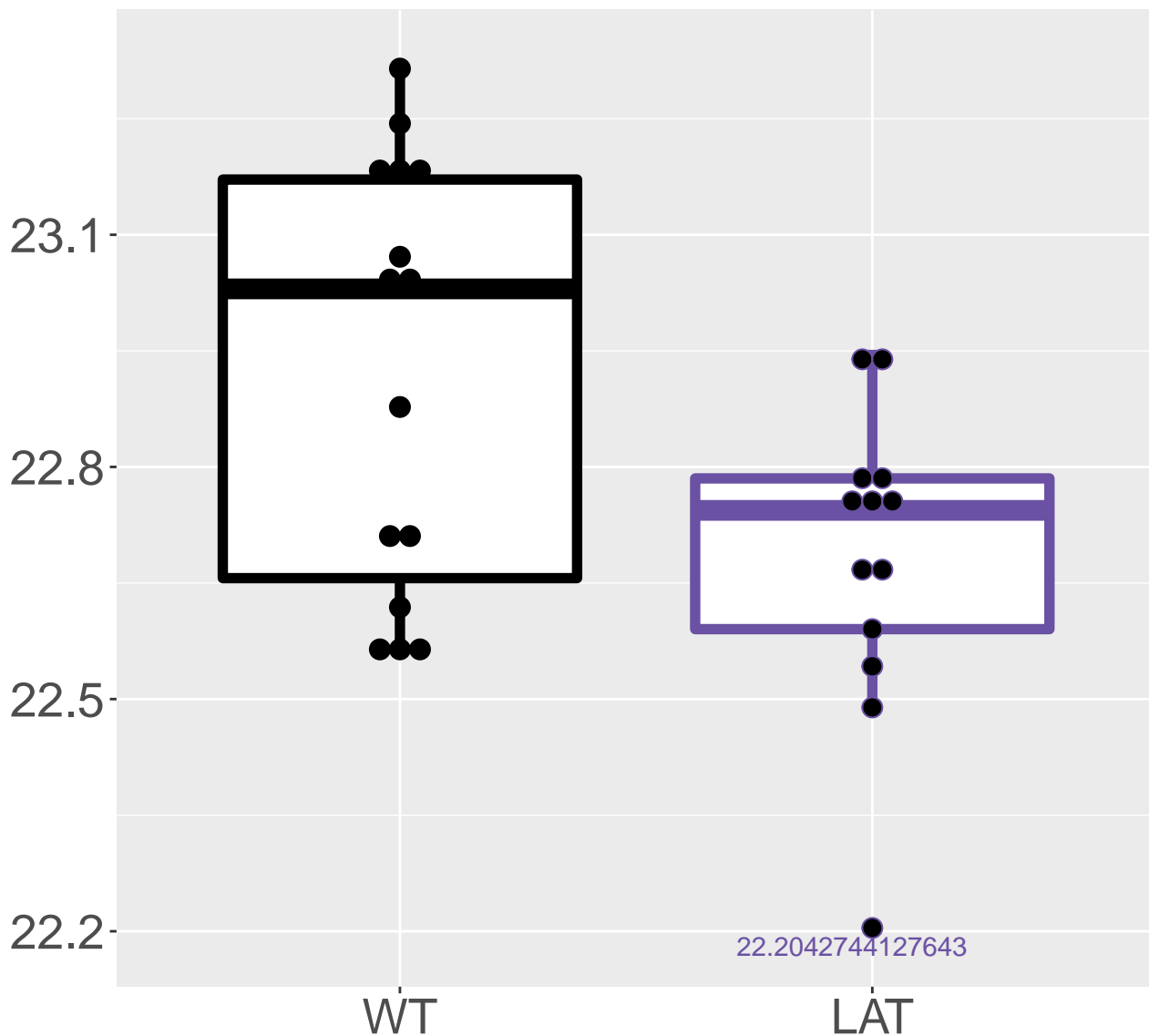


P62806_Histone H4

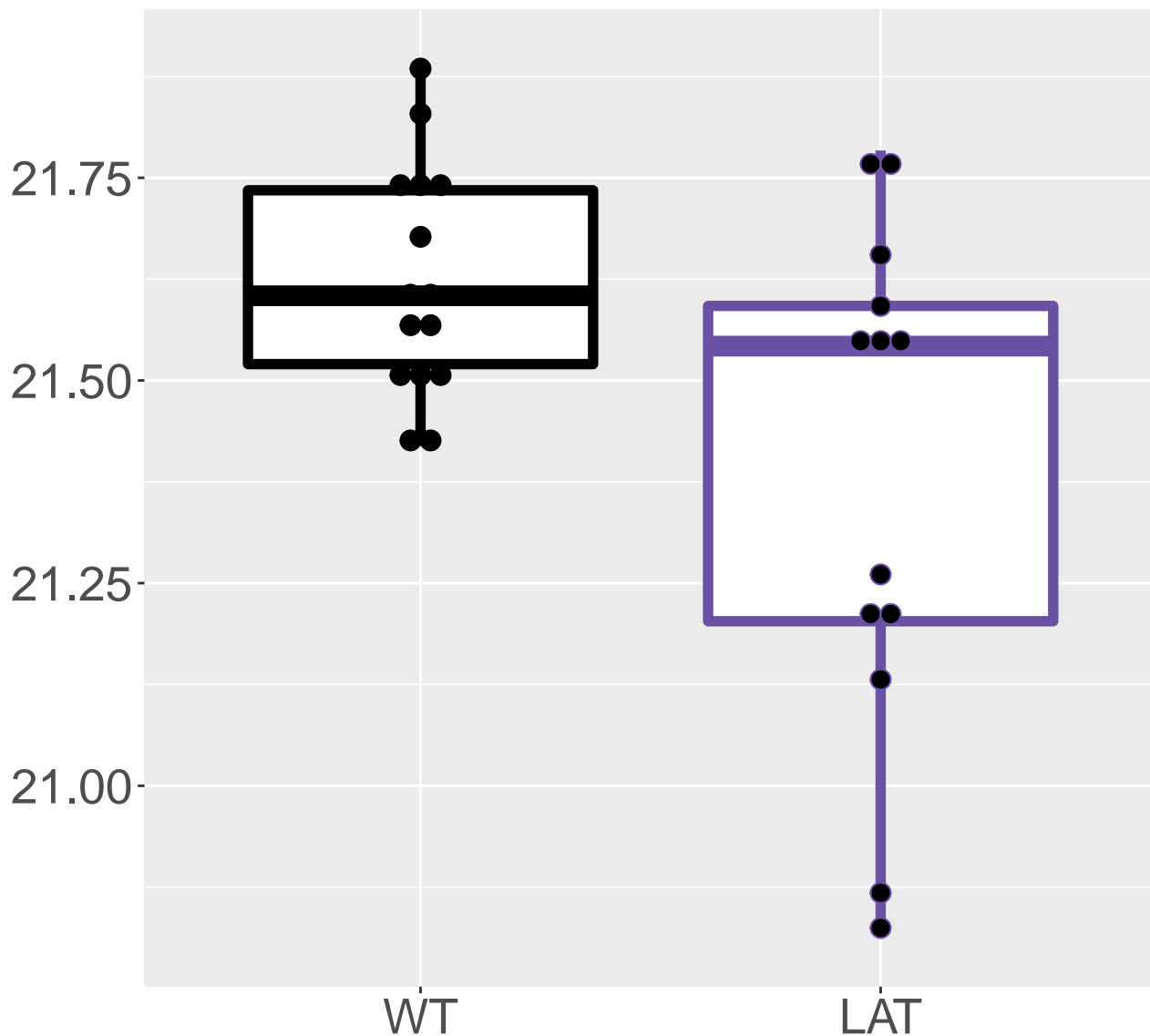
FDR = 0.0035, FC = -0.38, sex***



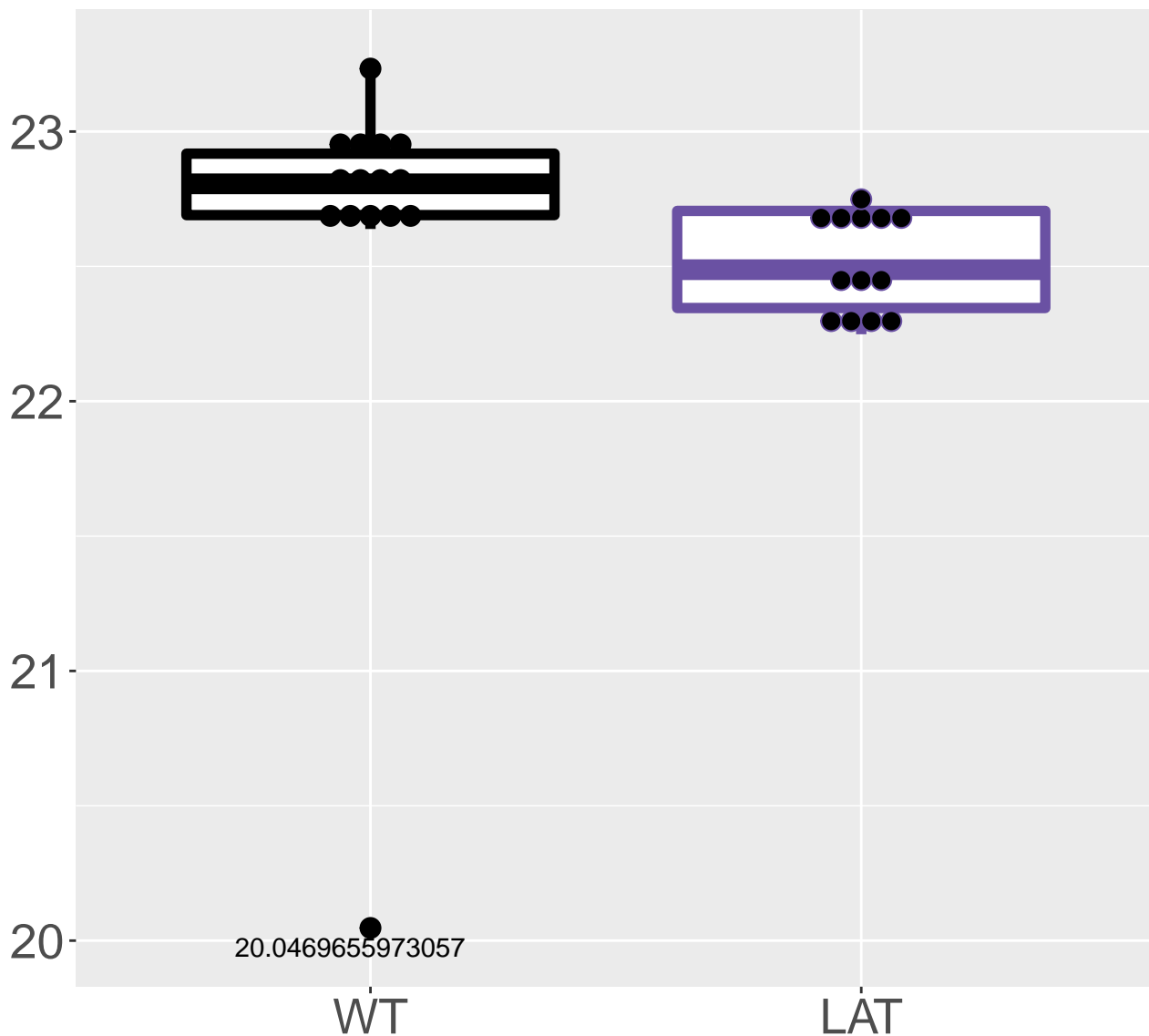
P68510_14-3-3 protein eta
FDR = 0.0035, FC = -0.33, sex***



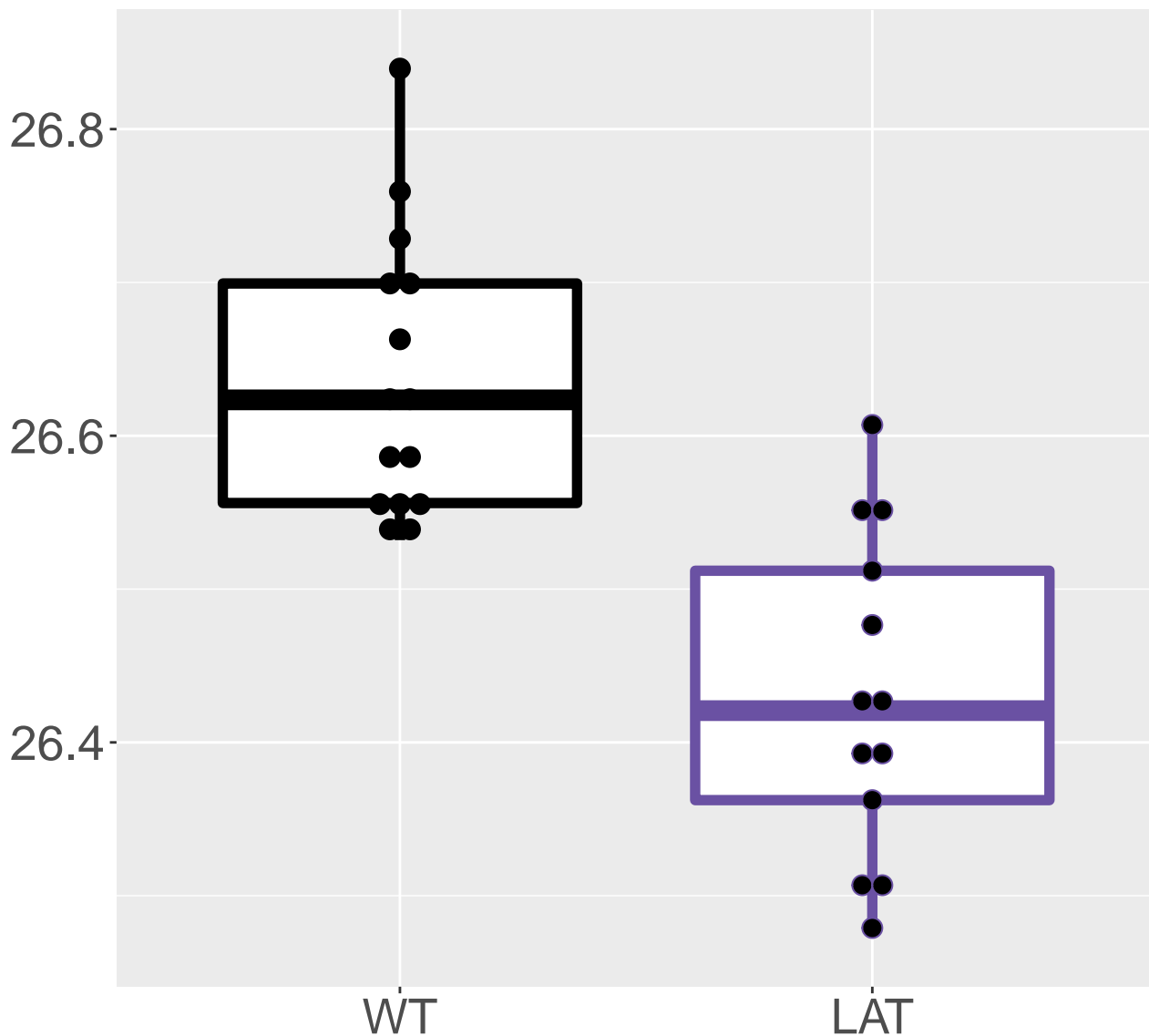
Q9CQ91_NADH dehydrogenase [ubiq.
FDR = 0.0035, FC = -0.53, sex**



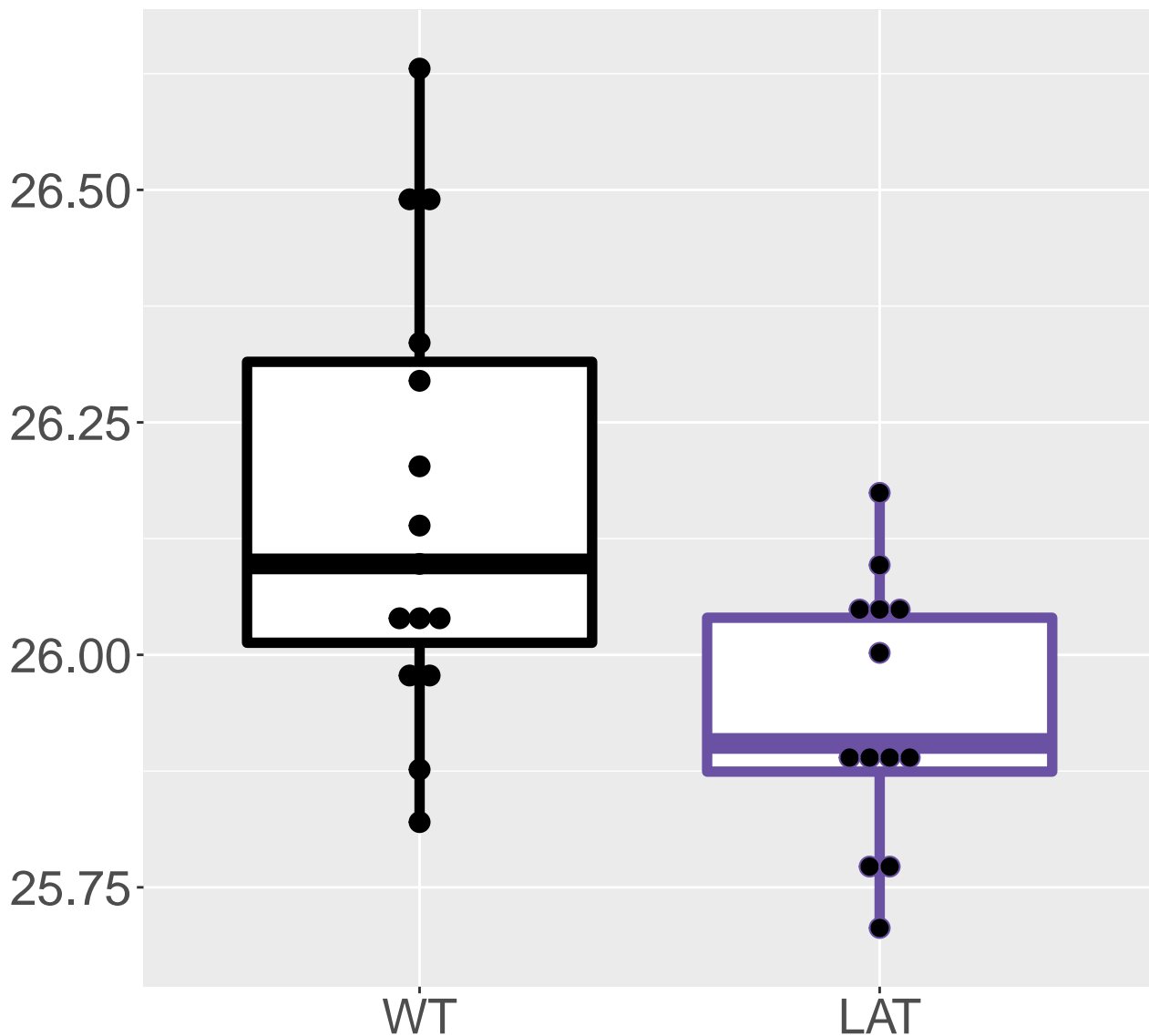
Q9D0T1_NHP2-like protein 1
FDR = 0.0035, FC = -0.49



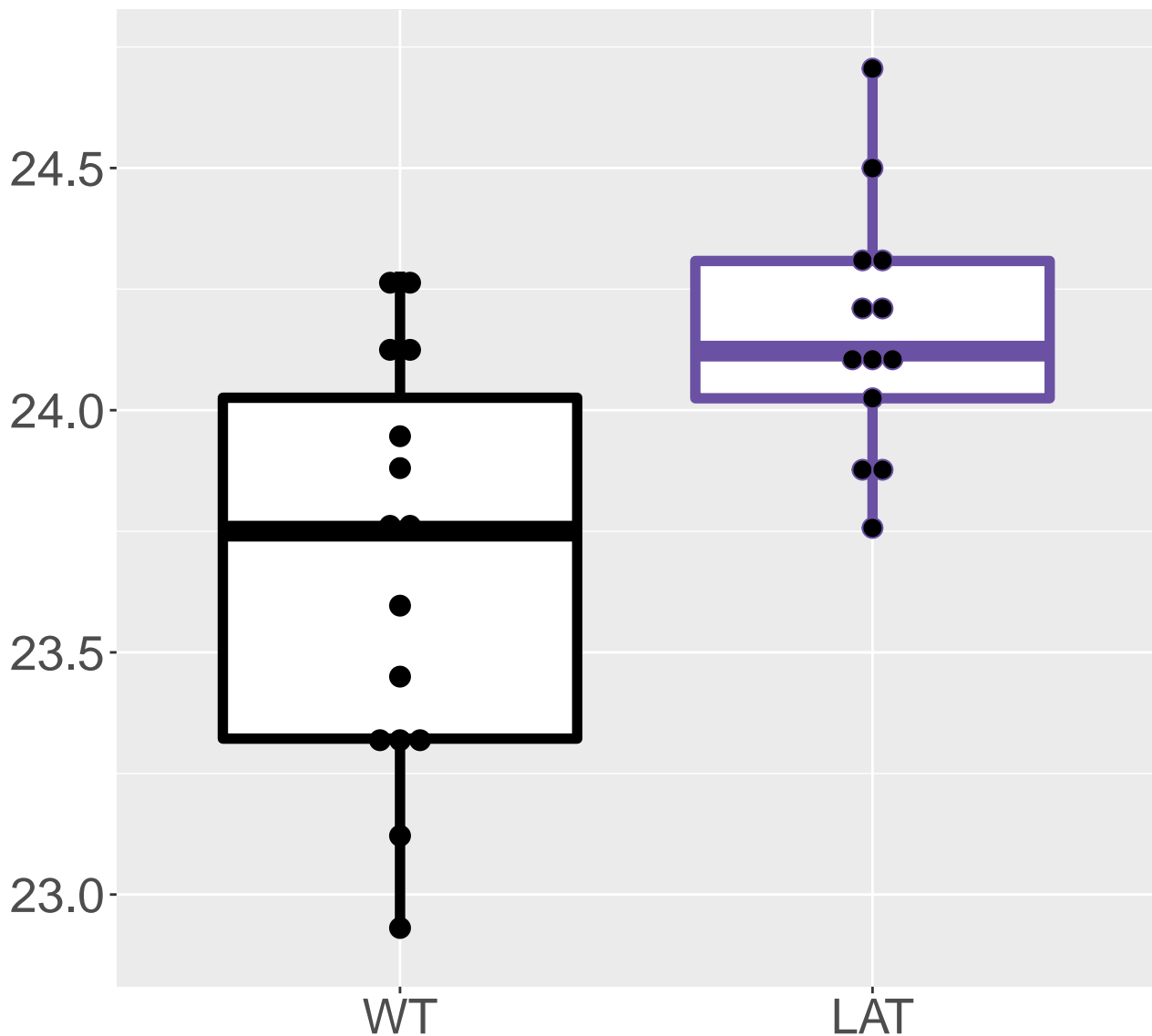
P62751_60S ribosomal protein L2.
FDR = 0.0035, FC = -0.29



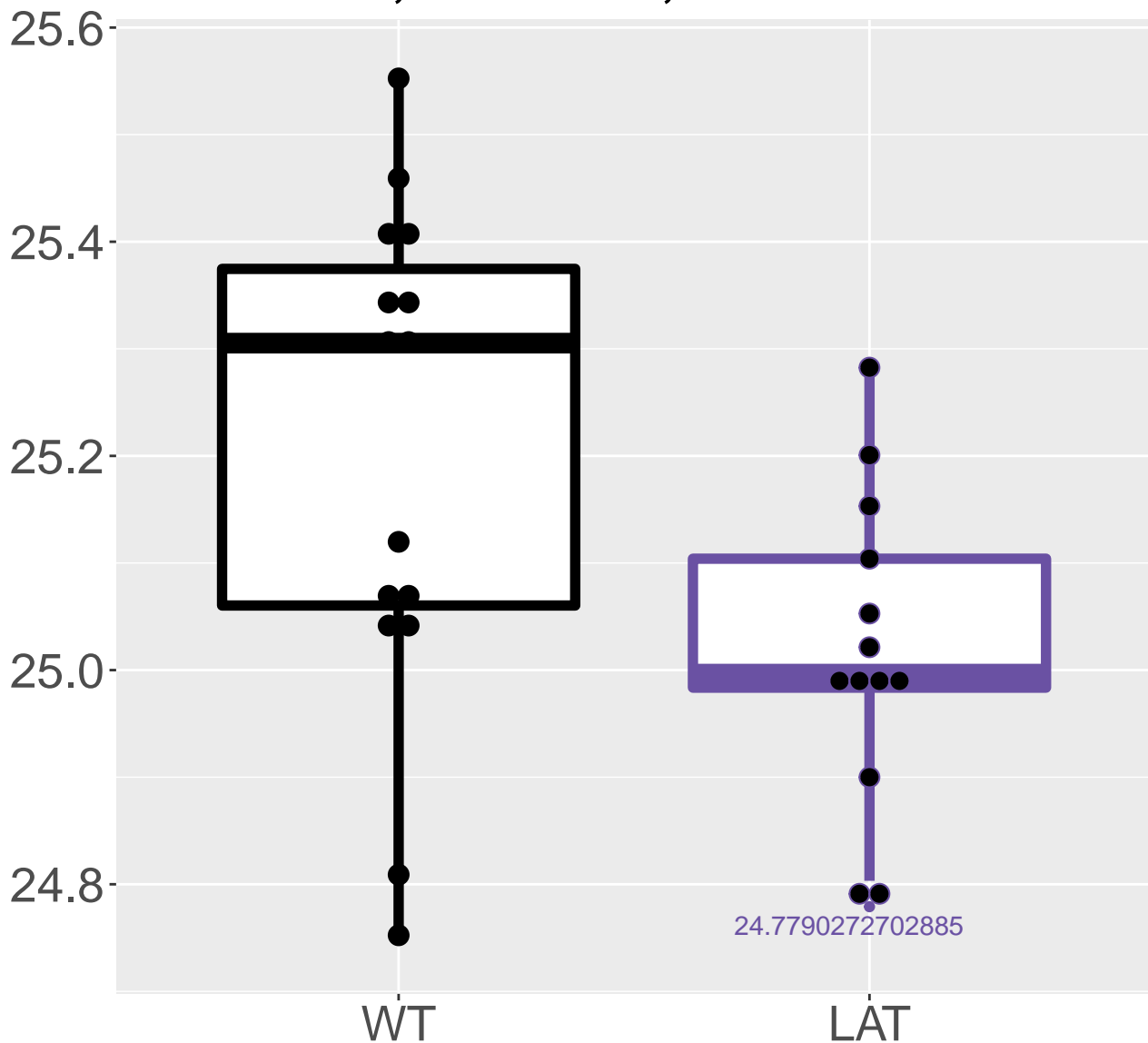
Q9CPQ1_Cytochrome c oxidase sub.
FDR = 0.0037, FC = -0.51, sex**



A2ATU0_Probable 2-oxoglutarate .
FDR = 0.0038, FC = 0.7, sex***

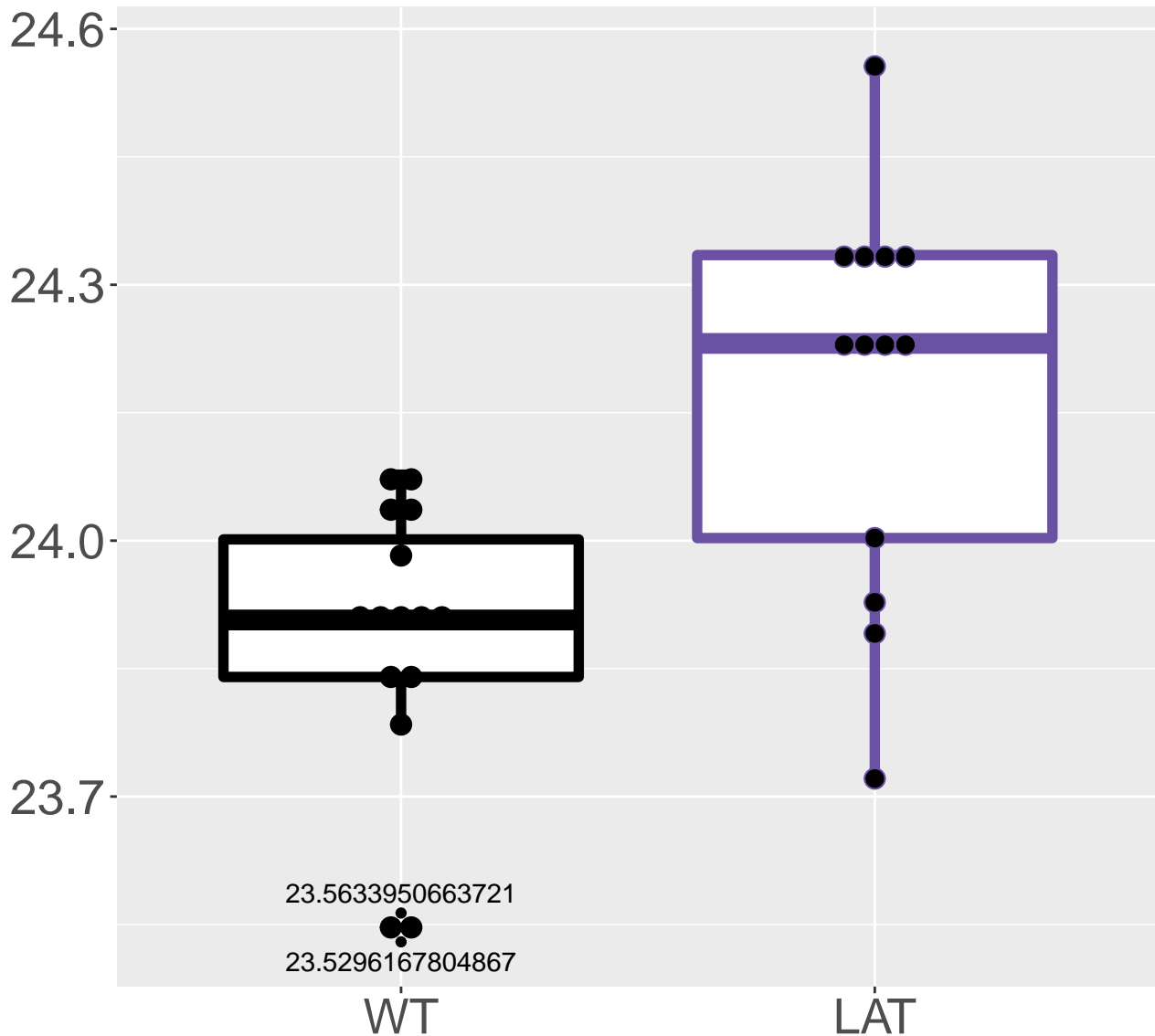


P56135_ATP synthase subunit f, .
FDR = 0.004, FC = -0.42, sex**

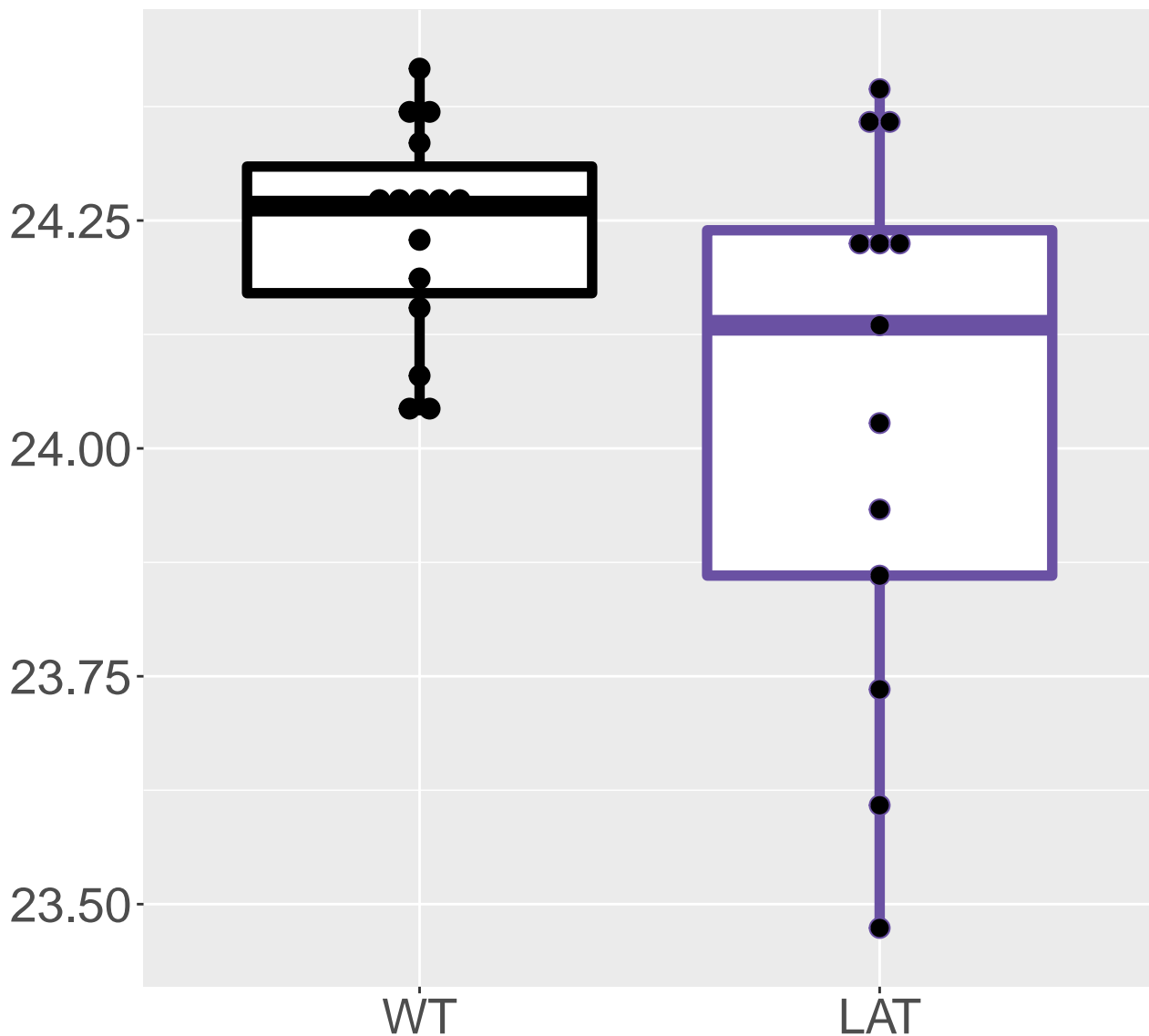


P70168_Importin subunit beta-1

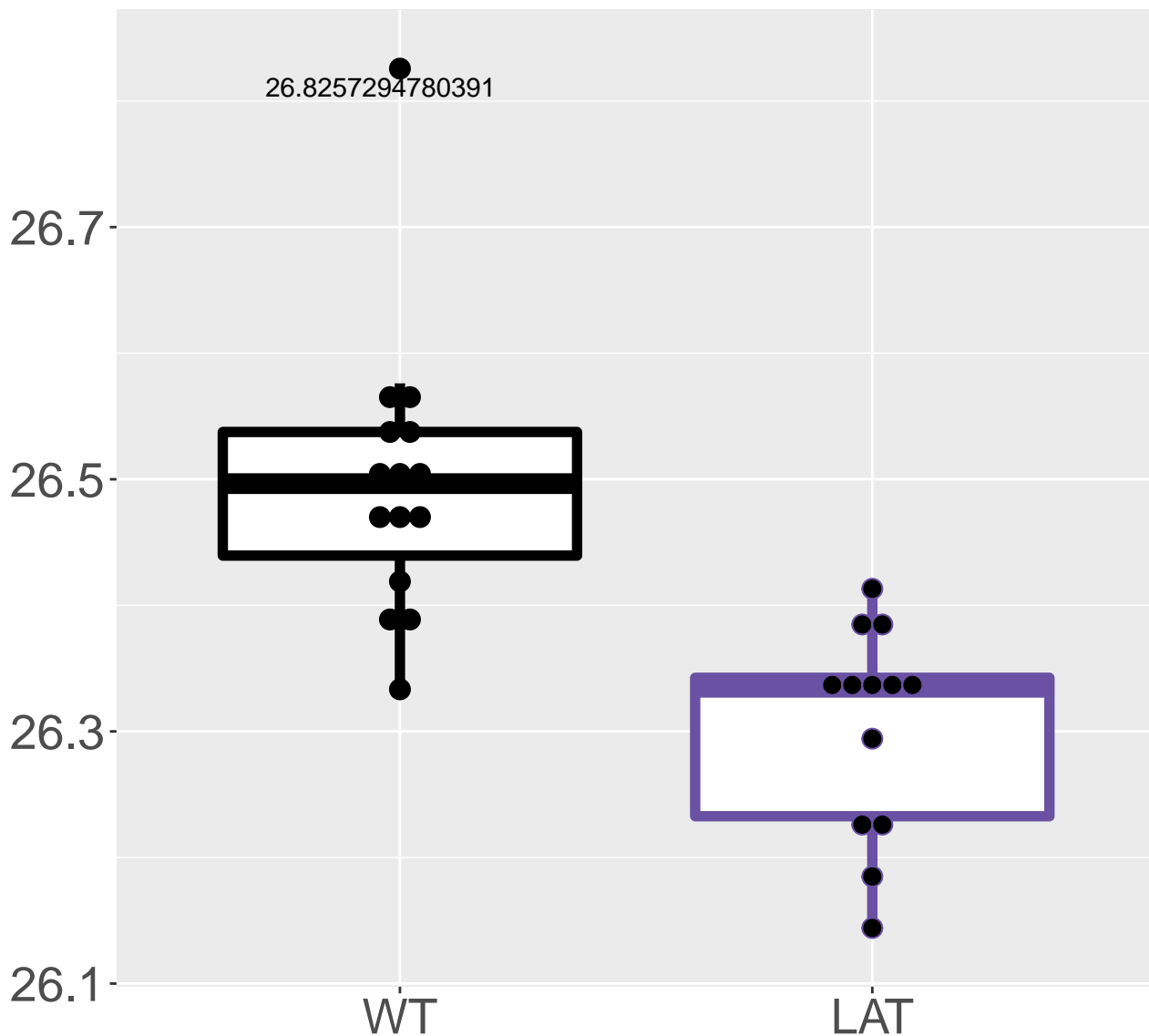
FDR = 0.0042, FC = 0.51



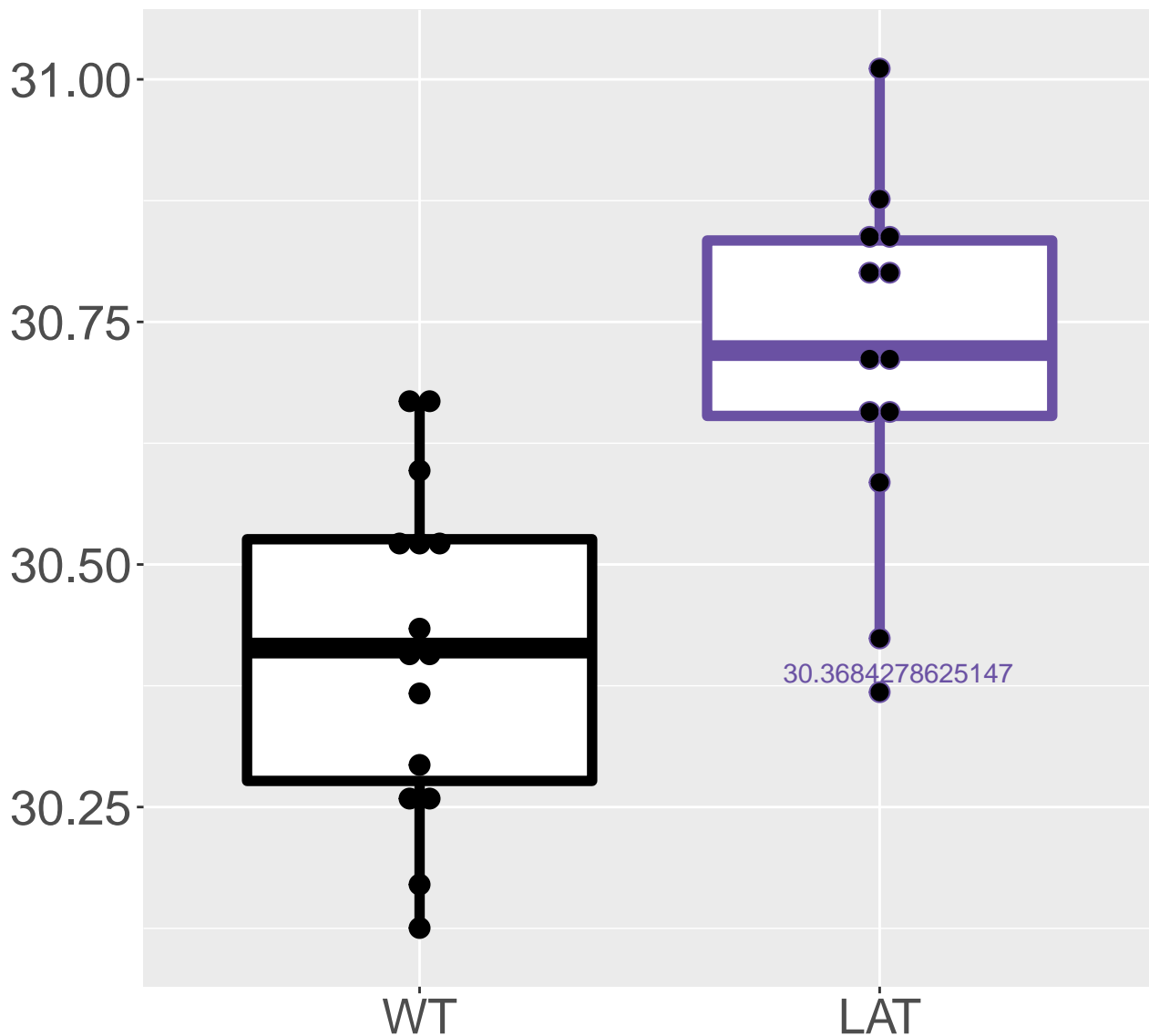
P52503_NADH dehydrogenase [ubiq.
FDR = 0.0042, FC = -0.5, sex**



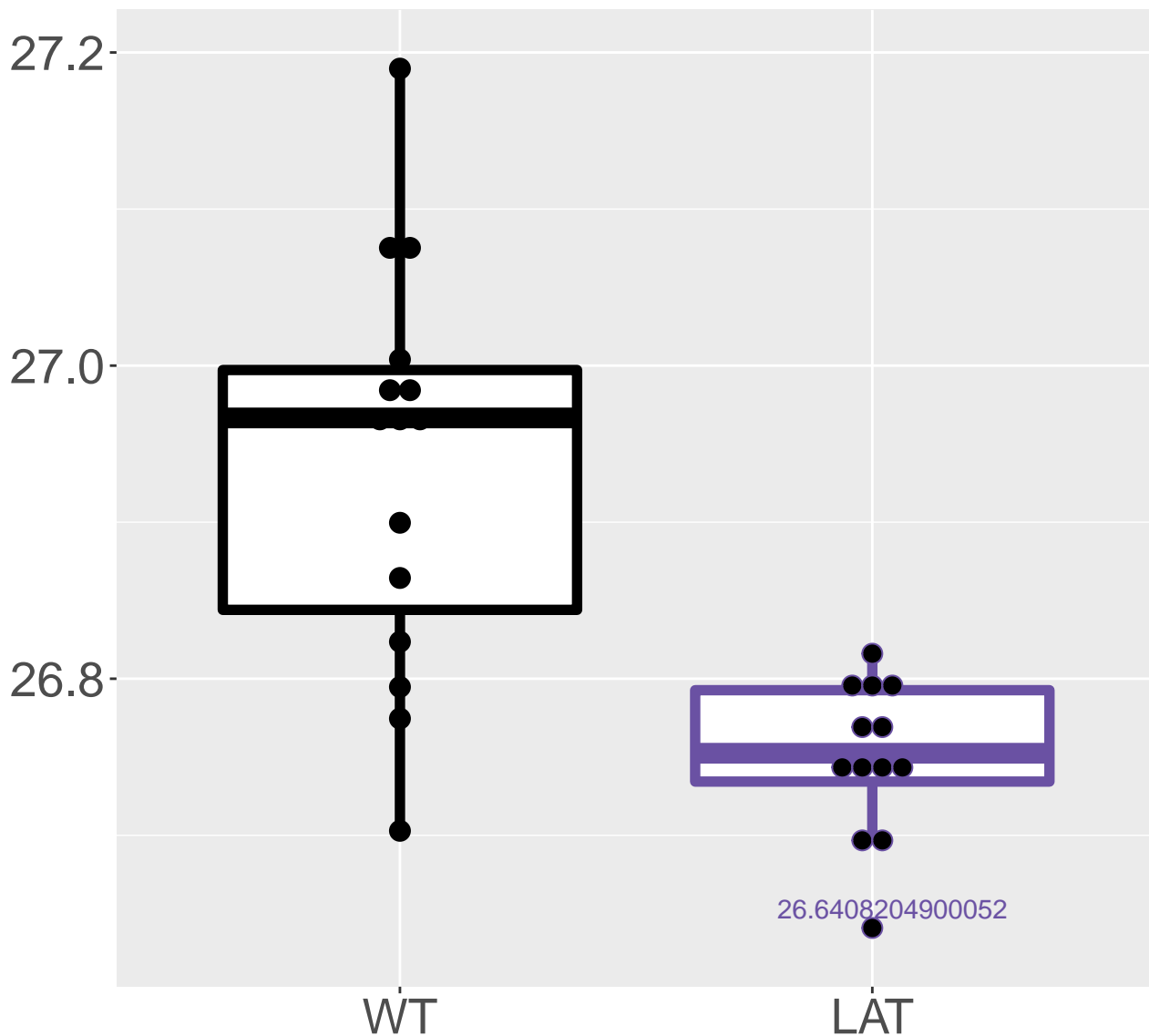
P62900_60S ribosomal protein L31
FDR = 0.0043, FC = -0.32, sex*



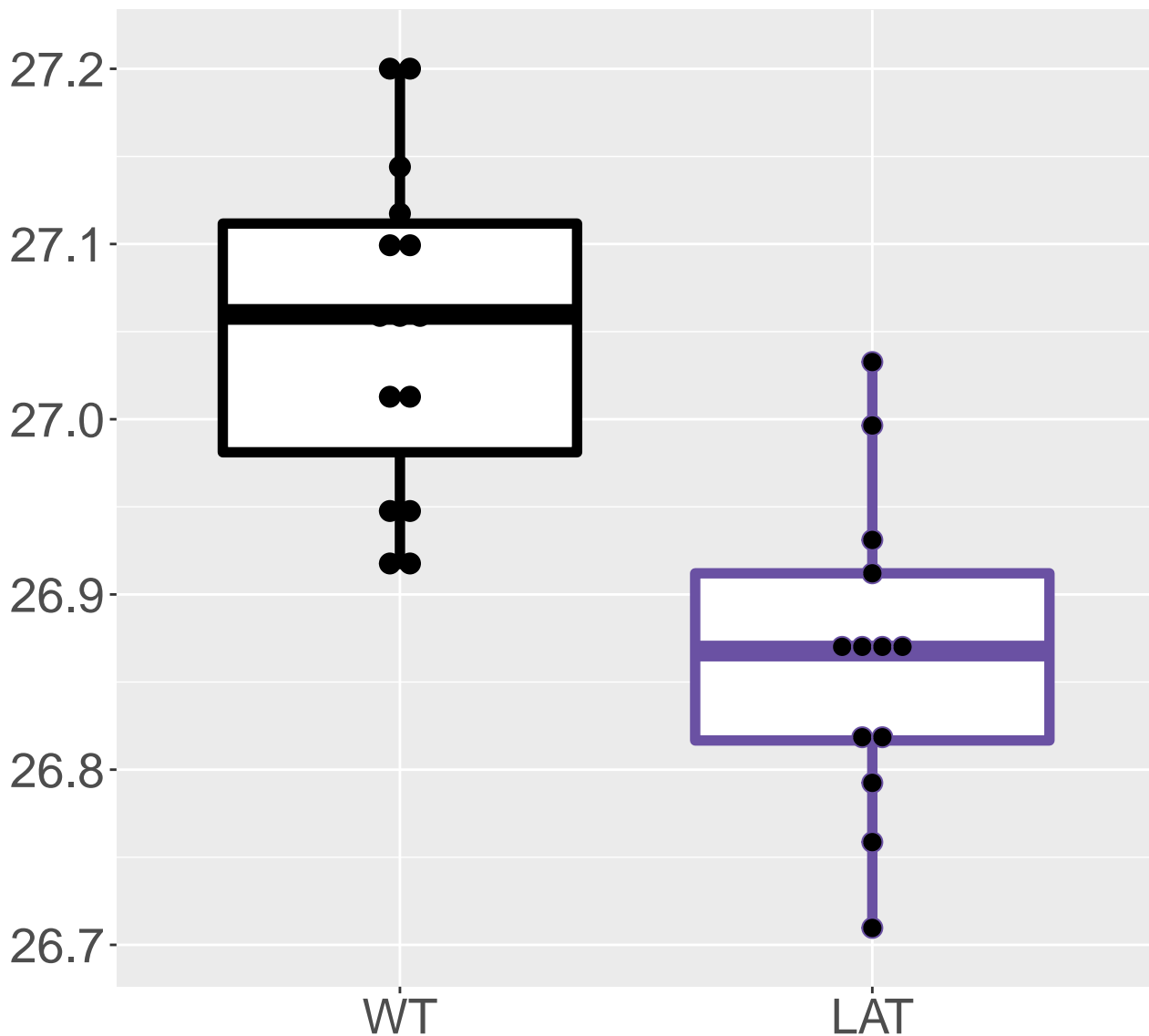
Q8R0Y6_Cytosolic 10-formyltetra.
FDR = 0.0043, FC = 0.42, sex*



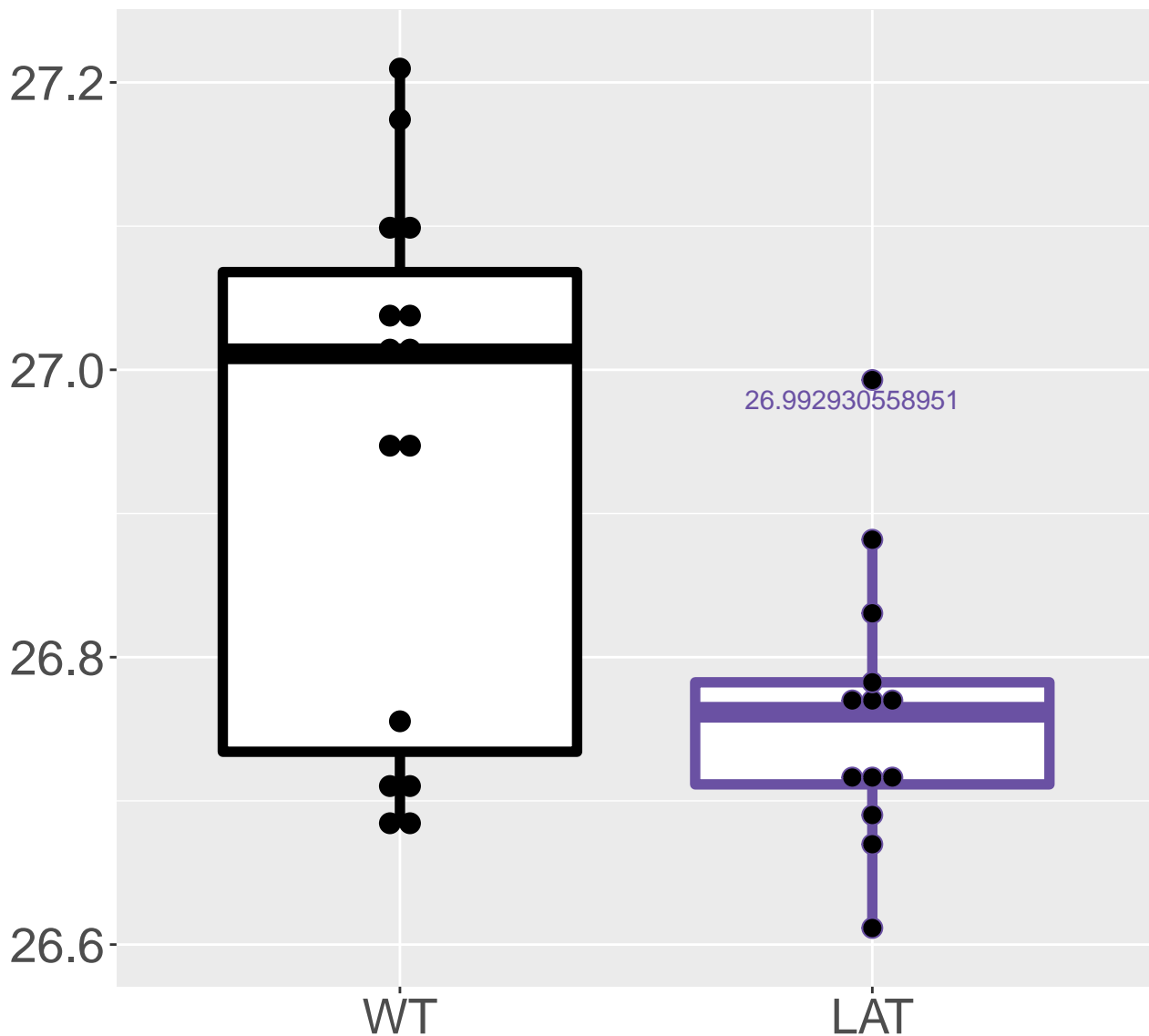
FDR = 0.0048, FC = -0.27



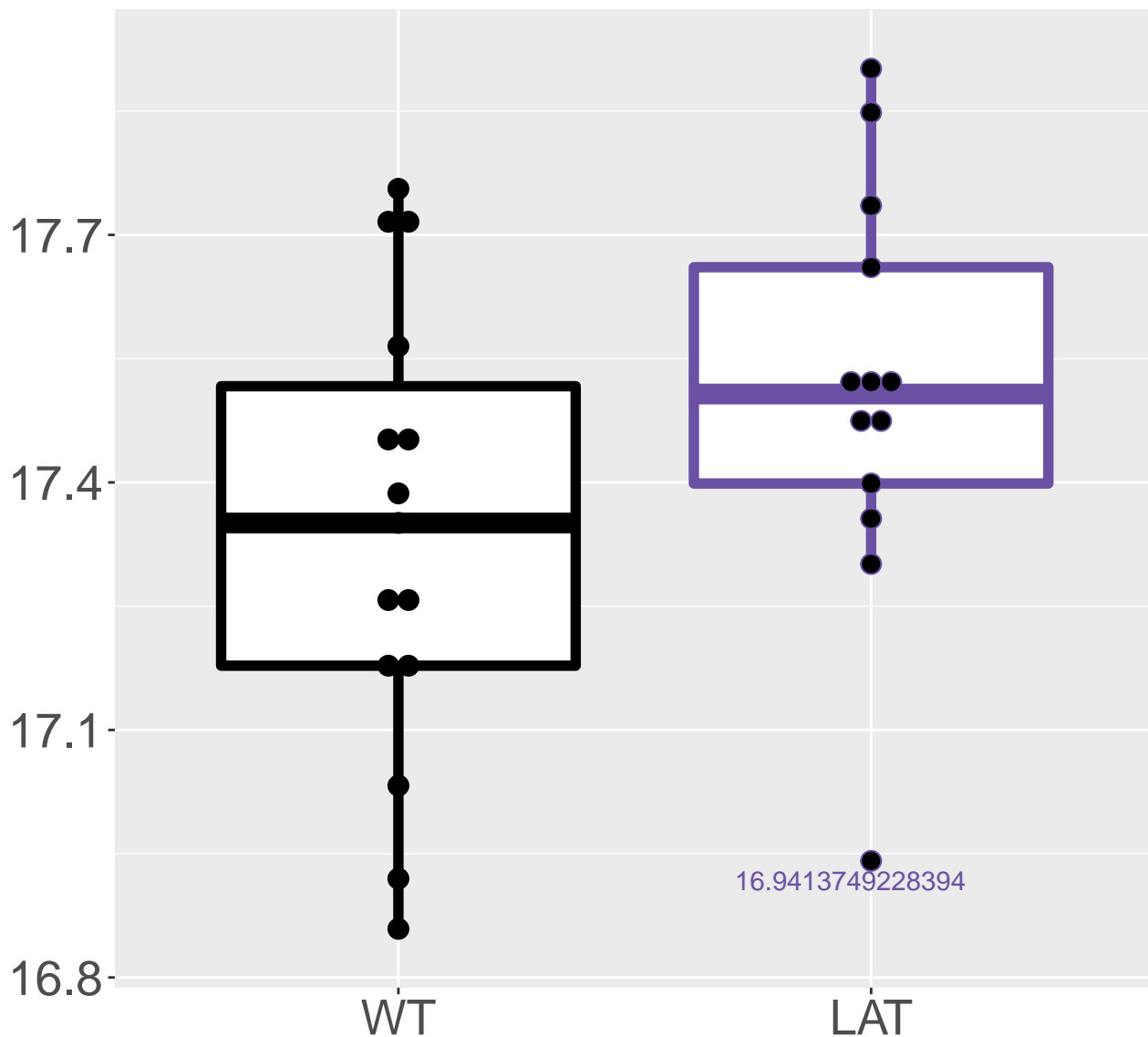
P62270_40S ribosomal protein S18
FDR = 0.0052, FC = -0.28, sex*



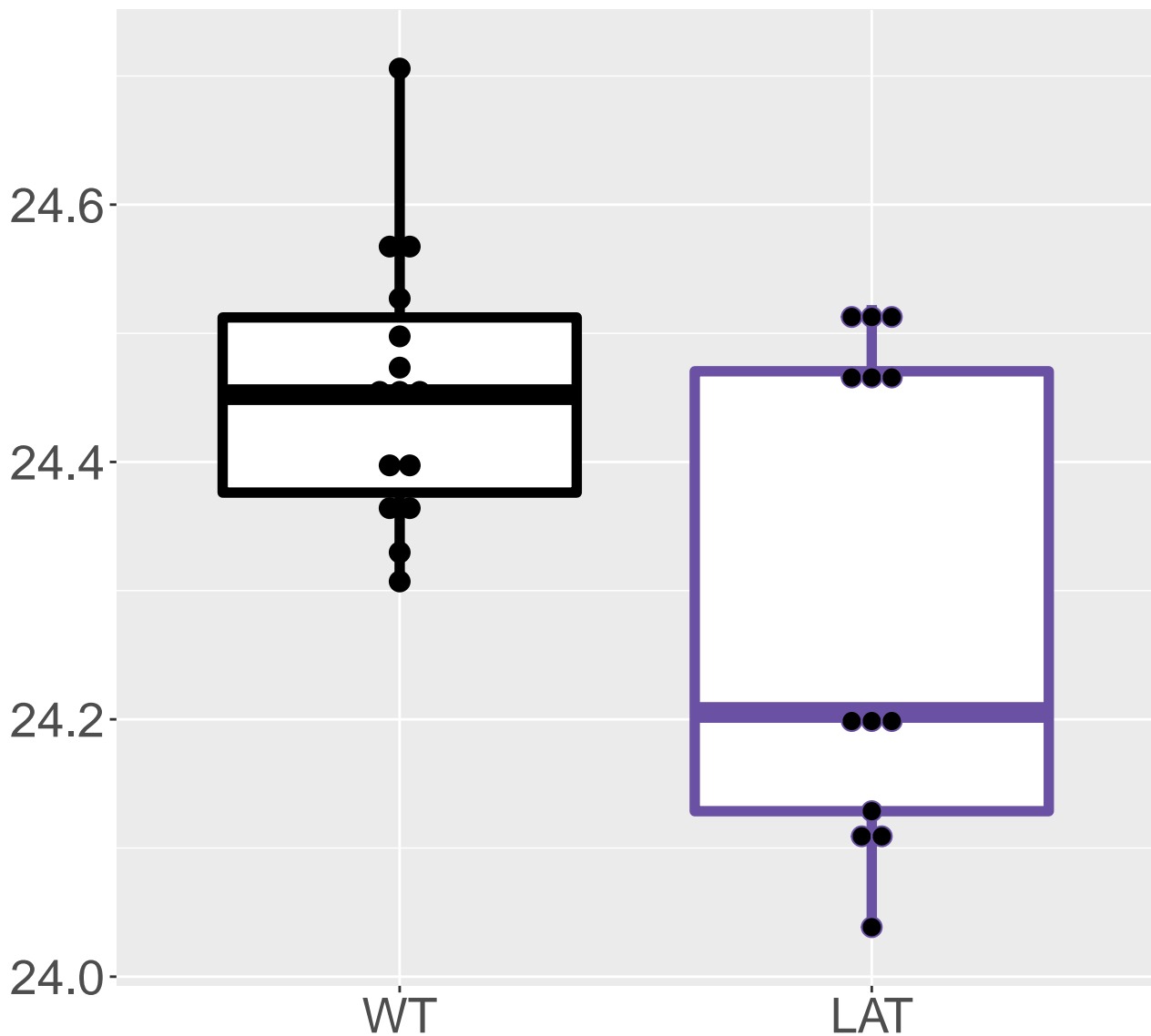
Q9DCQ2_Putative L-aspartate deh.
FDR = 0.0054, FC = -0.29, sex***



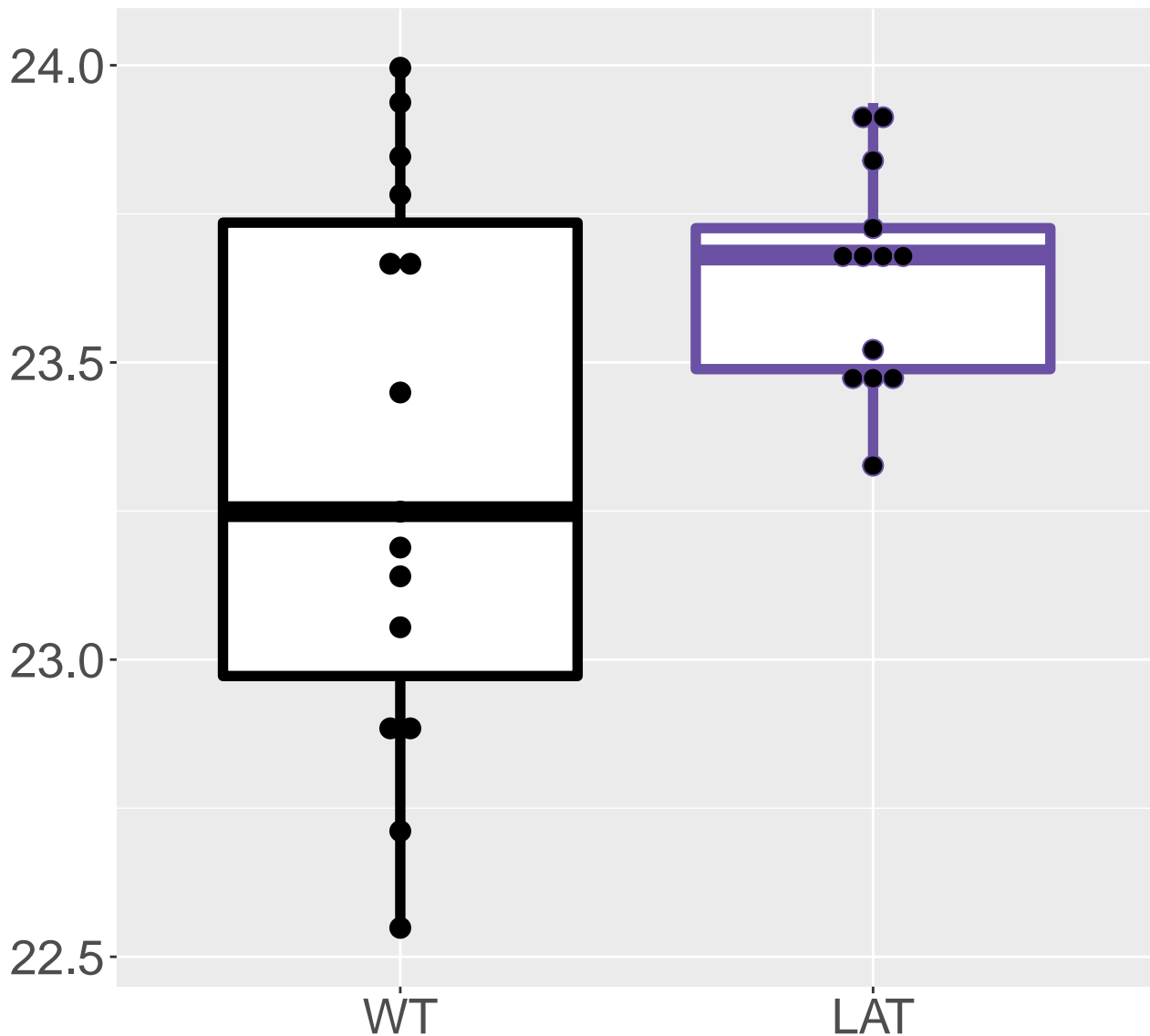
Q9D071_MMS19 nucleotide excisio.
FDR = 0.0056, FC = 0.52, sex**



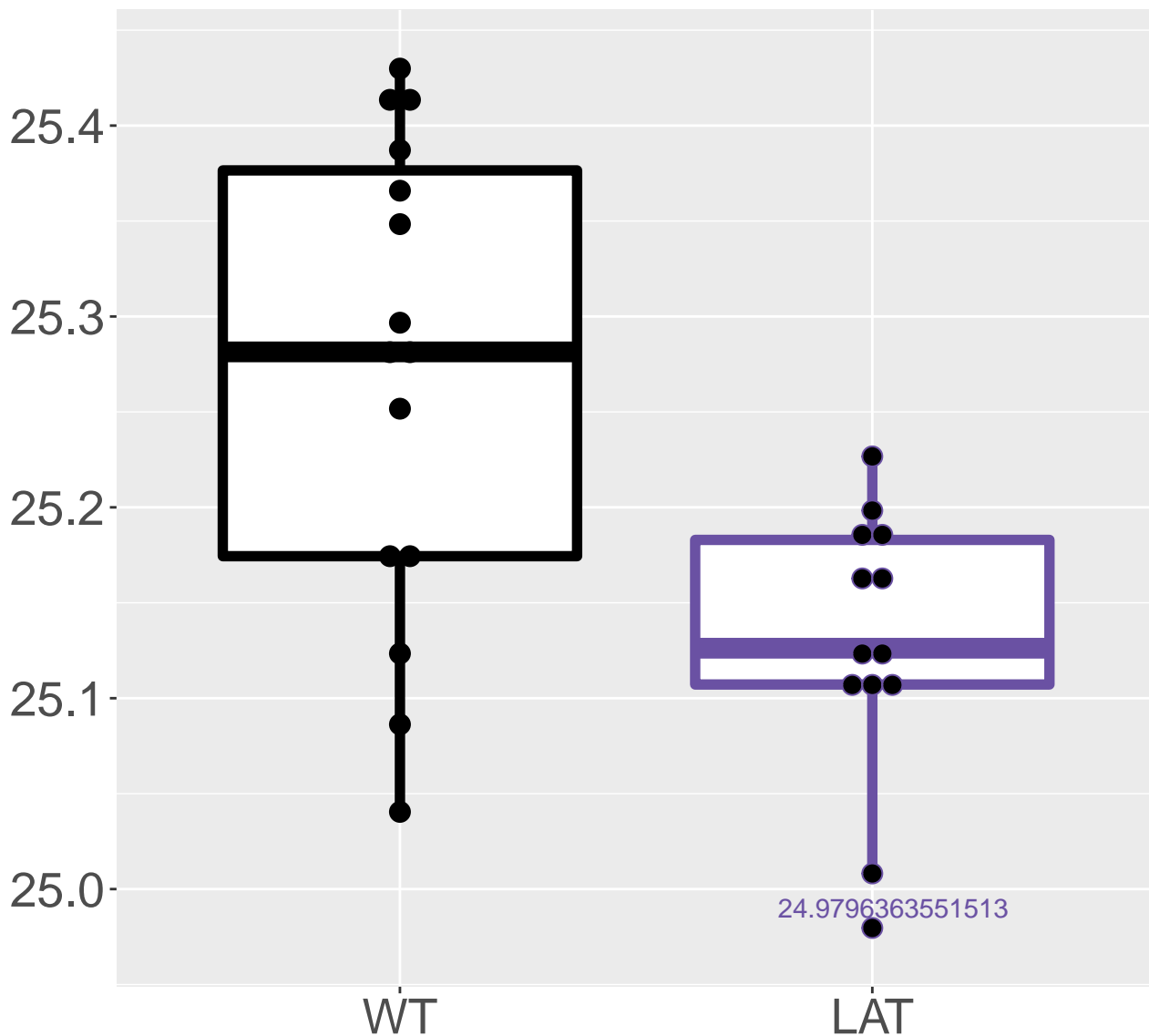
Q9CQ54_NADH dehydrogenase [ubiq.
FDR = 0.0058, FC = -0.33, sex**



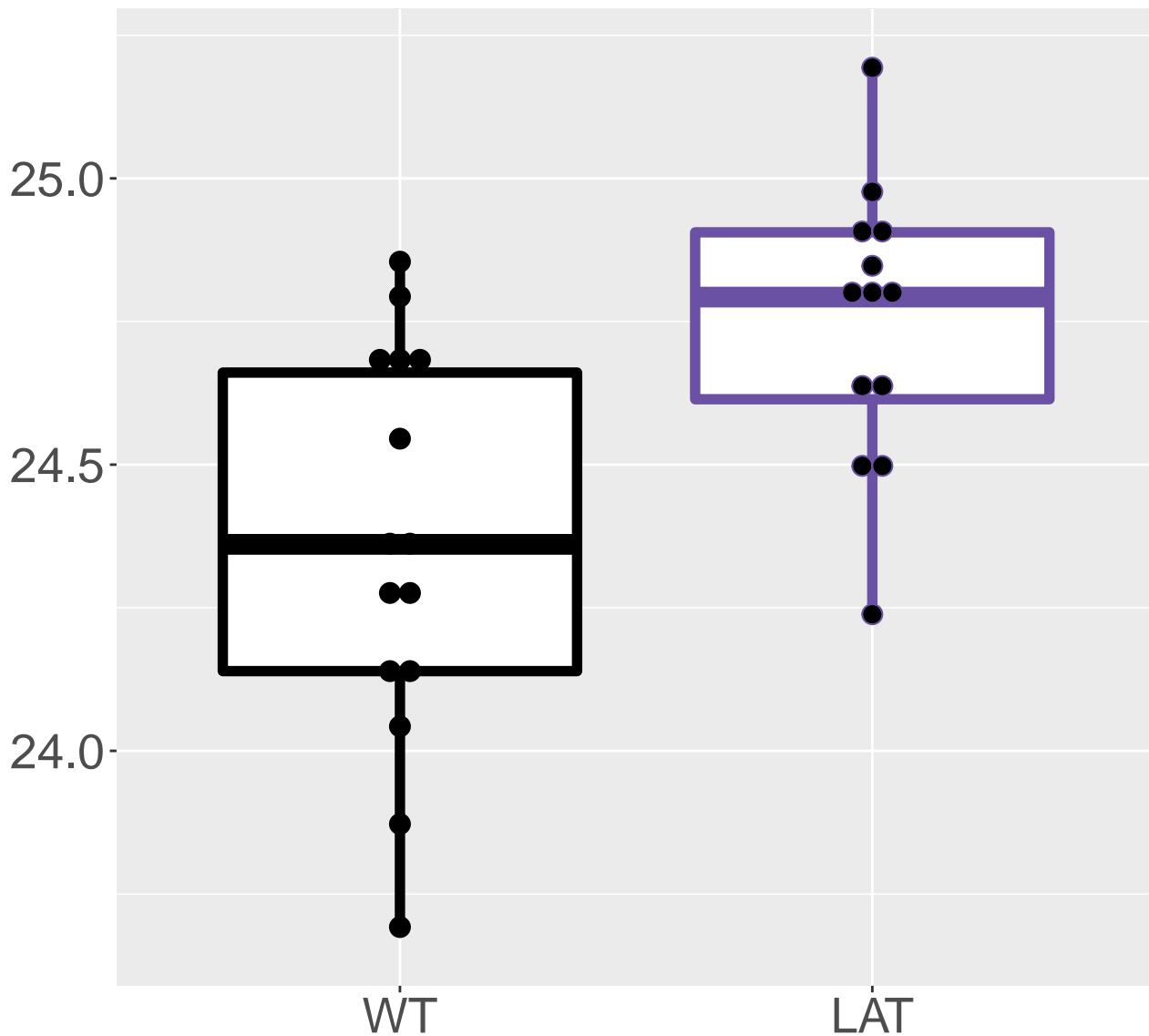
P35576_Glucose-6-phosphatase
FDR = 0.0059, FC = 0.65, sex***



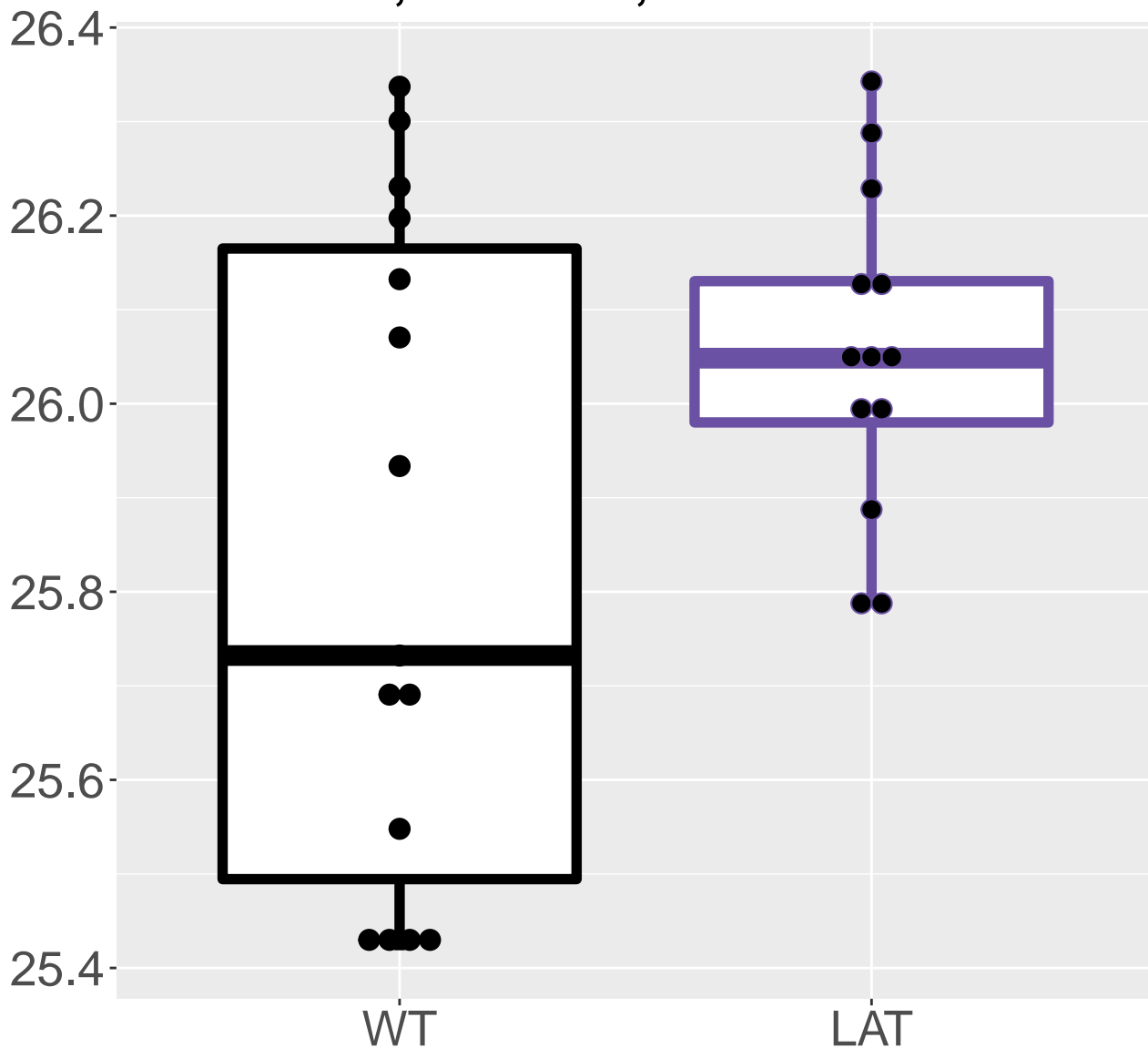
P61089_Ubiquitin-conjugating en.
FDR = 0.0059, FC = -0.2, sex**



Q6PB66_Leucine-rich PPR motif-c.
FDR = 0.0063, FC = 0.59, sex**

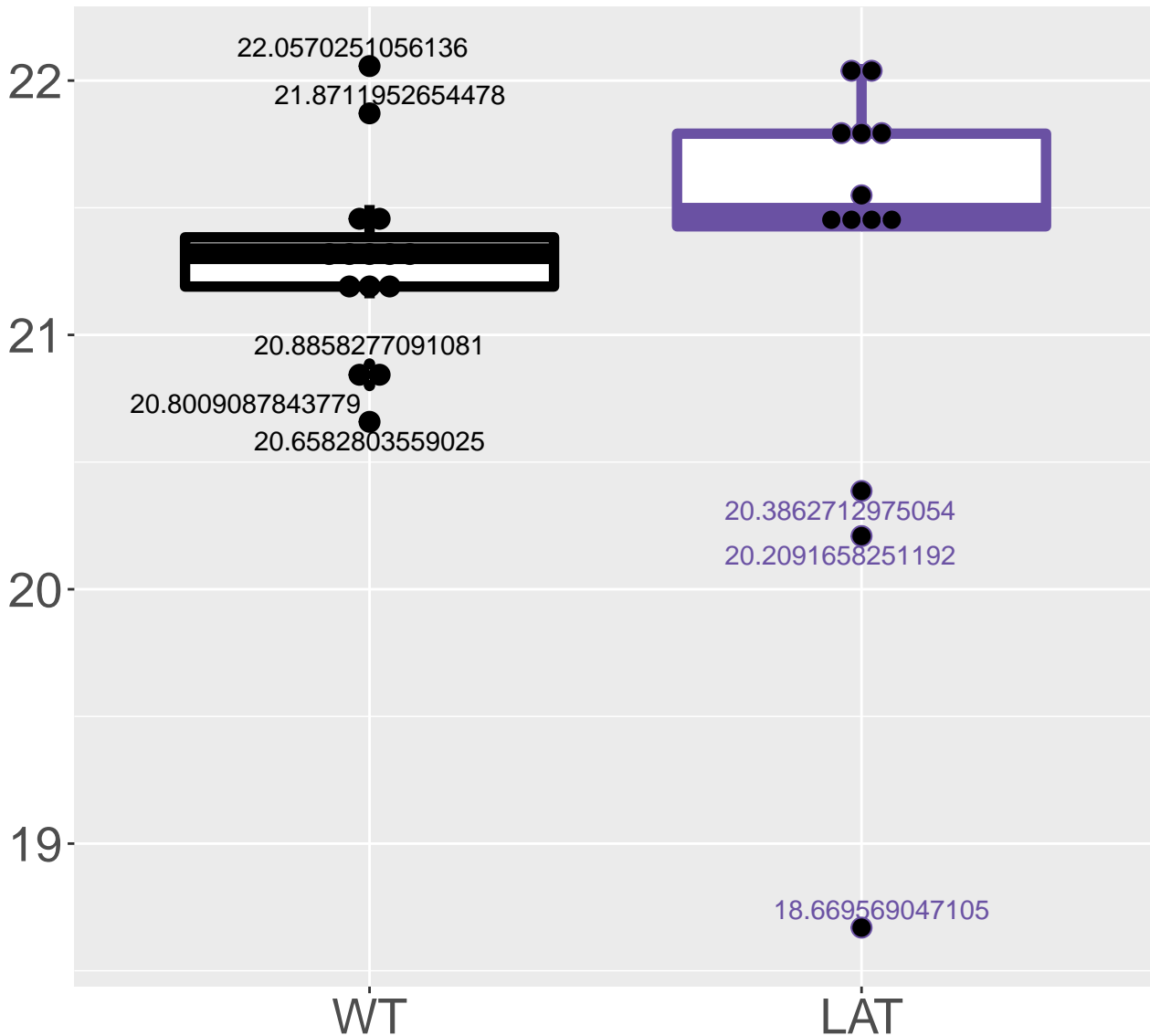


Q8VCB3_Glycogen [starch] syntha.
FDR = 0.007, FC = 0.37, sex***

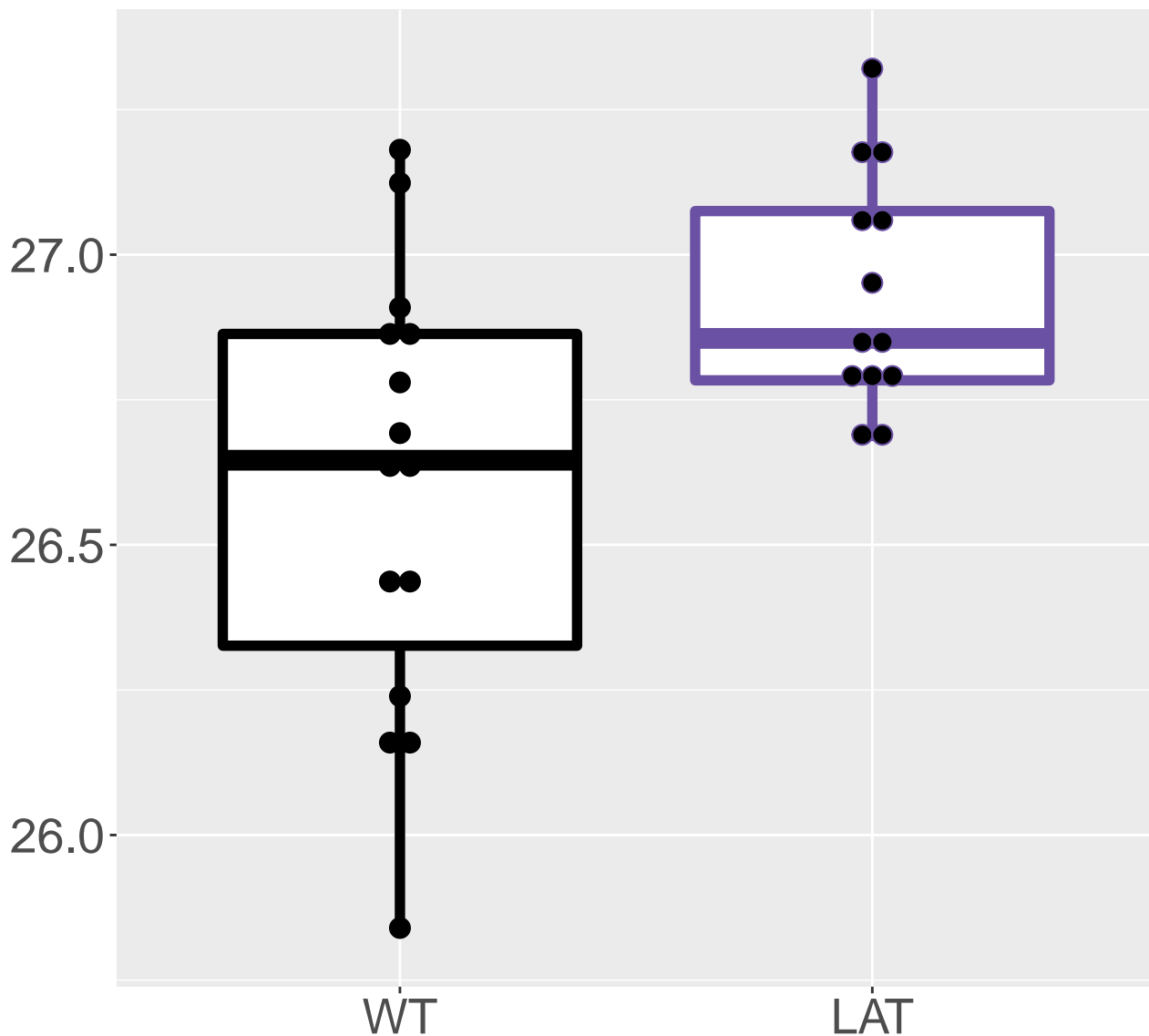


Q9WTI7_Unconventional myosin-Ic

FDR = 0.0073, FC = 0.7

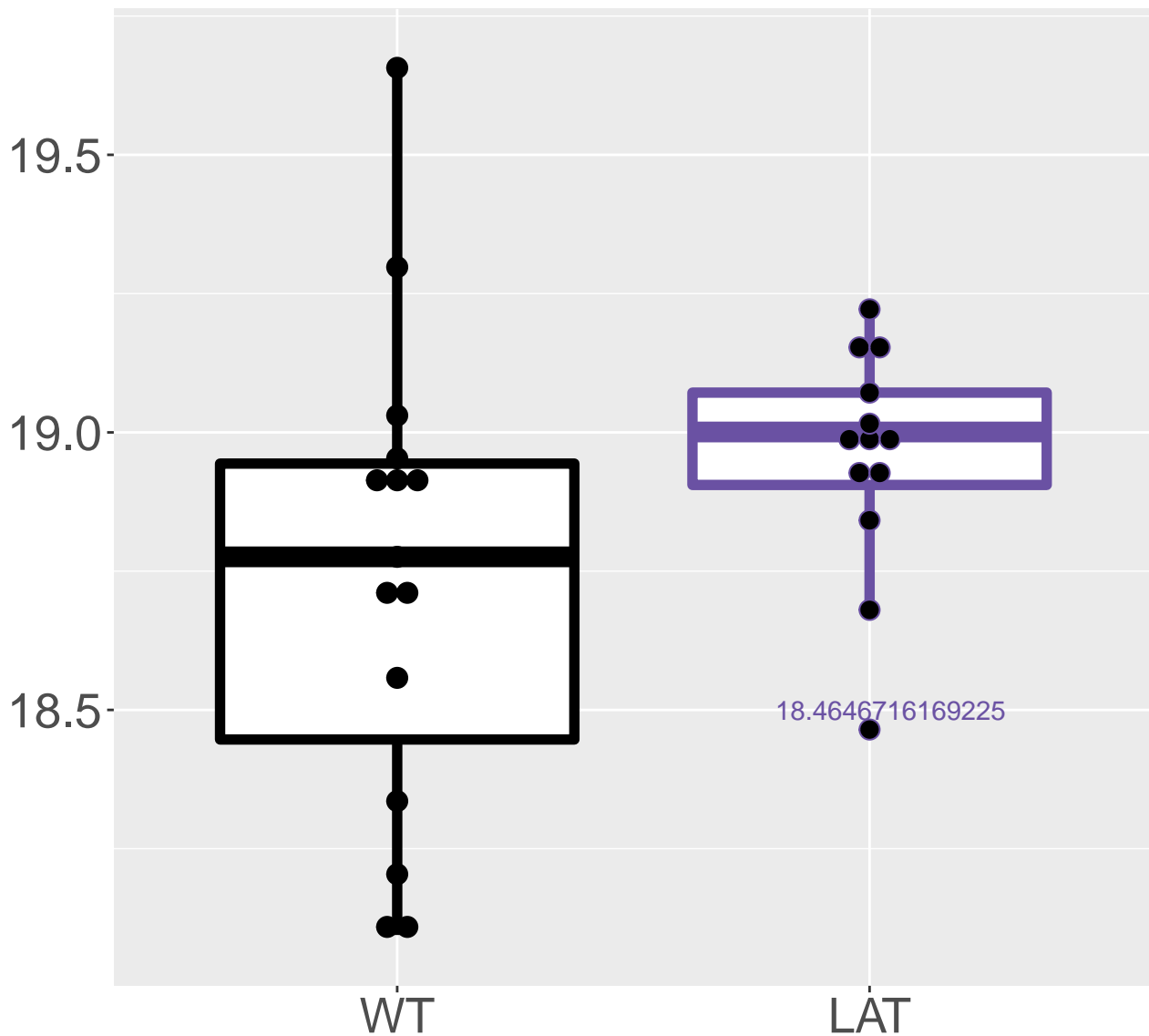


Q8QZR3_Pyrethroid hydrolase Ces.
FDR = 0.0073, FC = 0.63, sex**

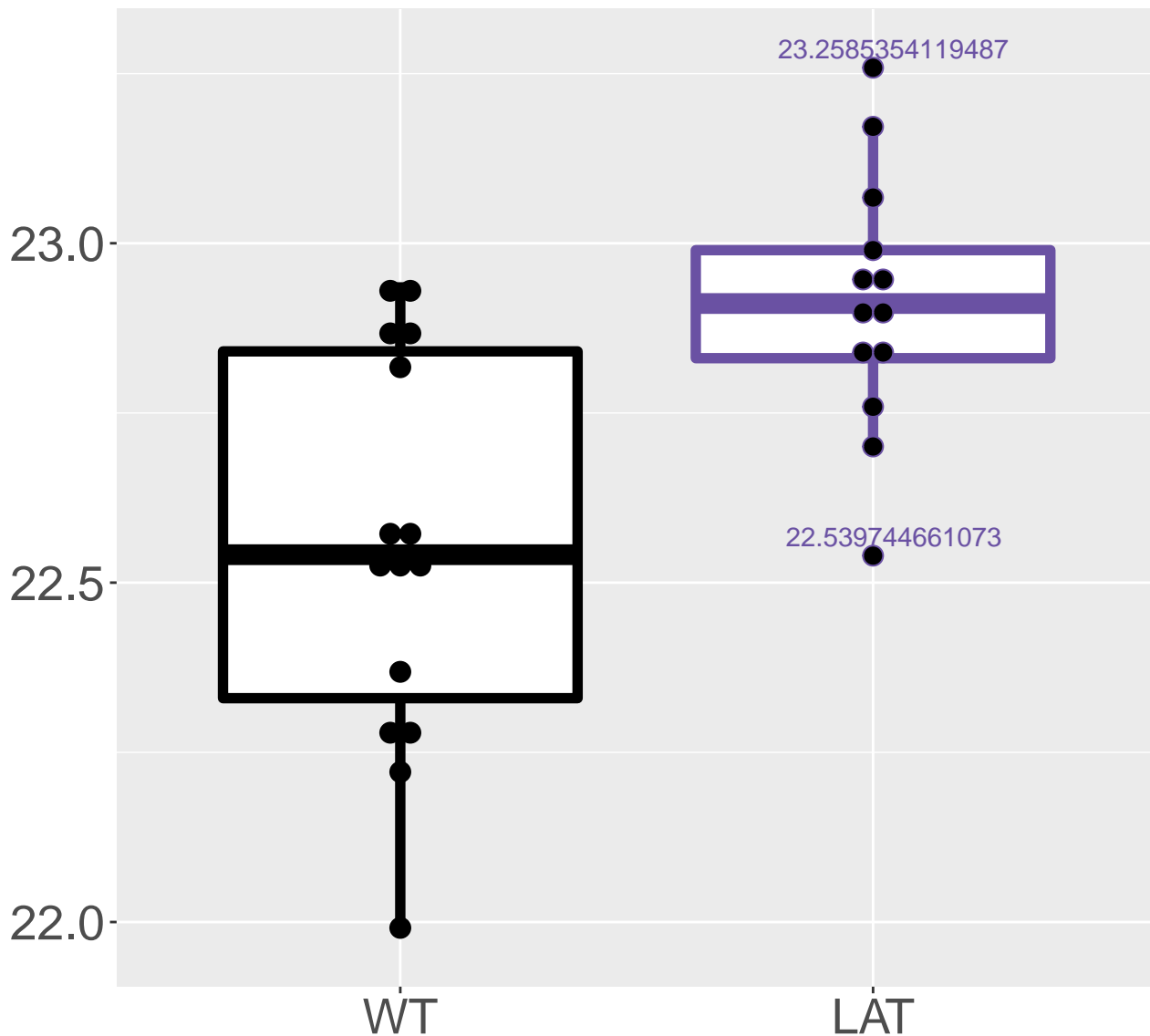


Q99J56_Derlin-1

FDR = 0.0073, FC = 0.63, sex**

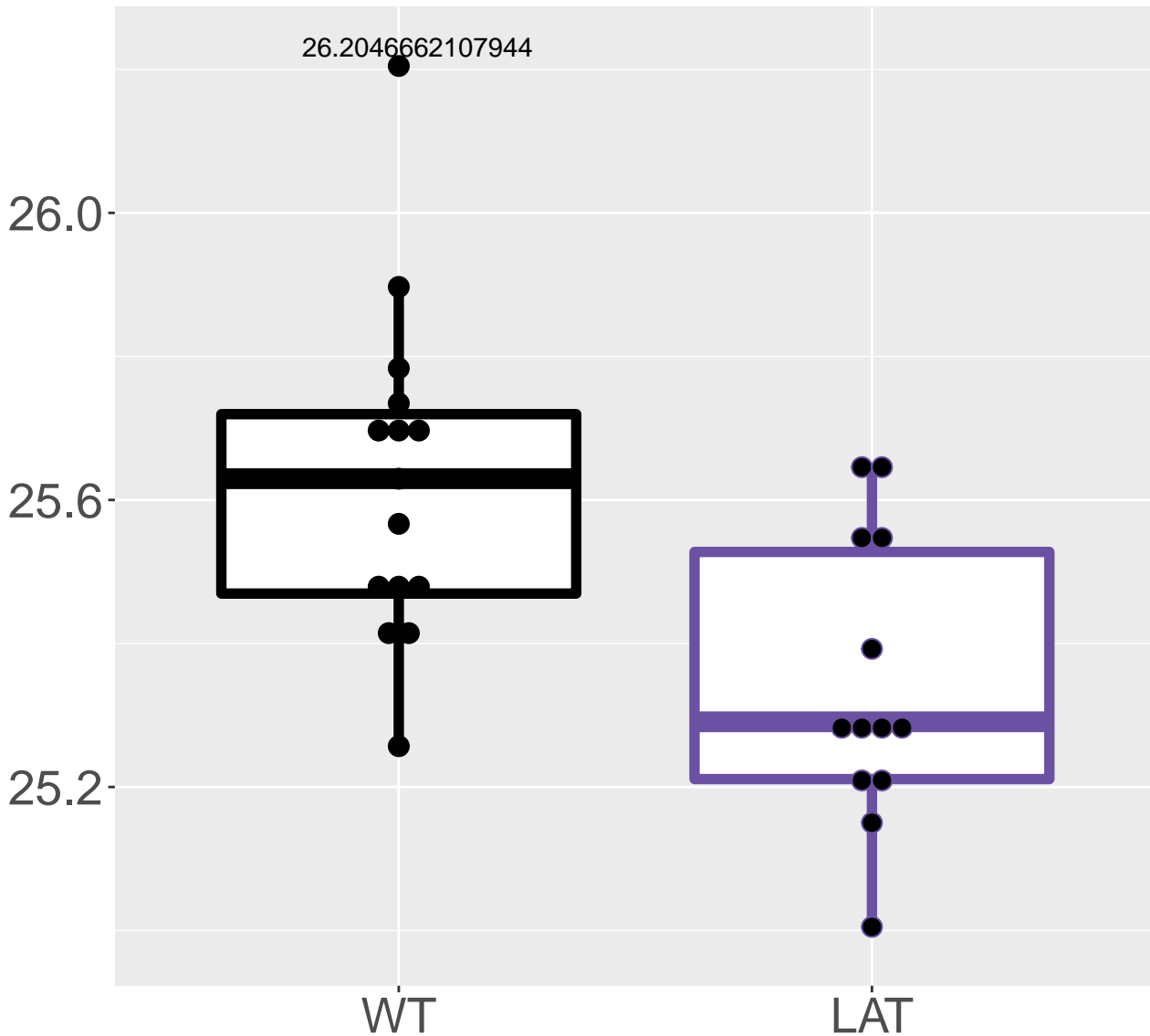


P03921_NADH-ubiquinone oxidored.
FDR = 0.0073, FC = 0.55, sex**

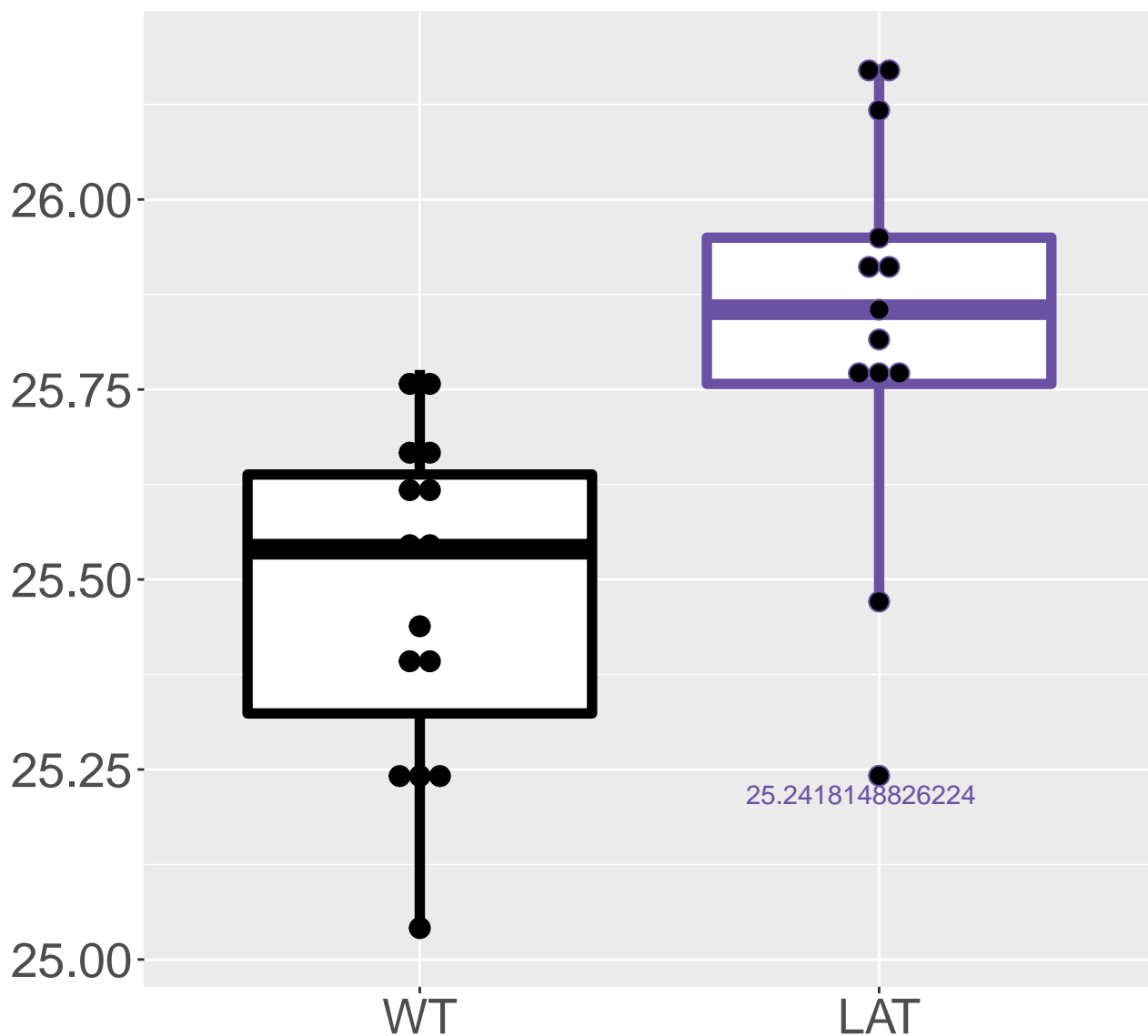


Q6Z WV7_60S ribosomal protein L35

FDR = 0.0073, FC = -0.52

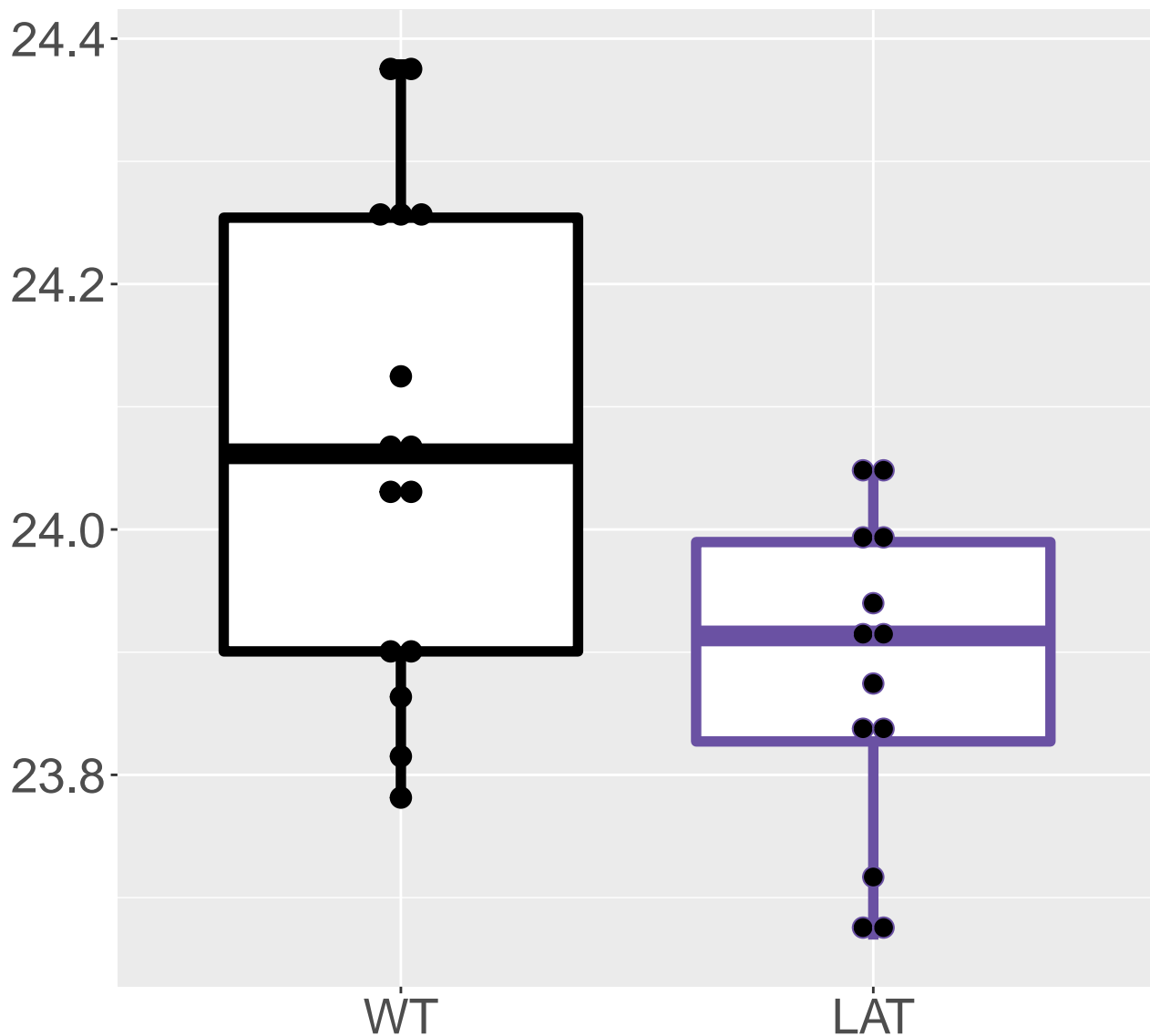


Q8CHR6_Dihydropyrimidine dehydr.
FDR = 0.0073, FC = 0.5

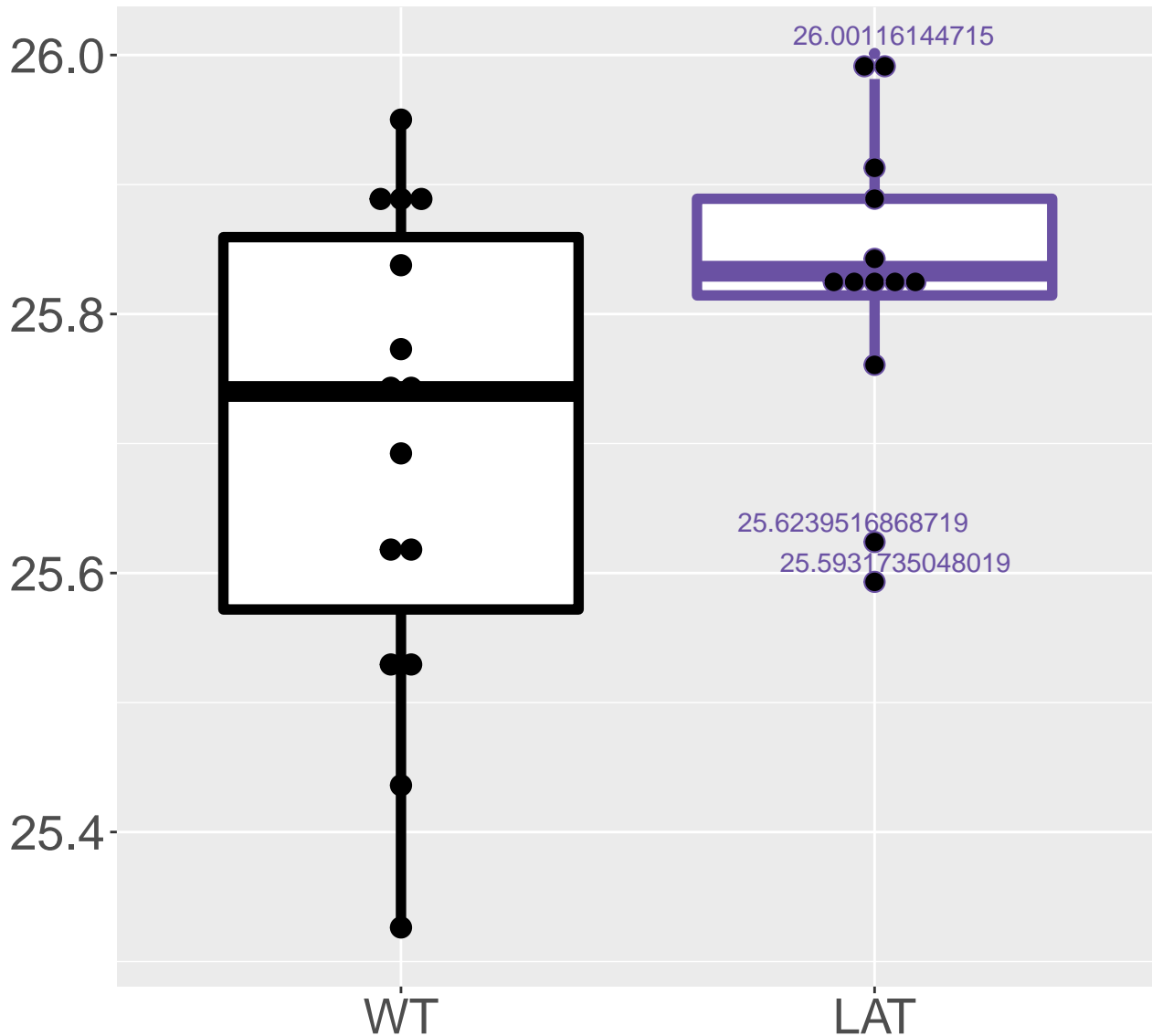


P0C0S6_Histone H2A.Z

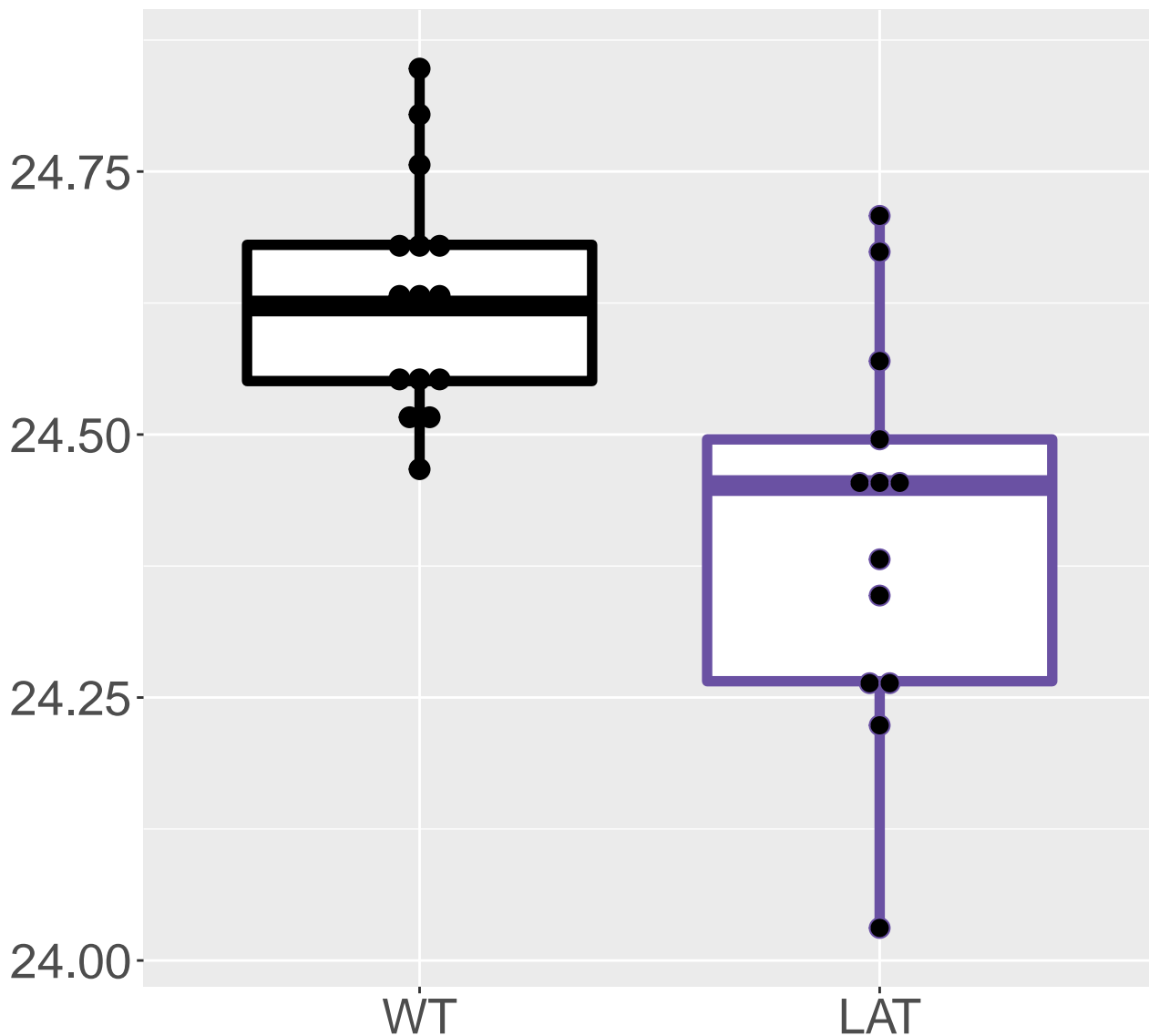
FDR = 0.0073, FC = -0.39, sex**



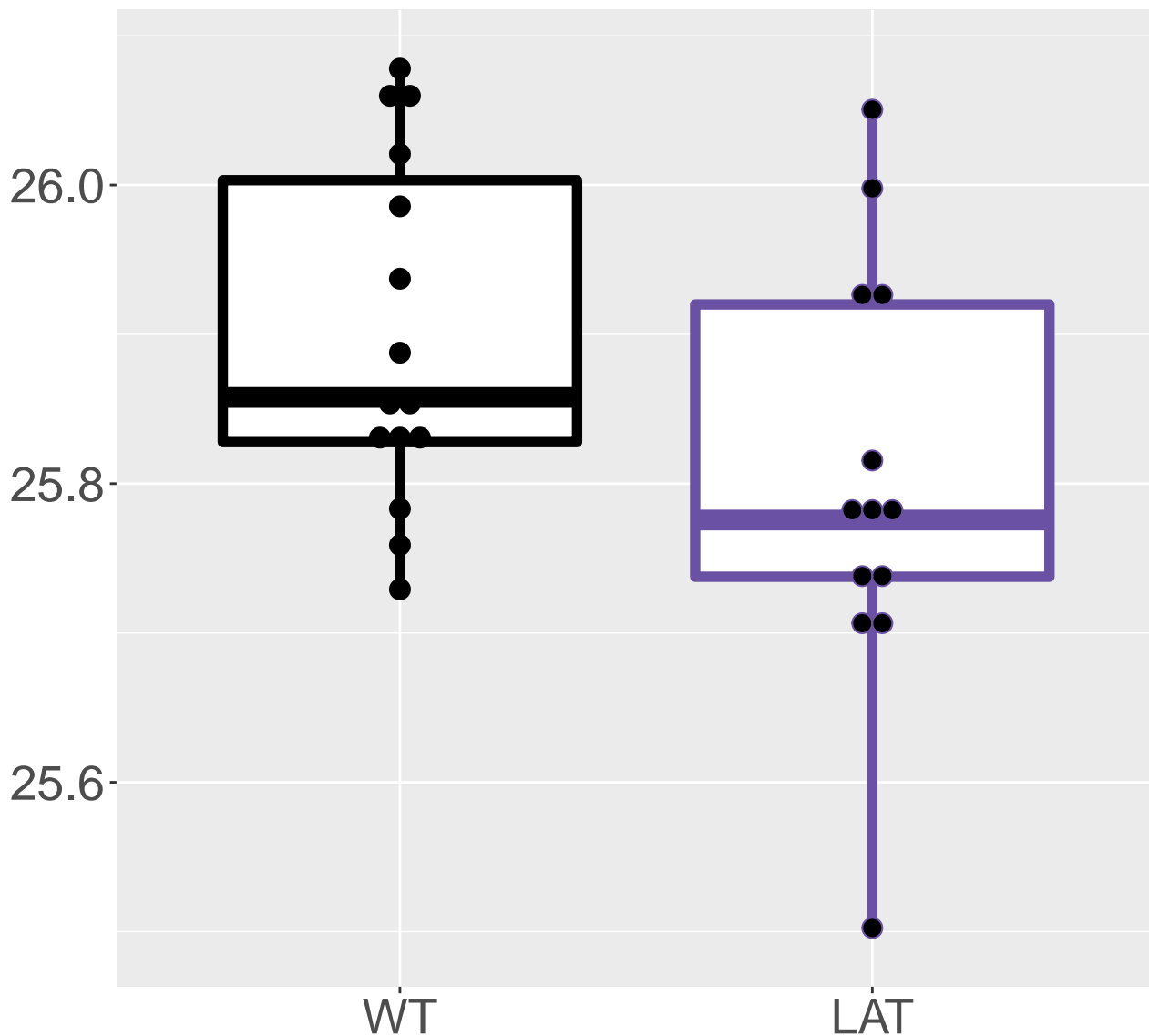
Q9D0R2_Threonine--tRNA ligase, .
FDR = 0.0073, FC = 0.34, sex**



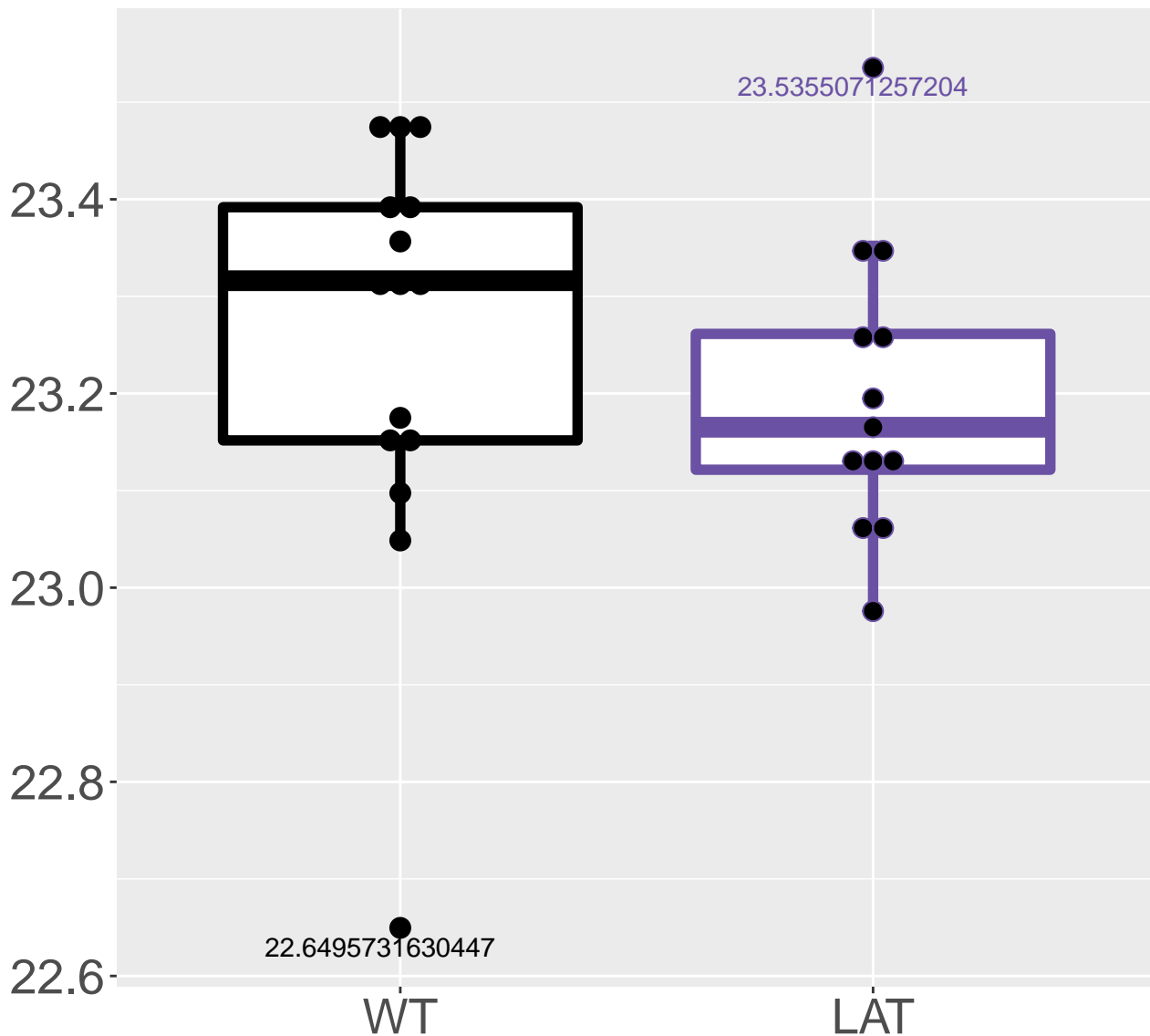
FDR = 0.0073, FC = -0.31



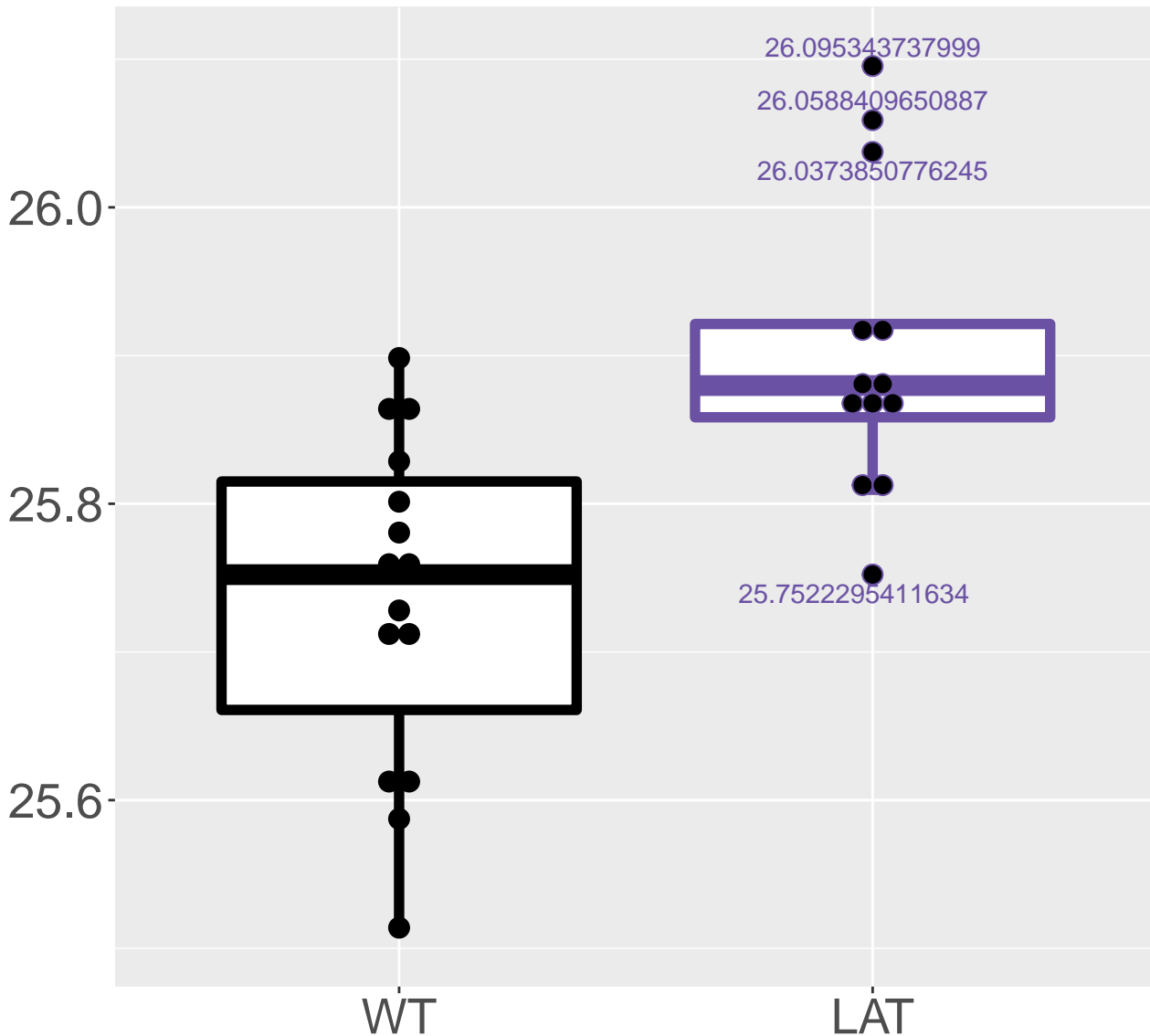
Q91WS0_CDGSH iron-sulfur domain.
FDR = 0.0073, FC = -0.28, sex**



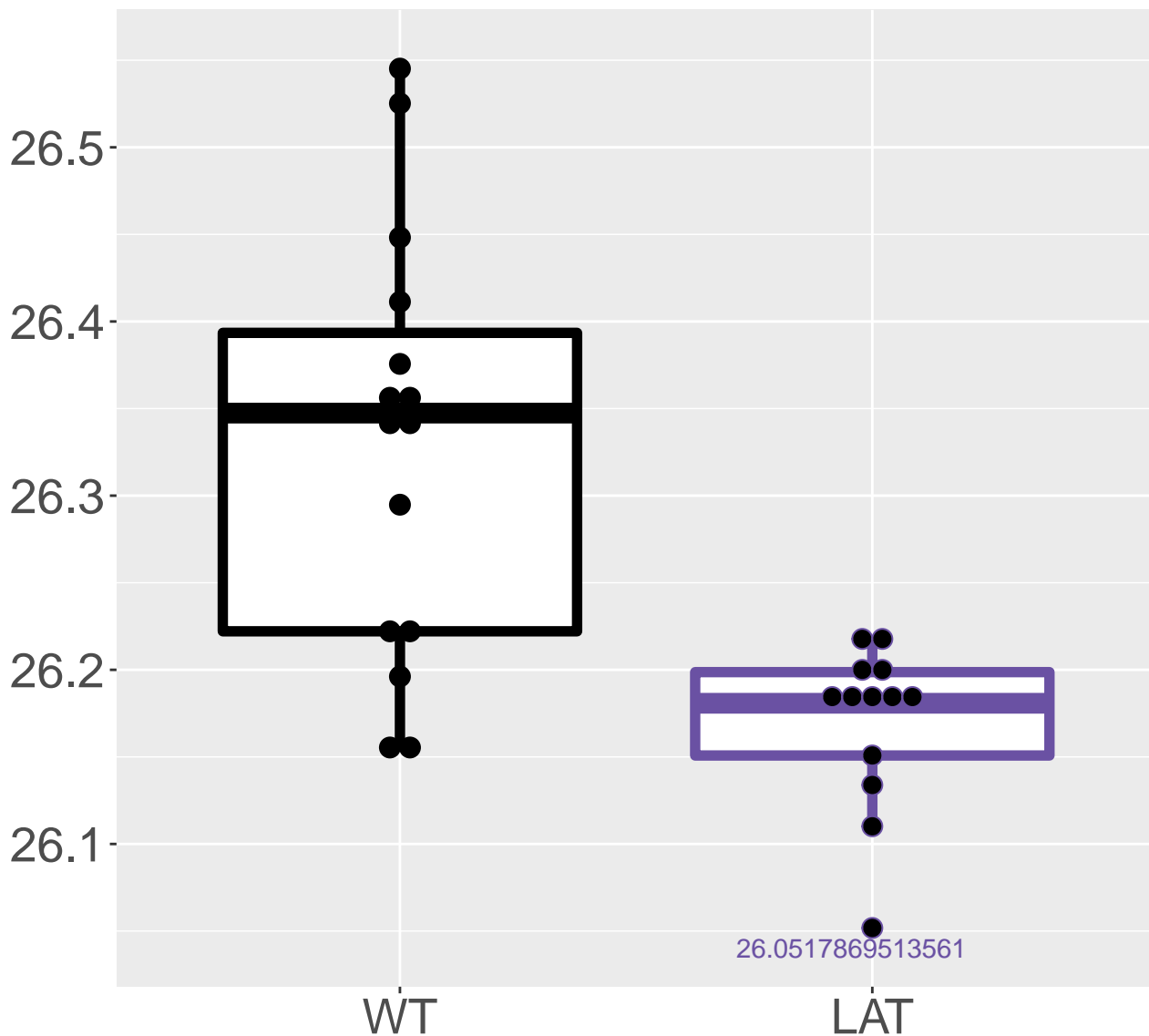
Q3ULJ0_Glycerol-3-phosphate deh.
FDR = 0.0073, FC = -0.26, sex**



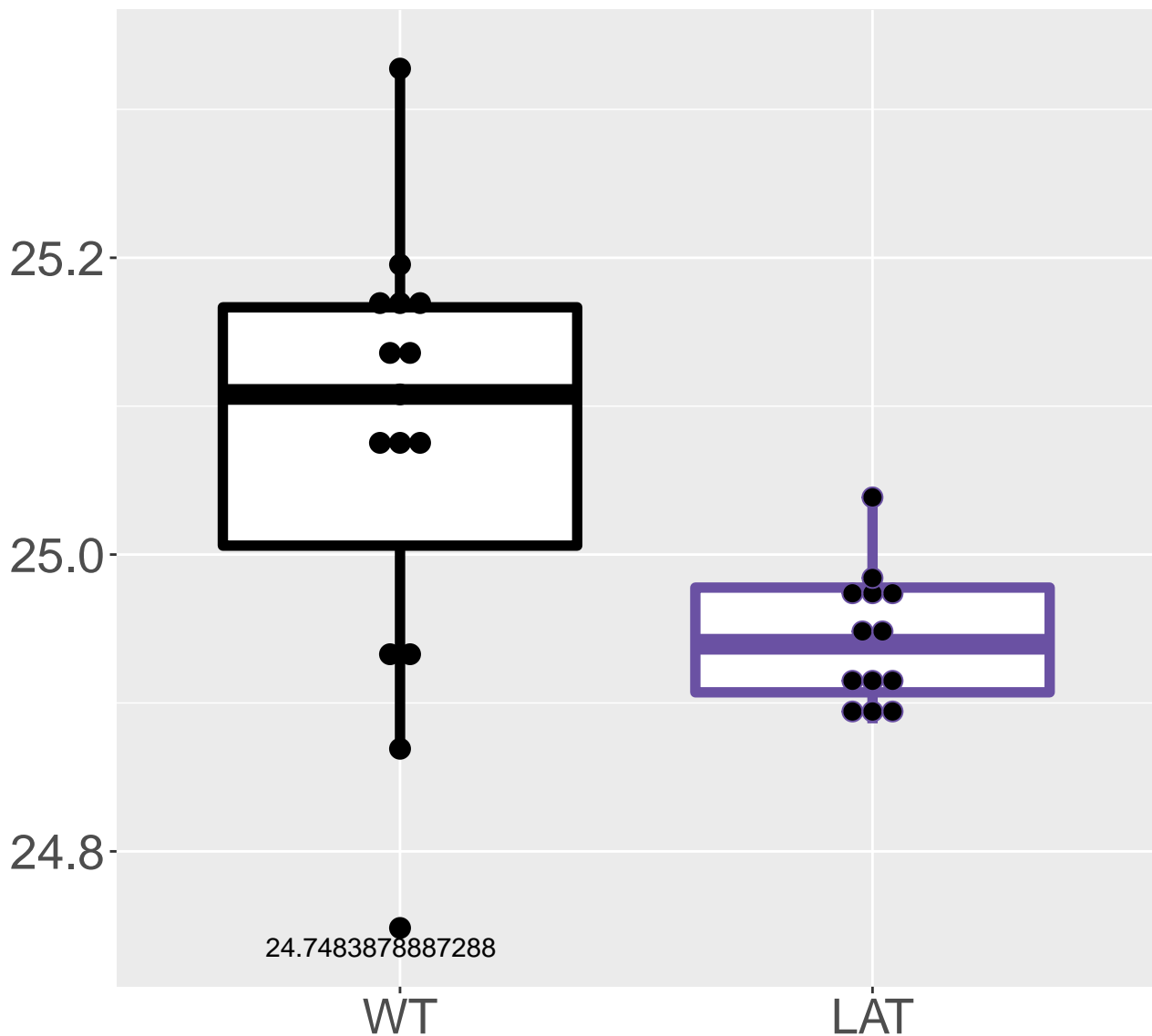
P16332_Methylmalonyl-CoA mutase.
FDR = 0.0073, FC = 0.26, sex*



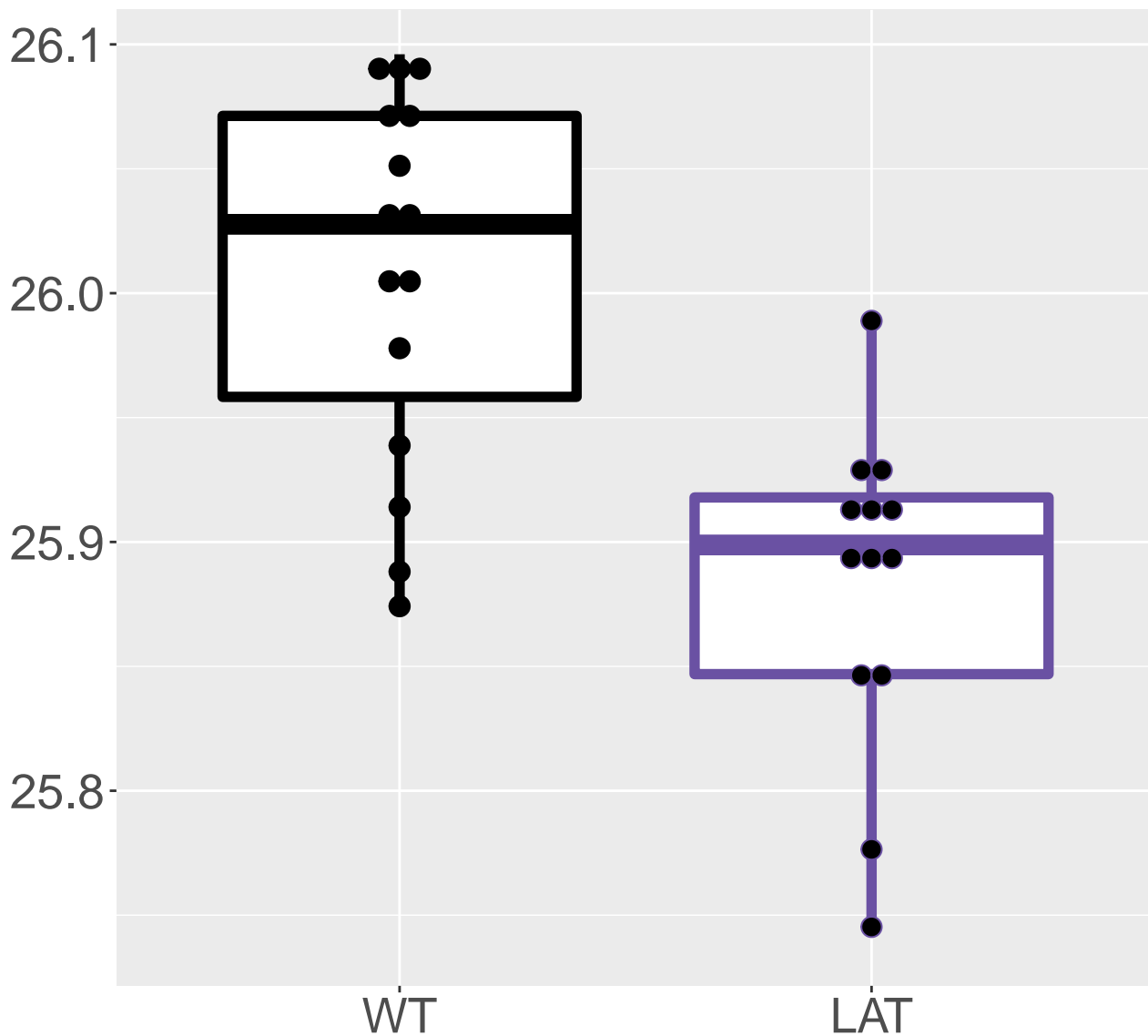
P47963_60S ribosomal protein L13
FDR = 0.0073, FC = -0.23



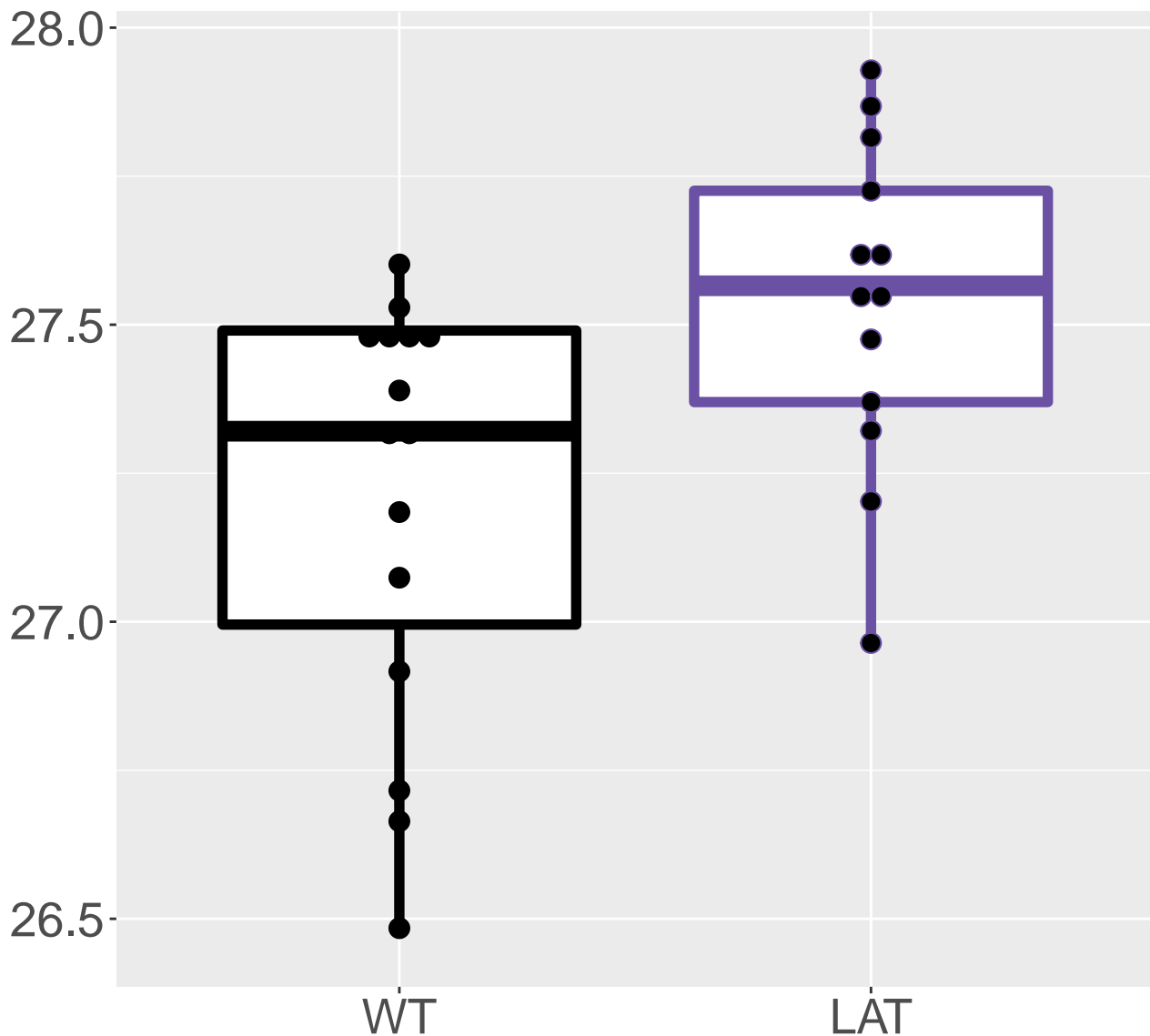
P46638_Ras-related protein Rab-.
FDR = 0.0073, FC = -0.22, sex*



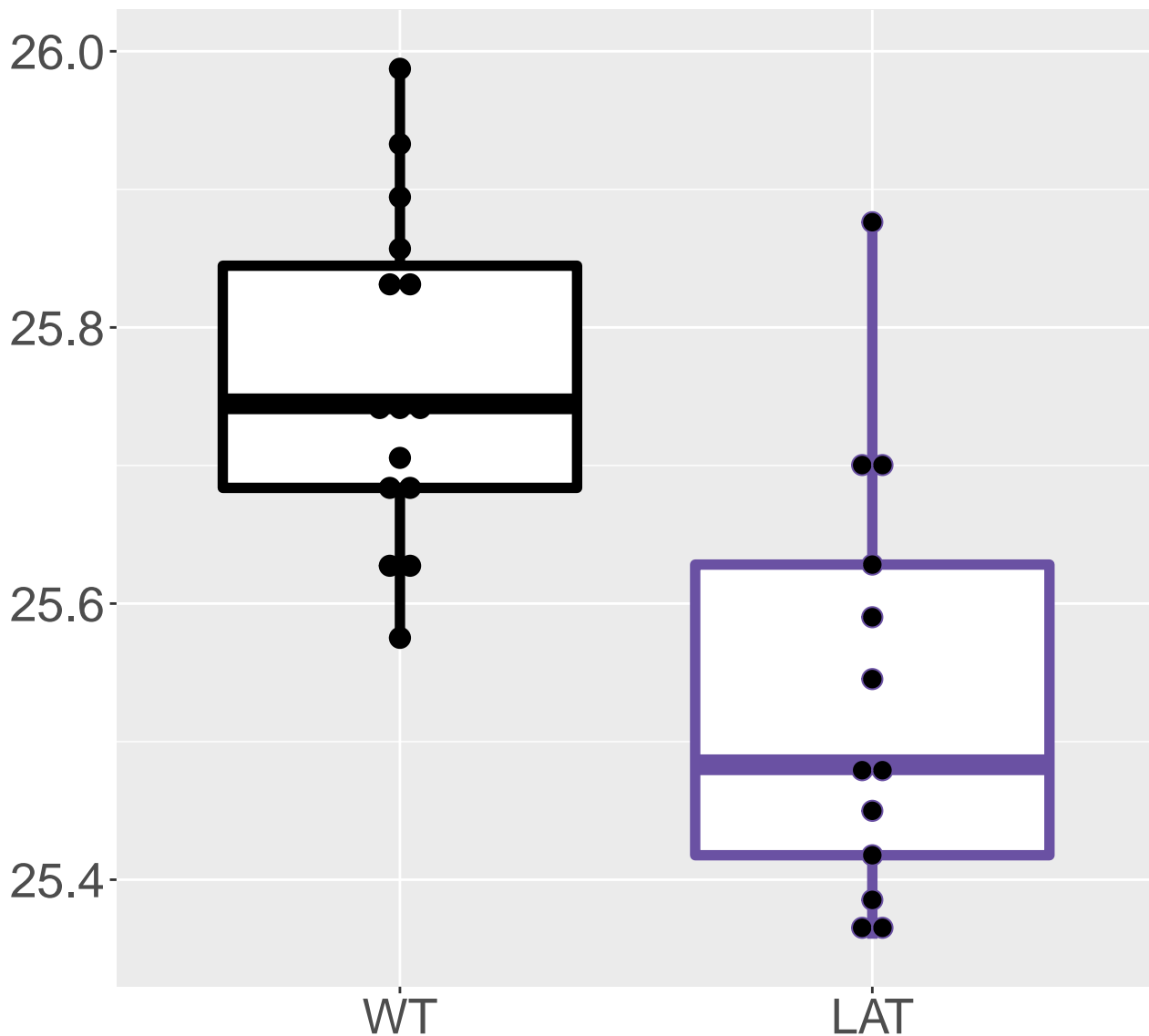
Q9QUM9_Proteasome subunit alpha.
FDR = 0.0073, FC = -0.2



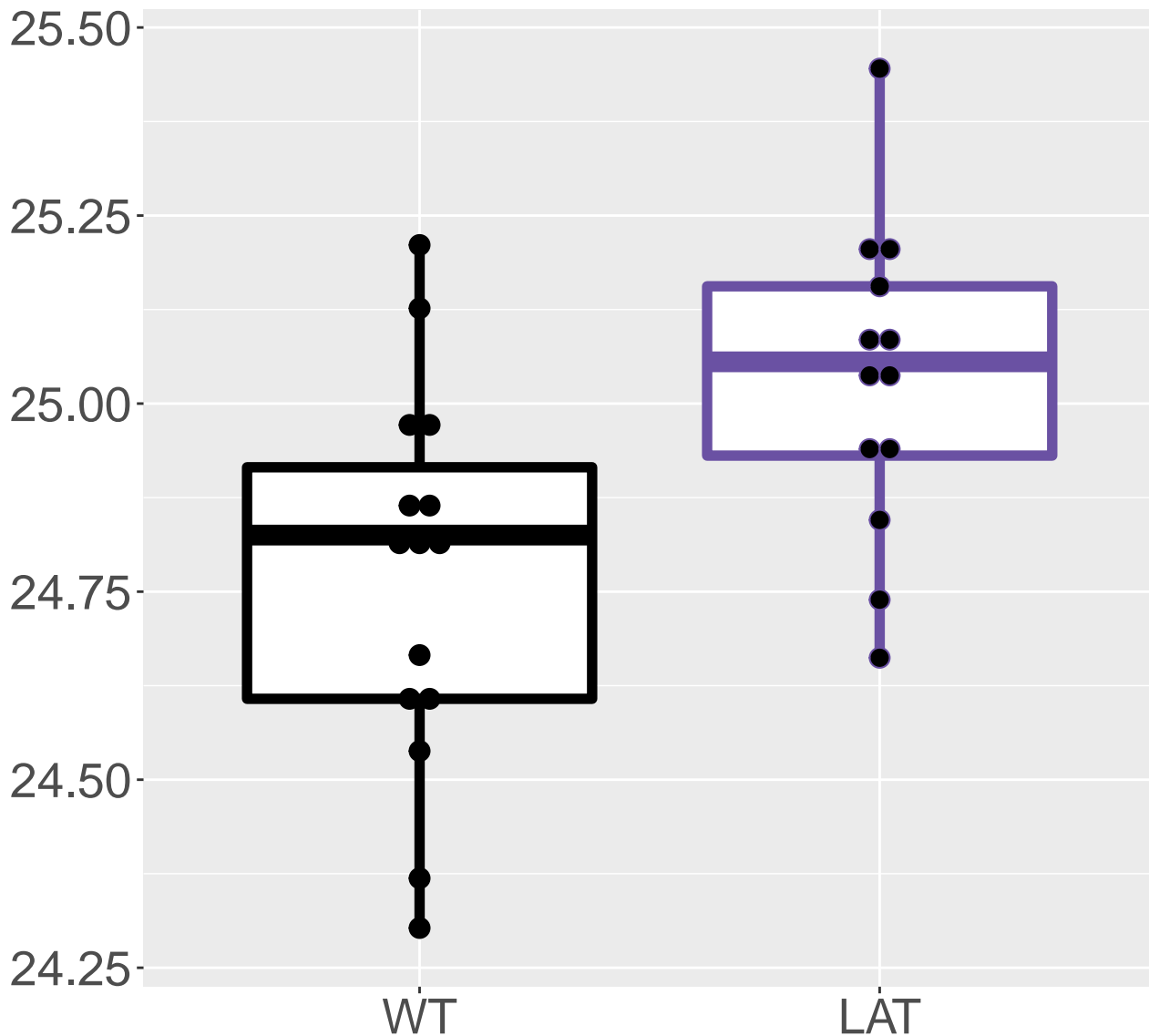
Q01853_Transitional endoplasmic.
FDR = 0.0076, FC = 0.68, sex**



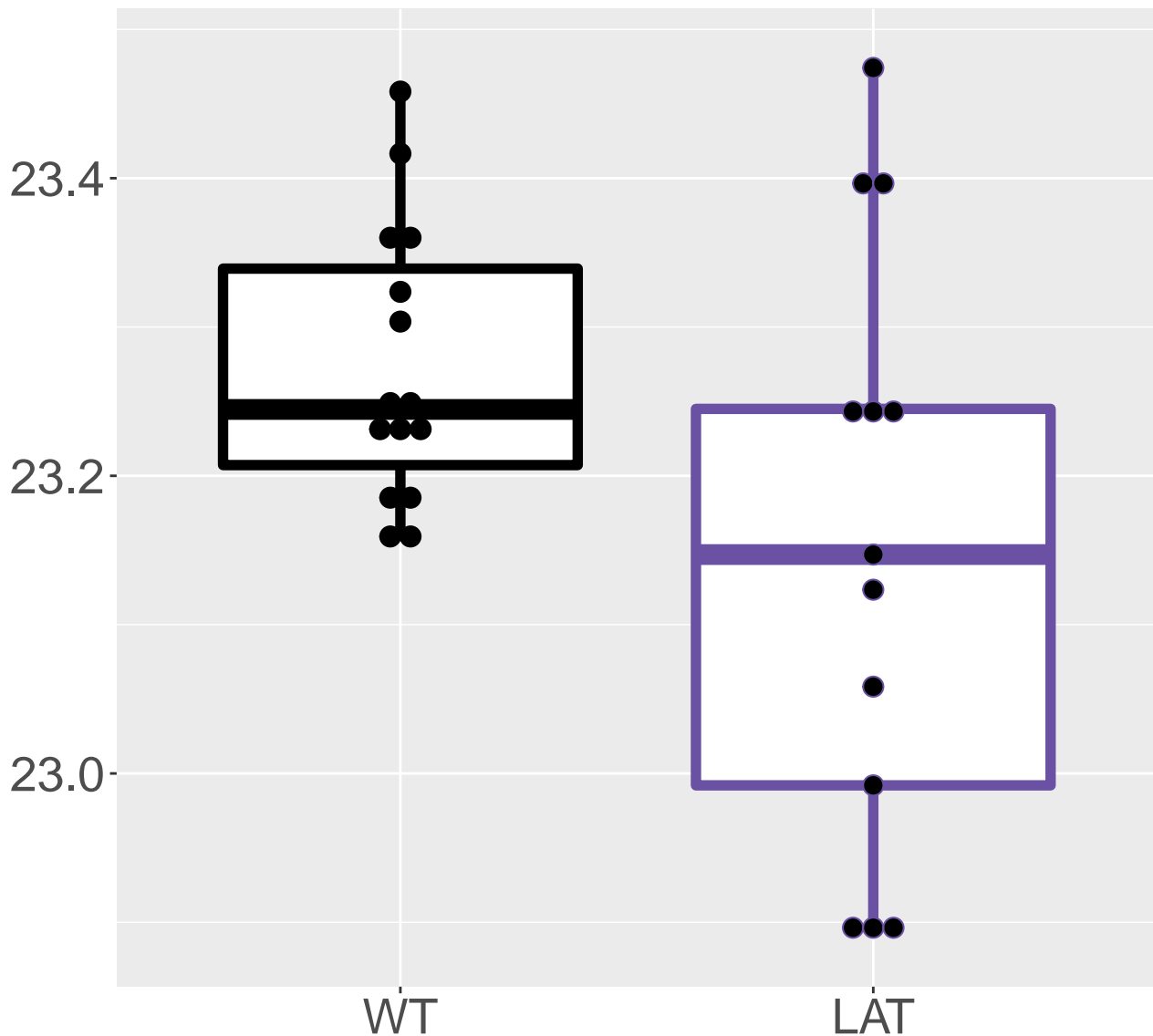
P15532_Nucleoside diphosphate k.
FDR = 0.0076, FC = -0.3



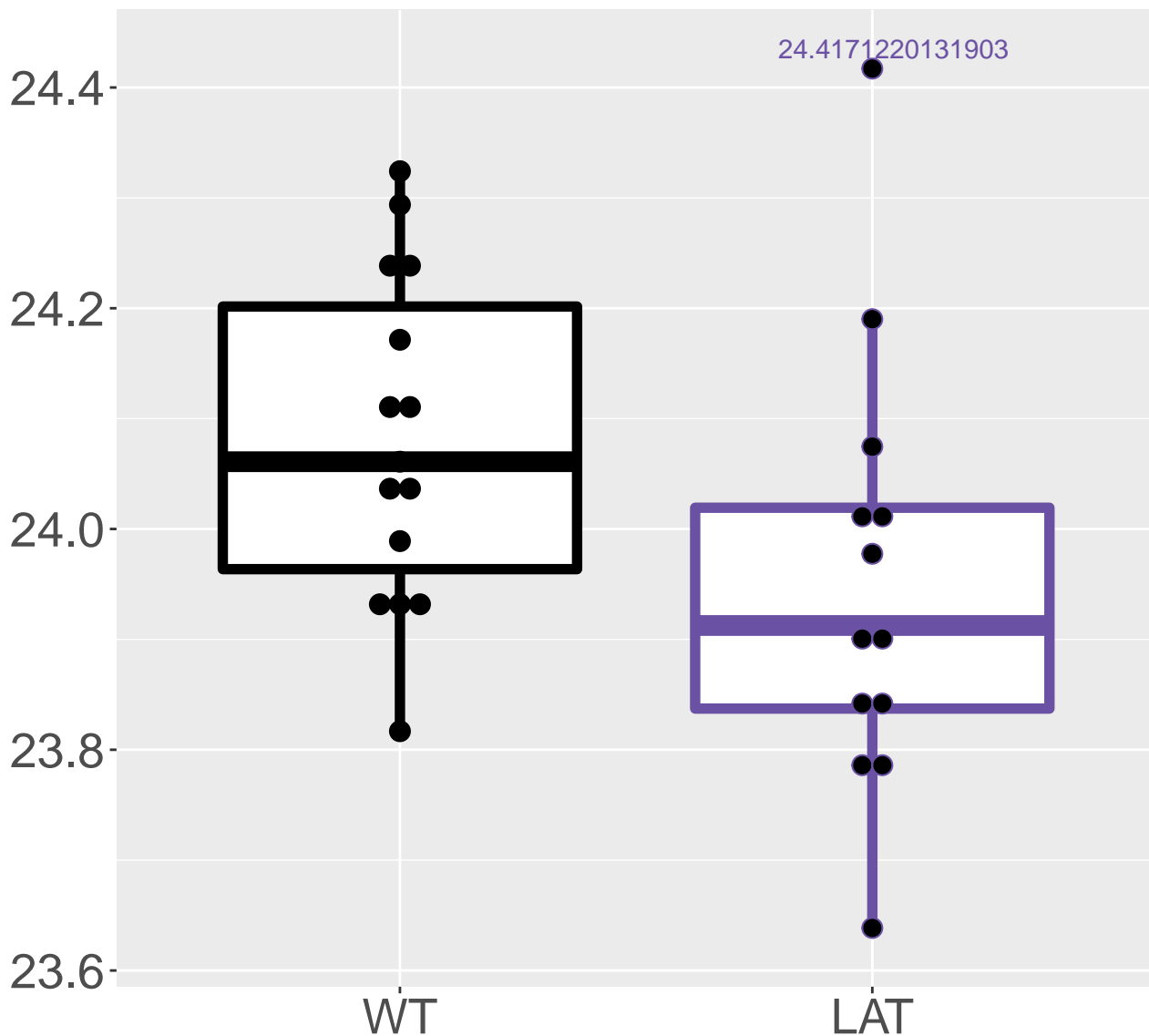
P00397_Cytochrome c oxidase sub.
FDR = 0.0078, FC = 0.52, sex*



Q6ZWY3_40S ribosomal protein S2.
FDR = 0.0079, FC = -0.25, sex**

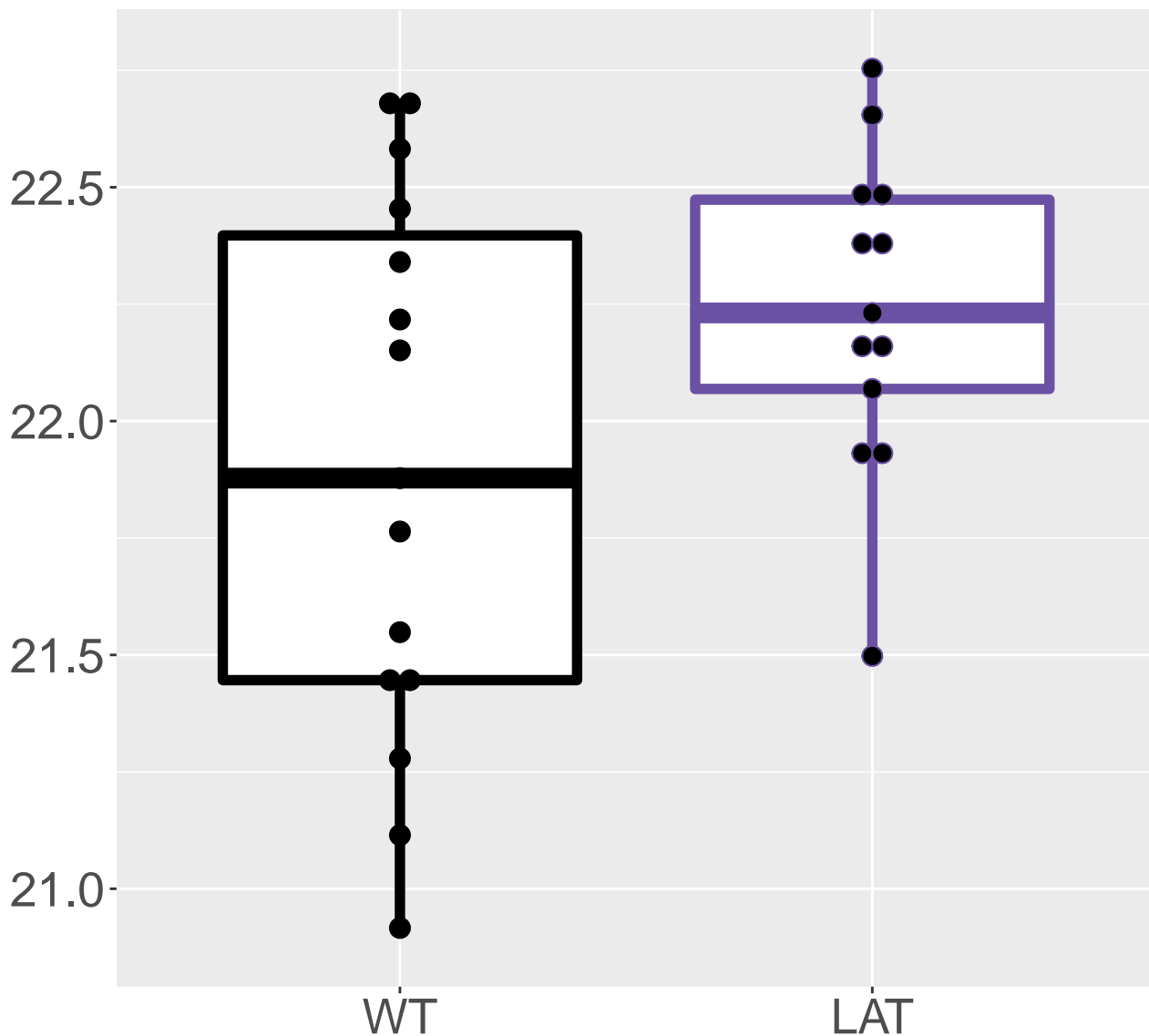


Q9CRB9_MICOS complex subunit Mi.
FDR = 0.0079, FC = -0.32

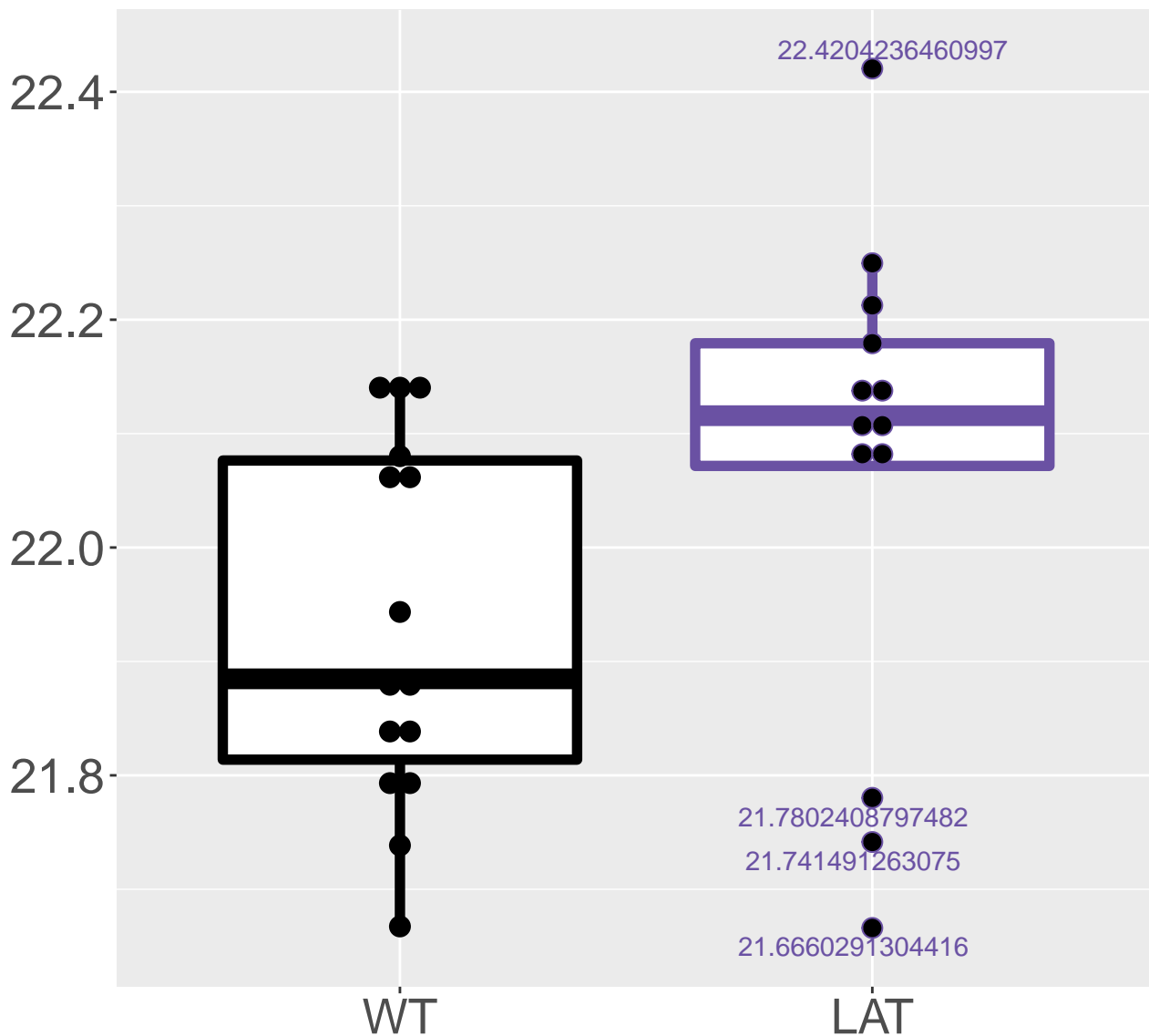


Q8R0W0_Epiplakin

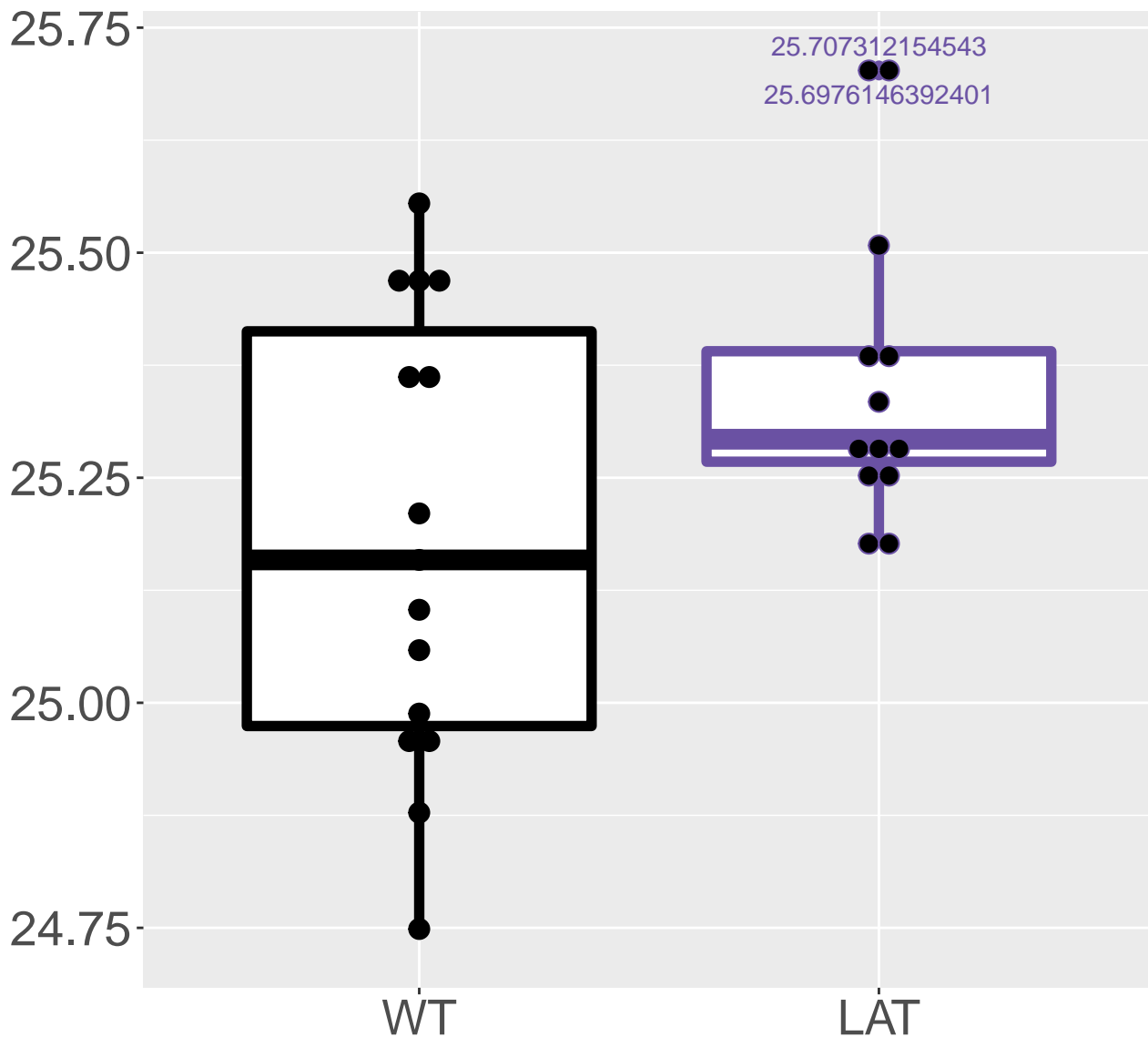
FDR = 0.0083, FC = 0.72, sex***



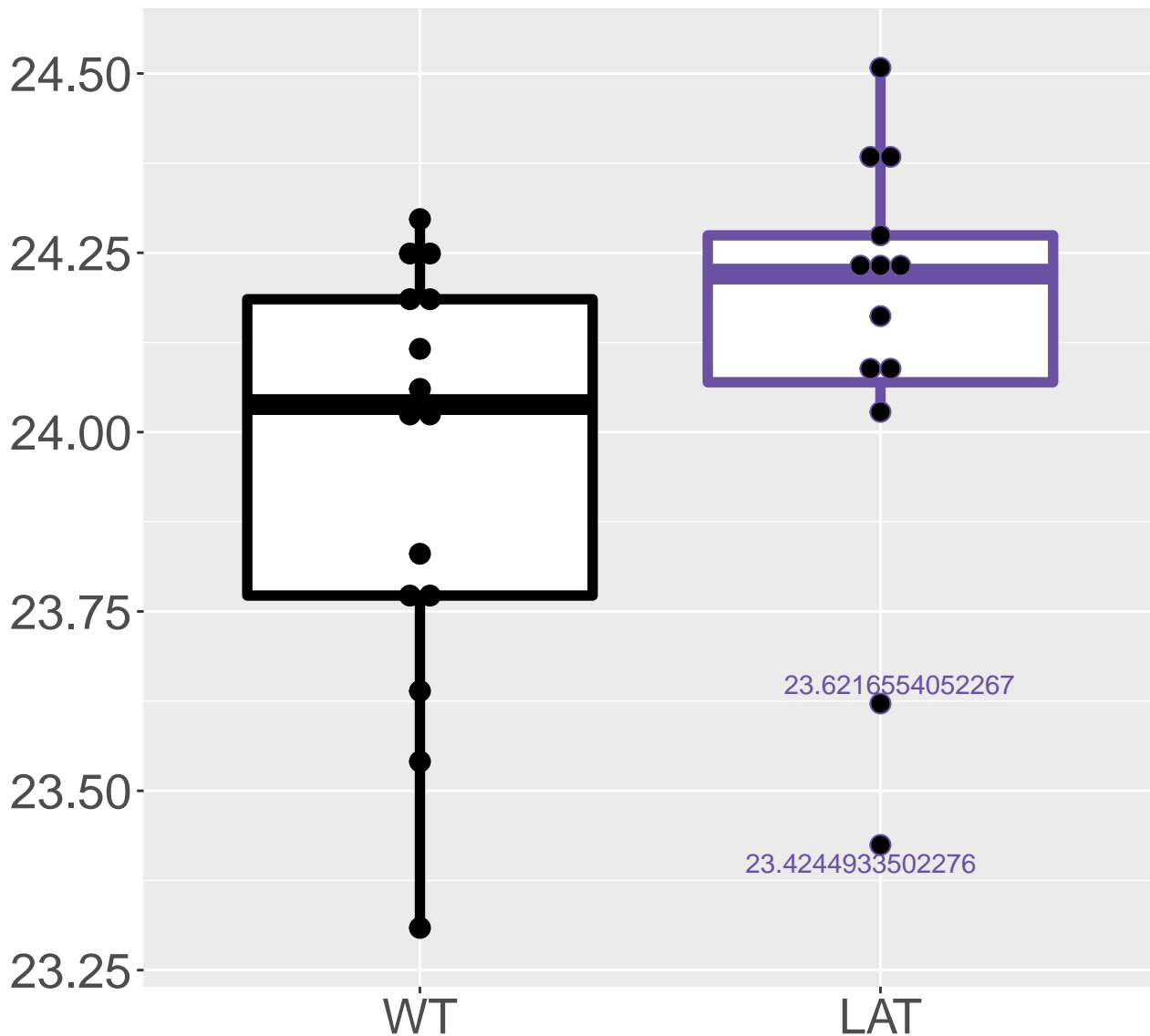
FDR = 0.0084, FC = 0.35, sex*



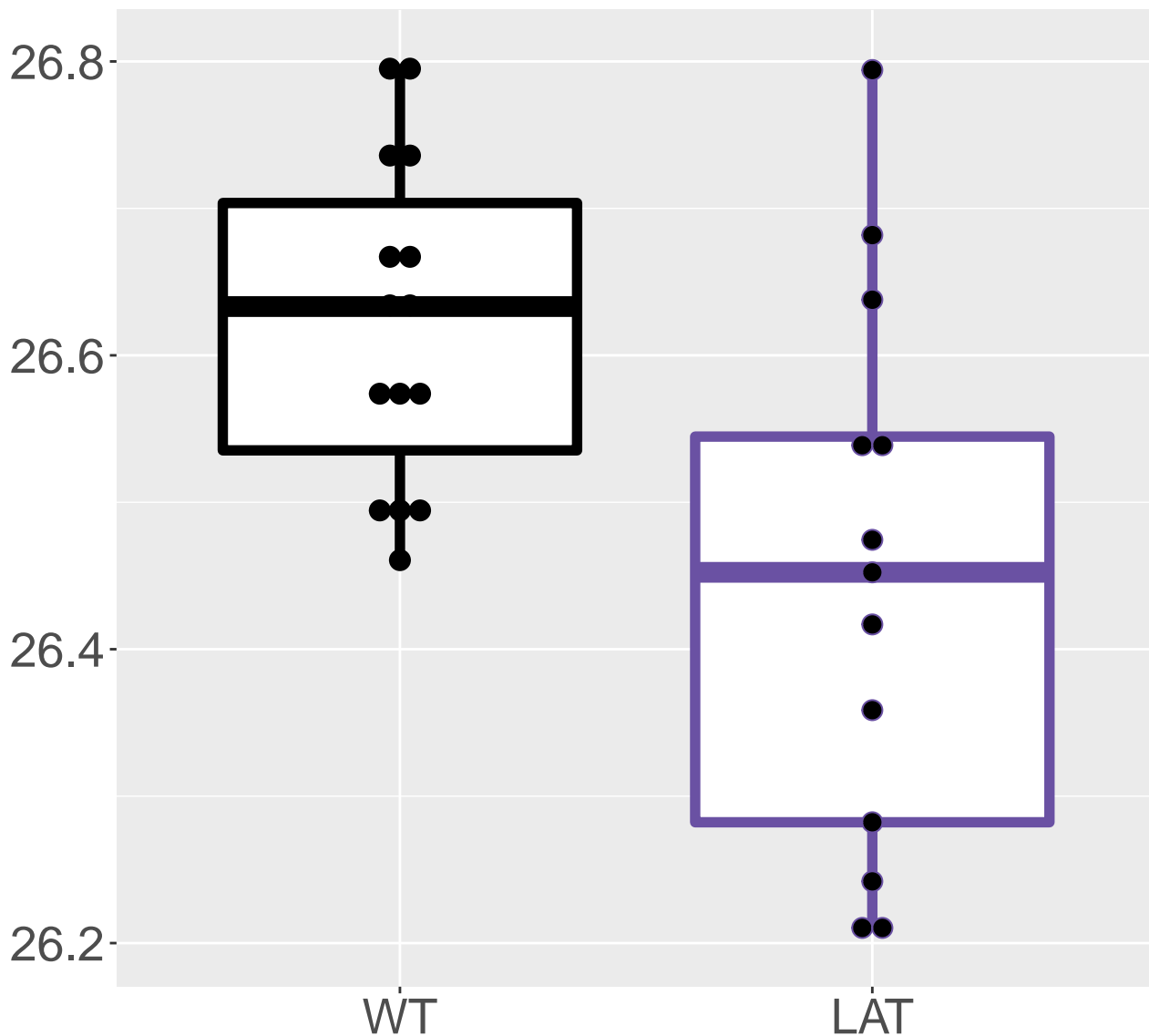
Q8CFX1_GDH/6PGL endoplasmic bif.
FDR = 0.0084, FC = 0.31, sex***



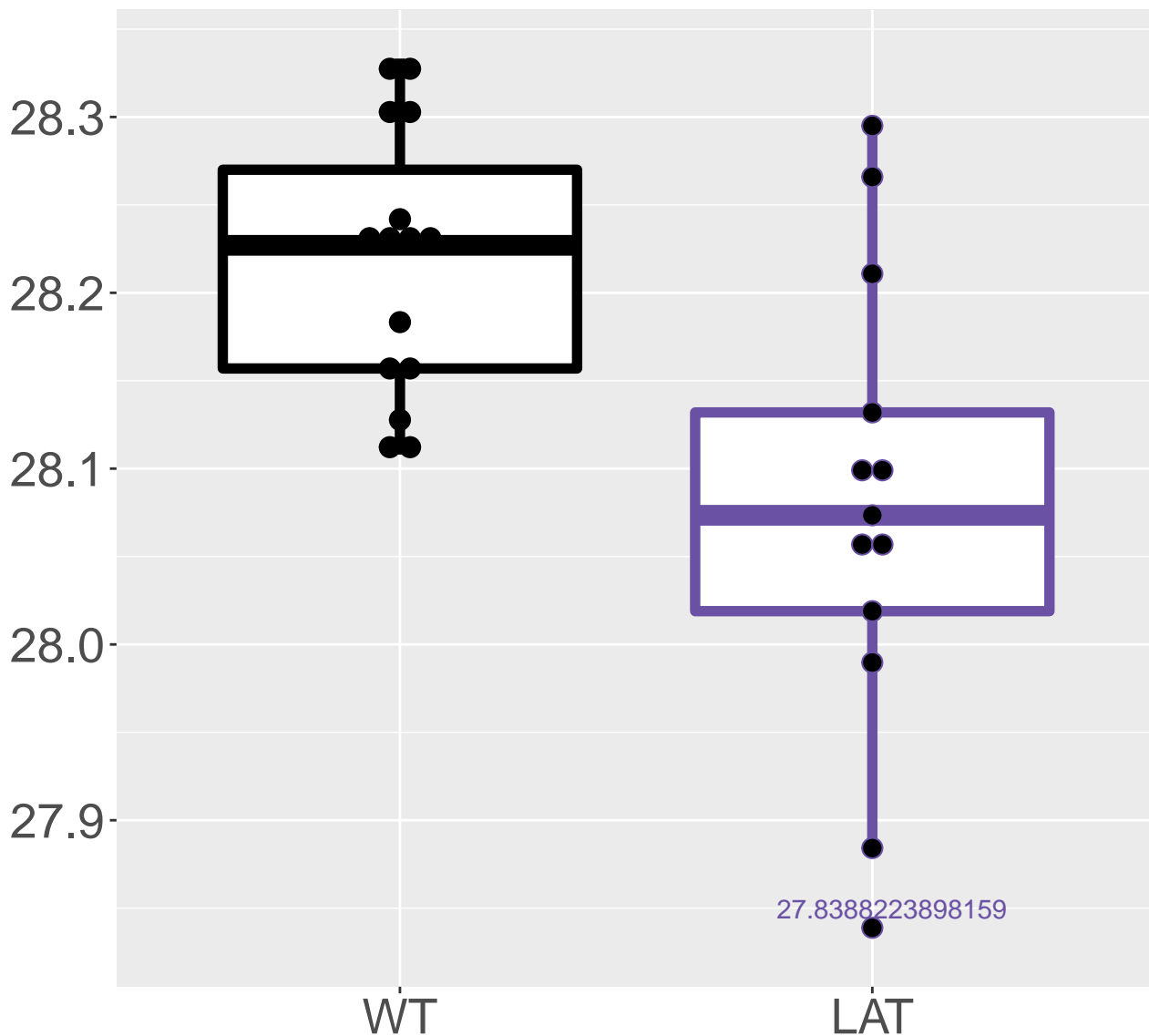
O55029_Coatomer subunit beta'
FDR = 0.0085, FC = 0.53, sex**



P62897_Cytochrome c, somatic
FDR = 0.0085, FC = -0.32, sex*

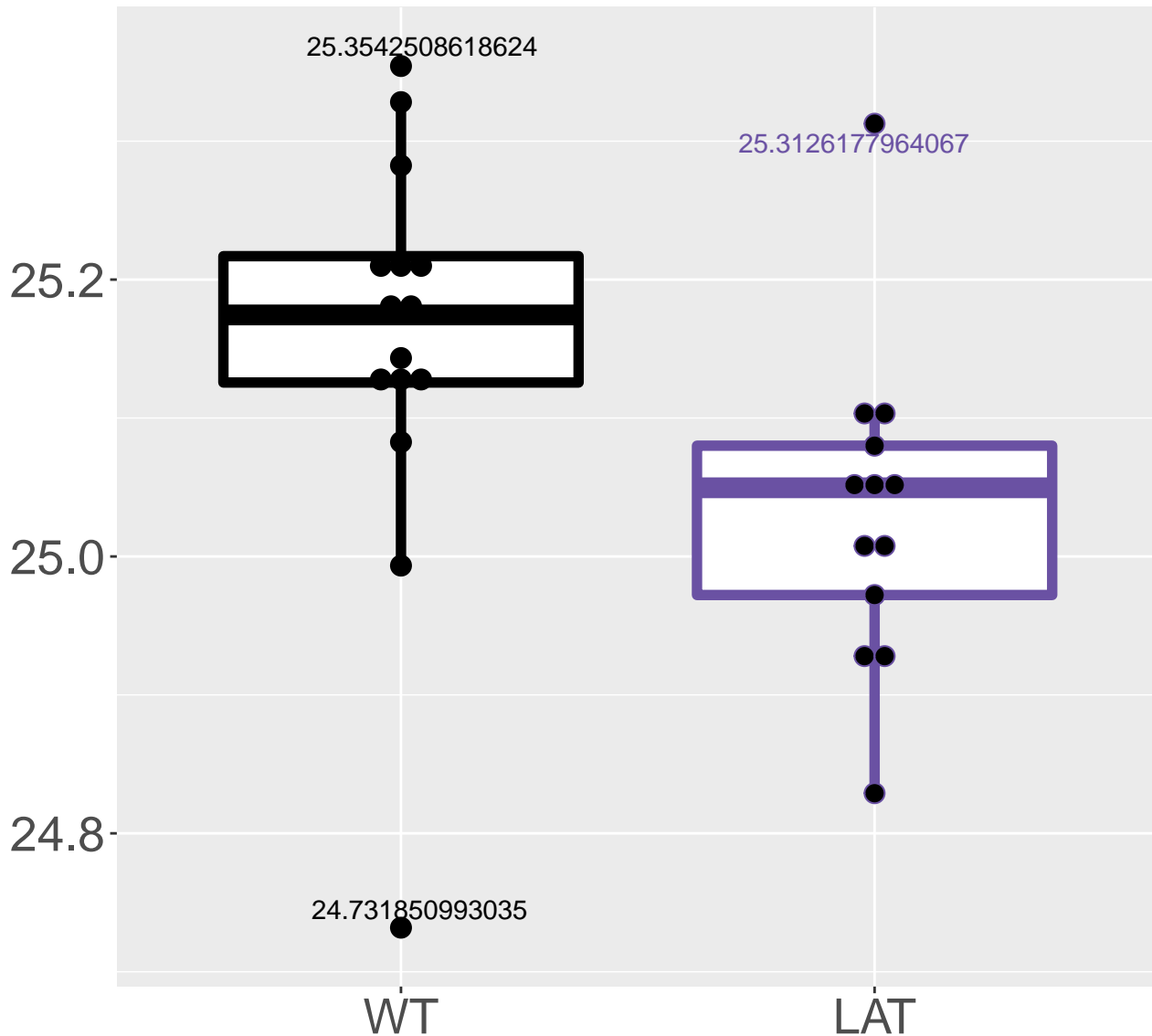


P62983_Ubiquitin-40S ribosomal .
FDR = 0.0085, FC = -0.26

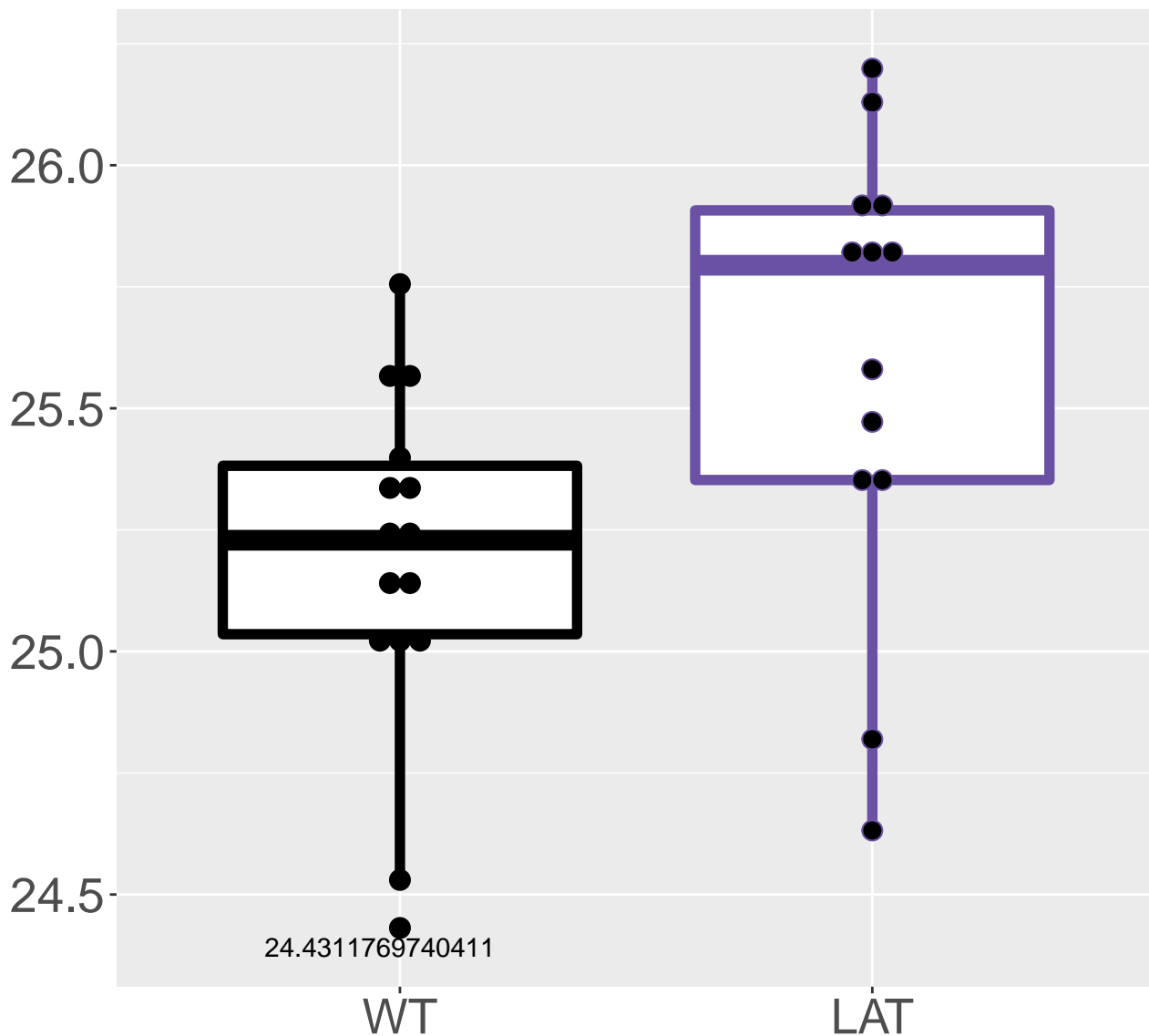


P62855_40S ribosomal protein S26

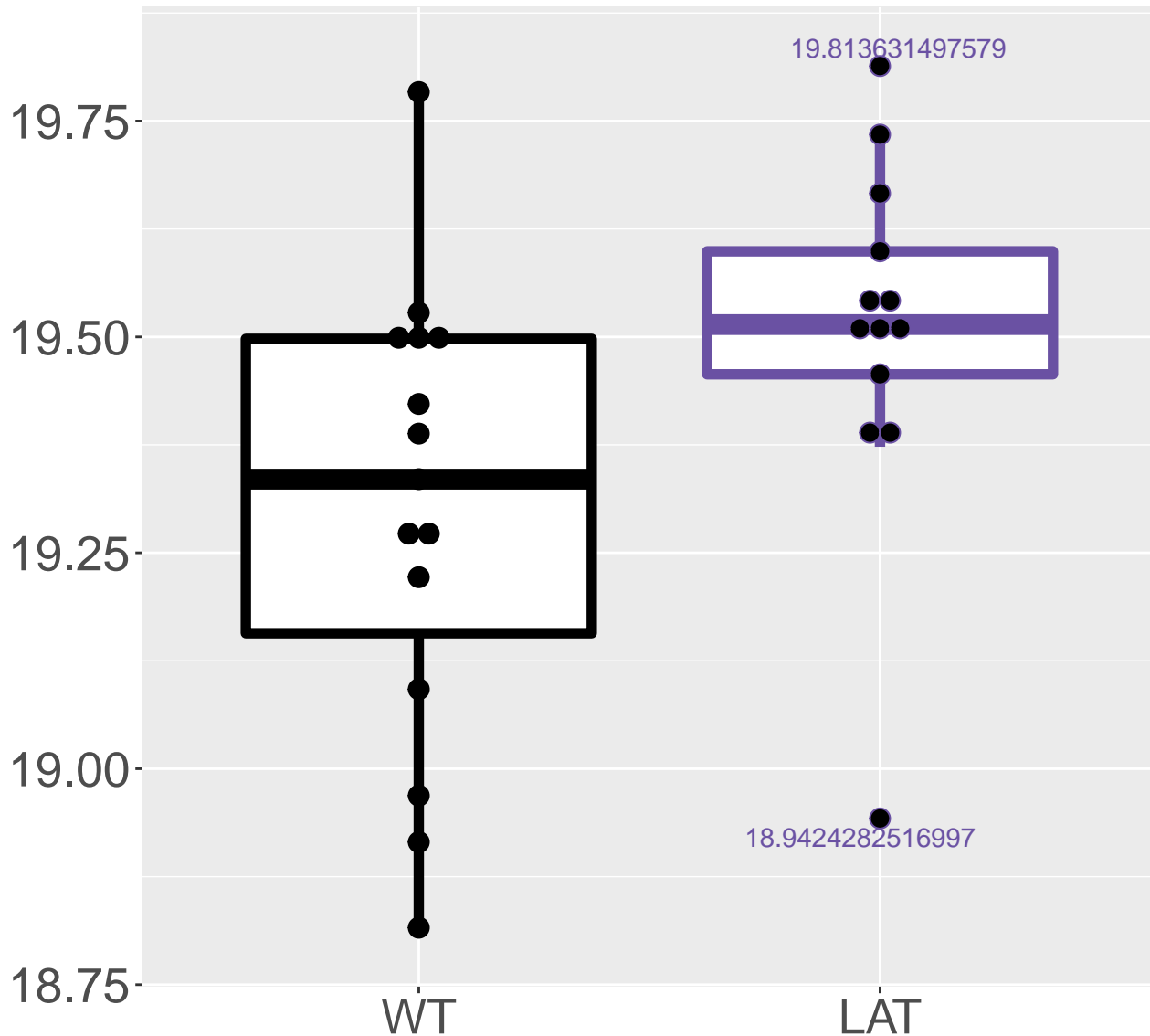
FDR = 0.0085, FC = -0.25



Q8BHN3_Neutral alpha-glucosidas.
FDR = 0.0089, FC = 0.84

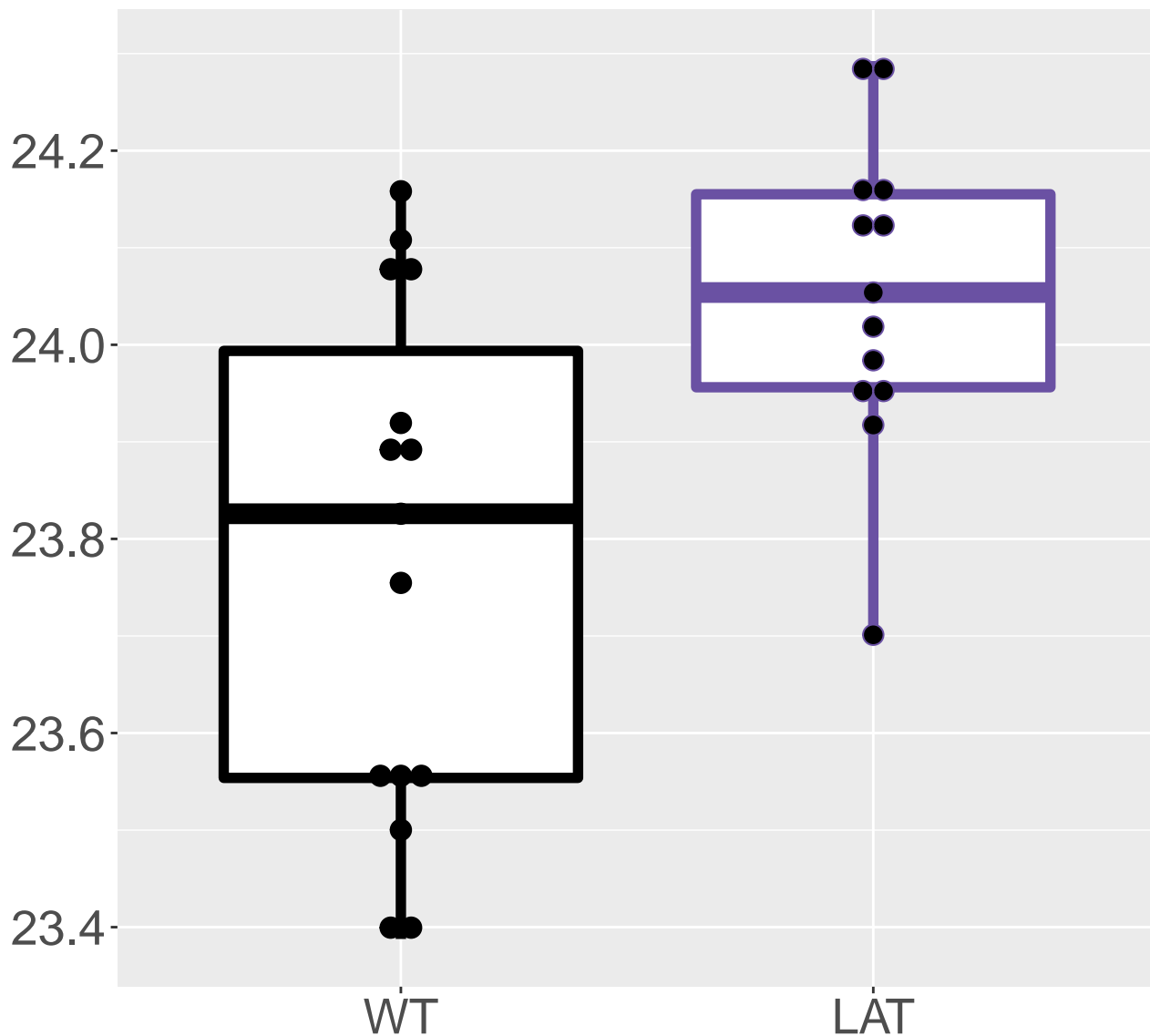


Q8R5H1_Ubiquitin carboxyl-termi.
FDR = 0.0089, FC = 0.49, sex*



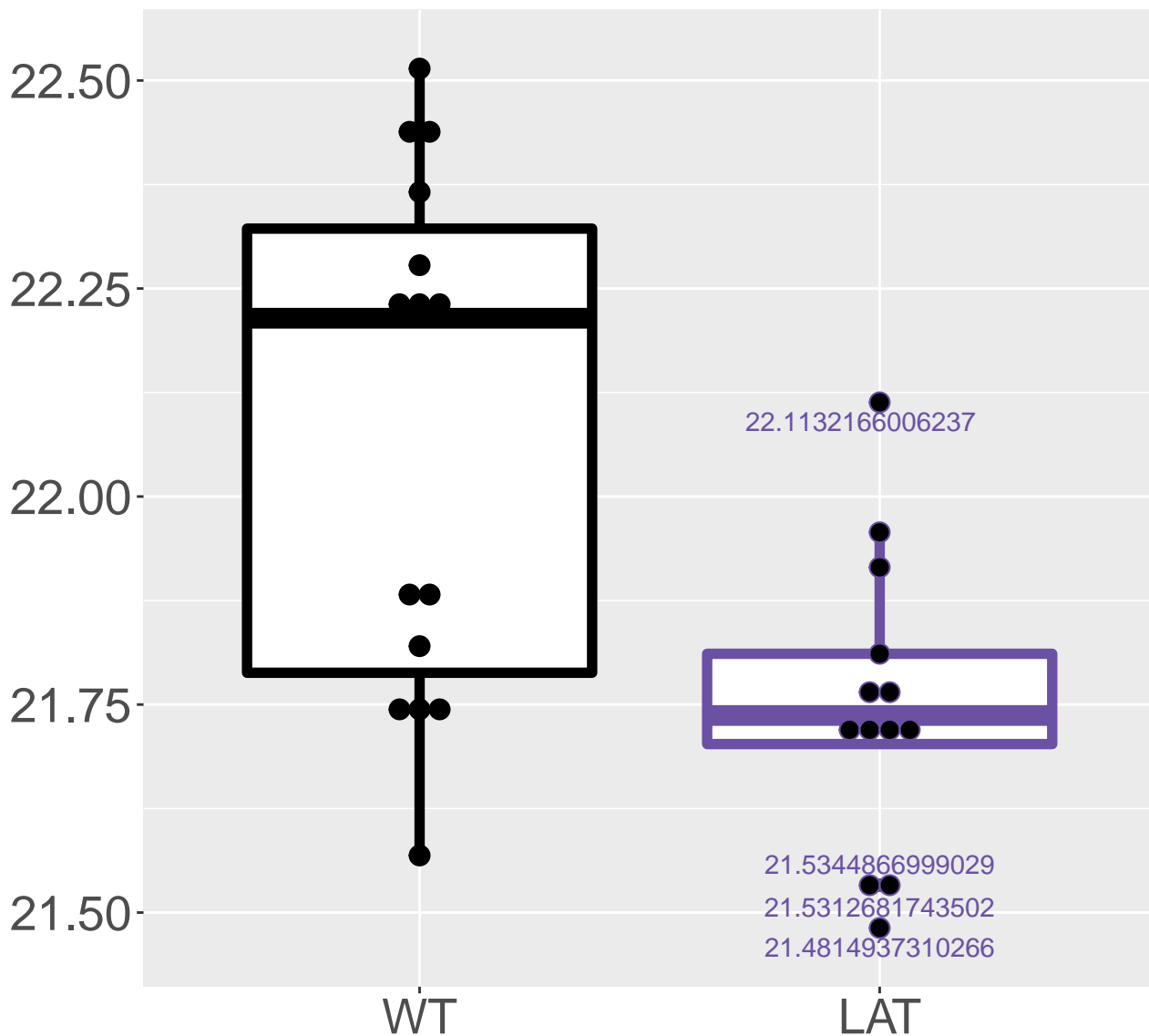
Q9CQN6_Transmembrane protein 14C

FDR = 0.0089, FC = 0.41, sex***

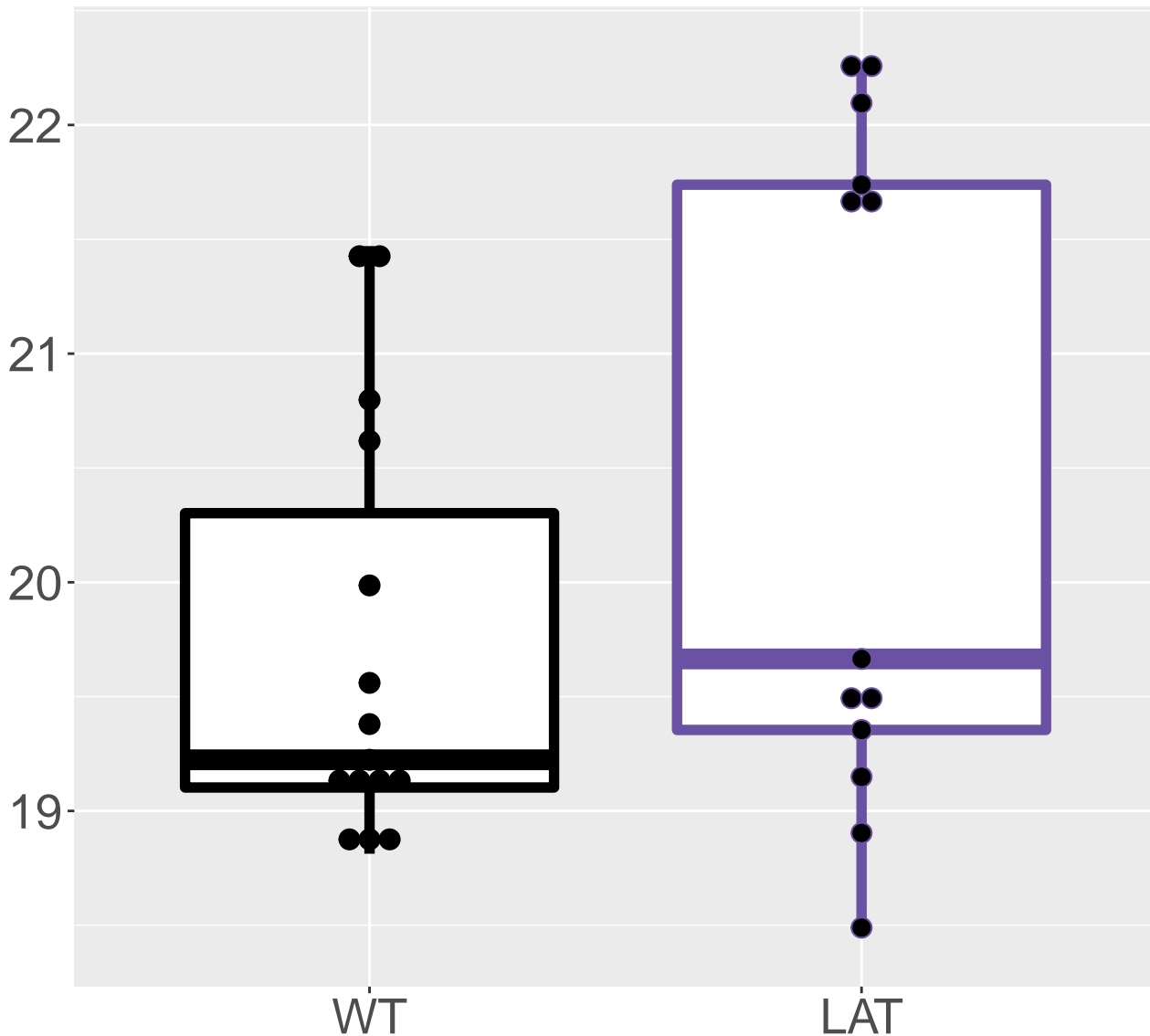


FDR = 0.0094, FC = -0.53, sex**

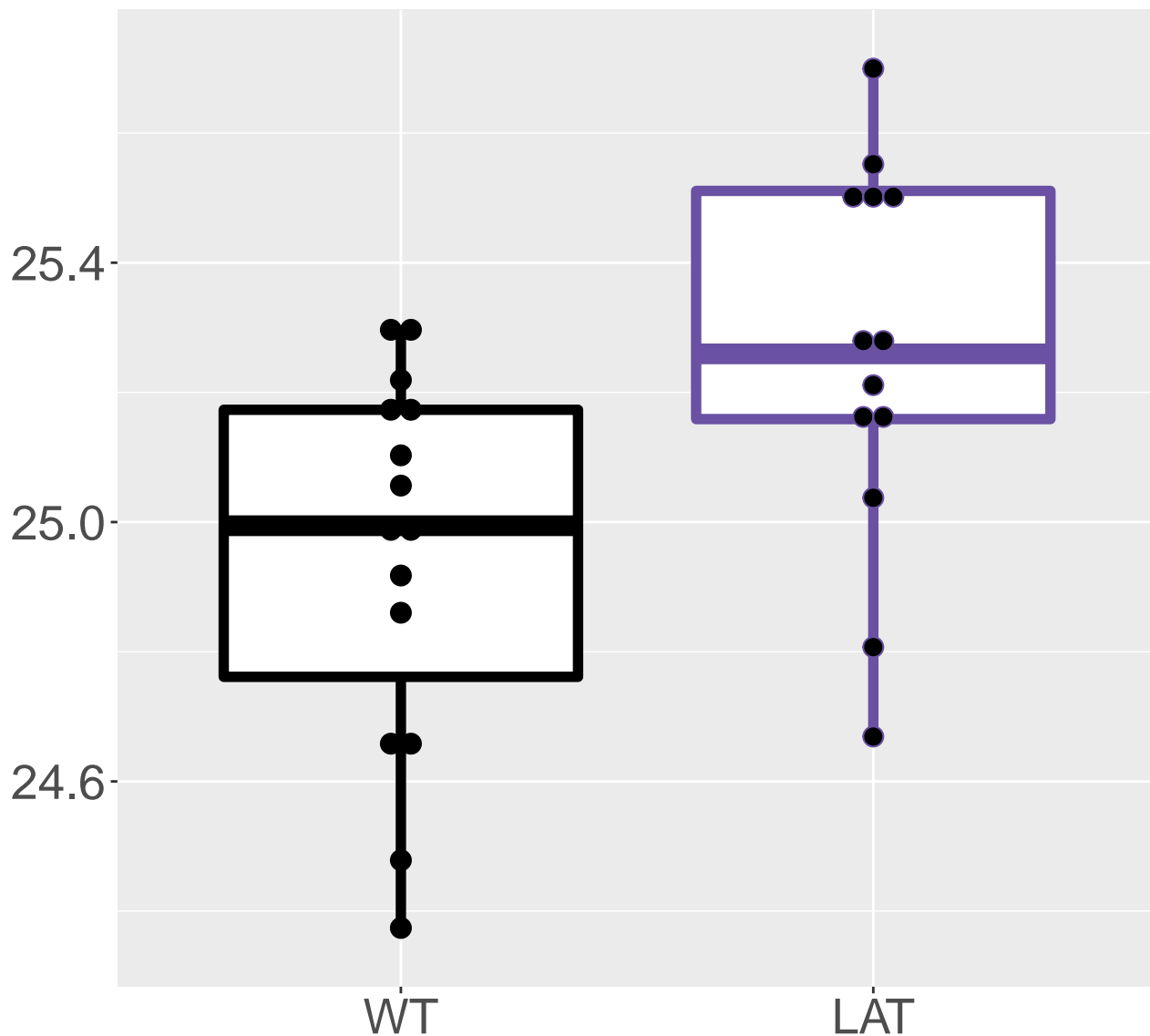
FDR = 0.0094, FC = -0.53, sex**



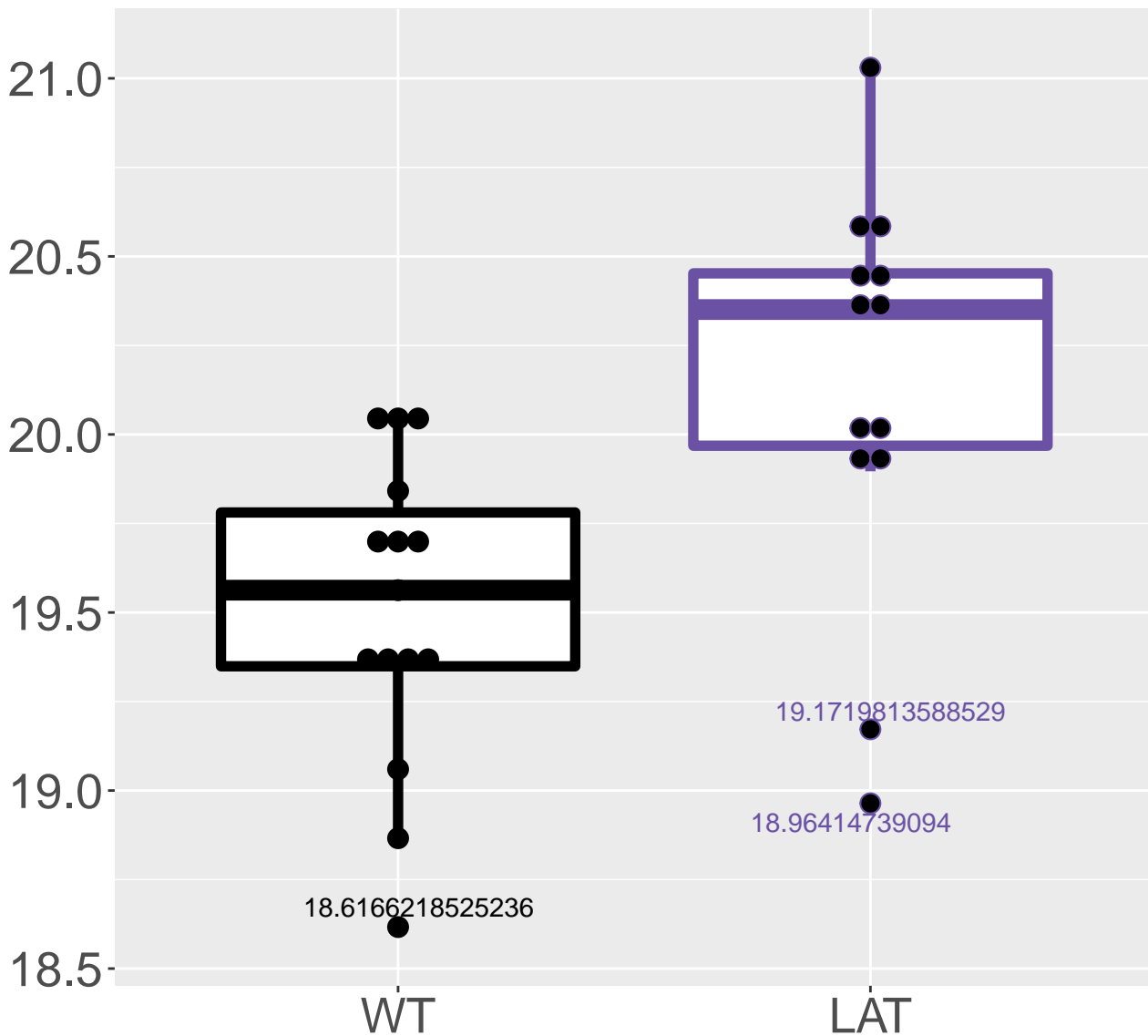
P11438_Lysosome-associated memb.
FDR = 0.0098, FC = 2.1, sex*



Q8CGK3_Lon protease homolog, mi.
FDR = 0.0099, FC = 0.6, sex*

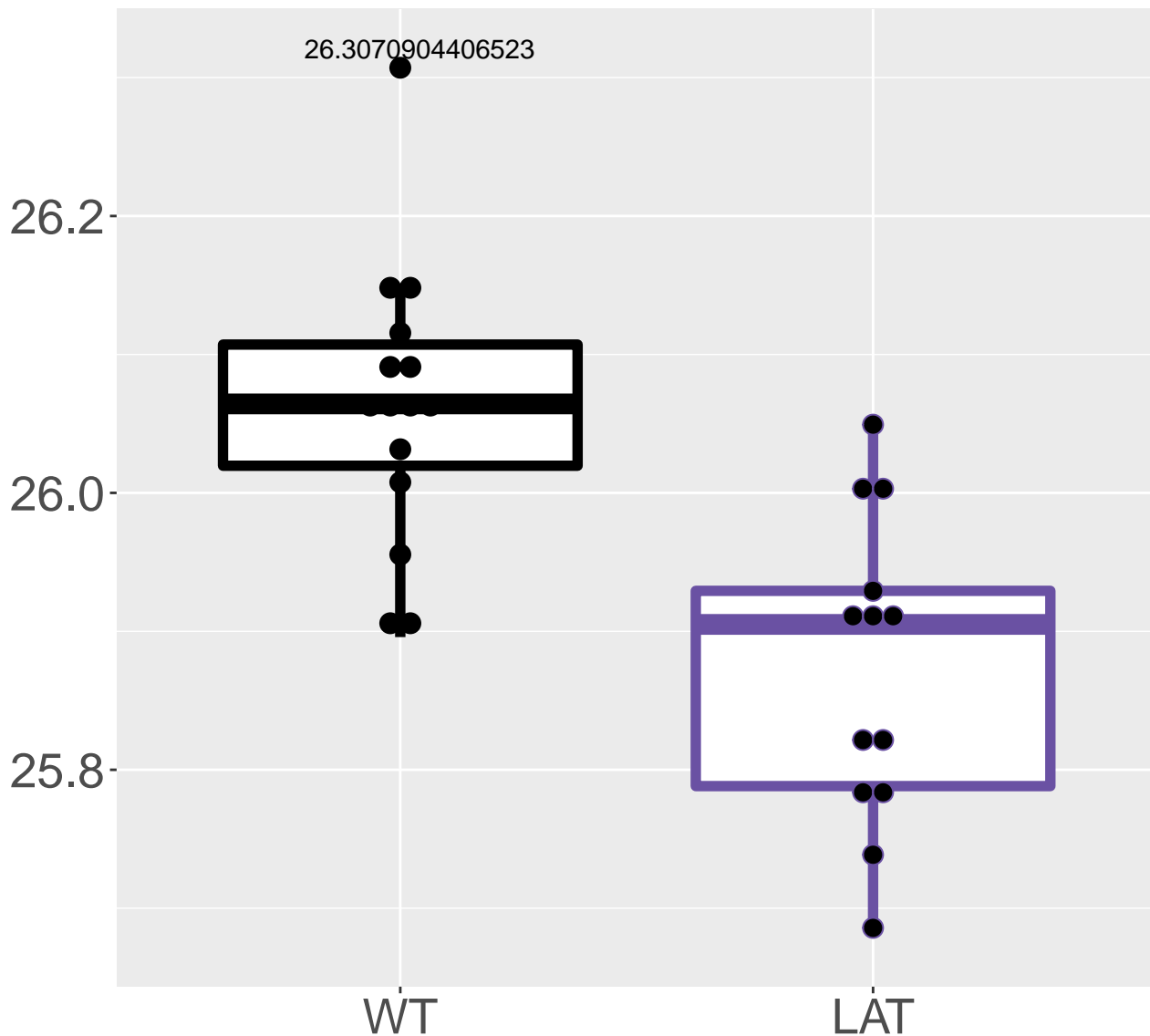


P09055_Integrin beta-1
FDR = 0.01, FC = 1

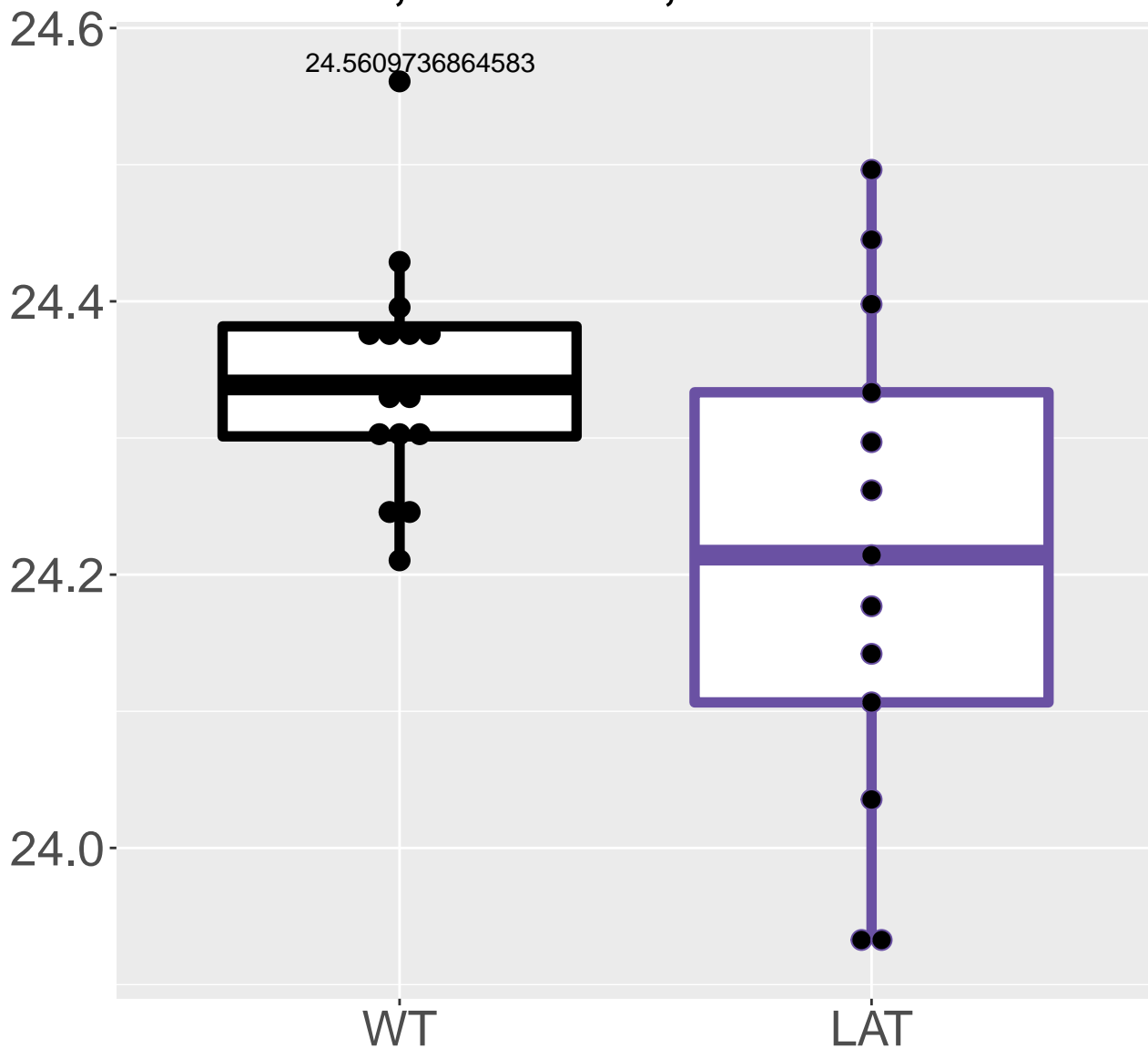


FDR = 0.011, FC = -0.28

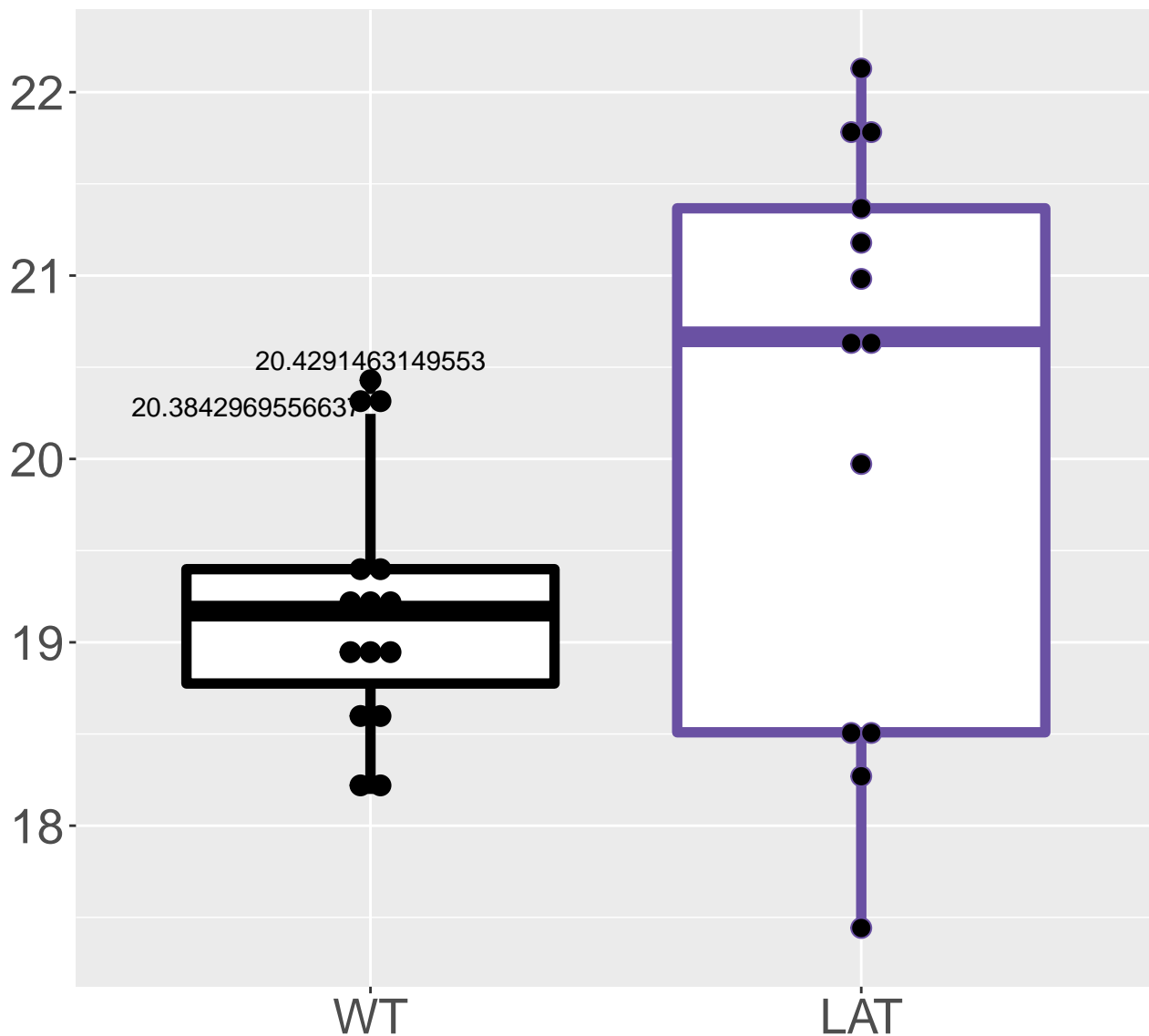
FDR = 0.011, FC = -0.28



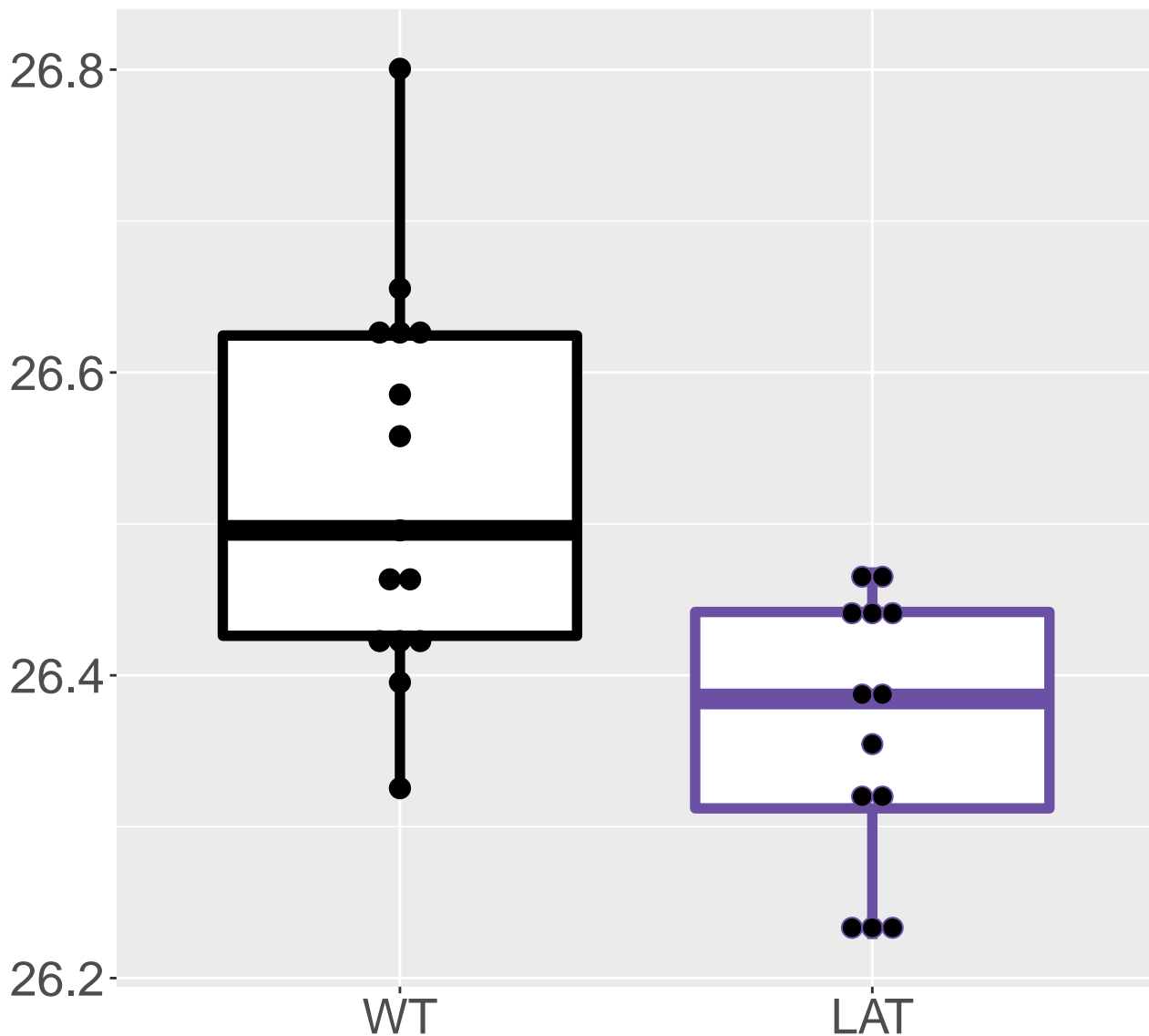
Q9CPP6_NADH dehydrogenase [ubiq.
FDR = 0.011, FC = -0.28, sex**



Q76LS9_Ubiquitin carboxyl-termi.
FDR = 0.011, FC = 2.1

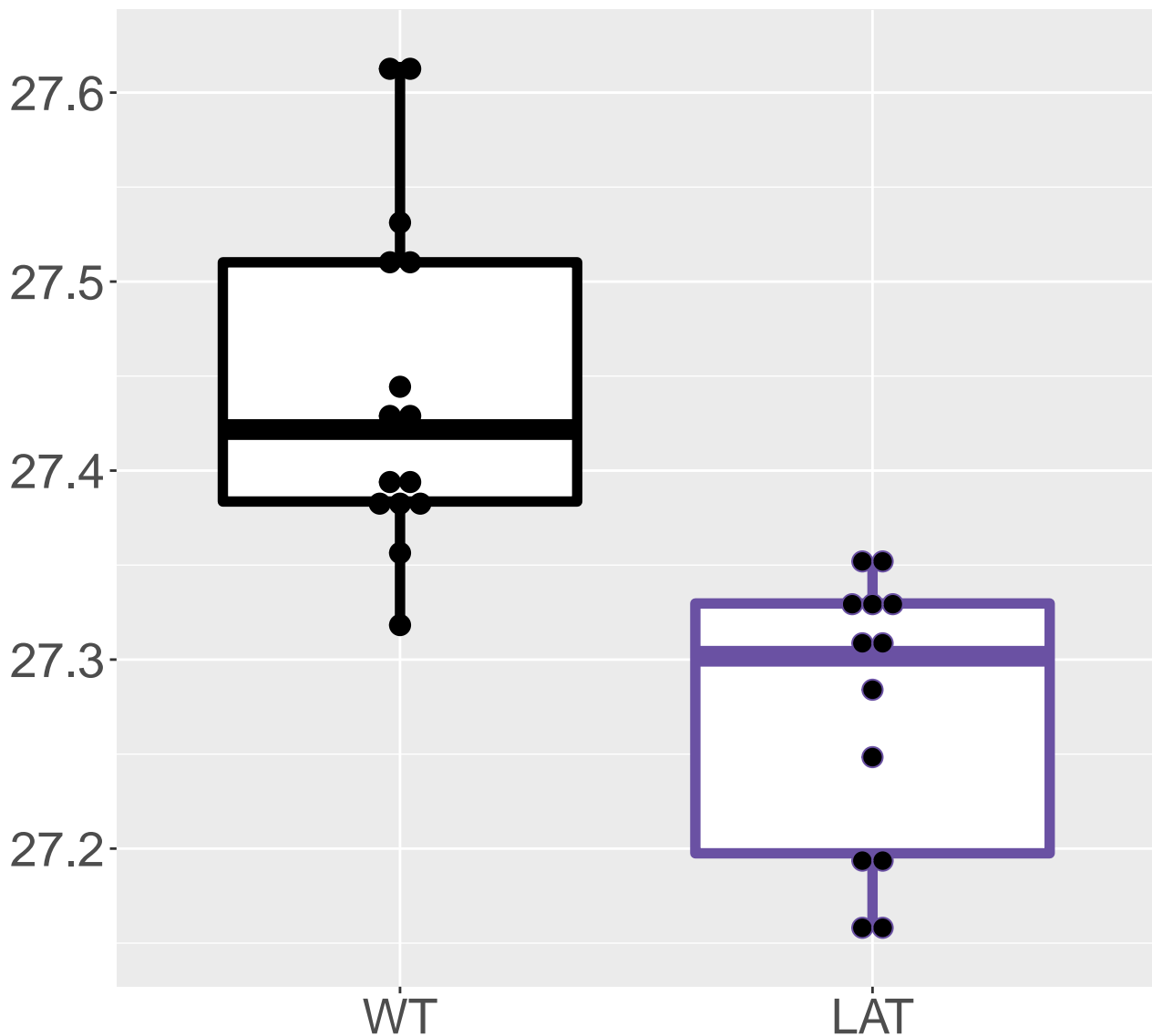


P62301_40S ribosomal protein S13
FDR = 0.011, FC = -0.24

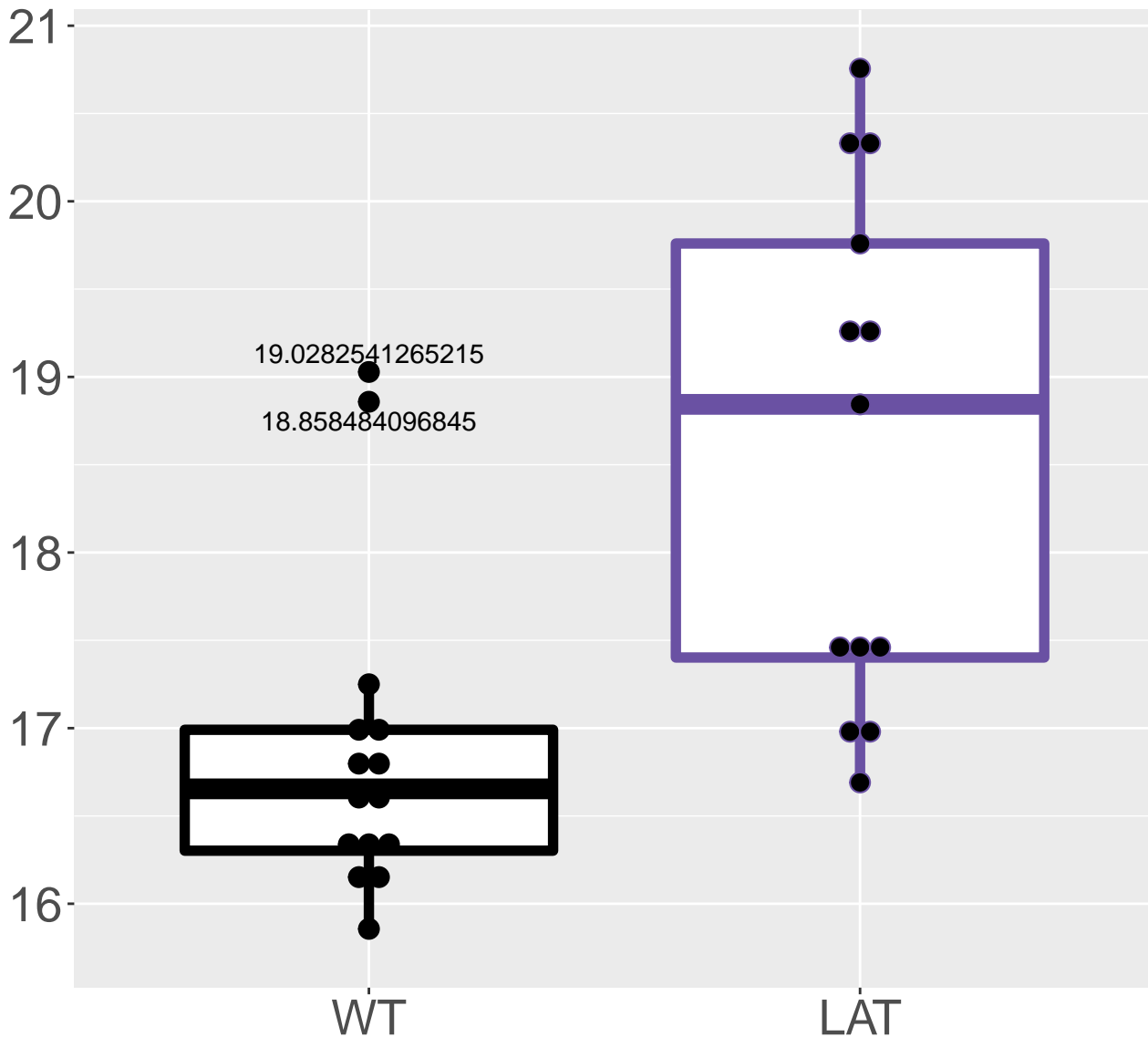


P62962_Profilin-1

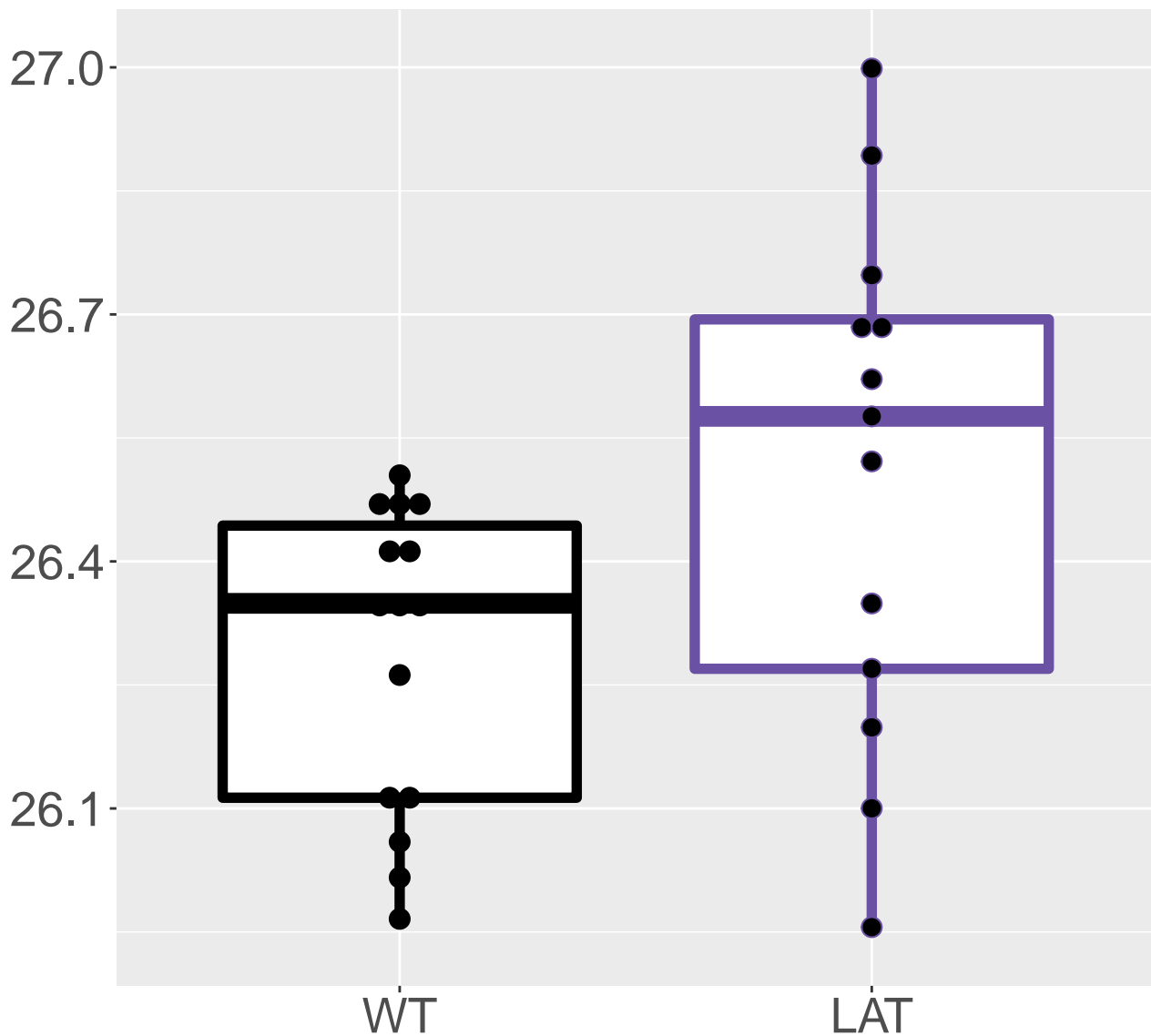
FDR = 0.011, FC = -0.23



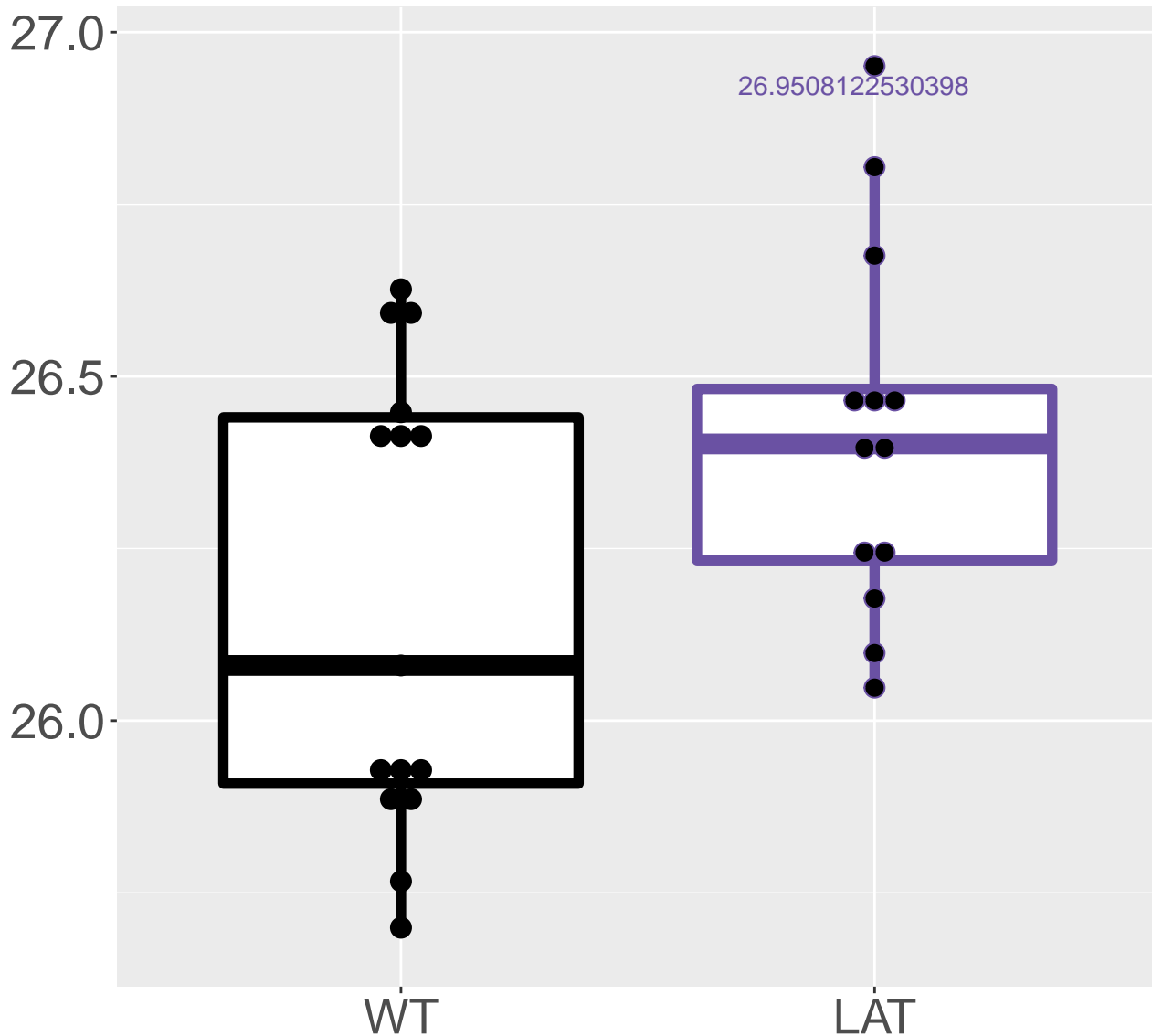
Q8R1S9_Sodium-coupled neutral a.
FDR = 0.012, FC = 2.5



P57780_Alpha-actinin-4
FDR = 0.012, FC = 0.47

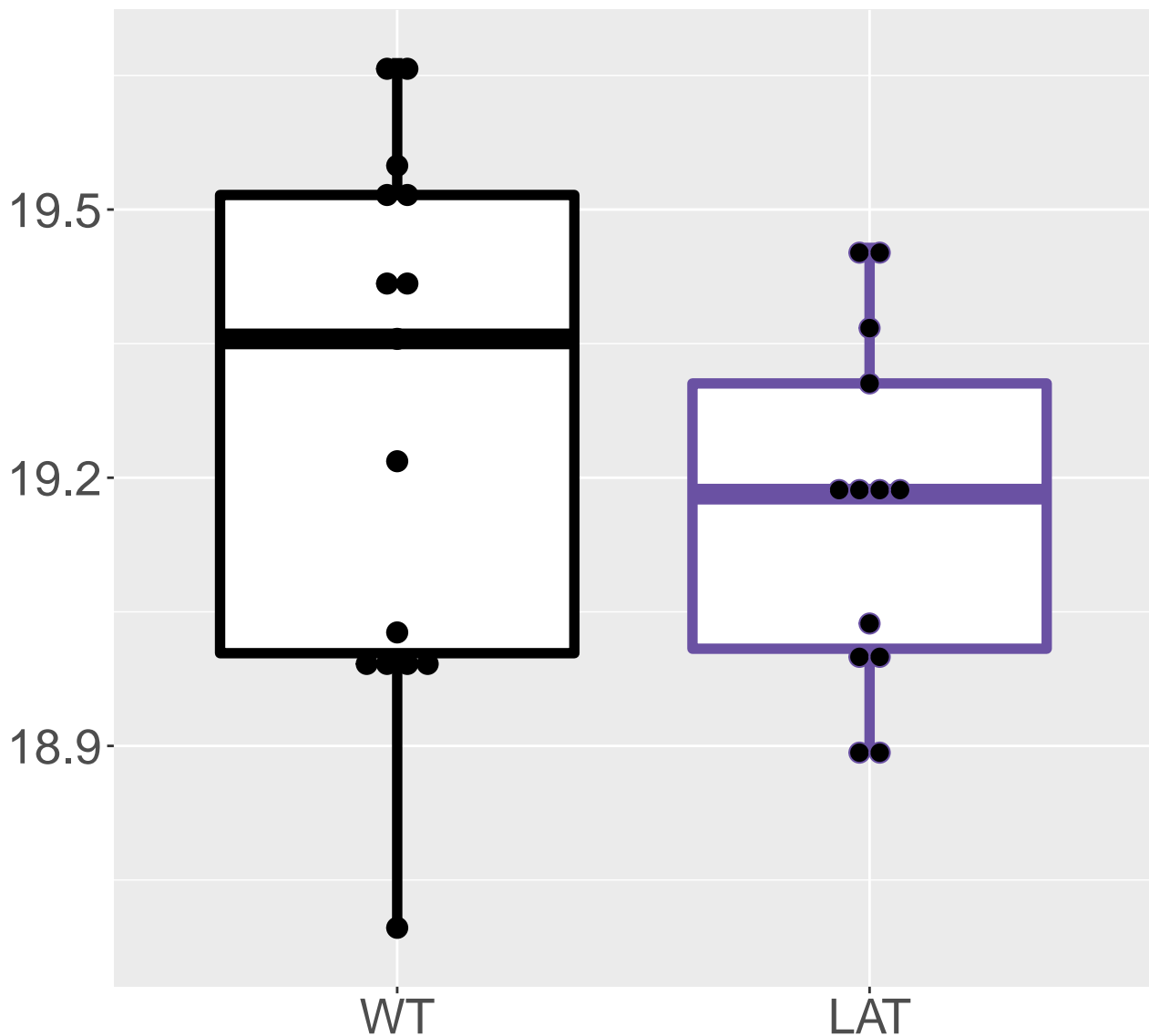


Q9QXD1_Peroxisomal acyl-coenzym.
FDR = 0.012, FC = 0.32, sex***

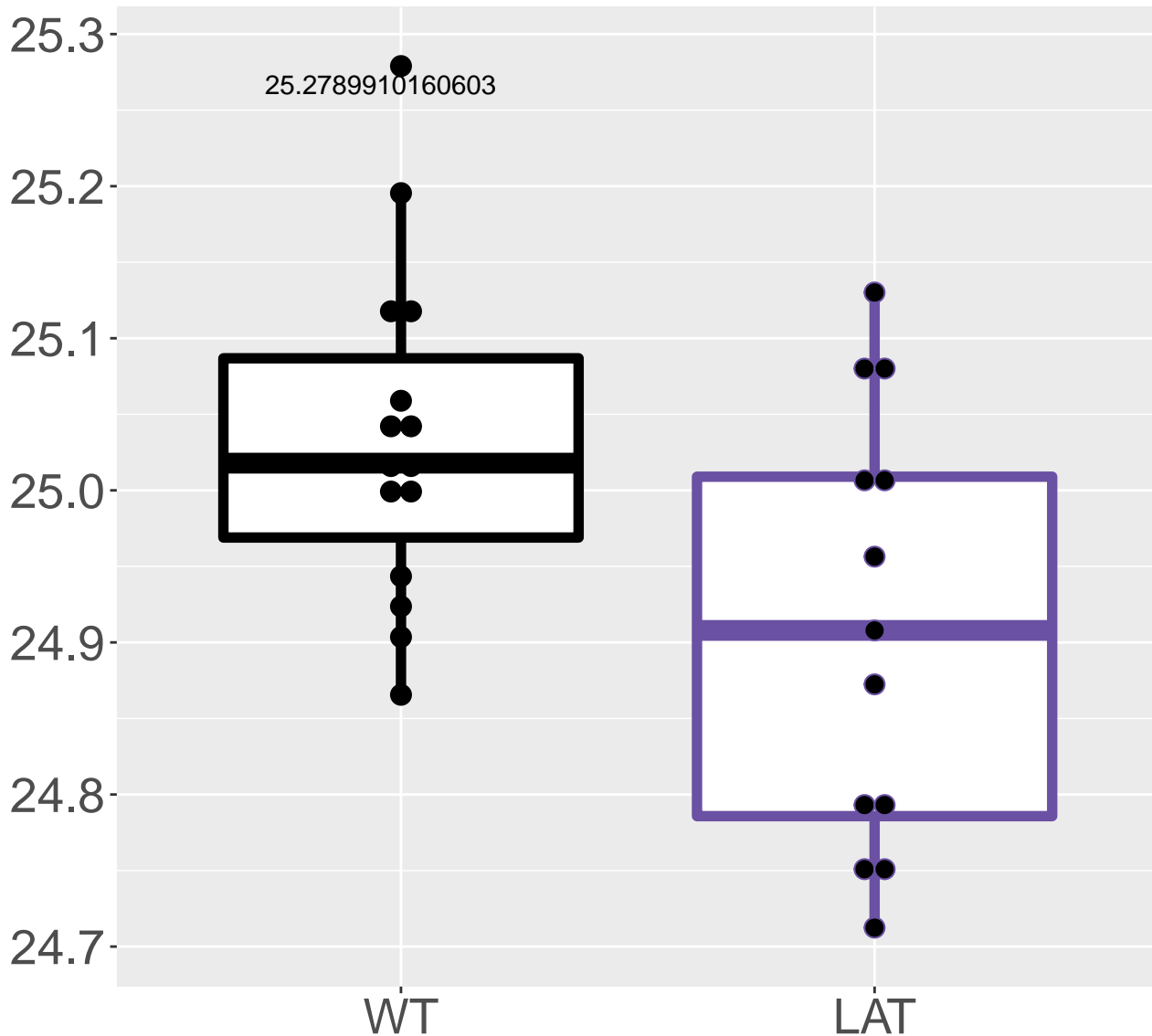


Q8VCH7_Retinol dehydrogenase 10

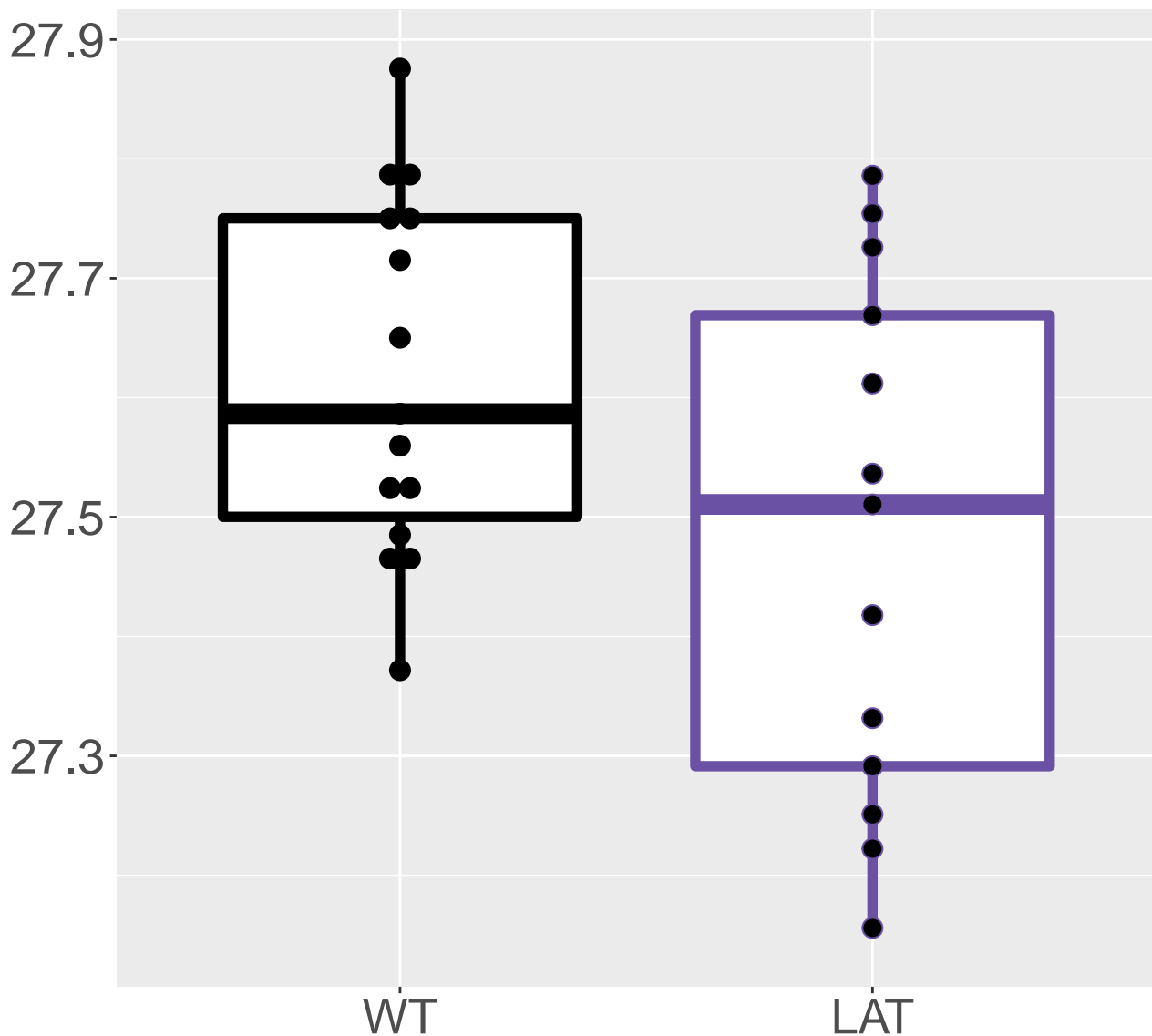
FDR = 0.012, FC = -0.3, sex***



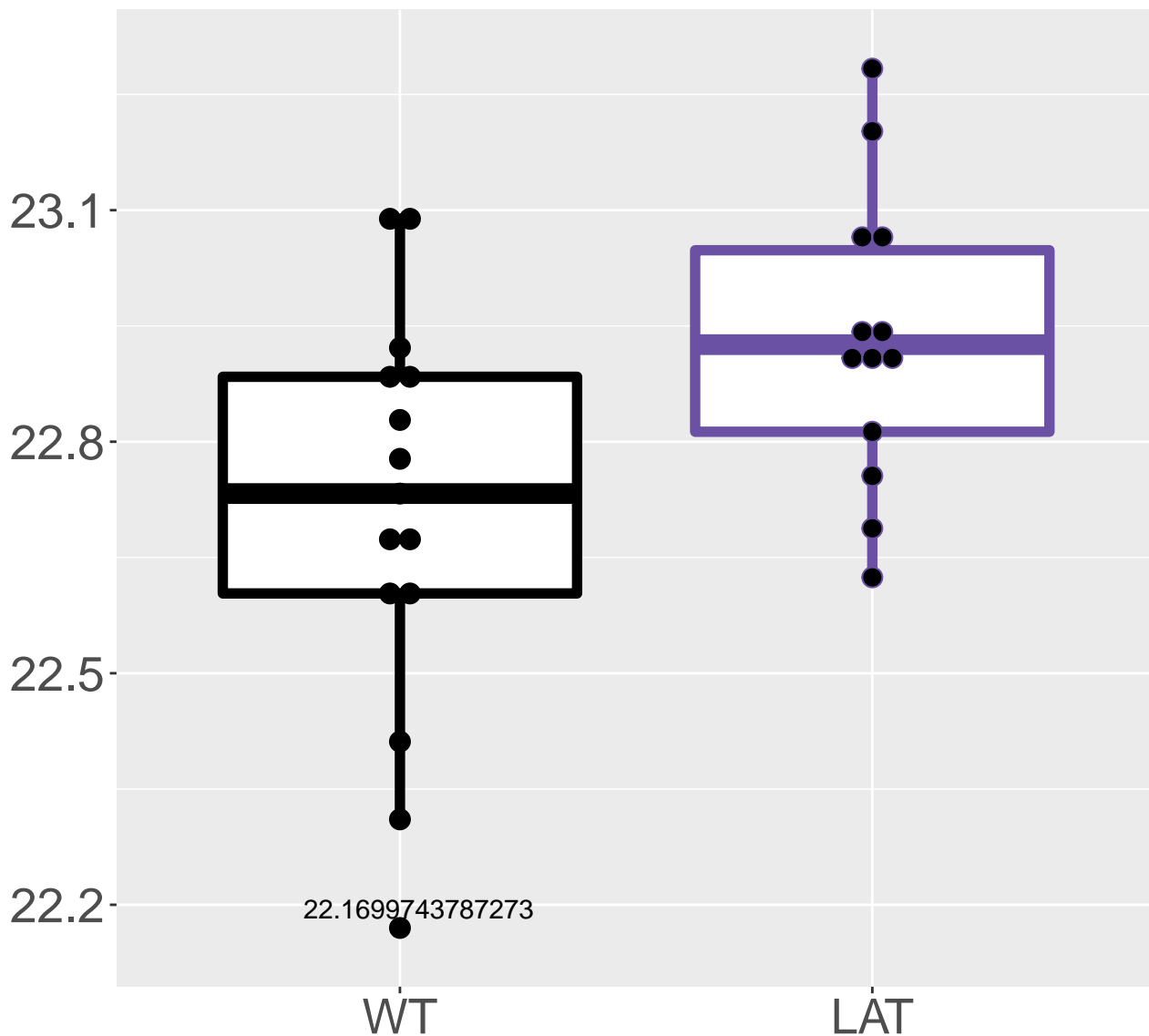
Q9CQZ5_NADH dehydrogenase [ubiq.
FDR = 0.012, FC = -0.28, sex**



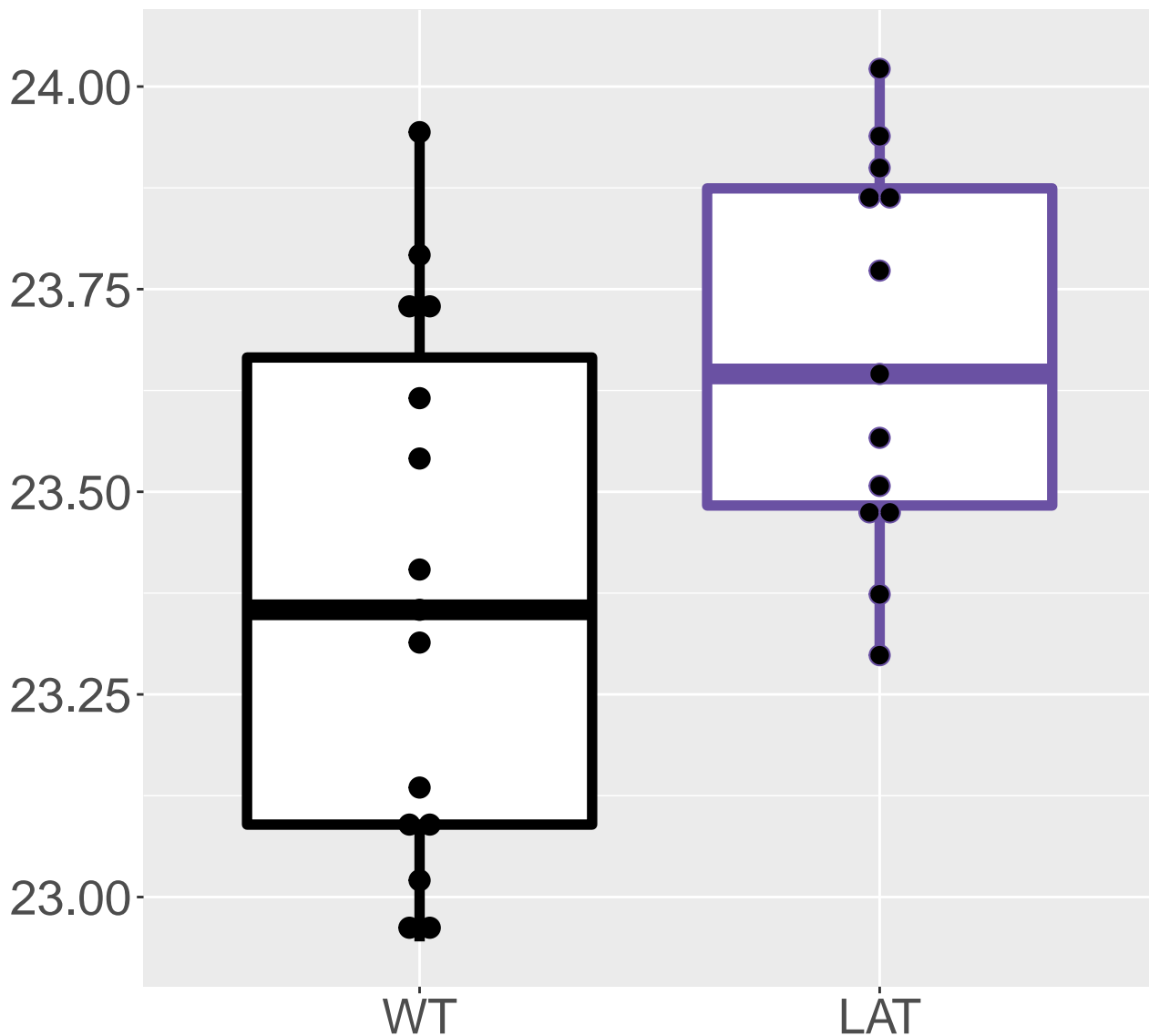
P99029_Peroxiredoxin-5, mitocho.
FDR = 0.012, FC = -0.22, sex***



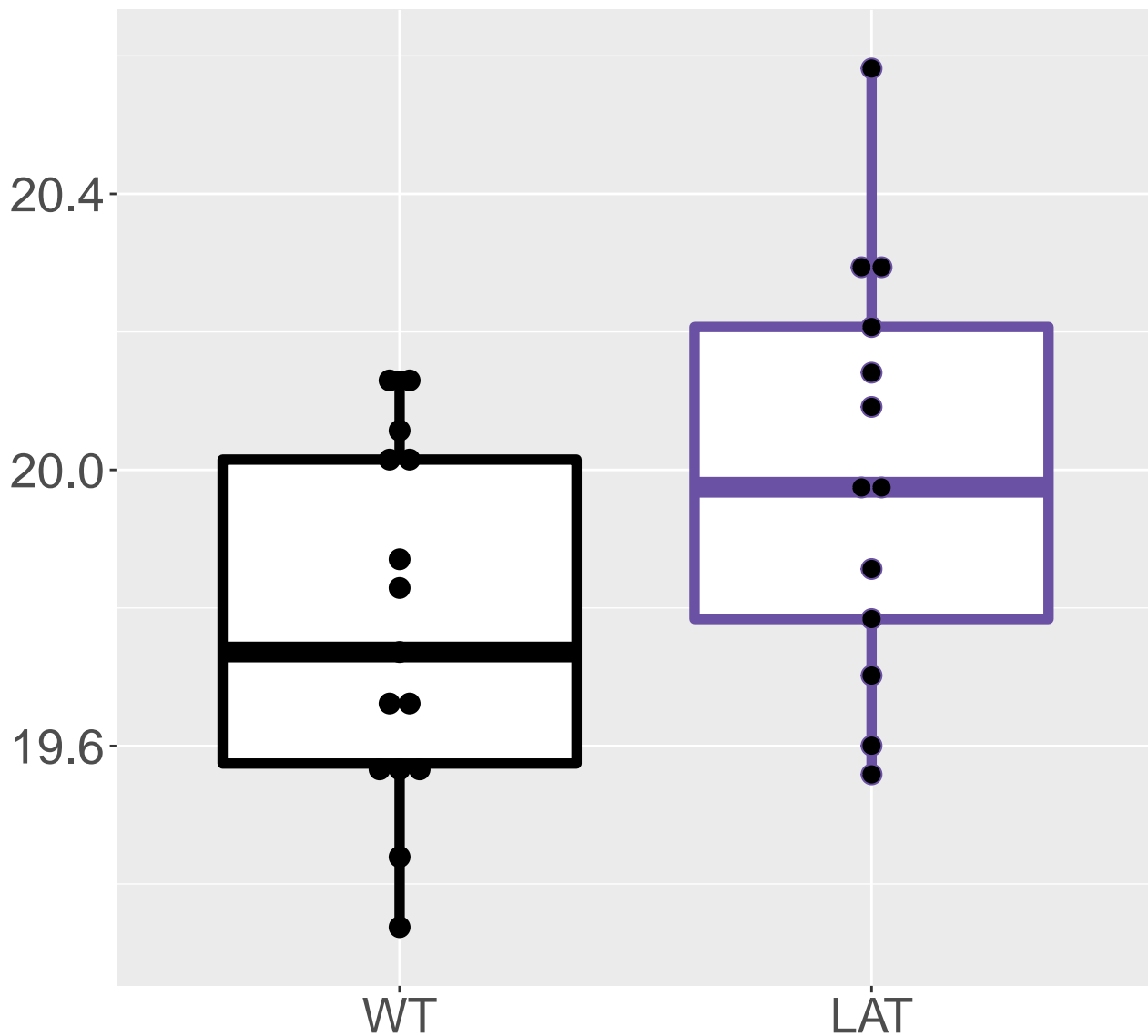
Q8BIJ6_Isoleucine--tRNA ligase,.
FDR = 0.012, FC = 0.43, sex**



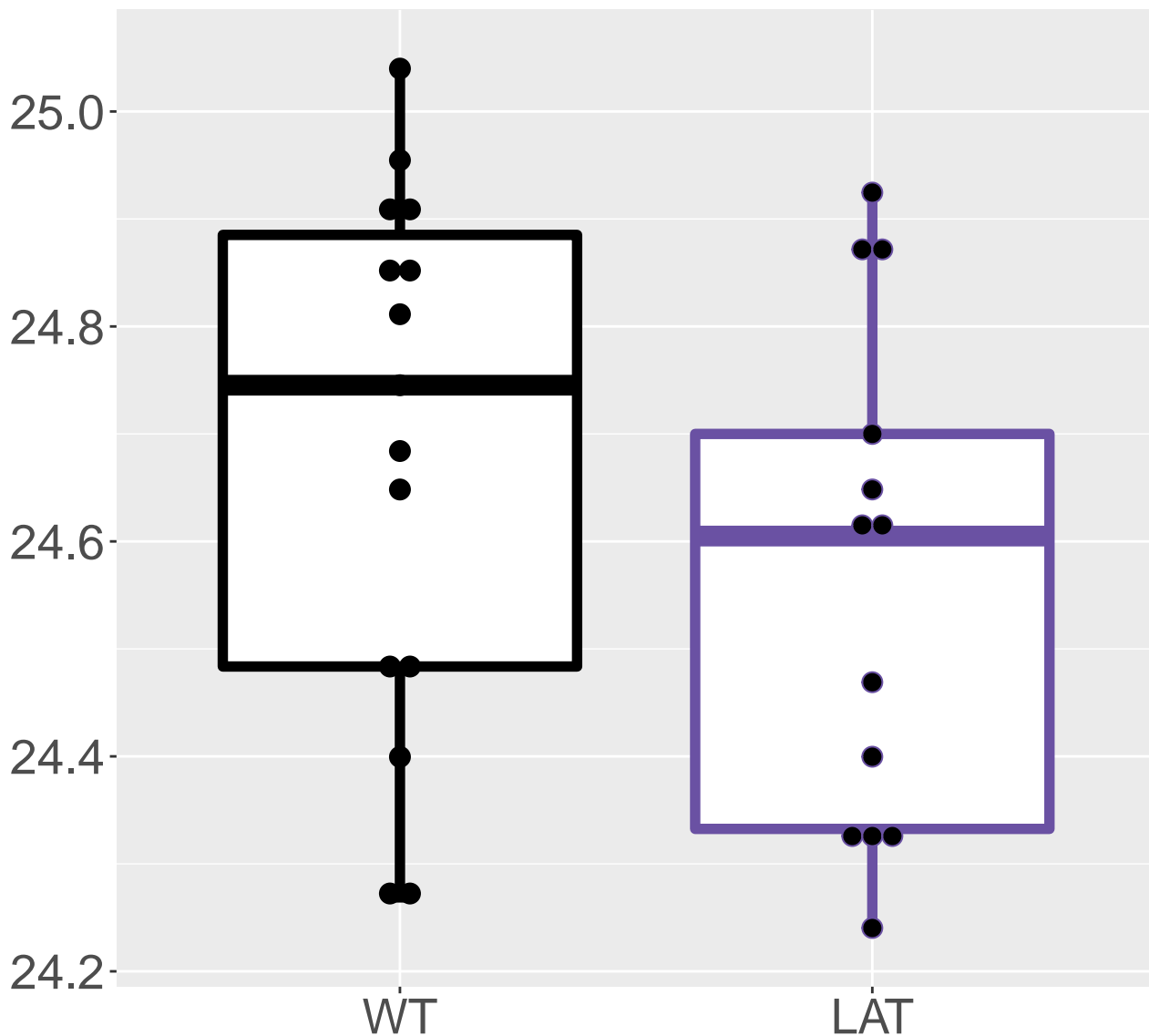
Q61586_Glycerol-3-phosphate acy.
FDR = 0.012, FC = 0.42, sex***



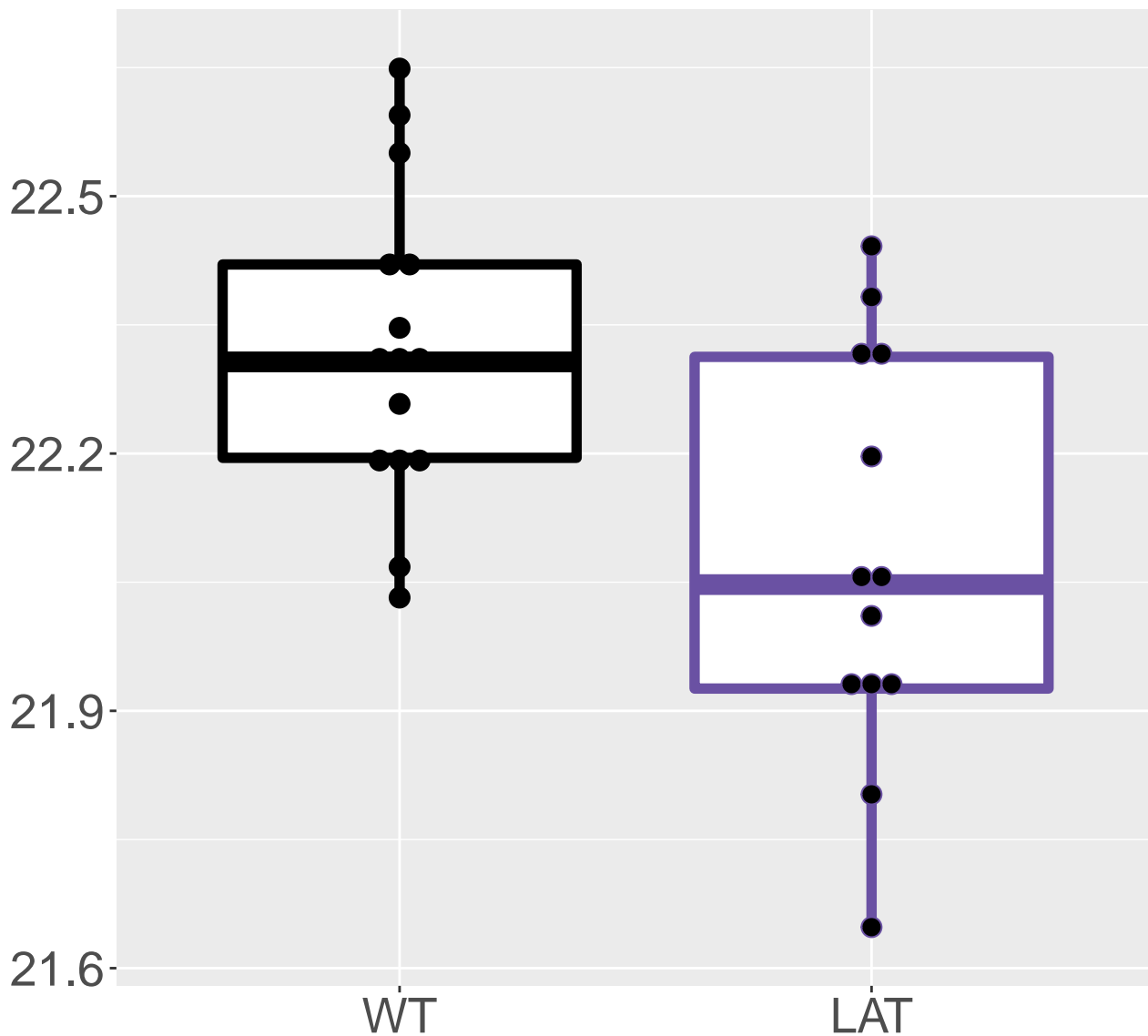
Q91VH2_Sorting nexin-9
FDR = 0.012, FC = 0.51, sex*



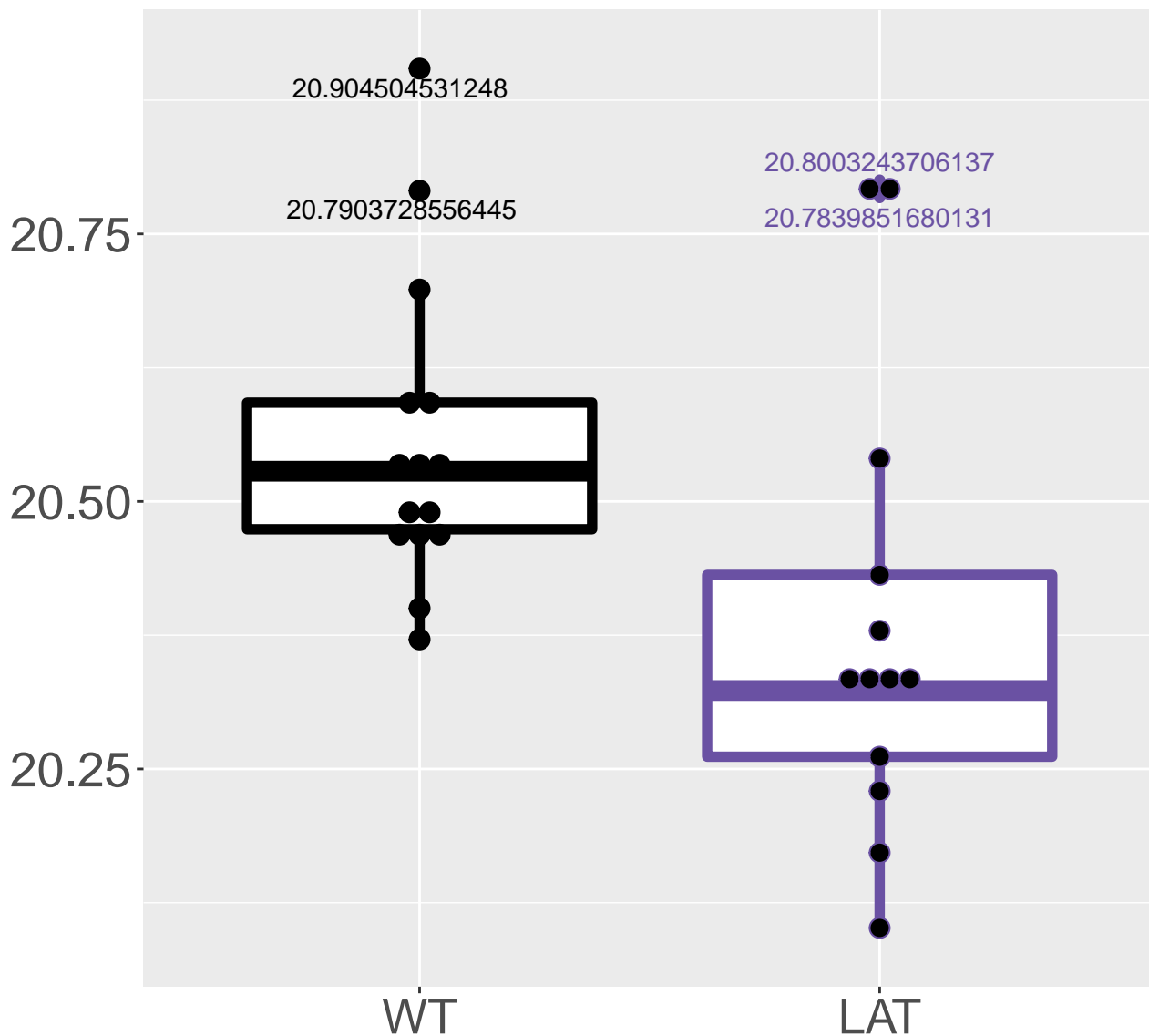
P48771_Cytochrome c oxidase sub.
FDR = 0.012, FC = -0.45, sex**



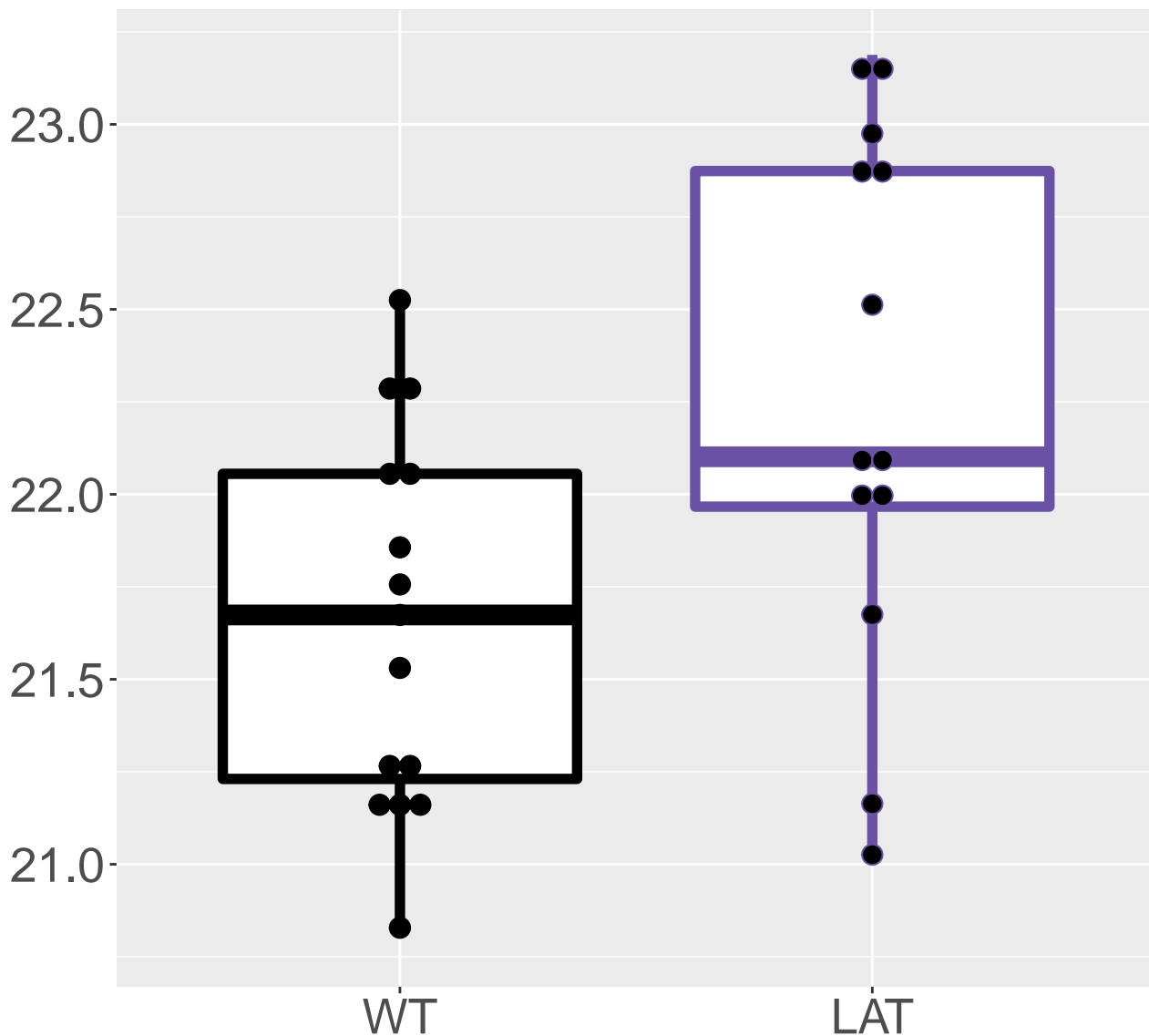
P49935_Pro-cathepsin H
FDR = 0.013, FC = -0.4



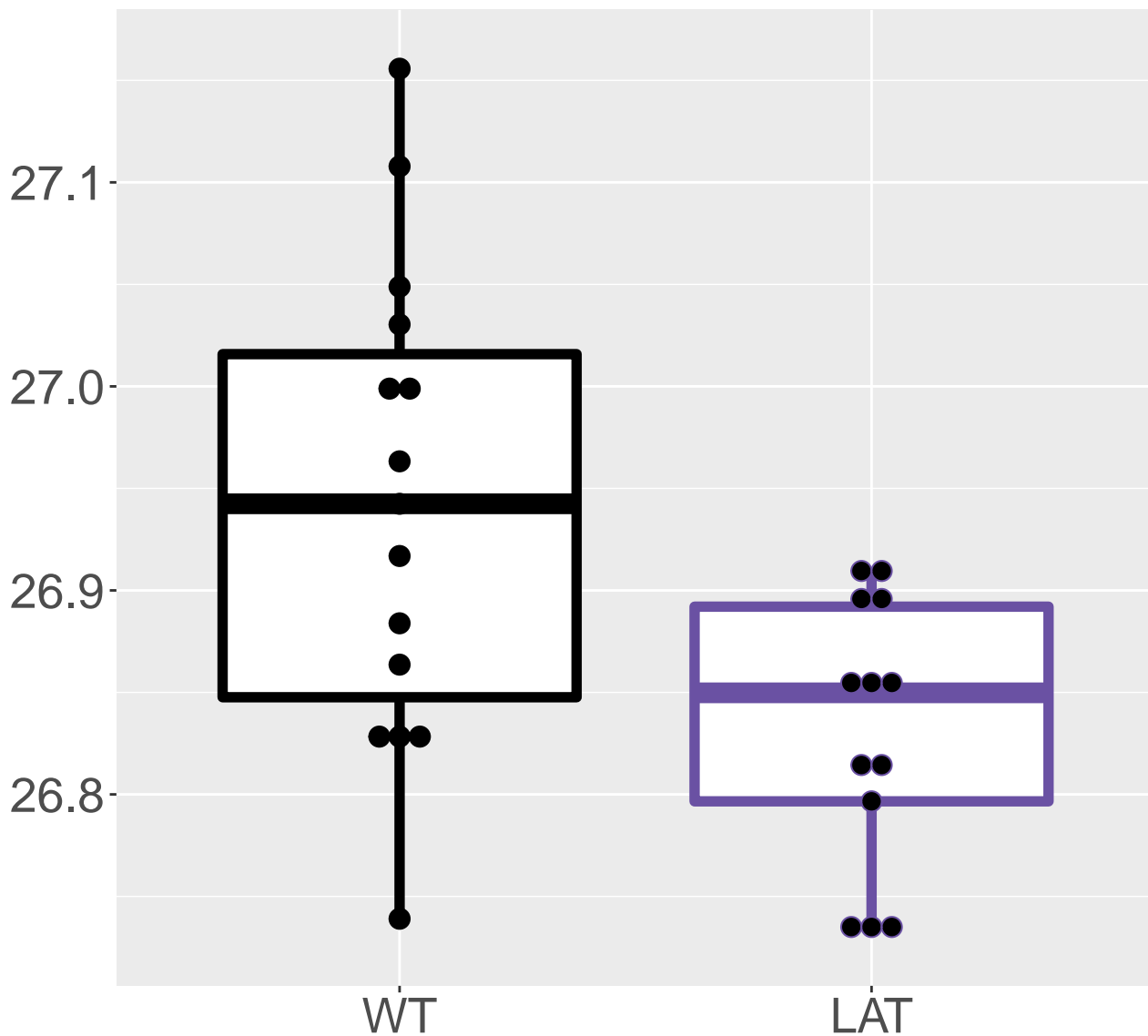
Q8BVA5_Lipid droplet-associated.
FDR = 0.013, FC = -0.36, sex*



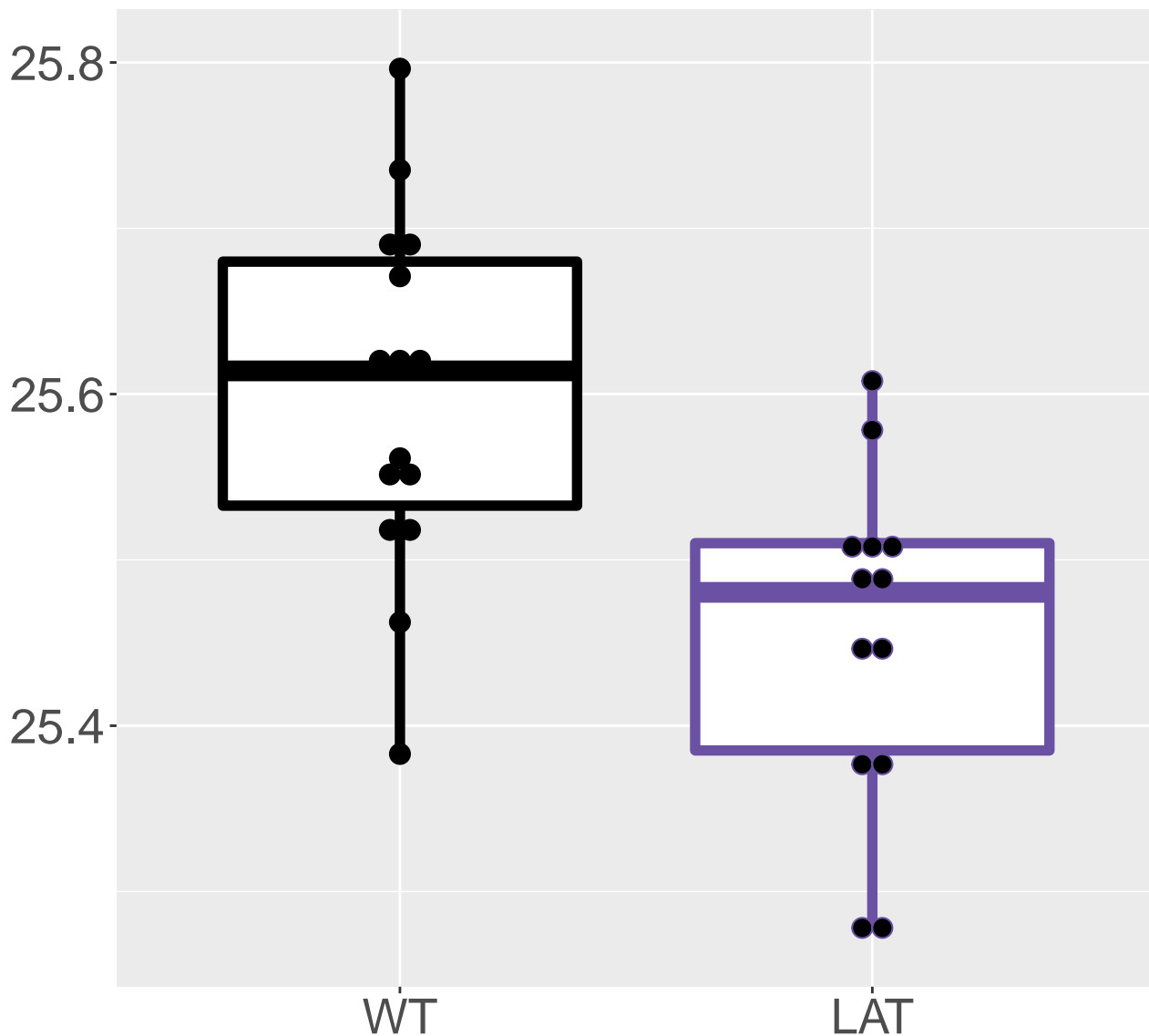
Q9ER72_Cysteine--tRNA ligase, c.
FDR = 0.013, FC = 1.1



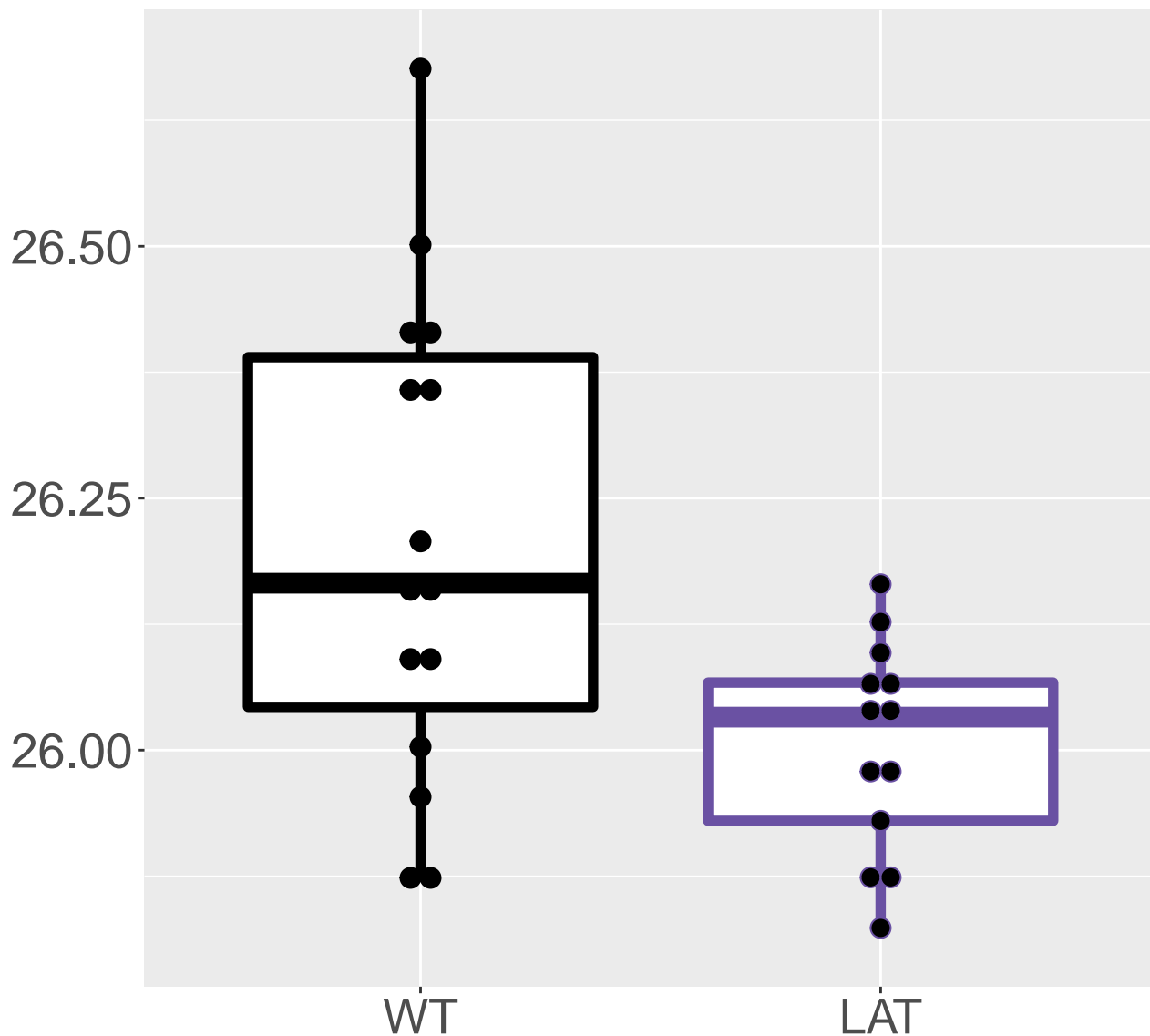
Q64105_Sepiapterin reductase
FDR = 0.014, FC = -0.22, sex**



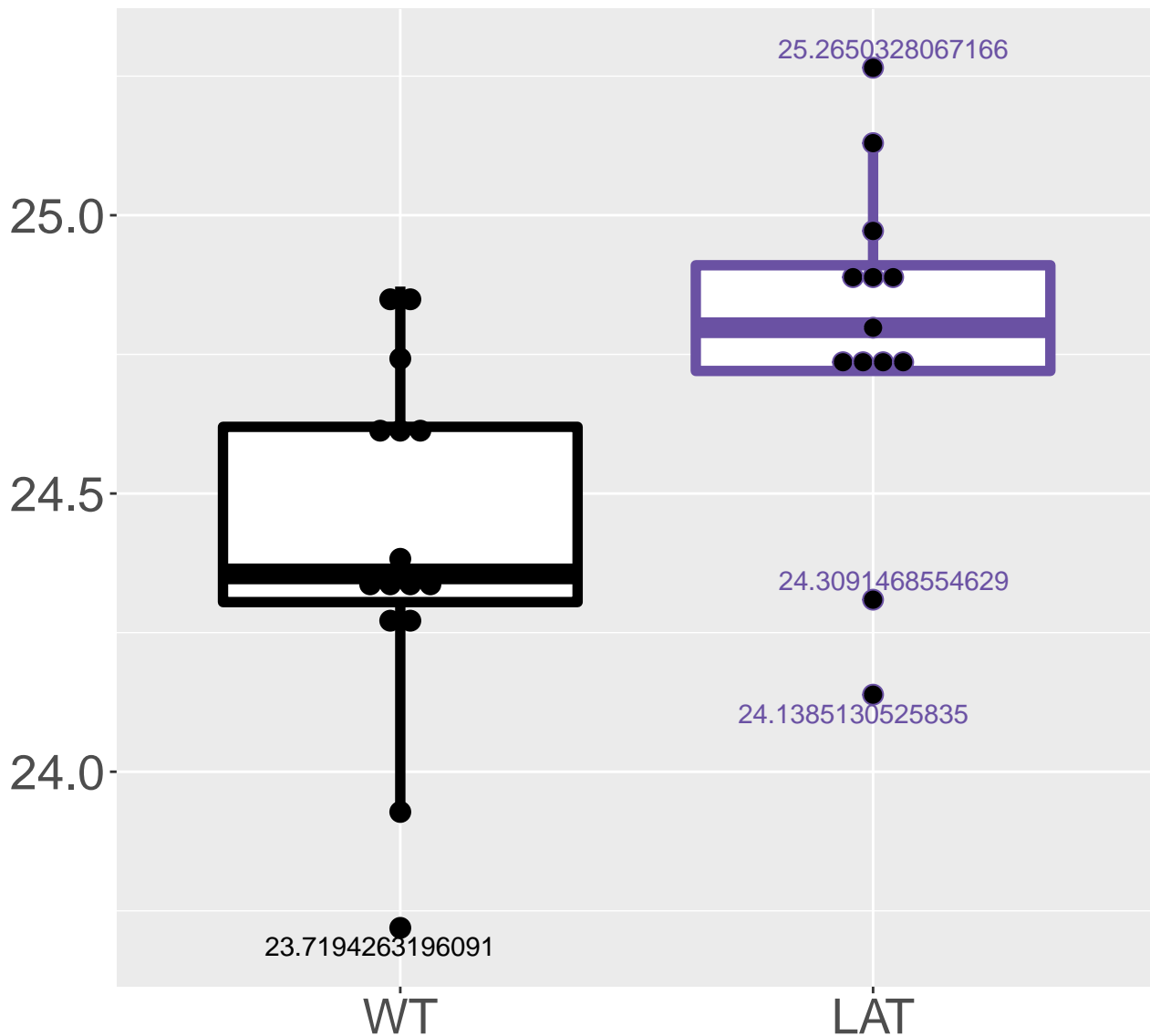
O70435_Proteasome subunit alpha.
FDR = 0.014, FC = -0.24



Q64523_Histone H2A type 2-C
FDR = 0.014, FC = -0.38, sex**

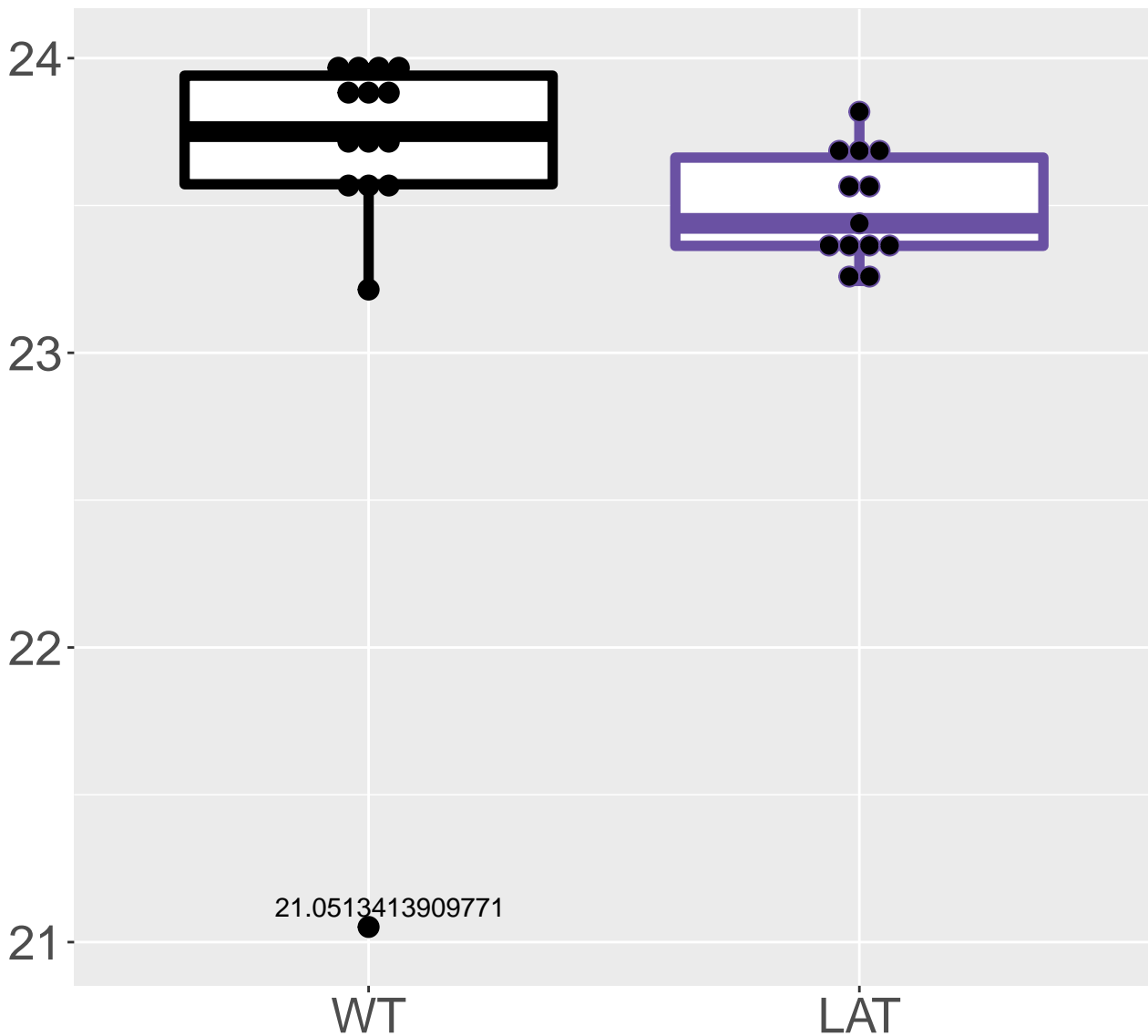


Q91W43_Glycine dehydrogenase (d. FDR = 0.014, FC = 0.56

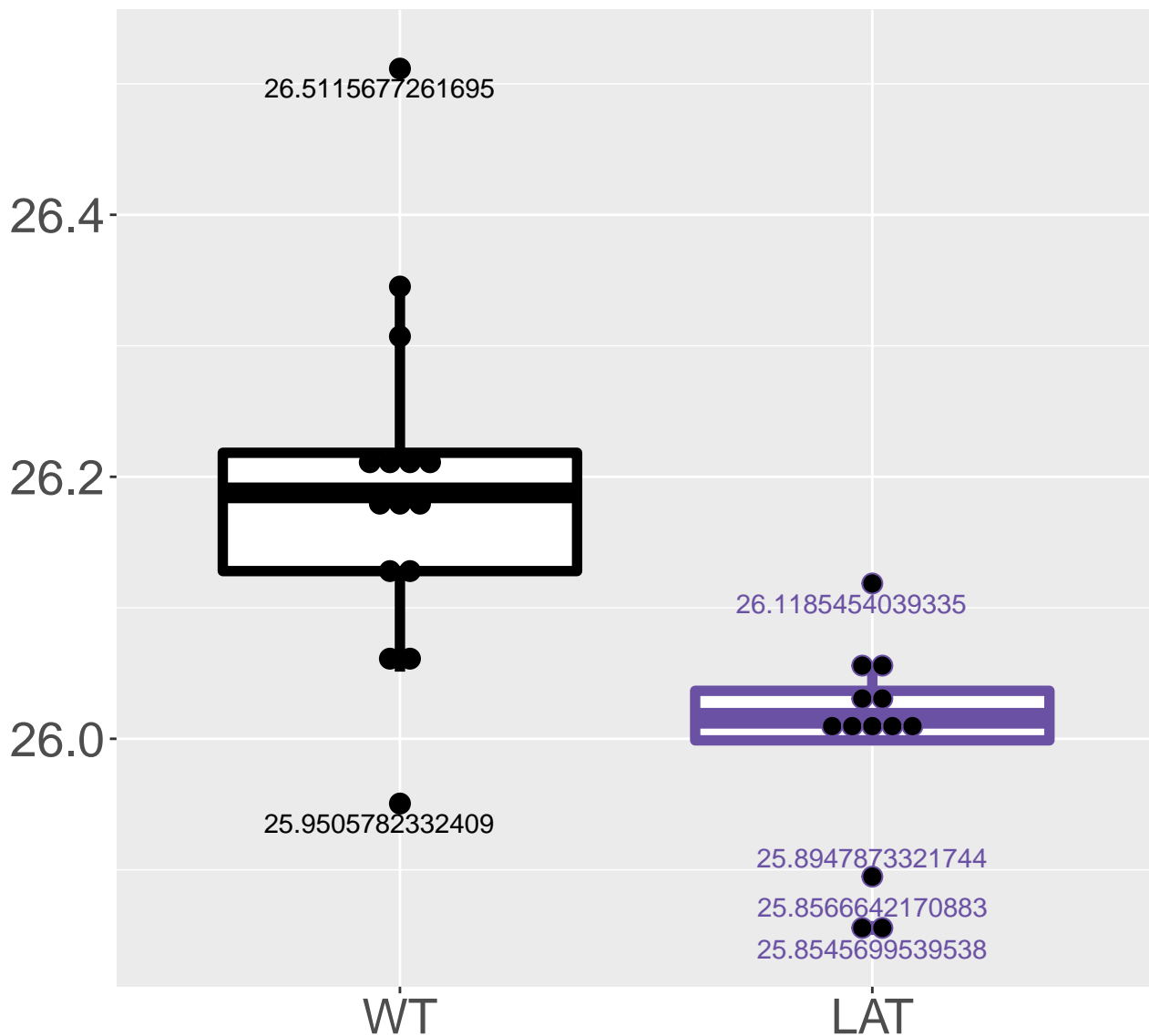


P47915_60S ribosomal protein L29

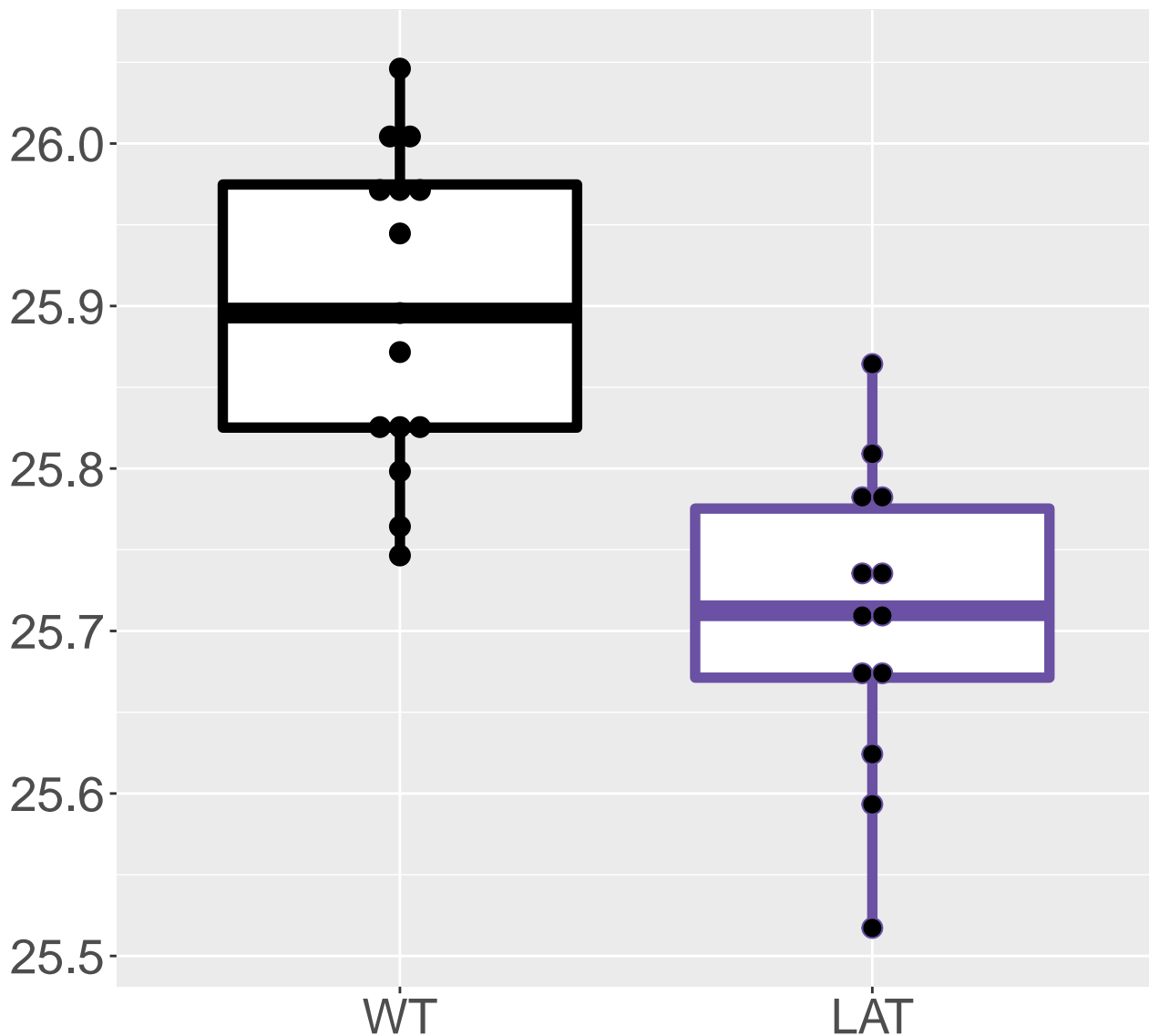
FDR = 0.014, FC = -0.38



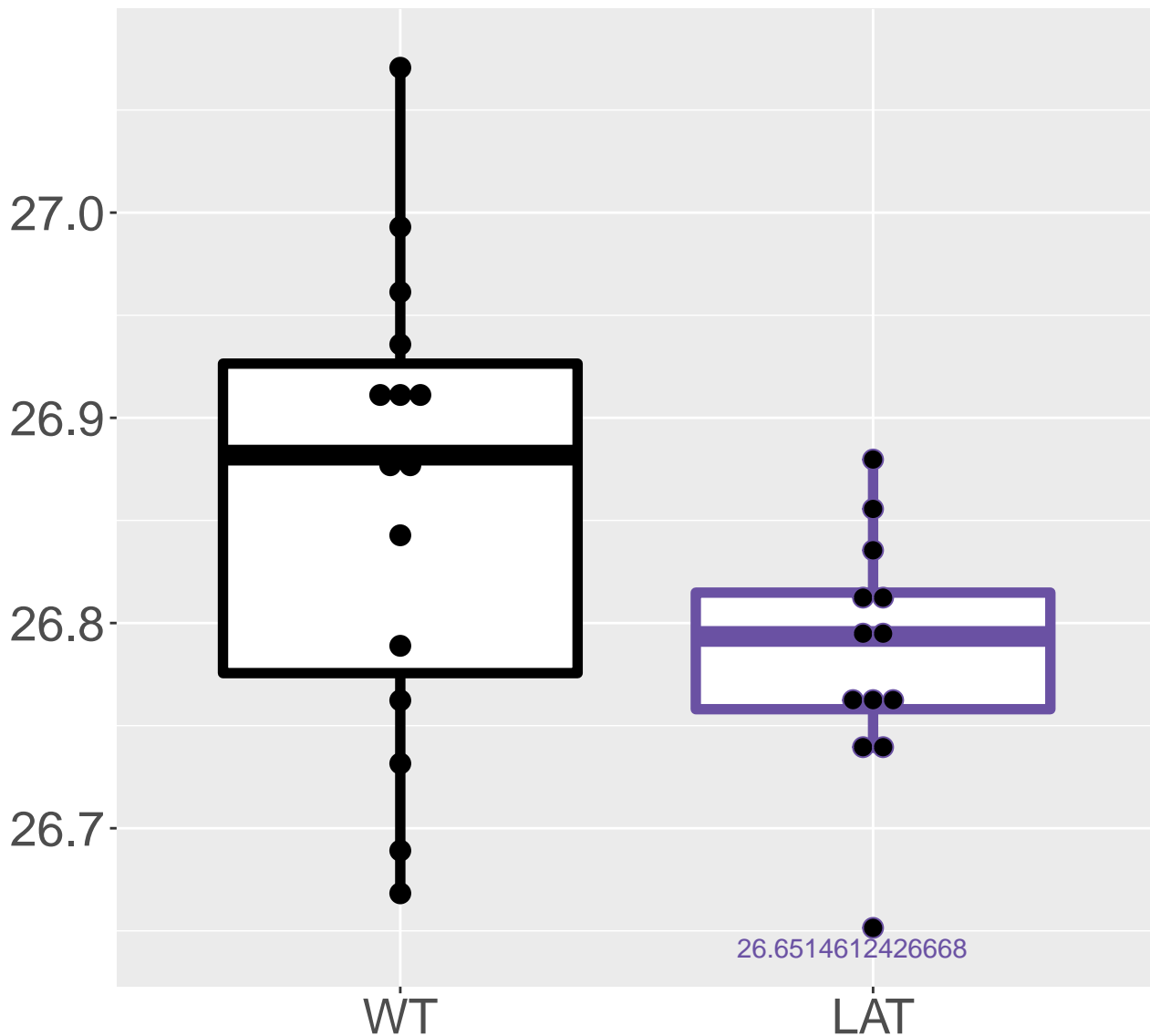
FDR = 0.014, FC = -0.28



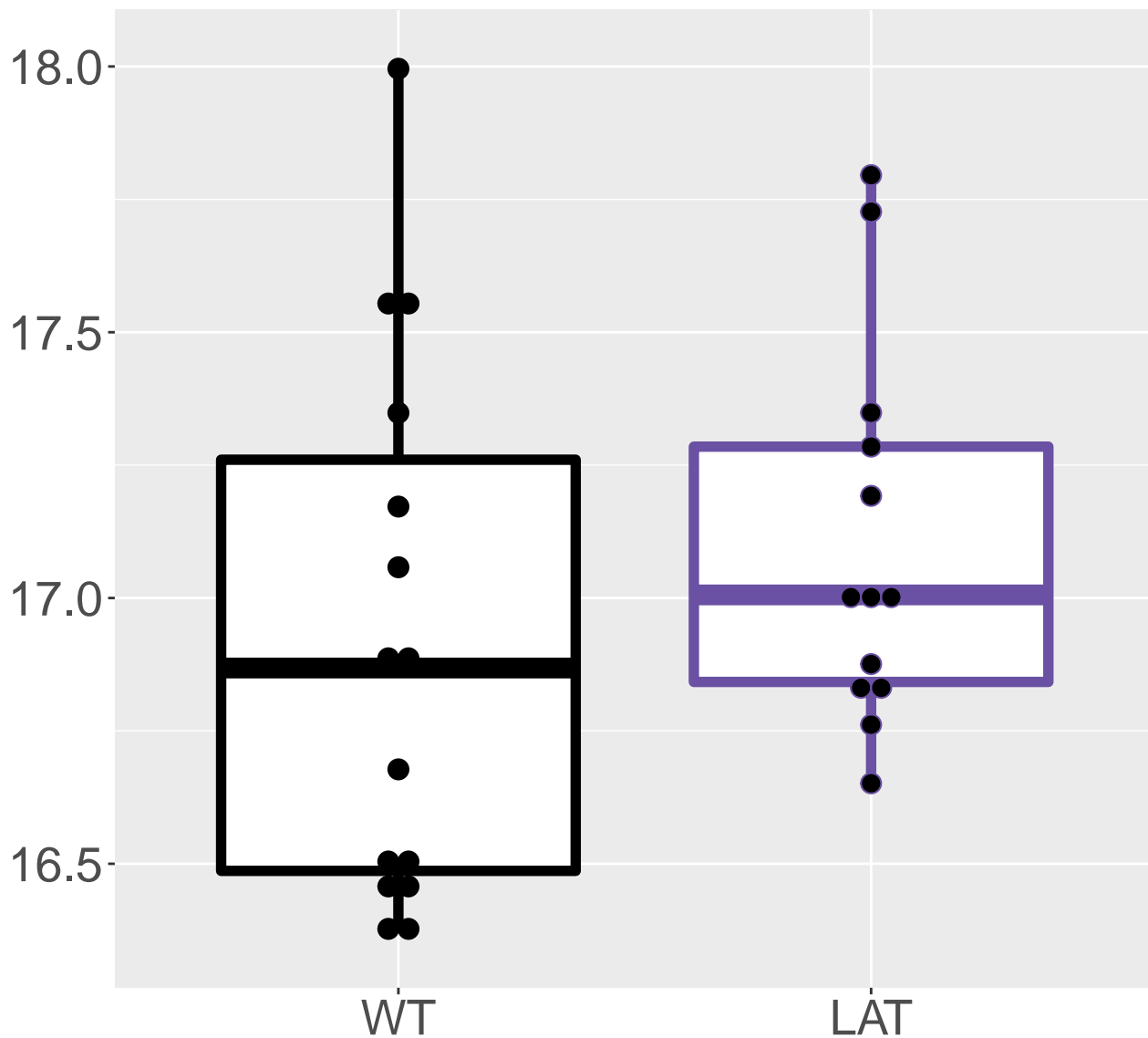
P63323_40S ribosomal protein S12
FDR = 0.014, FC = -0.25



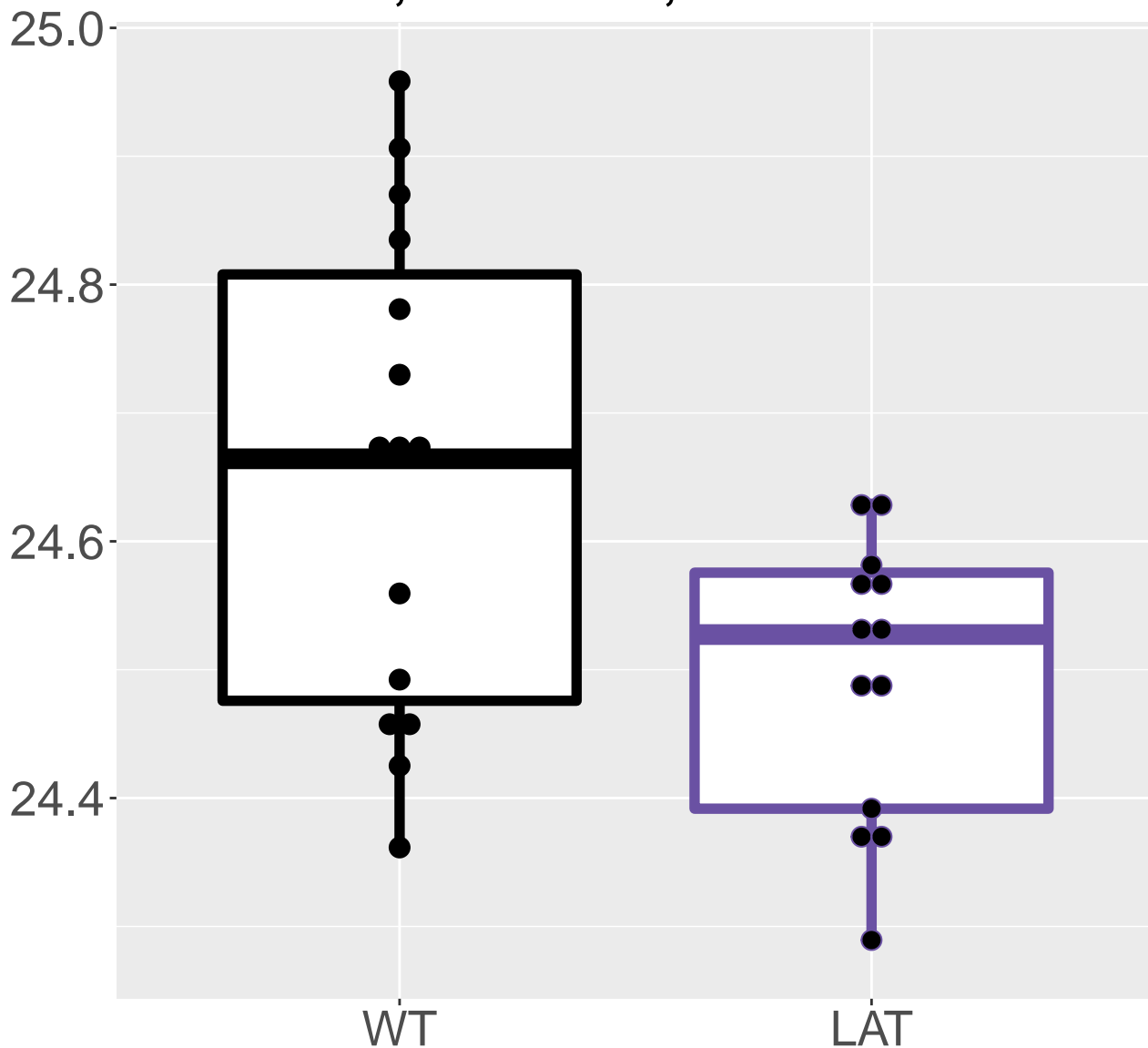
Q9WTP7_GTP:AMP phosphotransfera.
FDR = 0.014, FC = -0.19, sex**



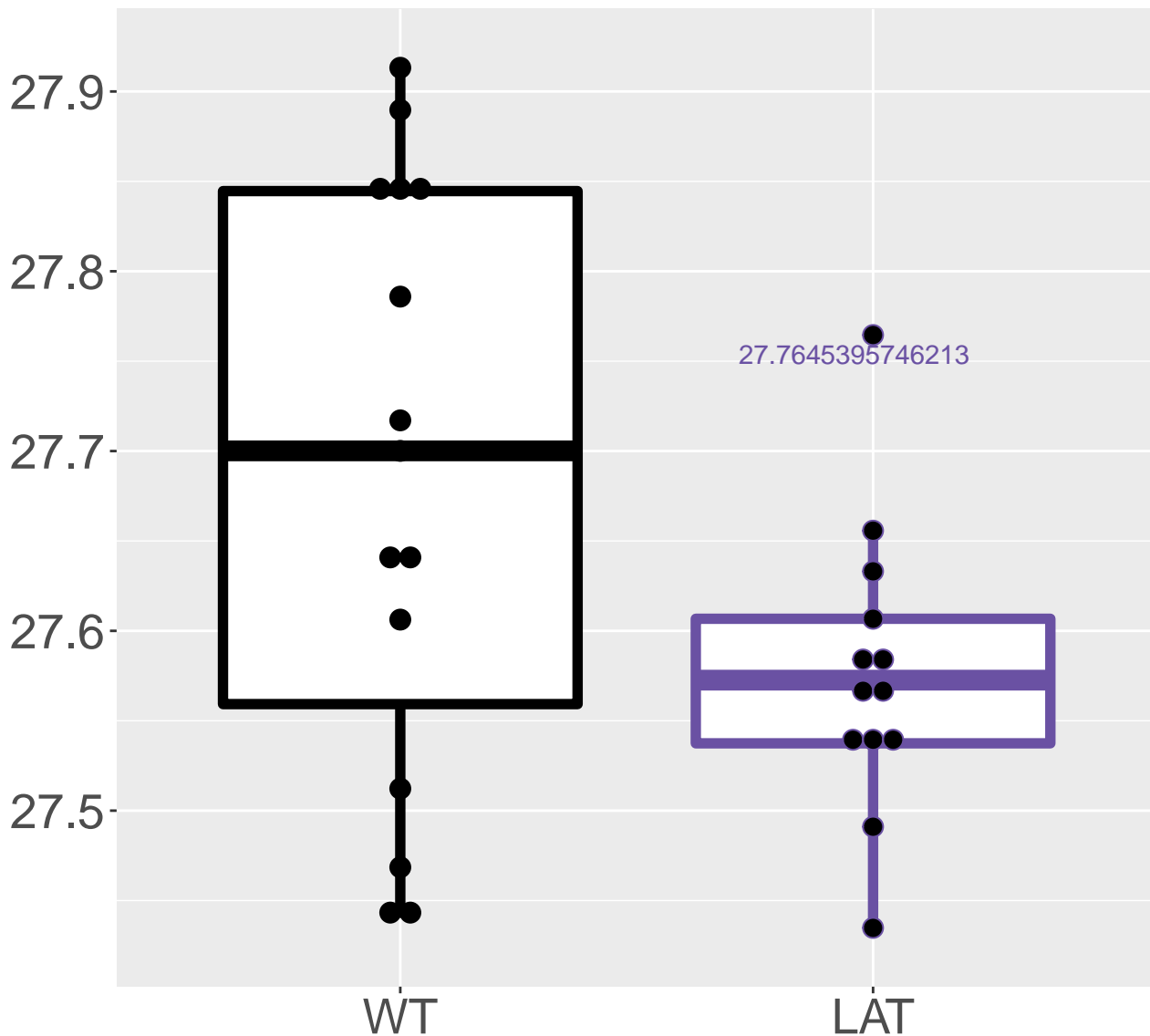
P70398_Probable ubiquitin carbo.
FDR = 0.014, FC = 0.6, sexNA



P63001_Ras-related C3 botulinum.
FDR = 0.014, FC = -0.23, sex***

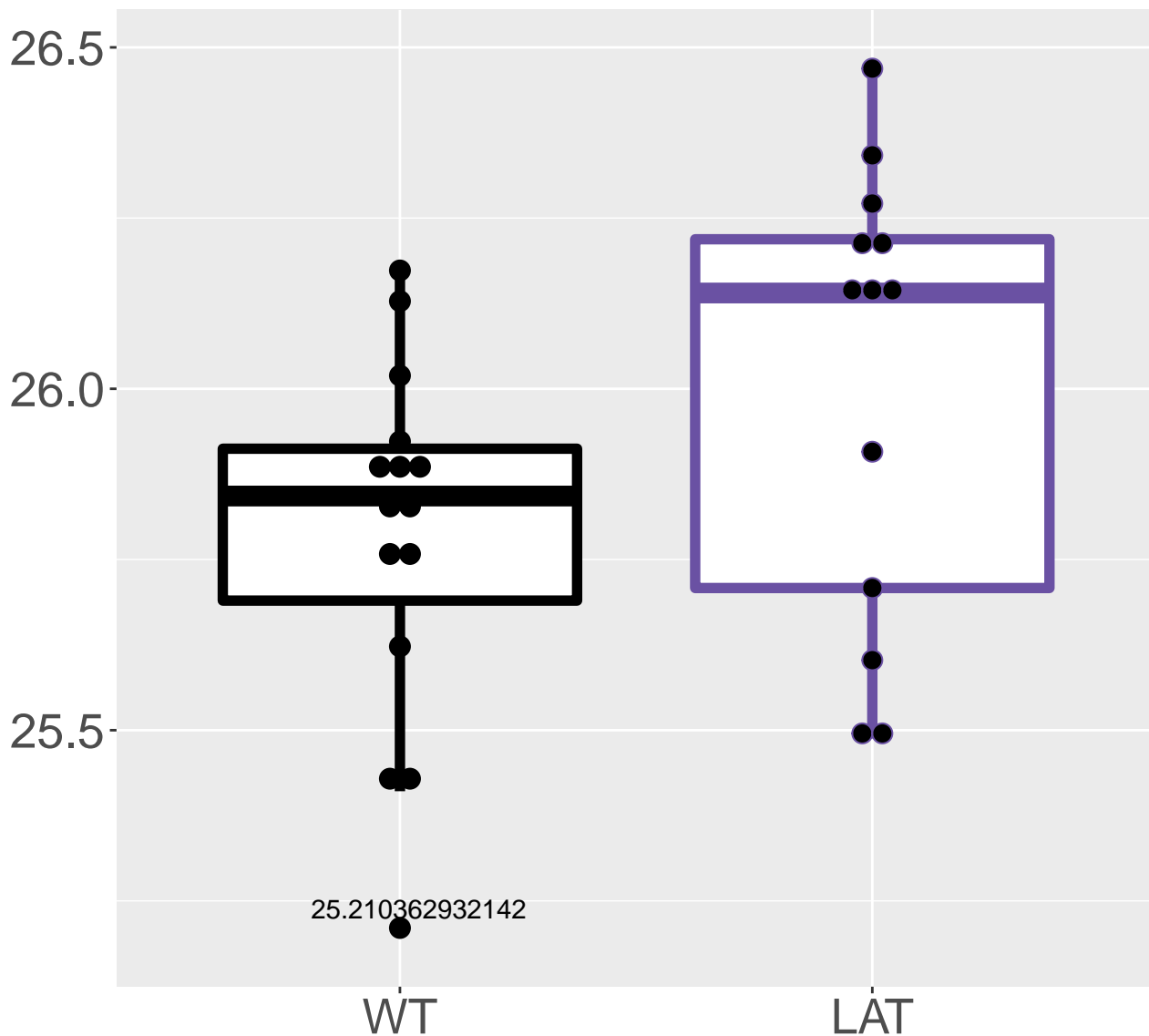


Q8BH95_Enoyl-CoA hydratase, mit.
FDR = 0.015, FC = -0.22, sex***

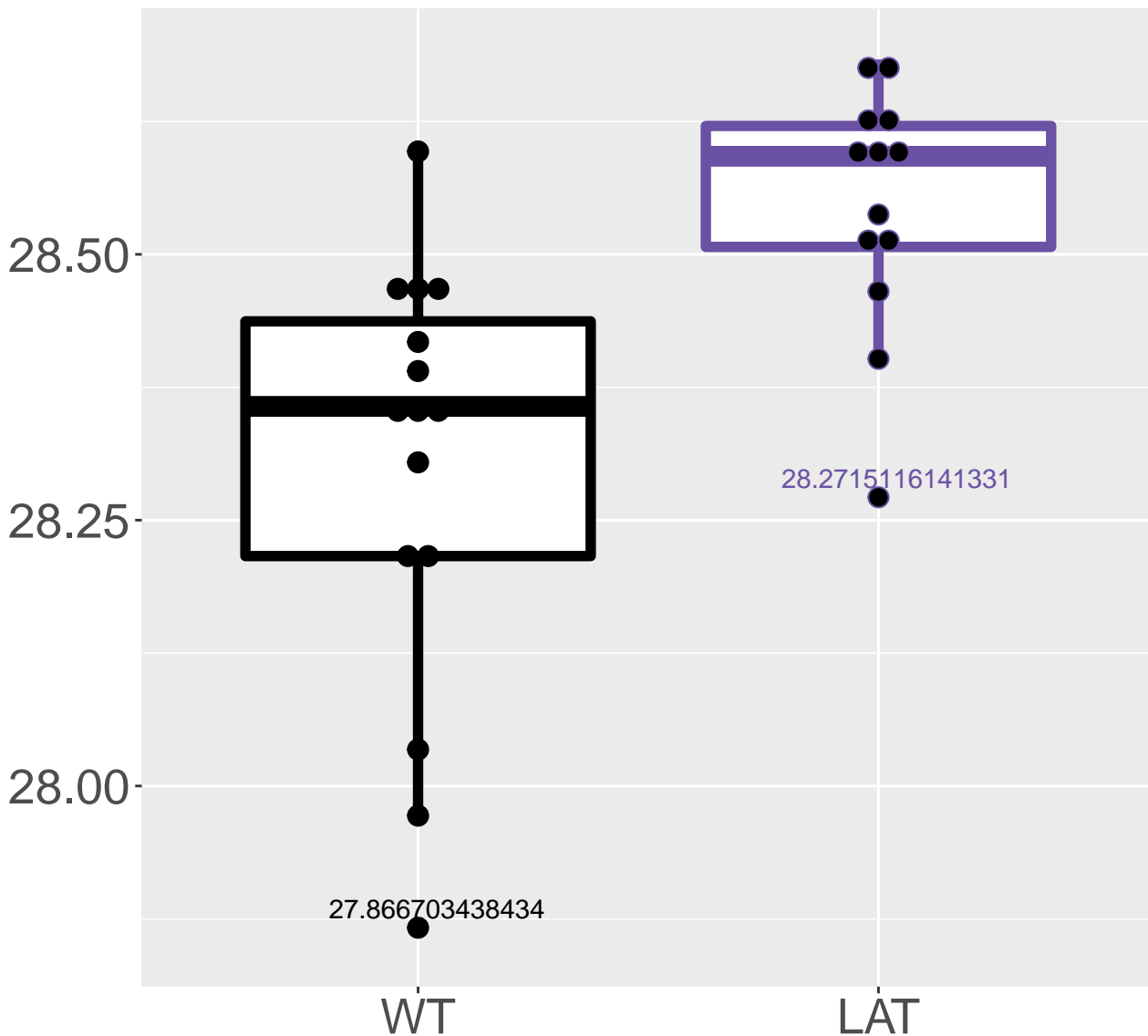


P35564_Calnexin

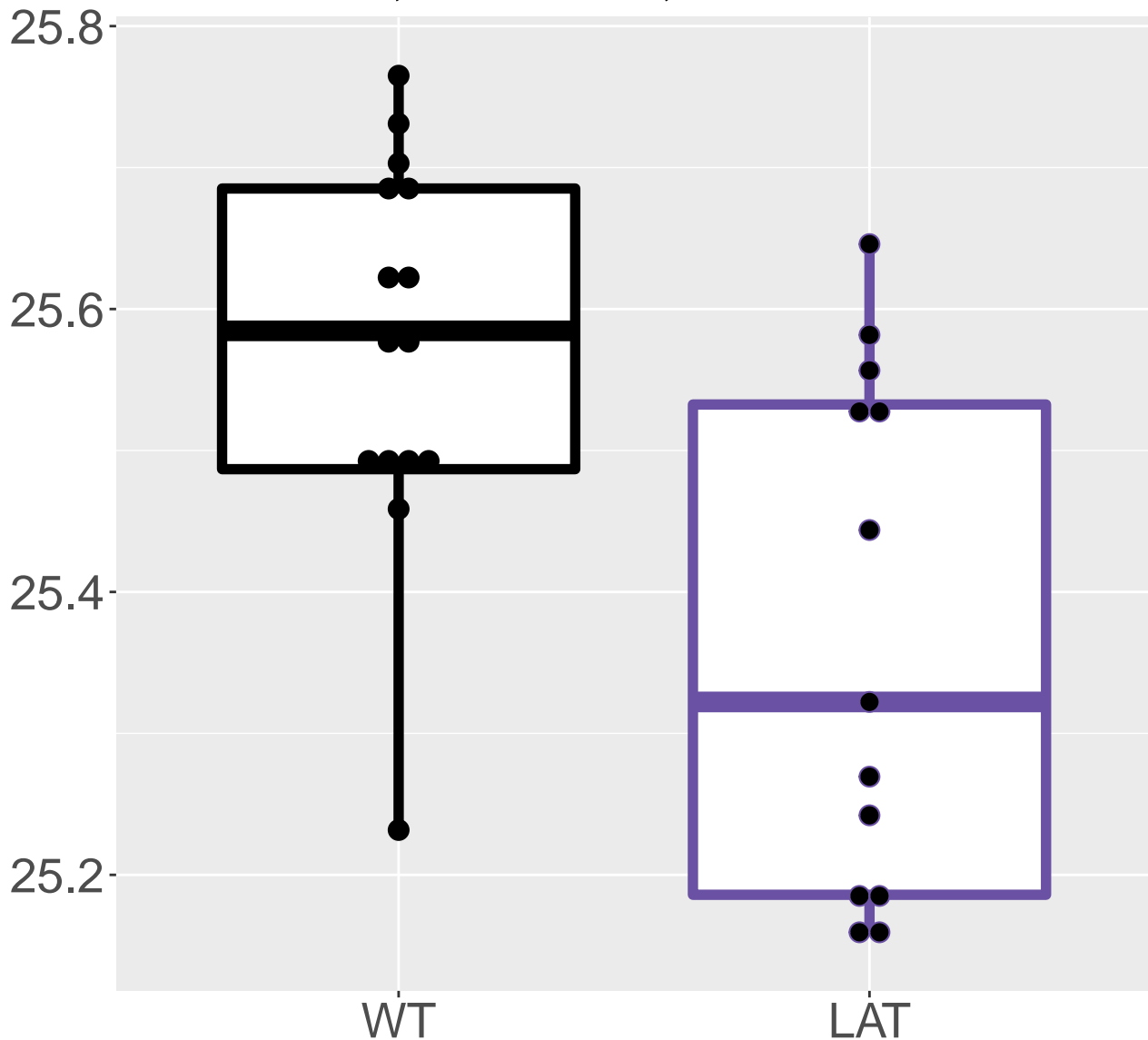
FDR = 0.015, FC = 0.54, sex*



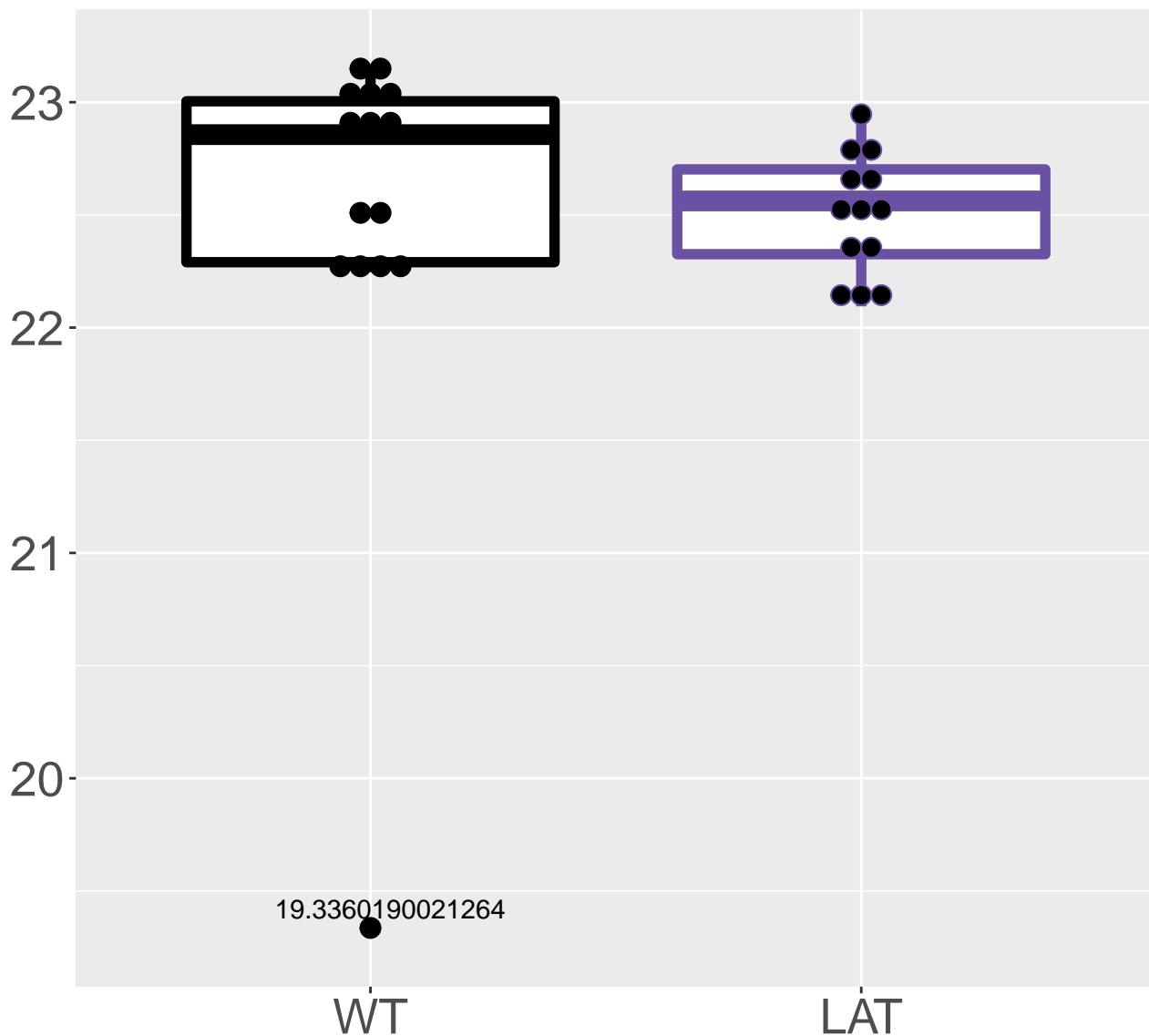
Q9DBT9_Dimethylglycine dehydrog.
FDR = 0.015, FC = 0.38



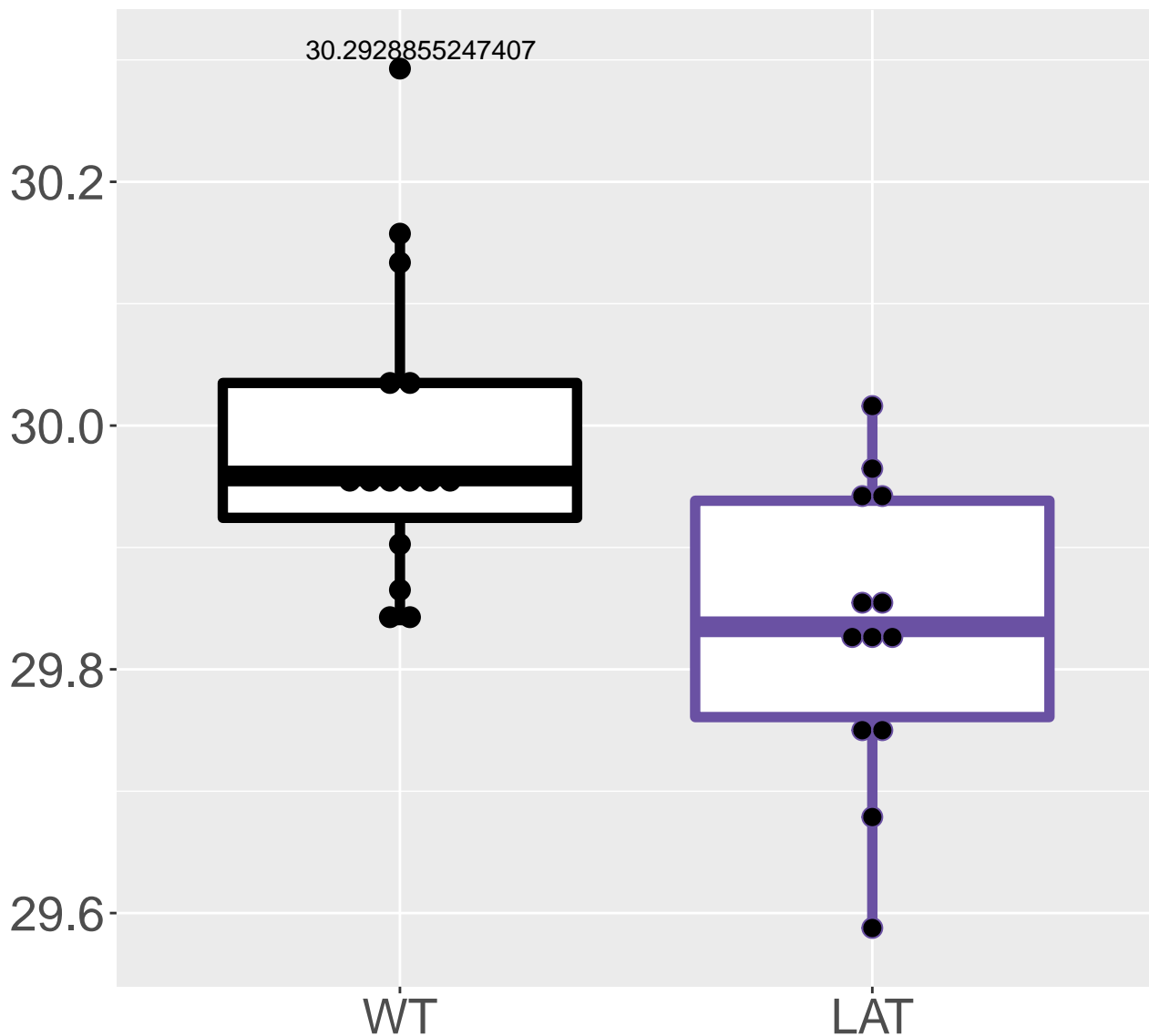
P56391_Cytochrome c oxidase sub.
FDR = 0.015, FC = -0.33, sex**



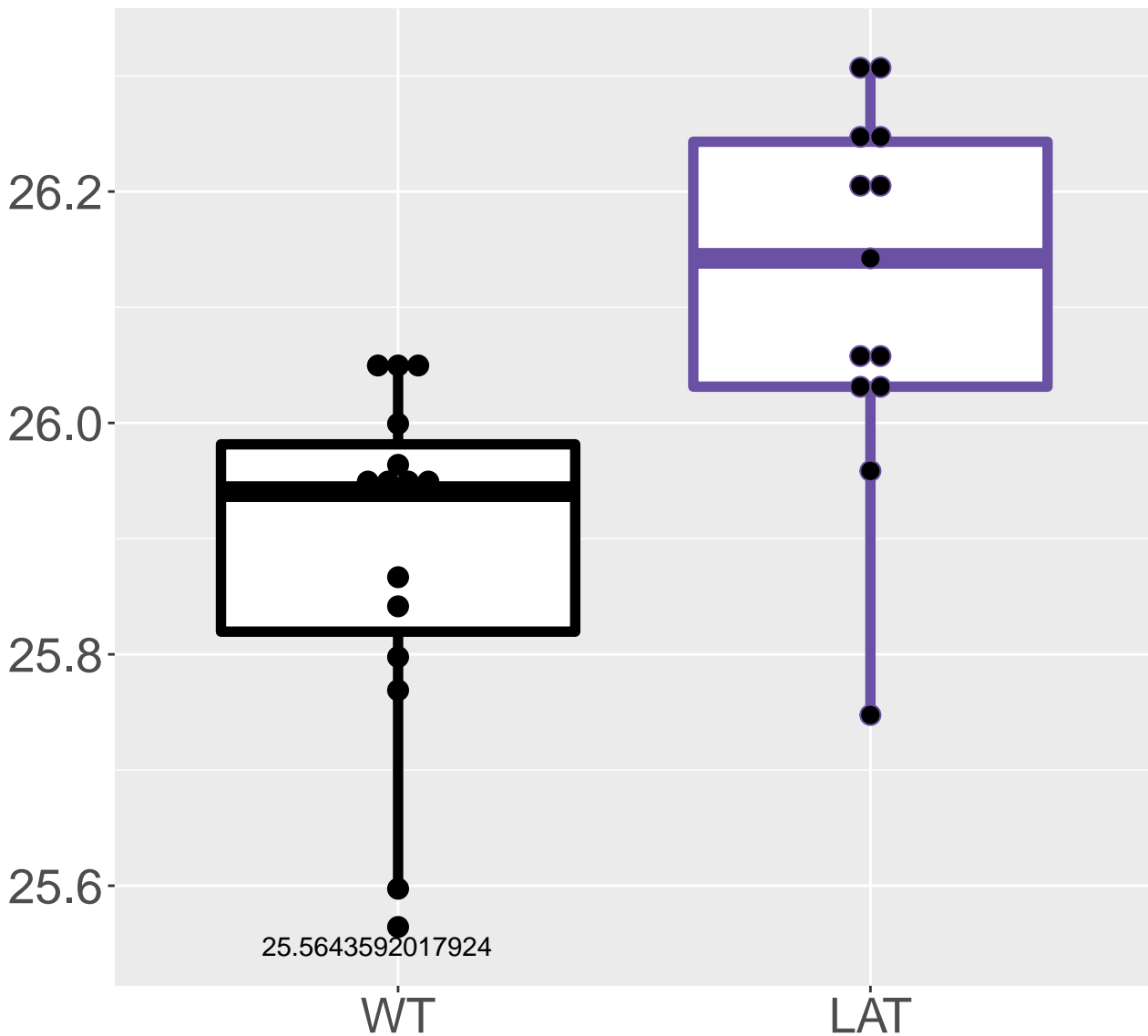
Q9D8Y0_EF-hand domain-containin.
FDR = 0.015, FC = -0.28, sex**



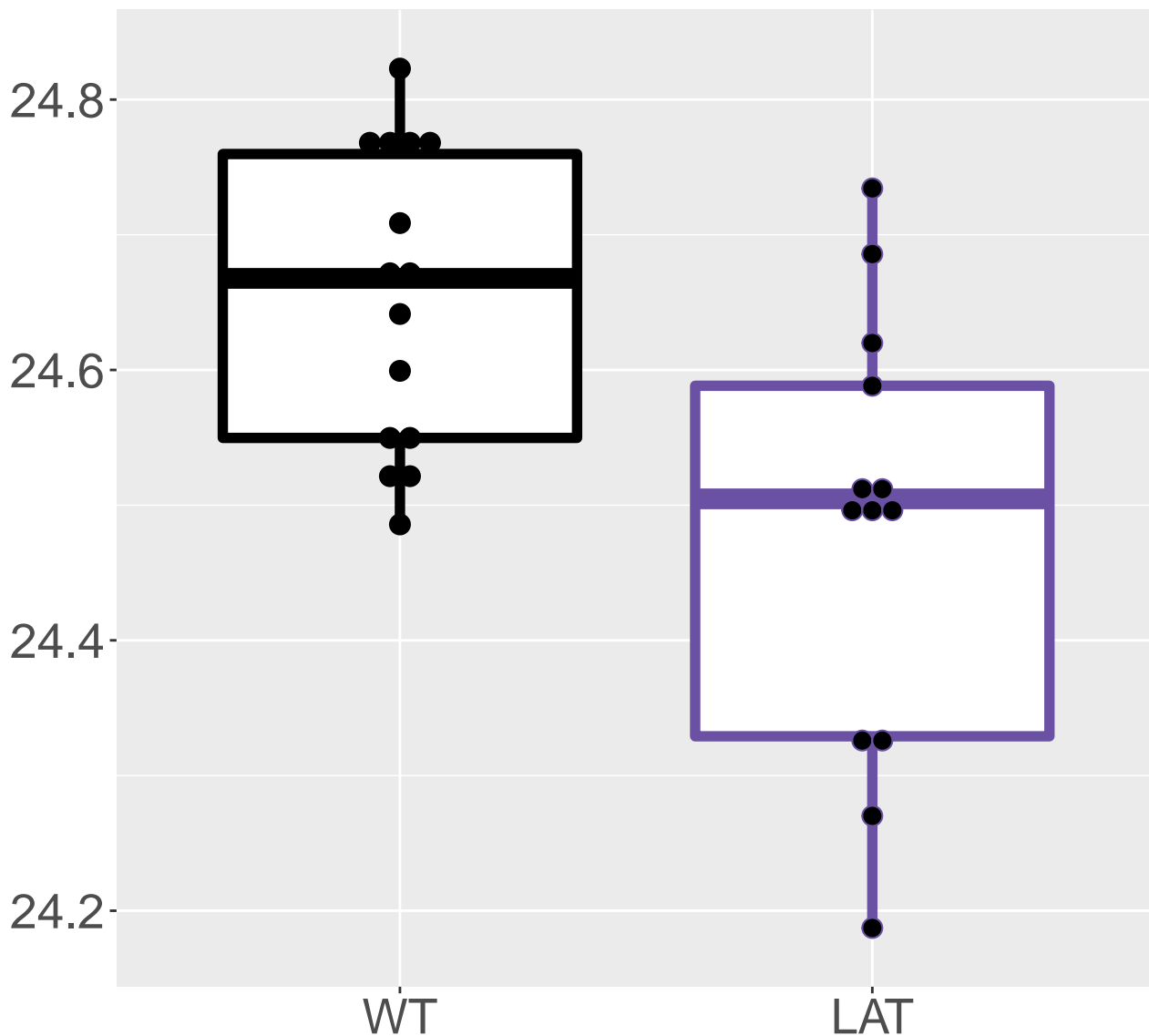
P08228_Superoxide dismutase [Cu.
FDR = 0.015, FC = -0.28



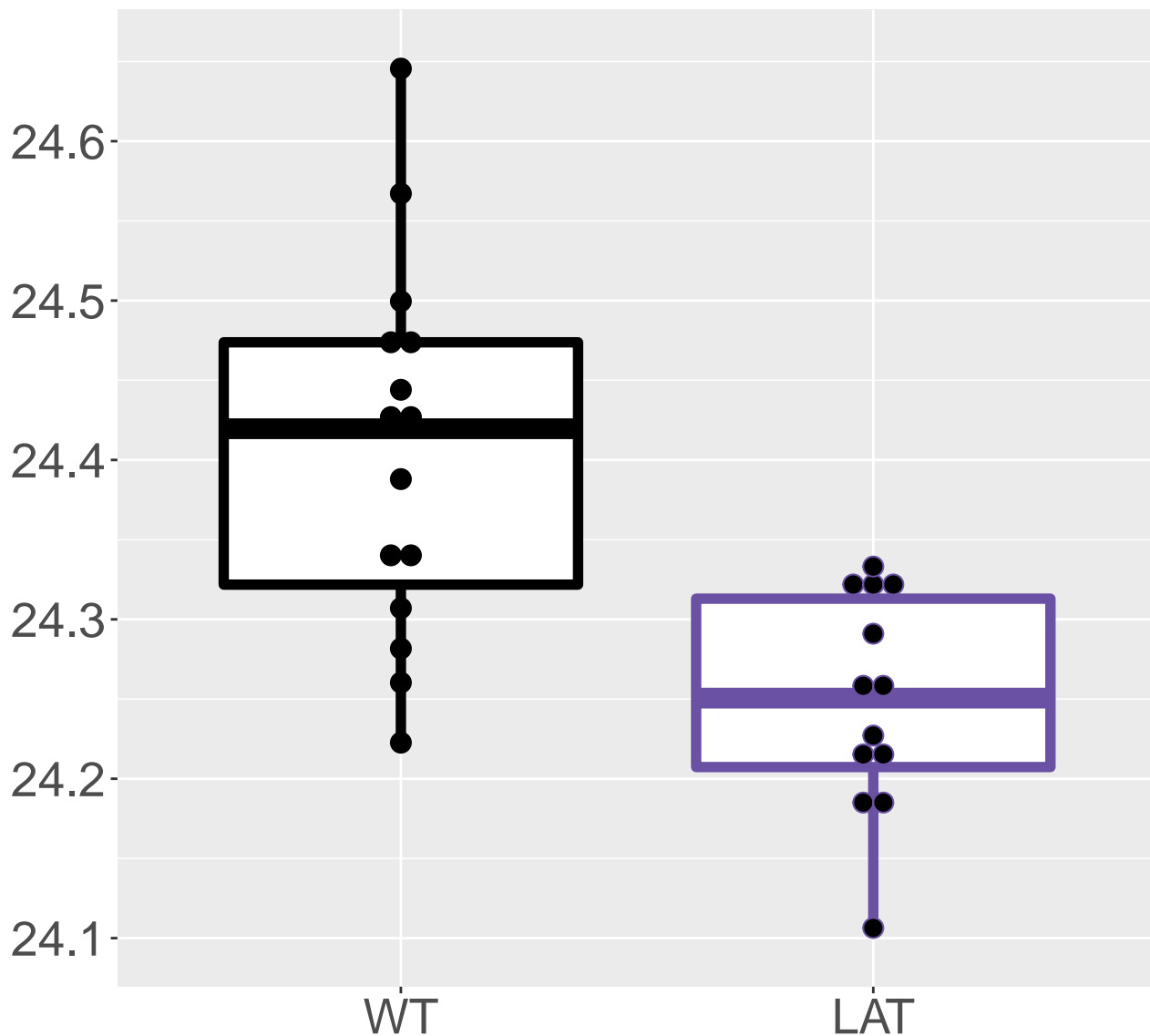
P97742_Carnitine O-palmitoyltra.
FDR = 0.016, FC = 0.36



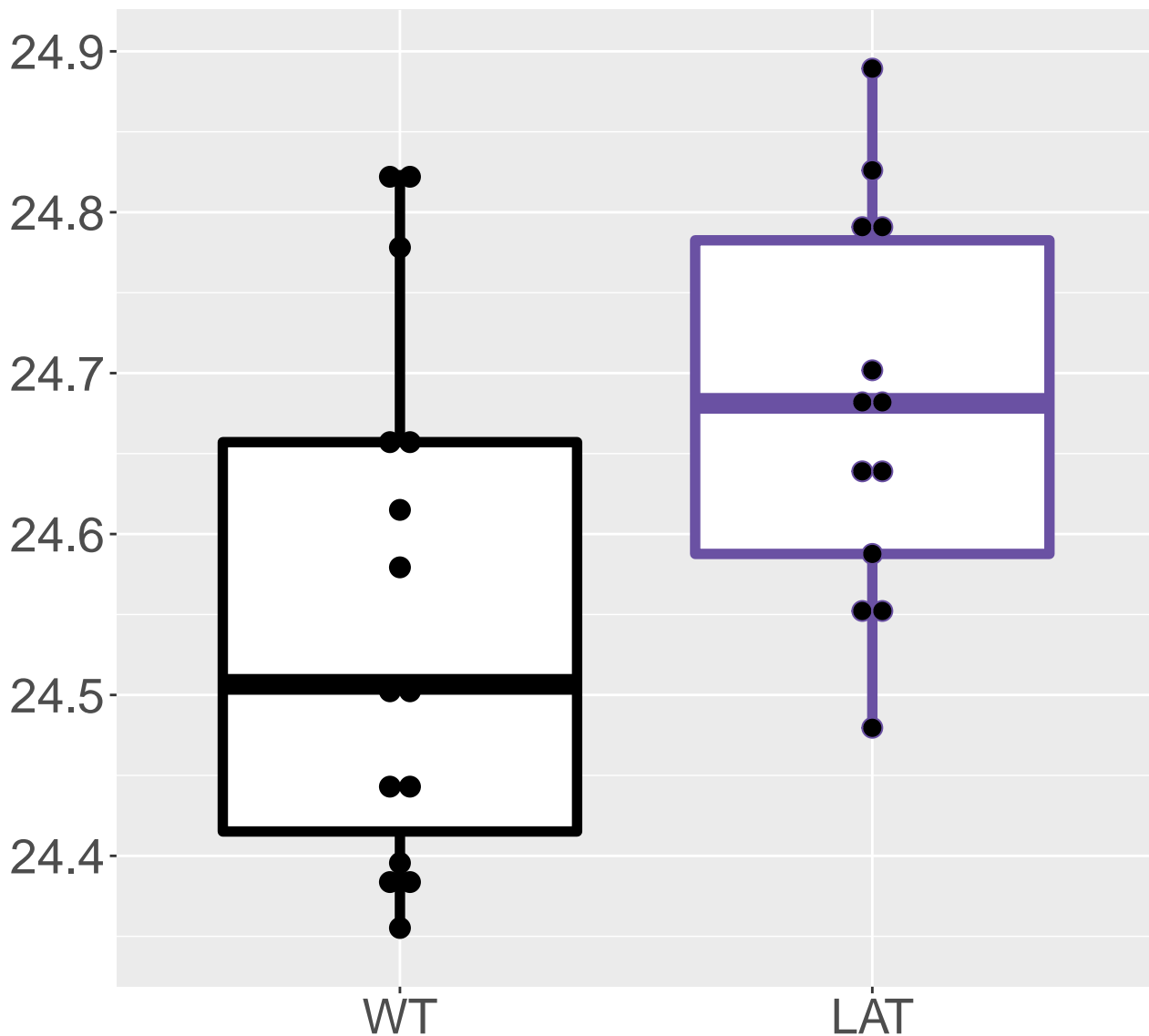
Q9DBB4_N-alpha-acetyltransferas.
FDR = 0.016, FC = -0.29, sex*



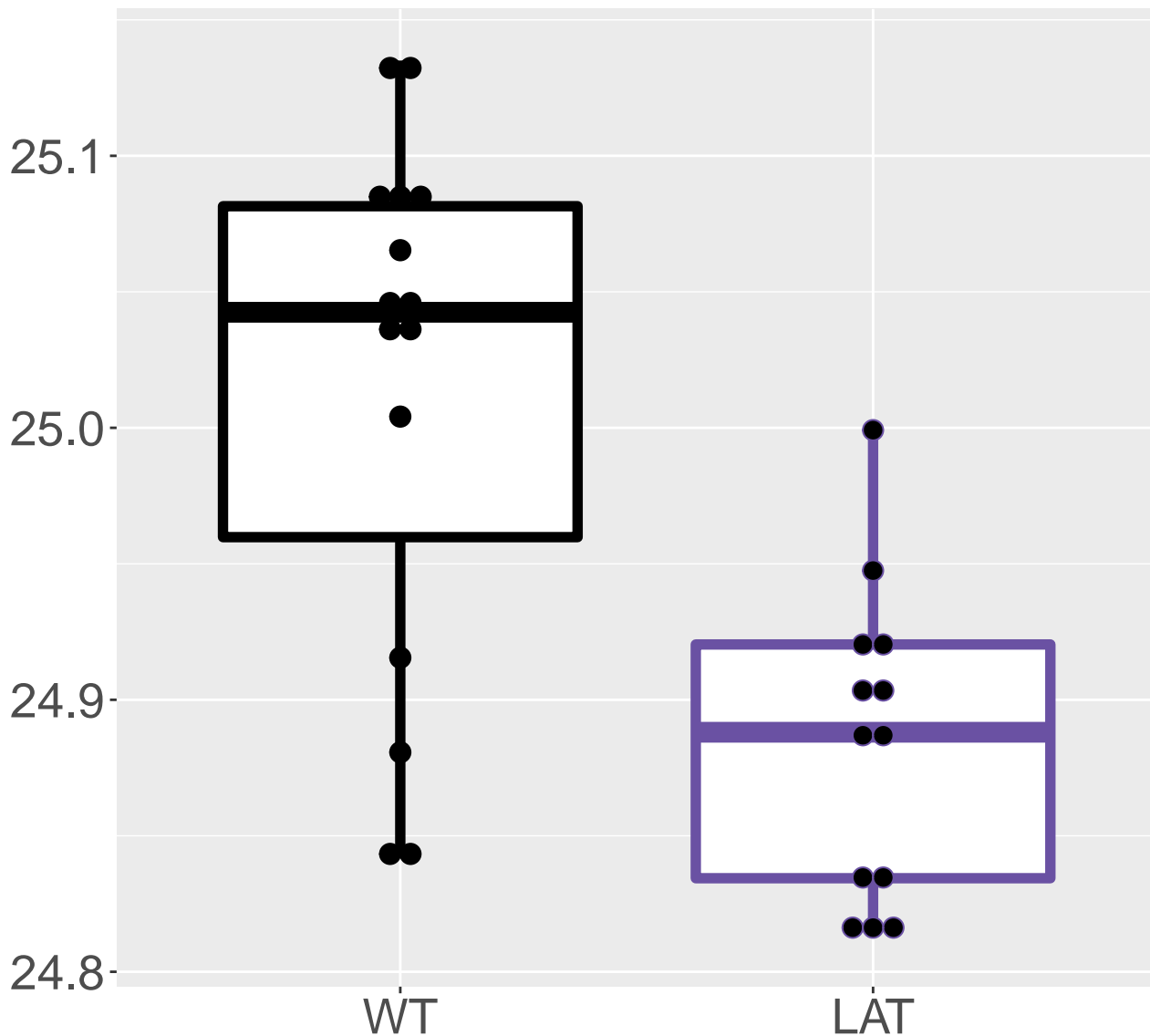
Q9QZD9_Eukaryotic translation i.
FDR = 0.016, FC = -0.21



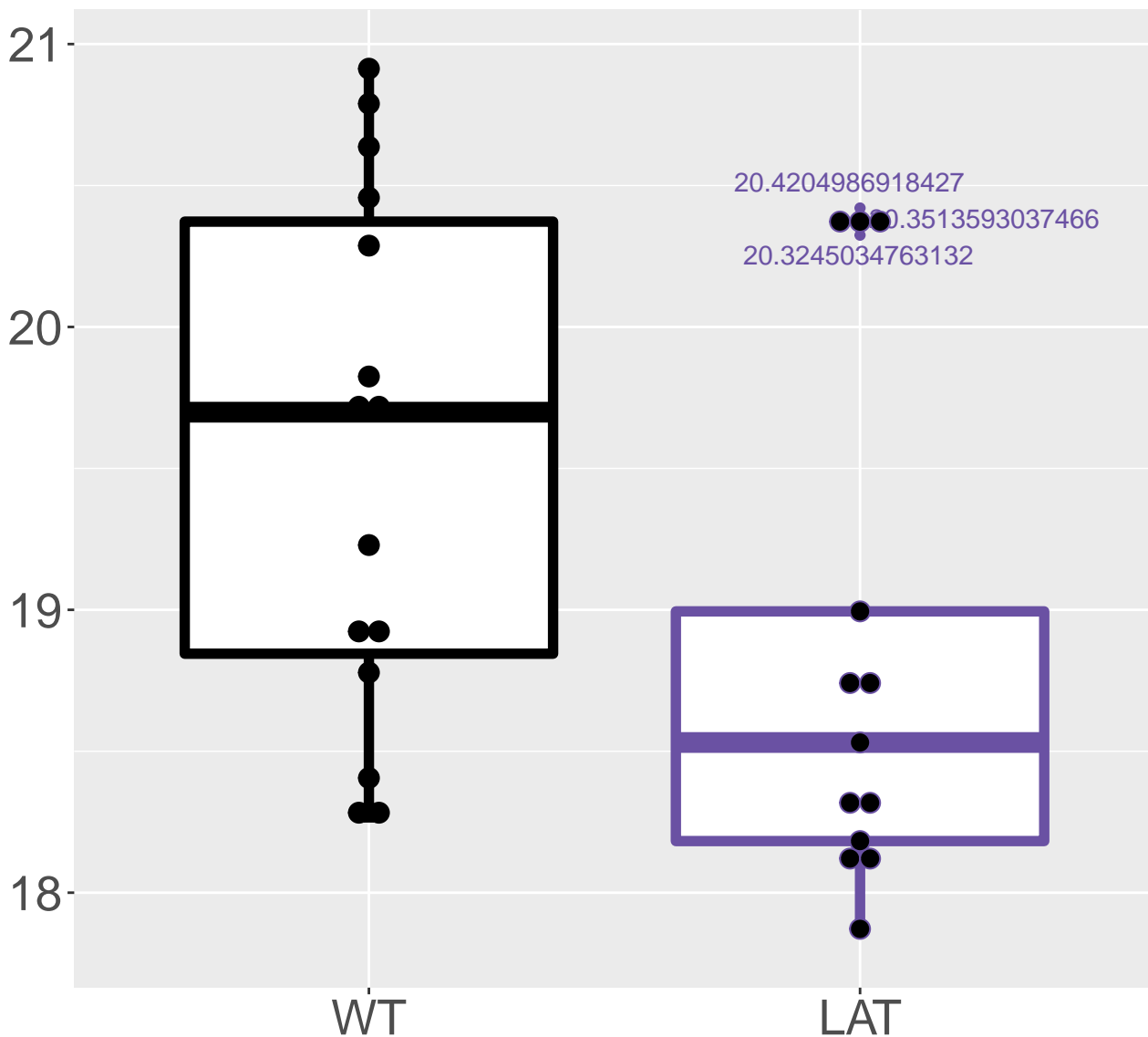
Q80W22_Threonine synthase-like 2
FDR = 0.016, FC = 0.19, sex***



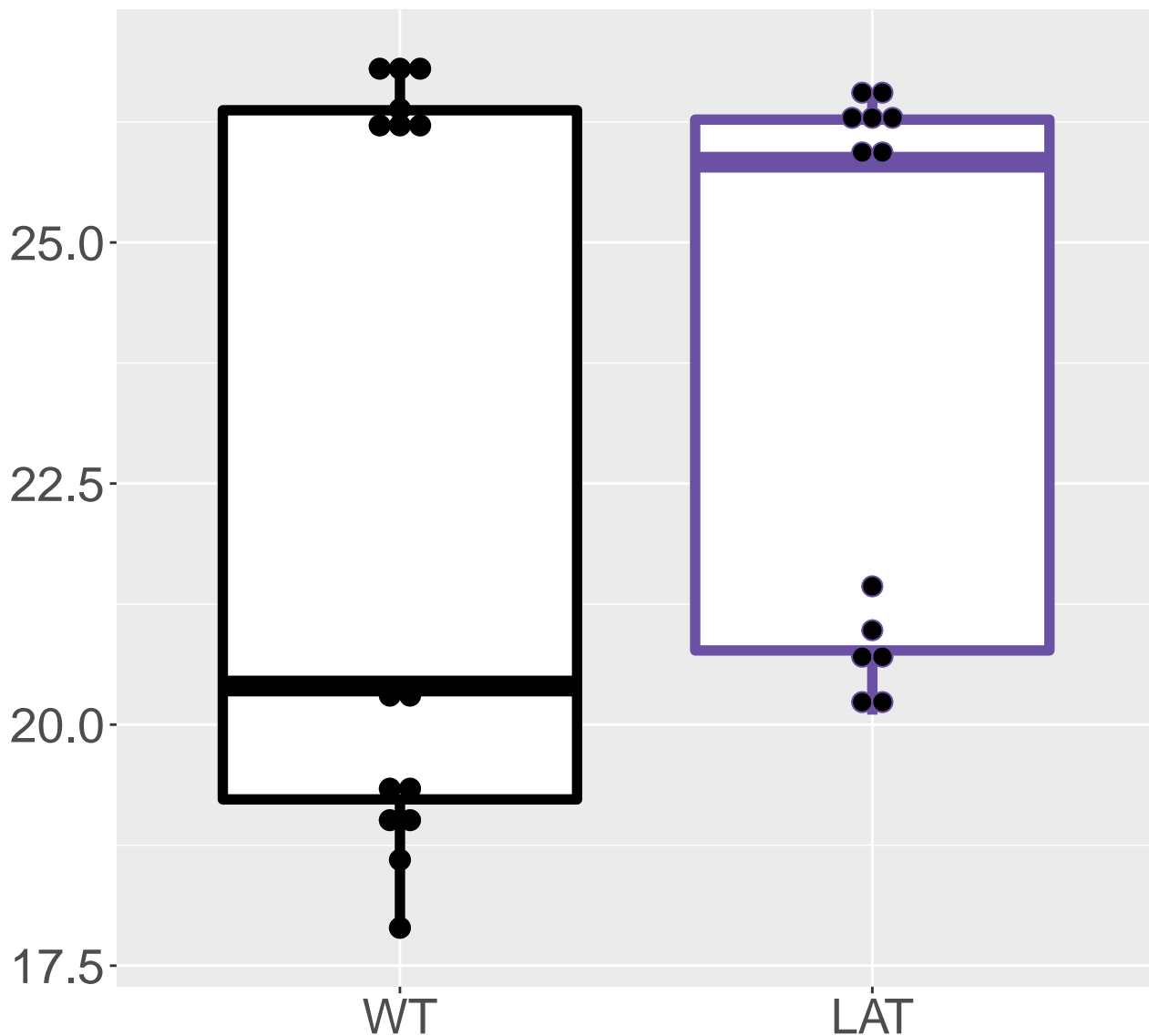
Q9D7B6_Isobutyryl-CoA dehydroge.
FDR = 0.016, FC = -0.18



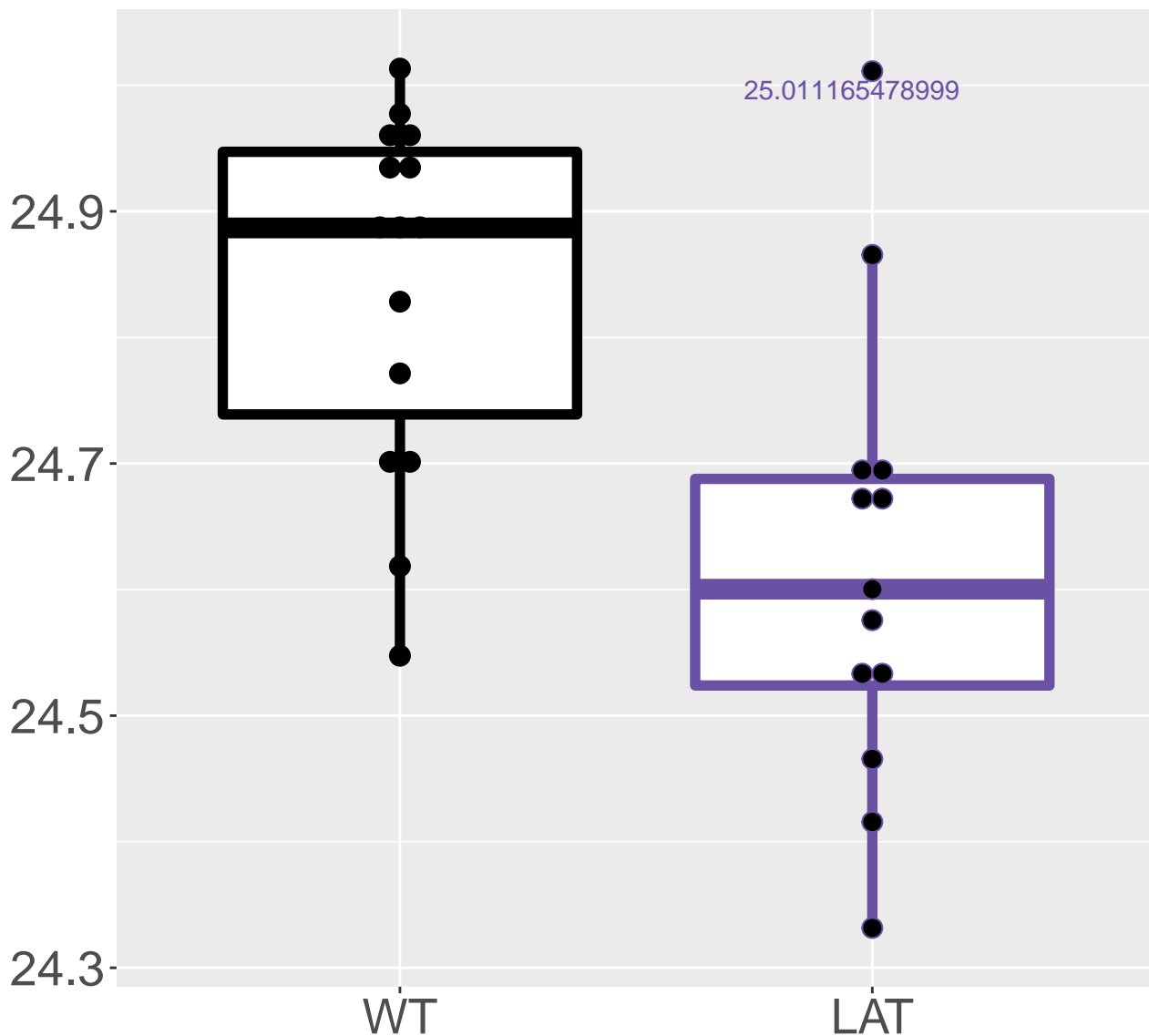
Q9WV98_Mitochondrial import inn.
FDR = 0.016, FC = -1.6, sex*



Q5FW60_Major urinary protein 20
FDR = 0.017, FC = 1.5, sex***

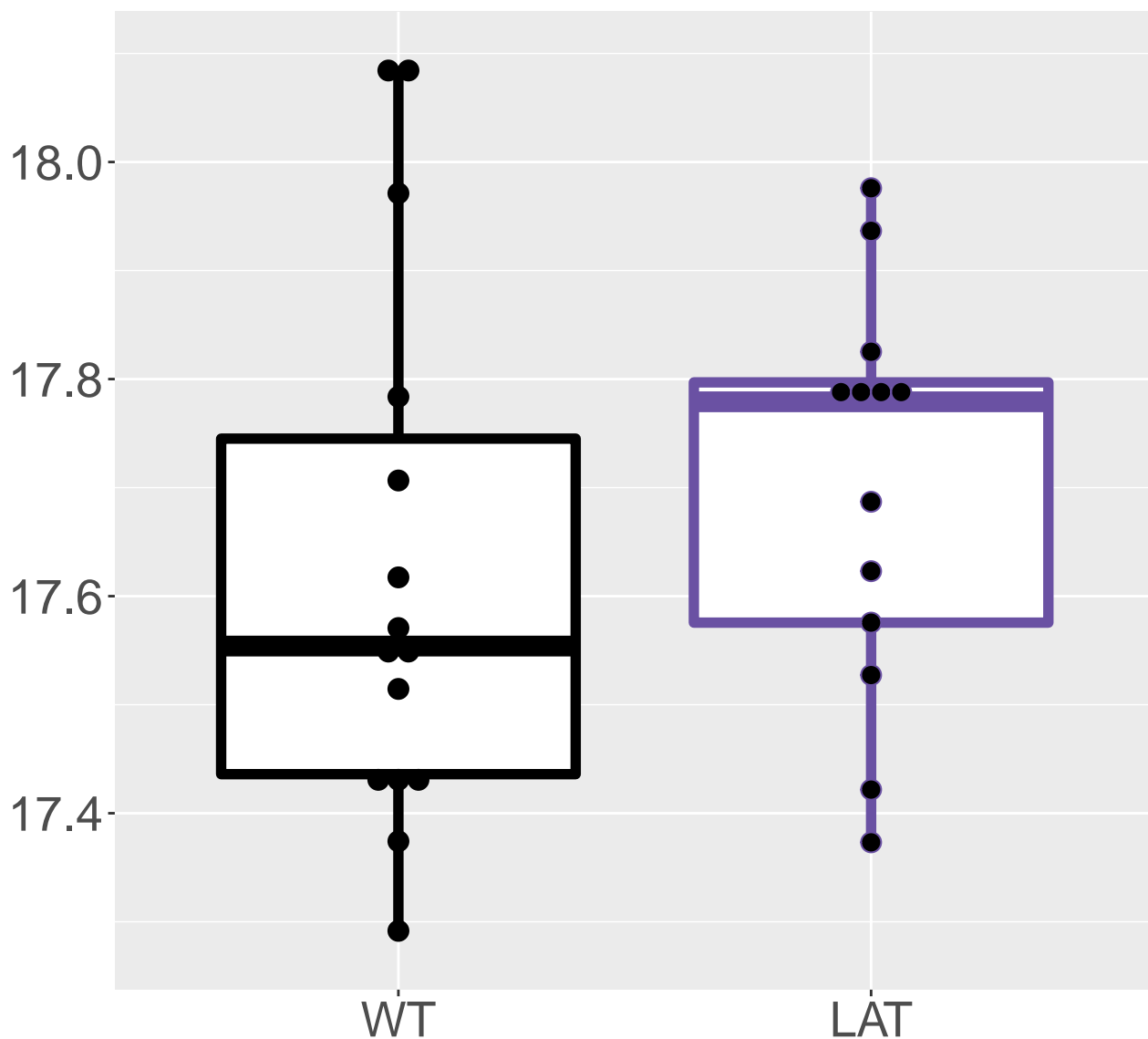


Q9JJI8_60S ribosomal protein L38
FDR = 0.017, FC = -0.34, sex*

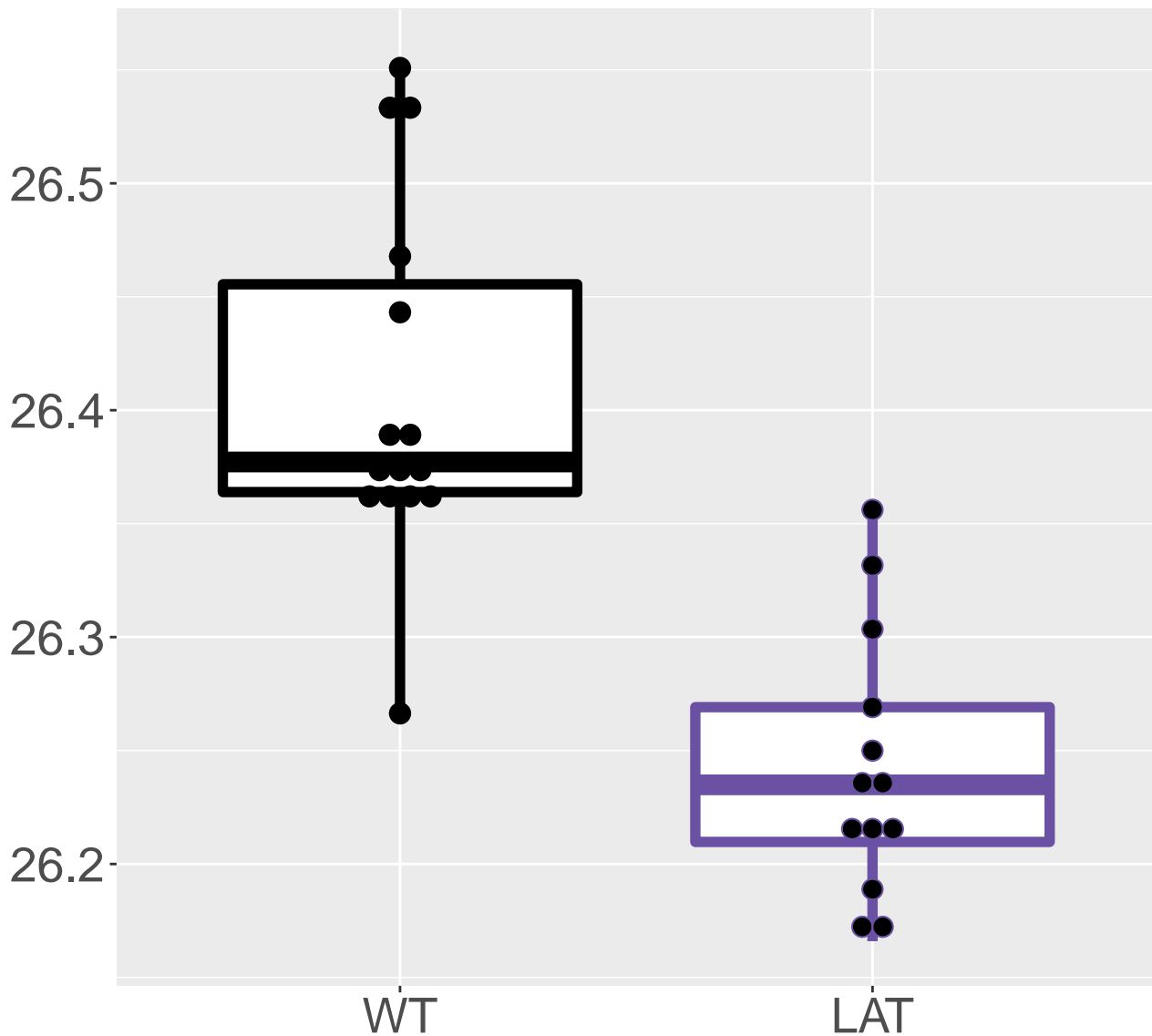


Q9JMF7_Dolichyldiphosphatase 1

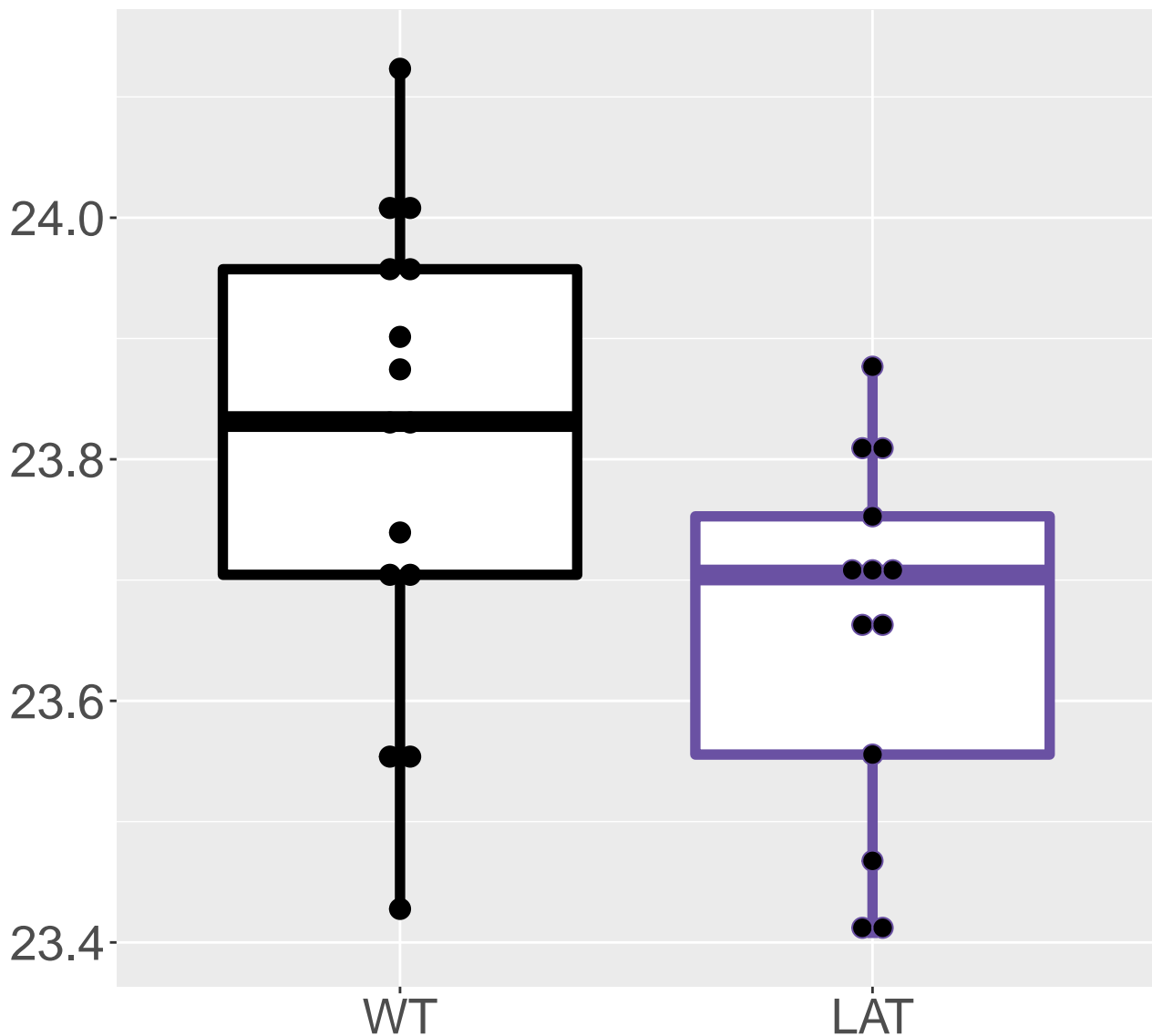
FDR = 0.017, FC = 0.3



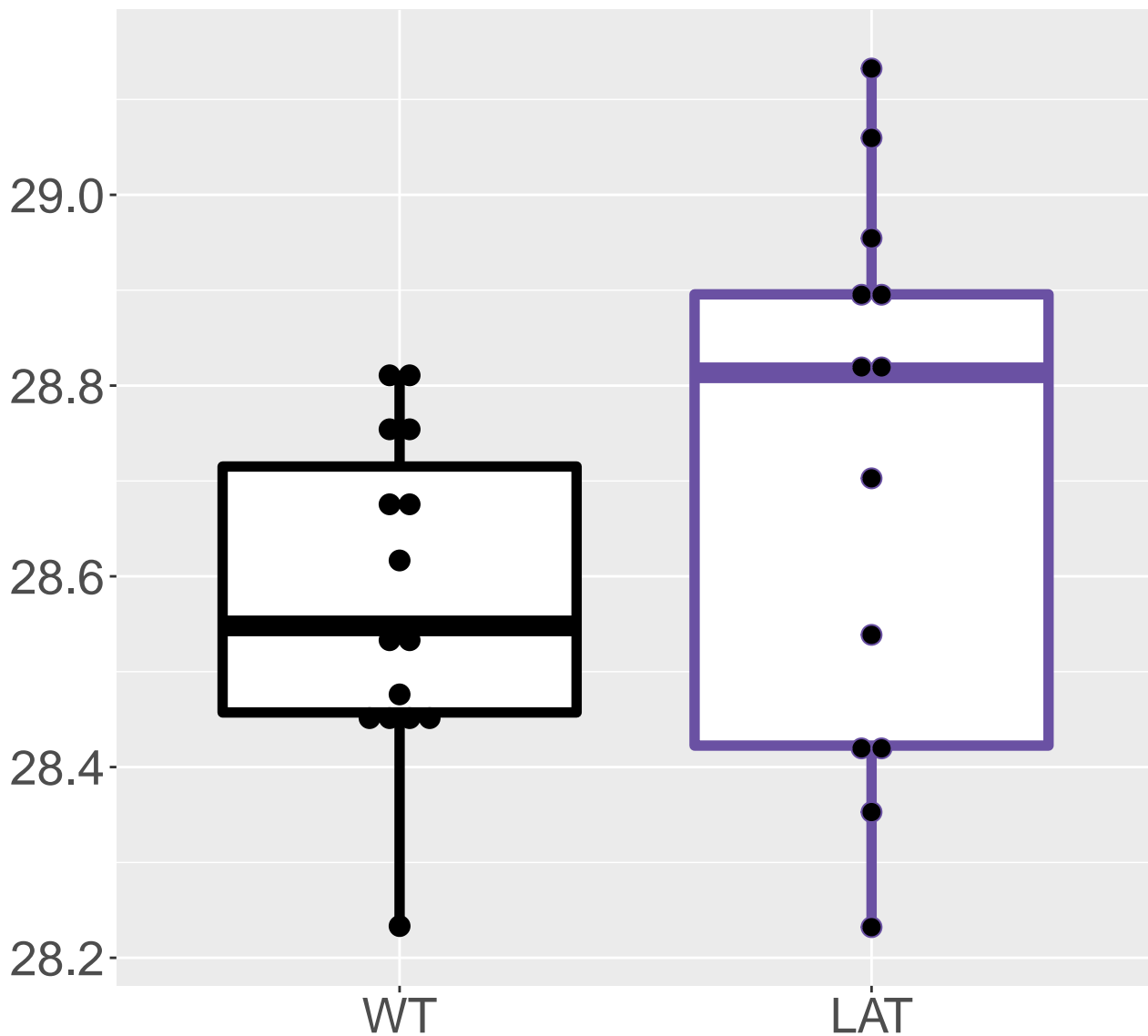
P35979_60S ribosomal protein L12
FDR = 0.017, FC = -0.17



Q9JM76_Actin-related protein 2/.
FDR = 0.017, FC = -0.33, sex*

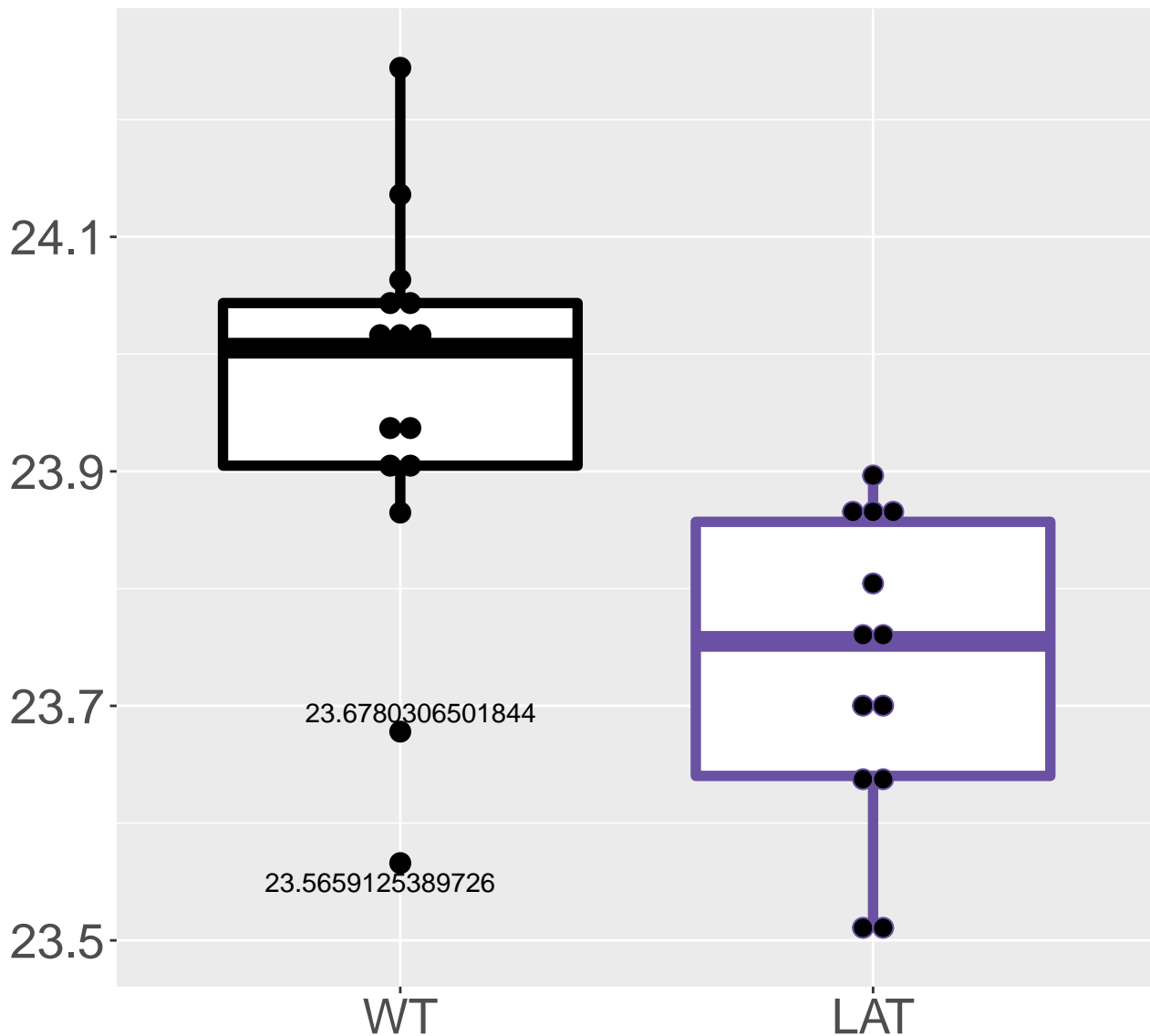


P08113_Endoplasmin
FDR = 0.017, FC = 0.38, sex*



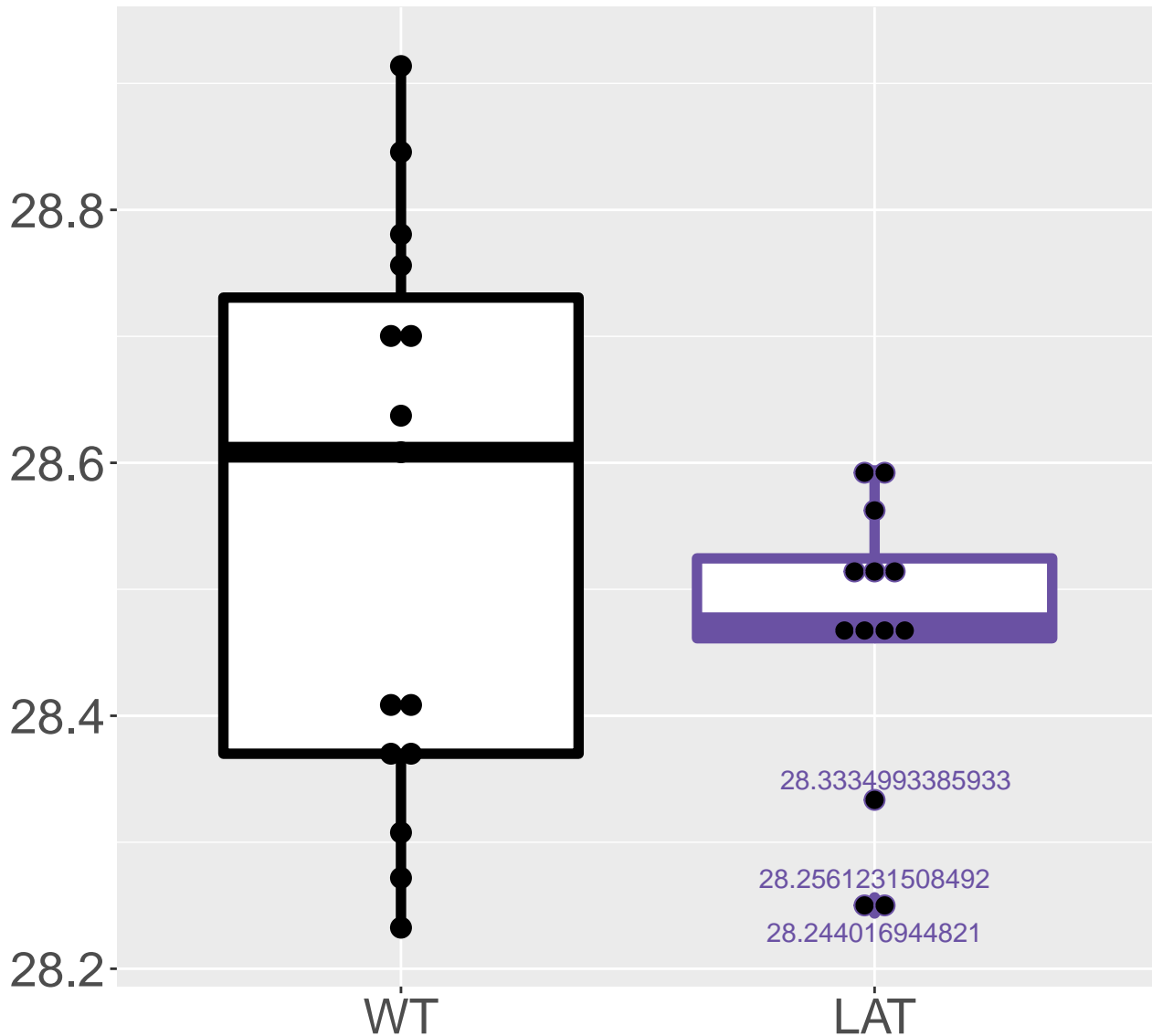
Q9QXT0_Protein canopy homolog 2

FDR = 0.017, FC = -0.25

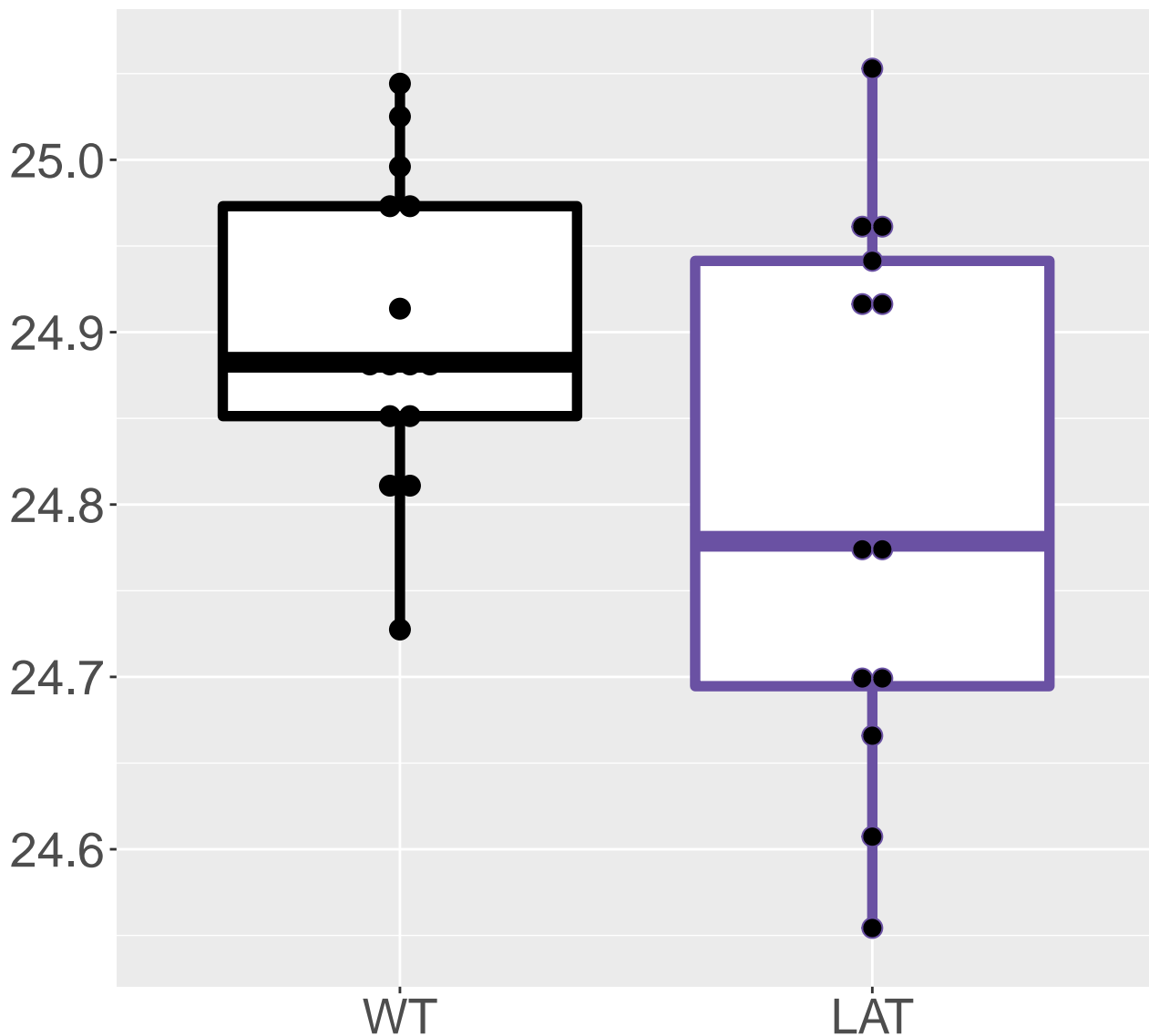


P11352_Glutathione peroxidase 1

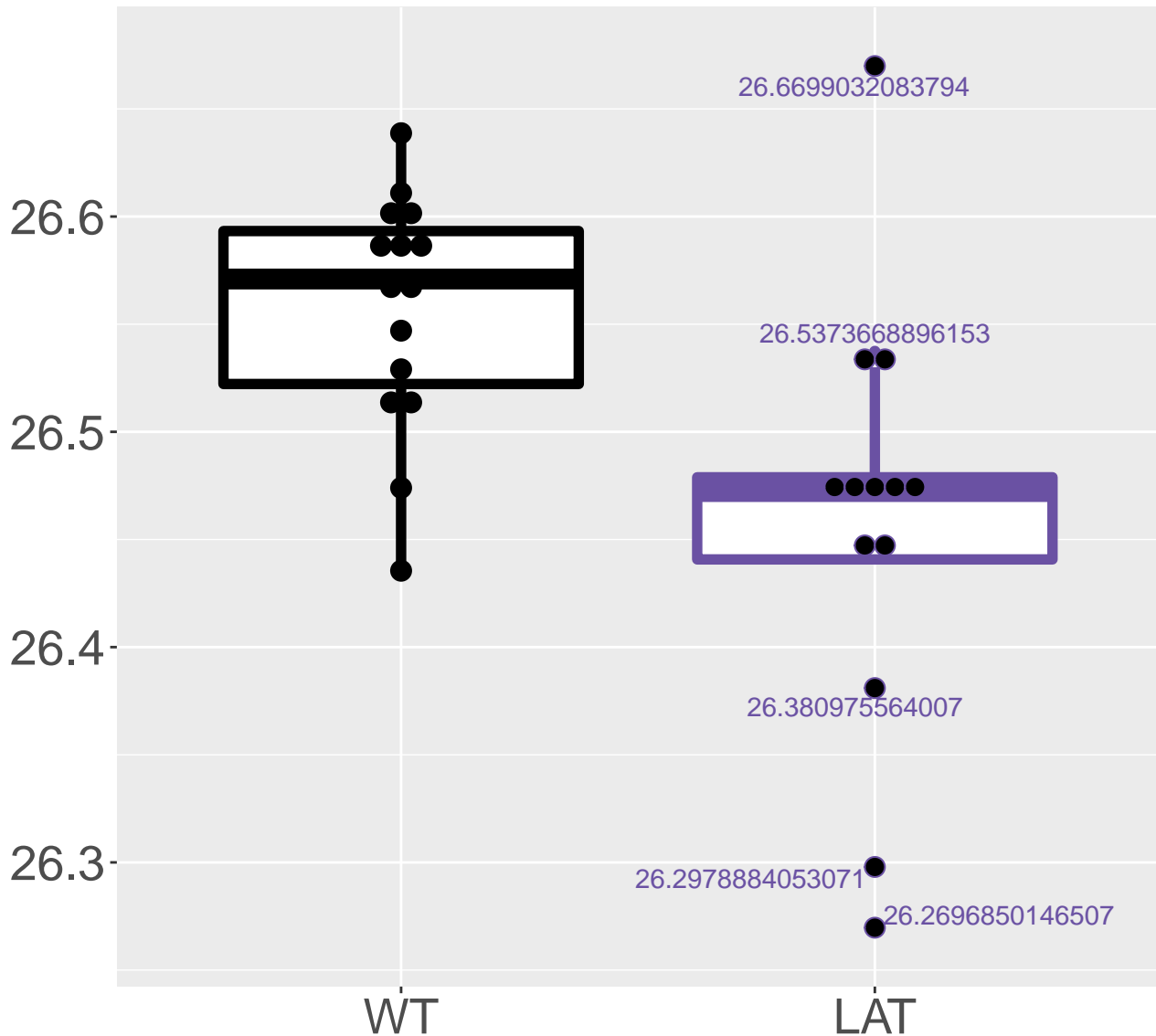
FDR = 0.017, FC = -0.22, sex***



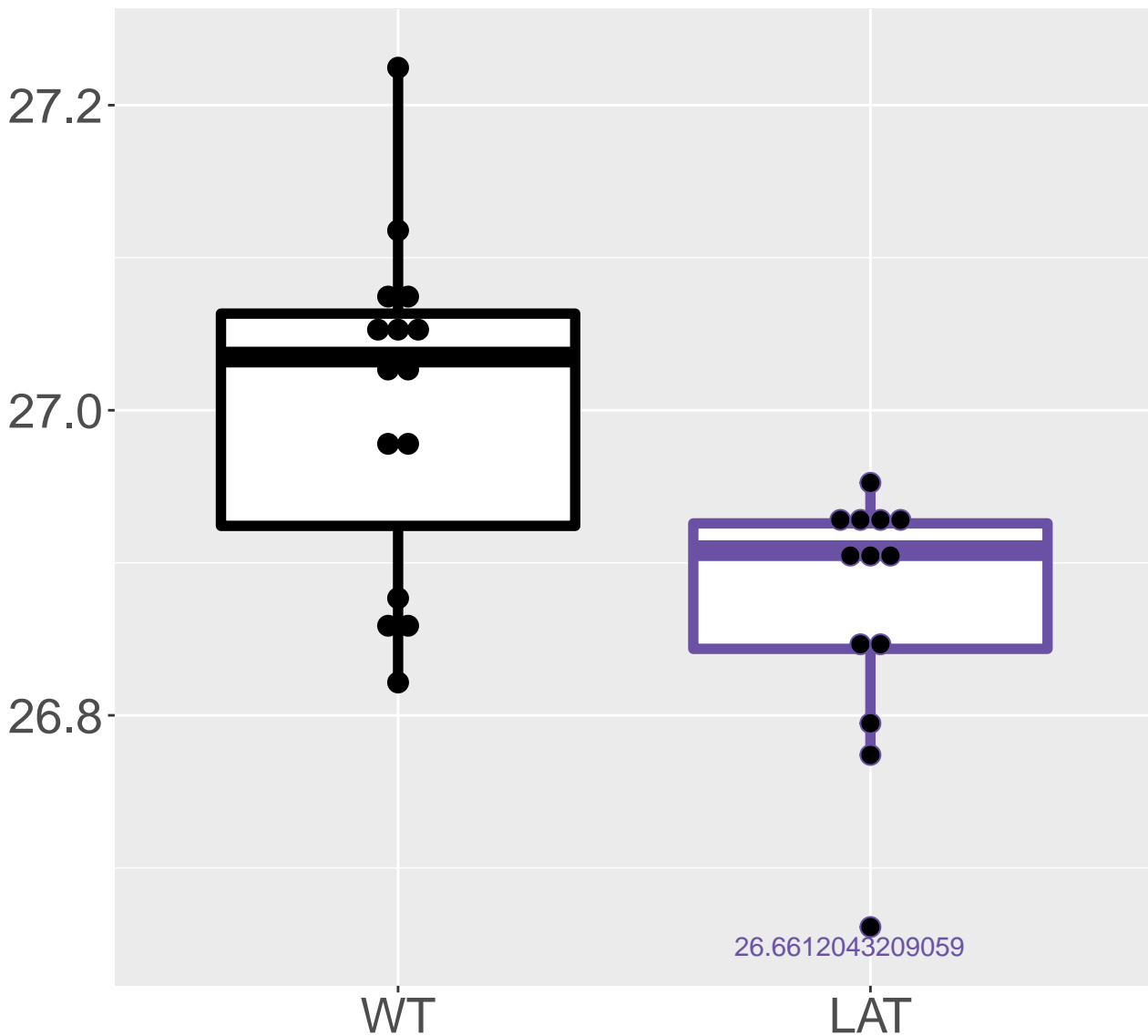
Q9DCJ5_NADH dehydrogenase [ubiq.
FDR = 0.017, FC = -0.18, sex**



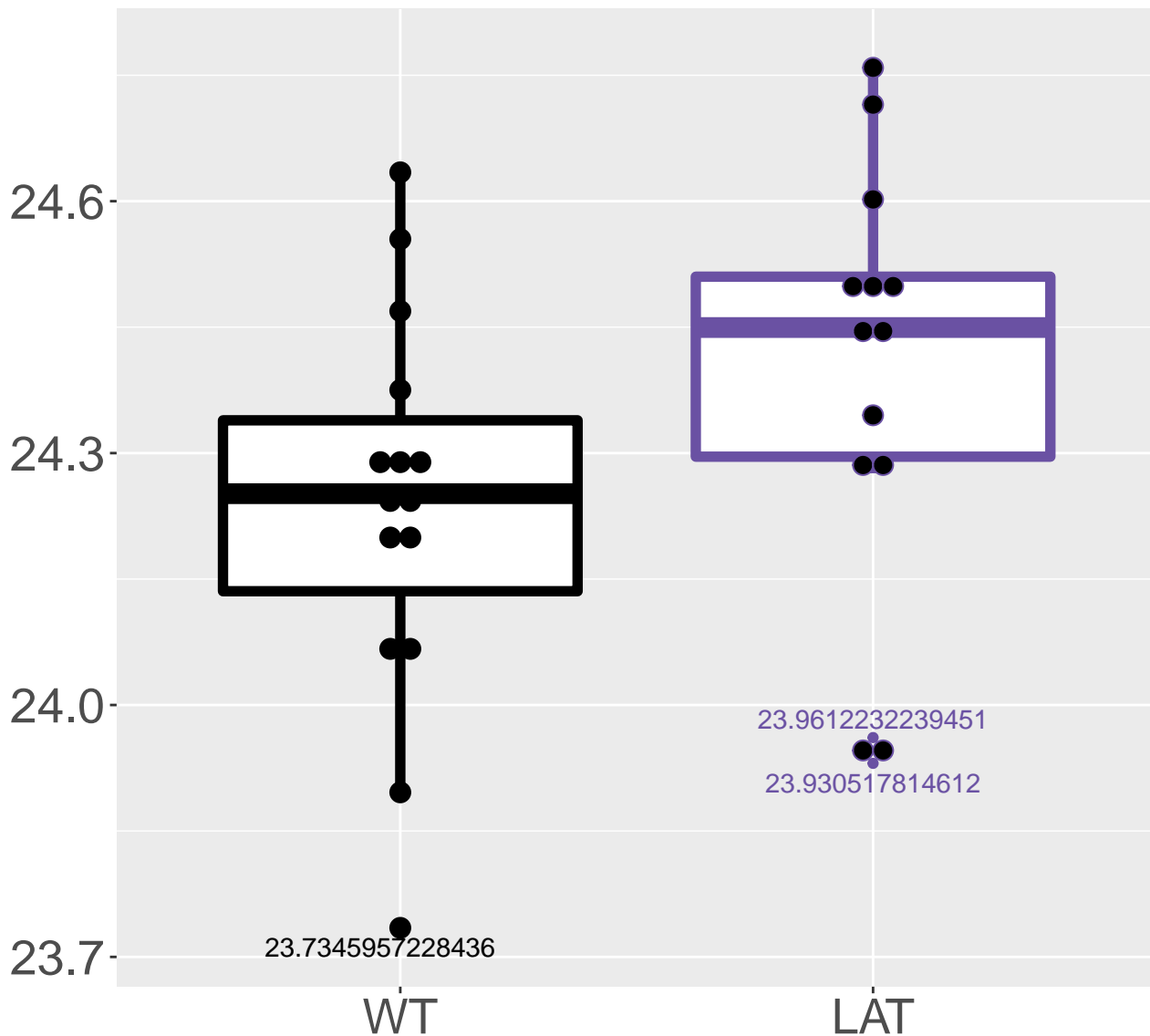
Q9R1P4_Proteasome subunit alpha.
FDR = 0.018, FC = -0.18



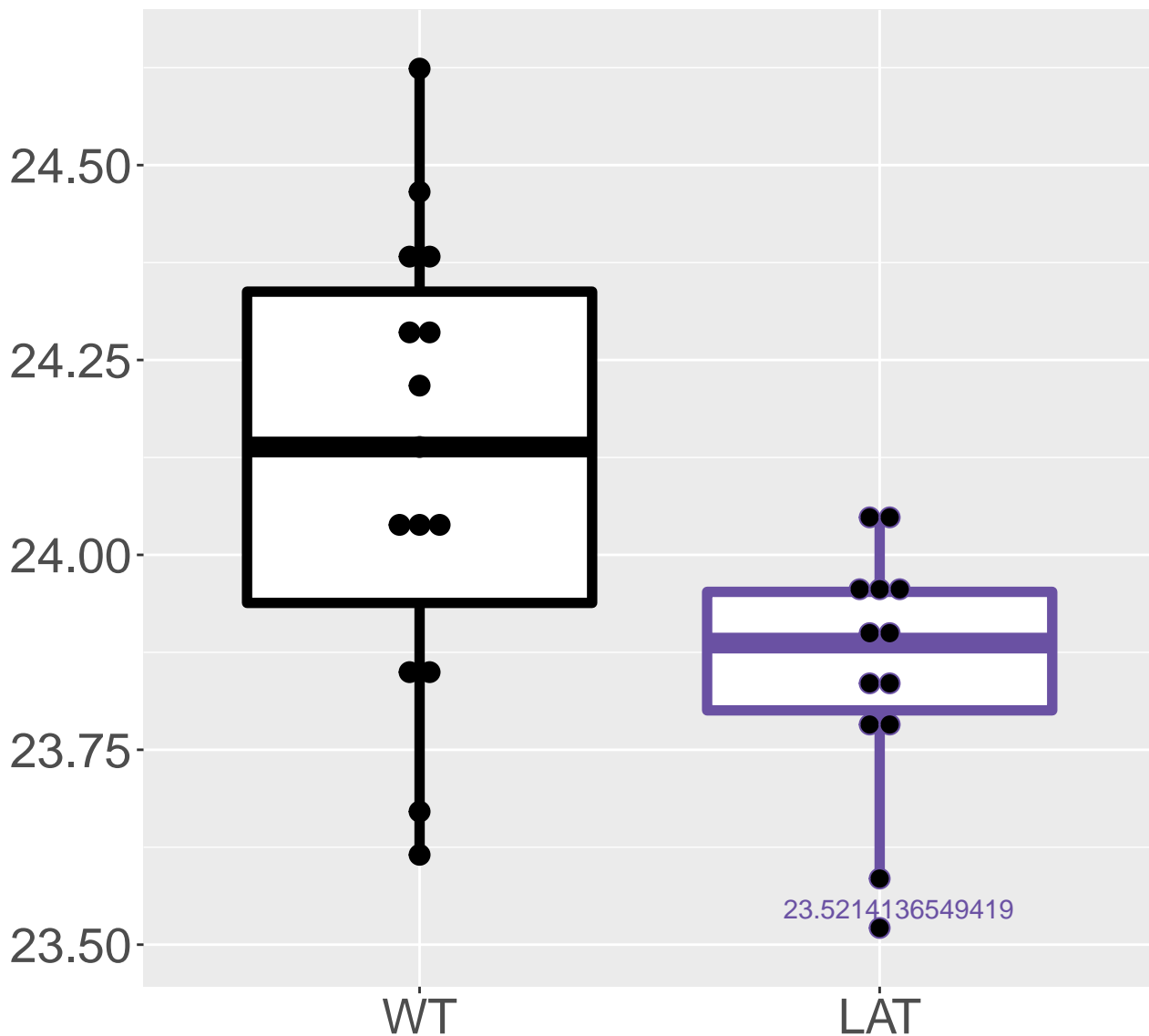
P62259_14-3-3 protein epsilon
FDR = 0.018, FC = -0.18, sex*



Q6ZQ38_Cullin-associated NEDD8-.
FDR = 0.019, FC = 0.39, sex**

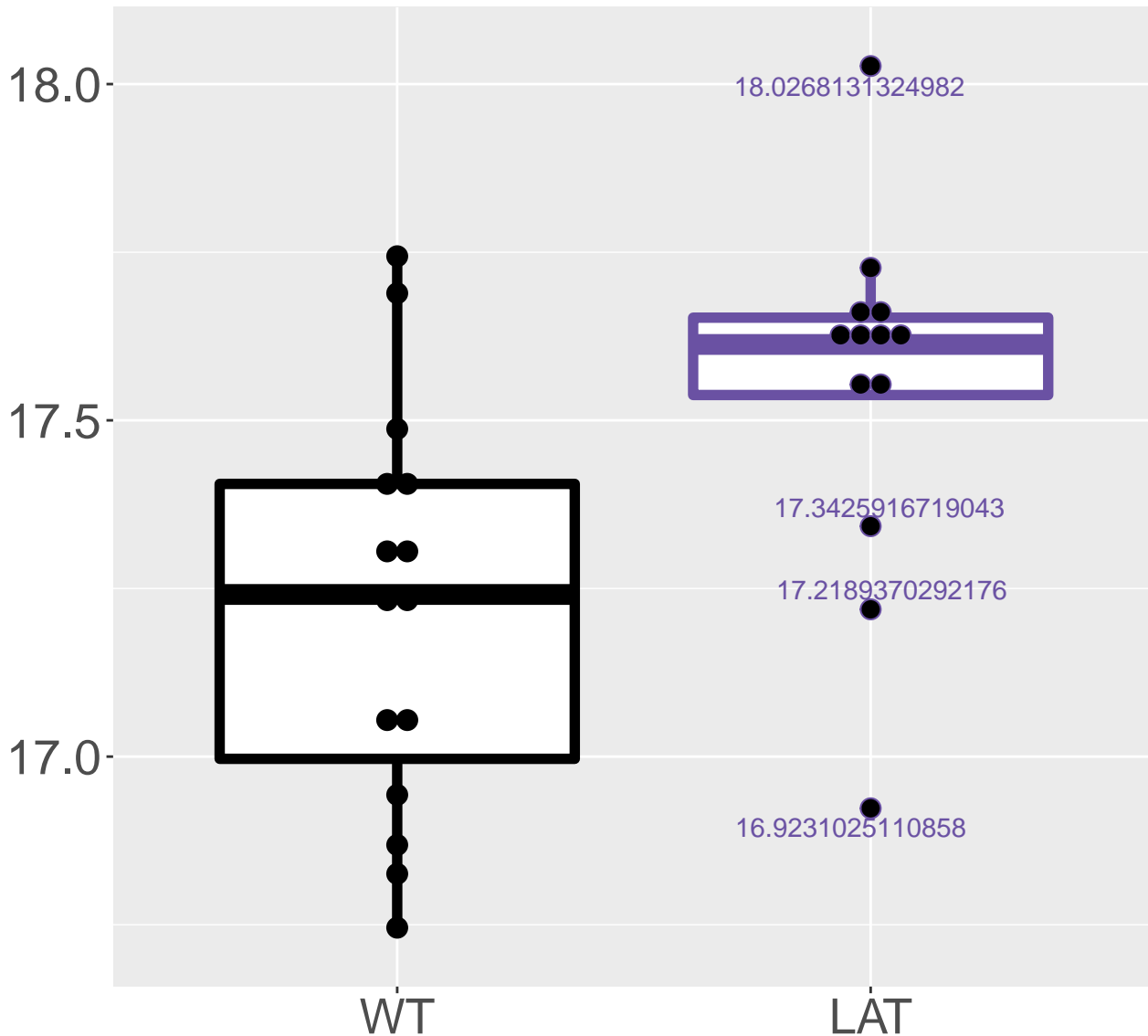


P56379_6.8 kDa mitochondrial pr.
FDR = 0.019, FC = -0.52, sex*

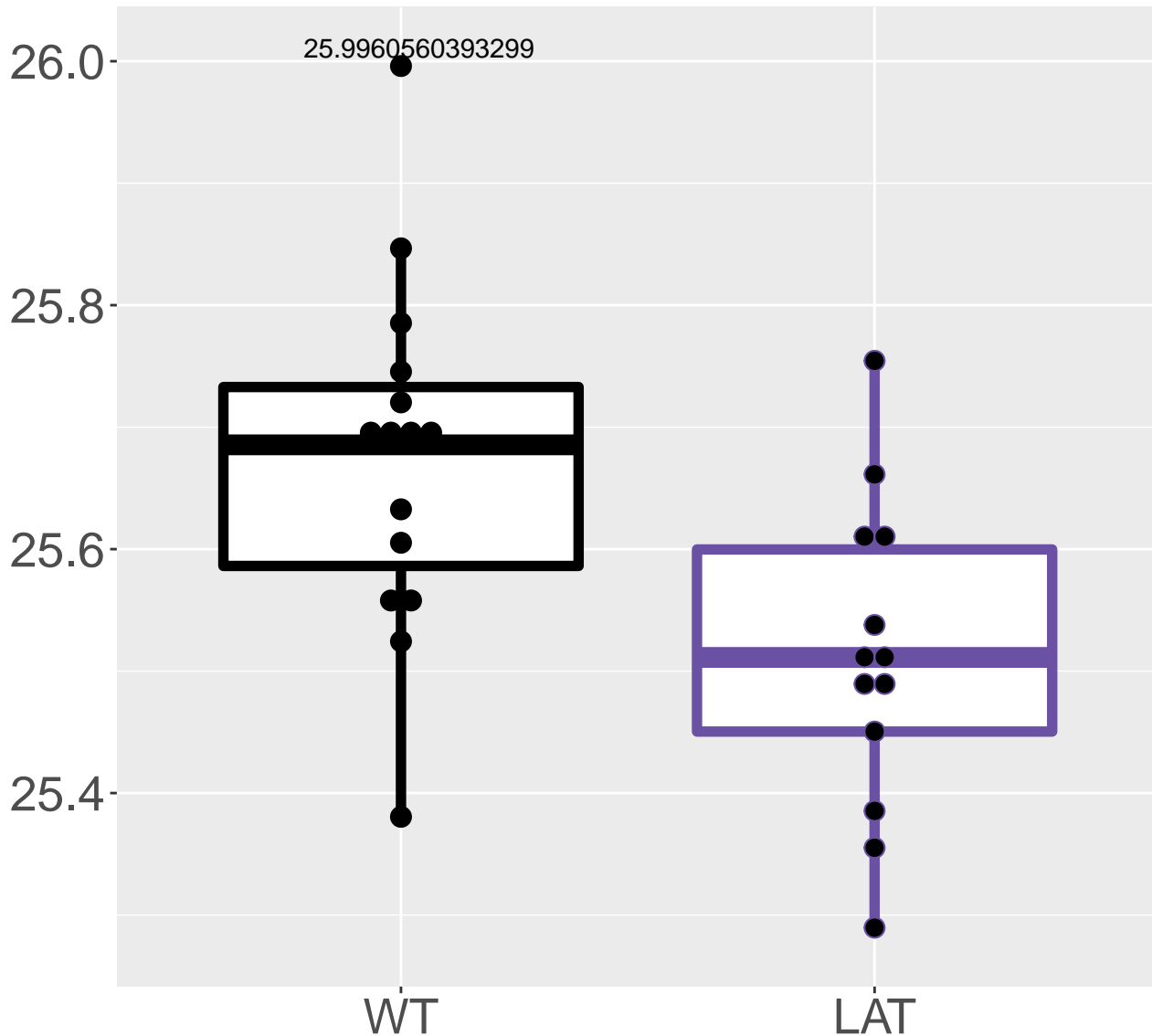


Q7TSH2_Phosphorylase b kinase r.

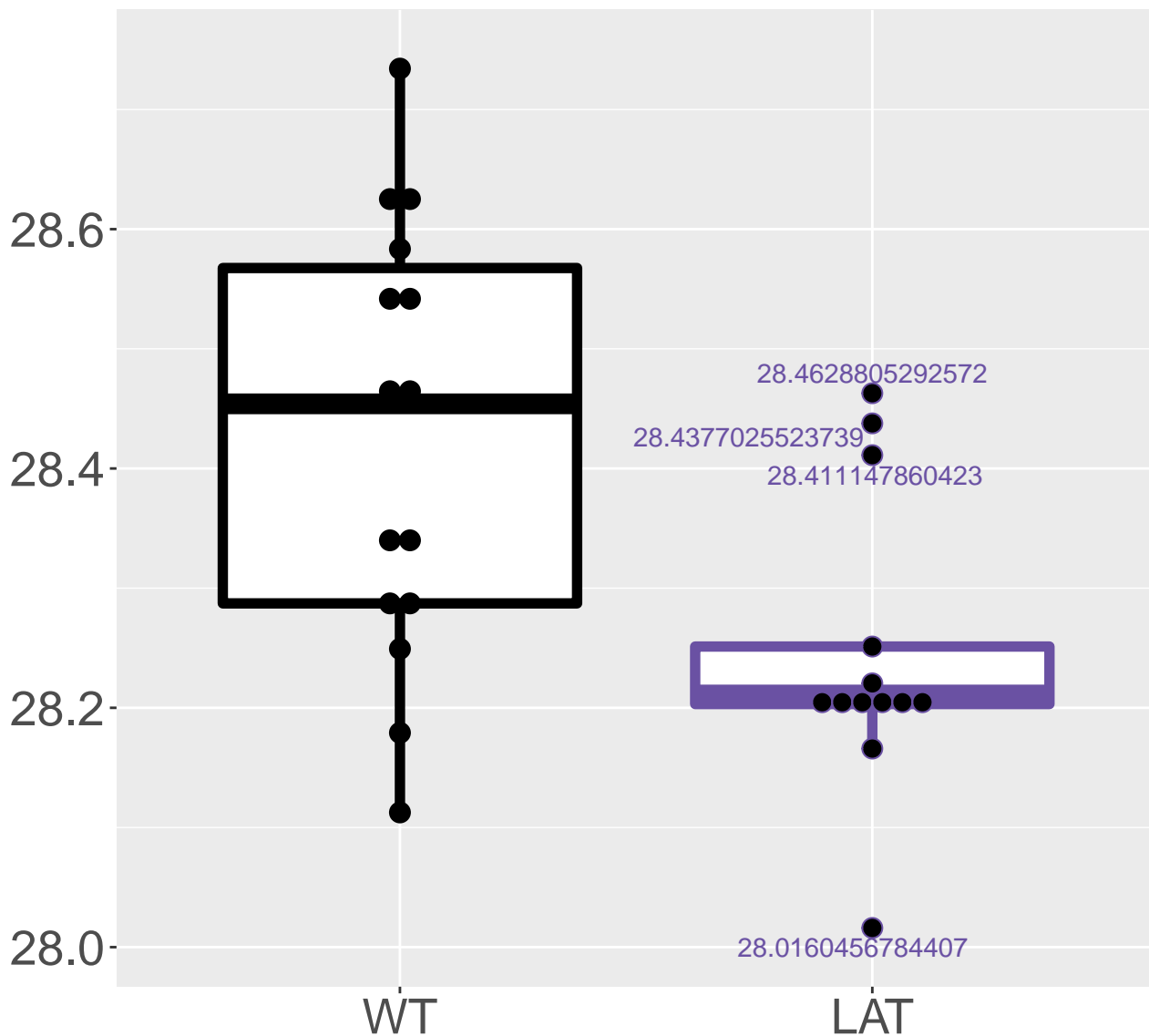
FDR = 0.02, FC = 0.54, sex*



P84099_60S ribosomal protein L19
FDR = 0.02, FC = -0.24

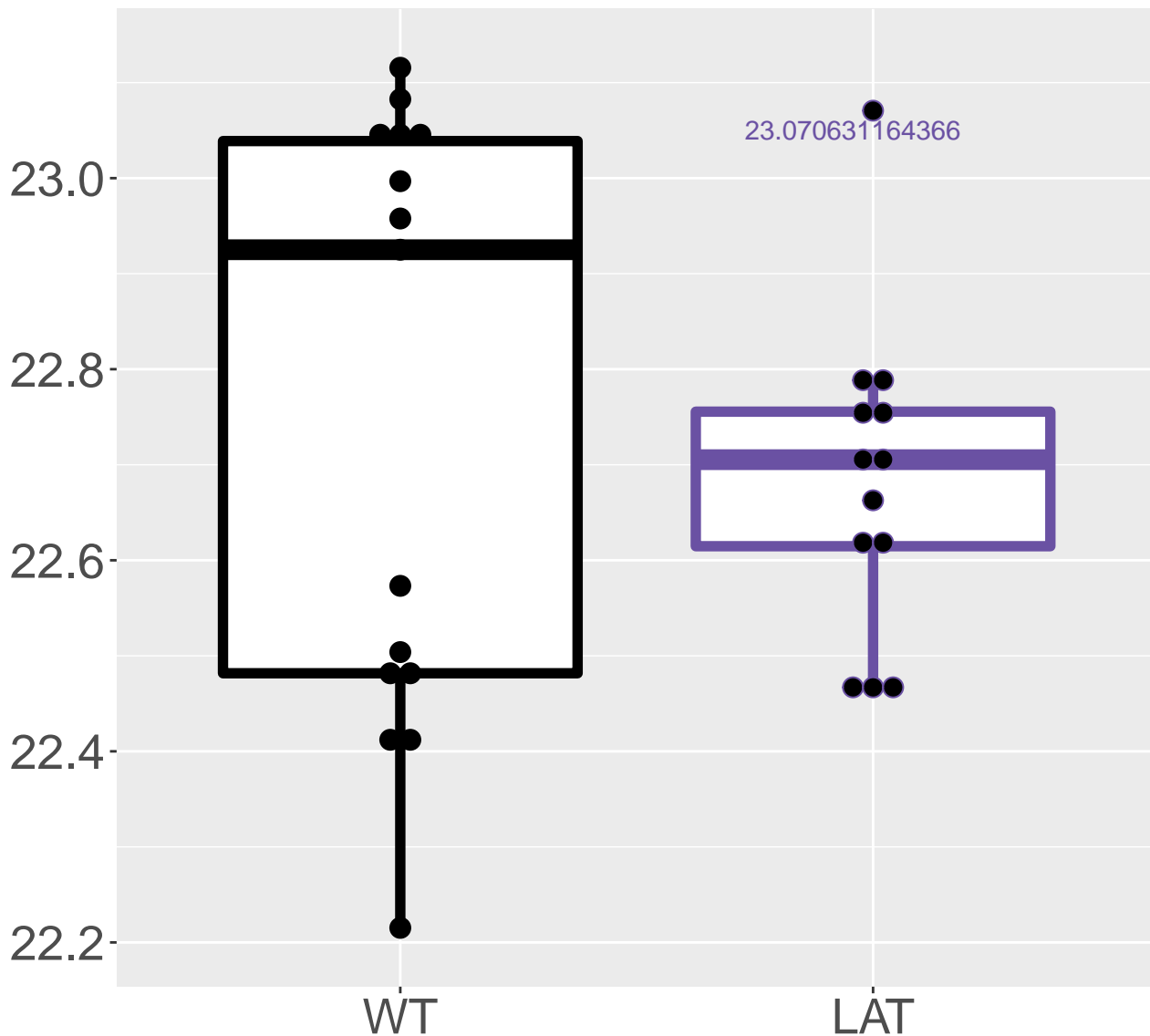


P10854_Histone H2B type 1-M
FDR = 0.02, FC = -0.25, sex***

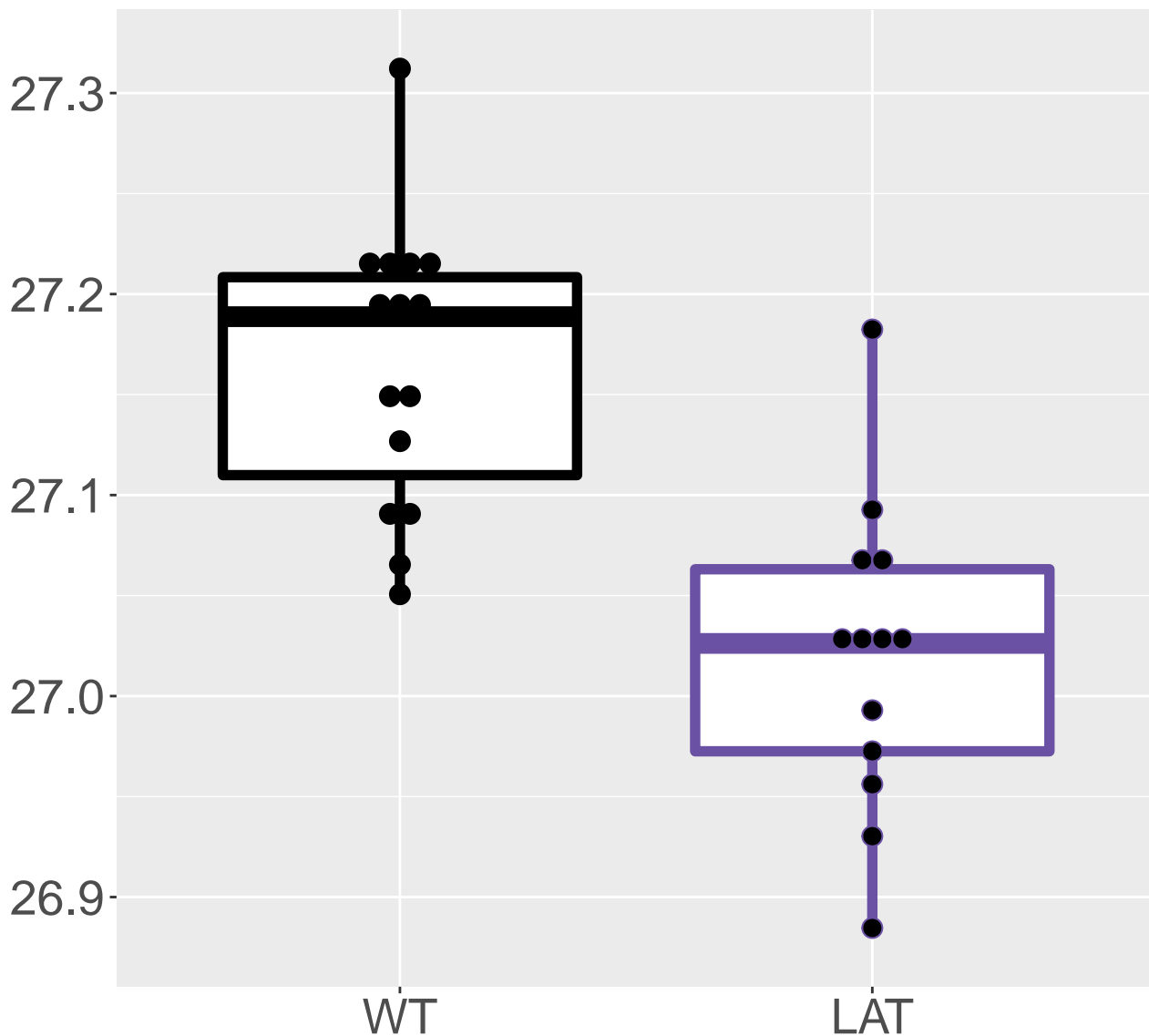


Q9Z2G9_Oxidoreductase HTATIP2

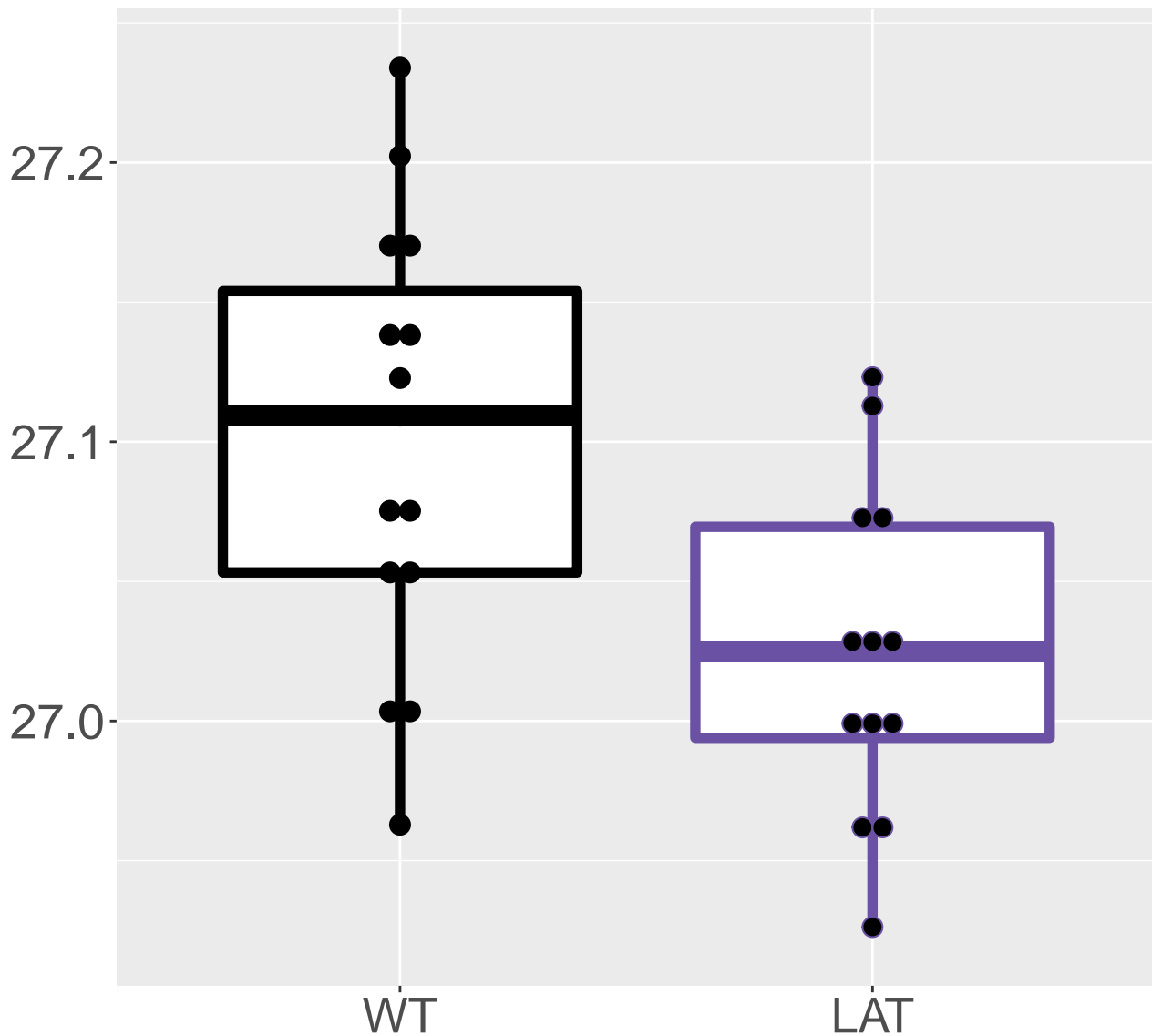
FDR = 0.021, FC = -0.27, sex***



P14148_60S ribosomal protein L7
FDR = 0.021, FC = -0.17

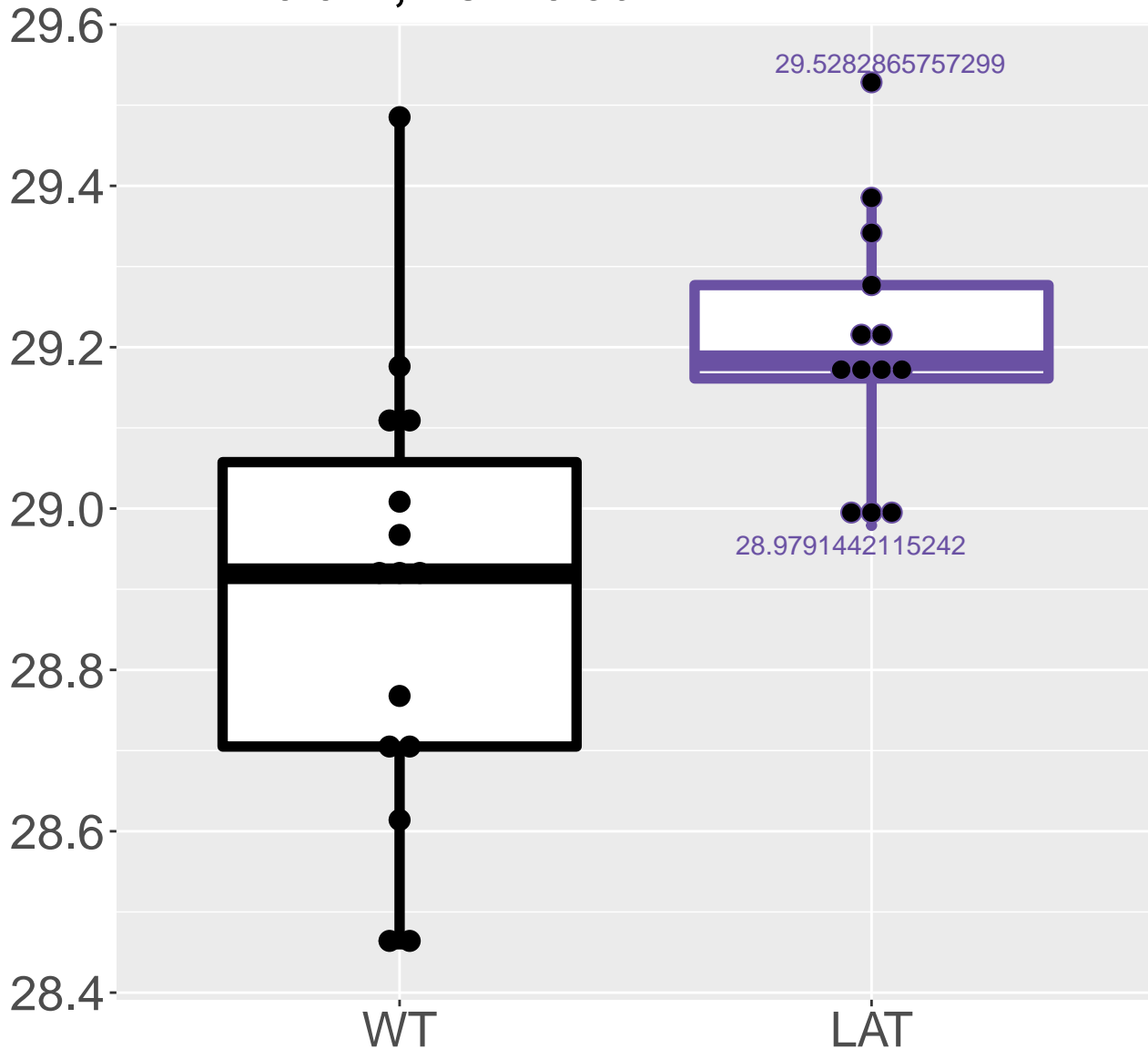


P14131_40S ribosomal protein S16
FDR = 0.022, FC = -0.16

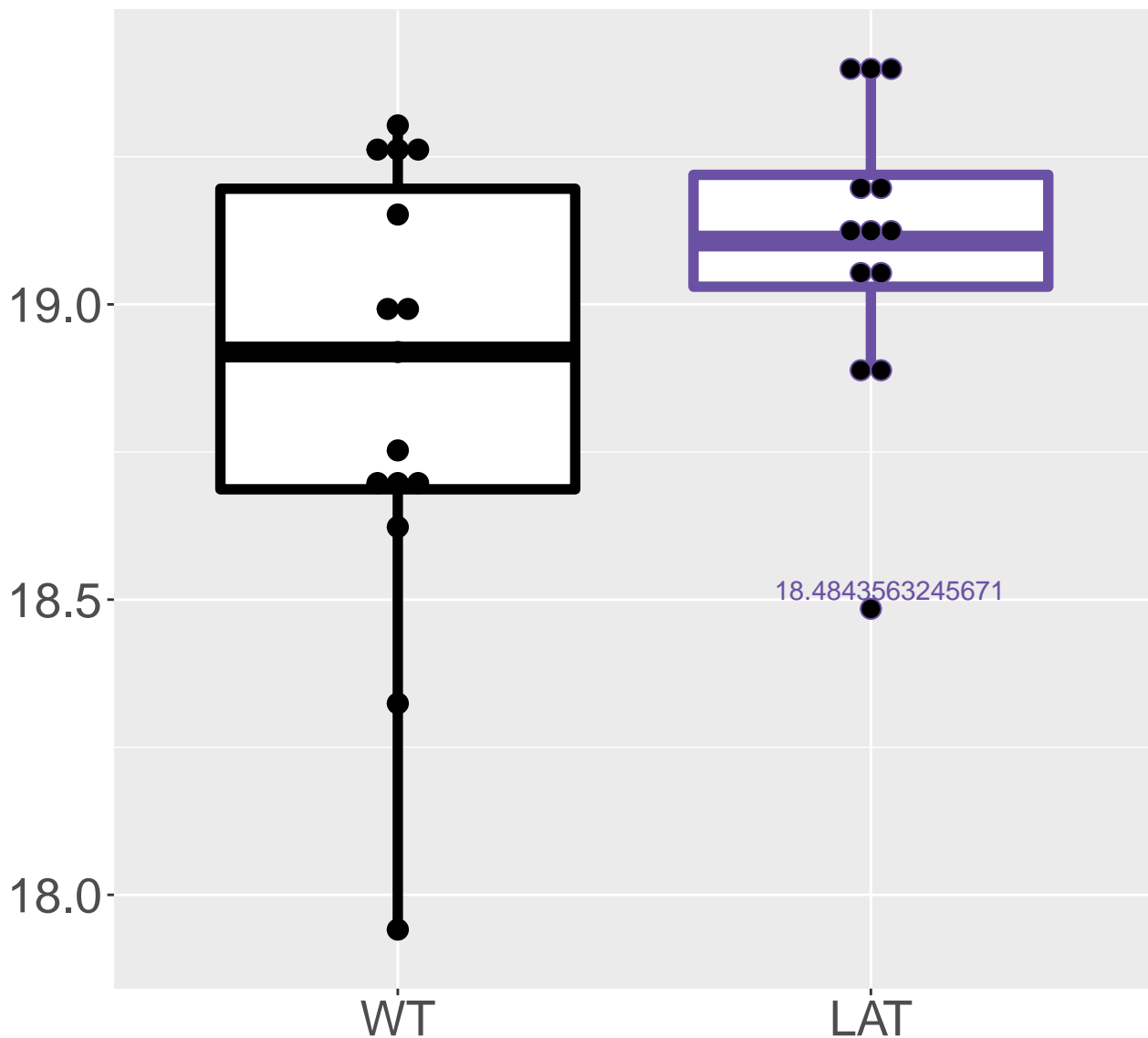


Q9DBM2_Peroxisomal bifunctional.

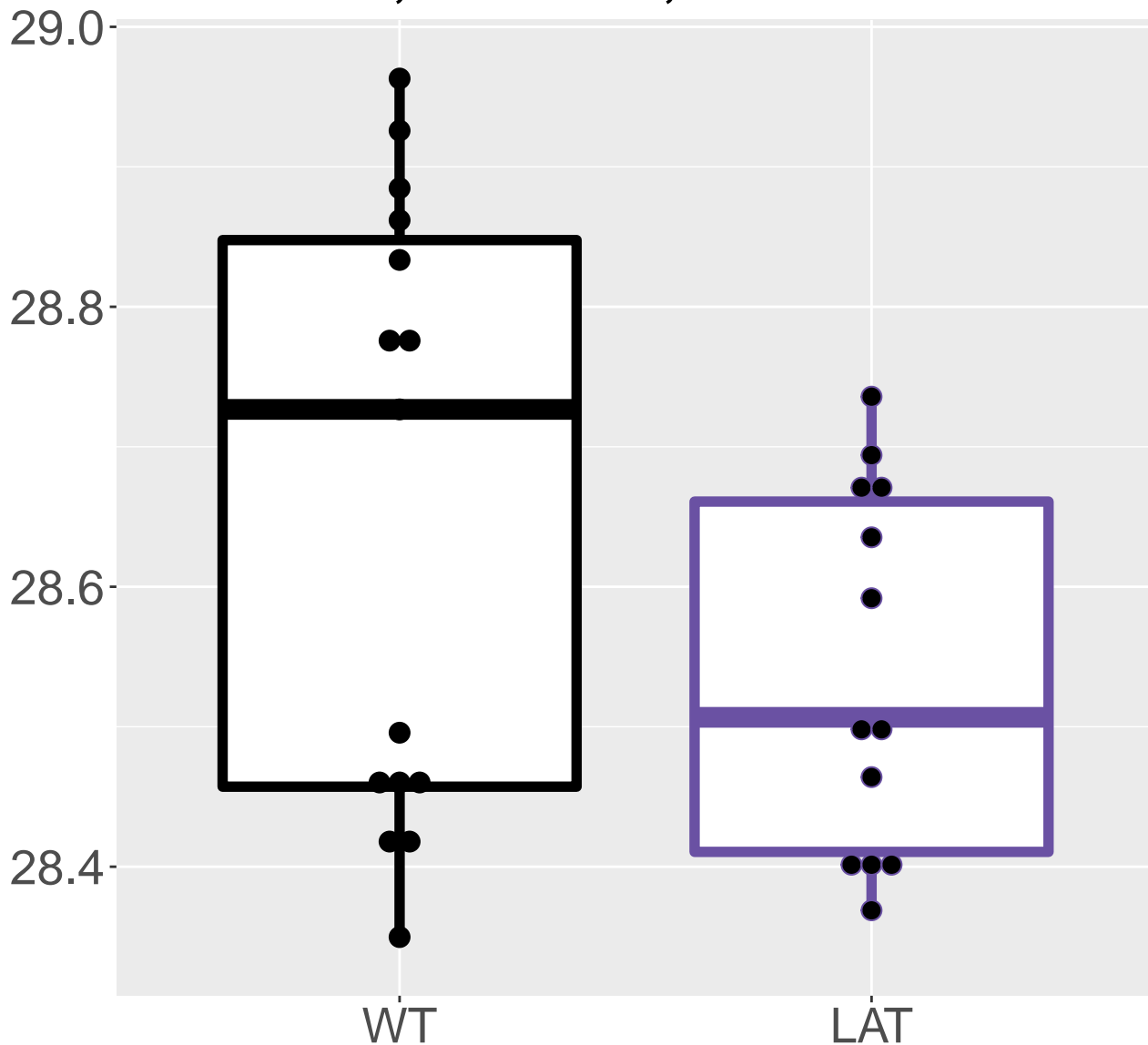
FDR = 0.022, FC = 0.36



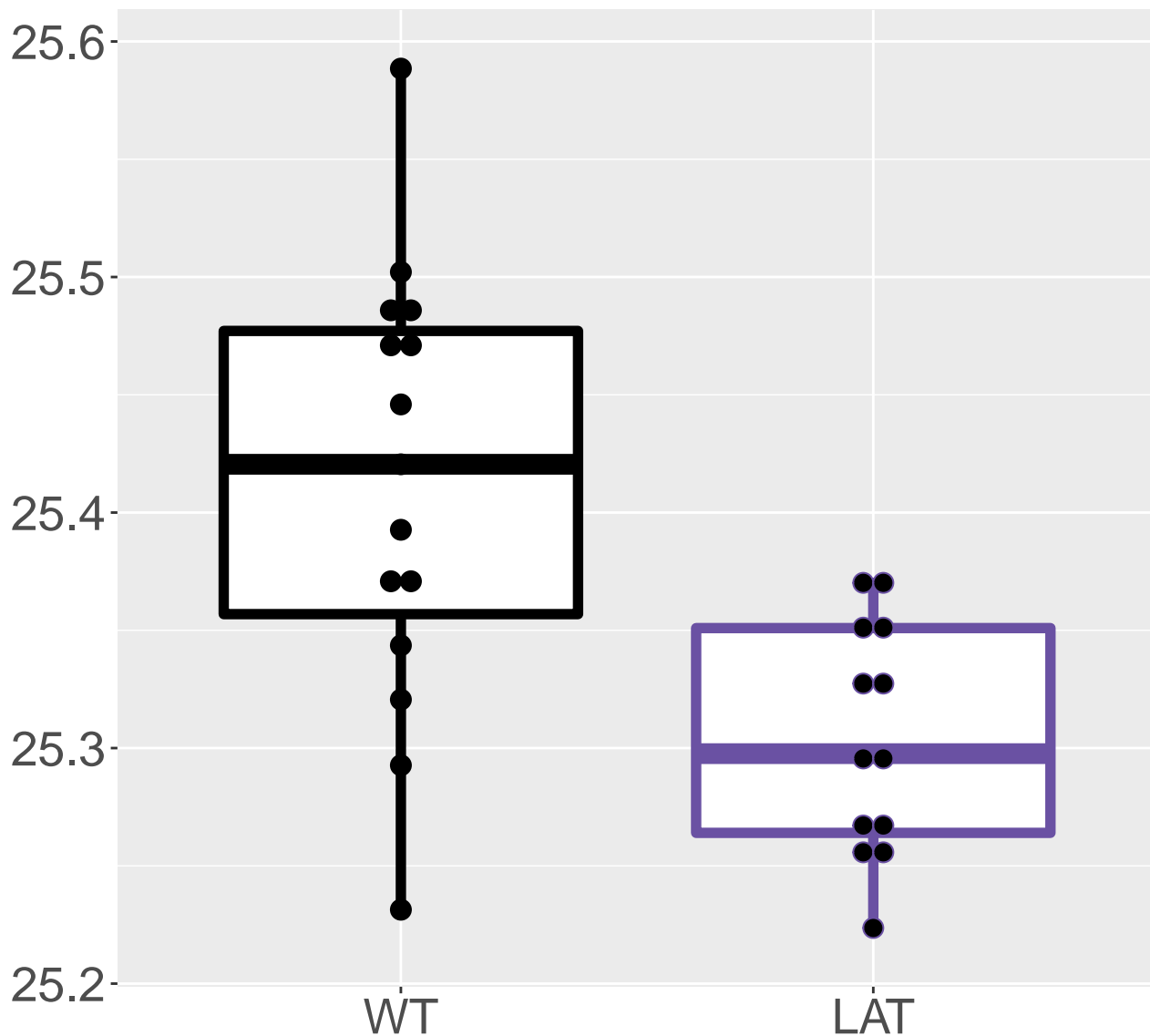
Q8BSY0_Asparyl/asparaginyl bet.
FDR = 0.022, FC = 0.58, sex*



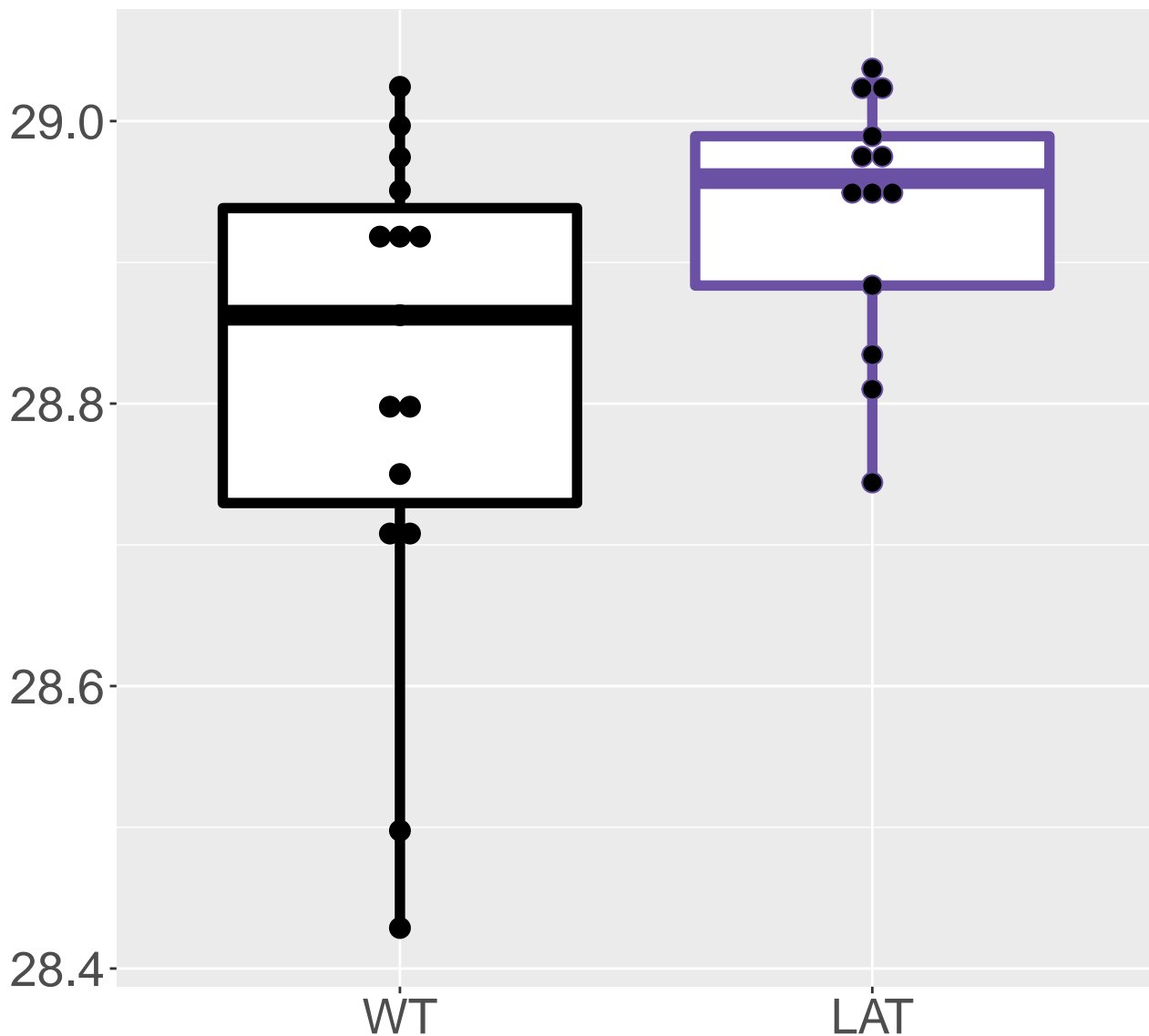
P52196_Thiosulfate sulfurtransf.
FDR = 0.024, FC = -0.18, sex***



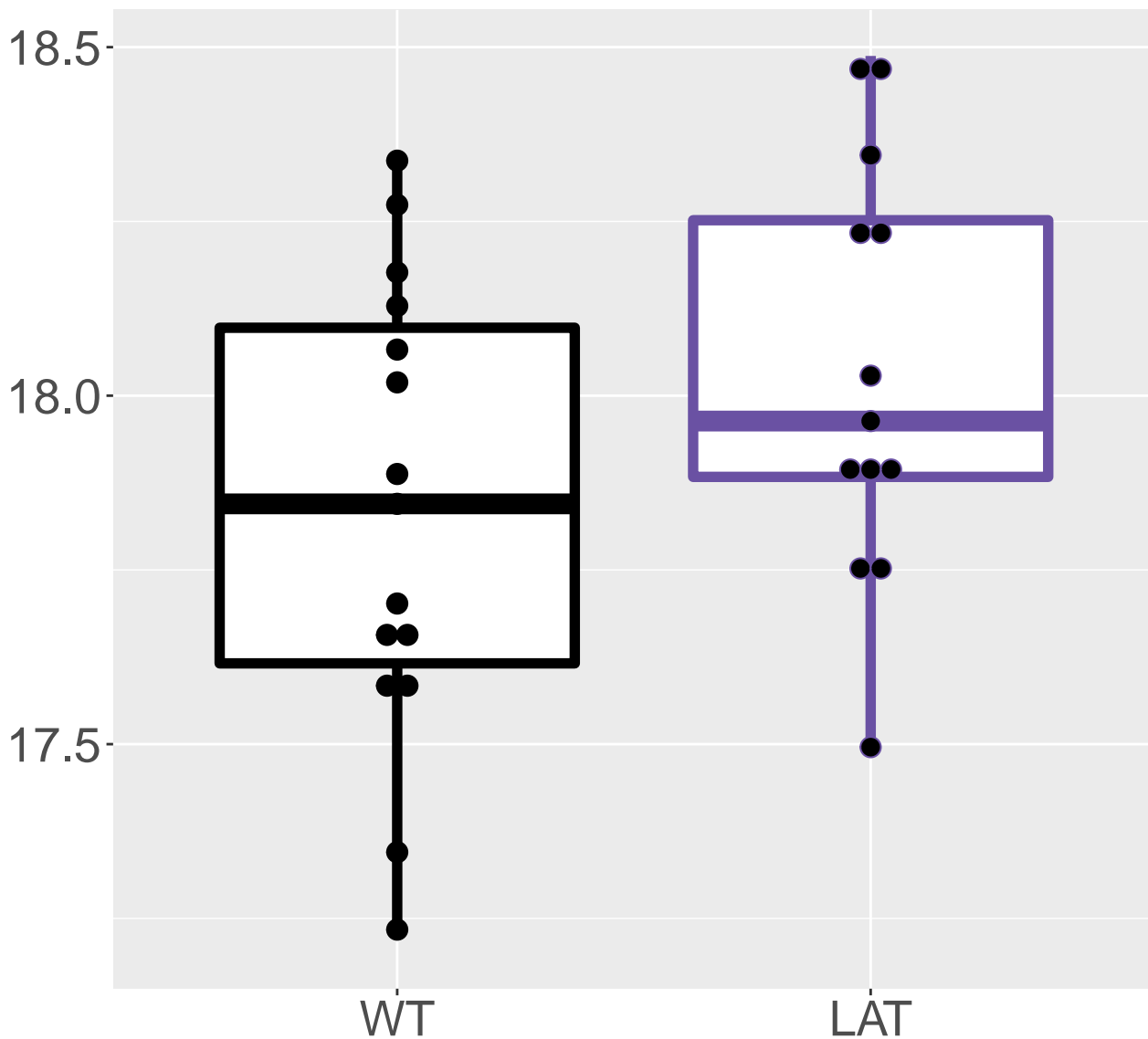
Q6ZWX6_Eukaryotic translation i.
FDR = 0.024, FC = -0.16



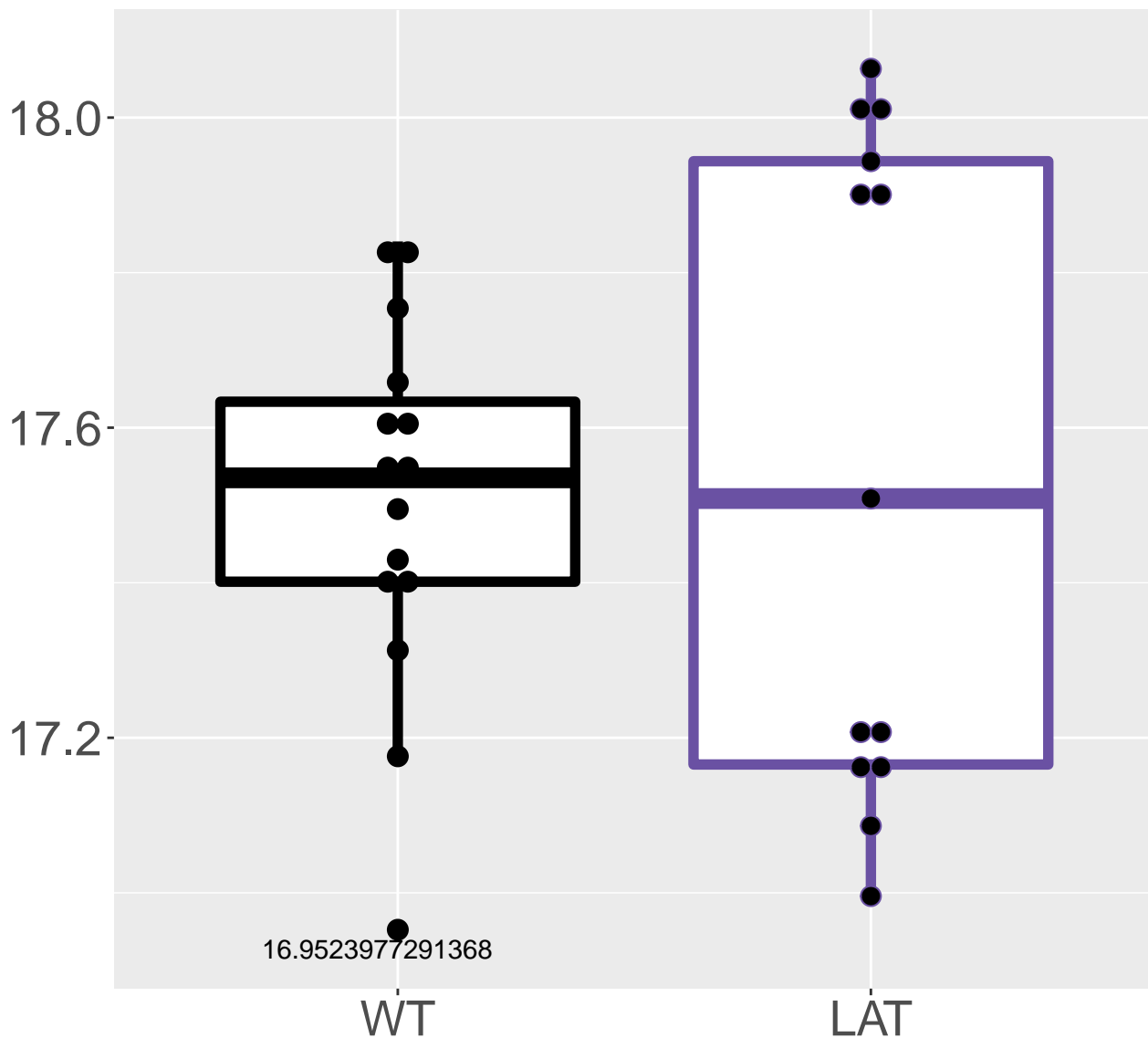
P58252_Elongation factor 2
FDR = 0.024, FC = 0.28, sex**



Q9QXK3_Coatomer subunit gamma-2
FDR = 0.024, FC = 0.52, sex*

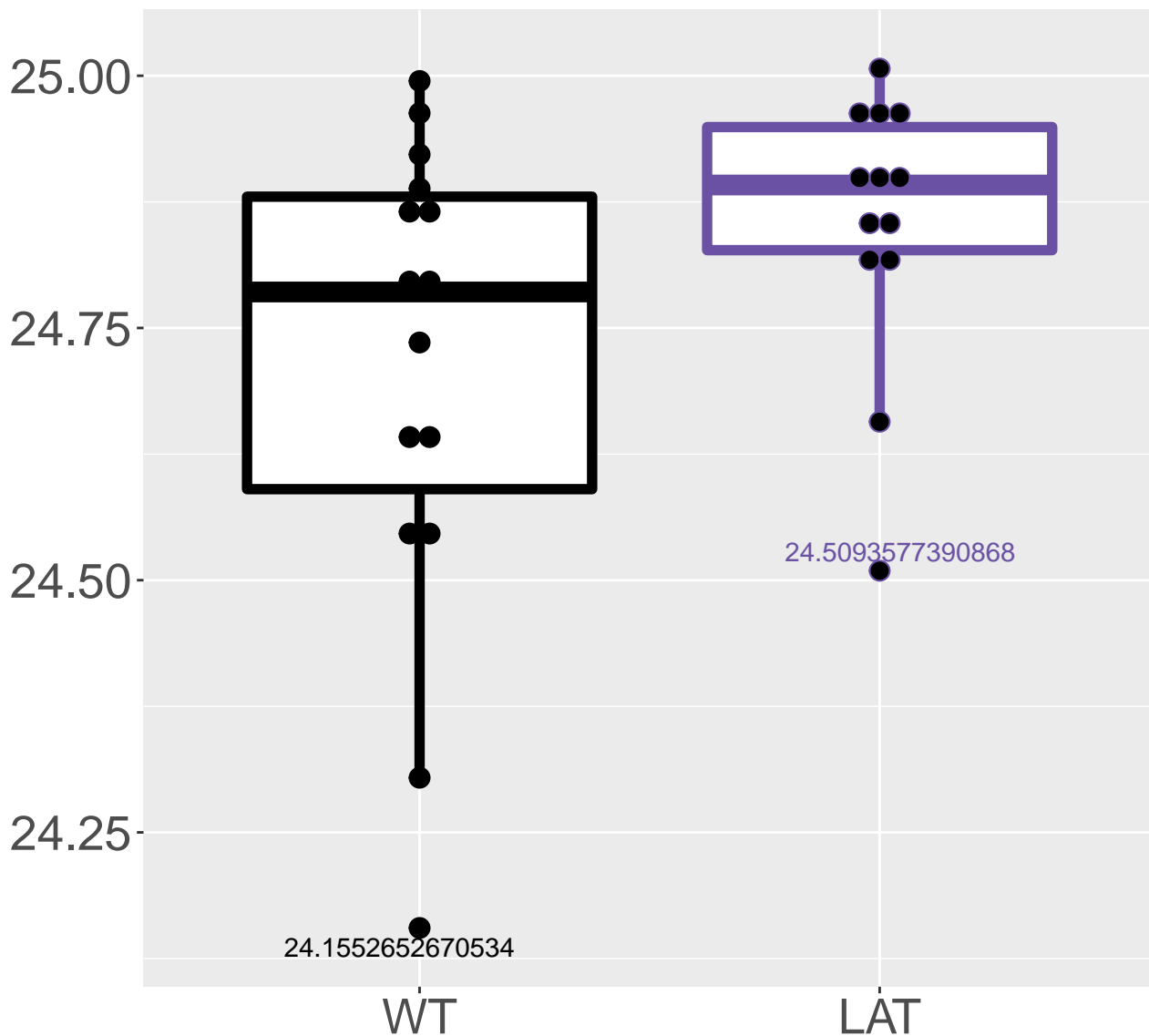


Q9JIG7_Coiled-coil domain-conta.
FDR = 0.024, FC = 0.5, sex**

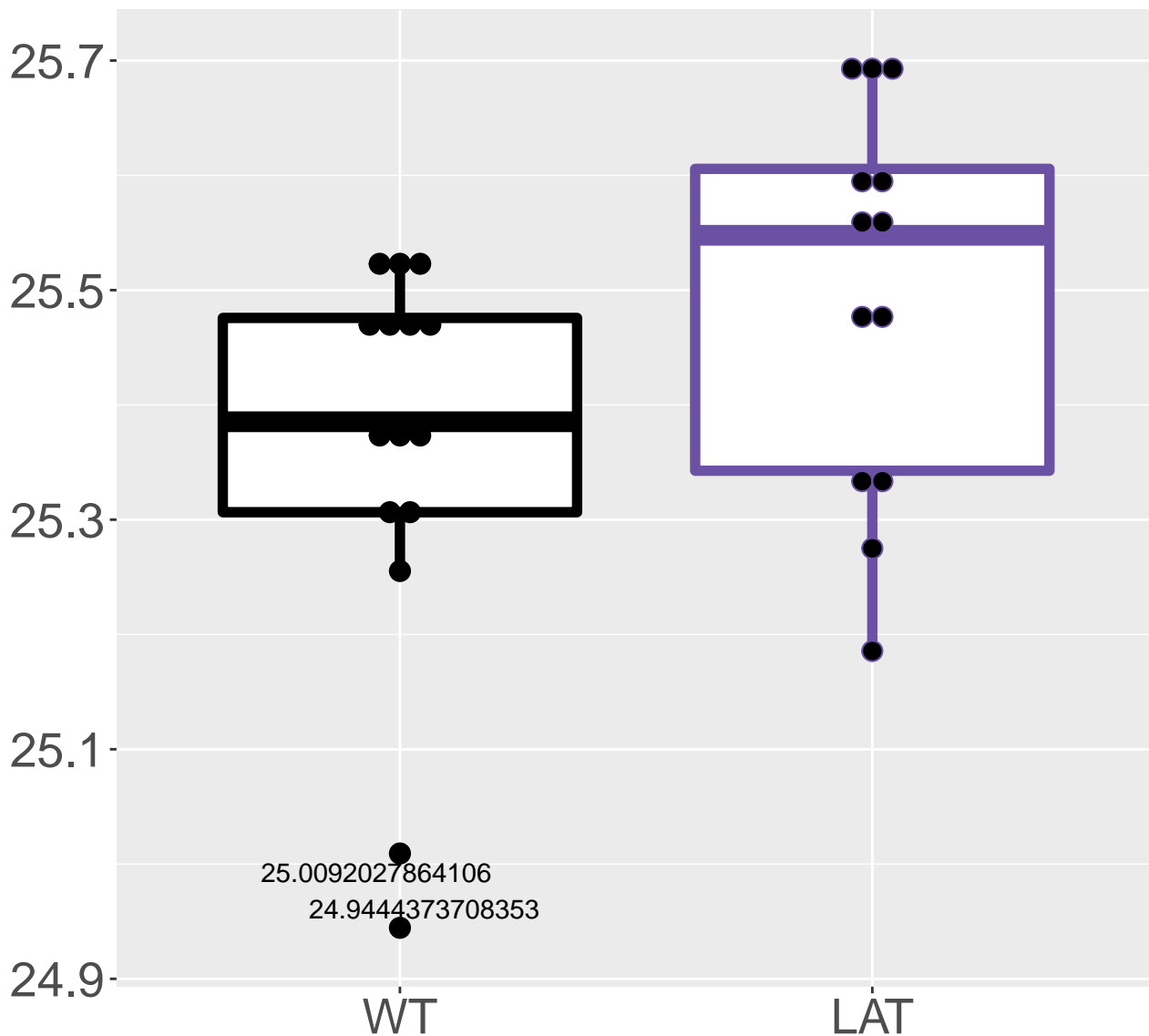


Q9JIF7_Coatomer subunit beta

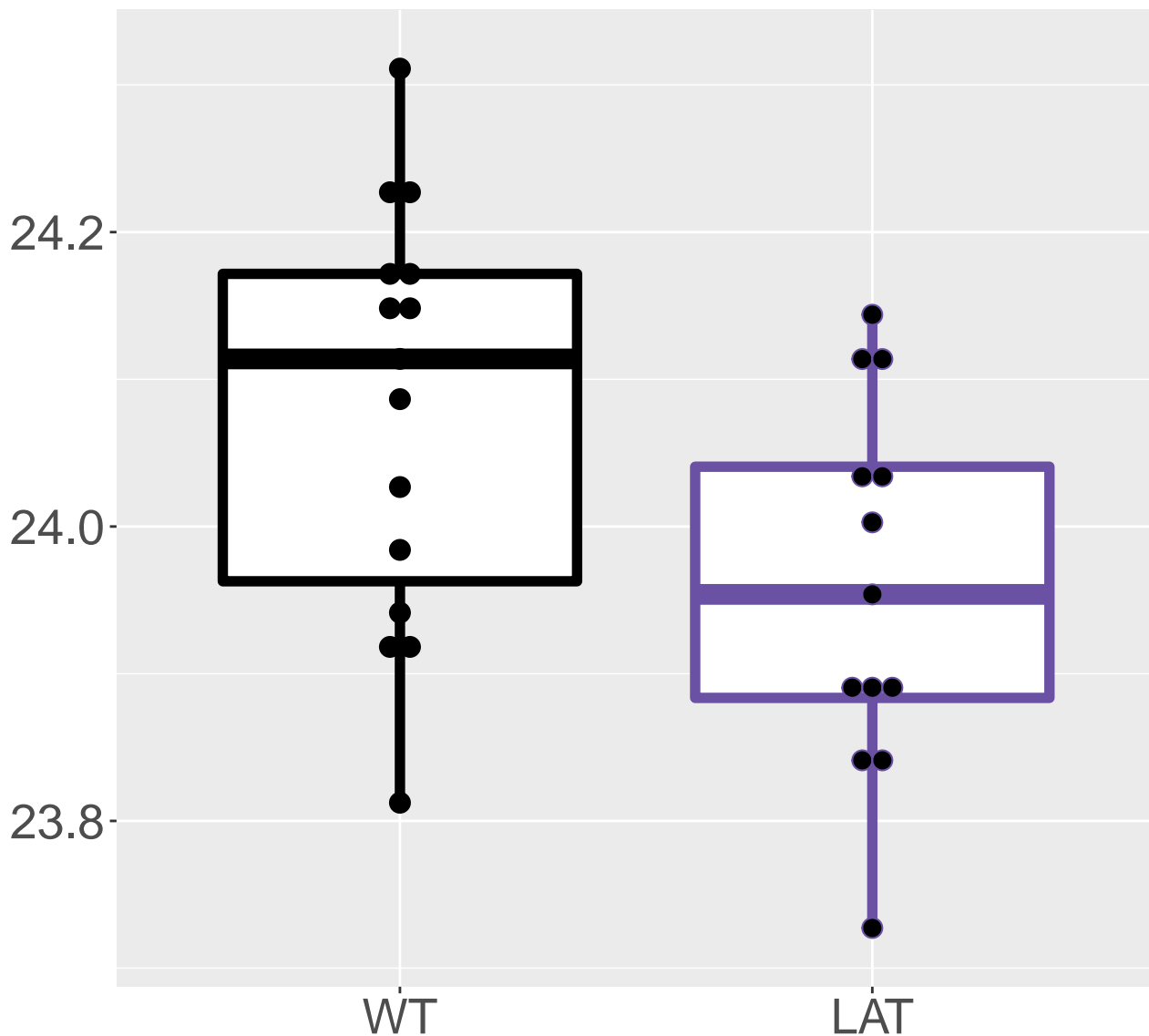
FDR = 0.024, FC = 0.37, sex**



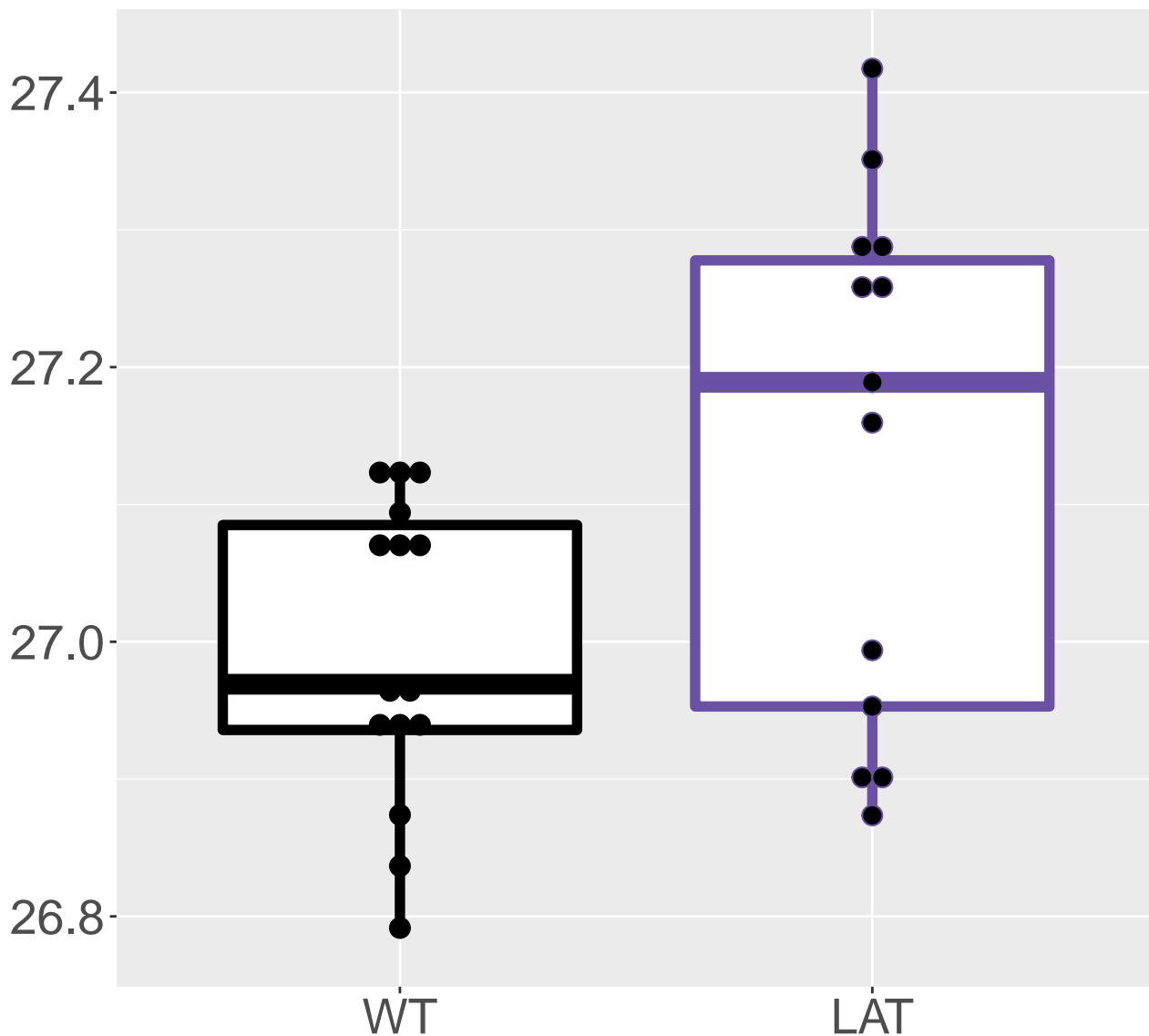
Q9EQH3_Vacuolar protein sorting.
FDR = 0.024, FC = 0.35, sex*



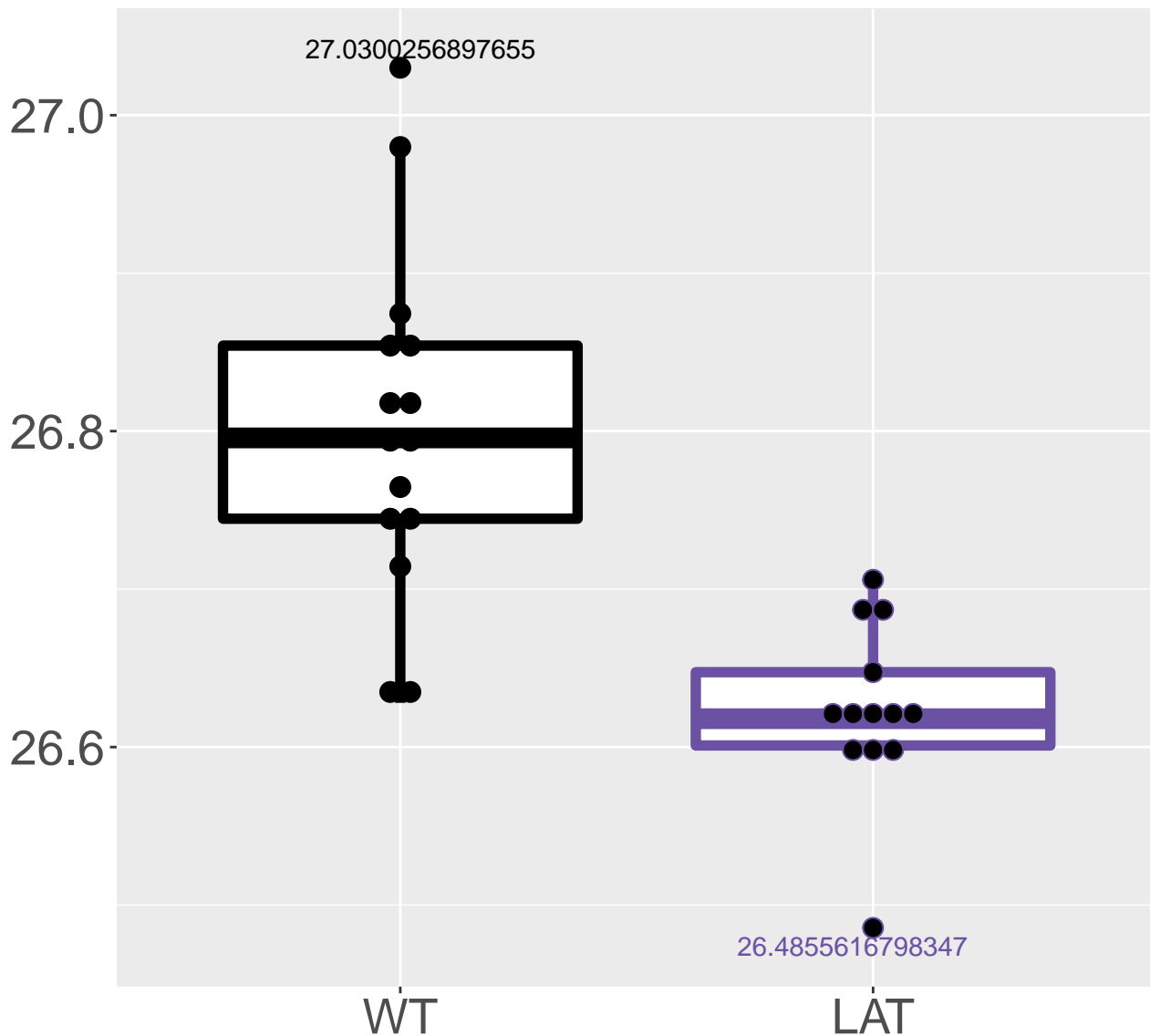
Q9JLZ3_Methylglutaconyl-CoA hyd.
FDR = 0.024, FC = -0.26, sex*



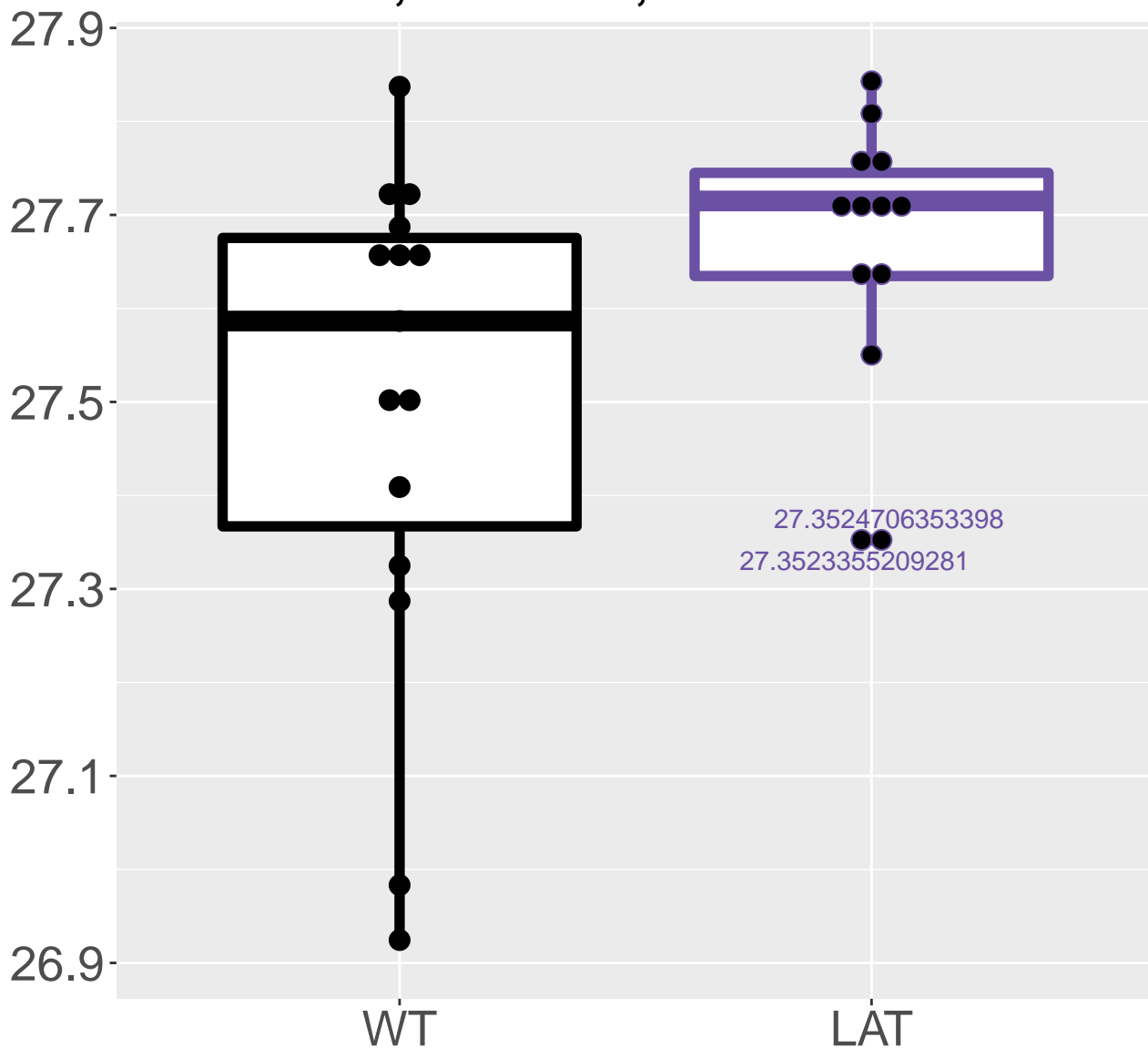
Q4LDG0_Bile acyl-CoA synthetase
FDR = 0.024, FC = 0.23, sex*



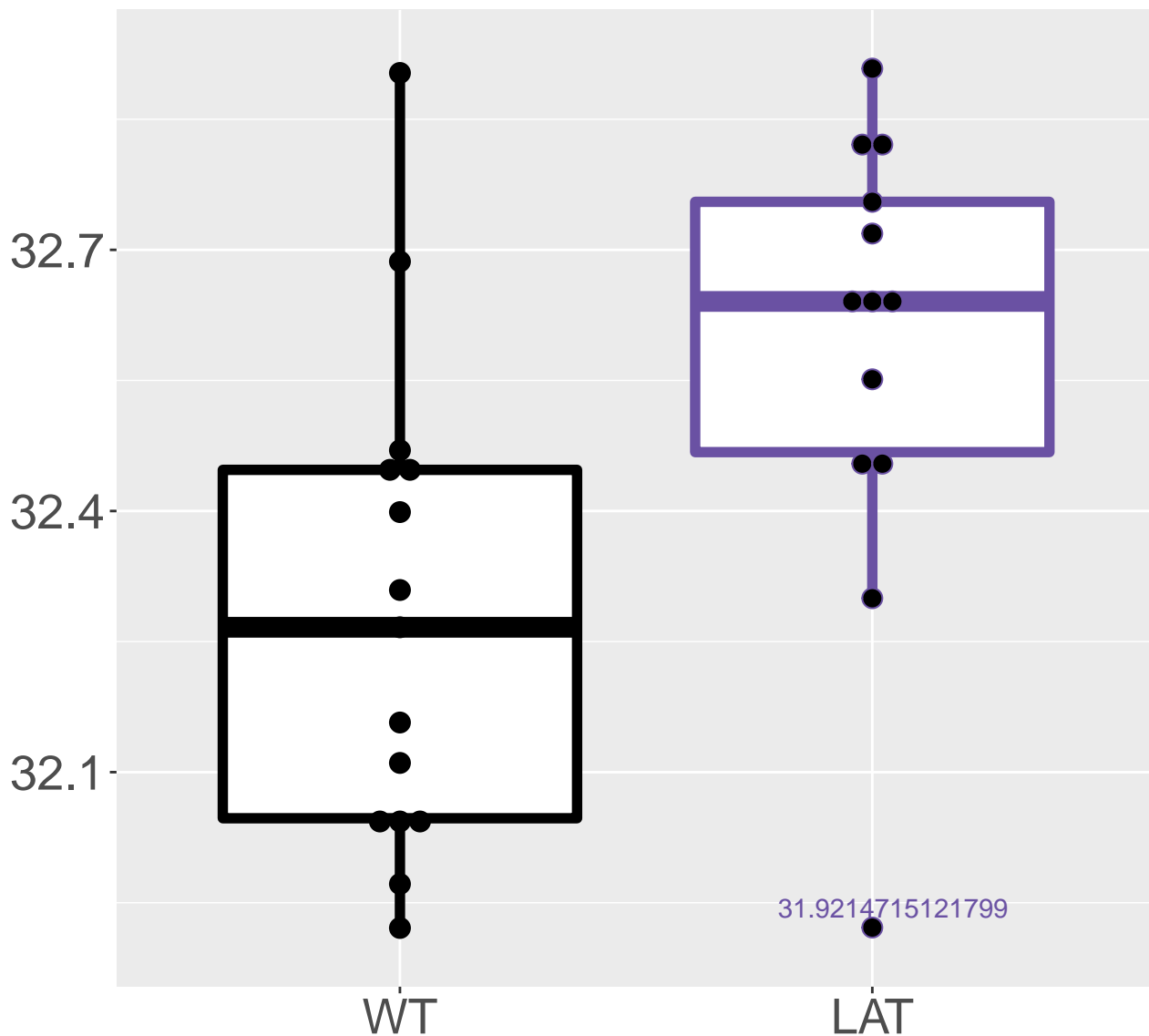
FDR = 0.024, FC = -0.2



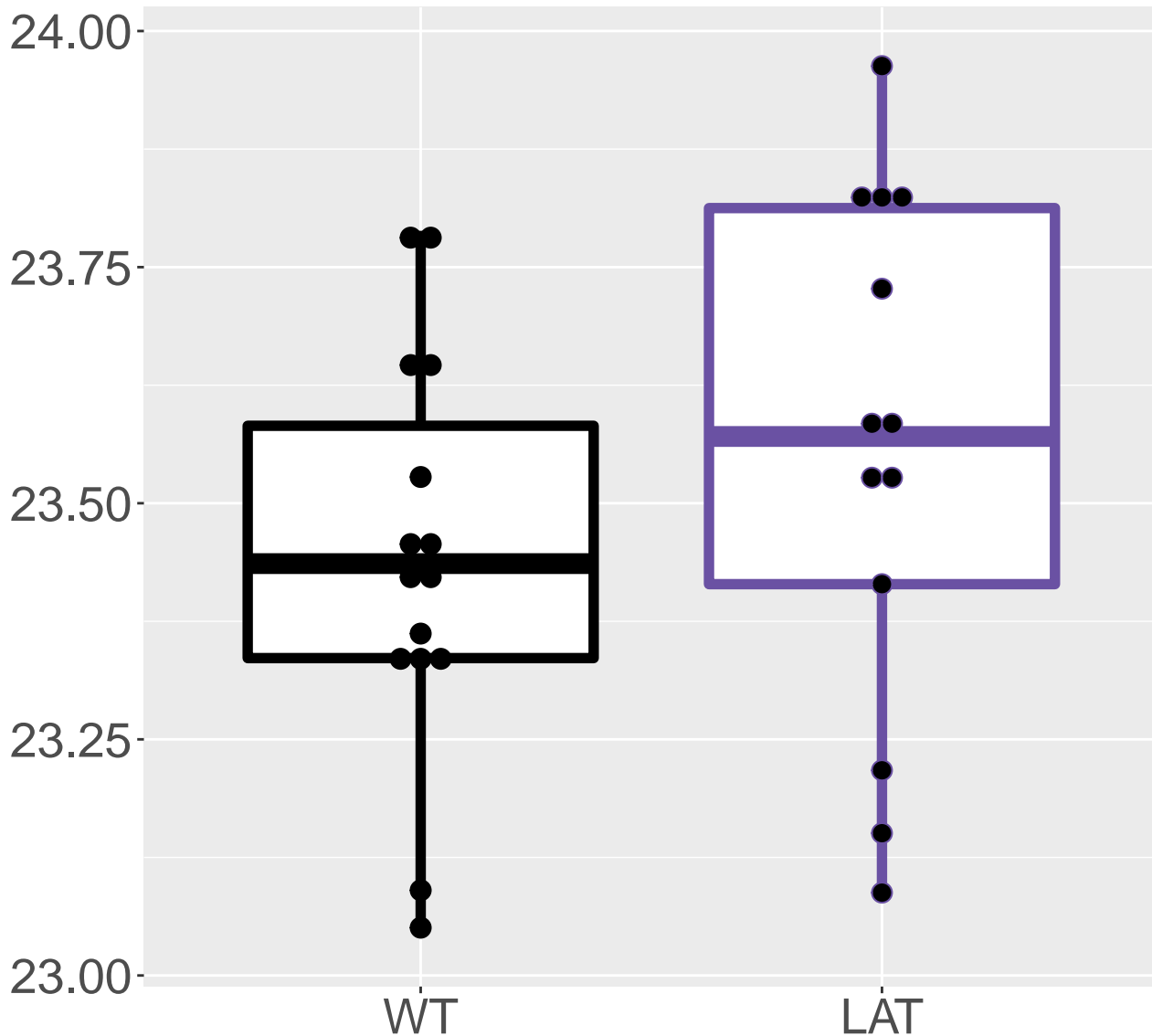
O08601_Microsomal triglyceride .
FDR = 0.025, FC = 0.41, sex**



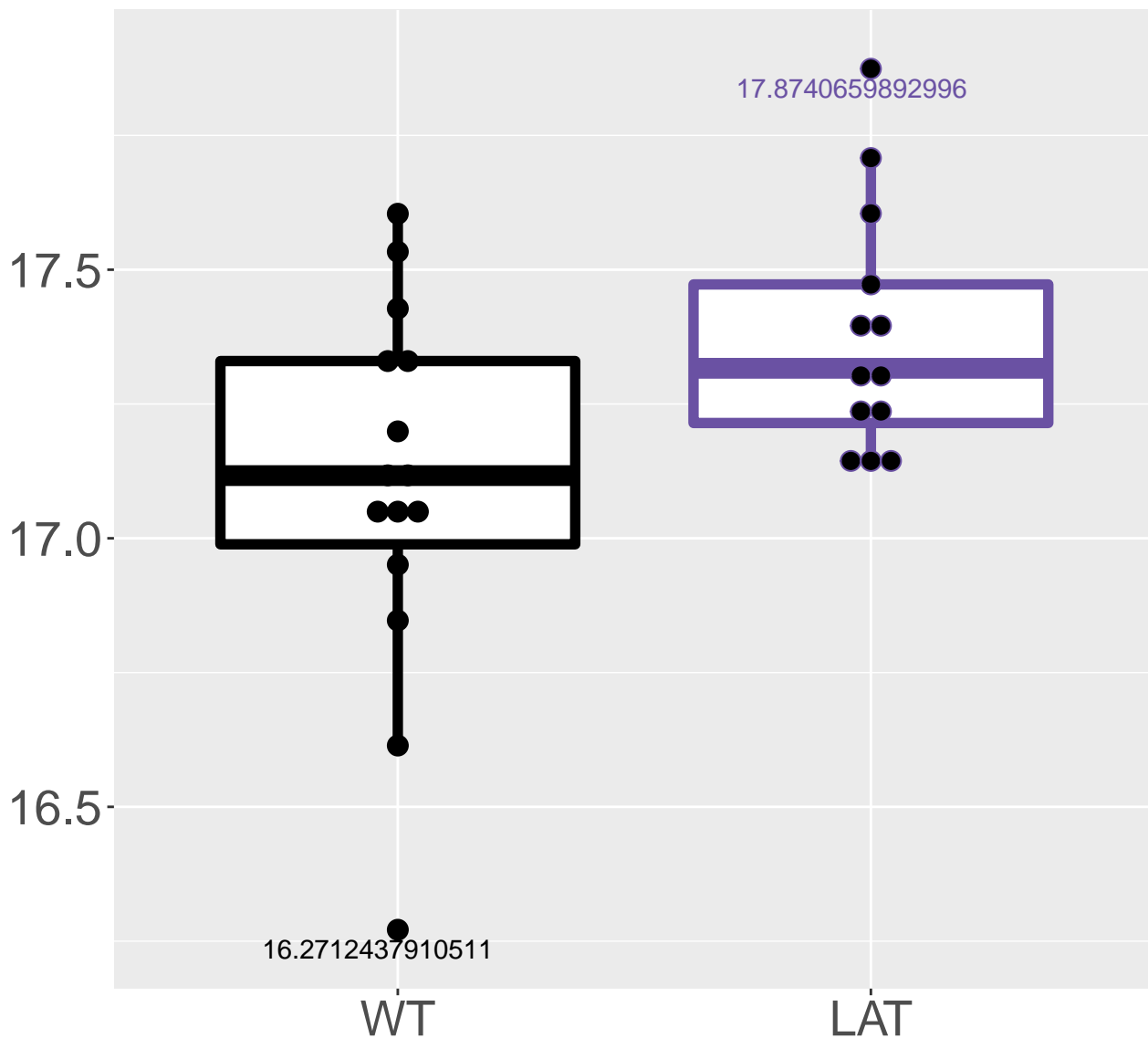
Q8C196_Carbamoyl-phosphate synt.
FDR = 0.025, FC = 0.43



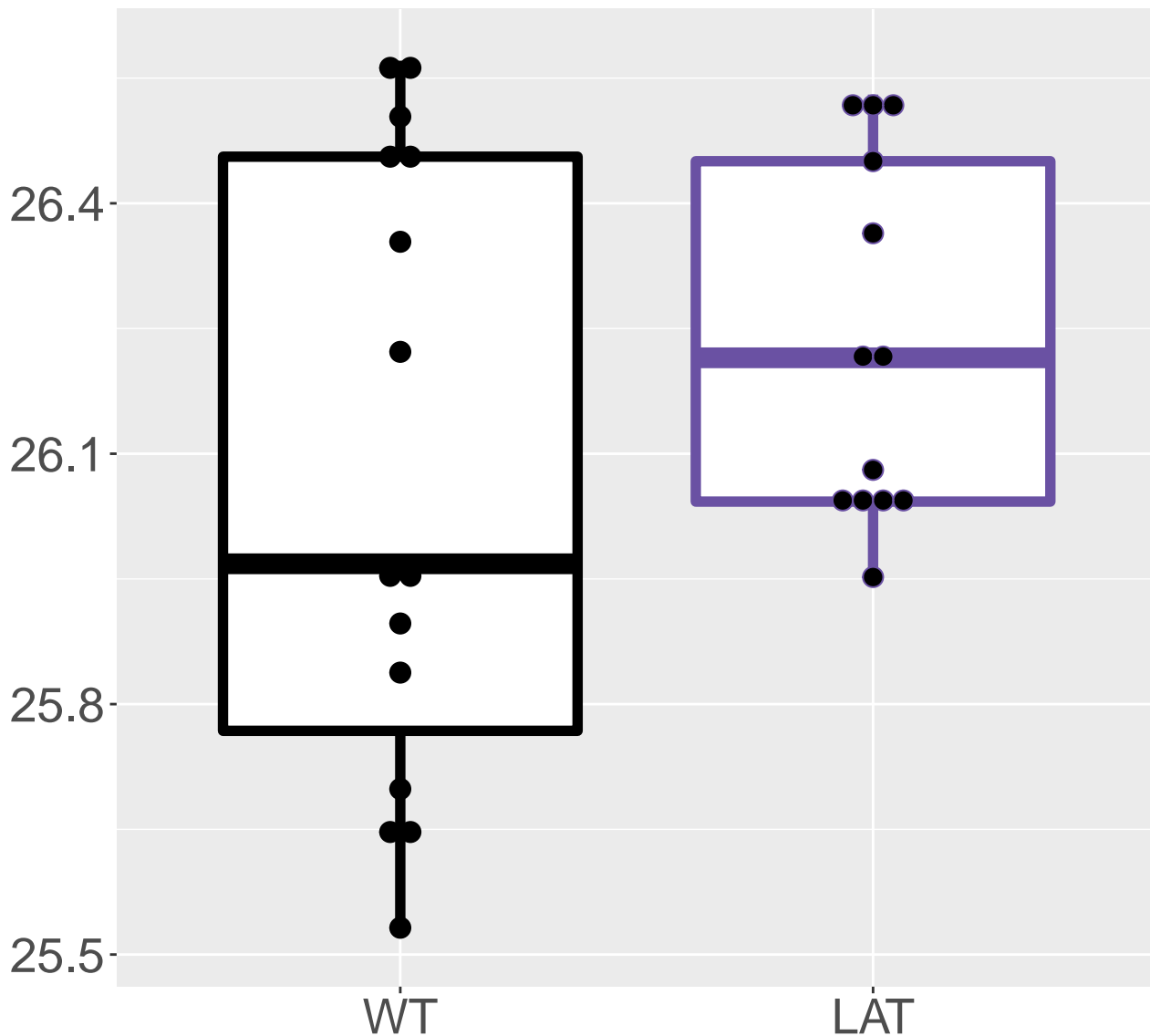
Q9EQH2_Endoplasmic reticulum am.
FDR = 0.025, FC = 0.4, sex*



Q8VI75_Importin-4
FDR = 0.025, FC = 0.54, sex*

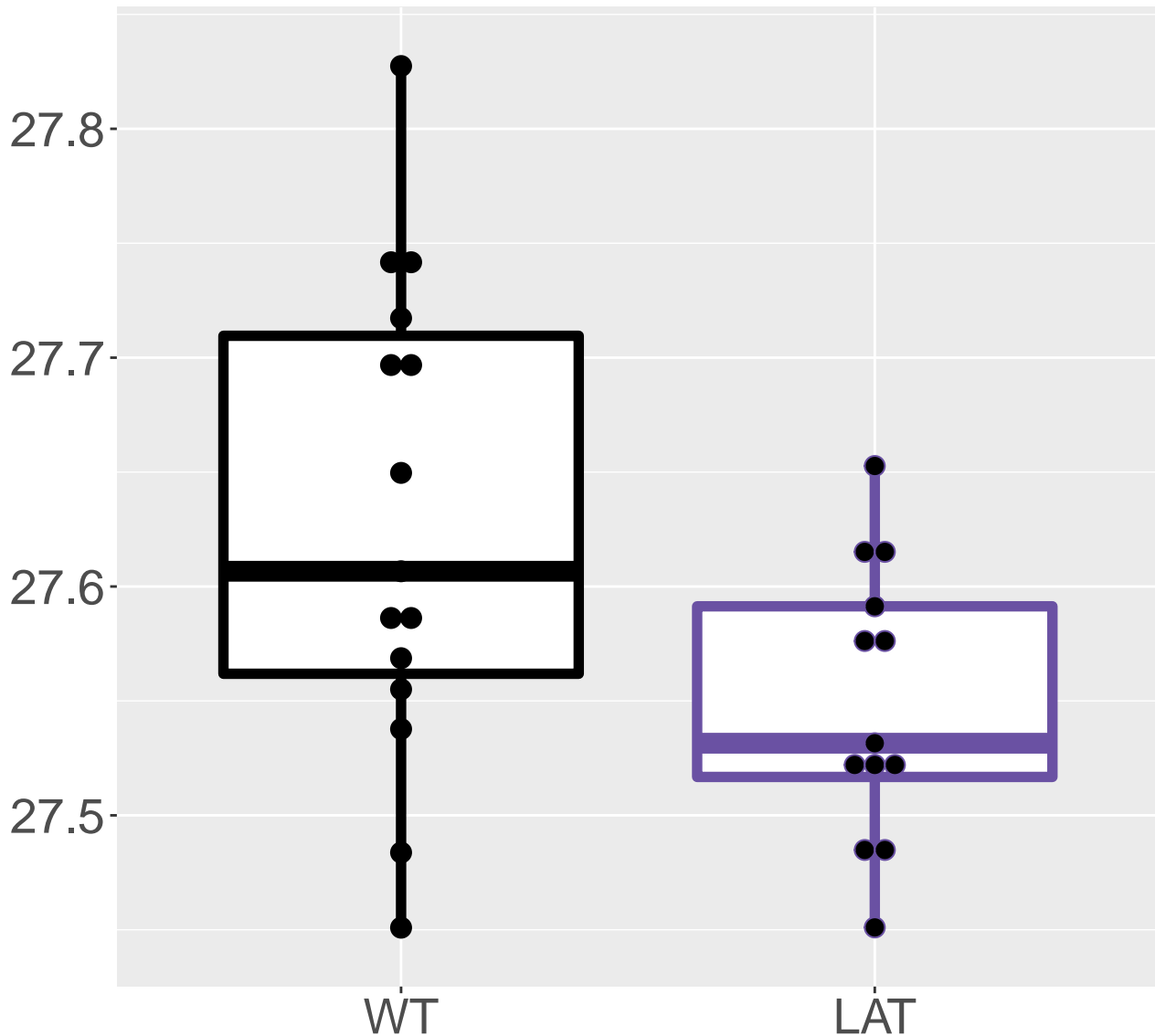


Q91X34_Bile acid-CoA:amino acid.
FDR = 0.025, FC = 0.29, sex***

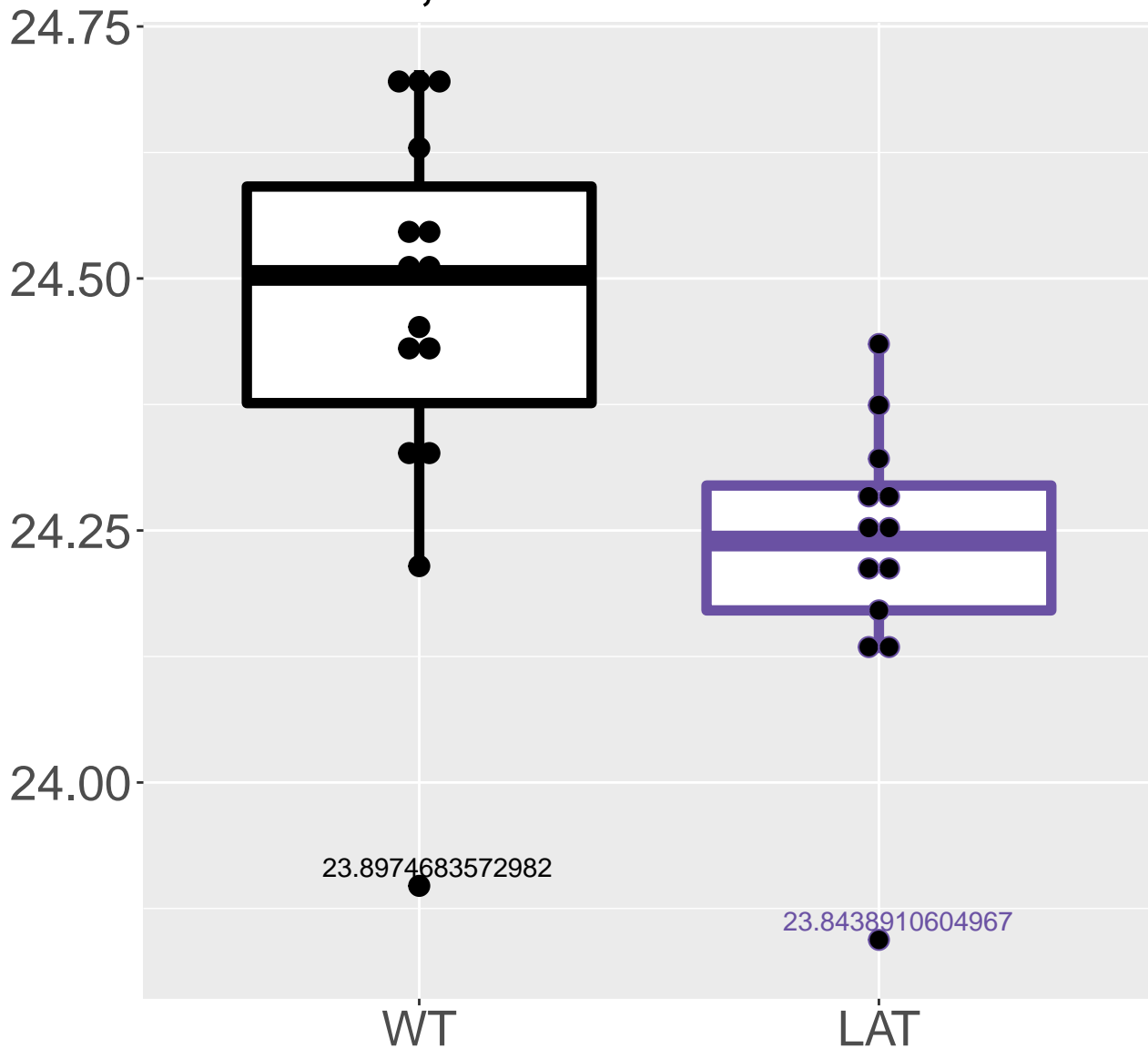


O35129_Prohibitin-2

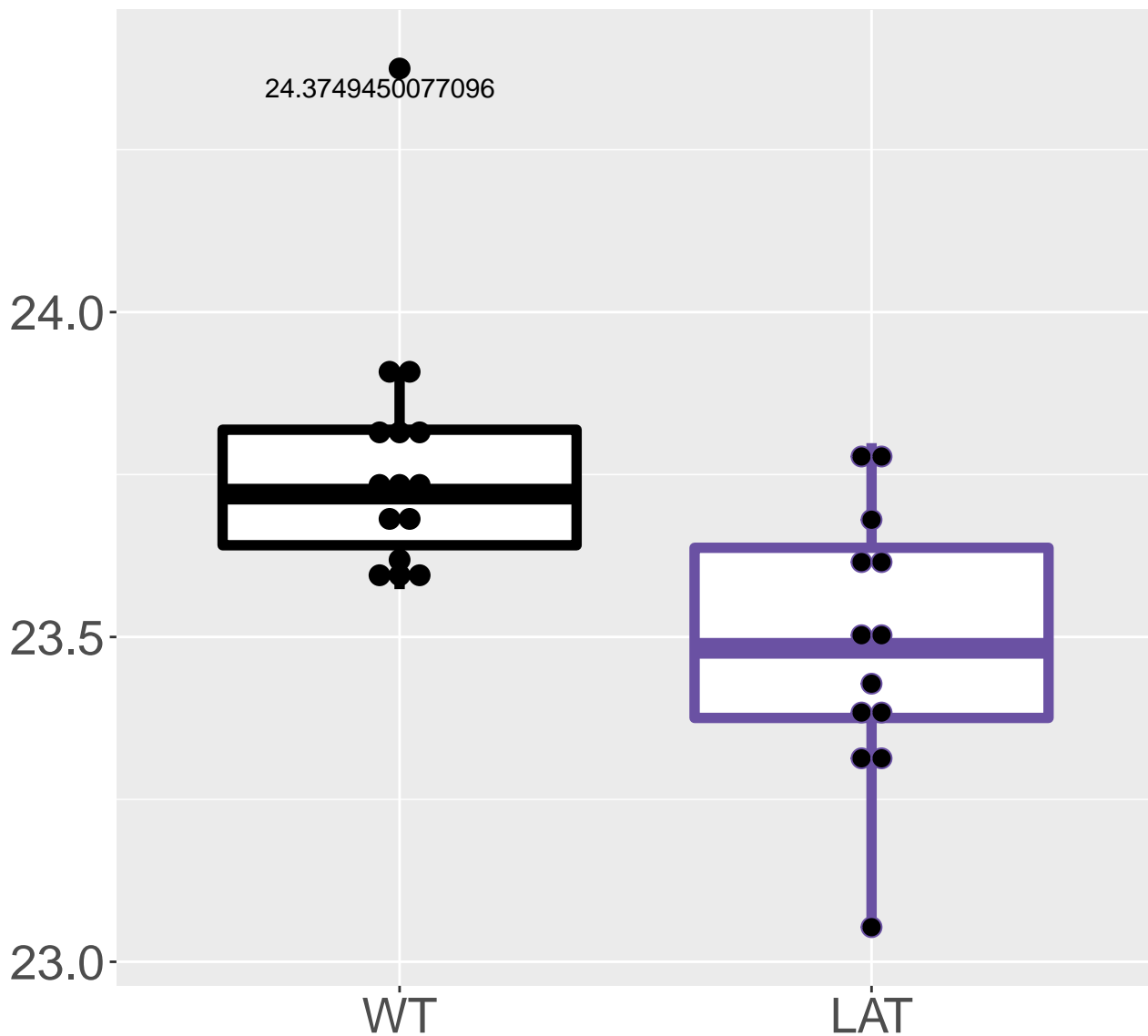
FDR = 0.026, FC = -0.17, sex**



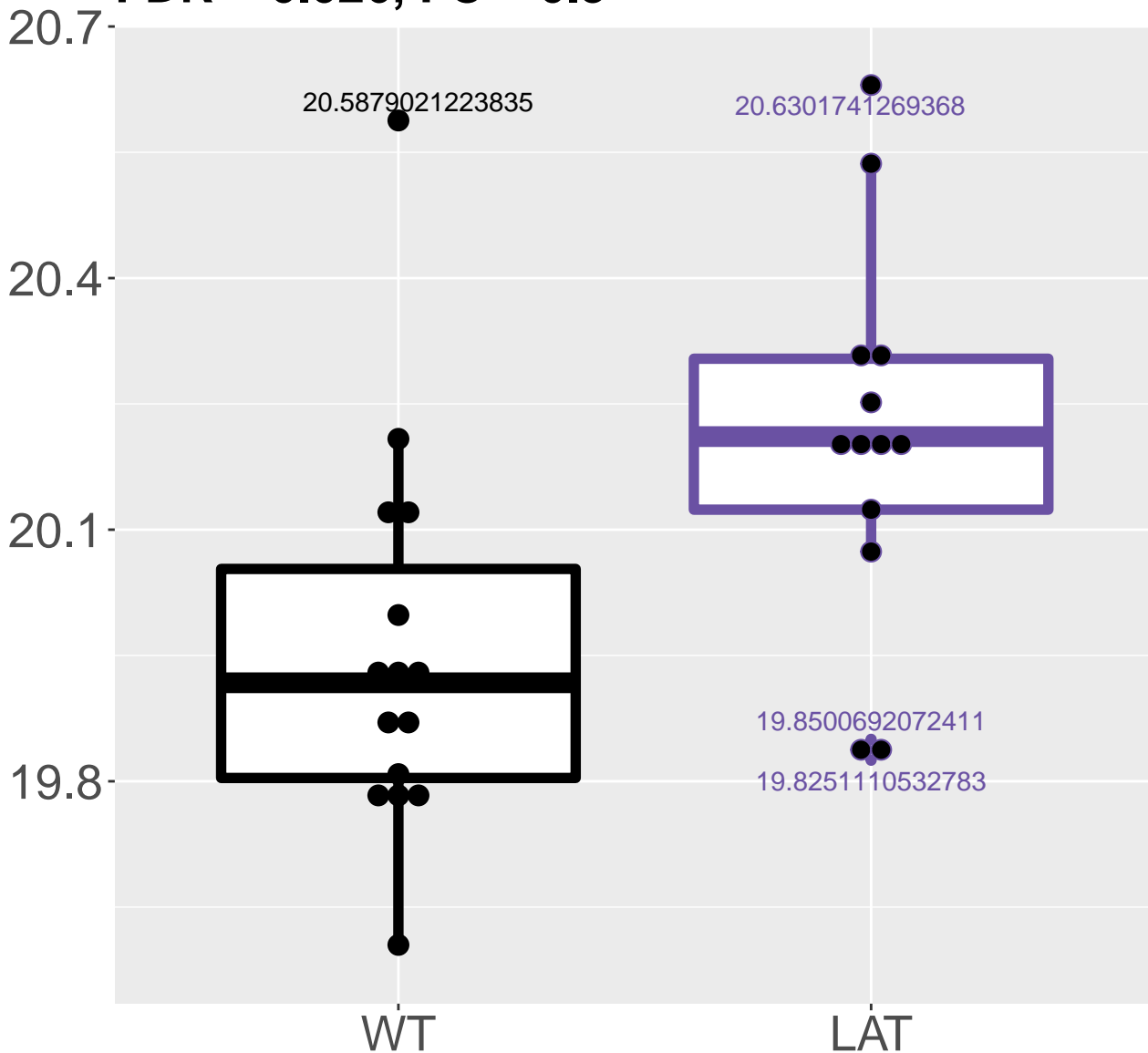
P99027_60S acidic ribosomal pro.
FDR = 0.026, FC = -0.34



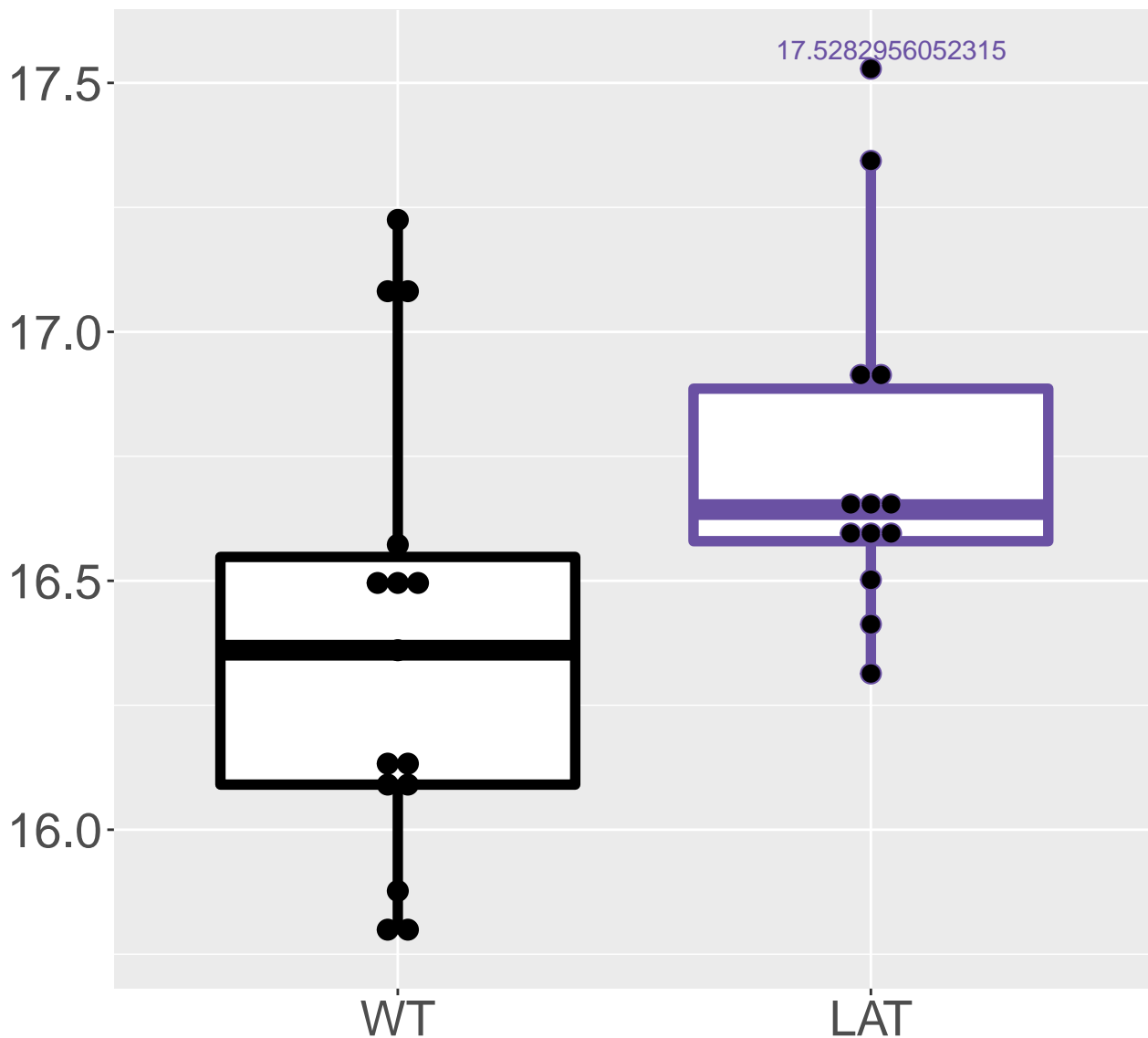
Q9WUU7_Cathepsin Z
FDR = 0.026, FC = -0.44, sex*



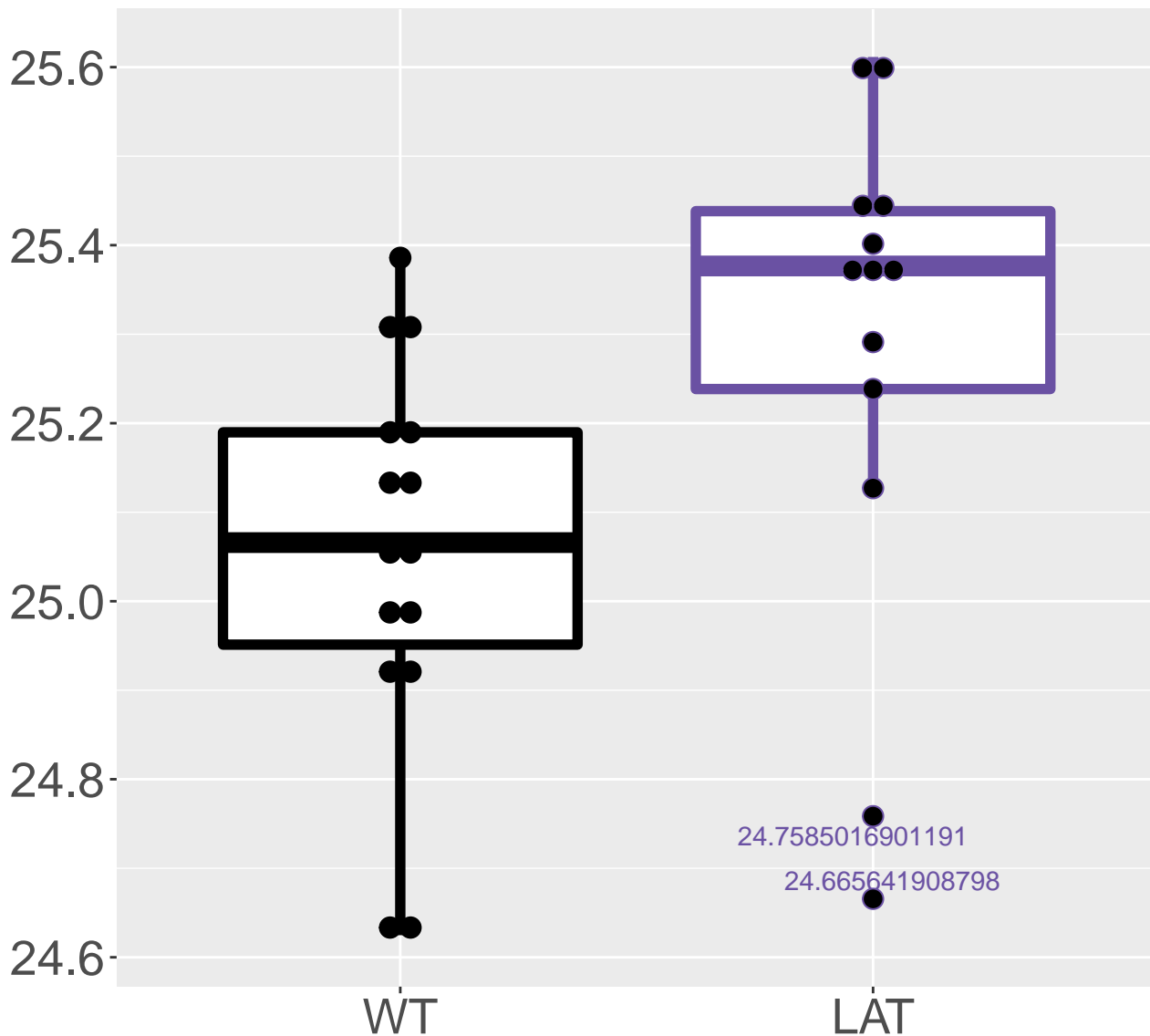
**Q6TEK5_Vitamin K epoxide reduct.
FDR = 0.026, FC = 0.3**



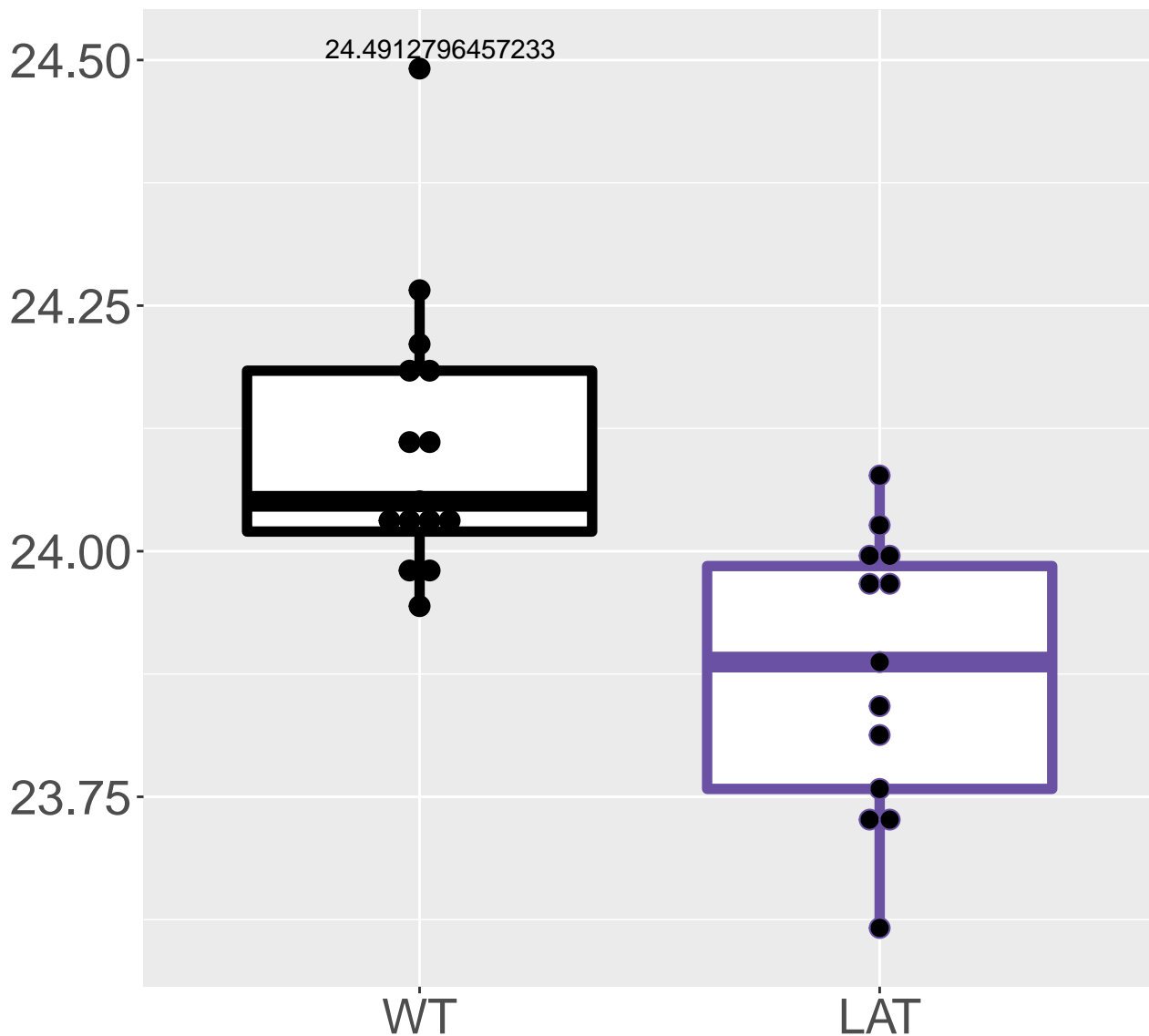
Q9DBA6_Peroxisomal leader pepti.
FDR = 0.027, FC = 0.6, sex*



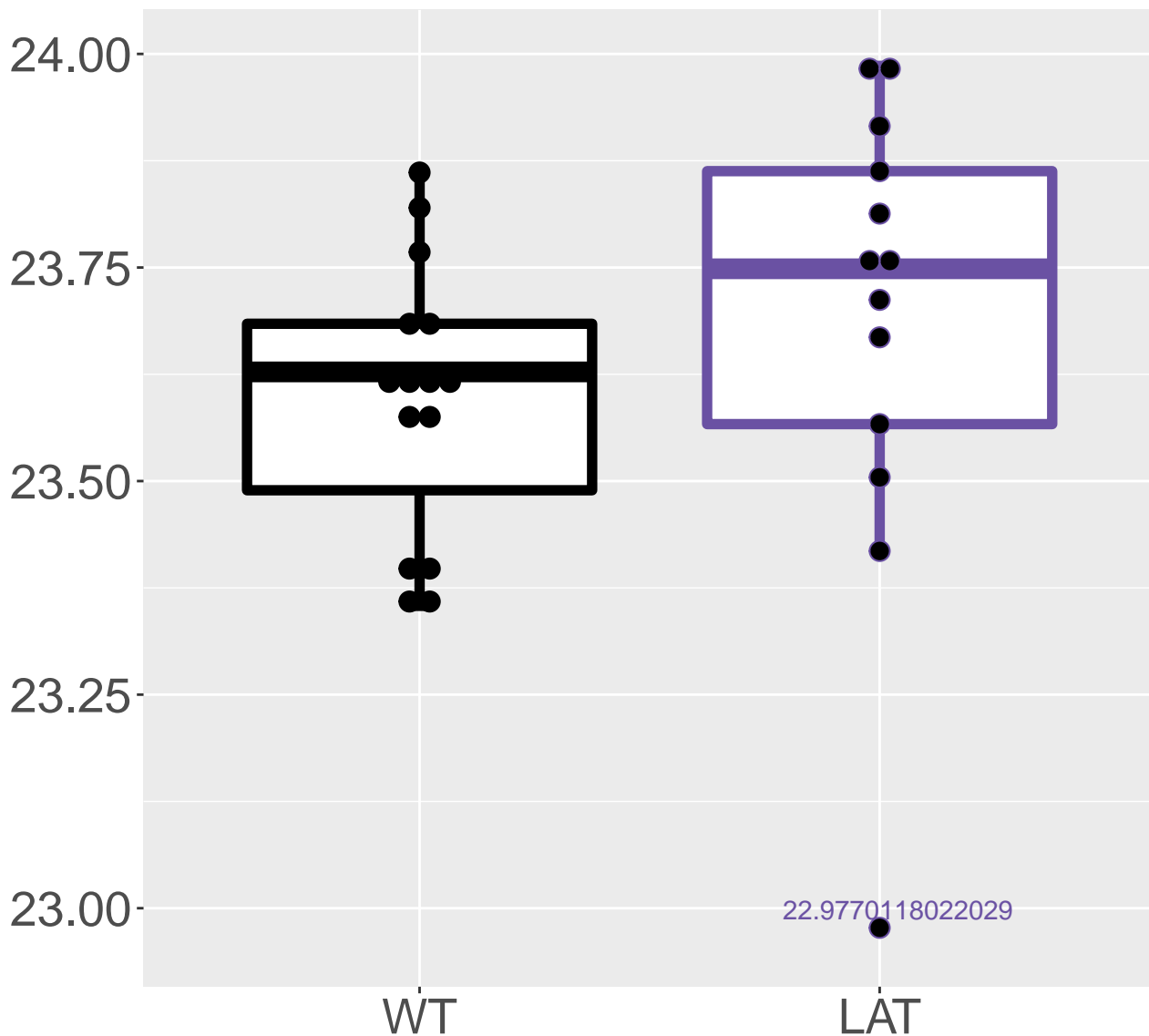
Q8VDN2_Sodium/potassium-transpo.
FDR = 0.027, FC = 0.42



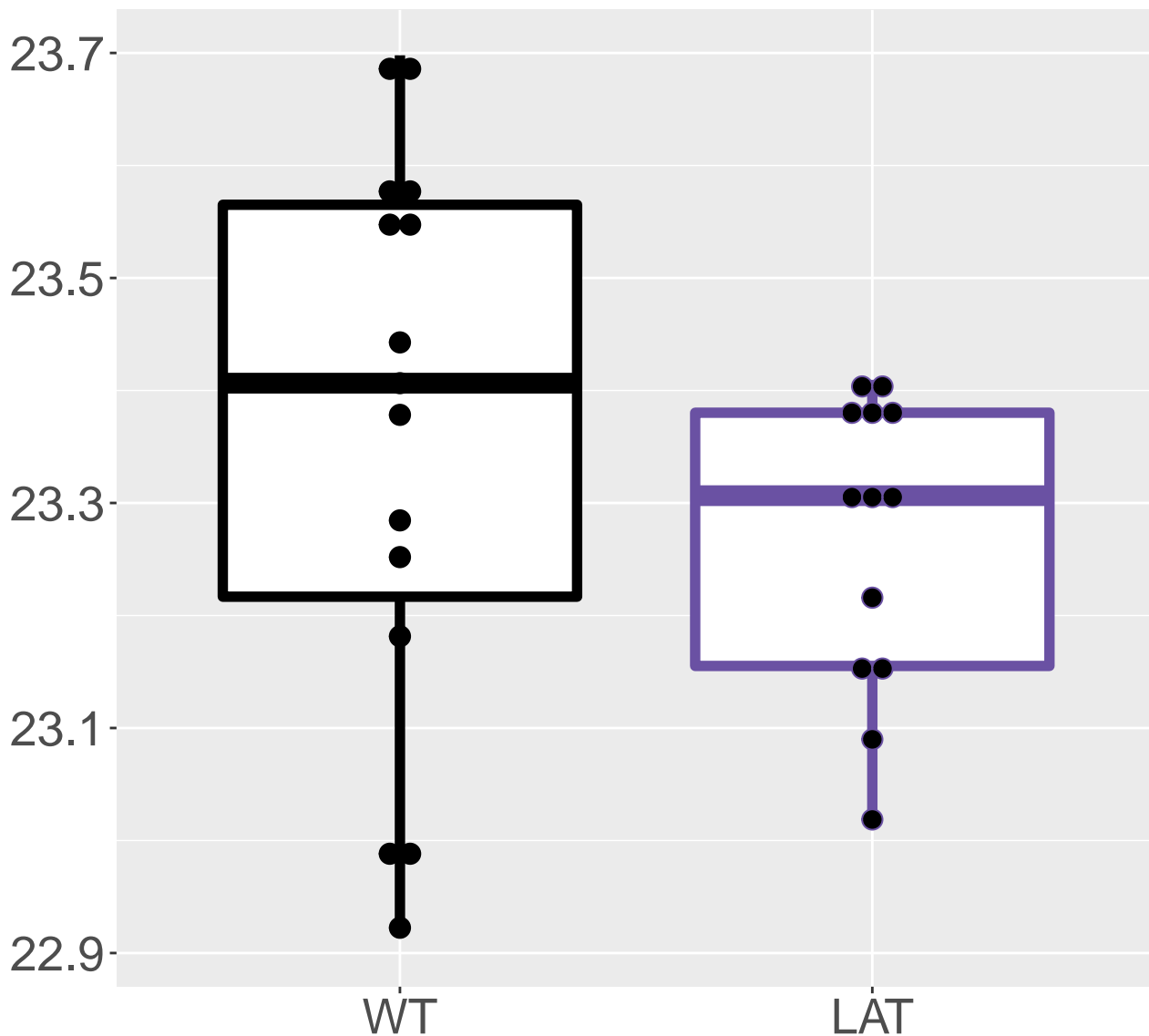
P83882_60S ribosomal protein L3.
FDR = 0.027, FC = -0.32



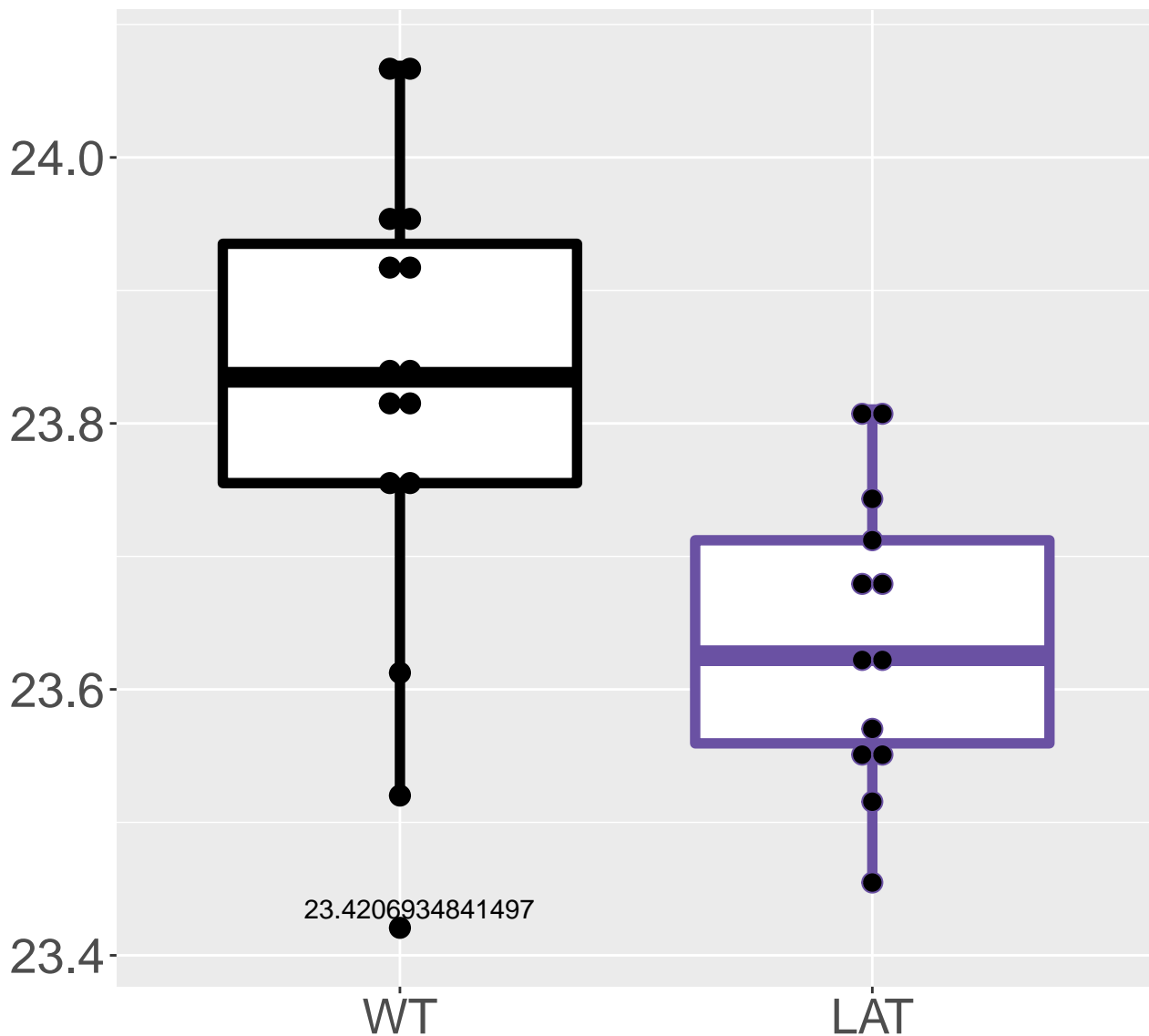
Q8R146_Acylamino-acid-releasing.
FDR = 0.027, FC = 0.31



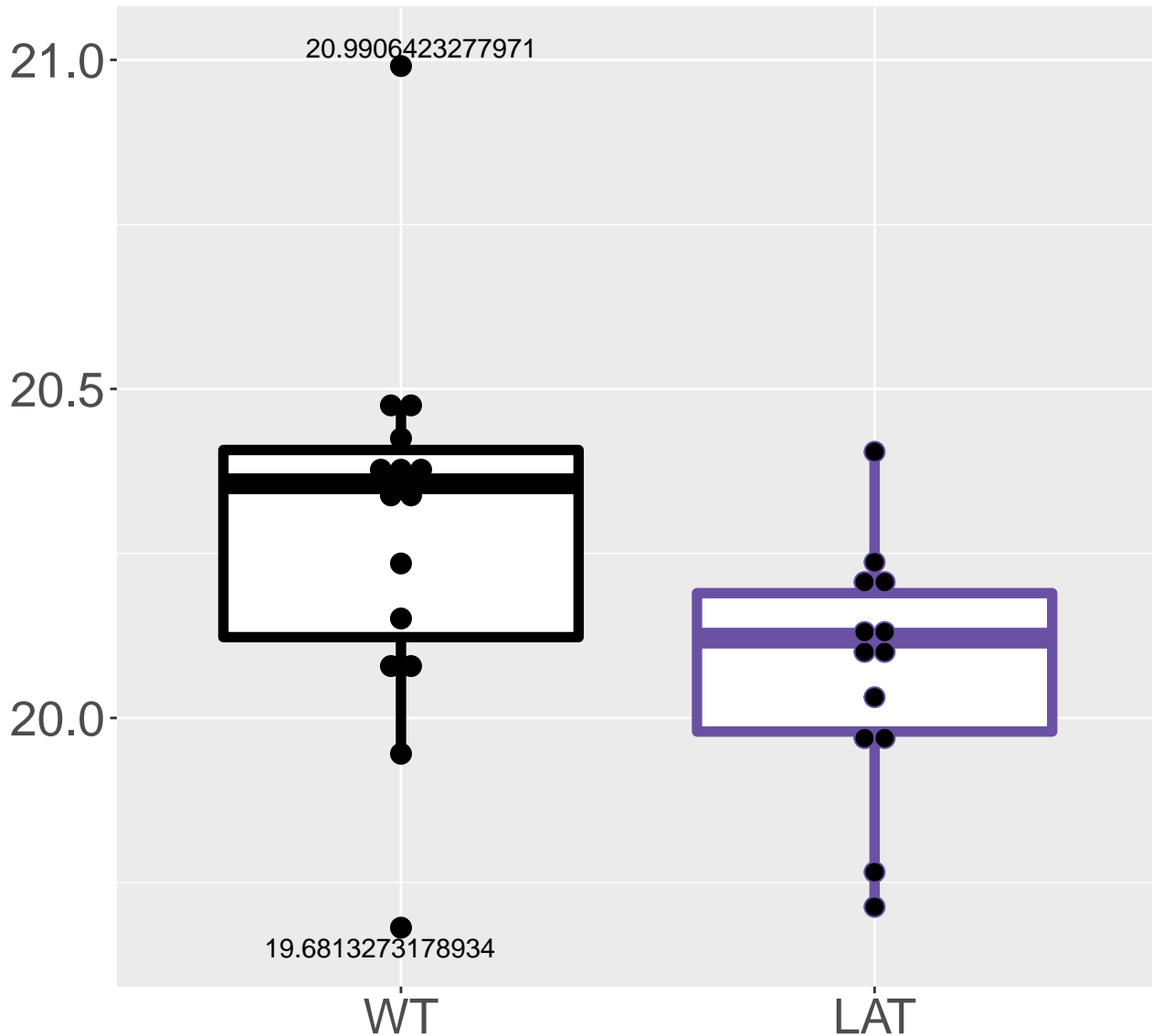
Q9D7G0_Ribose-phosphate pyropho.
FDR = 0.027, FC = -0.22, sex**



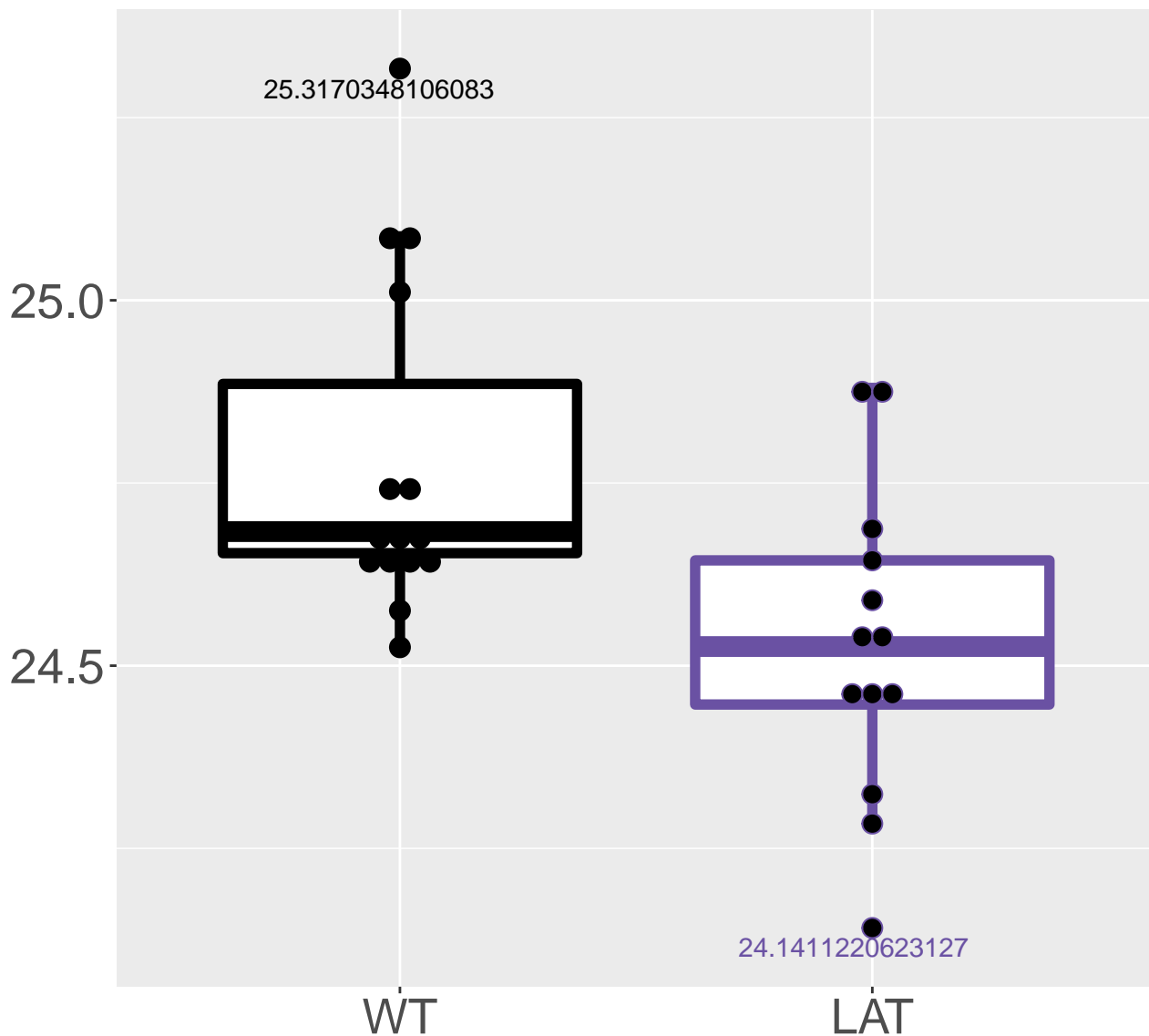
P61027_Ras-related protein Rab-.
FDR = 0.027, FC = -0.25, sex*



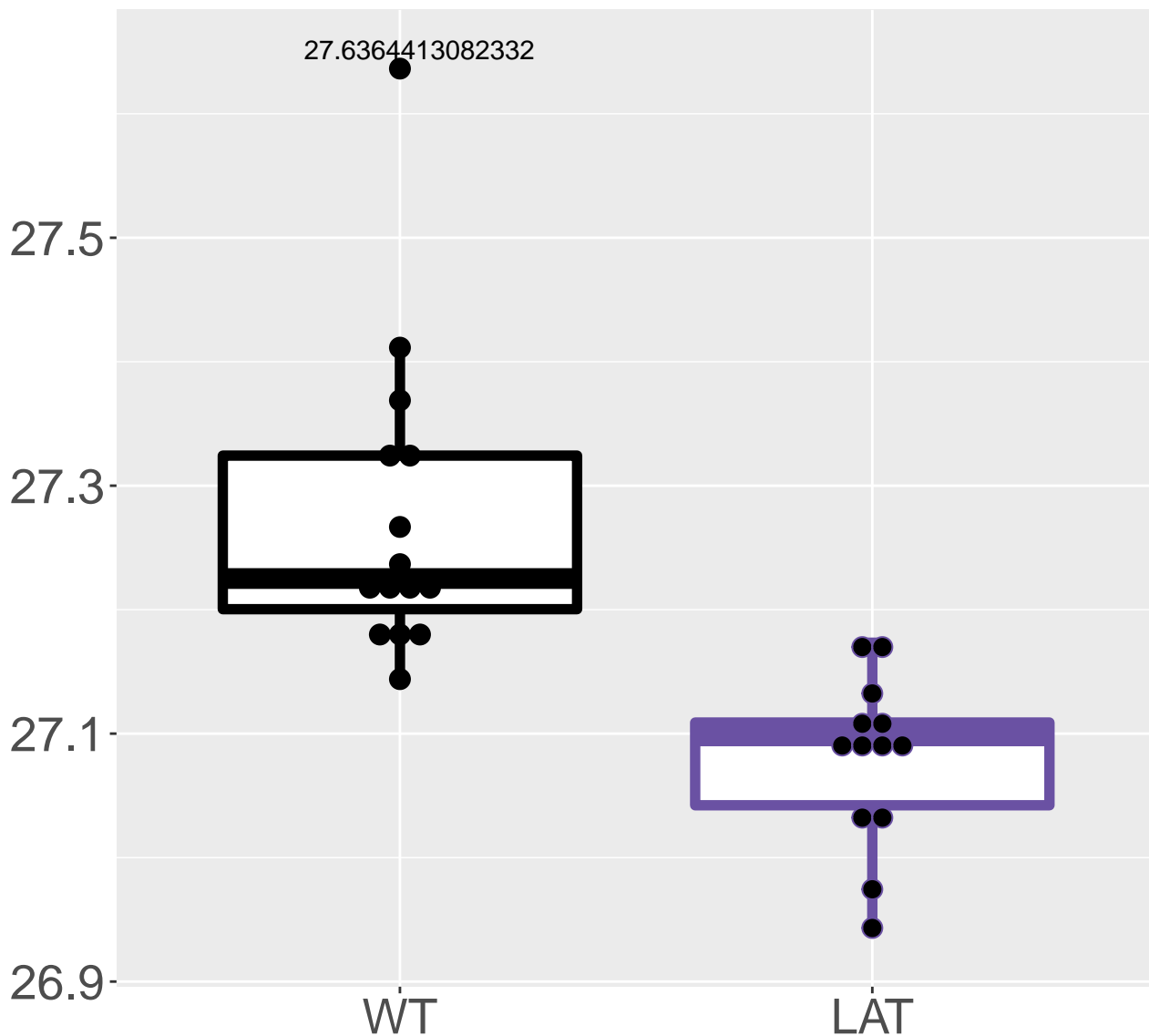
Q9CRD2_ER membrane protein comp.
FDR = 0.028, FC = -0.34, sex*



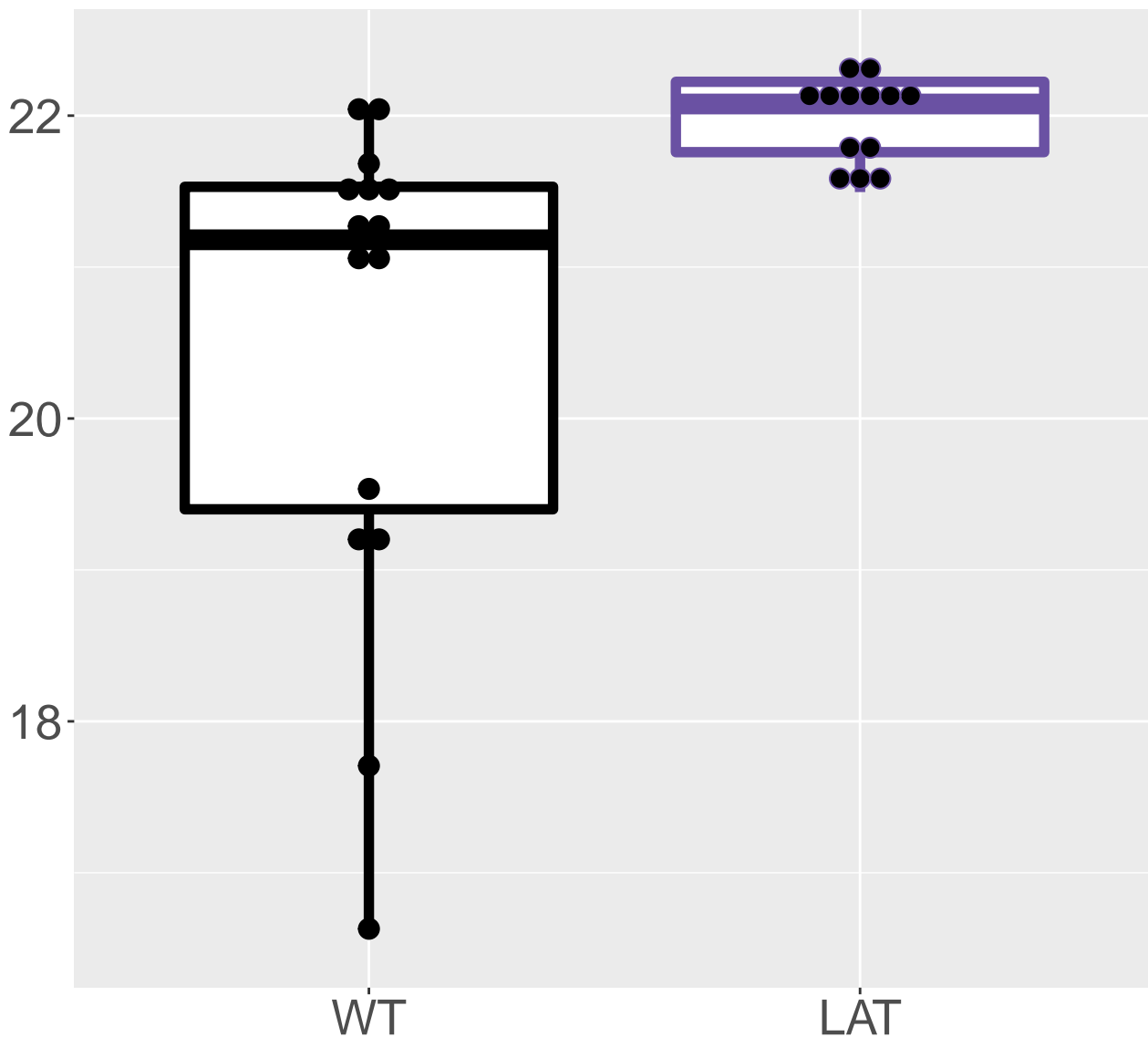
Q8BQ48_Centrosomal protein of 2.
FDR = 0.028, FC = -0.45



P62918_60S ribosomal protein L8
FDR = 0.028, FC = -0.27

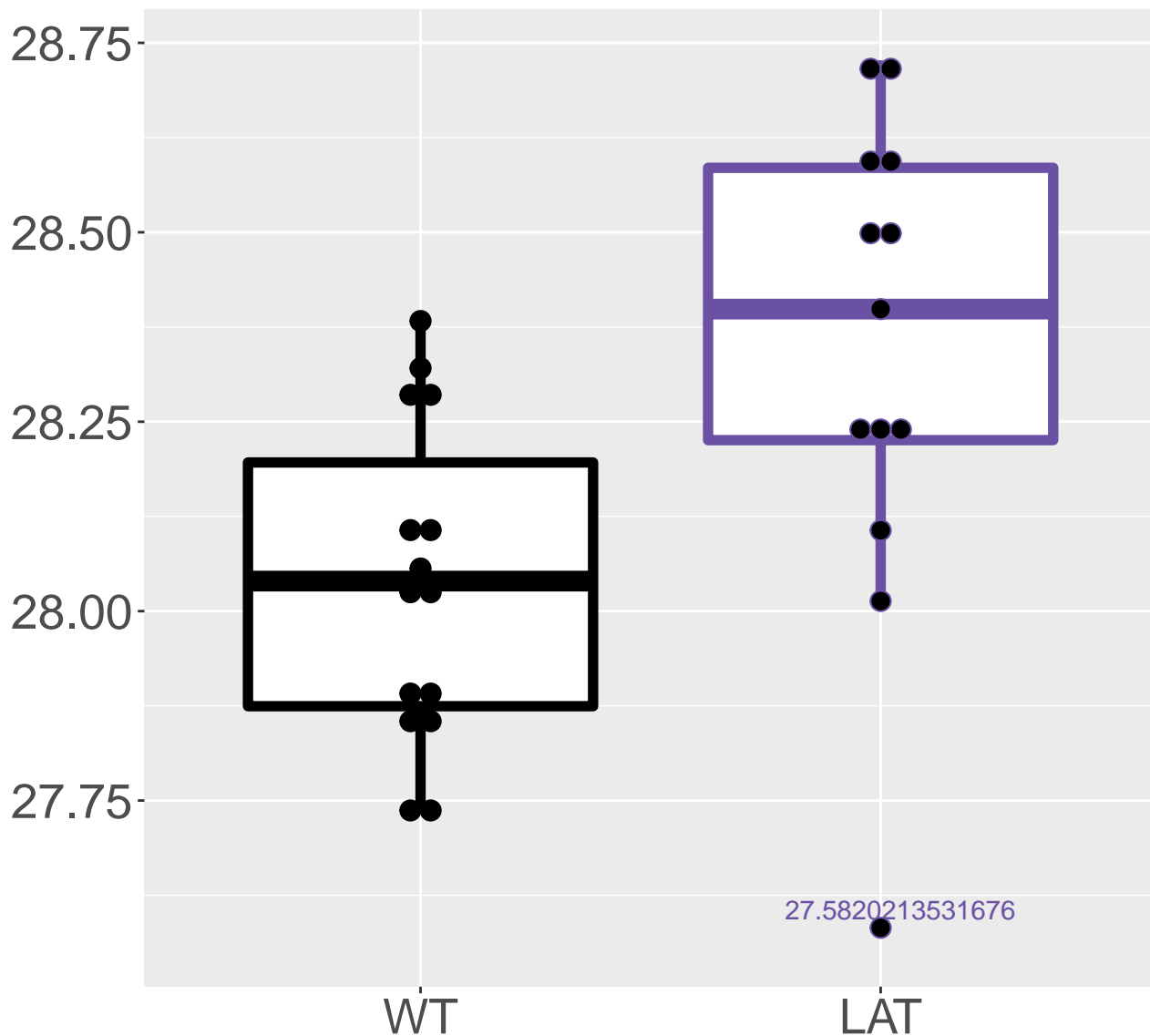


Q61009_Scavenger receptor class.
FDR = 0.028, FC = 1.6

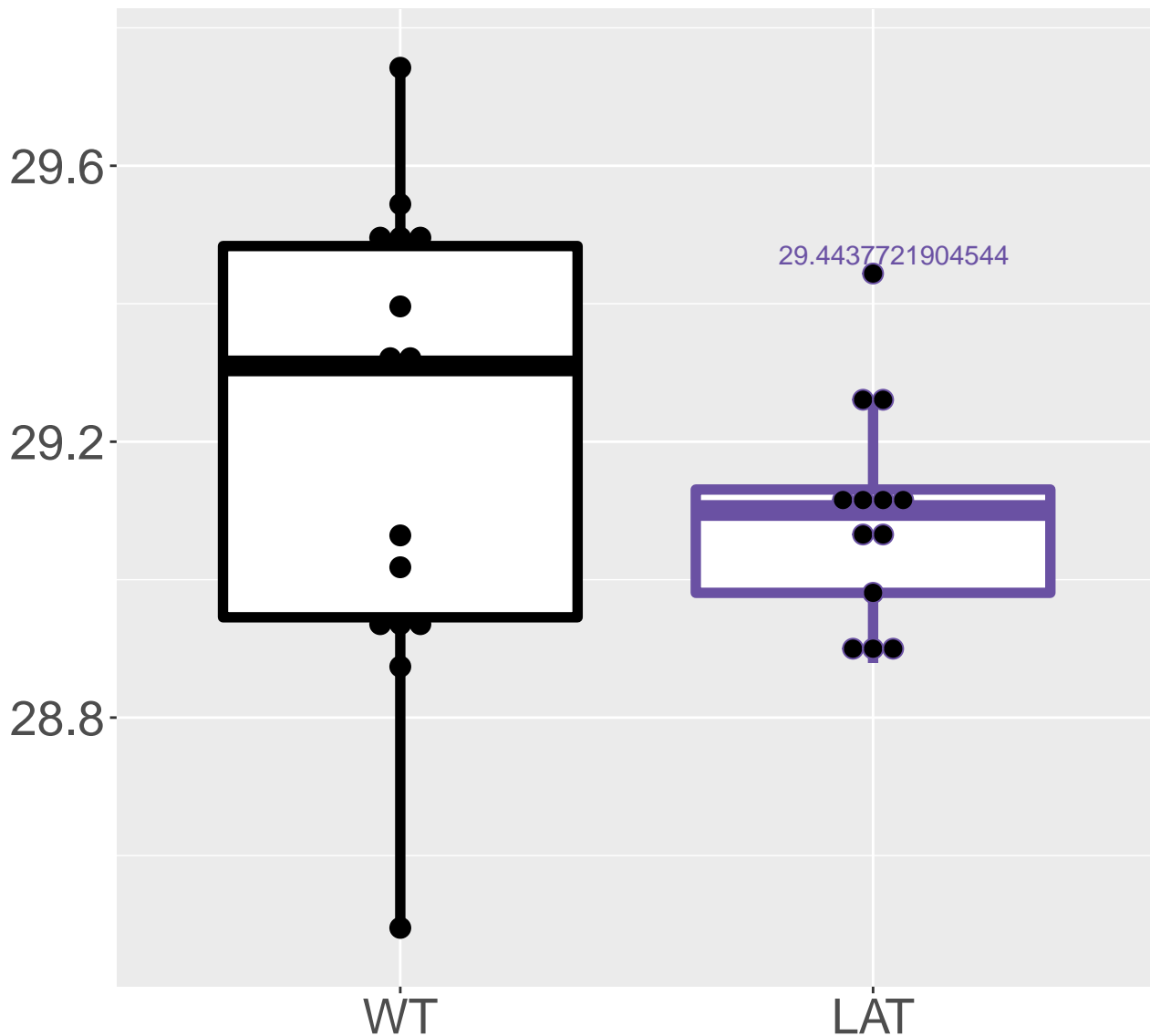


Q91V92_ATP-citrate synthase

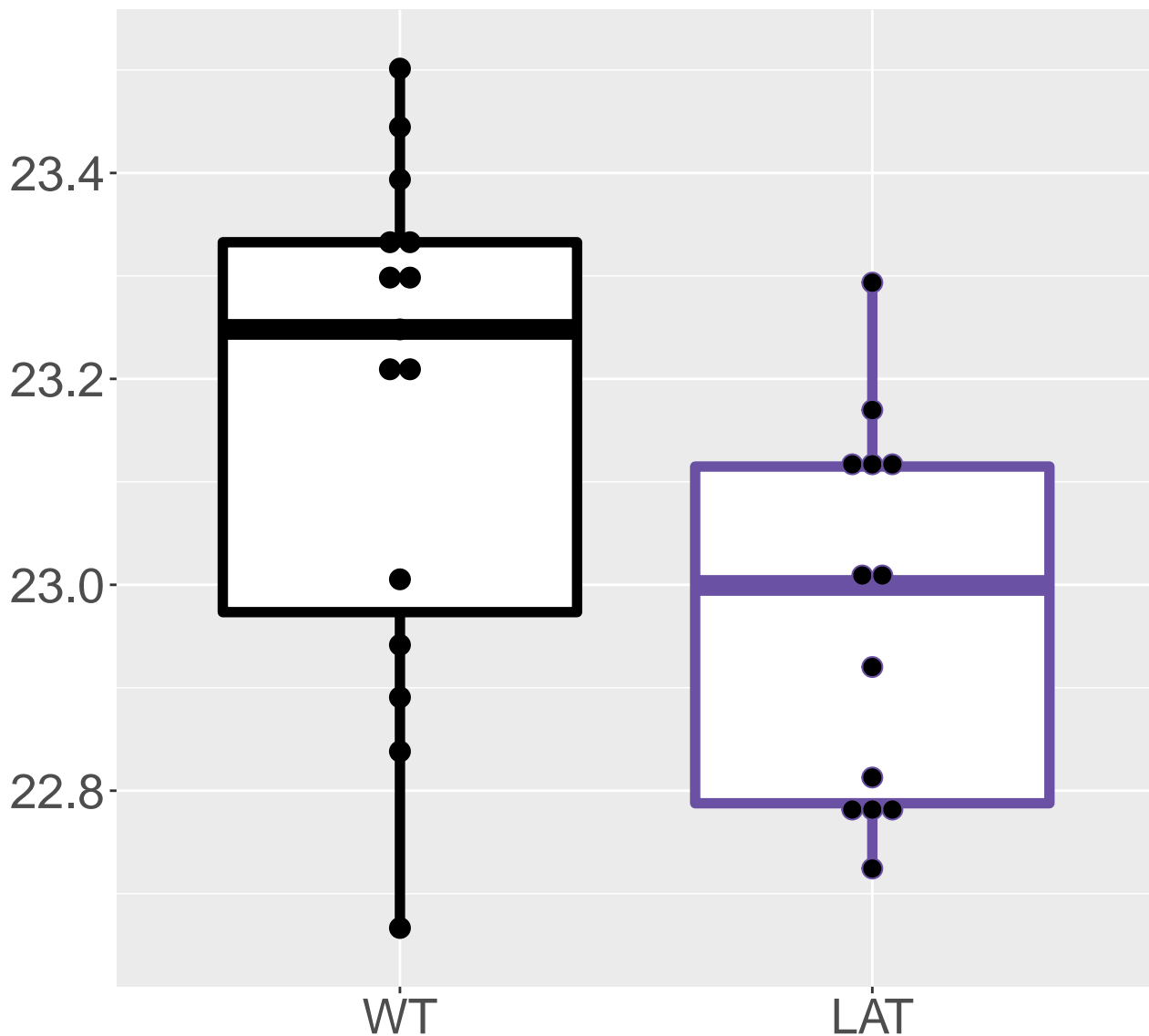
FDR = 0.028, FC = 0.4



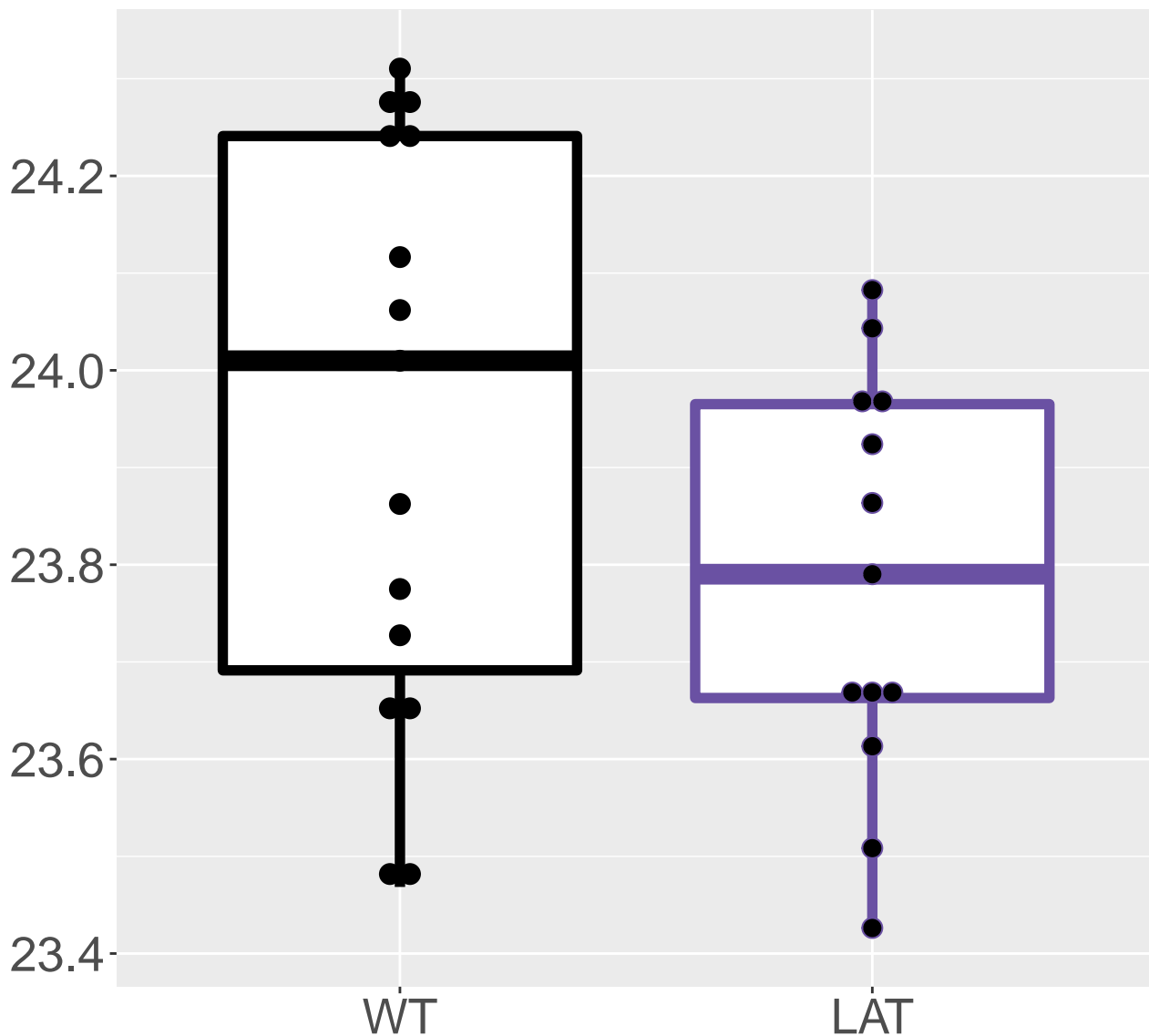
P52760_2-iminobutanoate/2-imino.
FDR = 0.029, FC = -0.32, sex***



Q99KV1_DnaJ homolog subfamily B.
FDR = 0.029, FC = -0.29, sex*

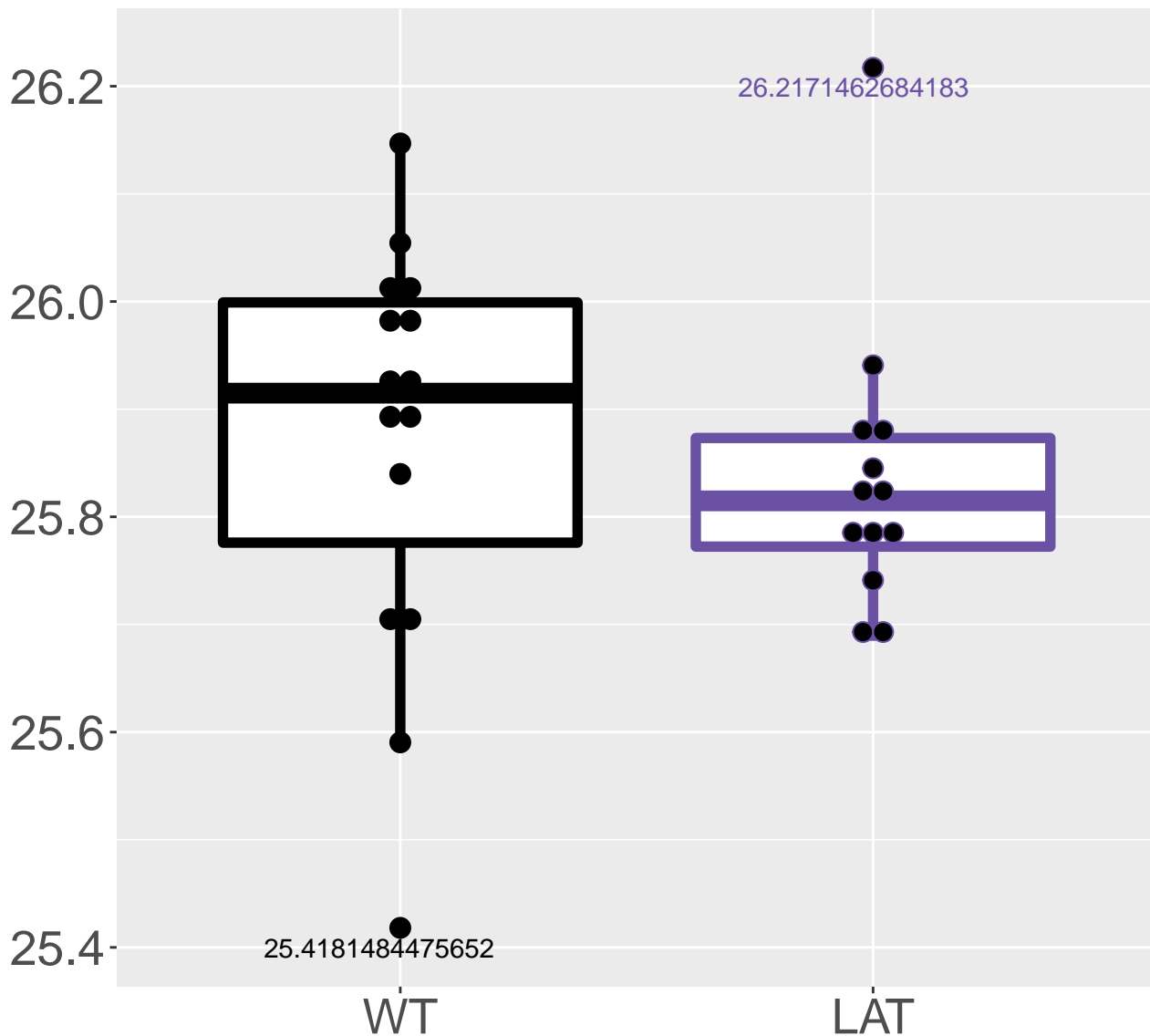


Q9JLI6_Selenocysteine lyase
FDR = 0.029, FC = -0.22, sex***

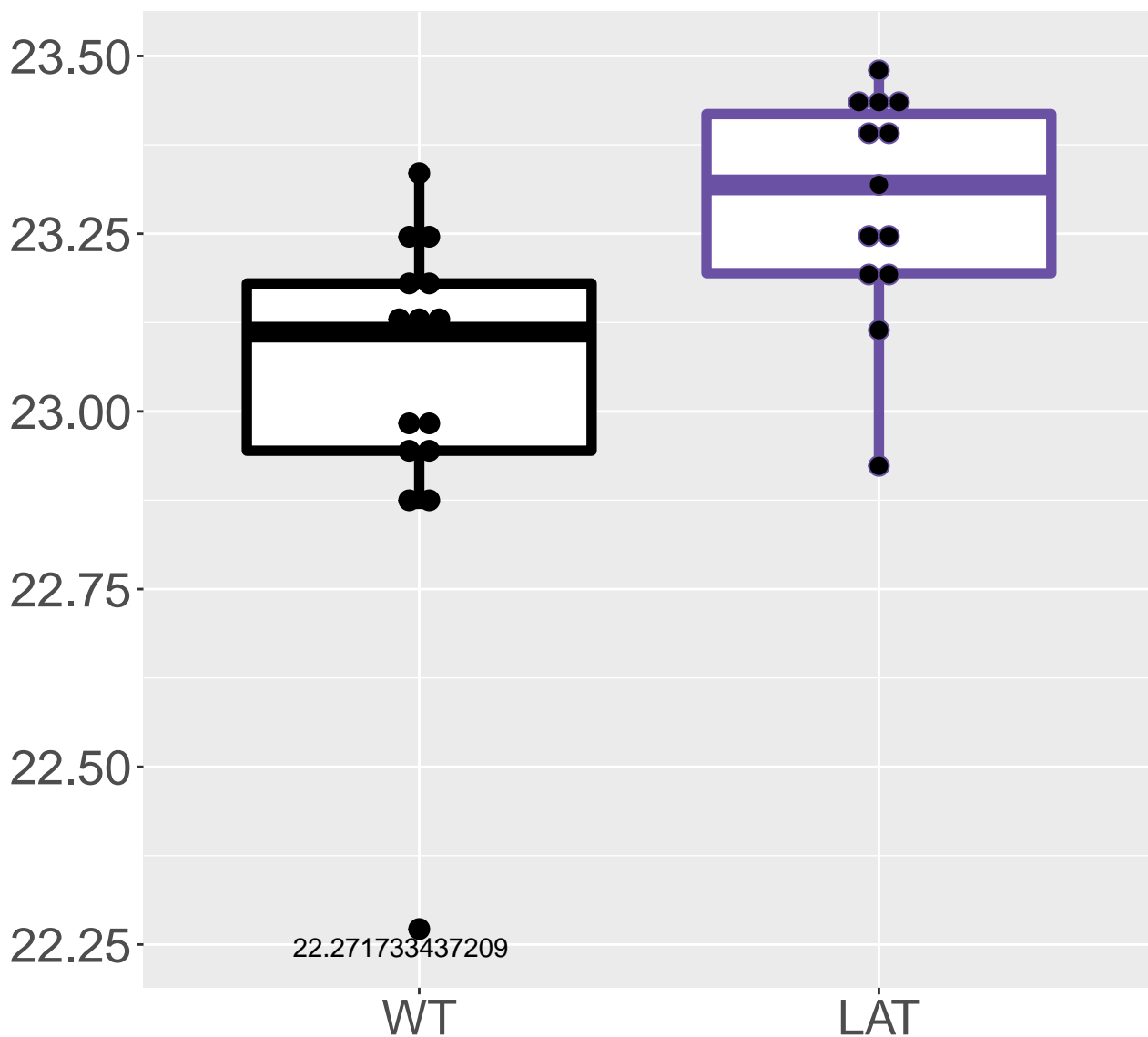


Q91X52_L-xylulose reductase

FDR = 0.029, FC = -0.18, sex*

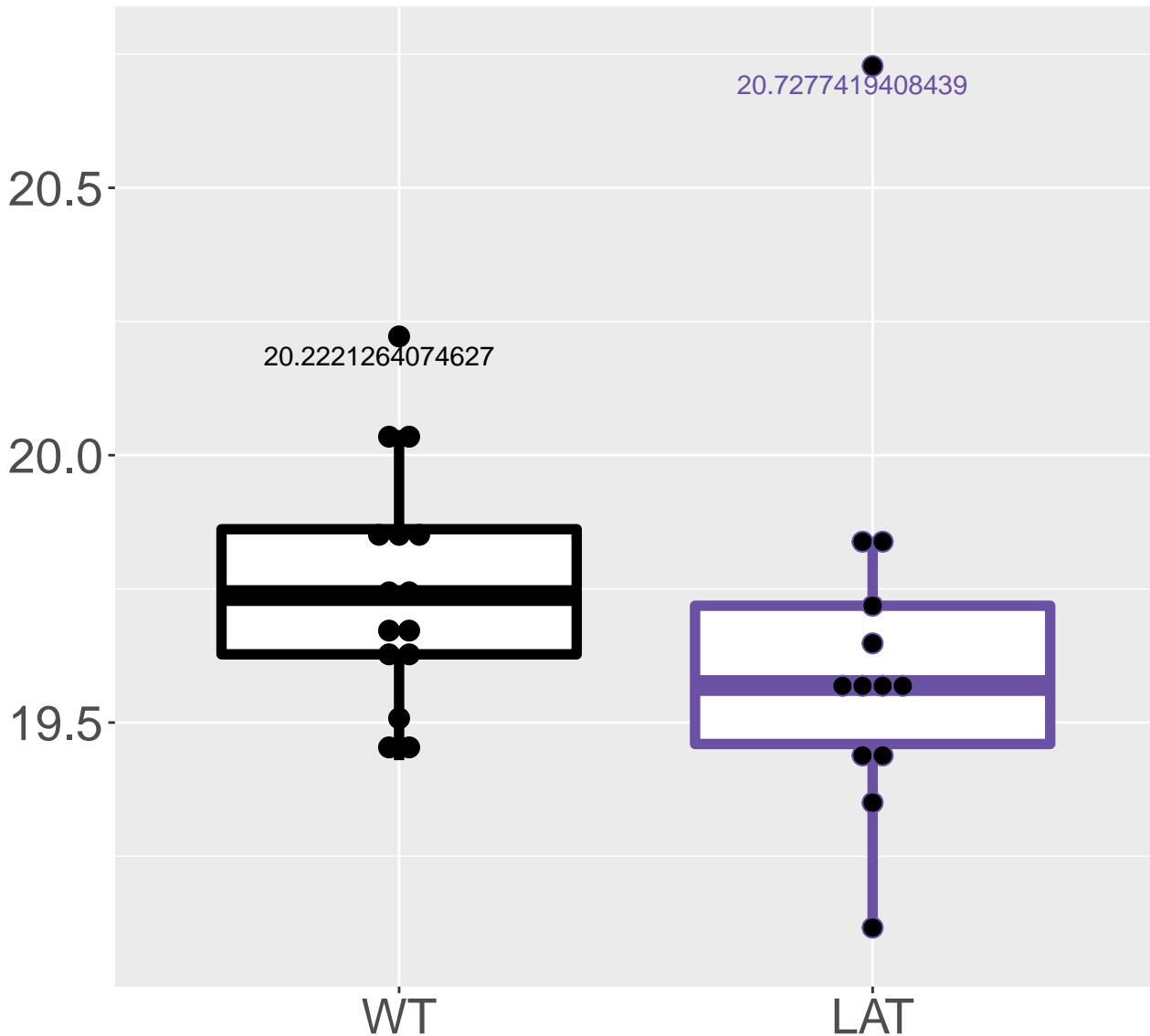


Q8BUV3_Gephyrin
FDR = 0.029, FC = 0.46

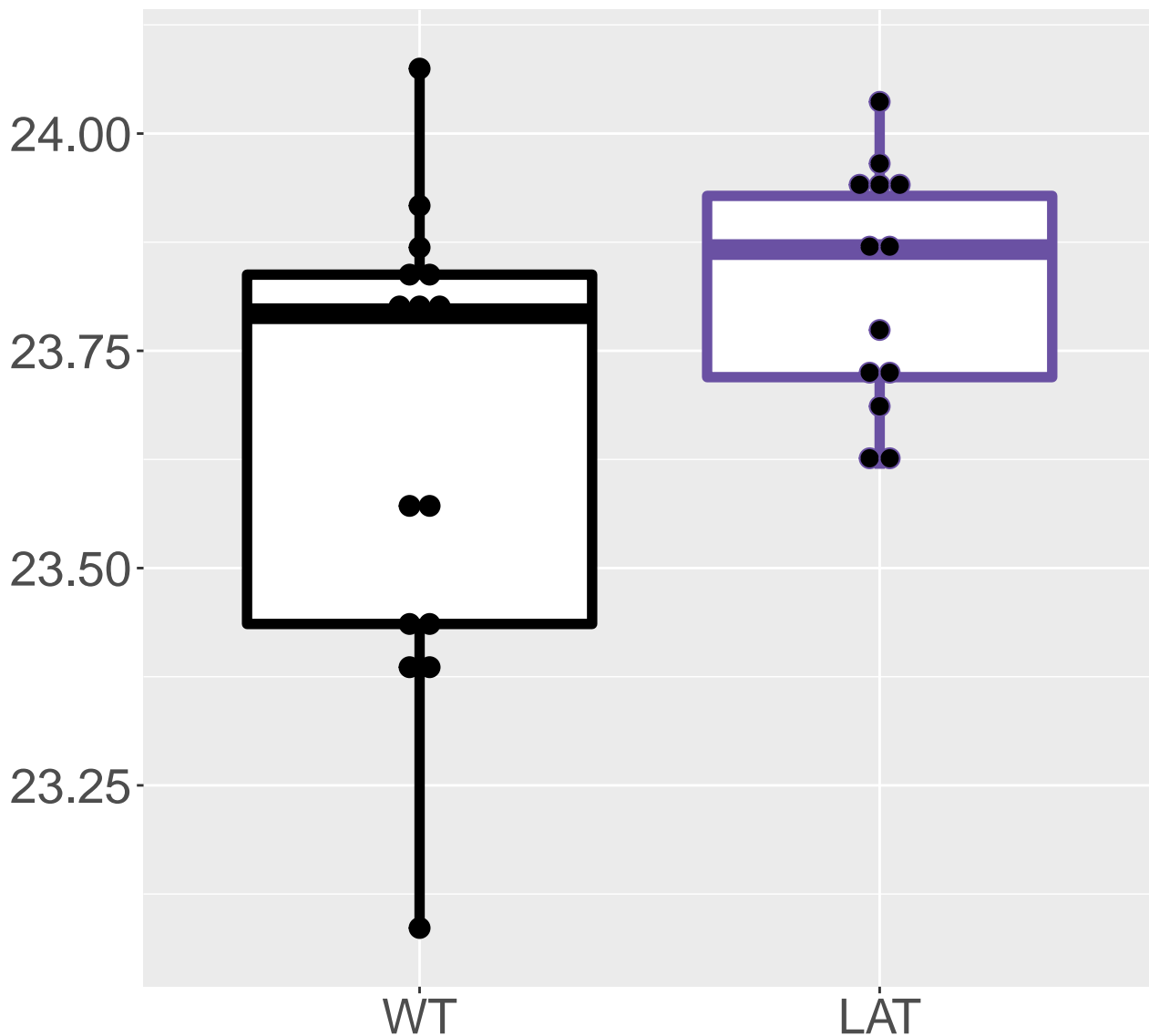


P70665_Sialate O-acetyltransferase

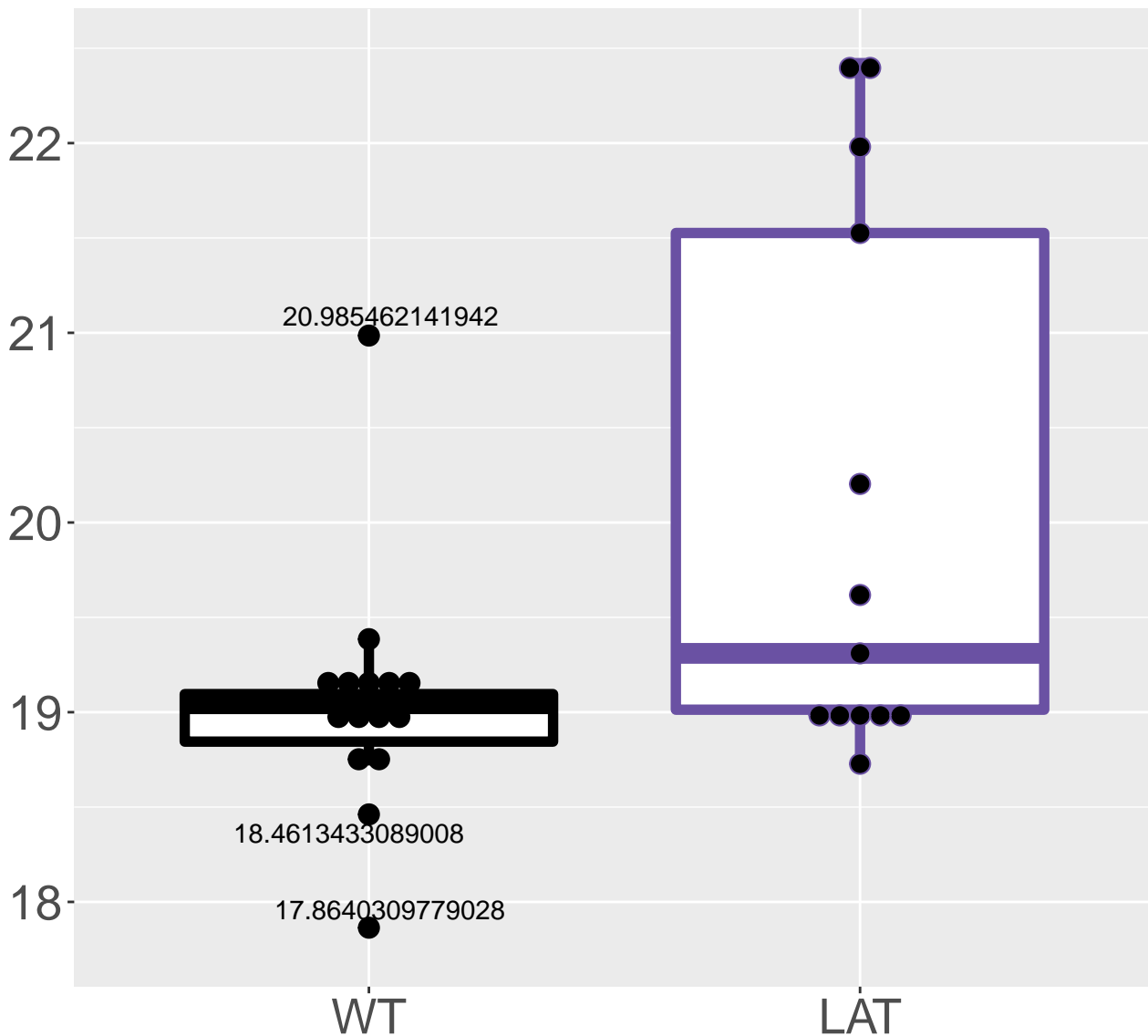
FDR = 0.029, FC = -0.43



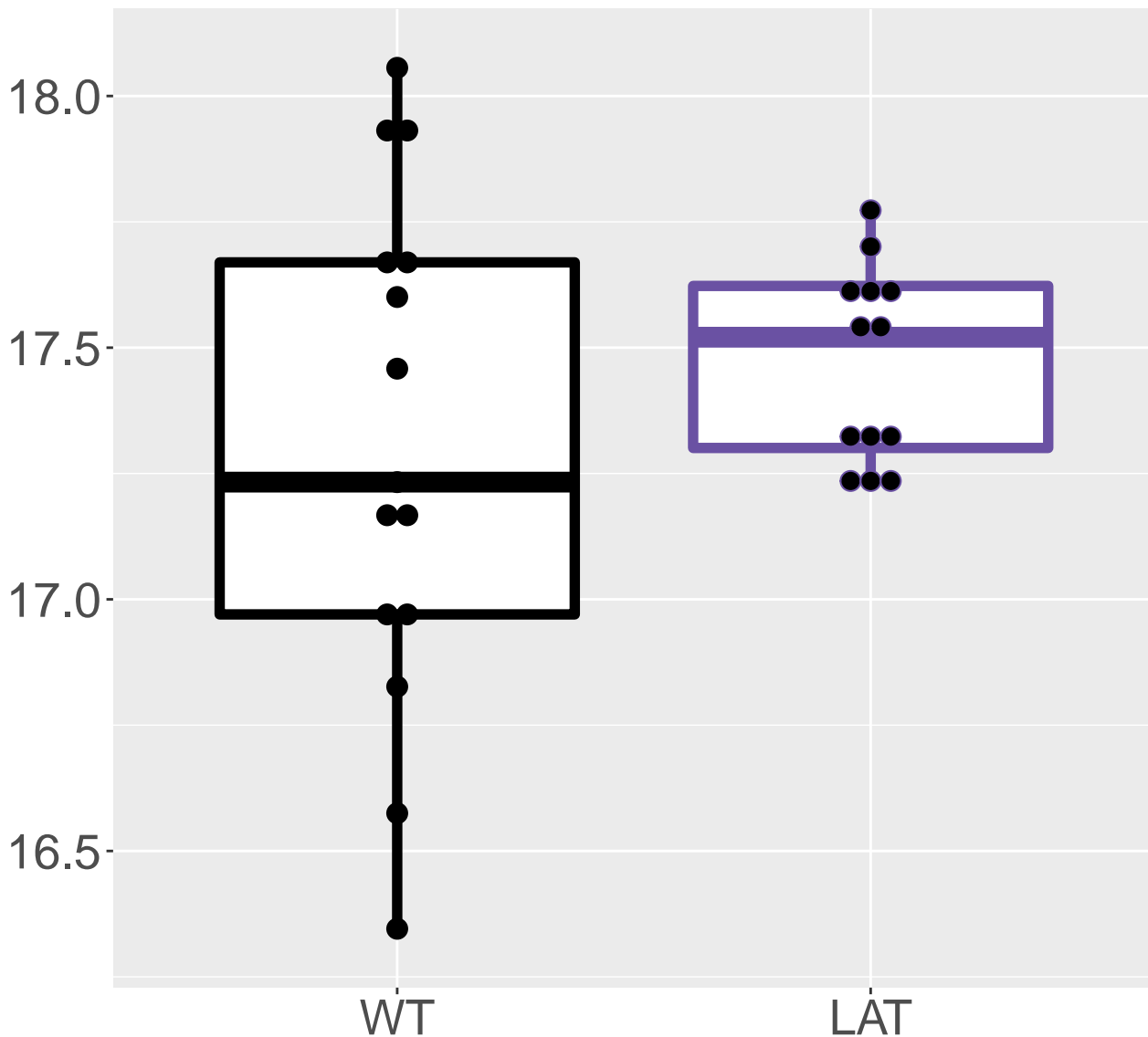
Q571I9_Aldehyde dehydrogenase f.
FDR = 0.029, FC = 0.39, sex*



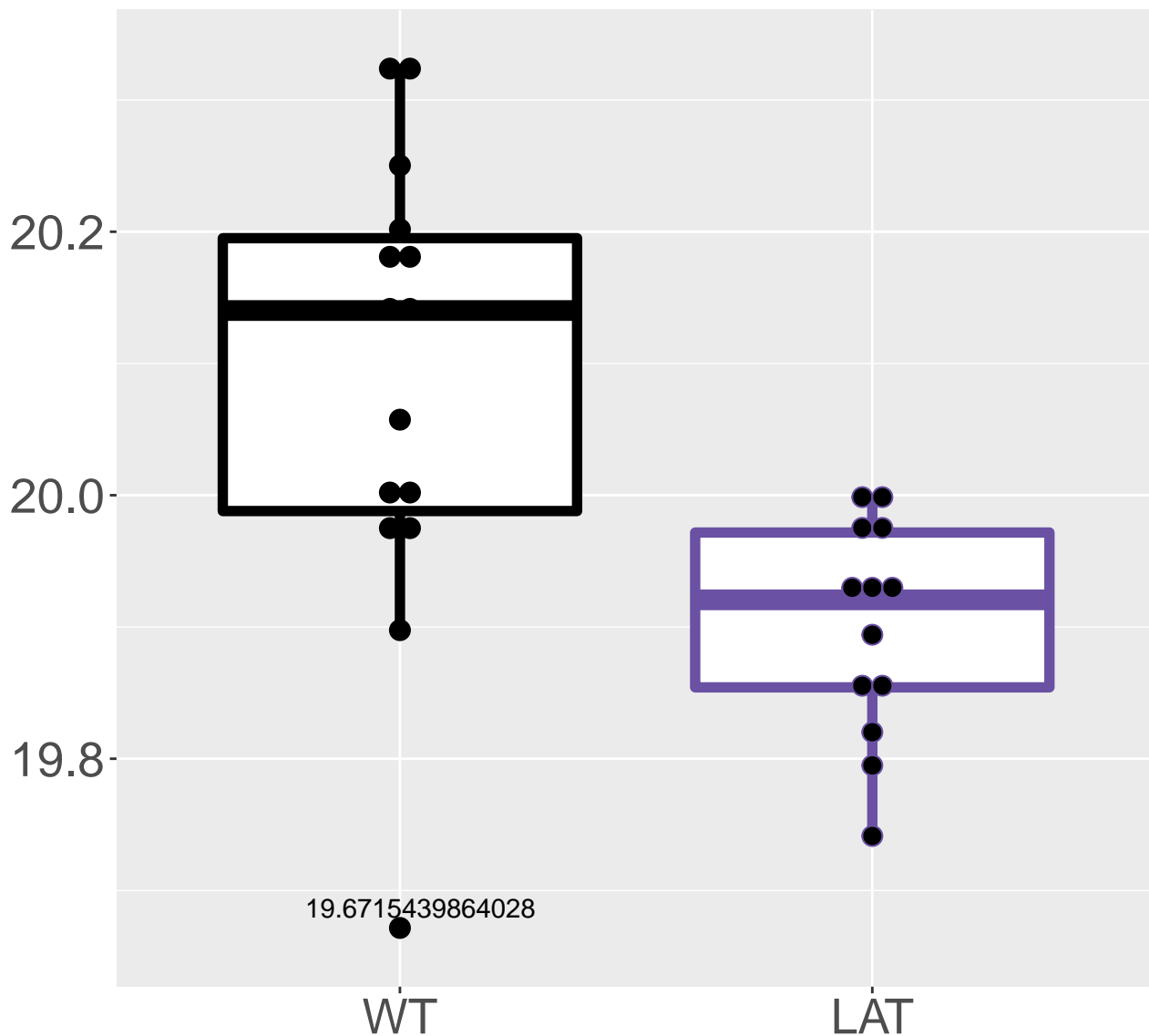
FDR = 0.029, FC = 1.9



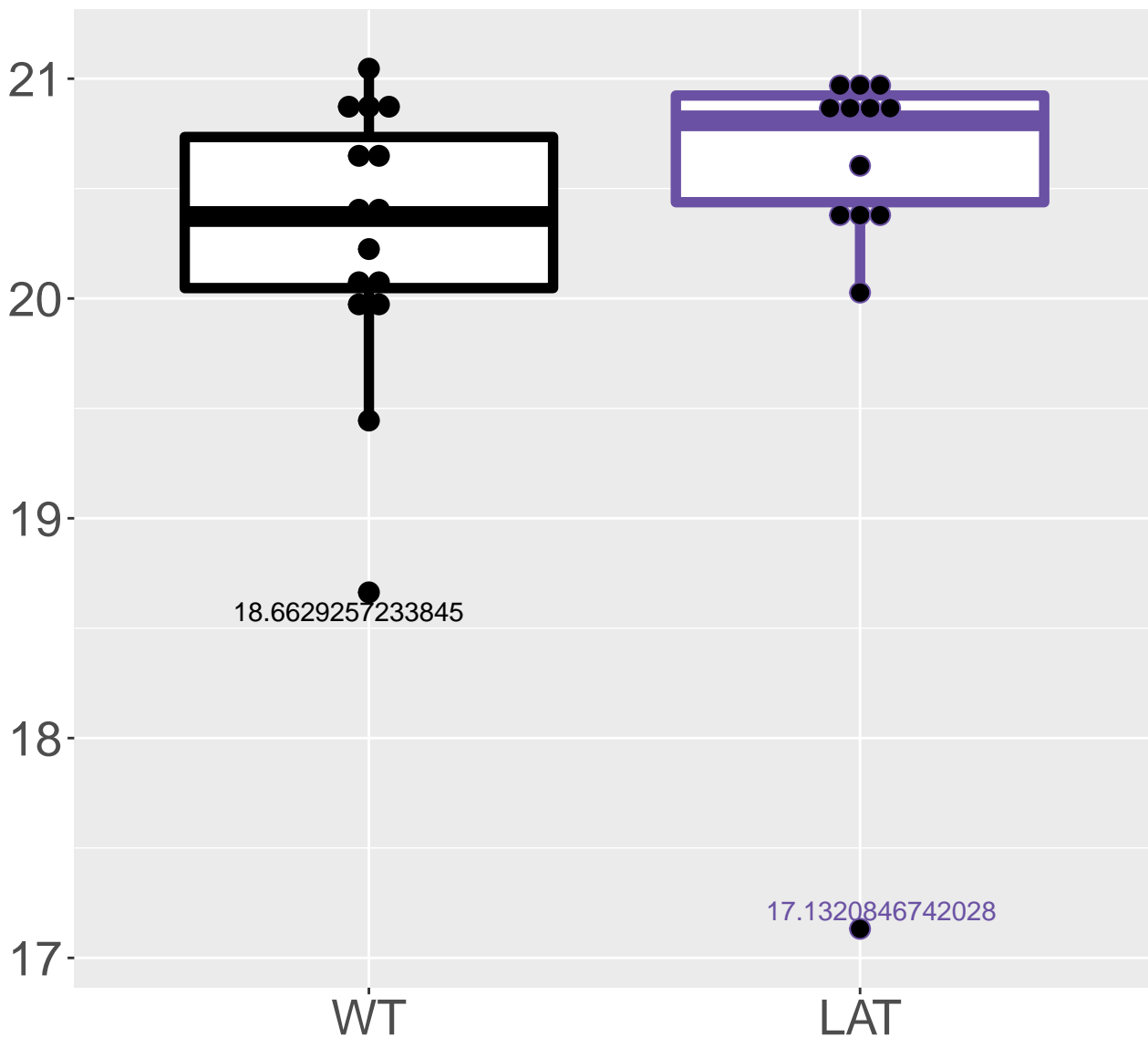
Q8QZY9_Splicing factor 3B subun.
FDR = 0.029, FC = 0.62, sex**



Q9Z1Q2_Protein ABHD16A
FDR = 0.029, FC = -0.27

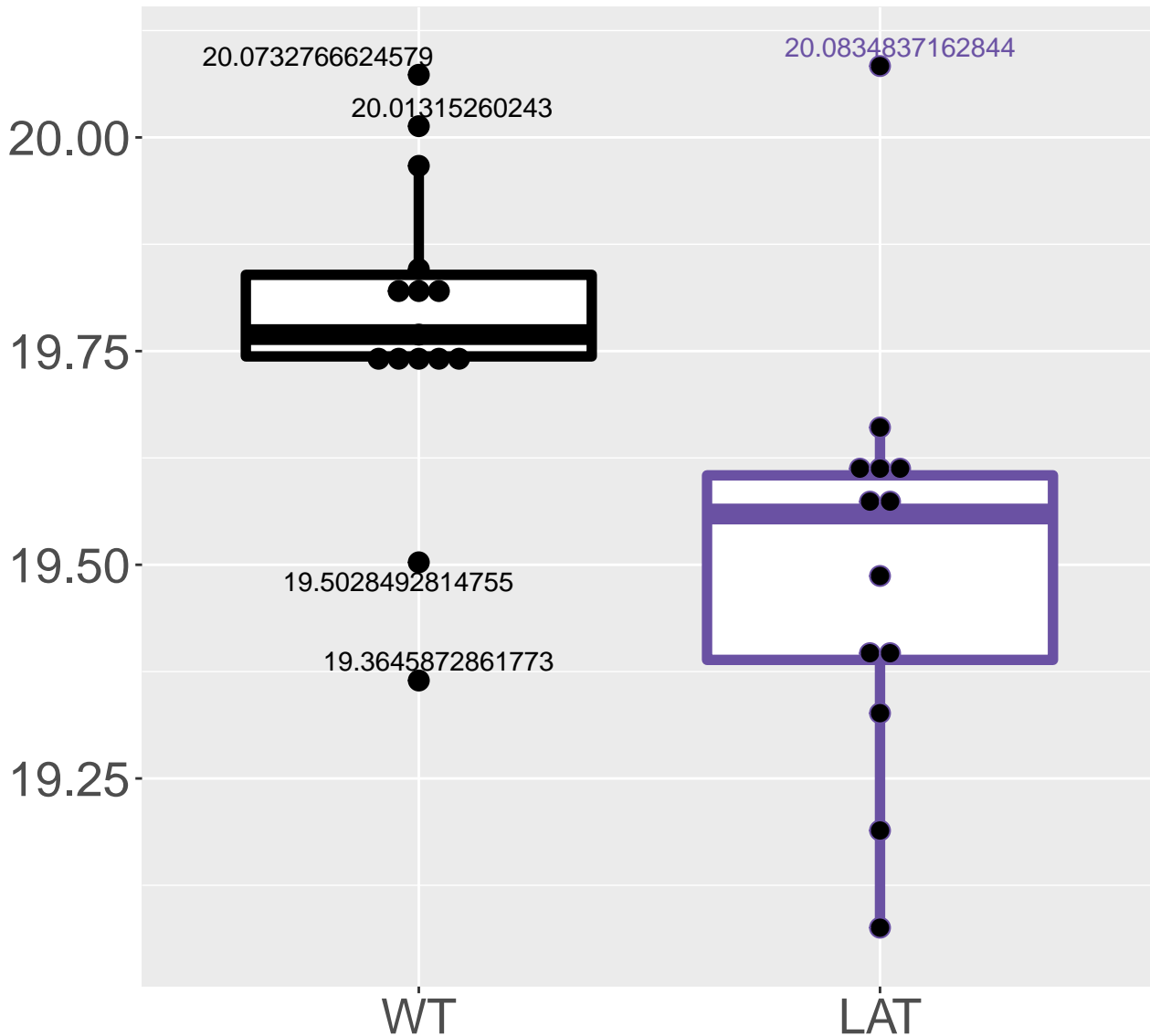


FDR = 0.03, FC = 0.87, sex**

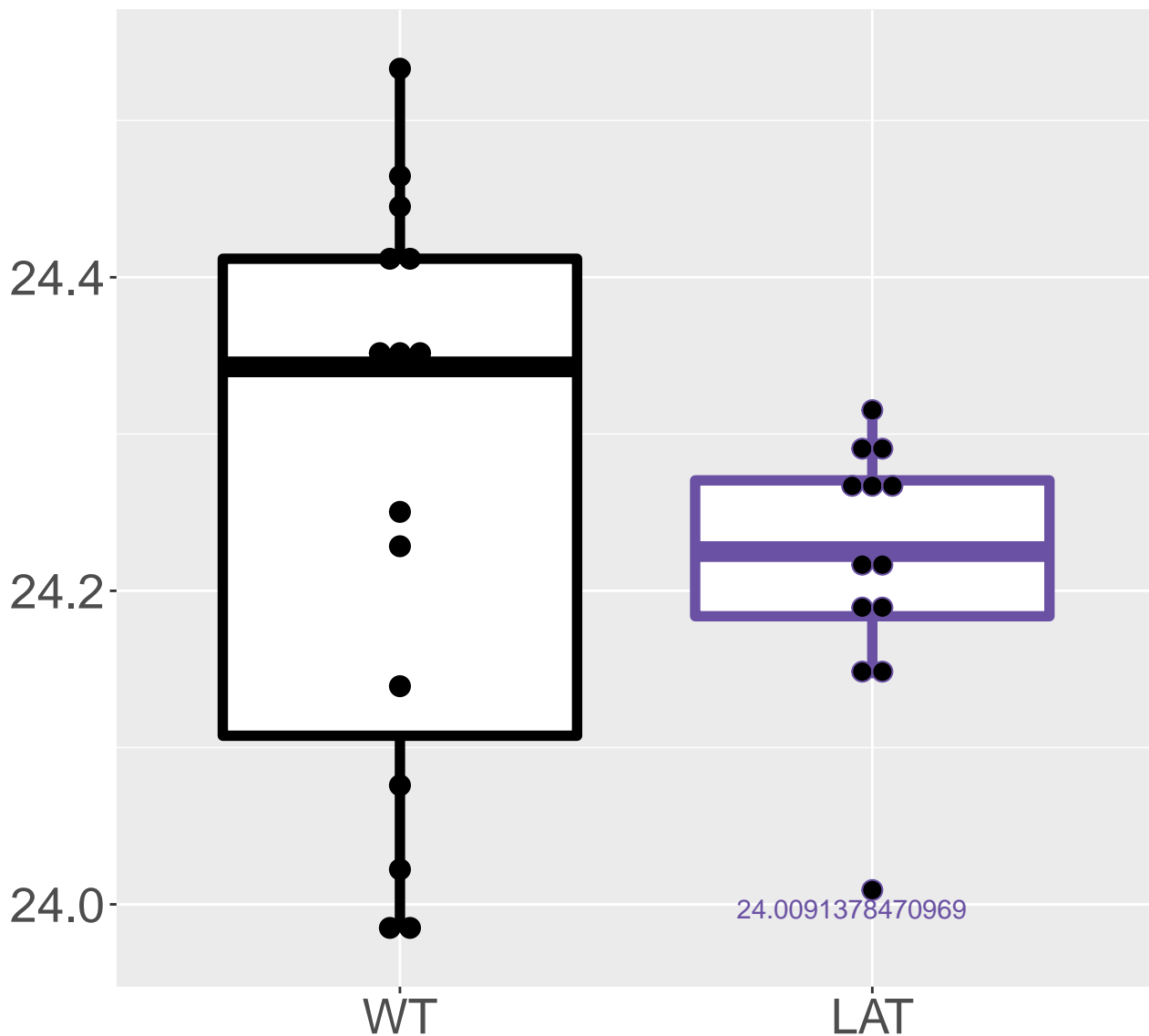


P60603_Reactive oxygen species .

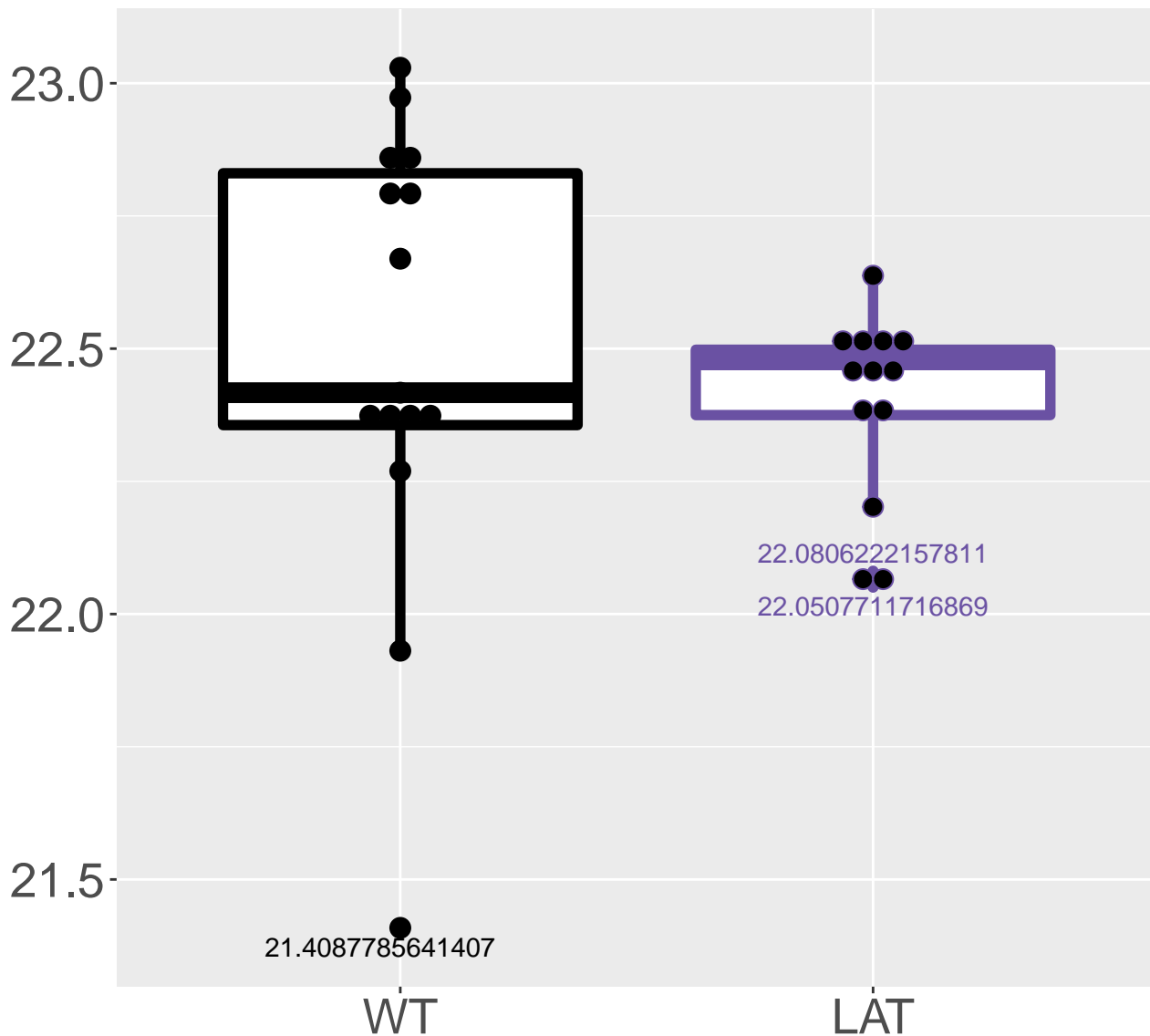
FDR = 0.03, FC = -0.4



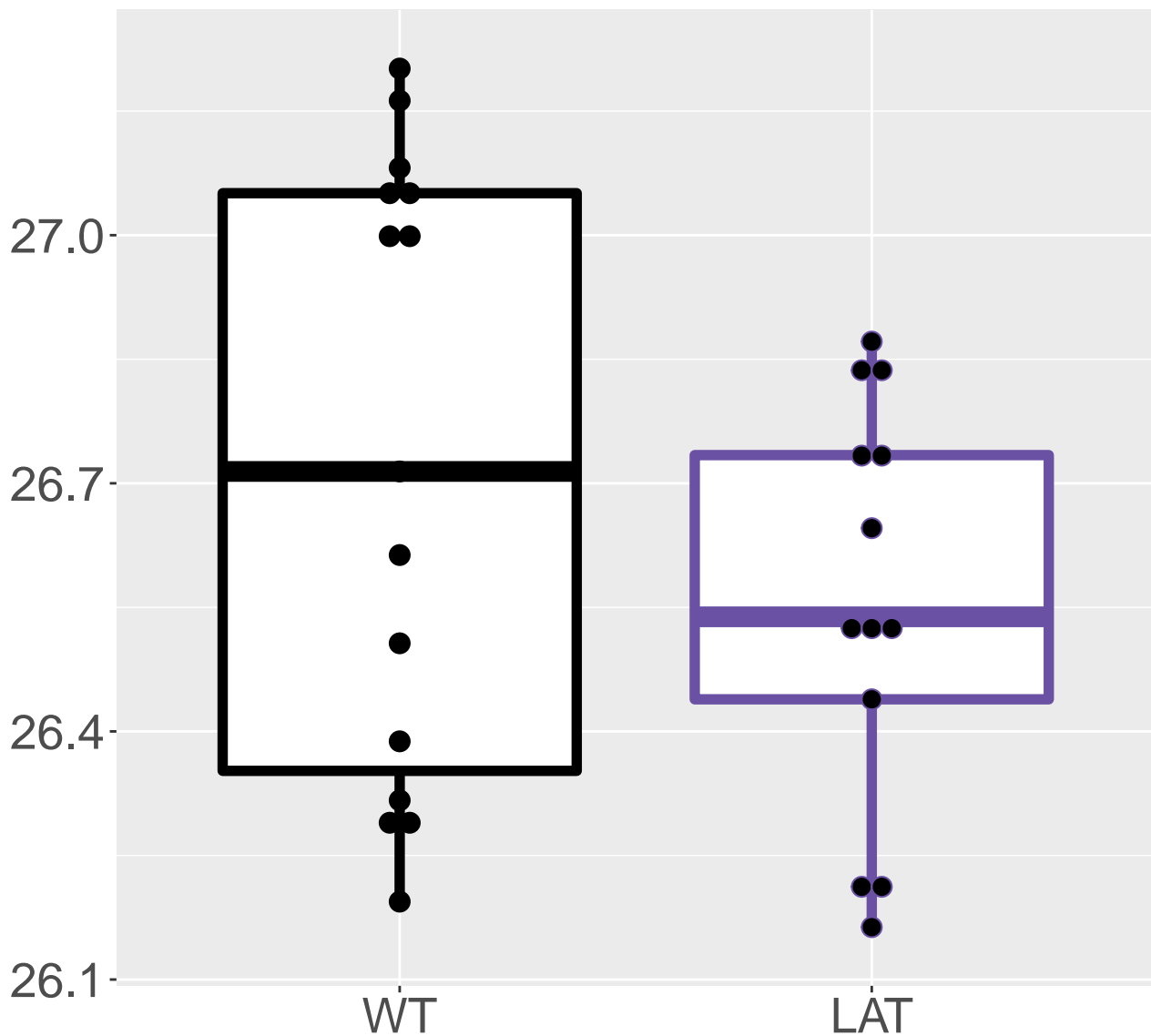
Q5U5V2_Hydroxylysine kinase
FDR = 0.03, FC = -0.18, sex**



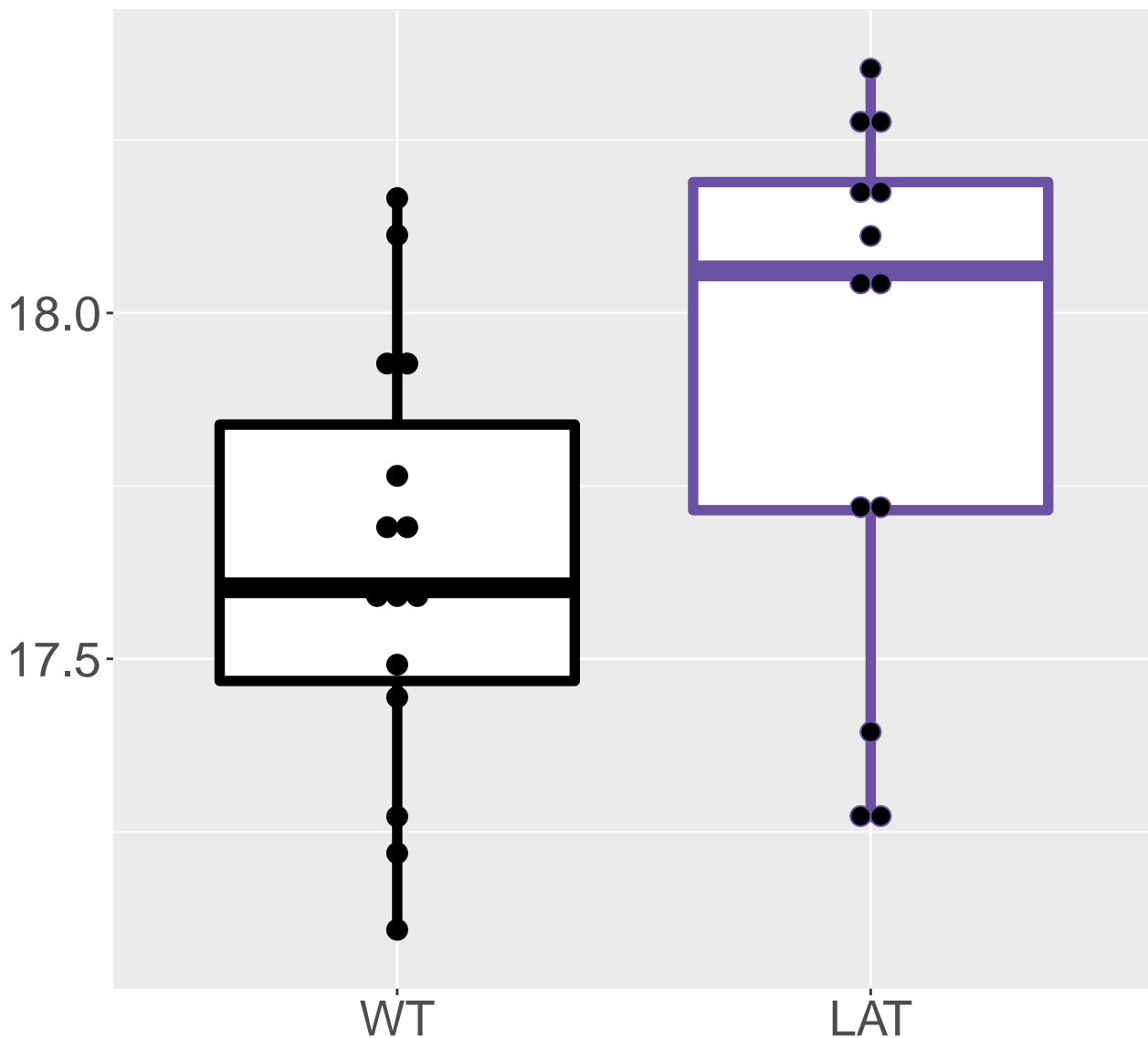
P40630_Transcription factor A, .
FDR = 0.031, FC = -0.35, sex**



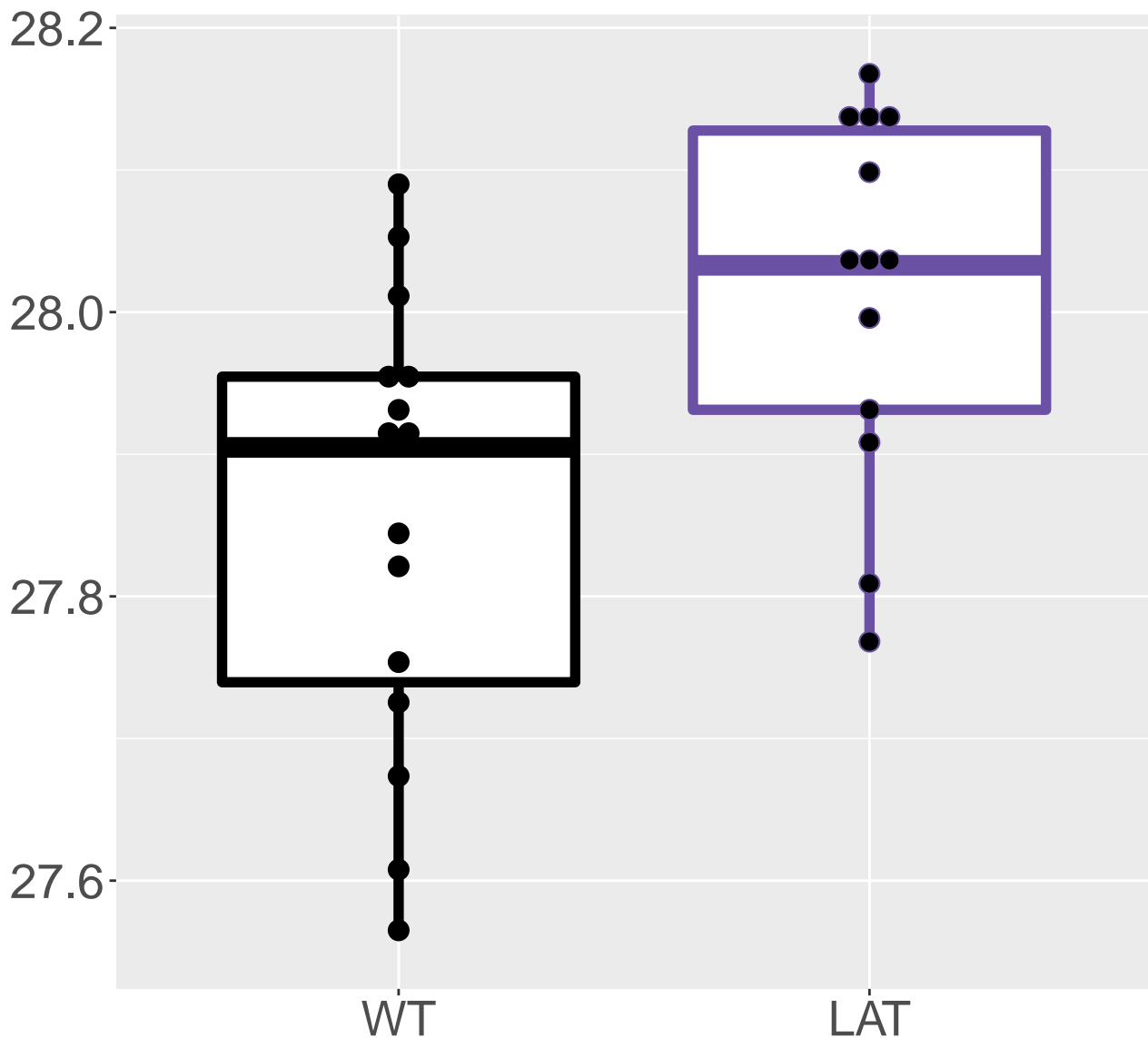
Q9D023_Mitochondrial pyruvate c.
FDR = 0.031, FC = -0.26, sex***



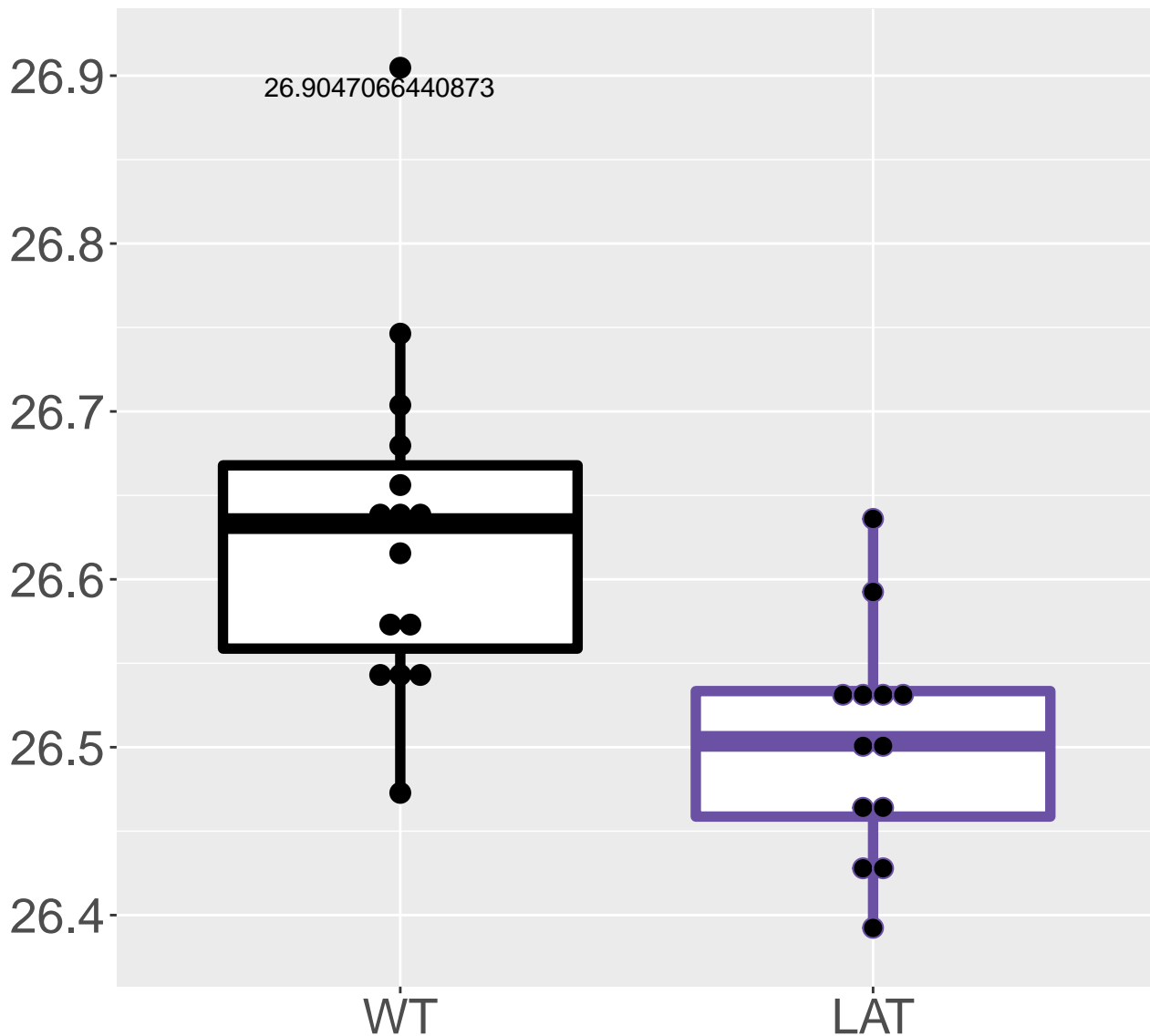
Q8BVG4_Dipeptidyl peptidase 9
FDR = 0.032, FC = 0.6



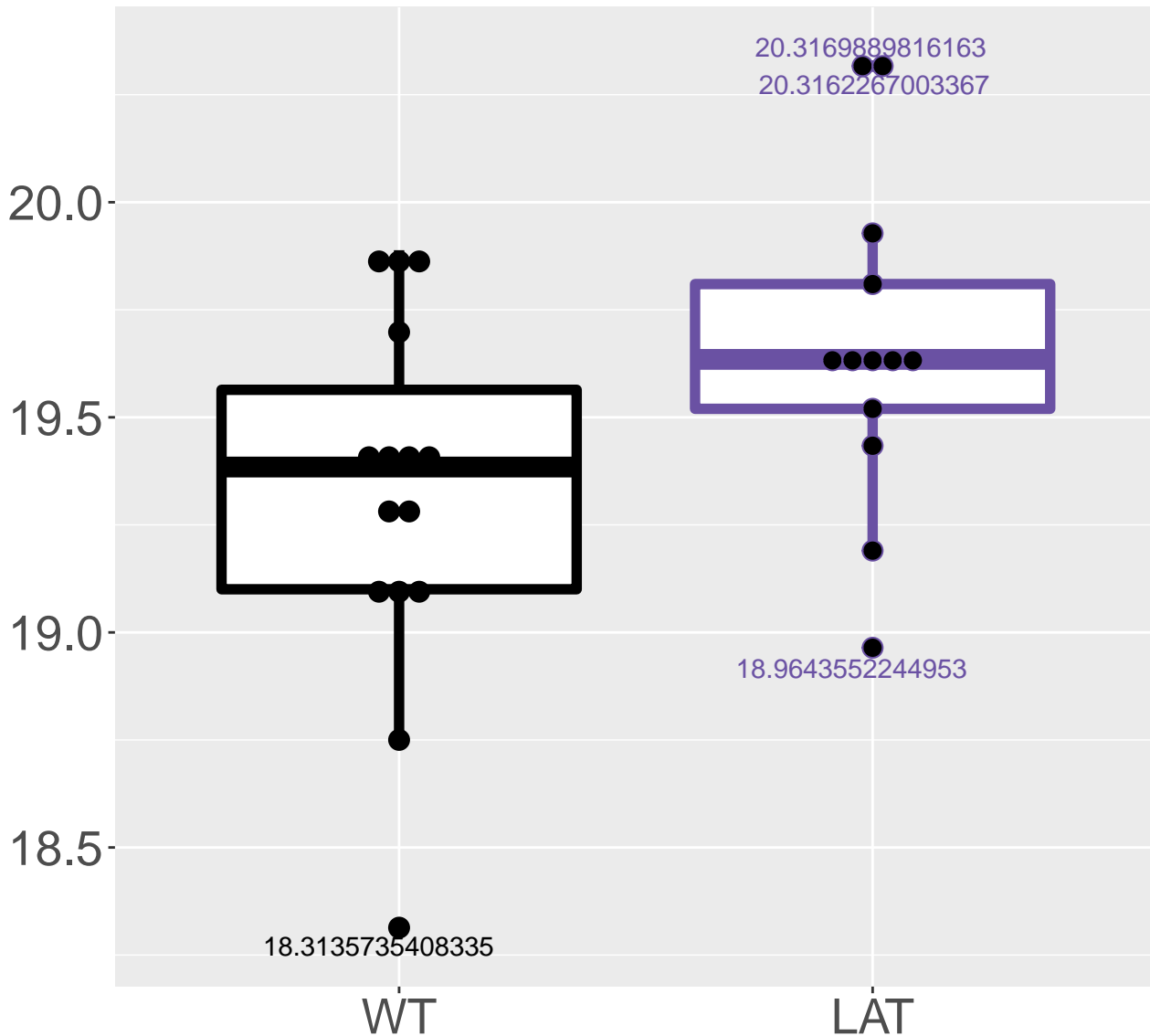
Q99KI0_Aconitate hydratase, mit.
FDR = 0.032, FC = 0.3, sex*



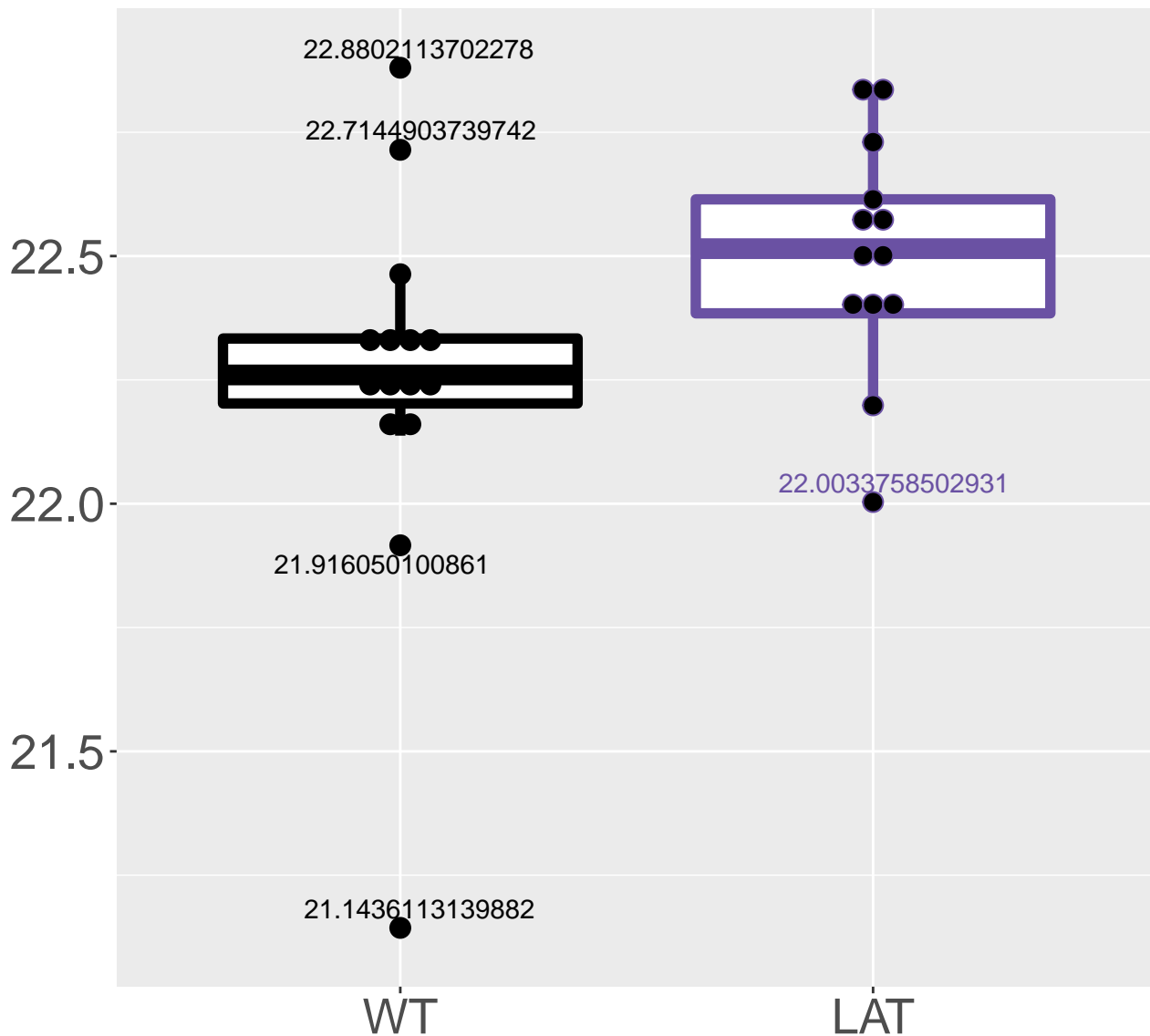
P62717_60S ribosomal protein L1.
FDR = 0.032, FC = -0.19



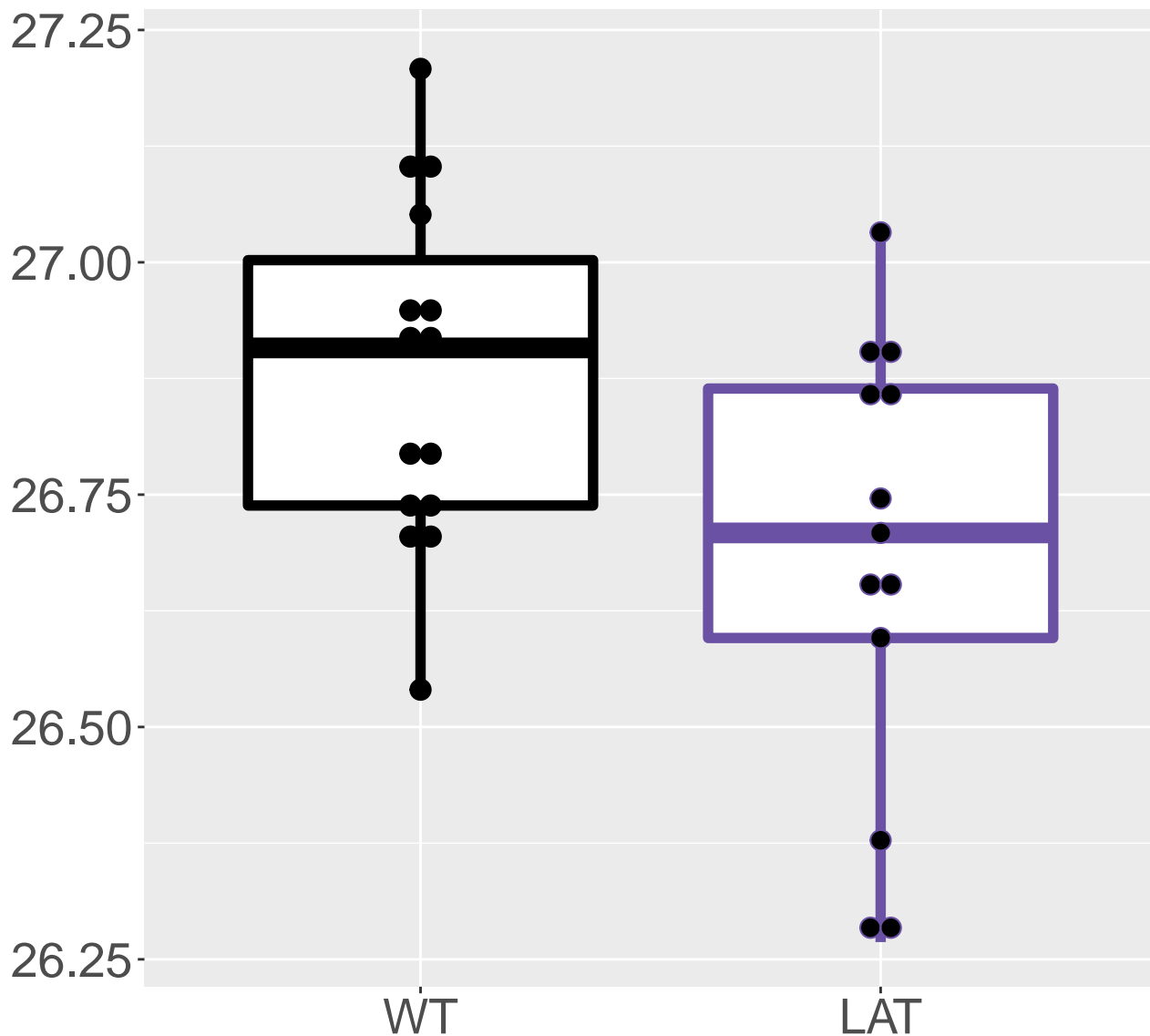
Q91ZX7_Prolow-density lipoprote.
FDR = 0.032, FC = 0.66, sex*



Q9CRC0_Vitamin K epoxide reduct.
FDR = 0.032, FC = 0.59

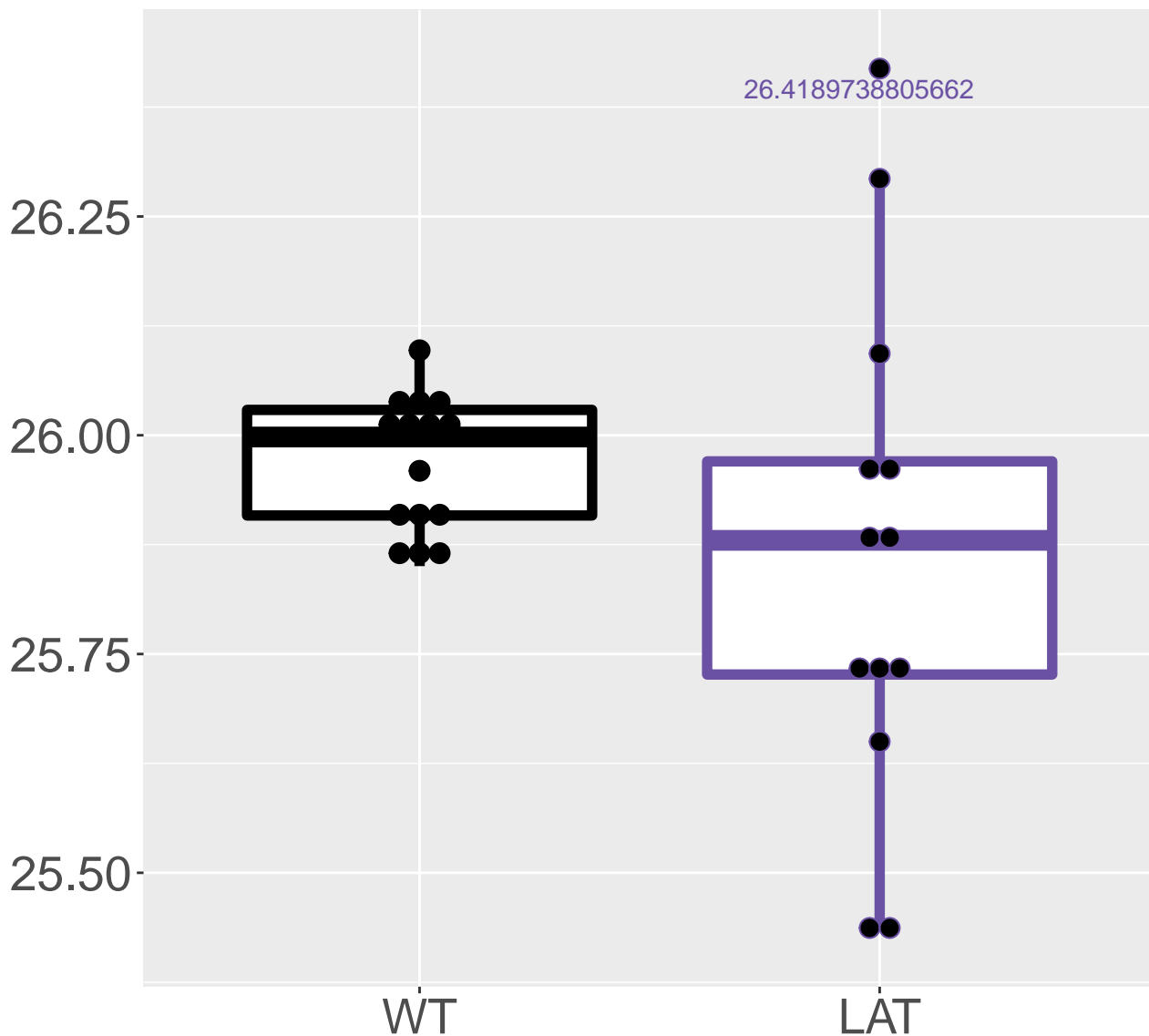


Q00623_Apolipoprotein A-I
FDR = 0.032, FC = -0.34, sex*

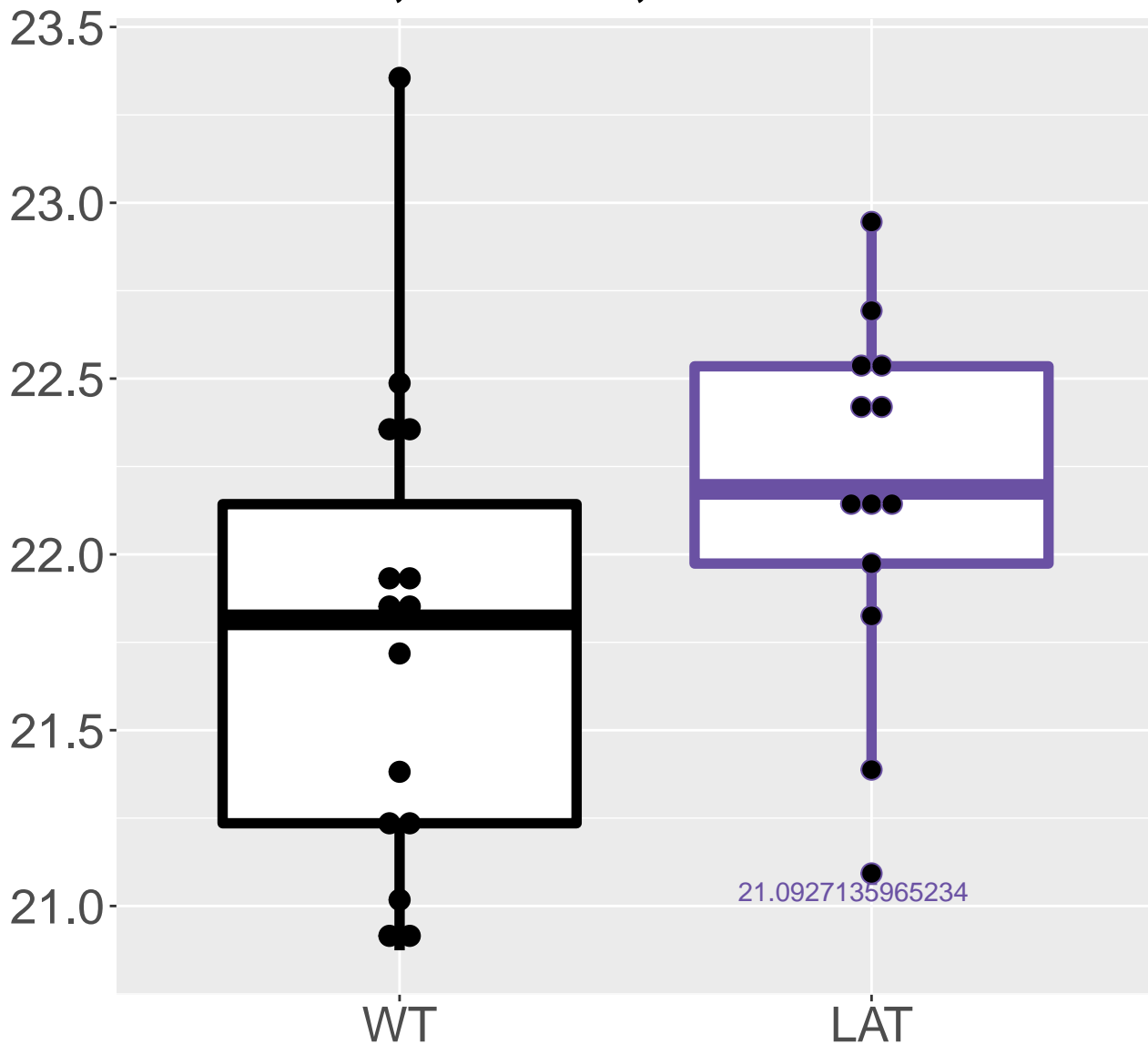


P61255_60S ribosomal protein L26

FDR = 0.032, FC = -0.29

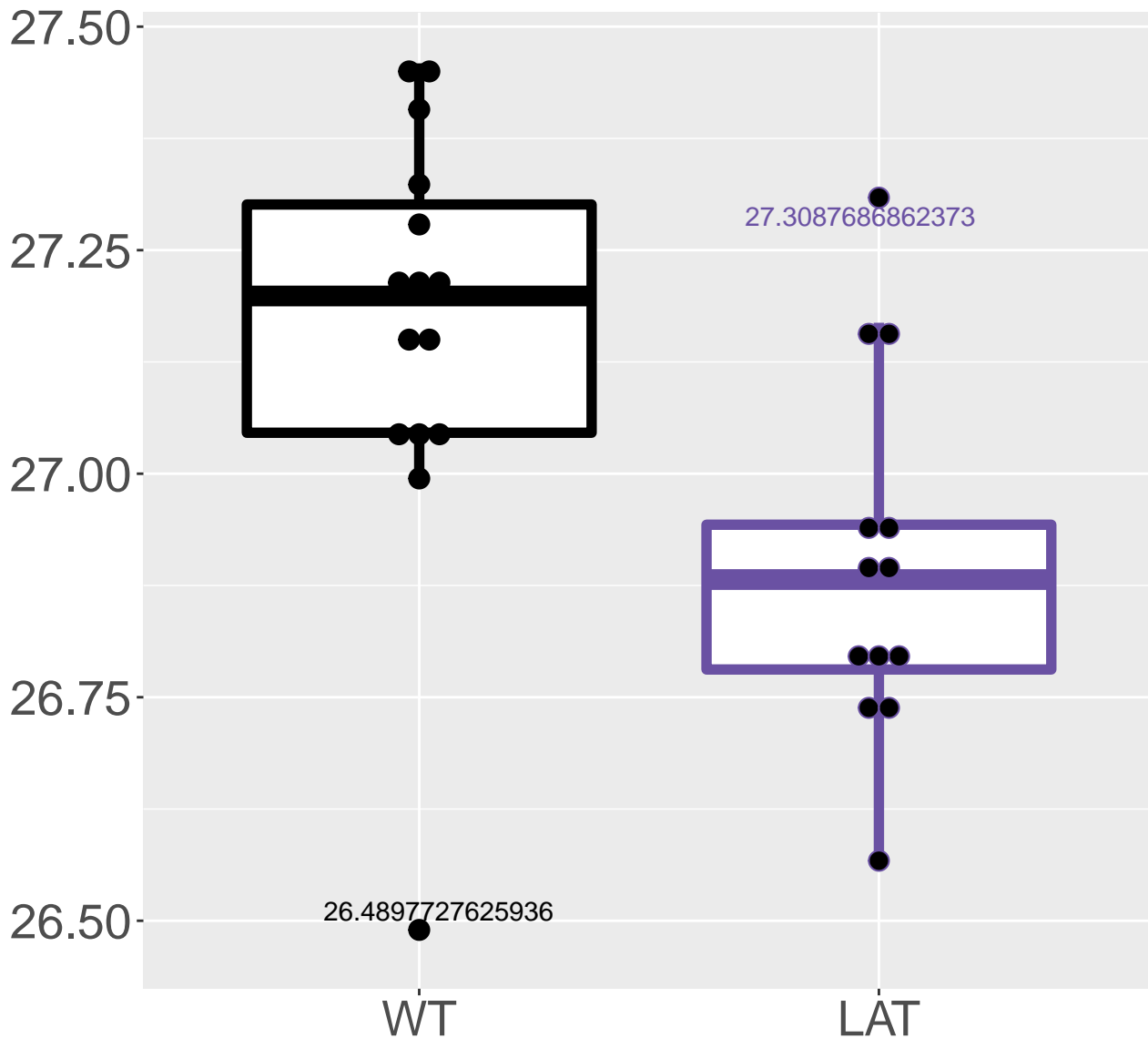


P17879_Heat shock 70 kDa protei.
FDR = 0.033, FC = 0.84, sex**



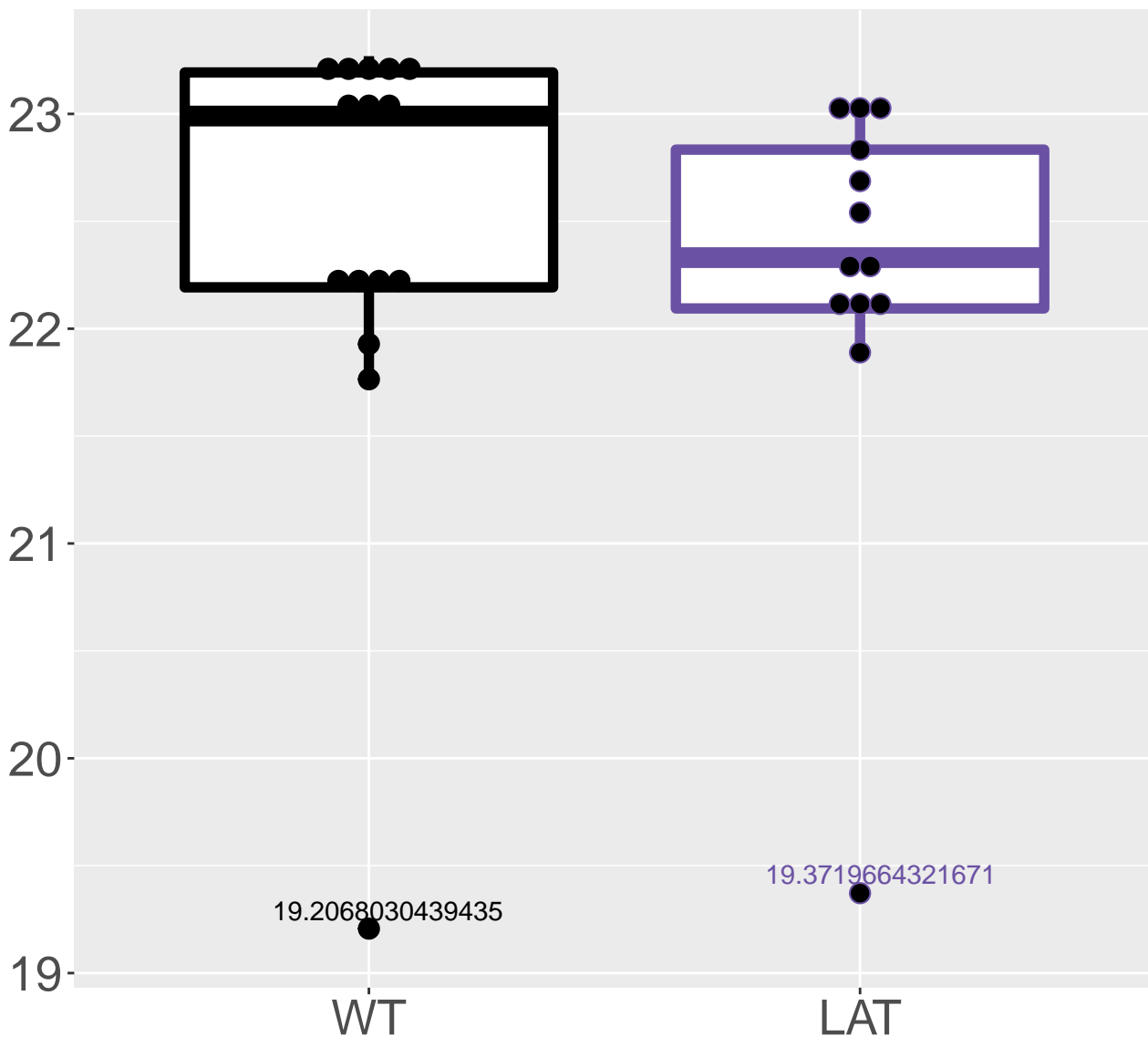
P31786_Acyl-CoA-binding protein

FDR = 0.033, FC = -0.45, sex*



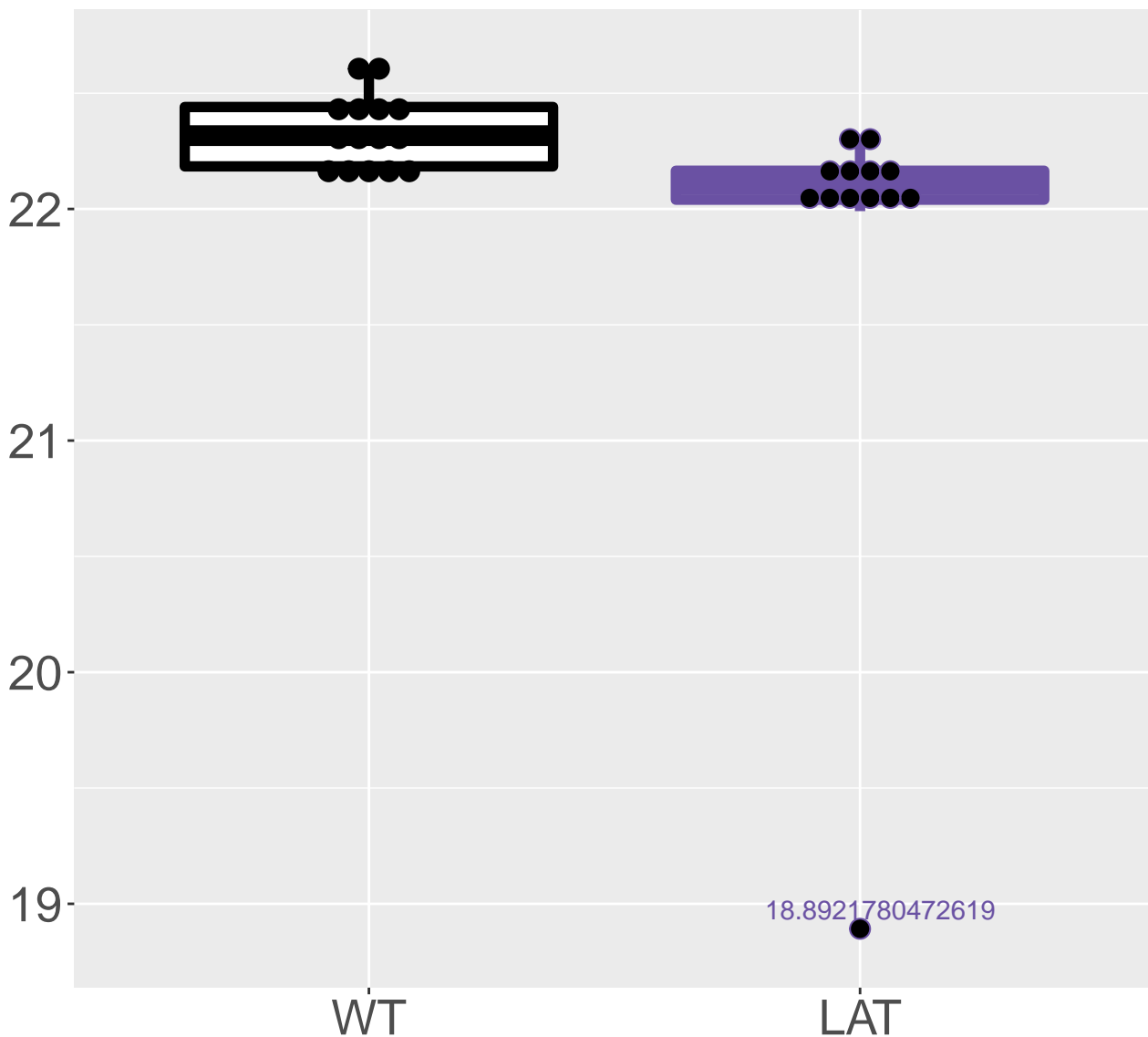
P97821_Dipeptidyl peptidase 1

FDR = 0.033, FC = -0.31, sex**

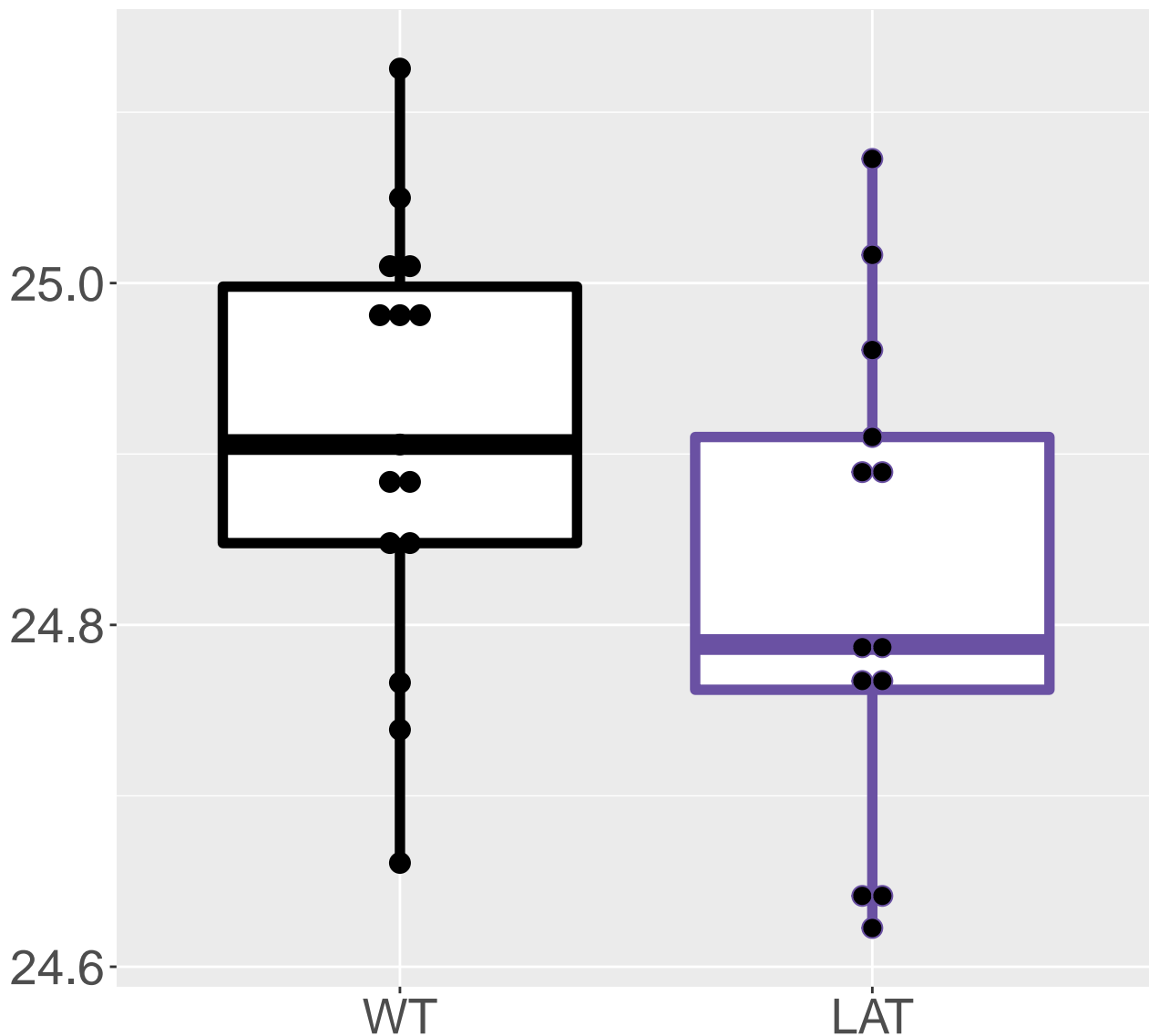


P56959_RNA-binding protein FUS

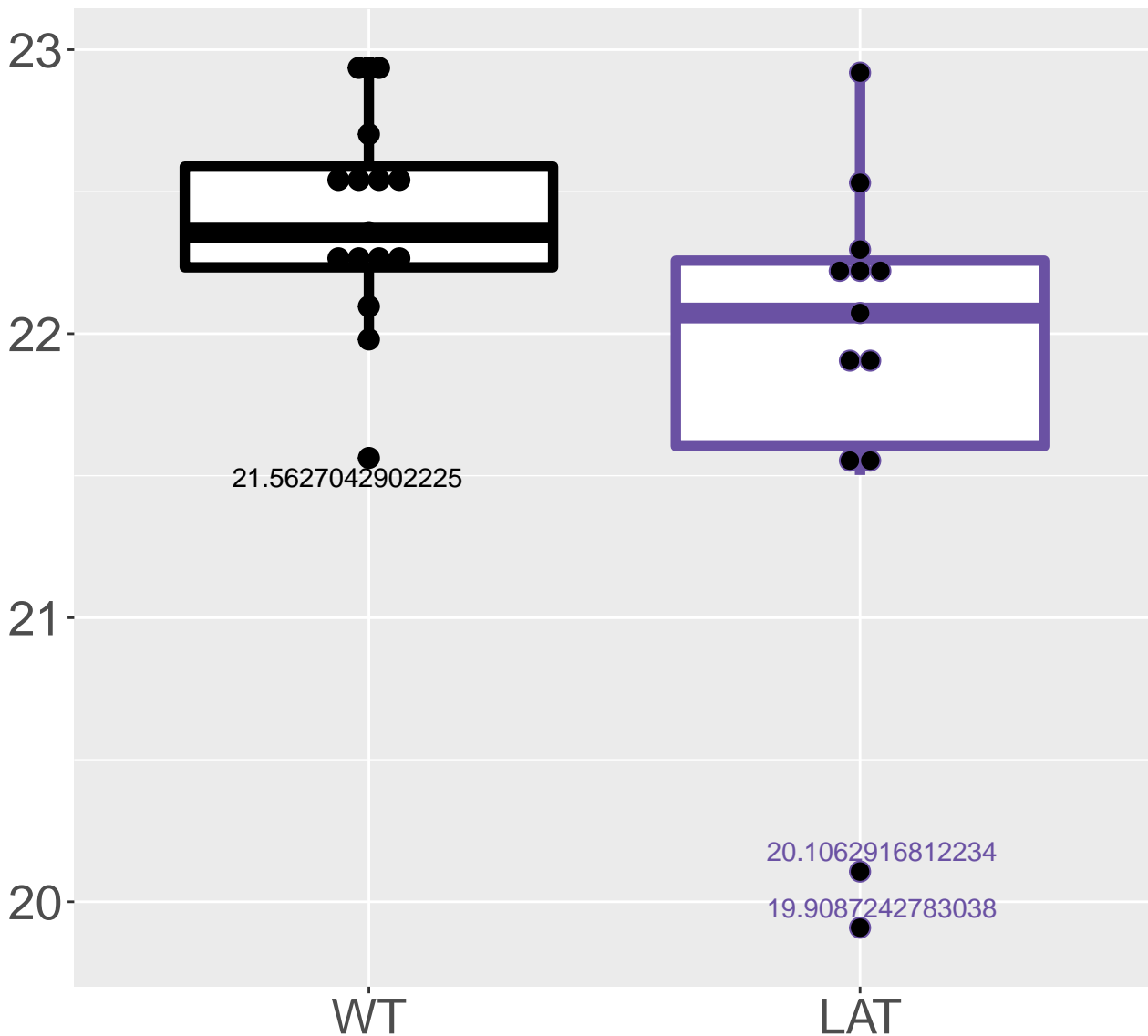
FDR = 0.033, FC = -0.26



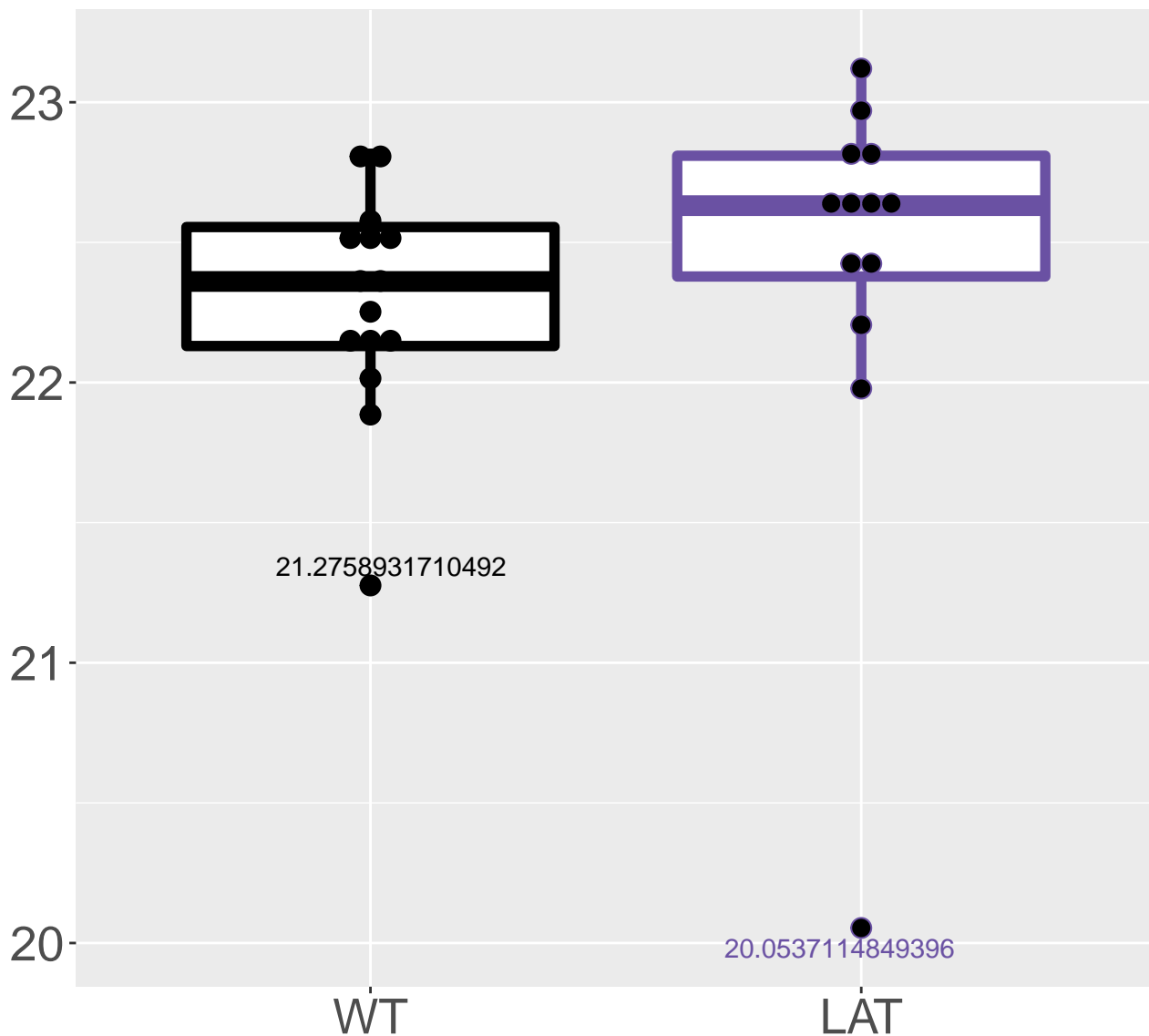
Q7TMF3_NADH dehydrogenase [ubiq.
FDR = 0.033, FC = -0.21, sex*



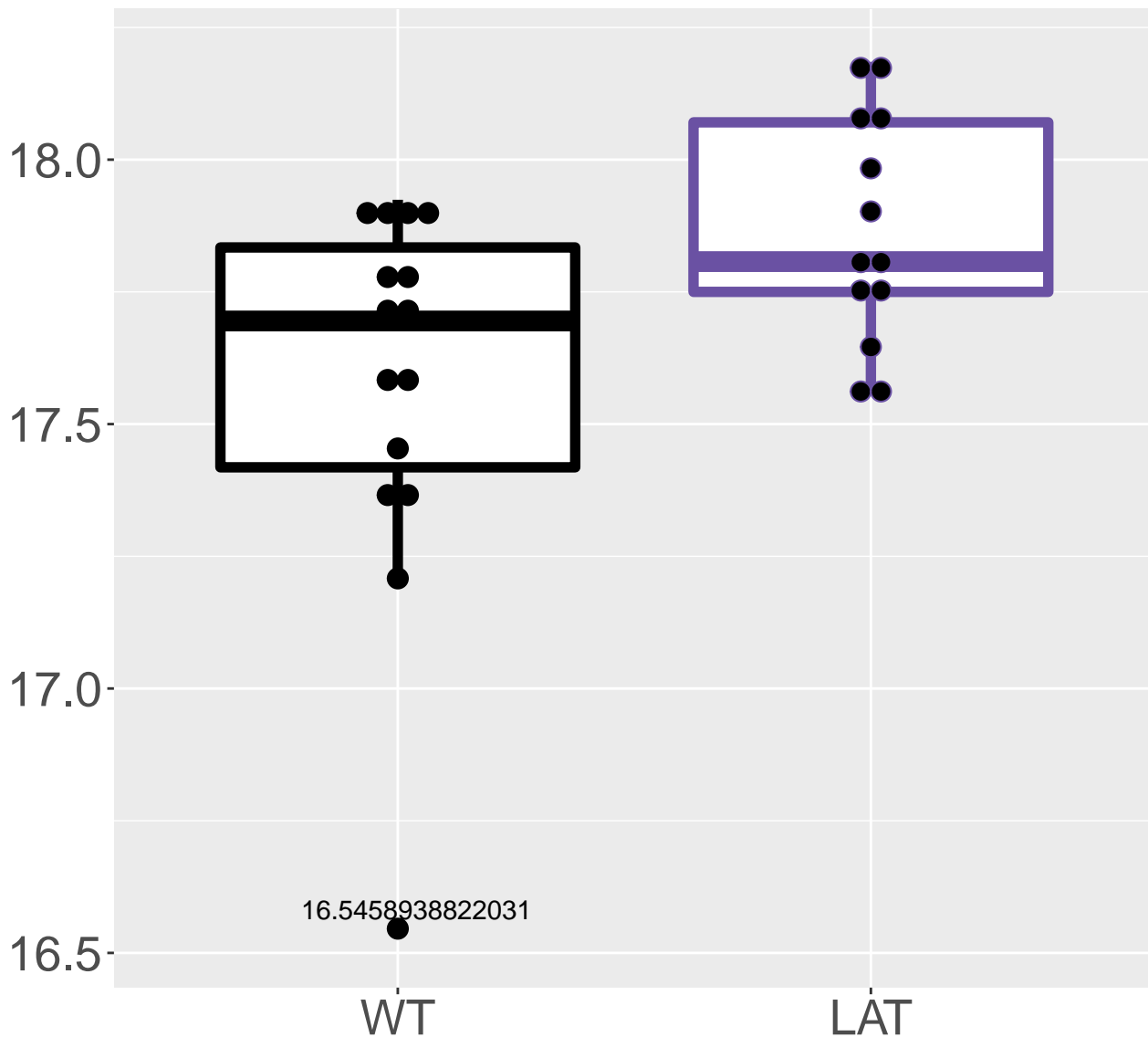
Q8BTZ7_Mannose-1-phosphate guan.
FDR = 0.033, FC = -1.2



Q8JZQ9_Eukaryotic translation i.
FDR = 0.033, FC = 0.61, sex*

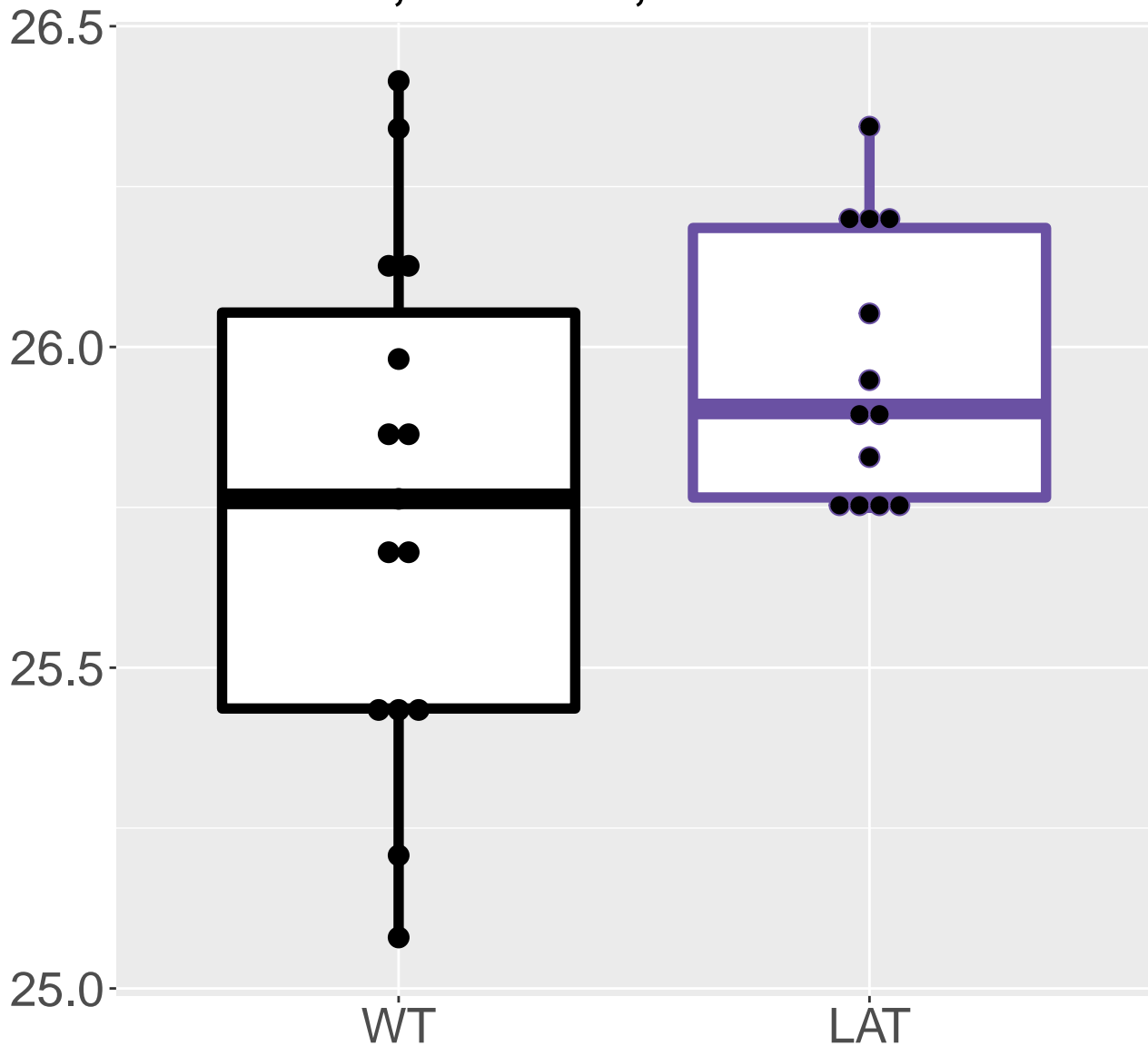


Q8BWQ6_UPF0505 protein C16orf62.
FDR = 0.033, FC = 0.58, sex*

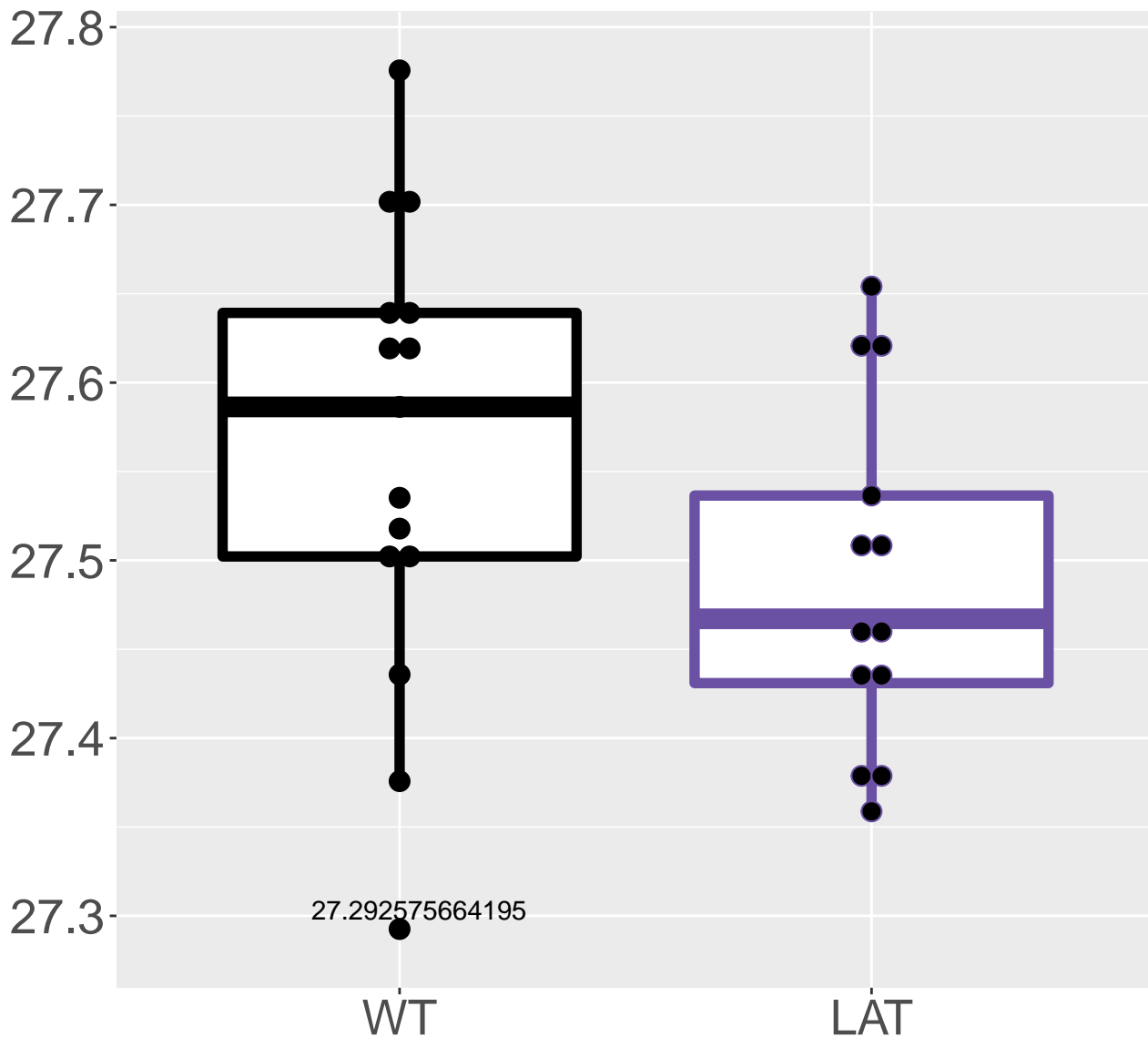


Q8VCU1_Carboxylesterase 3B

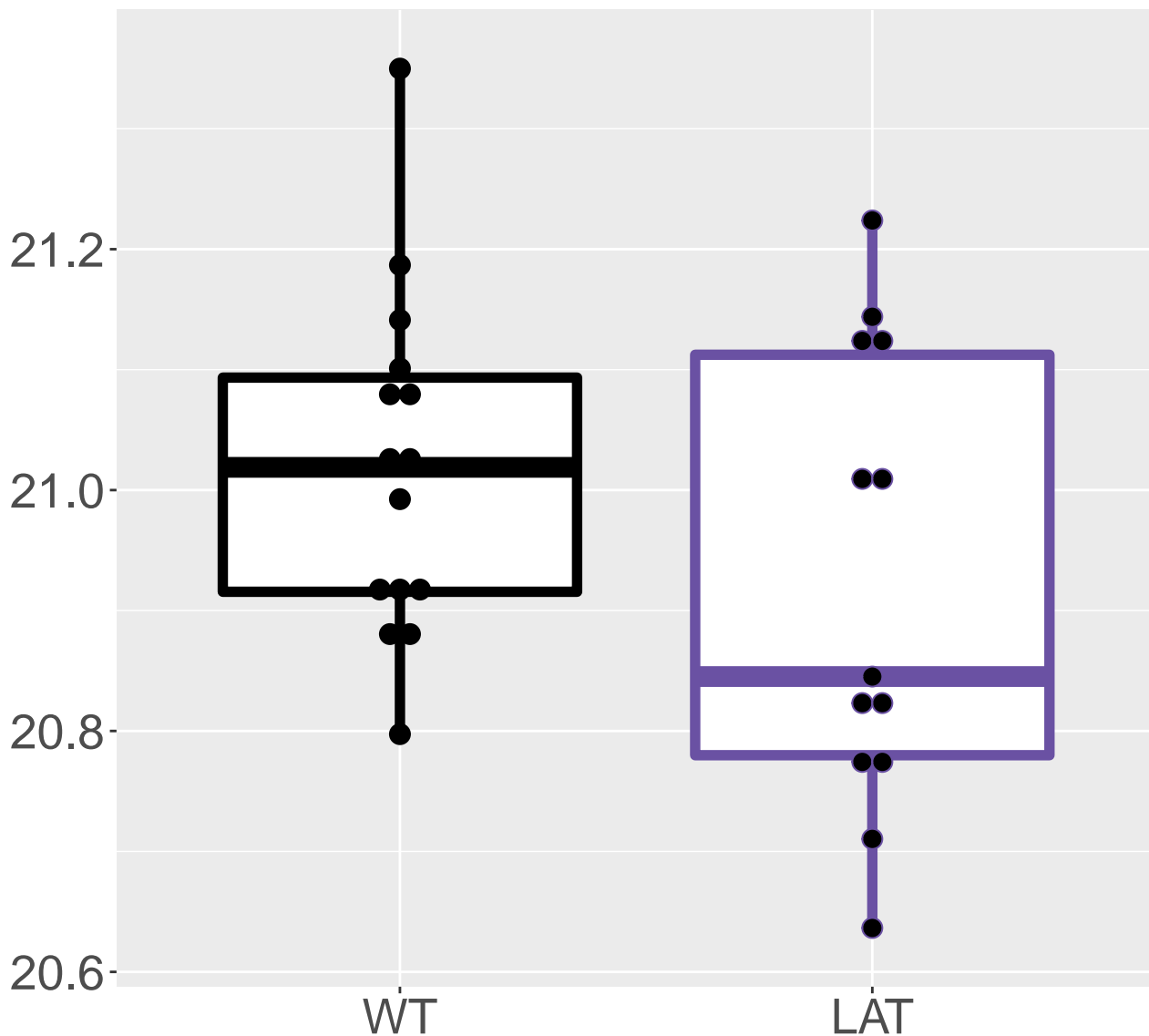
FDR = 0.033, FC = 0.43, sex**



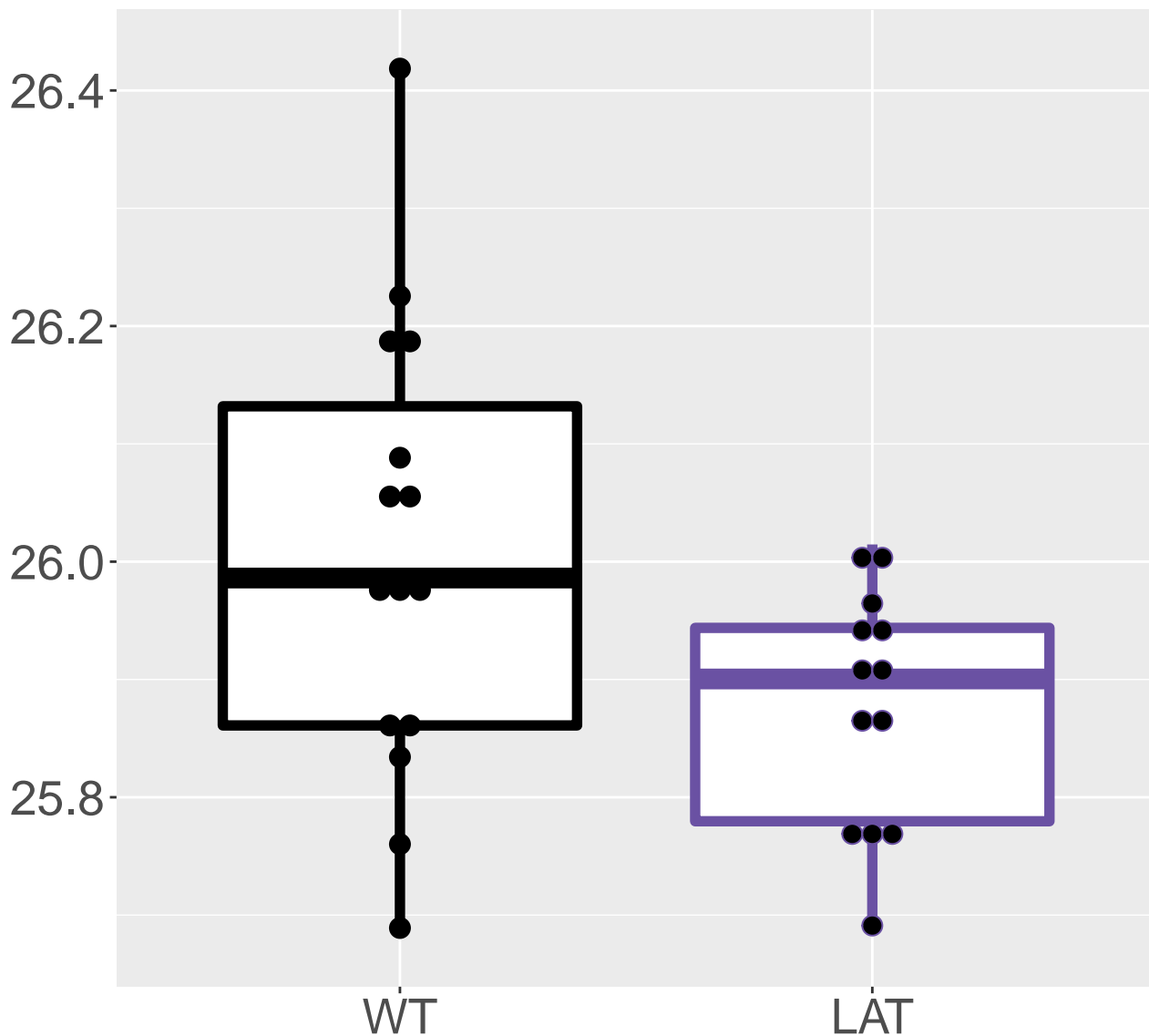
Q9DB20_ATP synthase subunit O, .
FDR = 0.033, FC = -0.2



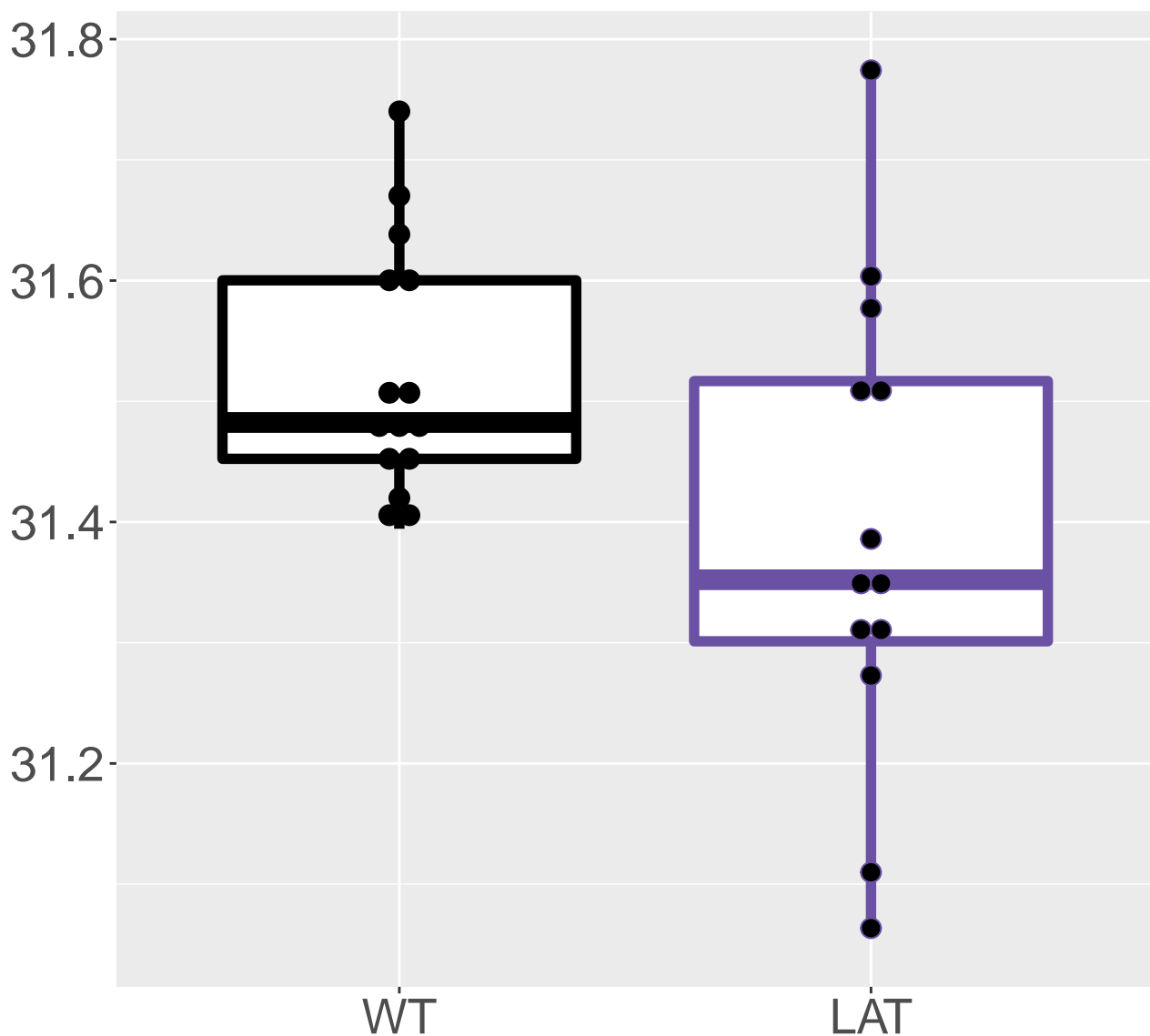
P03899_NADH-ubiquinone oxidored.
FDR = 0.033, FC = -0.19, sex**



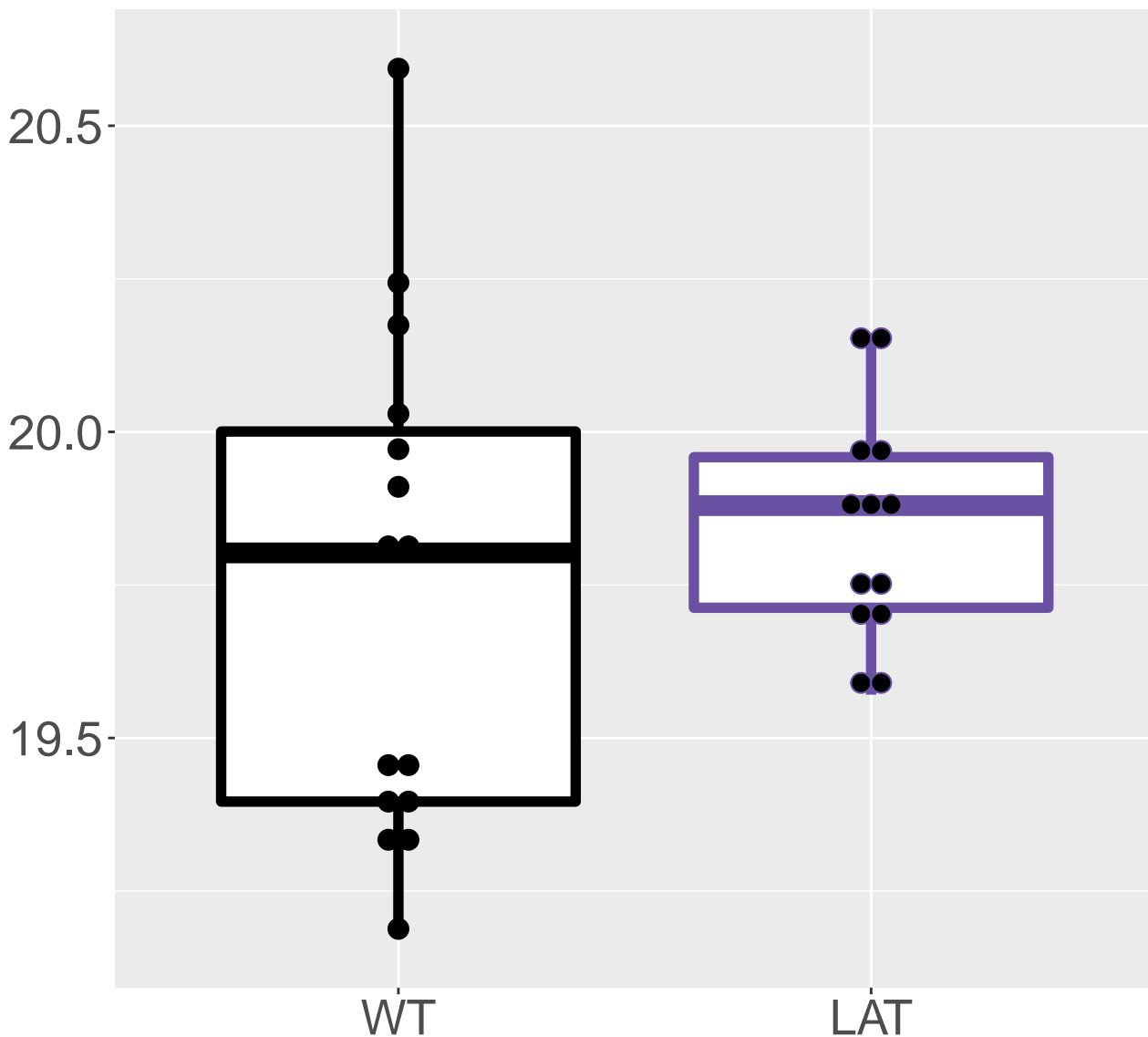
P61458_Pterin-4-alpha-carbinola.
FDR = 0.034, FC = -0.28, sex*



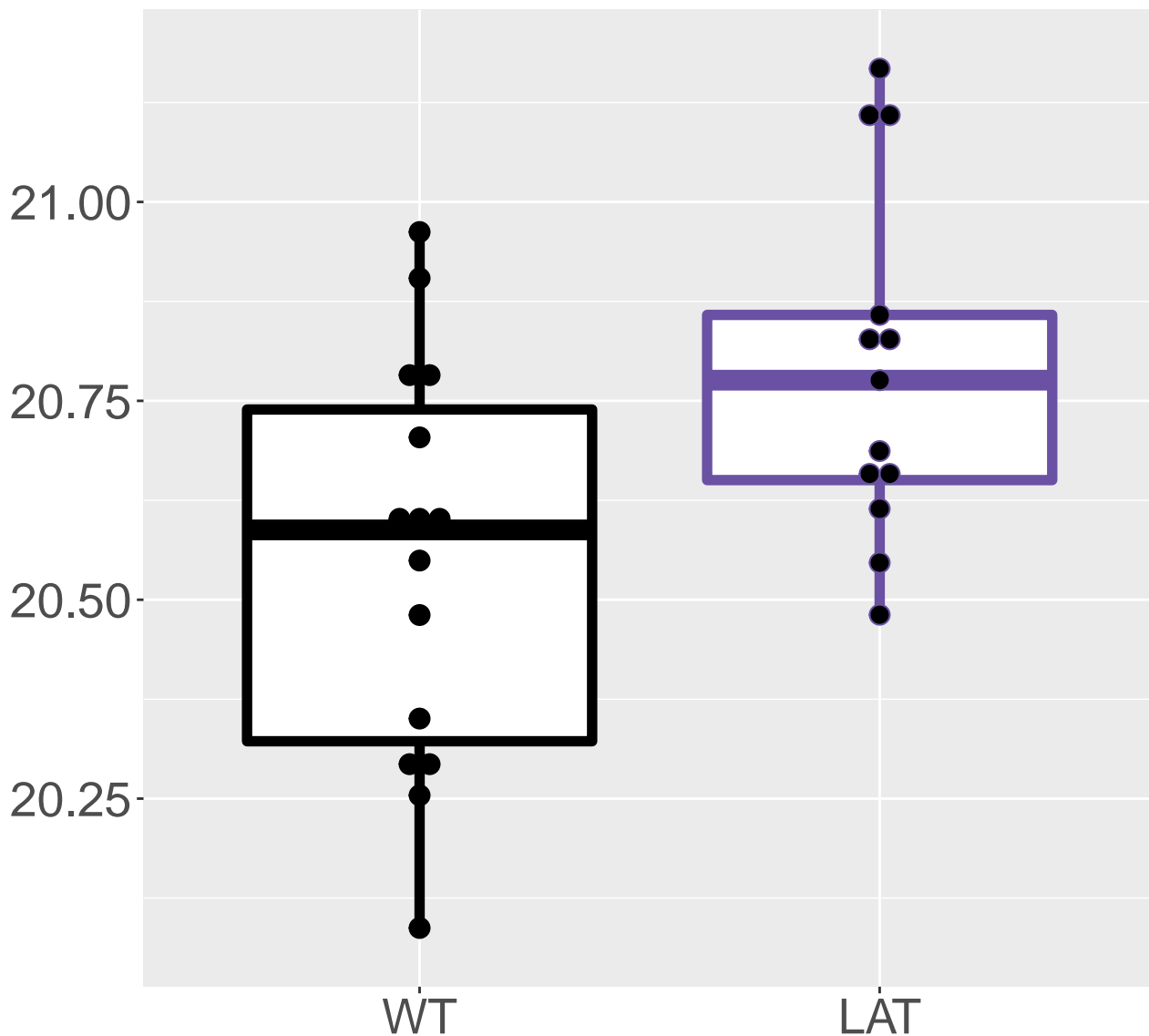
P12710_Fatty acid-binding prote.
FDR = 0.034, FC = -0.22



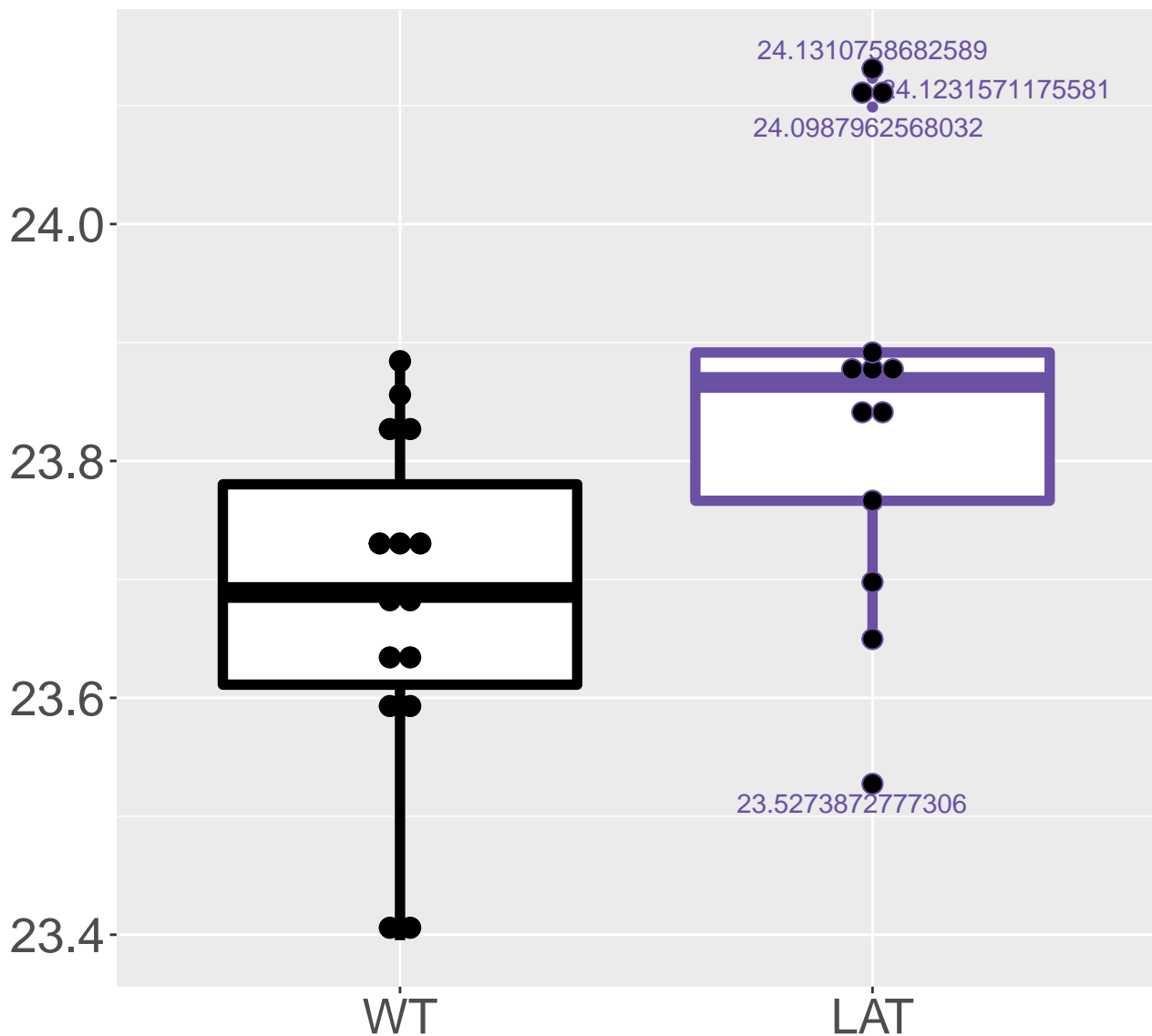
P57746_V-type proton ATPase sub.
FDR = 0.035, FC = 0.4, sex***



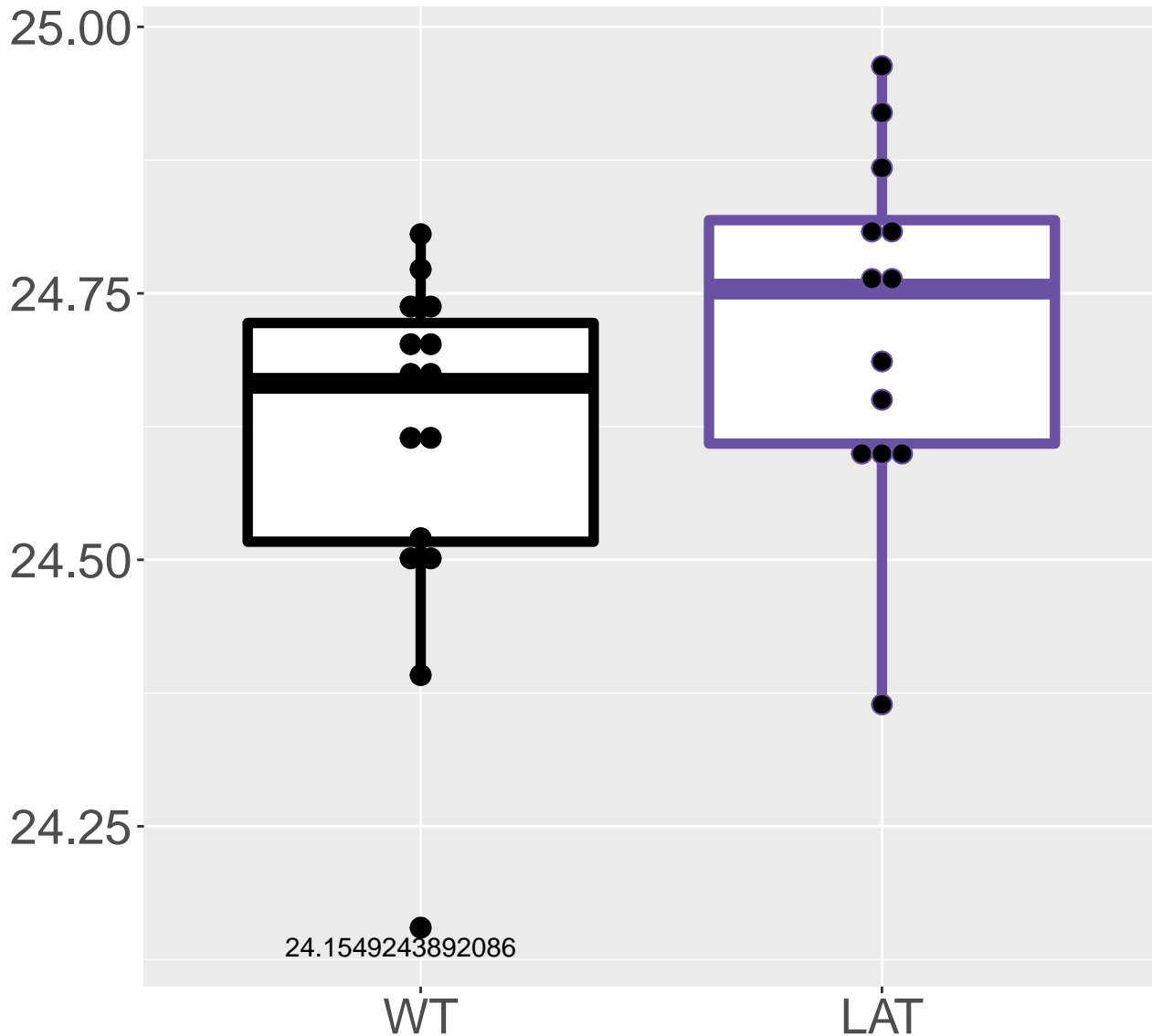
Q3URE1_Acyl-CoA synthetase fami.
FDR = 0.036, FC = 0.37, sex*



Q80UM7_Mannosyl-oligosaccharide.
FDR = 0.036, FC = 0.3

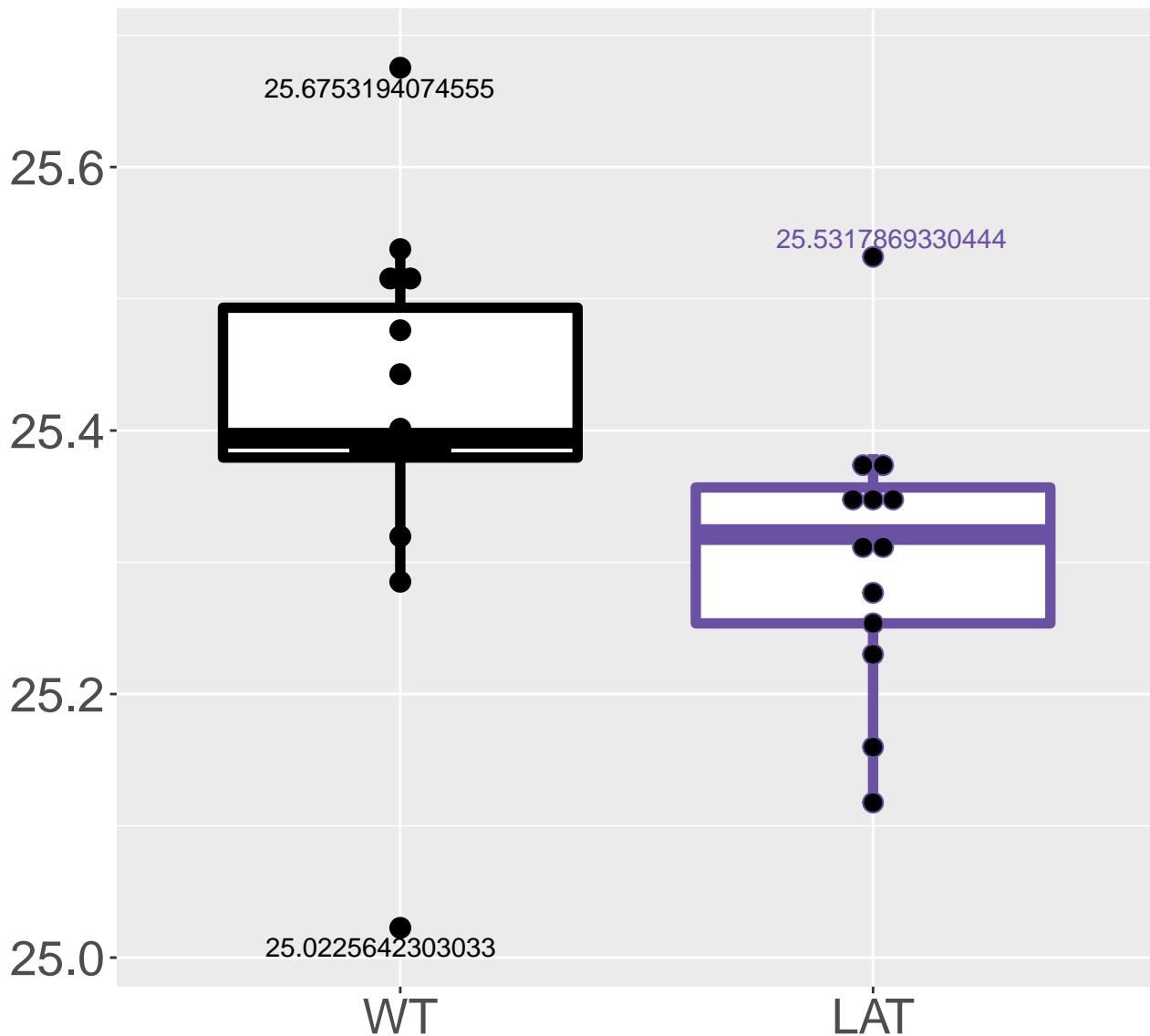


Q9QZE5_Coatomer subunit gamma-1
FDR = 0.036, FC = 0.28, sex*



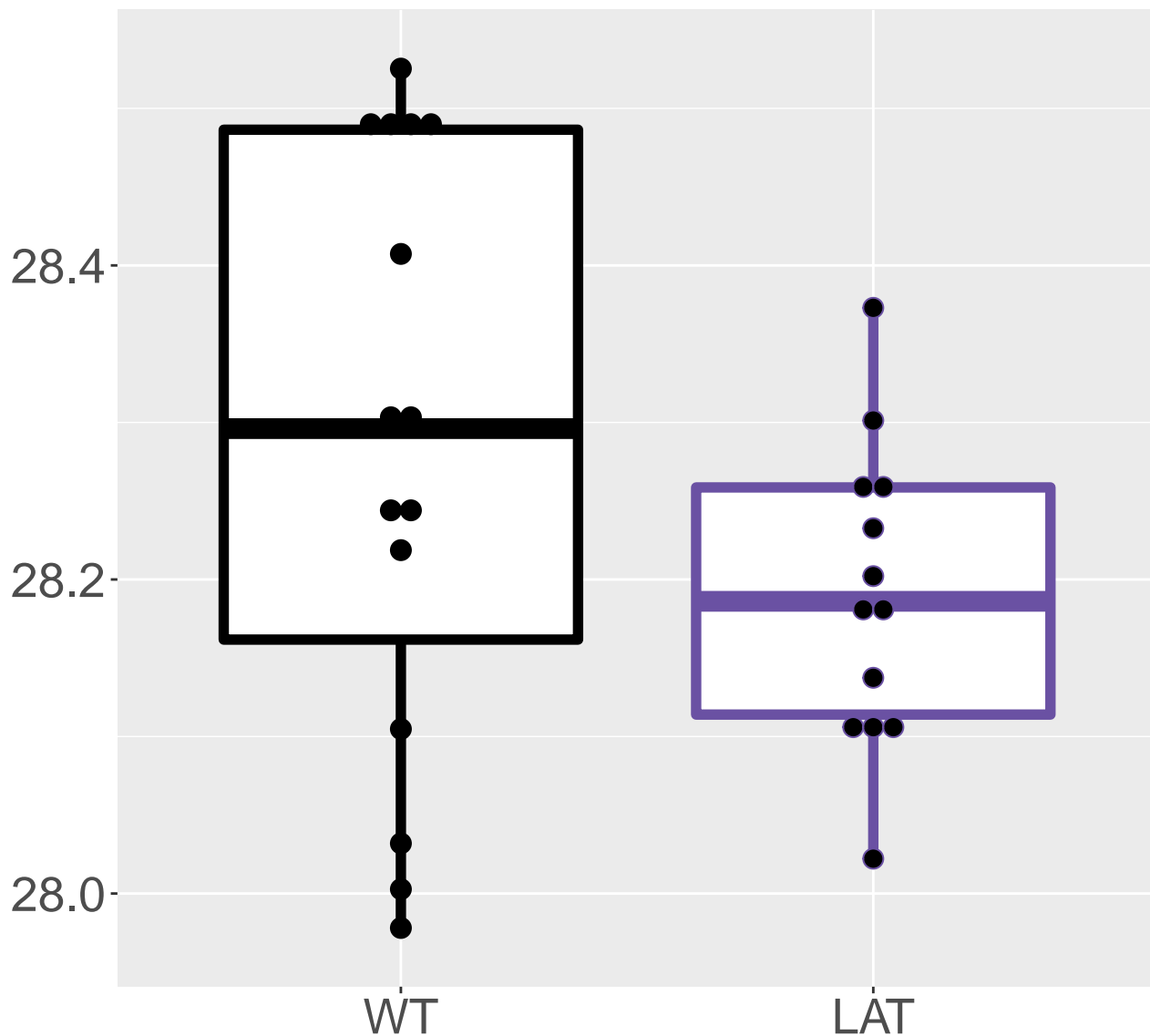
Q9D0S9_Histidine triad nucleoti.

FDR = 0.036, FC = -0.21



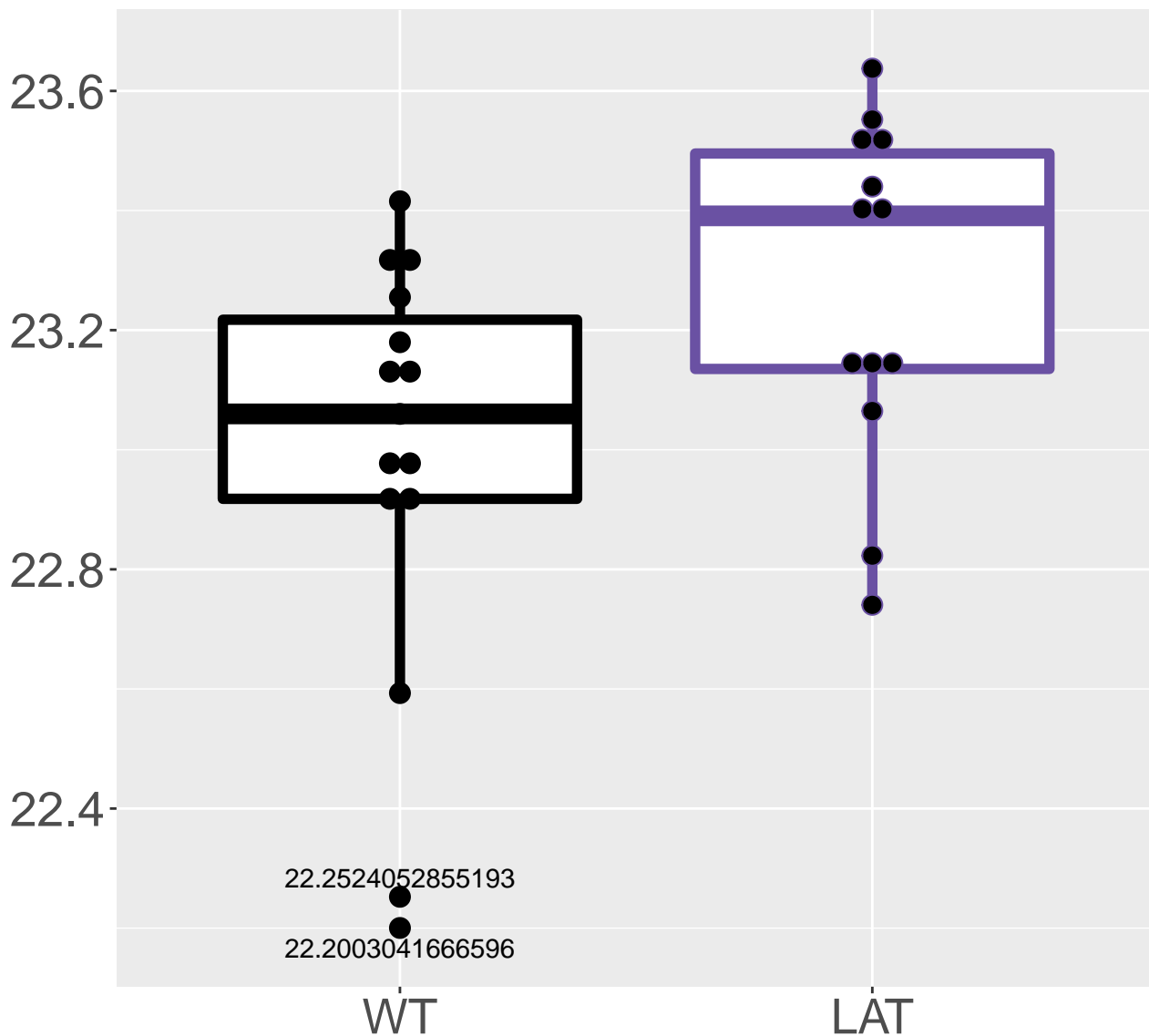
Q8CIM7_Cytochrome P450 2D26

FDR = 0.036, FC = -0.18, sex***

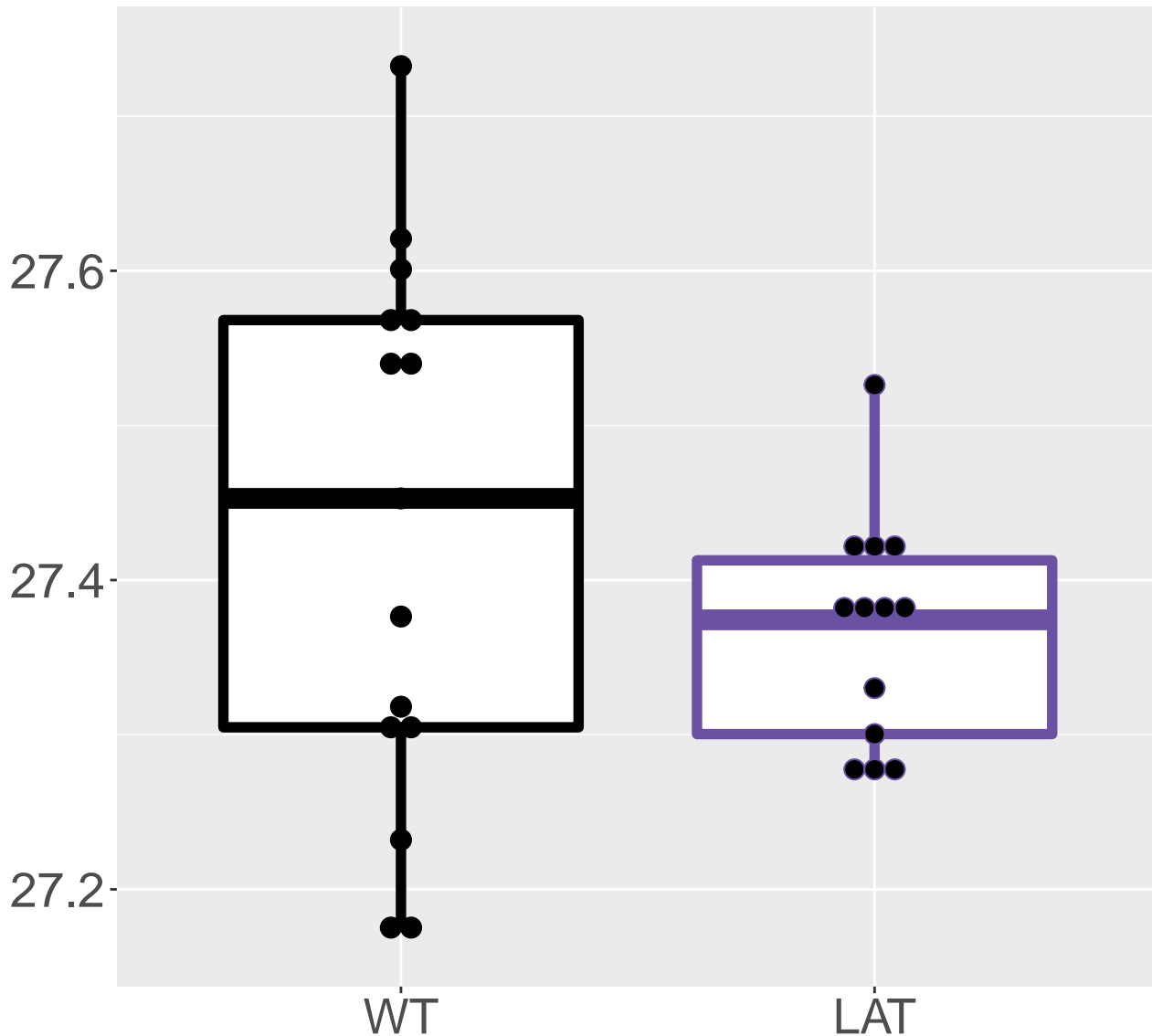


Q8BML9_Glutamine--tRNA ligase

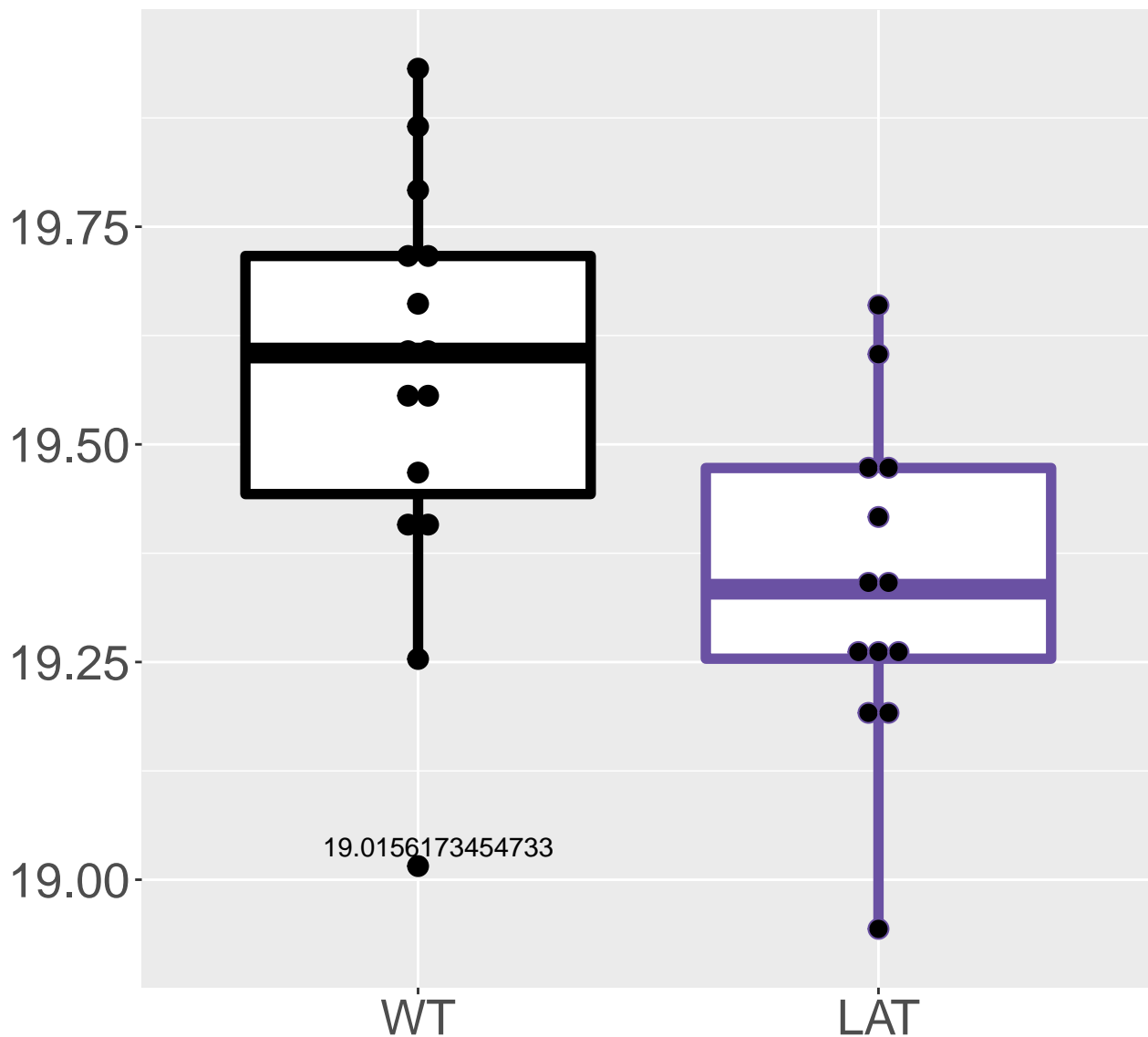
FDR = 0.037, FC = 0.64



Q8R164_Valacyclovir hydrolase
FDR = 0.037, FC = -0.16, sex***

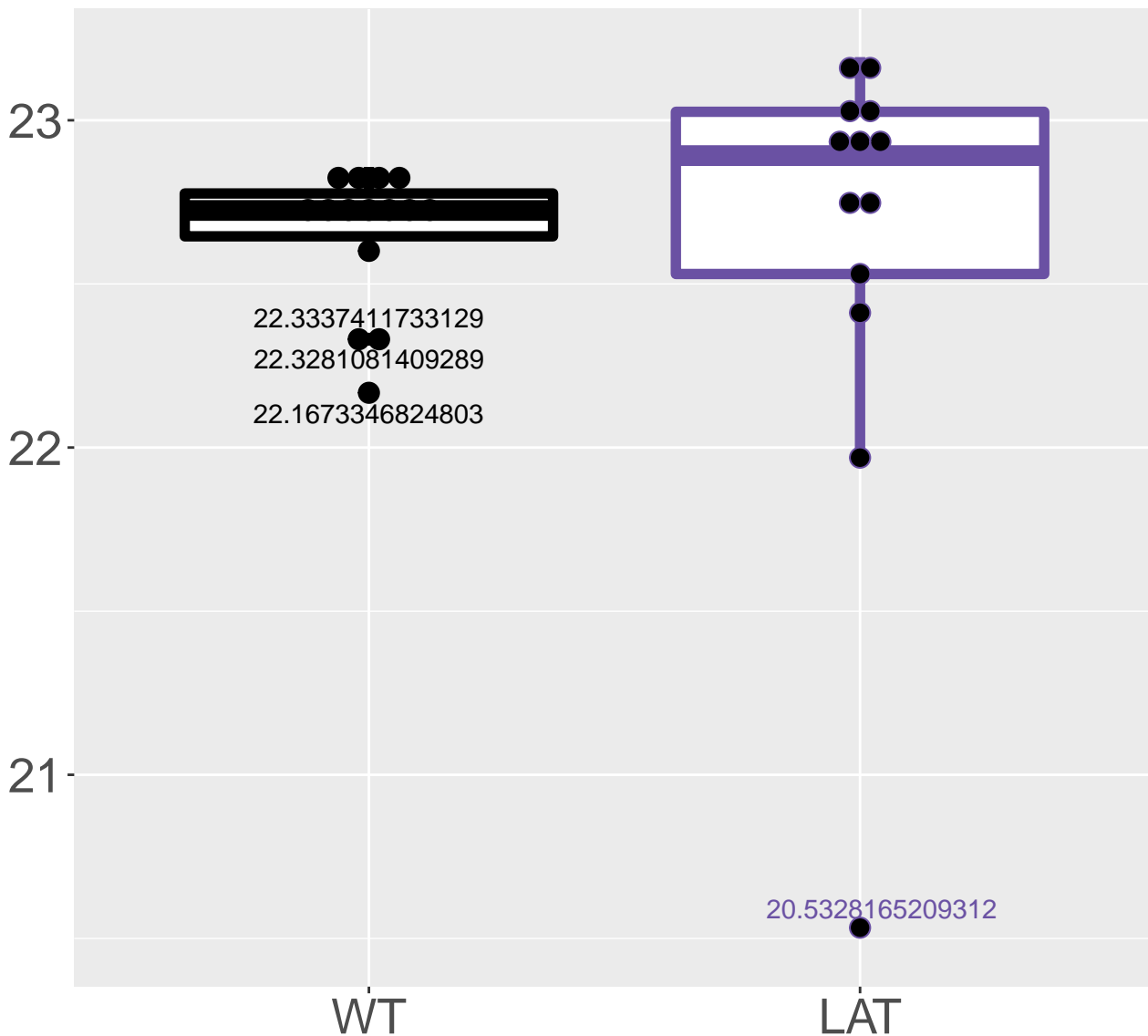


O88653_Ragulator complex protei.
FDR = 0.037, FC = -0.3, sex*

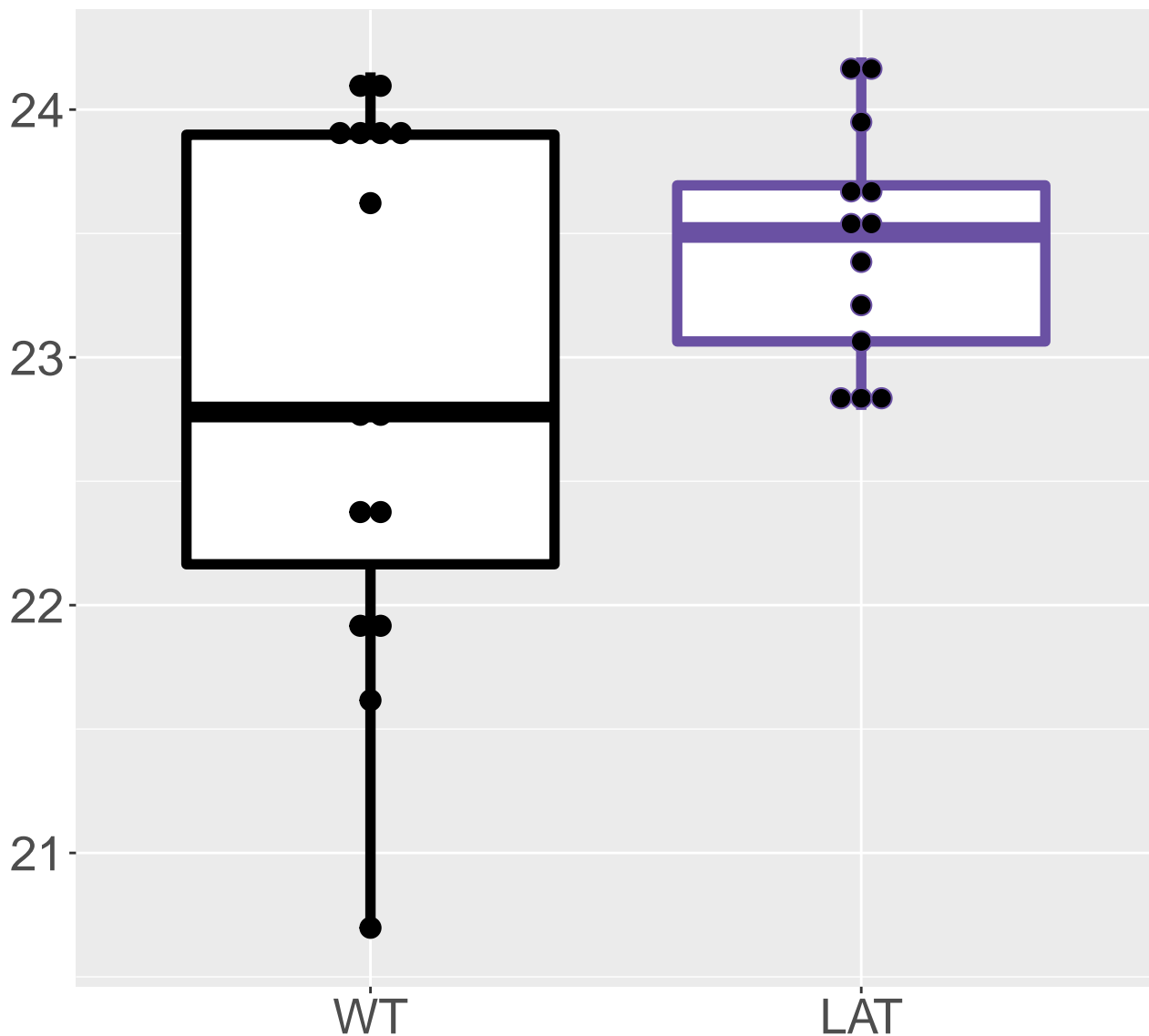


P26231_Catenin alpha-1

FDR = 0.037, FC = 0.42

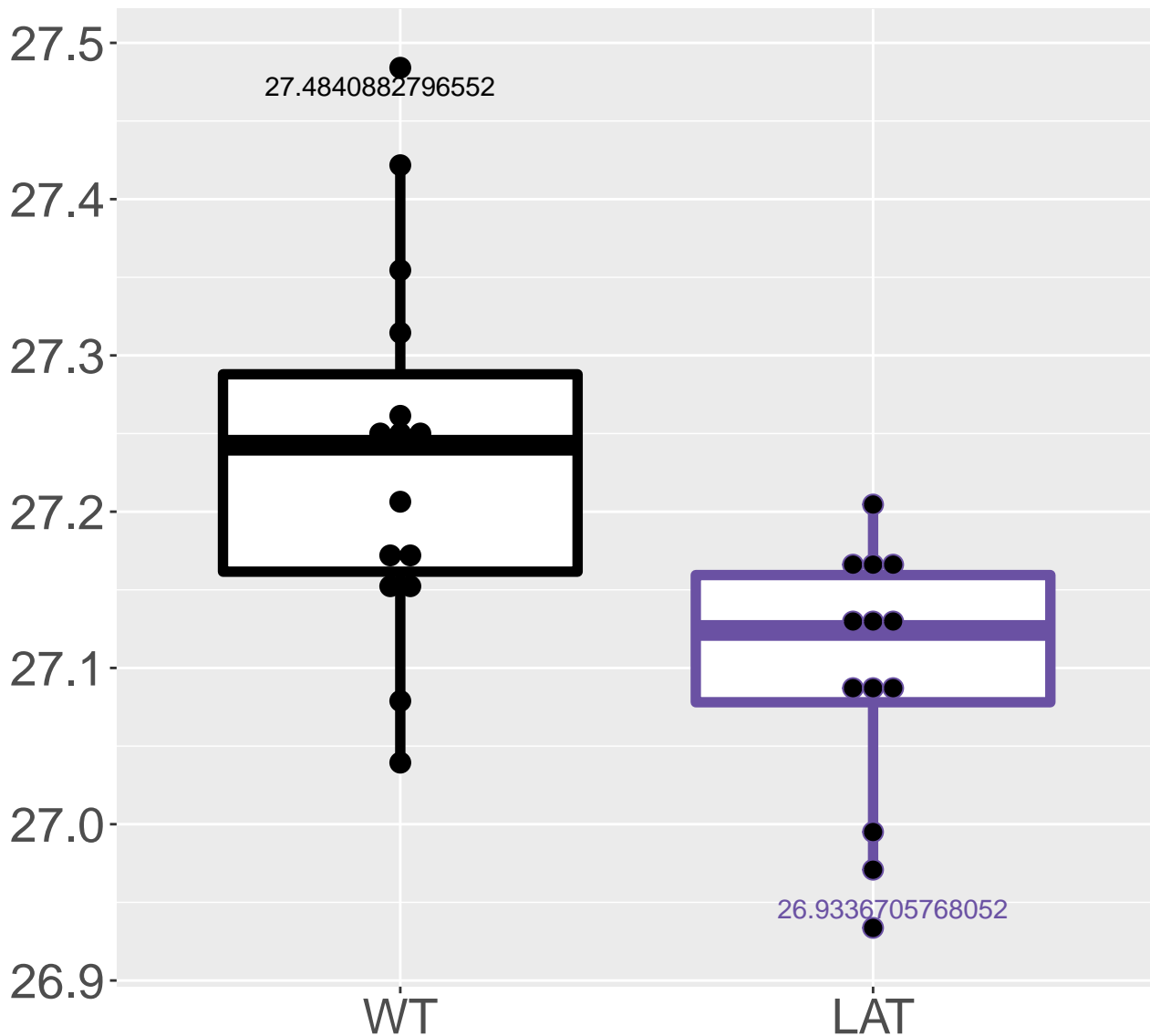


Q9DBN5_Lon protease homolog 2, .
FDR = 0.037, FC = 1, sex***



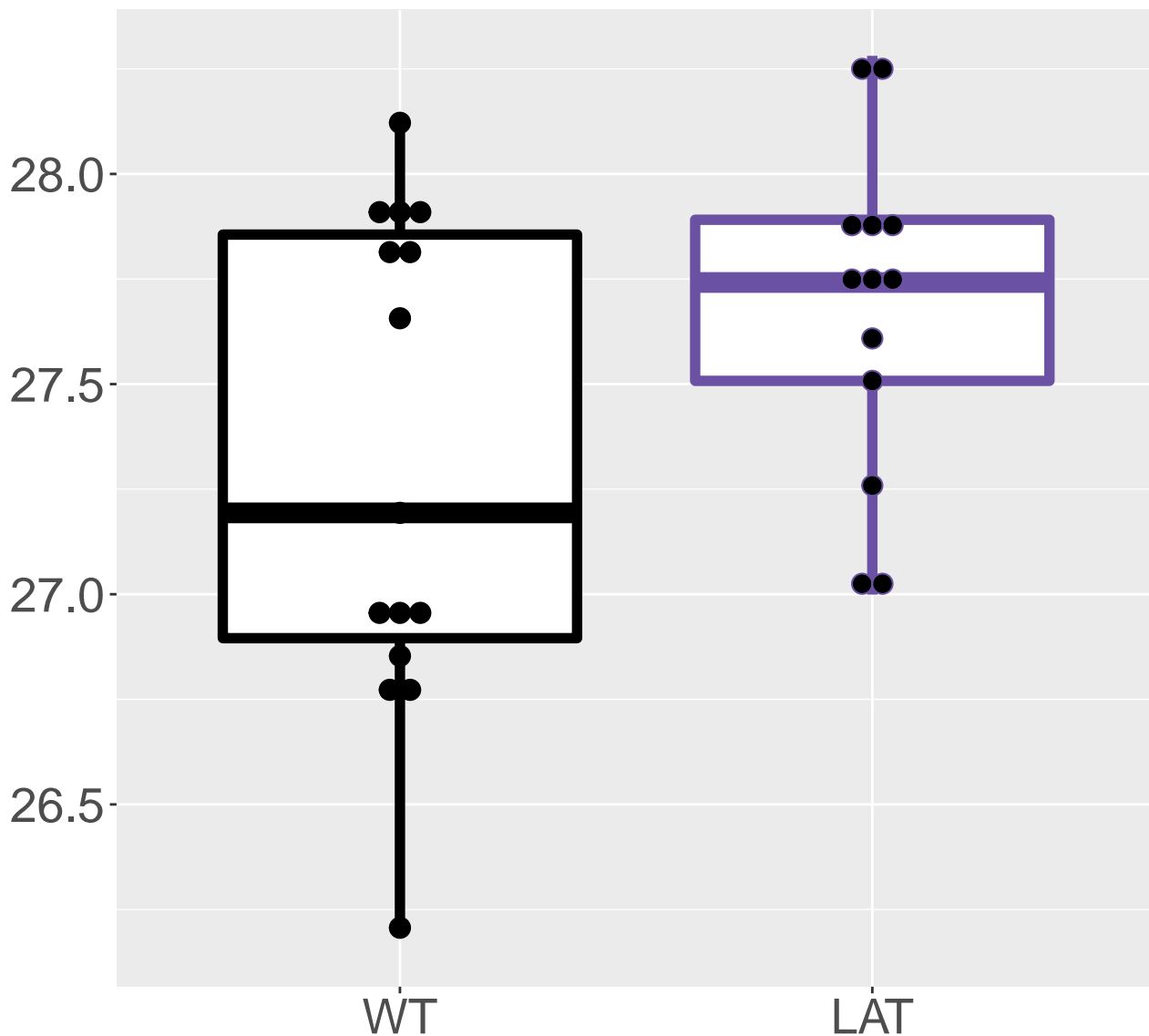
P12970_60S ribosomal protein L7a

FDR = 0.037, FC = -0.18

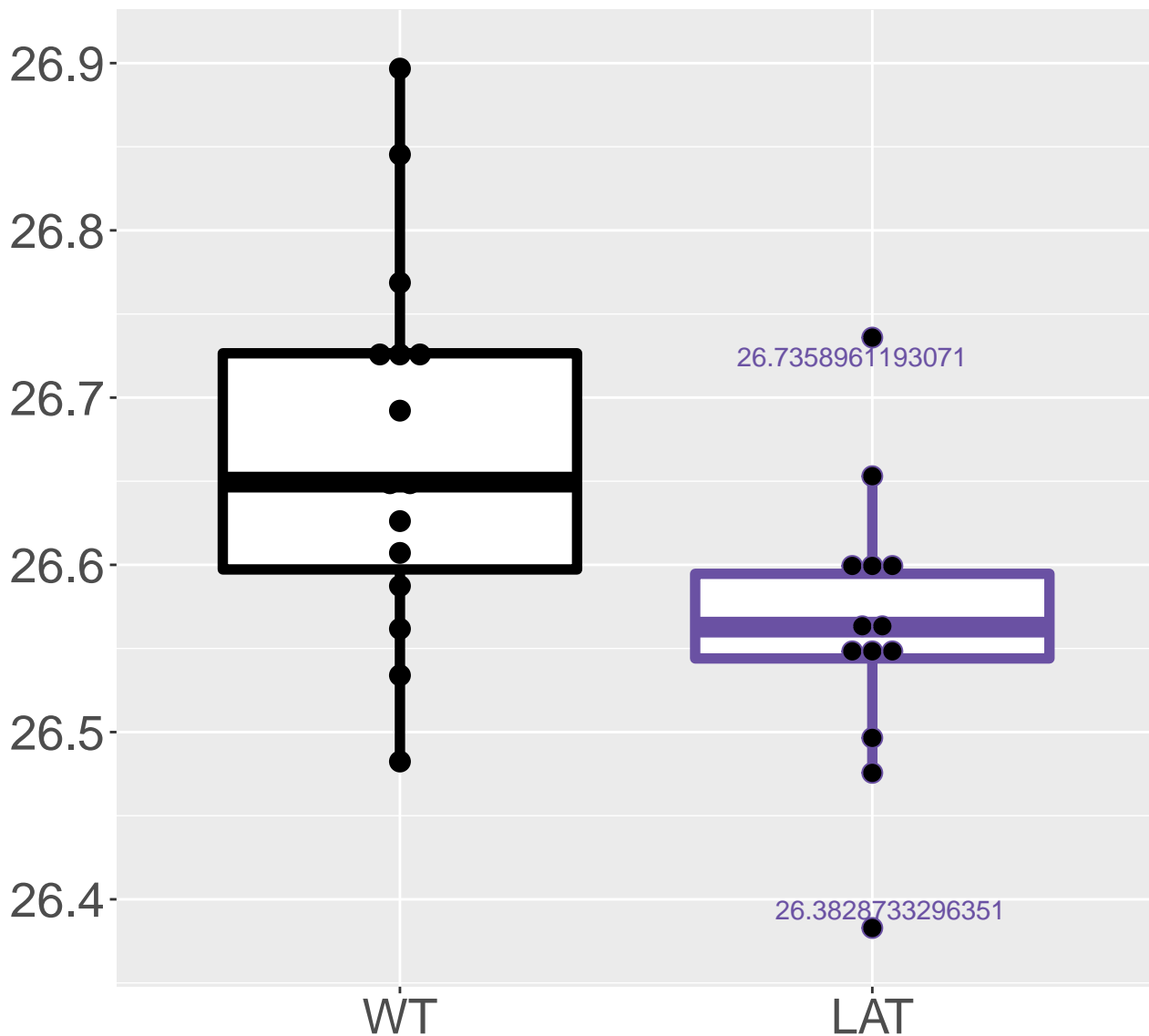


Q64458_Cytochrome P450 2C29

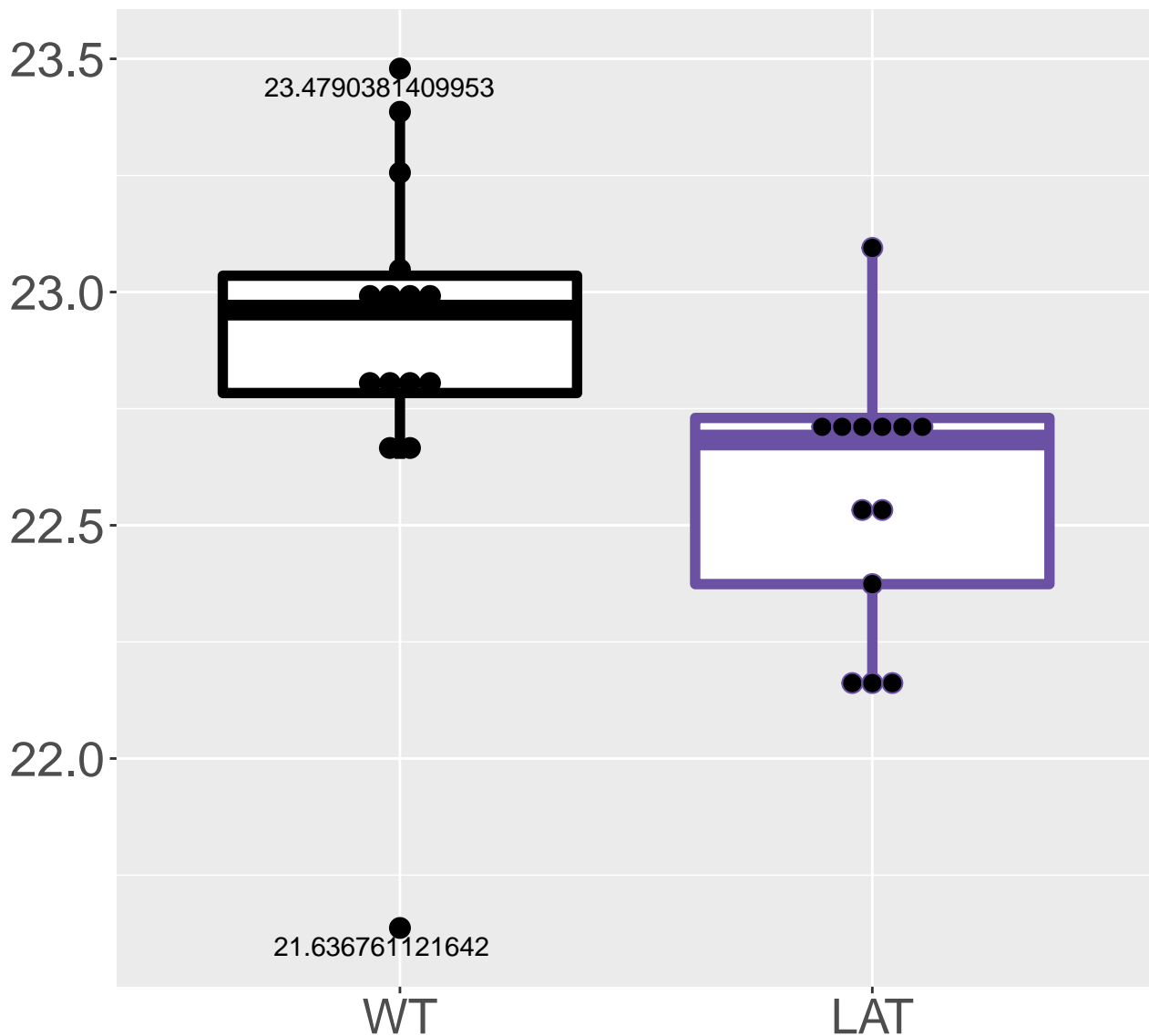
FDR = 0.039, FC = 0.53, sex***



Q9D6Y7_Mitochondrial peptide me.
FDR = 0.039, FC = -0.17

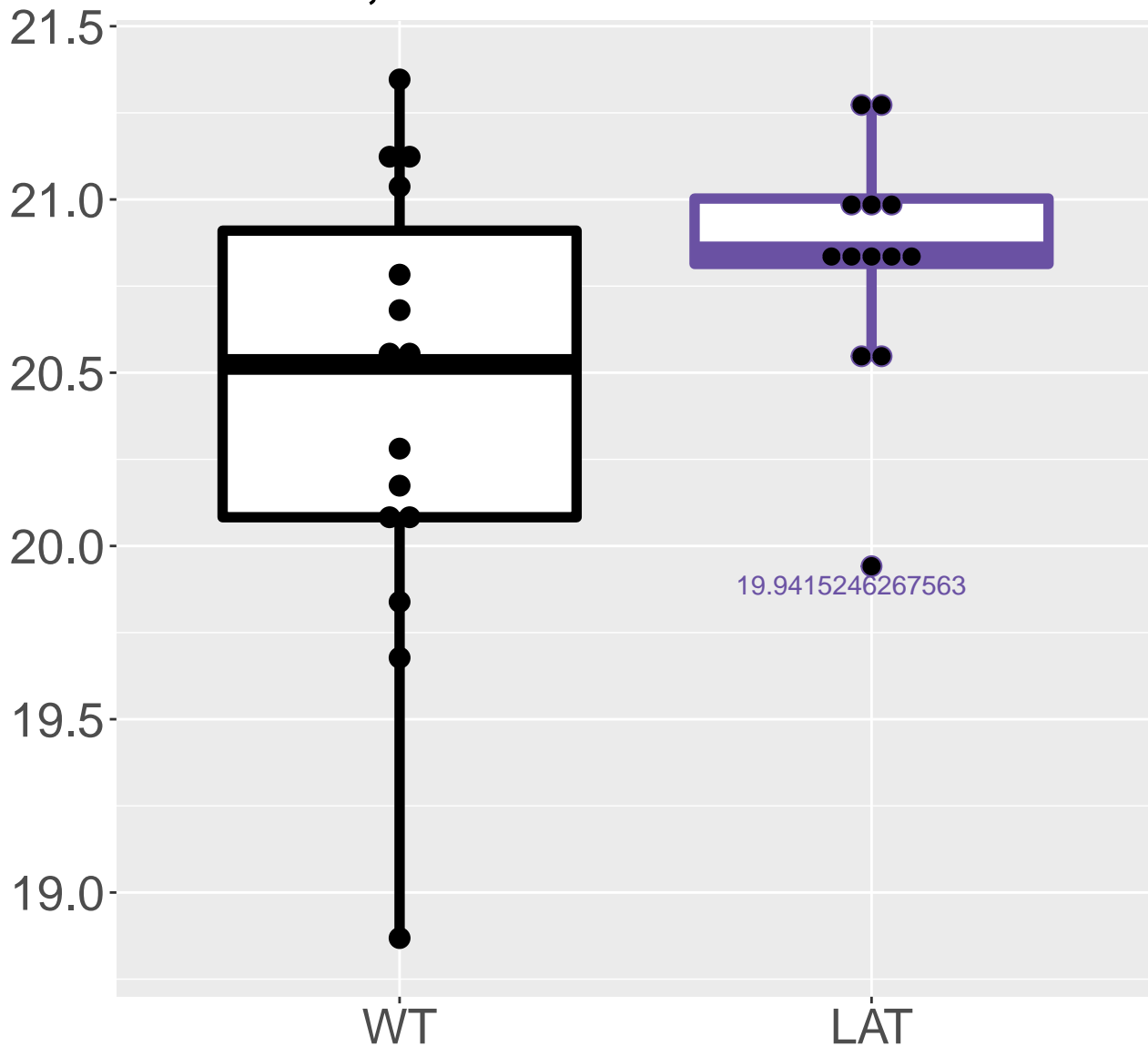


P62317_Small nuclear ribonucleo.
FDR = 0.039, FC = -0.46

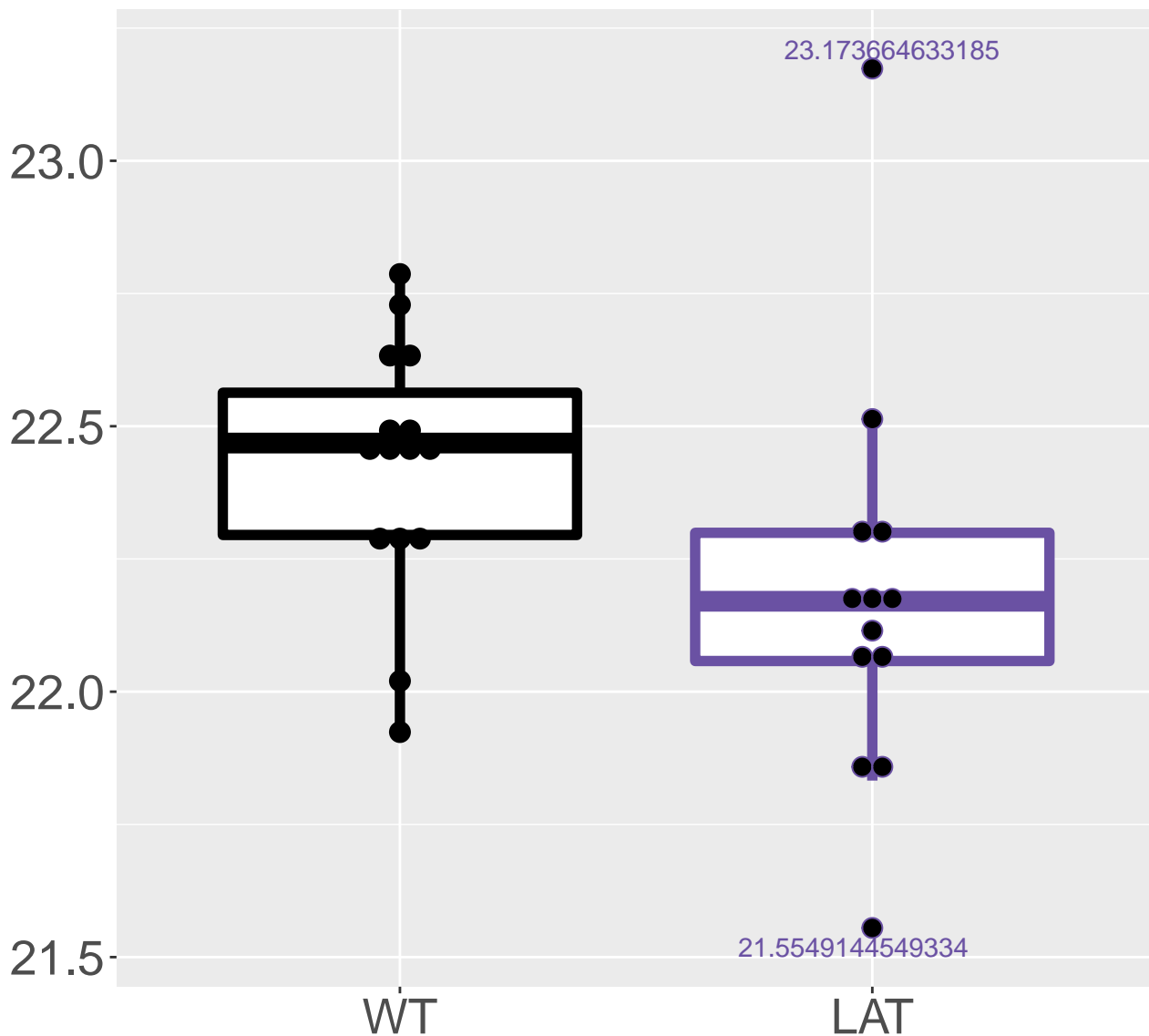


Q60960_Importin subunit alpha-5

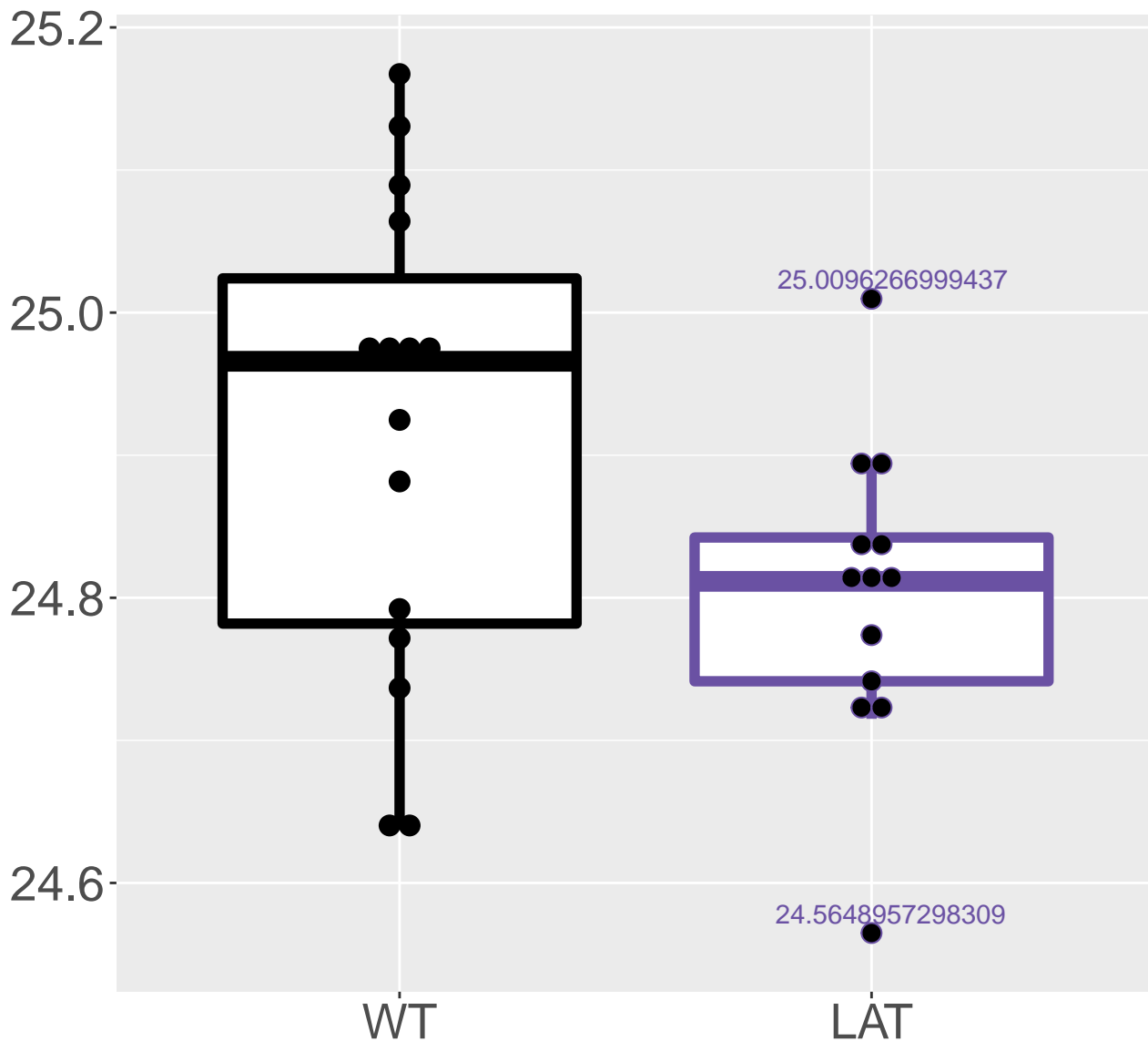
FDR = 0.04, FC = 0.58



P16045_Galectin-1
FDR = 0.04, FC = -0.43

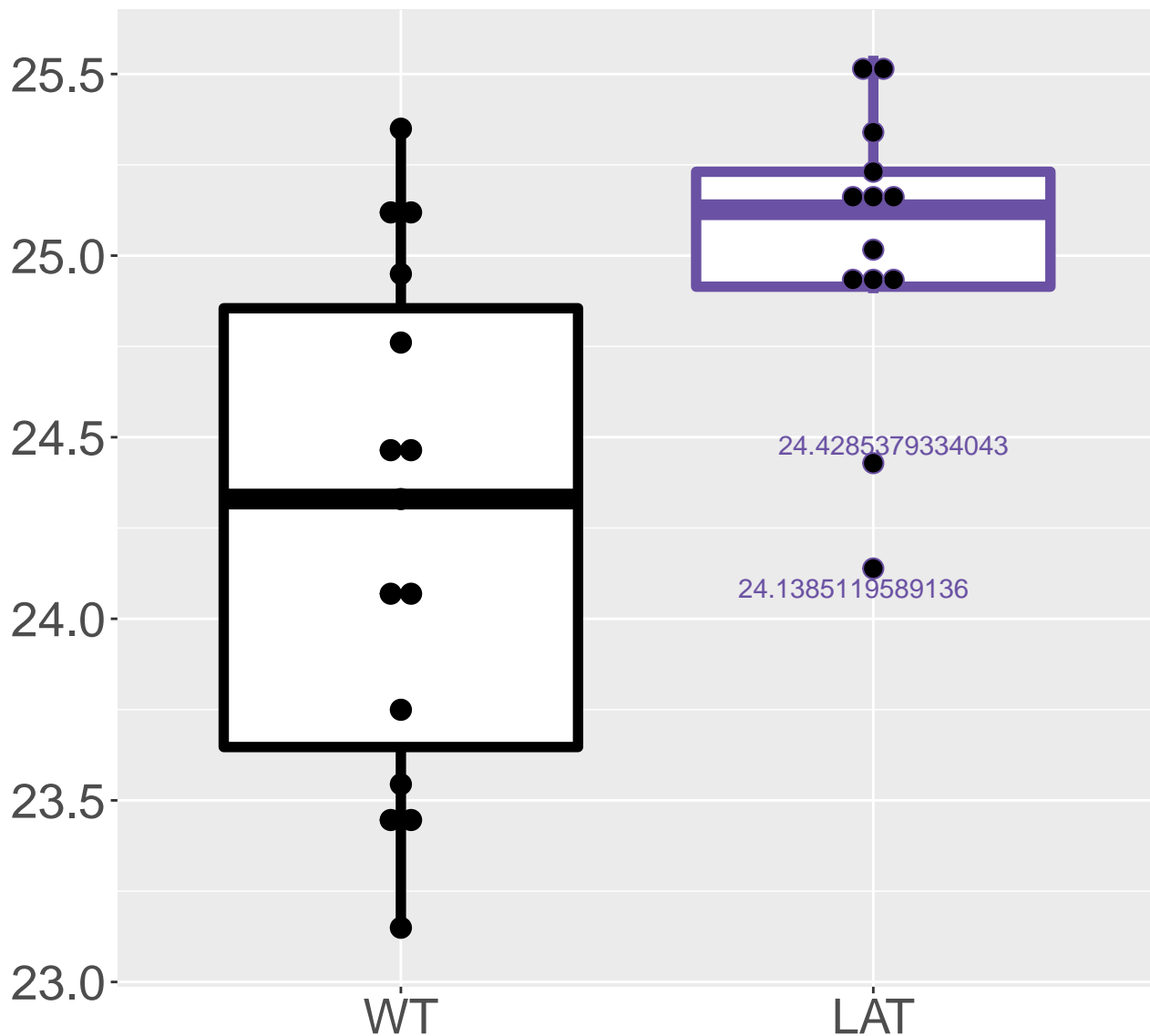


Q99JL6_Ras-related protein Rap-.
FDR = 0.04, FC = -0.19, sex**

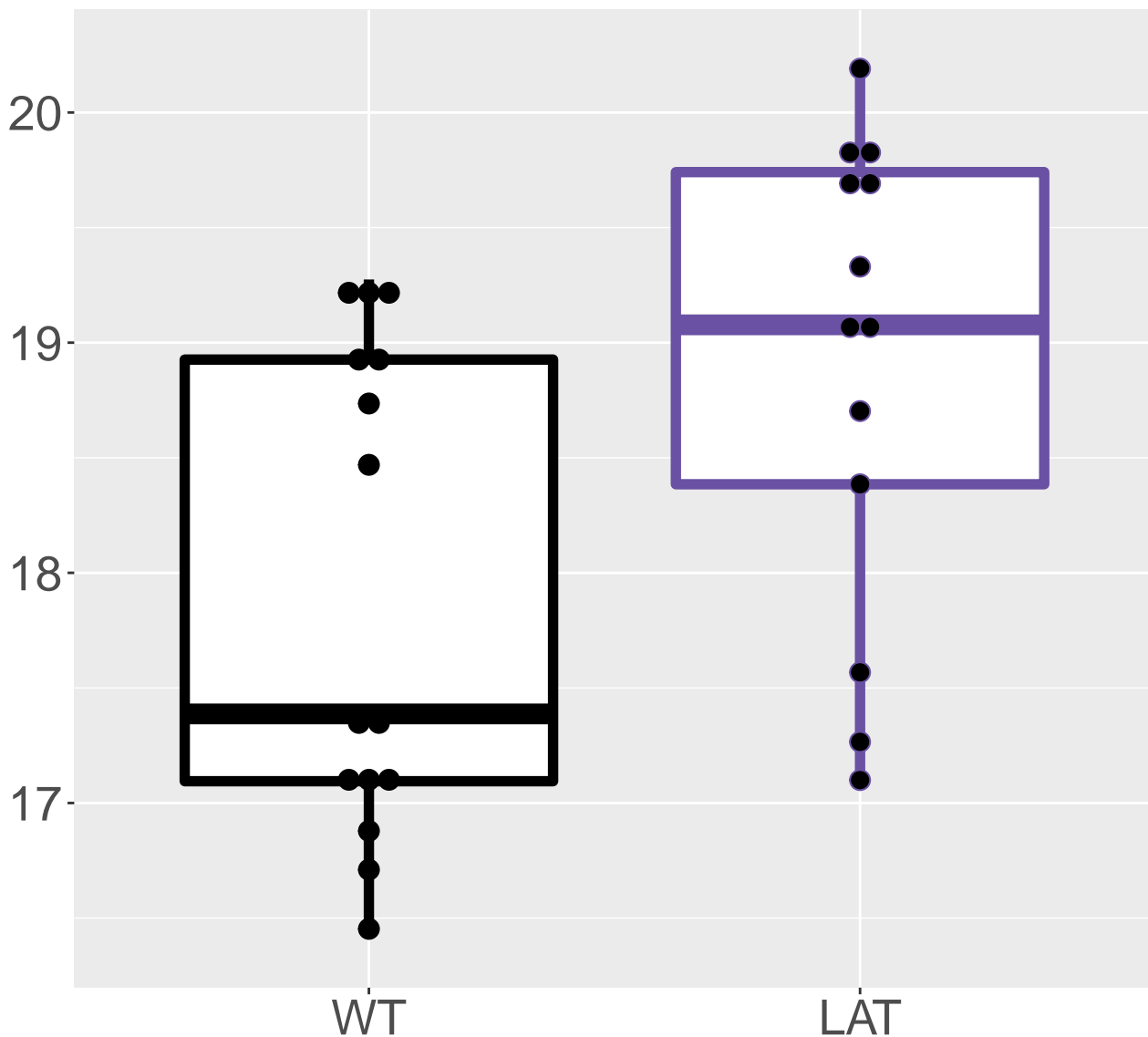


P13516_Acyl-CoA desaturase 1

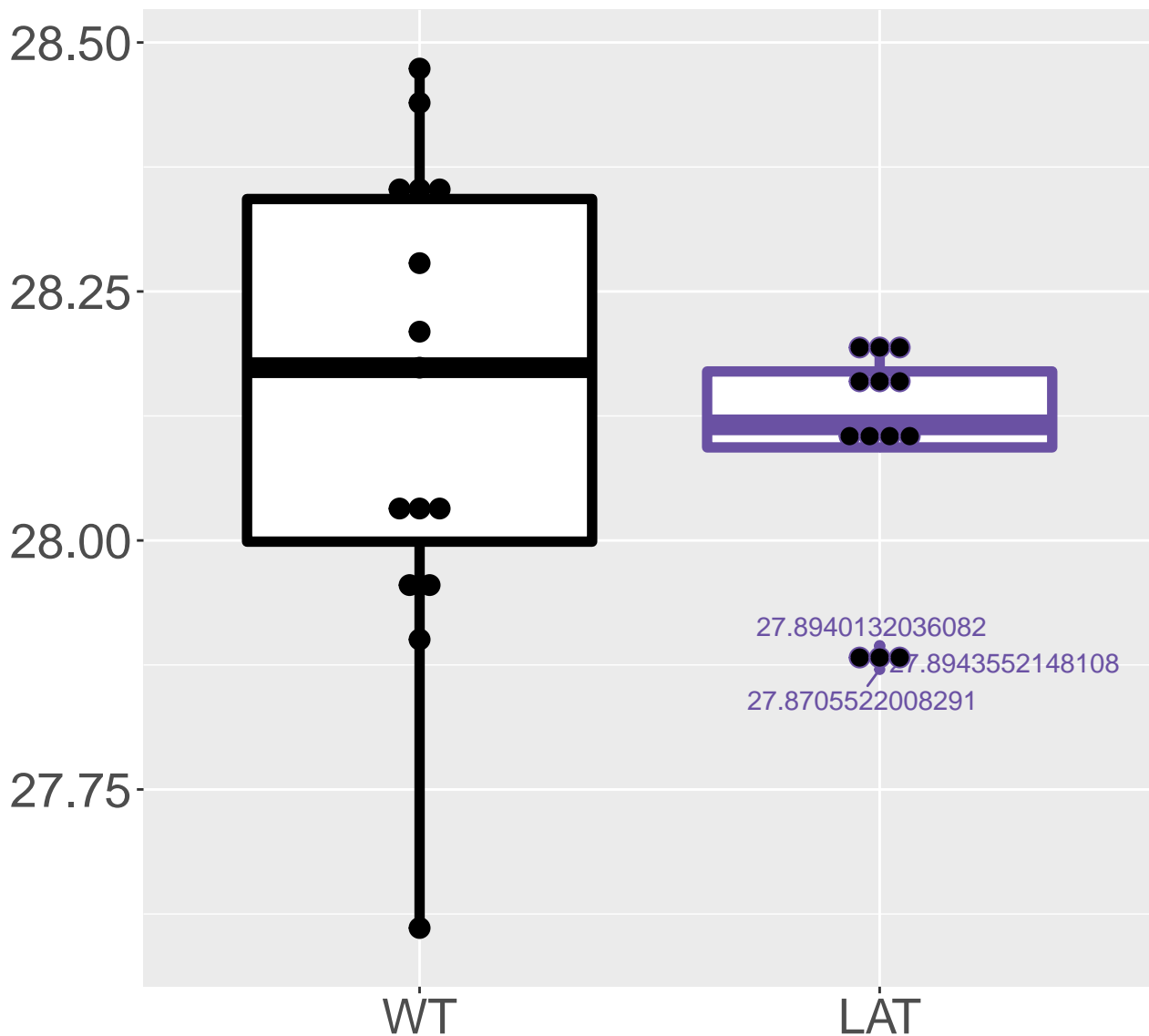
FDR = 0.041, FC = 0.93, sex*



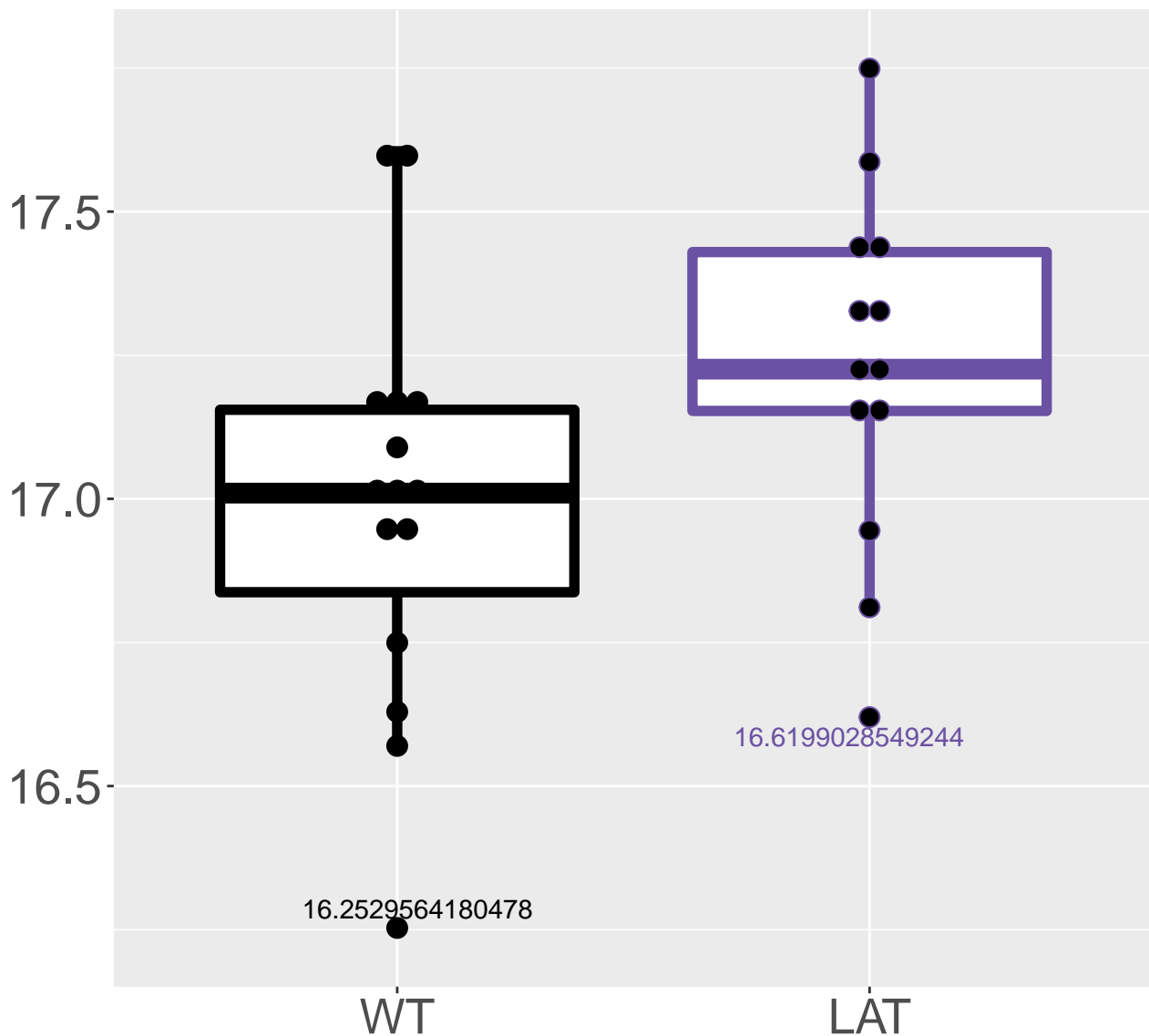
P28230_Gap junction beta-1 prot.
FDR = 0.041, FC = 1.6



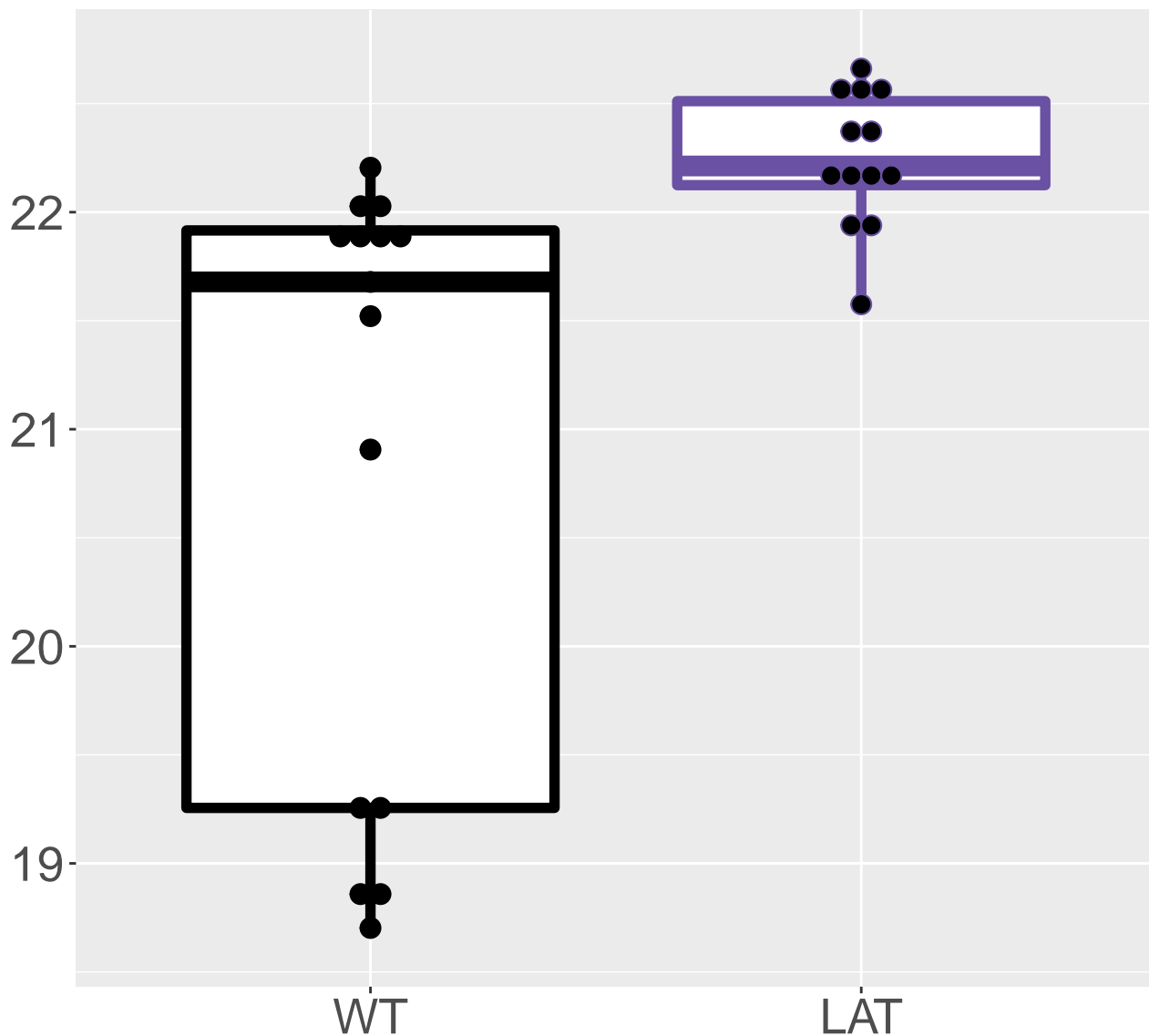
P51174_Long-chain specific acyl.
FDR = 0.042, FC = -0.17, sex***



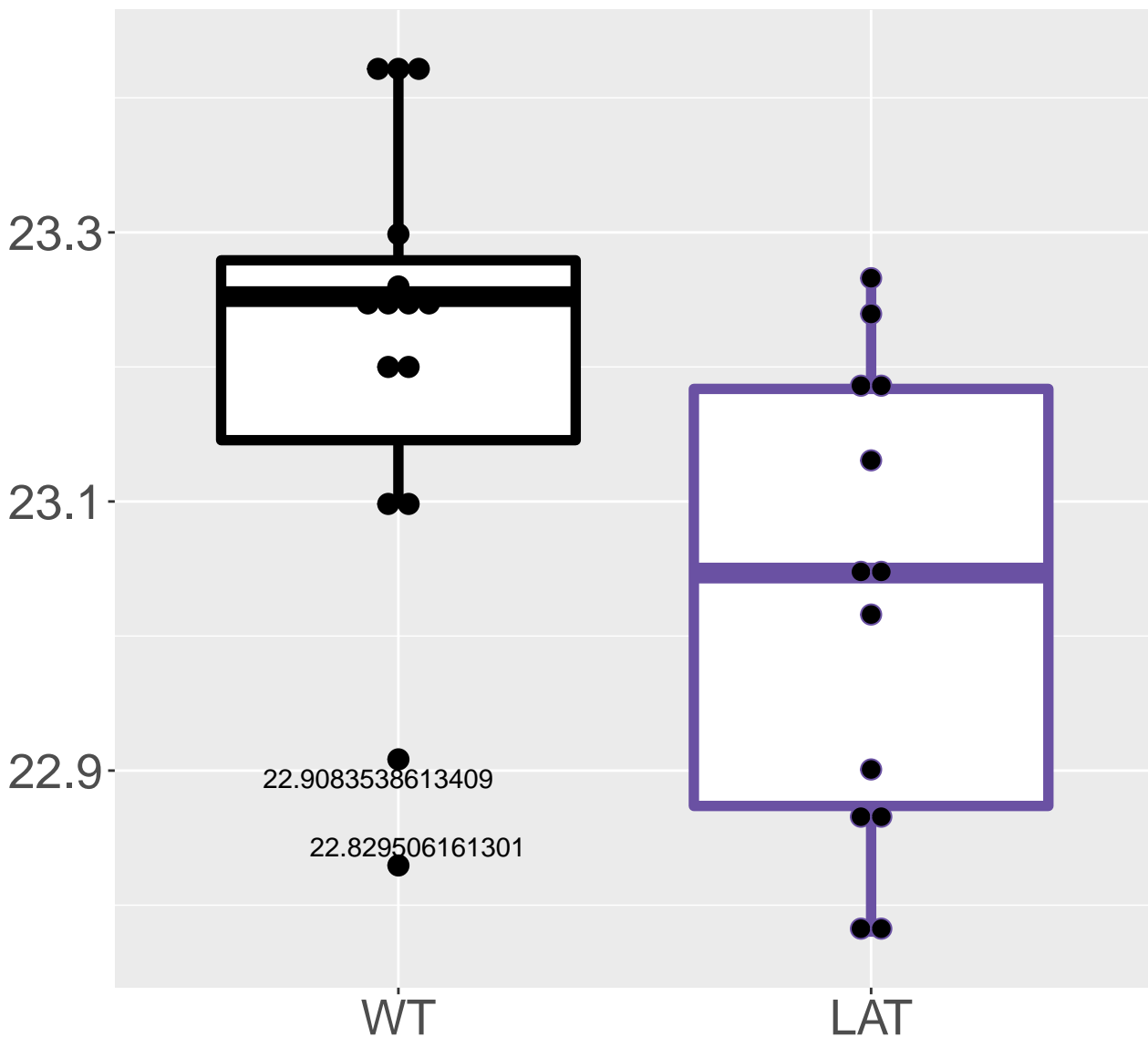
Q7TNP2_Serine/threonine-protein.
FDR = 0.042, FC = 0.39



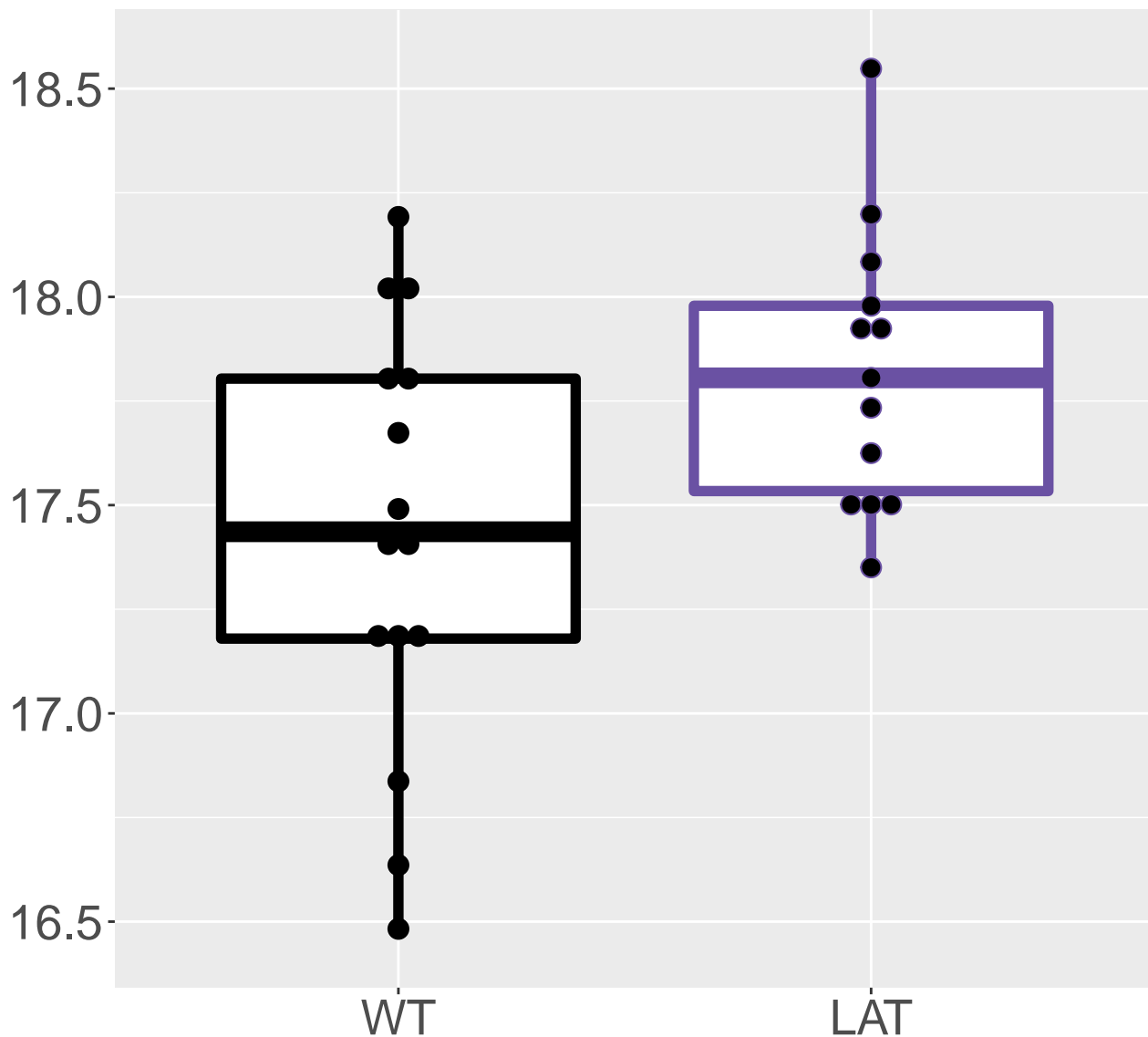
P00848_ATP synthase subunit a
FDR = 0.042, FC = 2



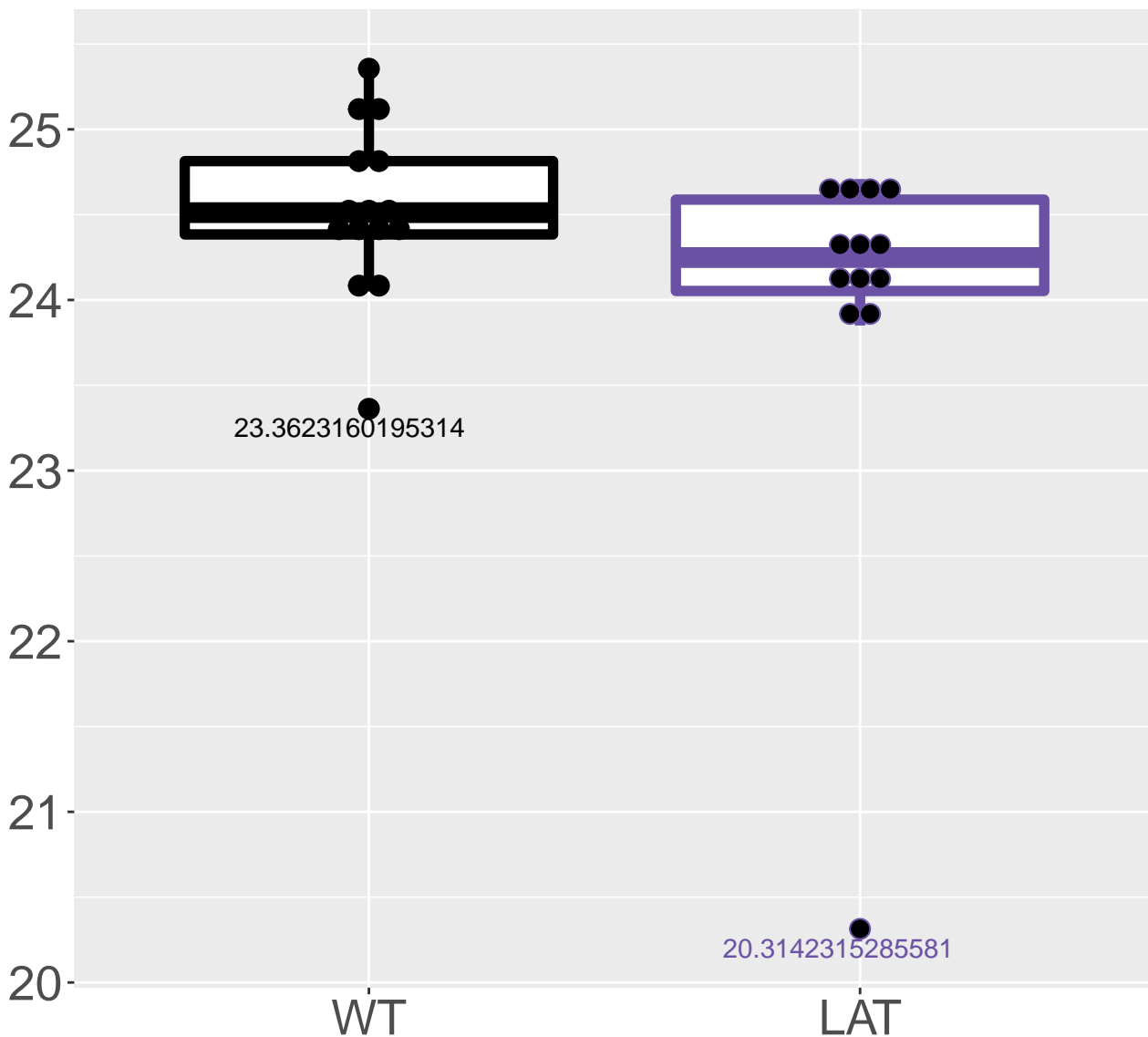
P47964_60S ribosomal protein L36
FDR = 0.042, FC = -0.31



O89051_Integral membrane protei.
FDR = 0.042, FC = 0.73

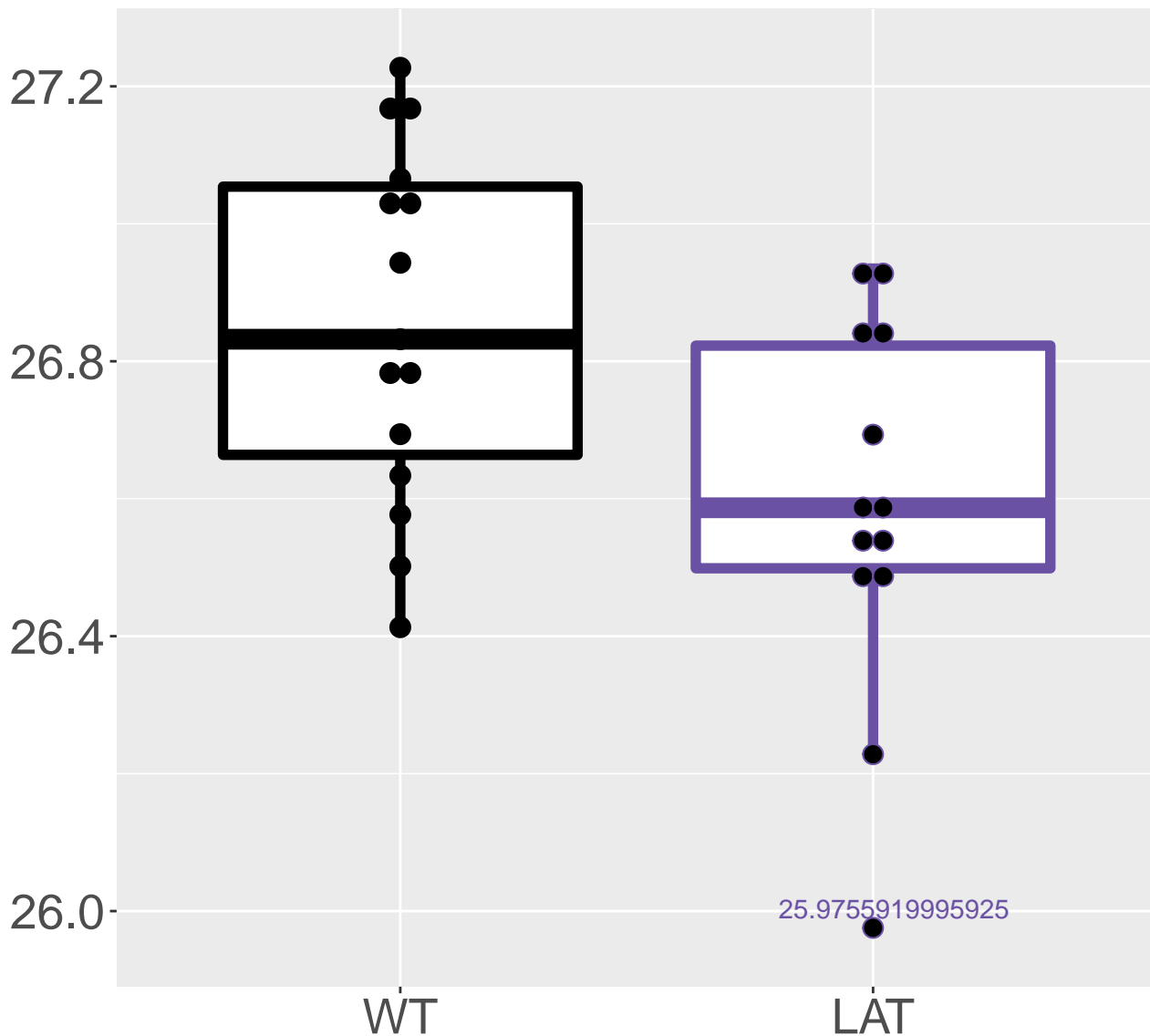


FDR = 0.044, FC = -0.6



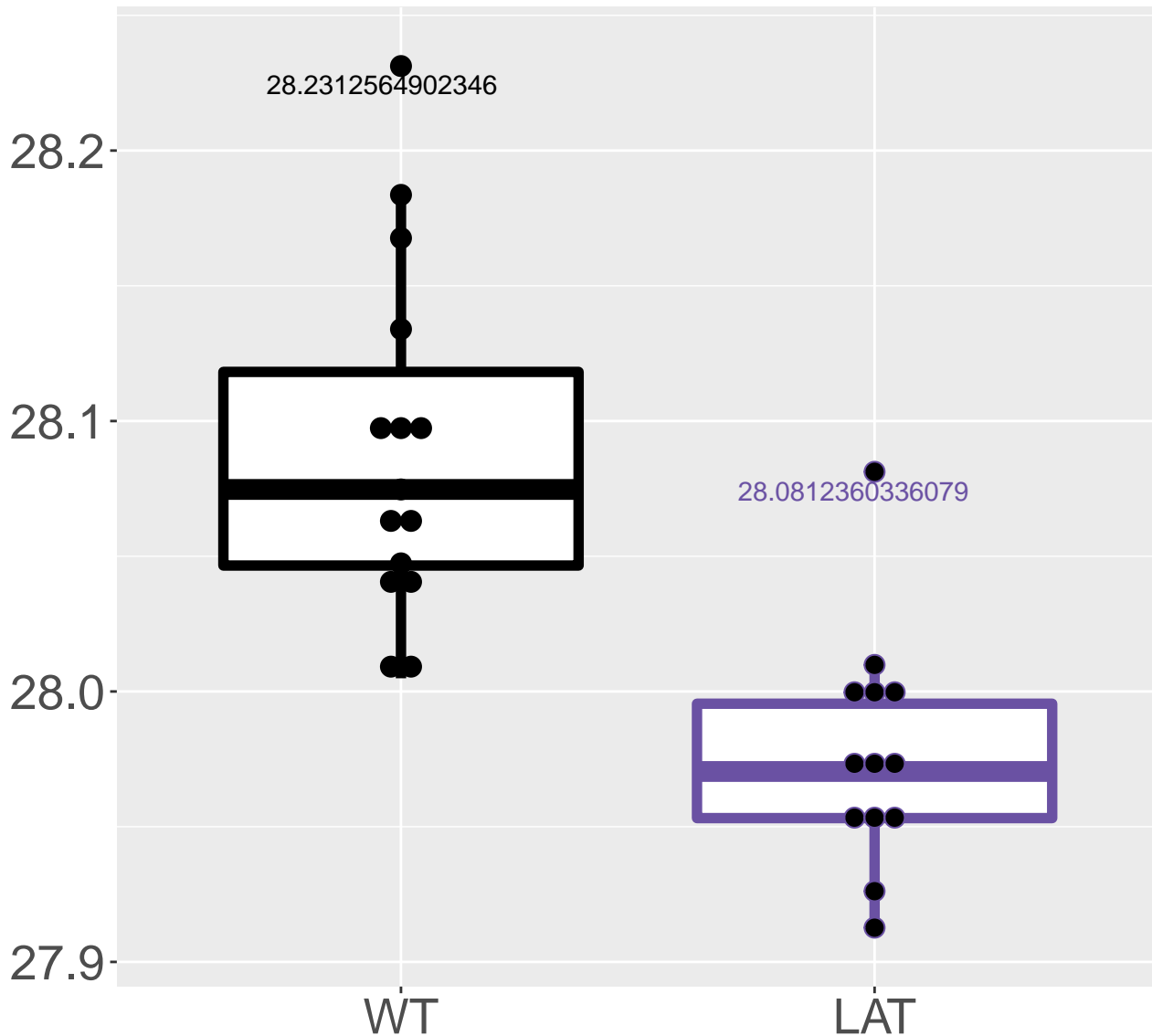
P10922_Histone H1.0

FDR = 0.044, FC = -0.27, sex***

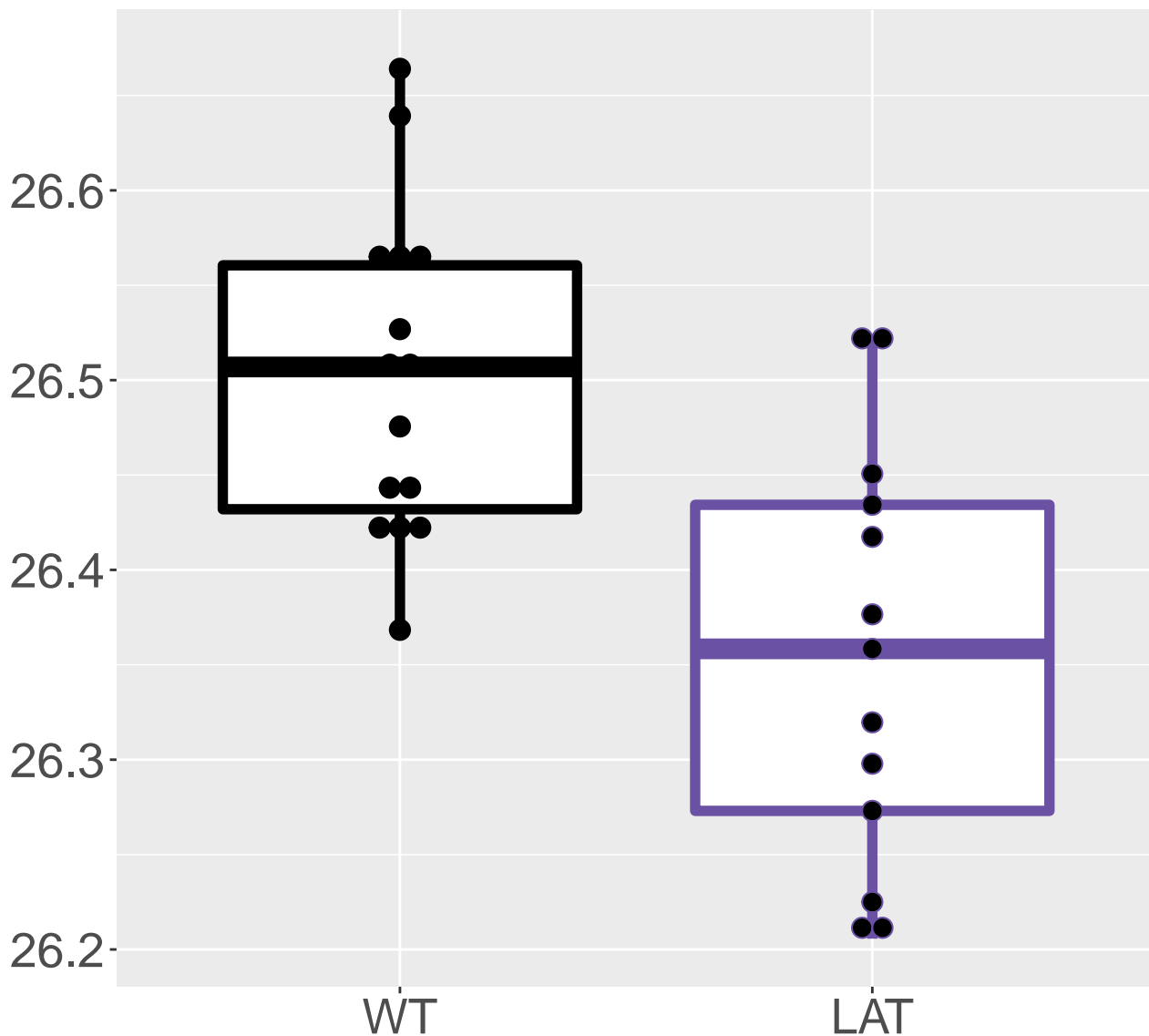


P62908_40S ribosomal protein S3

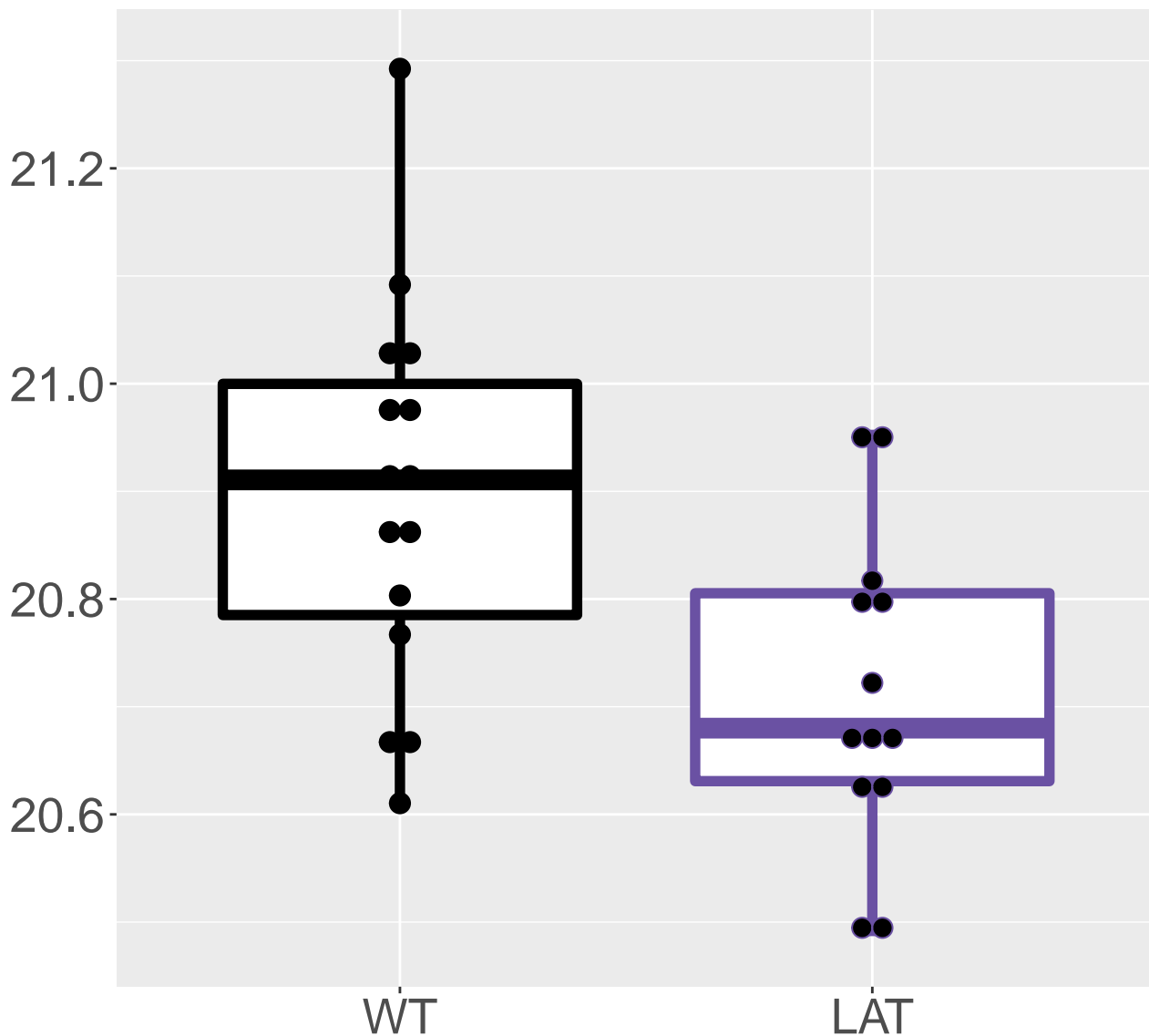
FDR = 0.044, FC = -0.11



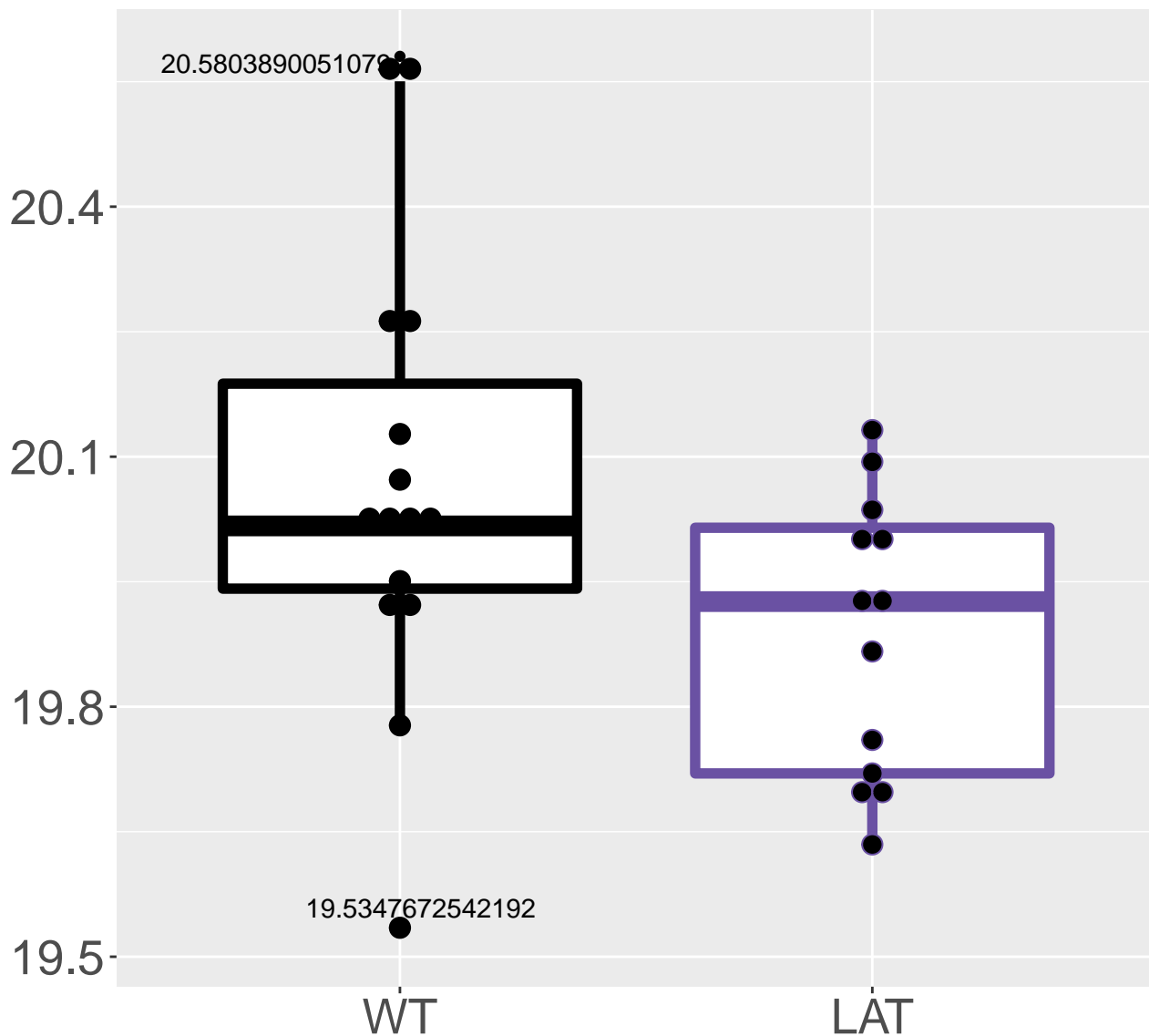
P62245_40S ribosomal protein S1.
FDR = 0.044, FC = -0.21



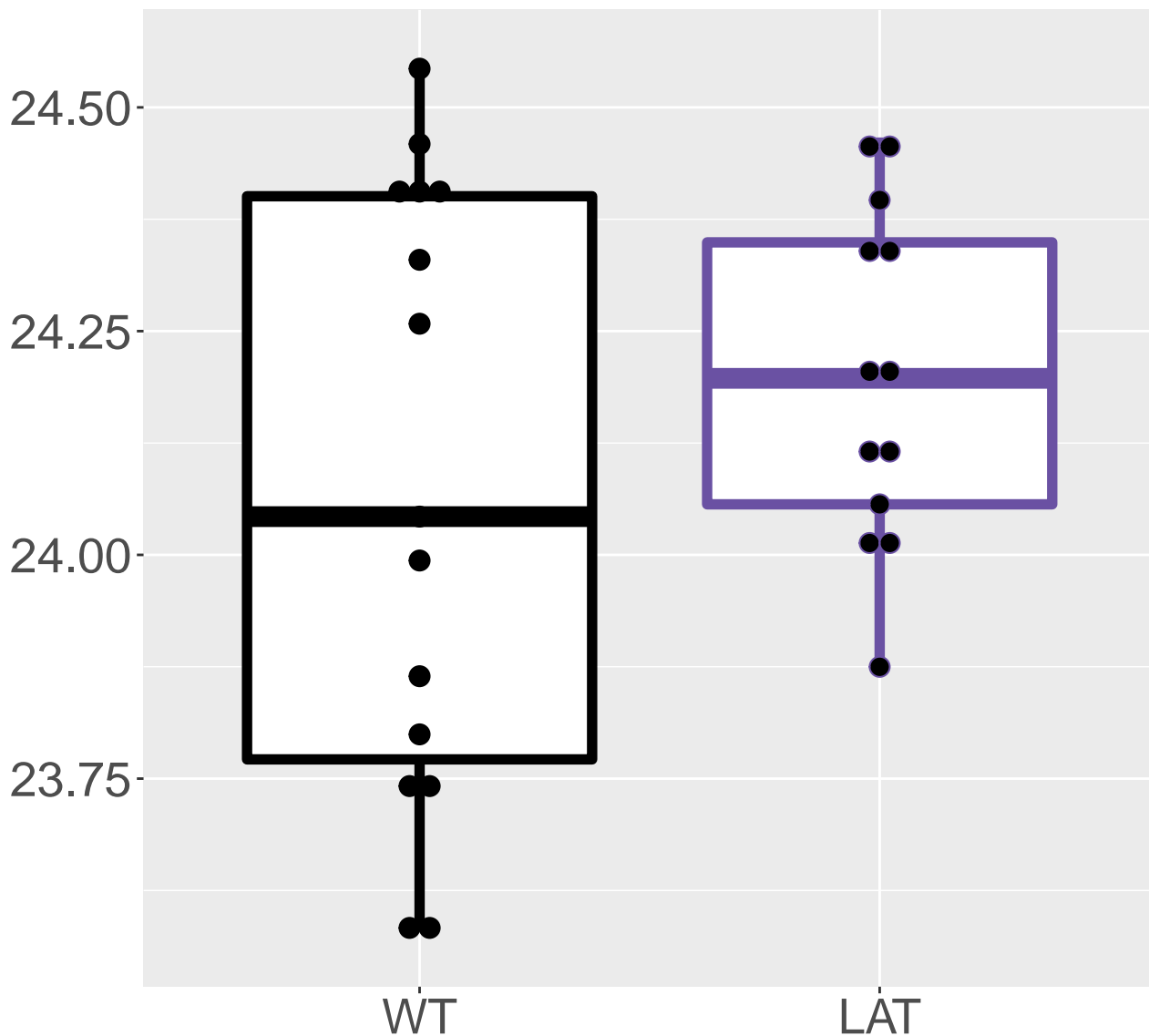
P63087_Serine/threonine-protein.
FDR = 0.044, FC = -0.2



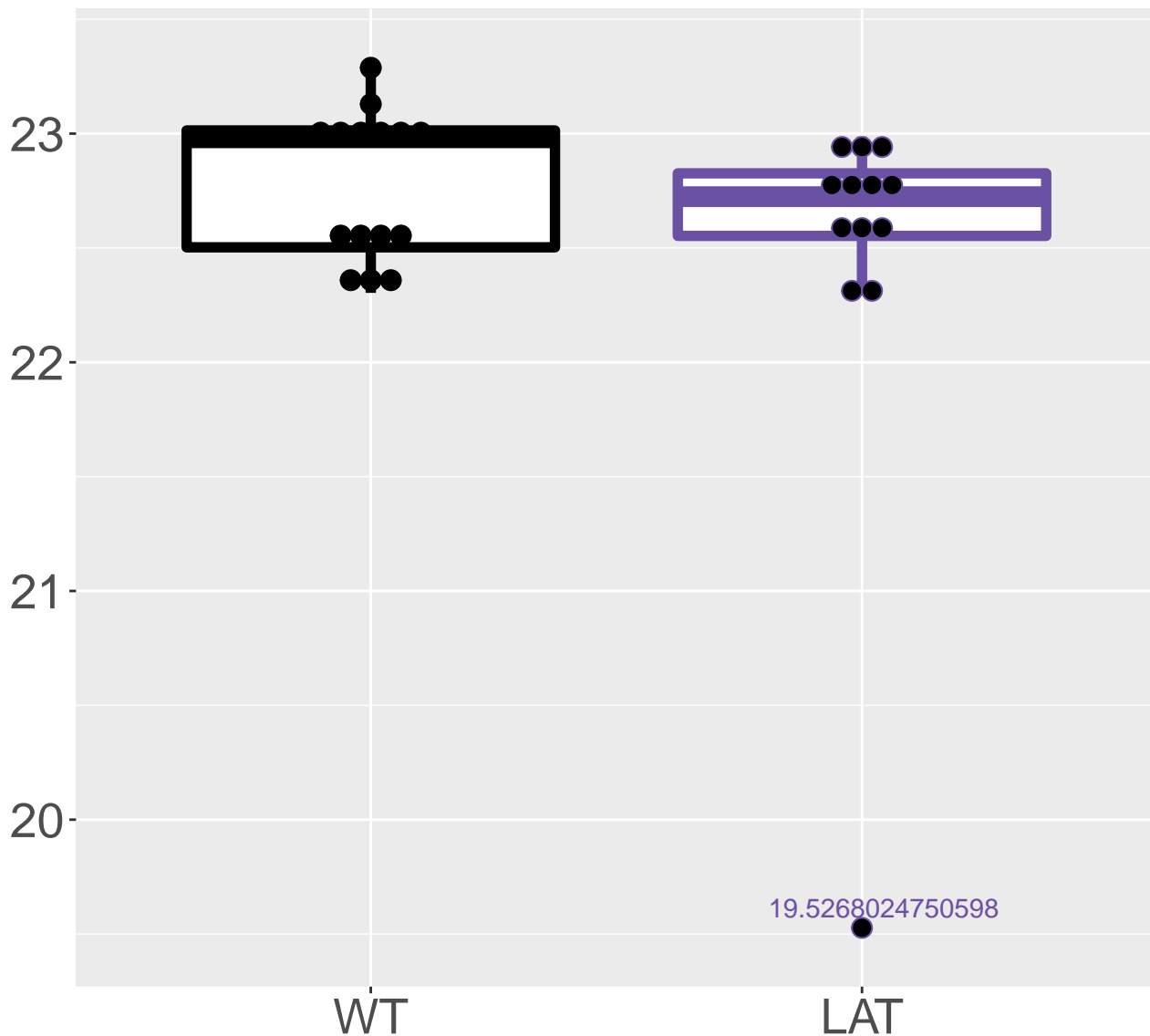
Q91XD6_Vacuolar protein-sorting.
FDR = 0.045, FC = -0.37, sex*



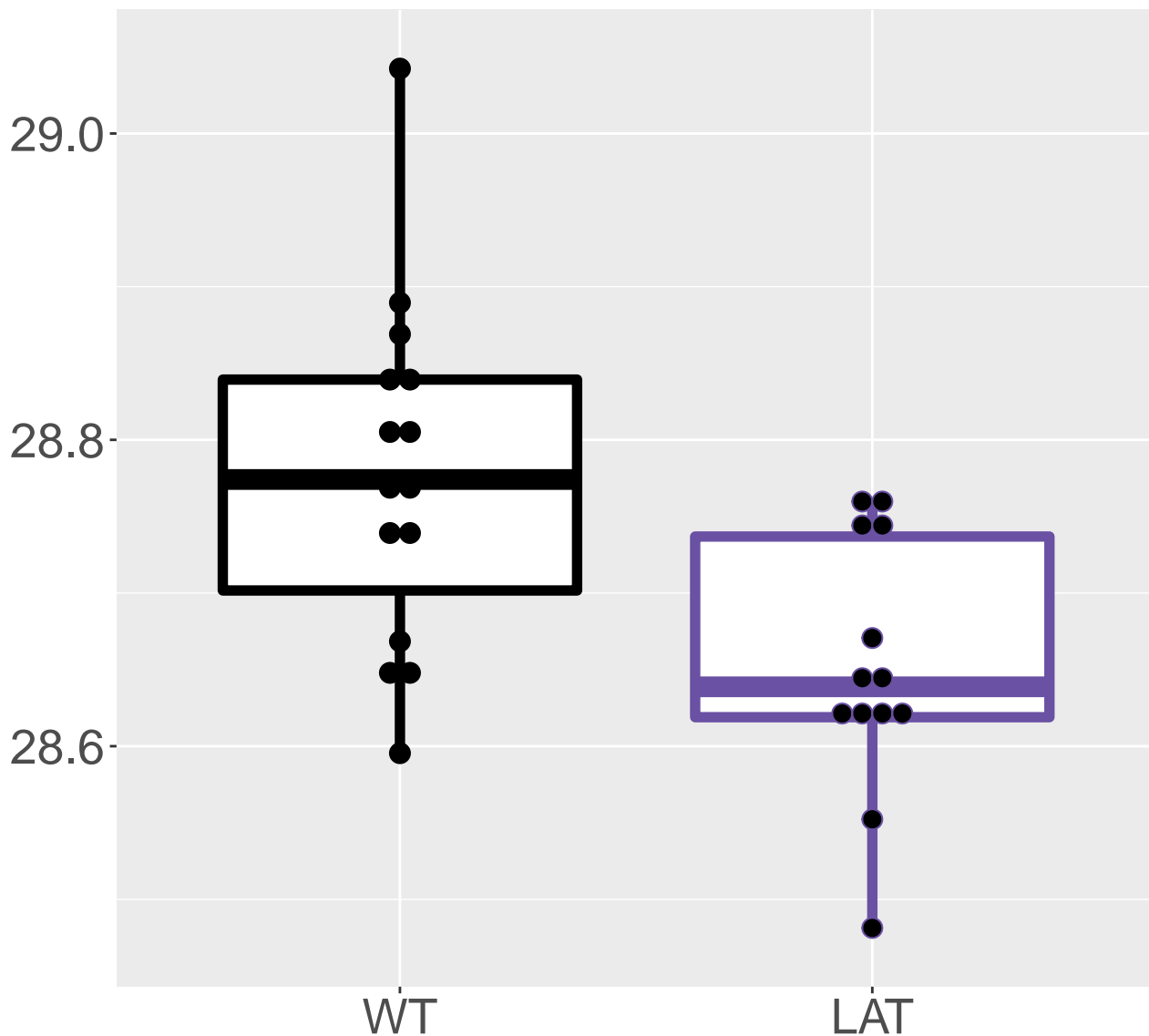
Q9QUR6_Prolyl endopeptidase
FDR = 0.045, FC = 0.28, sex***



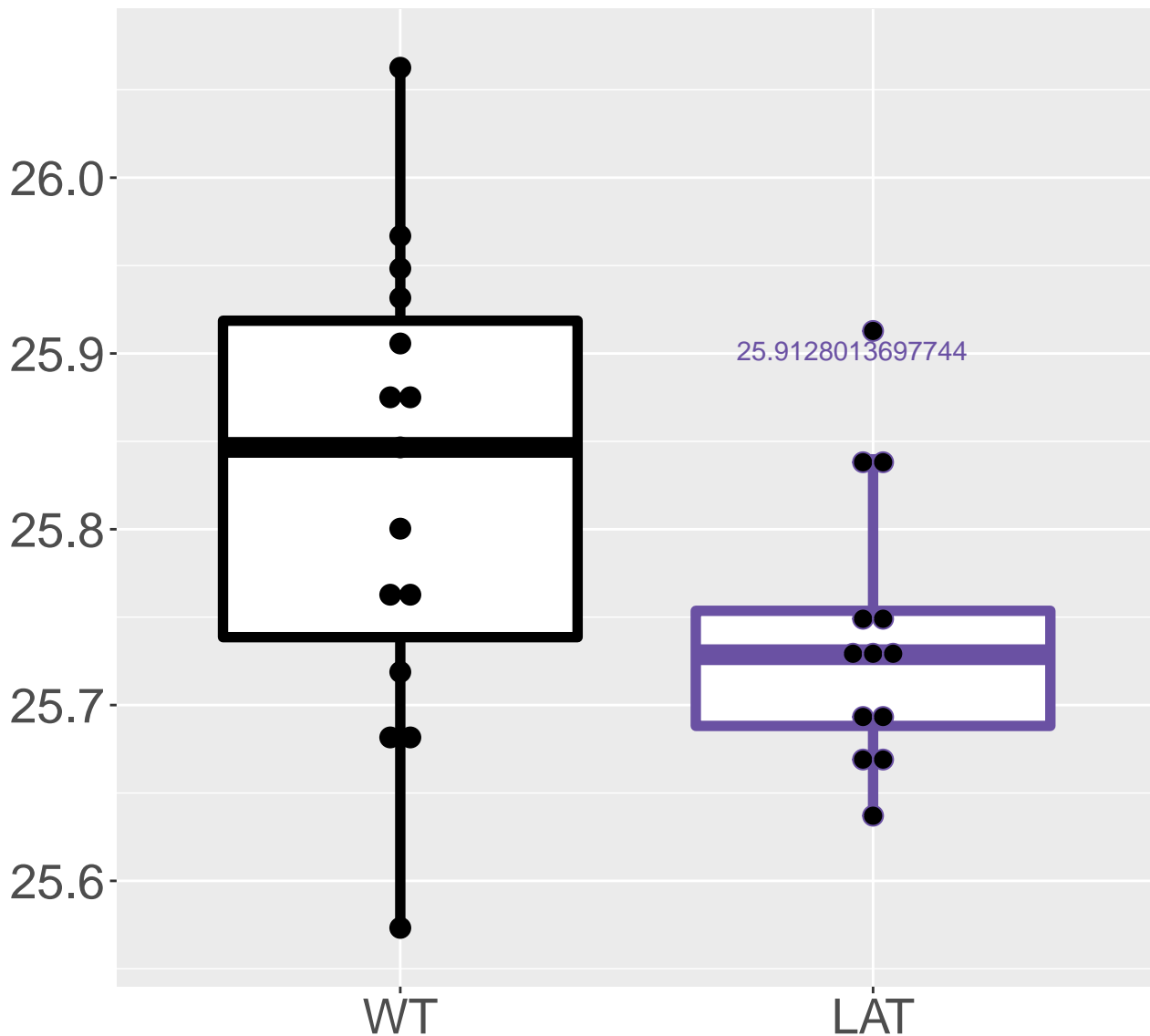
P13439_Uridine 5'-monophosphate.
FDR = 0.045, FC = -0.19, sex***



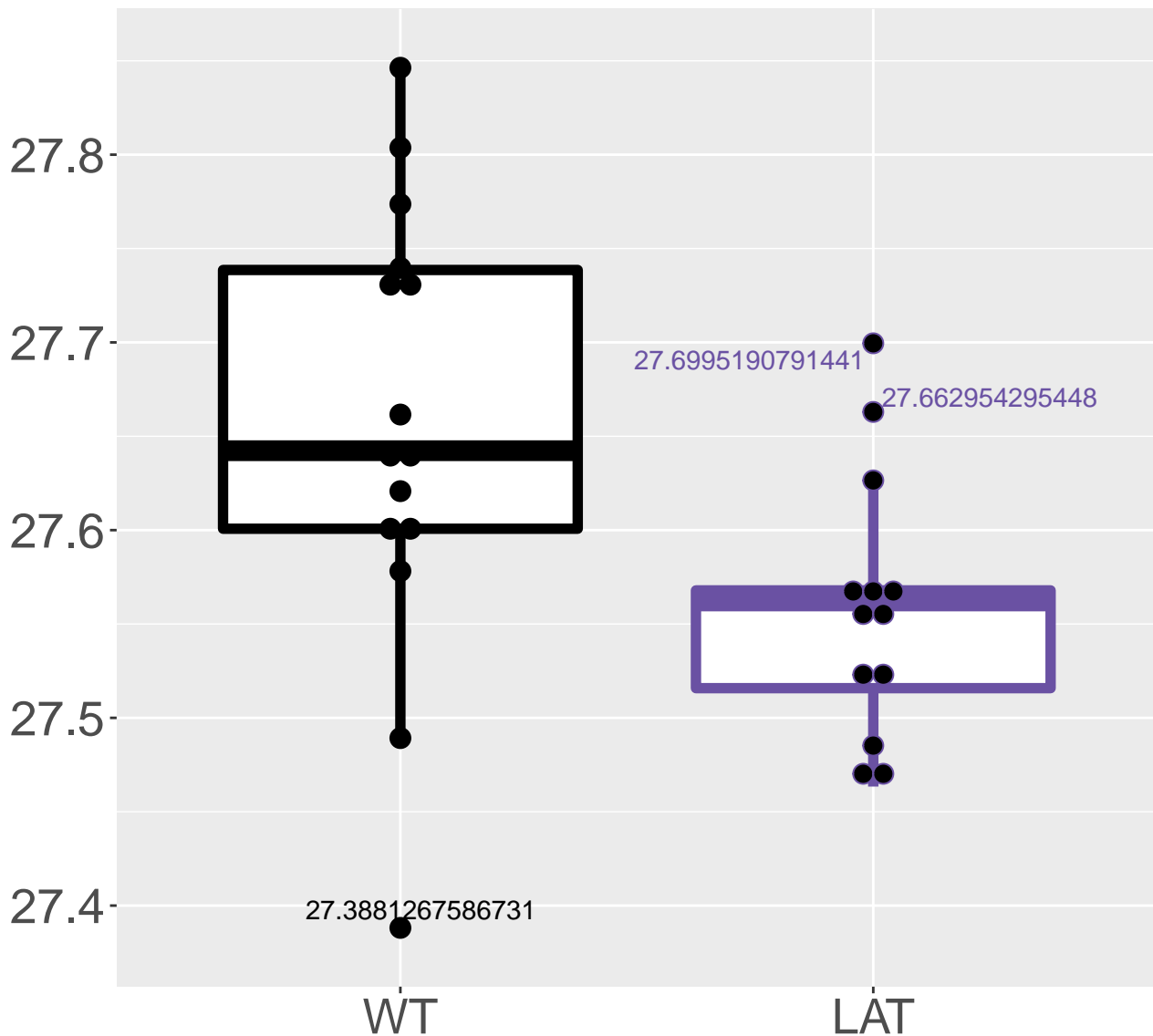
Q78JT3_3-hydroxyanthranilate 3,,
FDR = 0.045, FC = -0.18



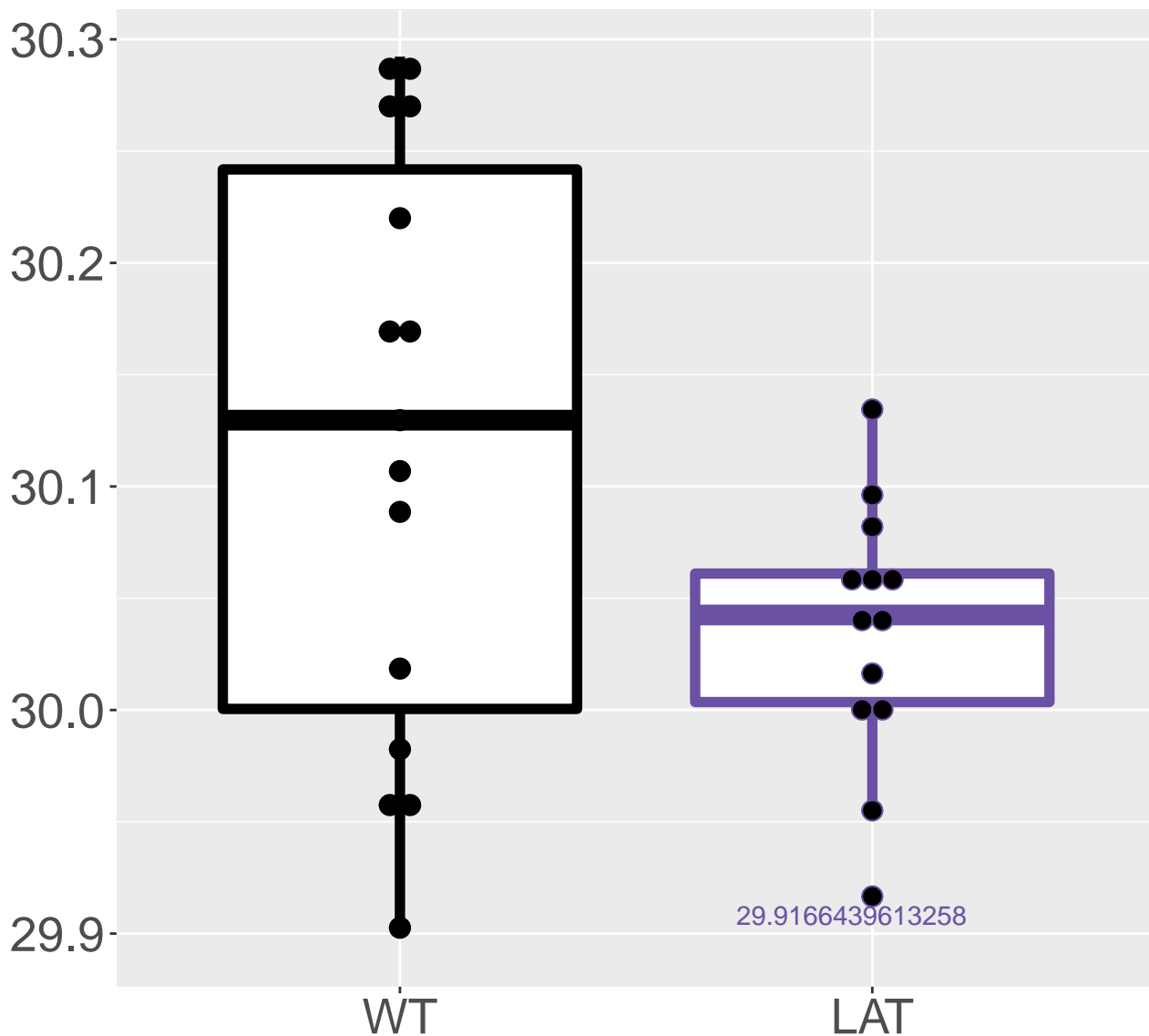
Q8C5H8_NAD kinase 2, mitochondr.
FDR = 0.045, FC = -0.16, sex**



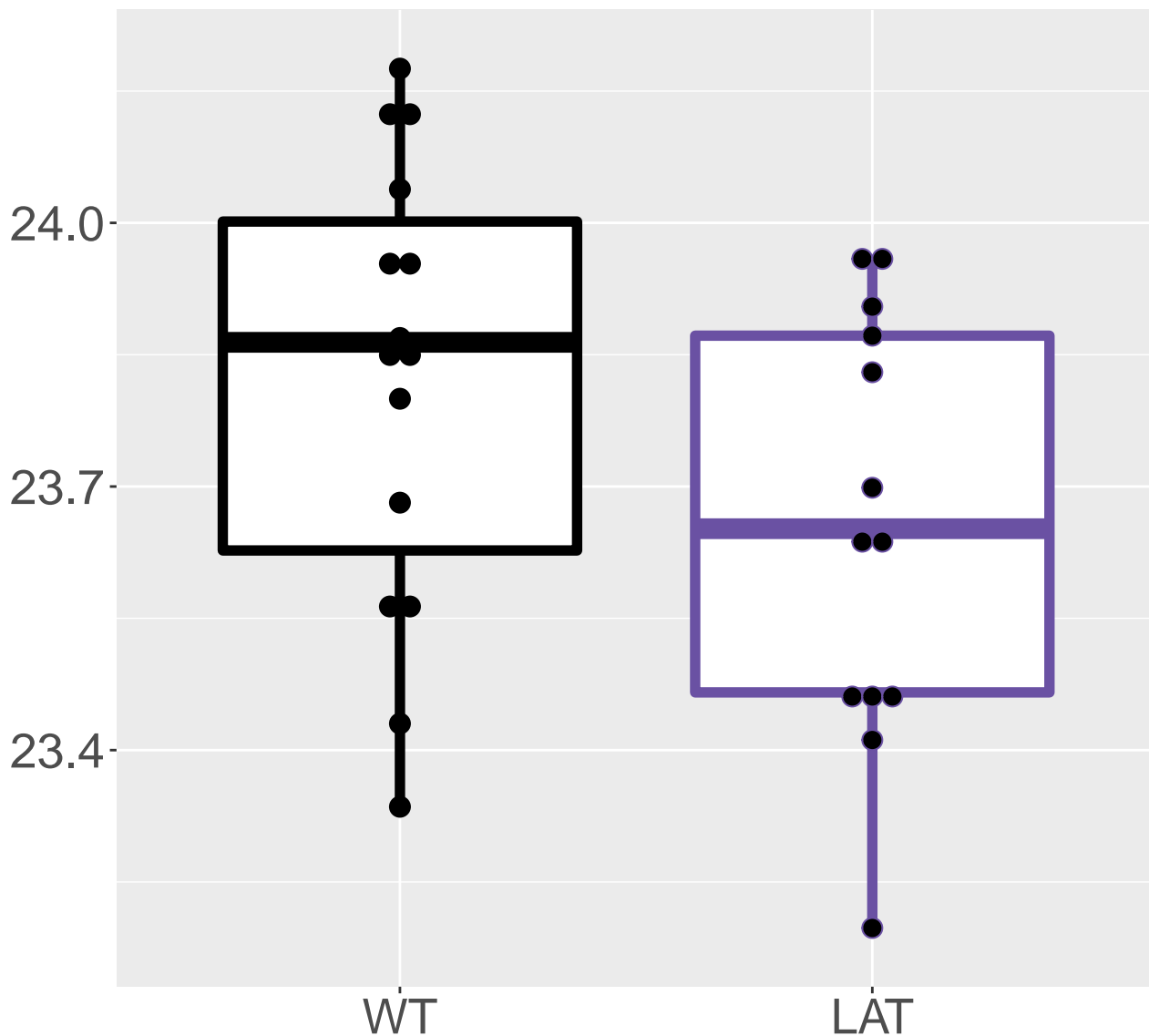
P09671_Superoxide dismutase [Mn.
FDR = 0.045, FC = -0.15, sex*



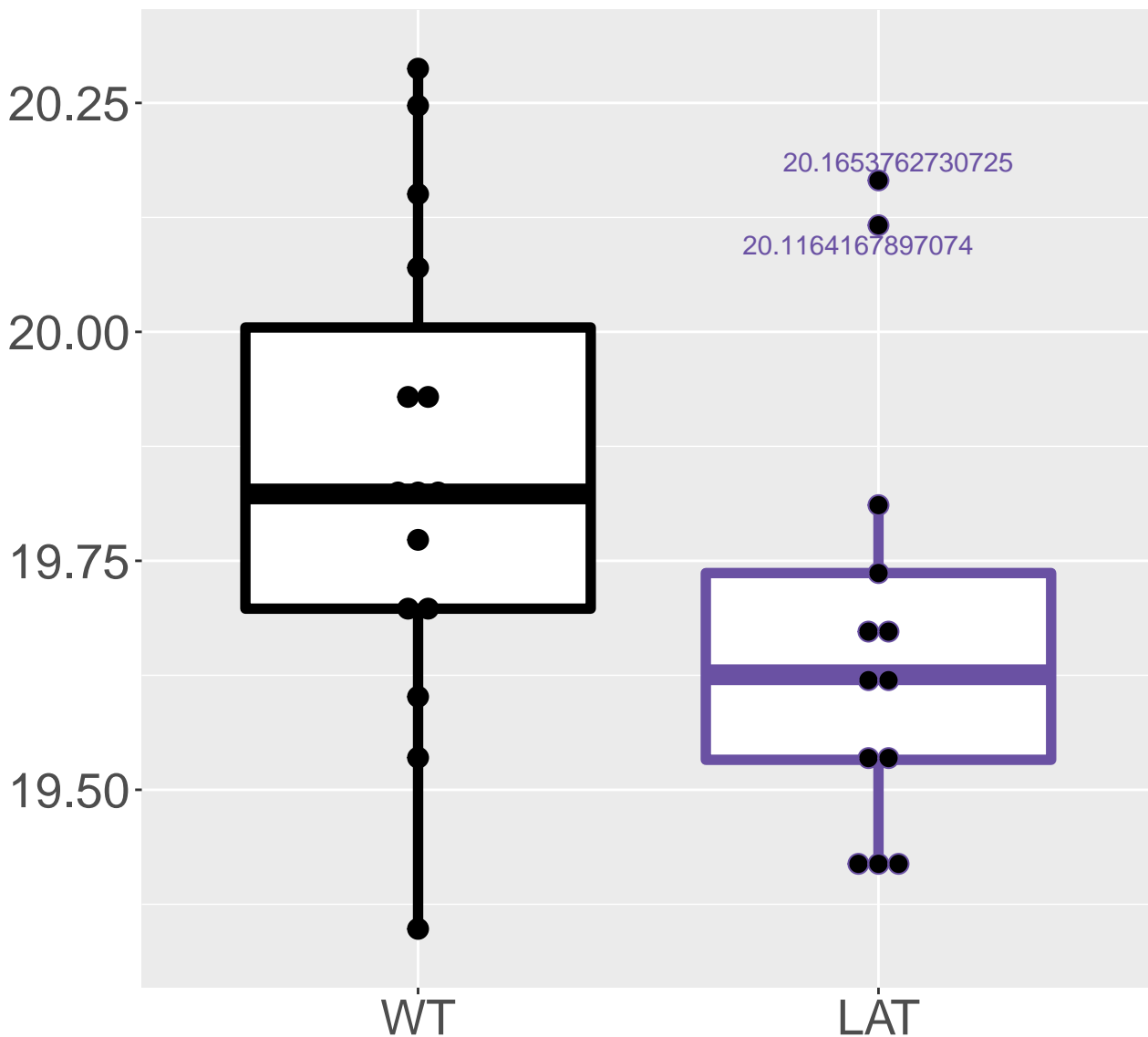
P35700_Peroxiredoxin-1
FDR = 0.045, FC = -0.14, sex**



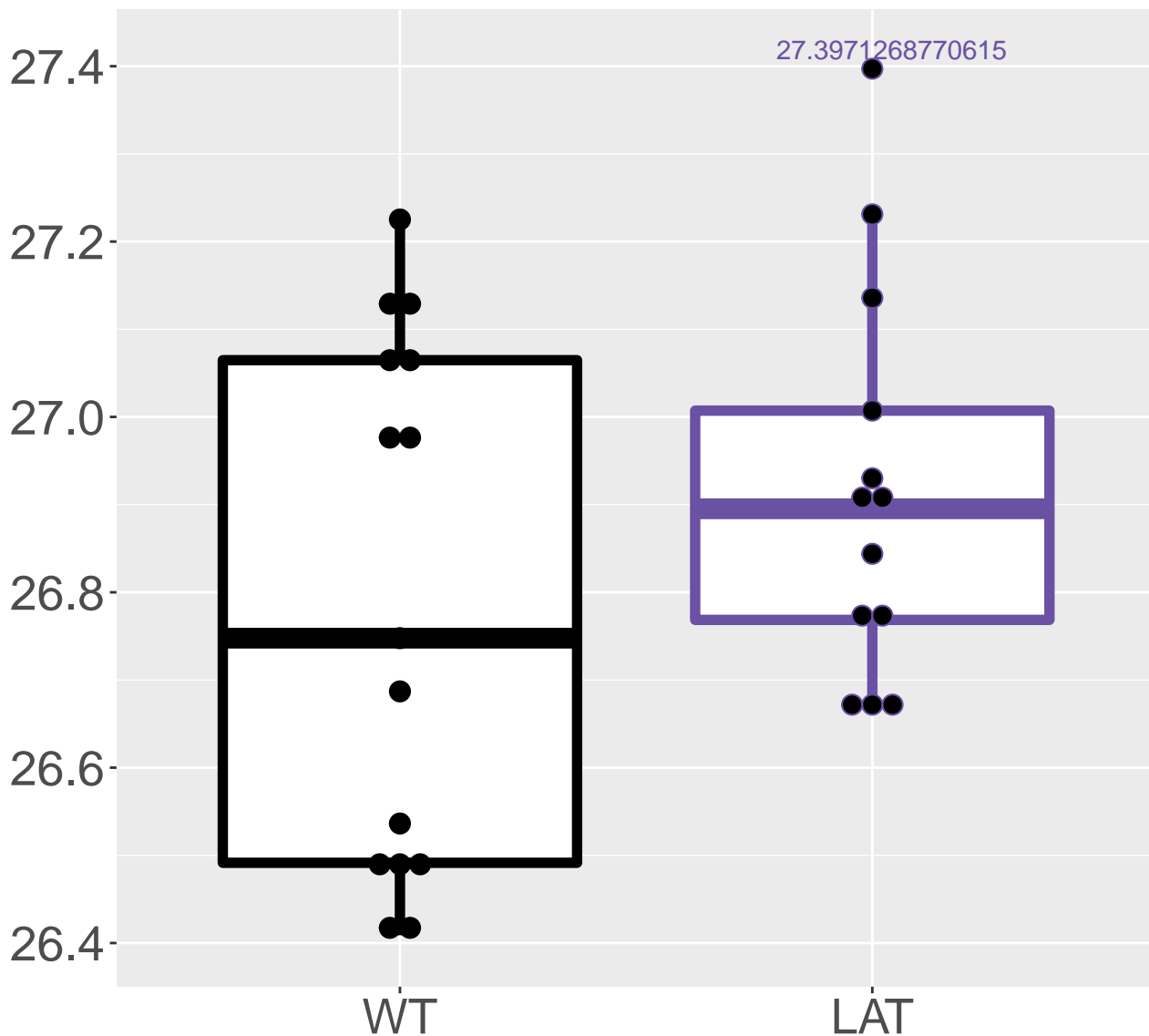
Q61207_Prosaposin
FDR = 0.045, FC = -0.34, sex**



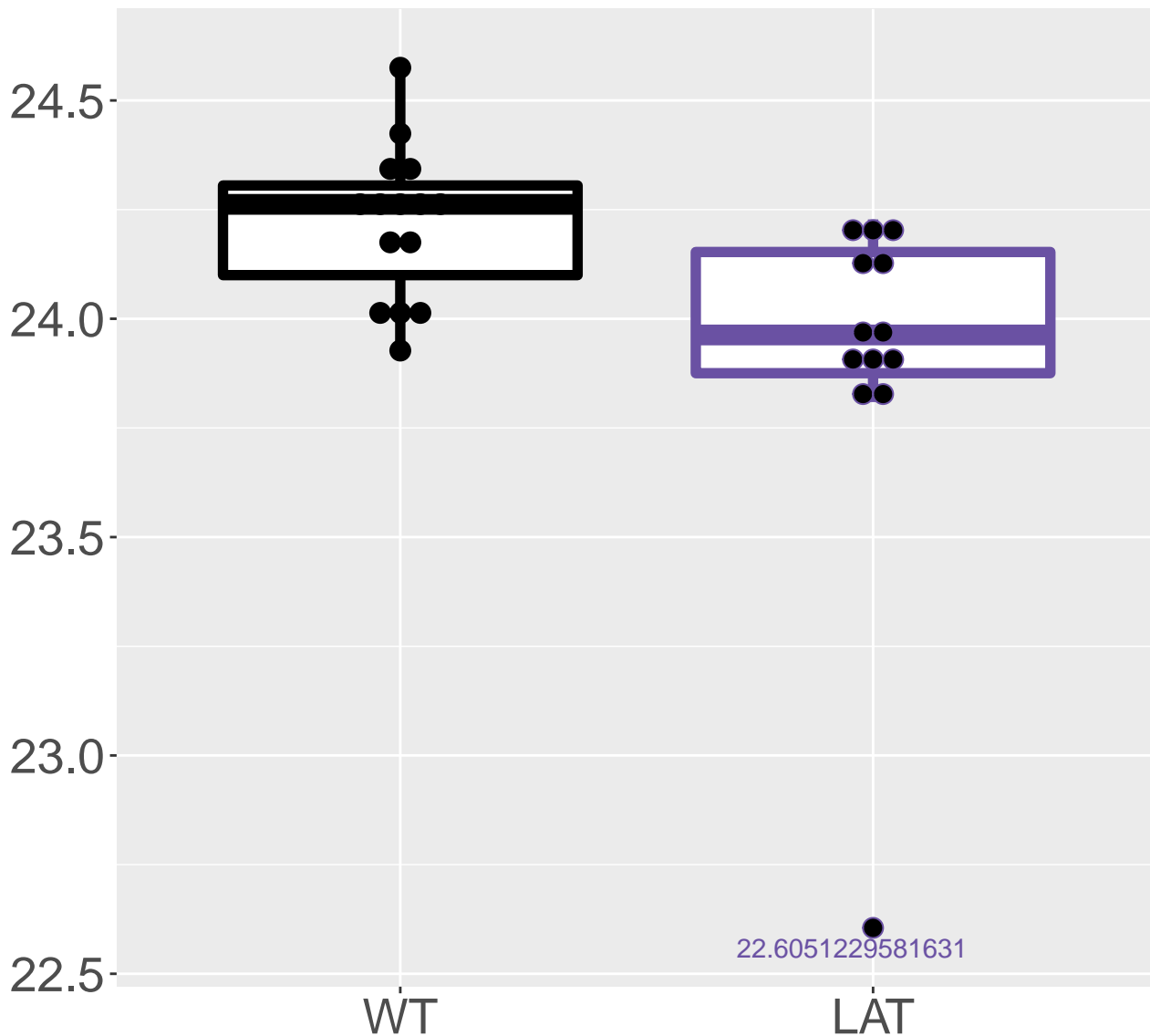
Q9CQ89_Protein CutA
FDR = 0.045, FC = -0.42



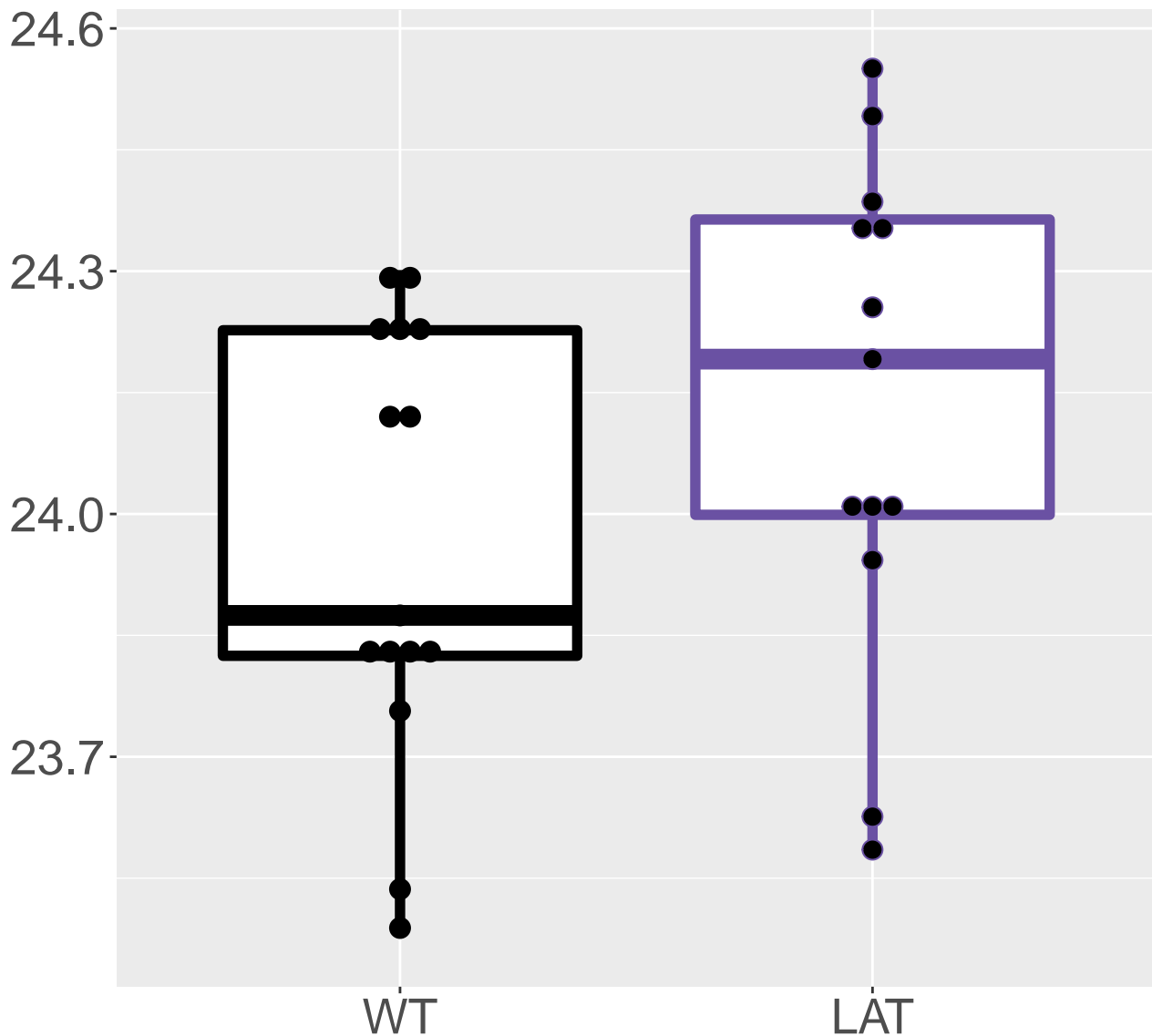
O70475_UDP-glucose 6-dehydrogen.
FDR = 0.045, FC = 0.21, sex***



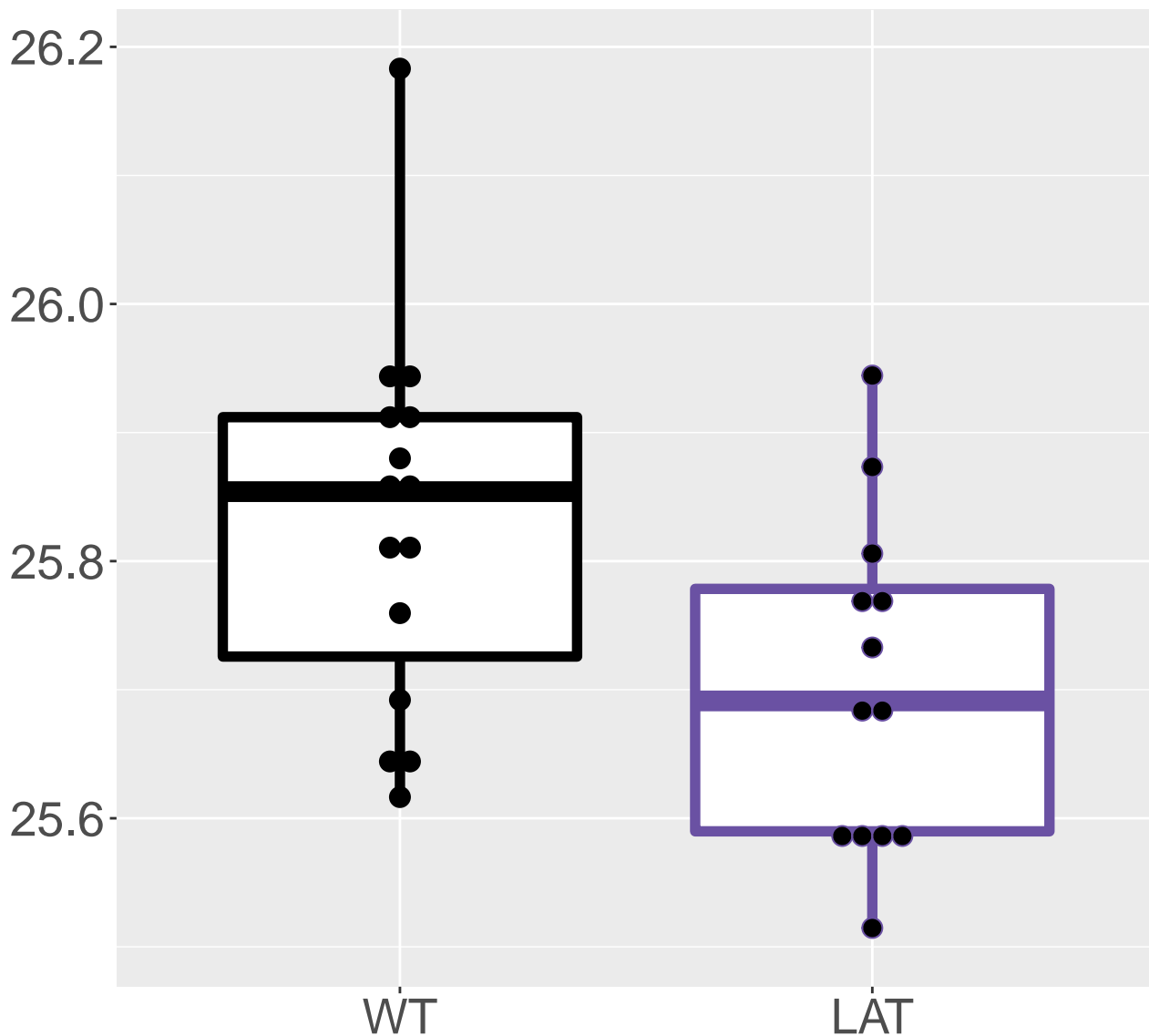
P62858_40S ribosomal protein S28
FDR = 0.046, FC = -0.63



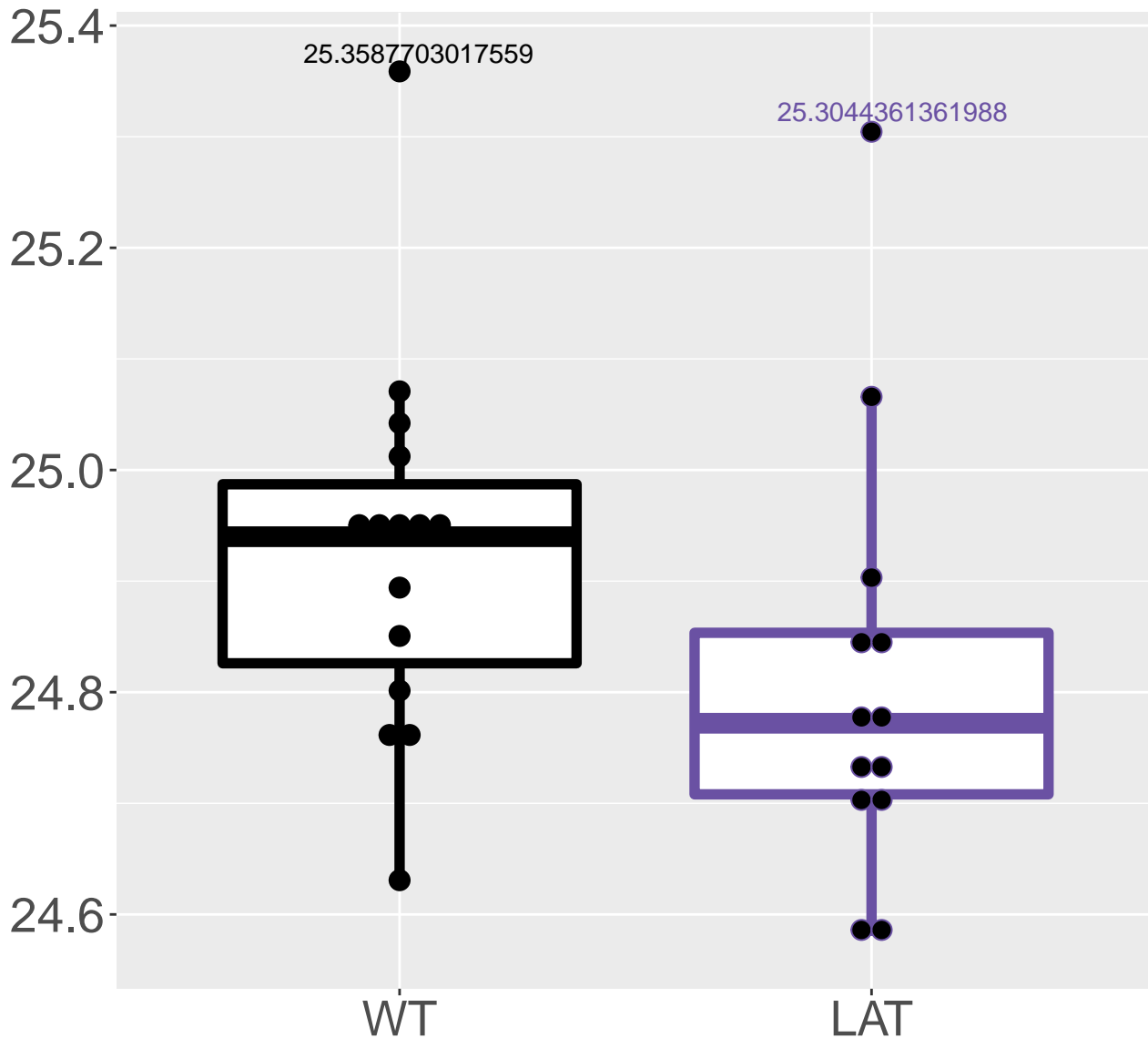
Q9EQK5_Major vault protein
FDR = 0.046, FC = 0.42



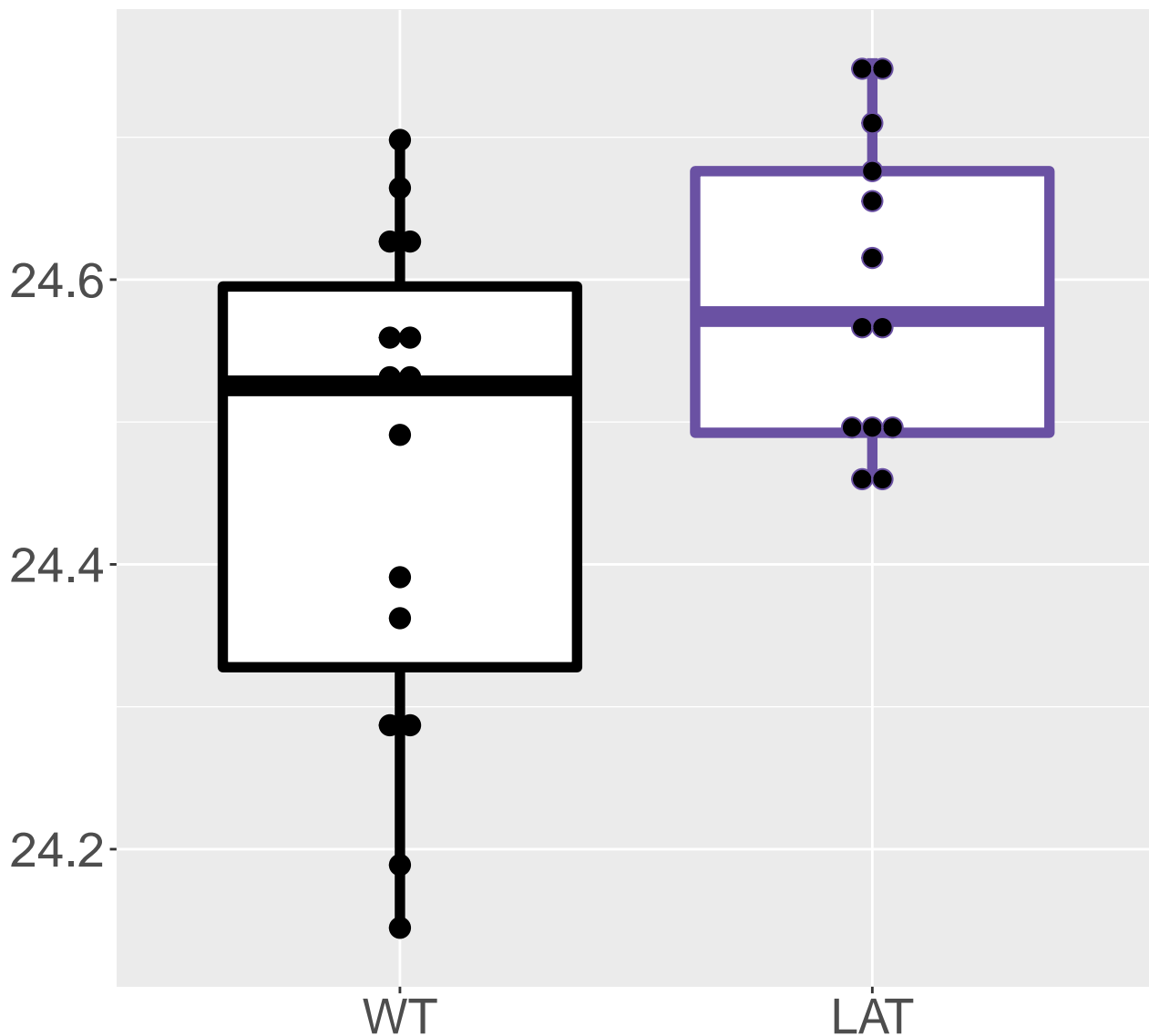
Q922Q8_Leucine-rich repeat-cont.
FDR = 0.046, FC = -0.19



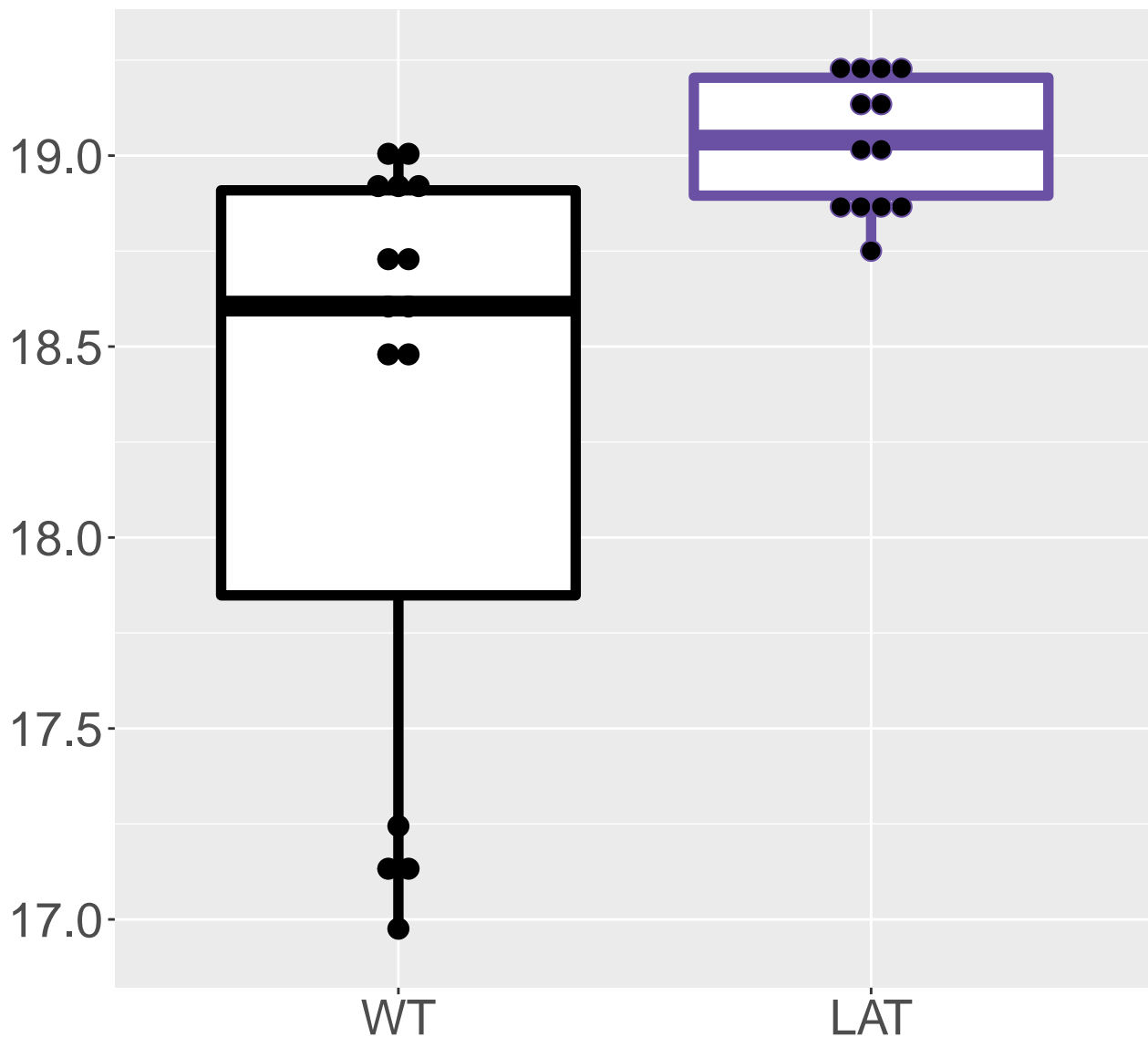
O88533_Aromatic-L-amino-acid de.
FDR = 0.046, FC = -0.16



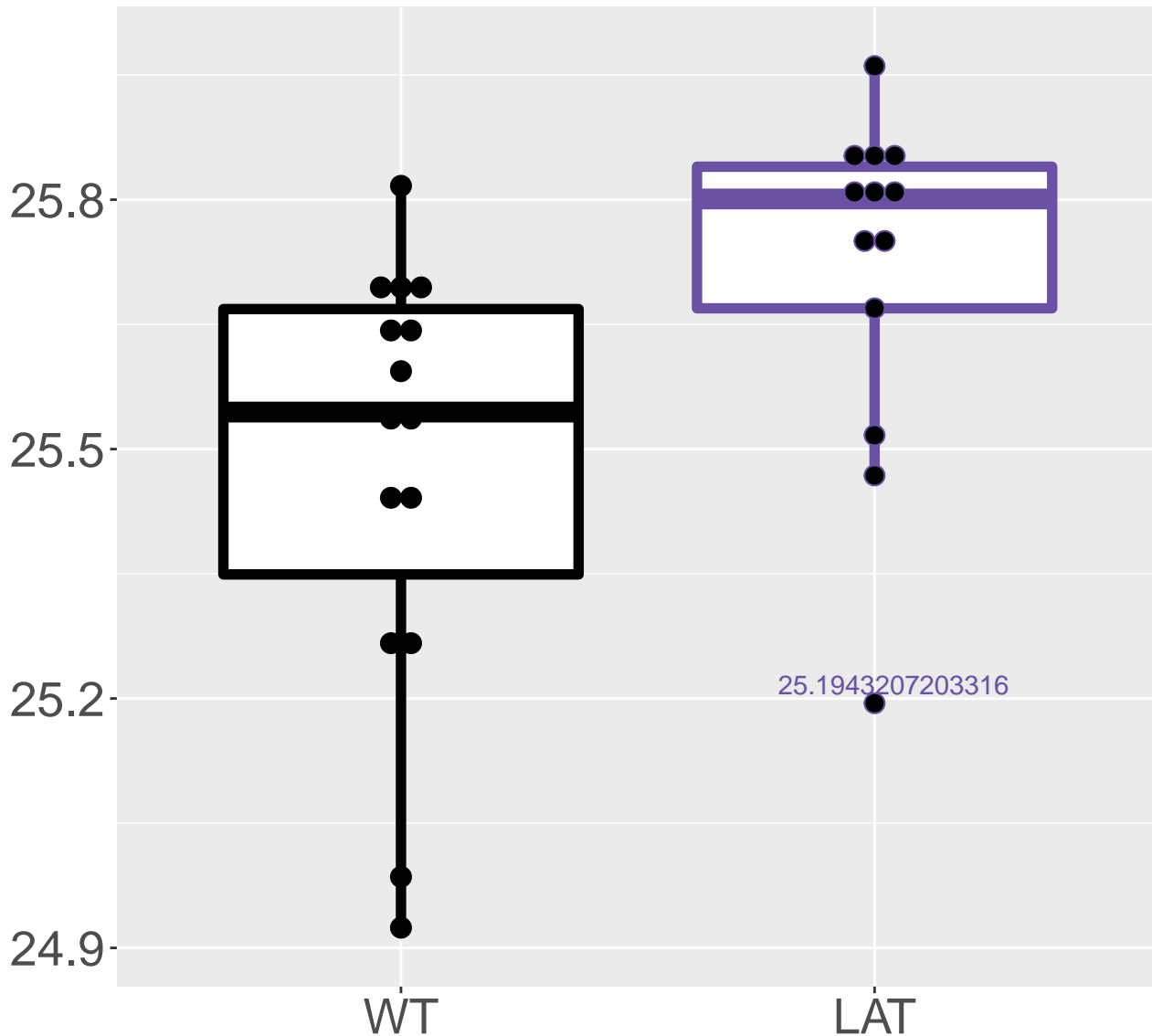
Q01405_Protein transport protei.
FDR = 0.046, FC = 0.25, sex*



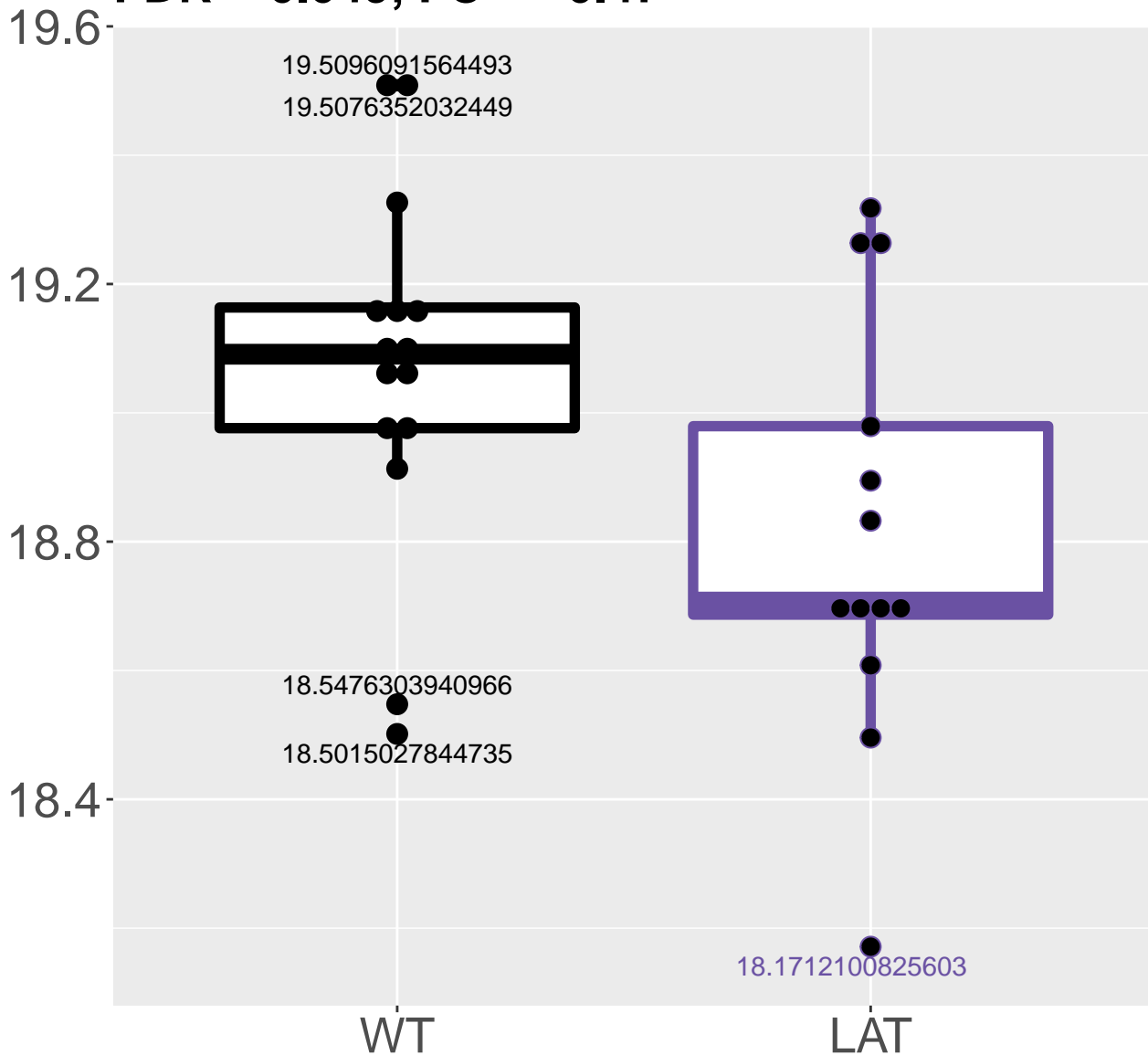
Q9D710_Thioredoxin-related tran.
FDR = 0.048, FC = 1.1



Q60597_2-oxoglutarate dehydroge.
FDR = 0.049, FC = 0.42



Q91W34_RUS1 family protein C16o.
FDR = 0.049, FC = -0.47



Q9Z2M7_Phosphomannomutase 2

FDR = 0.049, FC = -0.16, sex*

