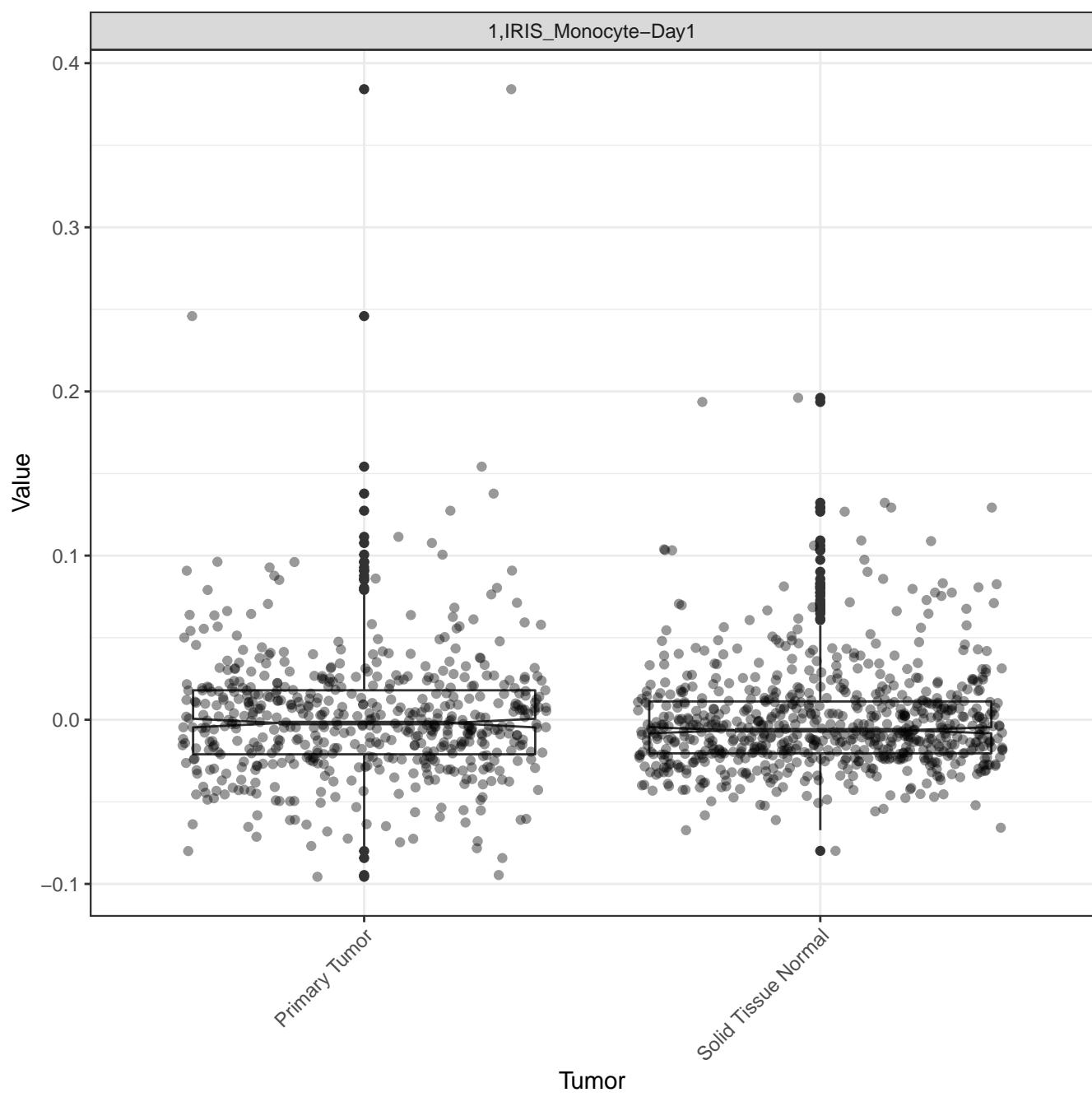
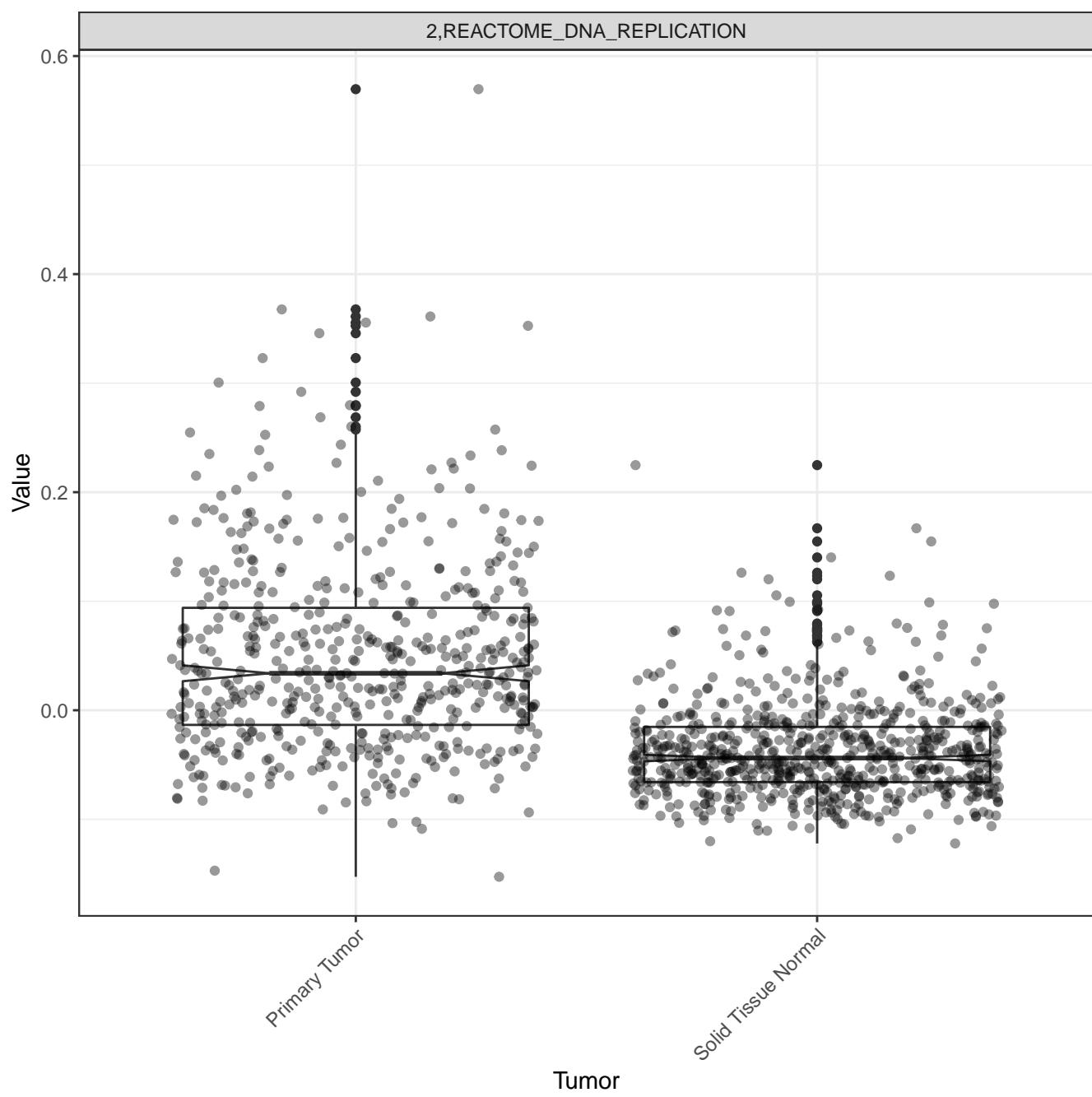


1,IRIS_Monocyte–Day1



2,REACTOME_DNA_REPLICATION



3,REACTOME_TCA_CYCLE_AND_RESPIRATORY_ELECTRON_TRANSPORT

Value

0.3

0.2

0.1

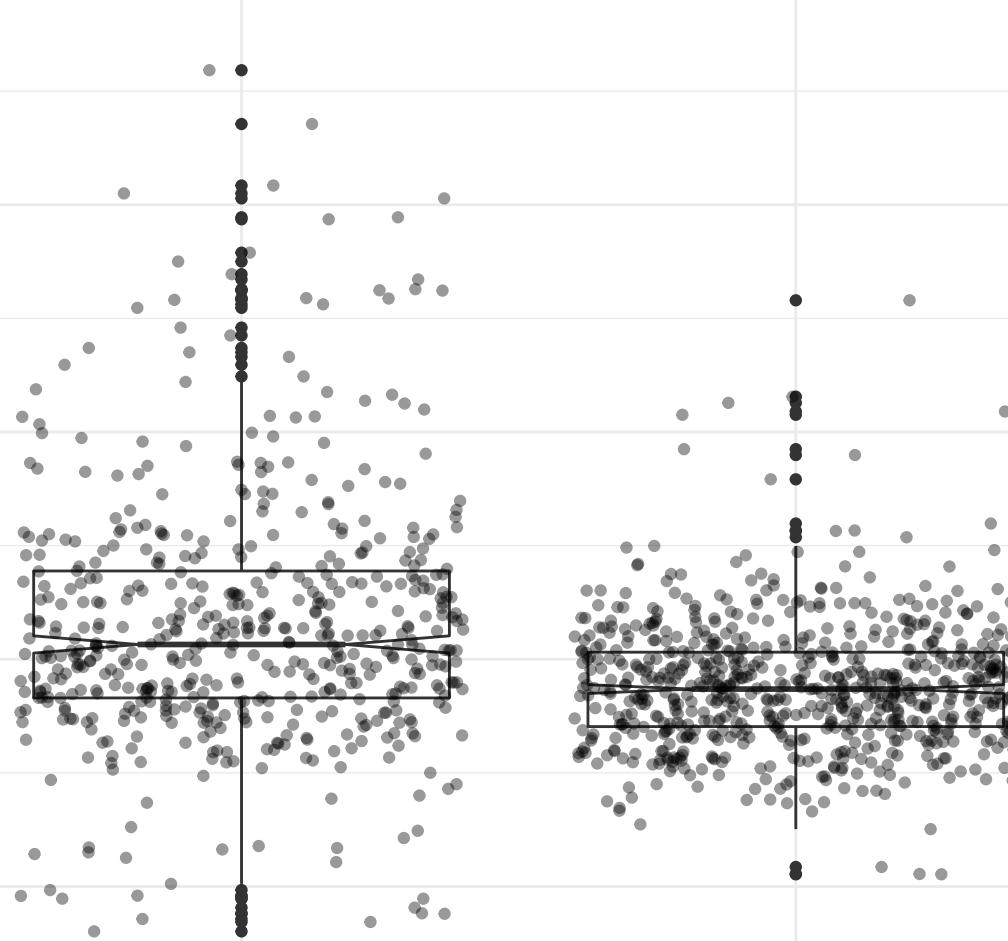
0.0

-0.1

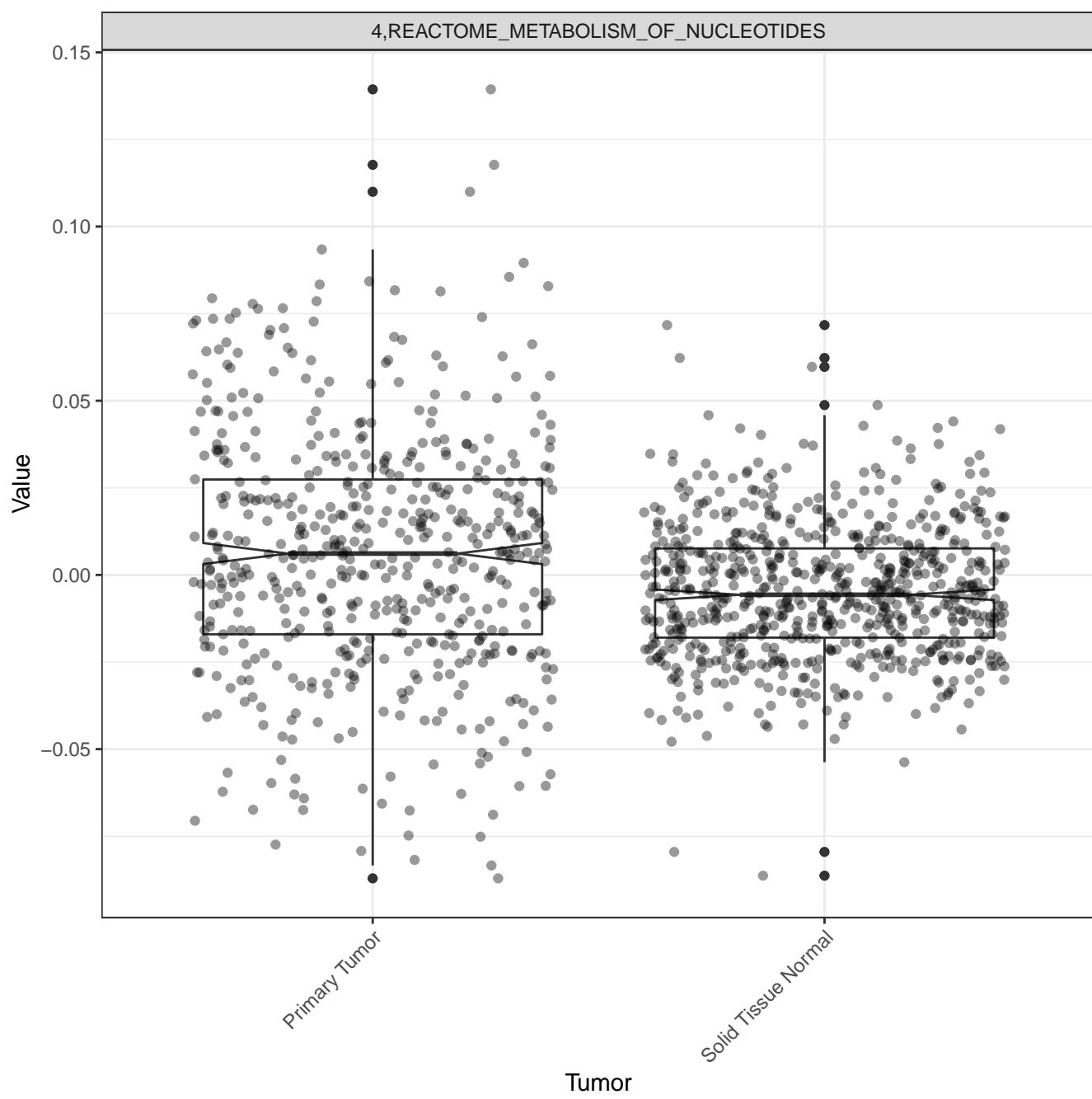
Primary Tumor

Solid Tissue Normal

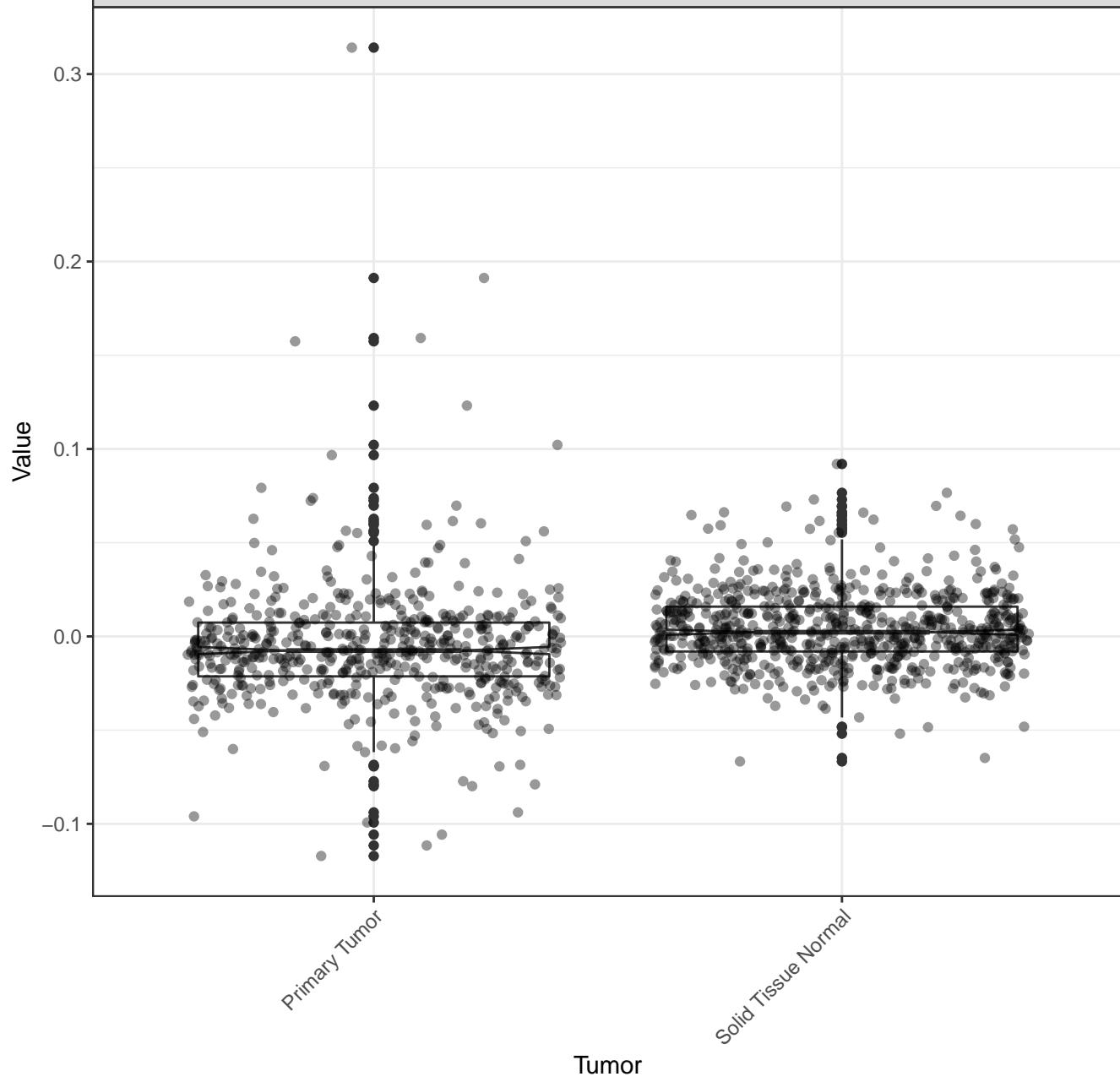
Tumor



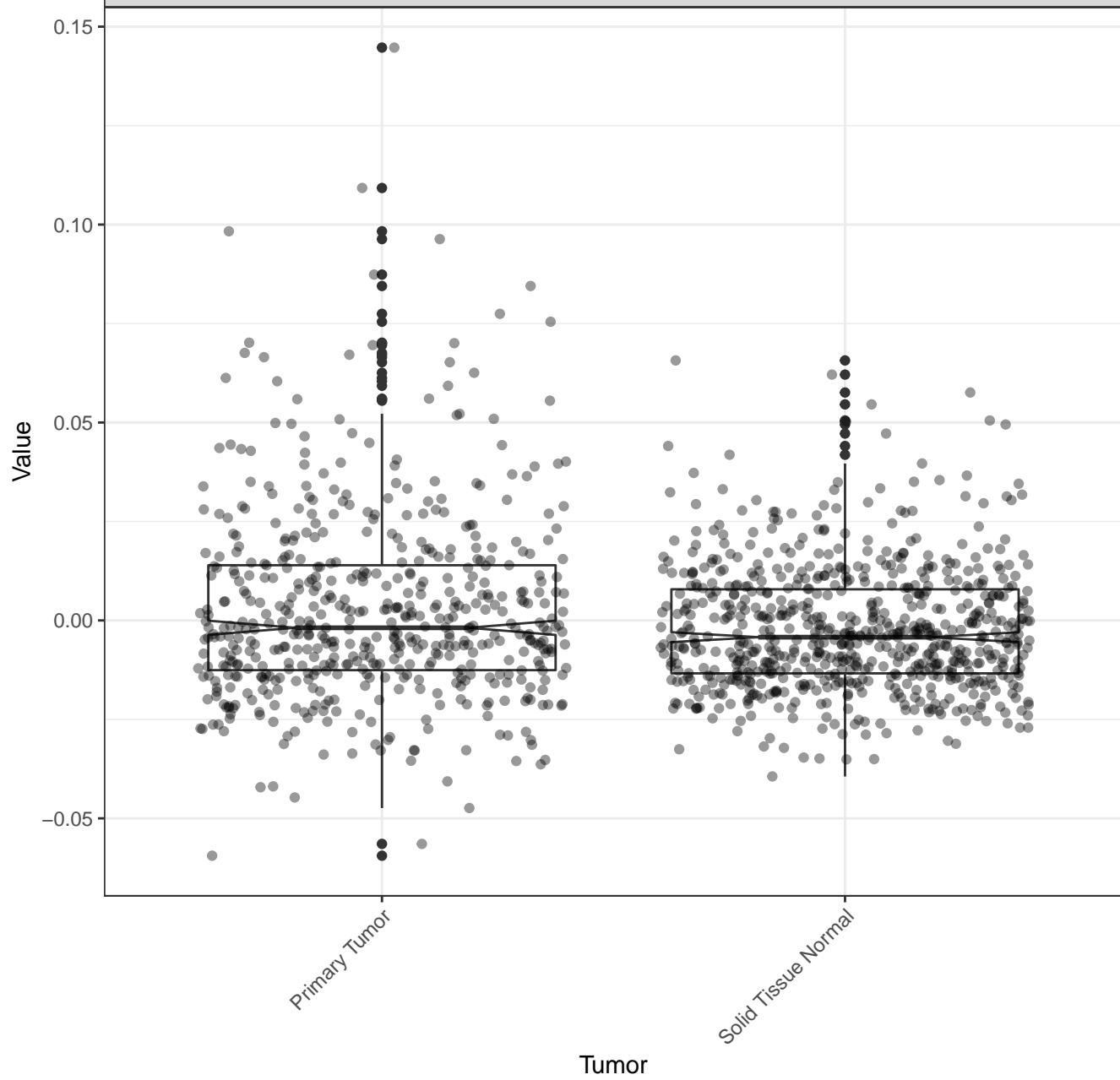
4,REACTOME_METABOLISM_OF_NUCLEOTIDES



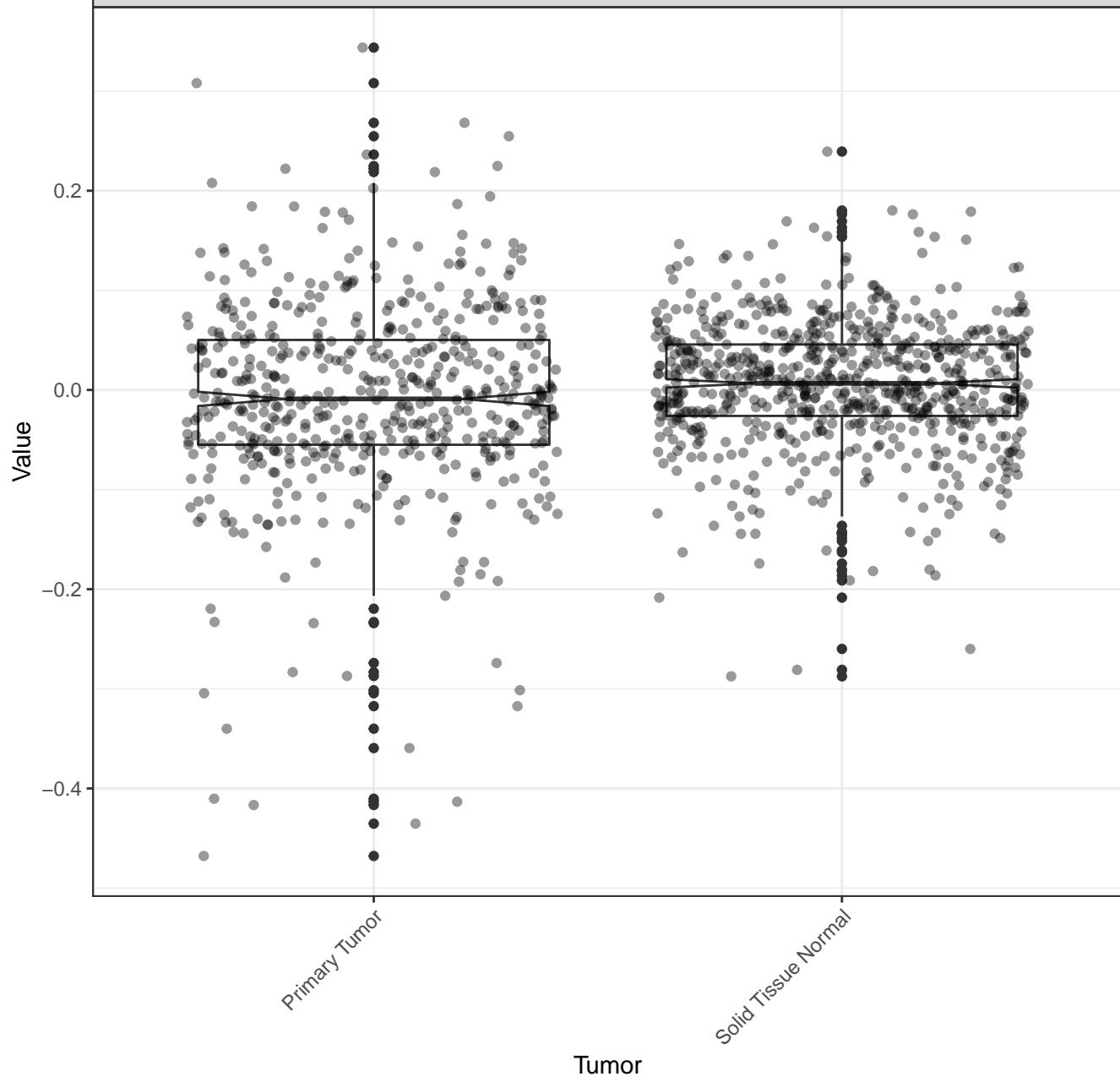
5,REACTOME_NEURONAL_SYSTEM



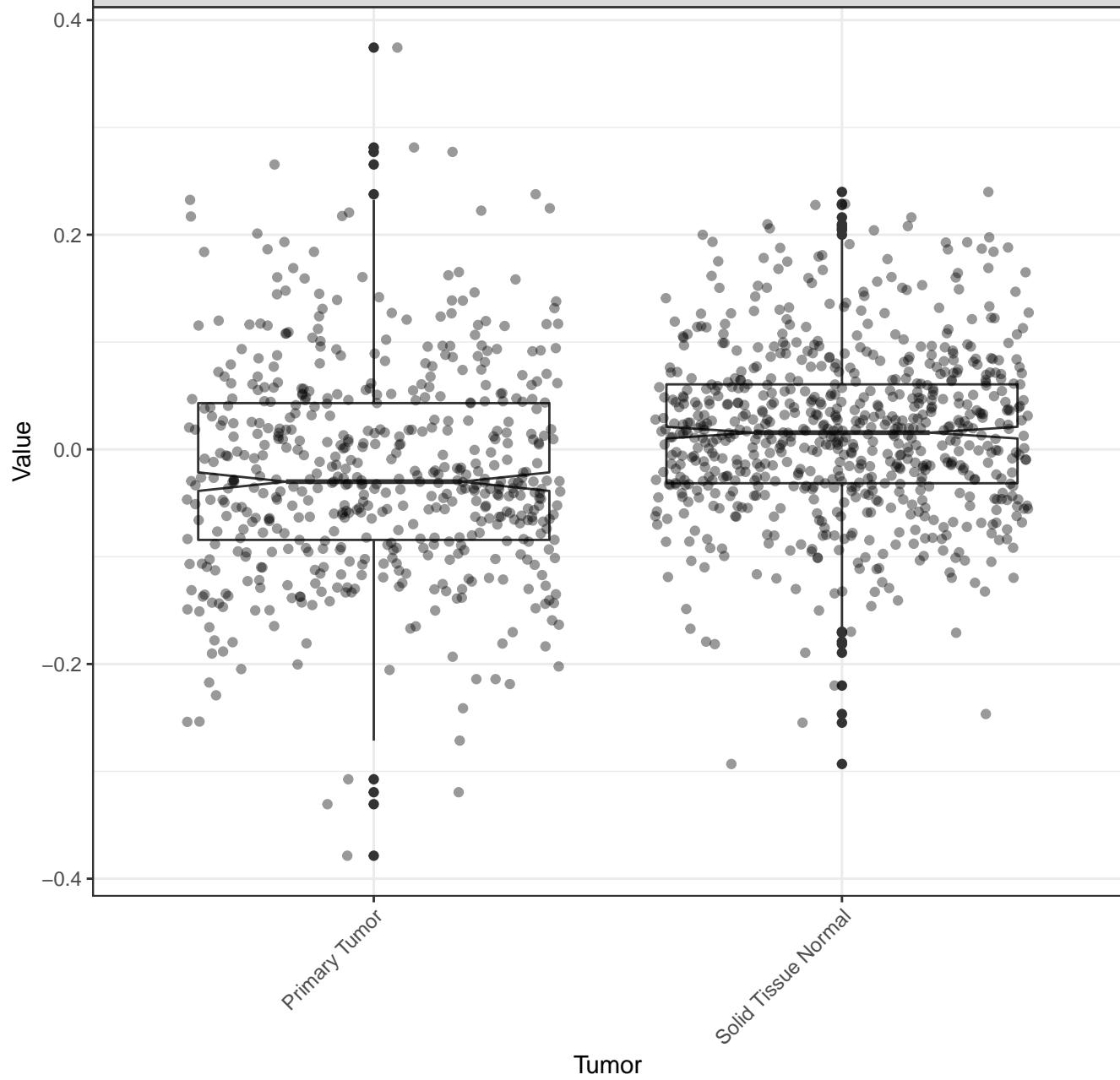
6,KEGG_HEDGEHOG_SIGNALING_PATHWAY



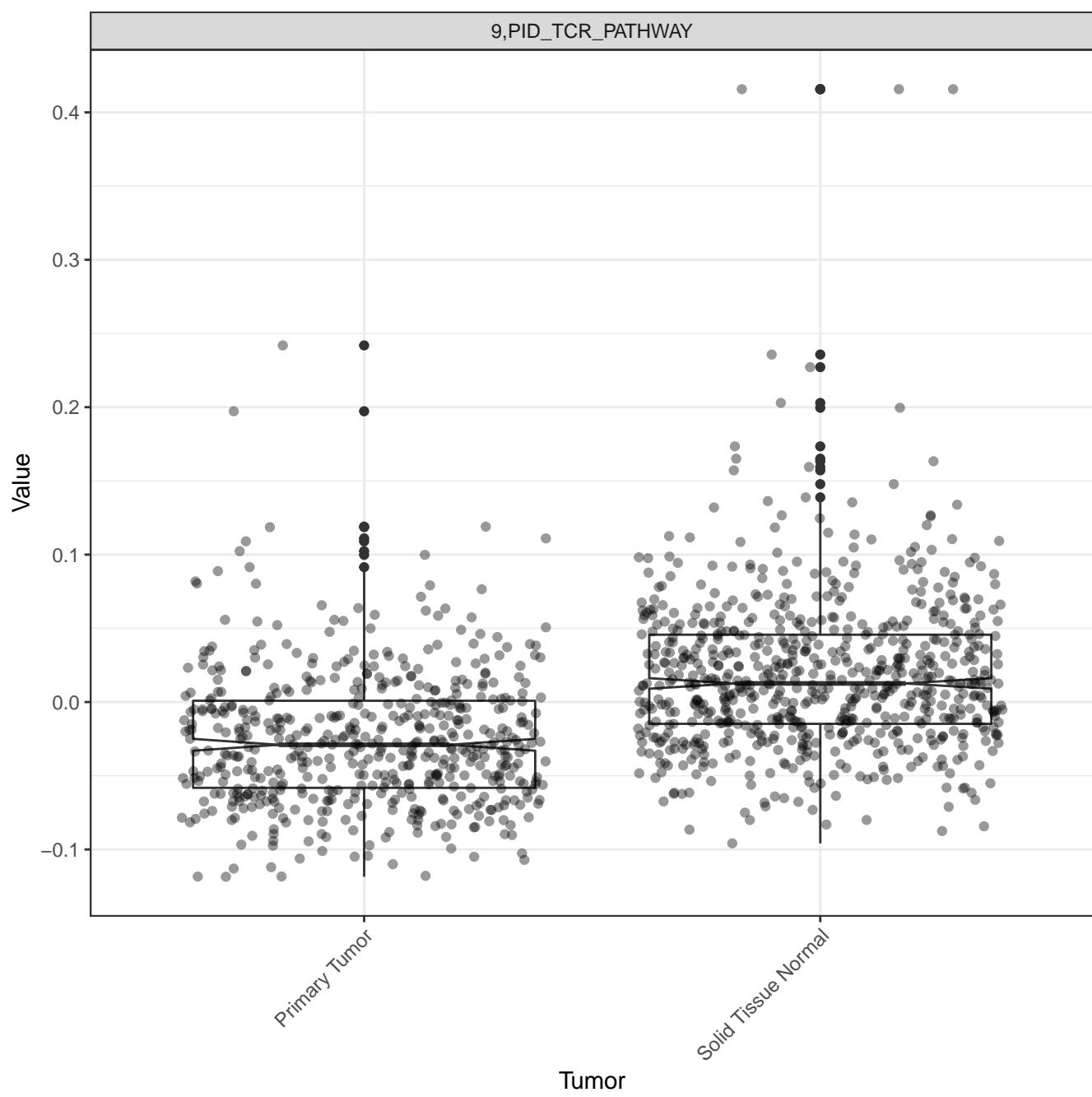
7,DMAP_ERY3



8,REACTOME_INFLUENZA_VIRAL_RNA_TRANSCRIPTION_AND_REPLICATION



9,PID_TCR_PATHWAY



10,SVM T cells CD4 memory resting

Value

0.2

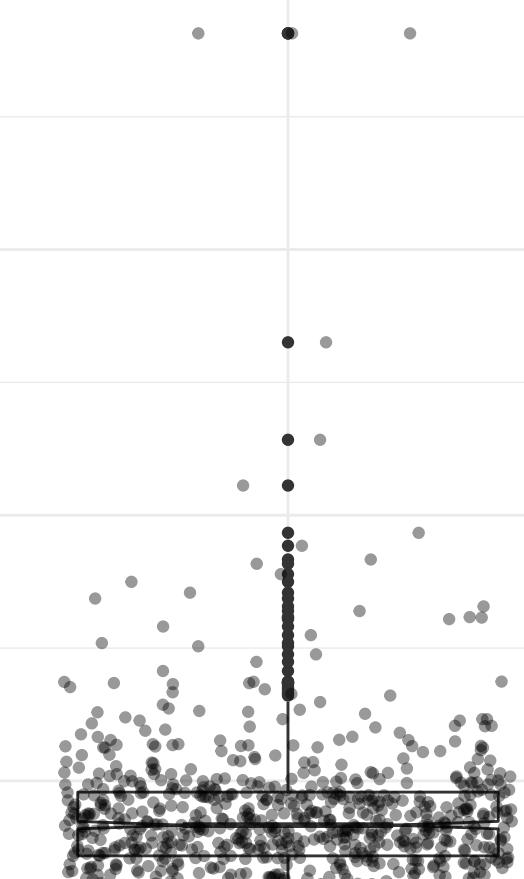
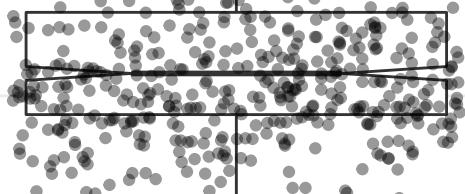
0.1

0.0

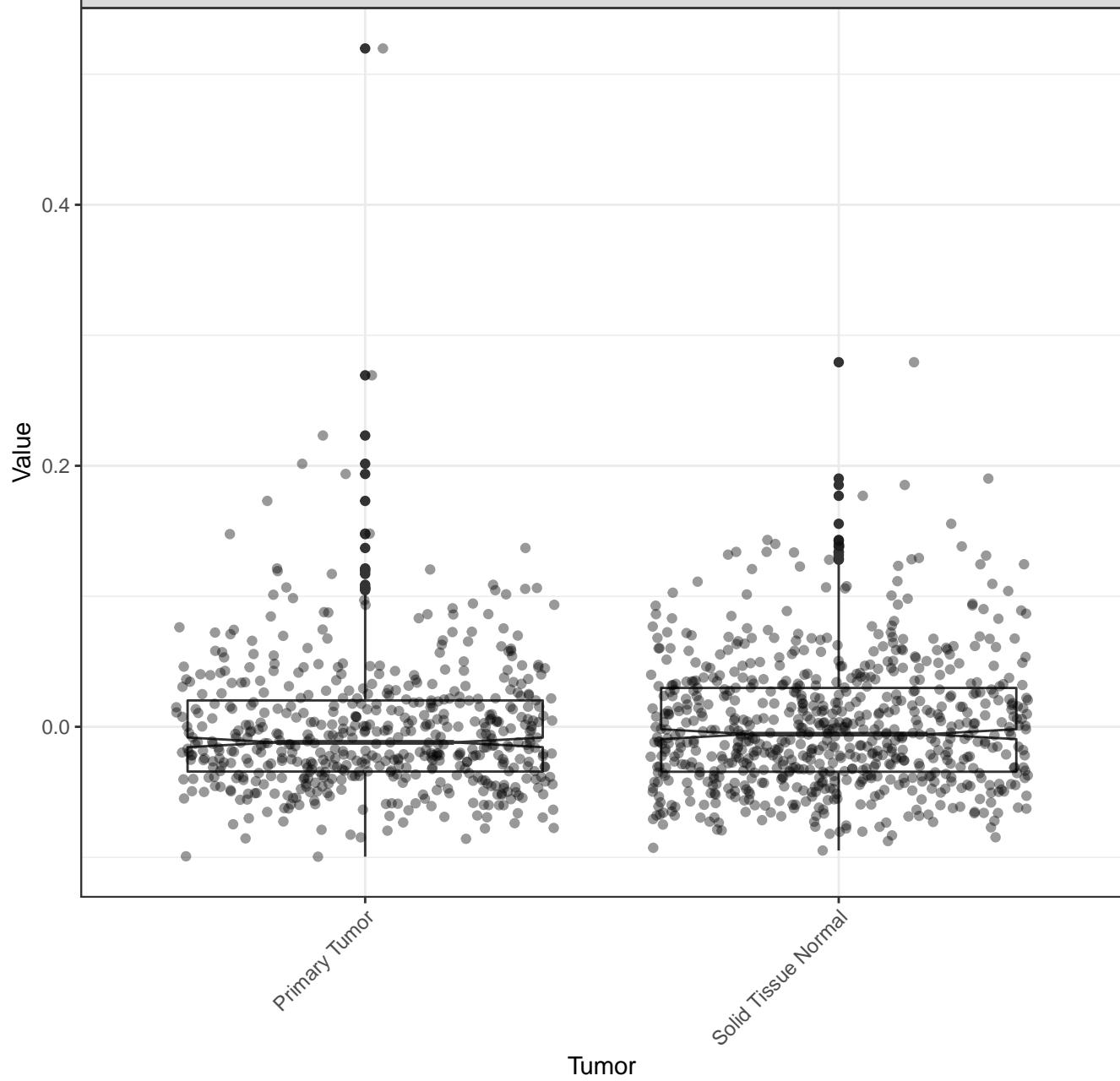
Primary Tumor

Solid Tissue Normal

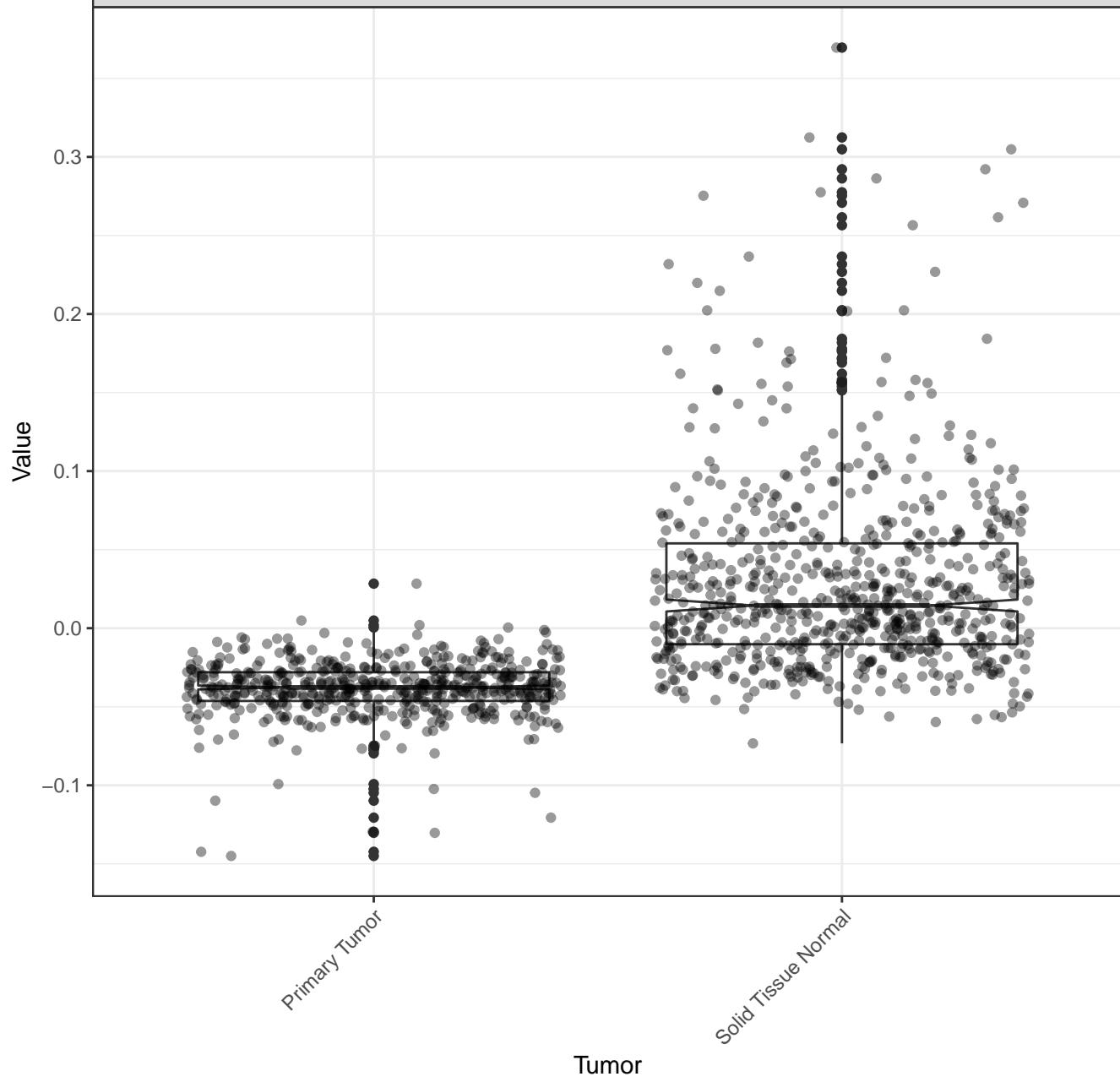
Tumor



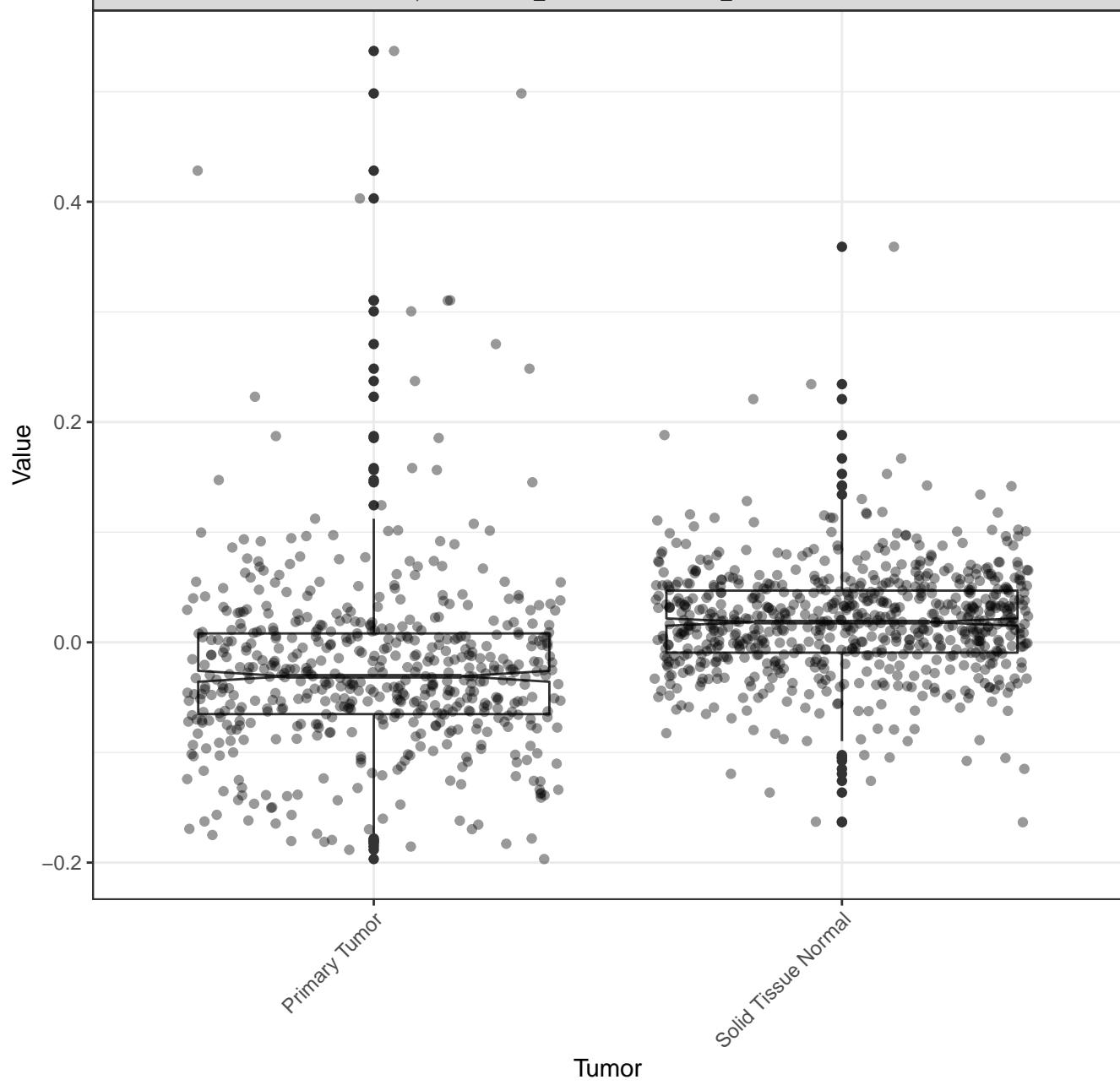
11,IRIS_DendriticCell-Control



12,SVM Neutrophils



13,REACTOME DEVELOPMENTAL BIOLOGY



14,KEGG_BASAL_CELL_CARCINOMA

Value

0.10

0.05

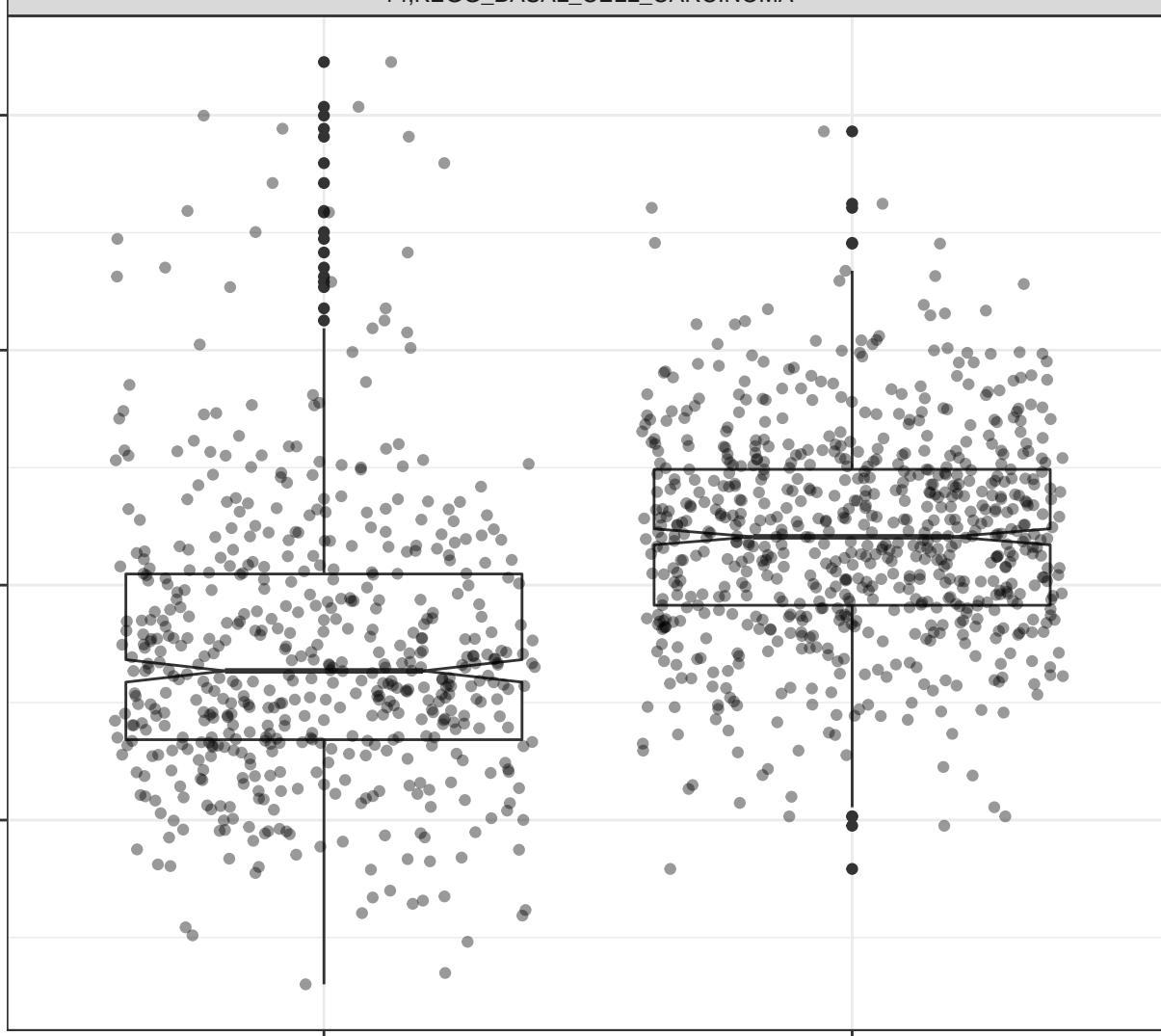
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



15,REACTOME_METABOLISM_OF_AMINO_ACIDS_AND_DERIVATIVES

Value

0.6

0.4

0.2

0.0

Primary Tumor

Solid Tissue Normal

Tumor

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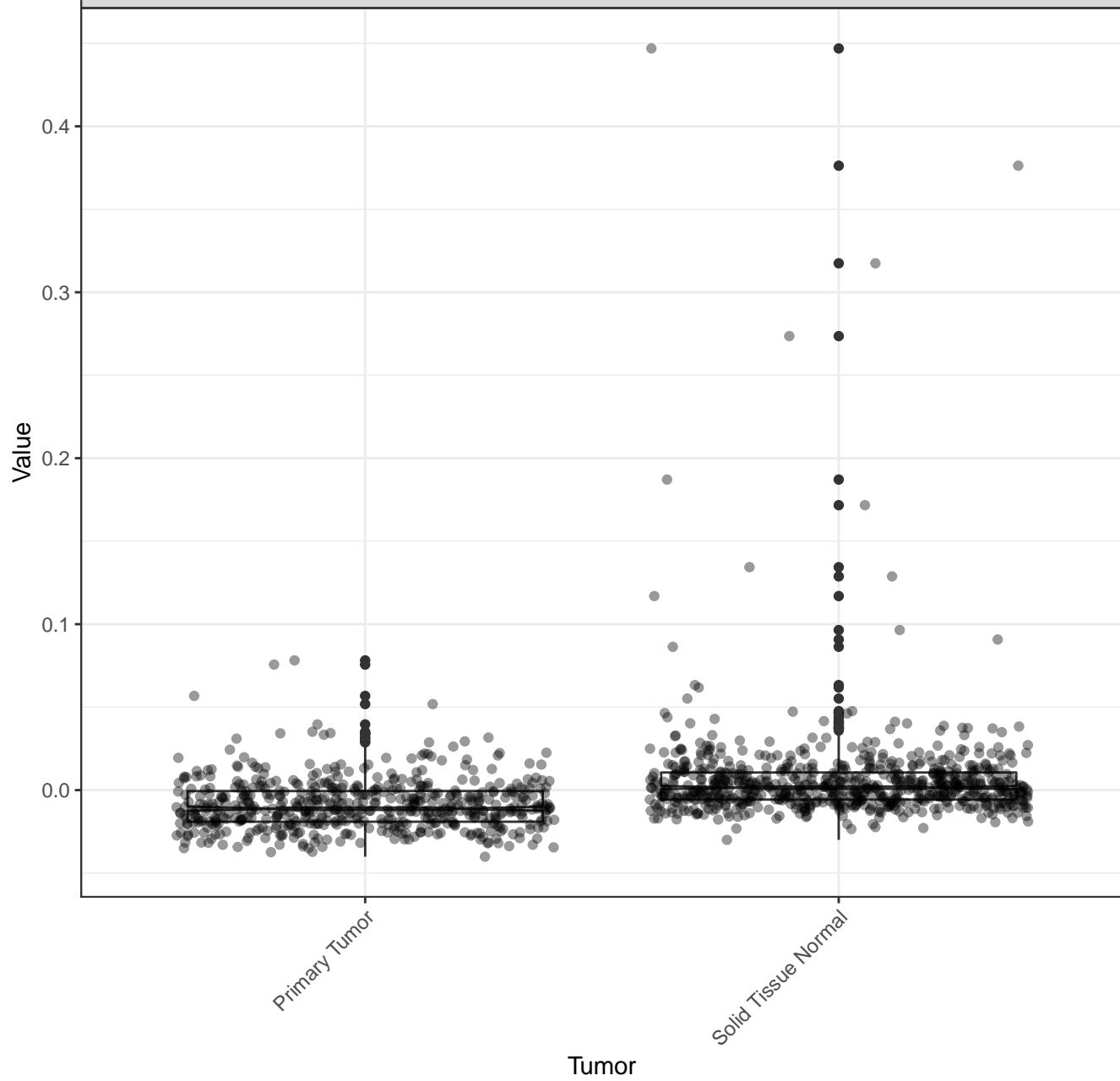
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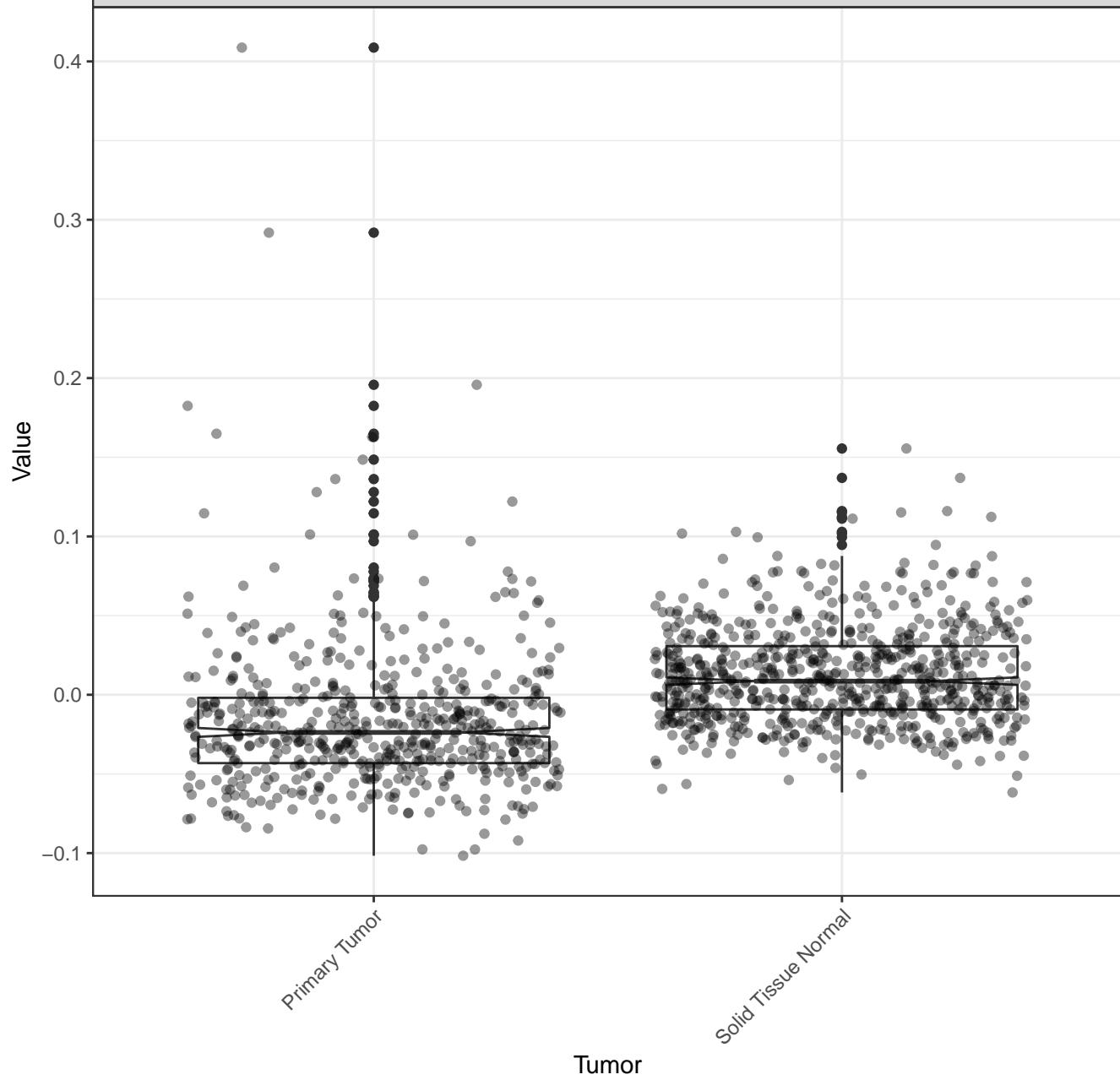
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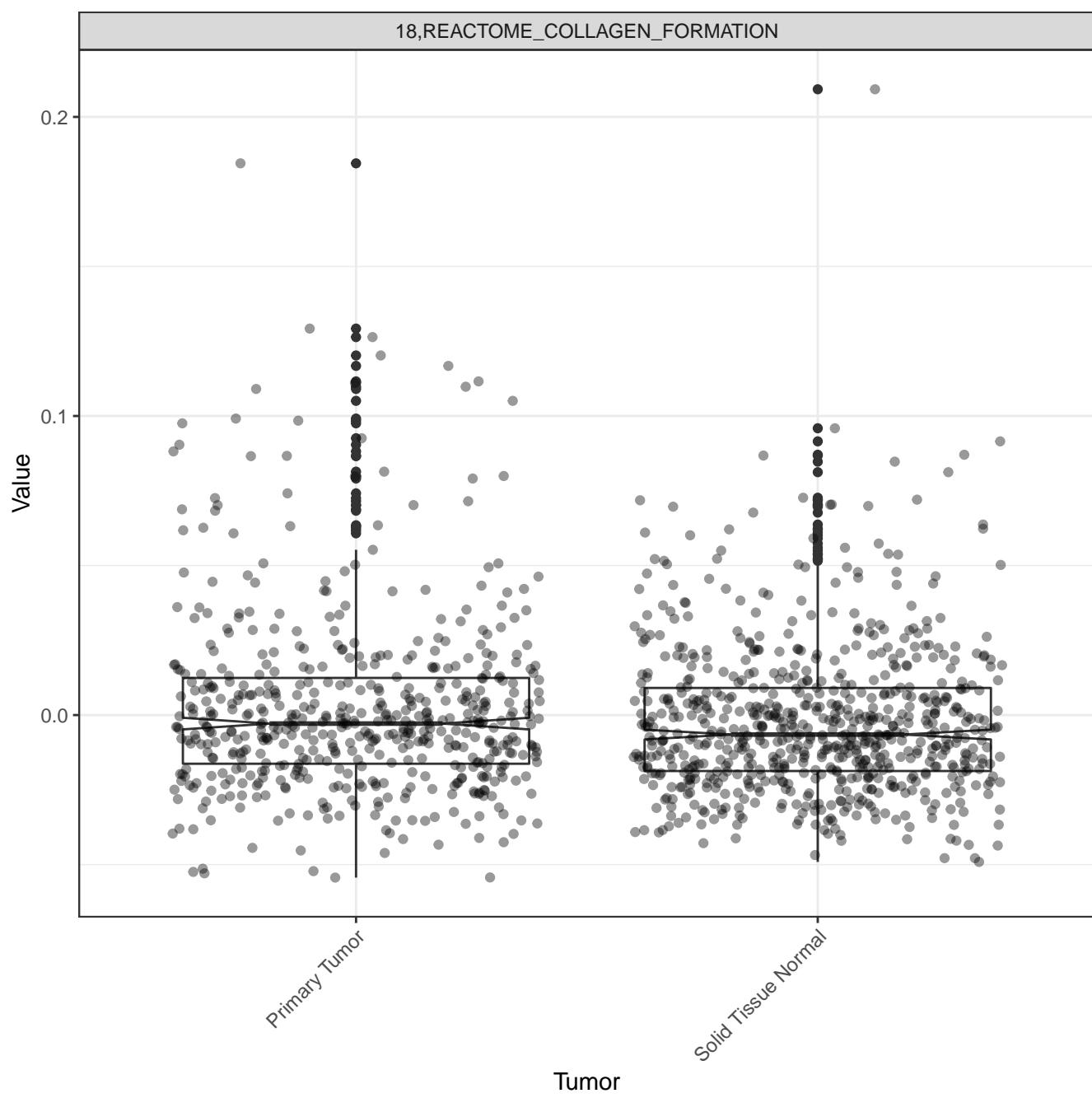
16,KEGG_CALCIUM_SIGNALING_PATHWAY



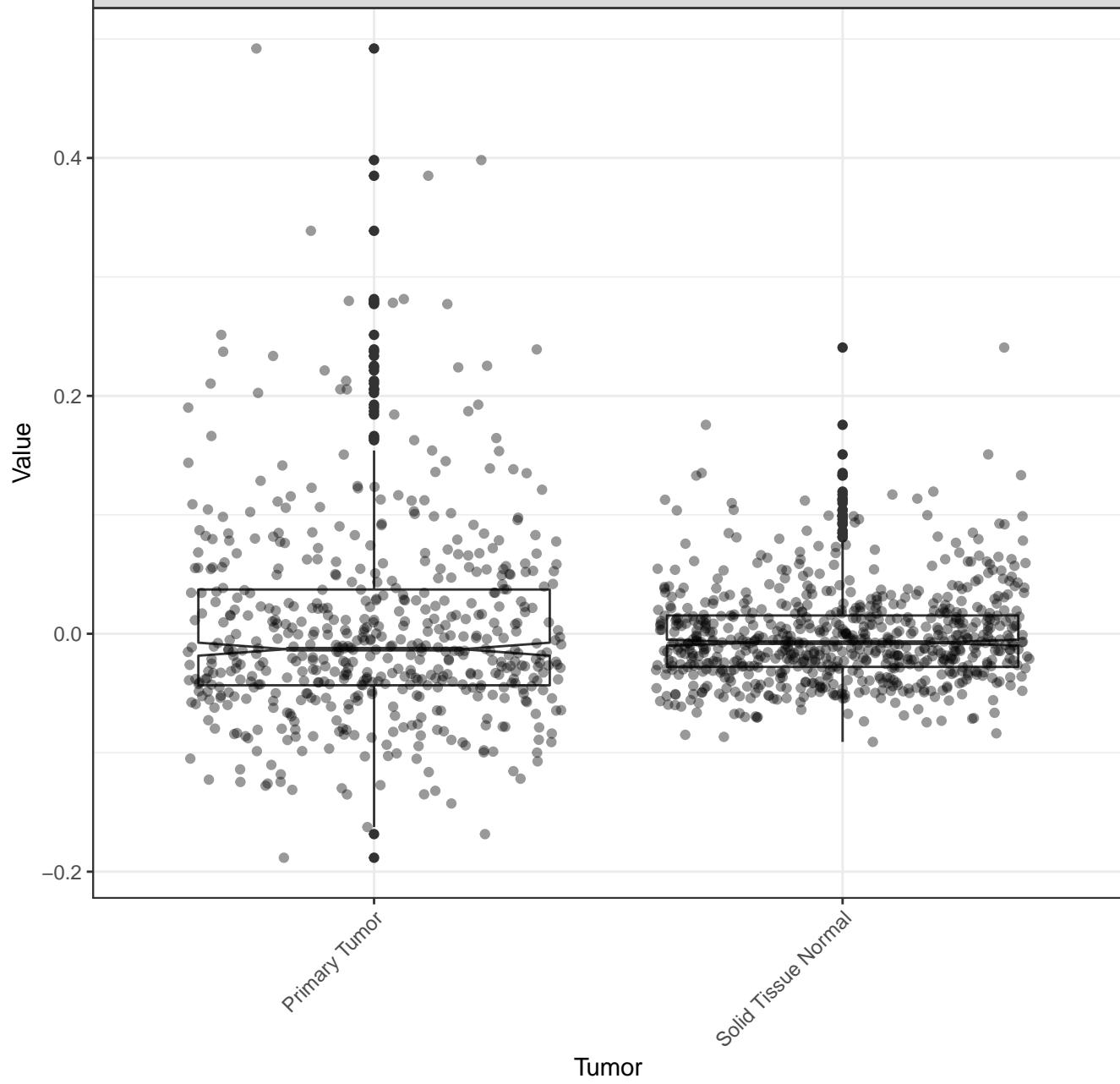
17,BIOCARTA_CHREBP2_PATHWAY



18,REACTOME_COLLAGEN_FORMATION



19,REACTOME_PEPTIDE_CHAIN_ELONGATION



20,KEGG_INTESTINAL_IMMUNE_NETWORK_FOR_IGA_PRODUCTION

Value

0.05

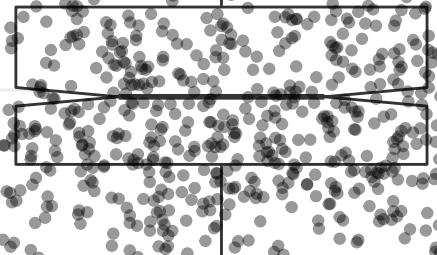
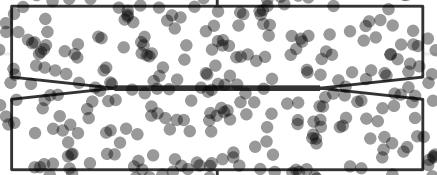
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



21,SVM Macrophages M1

Value

0.3

0.2

0.1

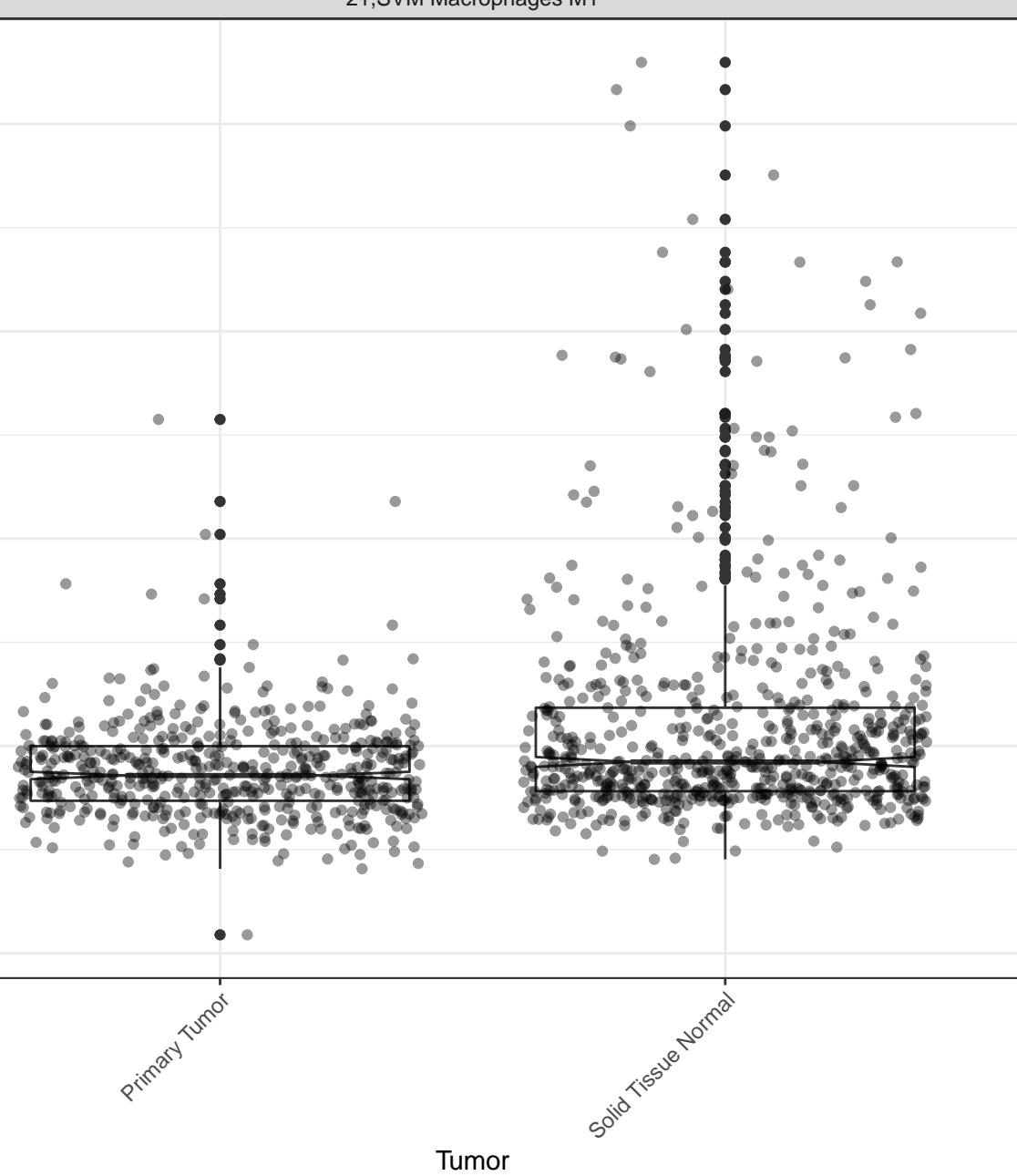
0.0

-0.1

Primary Tumor

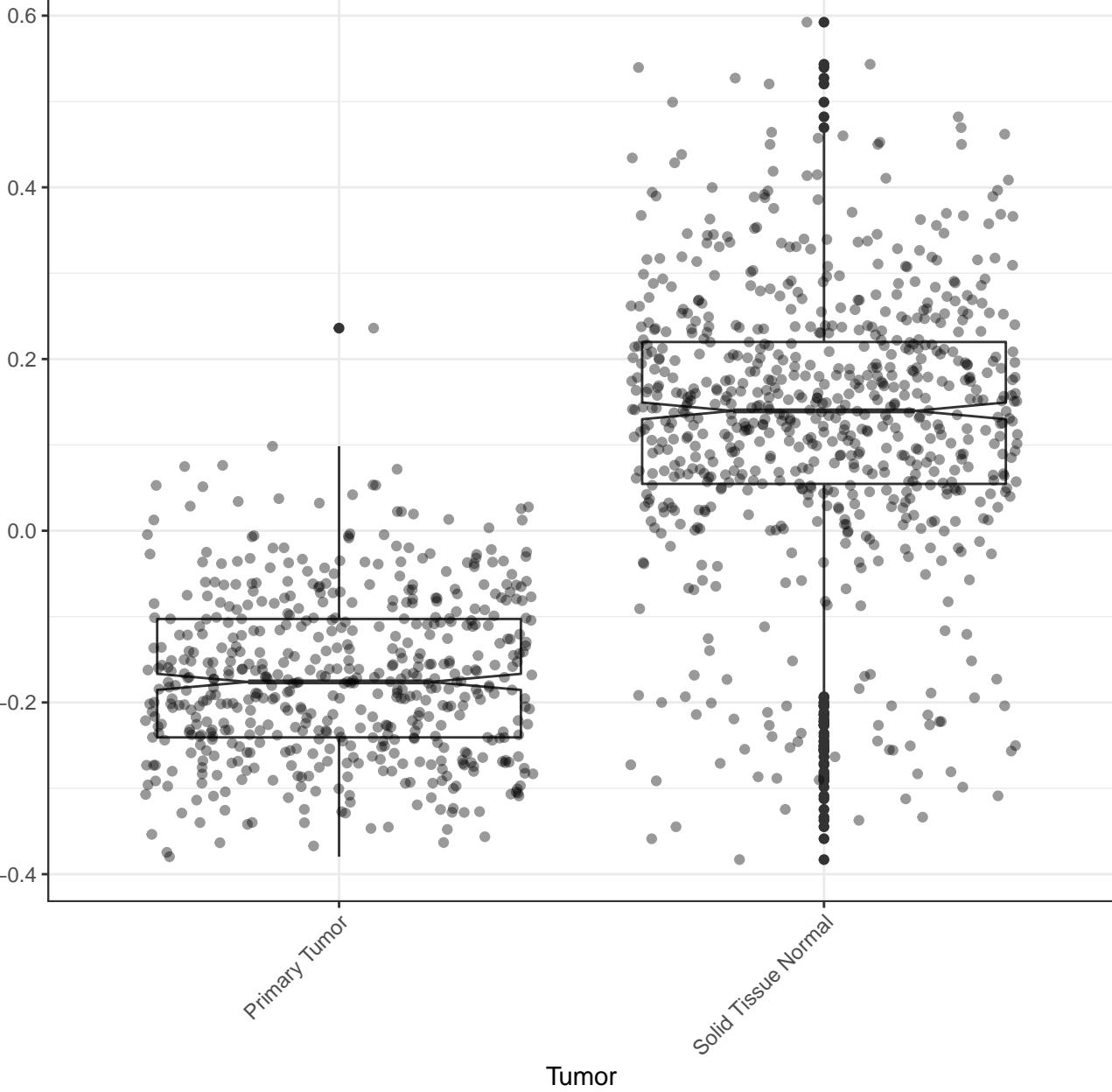
Solid Tissue Normal

Tumor

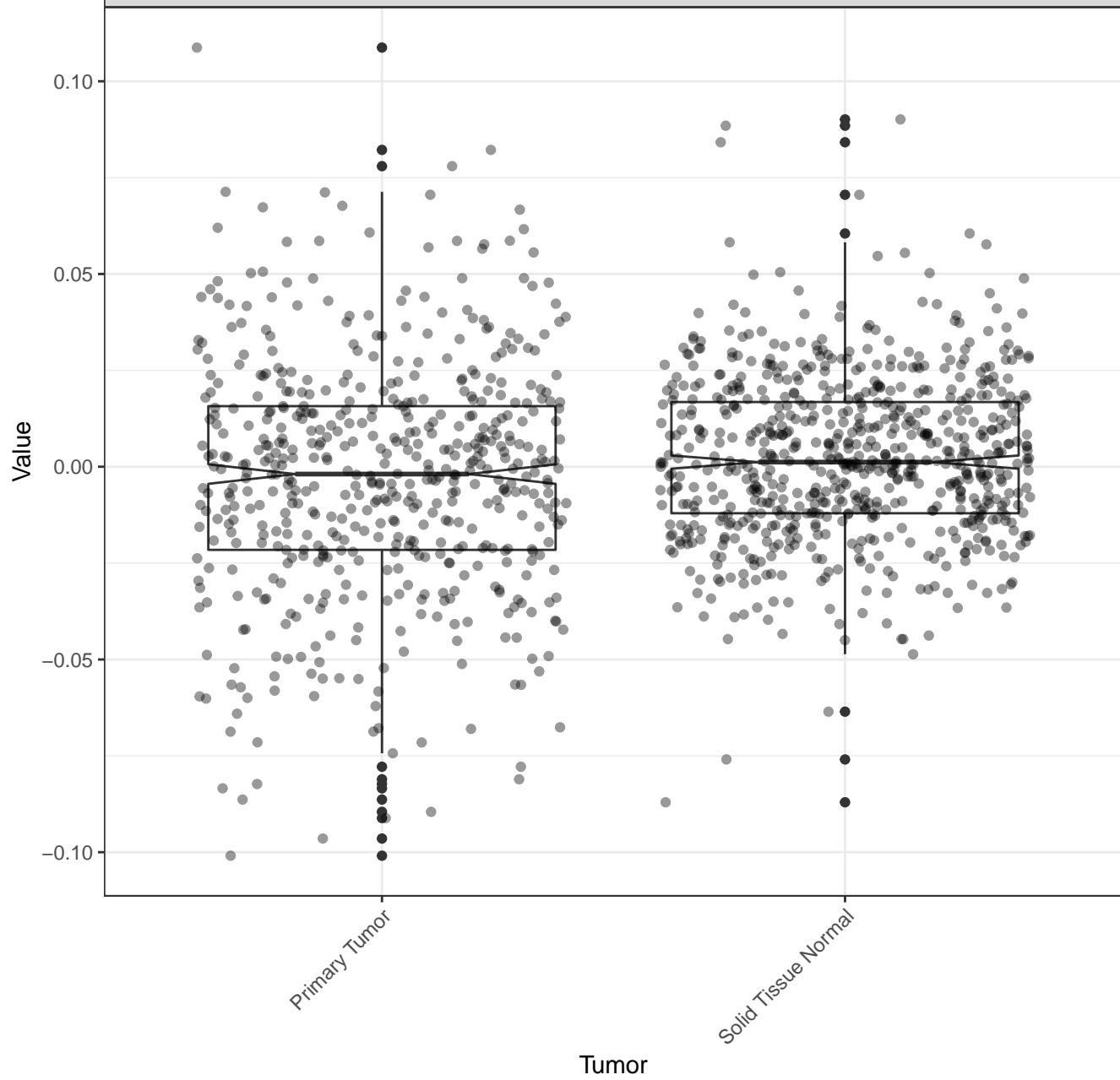


22,MIPS_SPLICEOSOME

Value



23,DMAP_ERY4



24,REACTOME_PLATELET_HOMEOSTASIS

Value

0.10

0.05

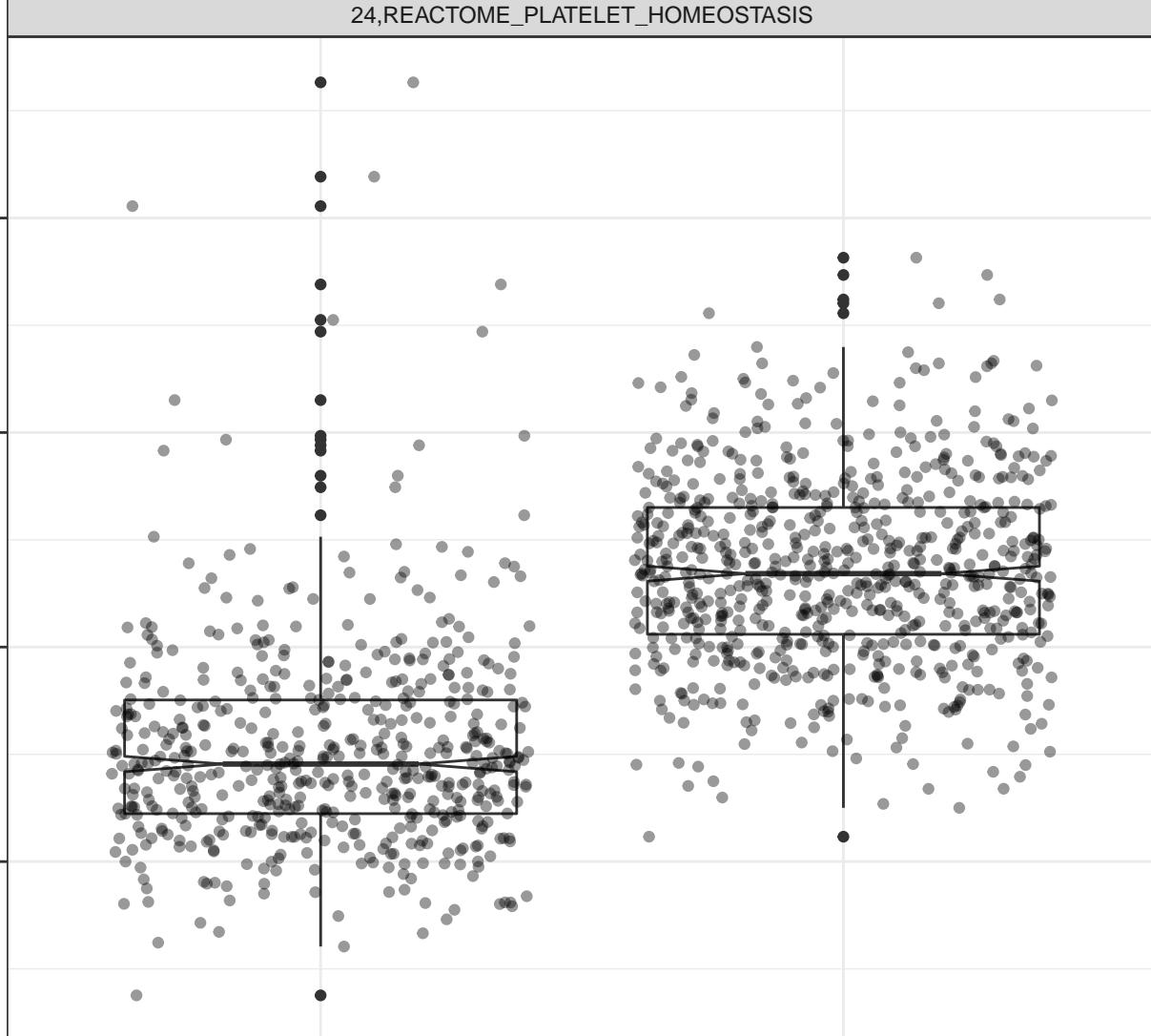
0.00

-0.05

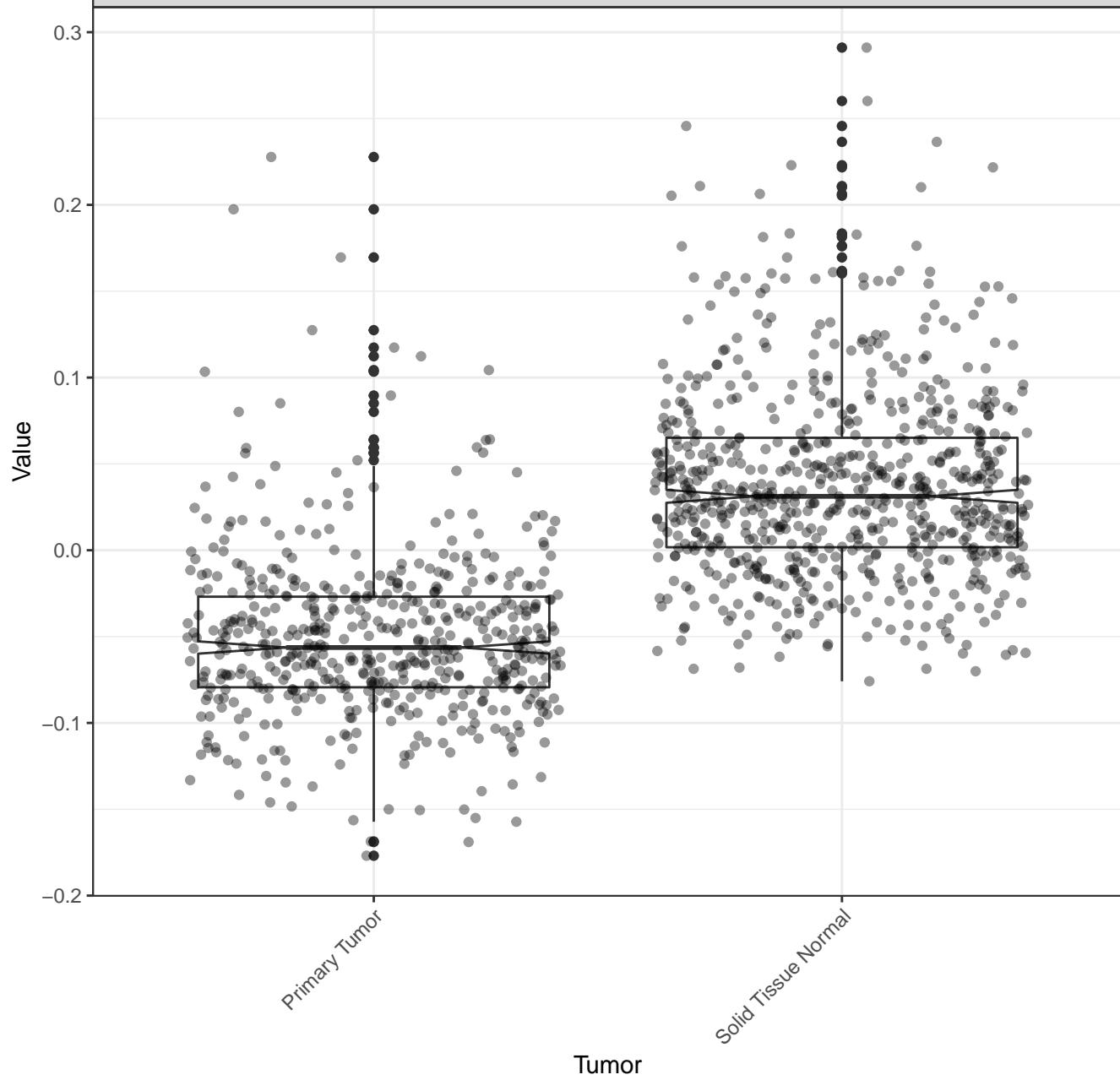
Primary Tumor

Solid Tissue Normal

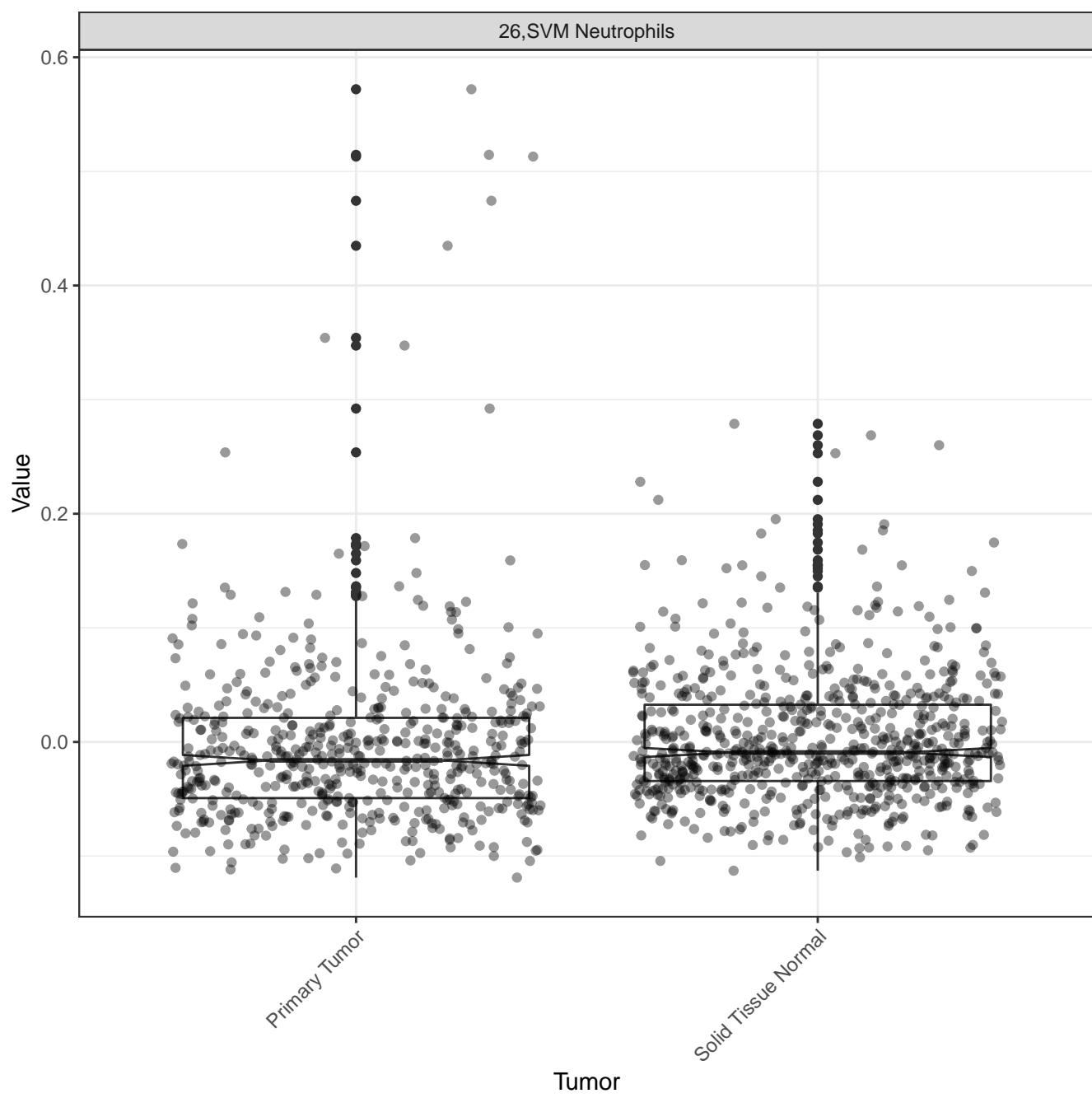
Tumor



25,PID_INTEGRIN1_PATHWAY



26,SVM Neutrophils



Value

0.10

0.05

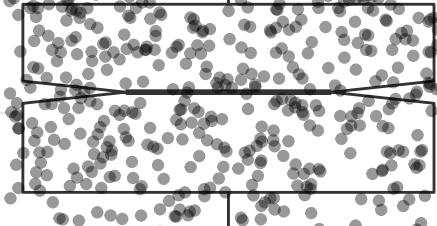
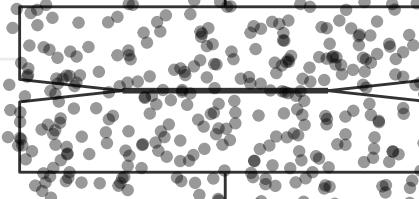
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



28,REACTOME_INTERFERON_ALPHA_BETA_SIGNALING

Value

0.4

0.3

0.2

0.1

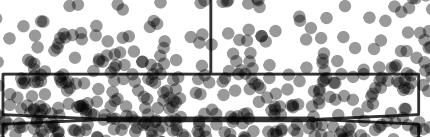
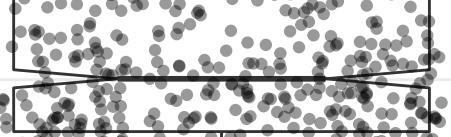
0.0

-0.1

Primary Tumor

Solid Tissue Normal

Tumor



29,REACTOME_E2F_MEDIATED_REGULATION_OF_DNA_REPLICATION

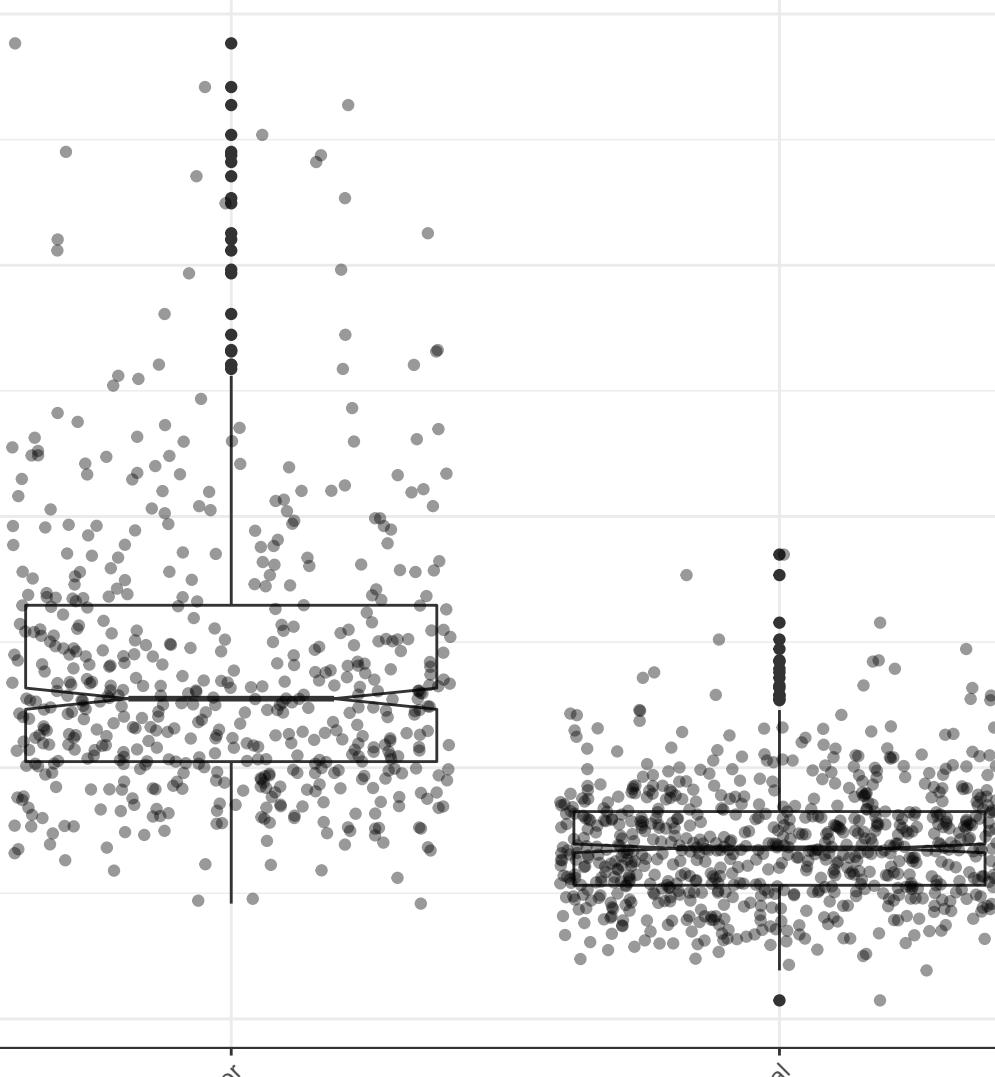
Value

0.15
0.10
0.05
0.00
-0.05

Primary Tumor

Solid Tissue Normal

Tumor



30,REACTOME_TRANSPORT_OF_INORGANIC_CATIONS_ANIONS_AND_AMINO_ACIDS_OLIGOPEPTIDES

Value

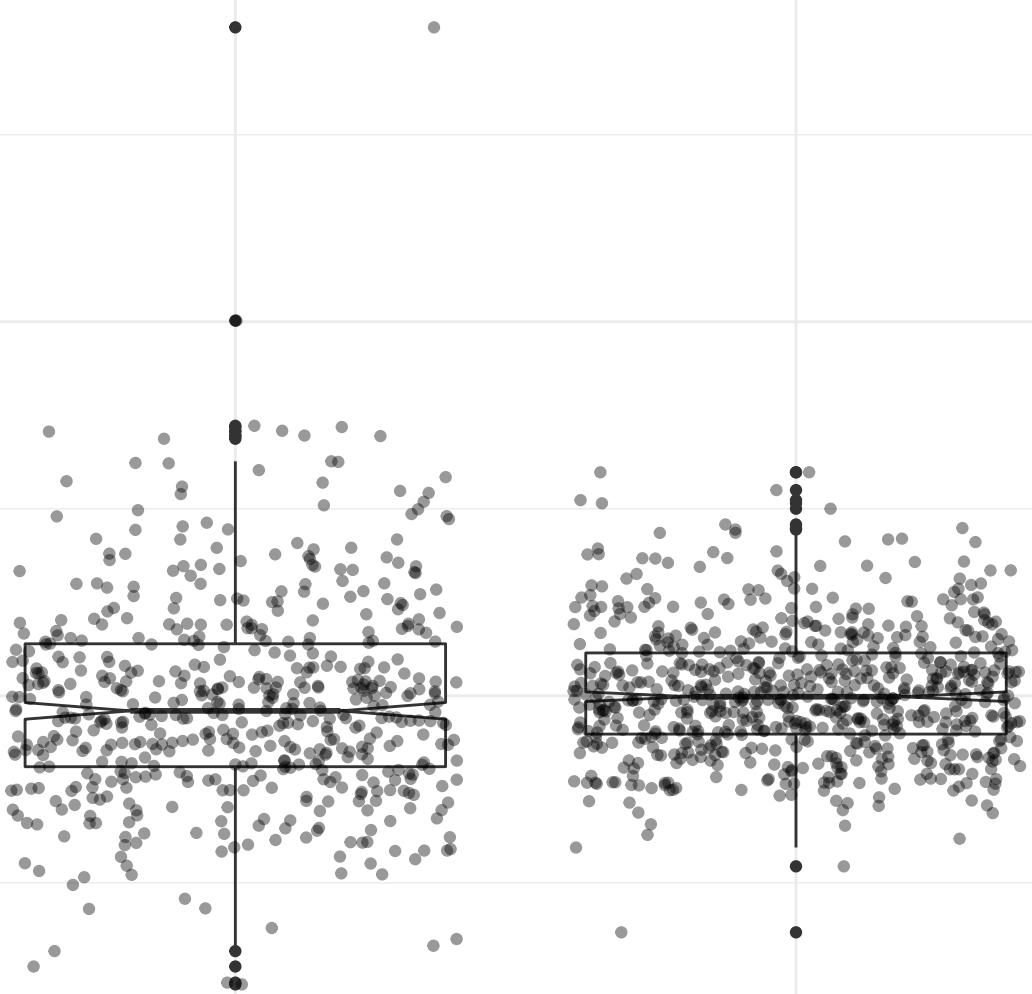
0.1

0.0

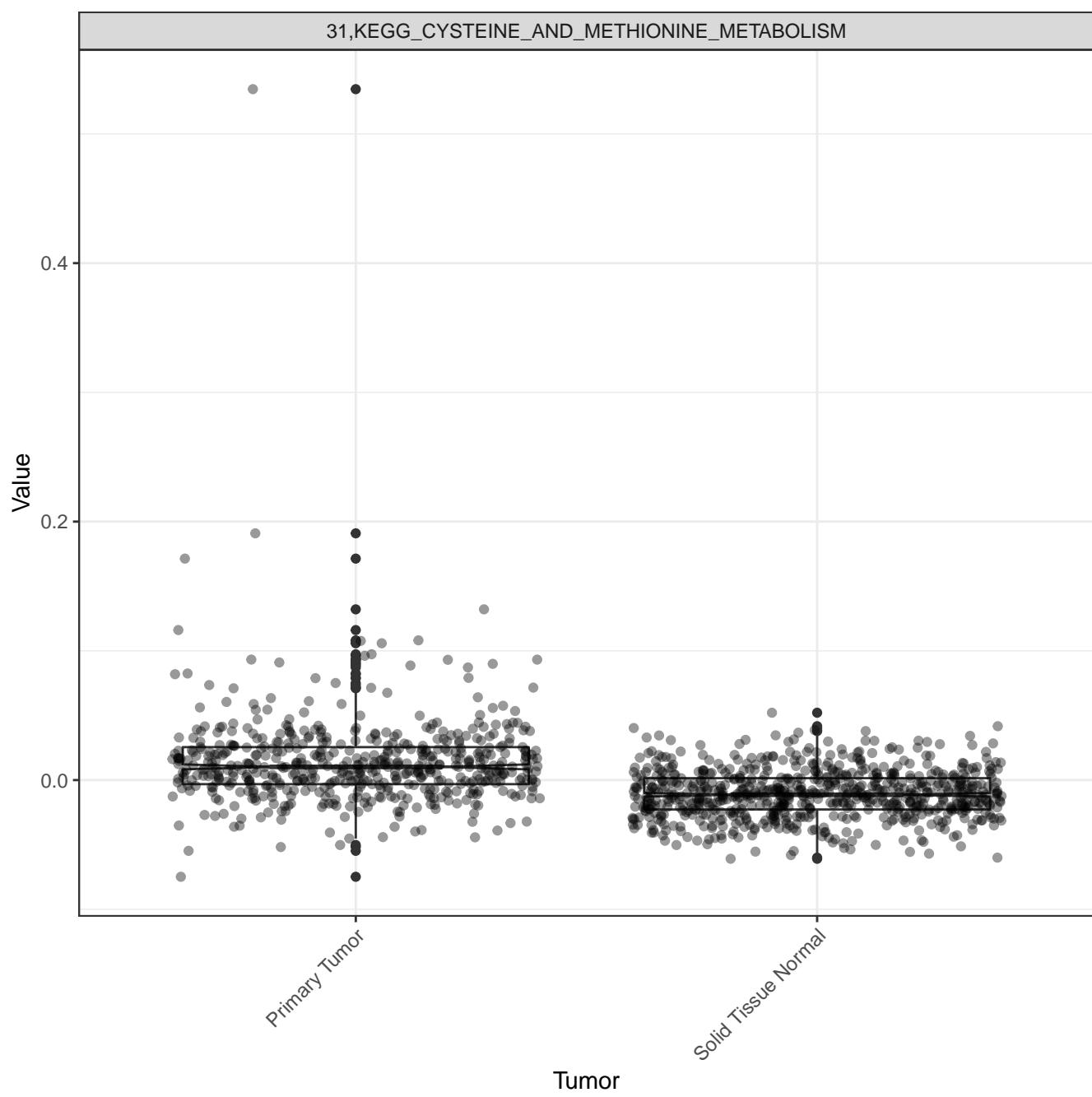
Primary Tumor

Tumor

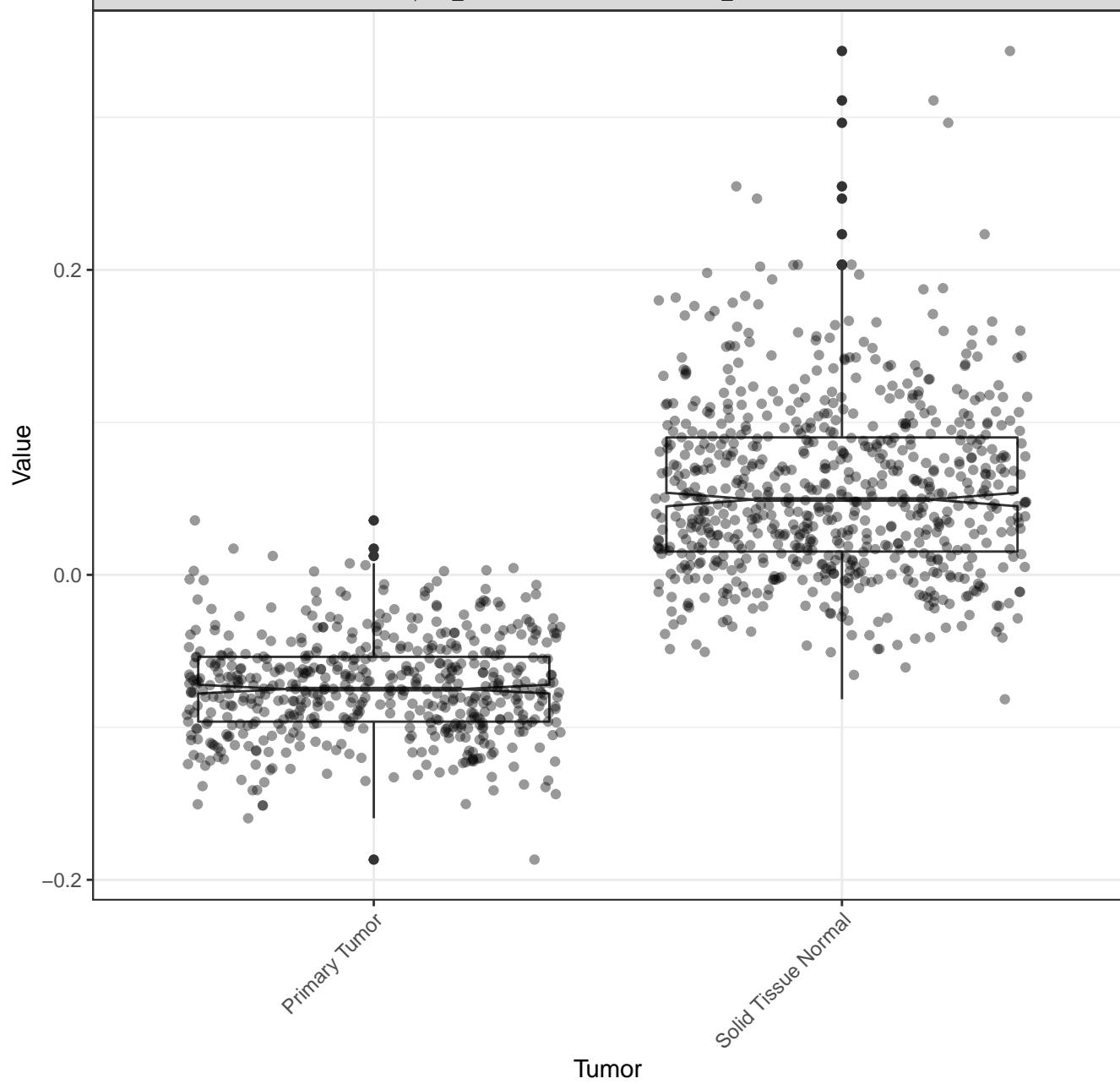
Solid Tissue Normal

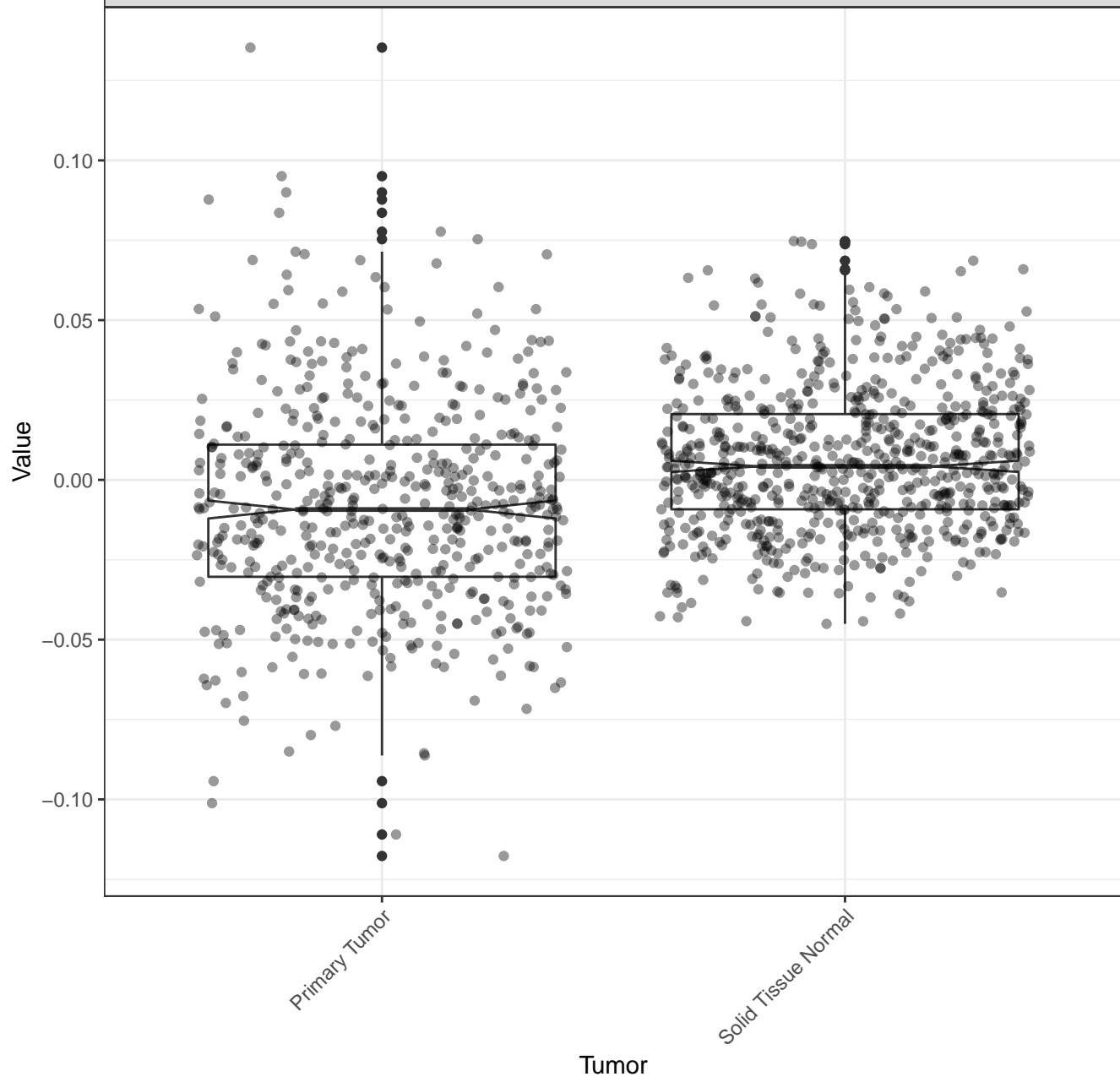


31,KEGG_CYSTEINE_AND_METHIONINE_METABOLISM

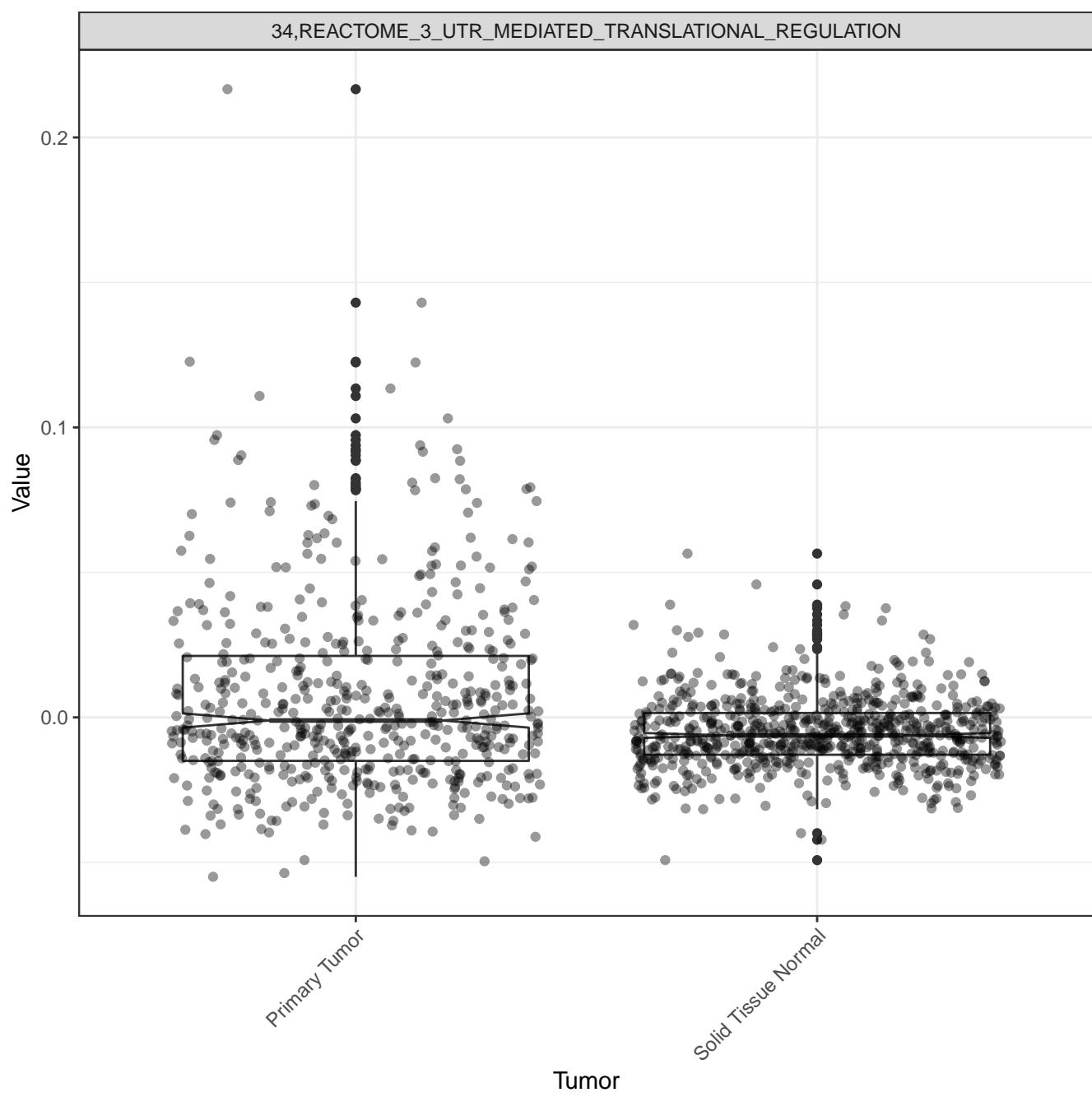


32,PID_ANGIOPOIETINRECEPTOR_PATHWAY





34,REACTOME_3_UTR_MEDIATED_TRANSLATIONAL_REGULATION



35,DMAP_MEGA2

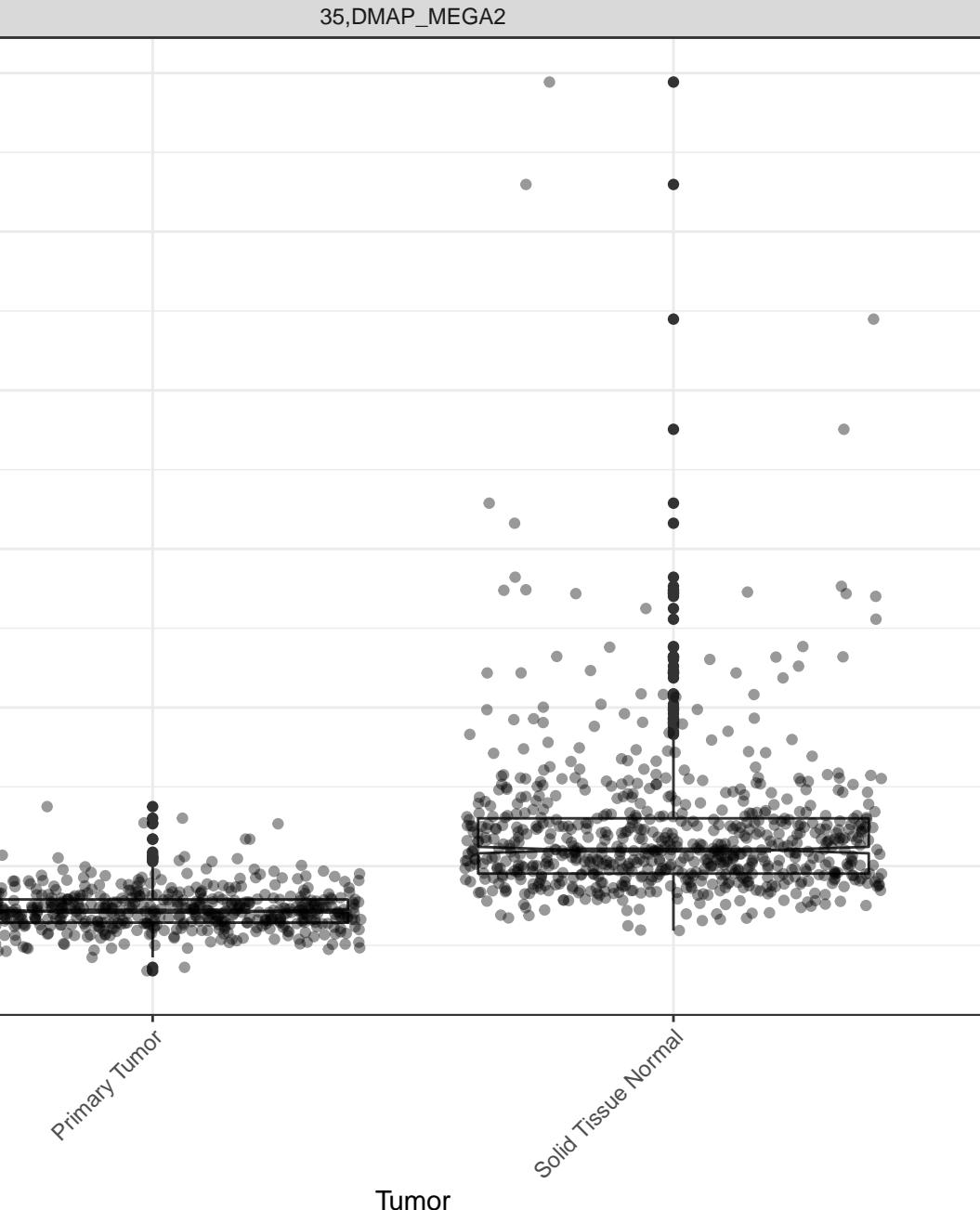
Value

0.5
0.4
0.3
0.2
0.1
0.0

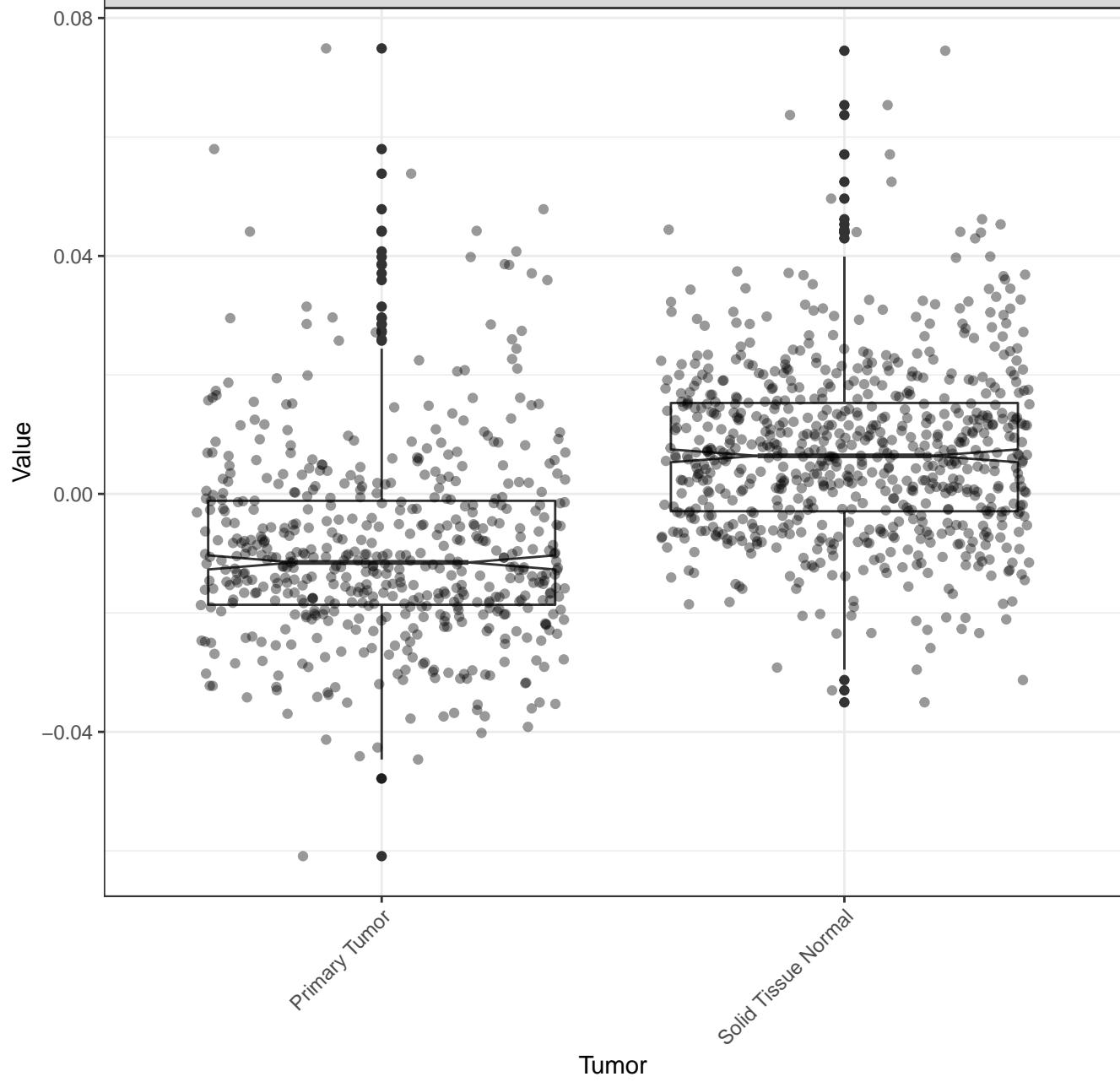
Primary Tumor

Solid Tissue Normal

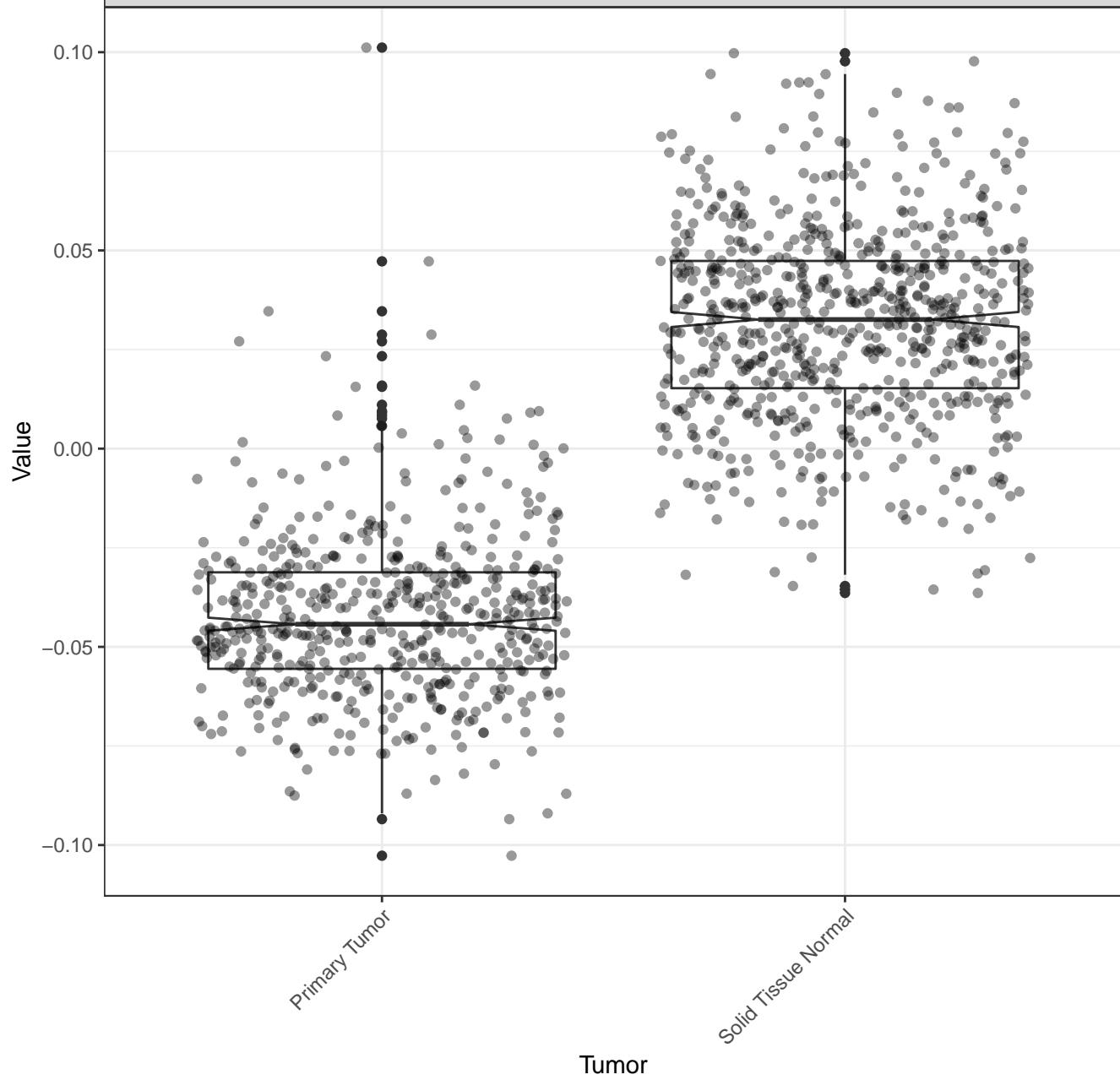
Tumor



36,DMAP_HSC3



37,KEGG_PPAR_SIGNALING_PATHWAY



38,KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT

Value

0.10

0.05

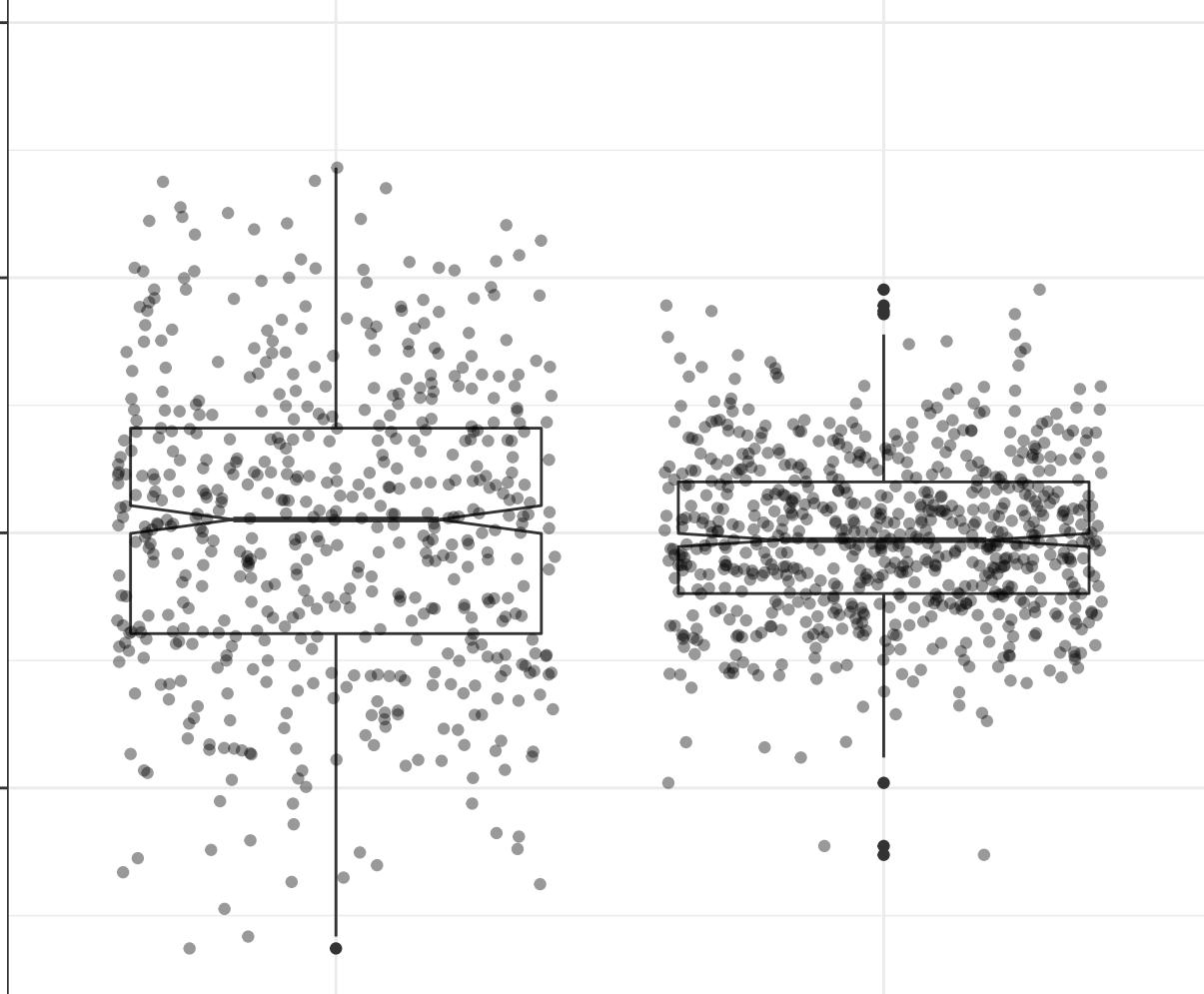
0.00

-0.05

Primary Tumor

Tumor

Solid Tissue Normal



39,KEGG_CELL_ADHESION_MOLECULES_CAMS

Value

0.2

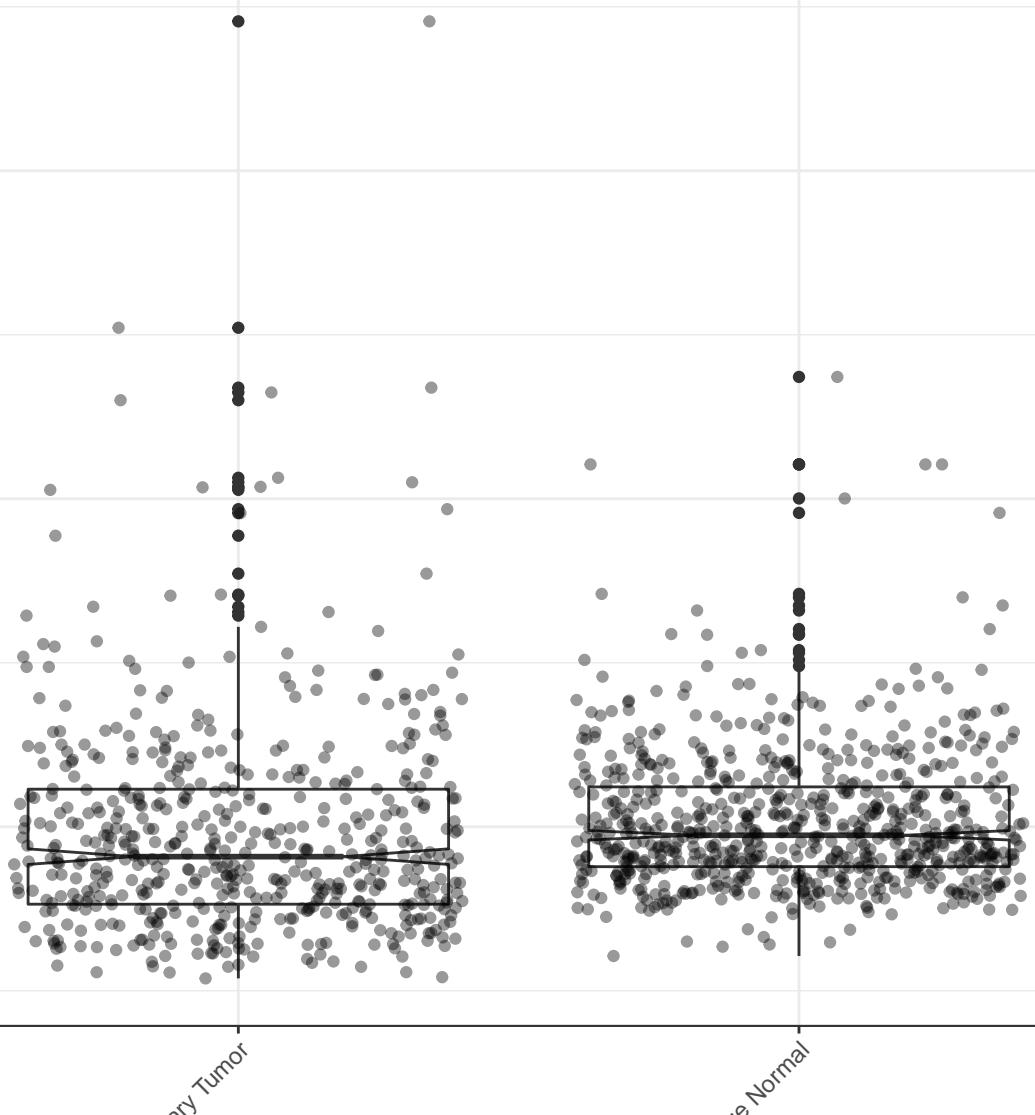
0.1

0.0

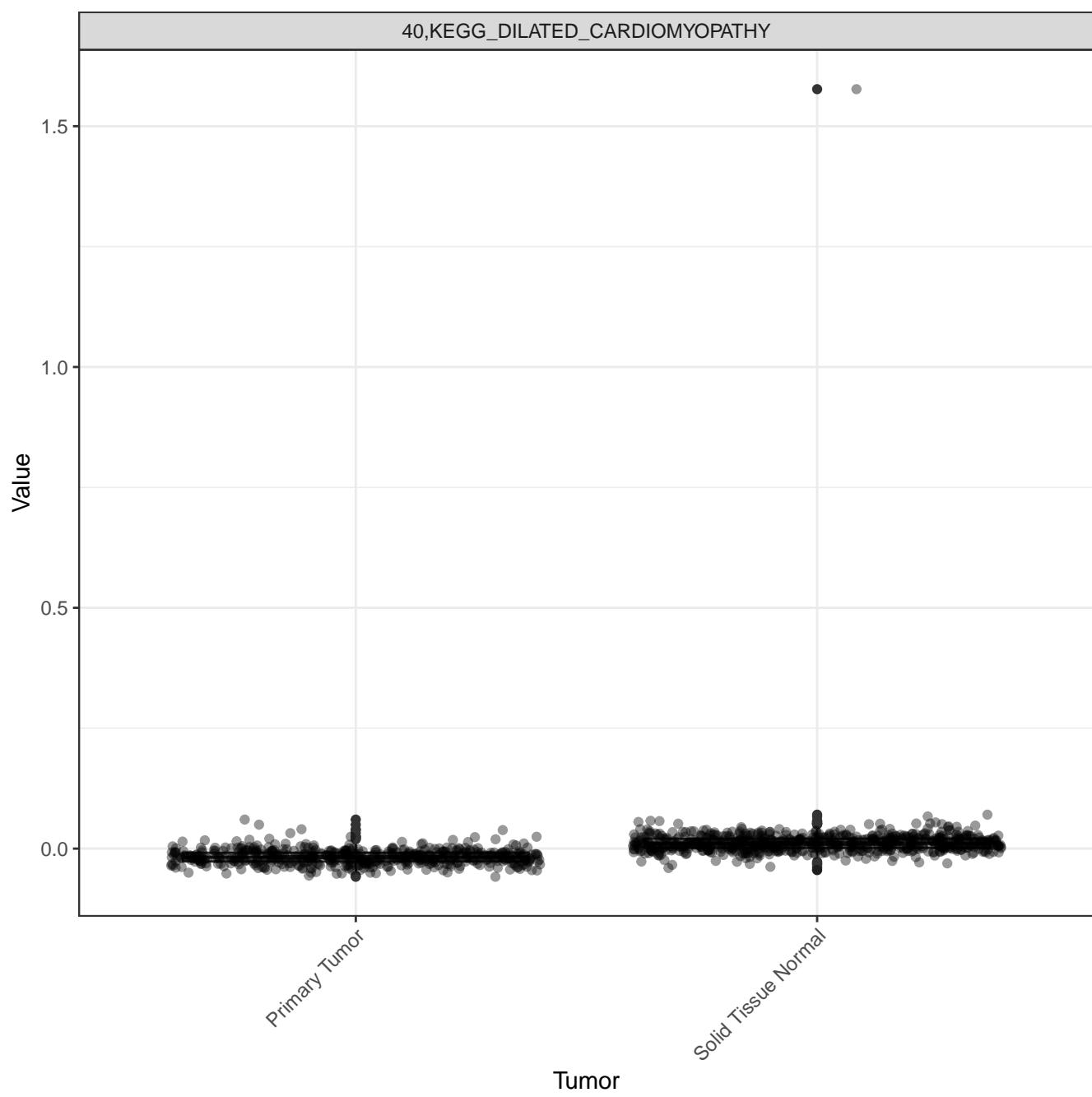
Primary Tumor

Solid Tissue Normal

Tumor



40,KEGG_DILATED_CARDIOMYOPATHY



41,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

Value

0.2

0.1

0.0

Primary Tumor

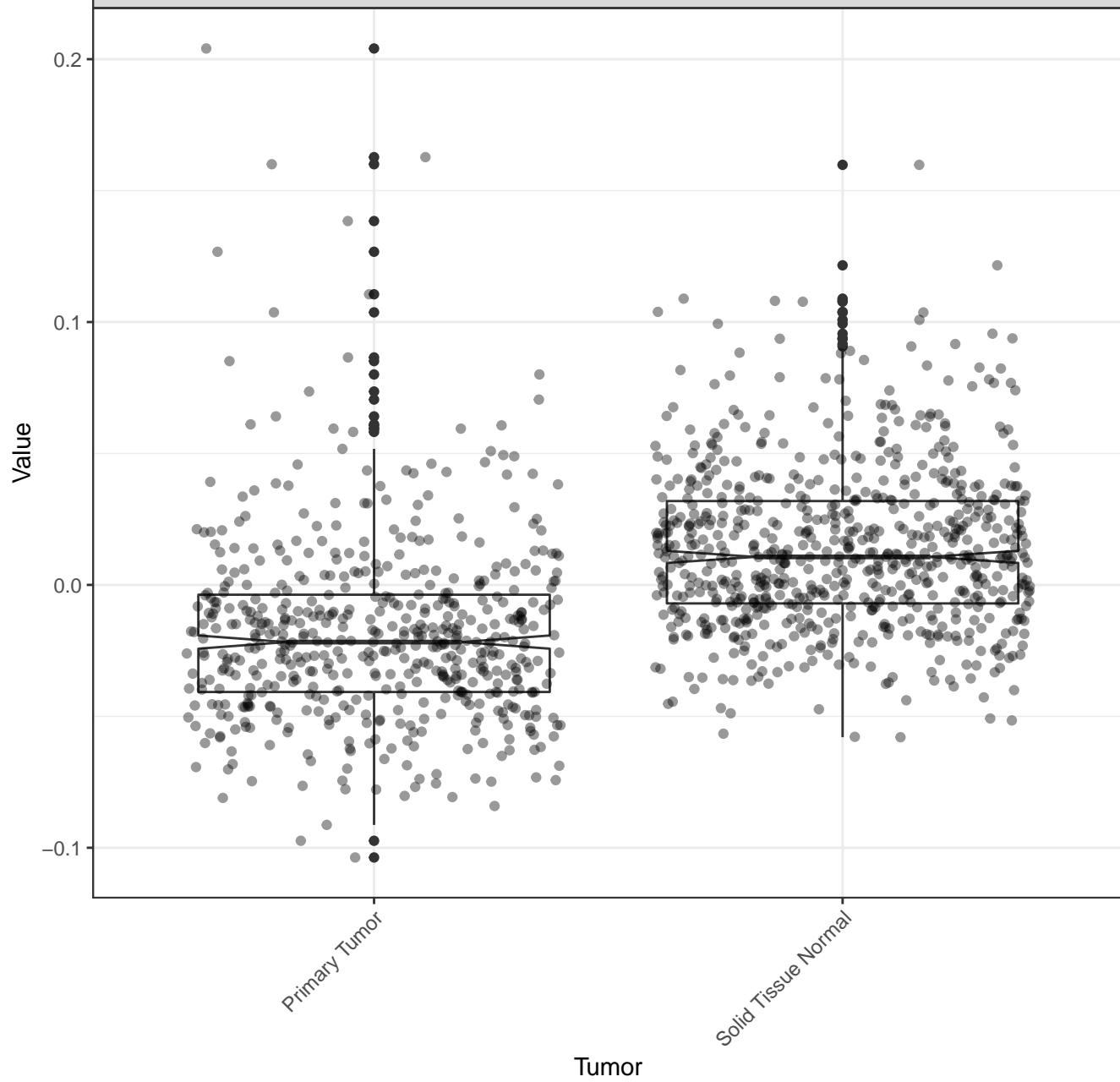
Solid Tissue Normal

Tumor

41,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

41,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

42,REACTOME DEVELOPMENTAL BIOLOGY



43,KEGG_BASAL_CELL_CARCINOMA

Value

0.6

0.4

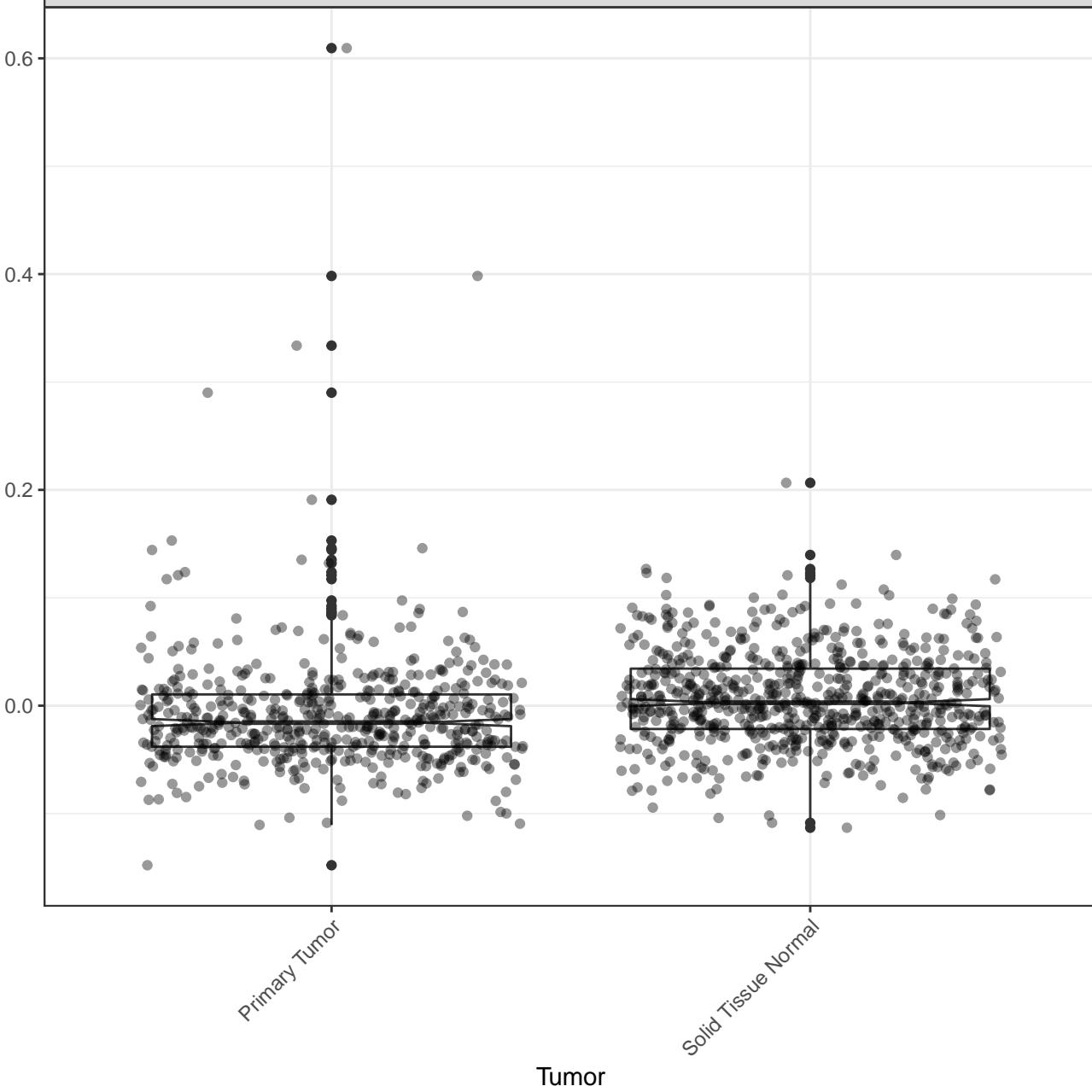
0.2

0.0

Primary Tumor

Solid Tissue Normal

Tumor



44,MIPS_55S_RIBOSOME MITOCHONDRIAL

Value

0.50

0.25

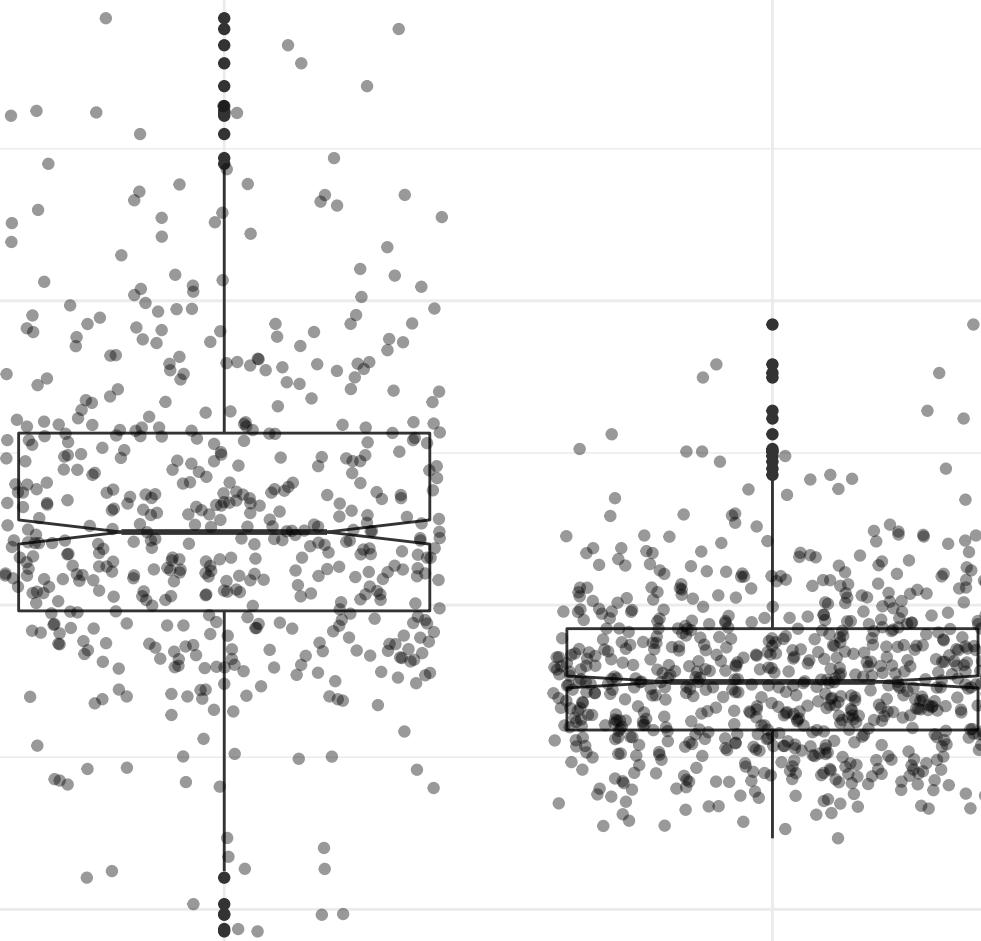
0.00

-0.25

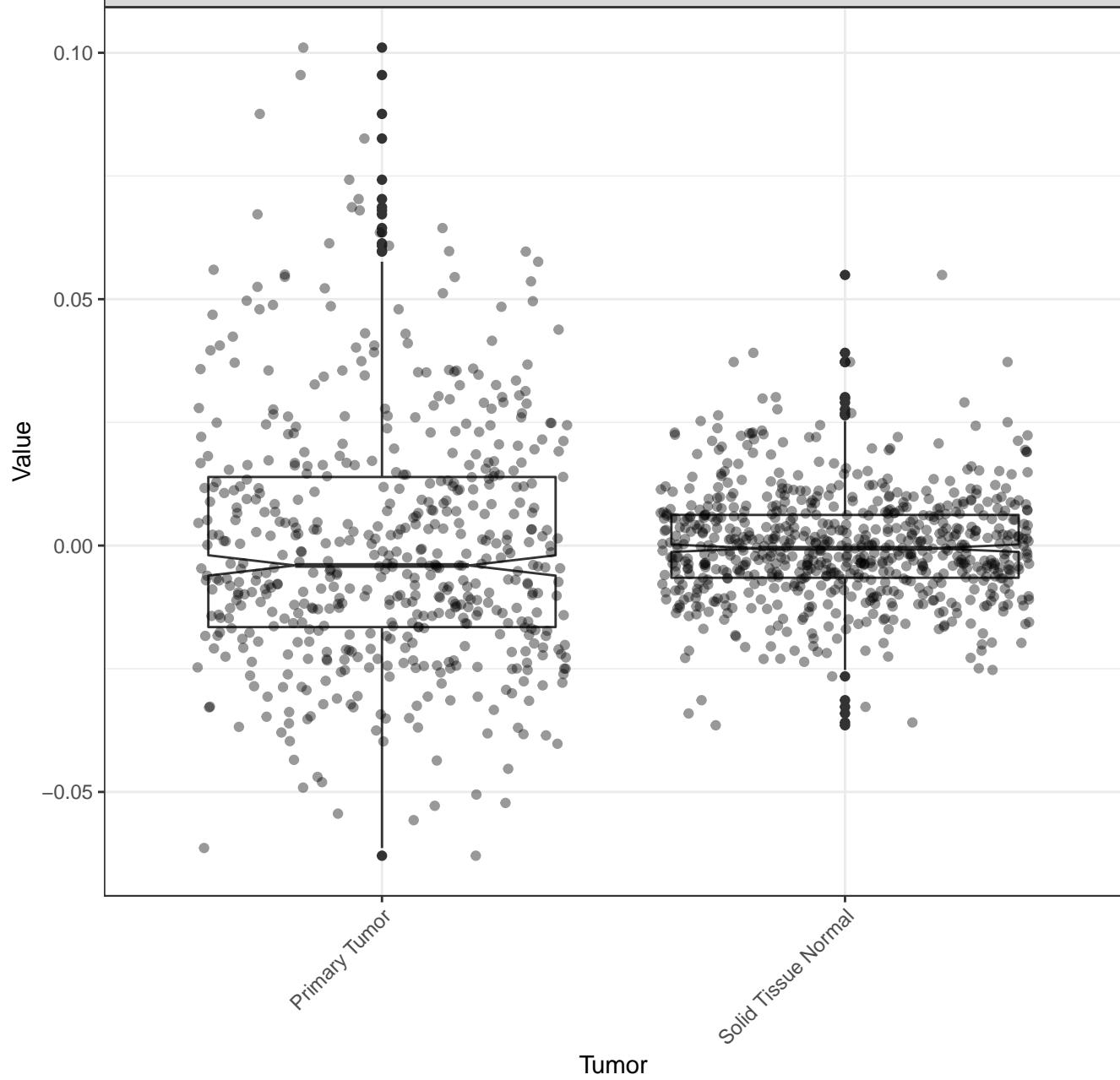
Primary Tumor

Solid Tissue Normal

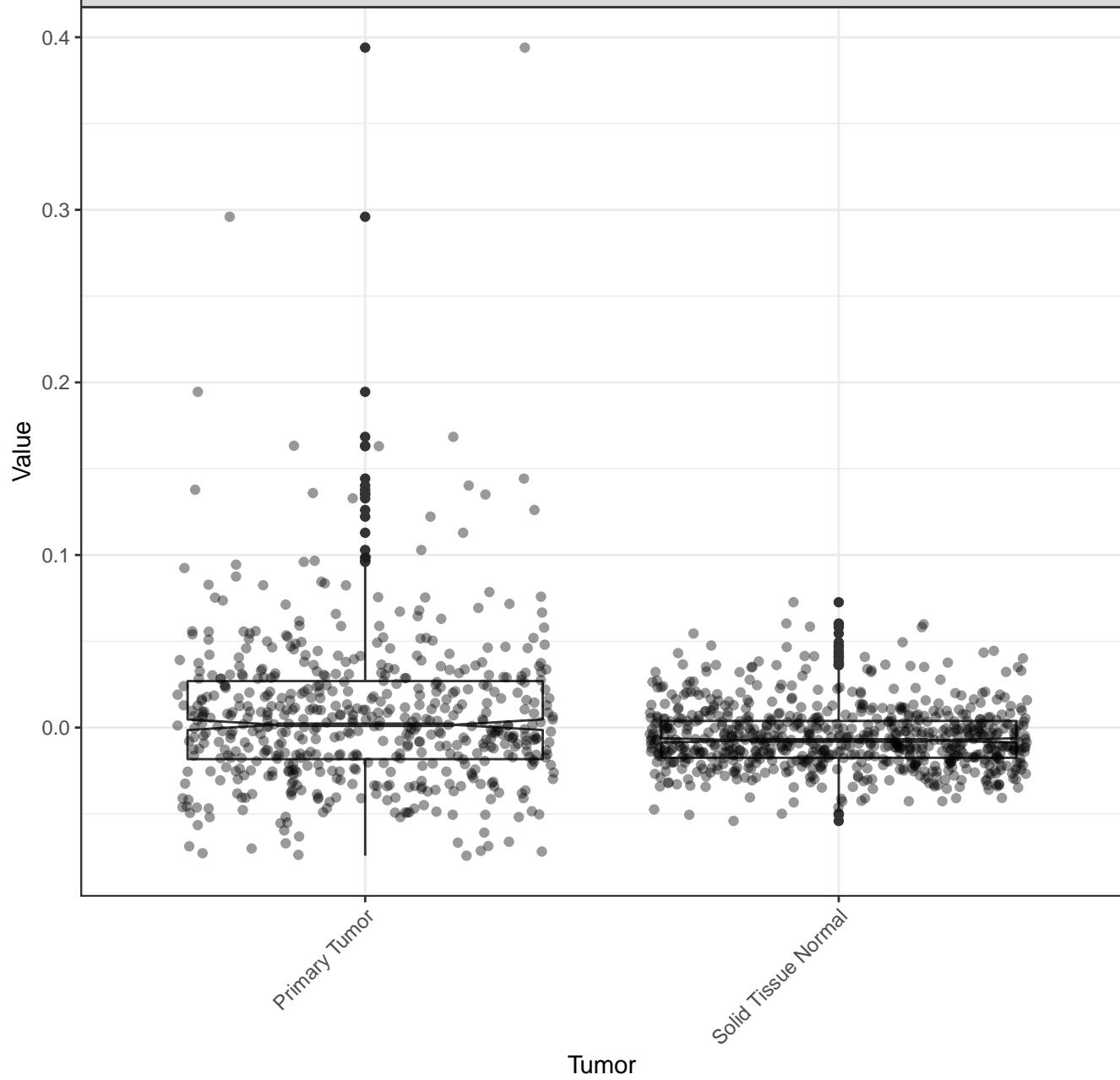
Tumor



45,PID_MYC_REPRESSPATHWAY

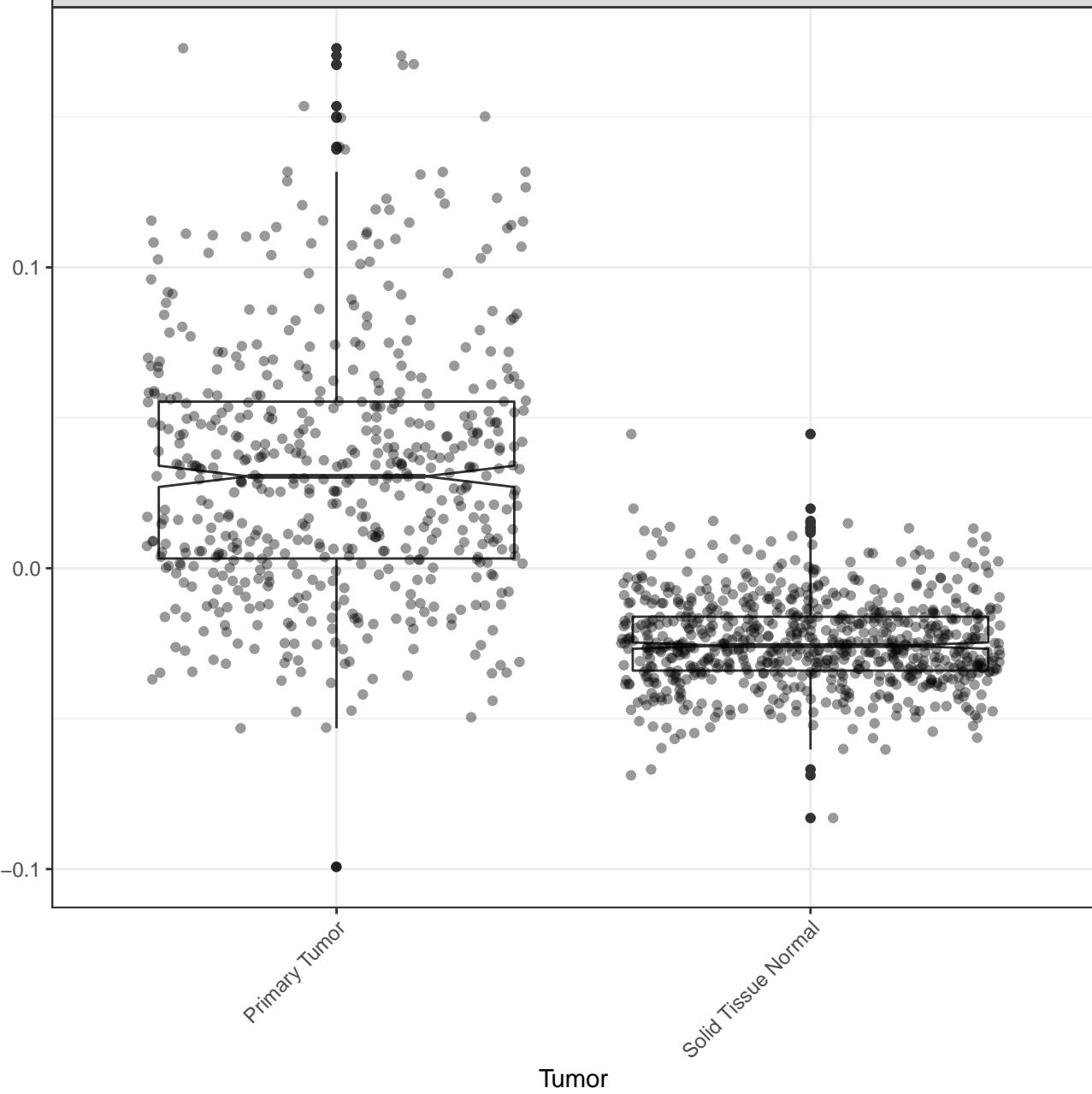


46,REACTOME_AXON_GUIDANCE

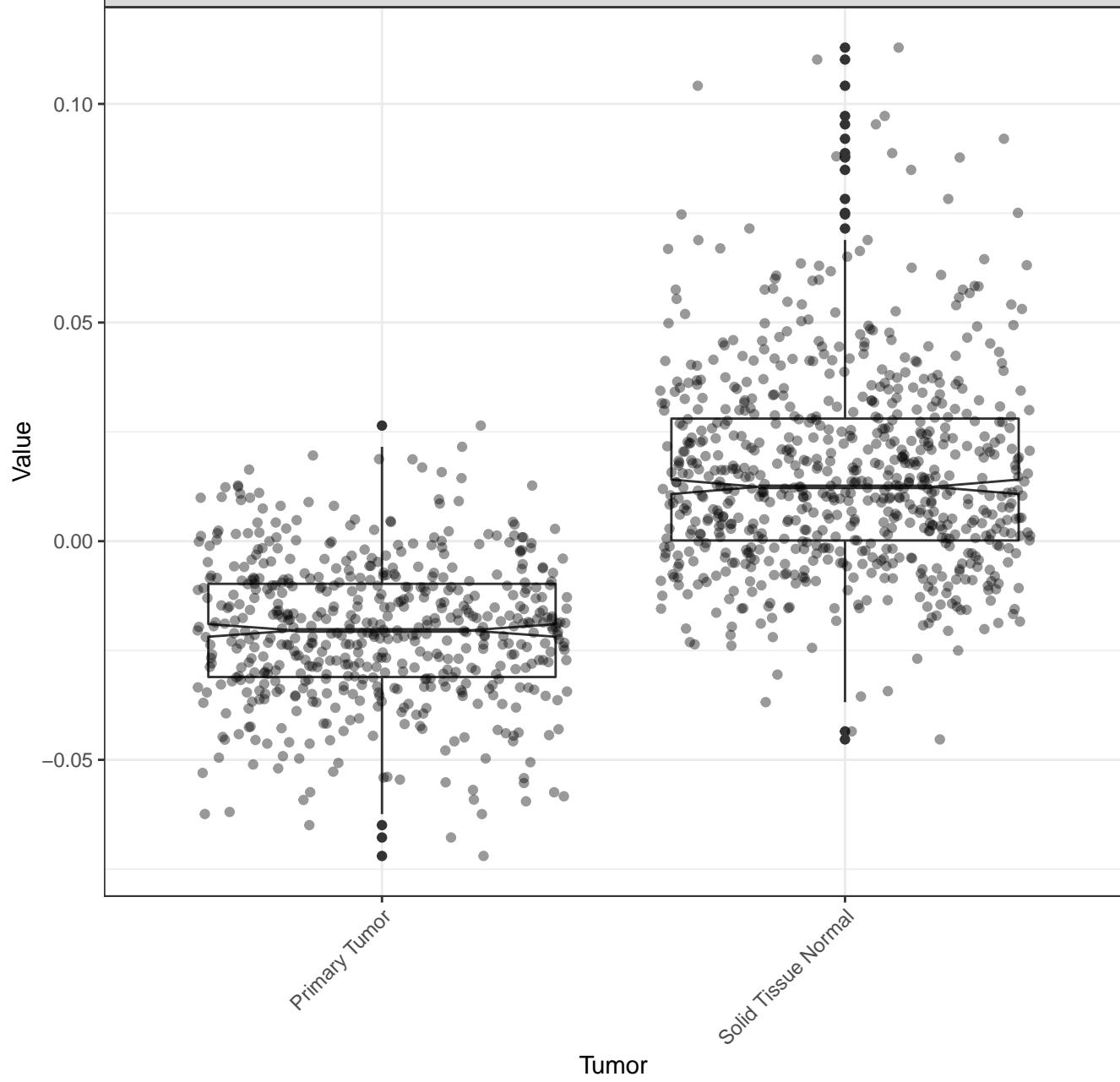


47,PID_A6B1_A6B4_INTEGRIN_PATHWAY

Value



48,IRIS_Neutrophil–Resting



49,SVM Neutrophils

Value

0.15

0.10

0.05

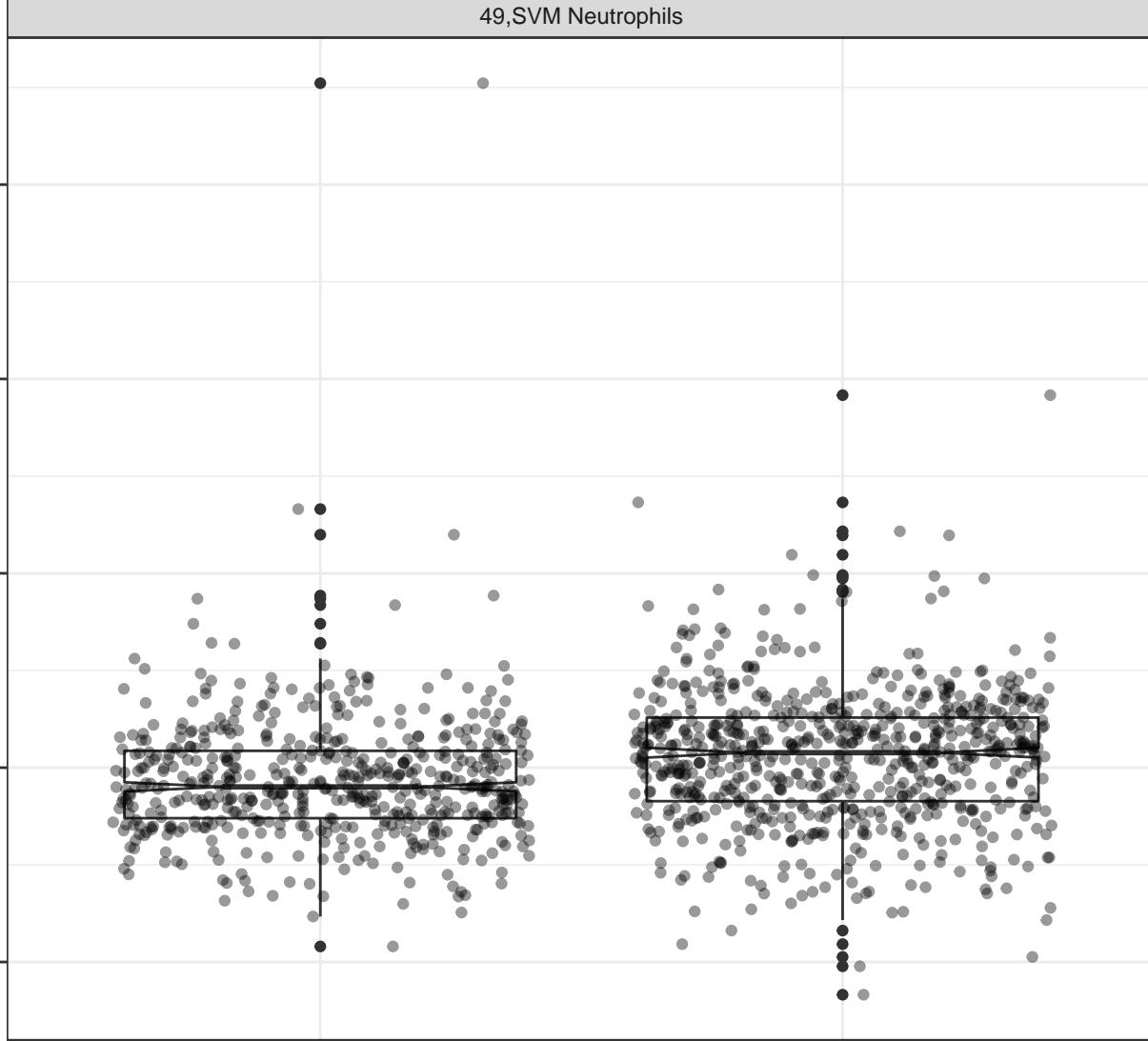
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.4

0.2

0.0

Primary Tumor

Solid Tissue Normal

Tumor



51,REACTOME_NUCLEAR_RECEPтор_TRANSCRIPTION_PATHWAY

Value

0.10

0.05

0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



52,REACTOME_RIG_I_MDA5_MEDIATED_INDUCTION_OF_IFN_ALPHA_BETA_PATHWAYS

Value

0.15

0.10

0.05

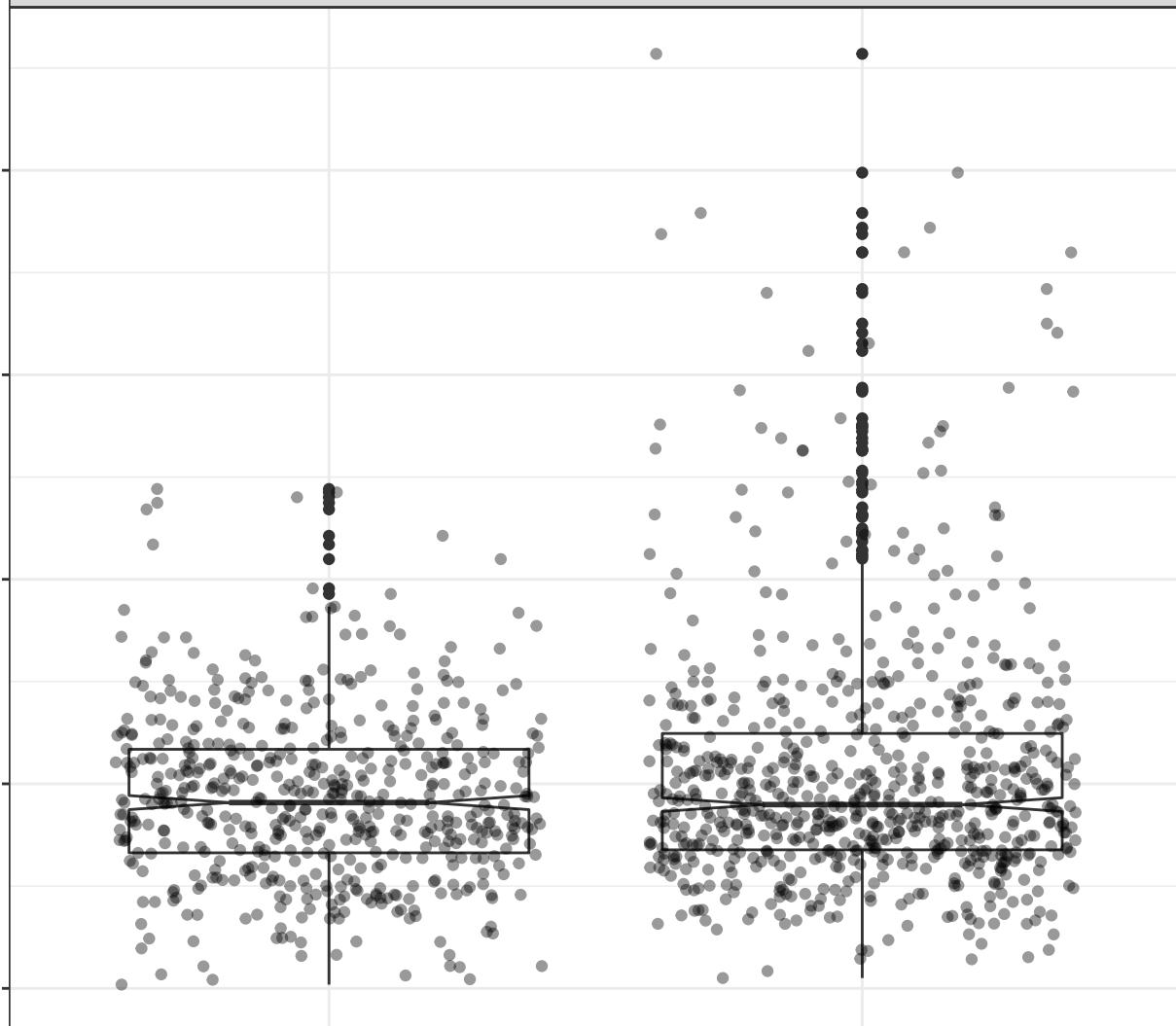
0.00

-0.05

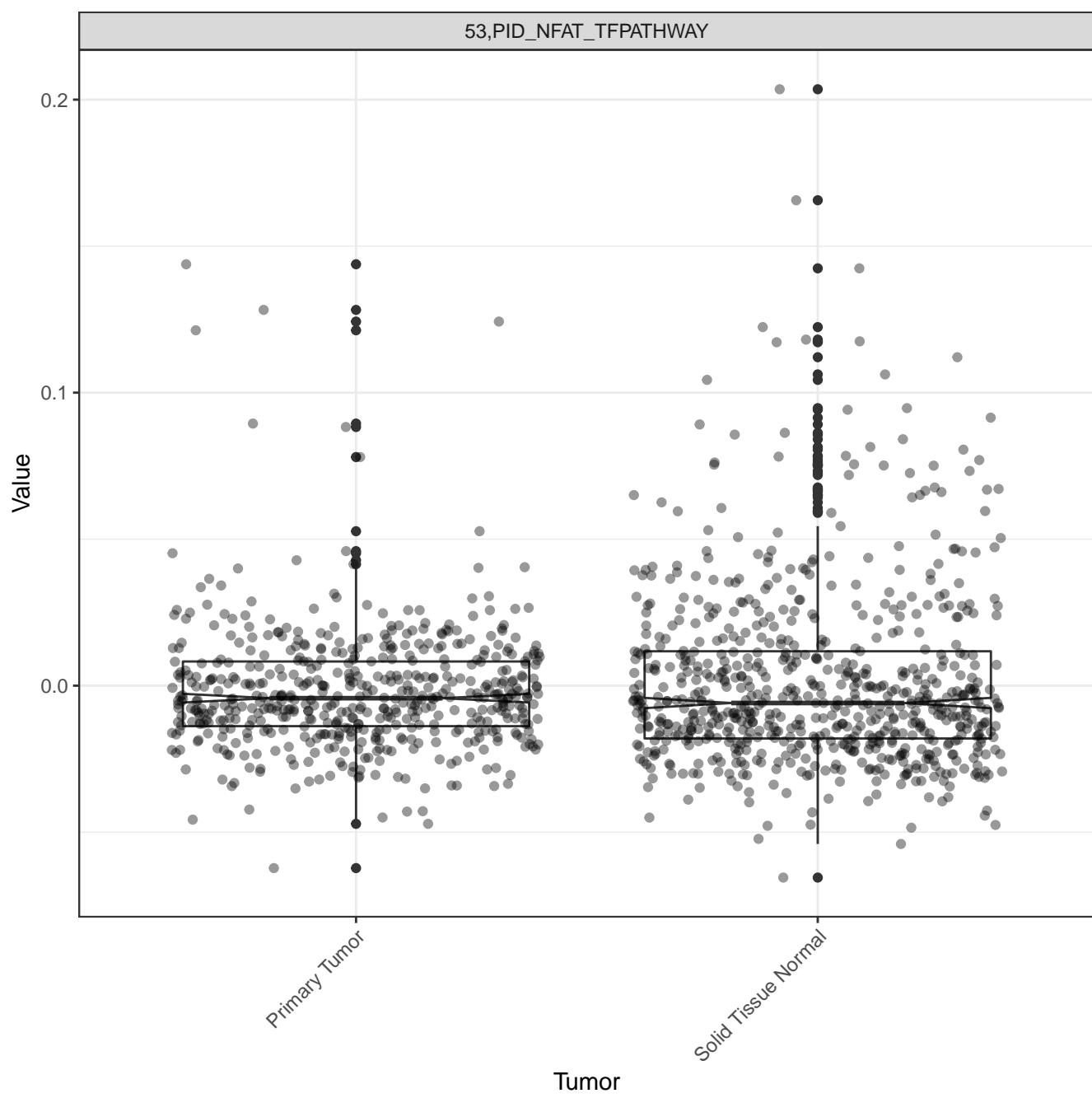
Primary Tumor

Solid Tissue Normal

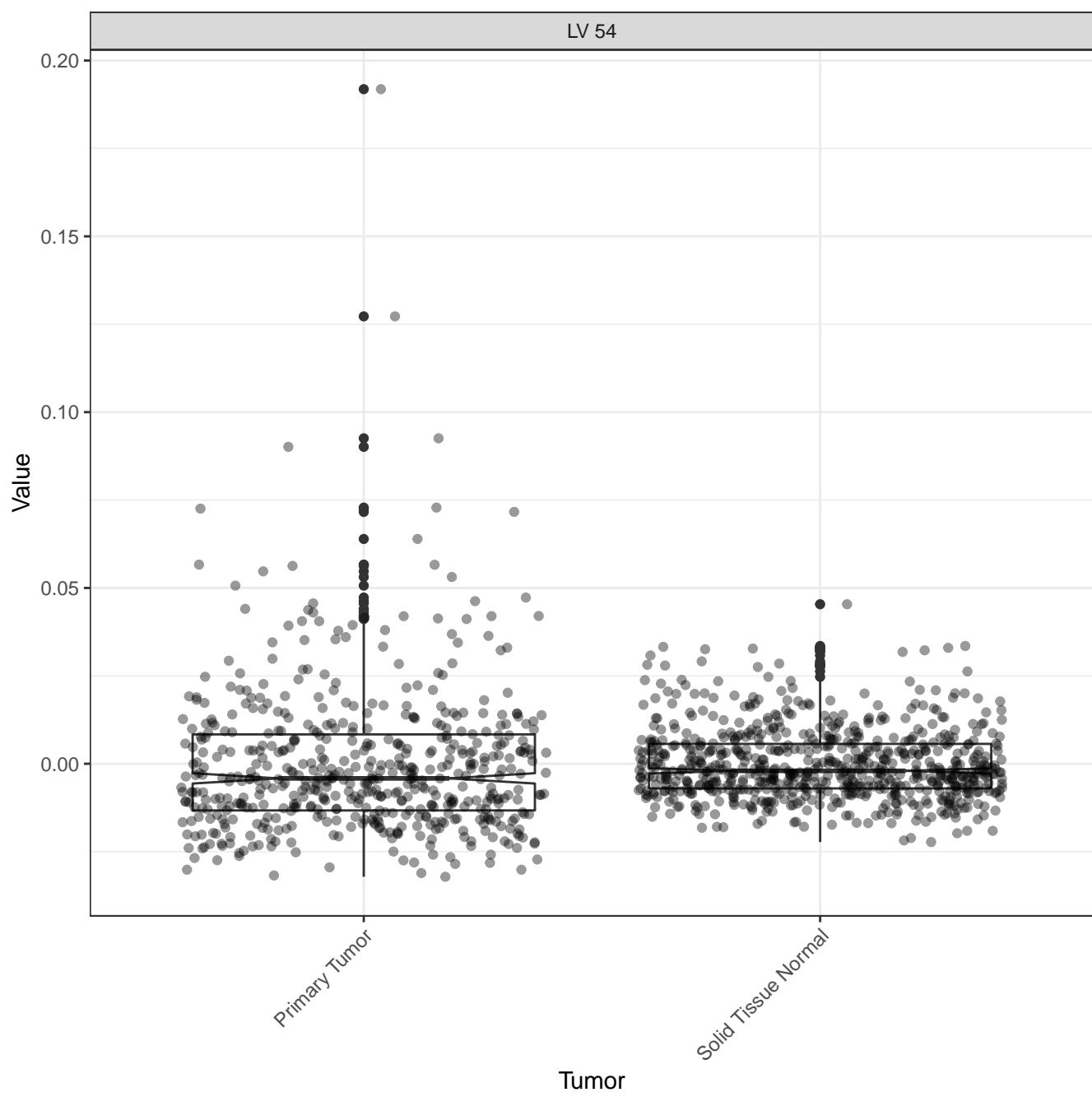
Tumor



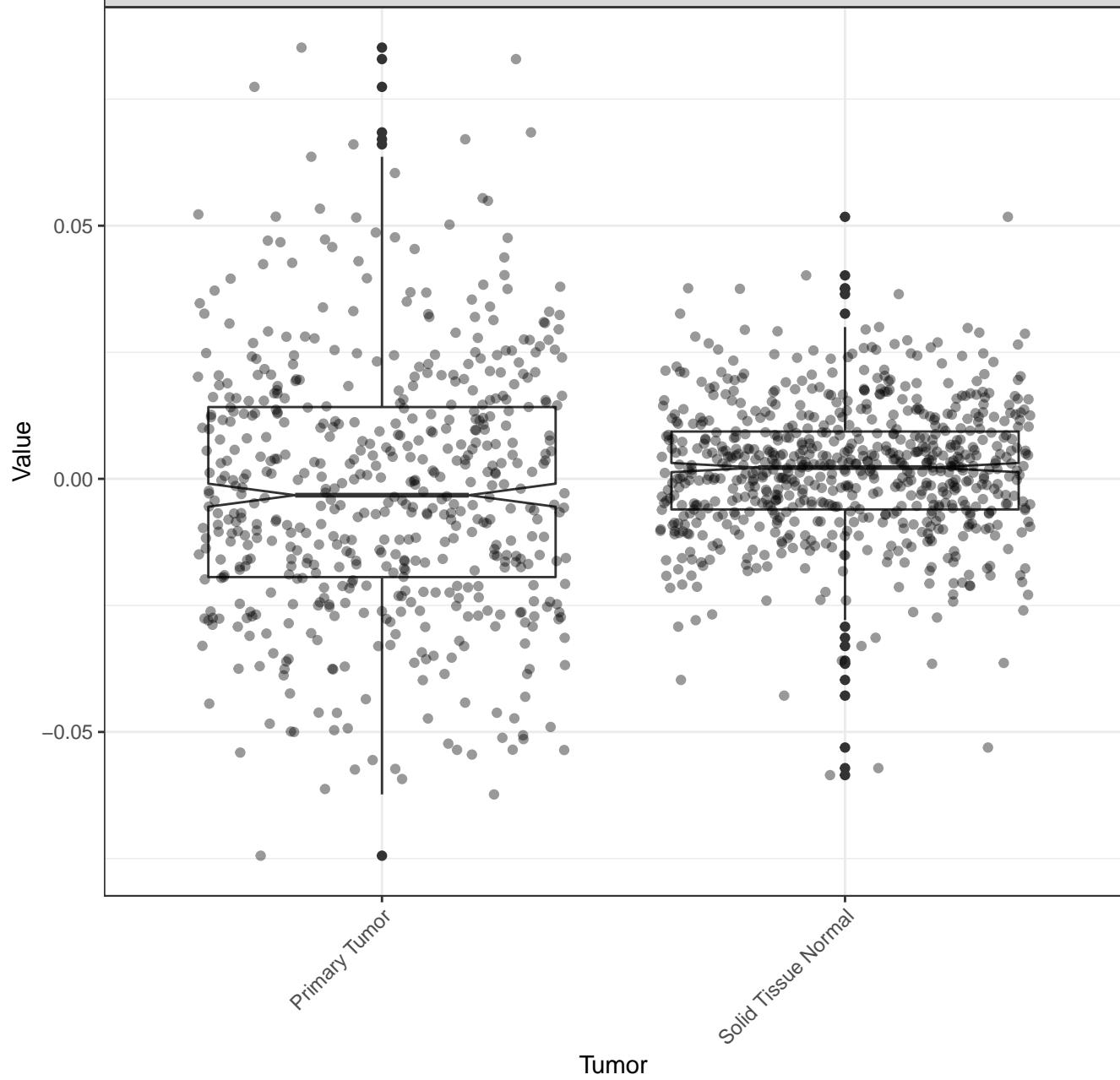
53,PID_NFAT_TFPATHWAY



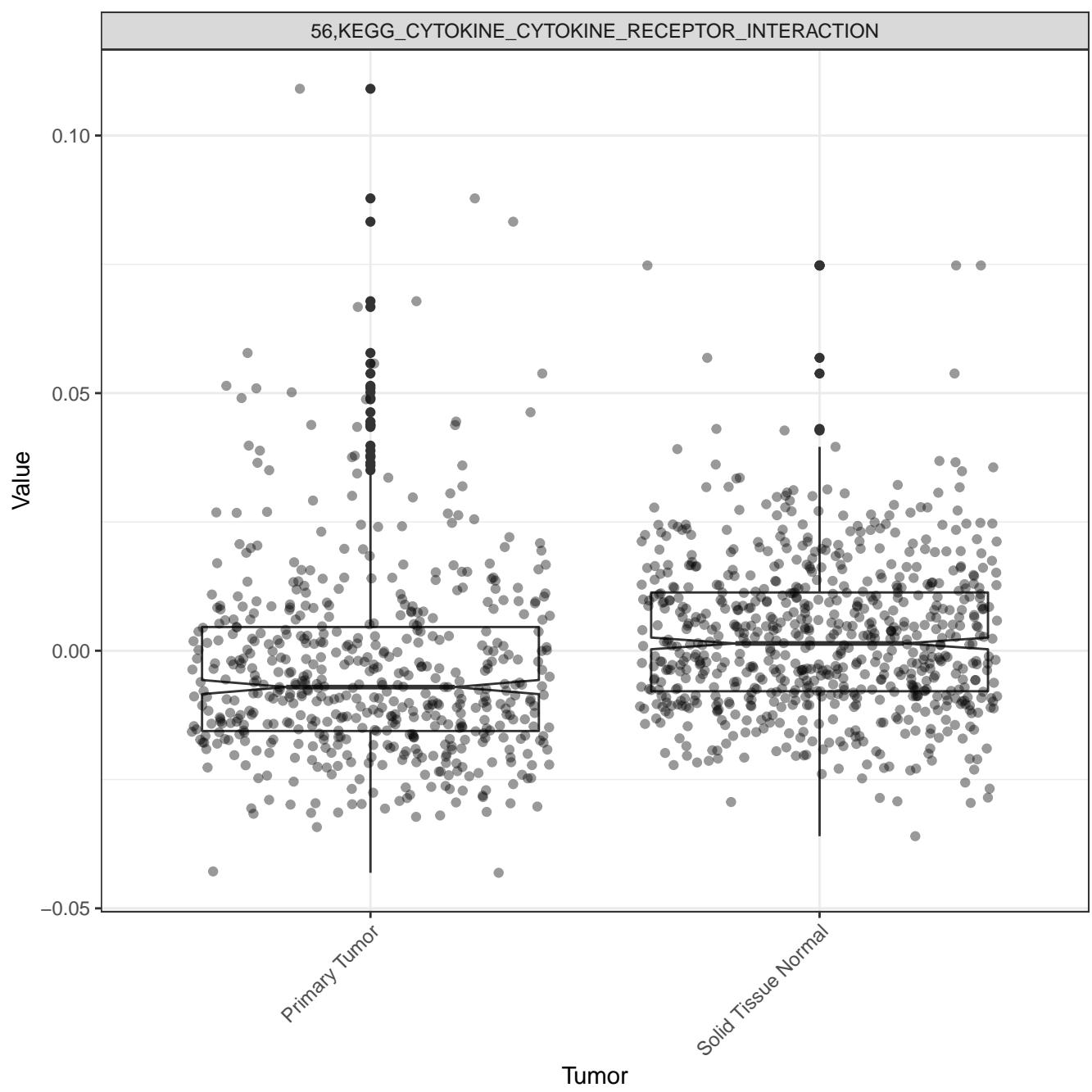
LV 54



55,PID_ERA_GENOMIC_PATHWAY



56,KEGG_CYTOKINE_CYTOKINE_RECECTOR_INTERACTION



57,KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS

Value

0.3

0.2

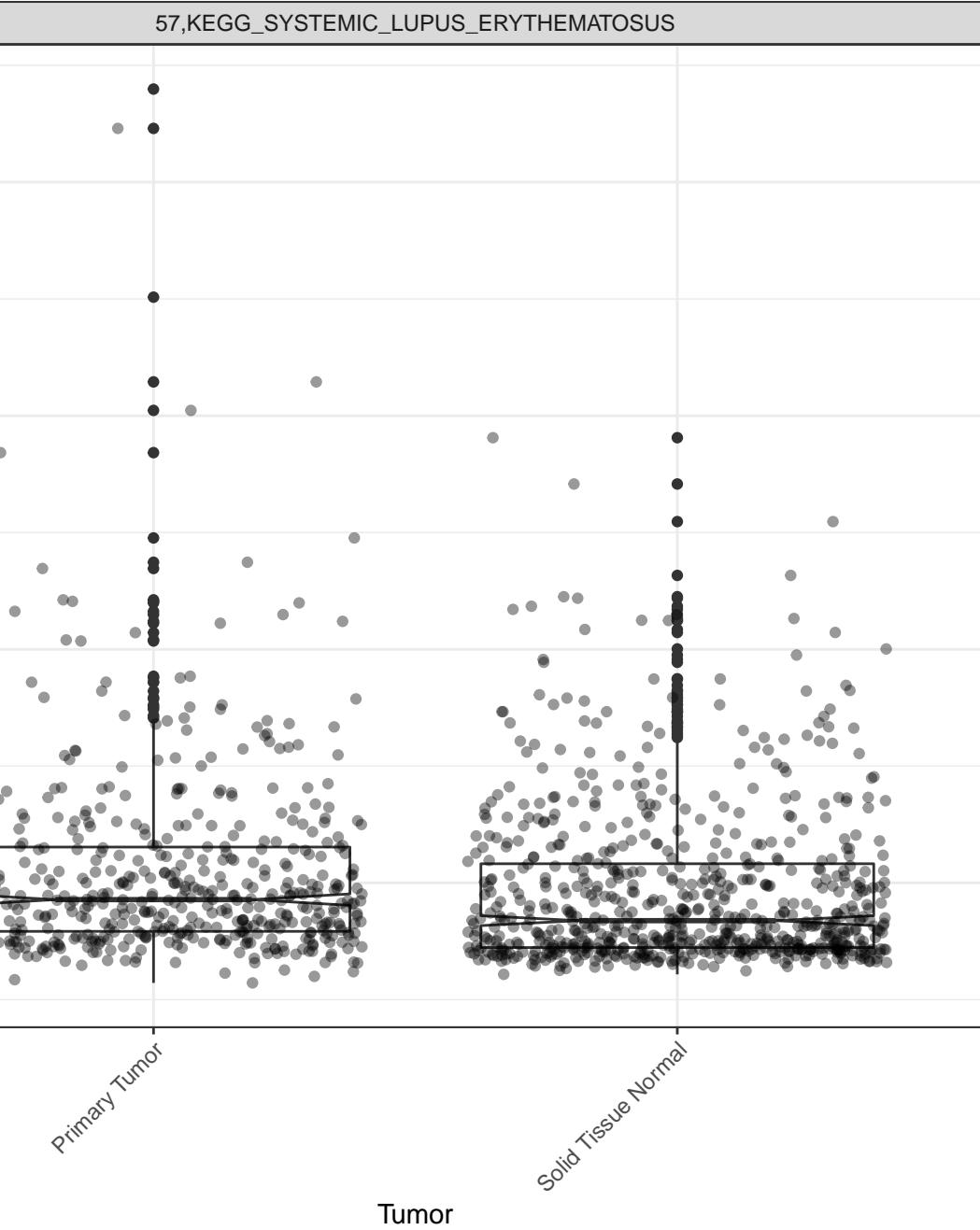
0.1

0.0

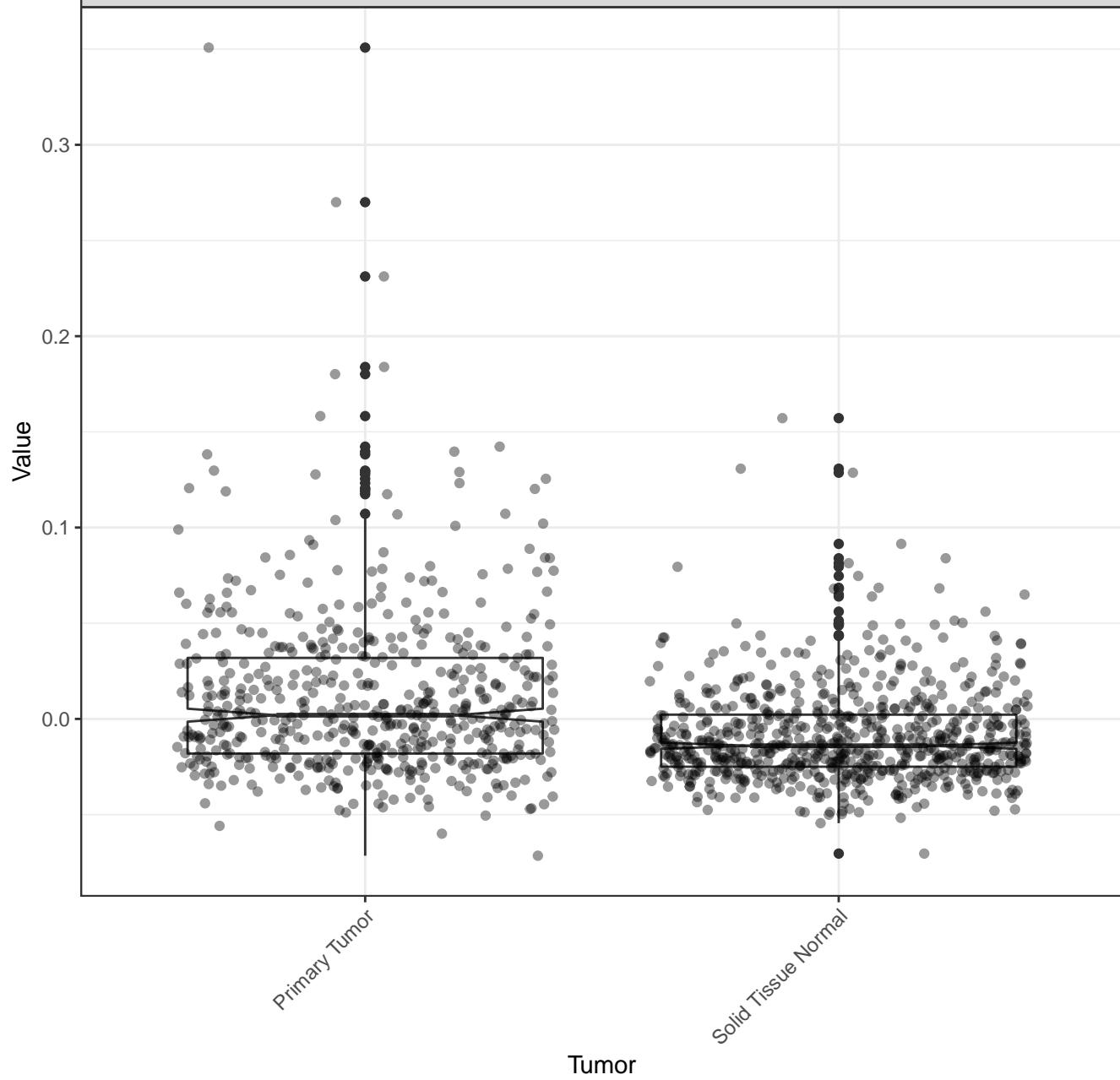
Primary Tumor

Solid Tissue Normal

Tumor



58,SVM Dendritic cells resting



59,REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE

Value

0.15

0.10

0.05

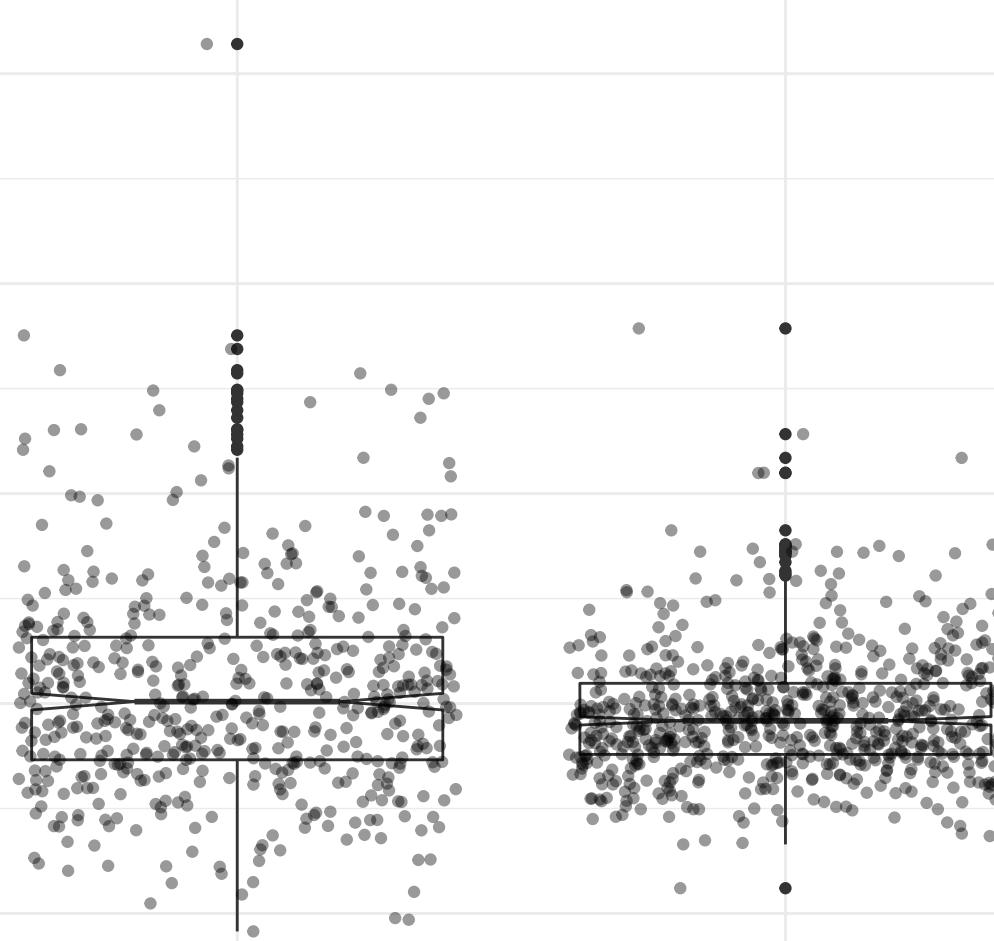
0.00

-0.05

Primary Tumor

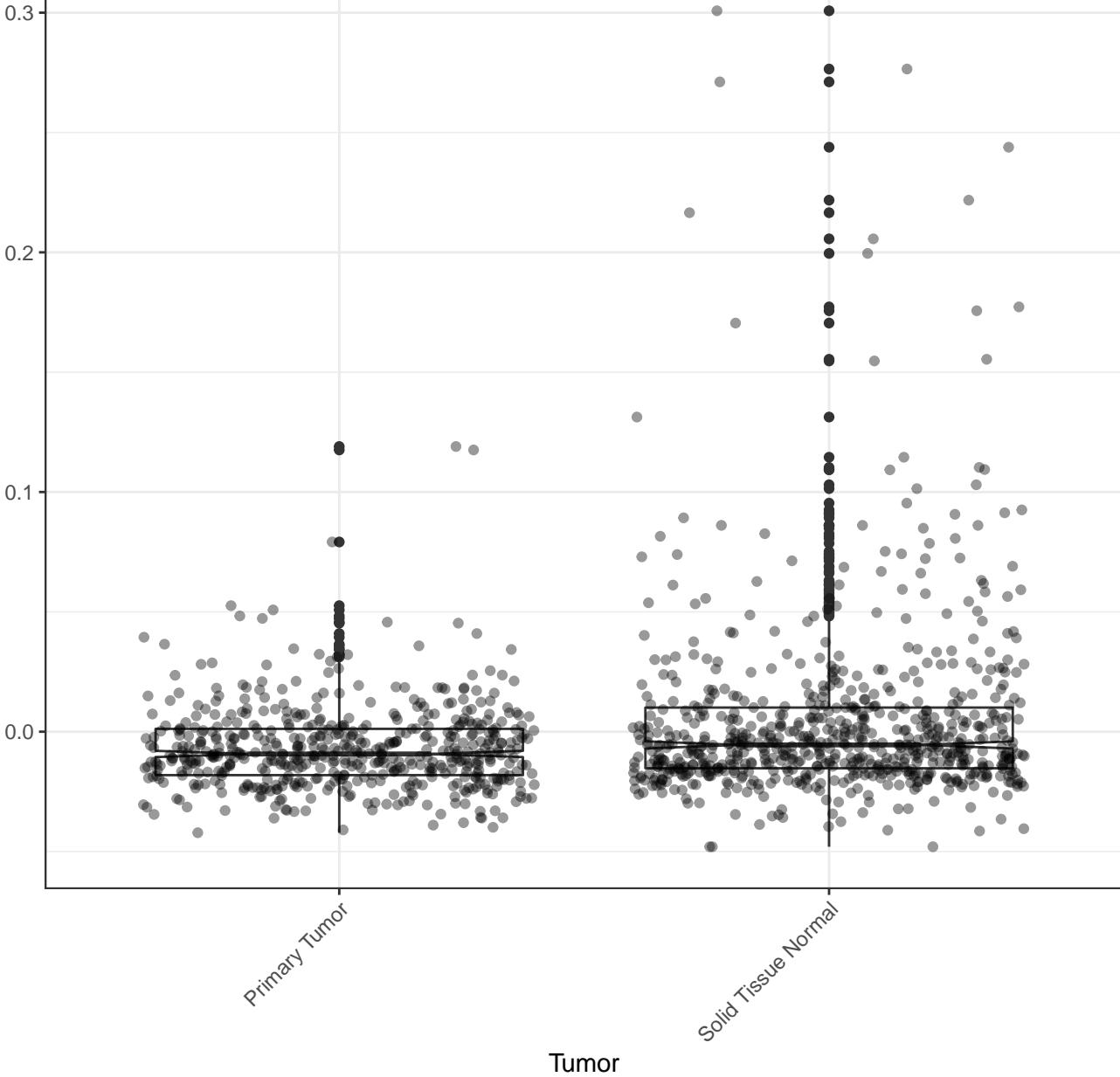
Solid Tissue Normal

Tumor

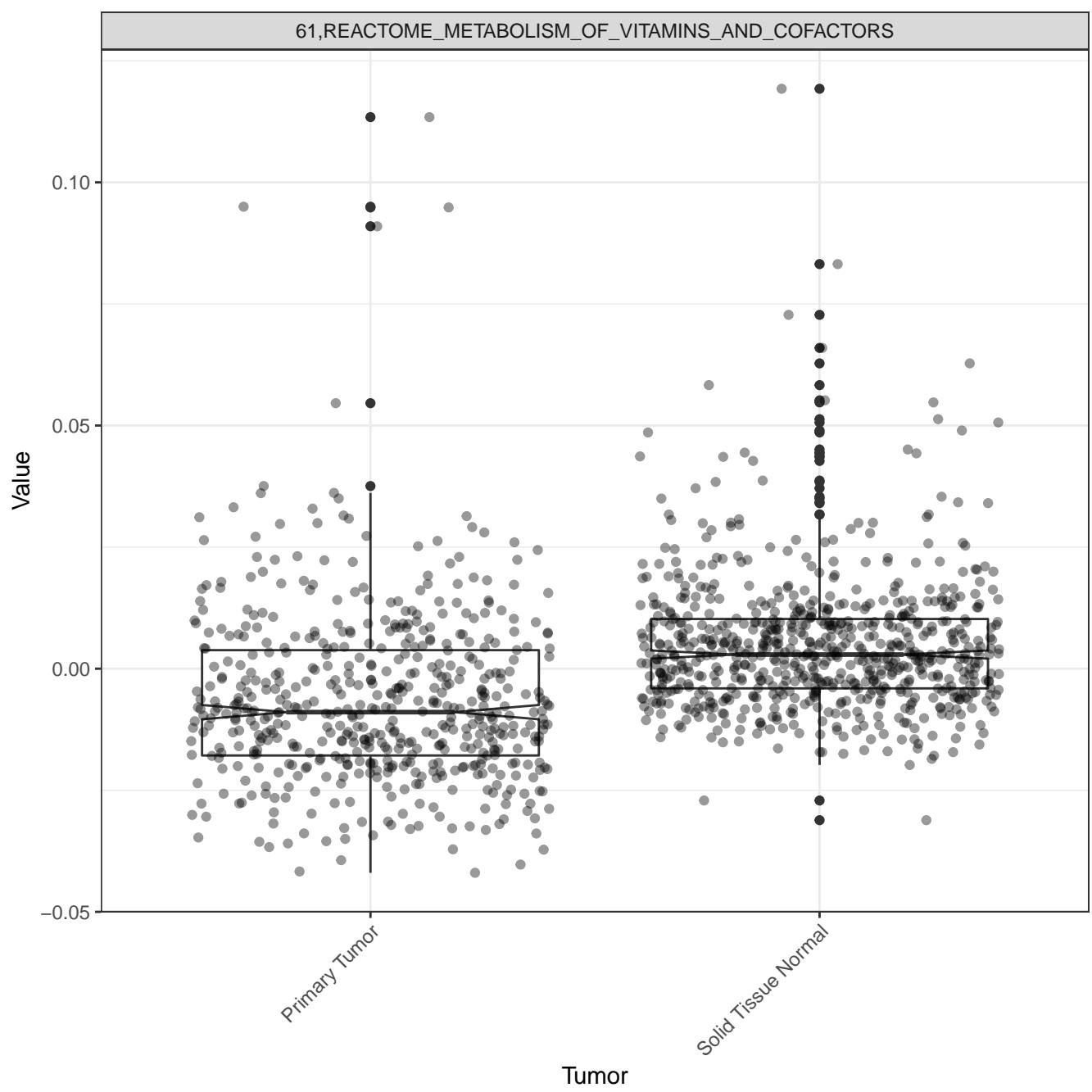


60,IRIS_NKcell-control

Value

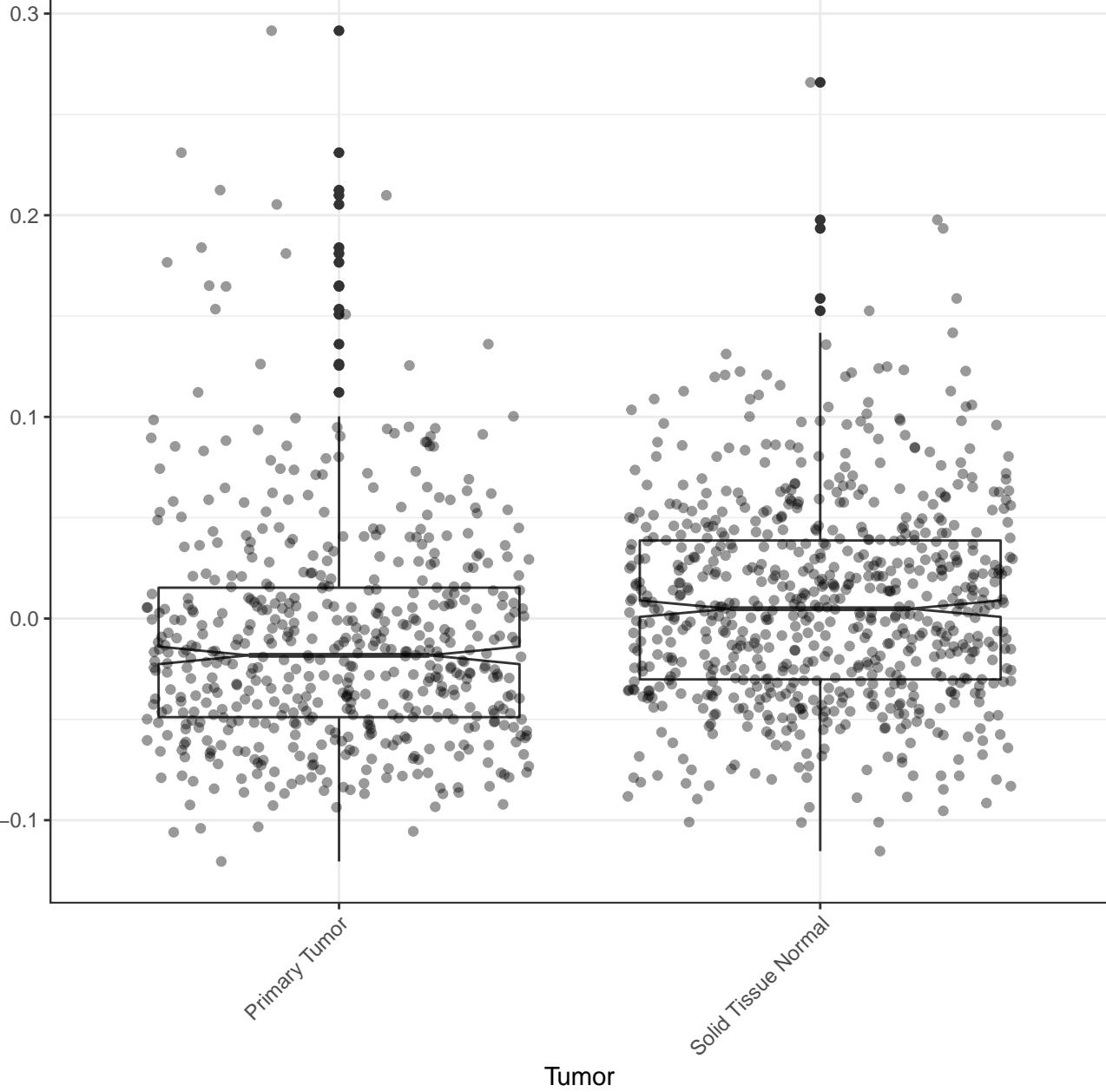


61,REACTOME_METABOLISM_OF_VITAMINS_AND_COFACTORS

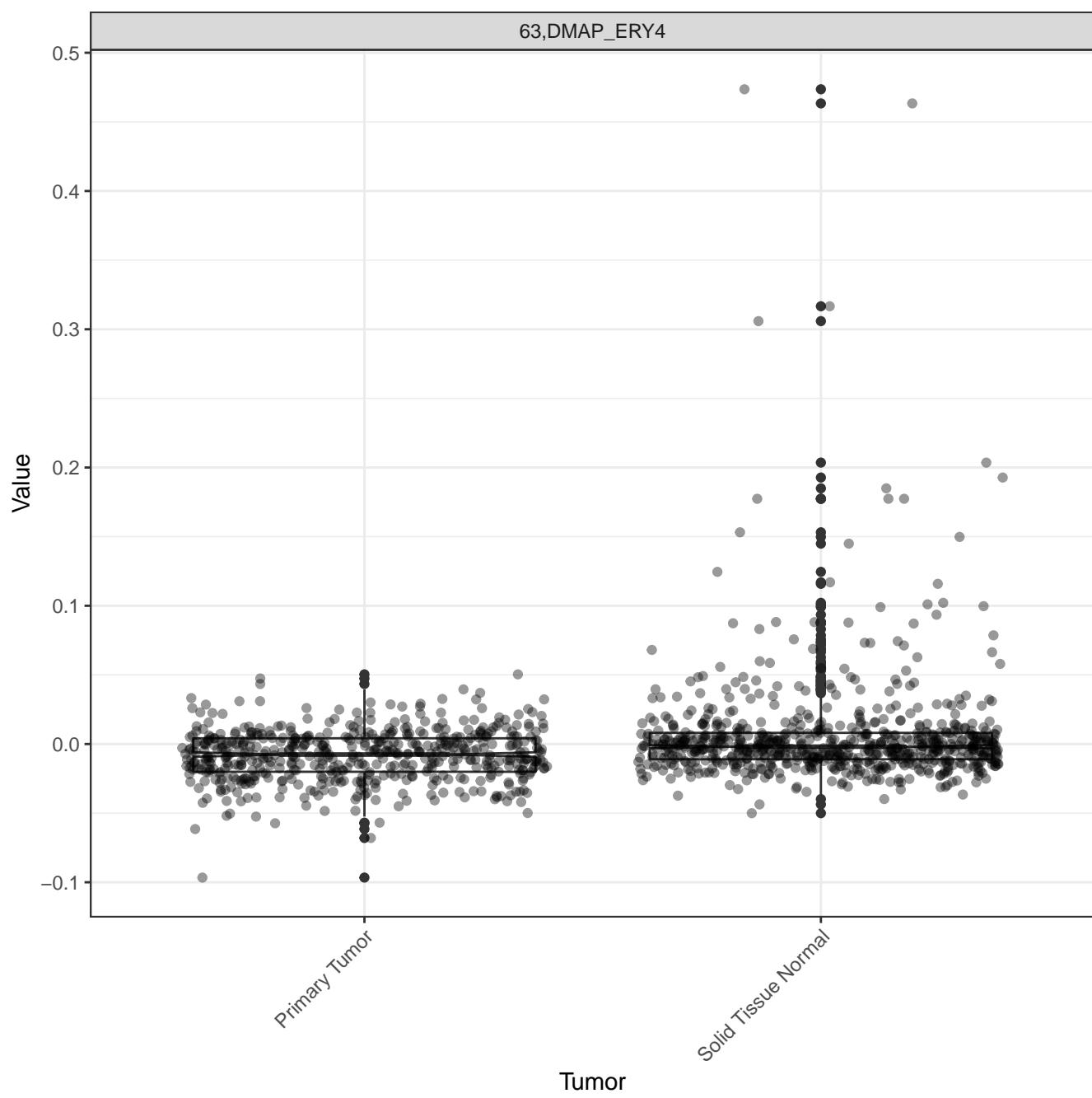


62,KEGG_LYSOSOME

Value

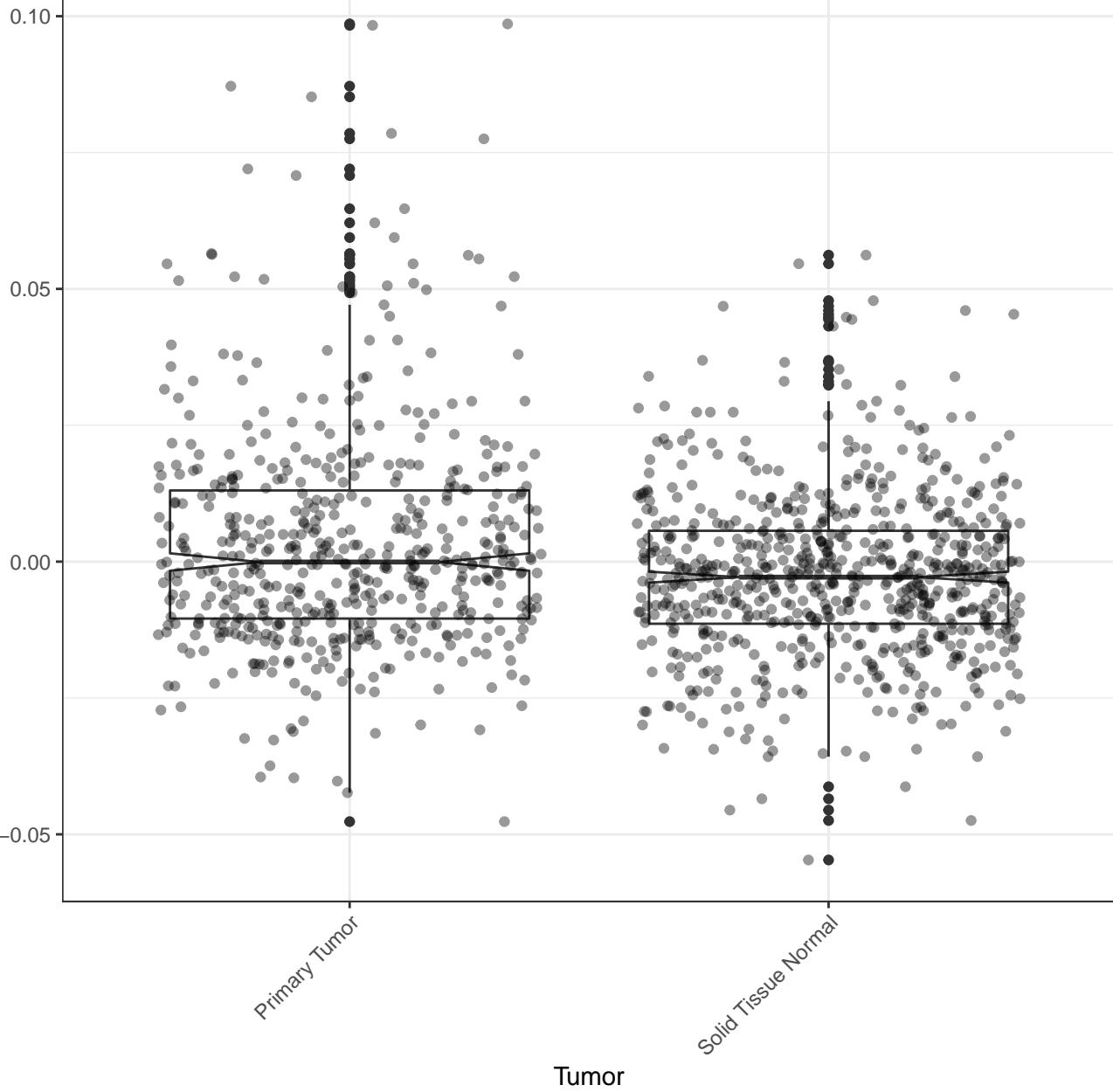


63,DMAP_ERY4

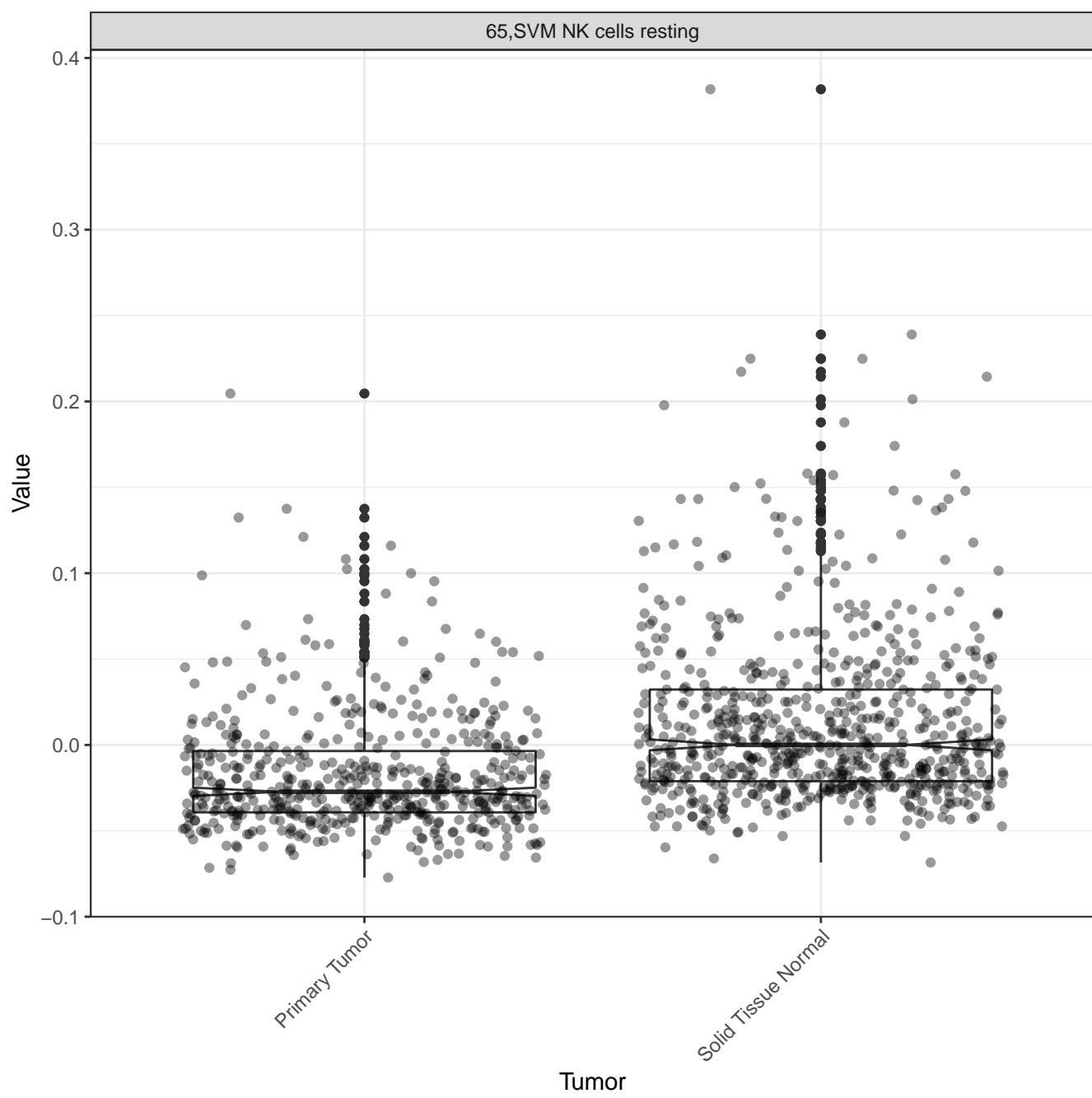


64,DMAP_DENDA2

Value



65,SVM NK cells resting



66,IRIS_Neutrophil–Resting

Value

0.2

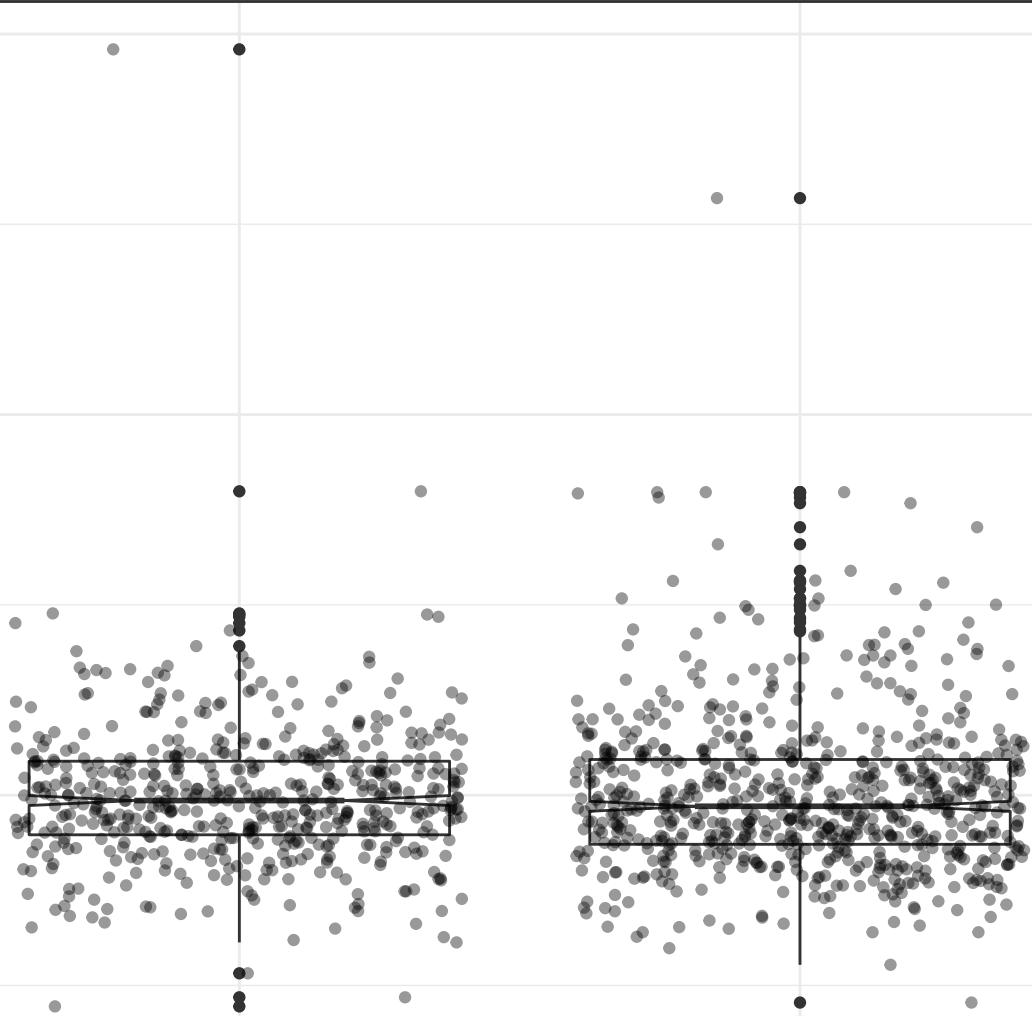
0.1

0.0

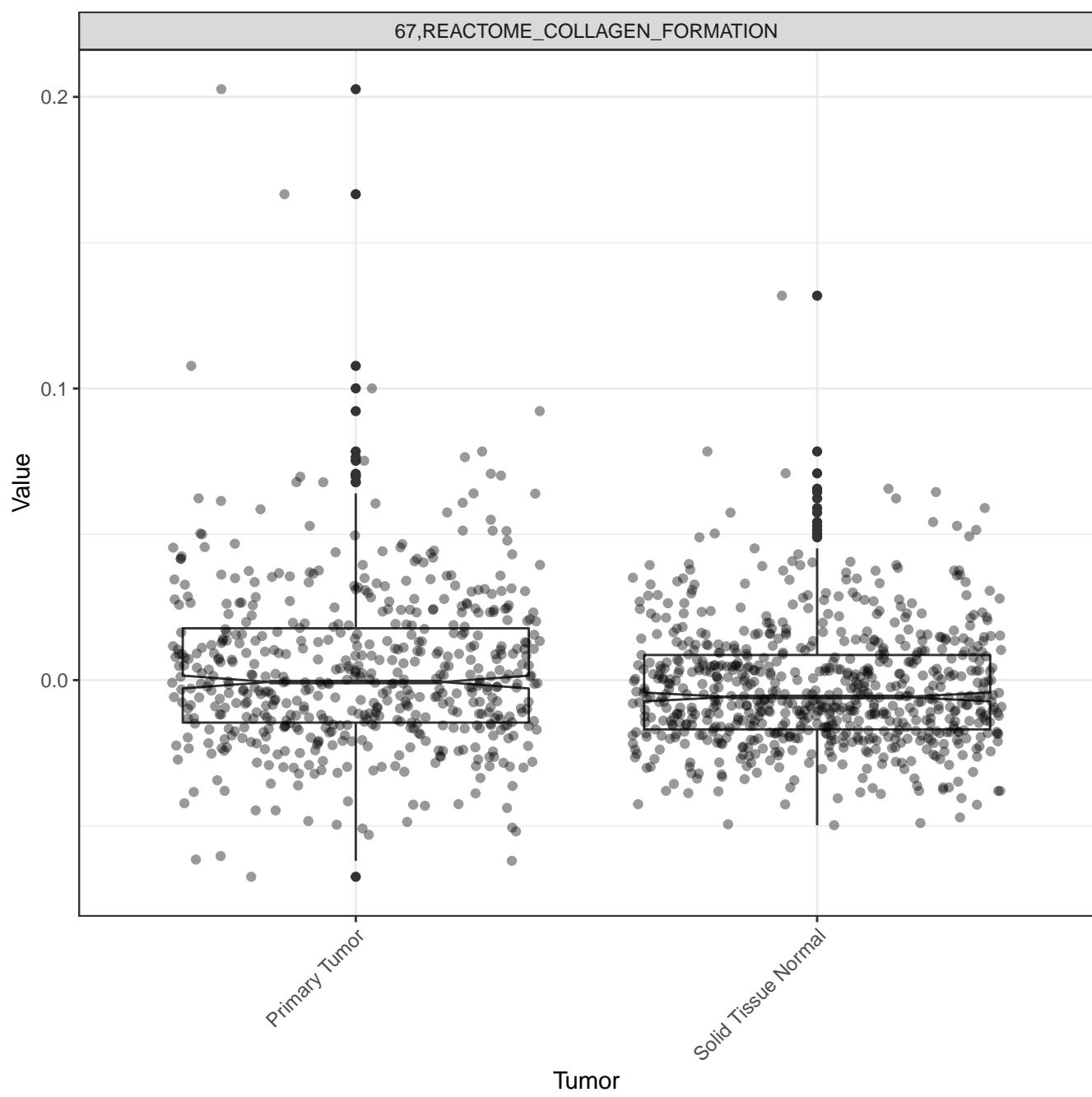
Primary Tumor

Solid Tissue Normal

Tumor



67,REACTOME_COLLAGEN_FORMATION



68,REACTOME_RESPIRATORY_ELECTRON_TRANSPORT

Value

0.20

0.15

0.10

0.05

0.00

-0.05

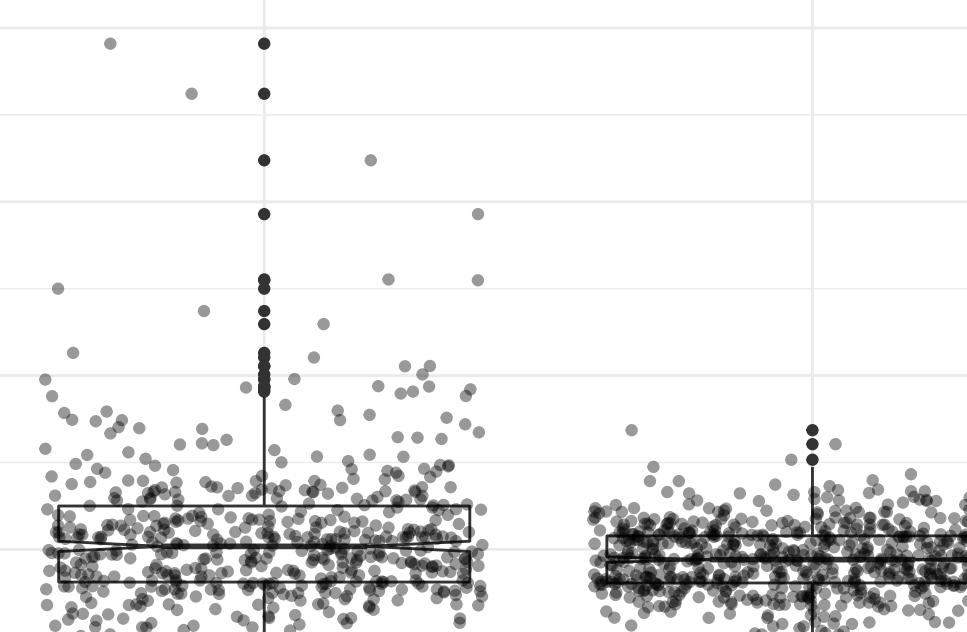
Primary Tumor

Solid Tissue Normal

Tumor

•

•



69,PID_EPHBFWDPATHWAY

Value

0.2

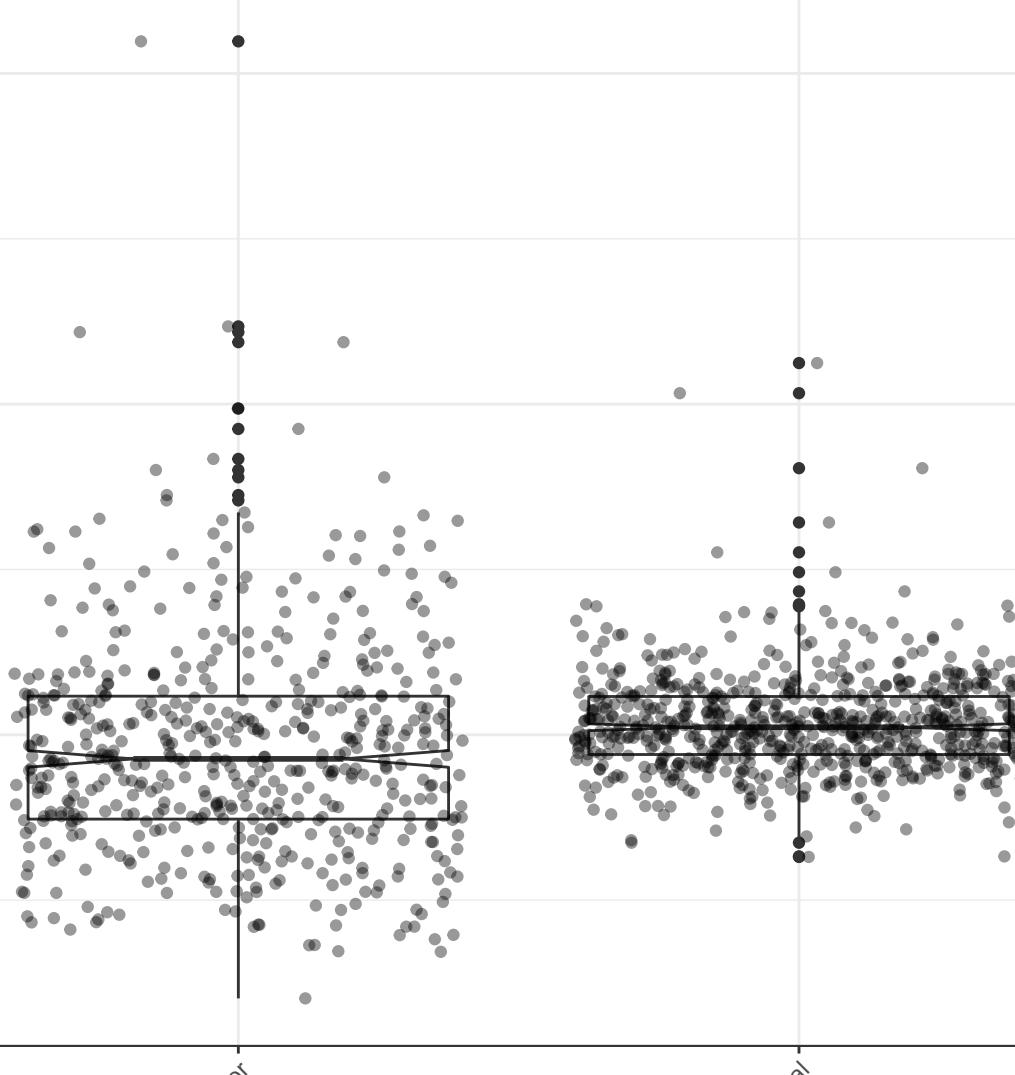
0.1

0.0

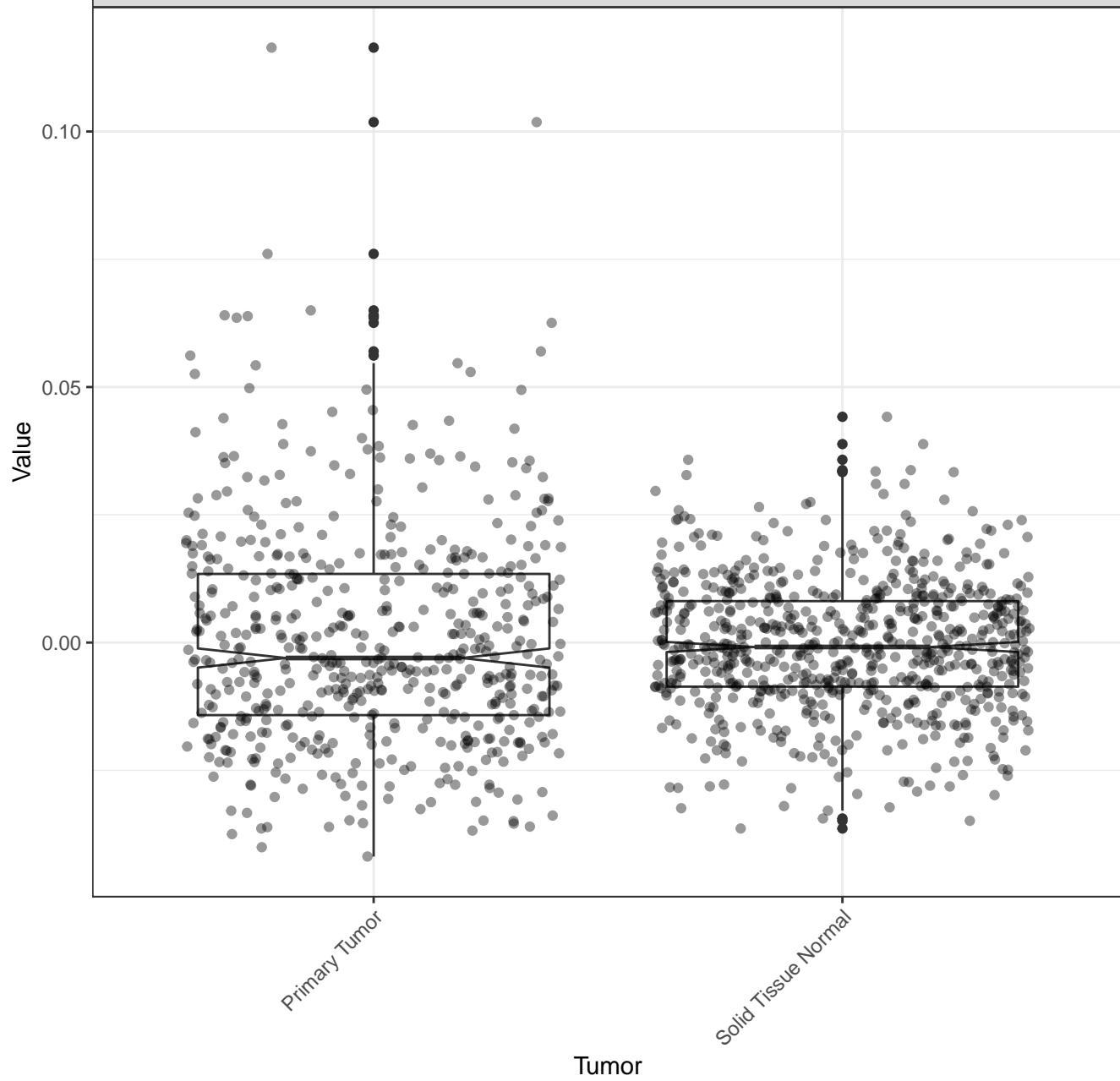
Primary Tumor

Solid Tissue Normal

Tumor



70,KEGG_OXIDATIVE_PHOSPHORYLATION



71,SIG_PIP3_SIGNALING_IN_CARDIAC_MYOCTES

Value

0.15

0.10

0.05

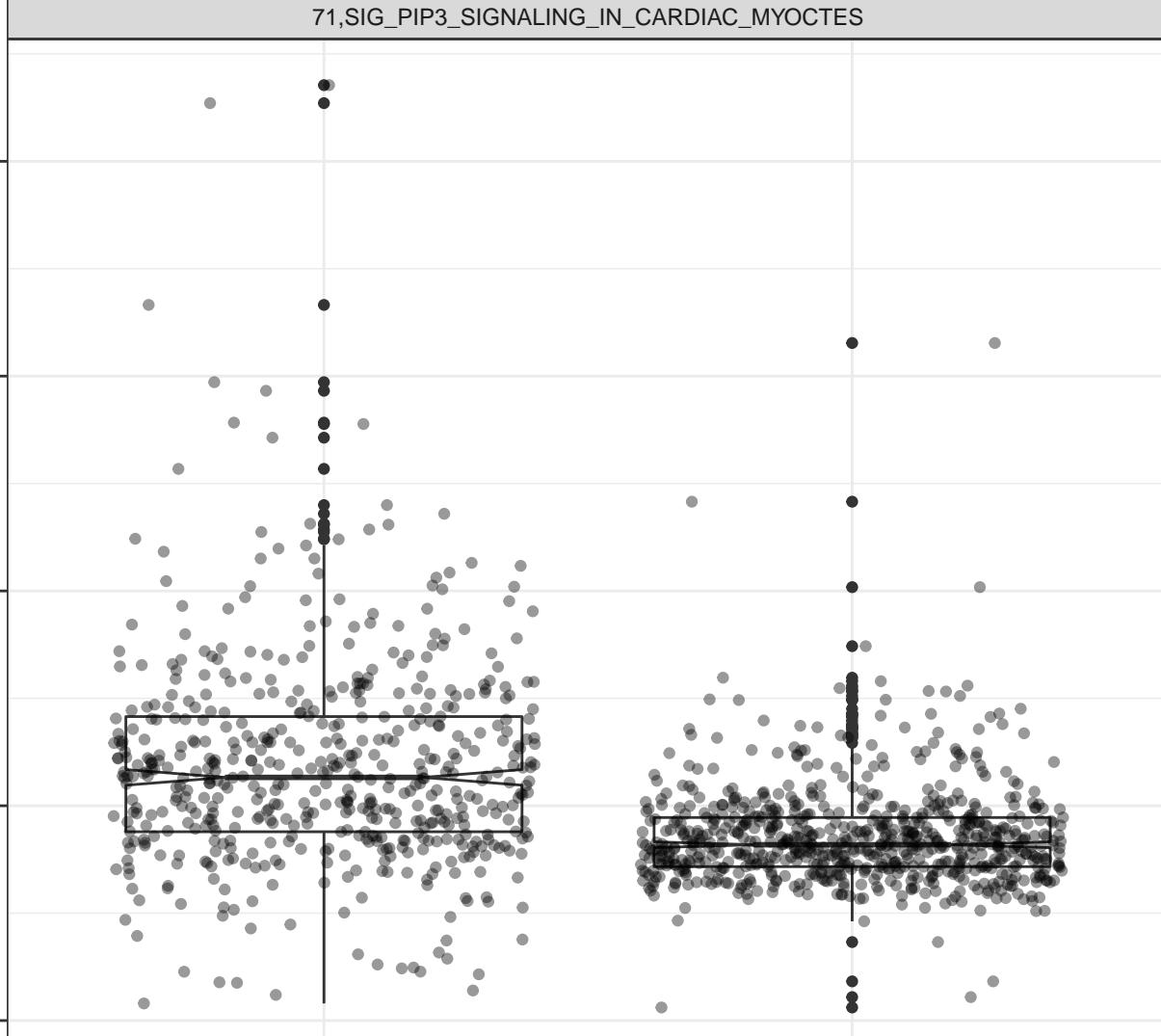
0.00

-0.05

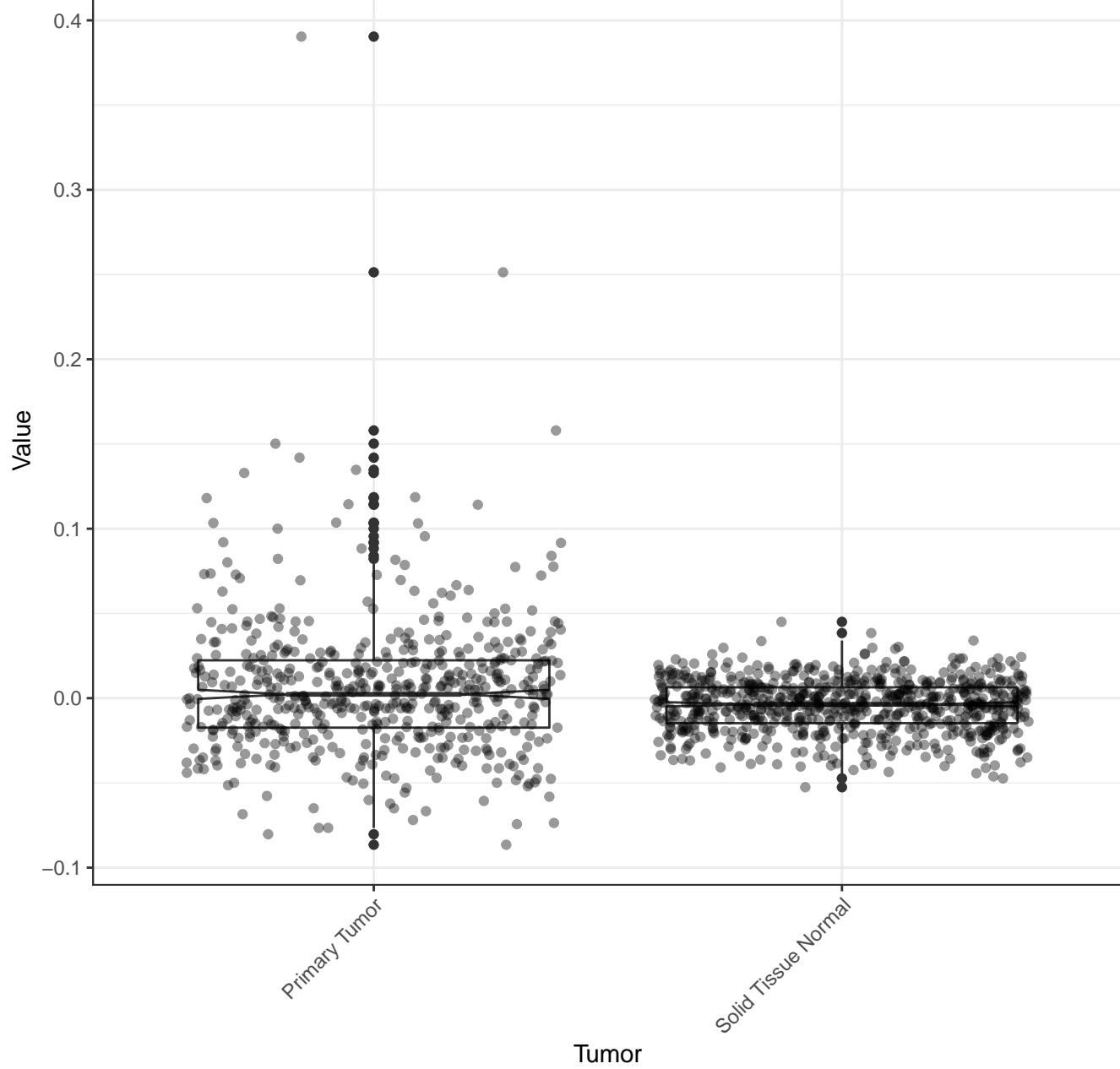
Primary Tumor

Solid Tissue Normal

Tumor



72,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES



73,REACTOME_GLYCEROPHOSPHOLIPID BIOSYNTHESIS

Value

0.05

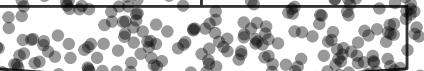
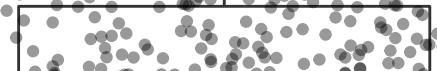
0.00

-0.05

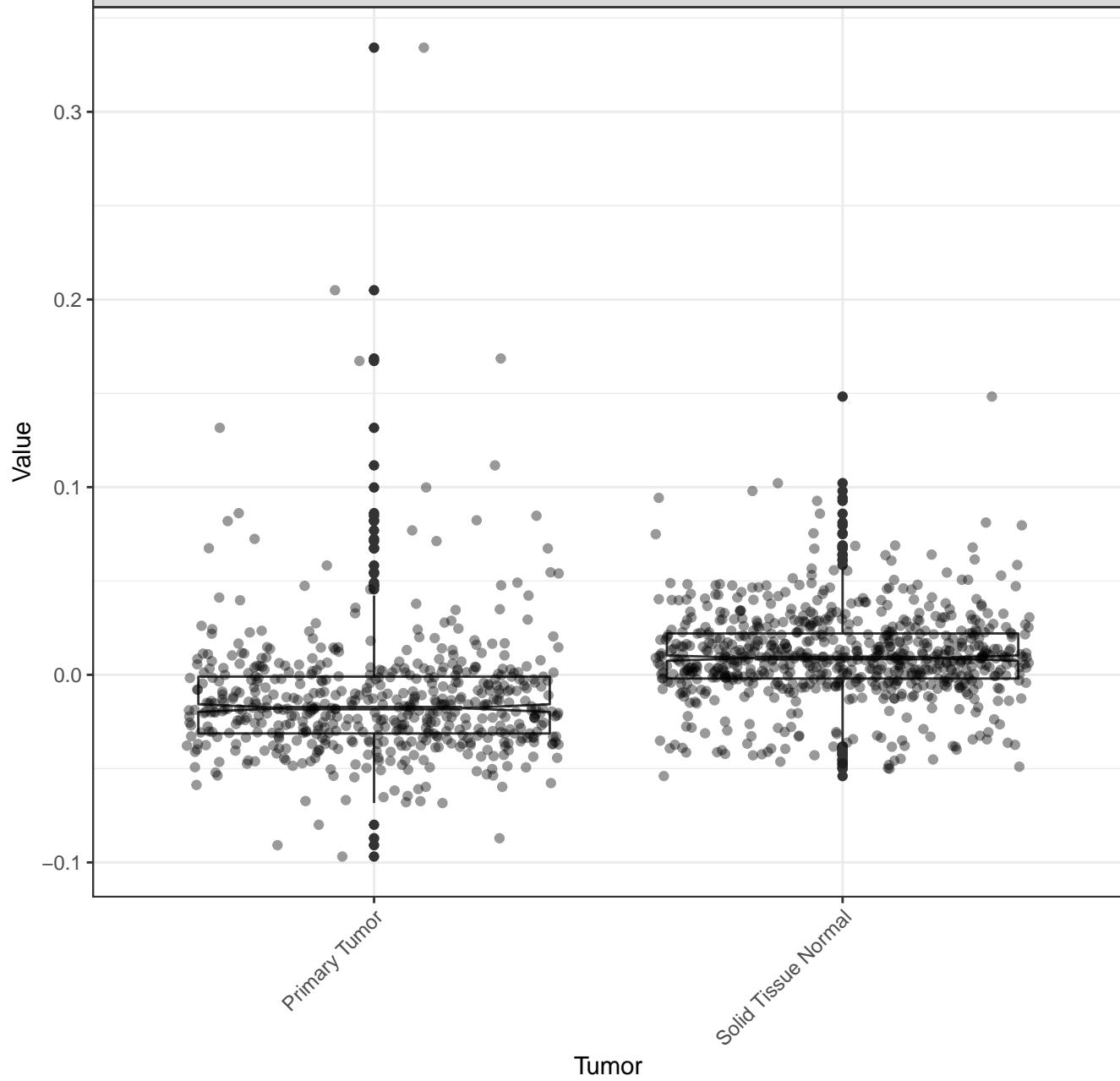
Primary Tumor

Solid Tissue Normal

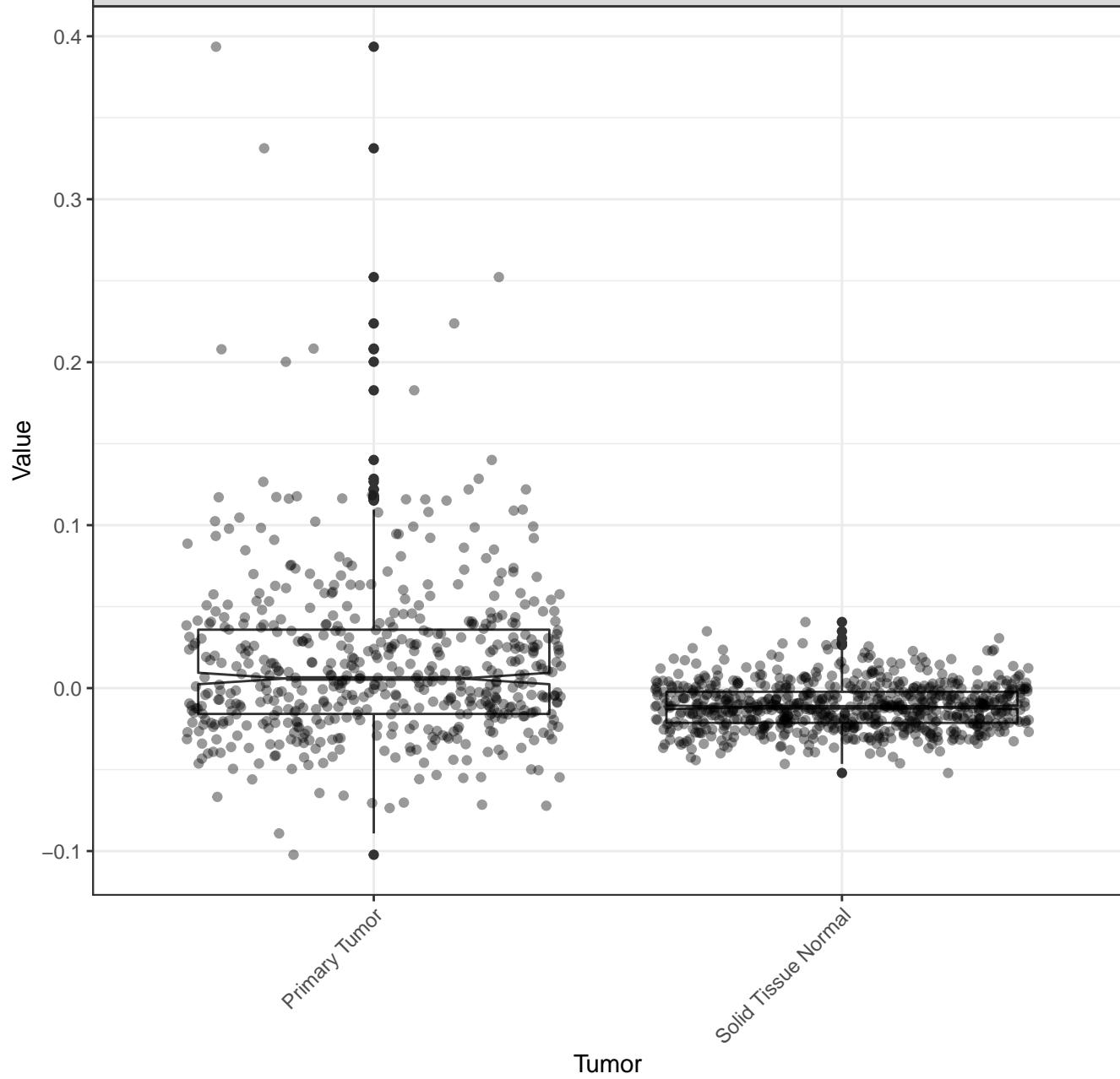
Tumor



74,KEGG_NOTCH_SIGNALING_PATHWAY



75,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

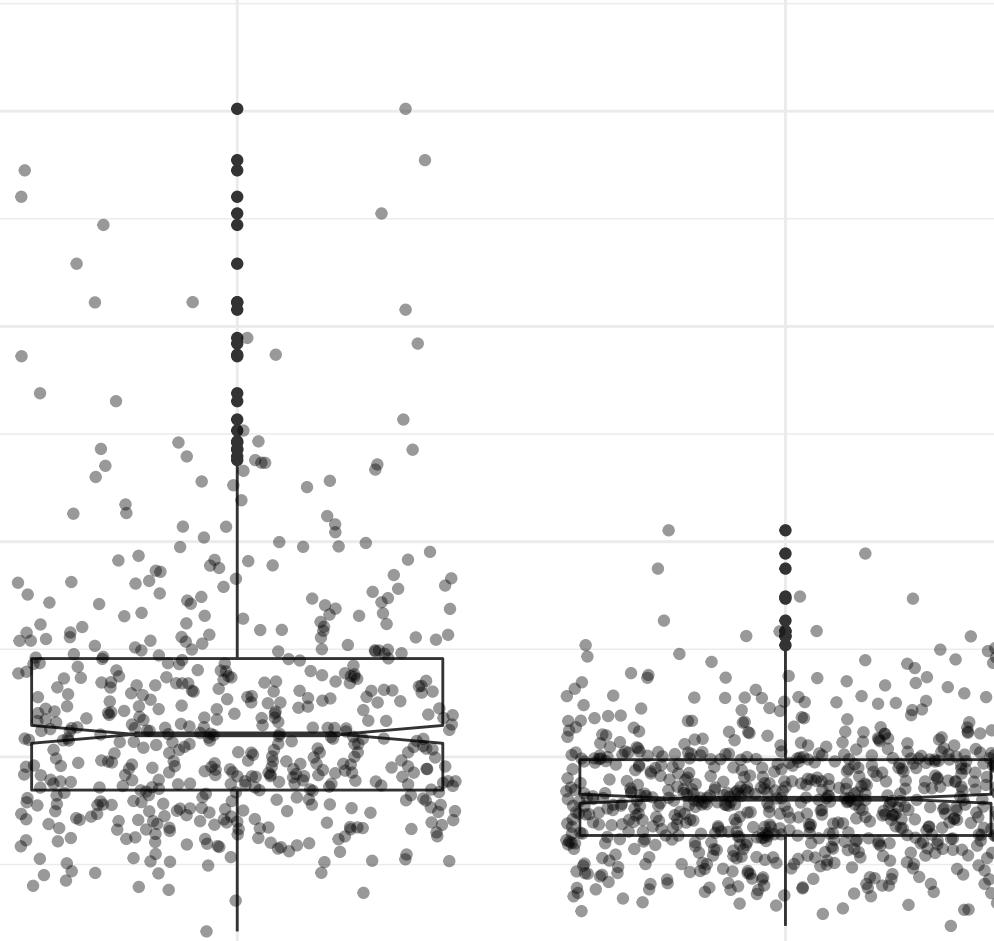


76,KEGG_CELL_CYCLE

Value

0.15
0.10
0.05
0.00
-0.05Primary Tumor
Solid Tissue Normal

Tumor



77,KEGG_METABOLISM_OF_XENOBIOTICS_BY_CYTOCHROME_P450

Value

0.4

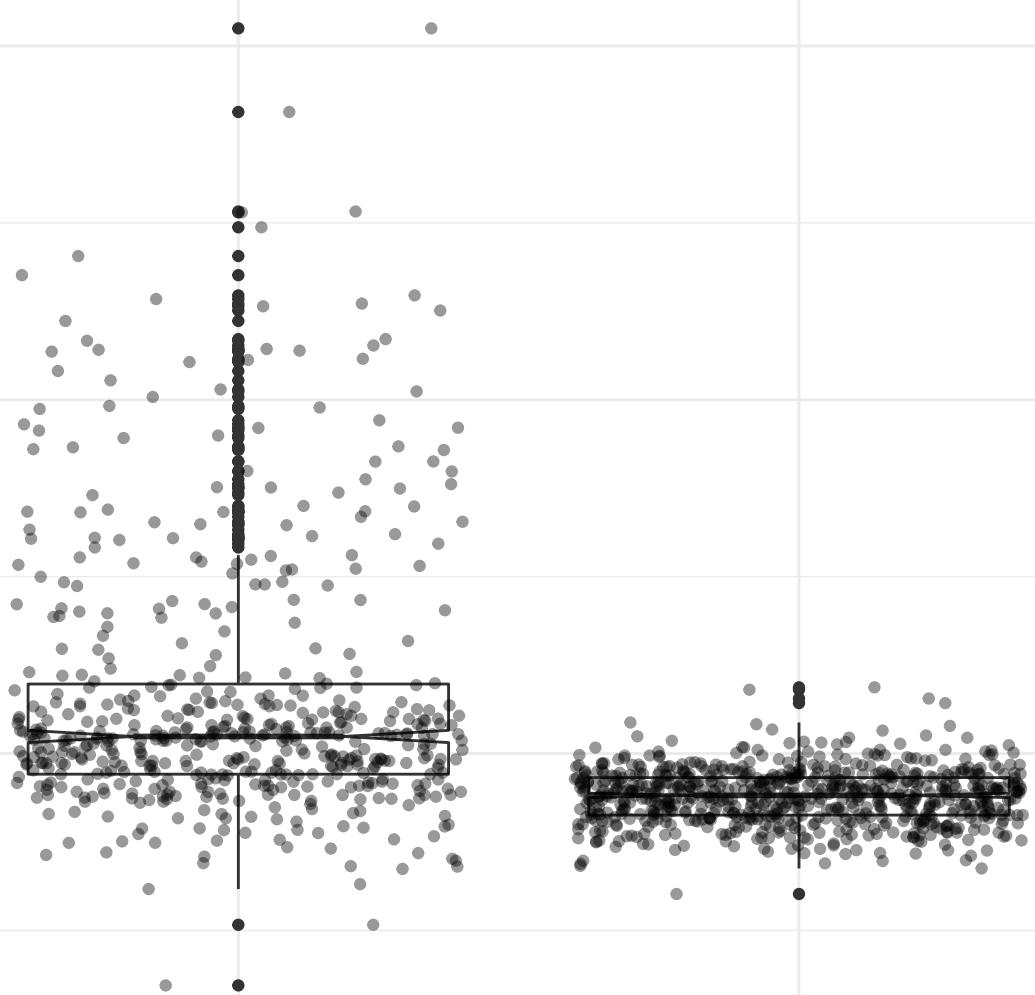
0.2

0.0

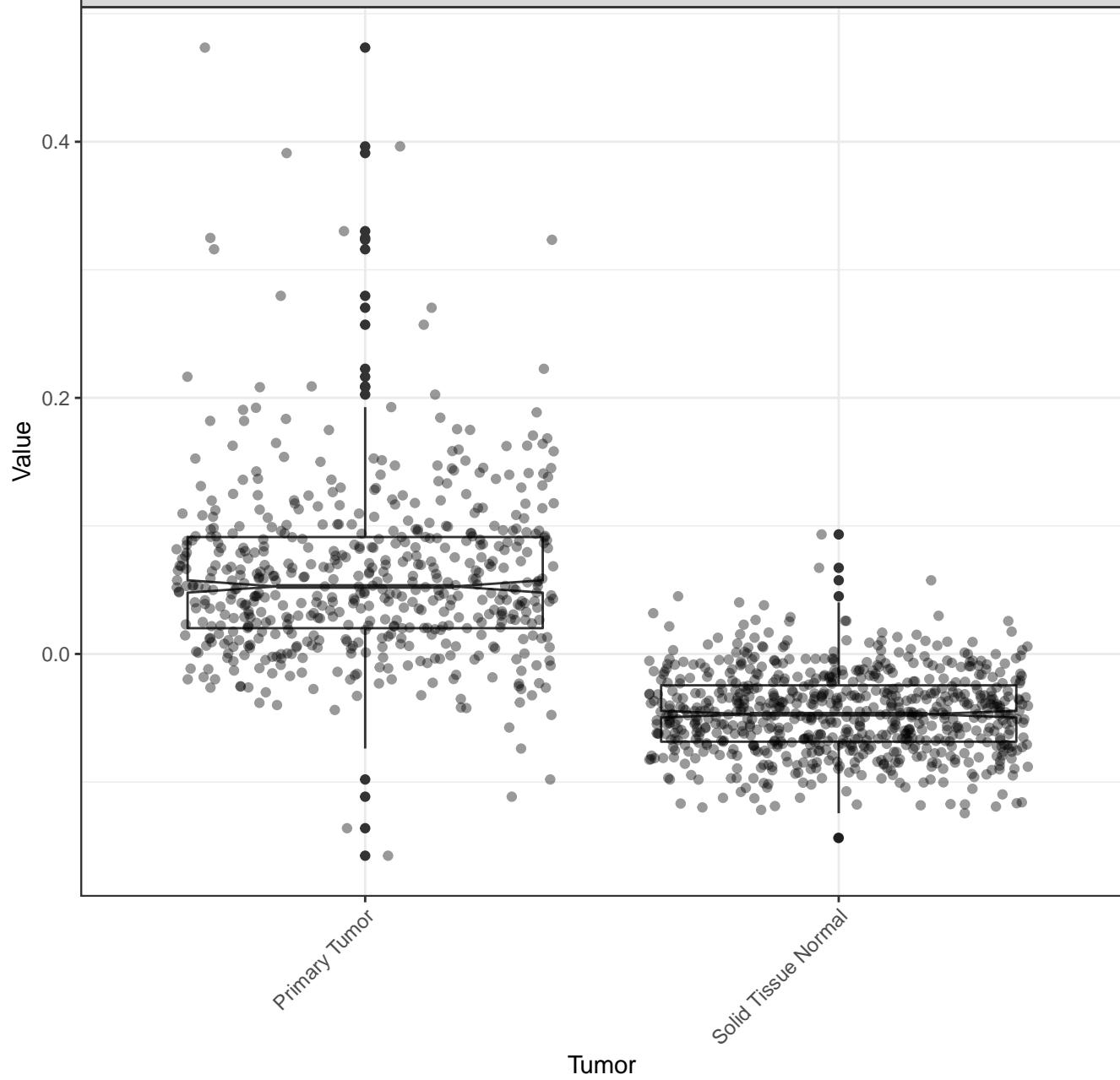
Primary Tumor

Tumor

Solid Tissue Normal



78,KEGG_TIGHT_JUNCTION



79,KEGG_METABOLISM_OF_XENOBIOTICS_BY_CYTOCHROME_P450

Value

0.2

0.1

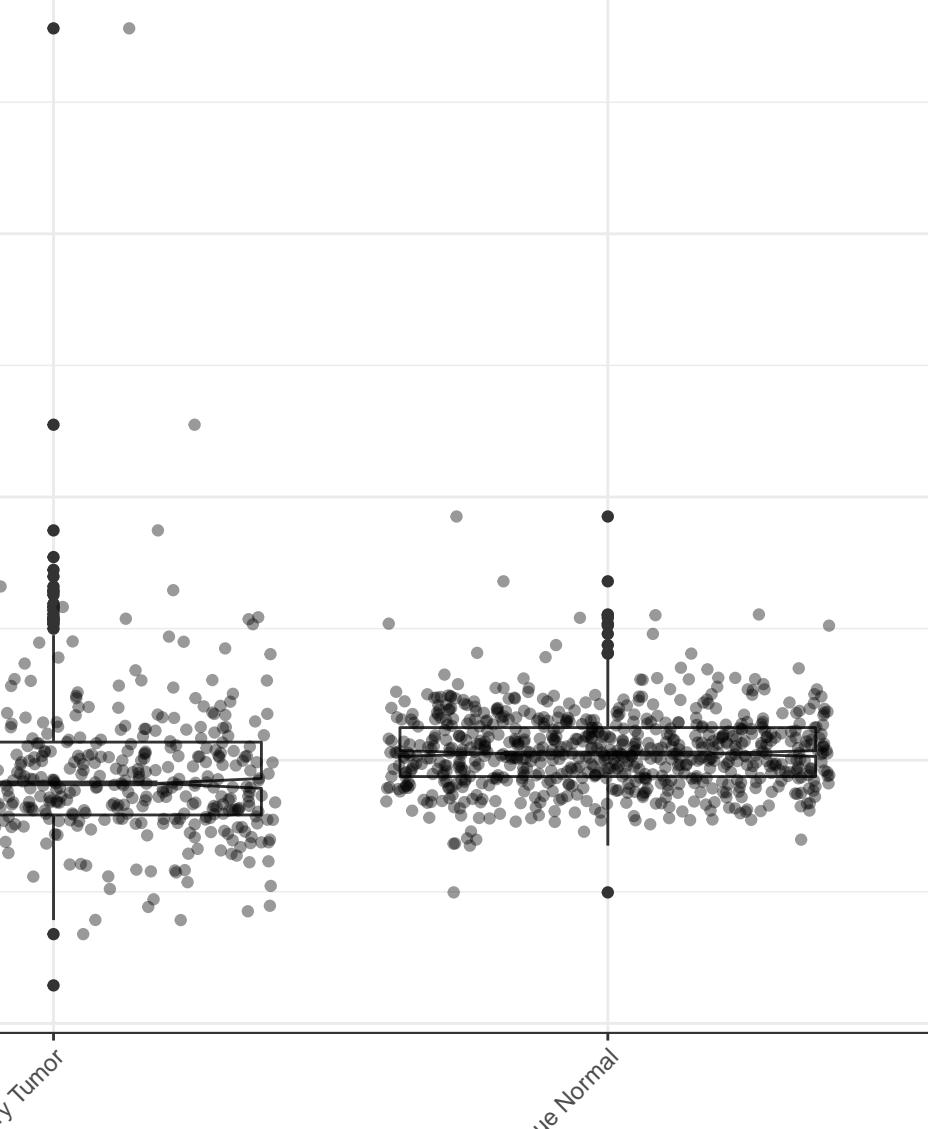
0.0

-0.1

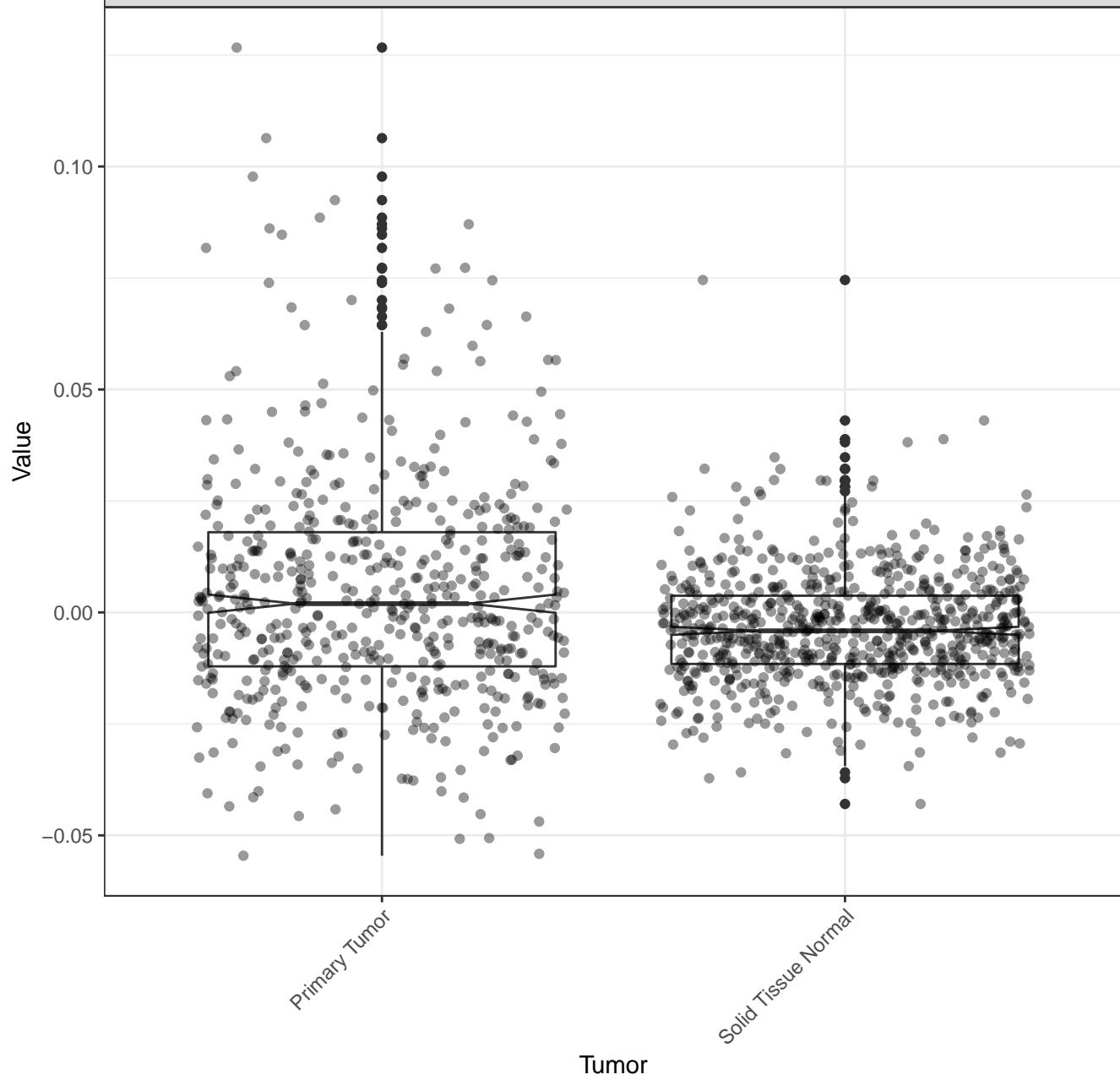
Primary Tumor

Solid Tissue Normal

Tumor



80,REACTOME_MEMBRANE_TRAFFICKING



81,MIPS_39S_RIBOSOMAL_SUBUNIT_MITOCHONDRIAL

Value

0.15

0.10

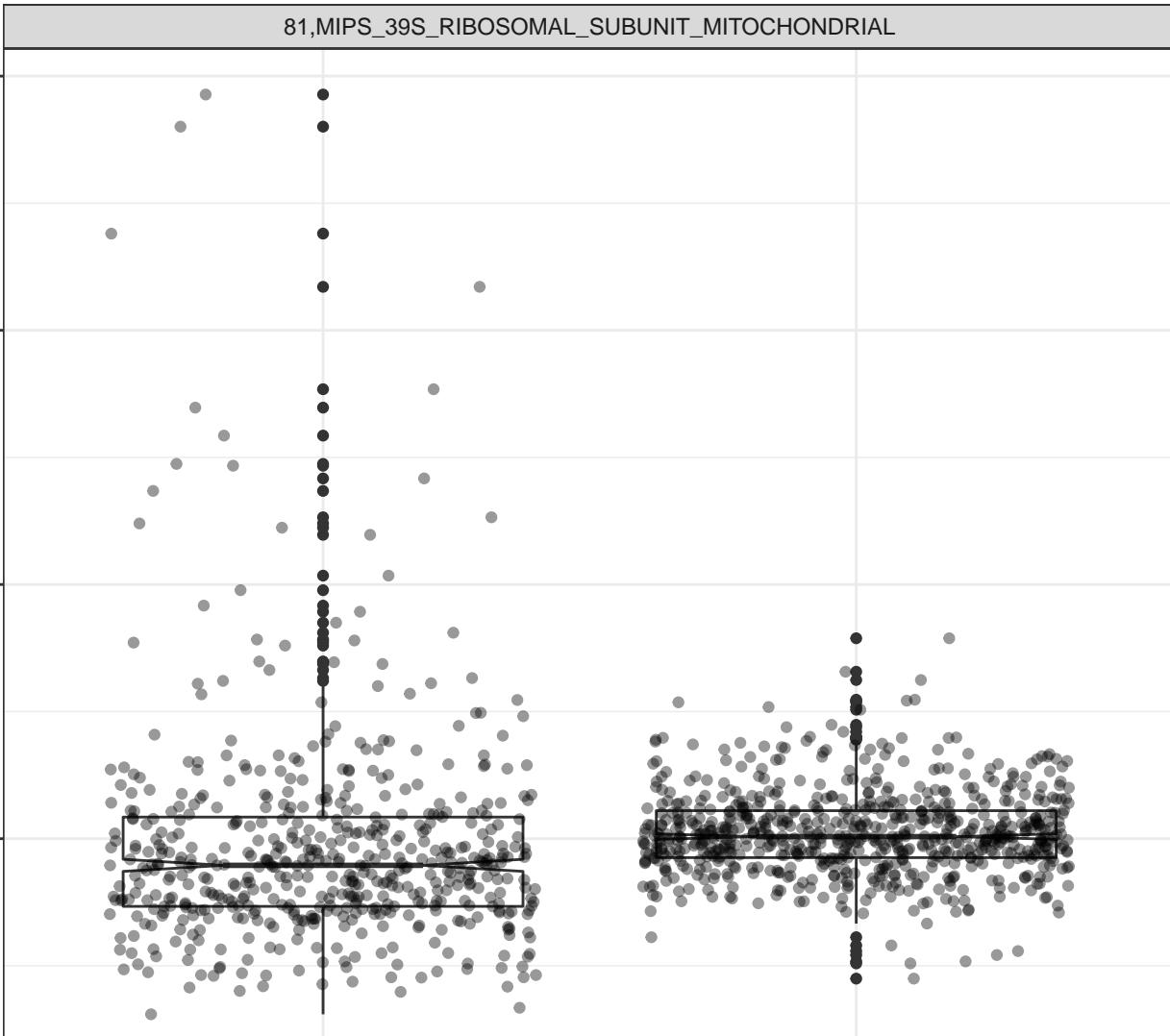
0.05

0.00

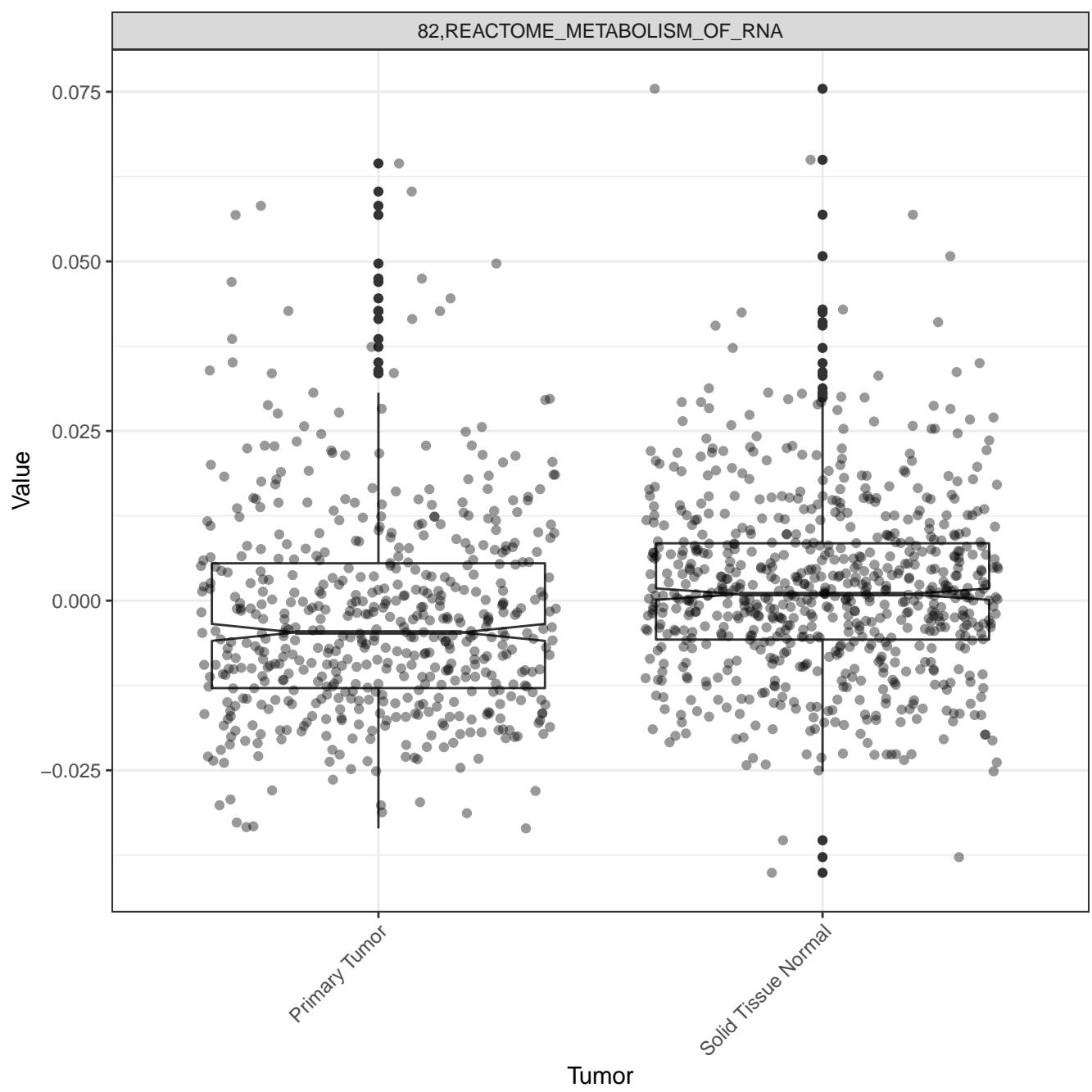
Primary Tumor

Solid Tissue Normal

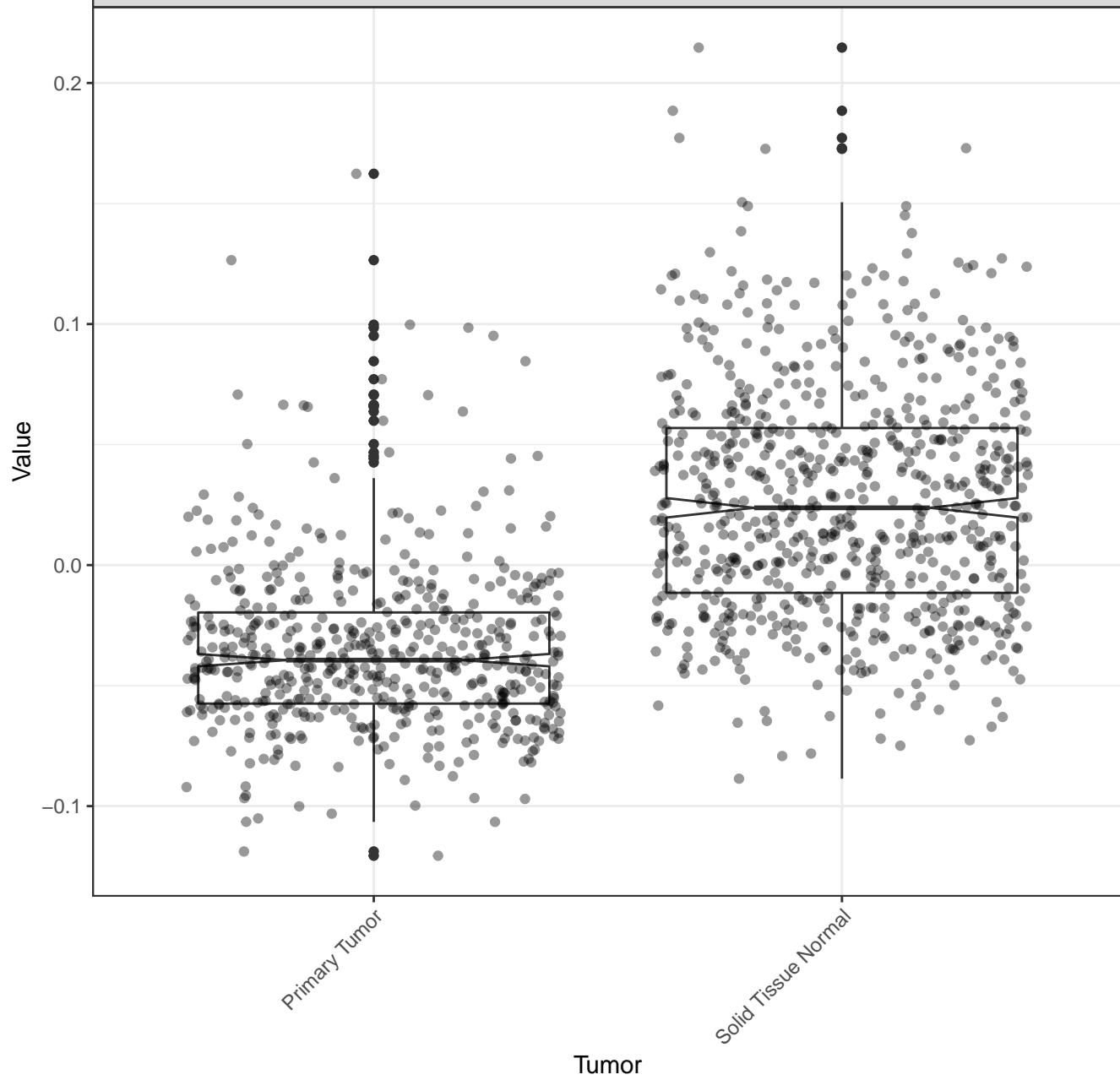
Tumor



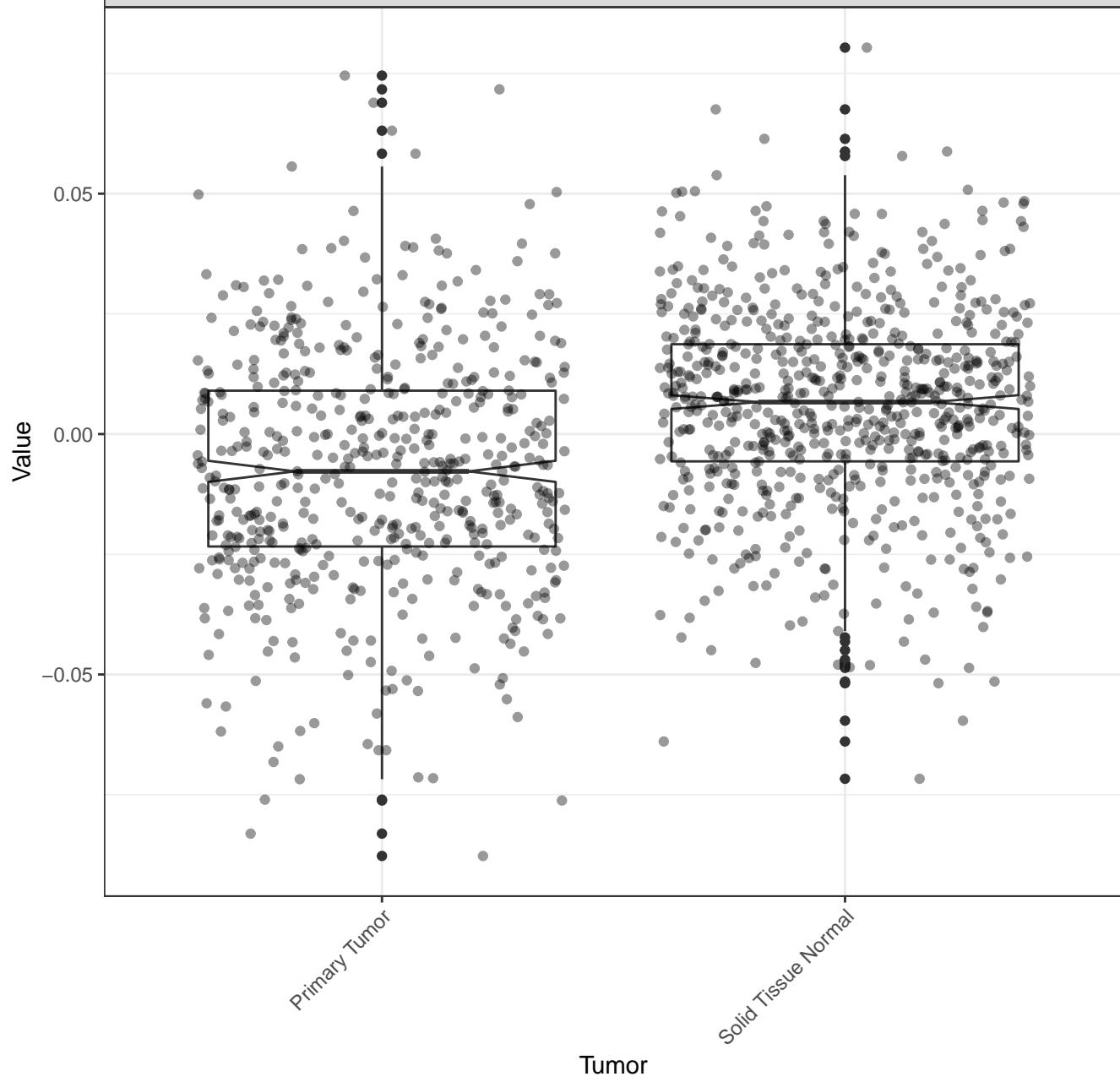
82,REACTOME_METABOLISM_OF_RNA



83,SVM Monocytes



84,REACTOME_RNA_POL_III_TRANSCRIPTION



85,REACTOME_CLASS_A1_RHODOPSIN_LIKE_RECEPTEORS

Value

0.05

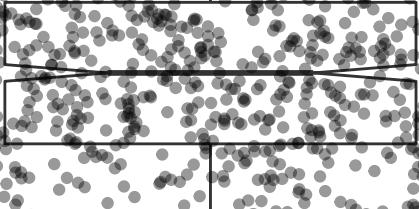
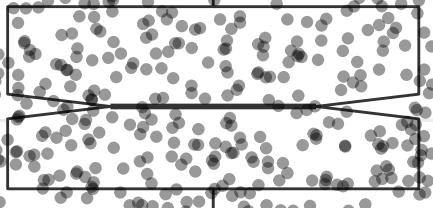
0.00

-0.05

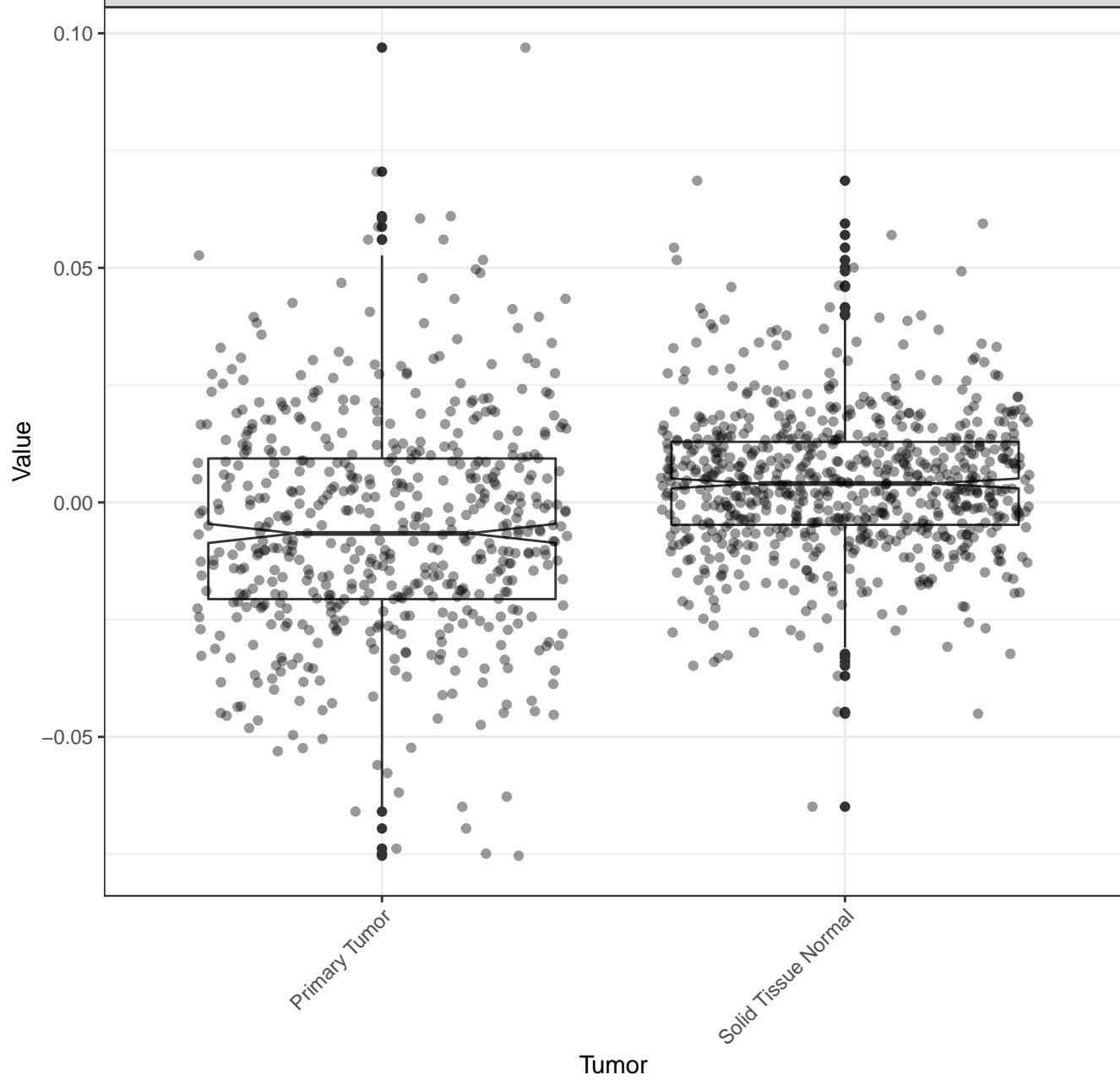
Primary Tumor

Solid Tissue Normal

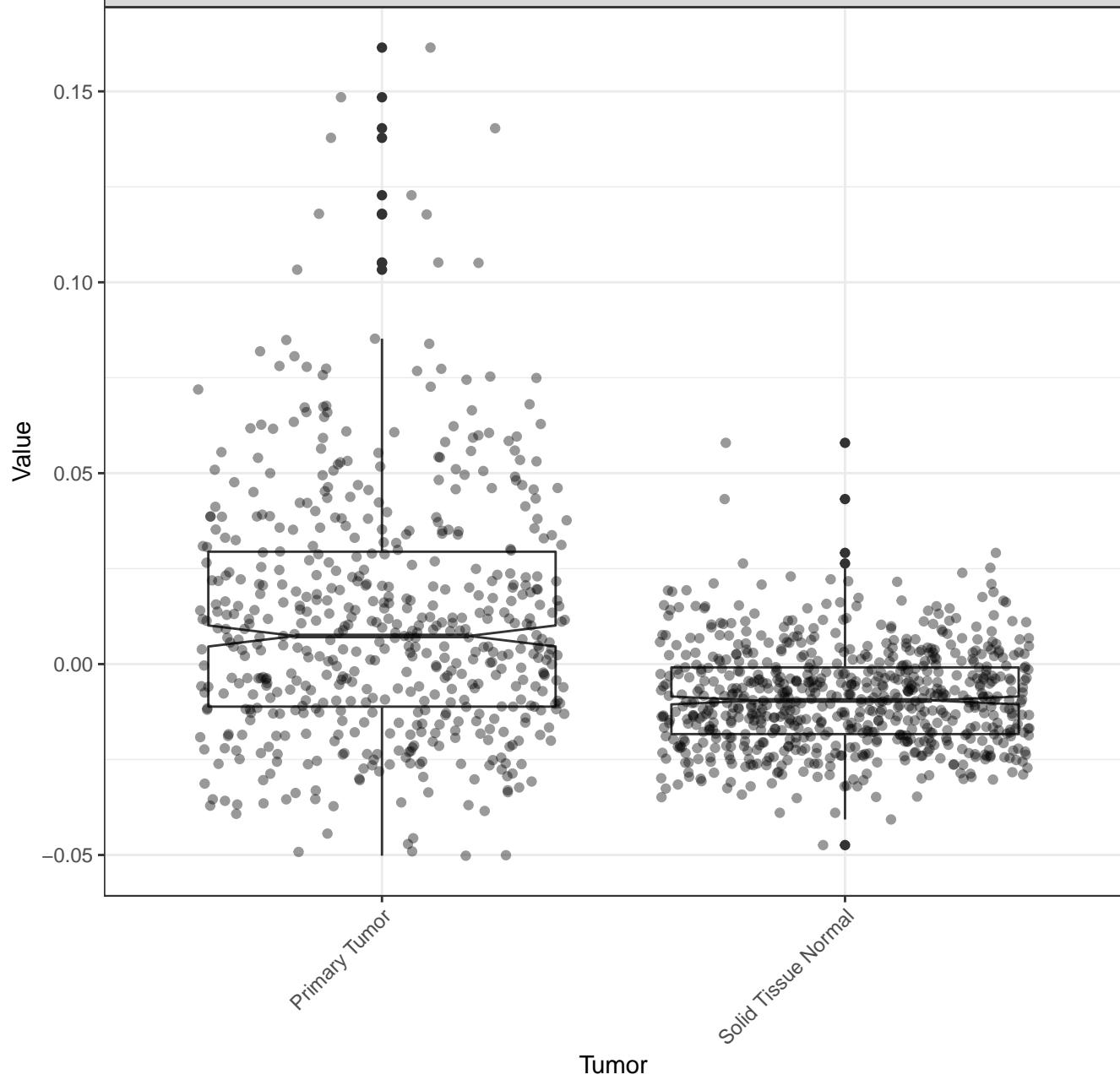
Tumor



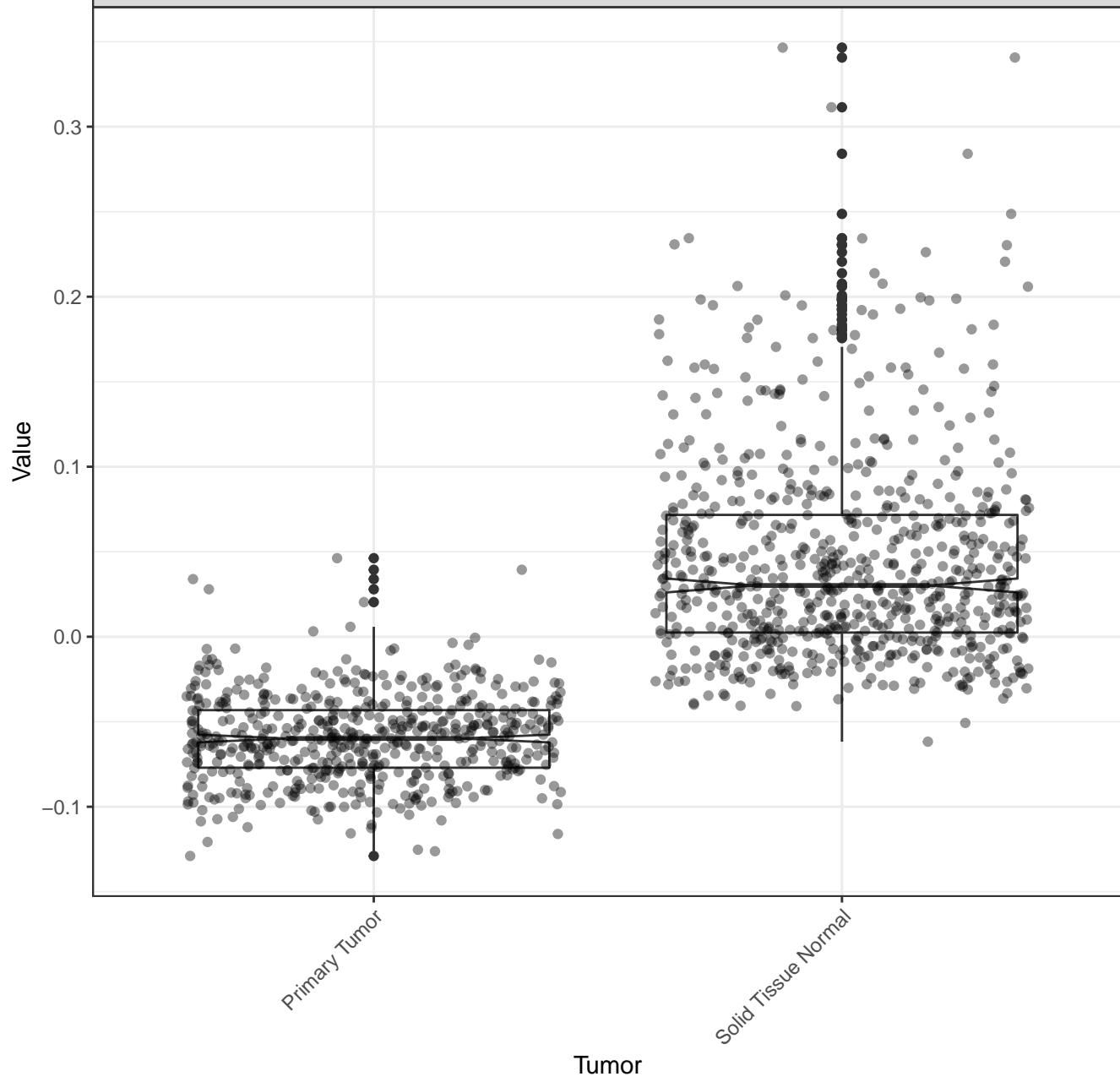
86,REACTOME_RNA_POL_II_PRE_TRANSCRIPTION_EVENTS



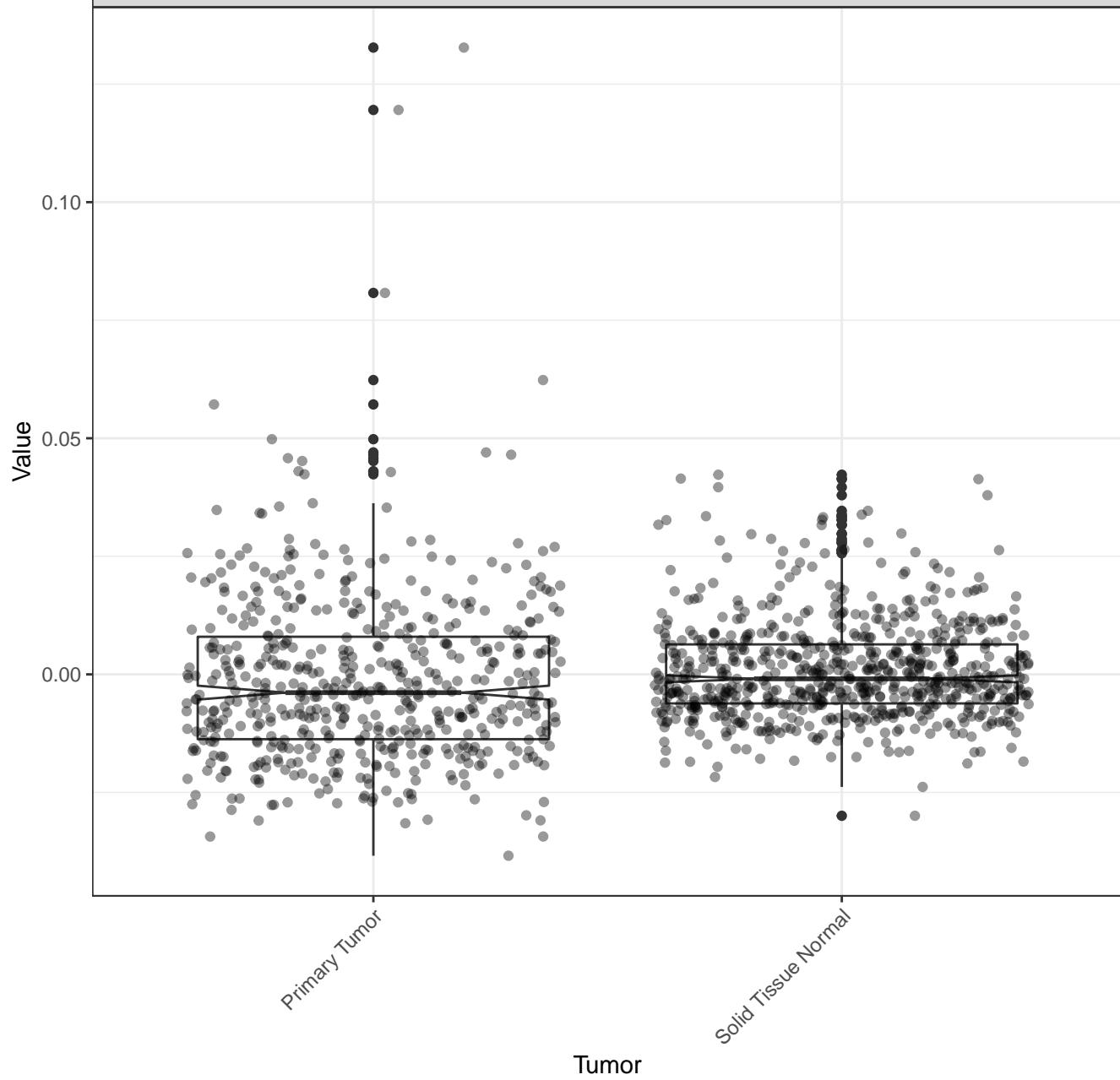
87,REACTOME_L1CAM_INTERACTIONS



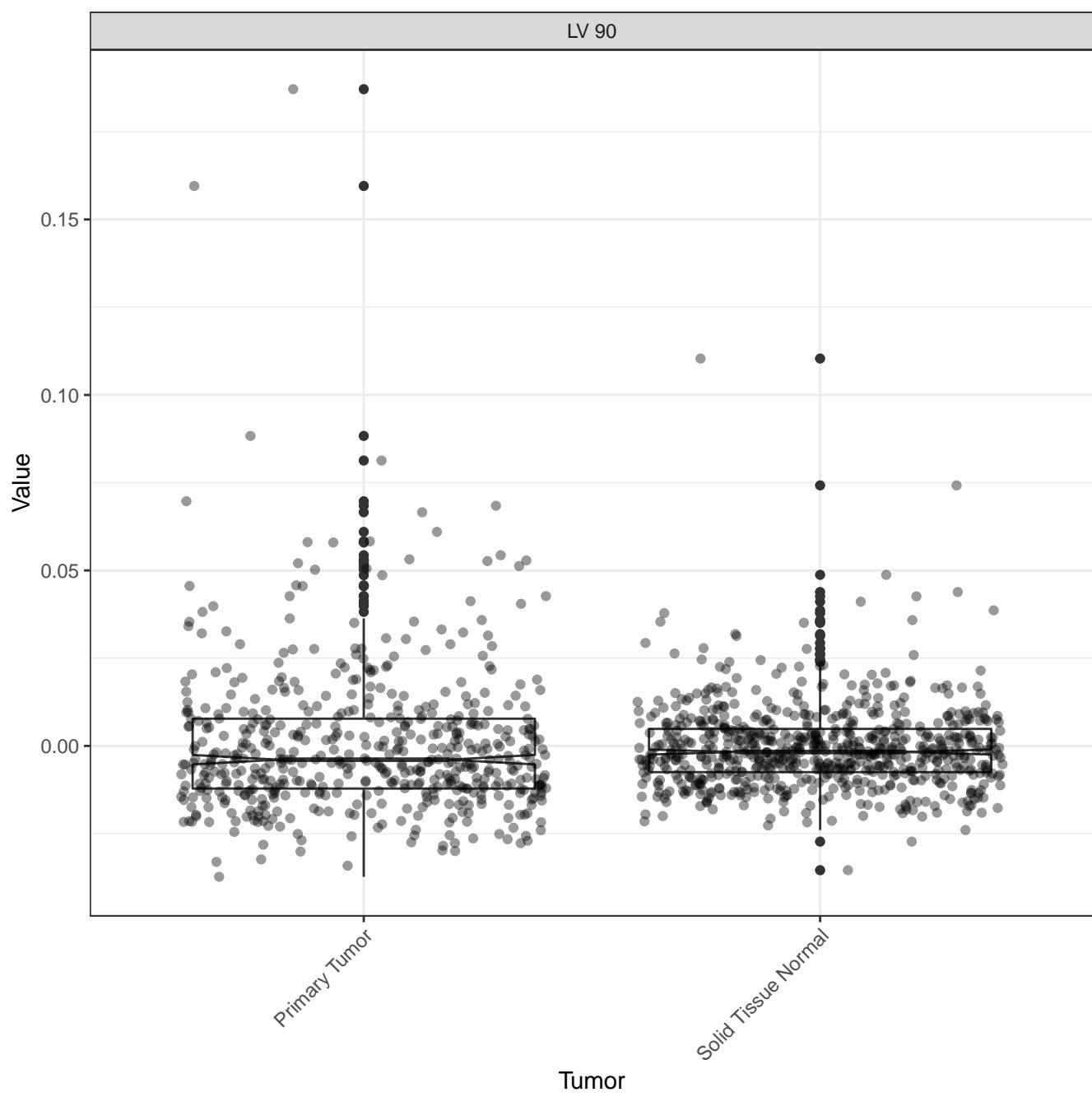
88,REACTOME_MUSCLE_CONTRACTION



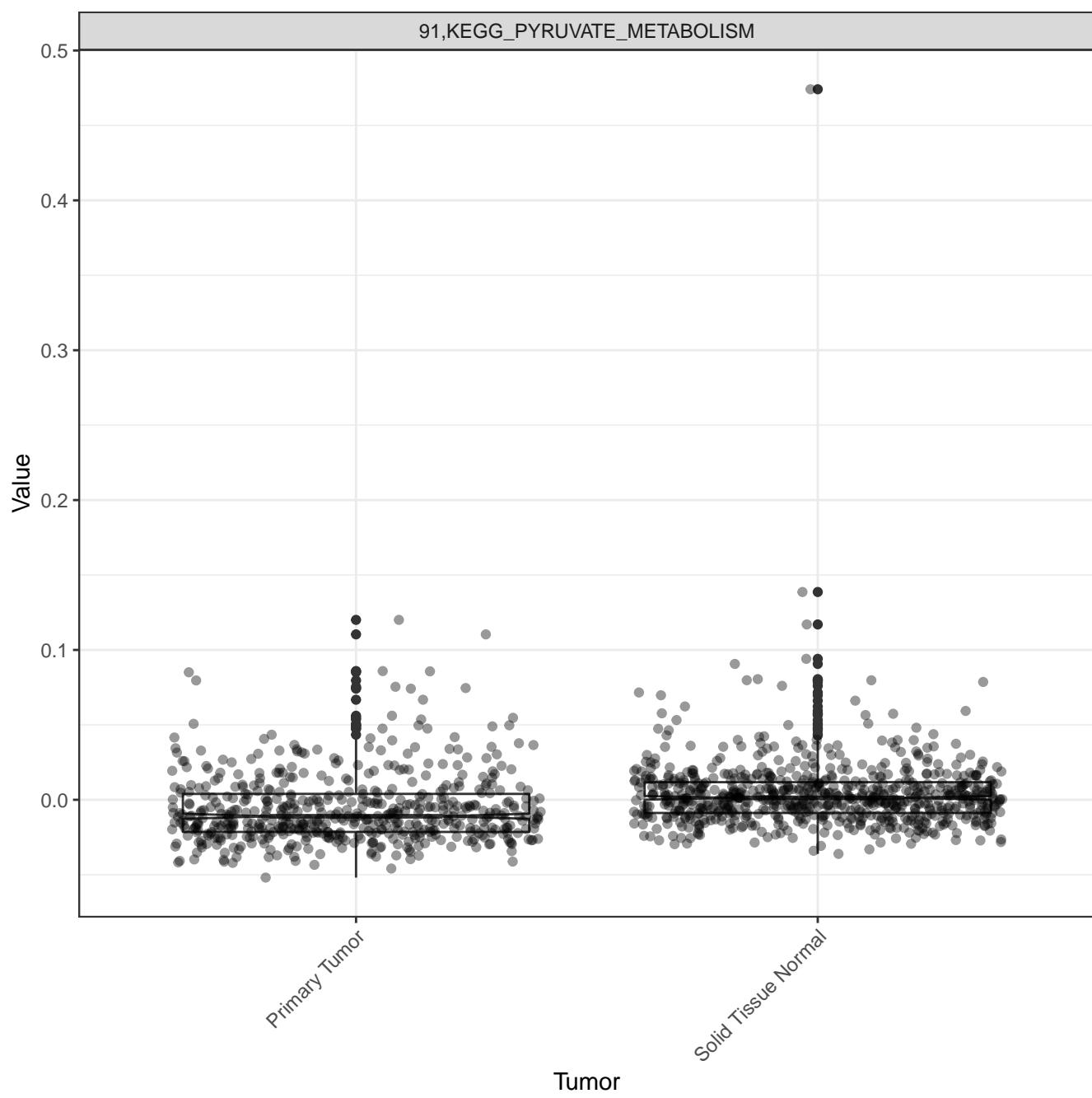
89,PID_BMPPATHWAY



LV 90



91,KEGG_PYRUVATE_METABOLISM



92,REACTOME_METABOLISM_OF_PROTEINS

Value

0.04

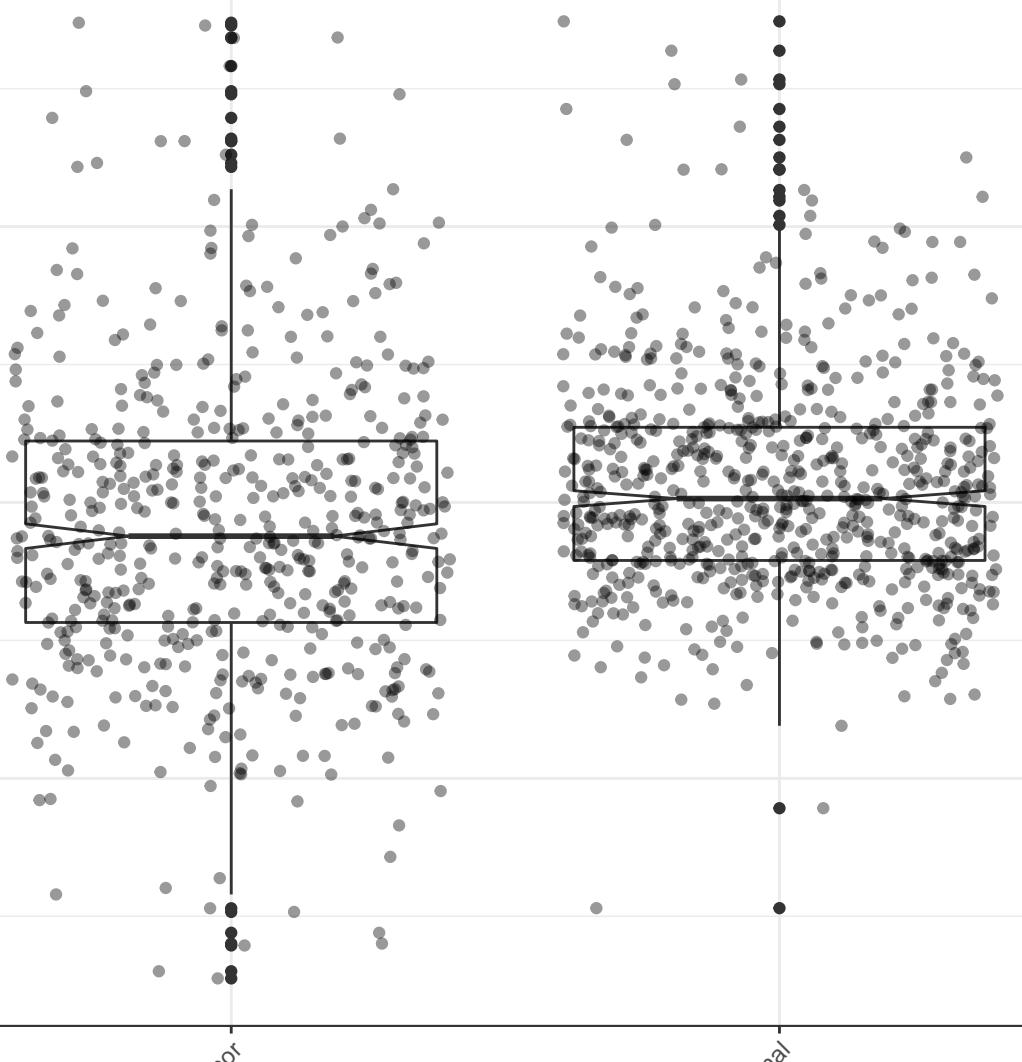
0.00

-0.04

Primary Tumor

Solid Tissue Normal

Tumor



93,REACTOME_SRP_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE

Value

0.1

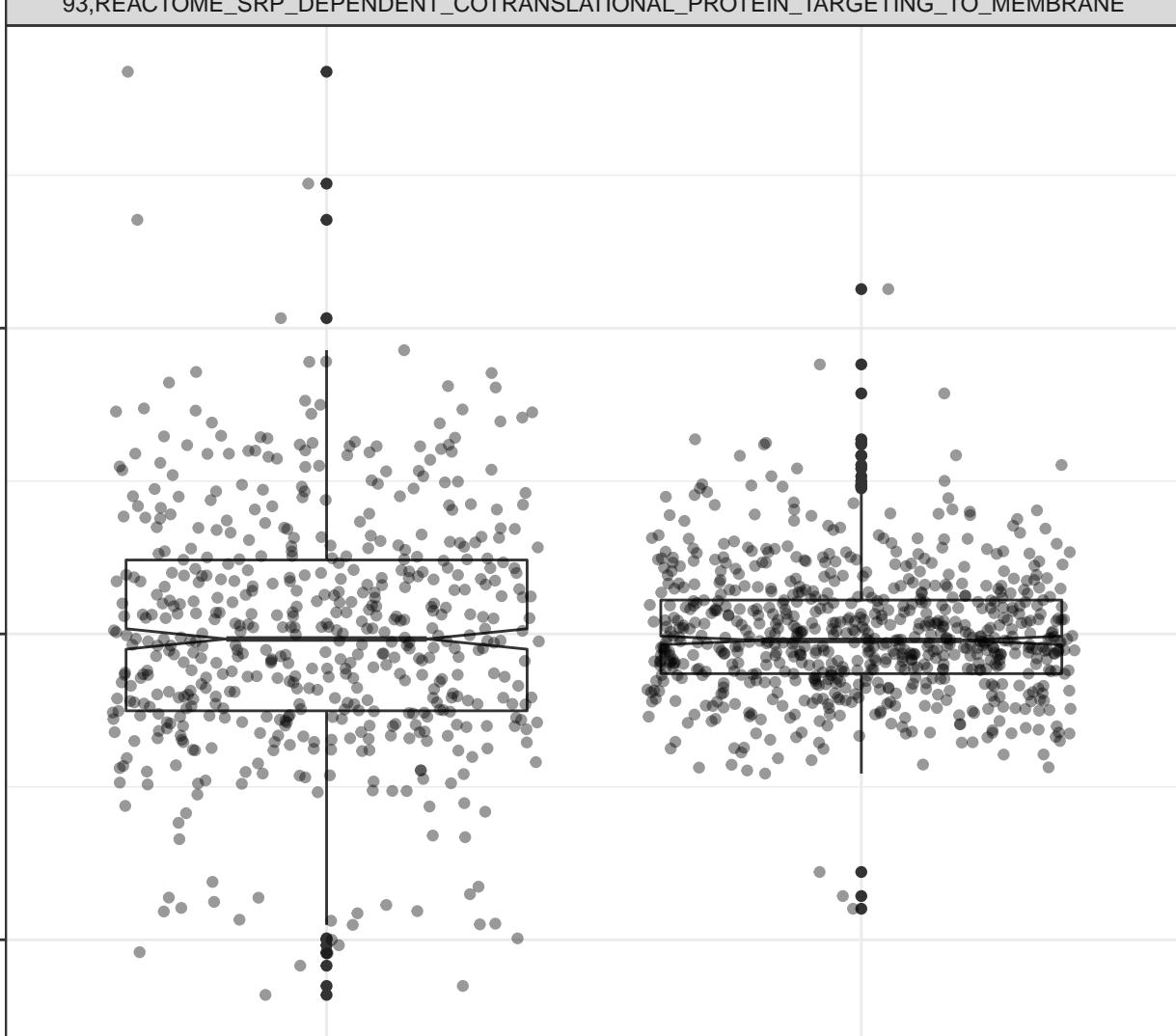
0.0

-0.1

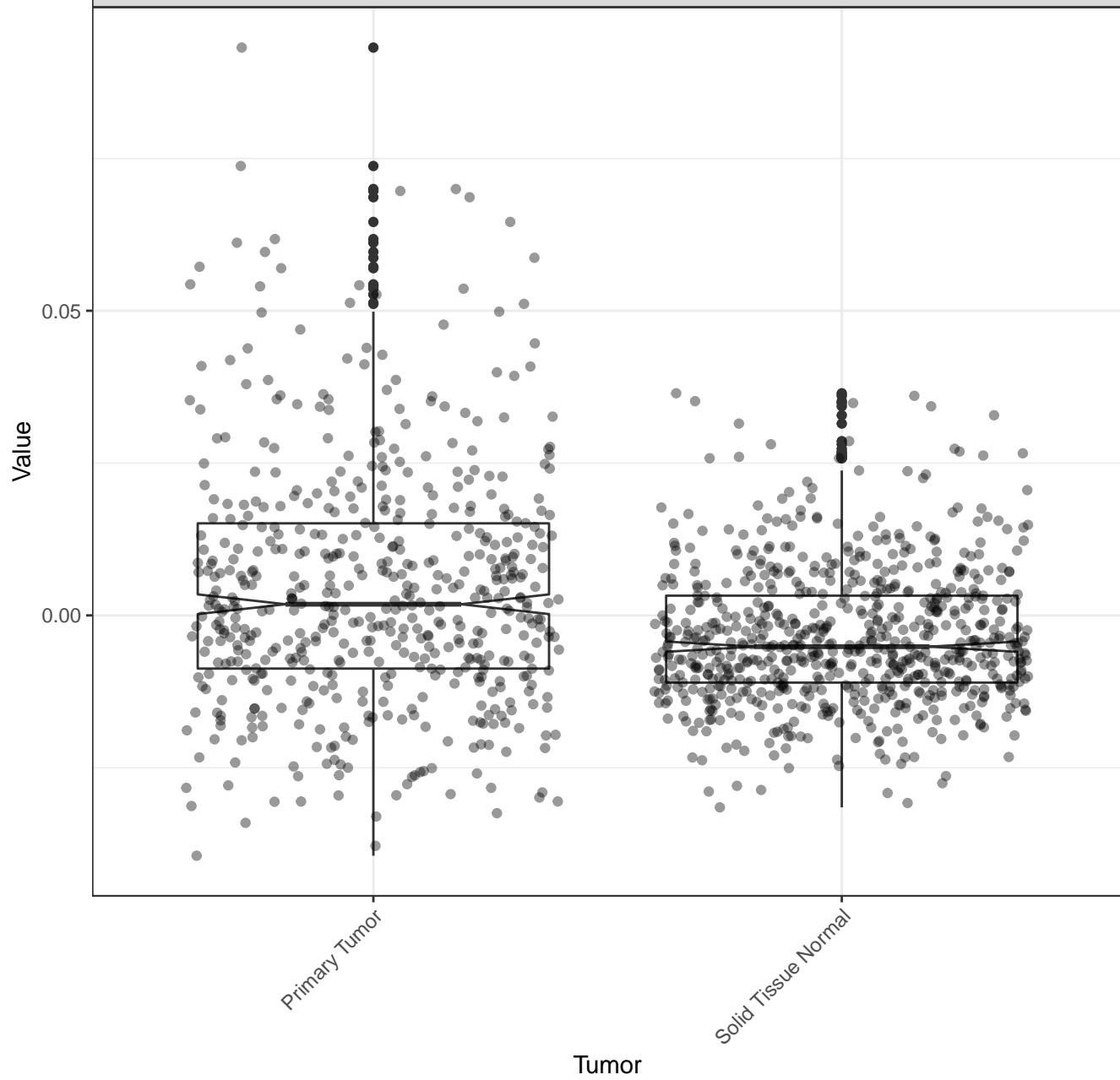
Primary Tumor

Tumor

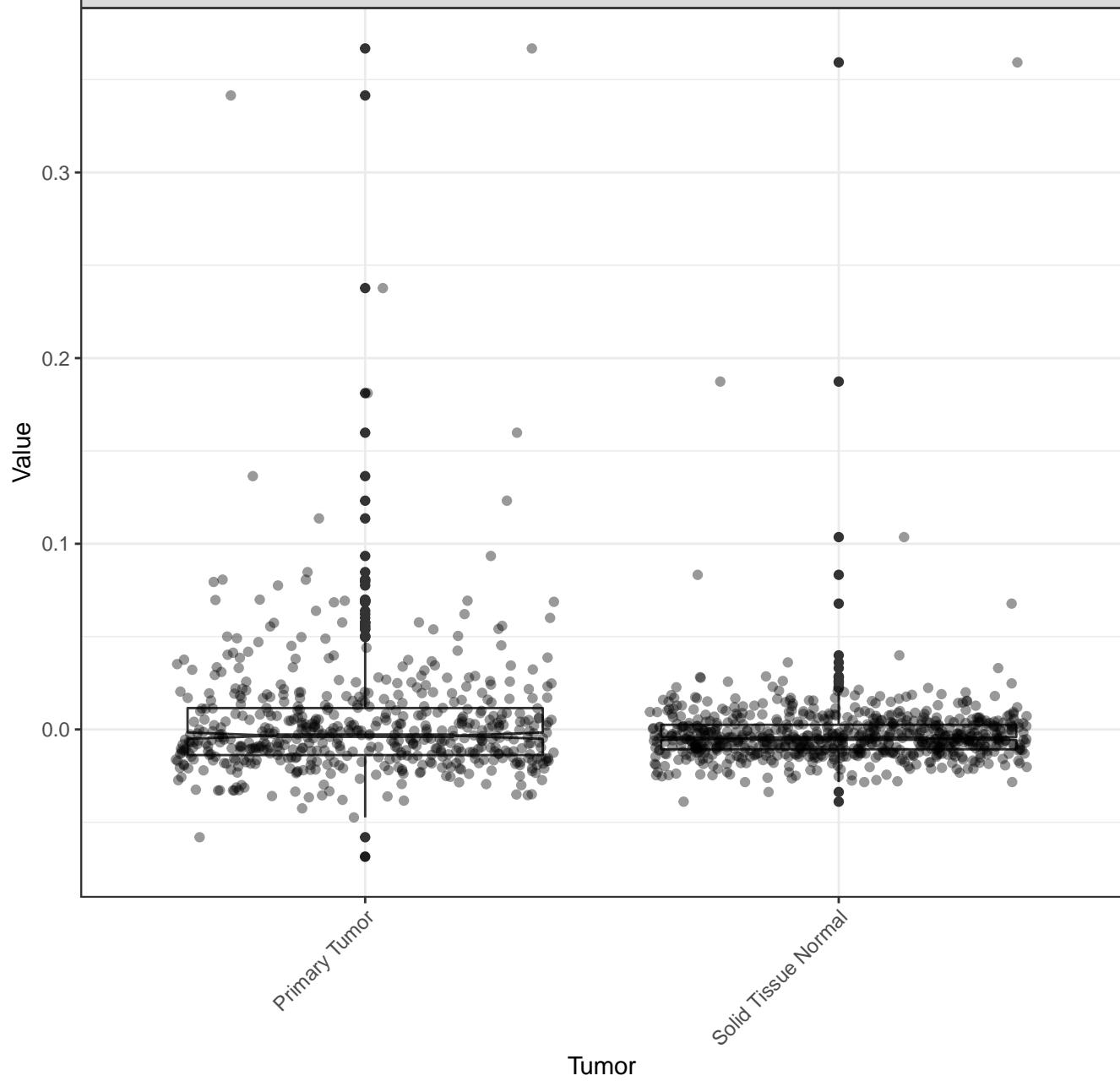
Solid Tissue Normal



94,REACTOME_MITOTIC_M_M_G1_PHASES



95,PID_DELTA_NP63PATHWAY



96,PID_AVB3_INTEGRIN_PATHWAY

Value

0.10

0.05

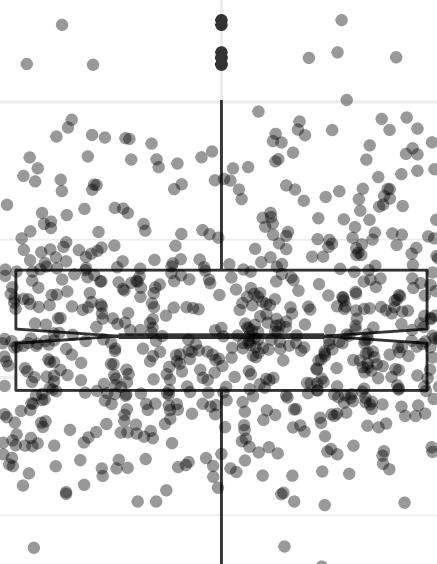
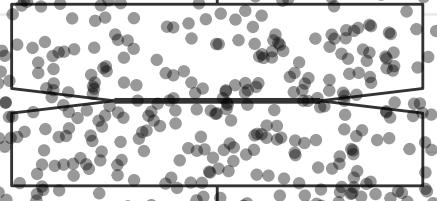
0.00

-0.05

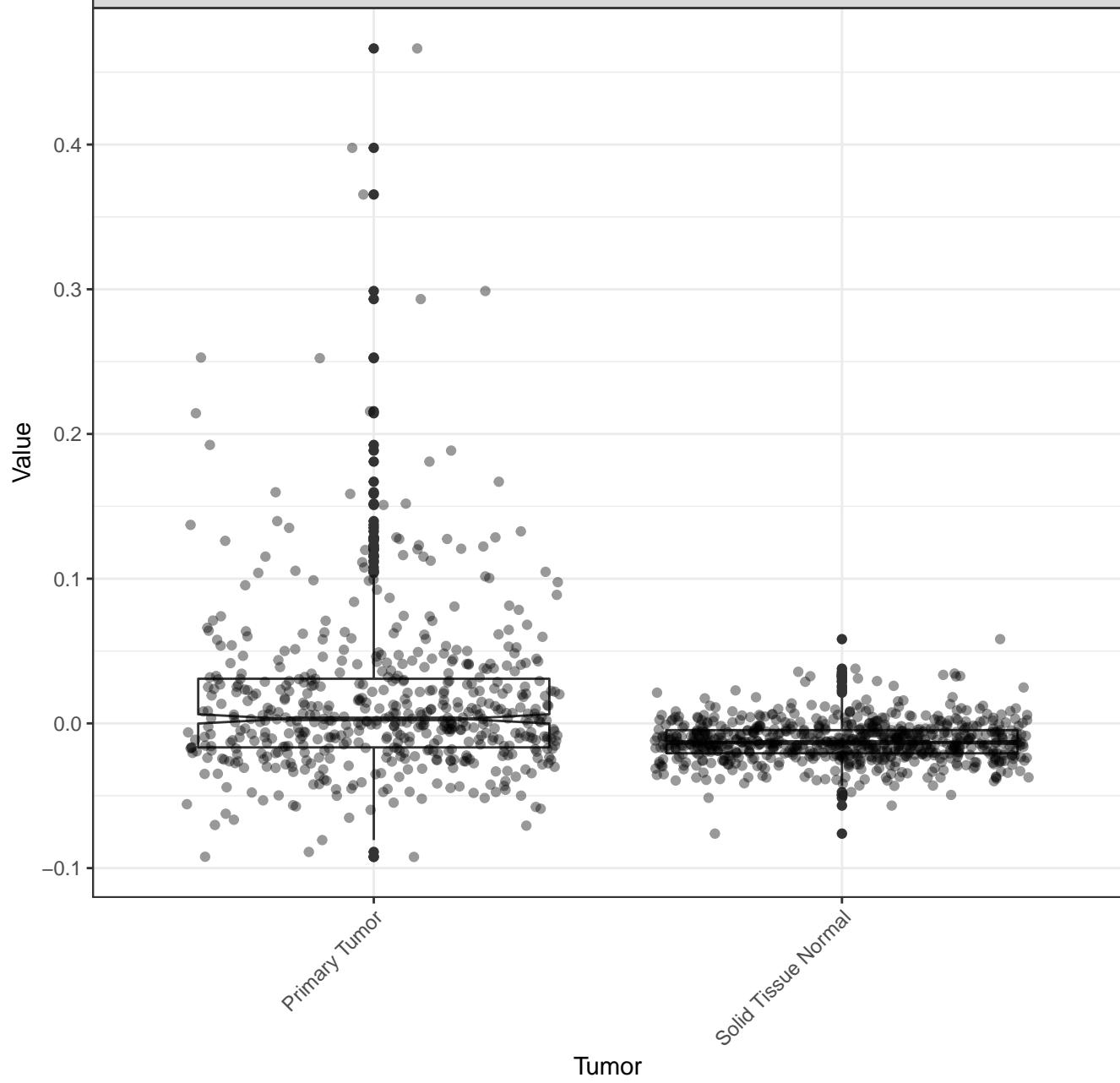
Primary Tumor

Solid Tissue Normal

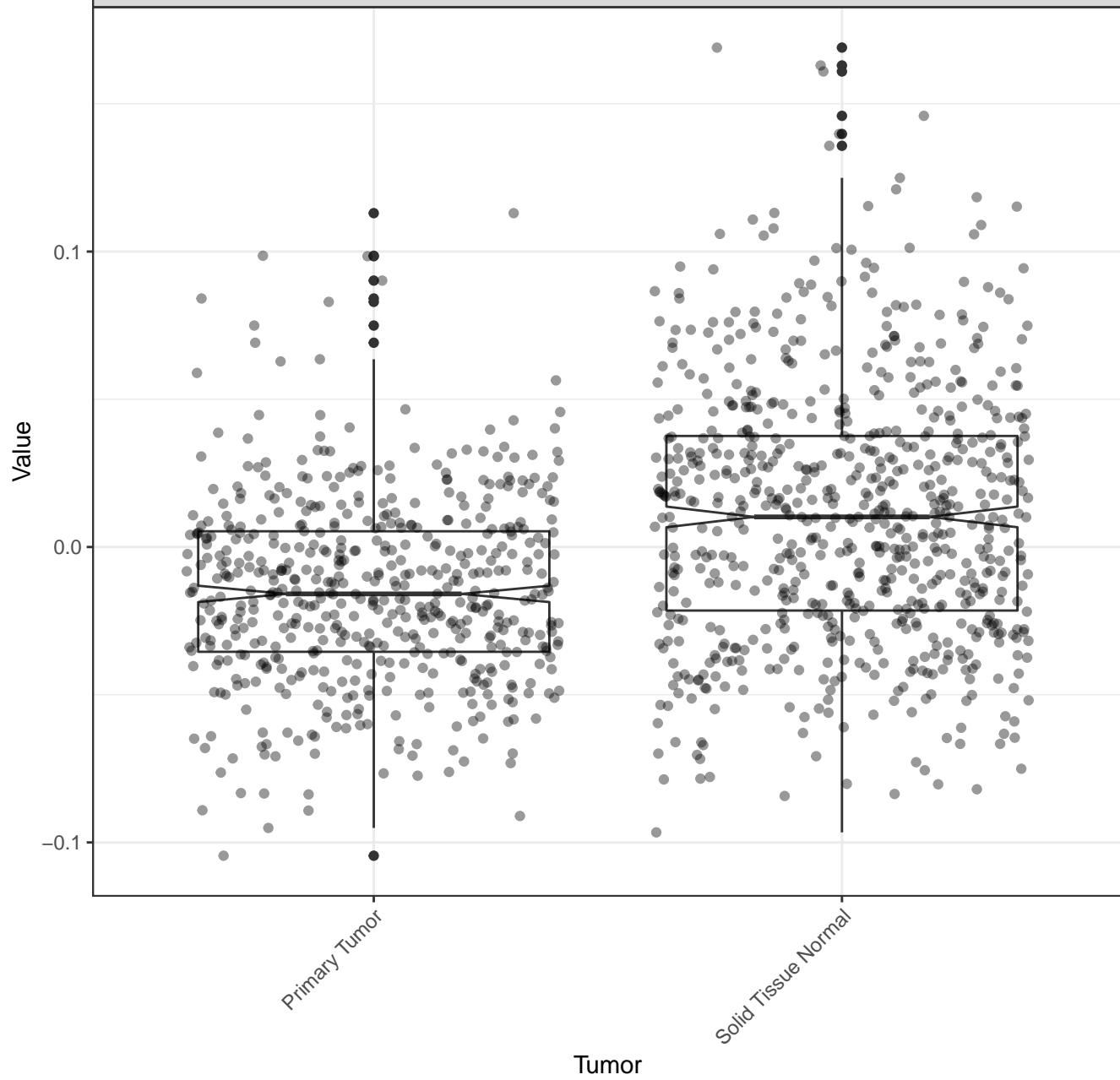
Tumor



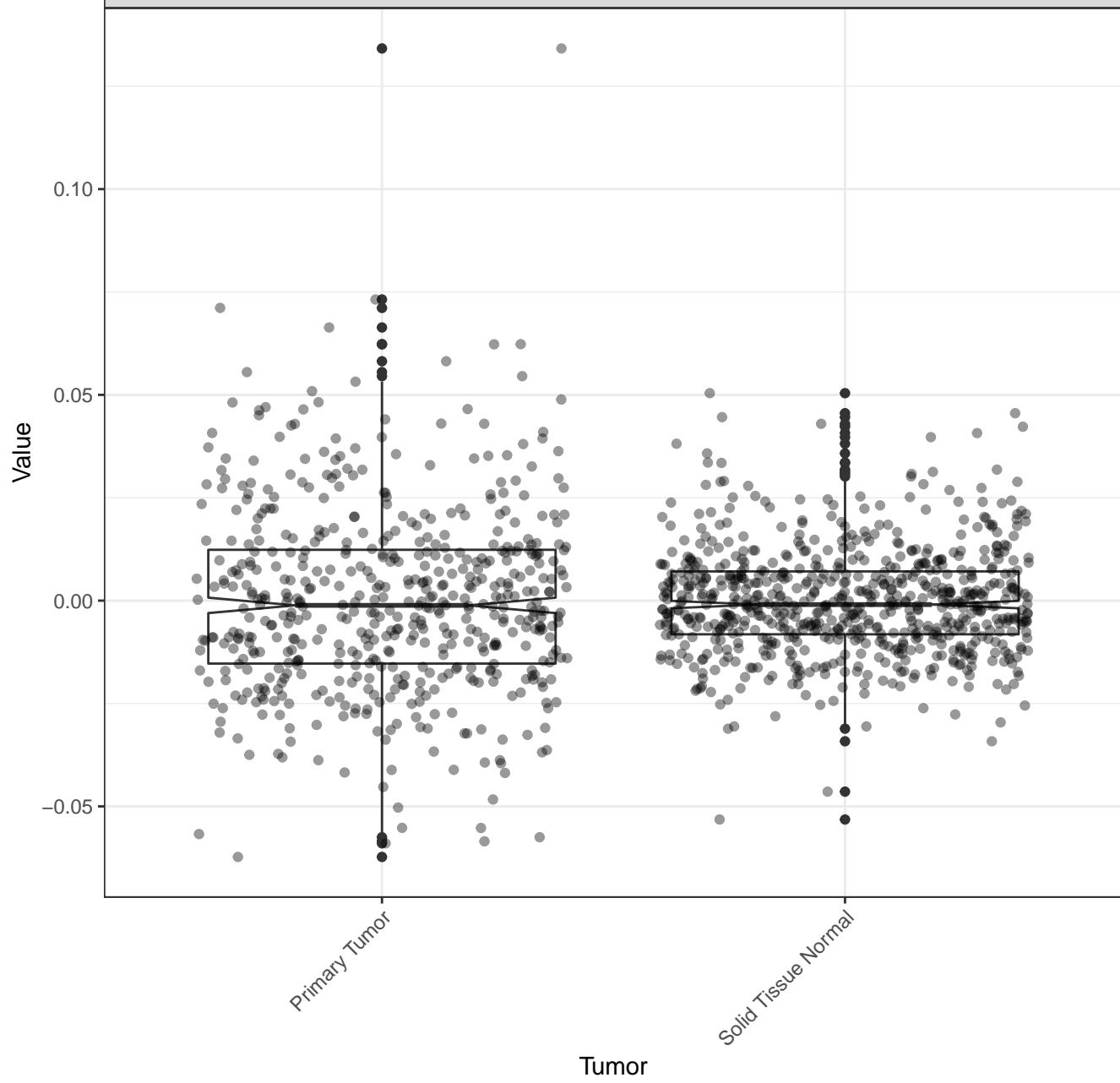
97,PID_DELTA_NP63PATHWAY



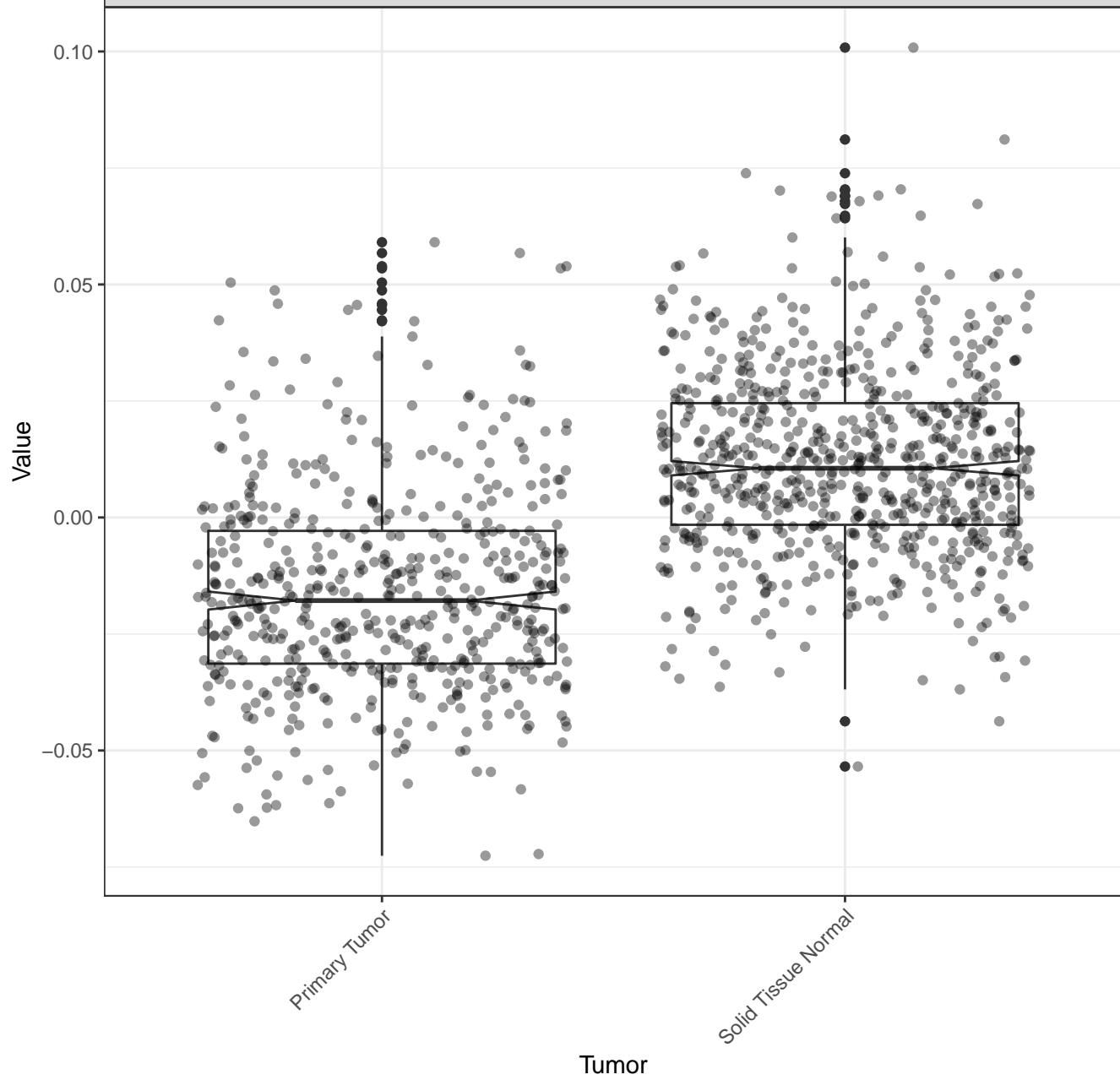
98,REACTOME_METABOLISM_OF_LIPIDS_AND_LIPOPROTEINS



99,PID_AURORA_A_PATHWAY



100,KEGG_RIG_I_LIKE_RECEPATOR_SIGNALING_PATHWAY



101,IRIS_Neutrophil–Resting

Value

0.4

0.3

0.2

0.1

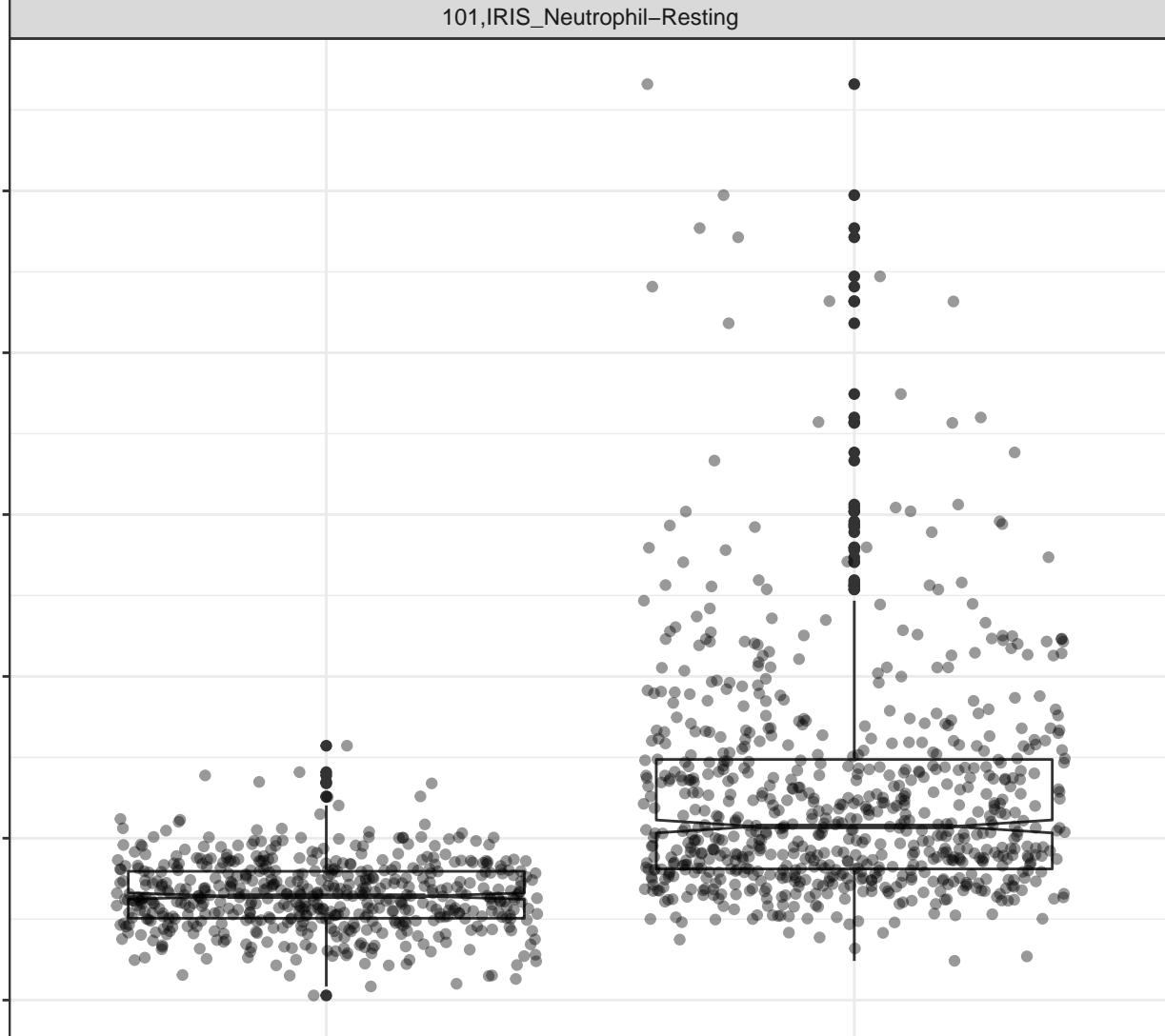
0.0

-0.1

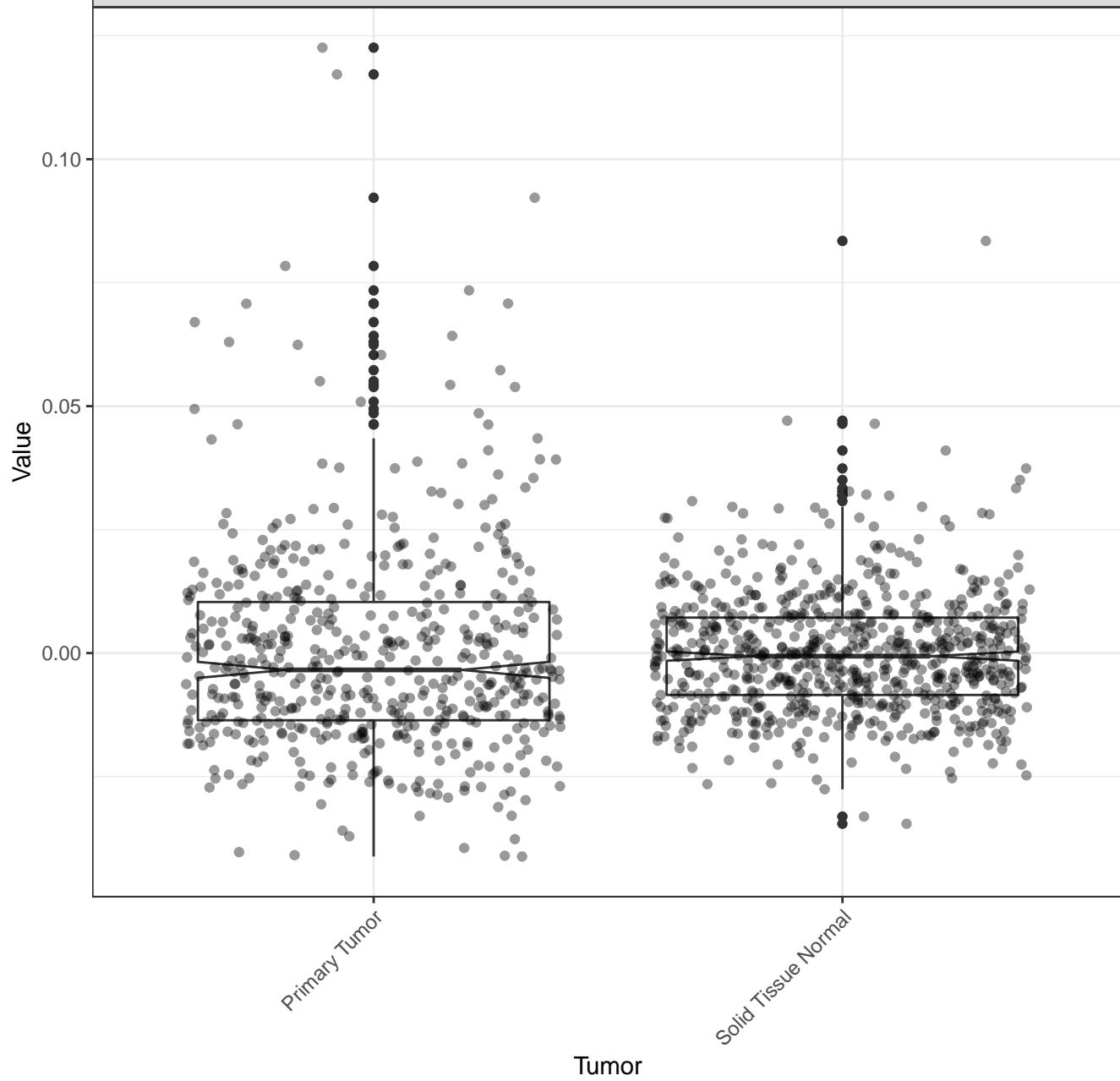
Primary Tumor

Solid Tissue Normal

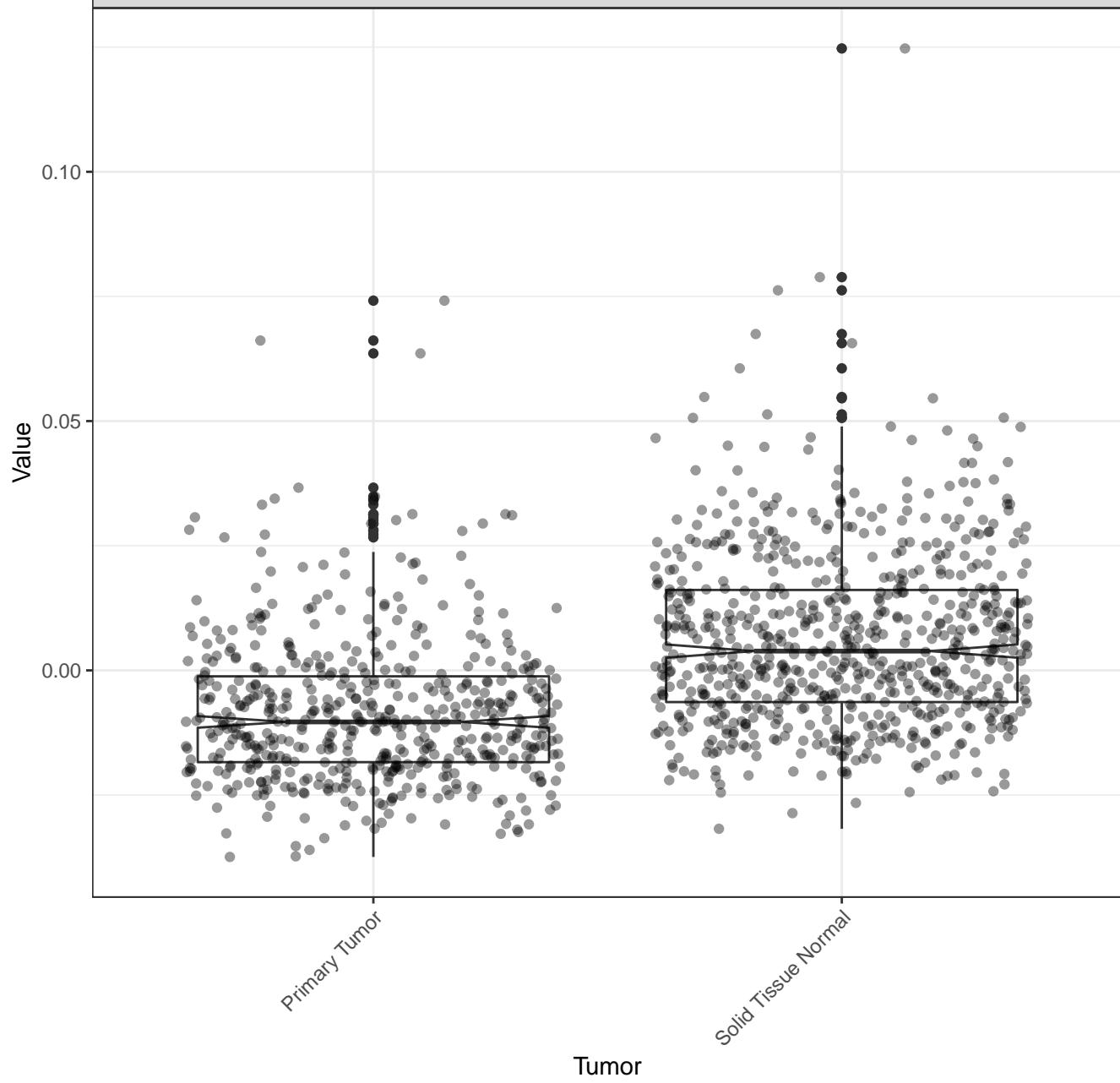
Tumor



102,REACTOME_METABOLISM_OF_CARBOHYDRATES



103,REACTOME_MRNA_PROCESSING



104,SVM Mast cells resting

Value

0.3

0.2

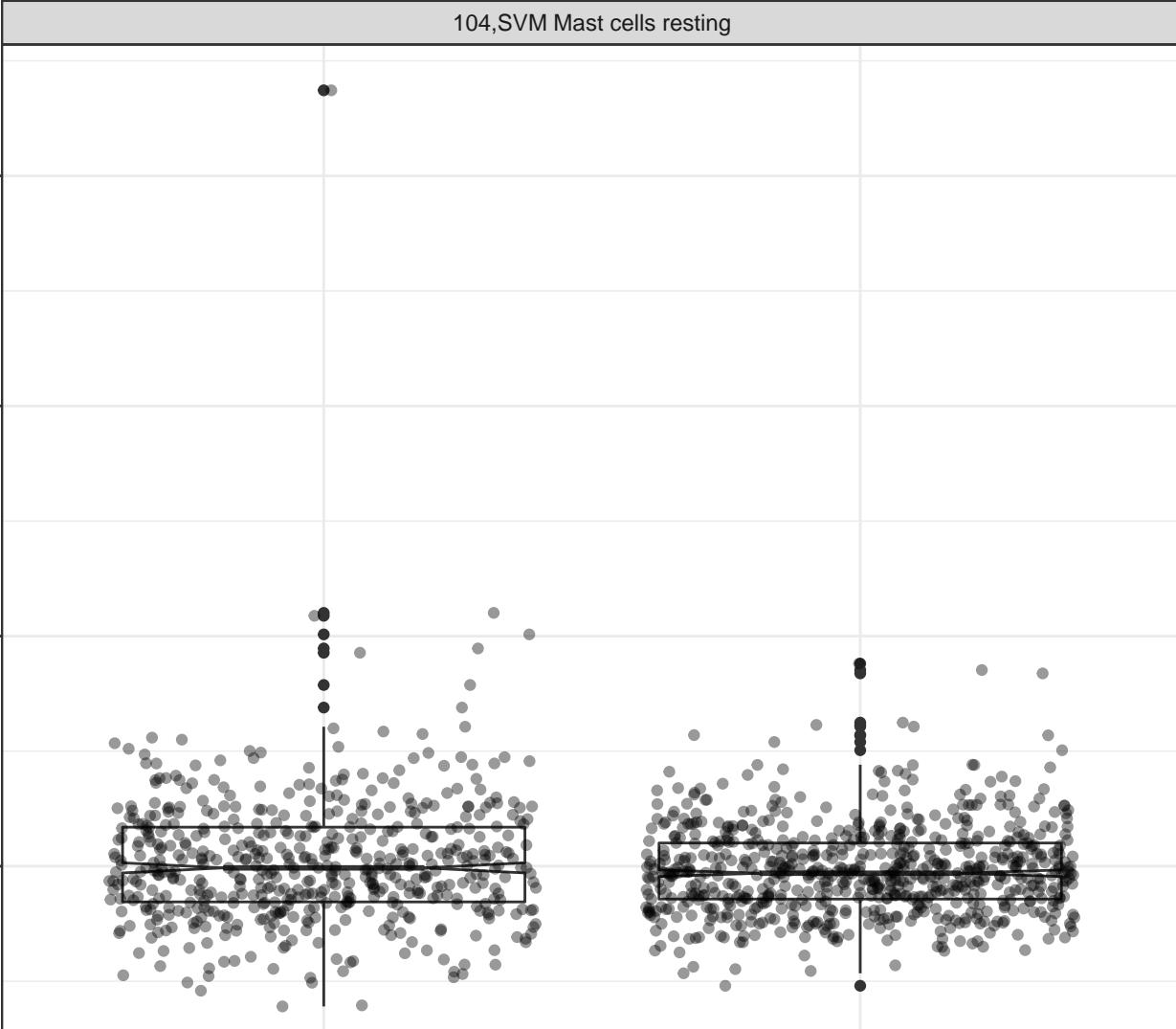
0.1

0.0

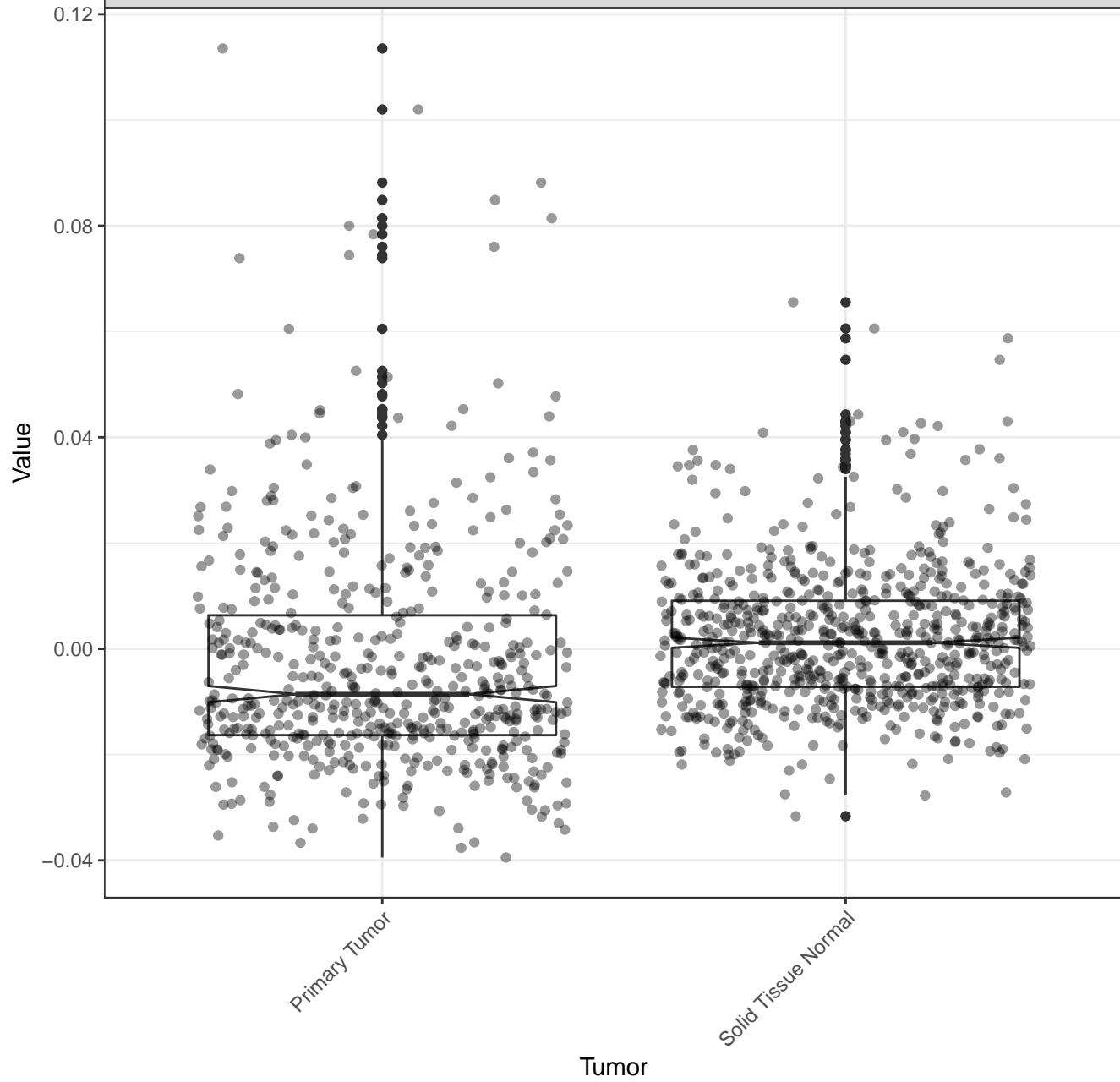
Primary Tumor

Solid Tissue Normal

Tumor



105,PID_ECADHERIN_STABILIZATION_PATHWAY



106,REACTOME_NCAM_SIGNALING_FOR_NEURITE_OUT_GROWTH

Value

0.10

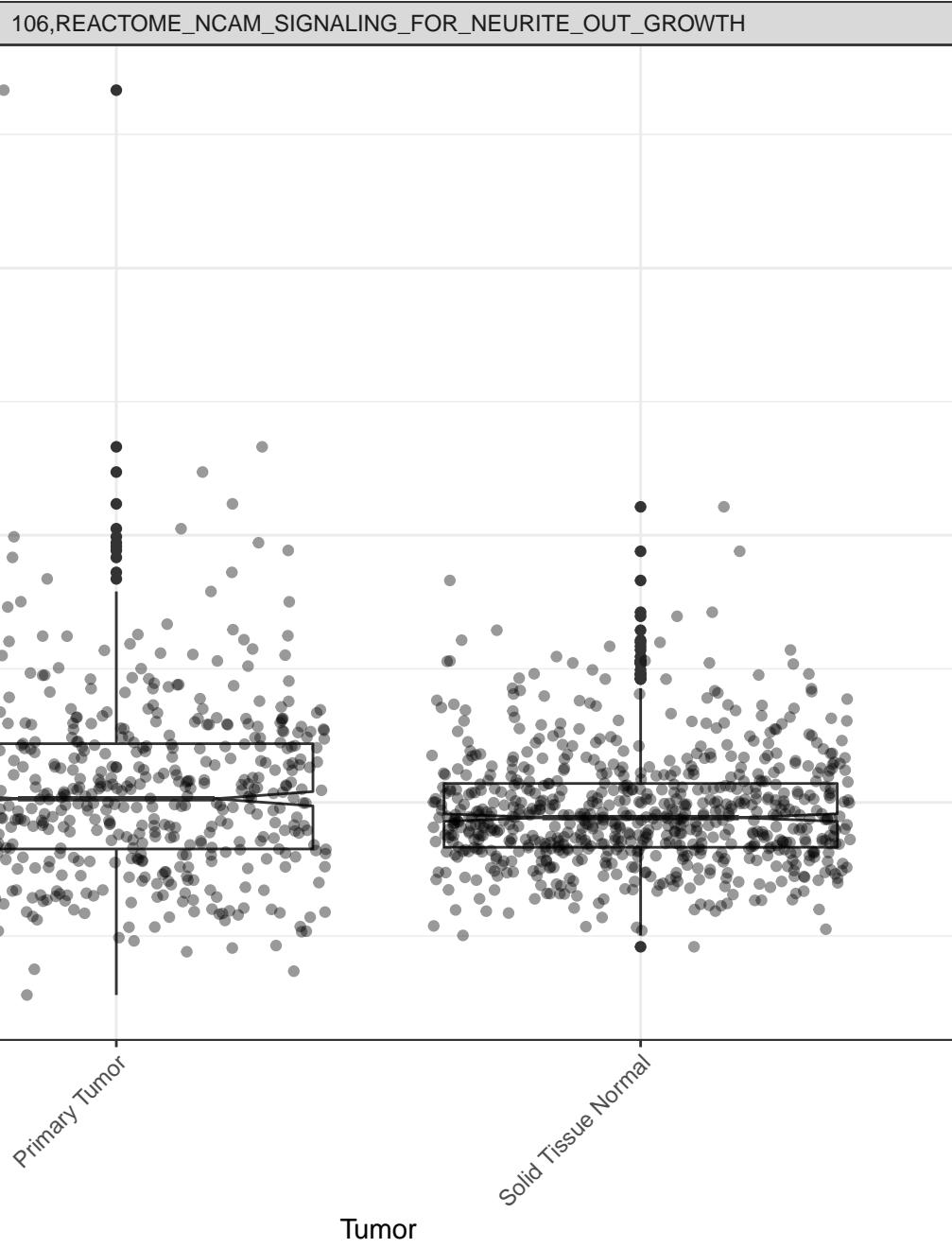
0.05

0.00

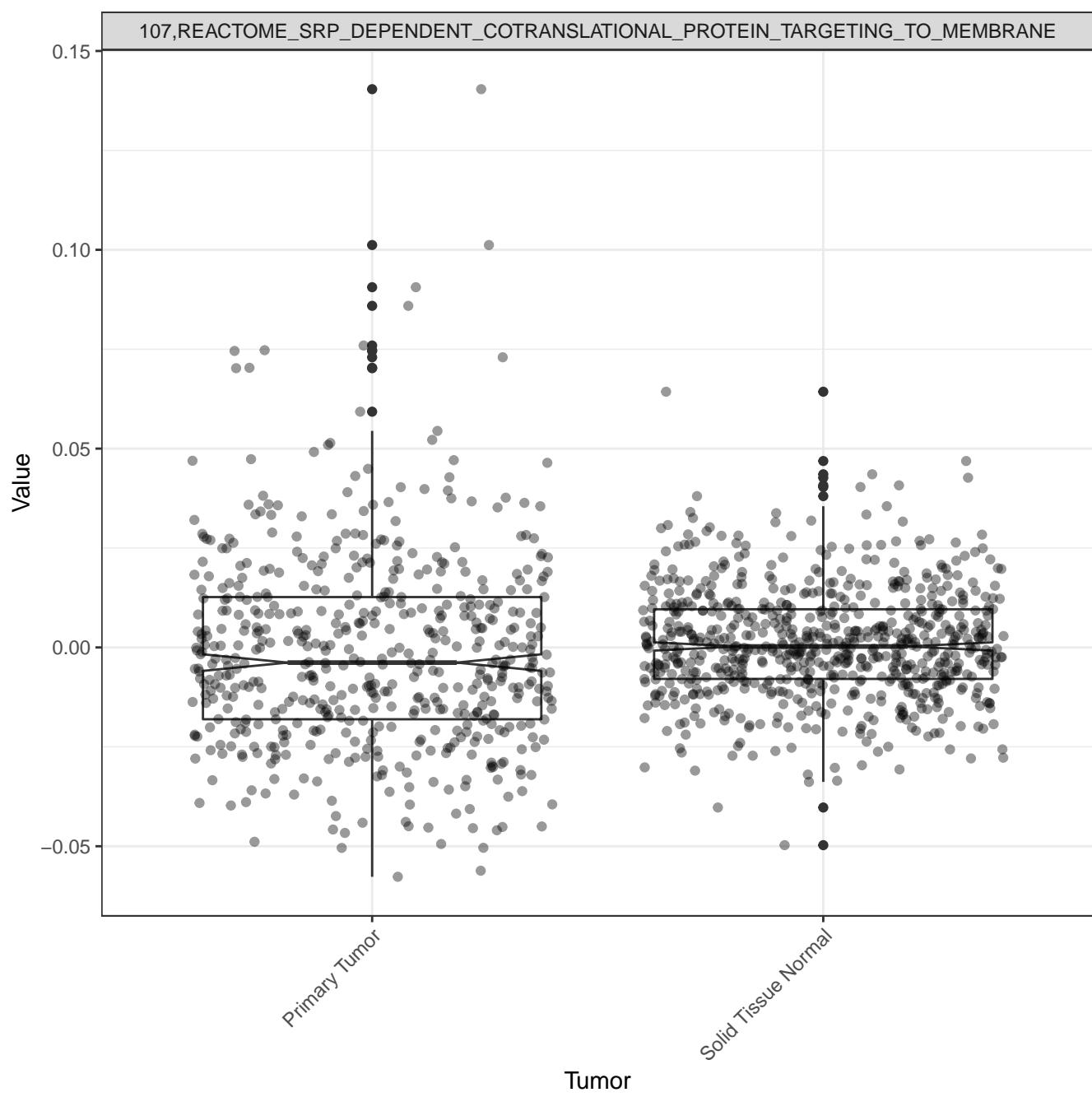
Primary Tumor

Tumor

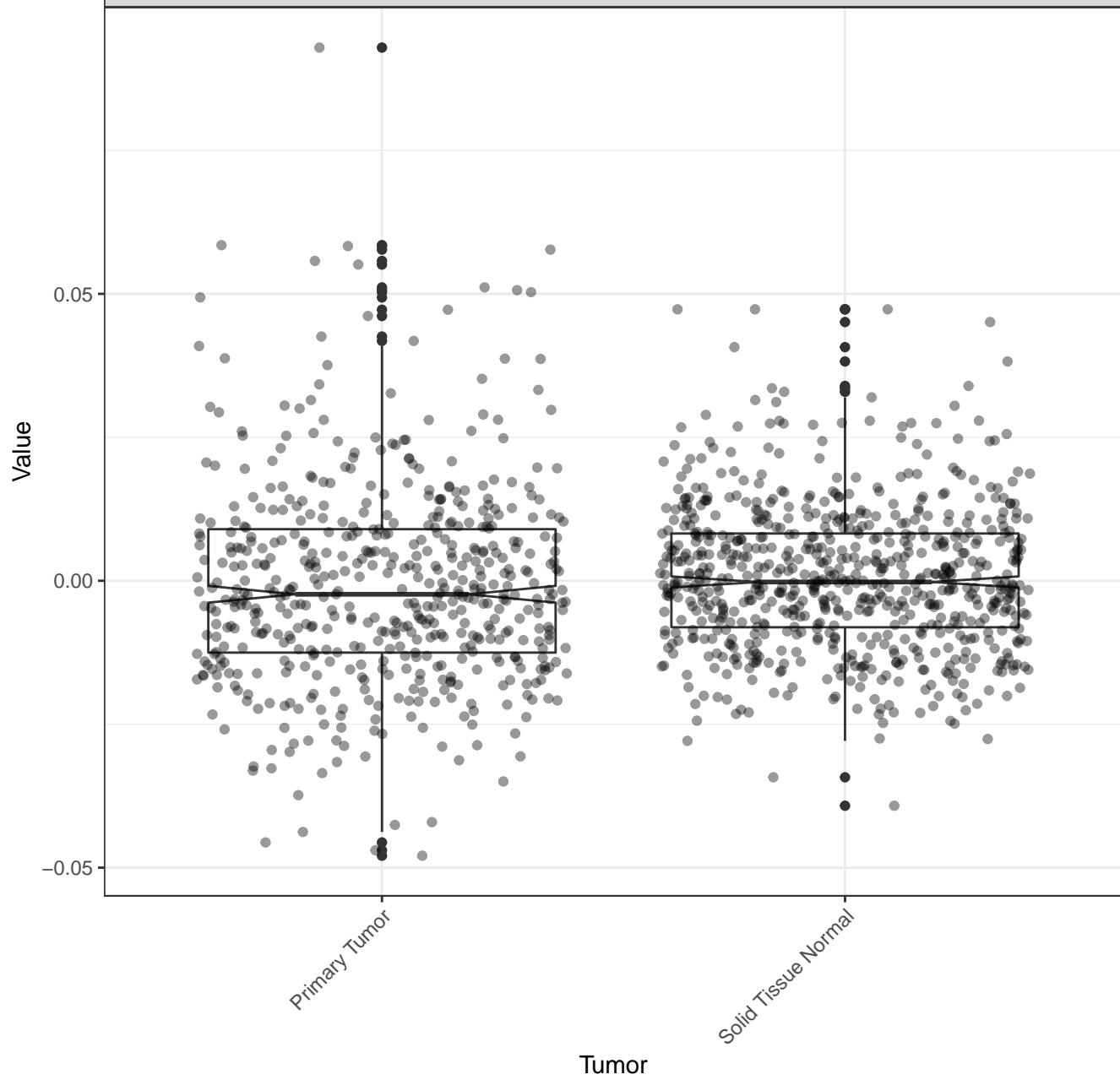
Solid Tissue Normal



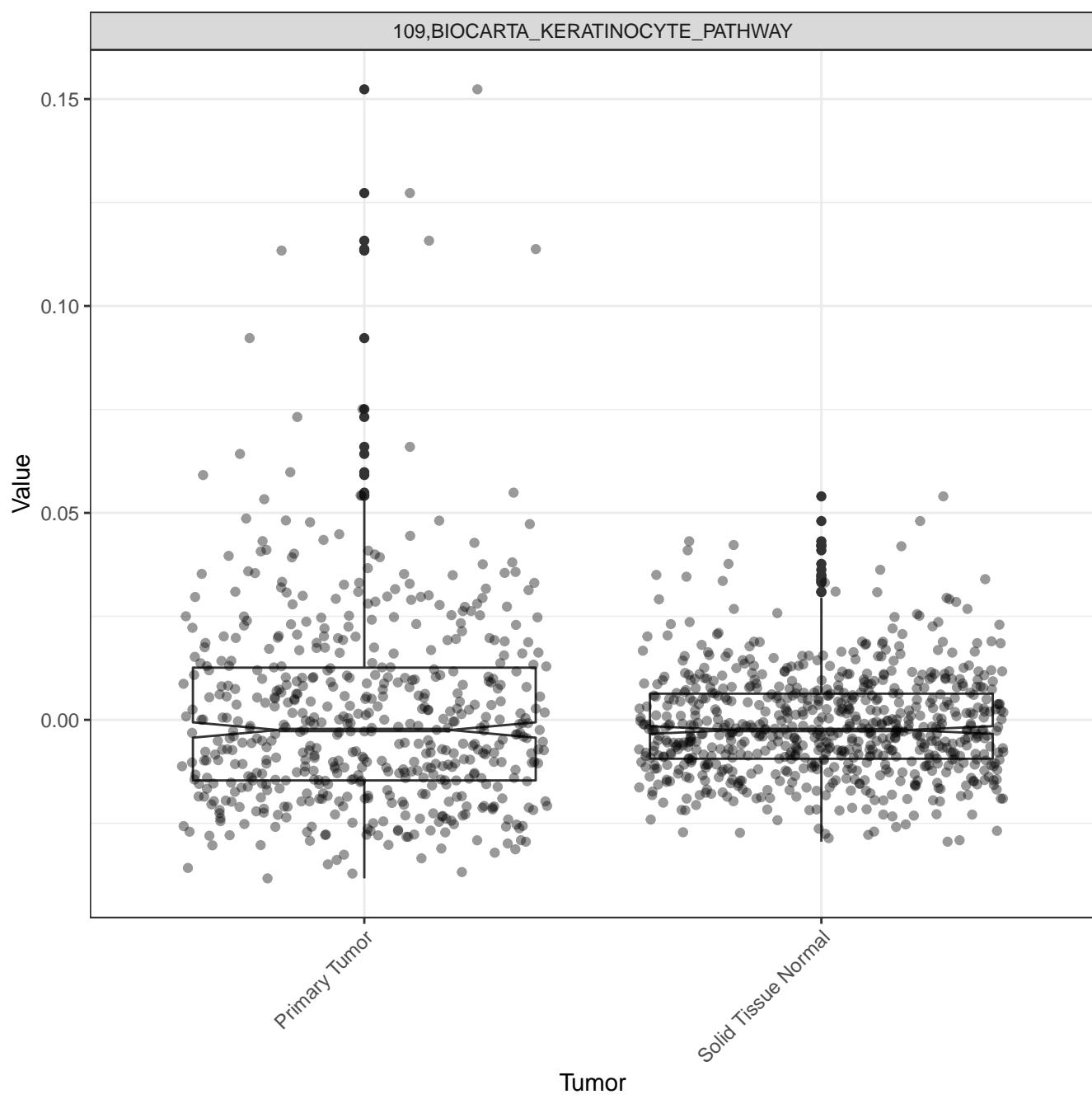
107,REACTOME_SR_P_DEPENDENT_COTRANSLATIONAL_PROTEIN_TARGETING_TO_MEMBRANE



108,REACTOME_MRNA_SPLICING



109,BIOCARTA_KERATINOCYTE_PATHWAY



110,REACTOME_PROCESSING_OF_CAPPED_INTRON_CONTAINING_PRE_MRNA

Value

0.10

0.05

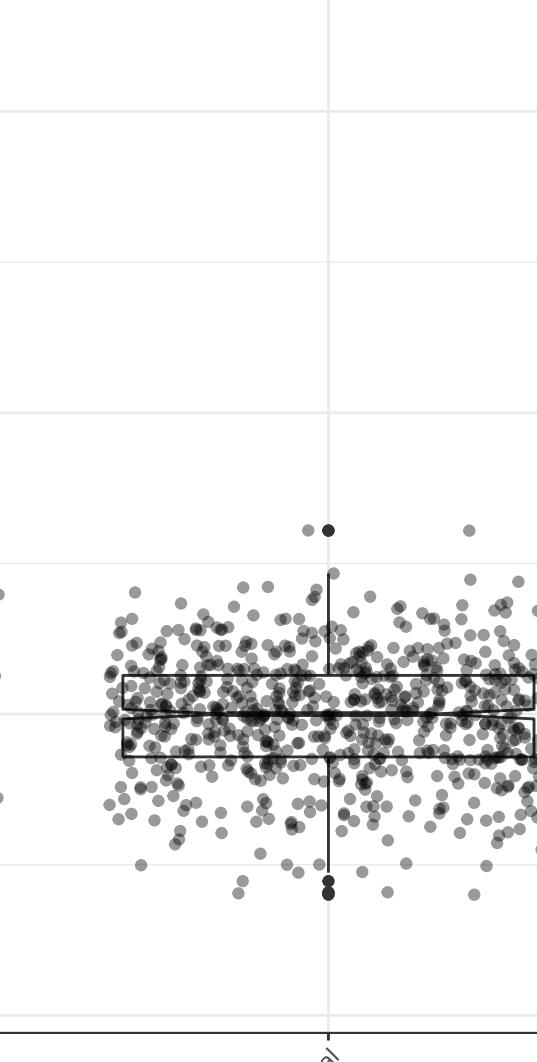
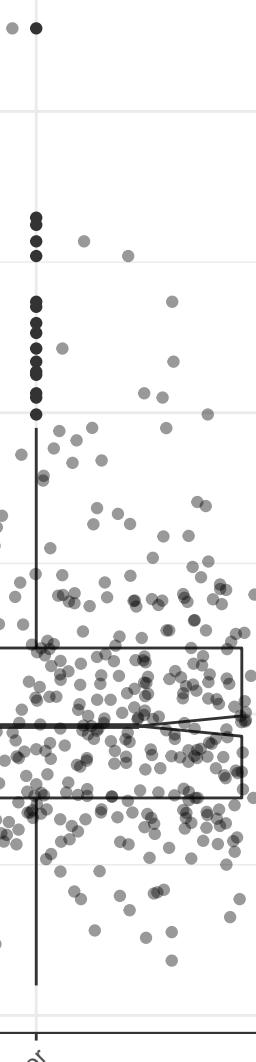
0.00

-0.05

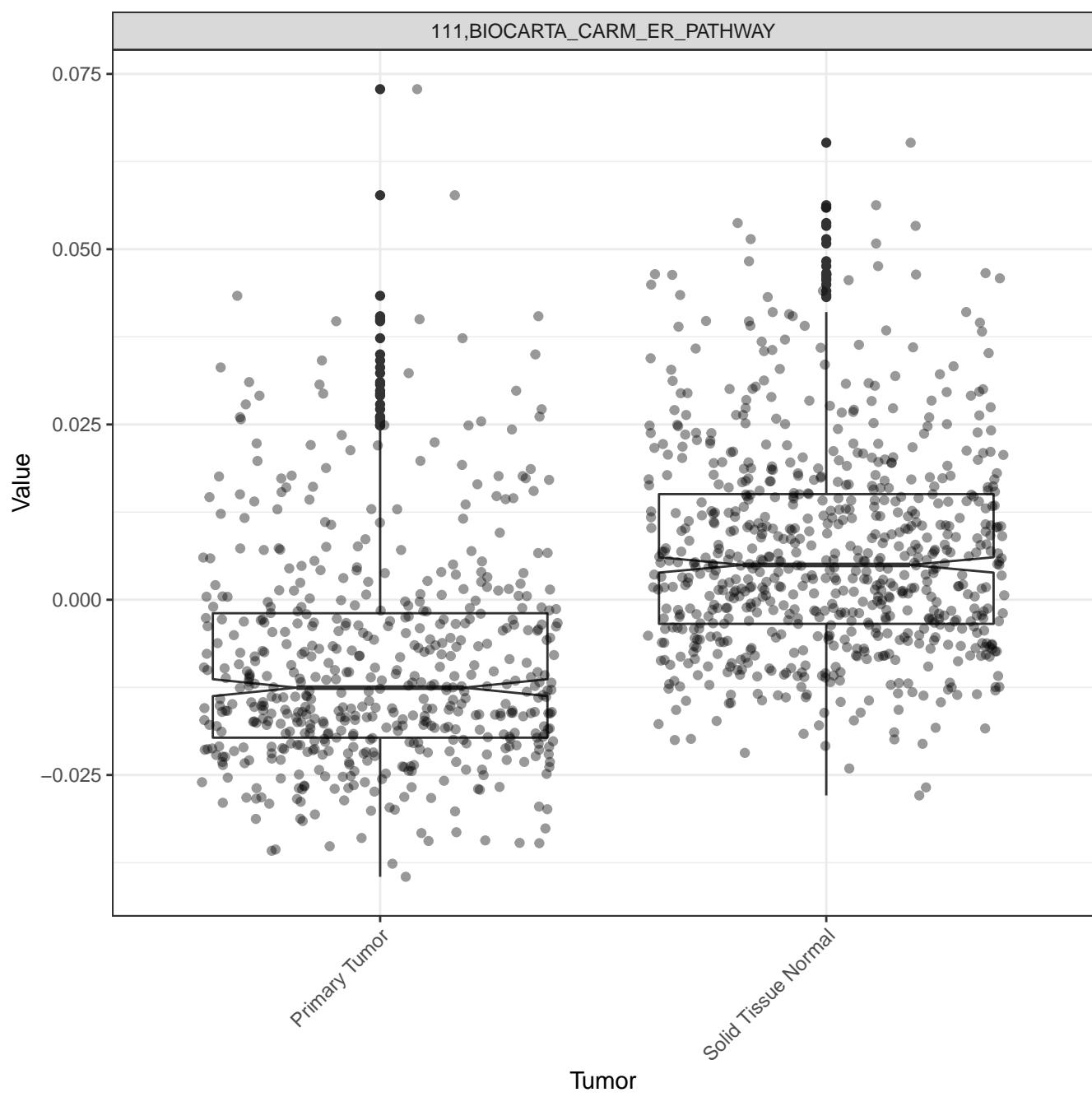
Primary Tumor

Tumor

Solid Tissue Normal



111,BIOCARTA_CARM_ER_PATHWAY



112,REACTOME_P75_NTR_RECECTOR_MEDIDATED_SIGNALLING

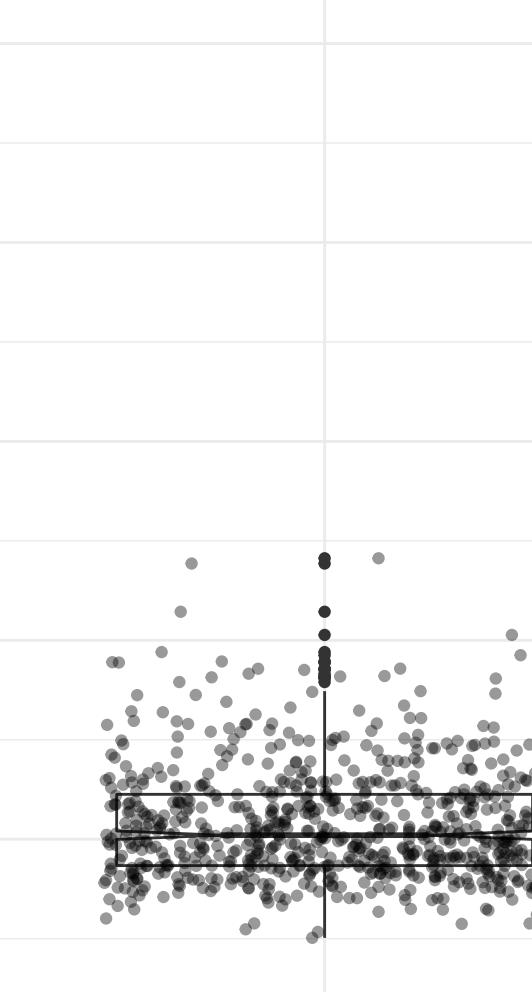
Value

0.20
0.15
0.10
0.05
0.00

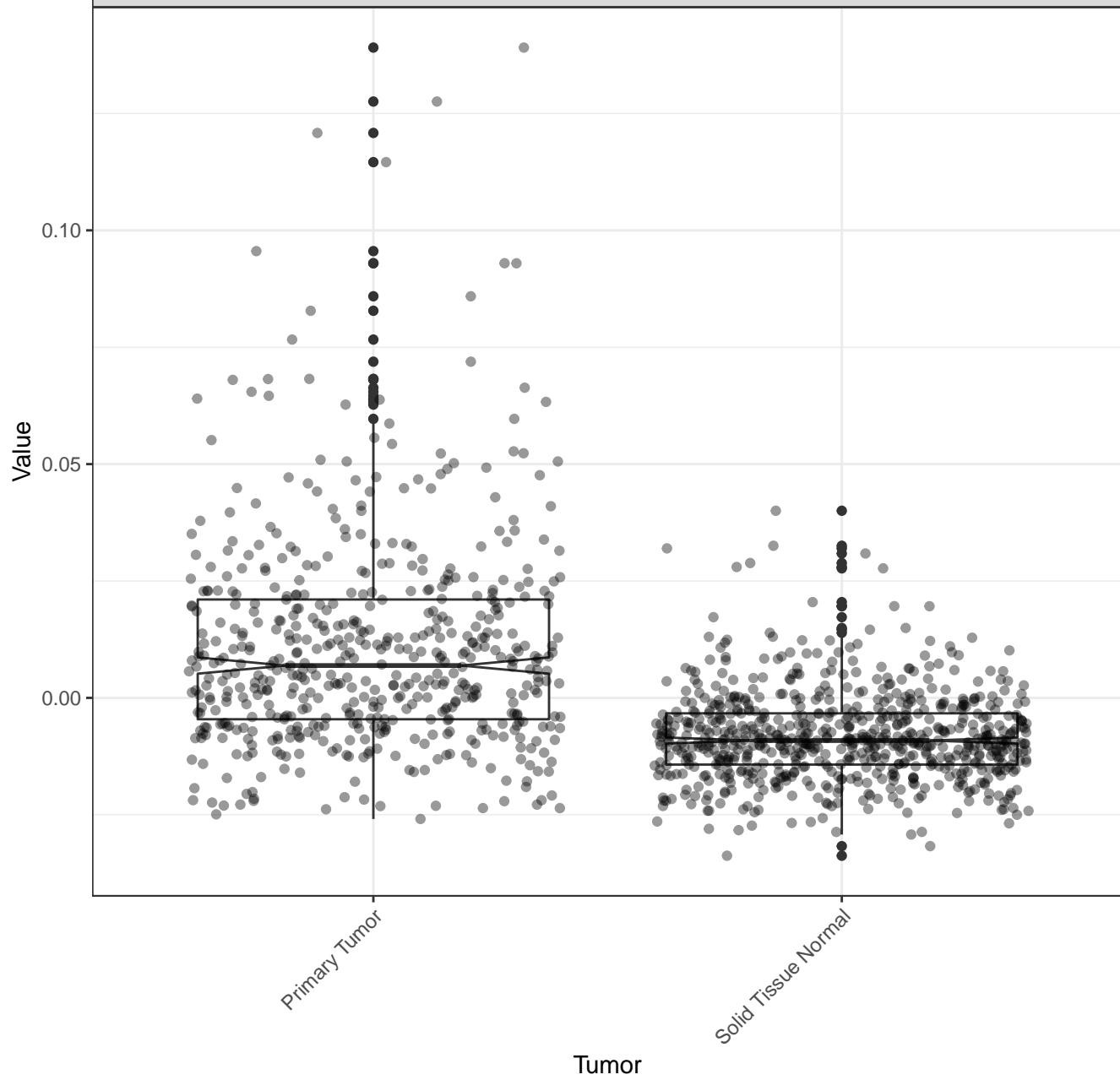
Primary Tumor

Solid Tissue Normal

Tumor



113,REACTOME_ACTIVATION_OF_ATR_IN_RESPONSE_TO_REPLICATION_STRESS



114,REACTOME_METABOLISM_OF_LIPIDS_AND_LIPOPROTEINS

Value

0.4

0.2

0.0

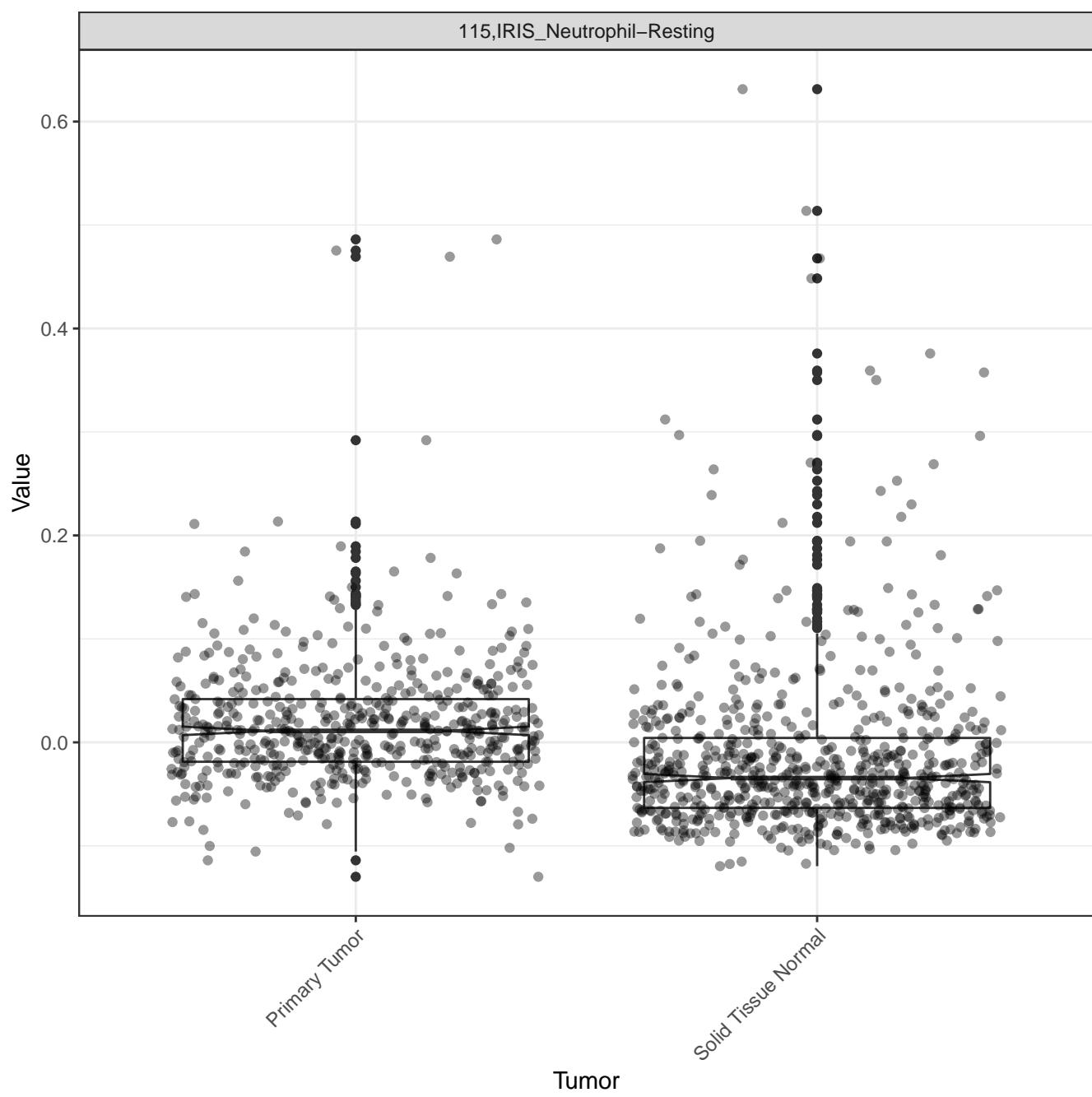
Primary Tumor

Solid Tissue Normal

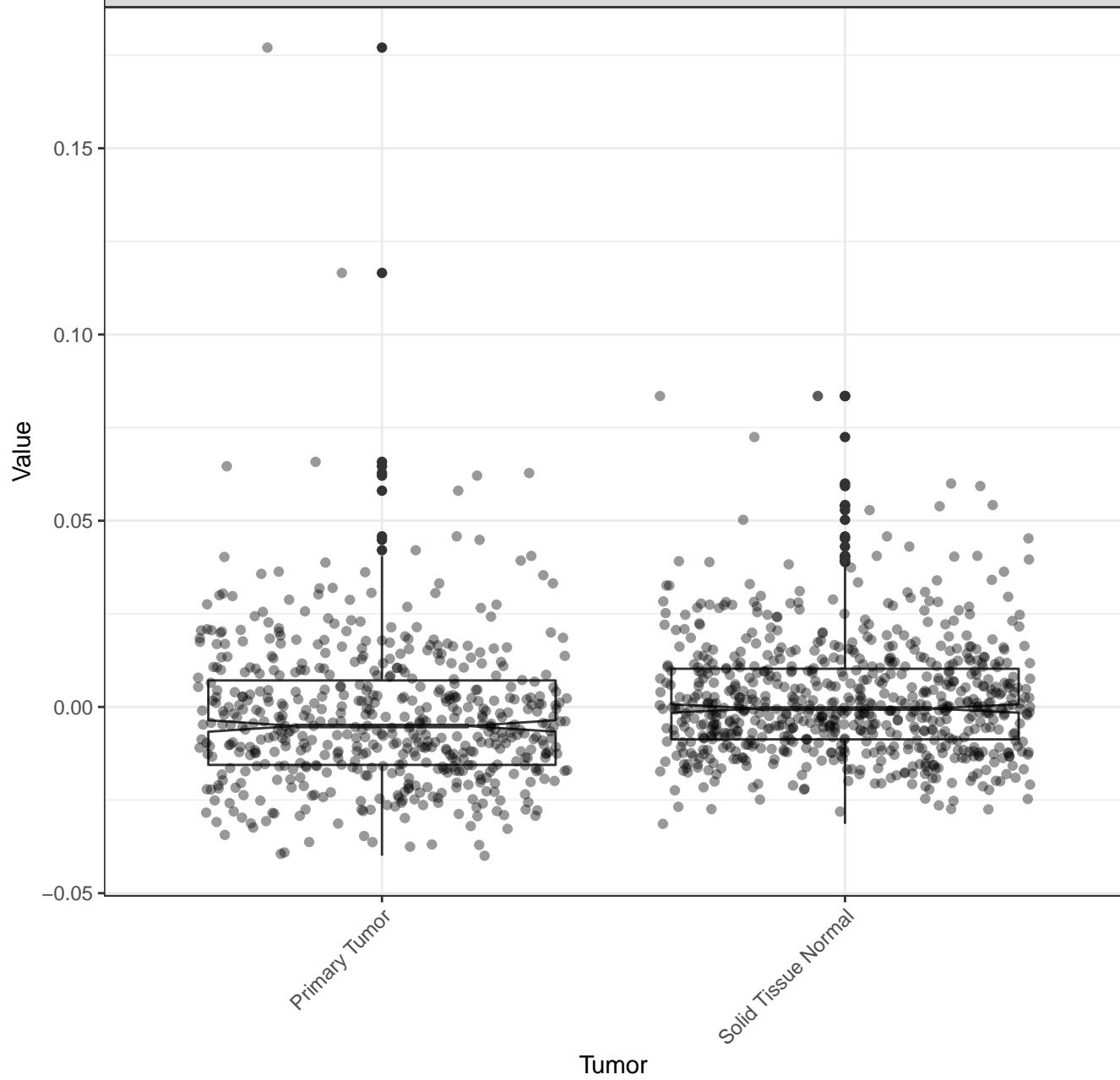
Tumor



115,IRIS_Neutrophil-Resting



116,KEGG_HEDGEHOG_SIGNALING_PATHWAY



117,REACTOME_PROCESSING_OF_CAPPED_INTRON_CONTAINING_PRE_MRNA

Value

0.15

0.10

0.05

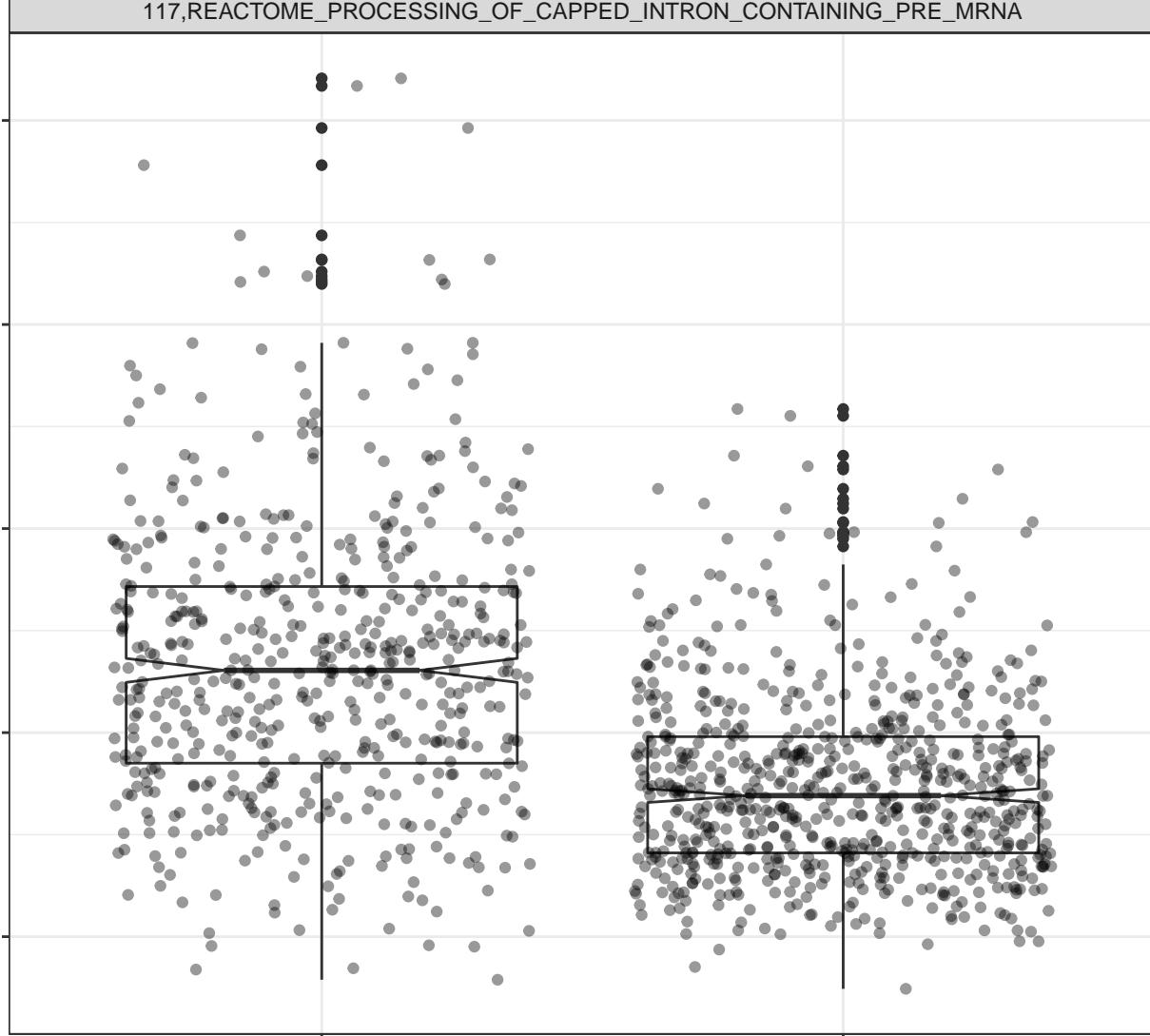
0.00

-0.05

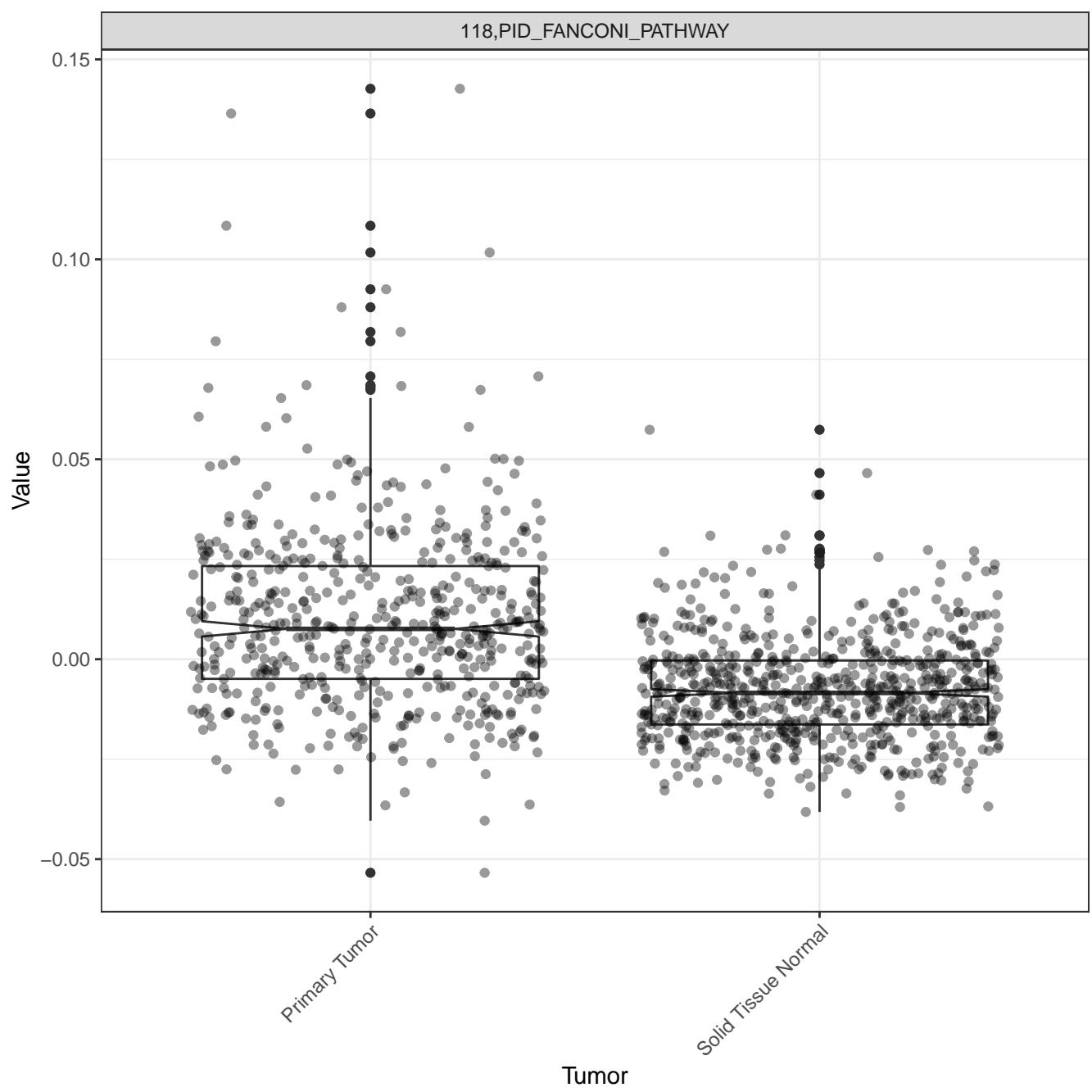
Primary Tumor

Solid Tissue Normal

Tumor



118,PID_FANCONI_PATHWAY



119,REACTOME_PHASE1_FUNCTIONALIZATION_OF_COMPOUNDS

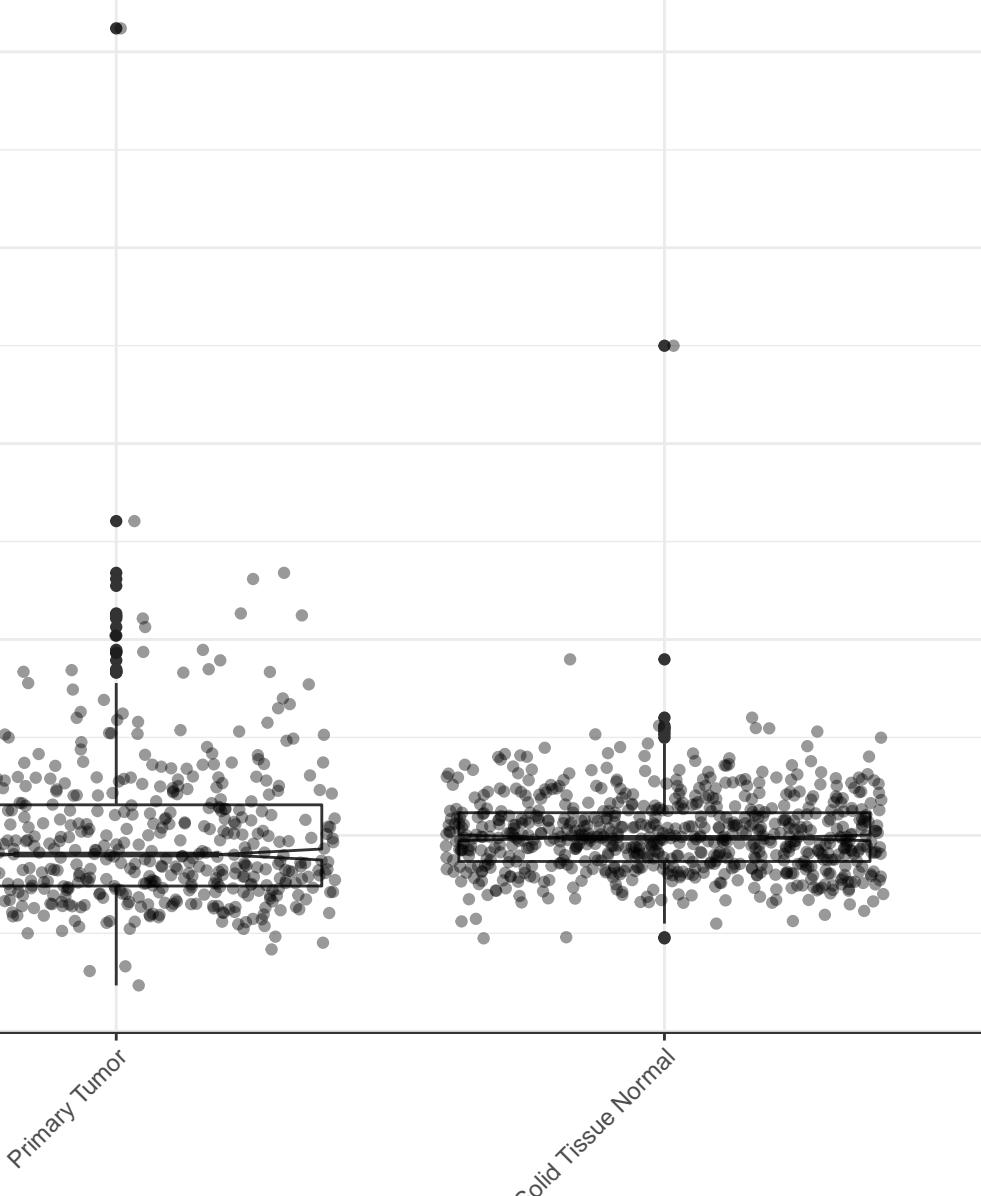
Value

0.20
0.15
0.10
0.05
0.00
-0.05

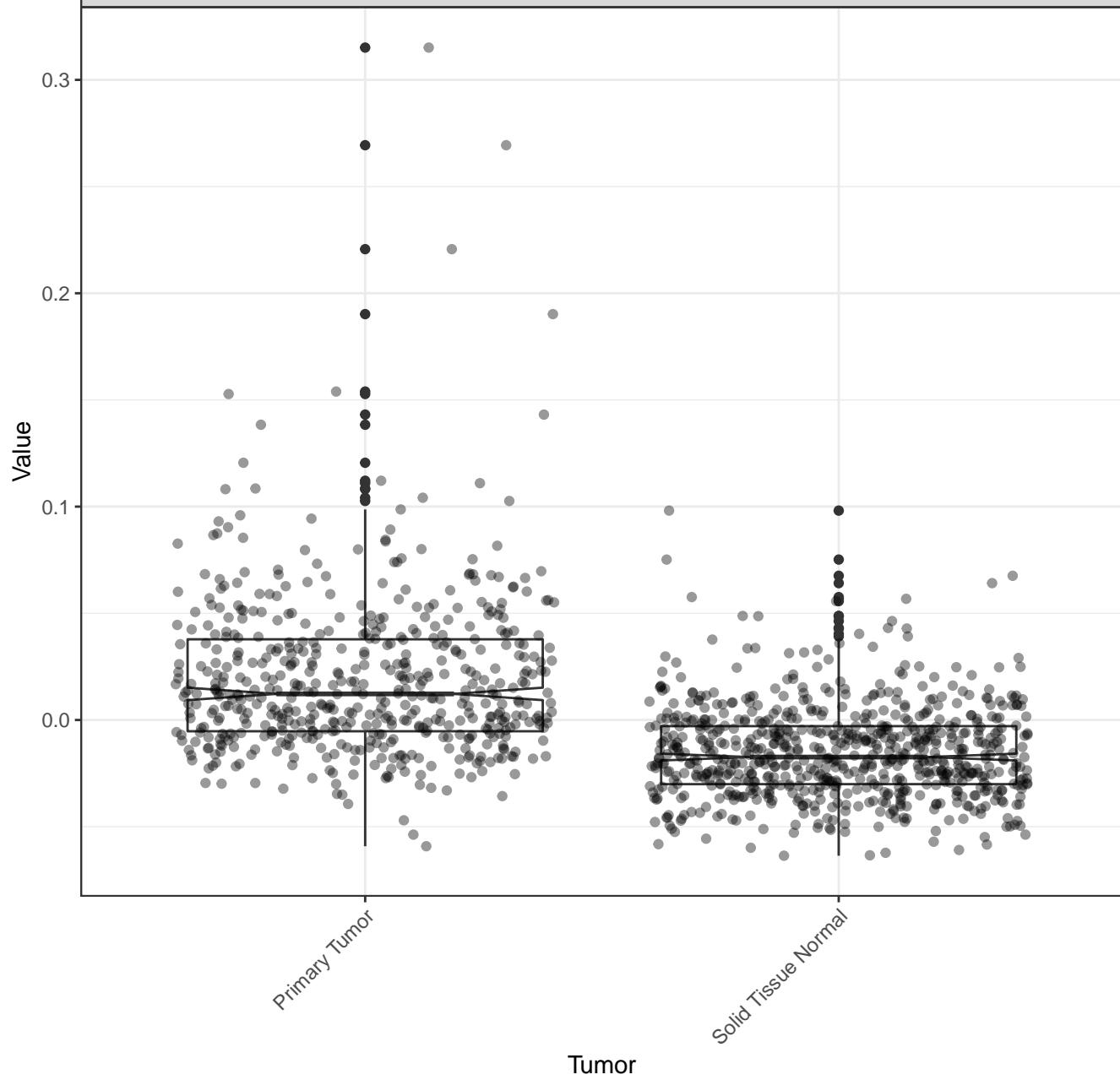
Primary Tumor

Solid Tissue Normal

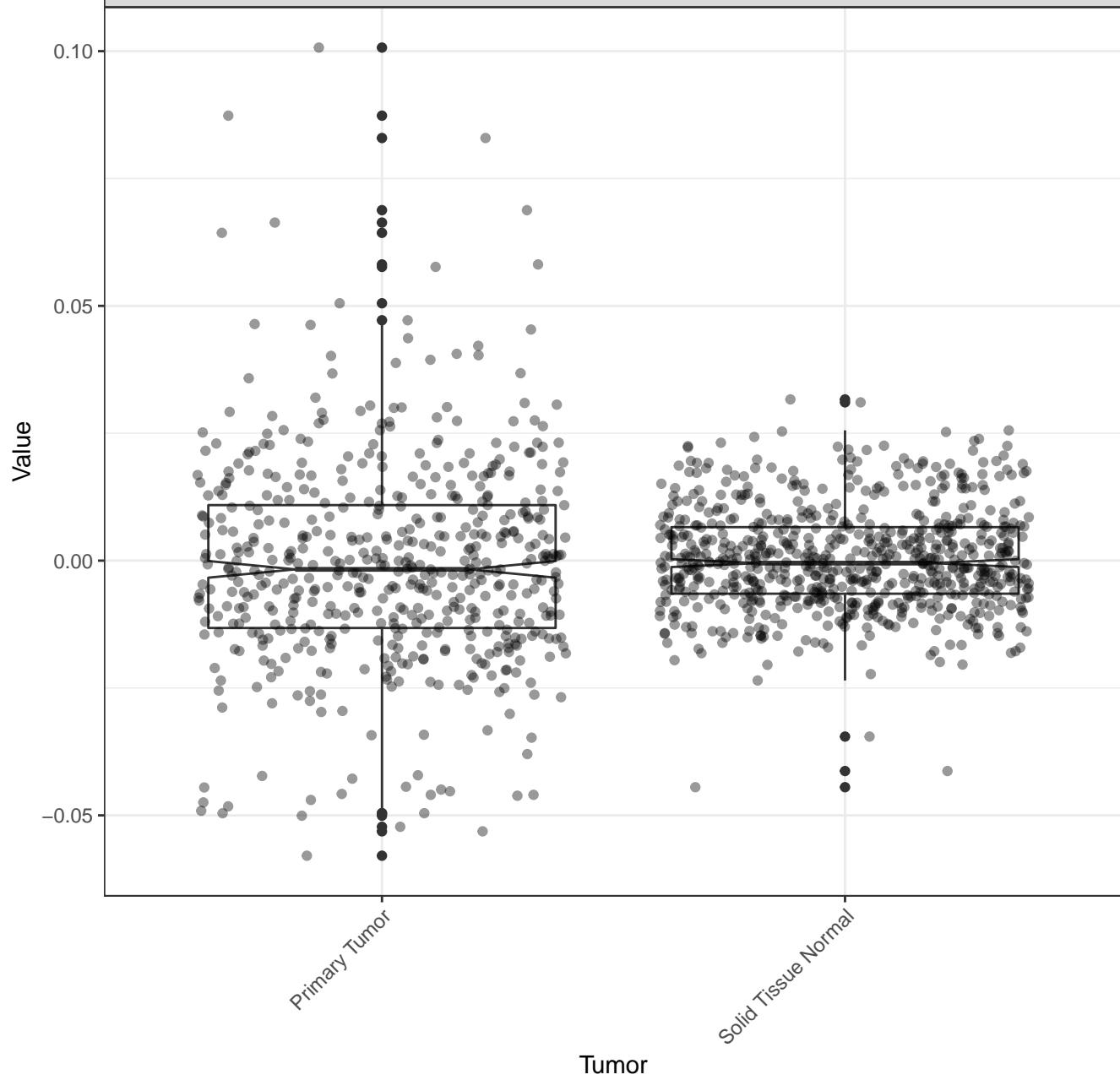
Tumor



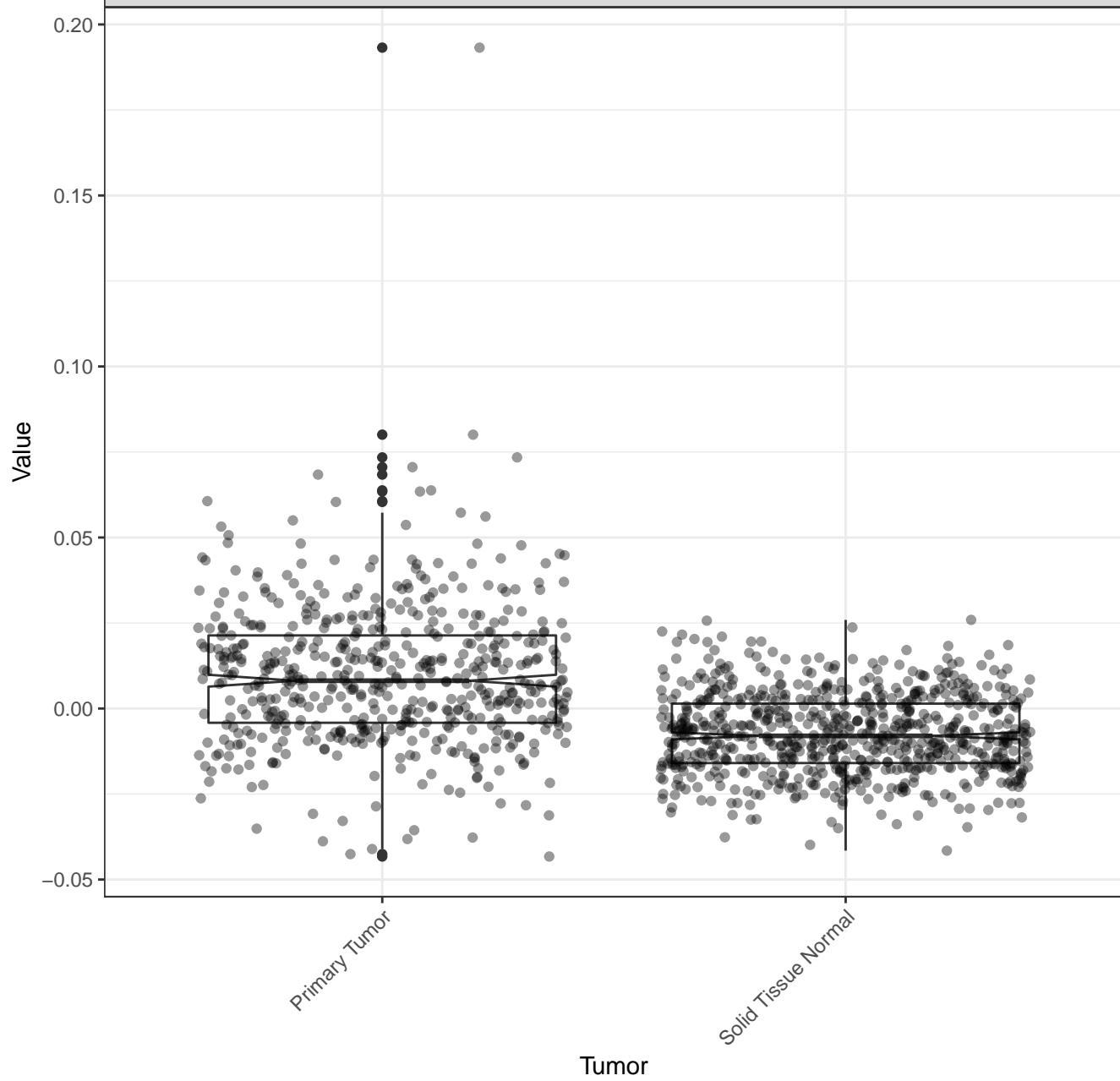
120,KEGG_PARKINSONS_DISEASE



121,REACTOME_METABOLISM_OF_PROTEINS



122,REACTOME DEVELOPMENTAL_BIOLOGY



123,KEGG_INOSITOL_PHOSPHATE_METABOLISM

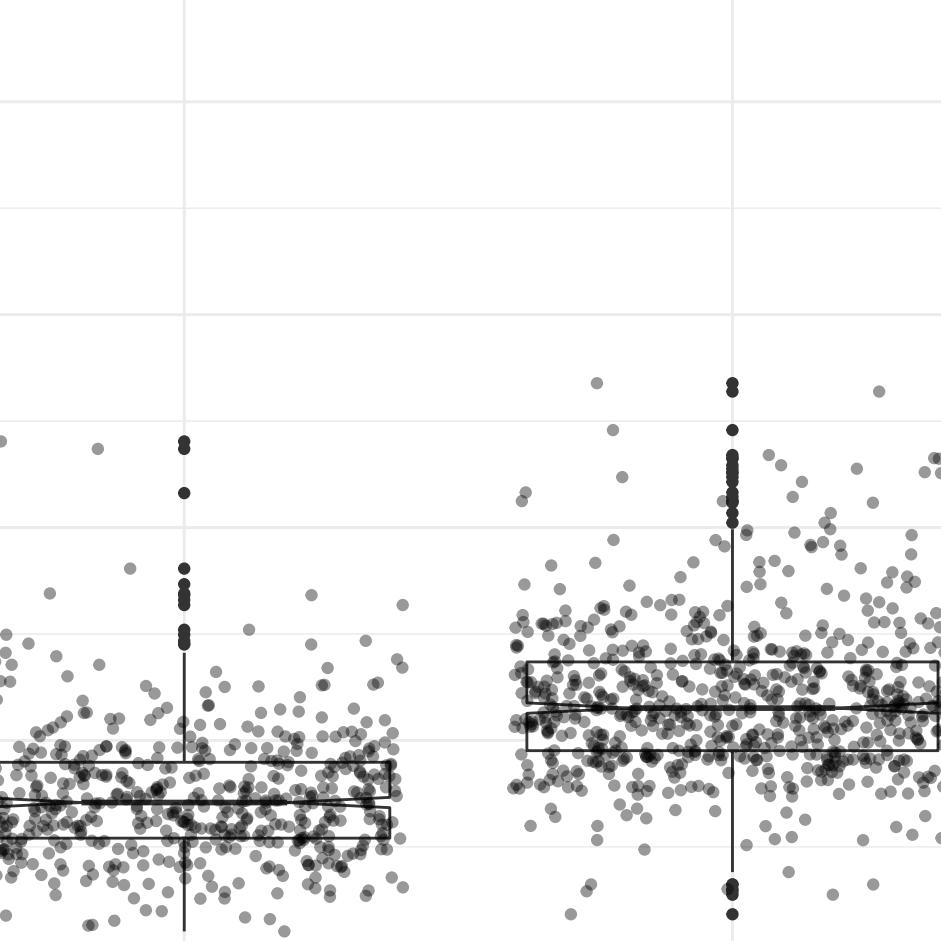
Value

0.15
0.10
0.05
0.00
-0.05

Primary Tumor

Solid Tissue Normal

Tumor



124,REACTOME_DEADENYLATION_DEPENDENT_MRNA_DECAY

Value

0.15

0.10

0.05

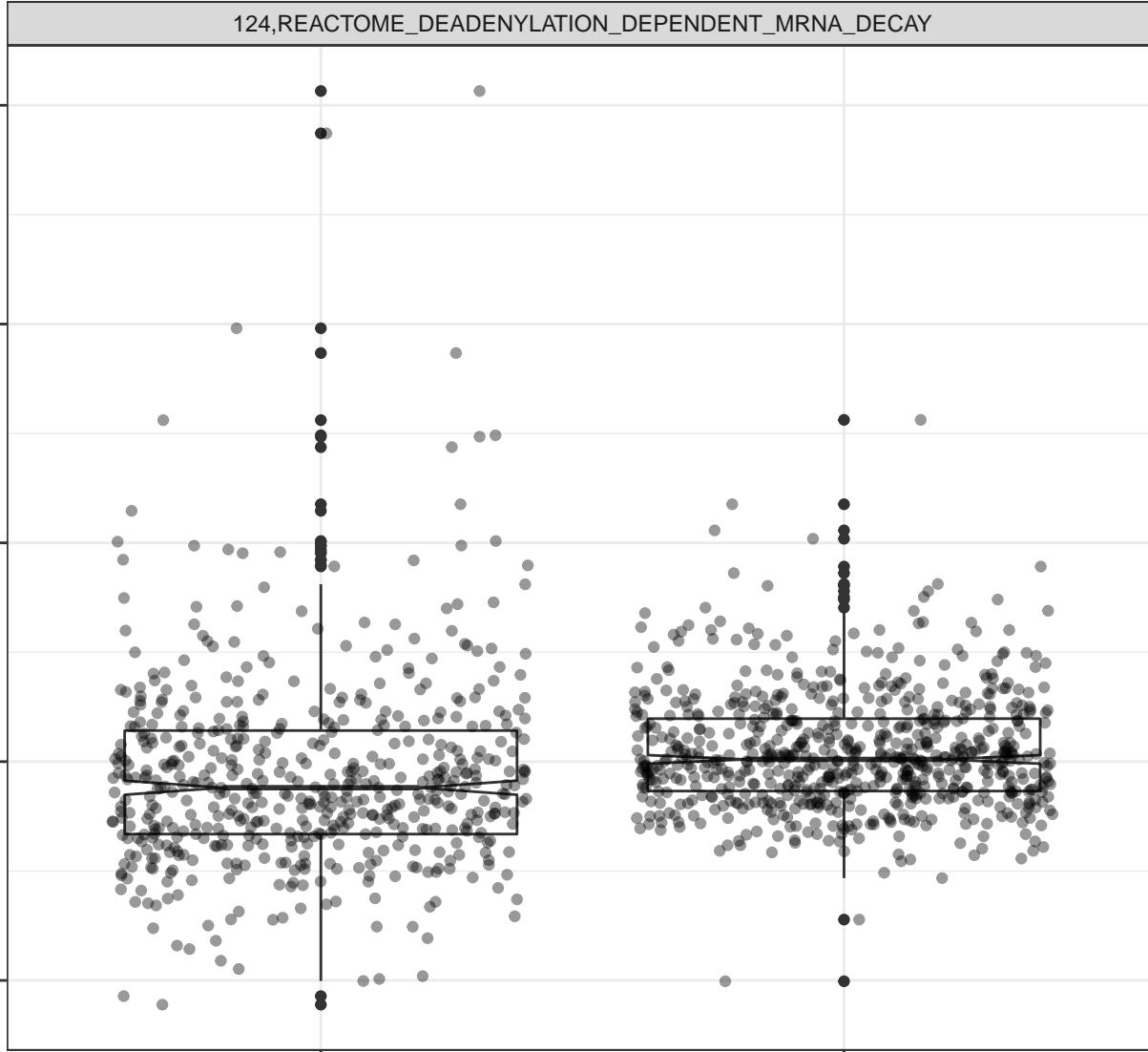
0.00

-0.05

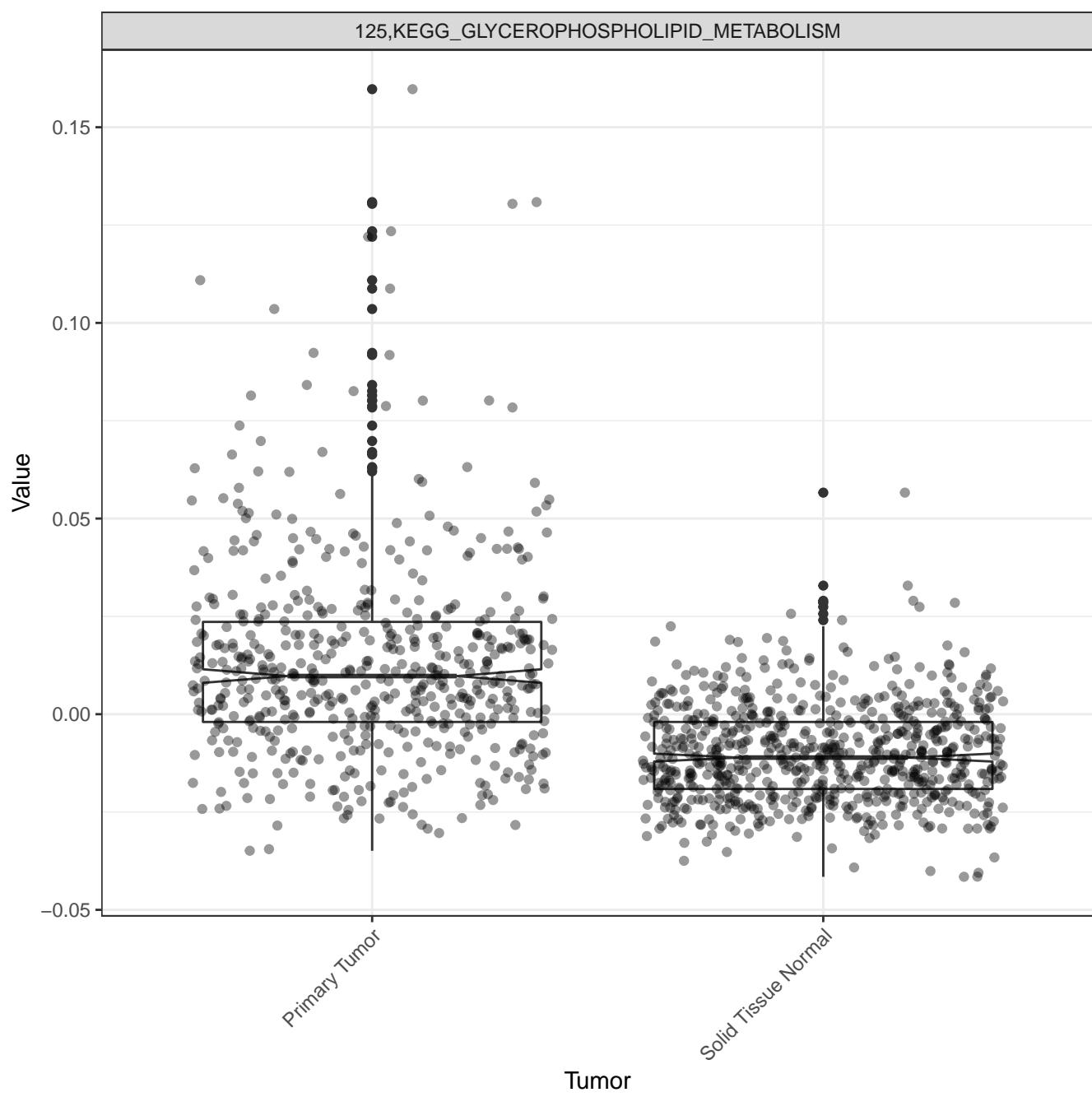
Primary Tumor

Solid Tissue Normal

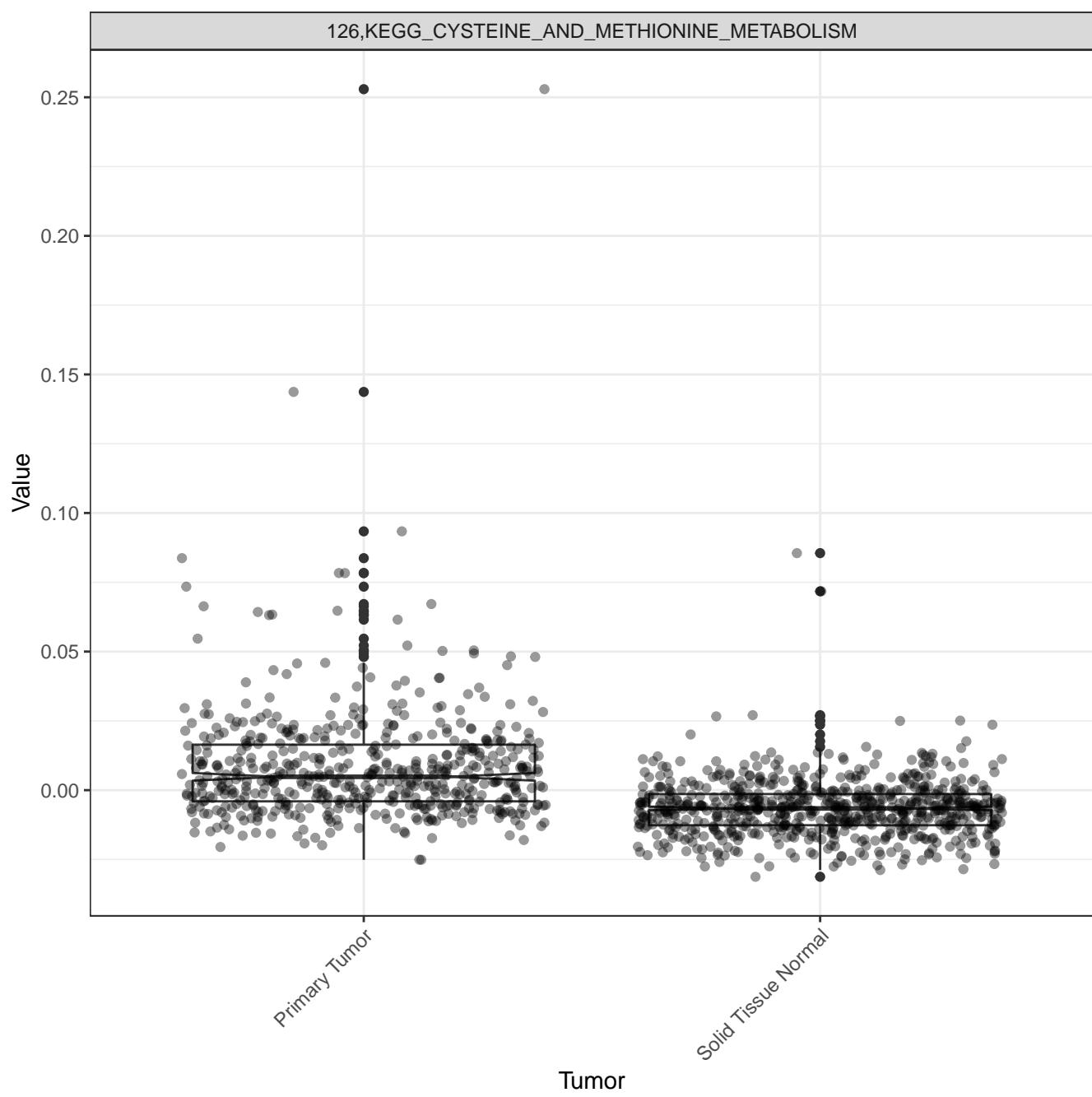
Tumor



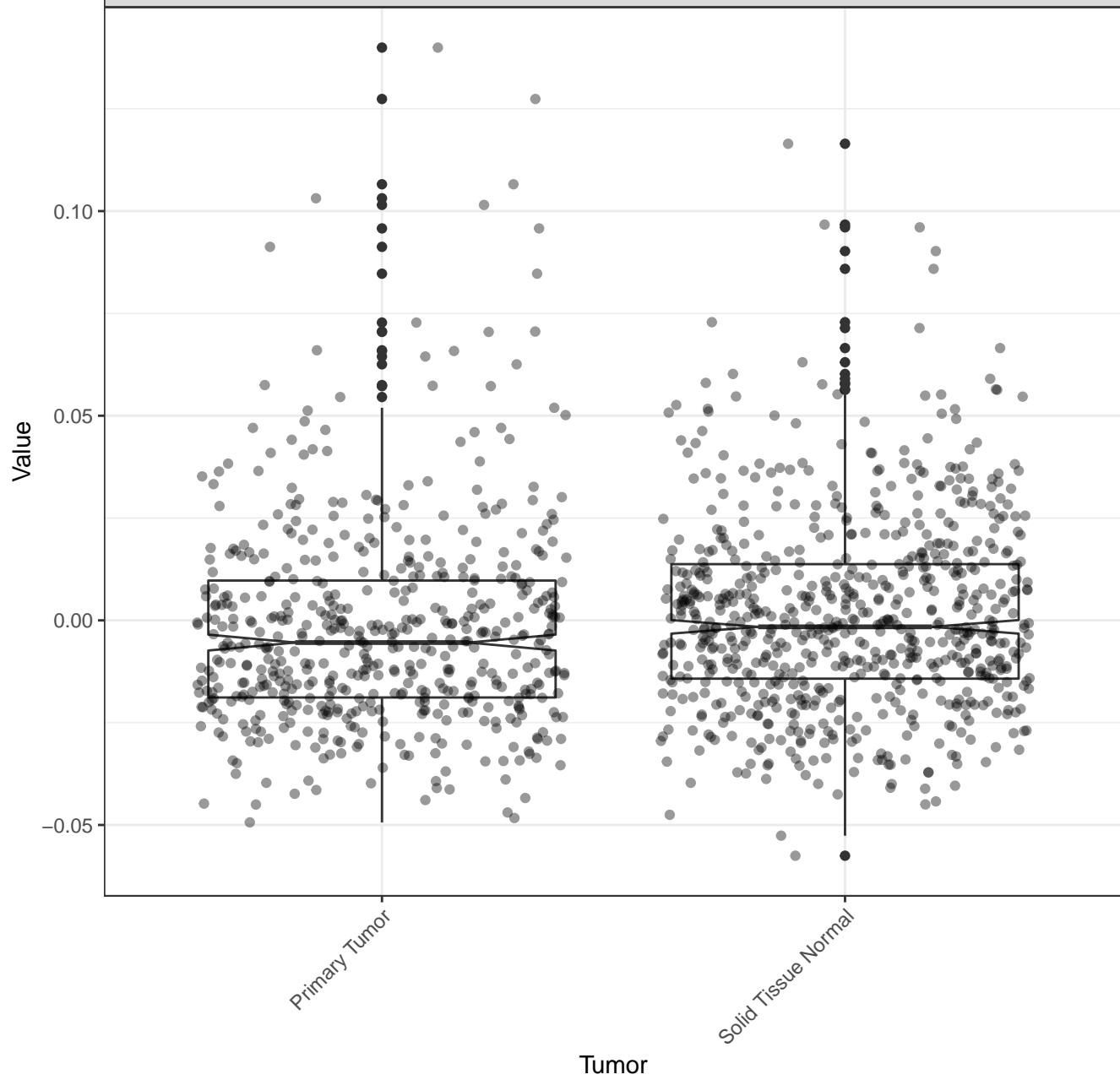
125,KEGG_GLYCEROPHOSPHOLIPID_METABOLISM



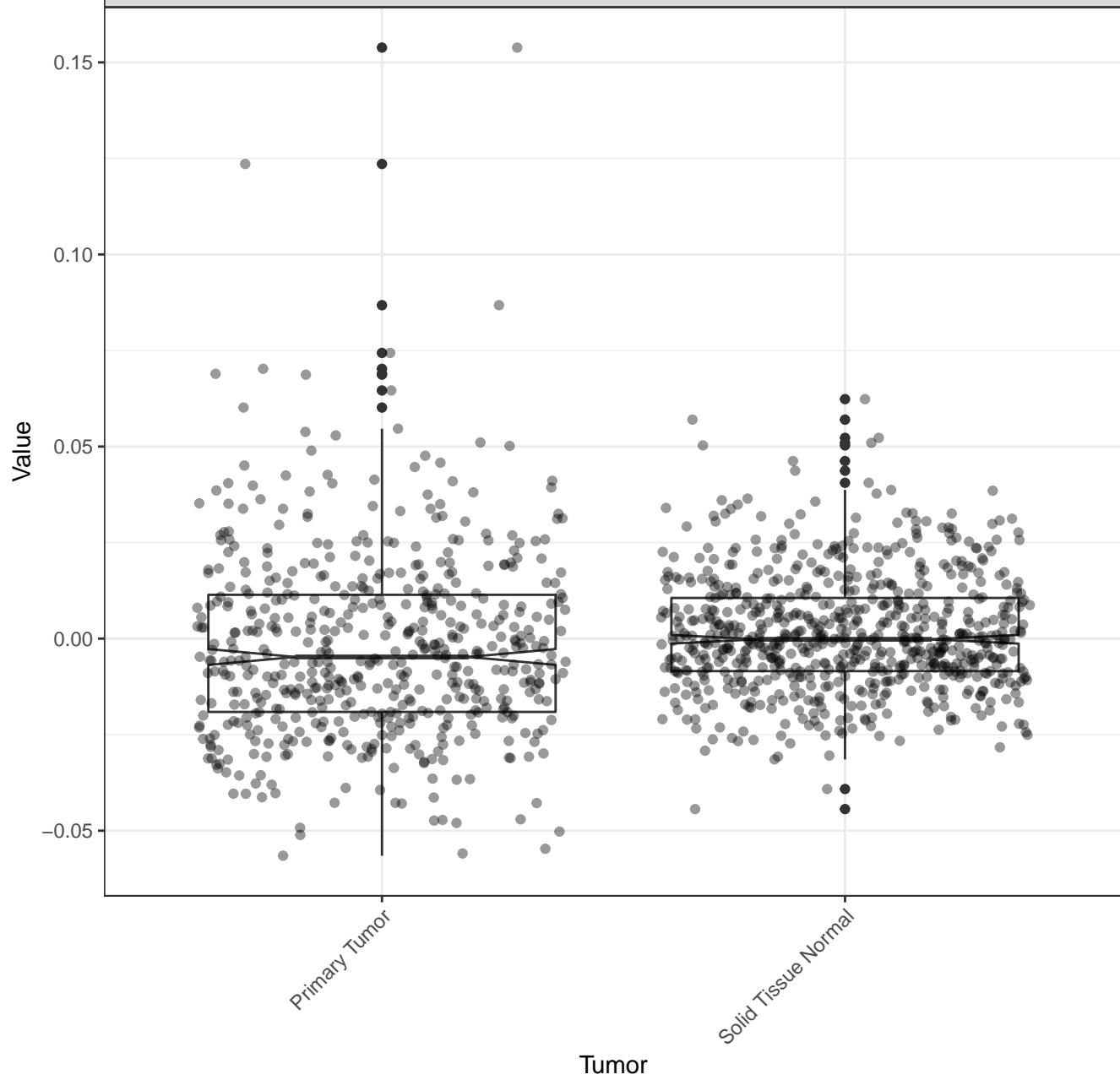
126,KEGG_CYSTEINE_AND_METHIONINE_METABOLISM



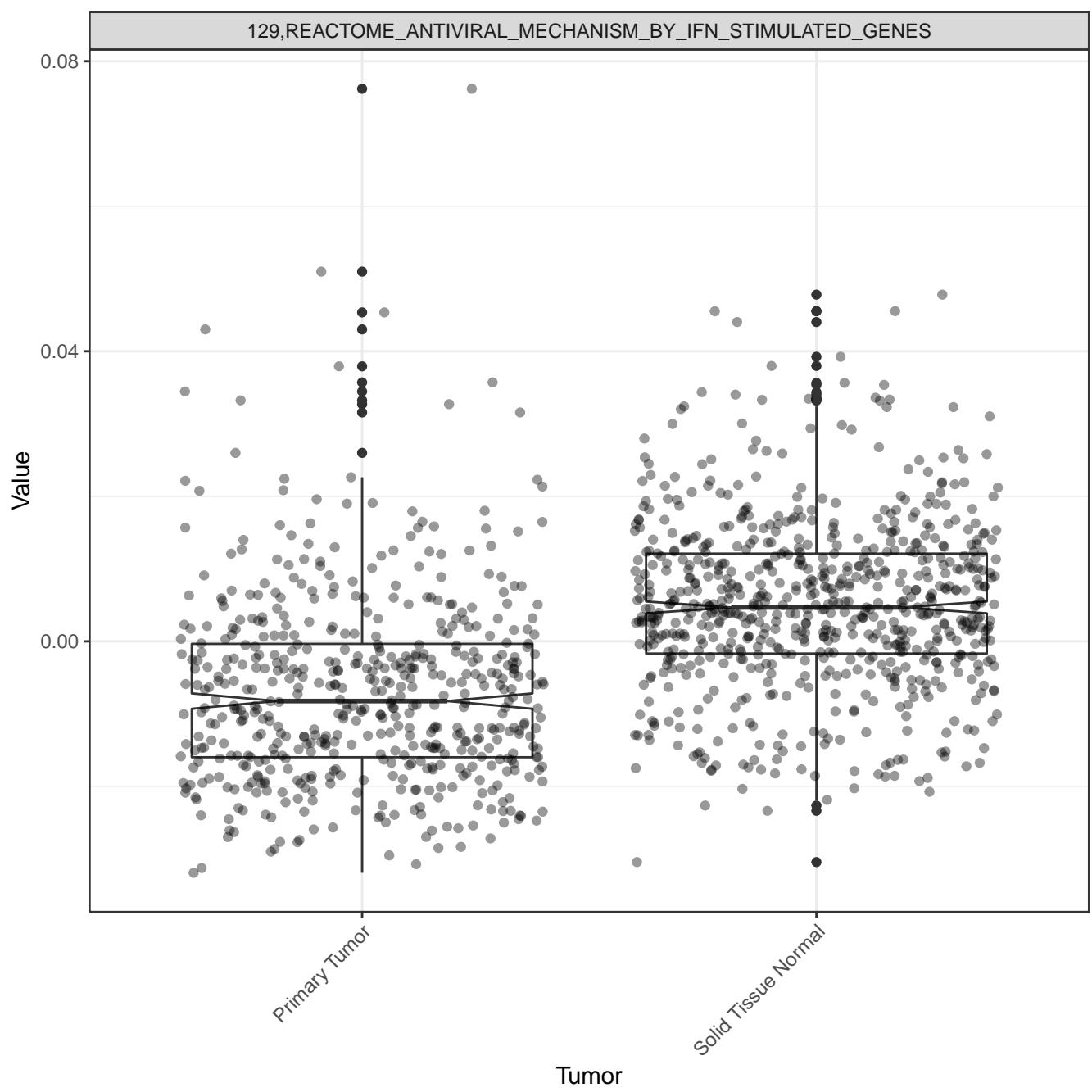
127,KEGG_NEUROACTIVE_LIGAND_RECEPTOR_INTERACTION



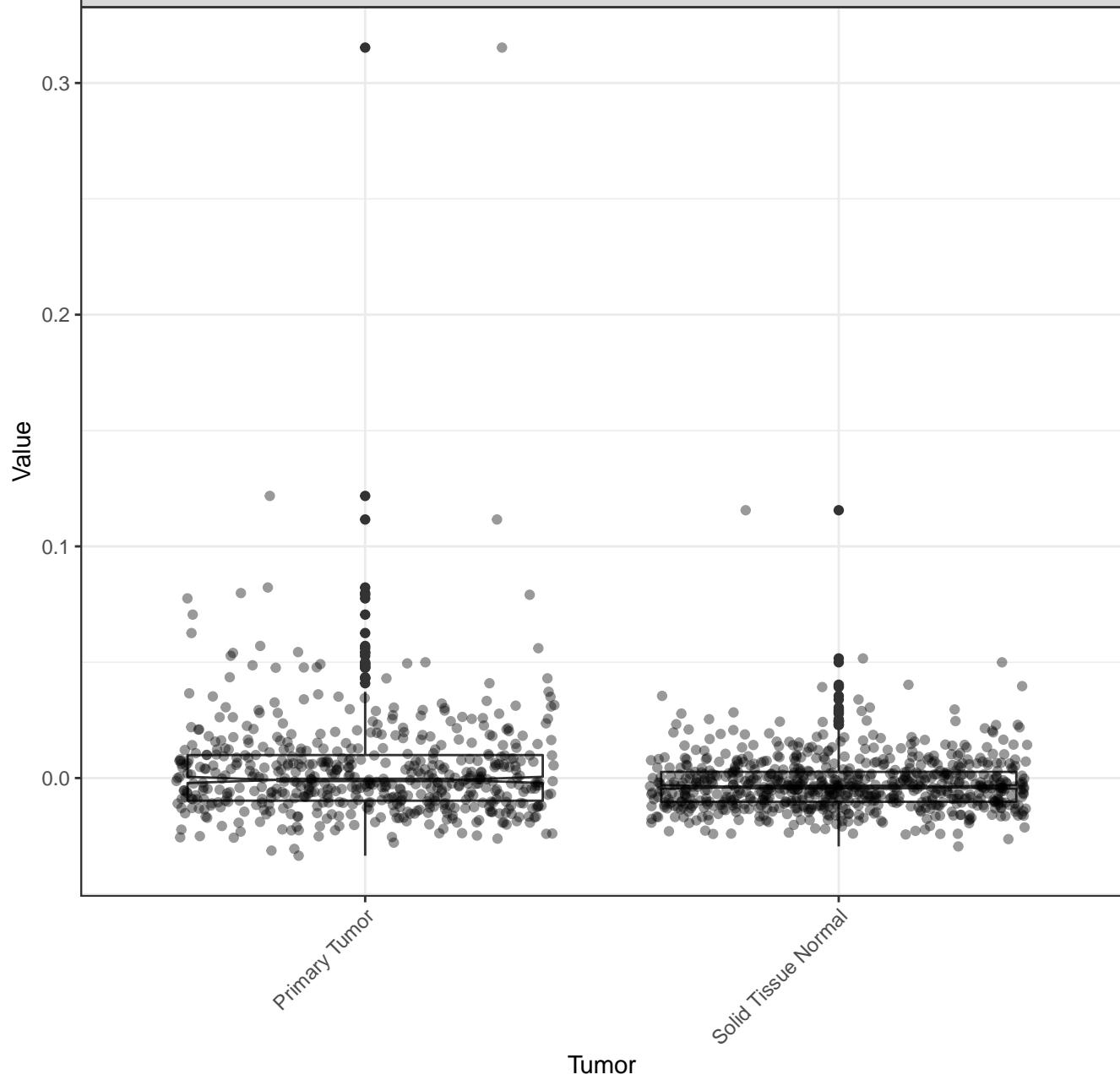
128,KEGG_PURINE_METABOLISM



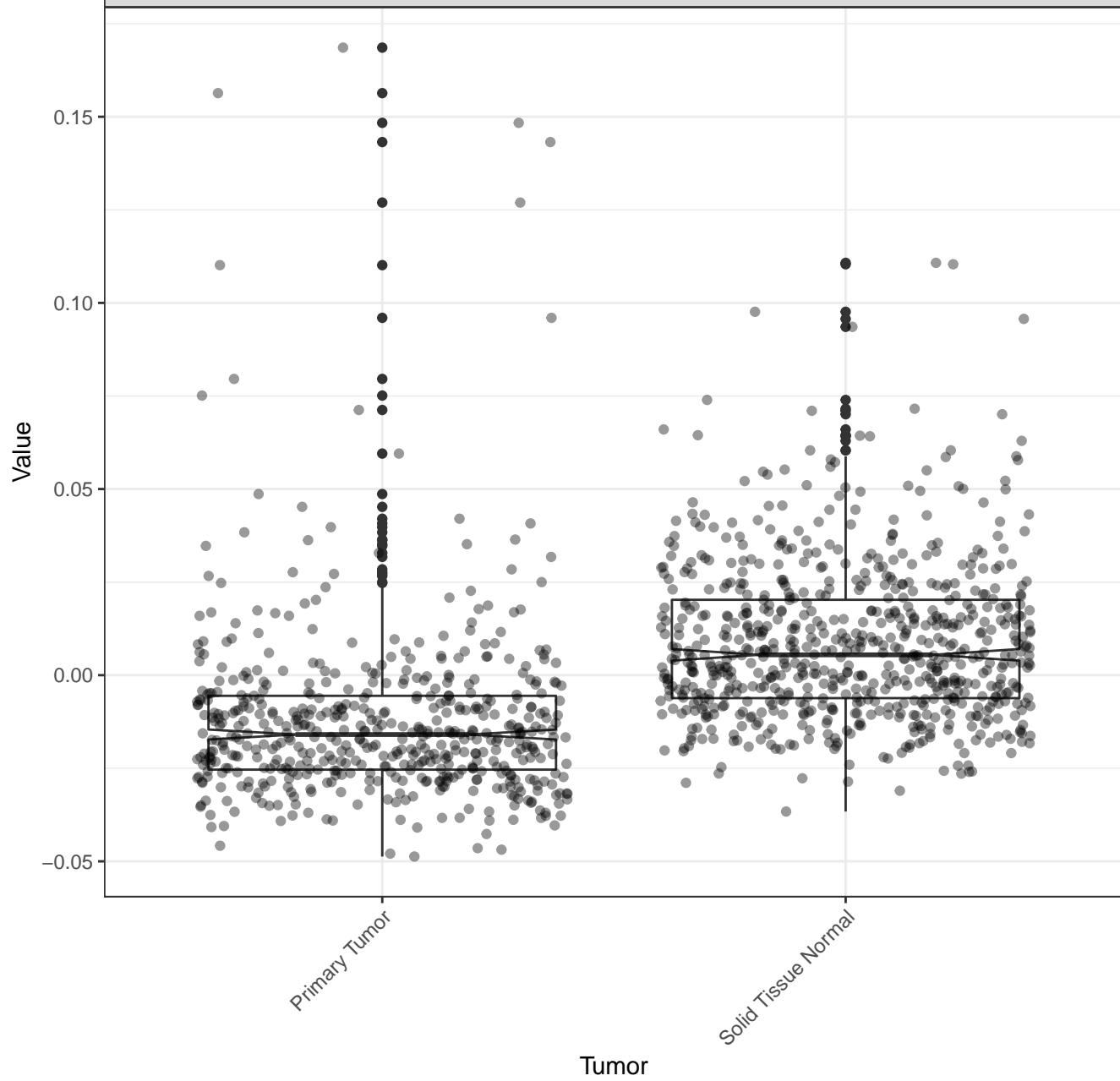
129,REACTOME_ANTIVIRAL_MECHANISM_BY_IFN_STIMULATED_GENES



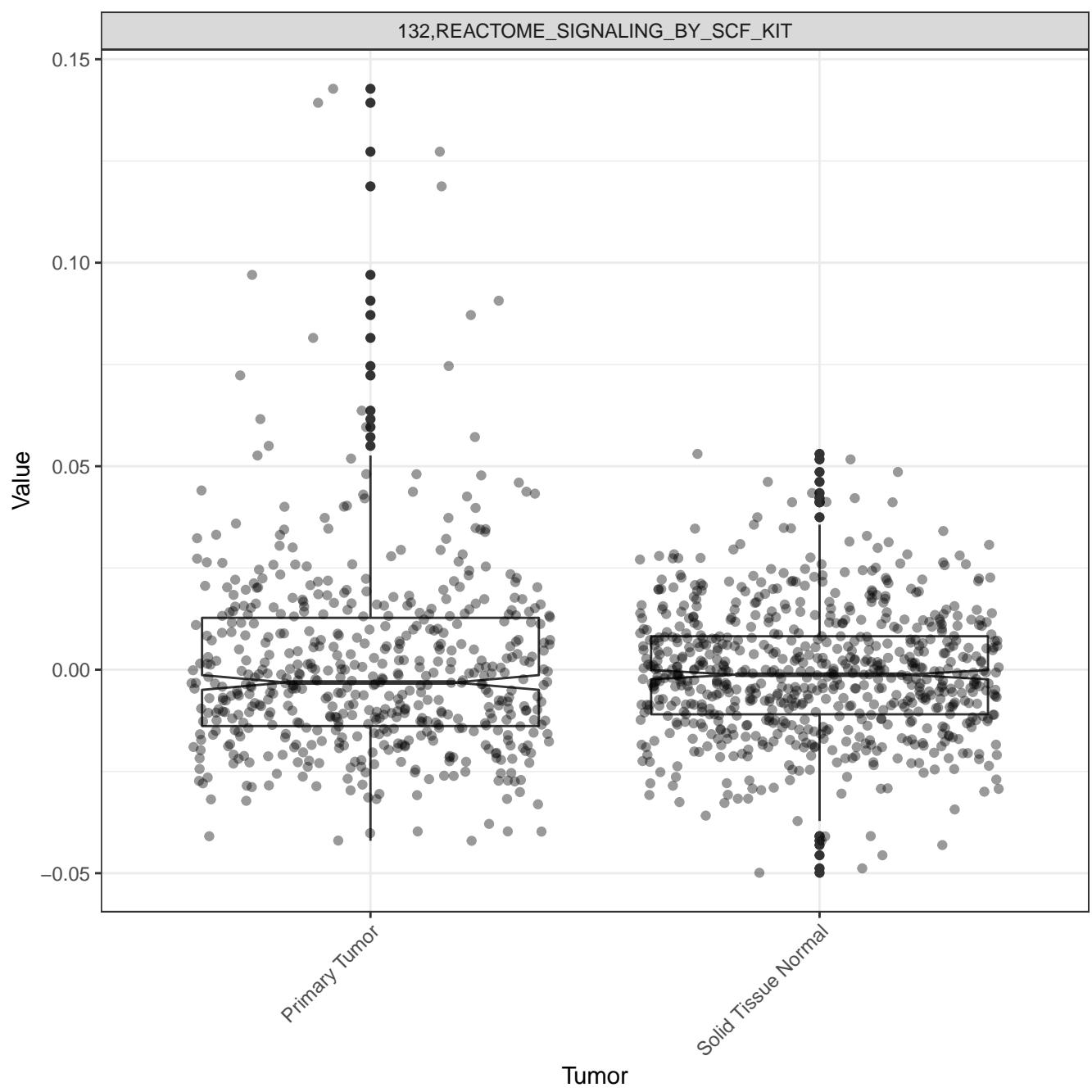
130,REACTOME_AXON_GUIDANCE



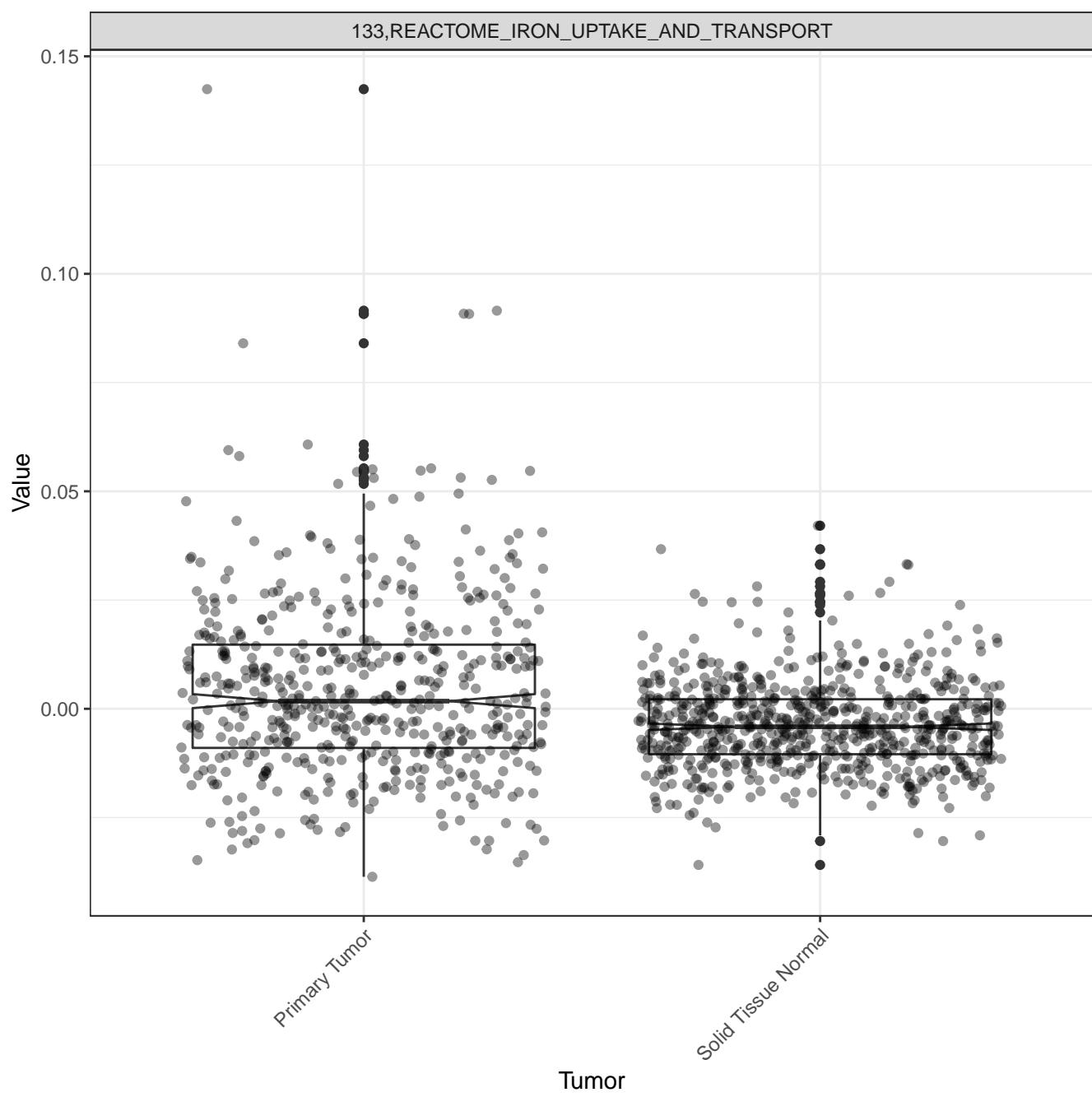
131,REACTOME_OPIOID_SIGNALLING



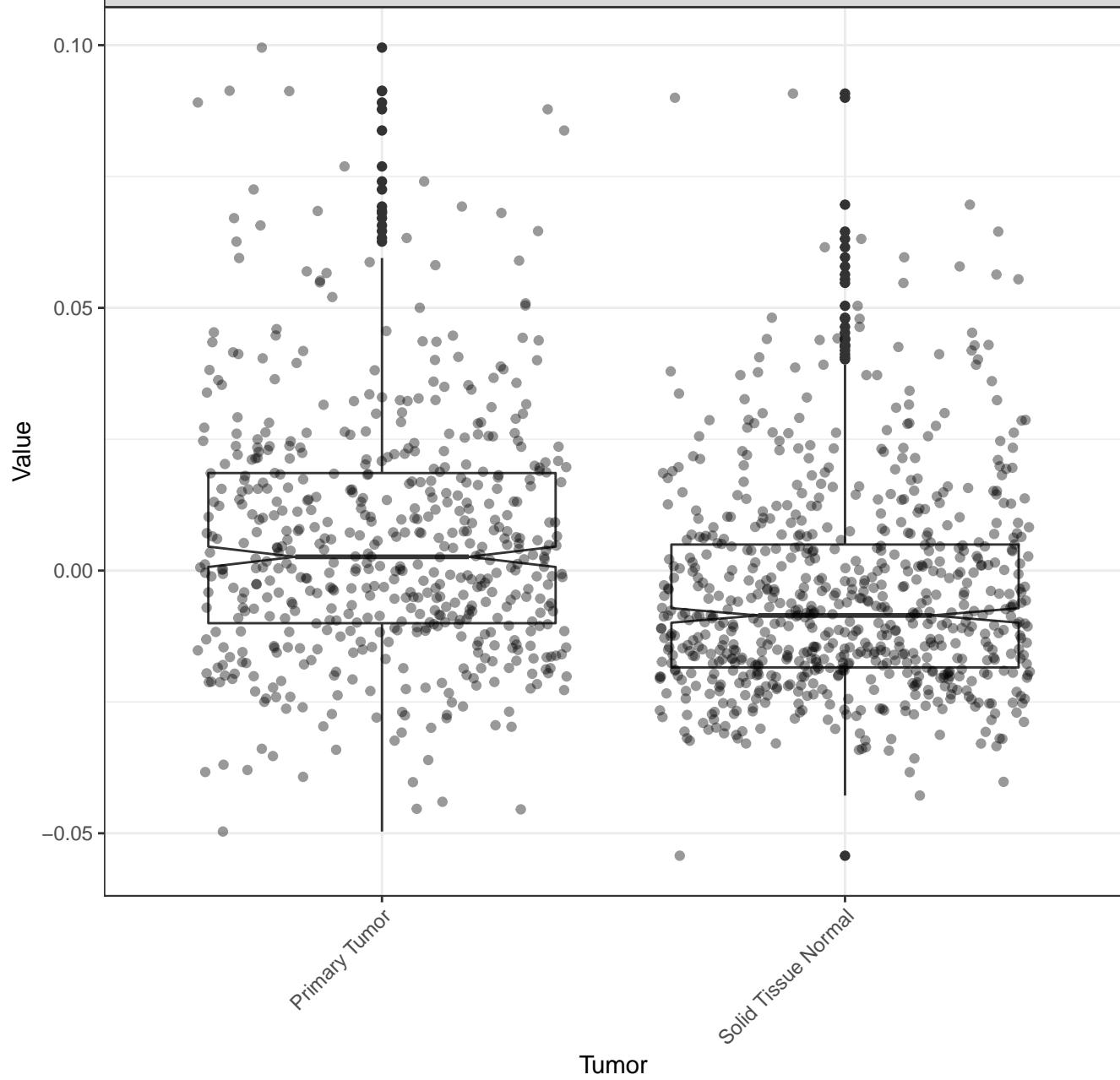
132,REACTOME_SIGNALING_BY_SCF_KIT



133,REACTOME_IRON_UPTAKE_AND_TRANSPORT



134,REACTOME_DIABETES_PATHWAYS



LV 135

Value

0.10

0.05

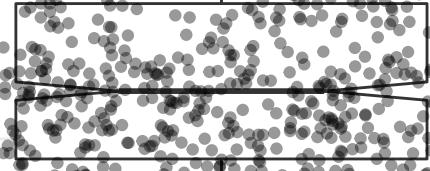
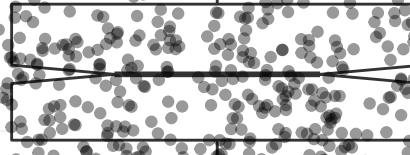
0.00

-0.05

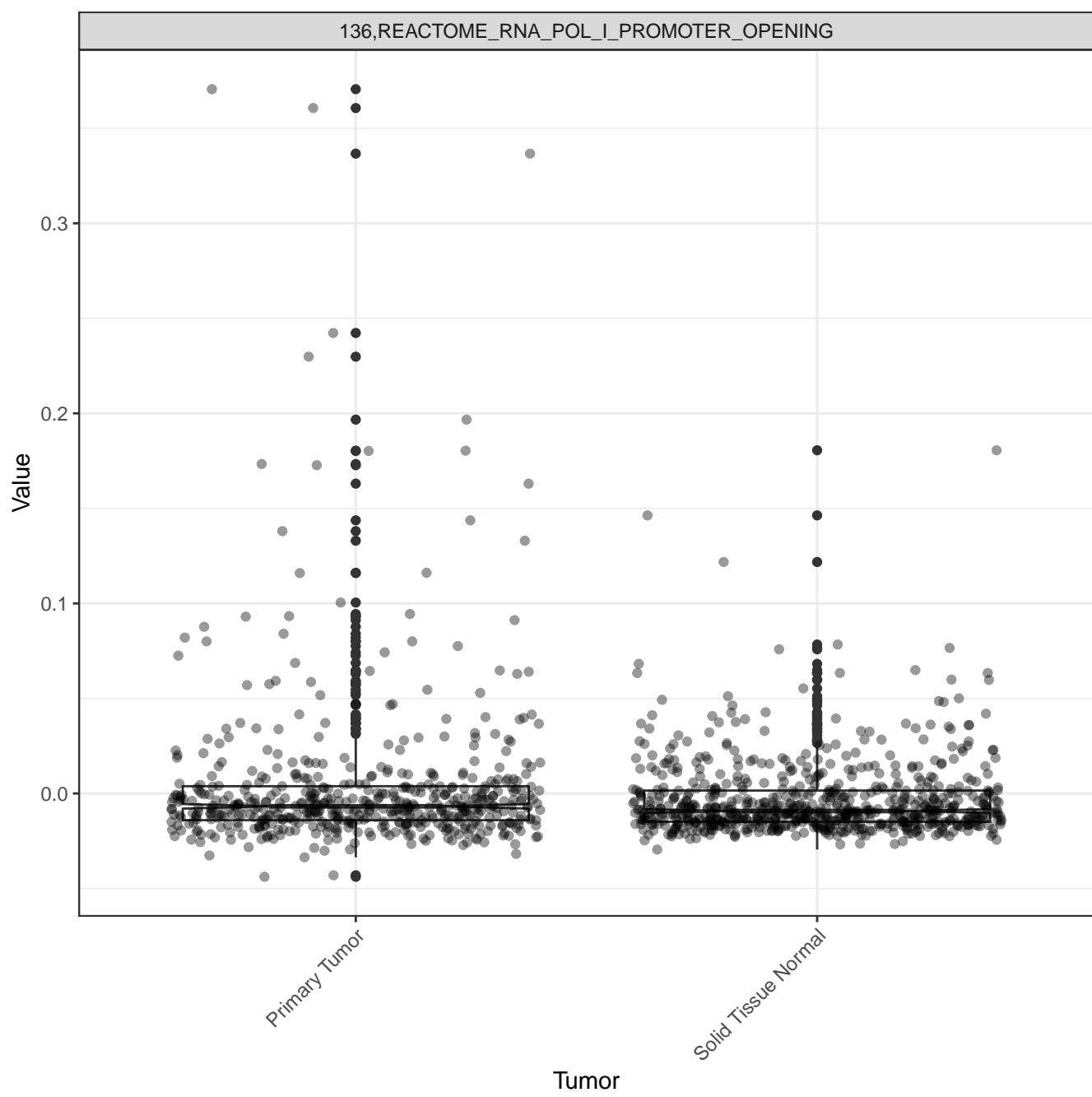
Primary Tumor

Solid Tissue Normal

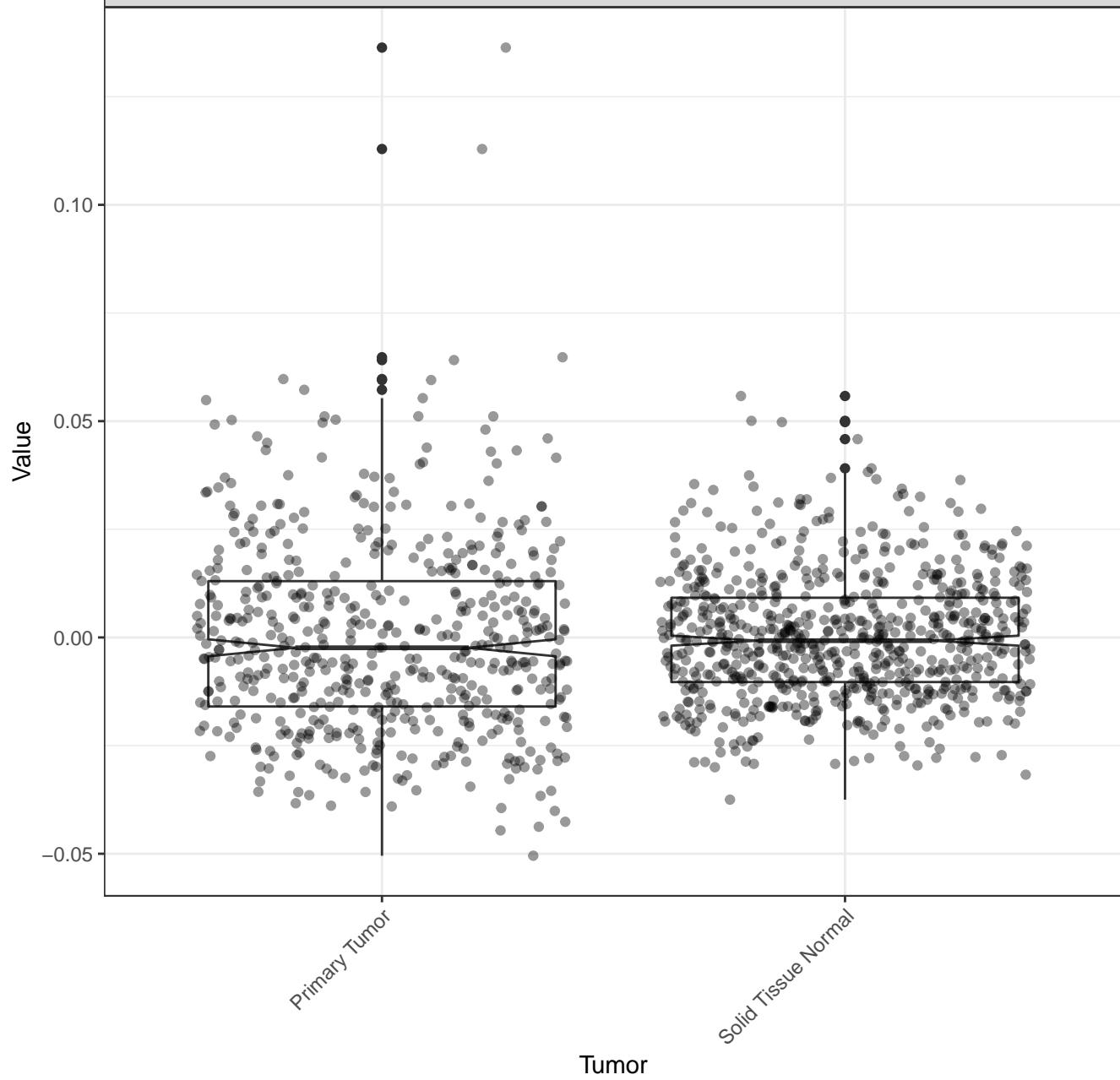
Tumor



136,REACTOME_RNA_POL_I_PROMOTER_OPENING



137,REACTOME_GOLGI_ASSOCIATED_VESICLE_BIOGENESIS



138,IRIS_CD4Tcell-Th2-restimulated12hour

Value

0.10

0.05

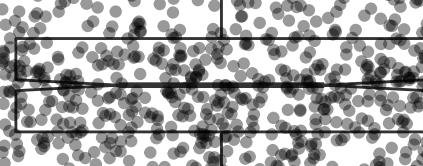
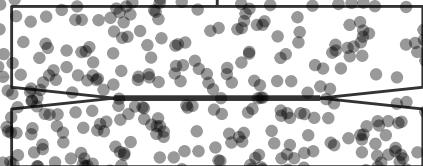
0.00

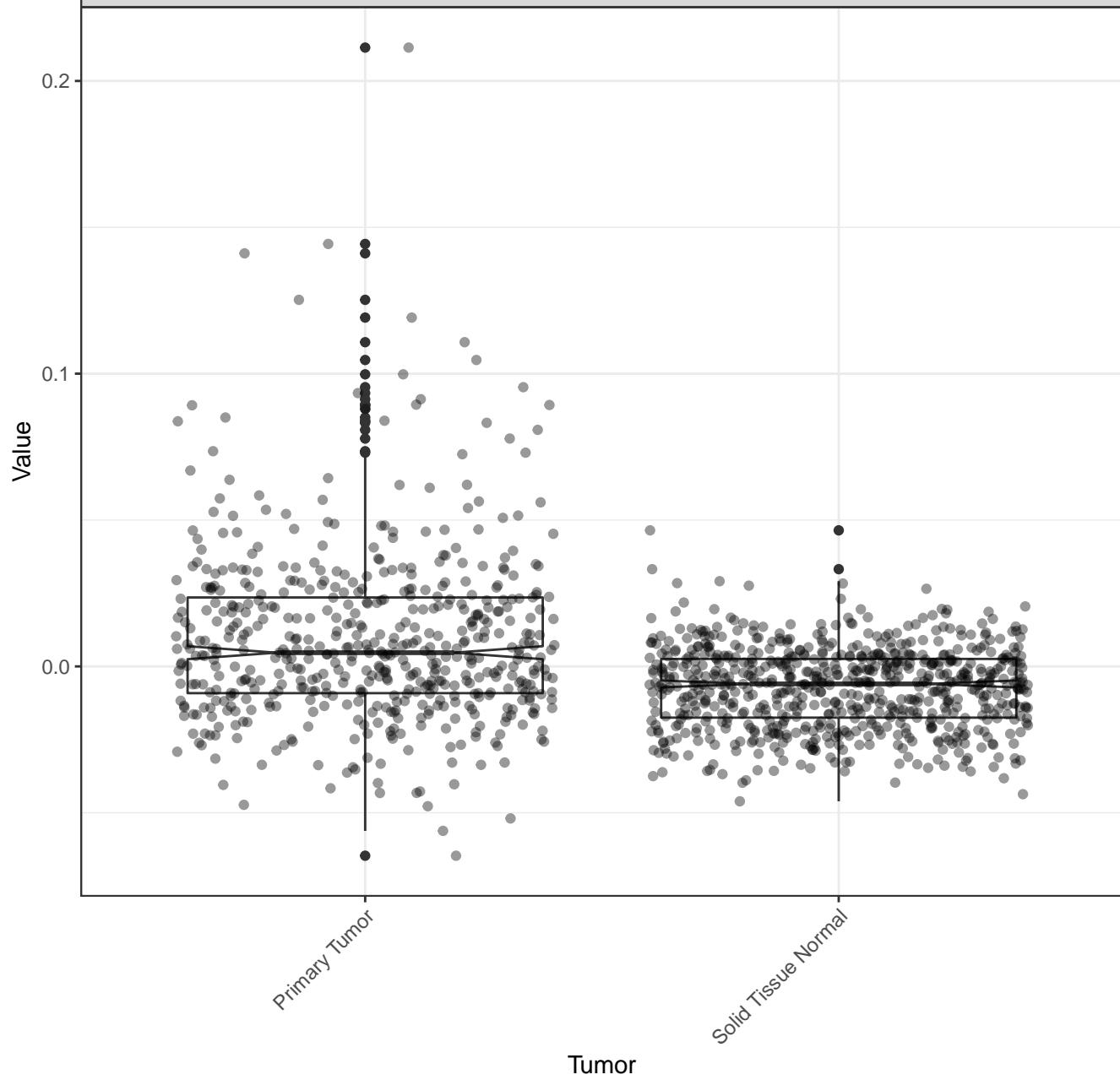
-0.05

Primary Tumor

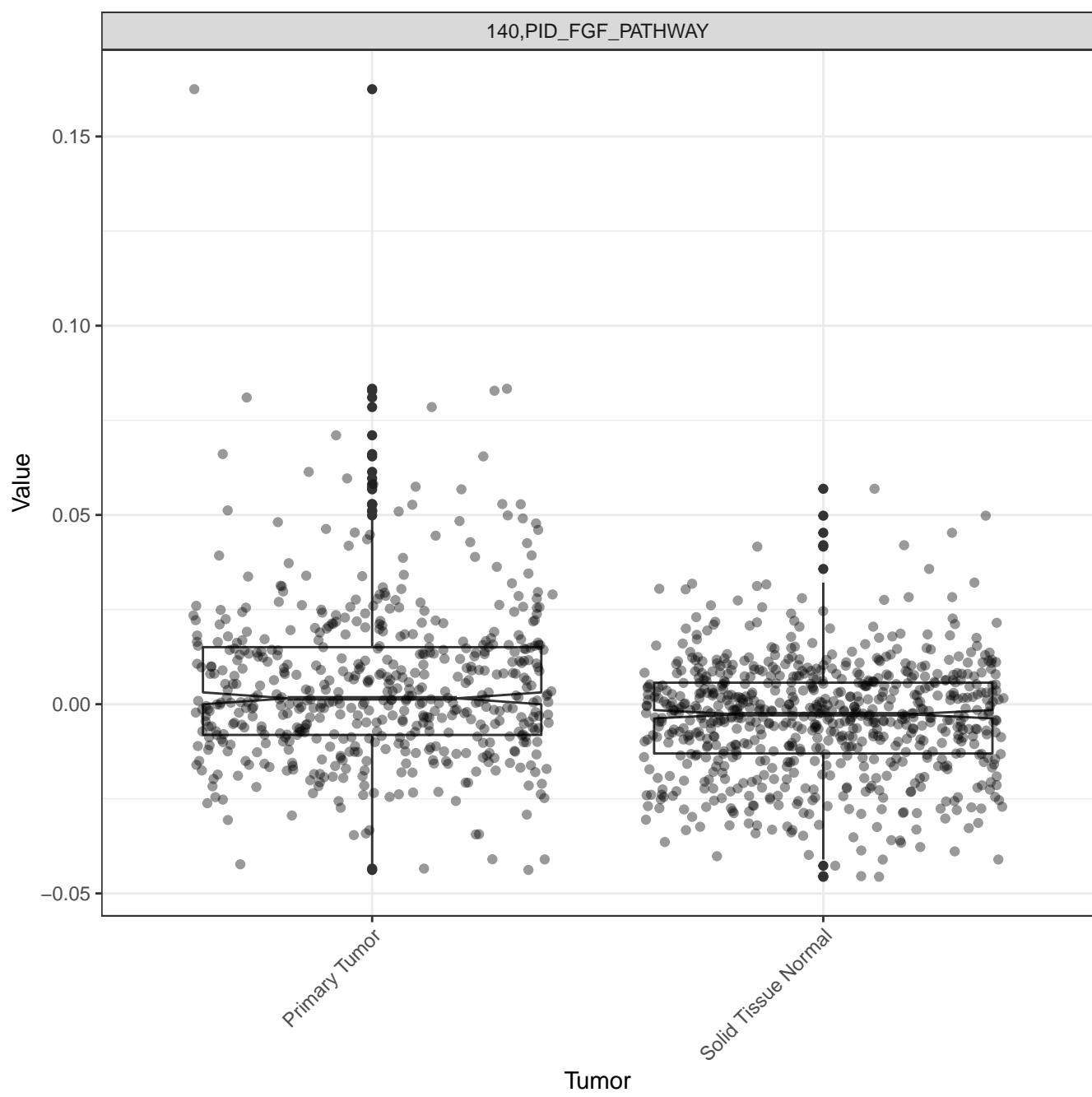
Solid Tissue Normal

Tumor

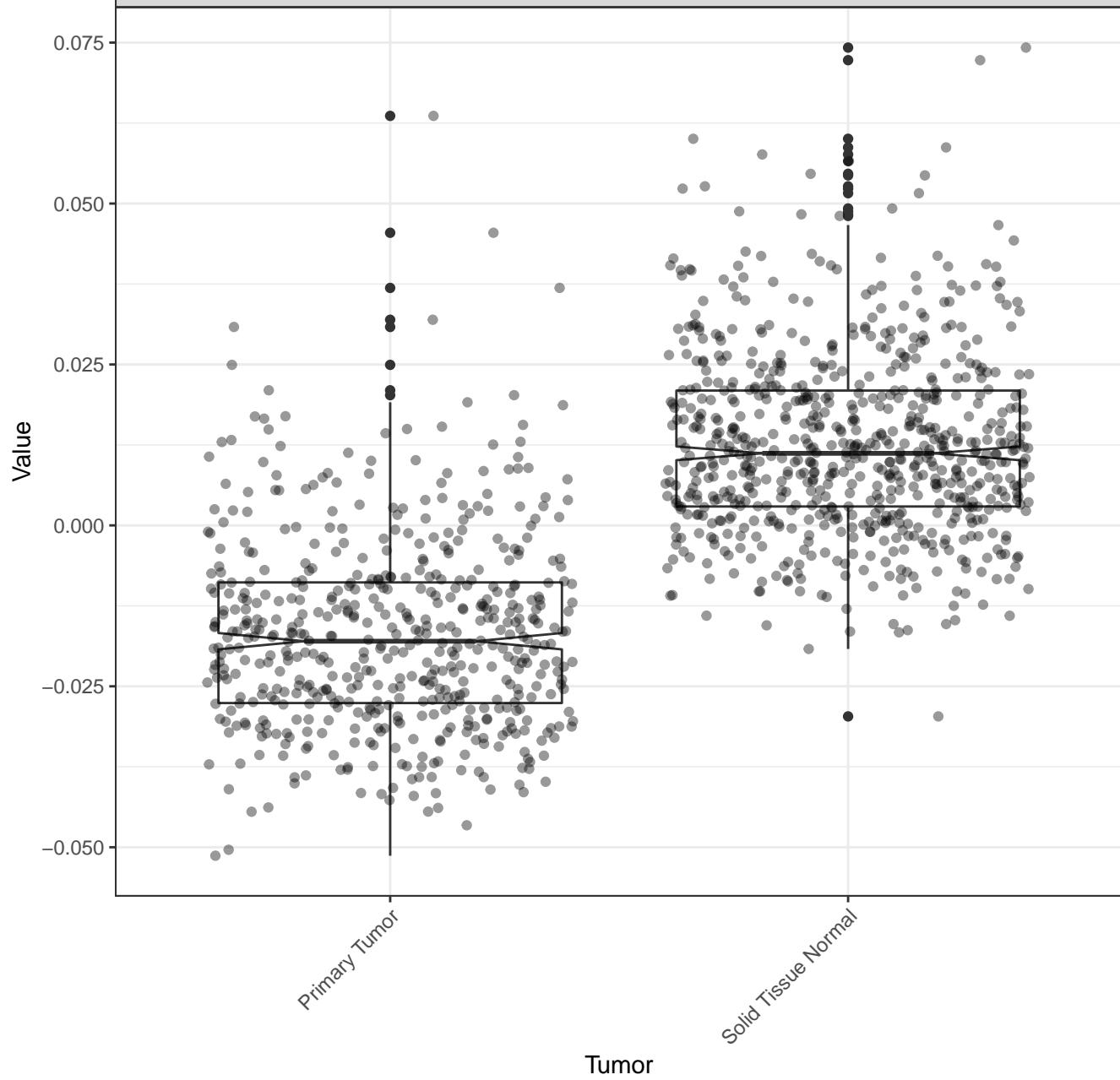




140,PID_FGF_PATHWAY



141,DMAP_ERY4



142,KEGG_METABOLISM_OF_XENOBIOTICS_BY_CYTOCHROME_P450

Value

0.10

0.05

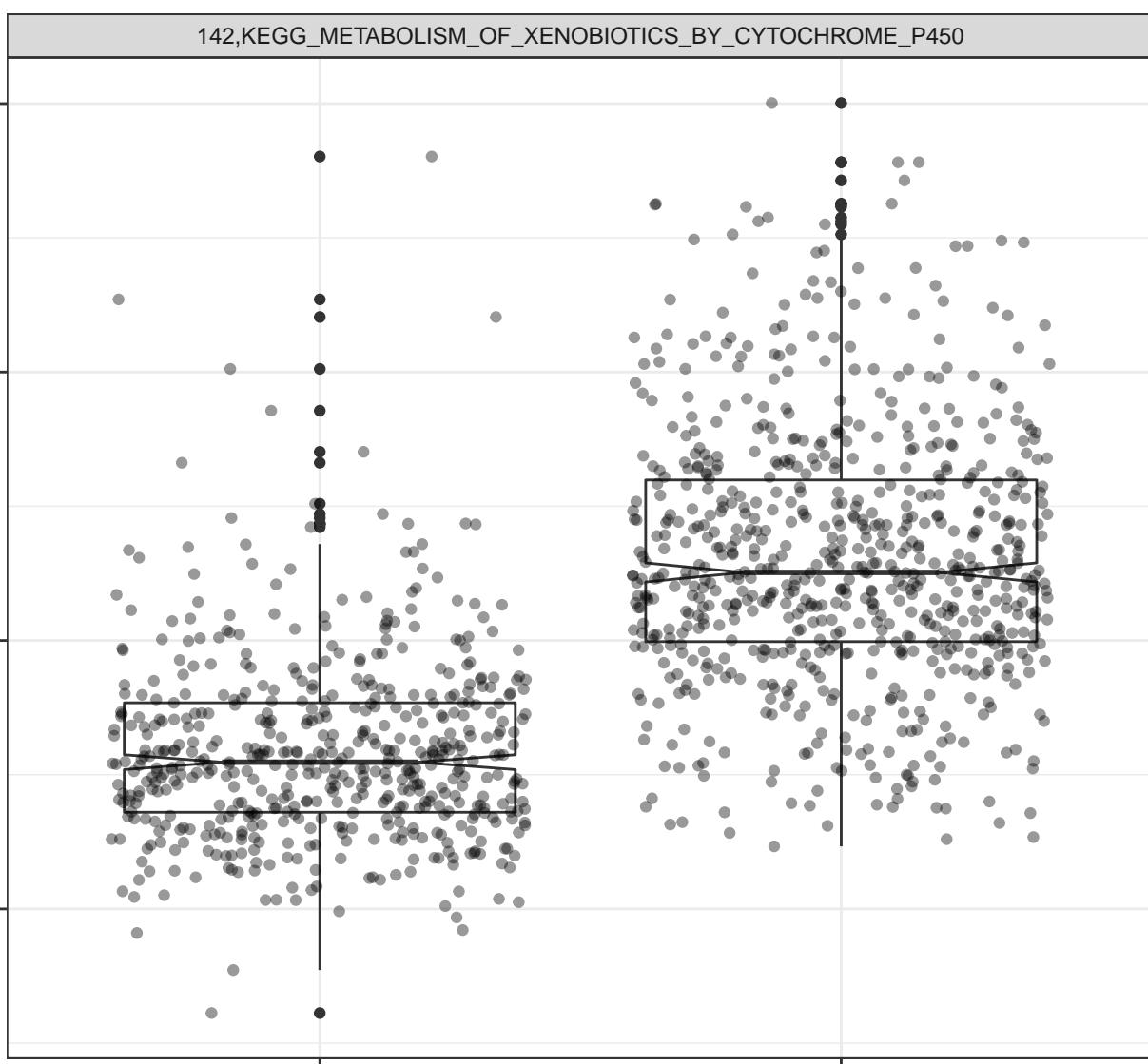
0.00

-0.05

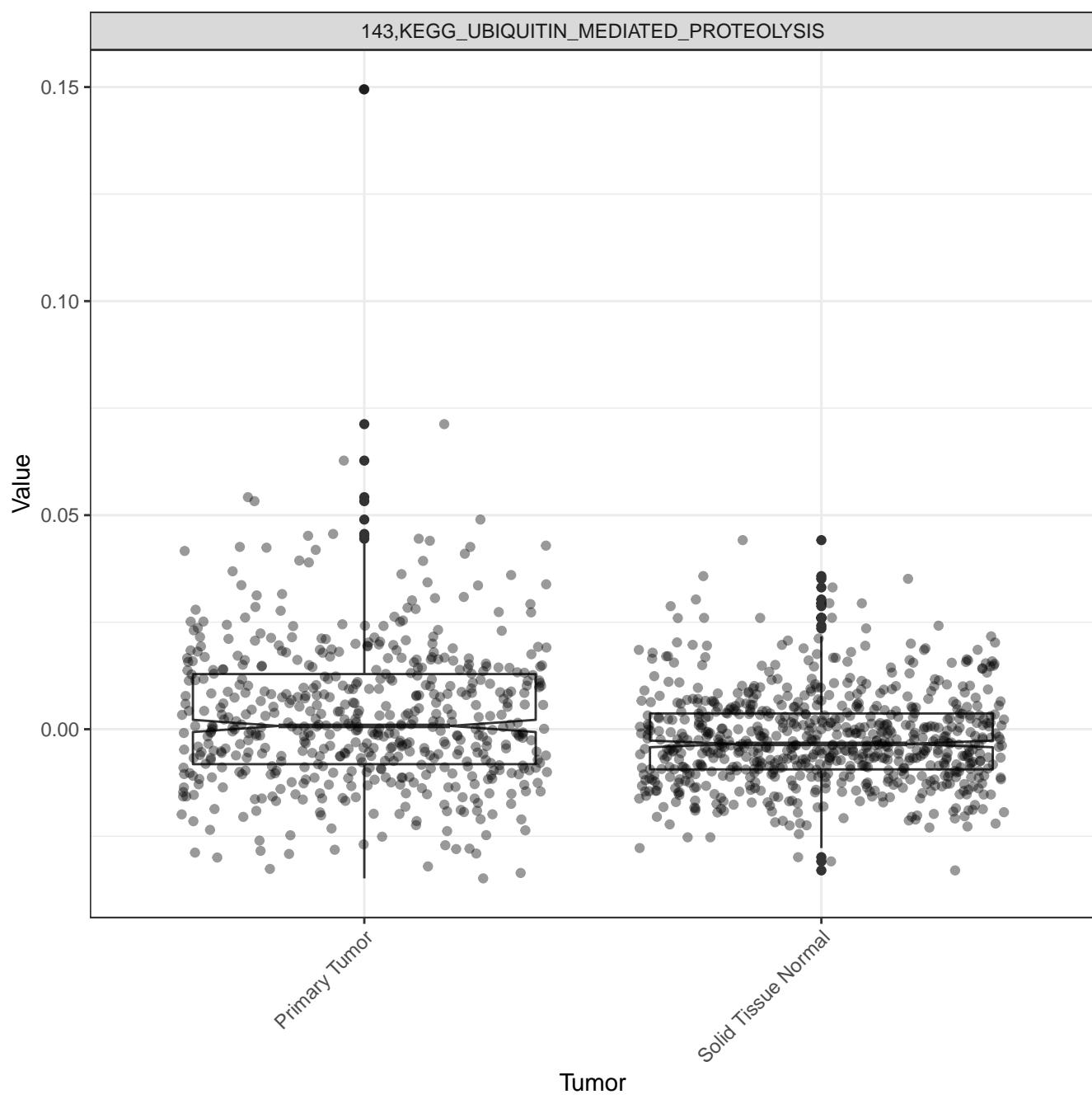
Primary Tumor

Solid Tissue Normal

Tumor



143,KEGG_UBIQUITIN_MEDiated_PROTEOLYSIS



144,REACTOME_SIGNALING_BY_RHO_GTPASES

Value

0.10

0.05

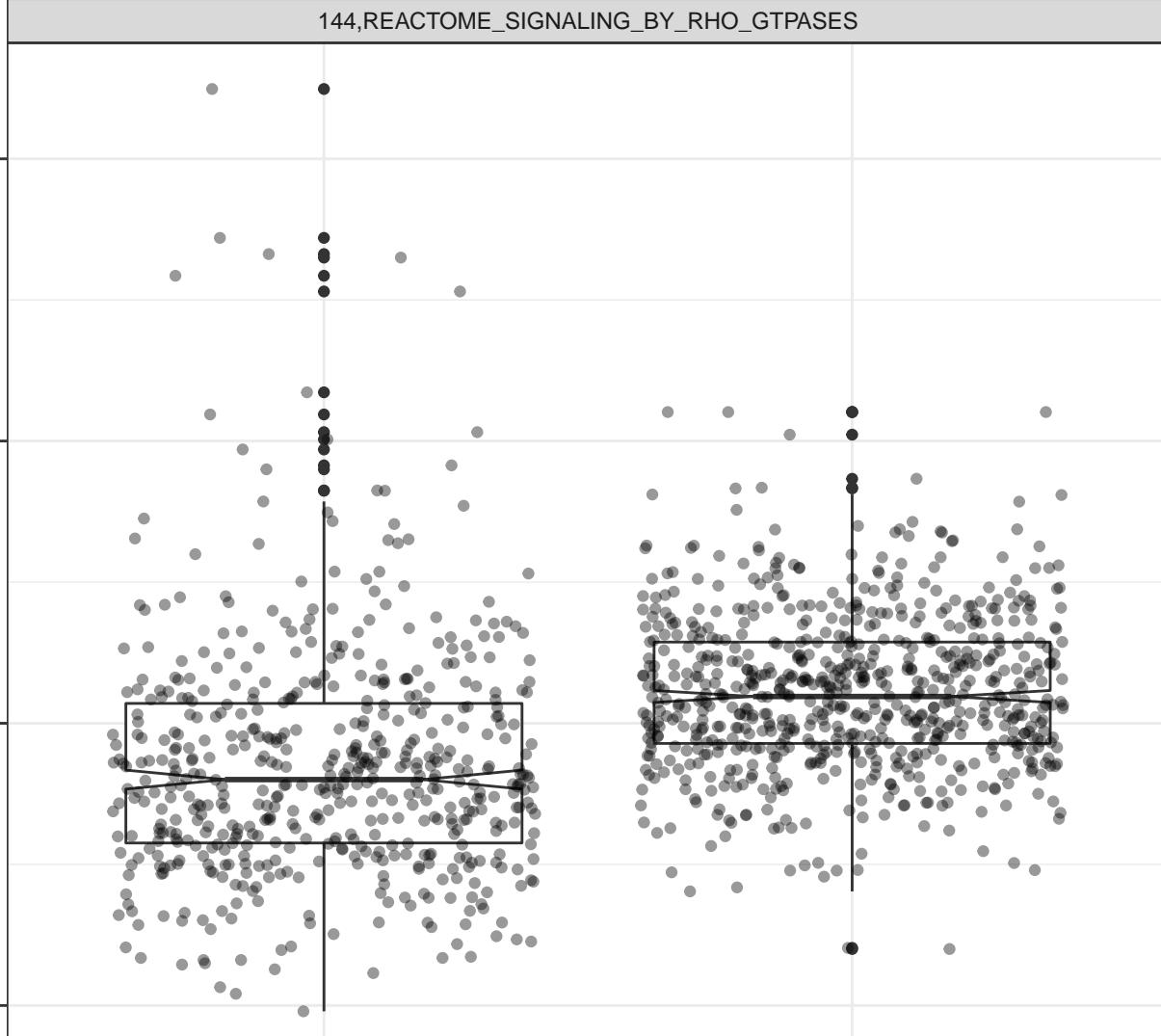
0.00

-0.05

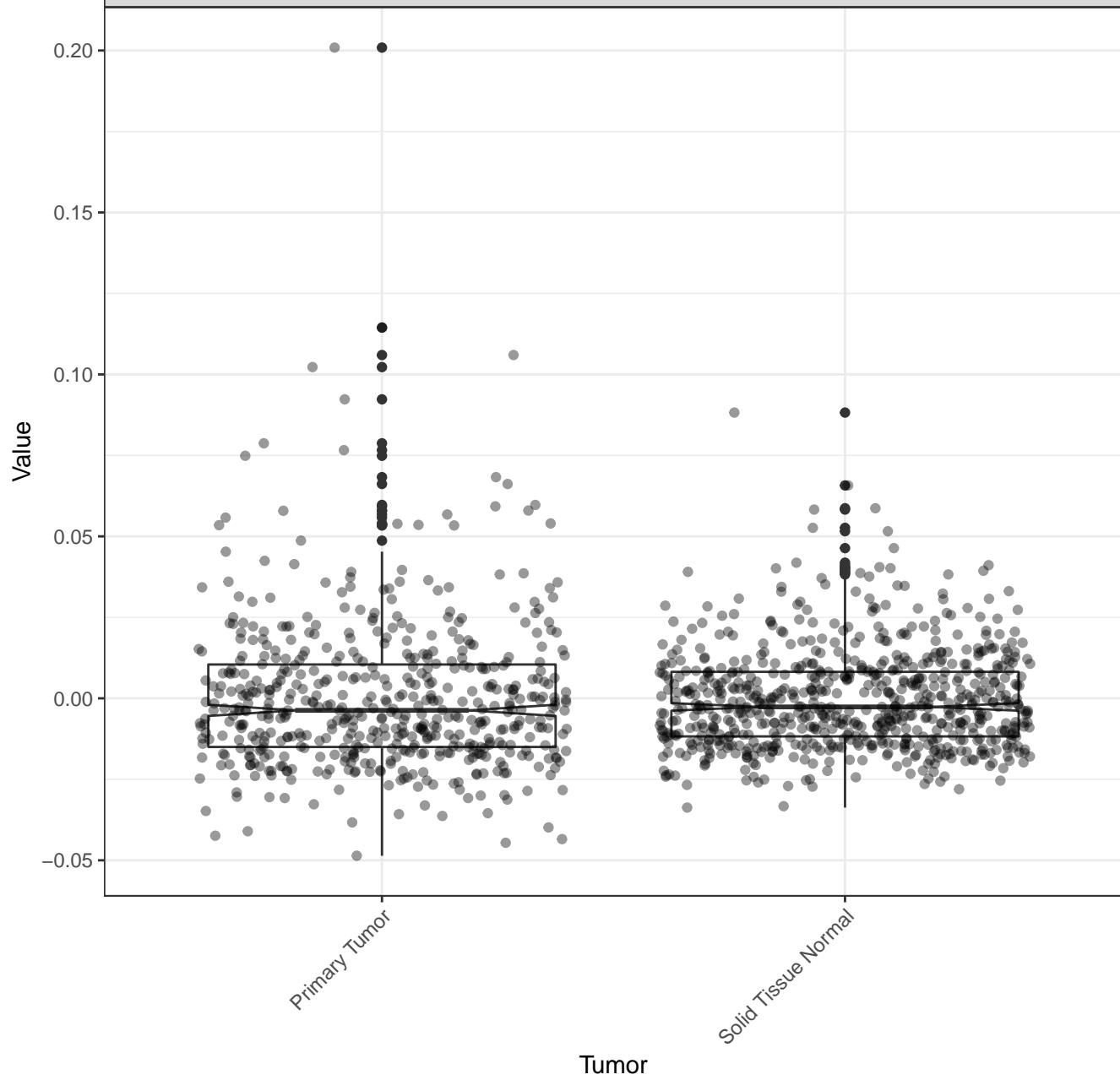
Primary Tumor

Solid Tissue Normal

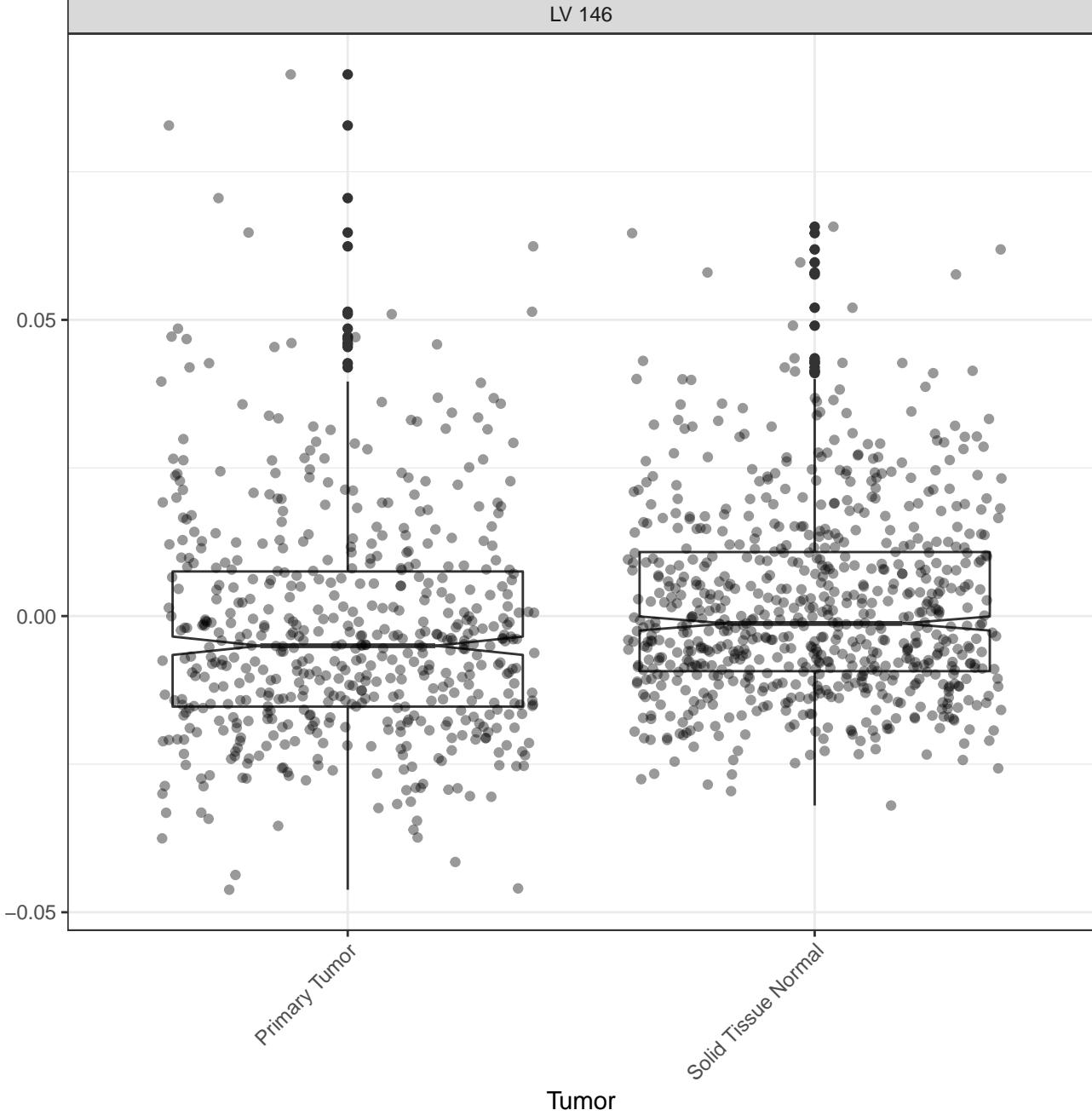
Tumor



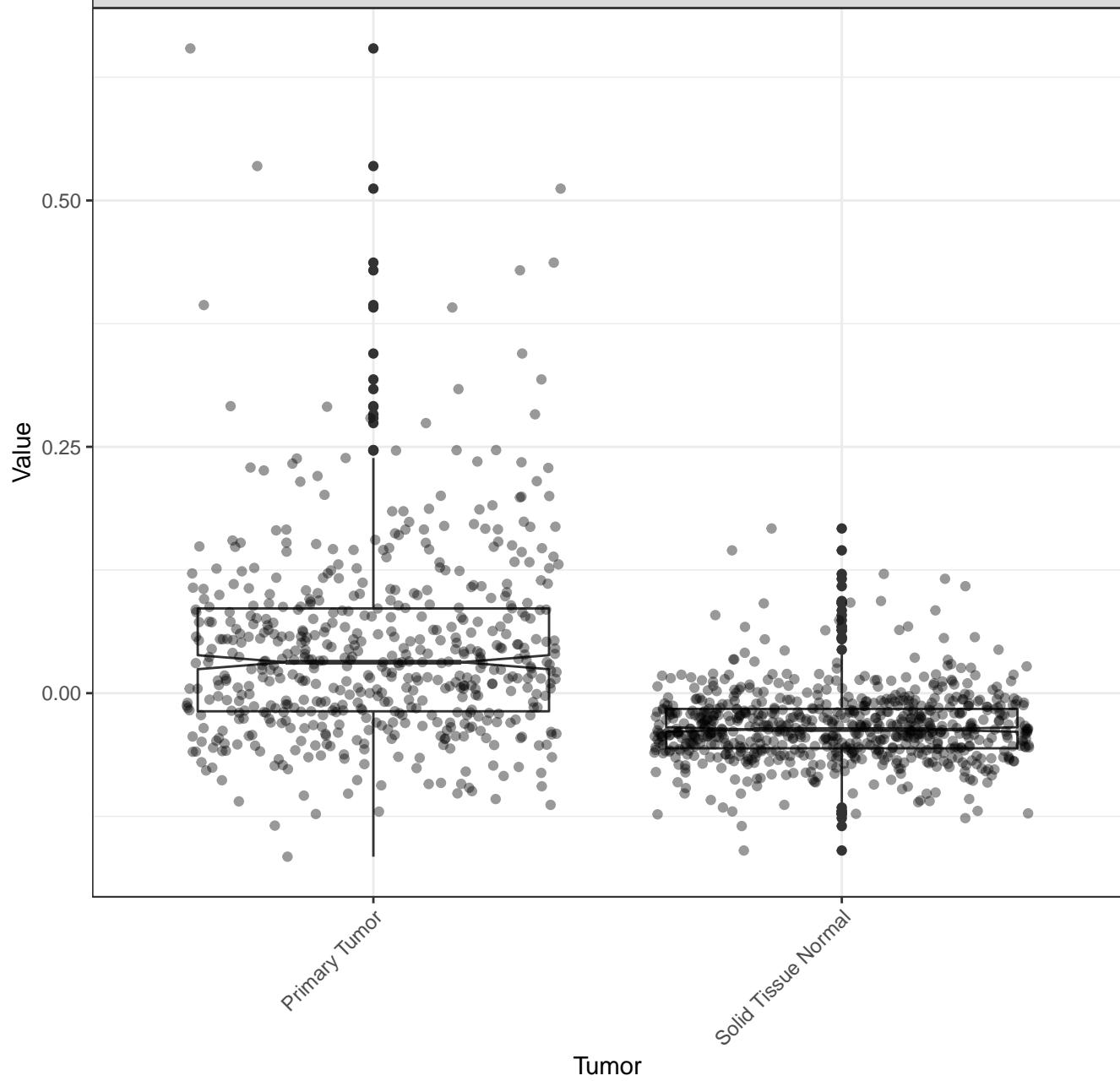
145,REACTOME_SIGNALING_BY_ILS

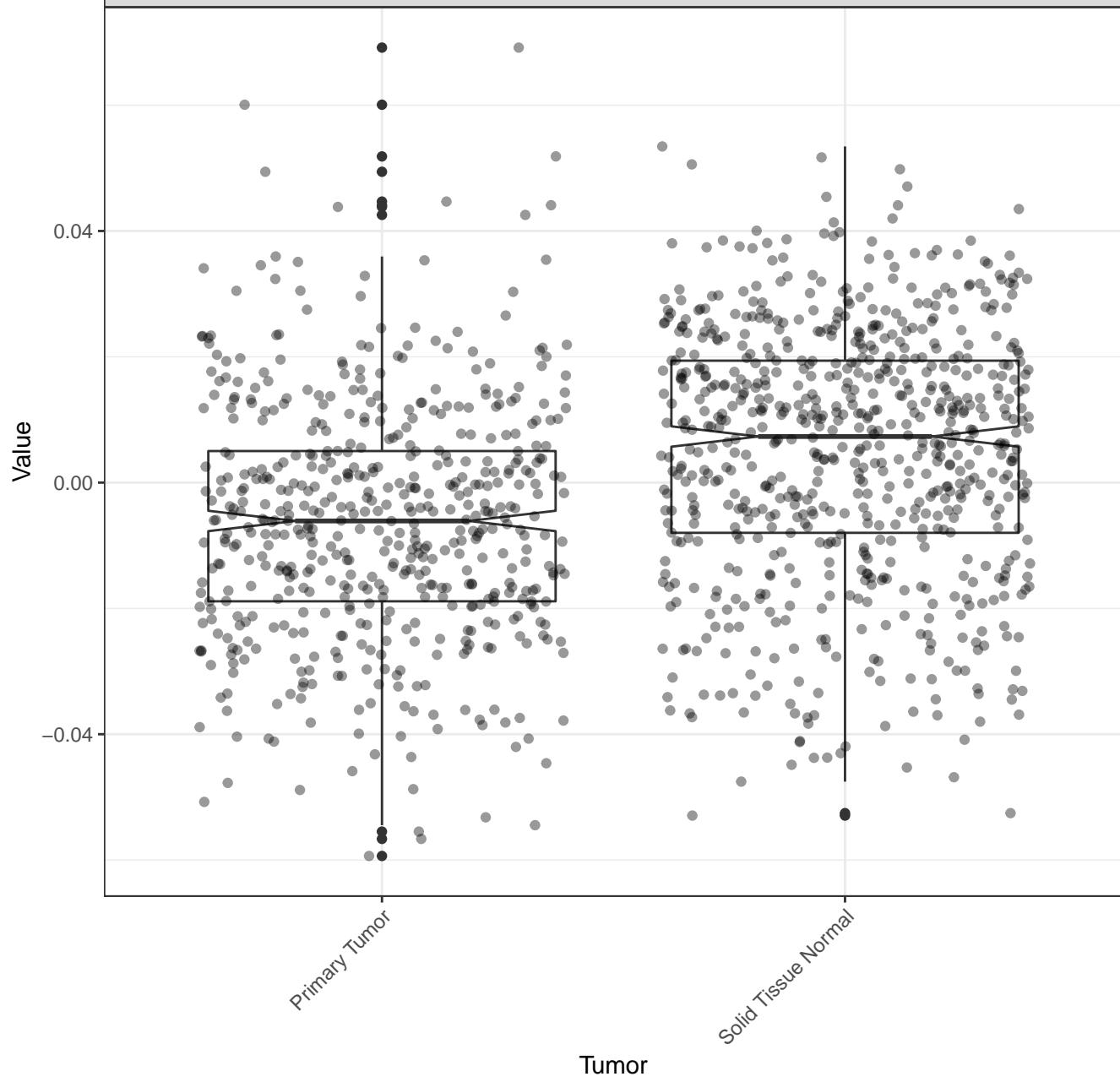


Value

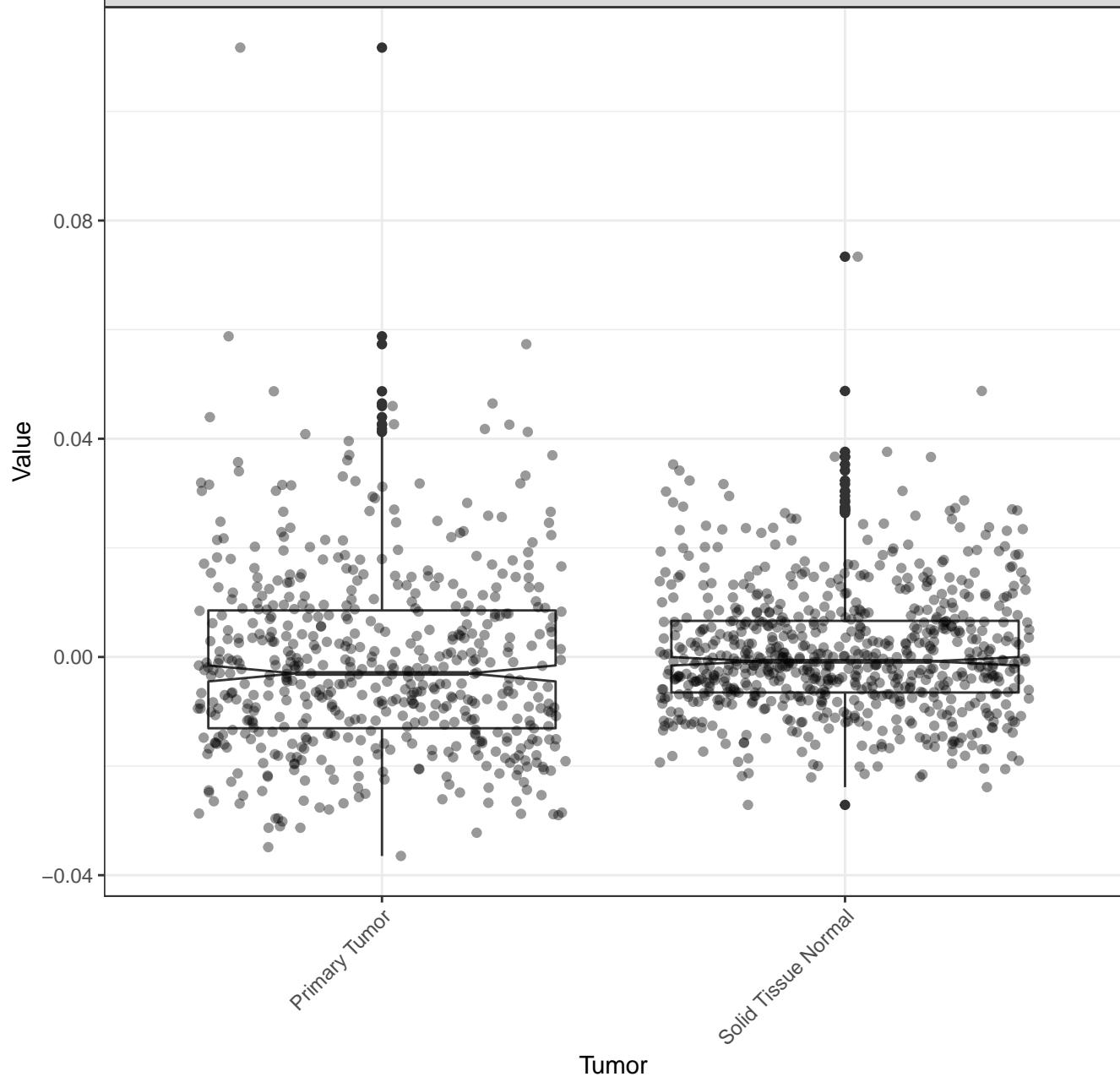


147,PID_AURORA_B_PATHWAY





149,SIG_CD40PATHWAYMAP



150,REACTOME_ANTIGEN_PROCESSING_UBIQUITINATION_PROTEASOME_DEGRADATION

Value

0.05

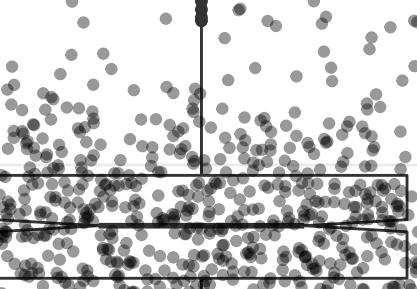
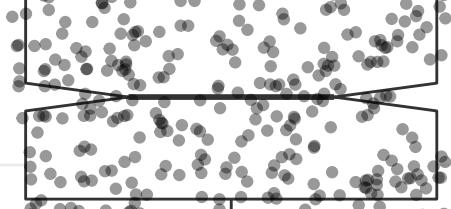
0.00

-0.05

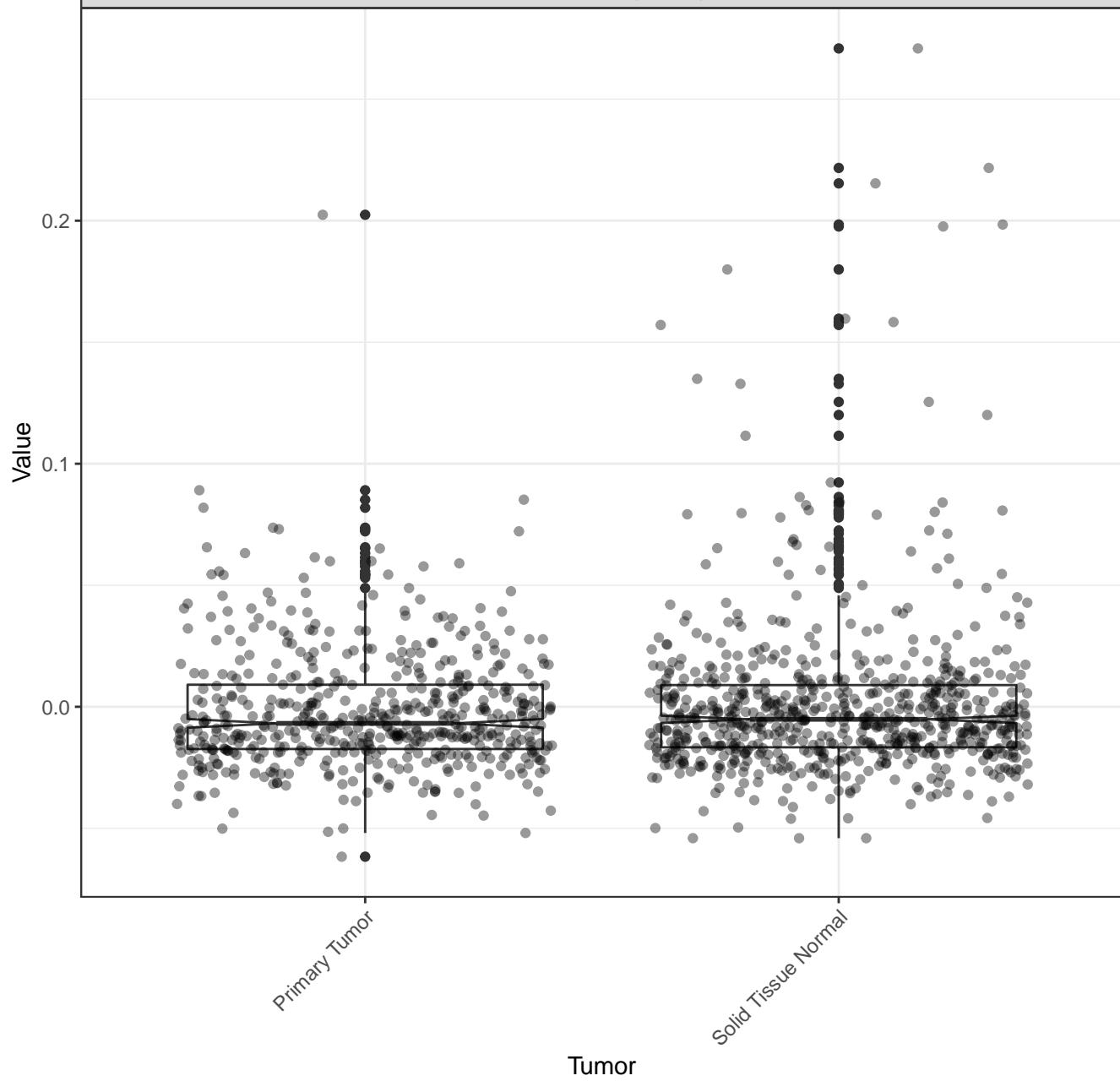
Primary Tumor

Tumor

Solid Tissue Normal



151,IRIS_Monocyte-Day1



152,KEGG_ARRHYTHMOGENIC_RIGHT_VENTRICULAR_CARDIOMYOPATHY_ARVC

Value

0.10

0.05

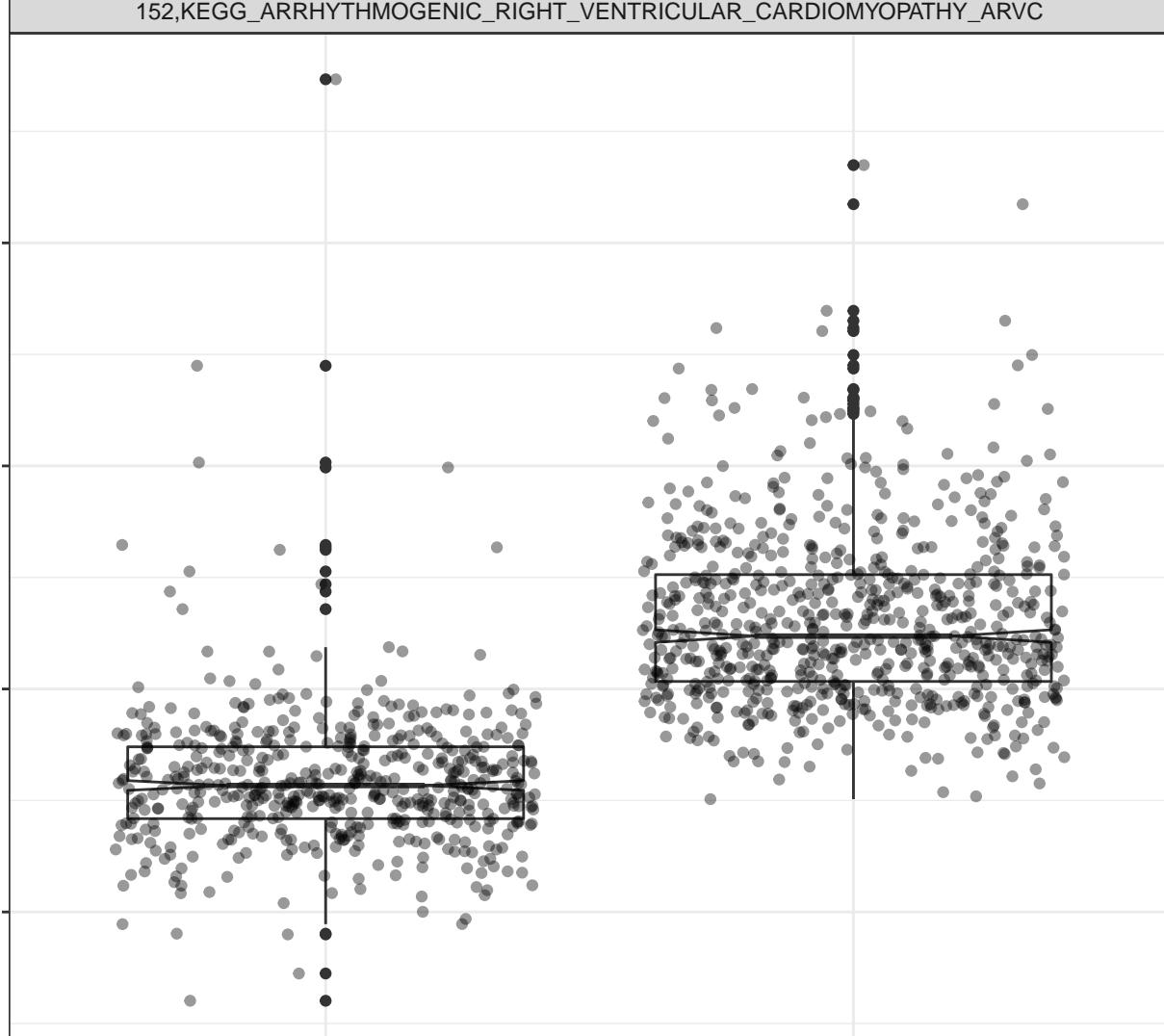
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 153

Value

0.15

0.10

0.05

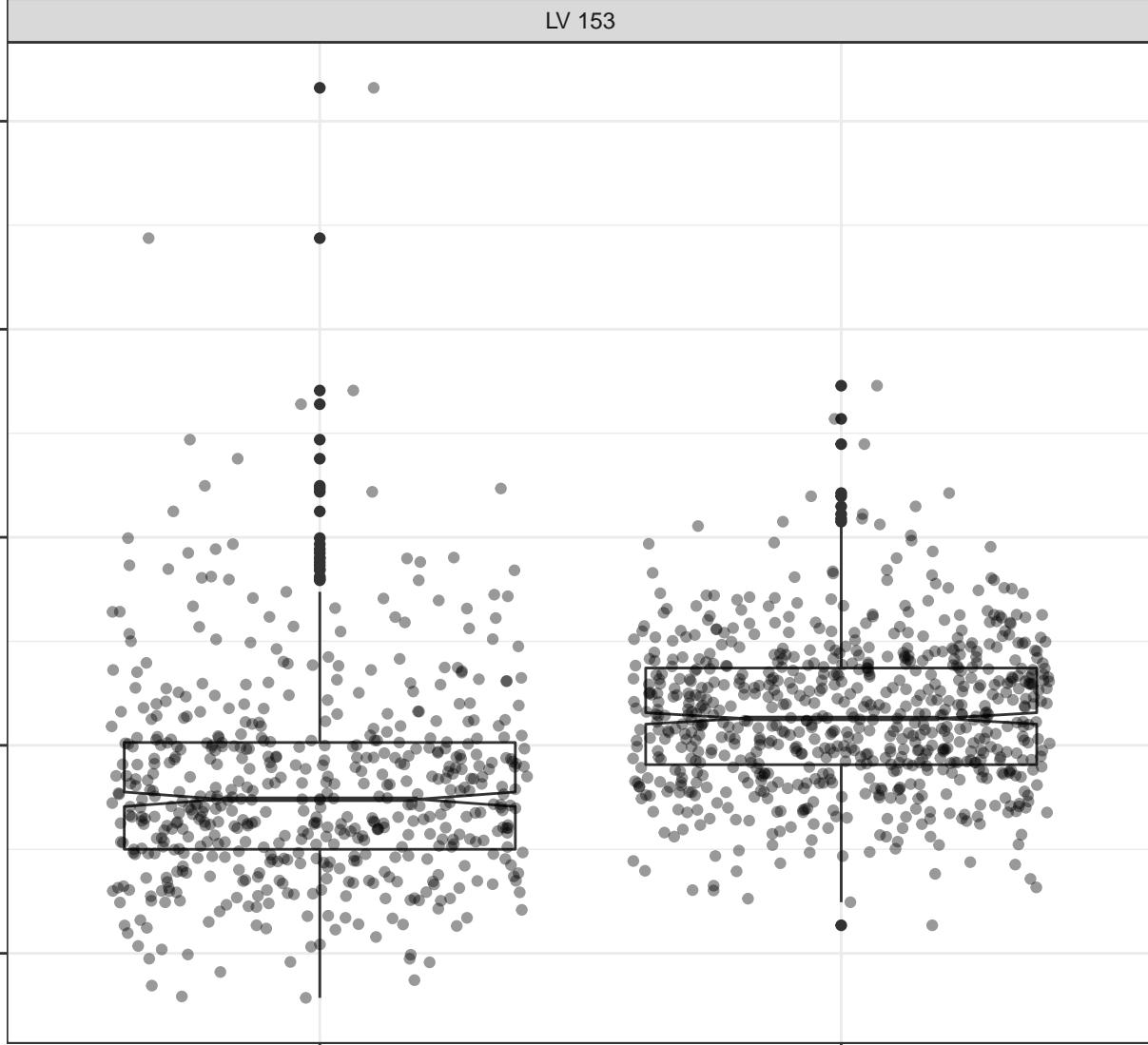
0.00

-0.05

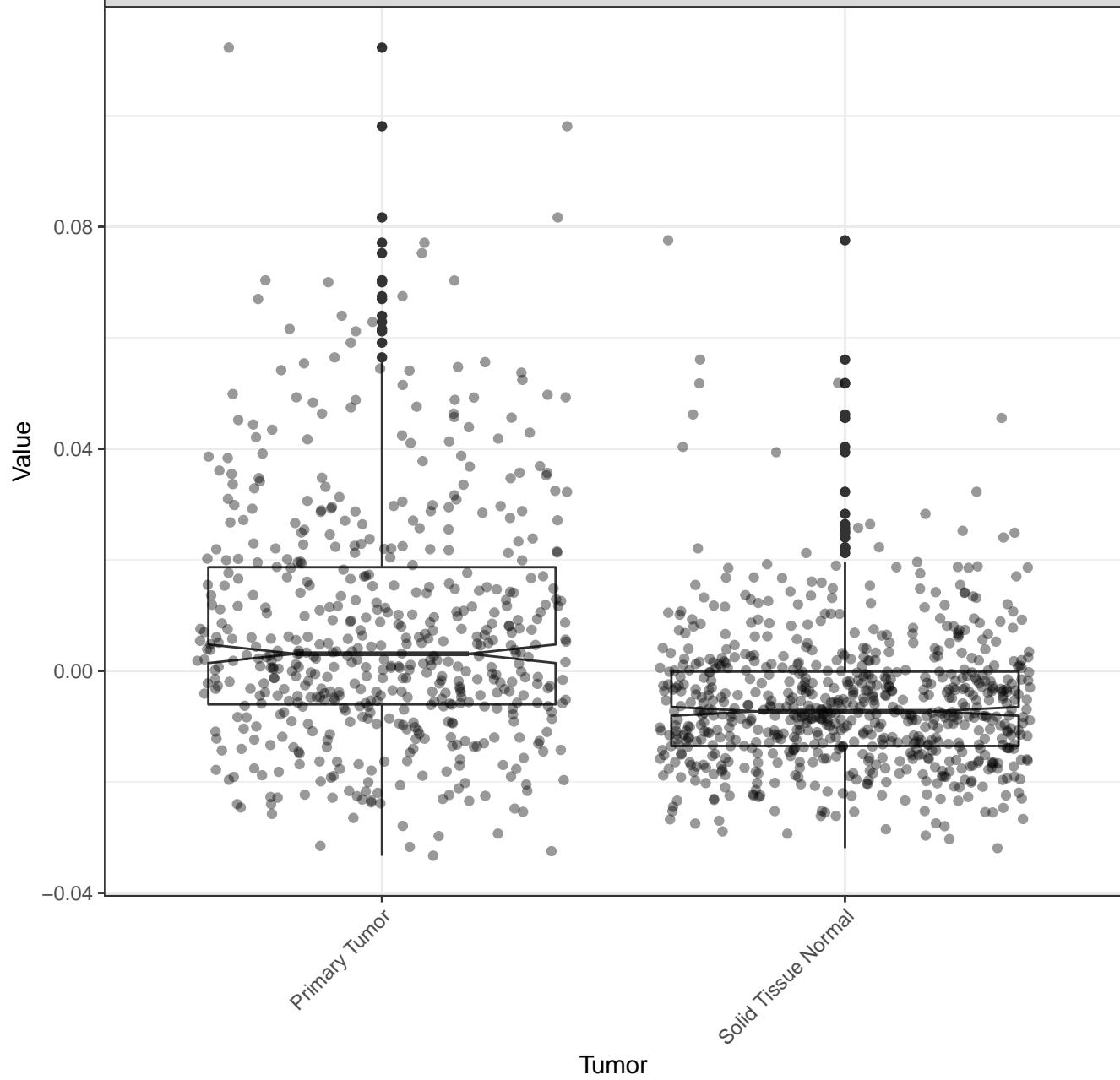
Primary Tumor

Solid Tissue Normal

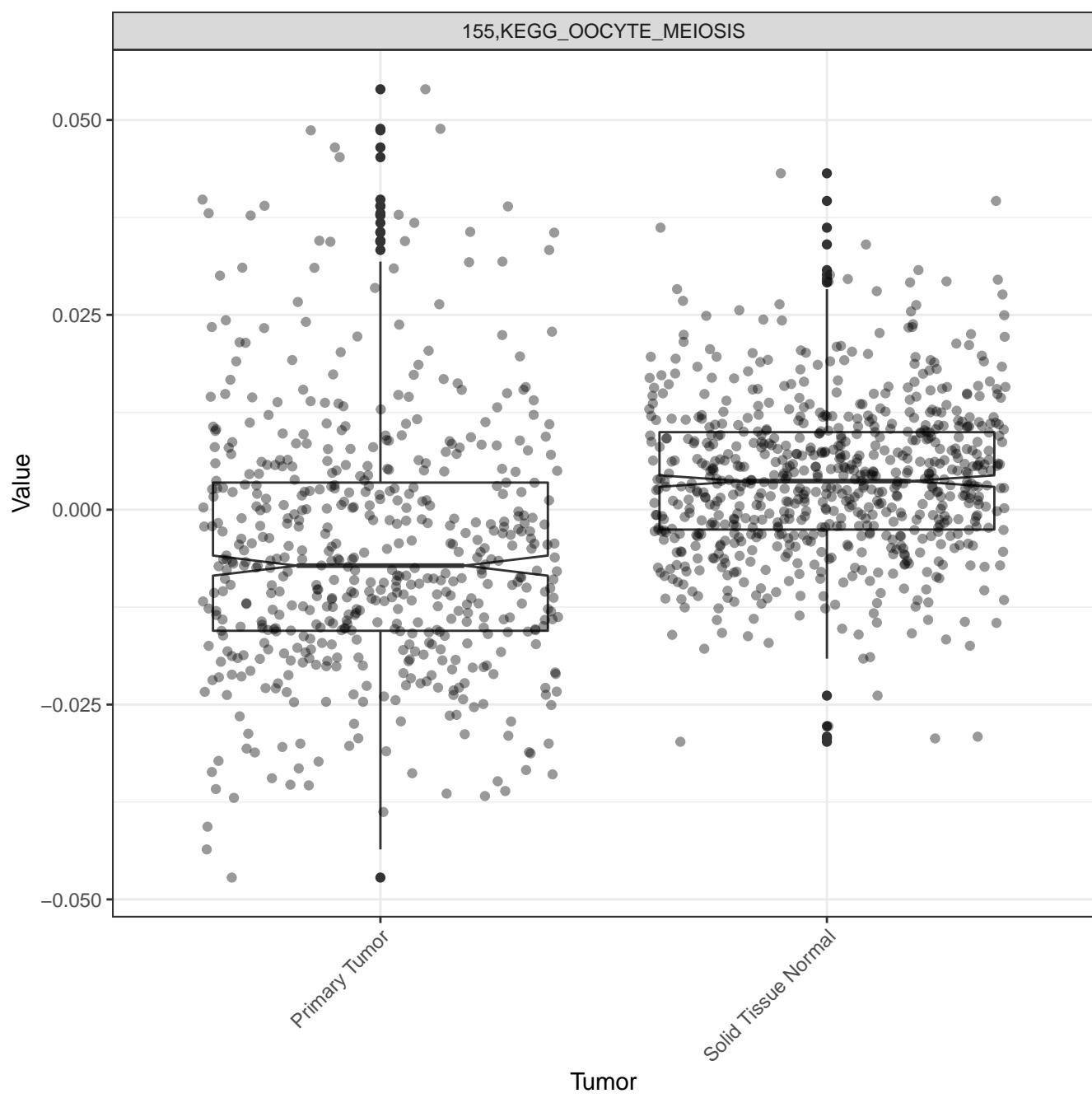
Tumor



154,KEGG_CYSTEINE_AND_METHIONINE_METABOLISM



155,KEGG_OOCYTE_MEIOSIS



156,DMAP_MONO2

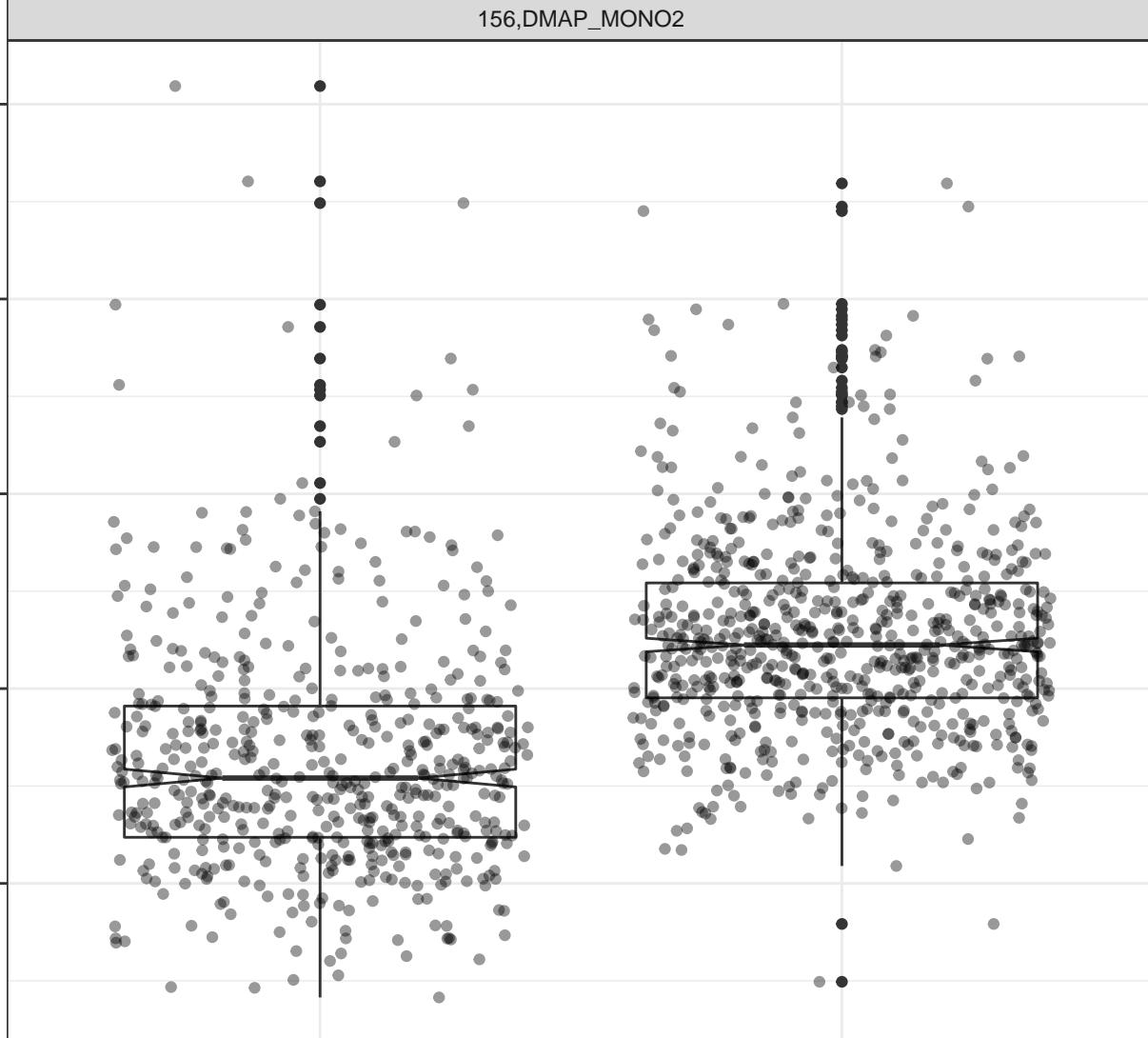
Value

0.15
0.10
0.05
0.00
-0.05

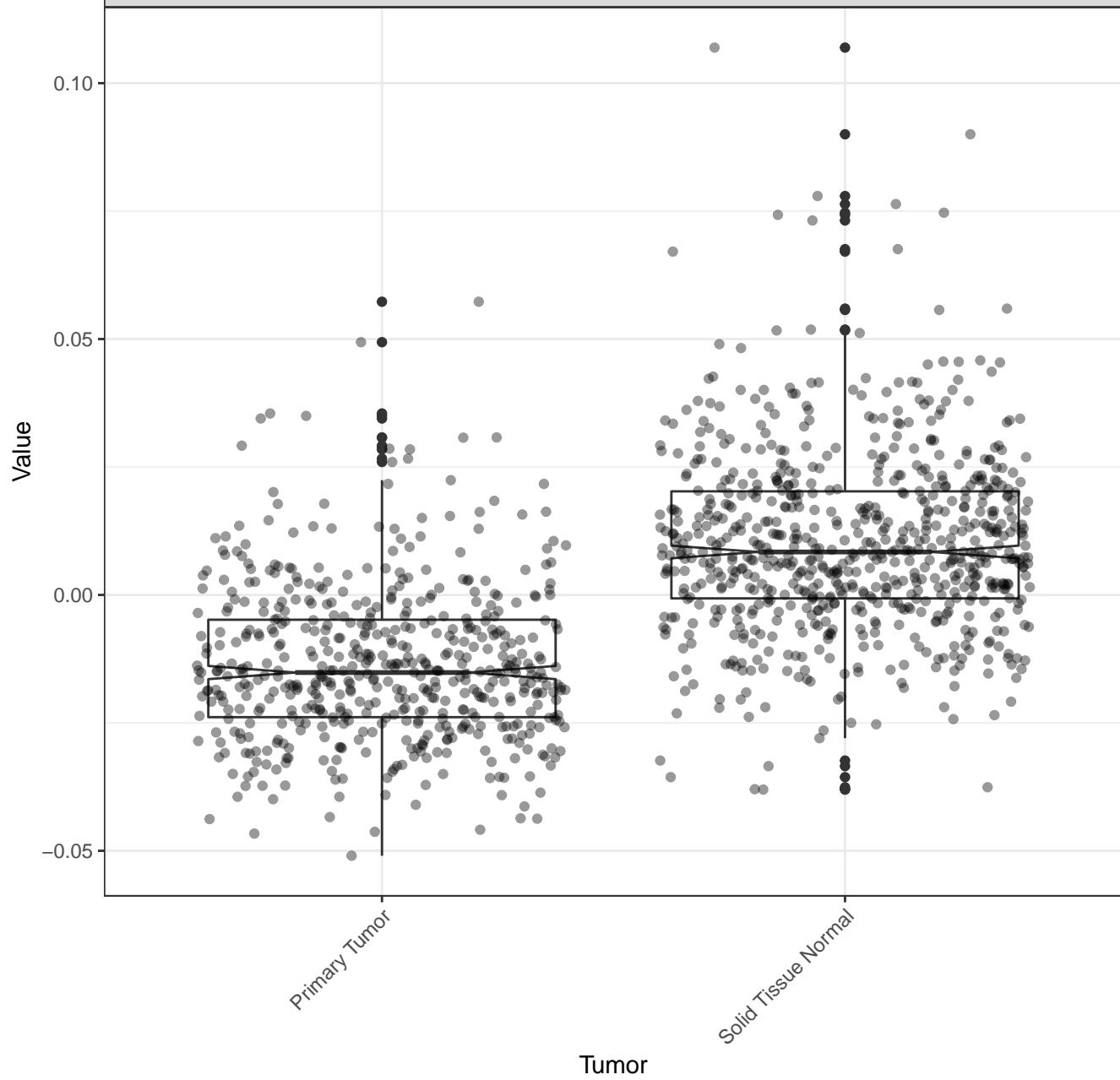
Primary Tumor

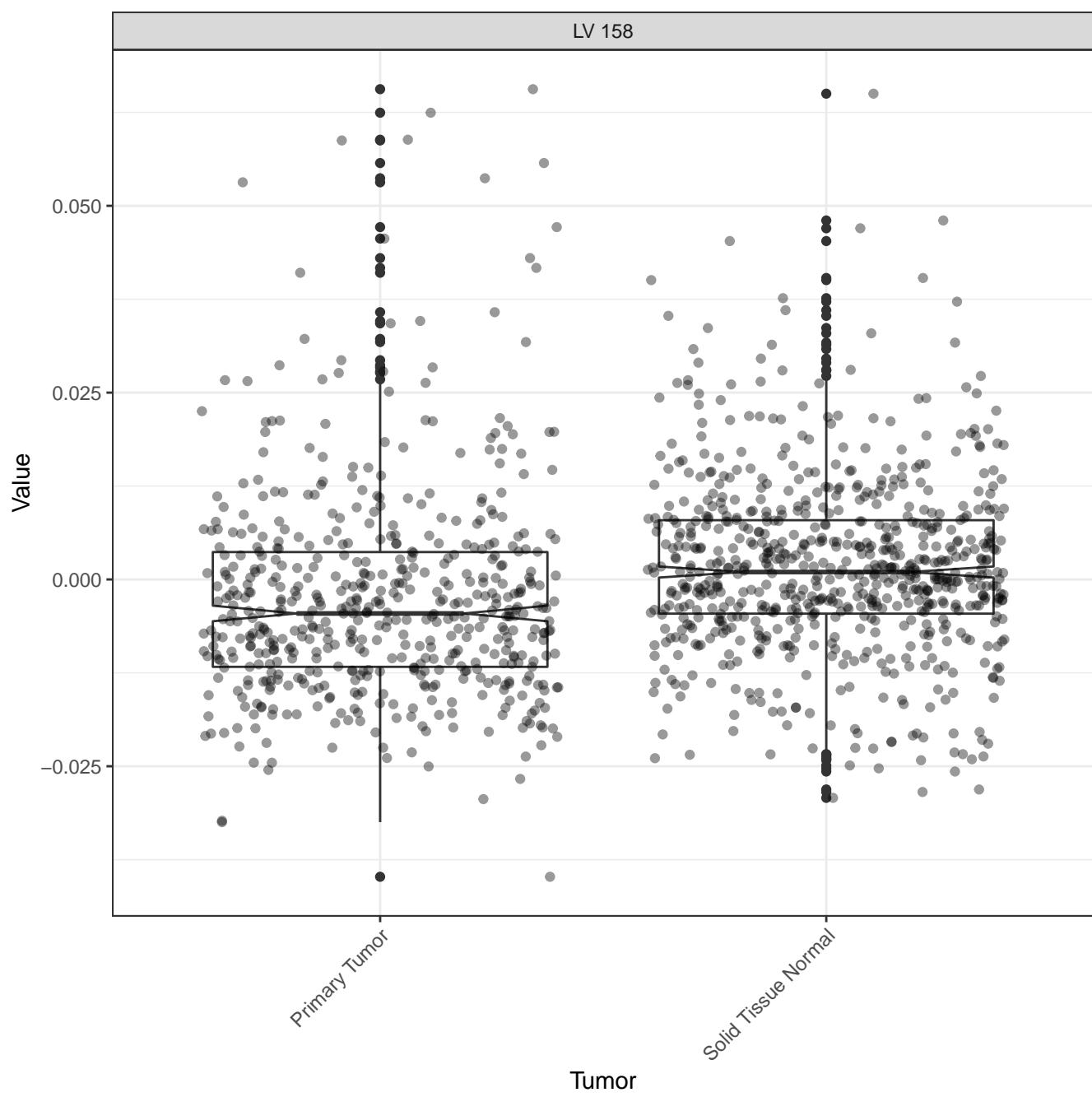
Solid Tissue Normal

Tumor

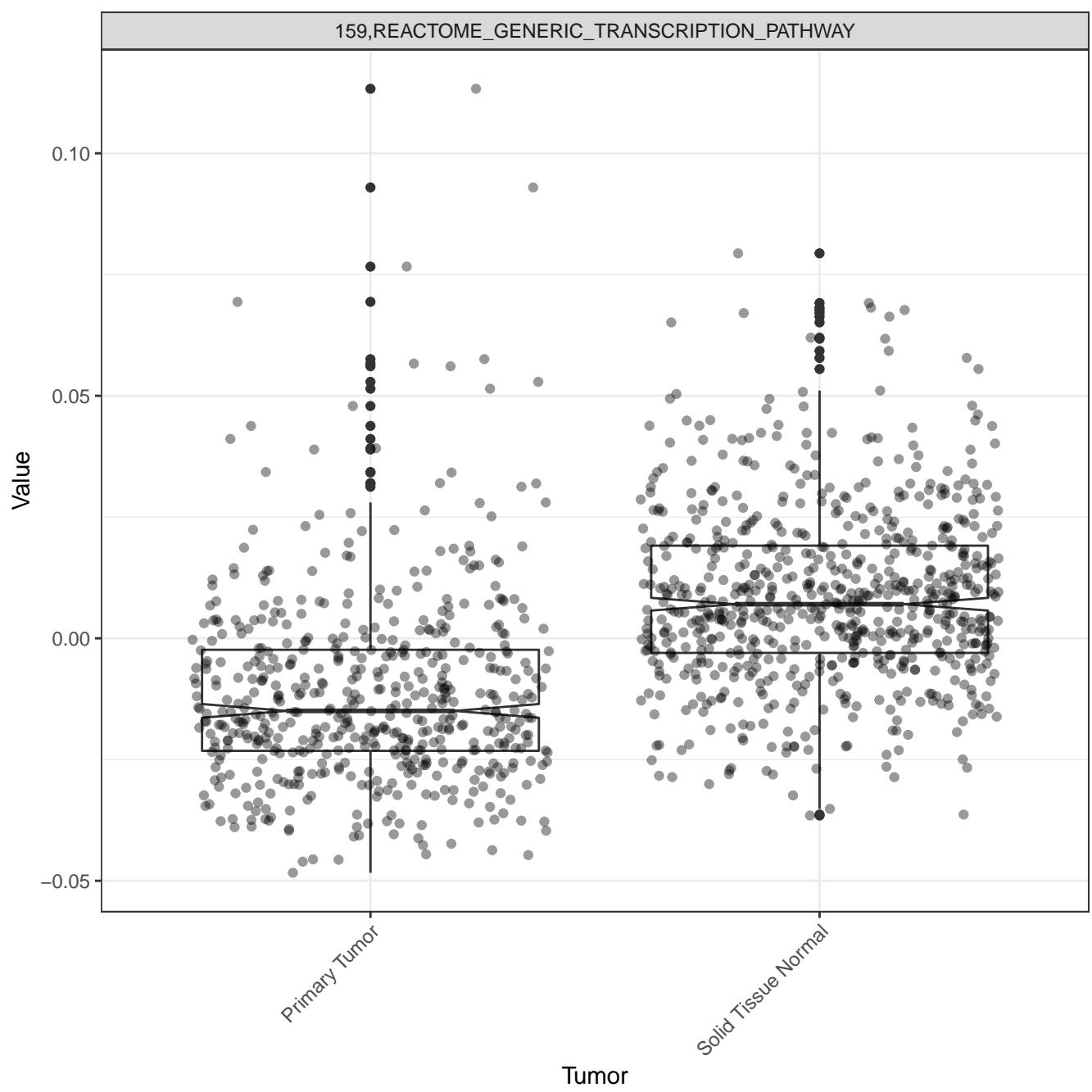


157,PID_HES_HEYPATHWAY

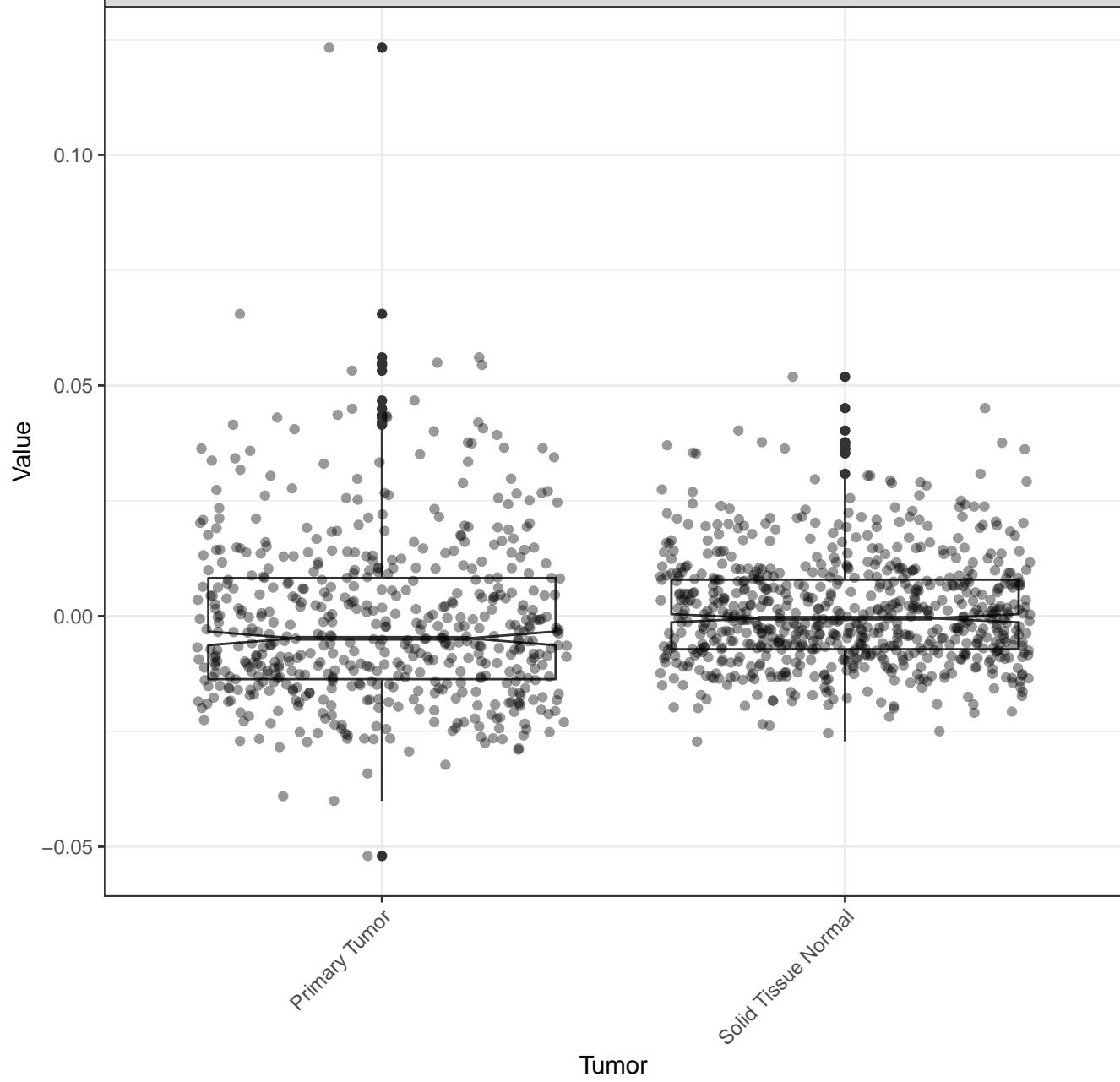




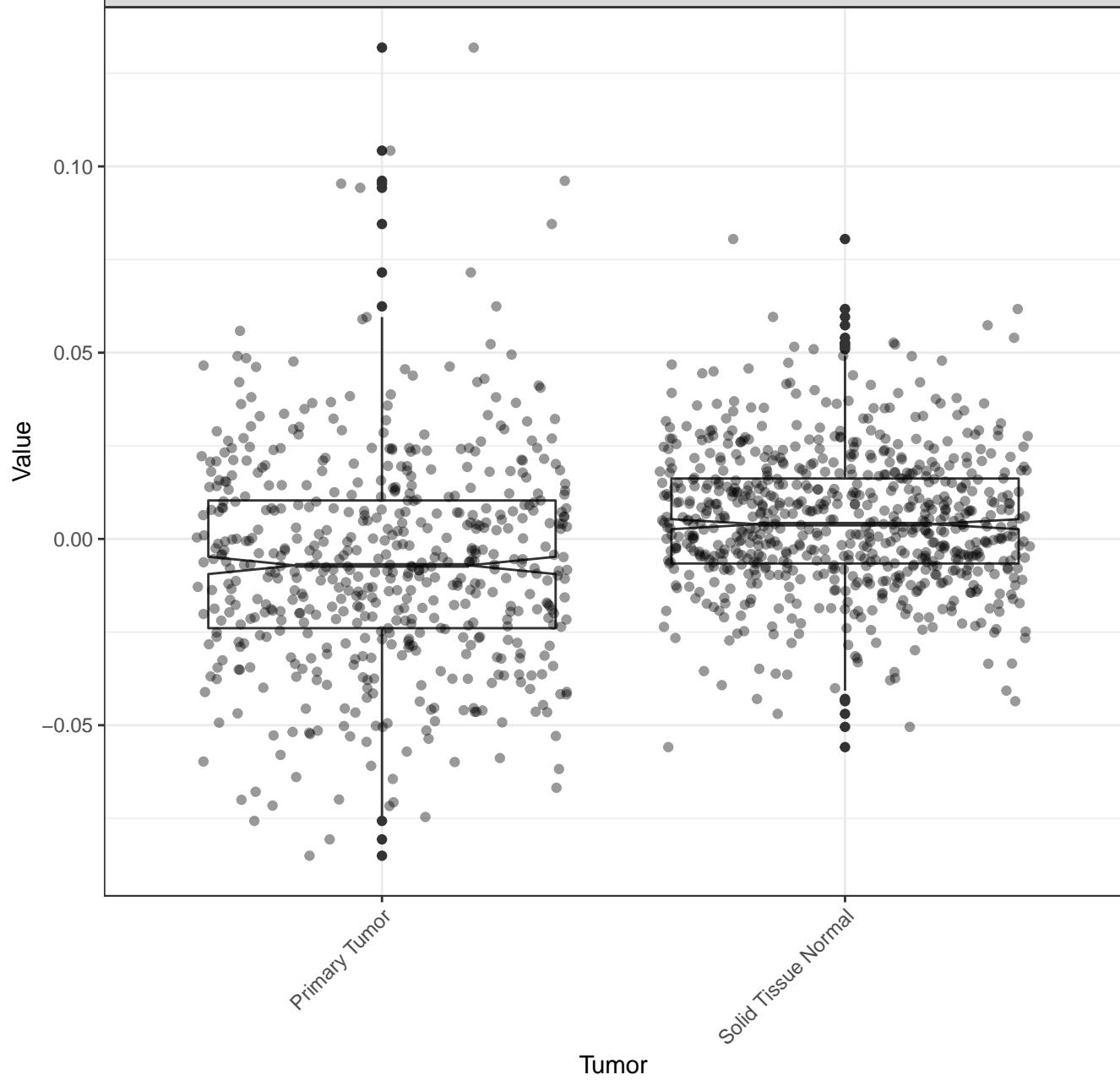
159,REACTOME_GENERIC_TRANSSCRIPTION_PATHWAY



160,REACTOME_GPCR_DOWNSTREAM_SIGNALING



161,PID_ARF6_TRAFFICKINGPATHWAY



162,KEGG_GLYCOLYSIS_GLUCONEOGENESIS

Value

0.2

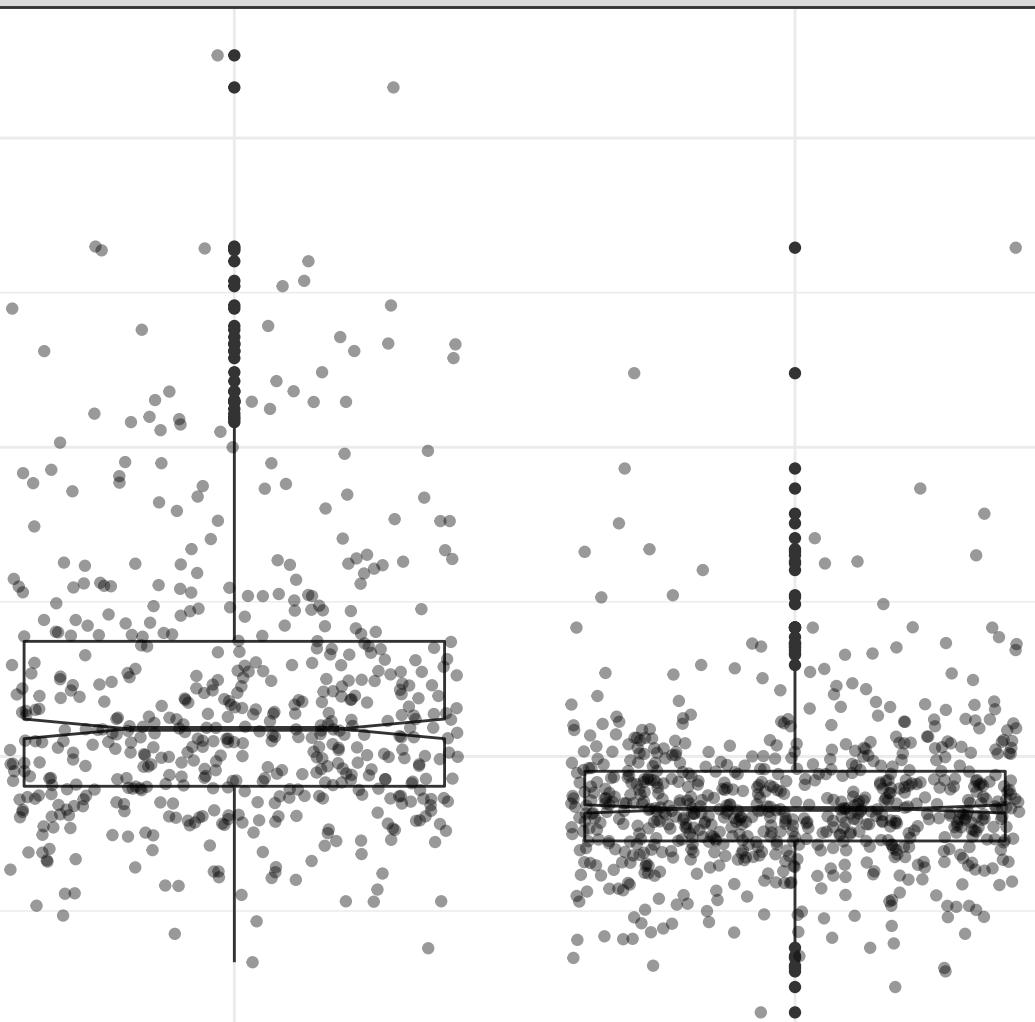
0.1

0.0

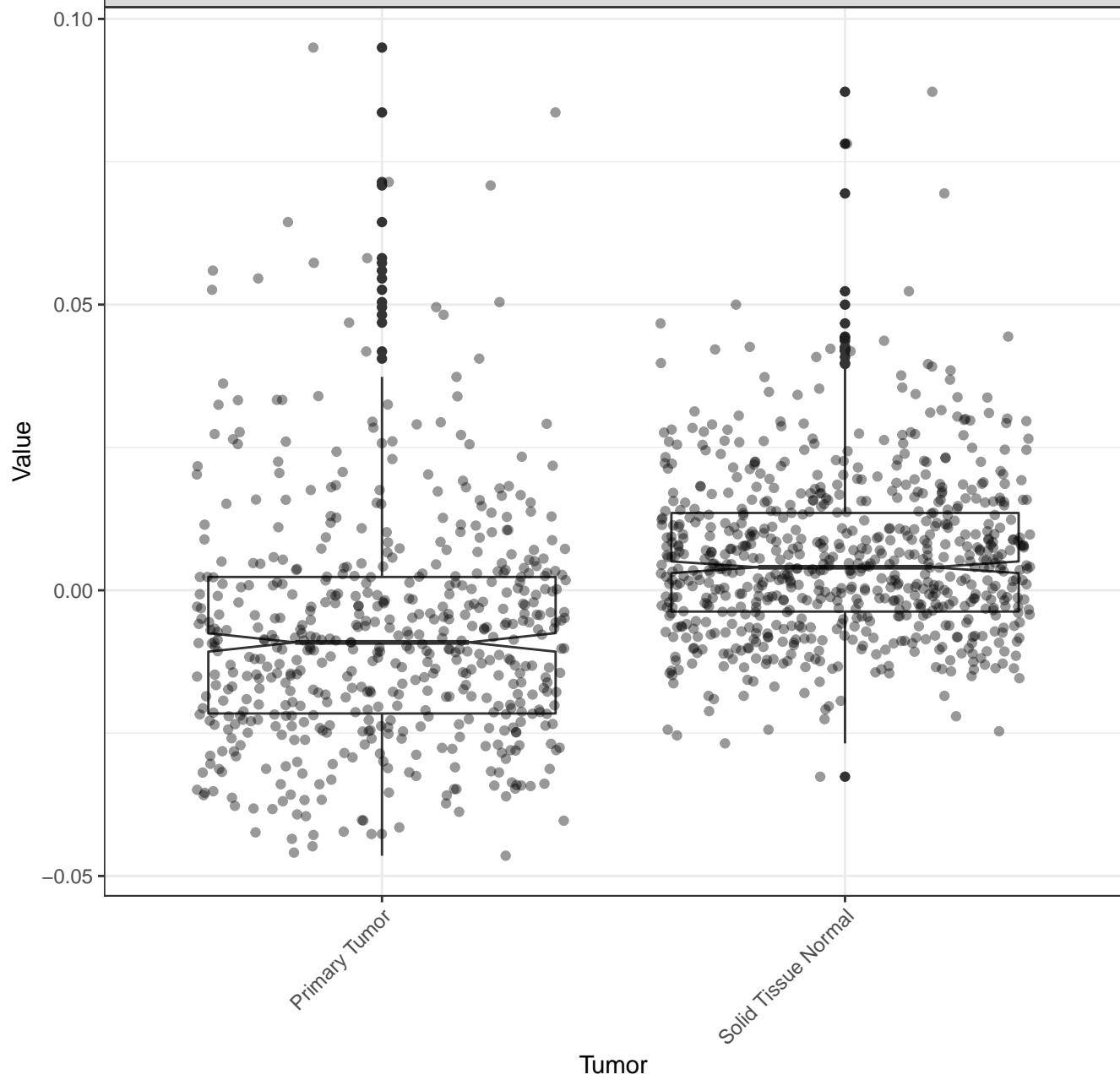
Primary Tumor

Solid Tissue Normal

Tumor



163,KEGG_COMPLEMENT_AND_COAGULATION_CASCADES



164,KEGG_LEISHMANIA_INFECTON

Value

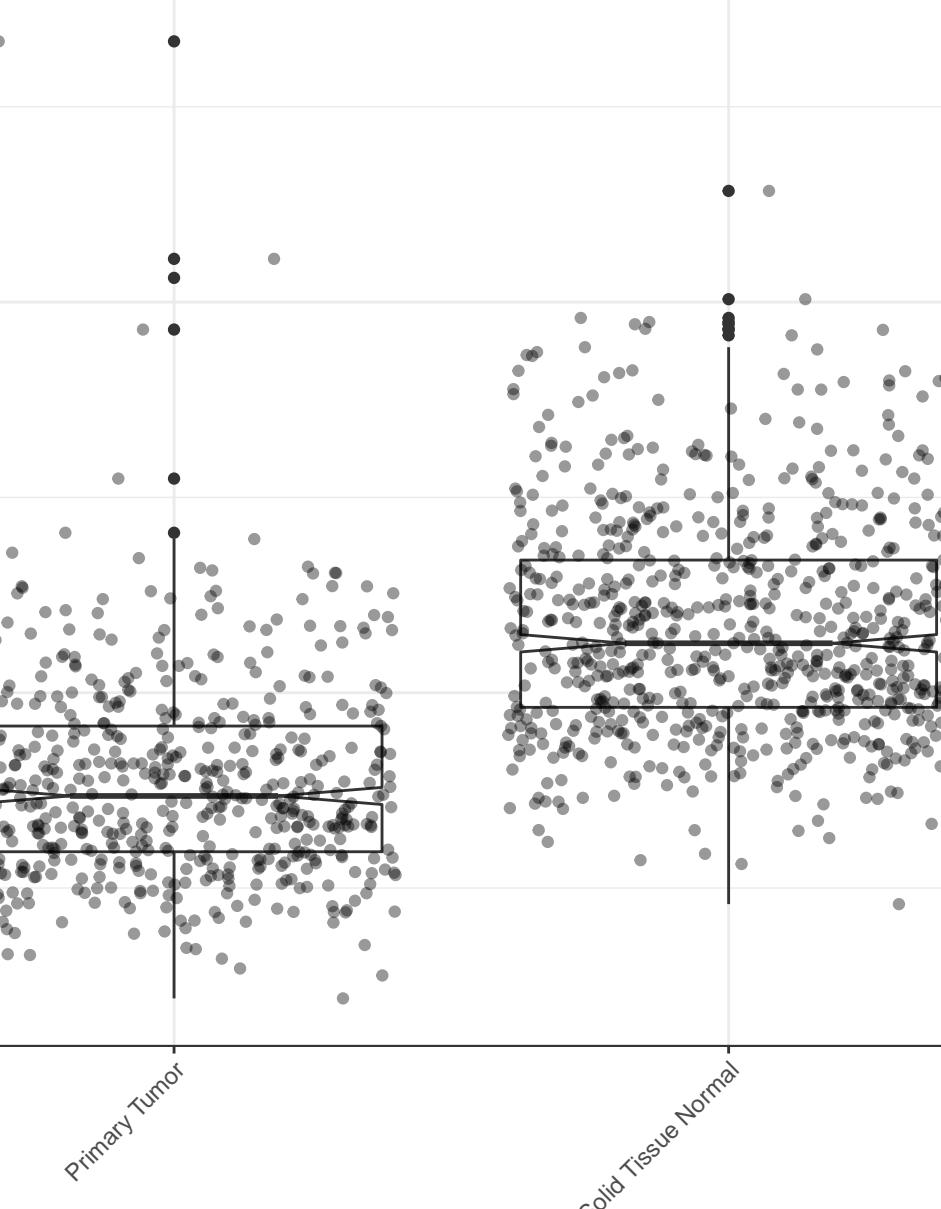
0.05

0.00

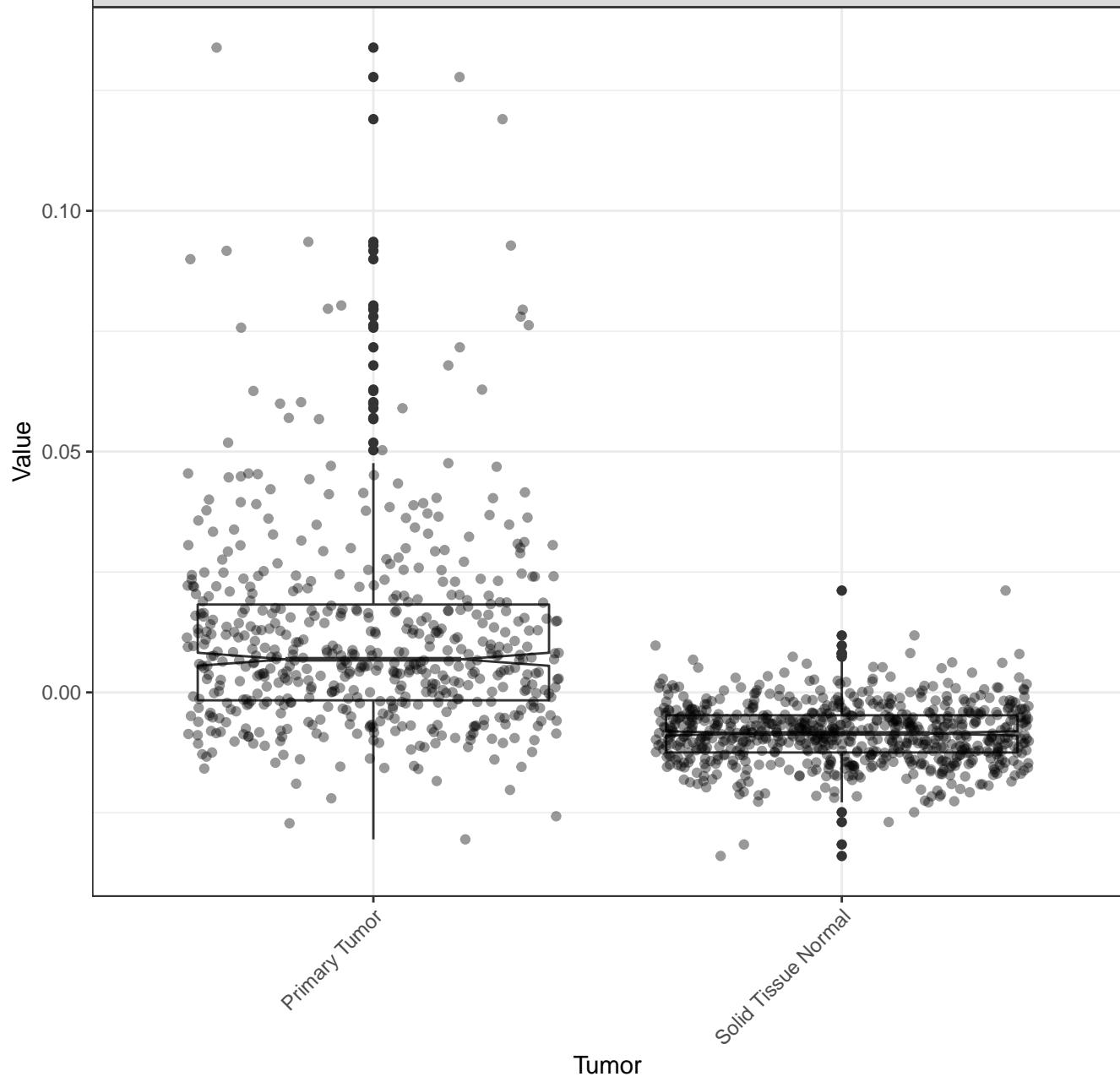
Primary Tumor

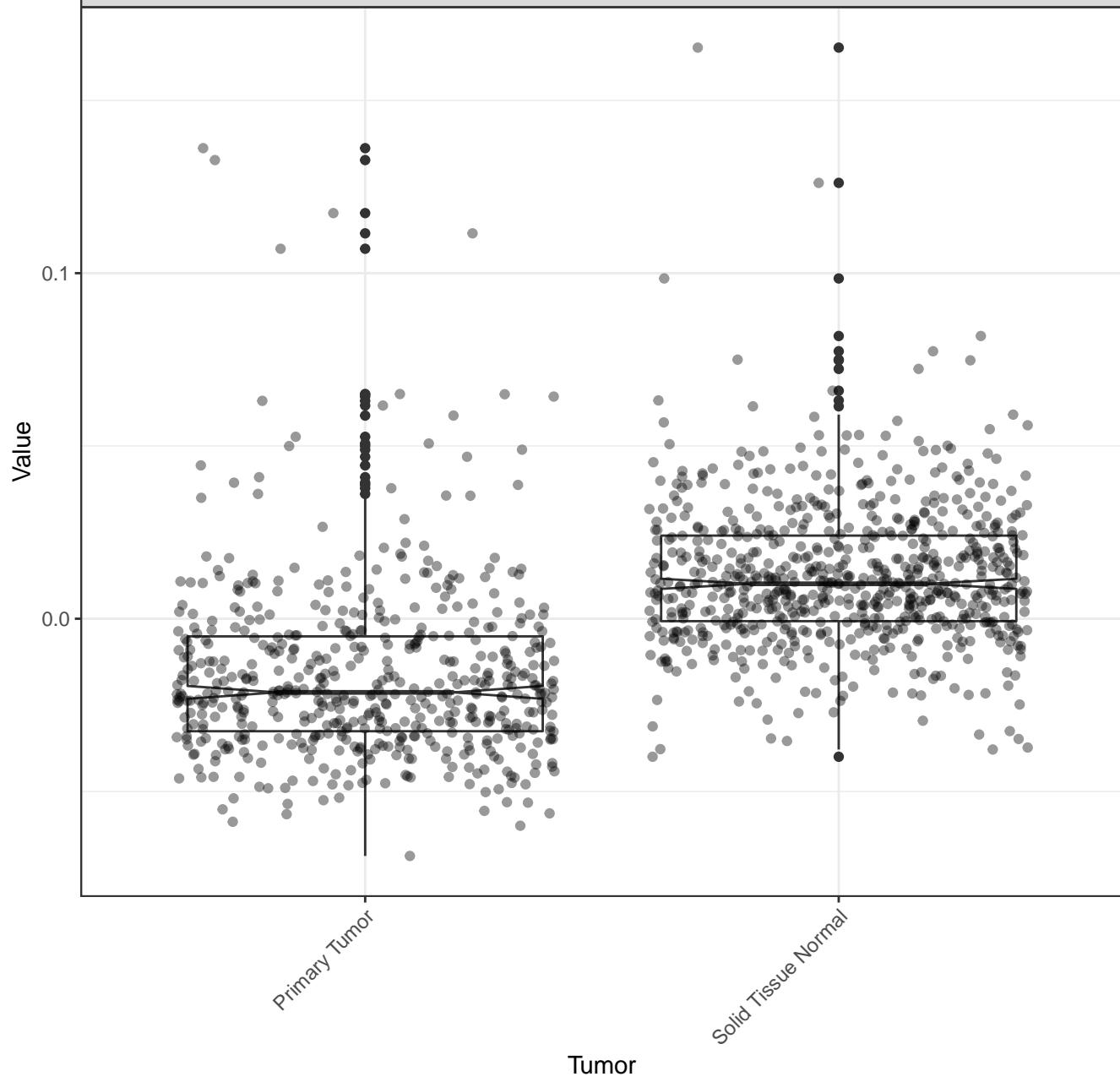
Tumor

Solid Tissue Normal

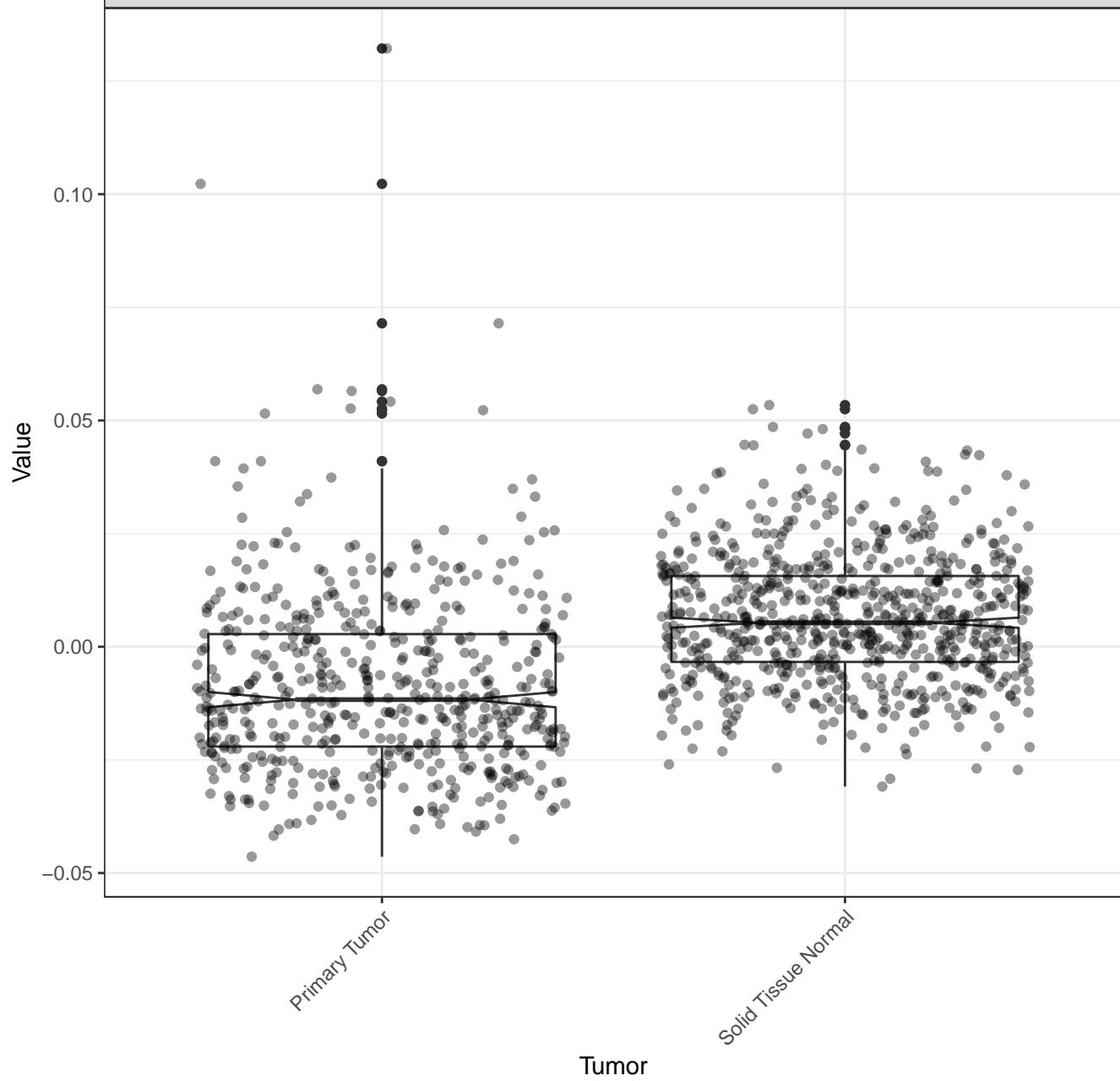


165,DMAP_MECA1





167,PID_LKB1_PATHWAY



168,REACTOME_DOWNSTREAM_SIGNALING_EVENTS_OF_B_CELL_RECECTOR_BCR

Value

0.10

0.05

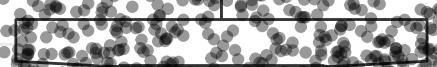
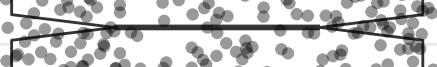
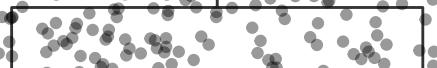
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



169,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

Value

0.1

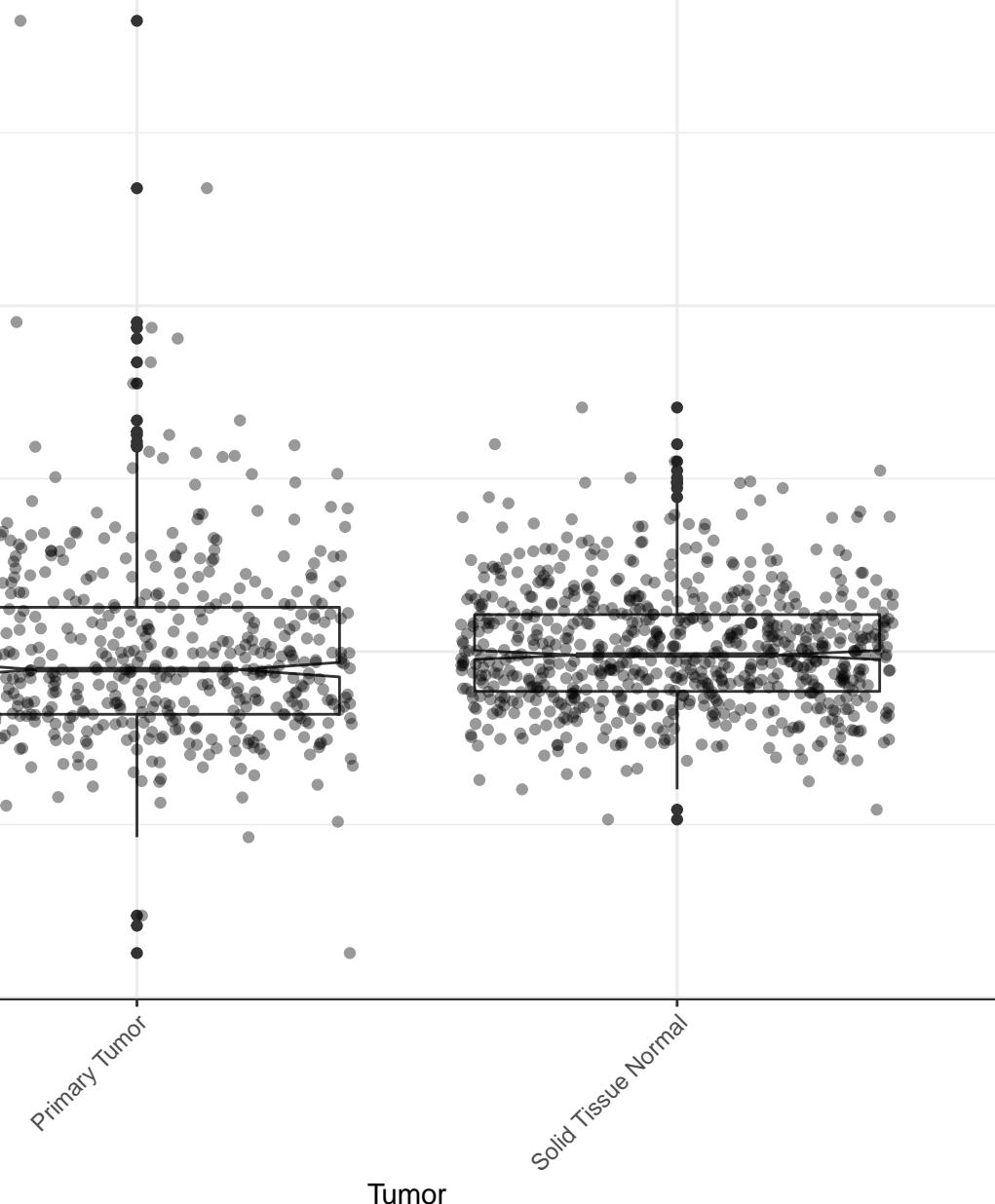
0.0

-0.1

Primary Tumor

Solid Tissue Normal

Tumor



170,KEGG_TOLL_LIKE_RECEPTOR_SIGNALING_PATHWAY

Value

0.10

0.05

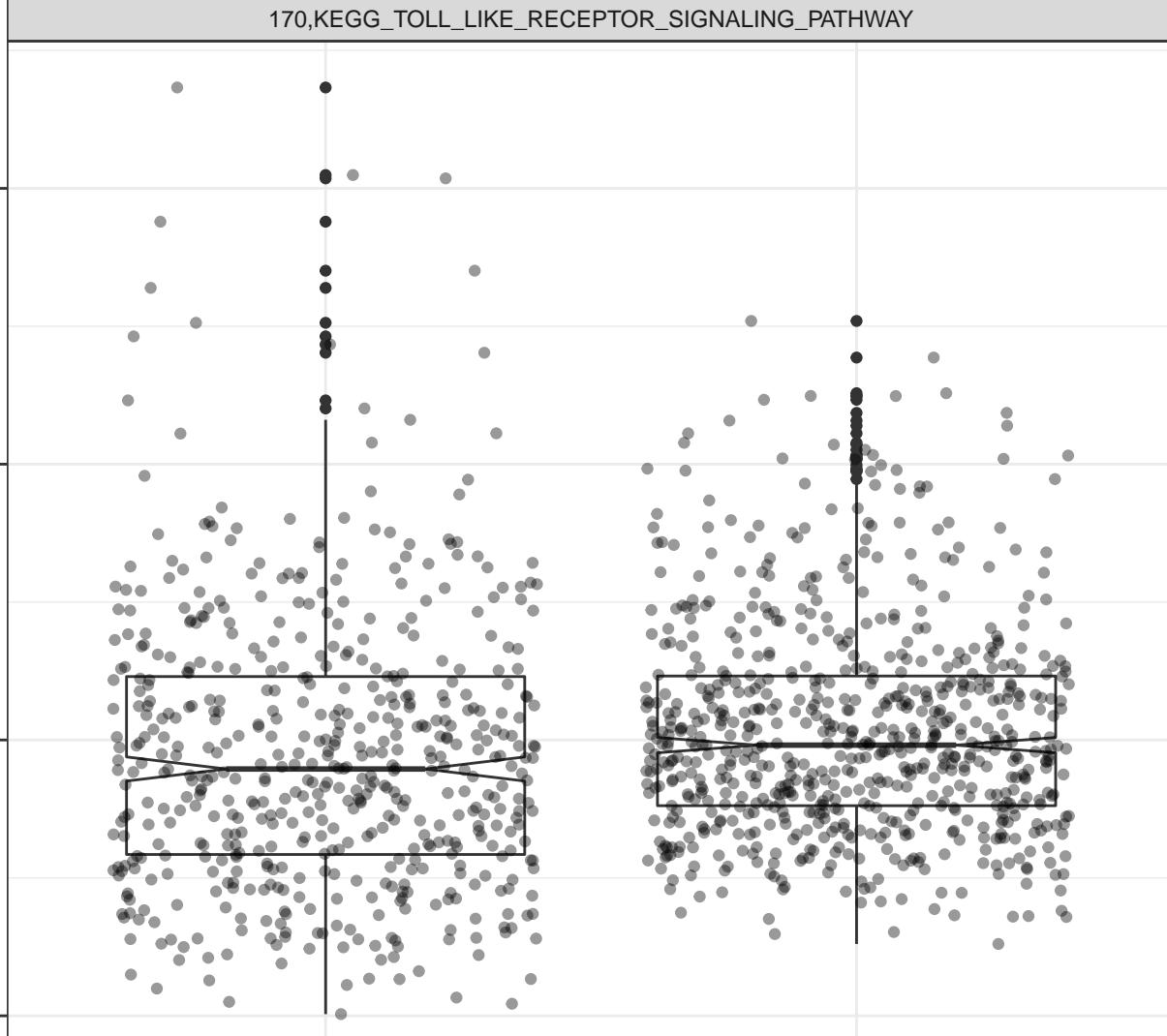
0.00

-0.05

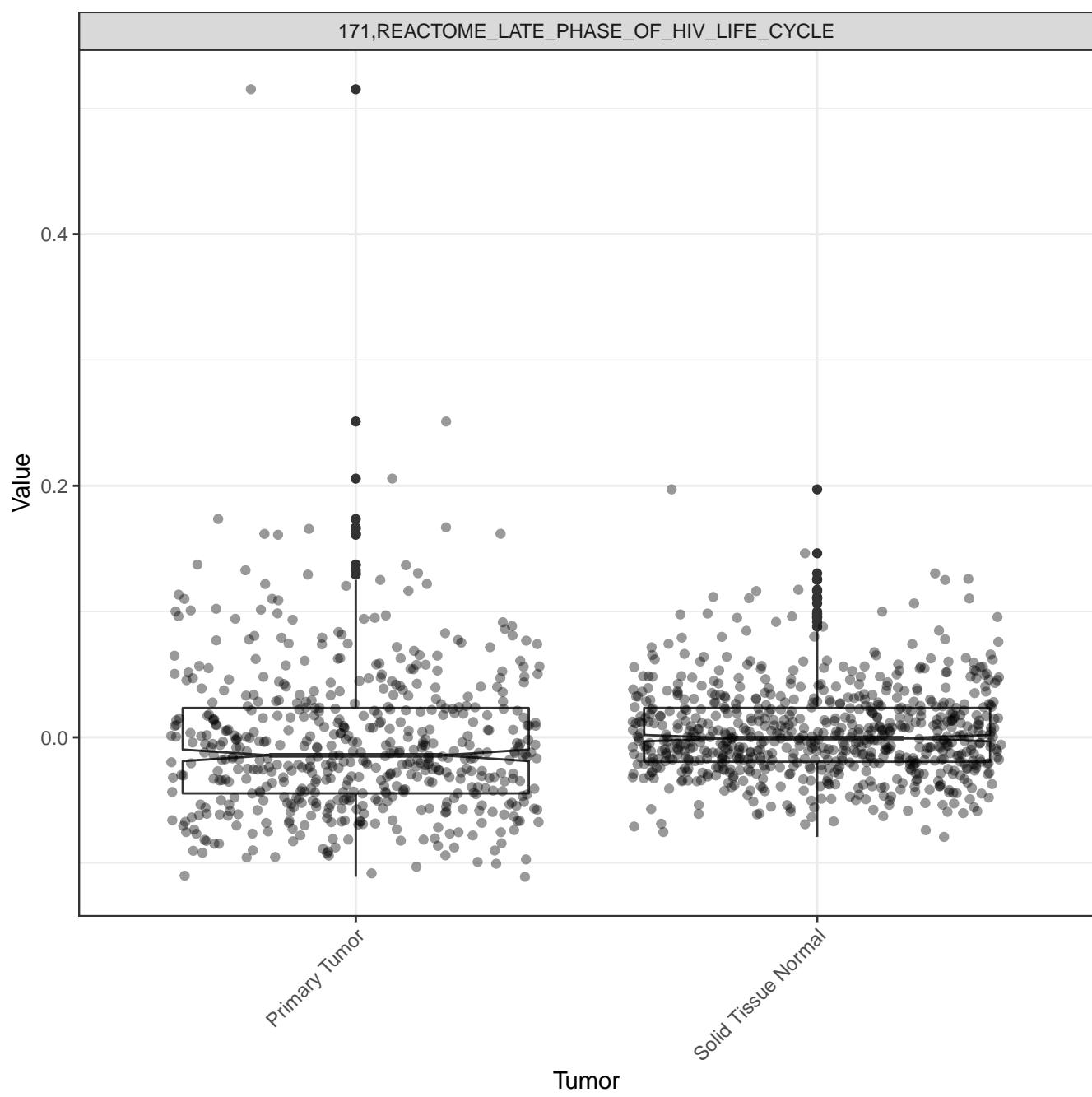
Primary Tumor

Solid Tissue Normal

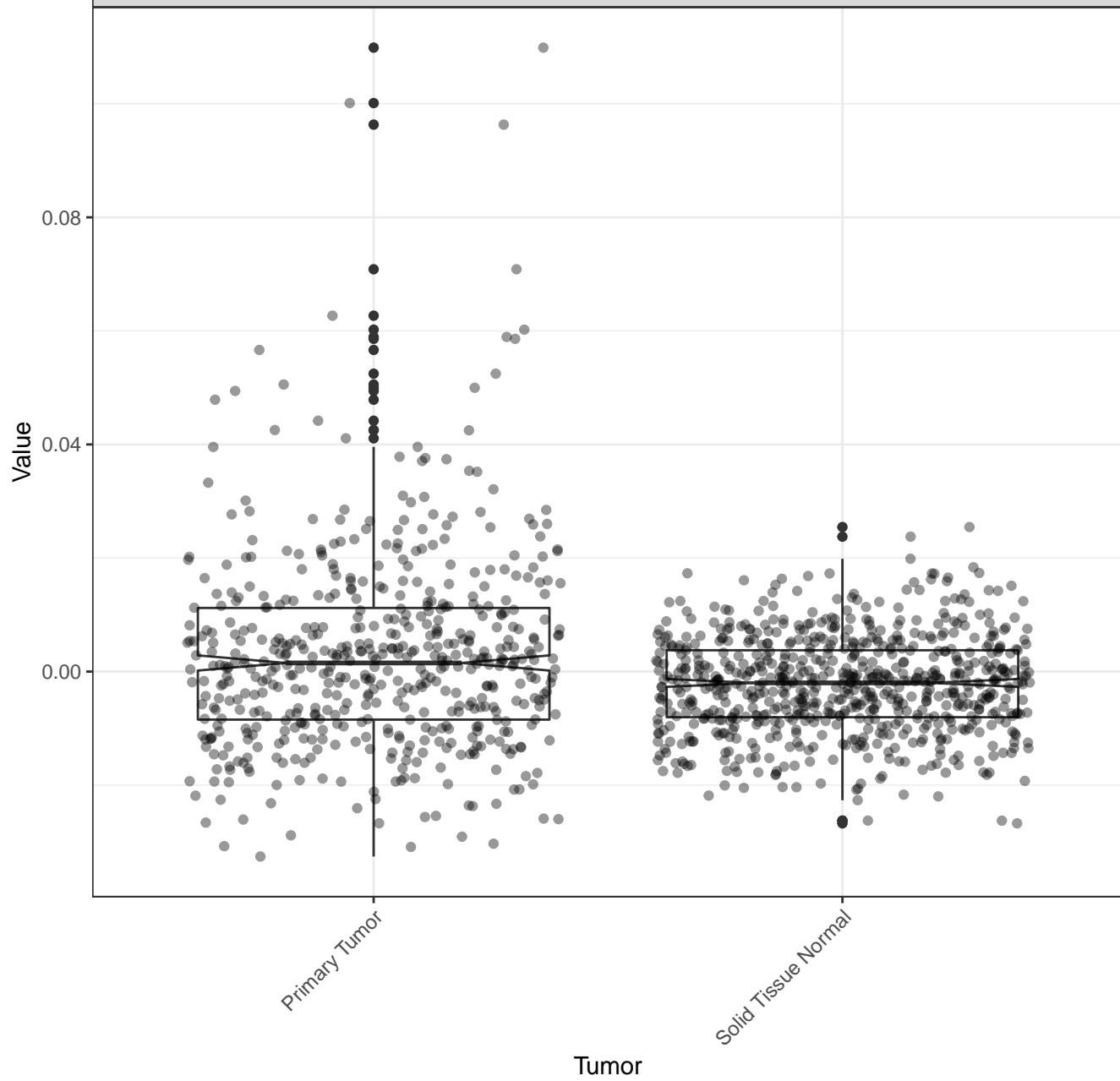
Tumor



171,REACTOME_LATE_PHASE_OF_HIV_LIFE_CYCLE



172,REACTOME_DEADENYLATION_DEPENDENT_MRNA_DECAY



Value

0.08

0.04

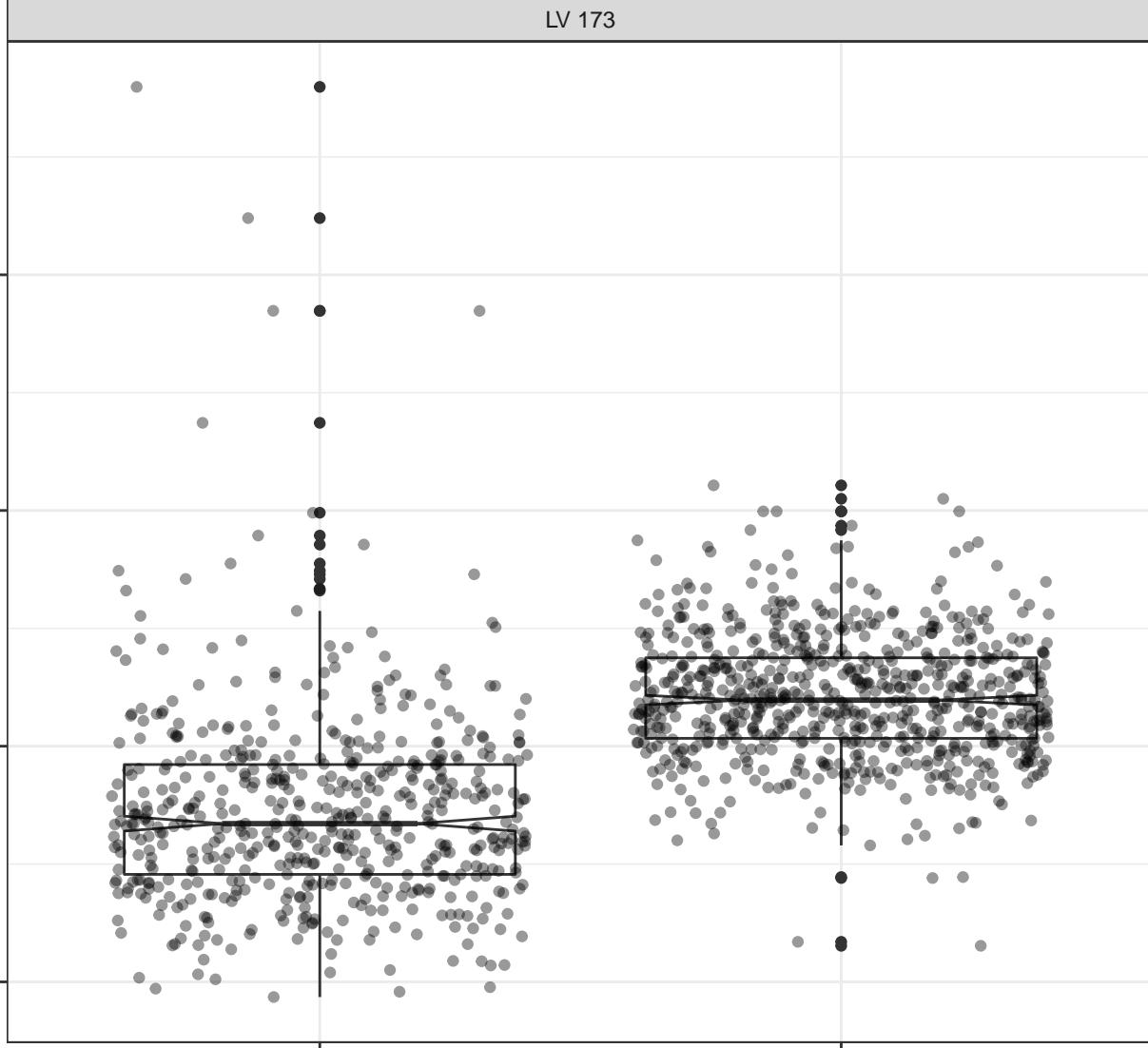
0.00

-0.04

Primary Tumor

Solid Tissue Normal

Tumor



174,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

Value

0.10

0.05

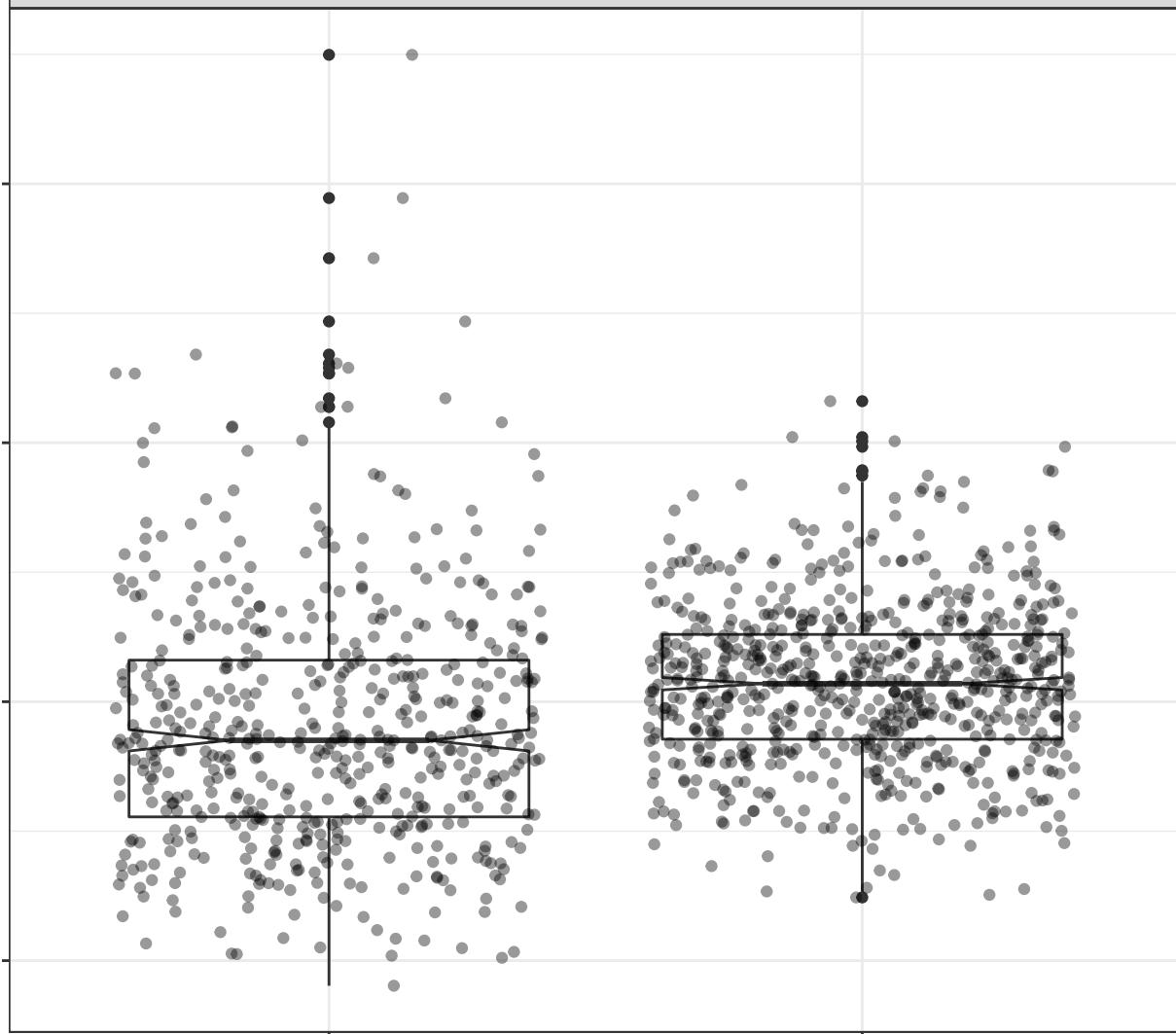
0.00

-0.05

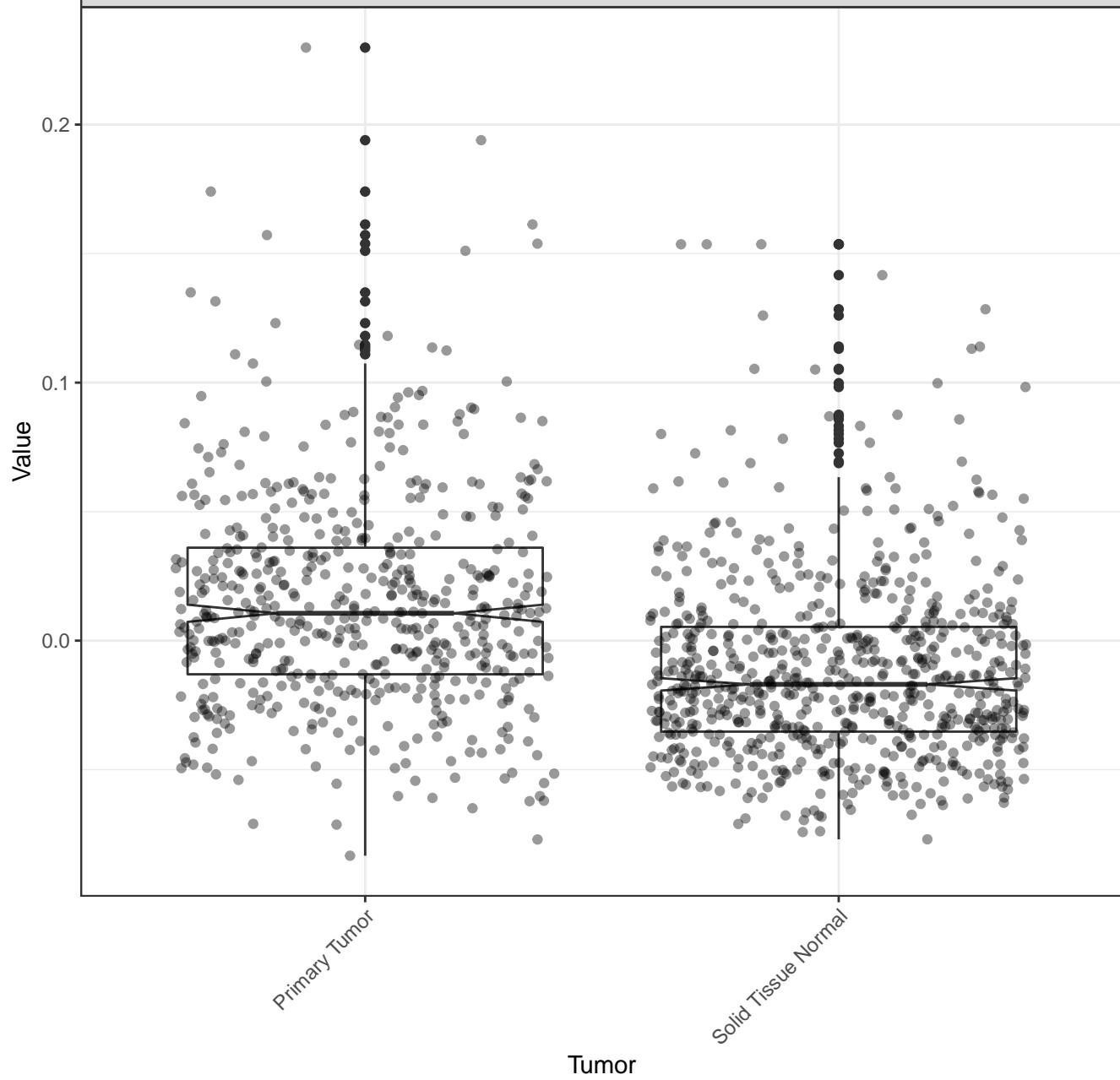
Primary Tumor

Solid Tissue Normal

Tumor



175,SVM T cells CD8



176,KEGG_UBIQUITIN_MEDiated_PROTEOLYSIS

Value

0.2

0.1

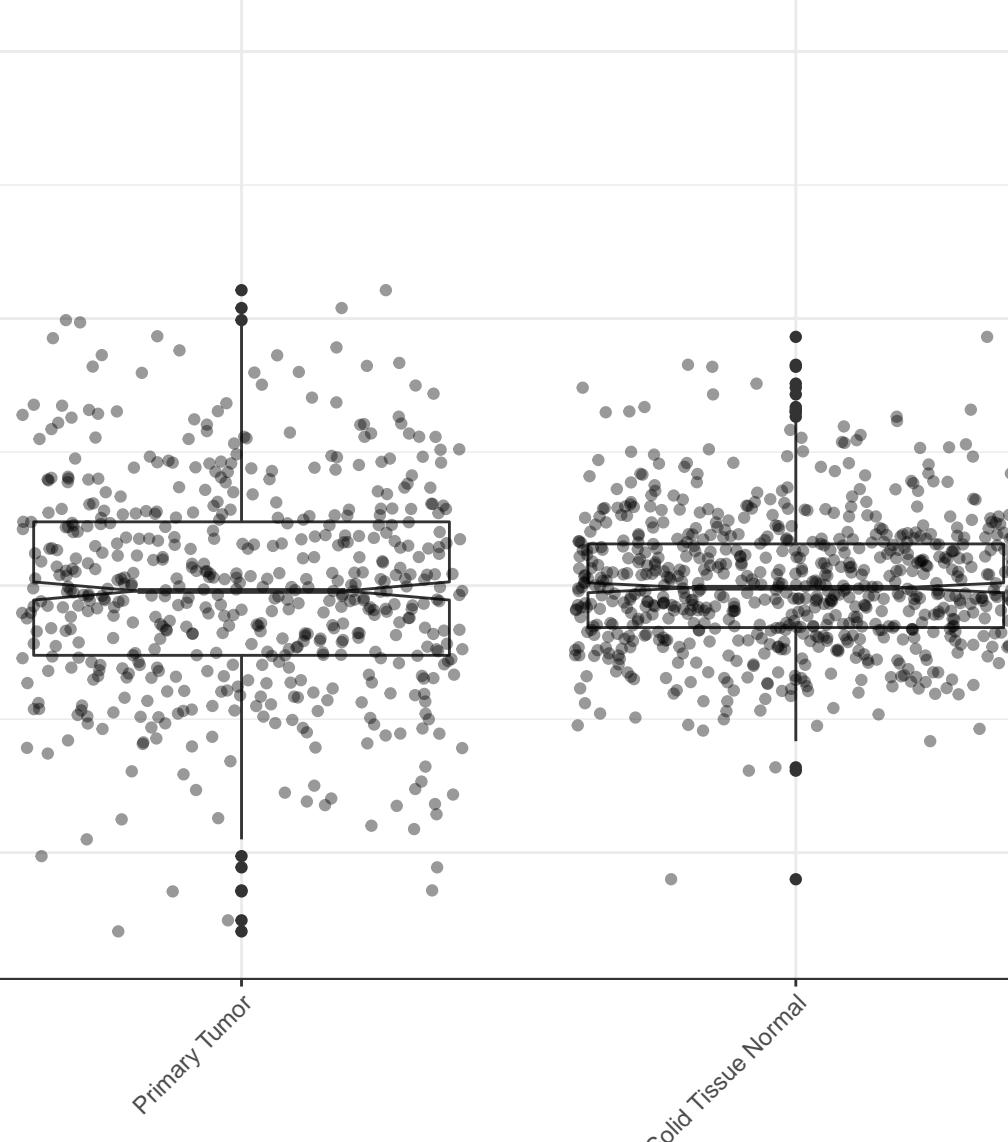
0.0

-0.1

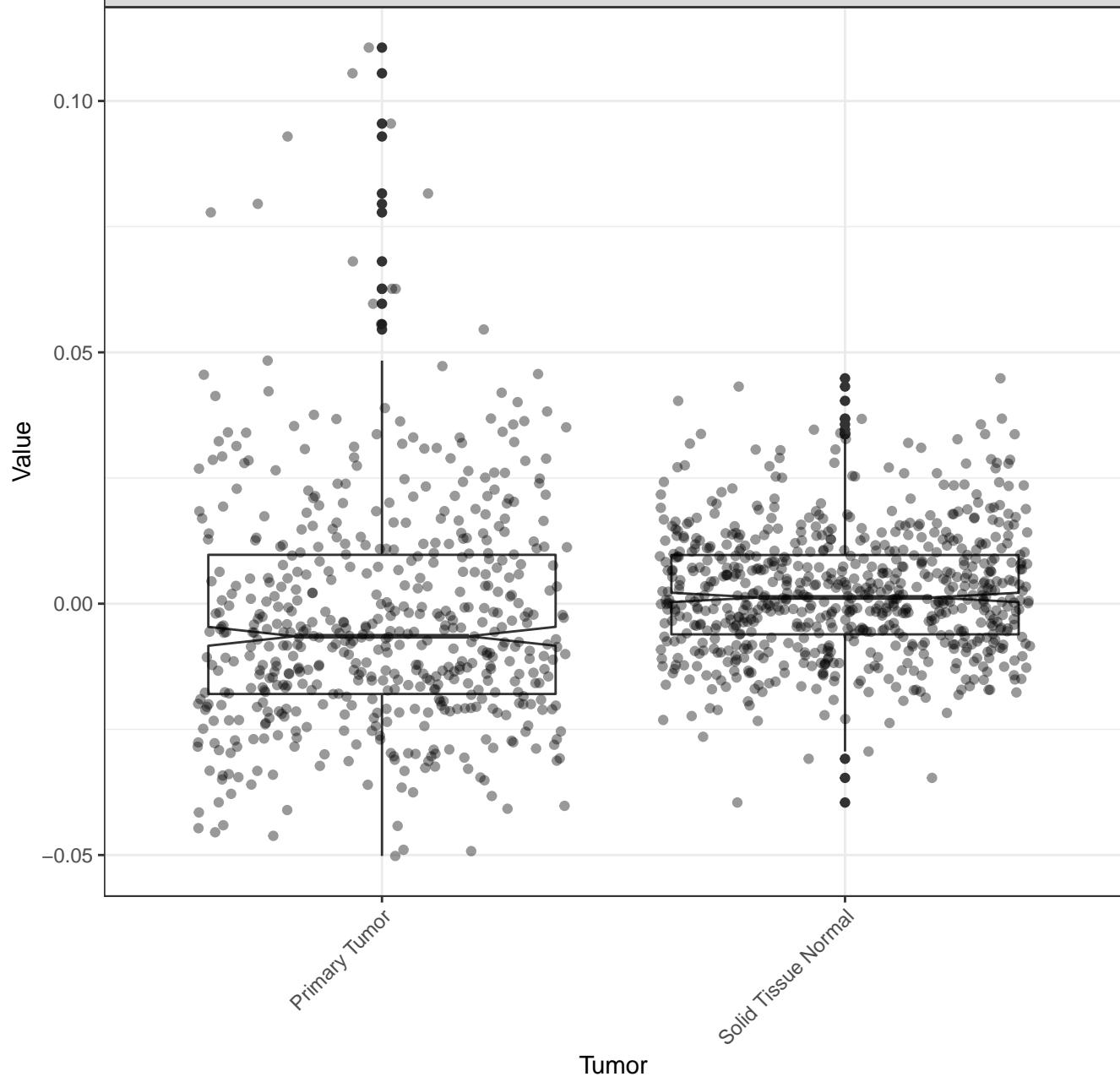
Primary Tumor

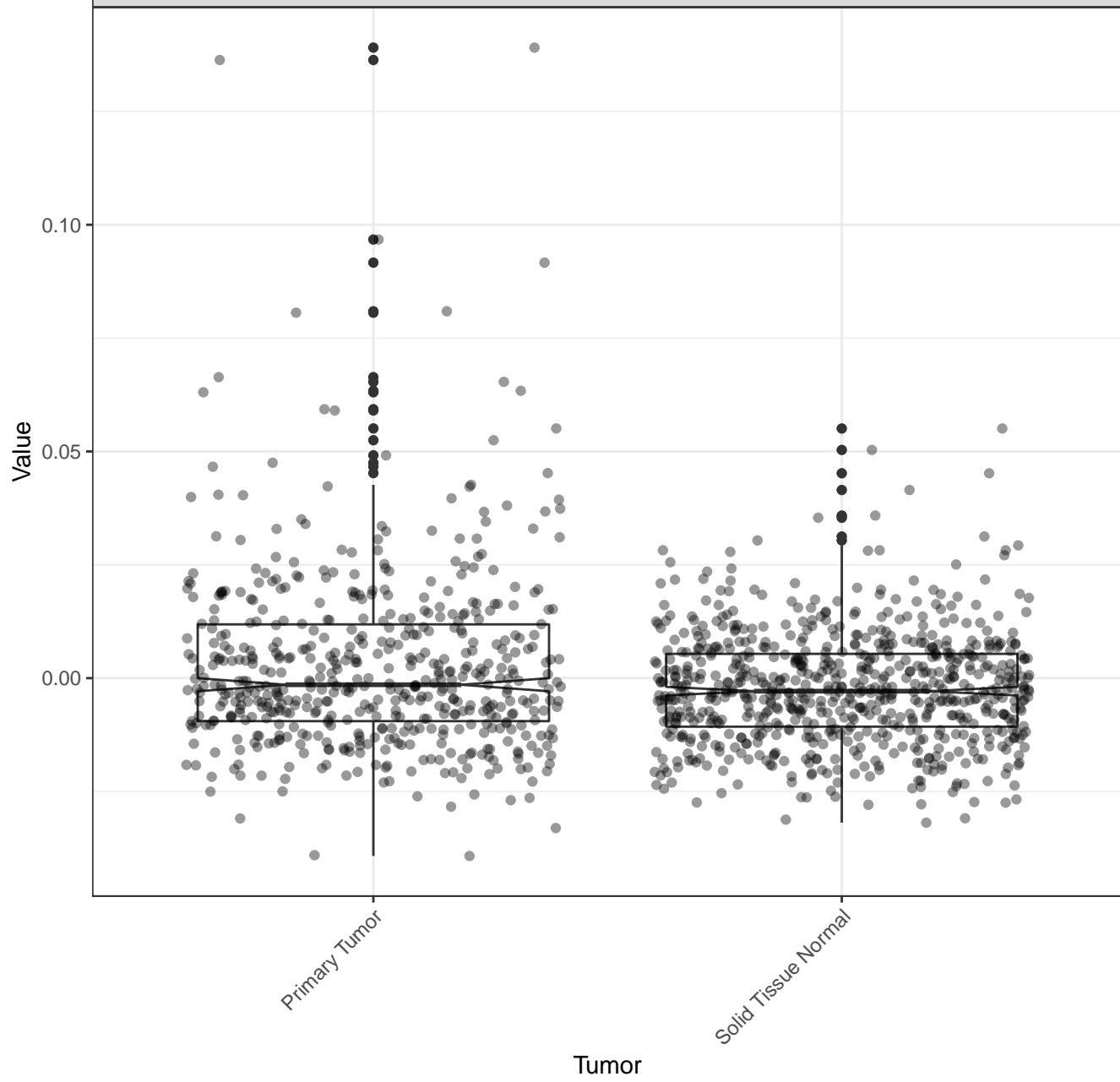
Solid Tissue Normal

Tumor

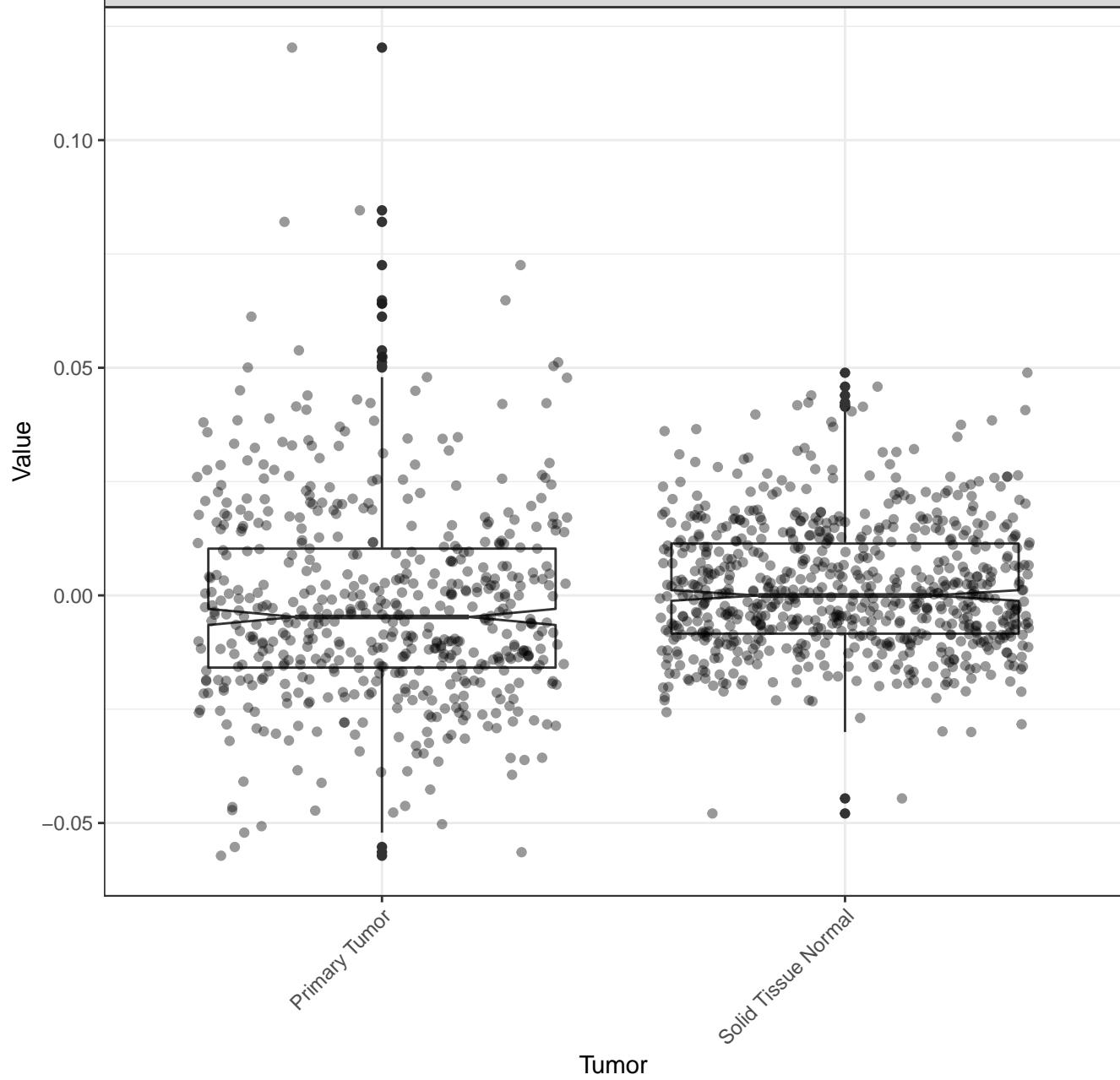


177,PID_CDC42_PATHWAY

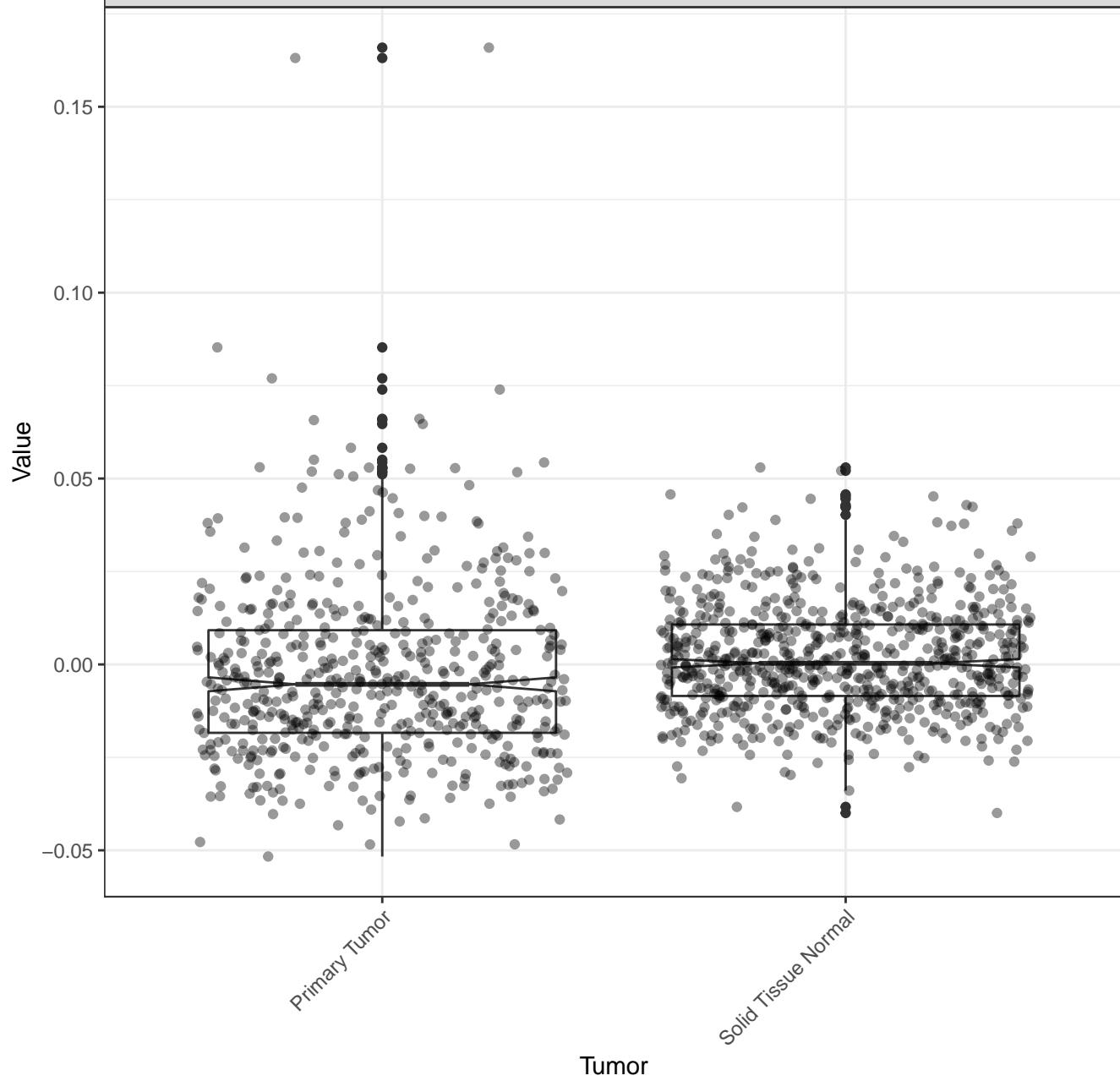


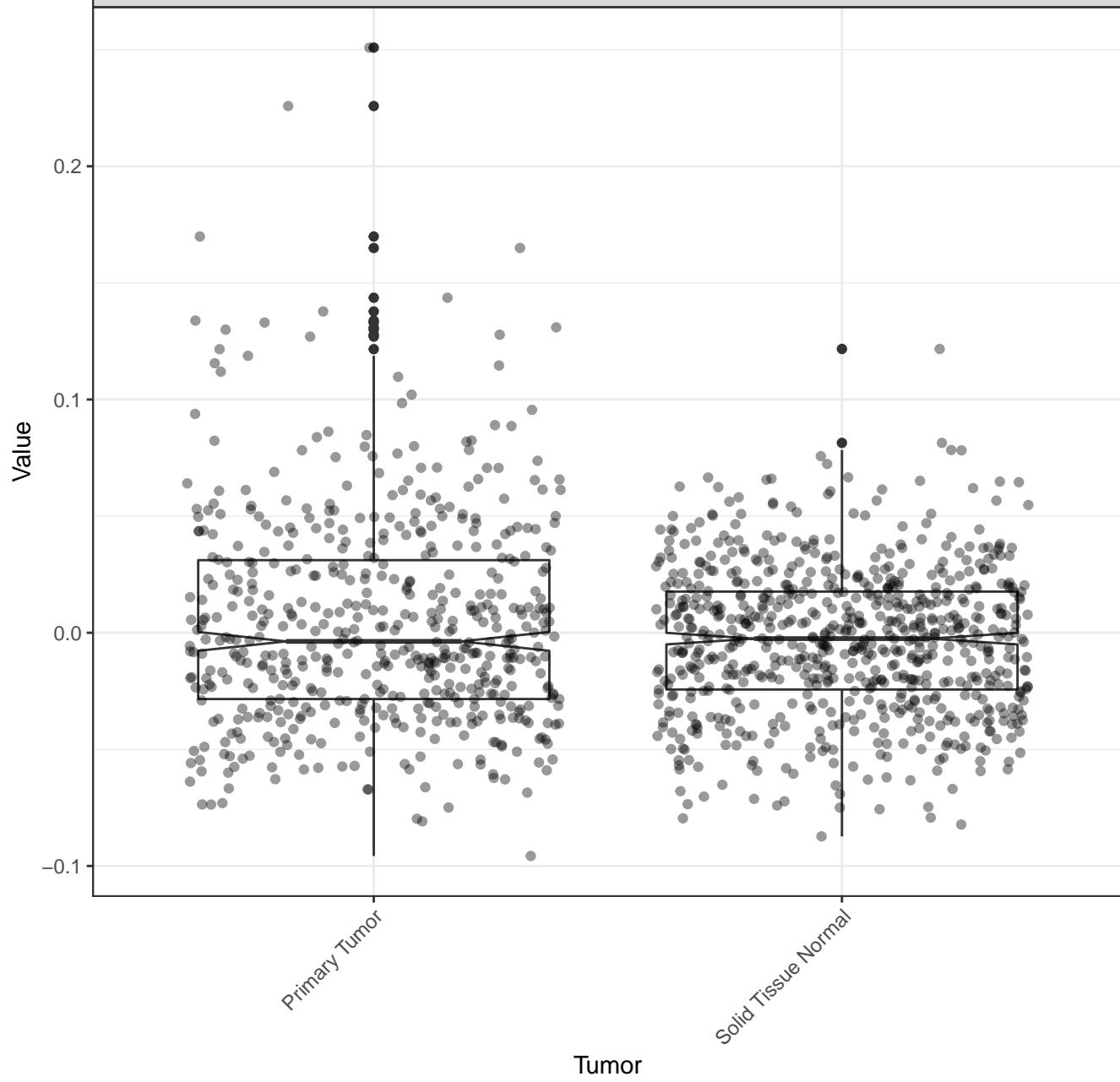


179,REACTOME_AMINO_ACID_AND_OLIGOPEPTIDE_SLC_TRANSPORTERS



180,REACTOME_PROTEIN_FOLDING





LV 182

Value

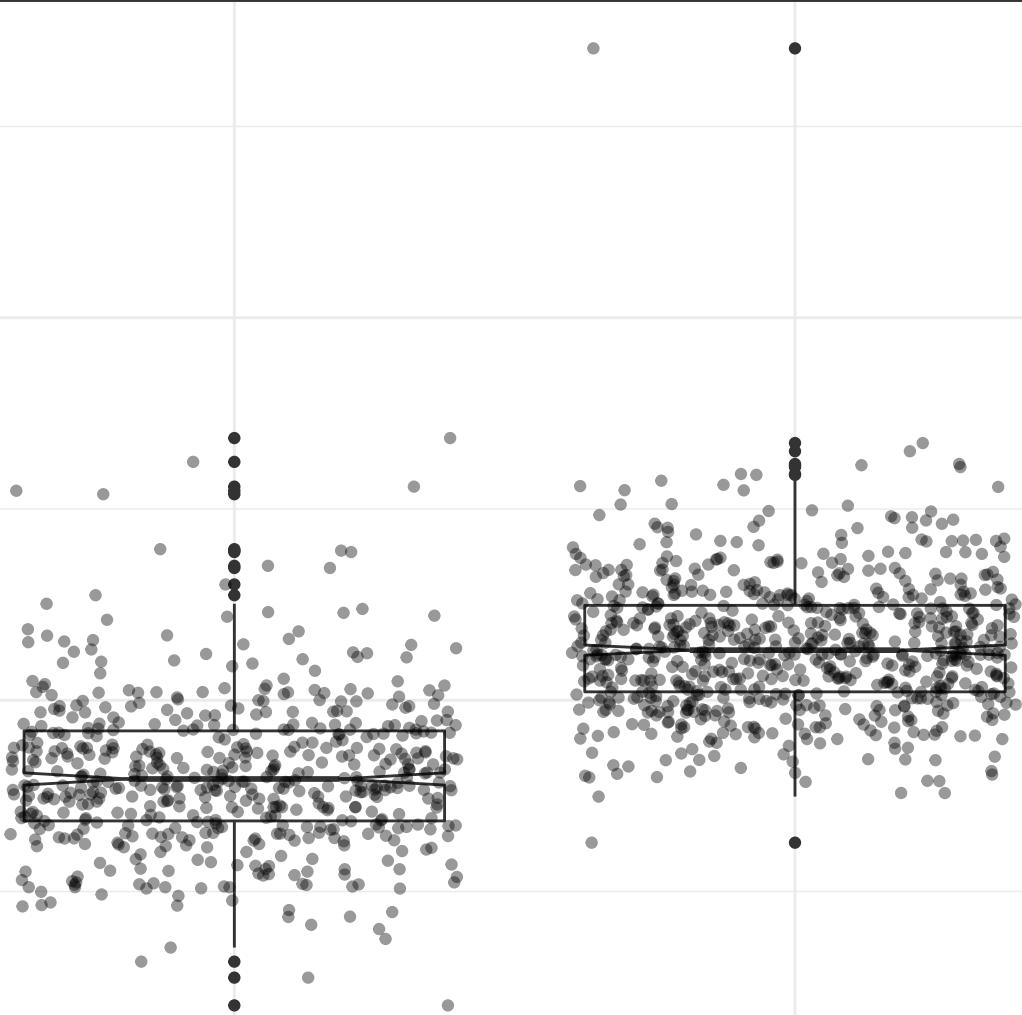
0.1

0.0

Primary Tumor

Solid Tissue Normal

Tumor



183,IRIS_DendriticCell-LPSstimulated

Value

0.10

0.05

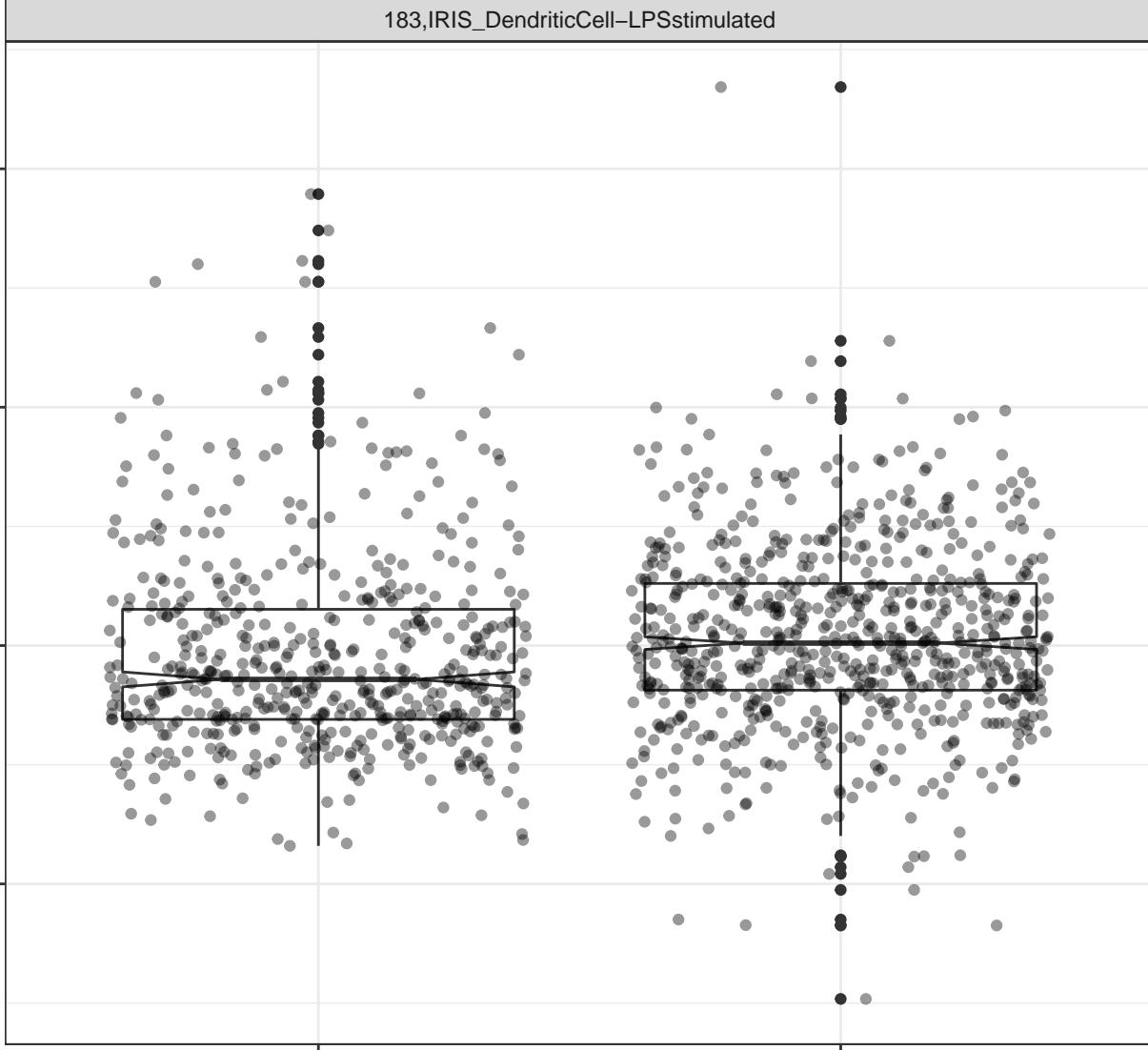
0.00

-0.05

Primary Tumor

Tumor

Solid Tissue Normal



184,REACTOME_TRNA_AMINOACYLATION

Value

0.15

0.10

0.05

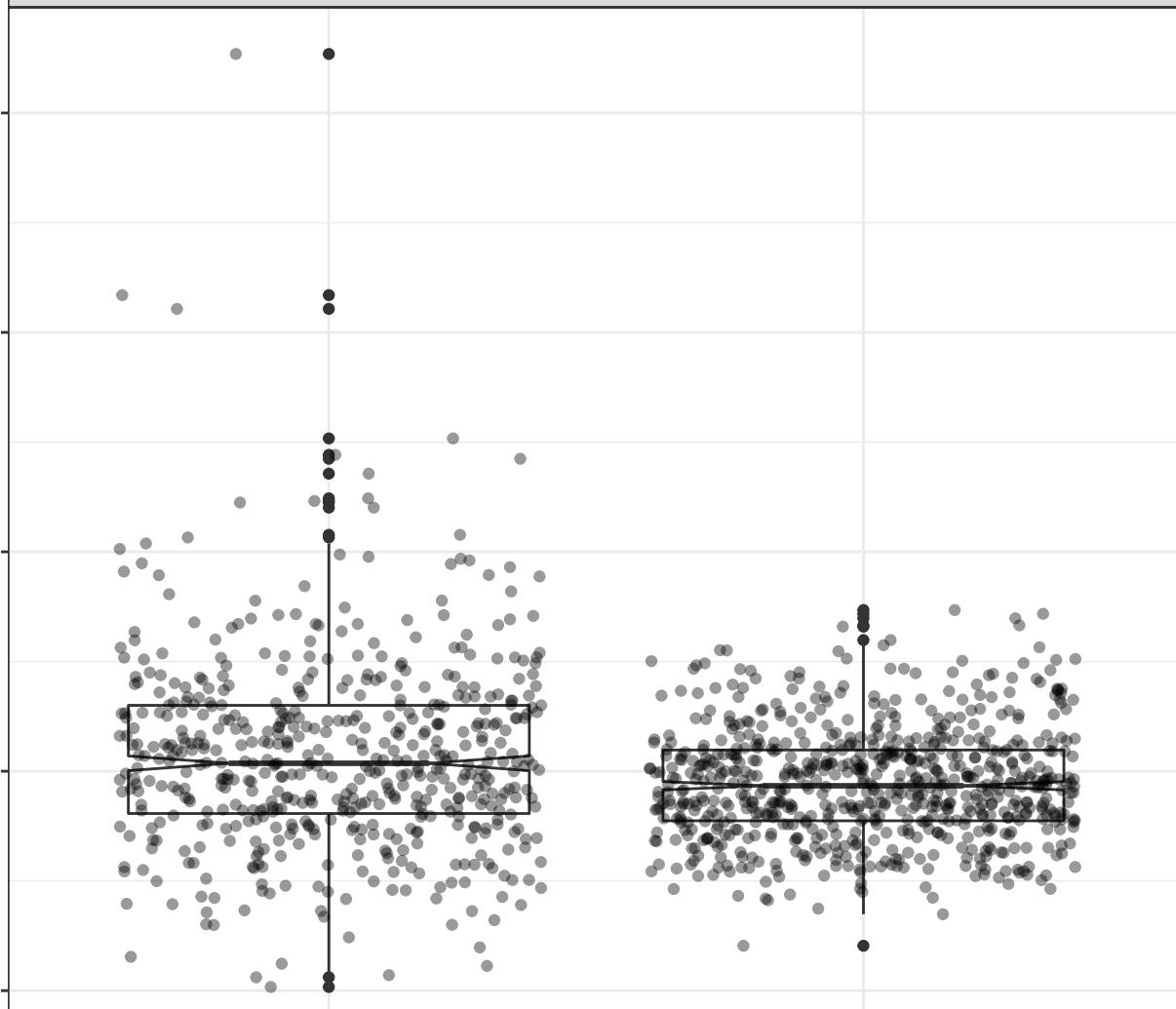
0.00

-0.05

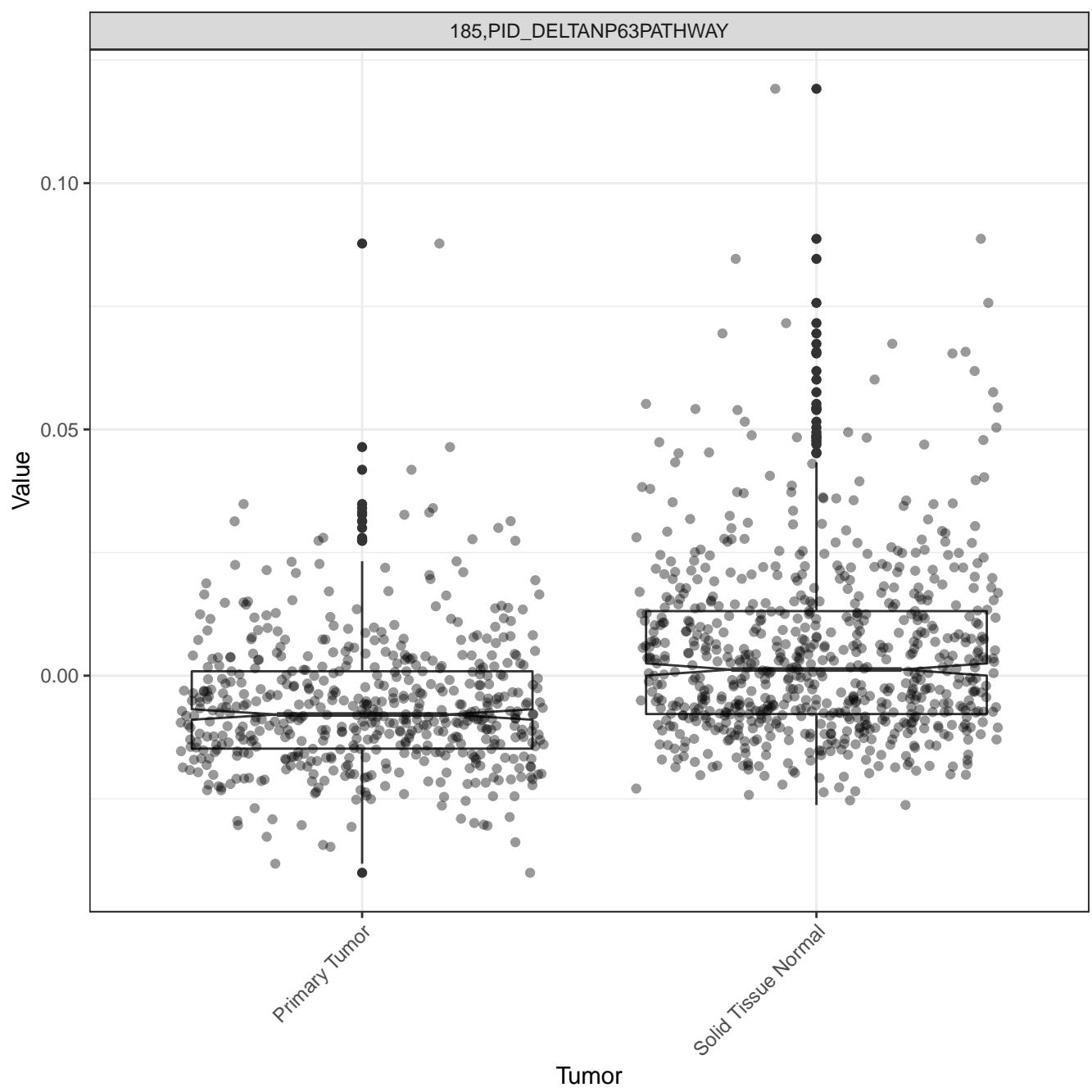
Primary Tumor

Solid Tissue Normal

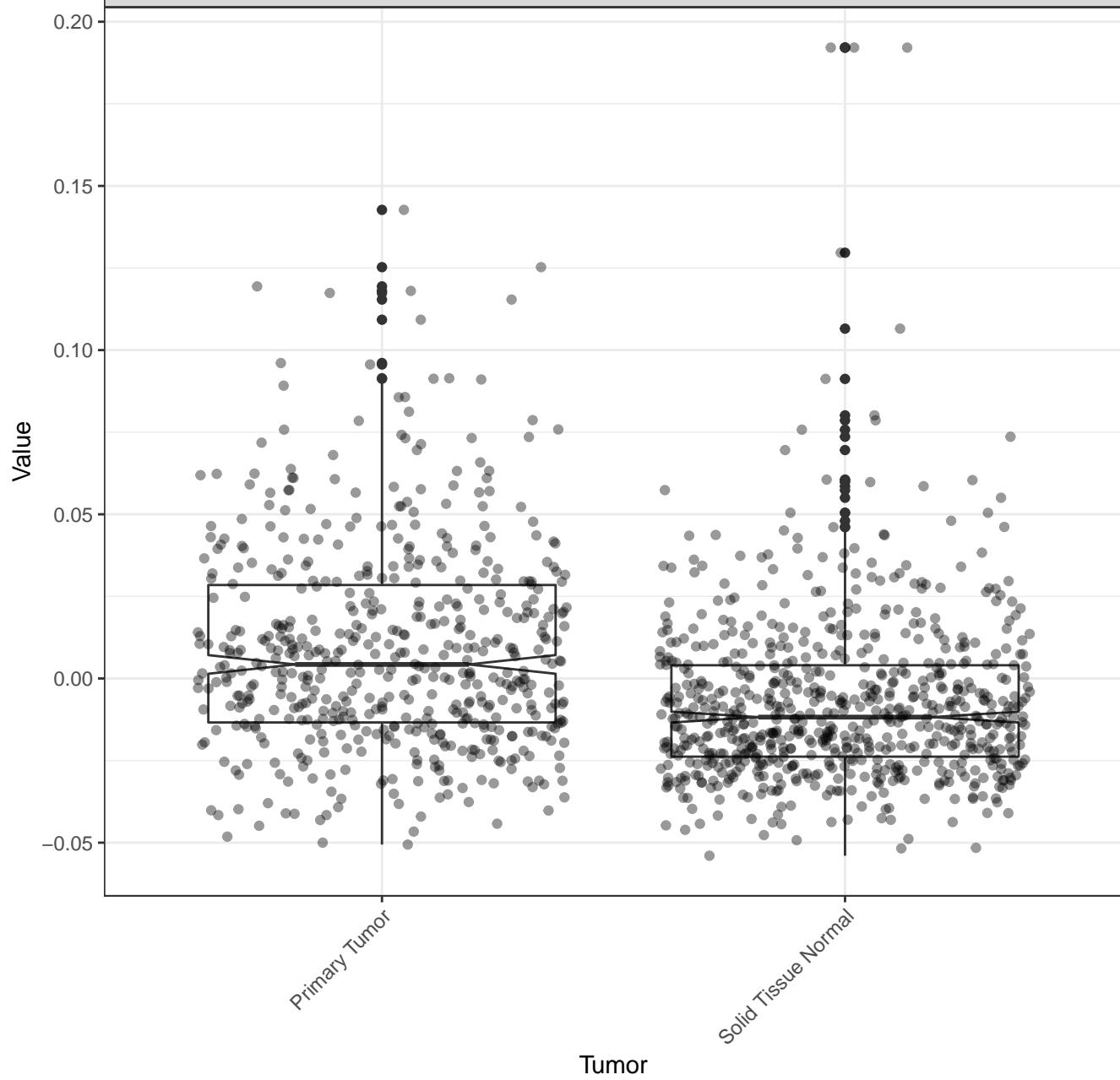
Tumor



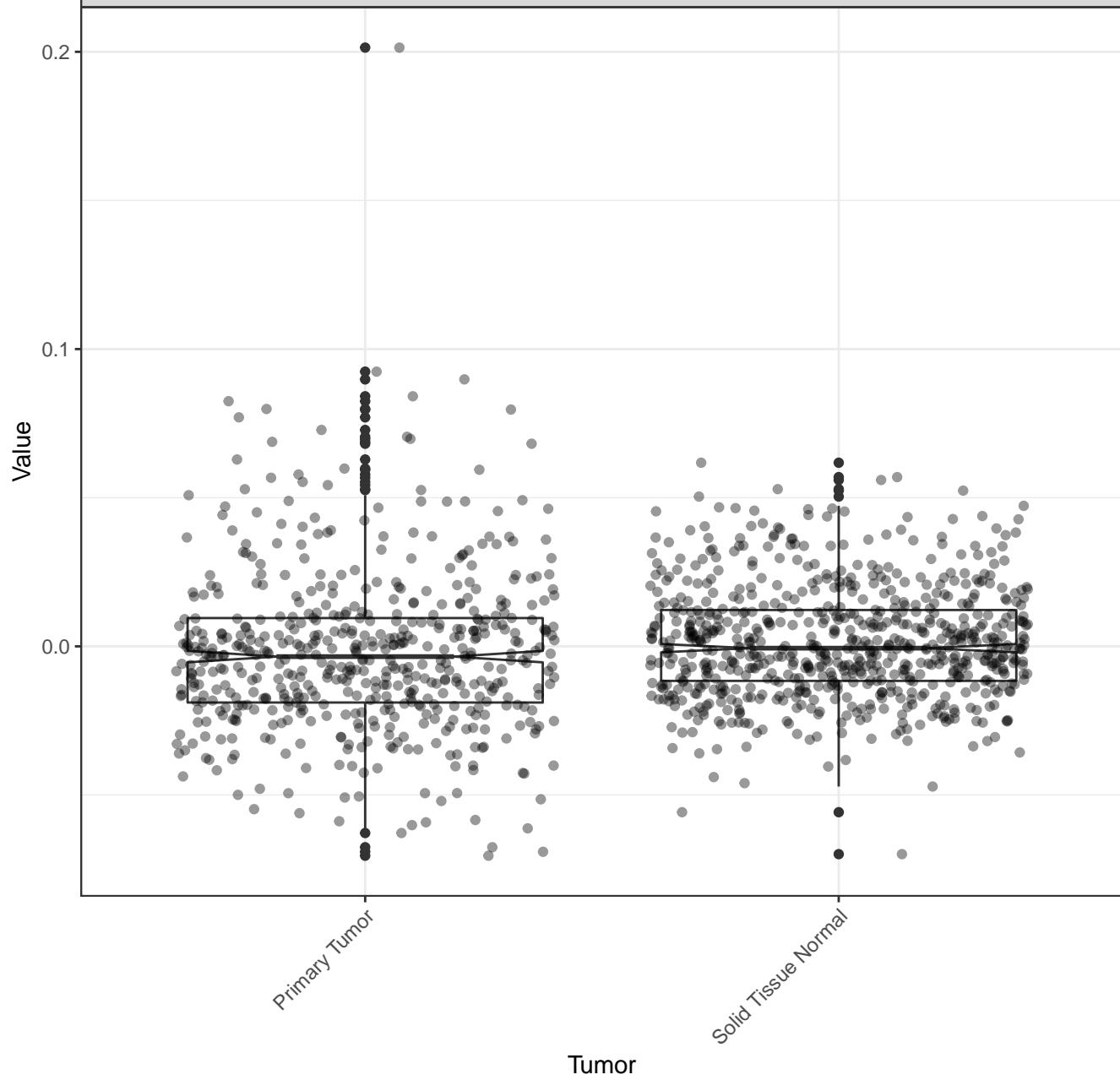
185,PID_DELTANP63PATHWAY



186,SVM T cells CD4 memory activated



187,KEGG_PHOSPHATIDYLINOSITOL_SIGNALING_SYSTEM



188,KEGG_PATHWAYS_IN_CANCER

Value

0.08

0.04

0.00

-0.04

Primary Tumor

Solid Tissue Normal

Tumor



189,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES

Value

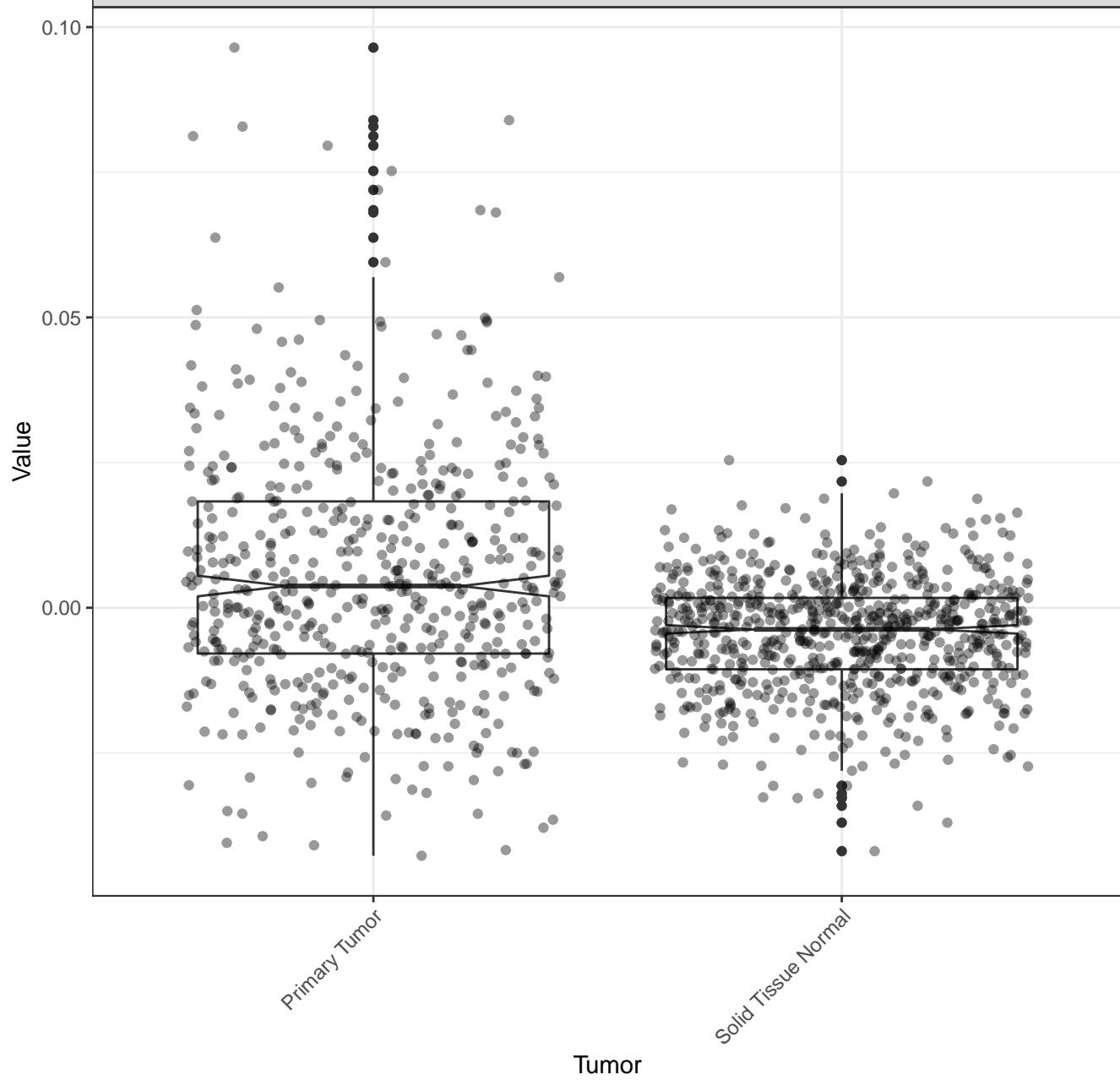
0.15
0.10
0.05
0.00
-0.05

Primary Tumor
Solid Tissue Normal

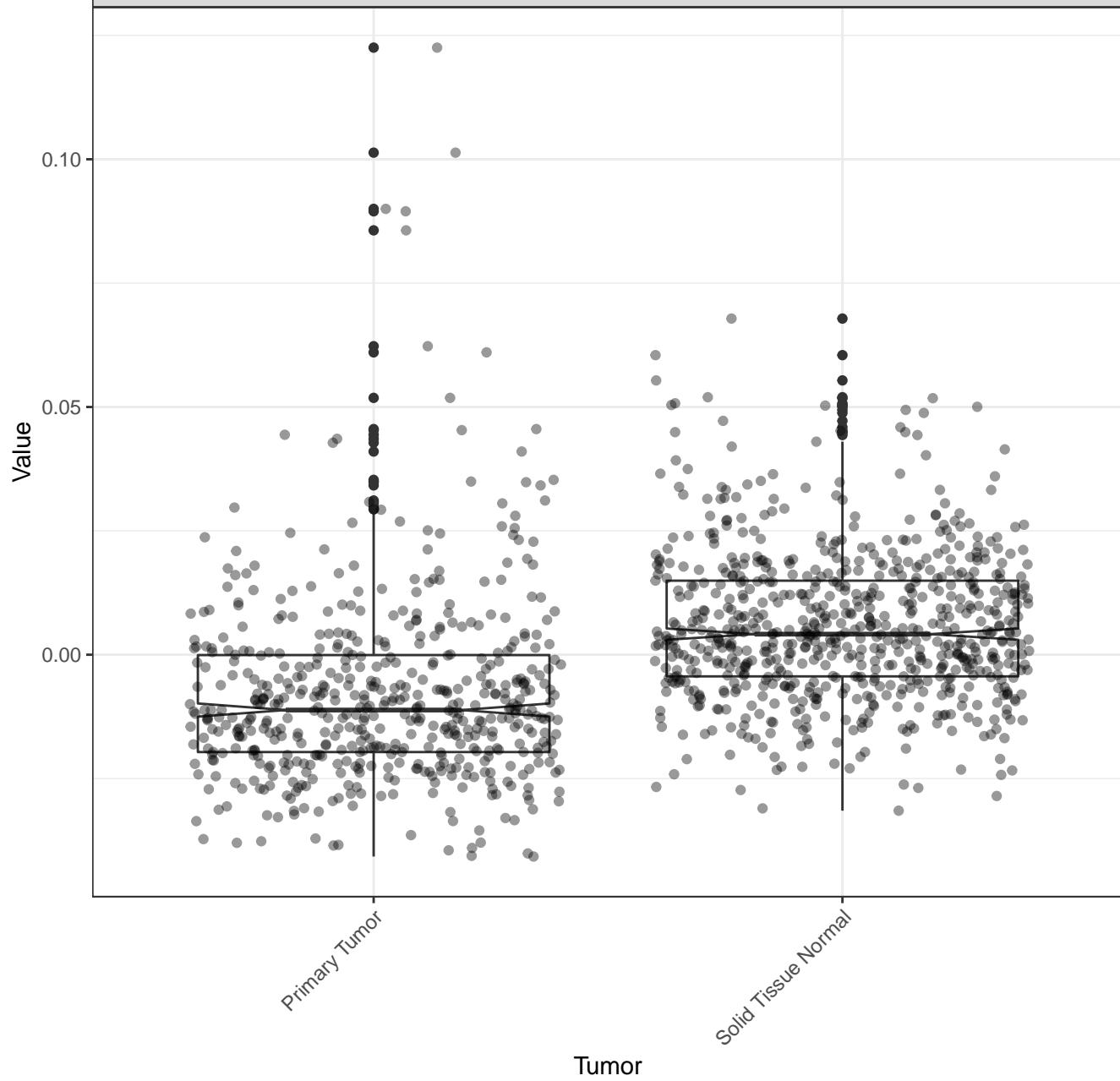
Tumor

The figure displays a box plot comparing gene expression values (Y-axis, ranging from -0.05 to 0.15) across two categories (X-axis): Primary Tumor and Solid Tissue Normal. The Y-axis is labeled 'Value' and has major ticks at -0.05, 0.00, 0.05, 0.10, and 0.15. The X-axis is labeled 'Tumor' and has two categories: 'Primary Tumor' and 'Solid Tissue Normal'. For each category, a box plot is shown with a black horizontal line for the median, a box for the interquartile range, and whiskers extending to the minimum and maximum values. Outliers are represented by individual gray dots. In the 'Primary Tumor' group, the median value is slightly above zero, with a range from approximately -0.03 to 0.06. In the 'Solid Tissue Normal' group, the median value is near zero, with a range from approximately -0.02 to 0.03.

190,MIPS_SPLICEOSOME



191,KEGG_PHOSPHATIDYLINOSITOL_SIGNALING_SYSTEM



192,REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION

Value

0.06

0.03

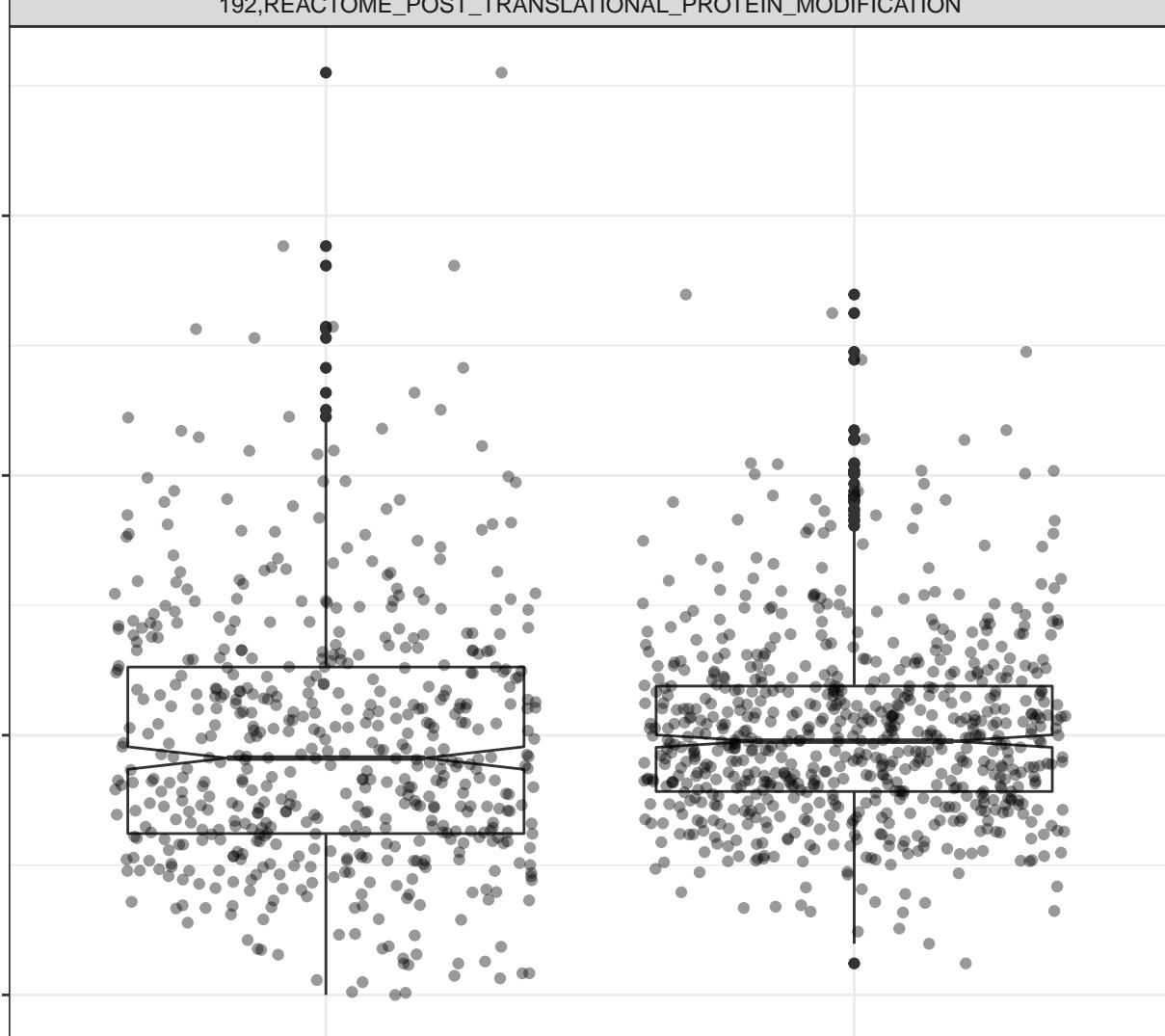
0.00

-0.03

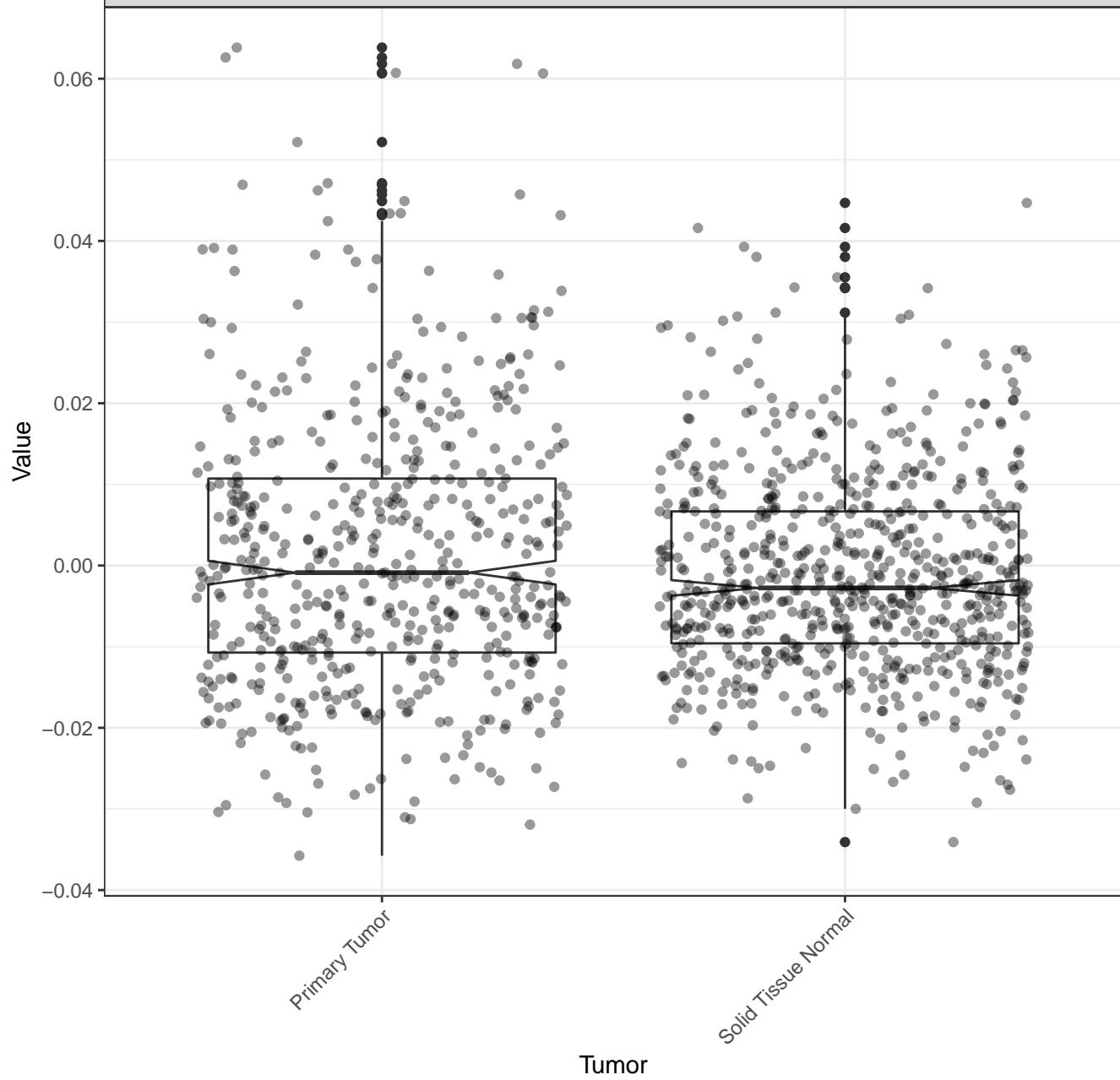
Primary Tumor

Solid Tissue Normal

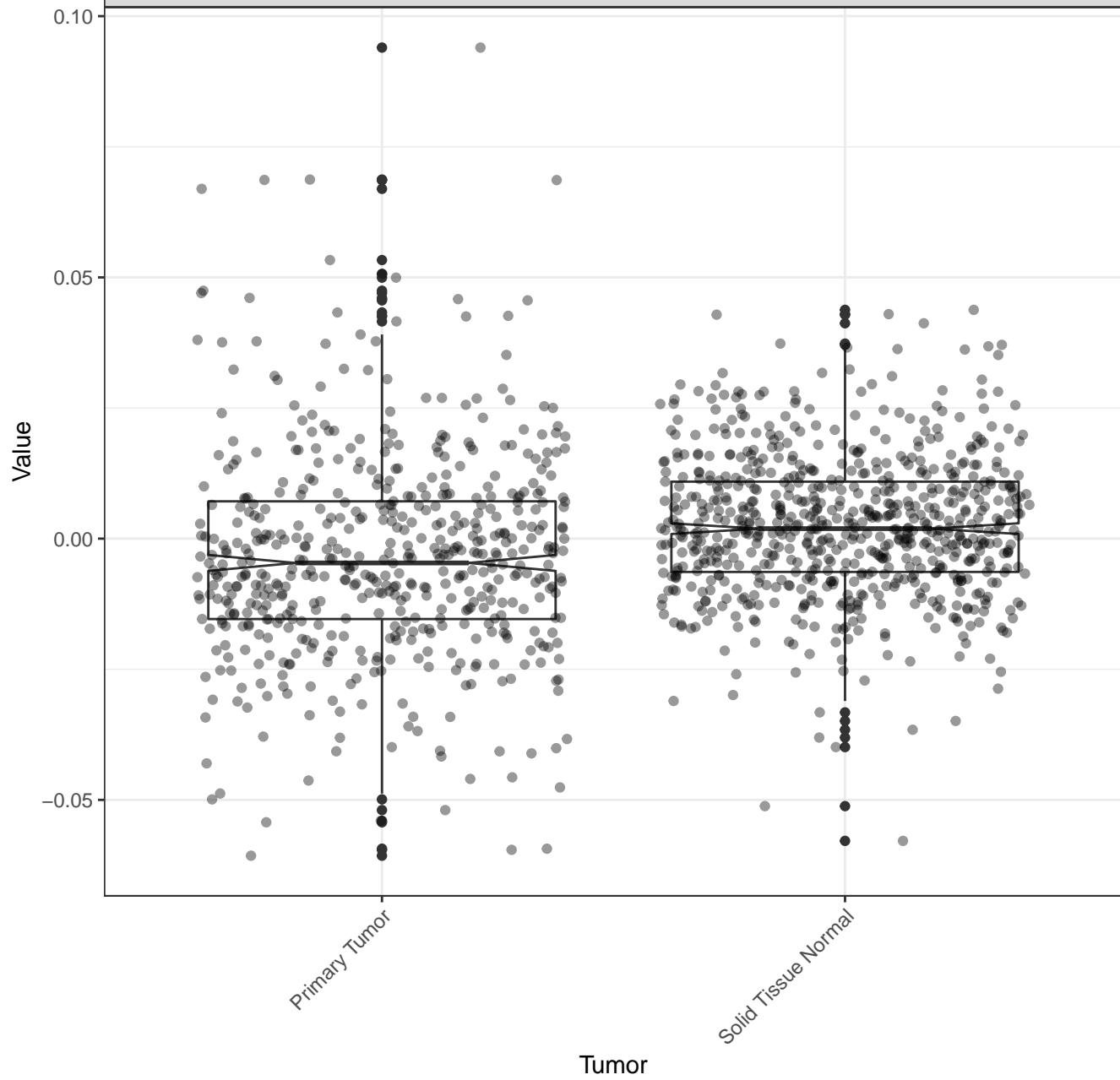
Tumor



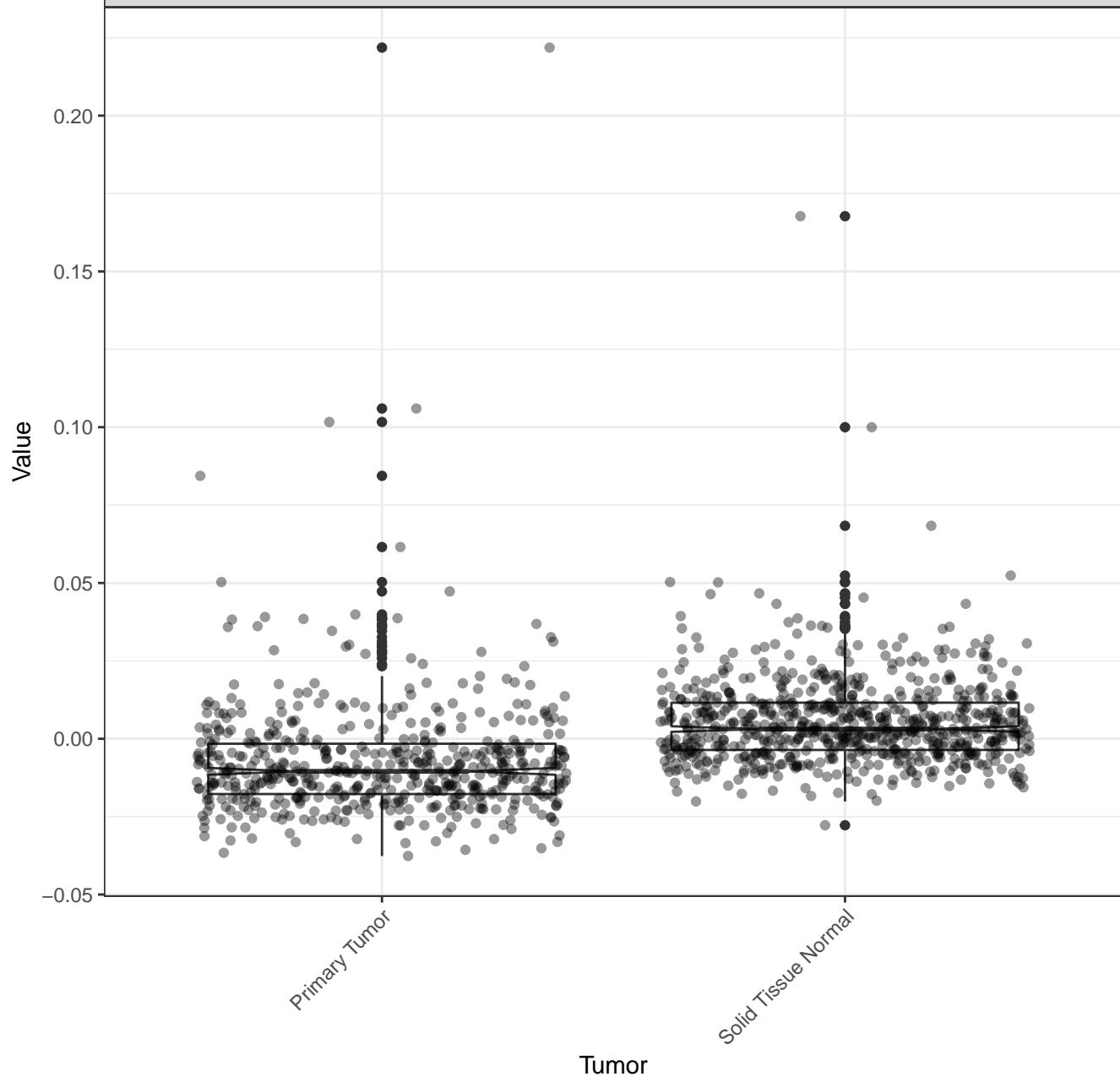
193,DMAP_ERY3



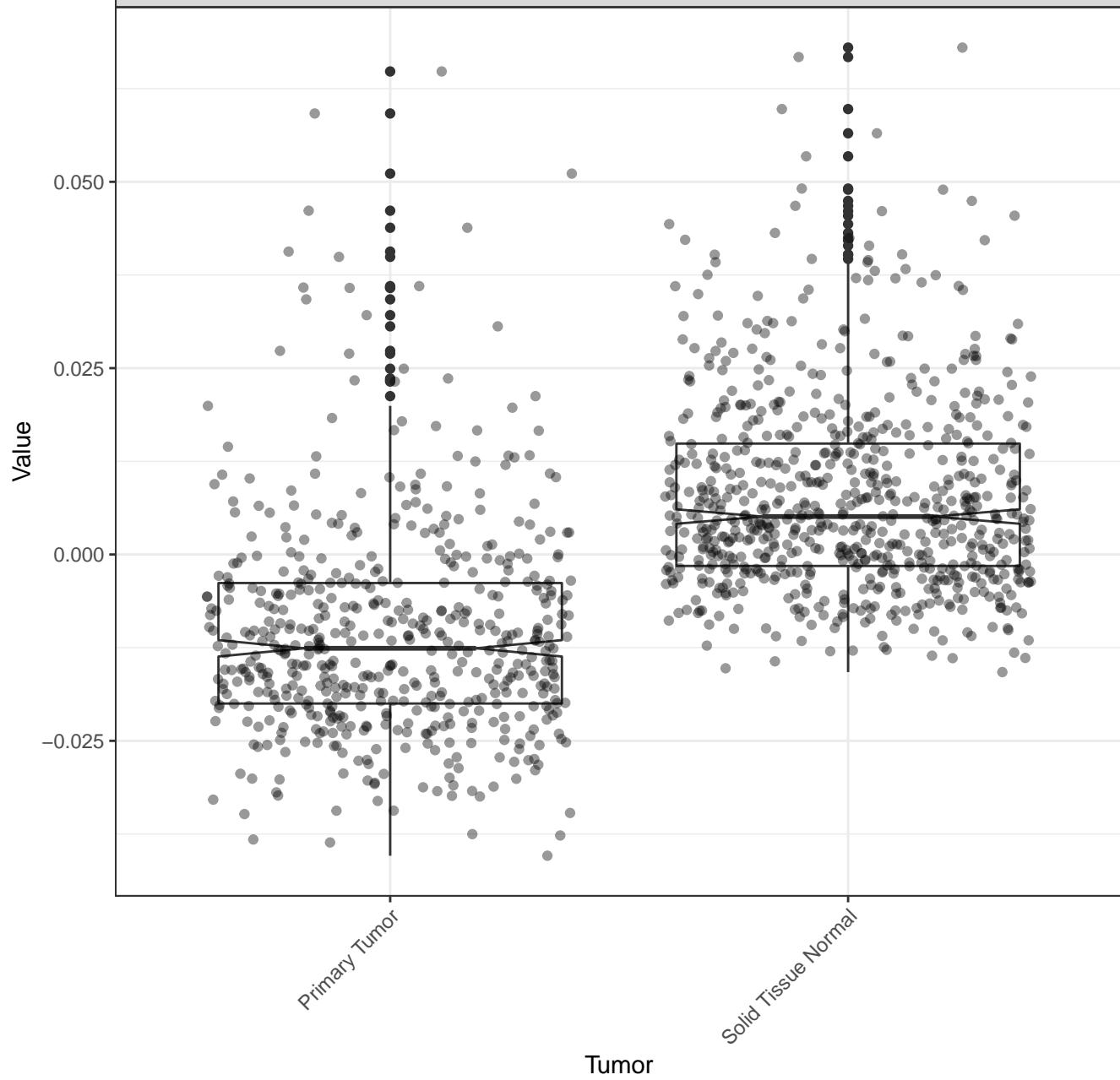
194,REACTOME_INTEGRATION_OF_ENERGY_METABOLISM



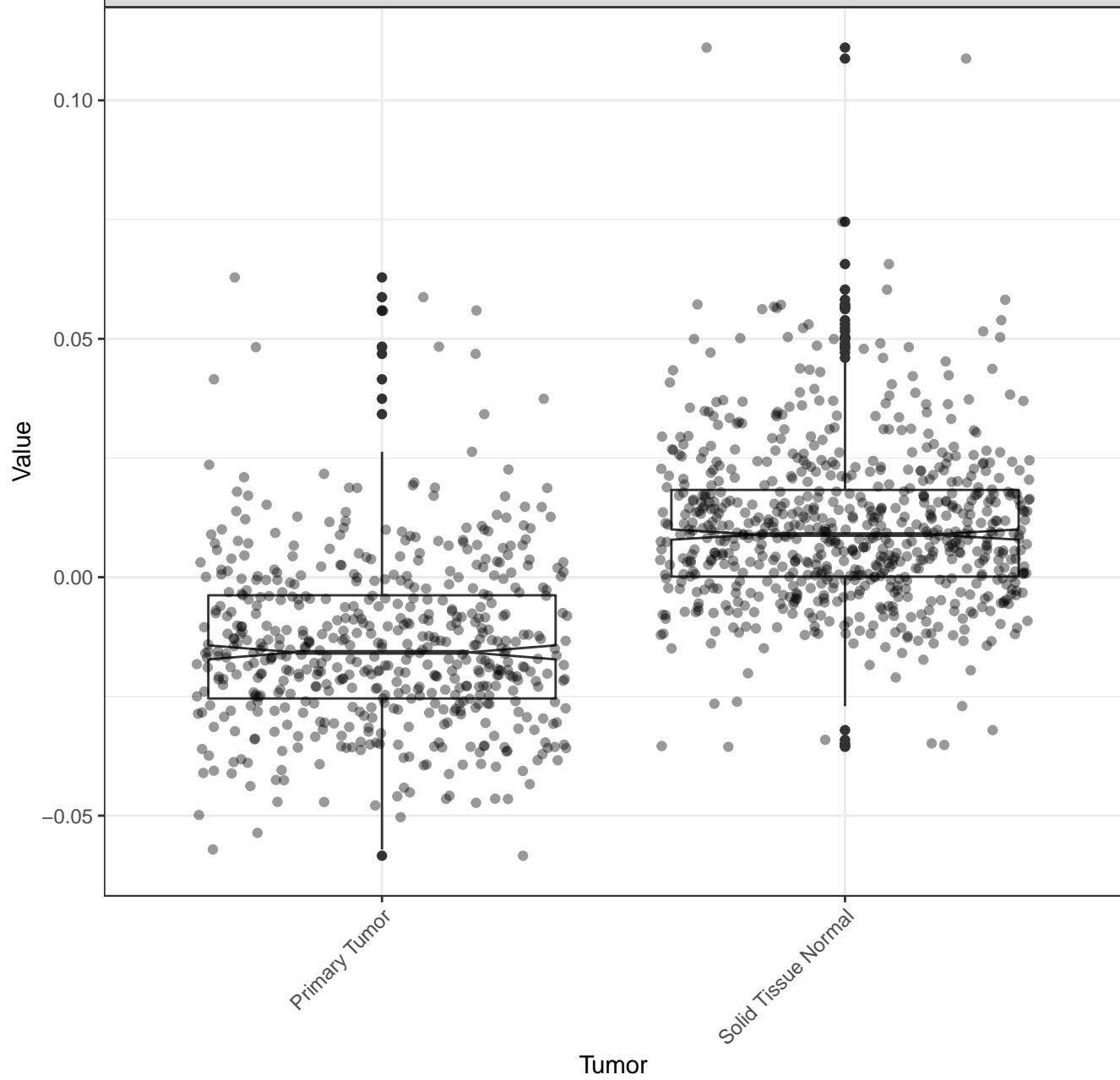
195,DMAP_MONO2



196,KEGG_ABC_TRANSPORTERS



197,REACTOME_NONSENSE_MEDIATED_DECAY_ENHANCED_BY_THE_EXON_JUNCTION_COMPLEX



LV 198

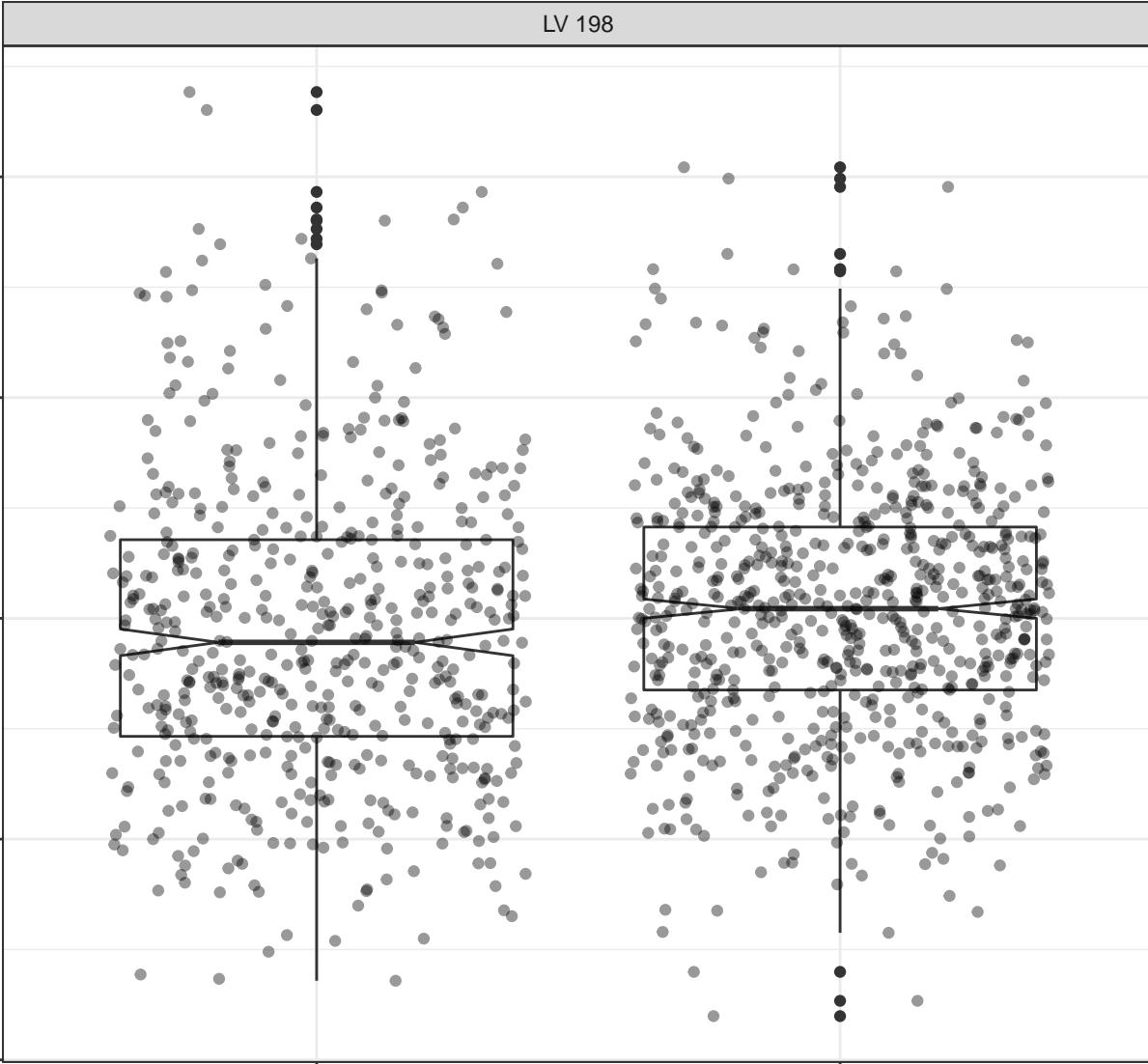
Value

0.050
0.025
0.000
-0.025
-0.050

Primary Tumor

Solid Tissue Normal

Tumor



199,KEGG_AUTOIMMUNE_THYROID_DISEASE

Value

0.6

0.4

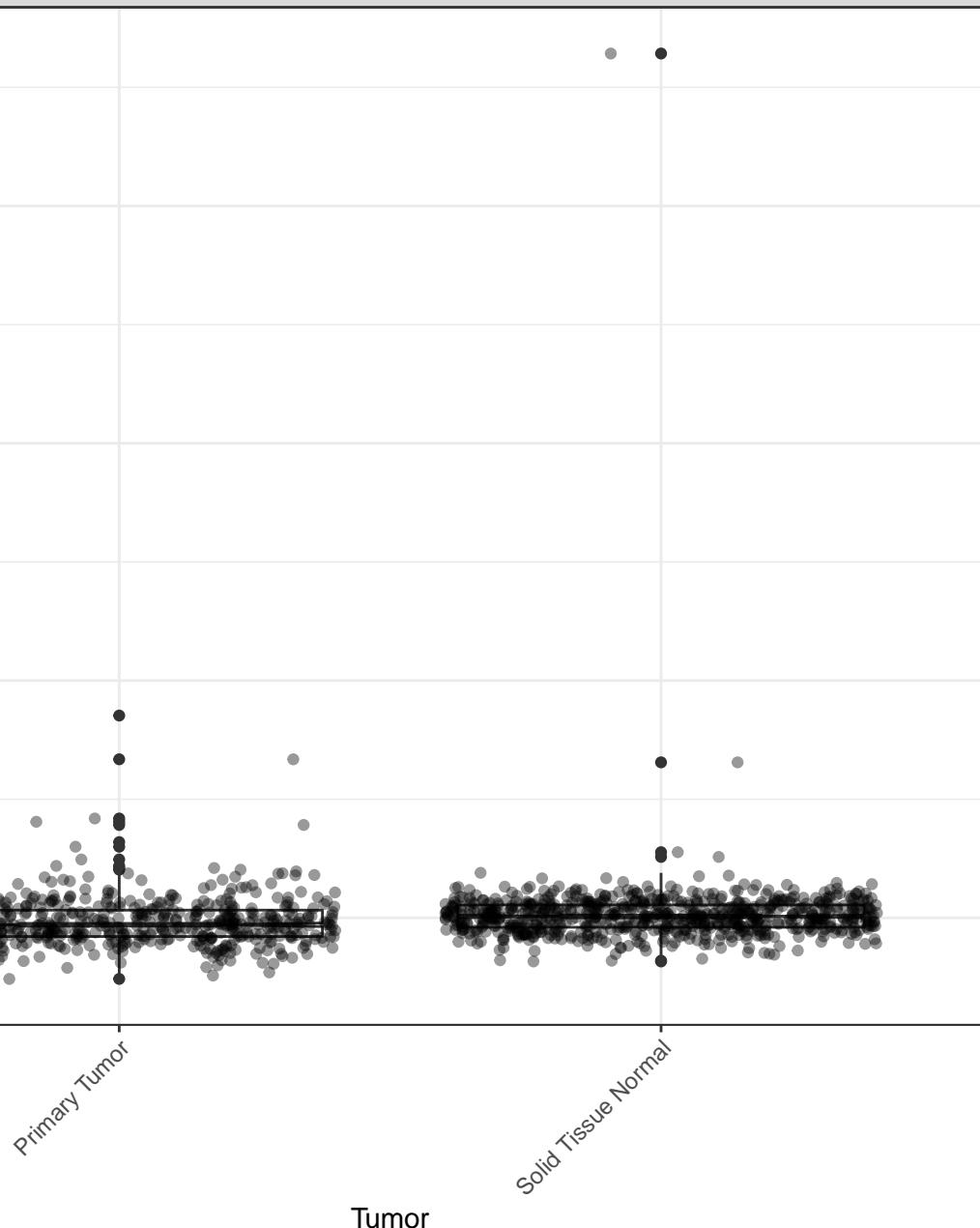
0.2

0.0

Primary Tumor

Solid Tissue Normal

Tumor



LV 200

Value

0.15

0.10

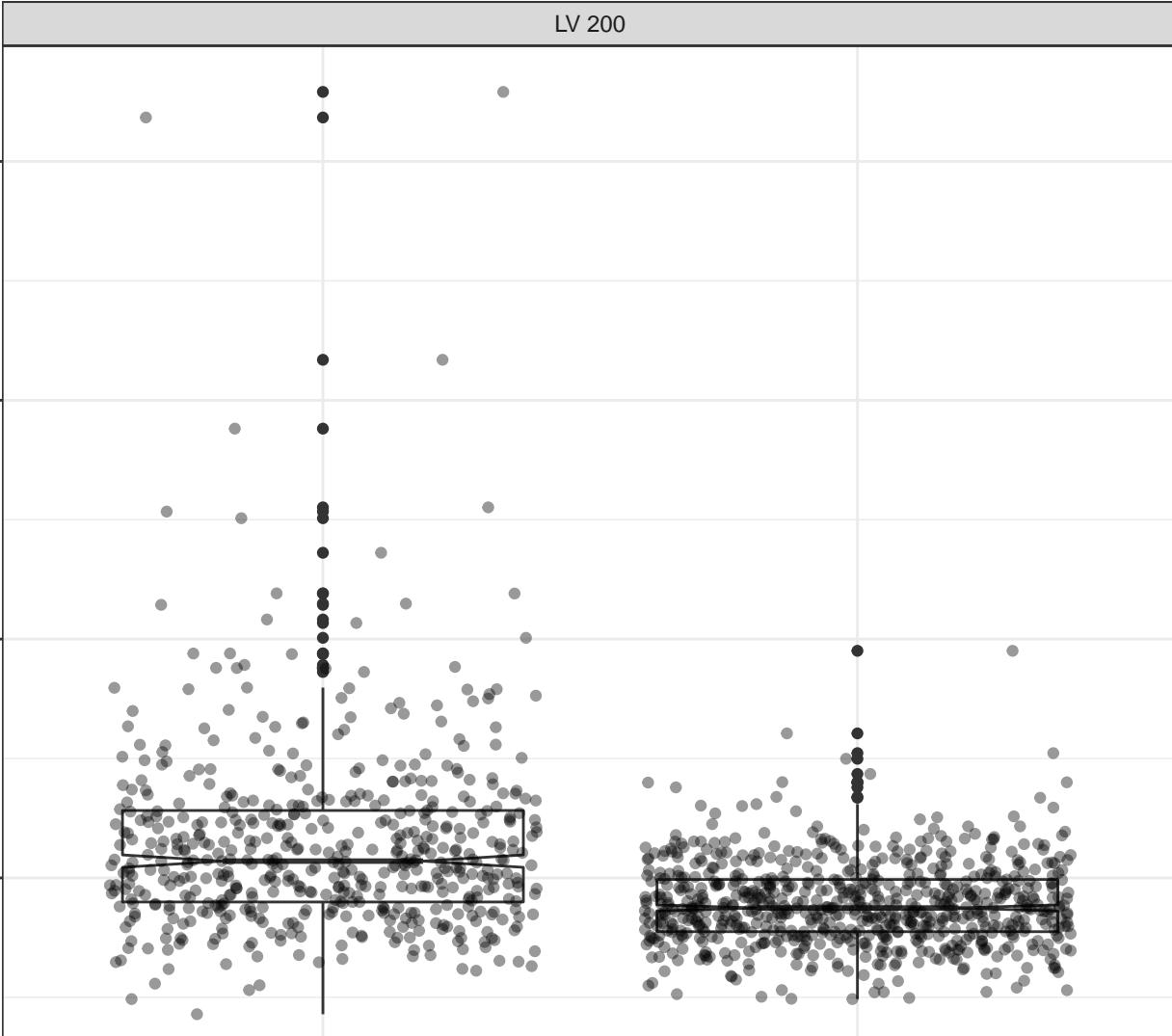
0.05

0.00

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.10

0.05

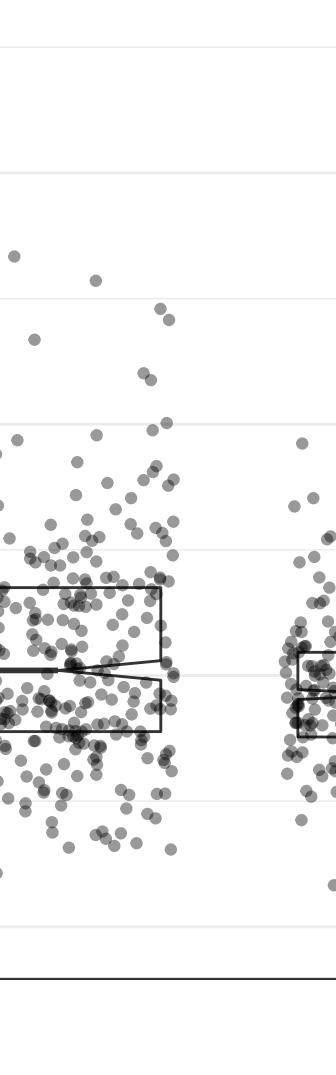
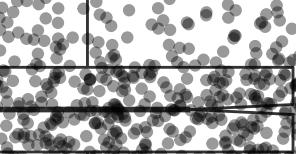
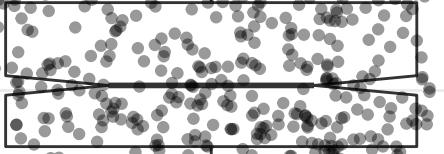
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 202

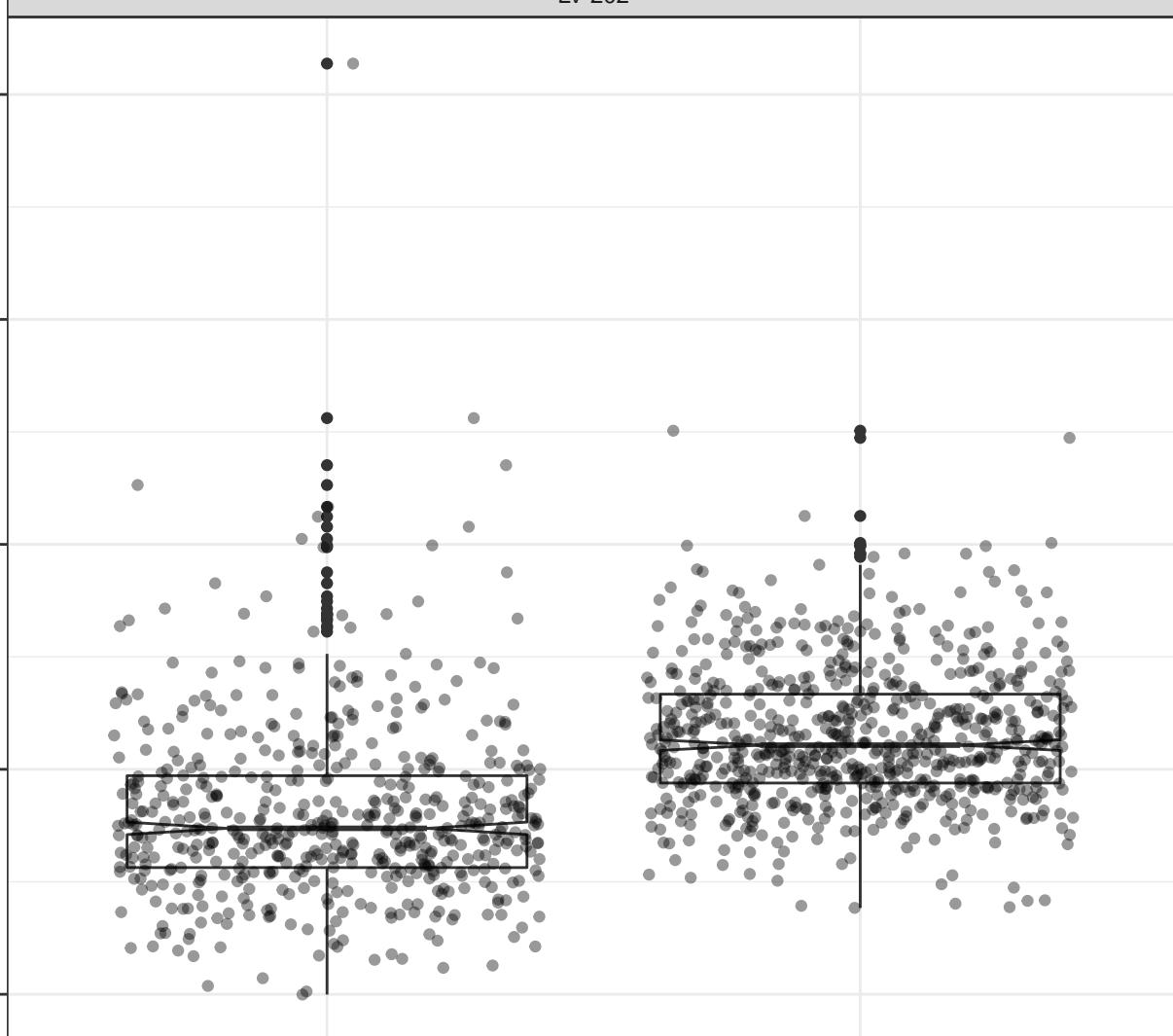
Value

0.15
0.10
0.05
0.00
-0.05

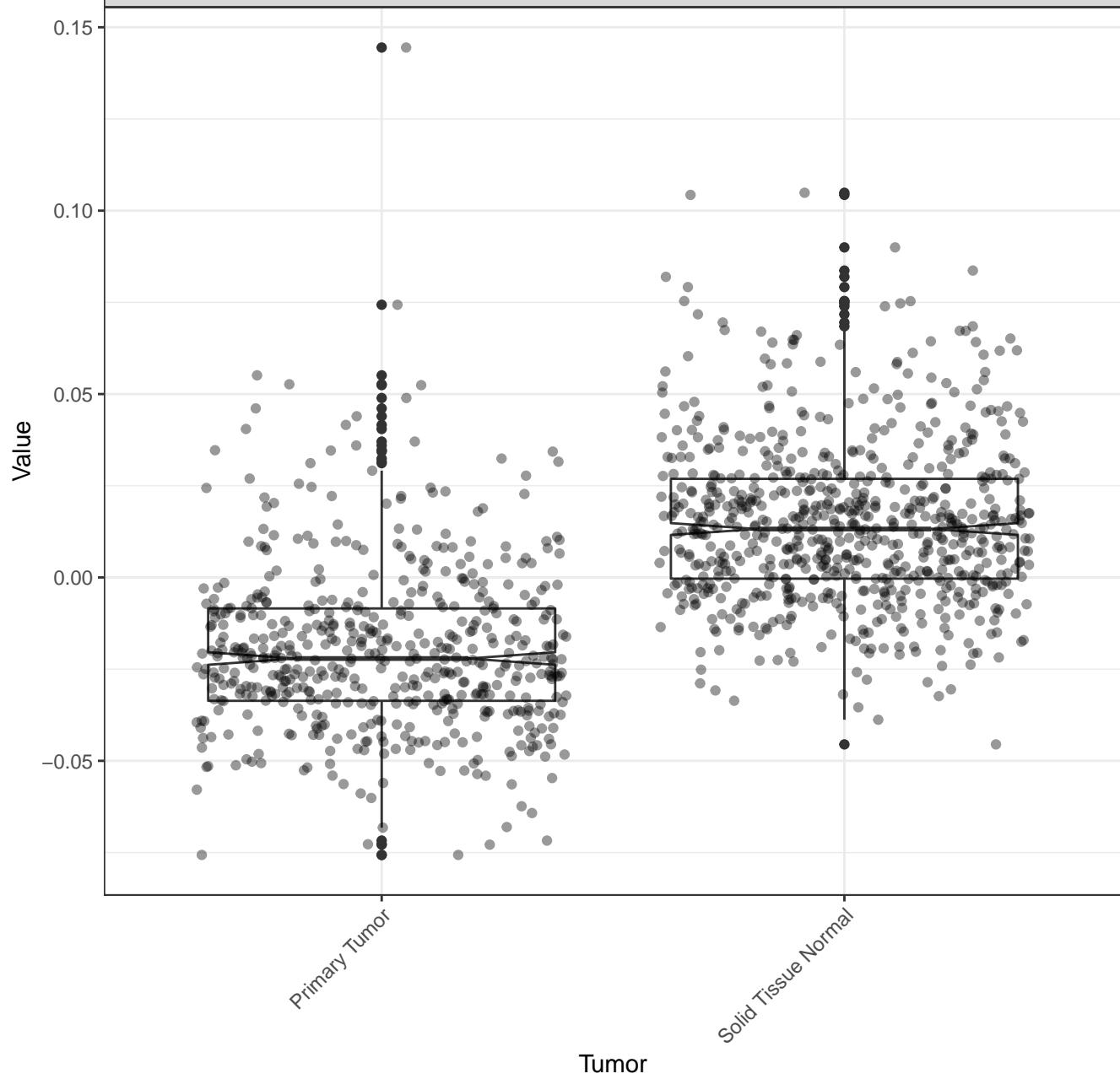
Primary Tumor

Solid Tissue Normal

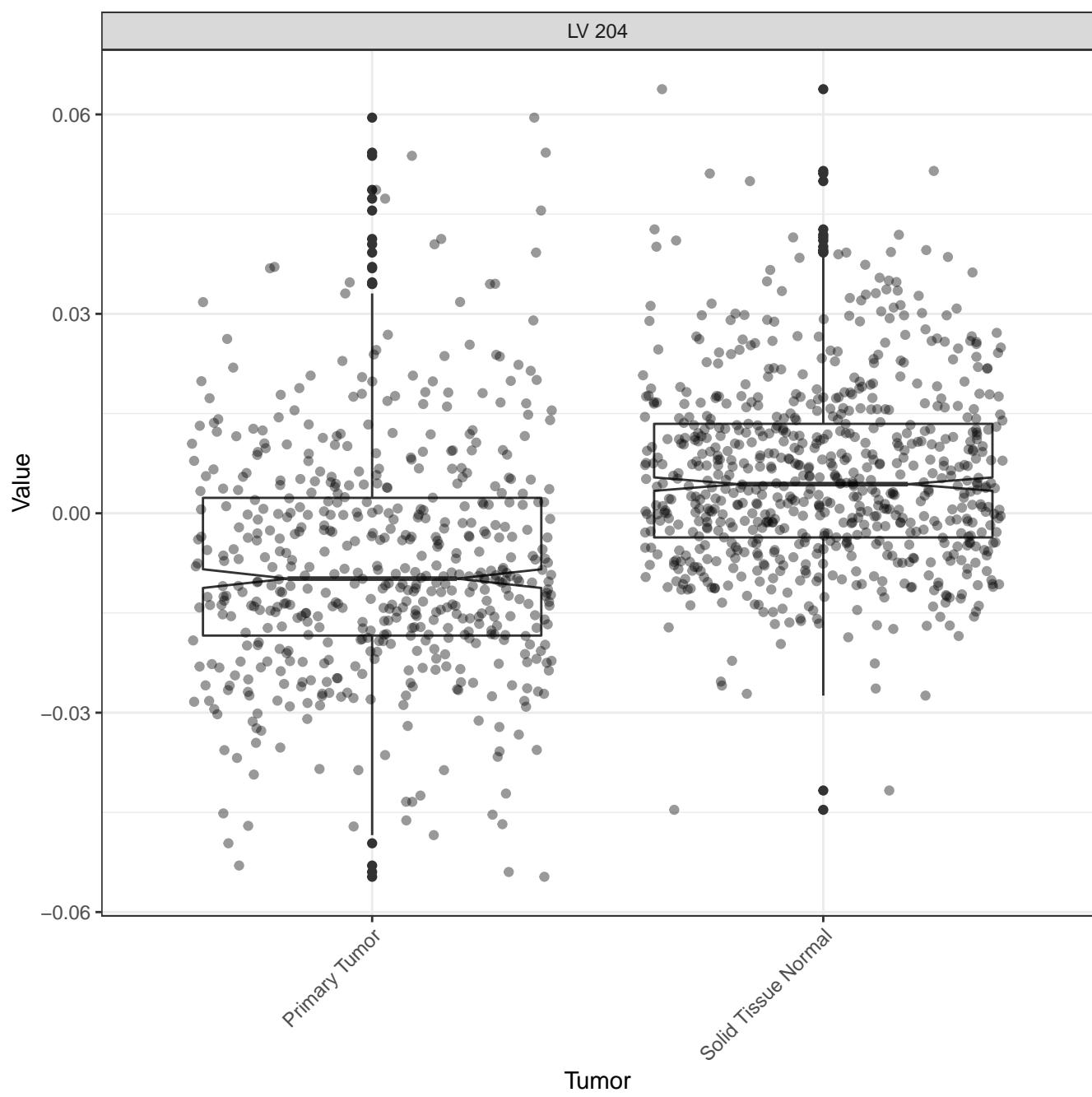
Tumor



E_NEUROTRANSMITTER_RECEPTOR_BINDING_AND_DOWNSTREAM_TRANSMISSION_IN_THE_POSTSY



LV 204



LV 205

Value

0.10

0.05

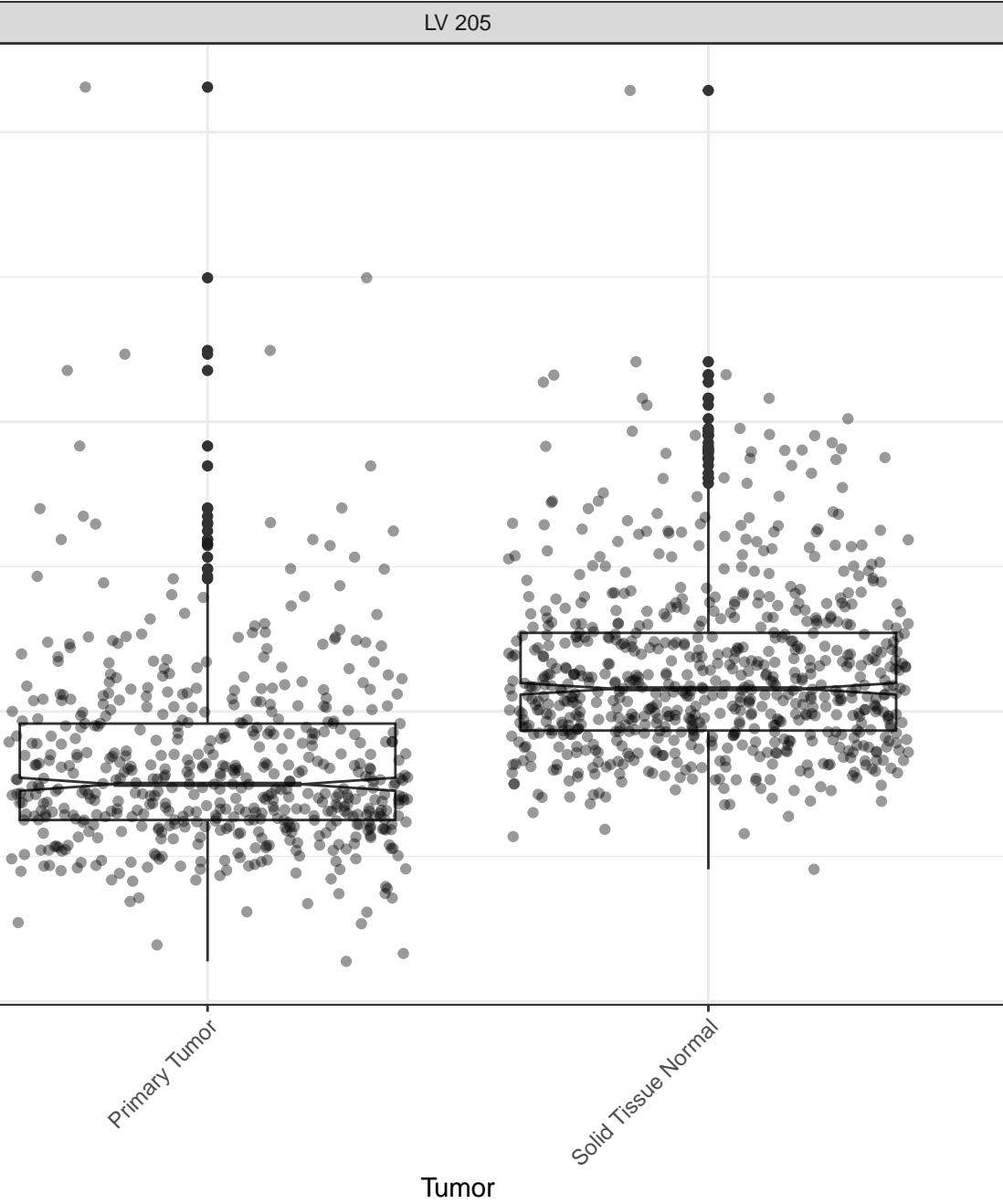
0.00

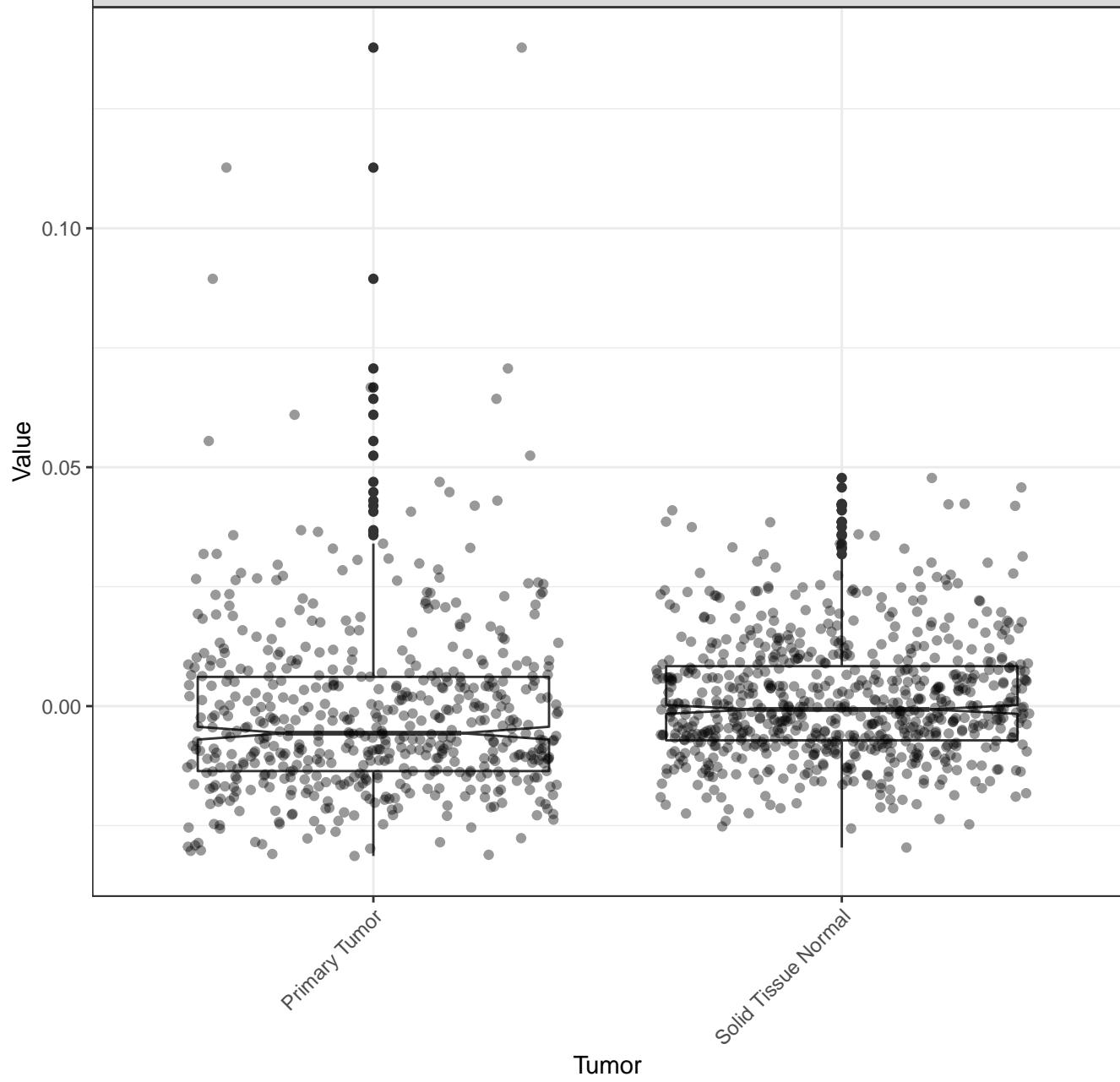
-0.05

Primary Tumor

Solid Tissue Normal

Tumor





207,REACTOME_GLOBAL_GENOMIC_NER_GG_NER

Value

0.15

0.10

0.05

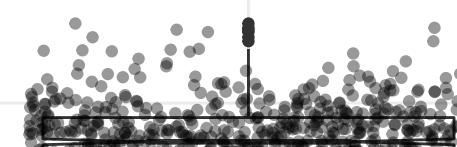
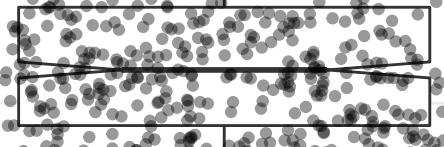
0.00

-0.05

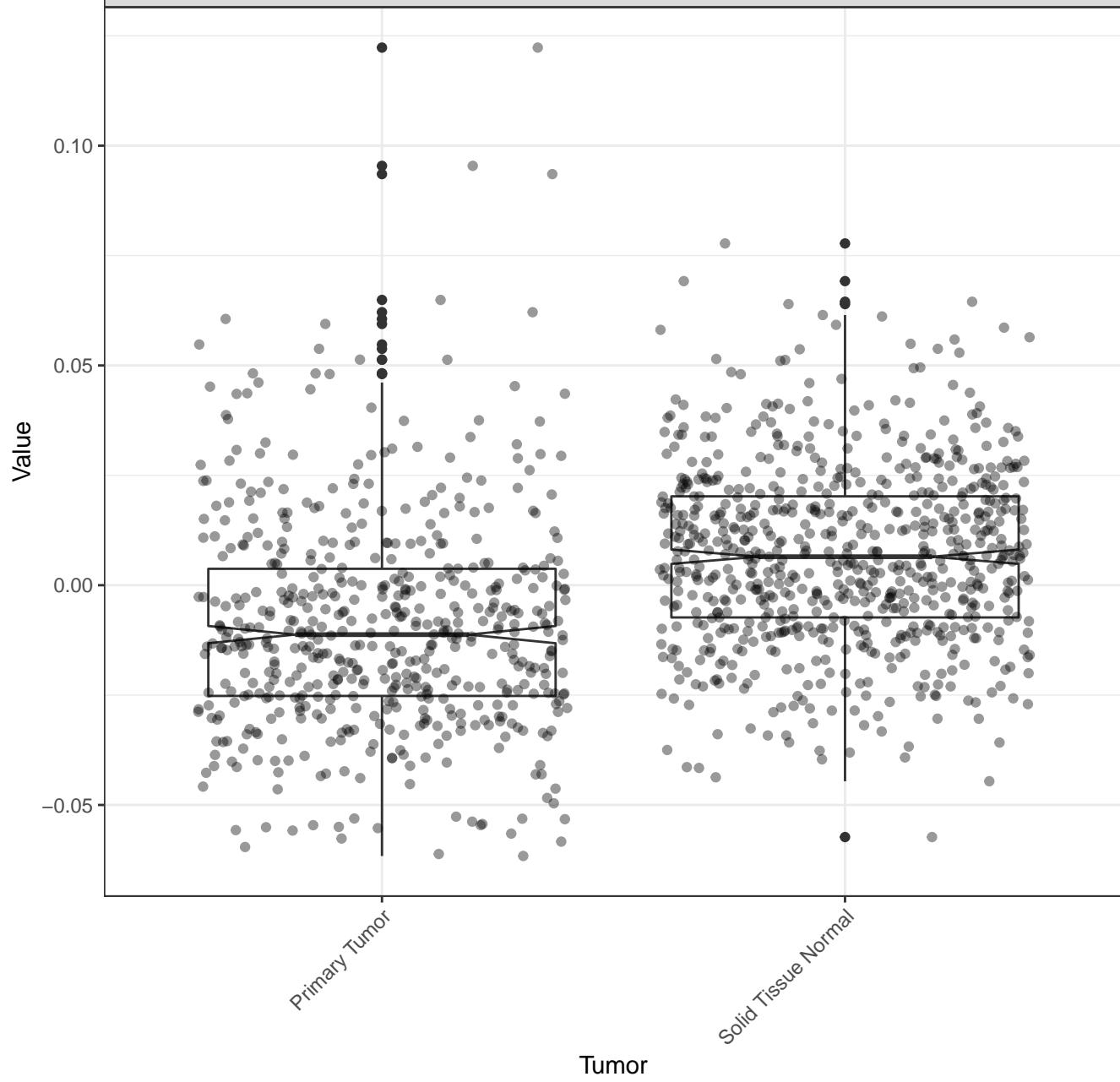
Primary Tumor

Solid Tissue Normal

Tumor



208,KEGG_BASAL_TRANSRIPTION_FACTORS



209,PID_CERAMIDE_PATHWAY

Value

0.05

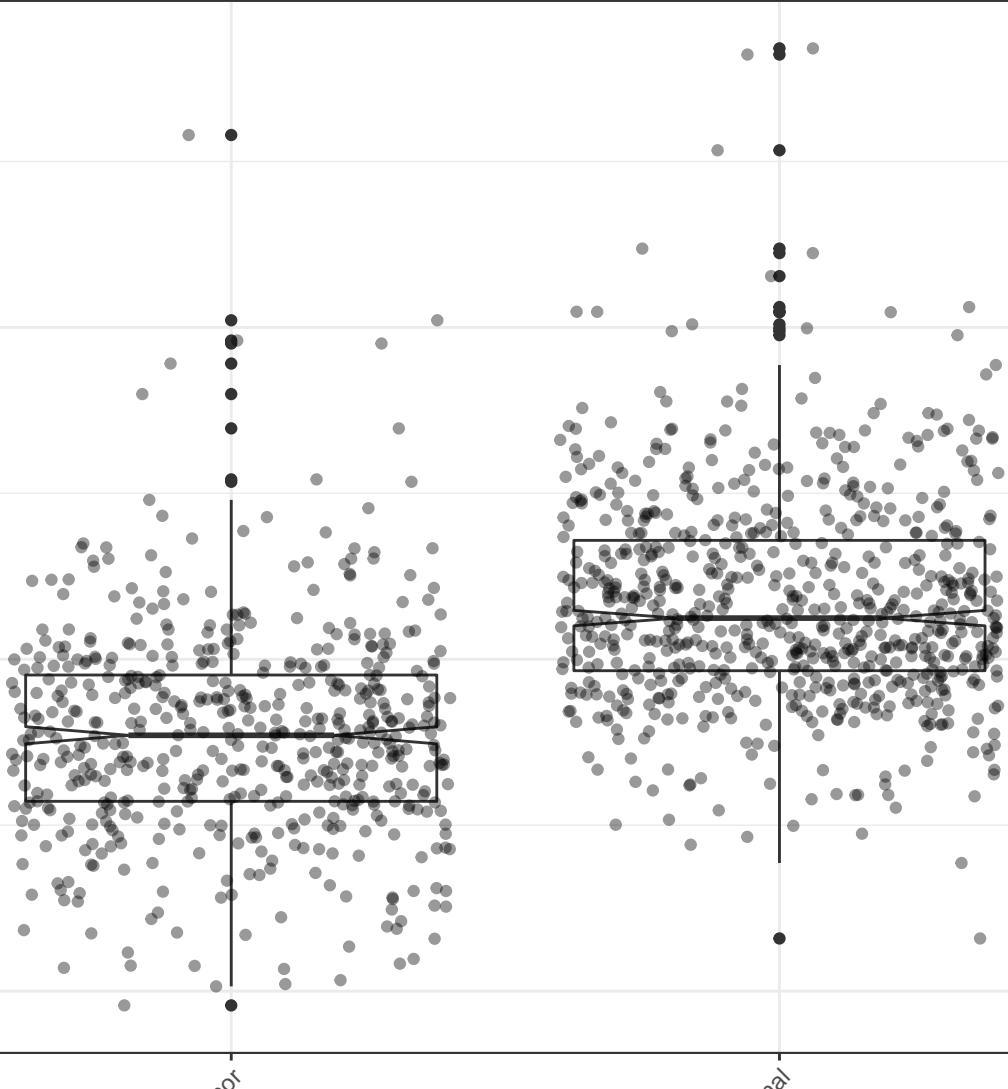
0.00

-0.05

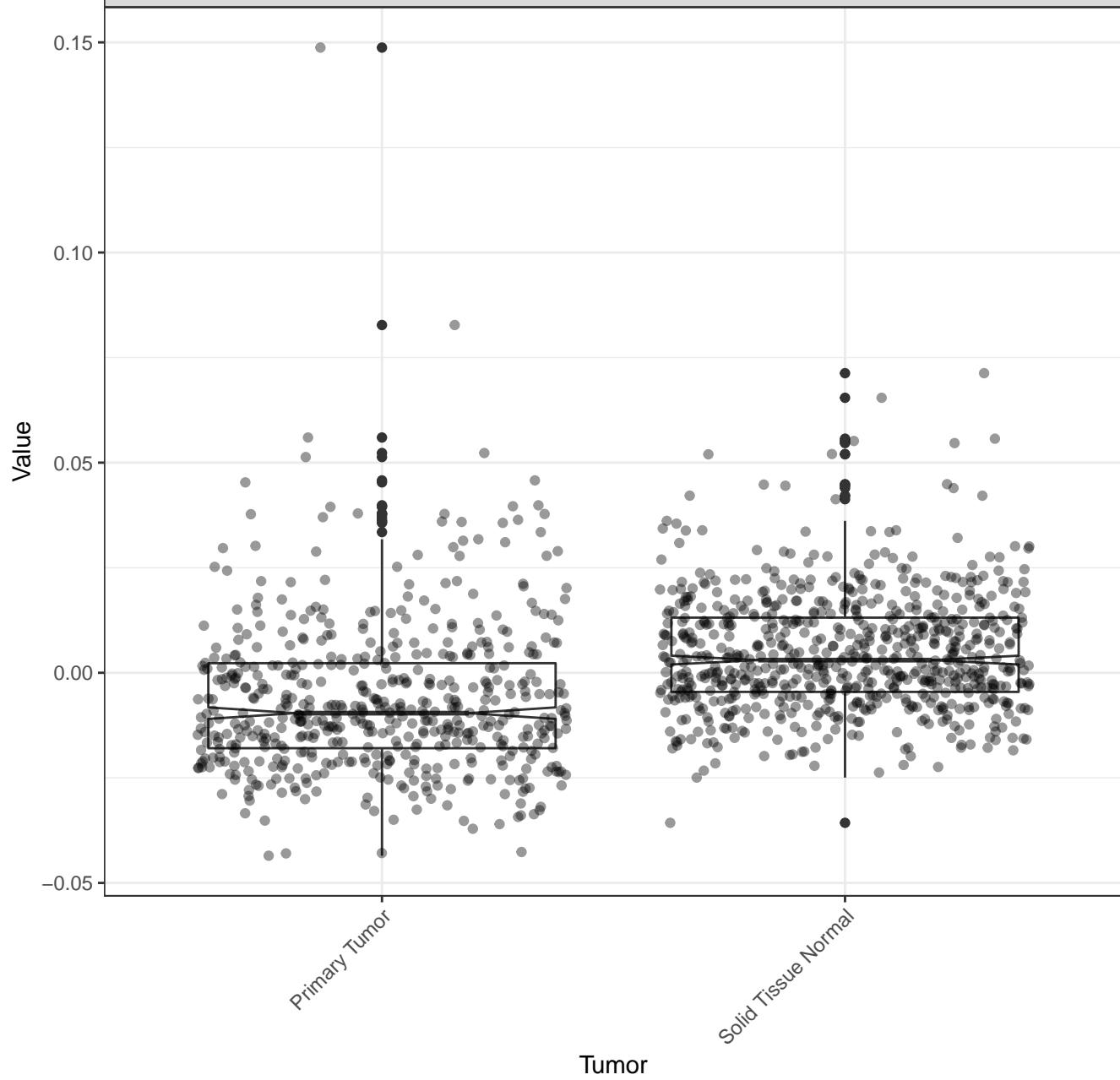
Primary Tumor

Solid Tissue Normal

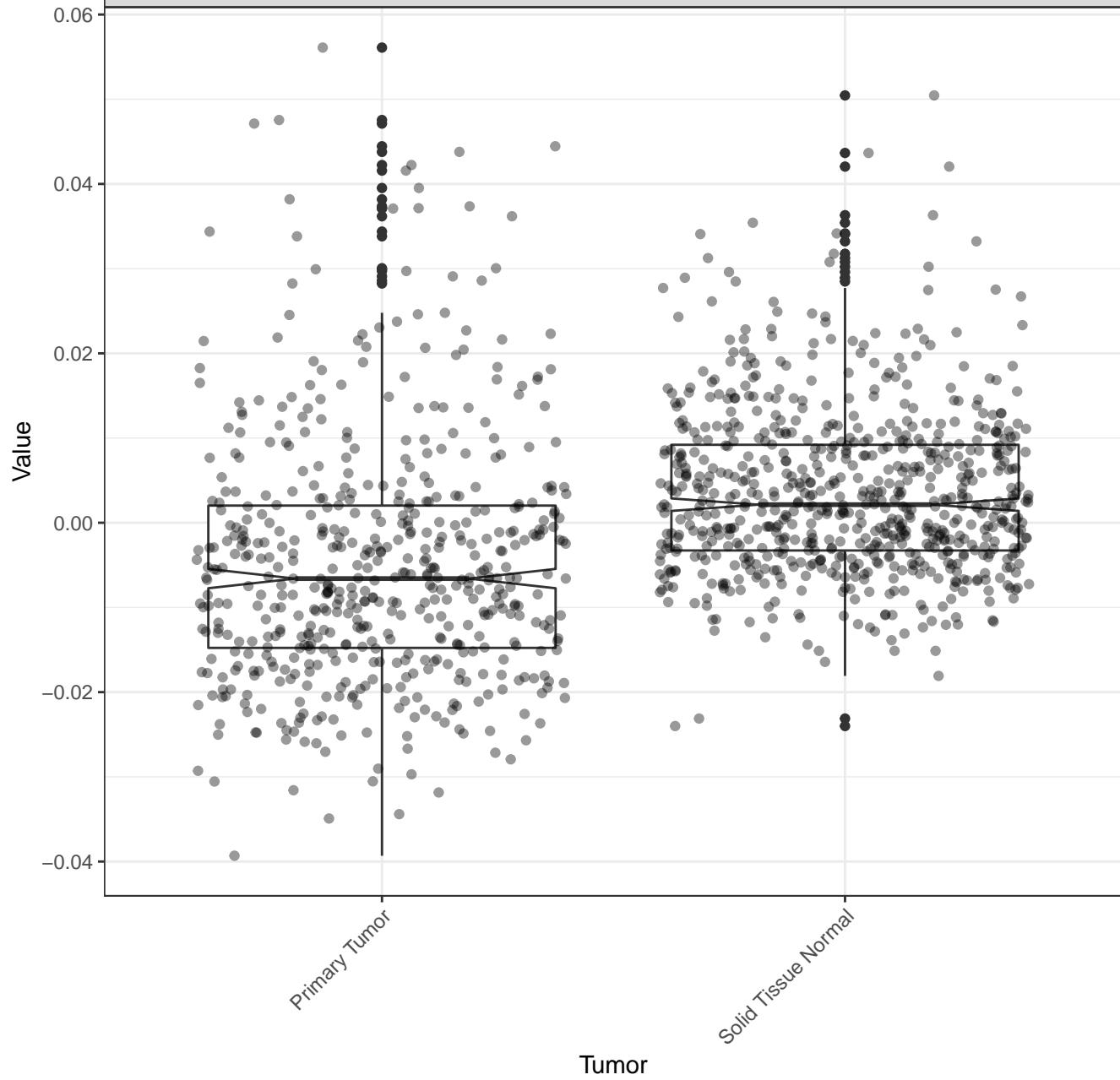
Tumor



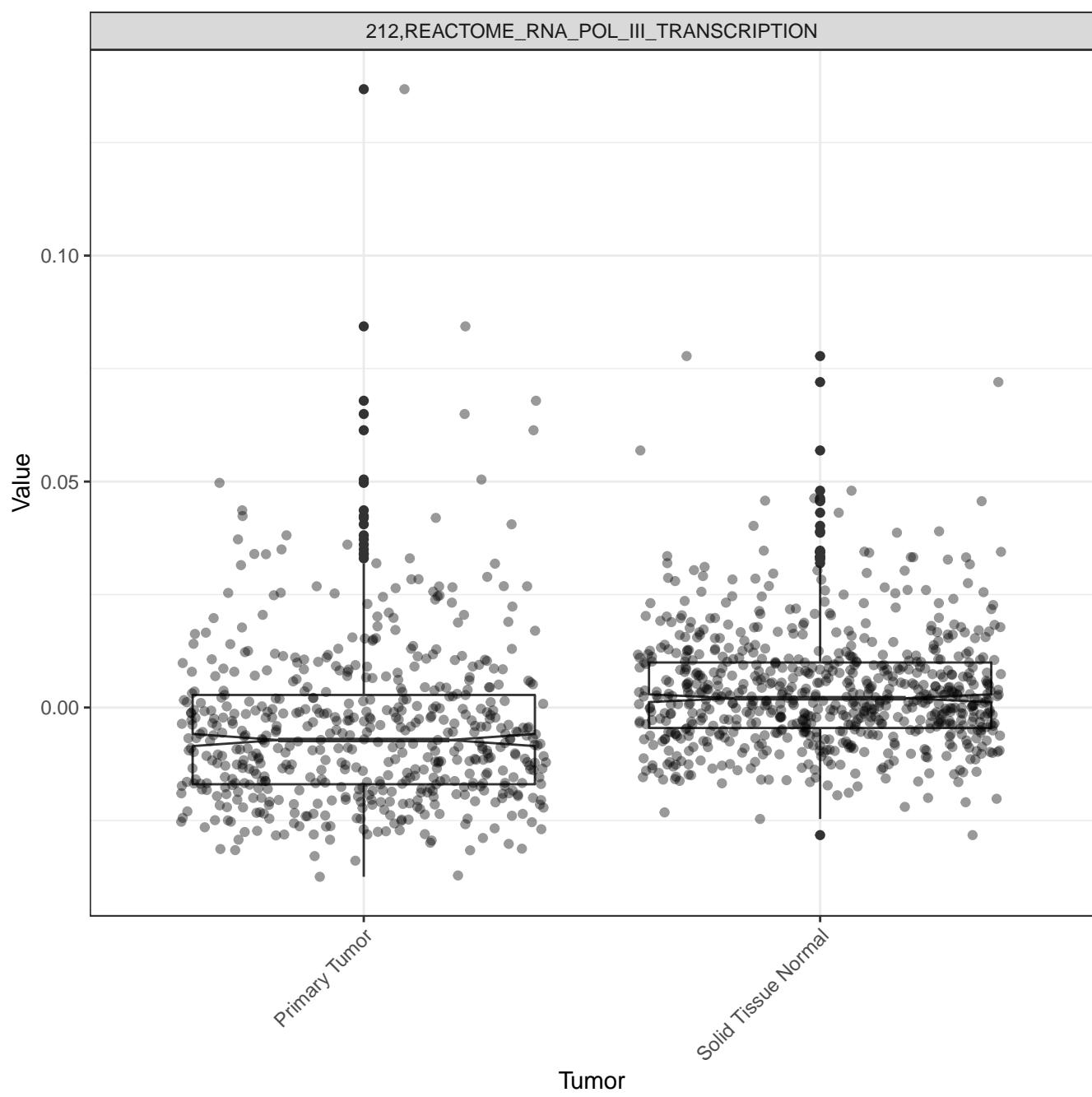
210,PID_HES_HEYPATHWAY



211,IRIS_PlasmaCell–FromBoneMarrow



212,REACTOME_RNA_POL_III_TRANSCRIPTION



213,PID_MYC_ACTIVPATHWAY

Value

0.05

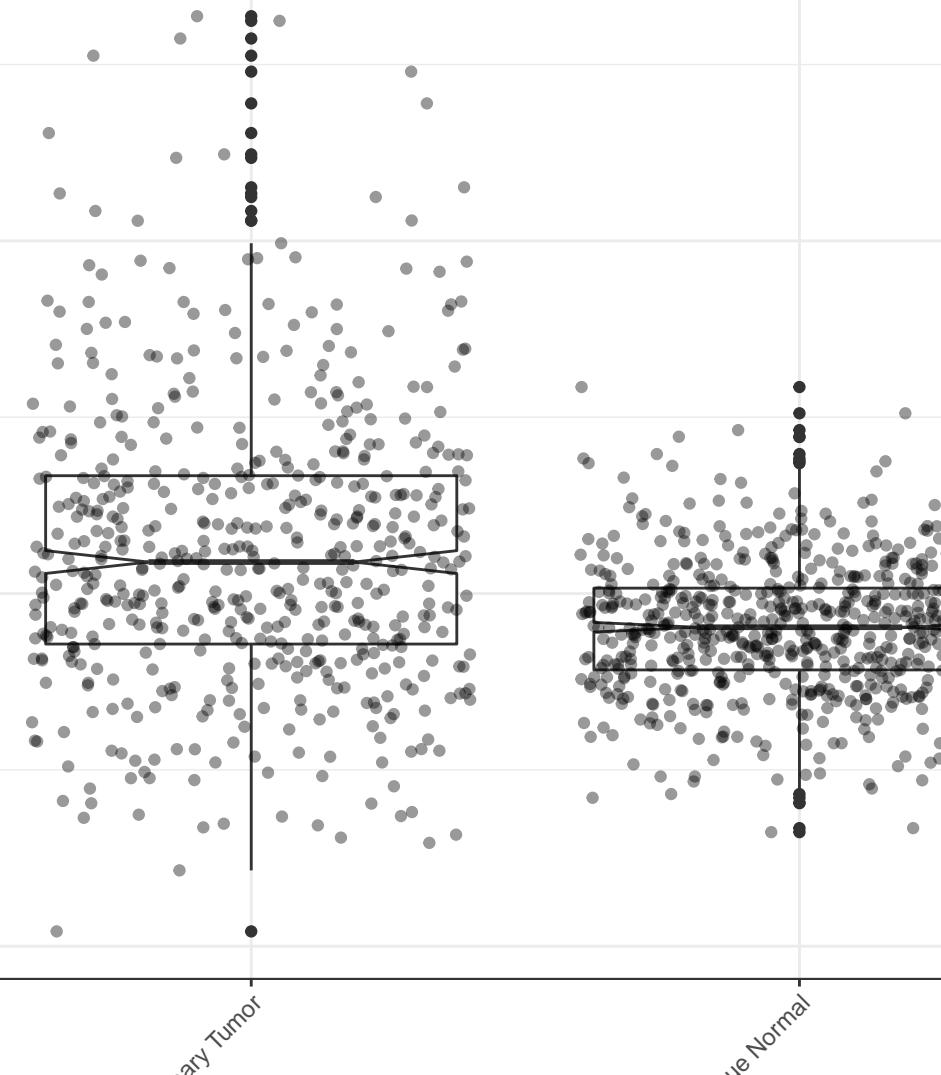
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 214

Value

0.15

0.10

0.05

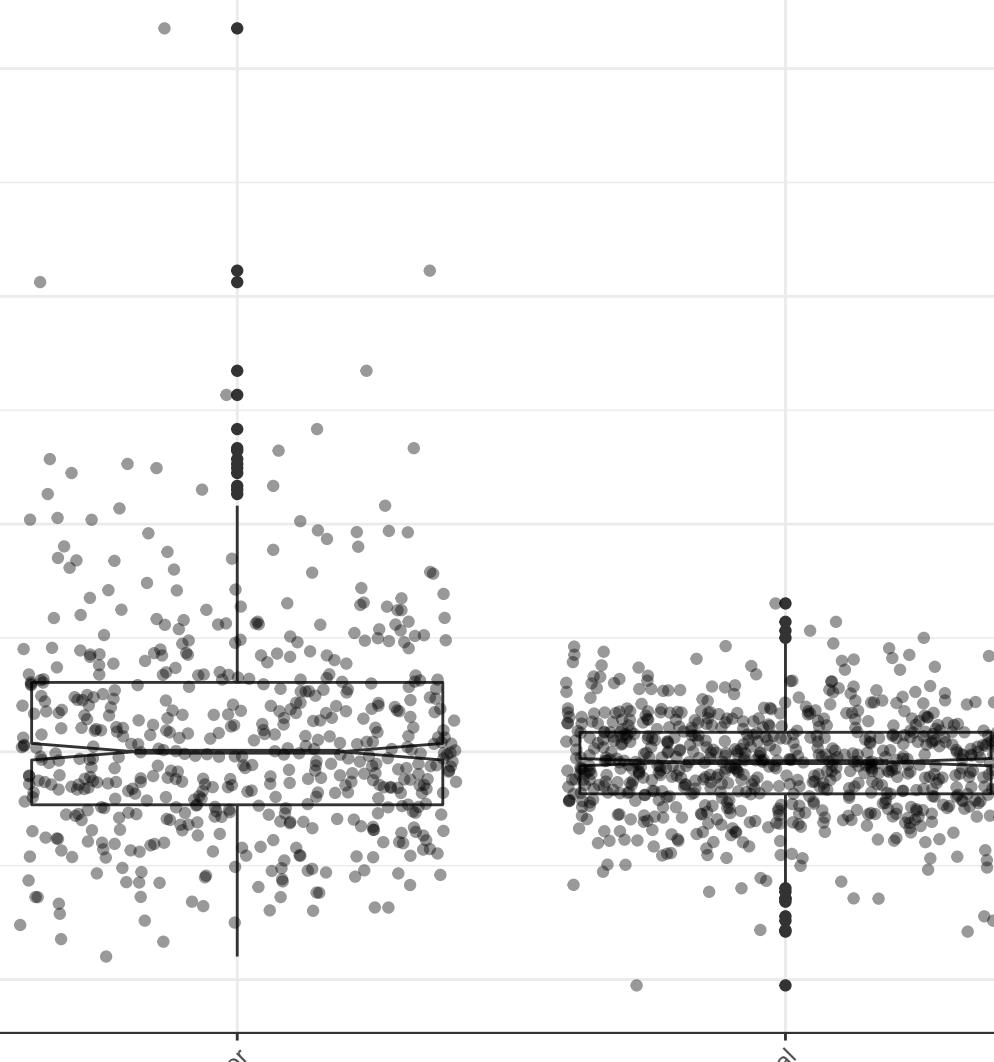
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.05

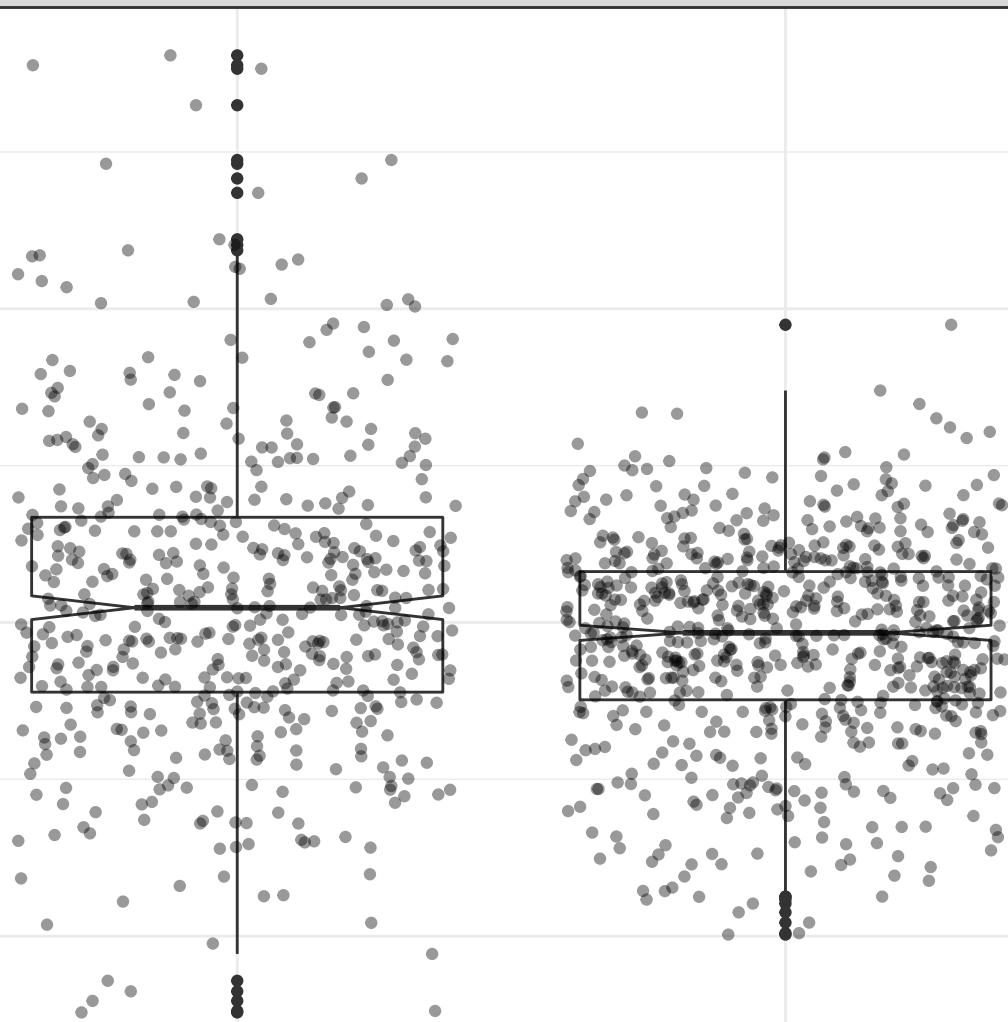
0.00

-0.05

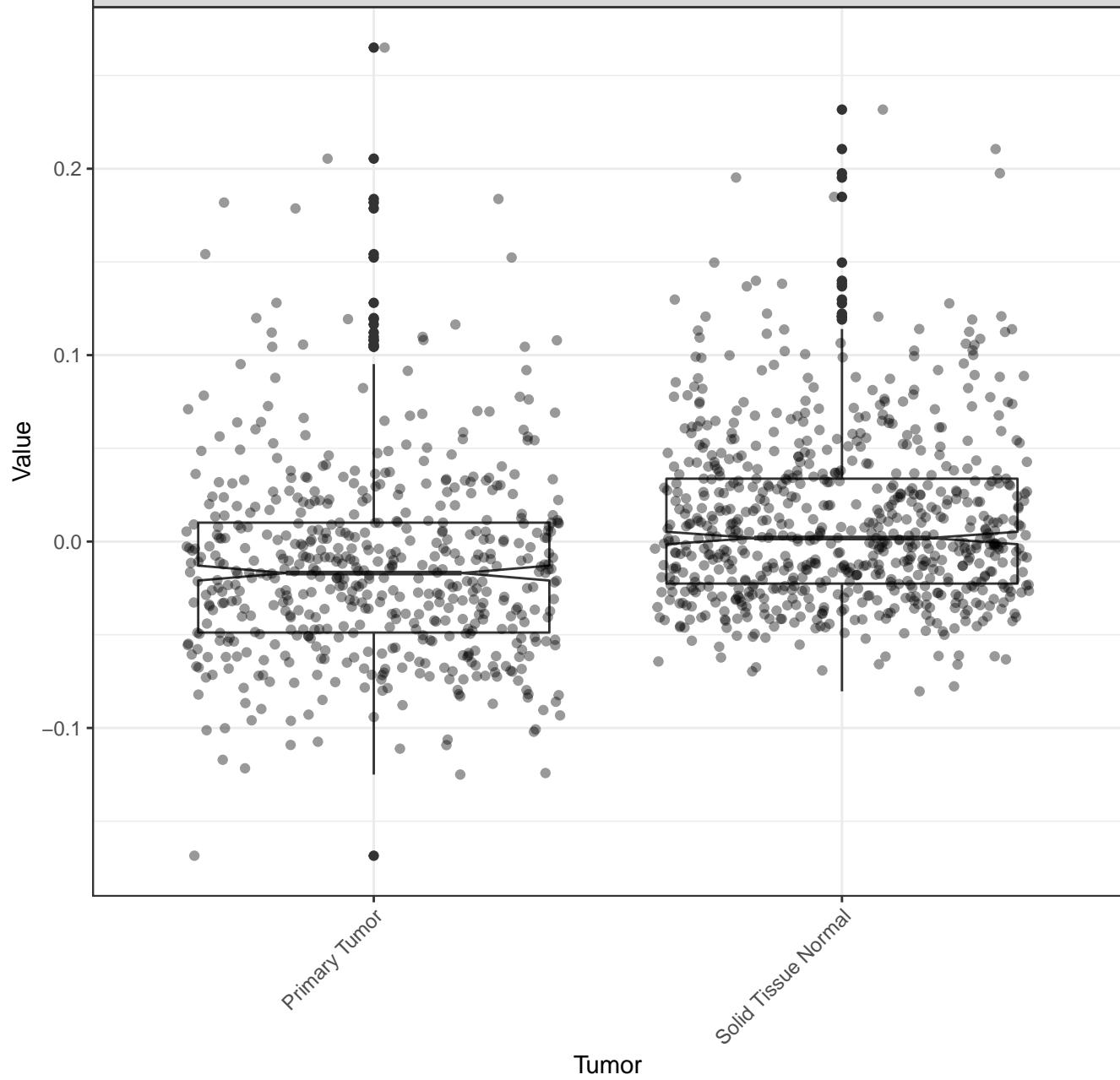
Primary Tumor

Solid Tissue Normal

Tumor



216,KEGG_PATHOGENIC_ESCHERICHIA_COLI_INFECT



LV 217

Value

0.10

0.05

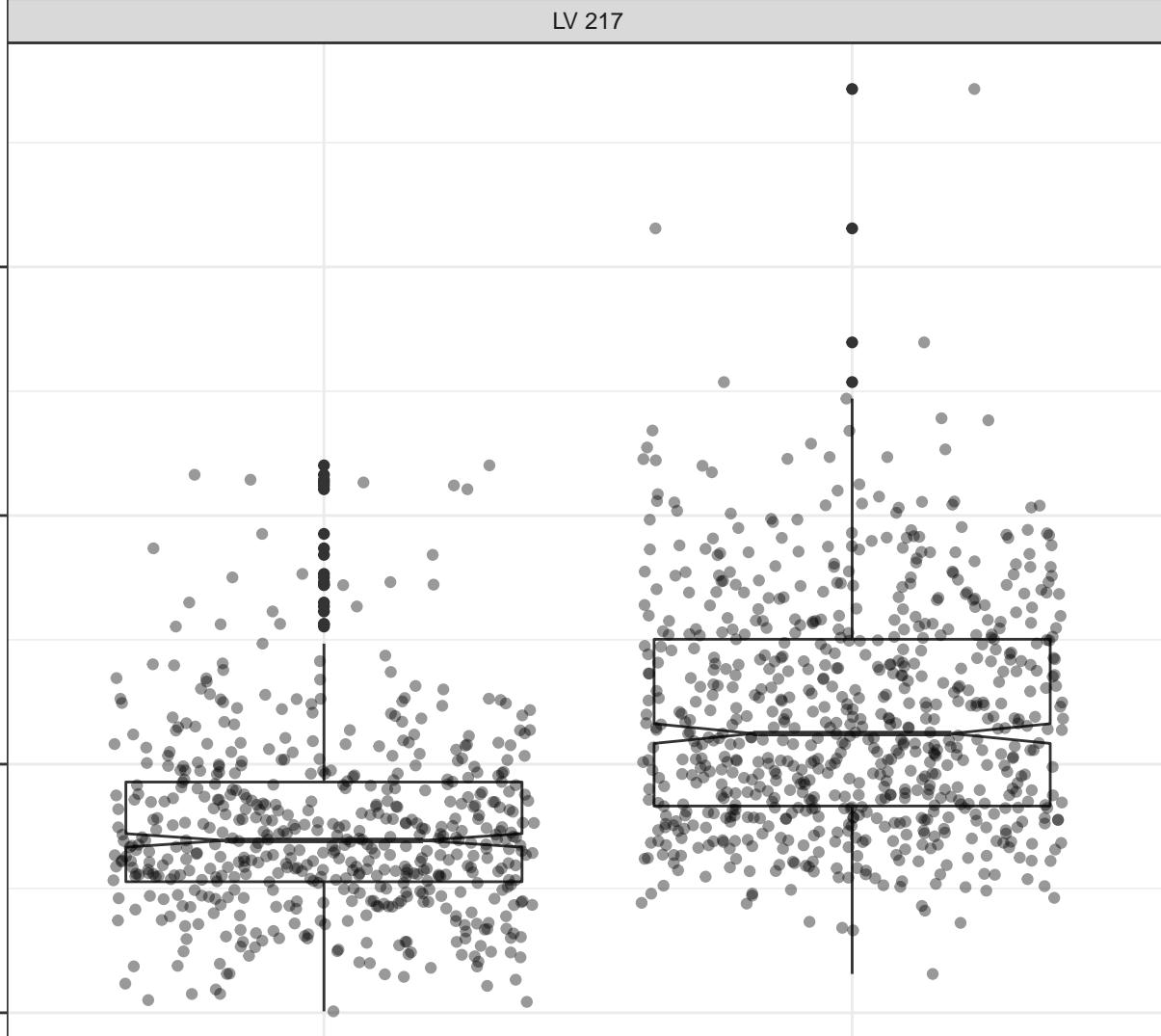
0.00

-0.05

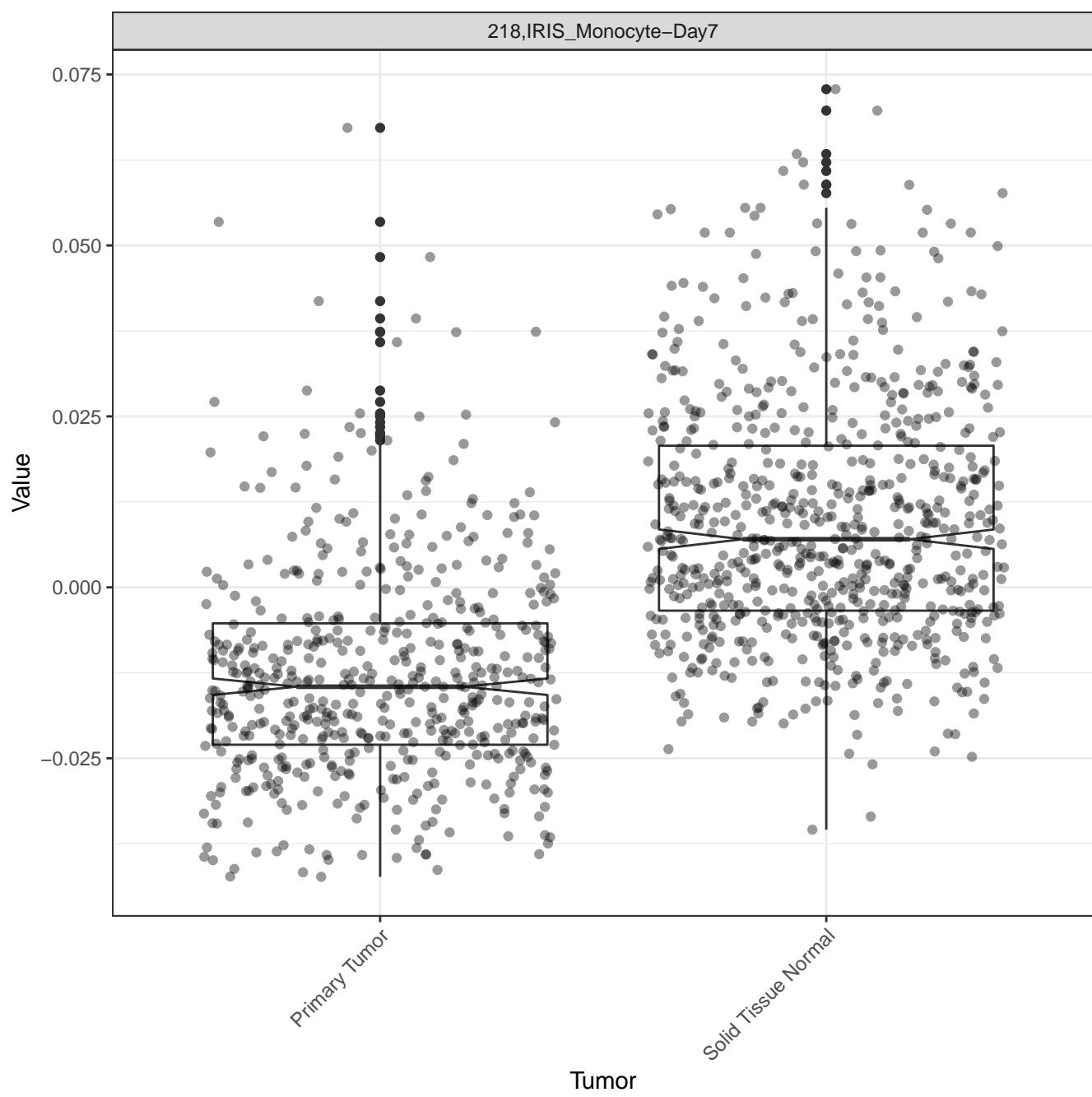
Primary Tumor

Solid Tissue Normal

Tumor



218,IRIS_Monocyte–Day7



219,REACTOME_SLC_MEDIATED_TRANSMEMBRANE_TRANSPORT

Value

0.05

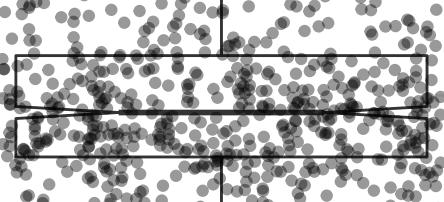
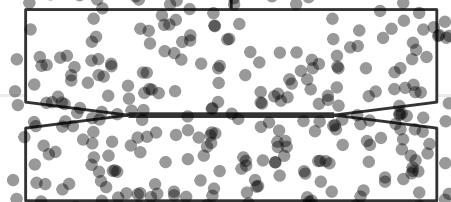
0.00

-0.05

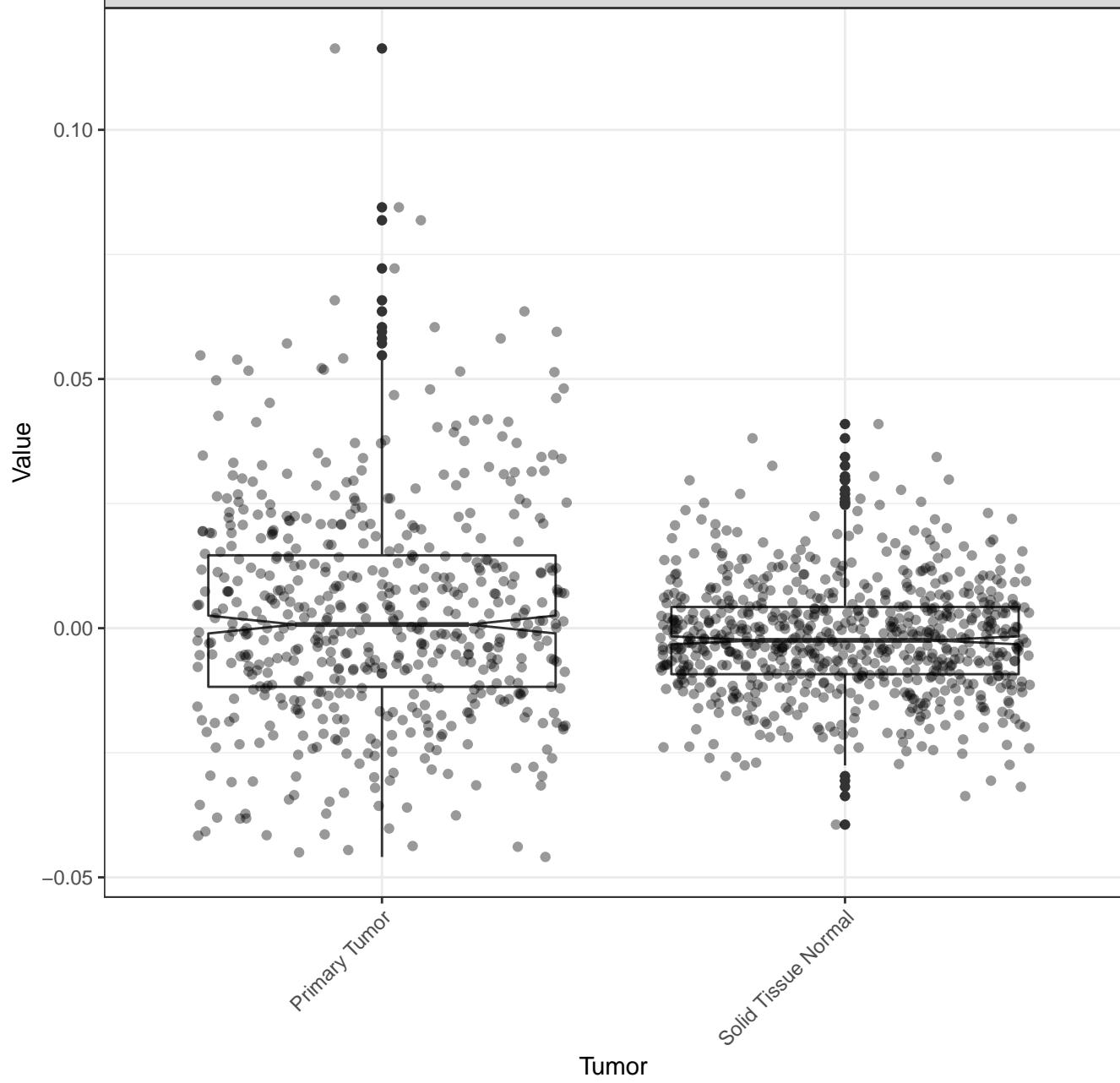
Primary Tumor

Tumor

Solid Tissue Normal

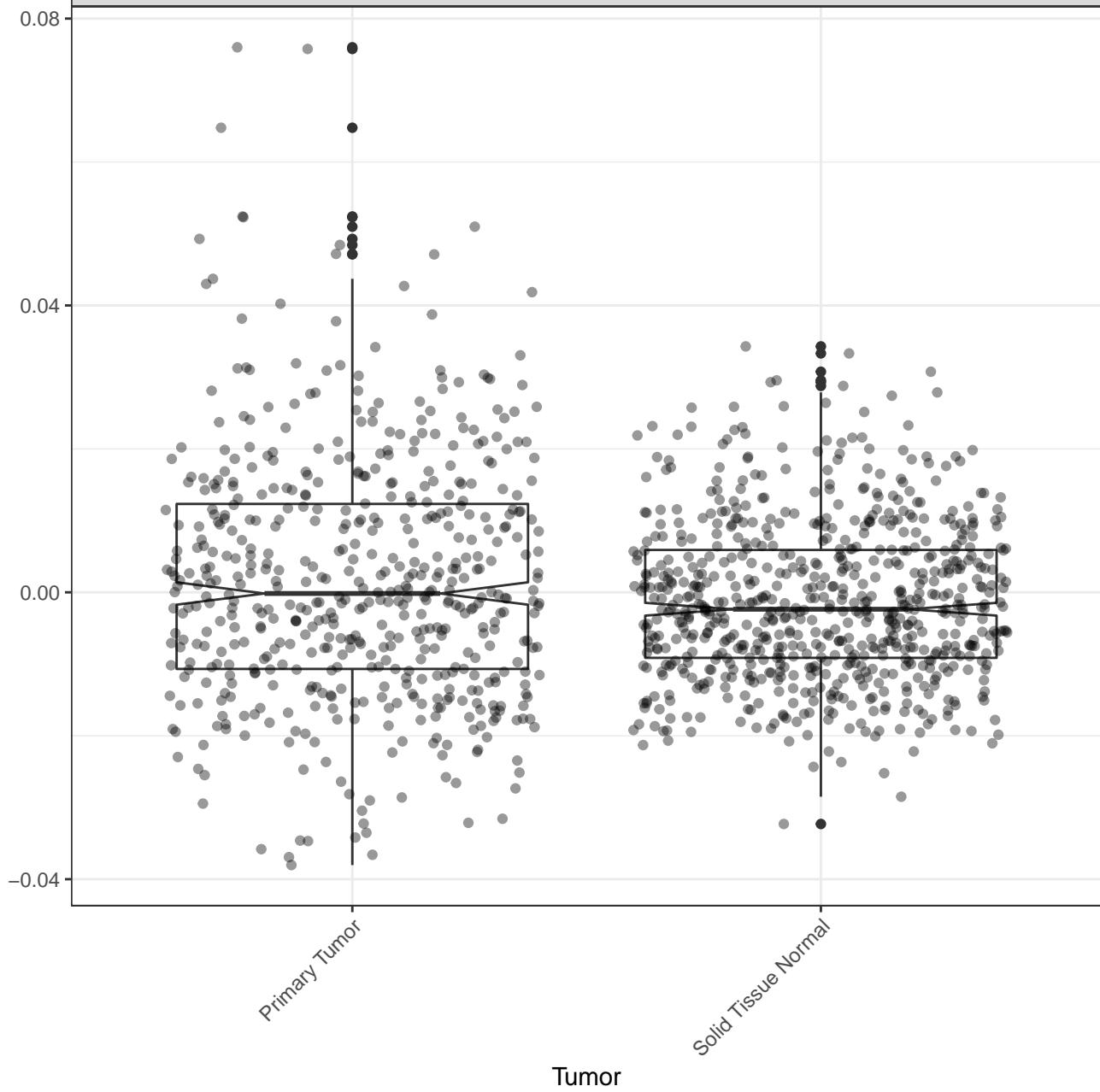


220,REACTOME_RIG_I_MDA5_MEDIATED_INDUCTION_OF_IFN_ALPHA_BETA_PATHWAYS



221,DMAP_ERY4

Value



222,REACTOME_METABOLISM_OF_LIPIDS_AND_LIPOPROTEINS

Value

0.10

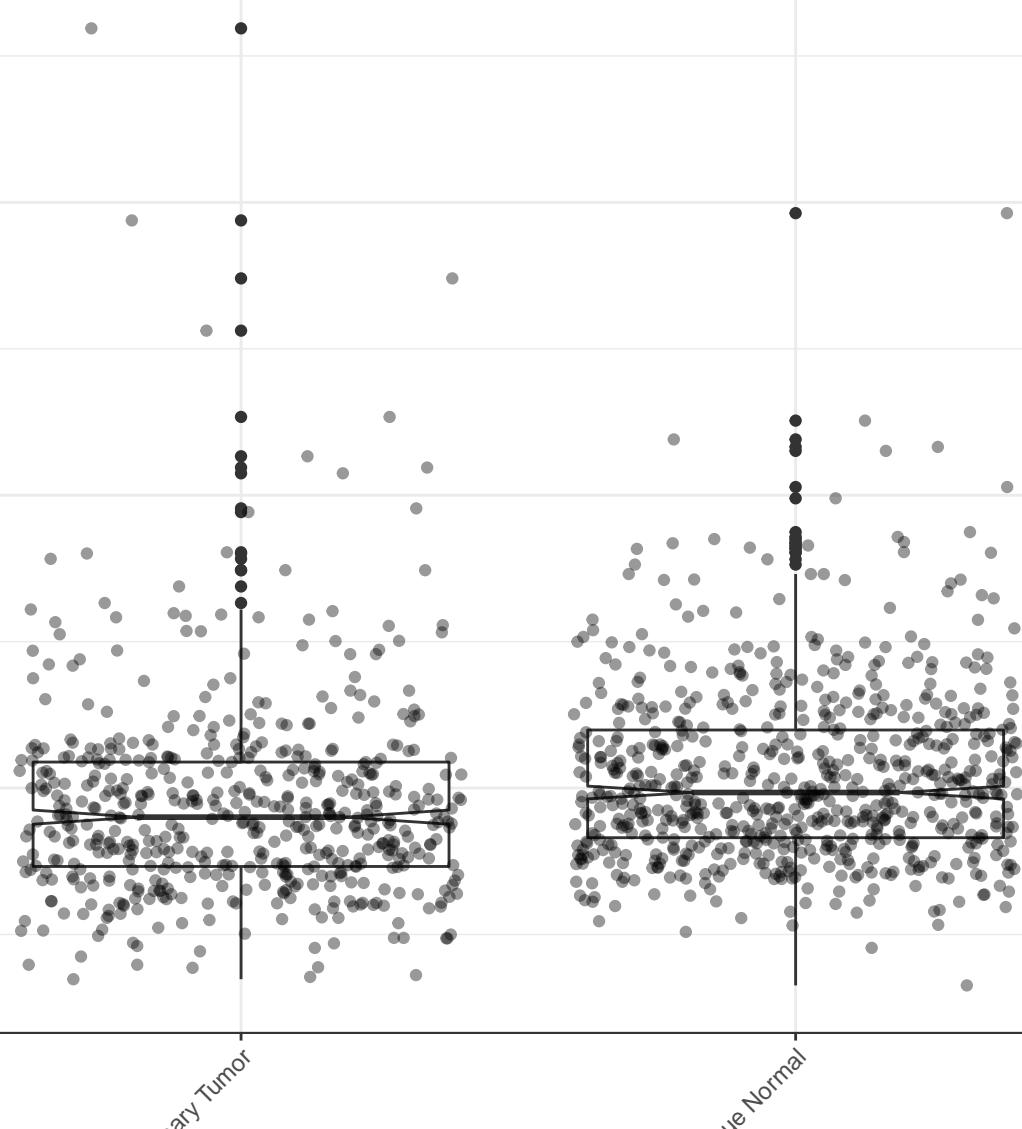
0.05

0.00

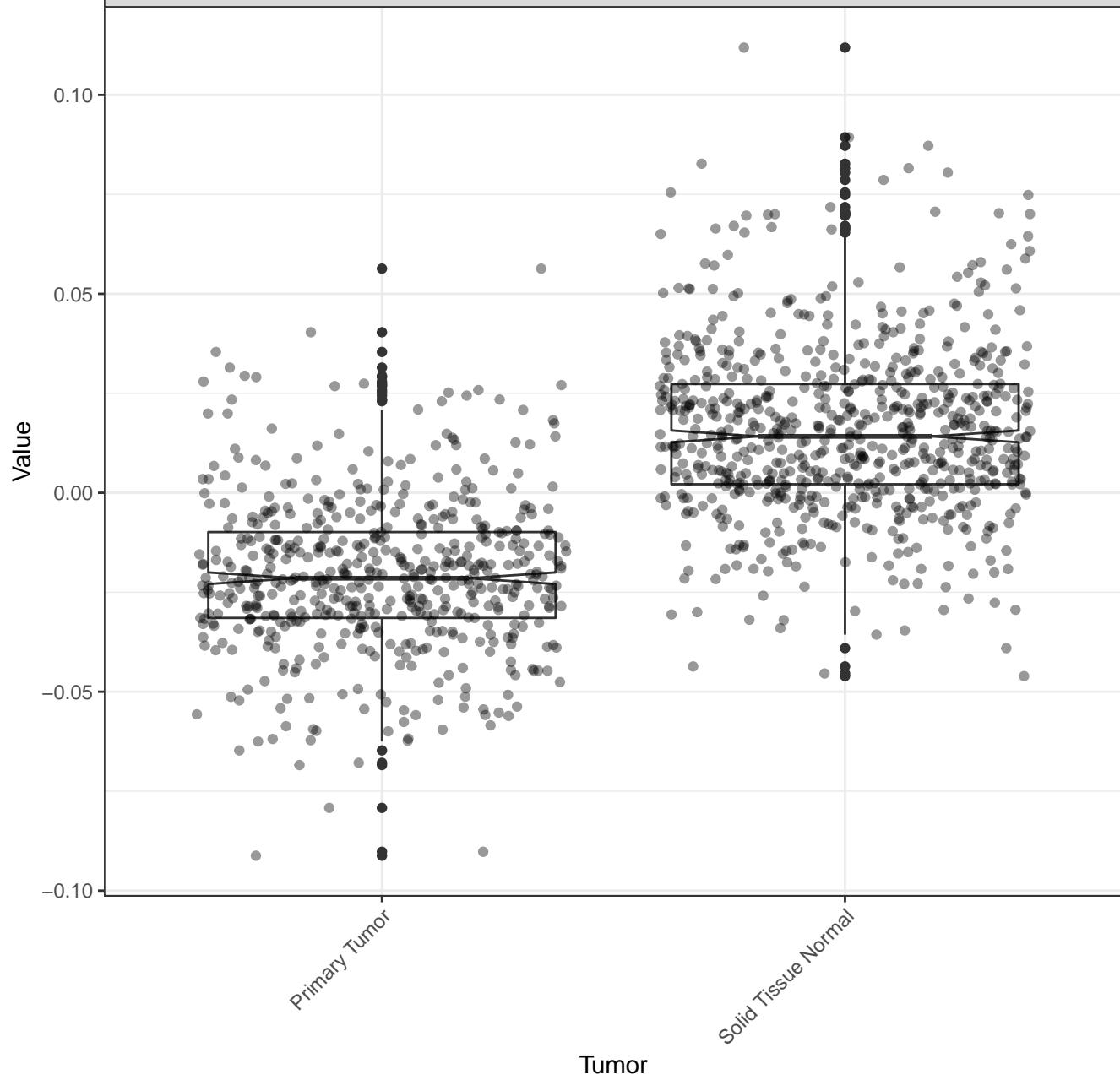
Primary Tumor

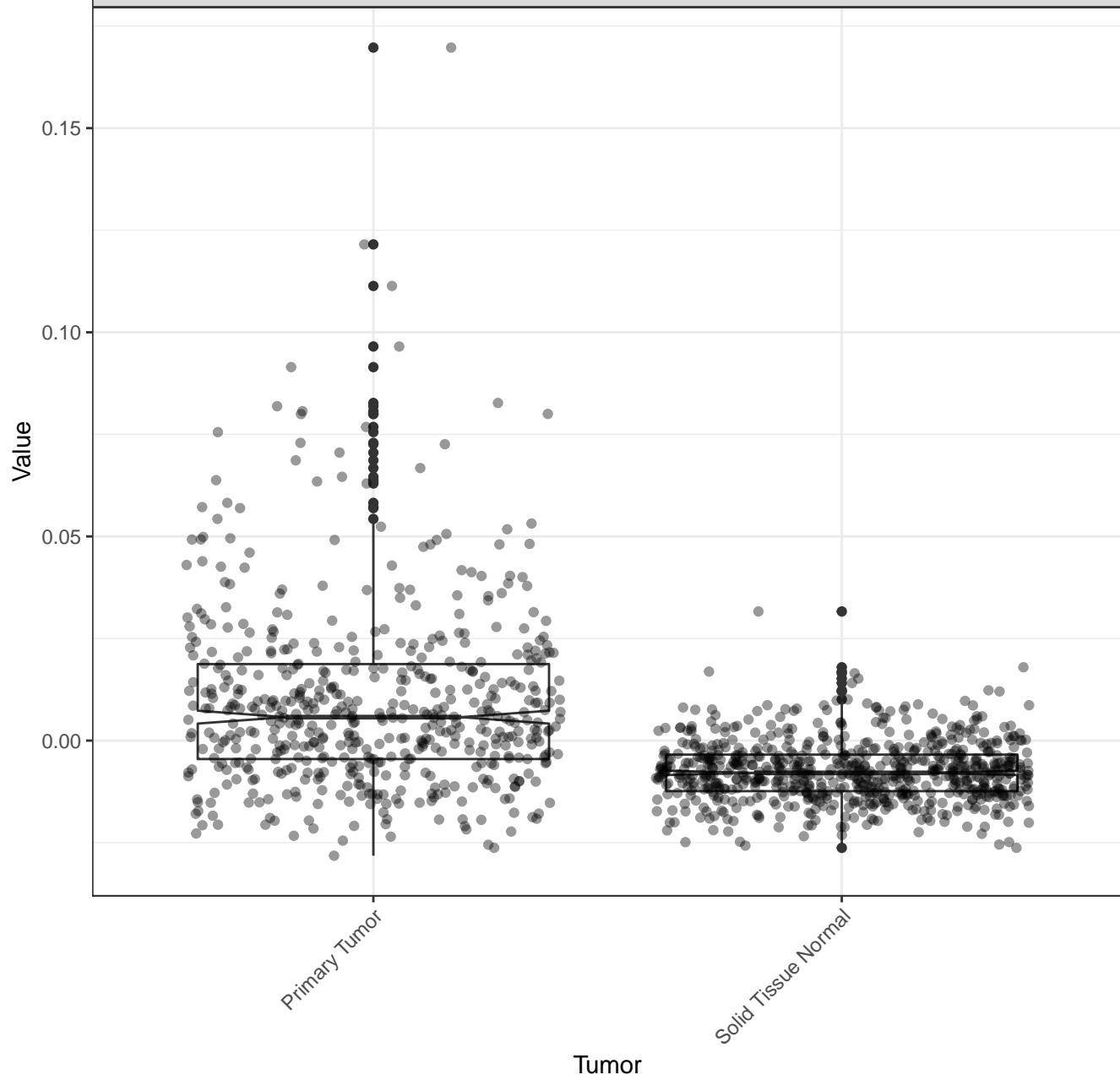
Solid Tissue Normal

Tumor



223,PID_P75NTRPATHWAY





225,KEGG_CYSTEINE_AND_METHIONINE_METABOLISM

Value

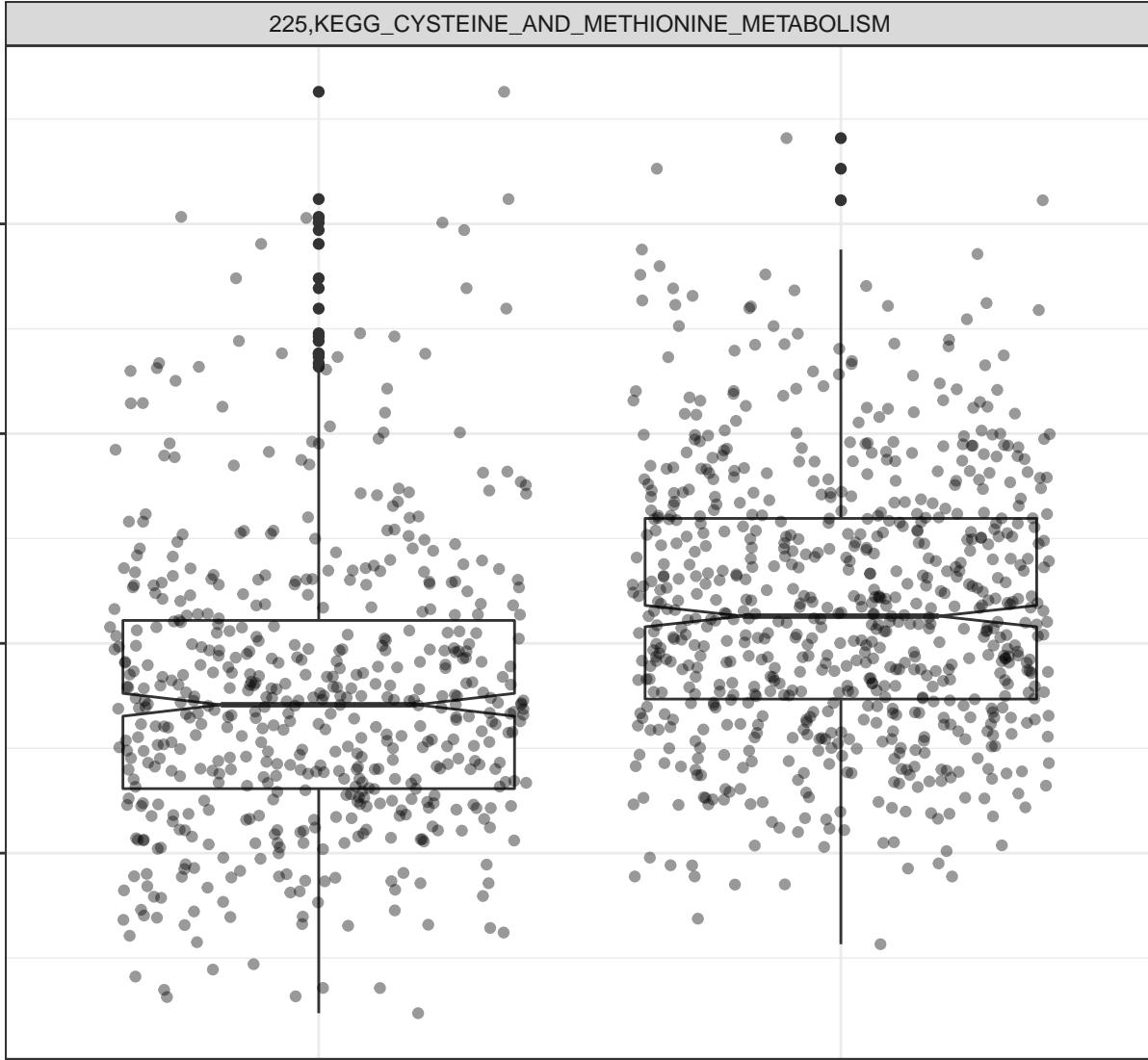
0.050
0.025
0.000

-0.025

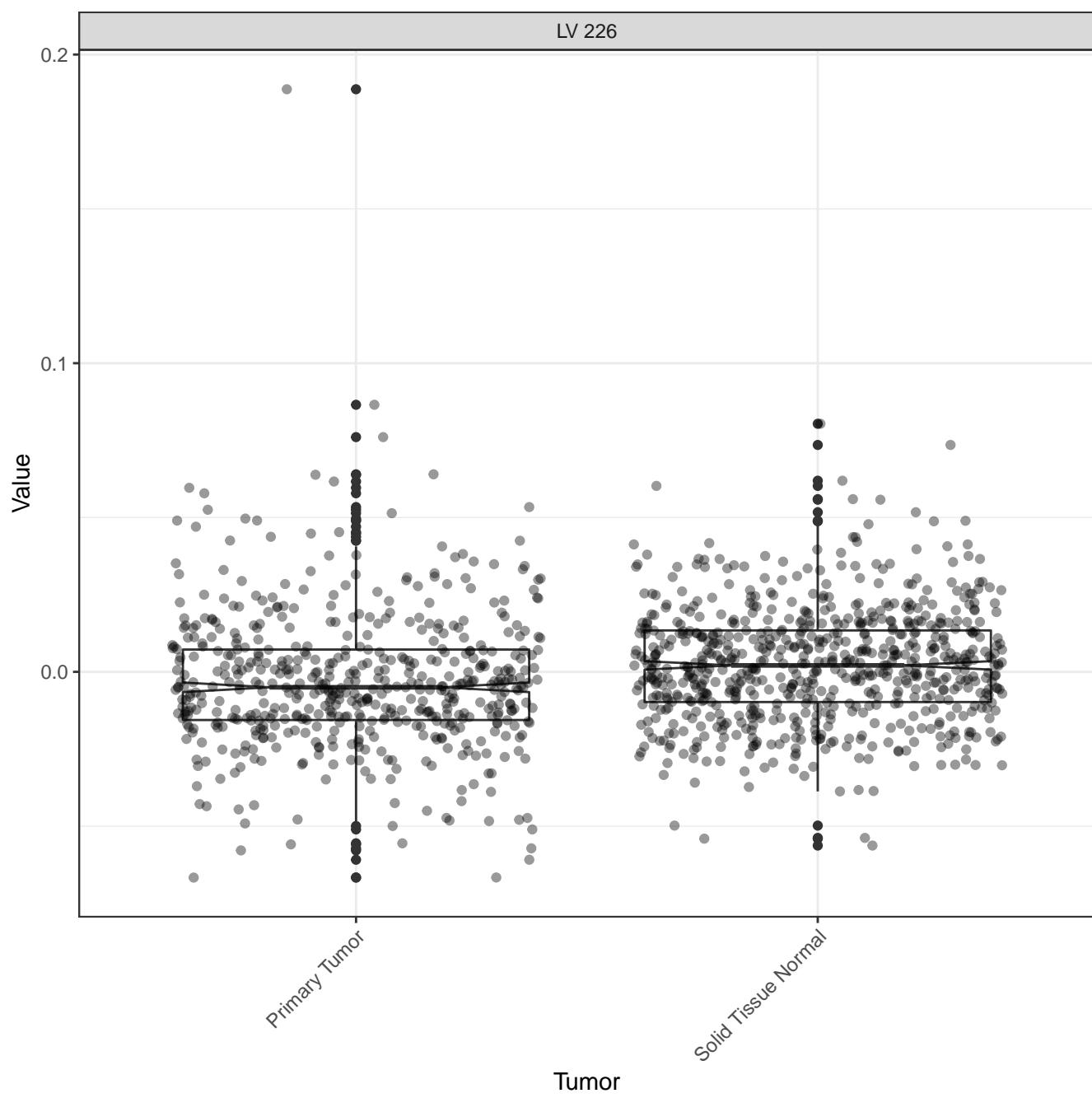
Primary Tumor

Solid Tissue Normal

Tumor



LV 226



227,REACTOME_INTERACTIONS_OF_VPR_WITH_HOST_CELLULAR_PROTEINS

Value

0.15

0.10

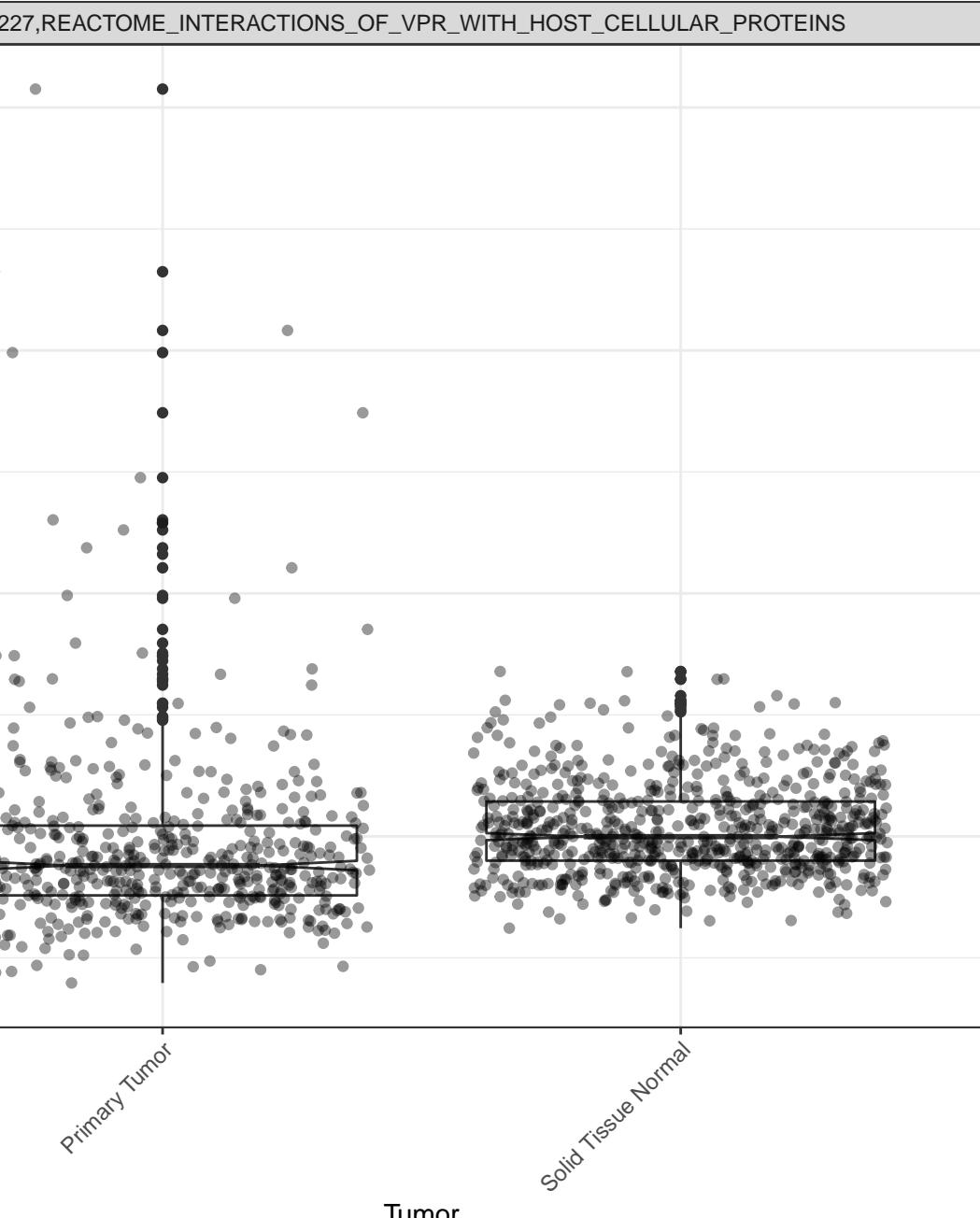
0.05

0.00

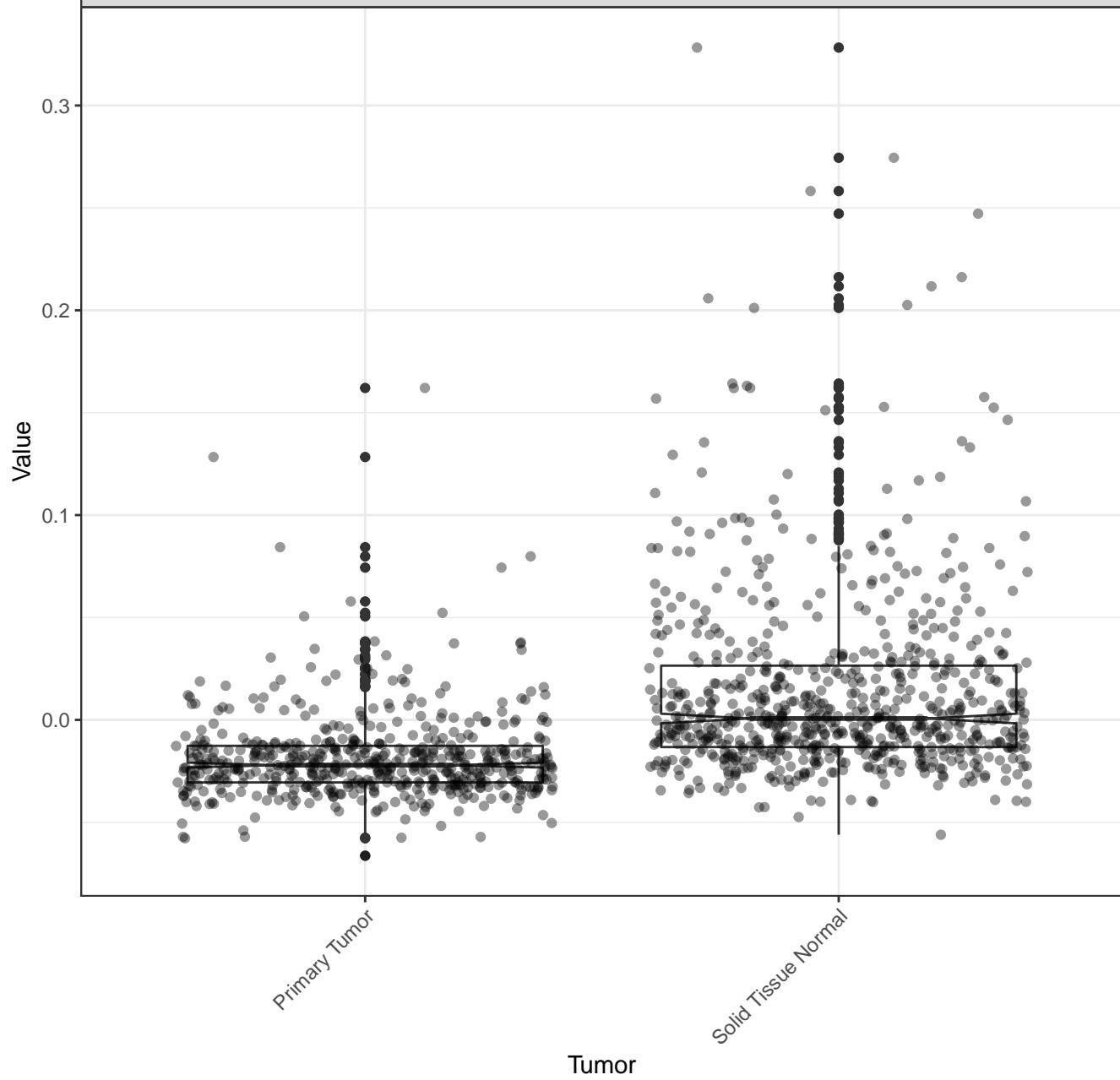
Primary Tumor

Solid Tissue Normal

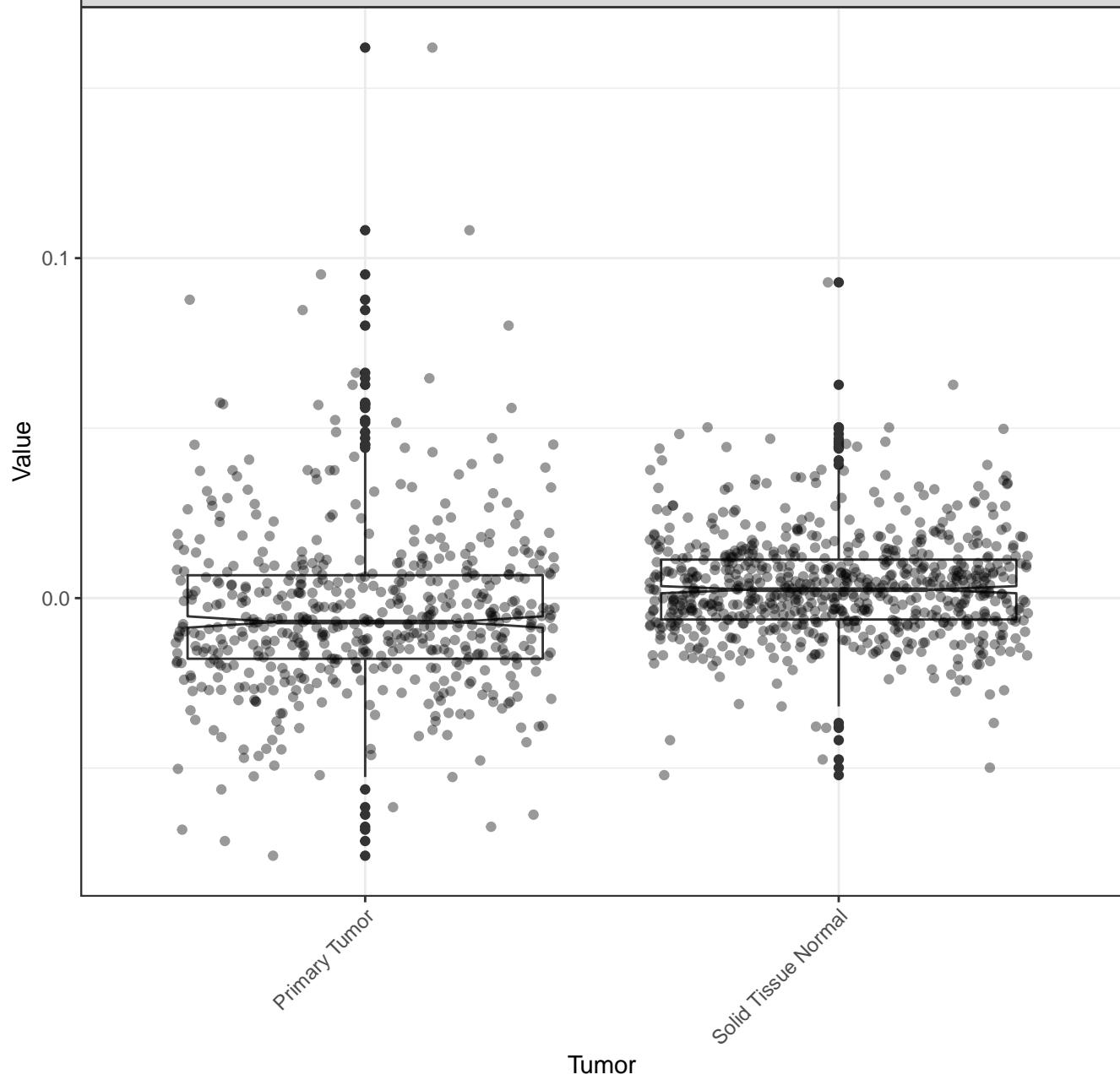
Tumor



228,DMAP_NKA3



229,PID_ATF2_PATHWAY



230,REACTOME_CELL_CYCLE

Value

0.05

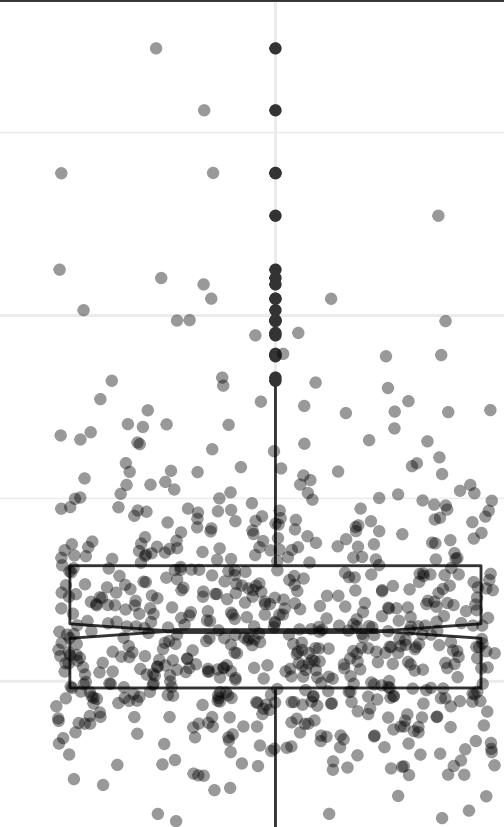
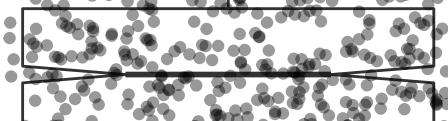
0.00

-0.05

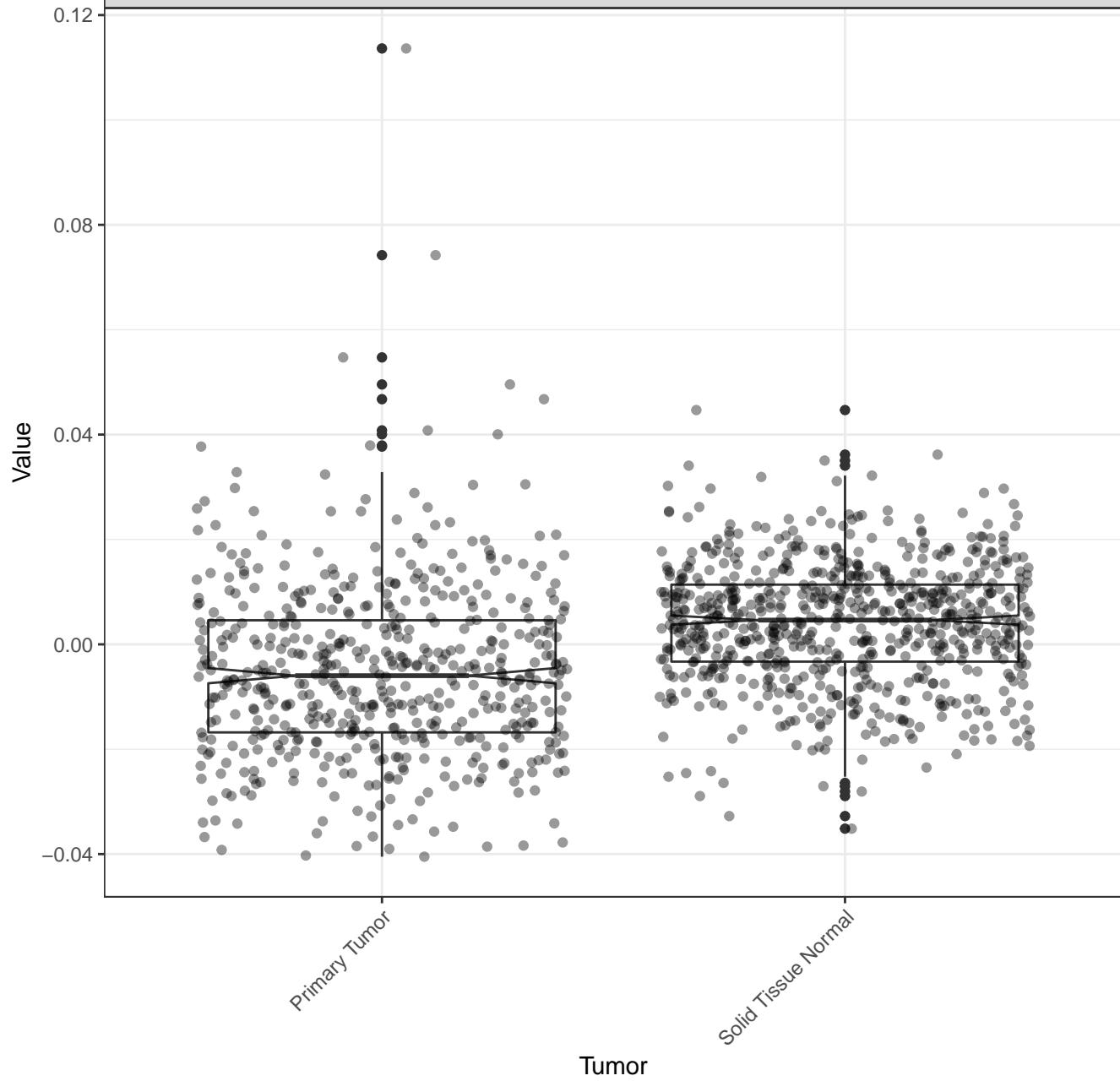
Primary Tumor

Solid Tissue Normal

Tumor

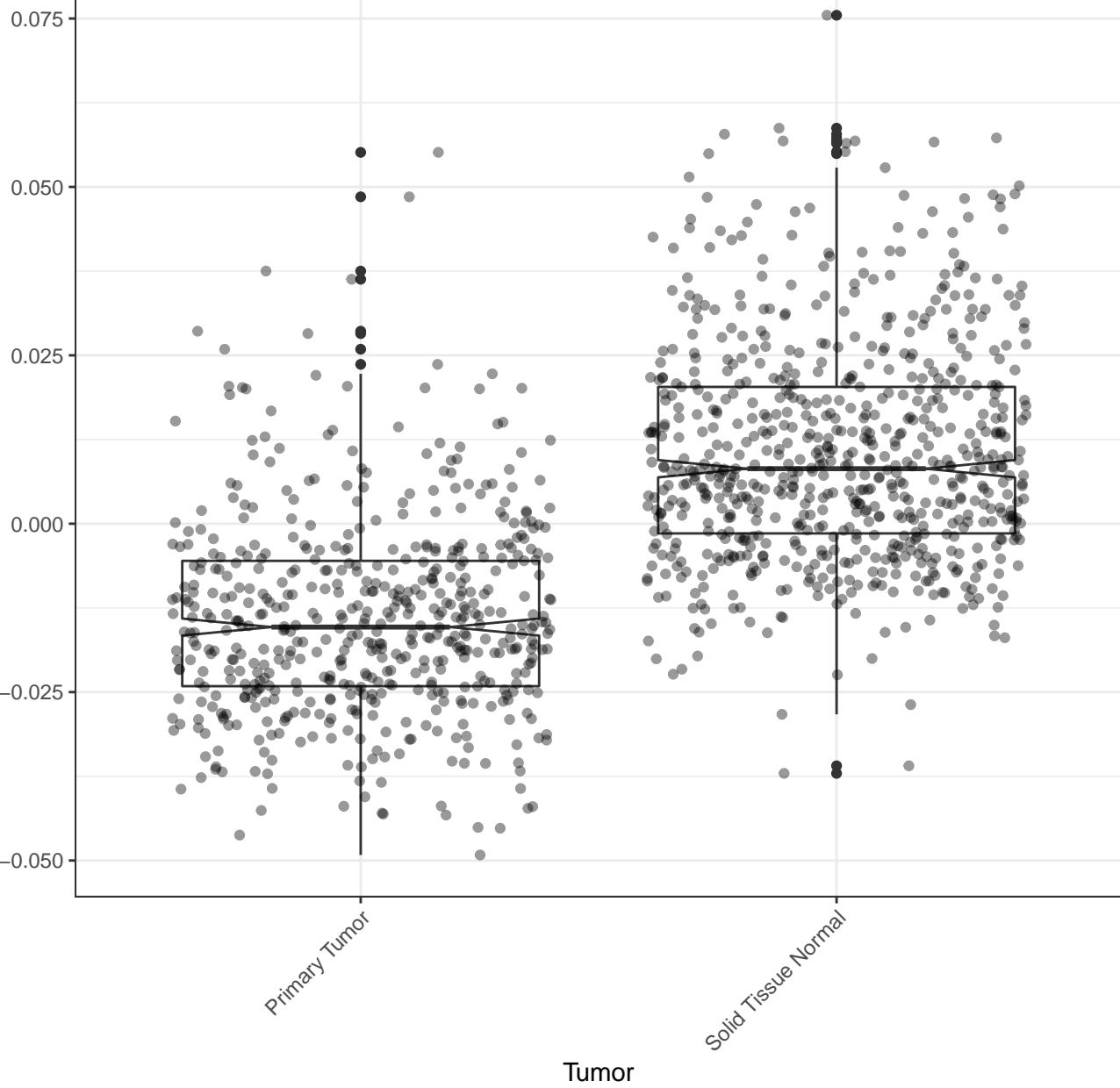


231,REACTOME_CELL_CYCLE

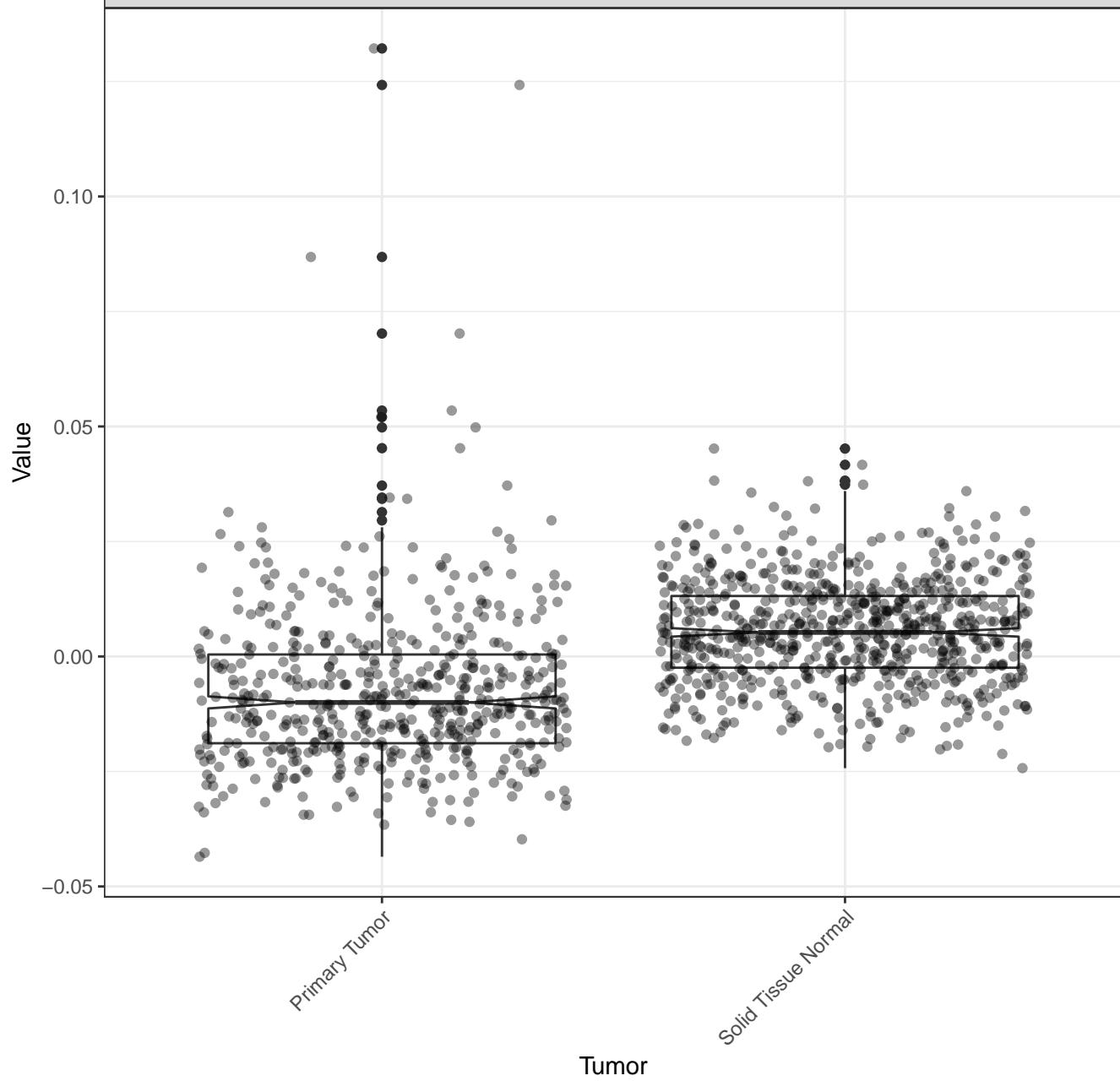


232,REACTOME_LOSS_OF_NLP_FROM_MITOTIC_CENTROSOMES

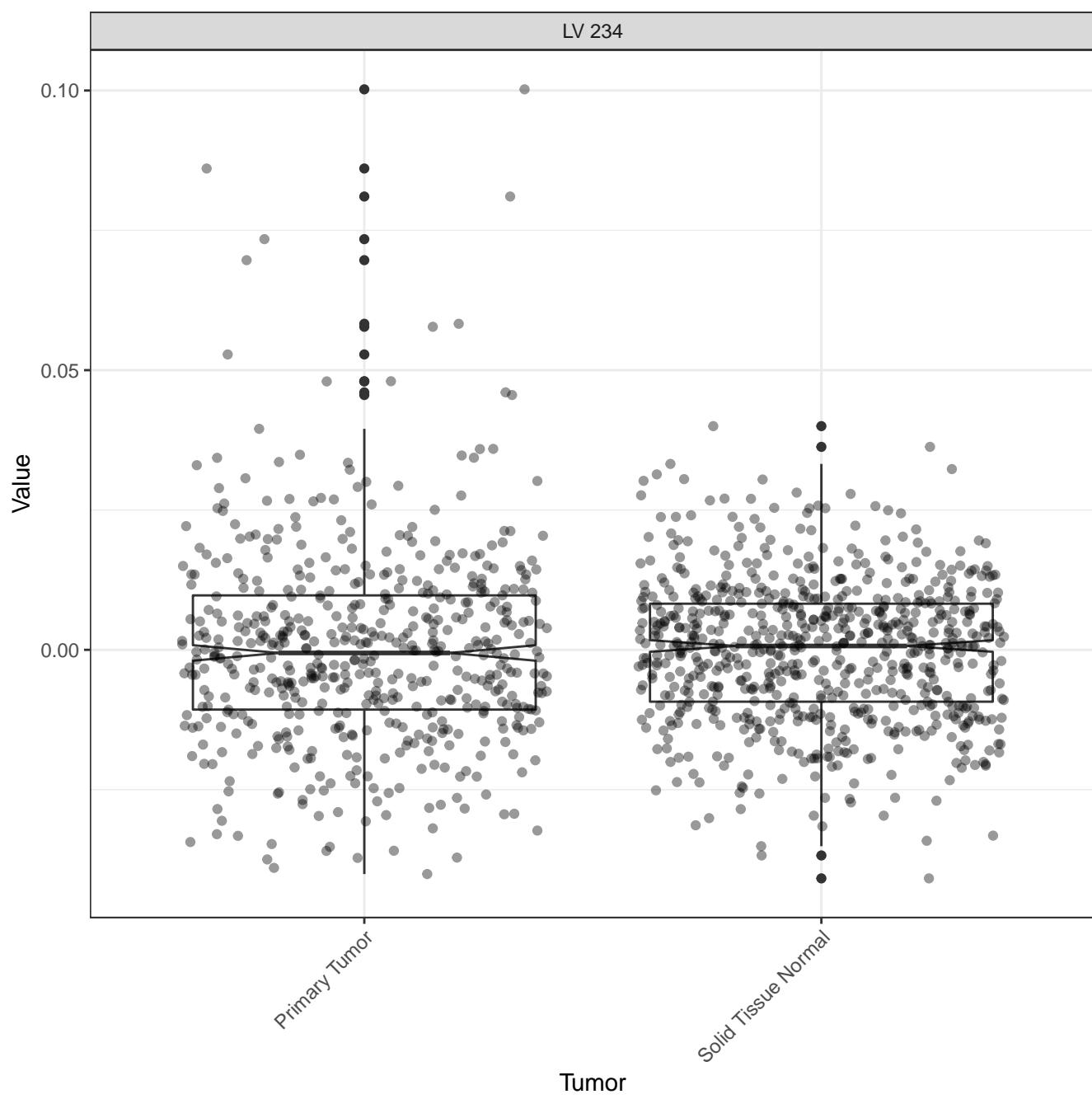
Value



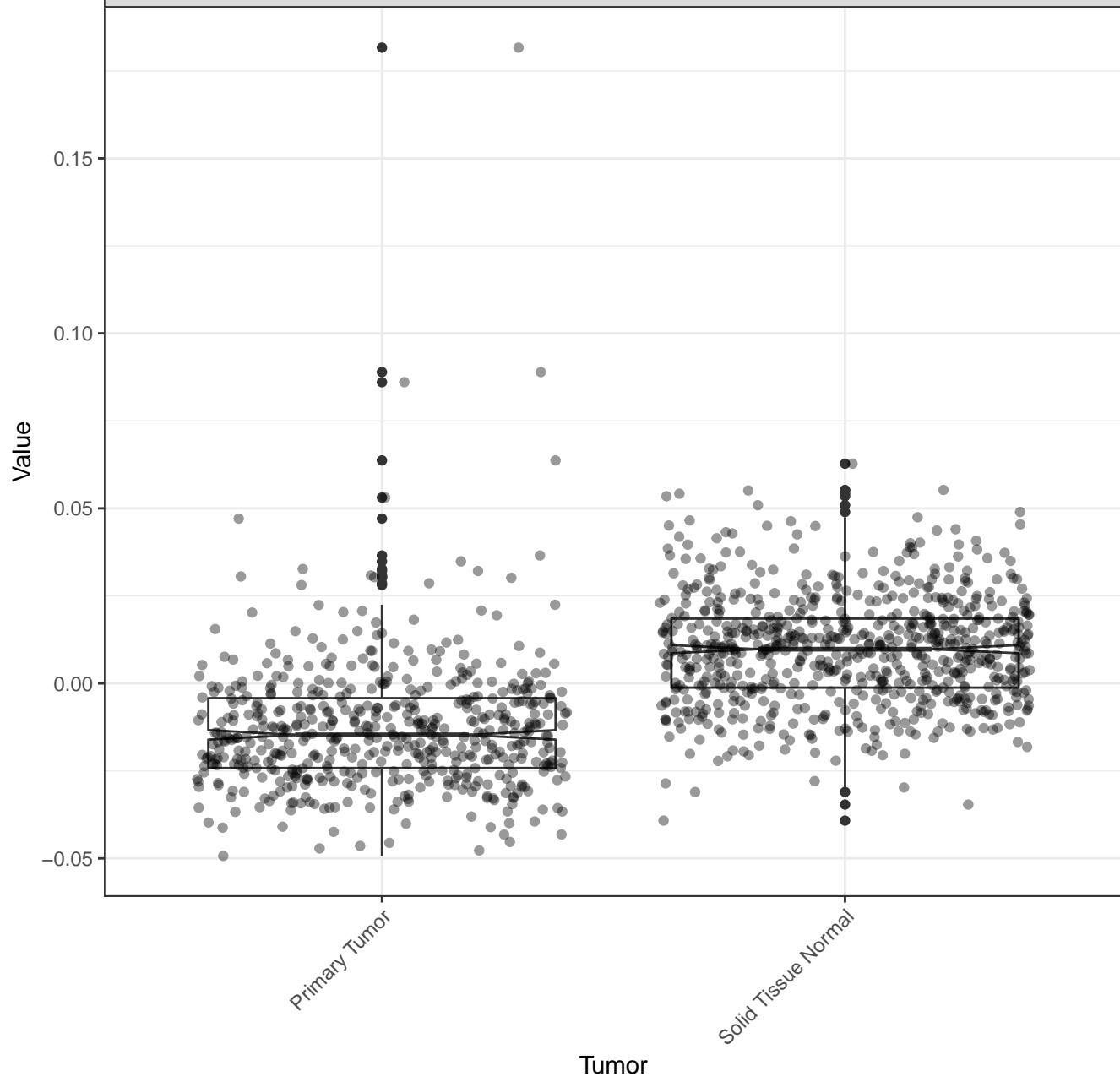
233,DMAP_ERY5



LV 234



235,IRIS_PlasmaCell–FromBoneMarrow



236,REACTOME_MEIOSIS

Value

0.10

0.05

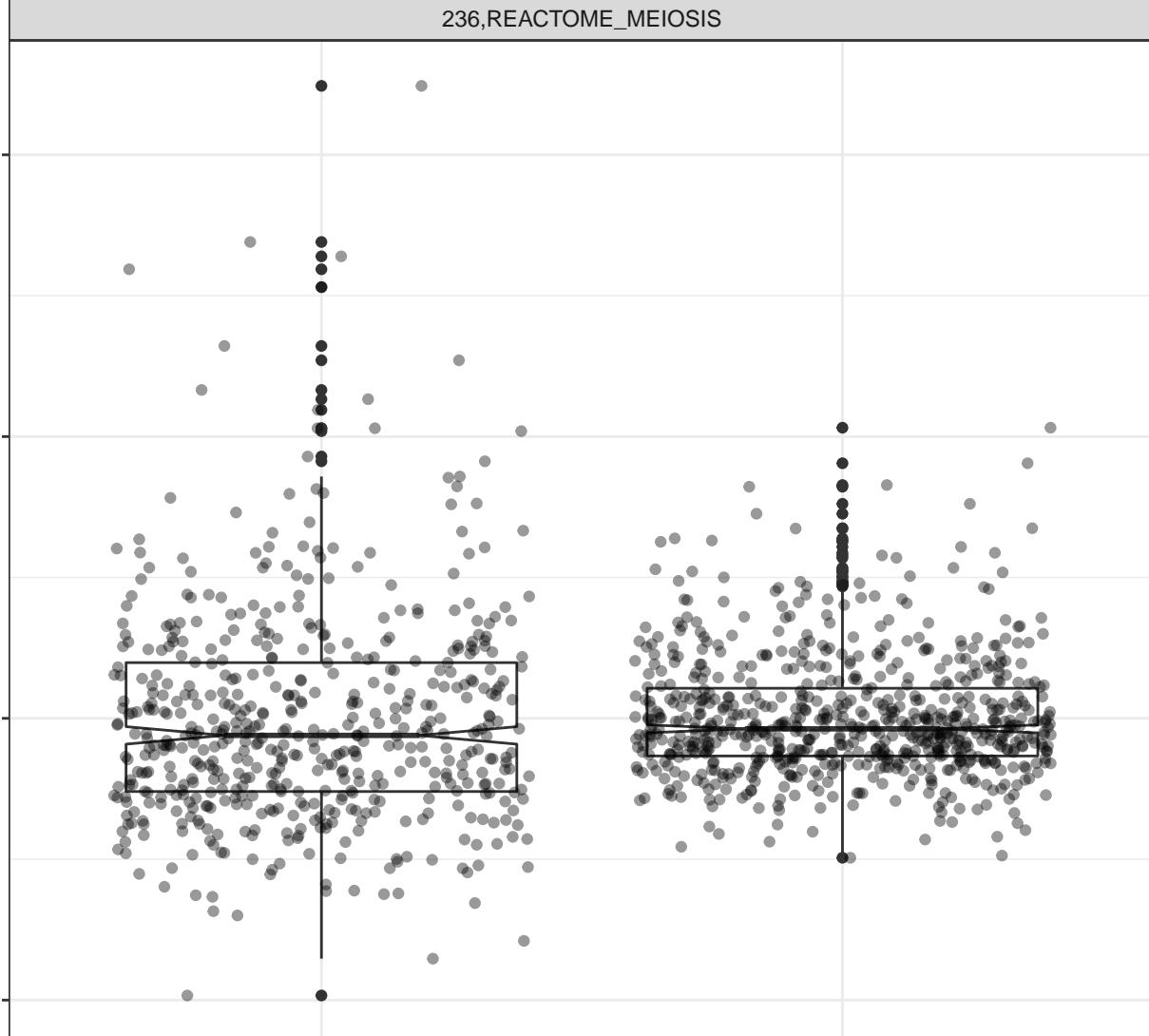
0.00

-0.05

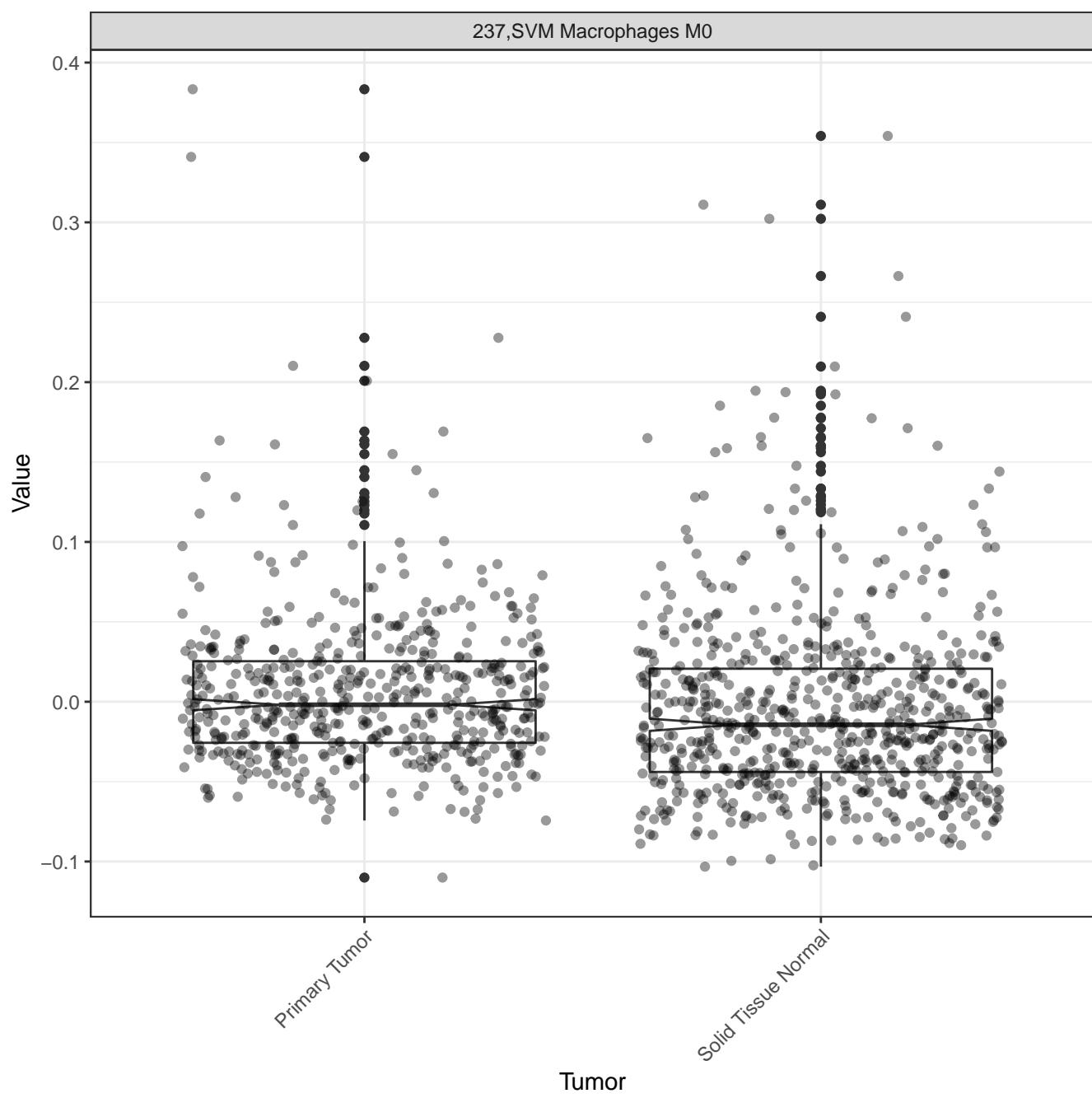
Primary Tumor

Solid Tissue Normal

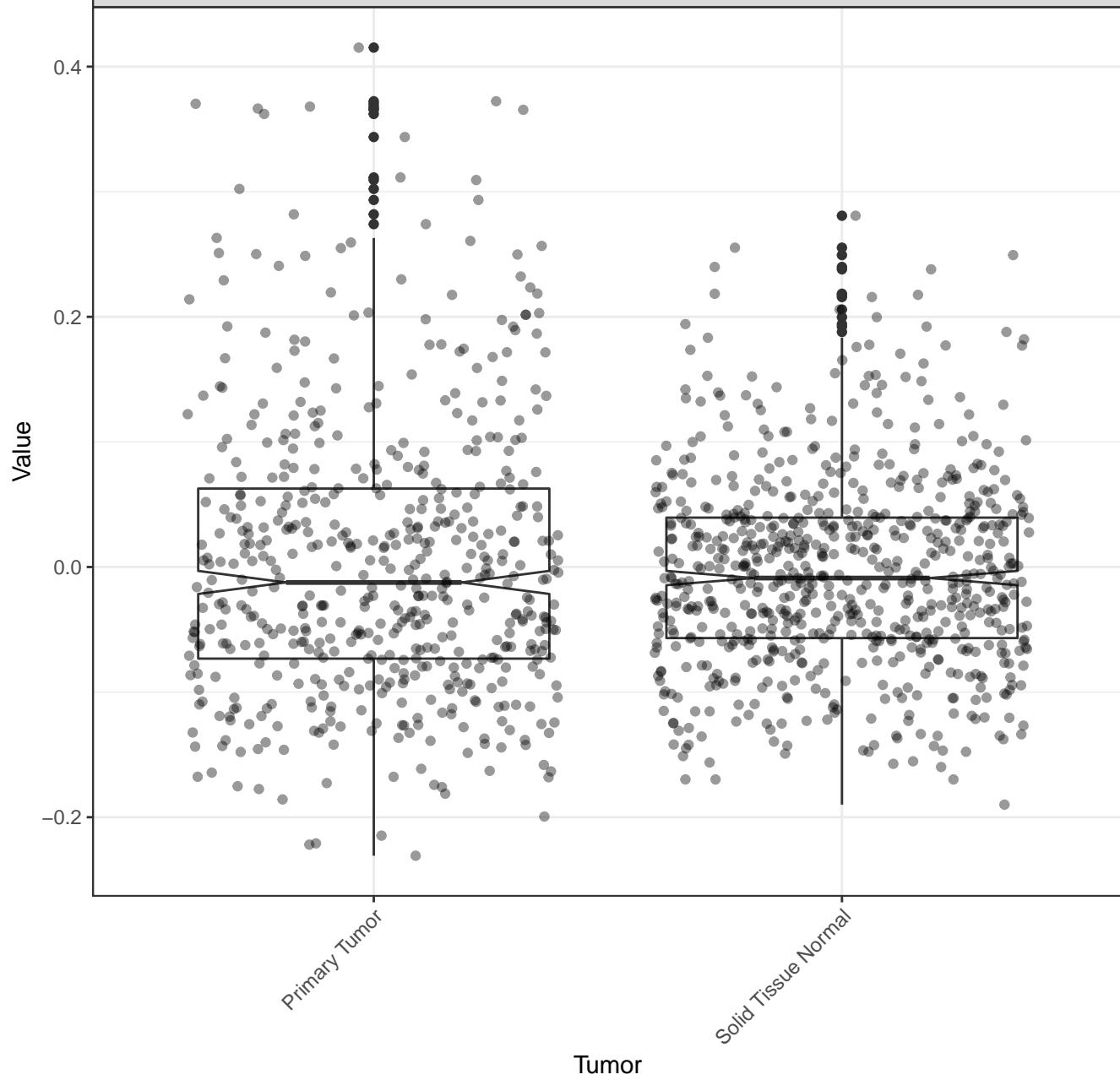
Tumor



237,SVM Macrophages M0



238,PID_FAK_PATHWAY



LV 239

Value

0.10

0.05

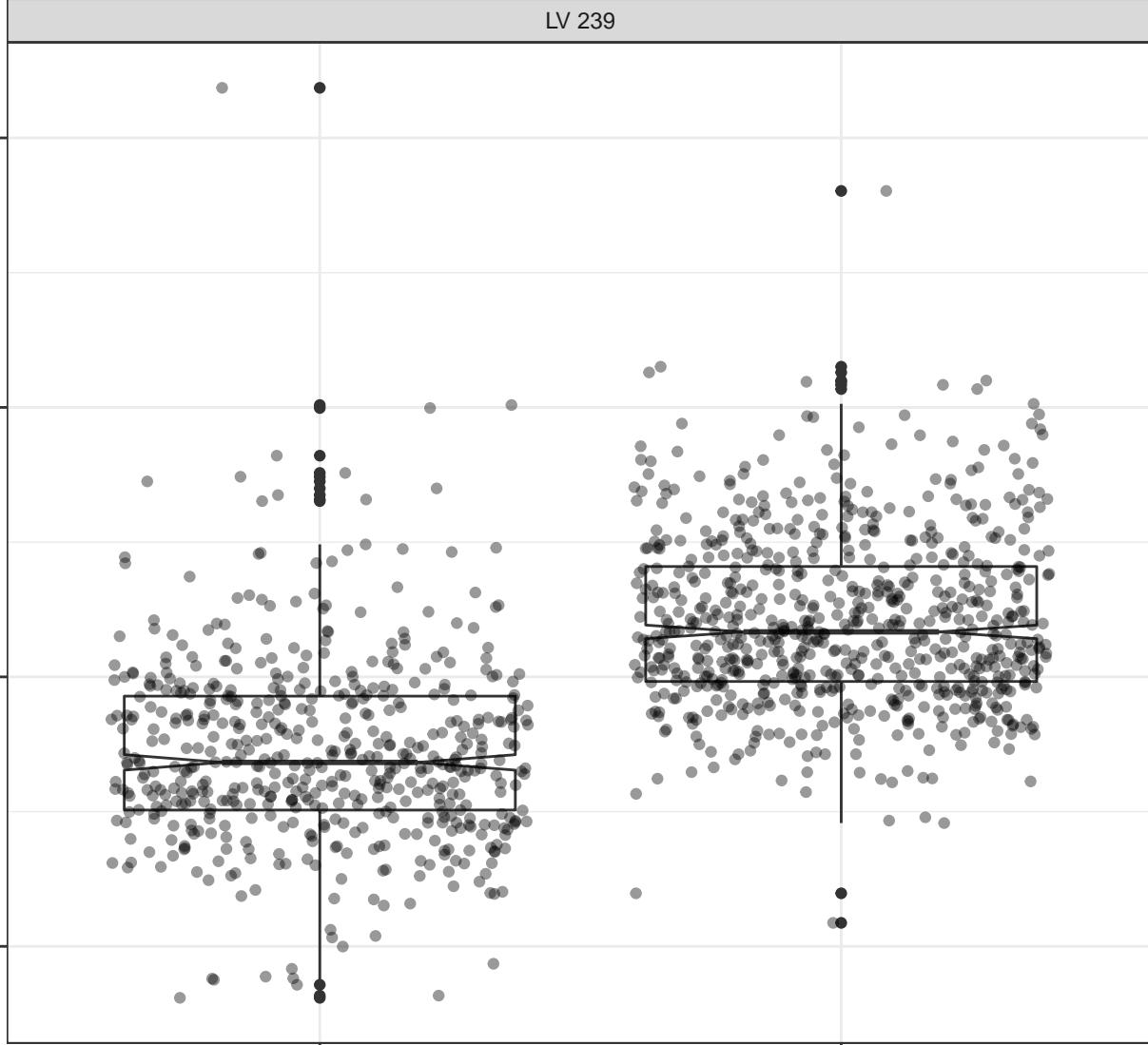
0.00

-0.05

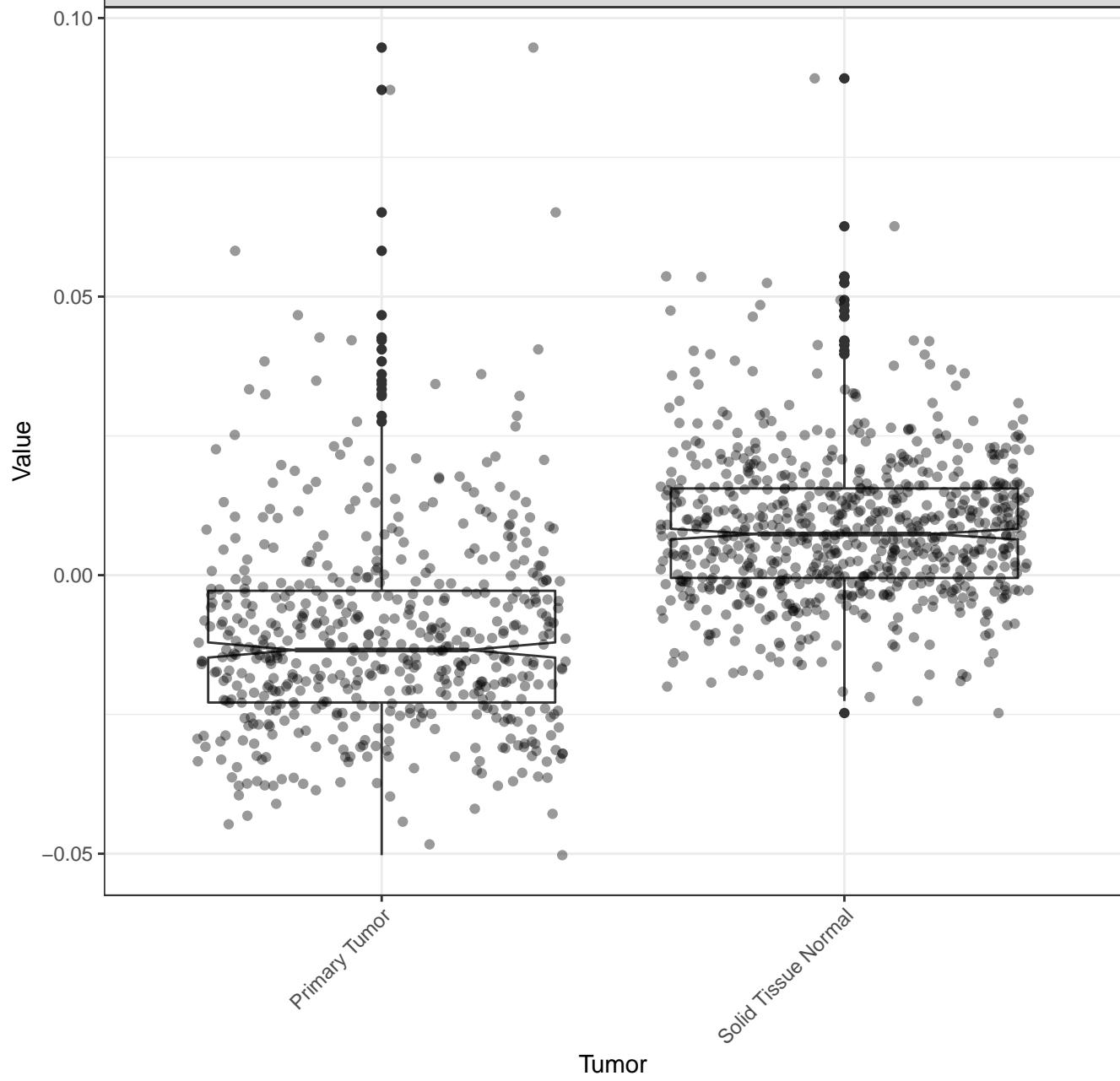
Primary Tumor

Solid Tissue Normal

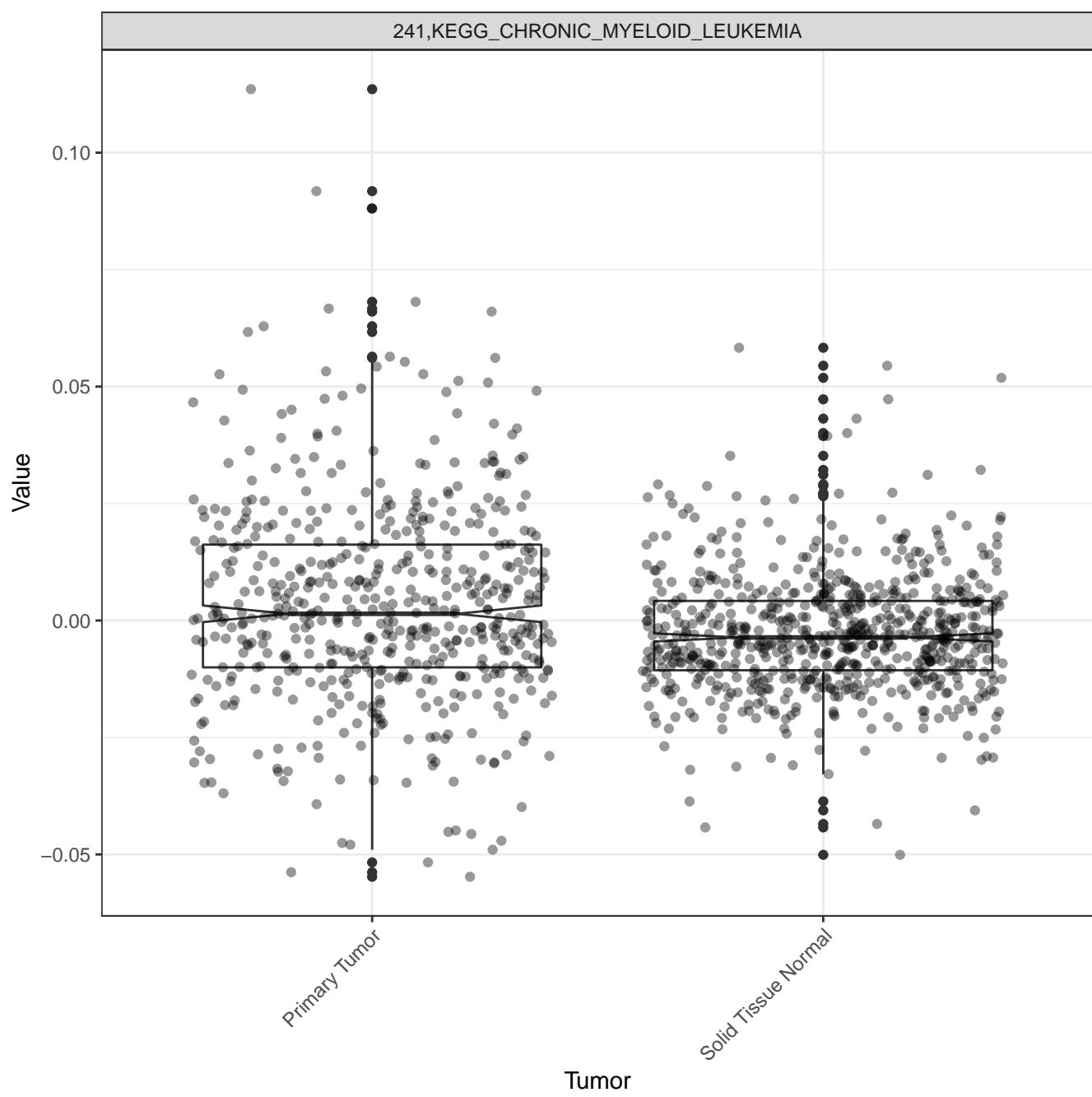
Tumor



240,SVM Dendritic cells activated



241,KEGG_CHRONIC_MYELOID_LEUKEMIA



LV 242

Value

0.05

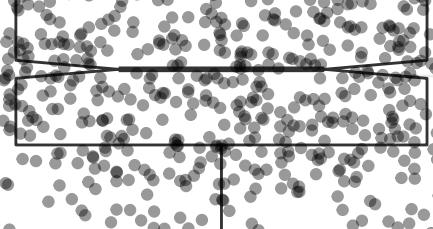
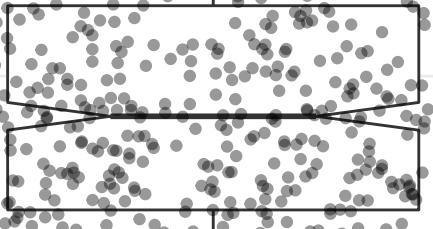
0.00

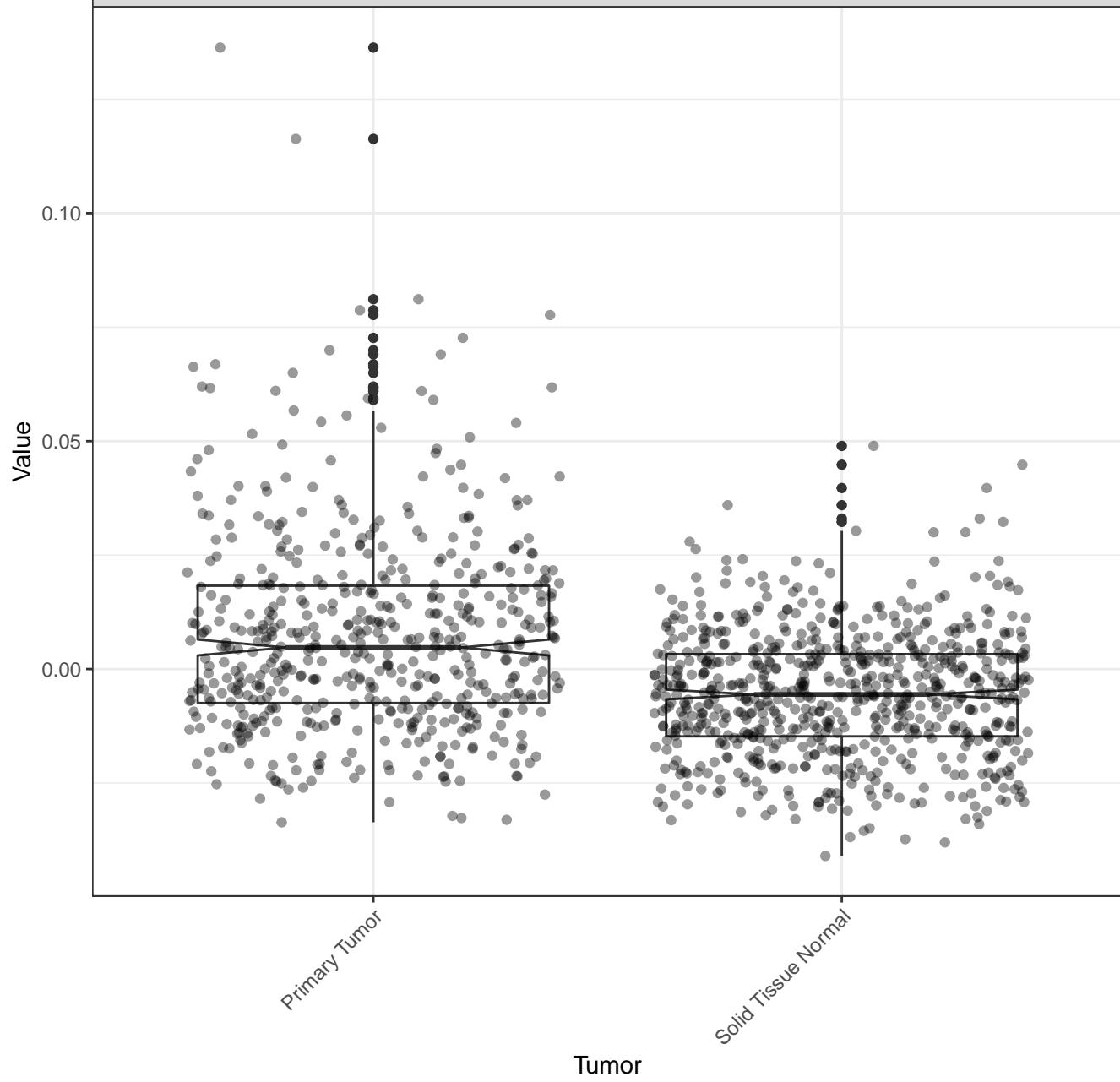
-0.05

Primary Tumor

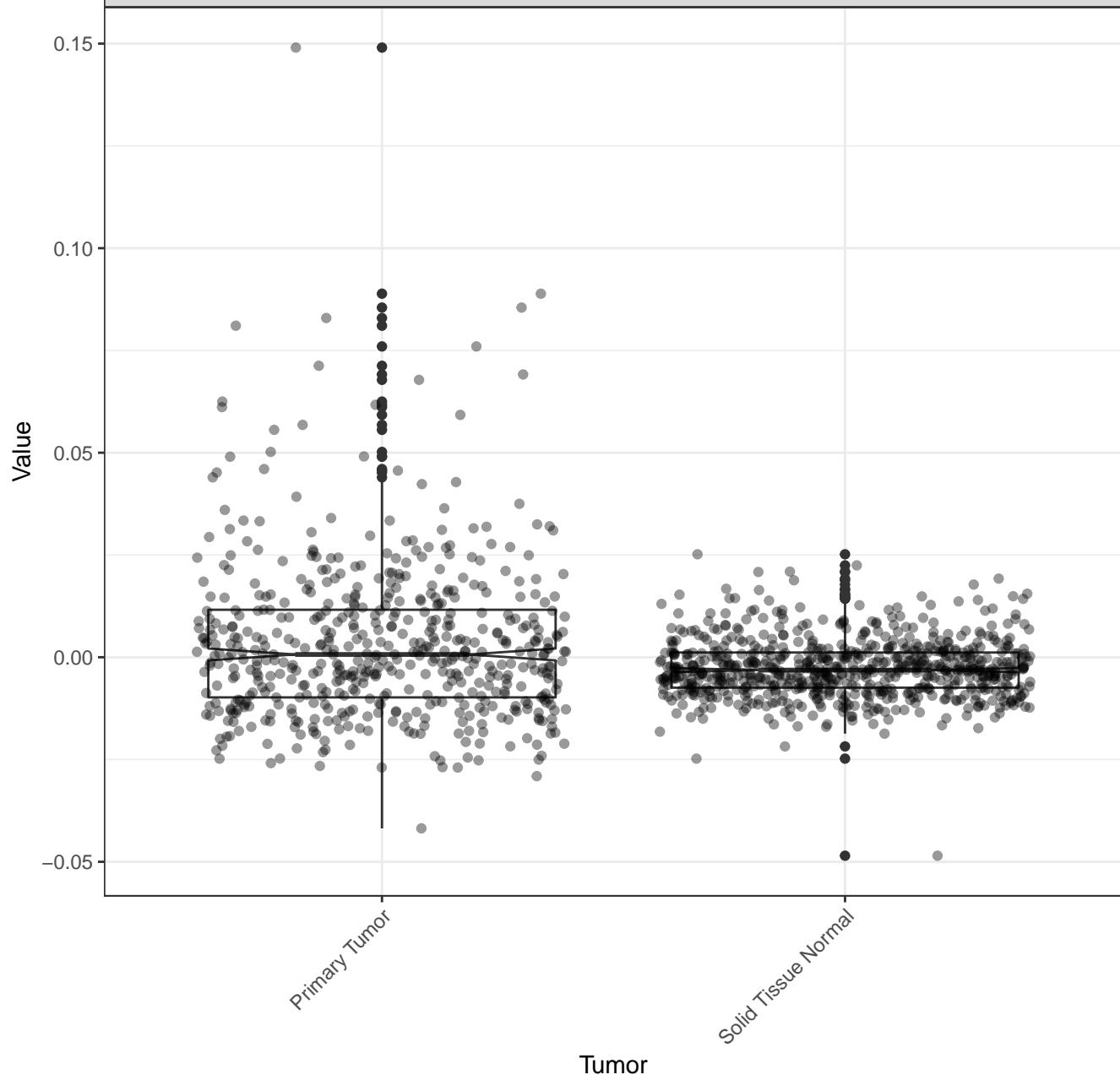
Solid Tissue Normal

Tumor

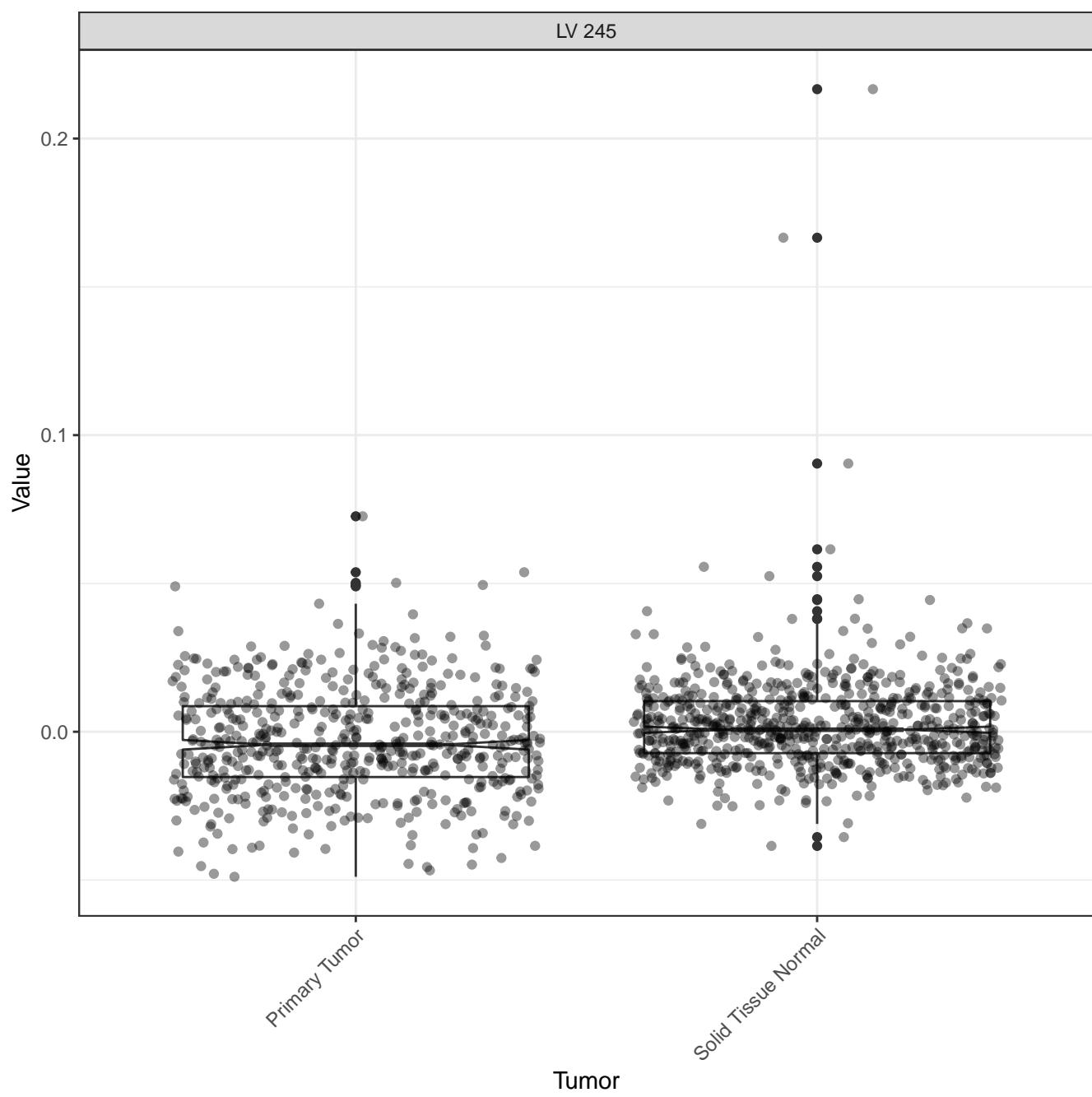




244,IRIS_Monocyte–Day7



LV 245



LV 246

Value

0.06

0.03

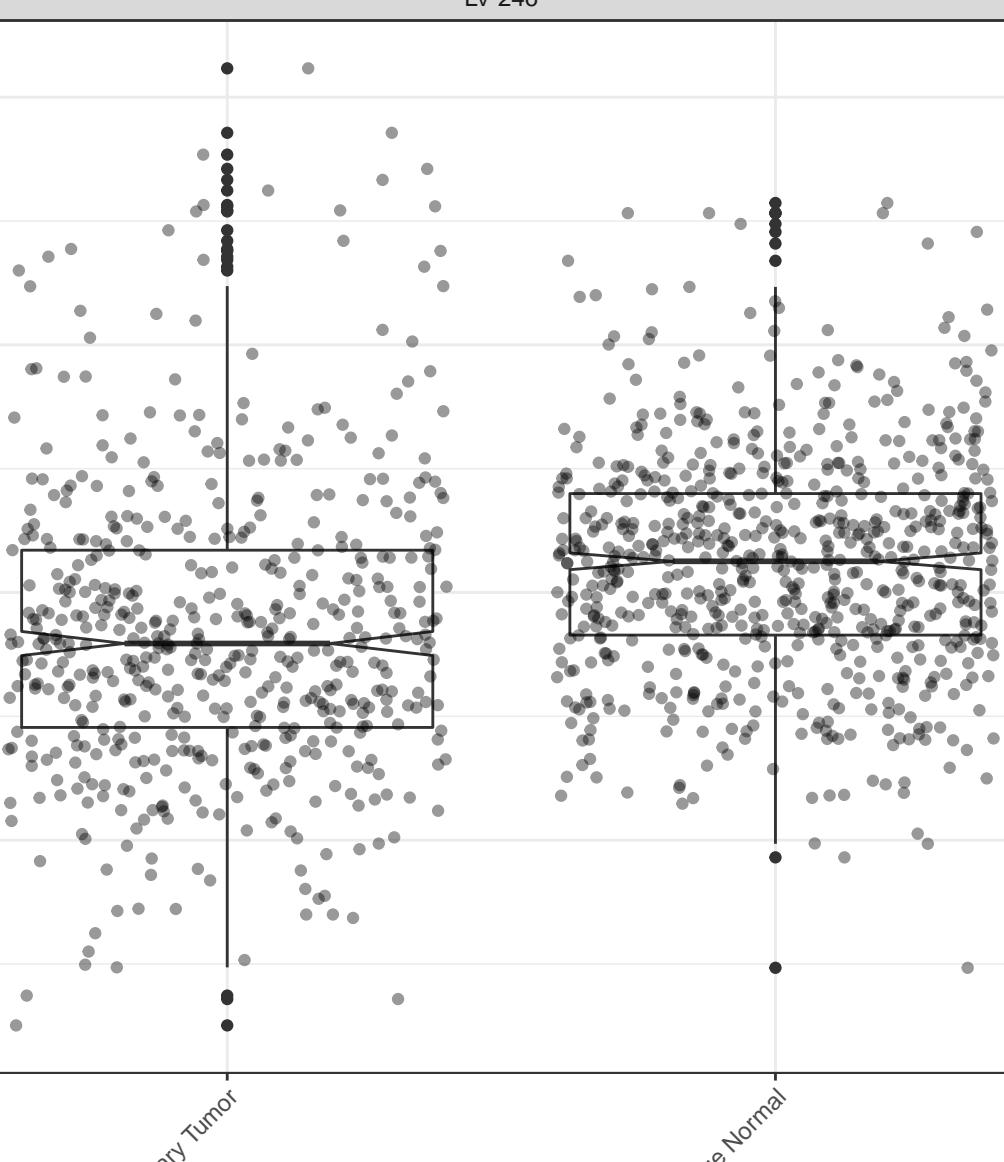
0.00

-0.03

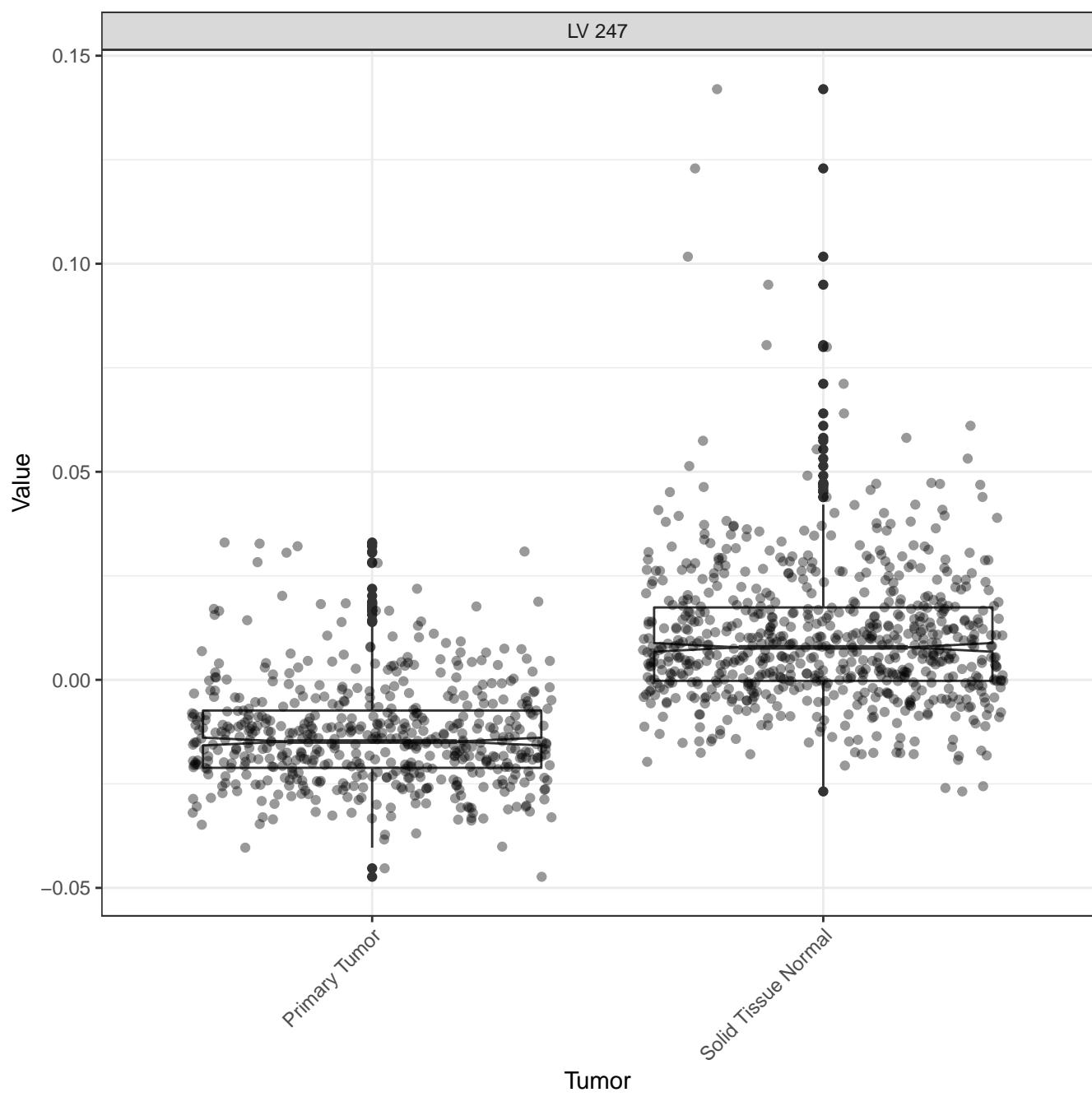
Primary Tumor

Solid Tissue Normal

Tumor



LV 247



248,REACTOME_EXTRACELLULAR_MATRIX_ORGANIZATION

Value

0.2

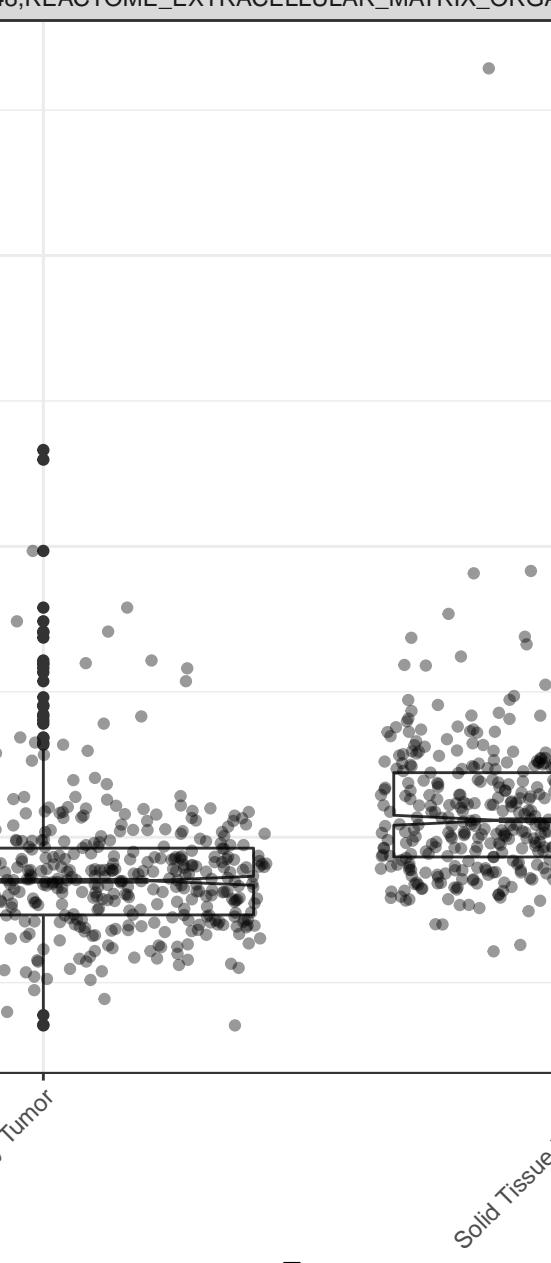
0.1

0.0

Primary Tumor

Solid Tissue Normal

Tumor



LV 249

Value

0.10

0.05

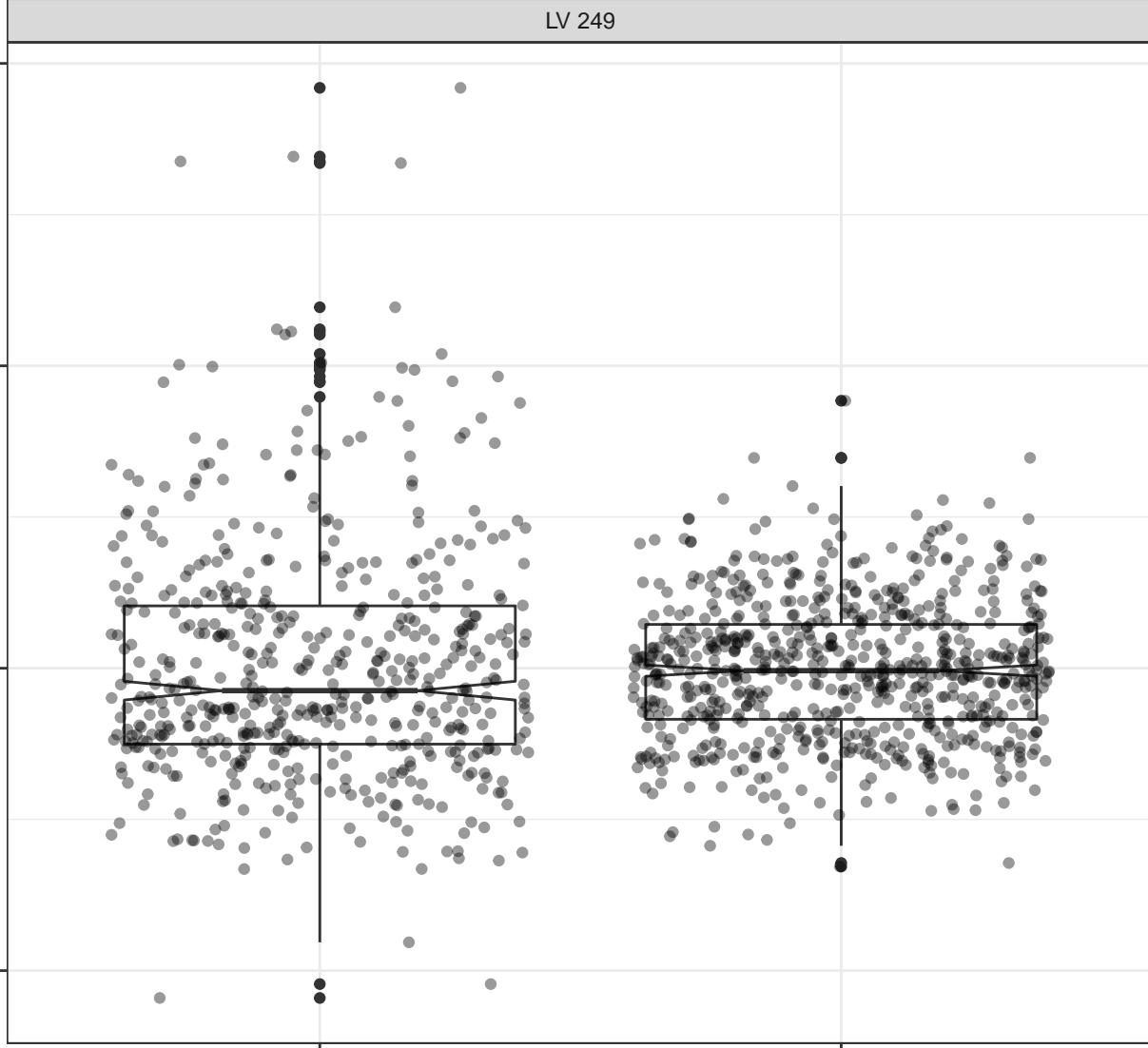
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



250,DMAP_HSC1

Value

0.10

0.05

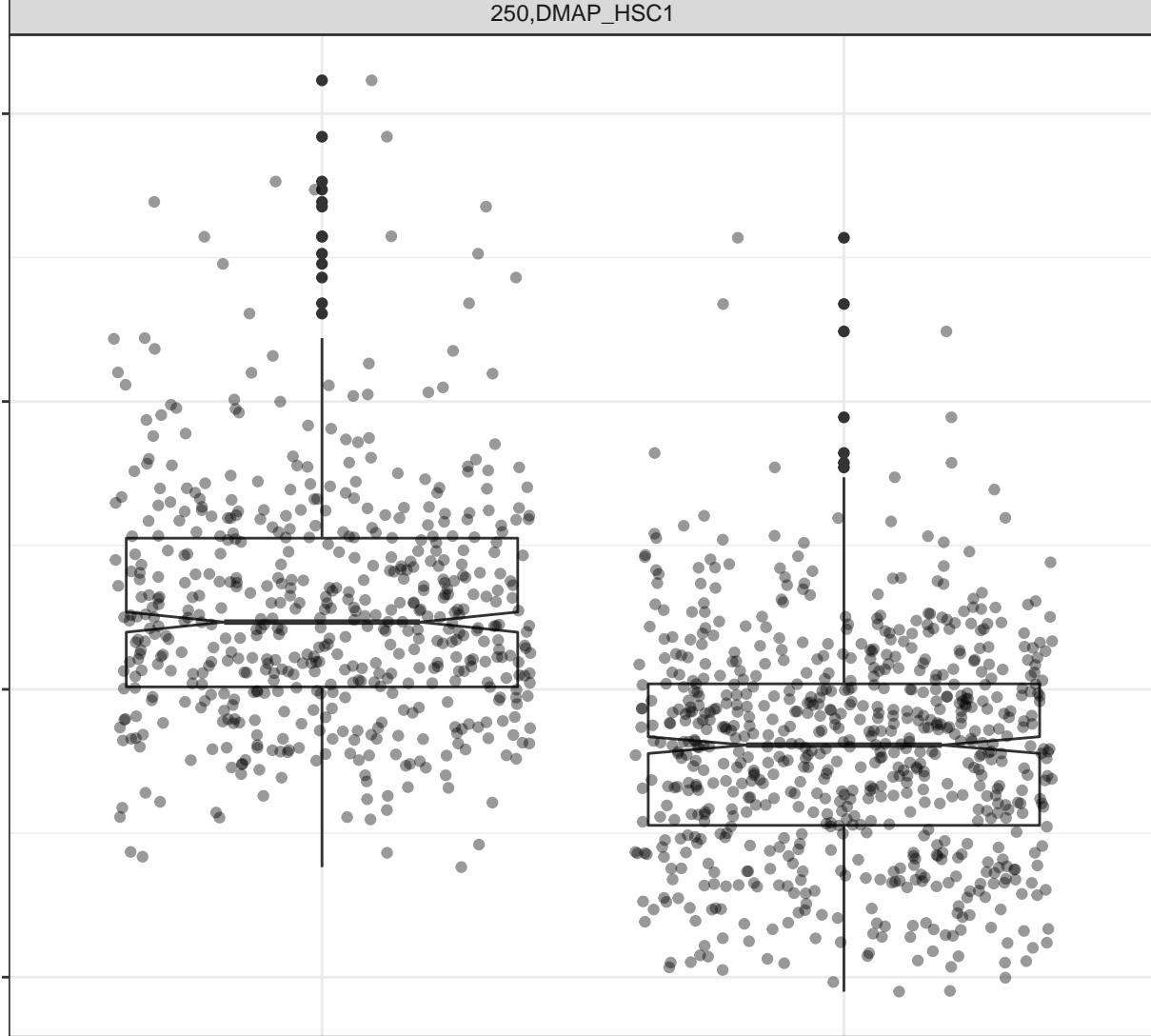
0.00

-0.05

Primary Tumor

Tumor

Solid Tissue Normal



Value

0.05

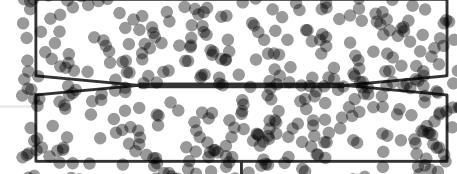
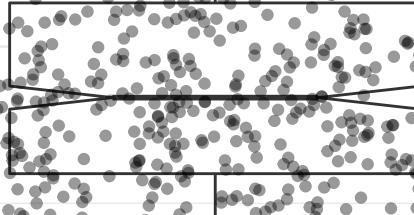
0.00

-0.05

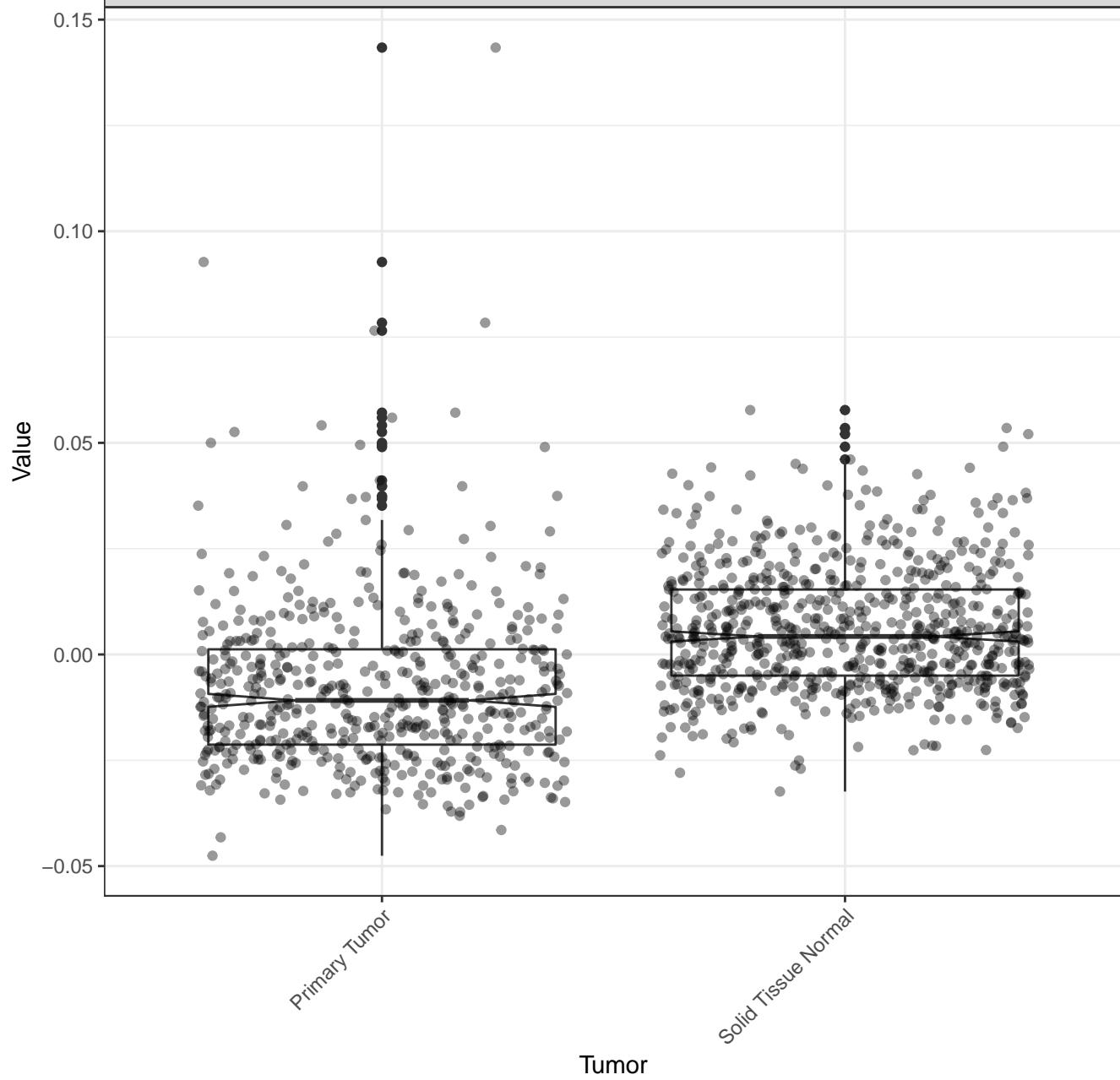
Primary Tumor

Solid Tissue Normal

Tumor



LV 252



253,KEGG_HEMATOPOIETIC_CELL_LINEAGE

Value

0.15

0.10

0.05

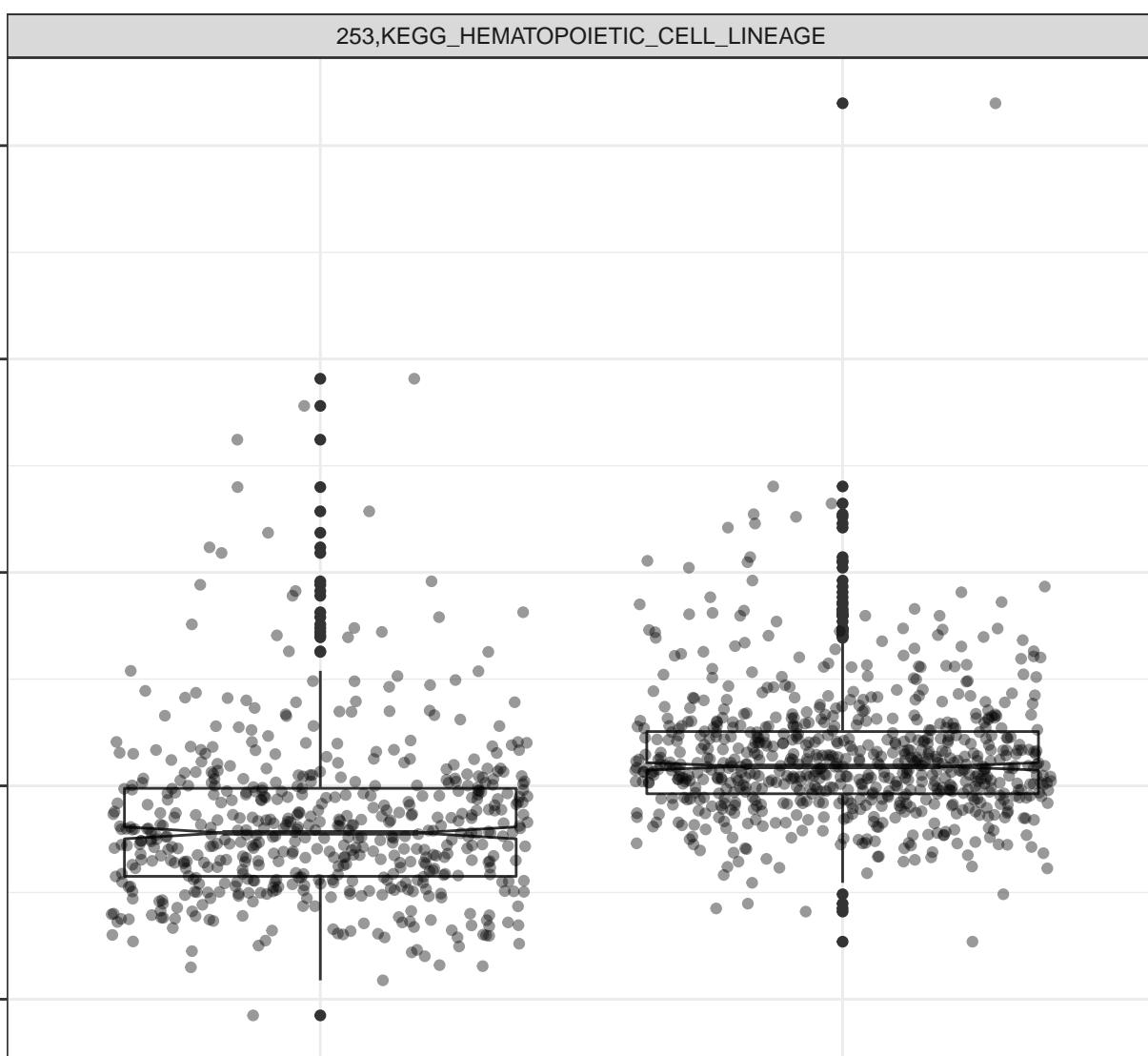
0.00

-0.05

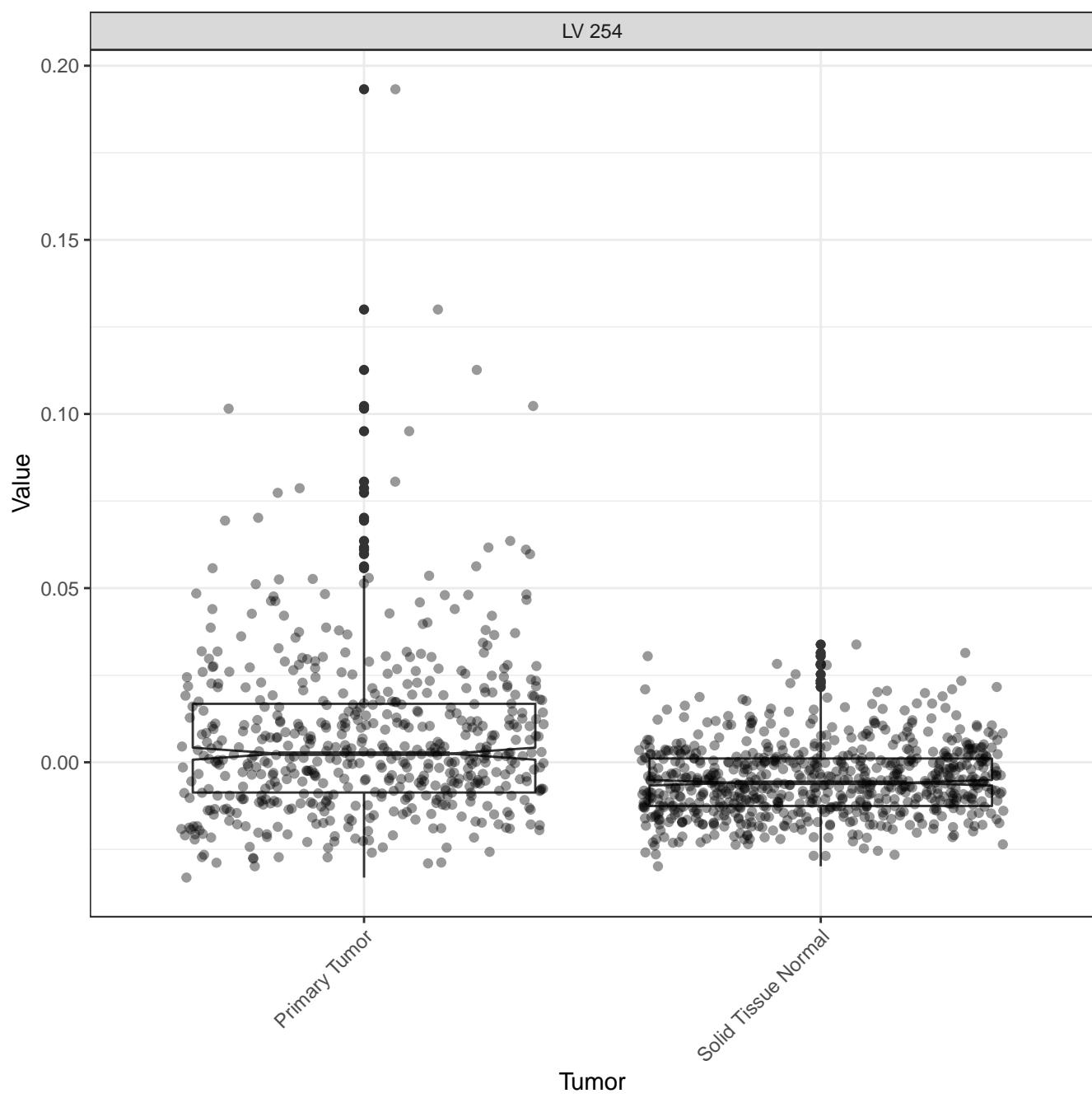
Primary Tumor

Solid Tissue Normal

Tumor



LV 254



255,KEGG_BASAL_CELL_CARCINOMA

Value

0.10

0.05

0.00

-0.05

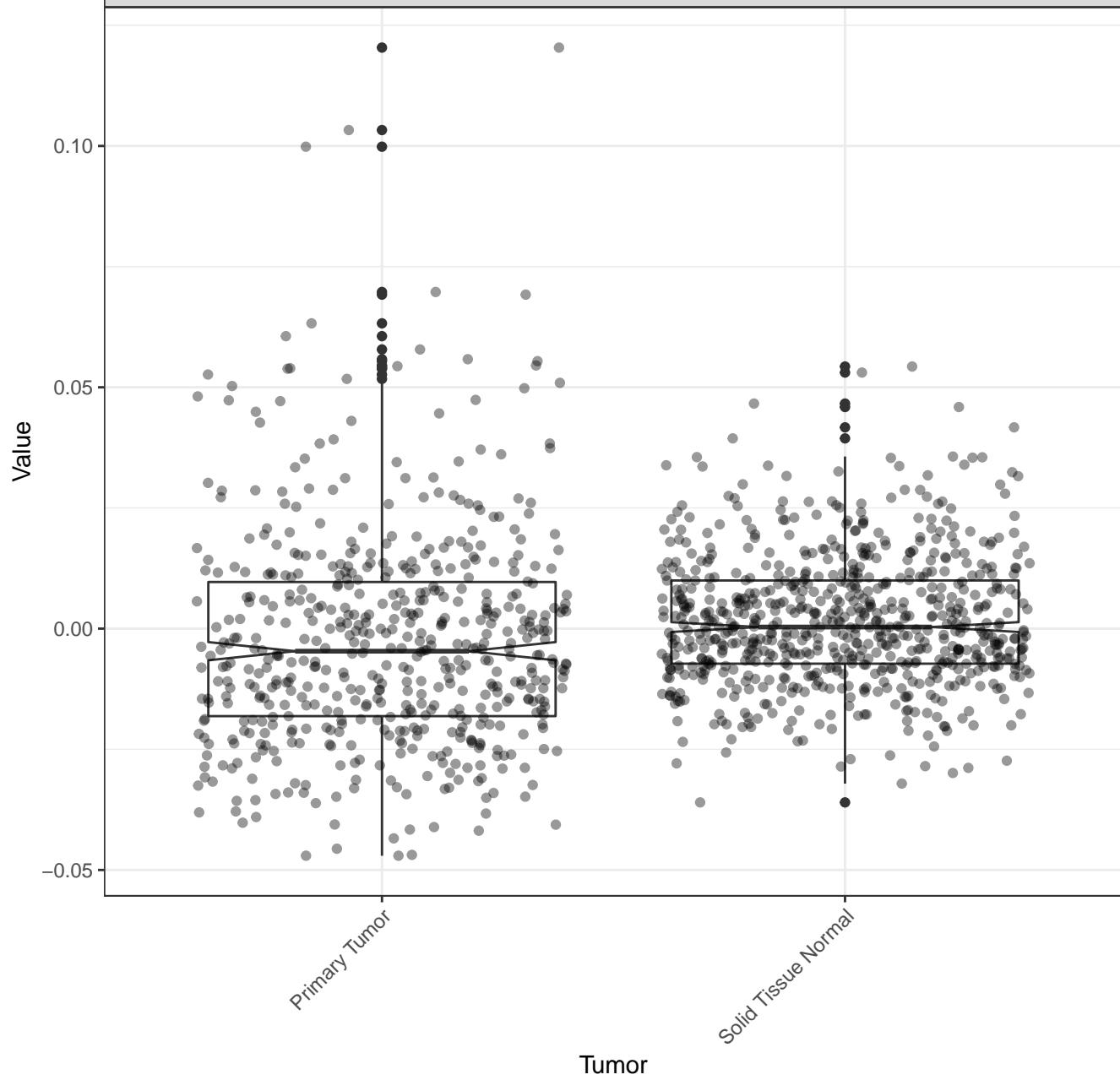
Primary Tumor

Solid Tissue Normal

Tumor



256,KEGG_ARGININE_AND_PROLINE_METABOLISM



Value

0.10

0.05

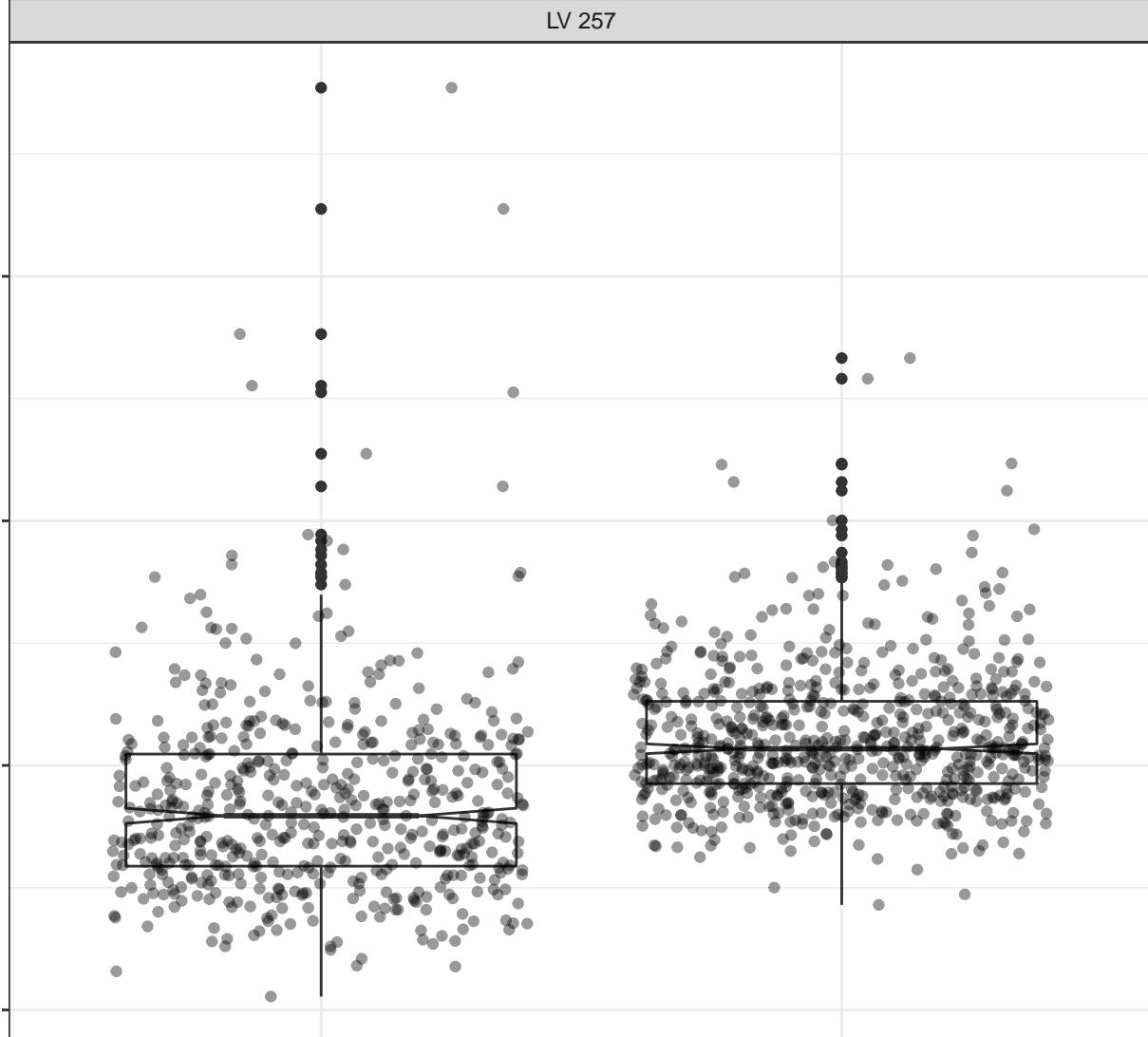
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 258

Value

0.10

0.05

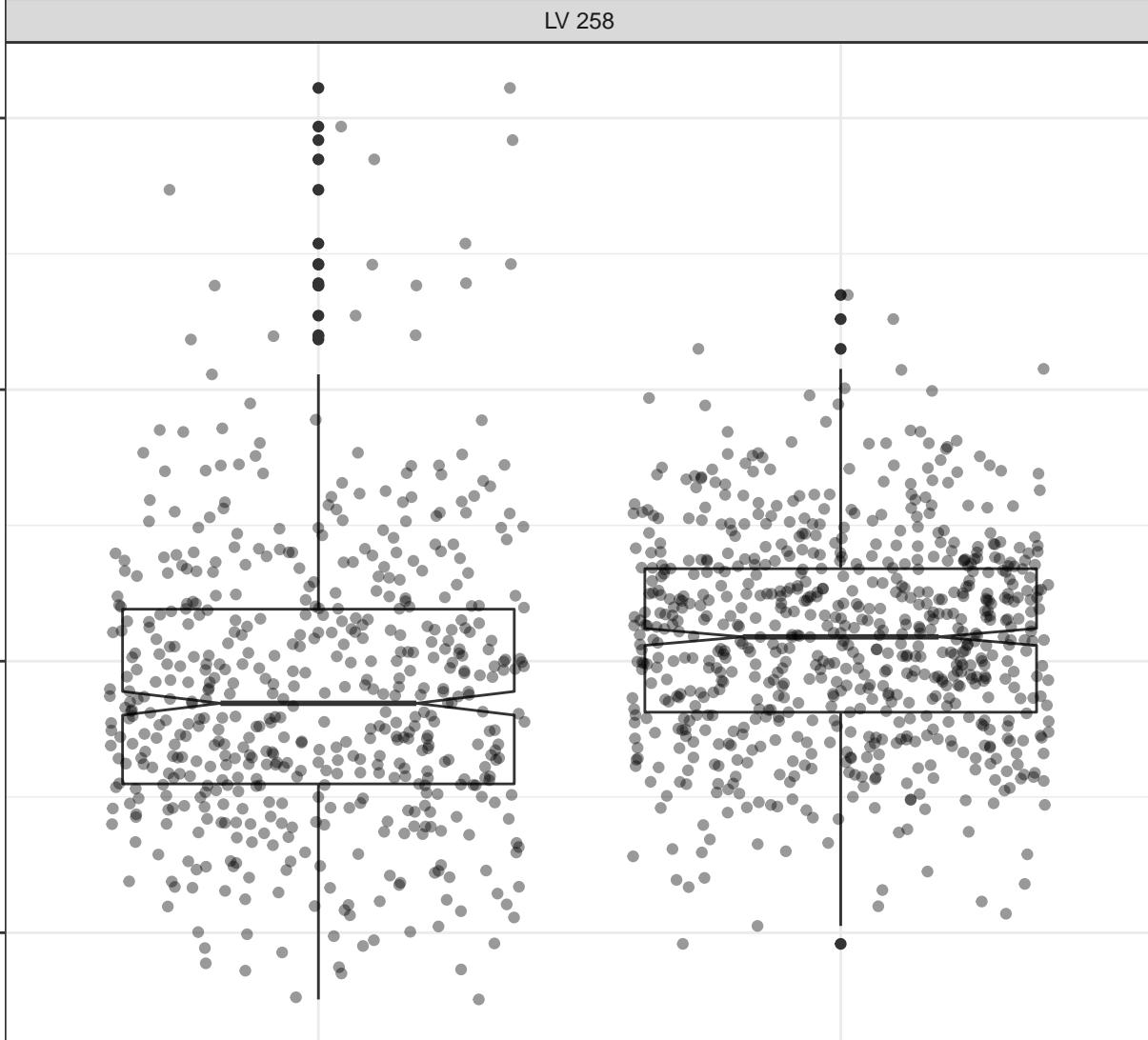
0.00

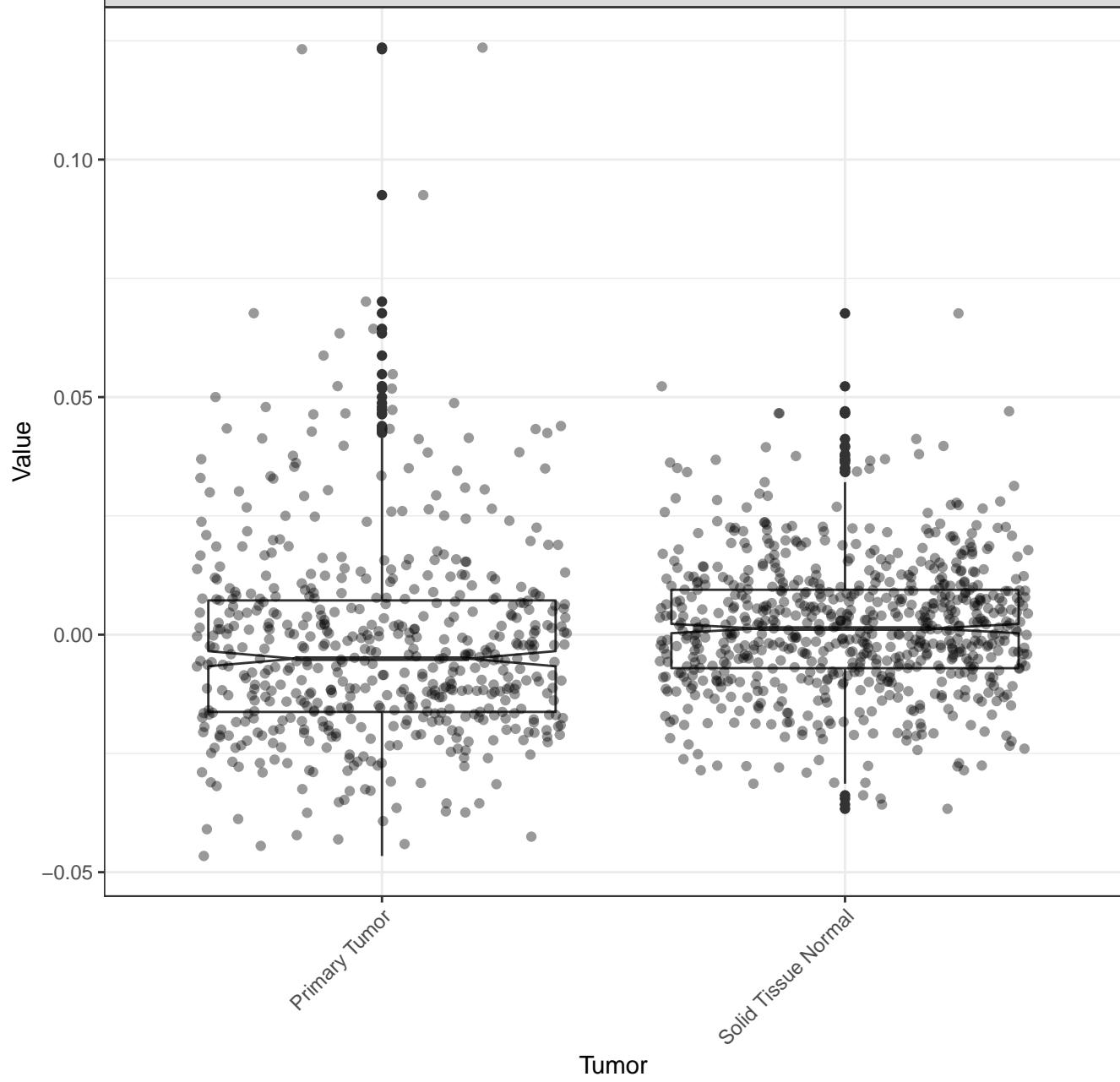
-0.05

Primary Tumor

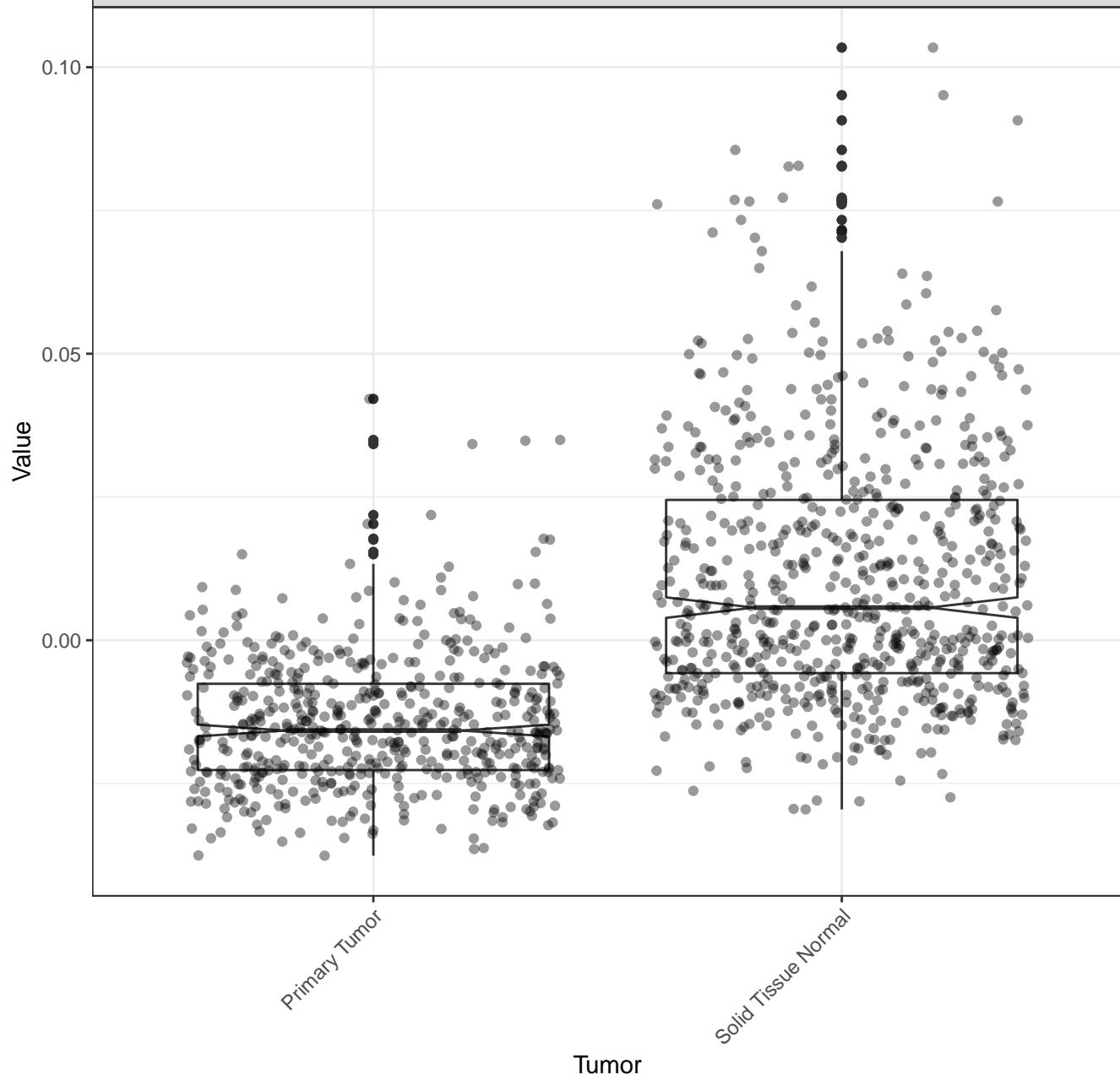
Solid Tissue Normal

Tumor

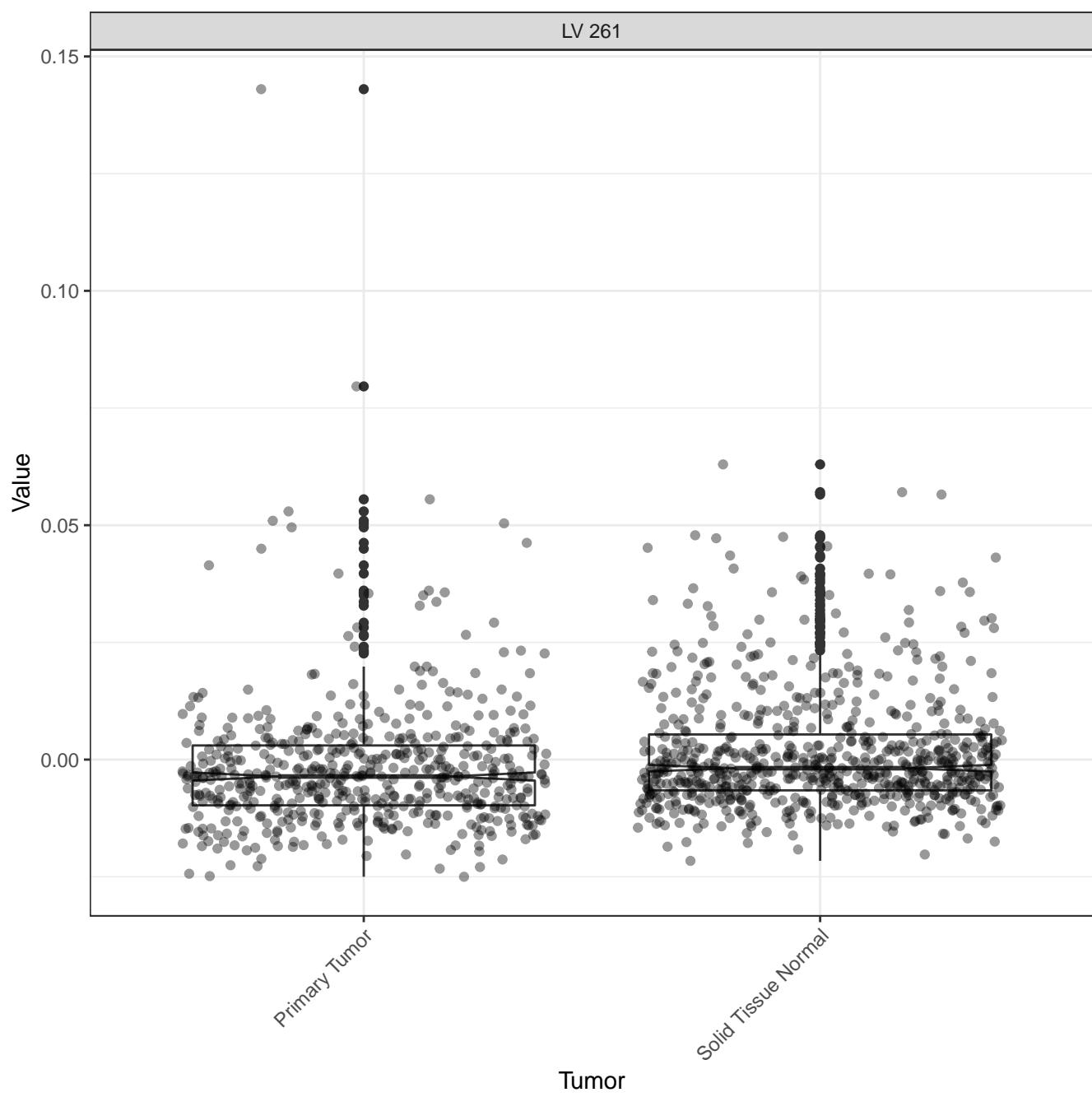




260, BIOCARTA_CHREBP2_PATHWAY



LV 261



LV 262

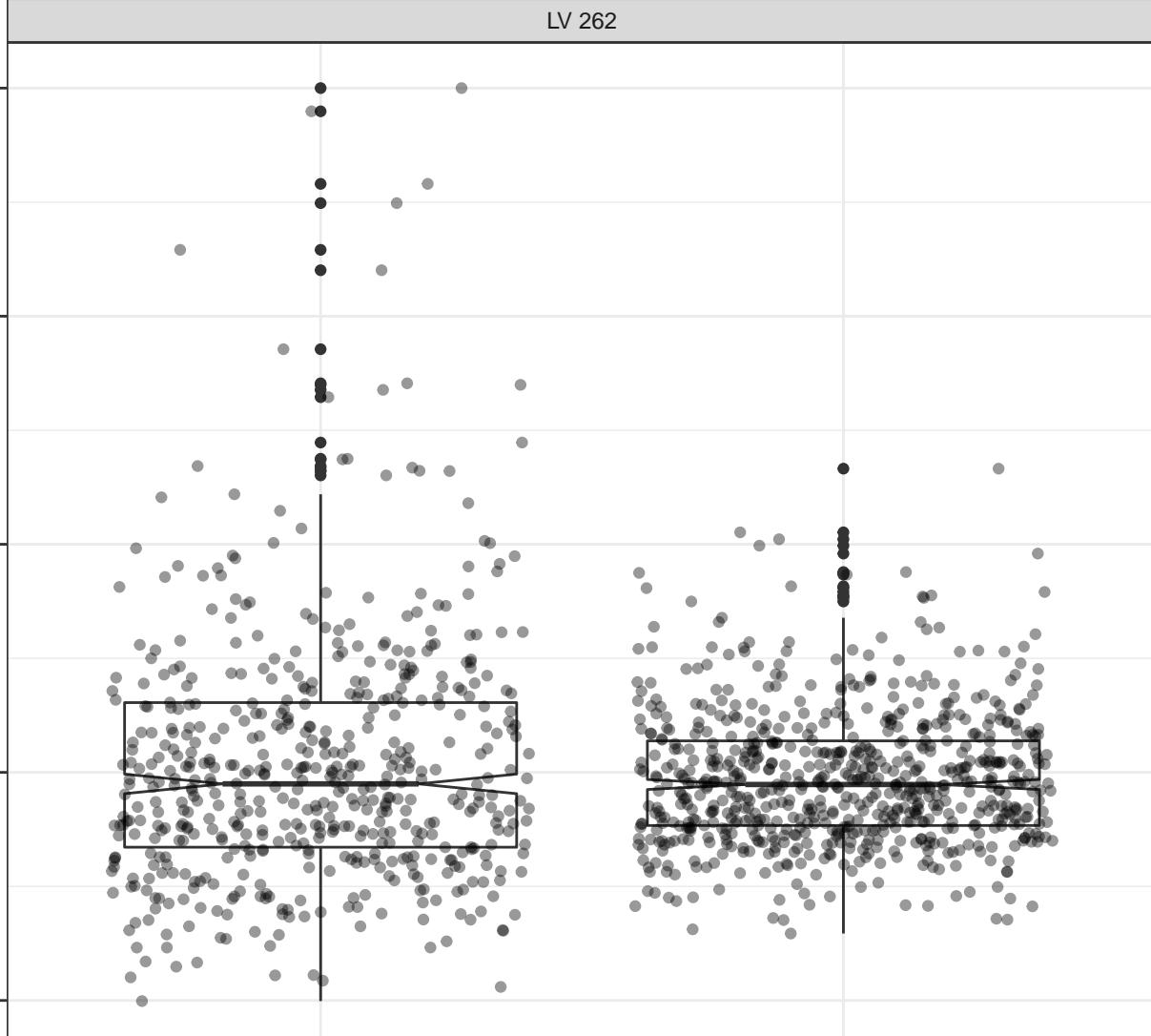
Value

0.15
0.10
0.05
0.00
-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 263

Value

0.3

0.2

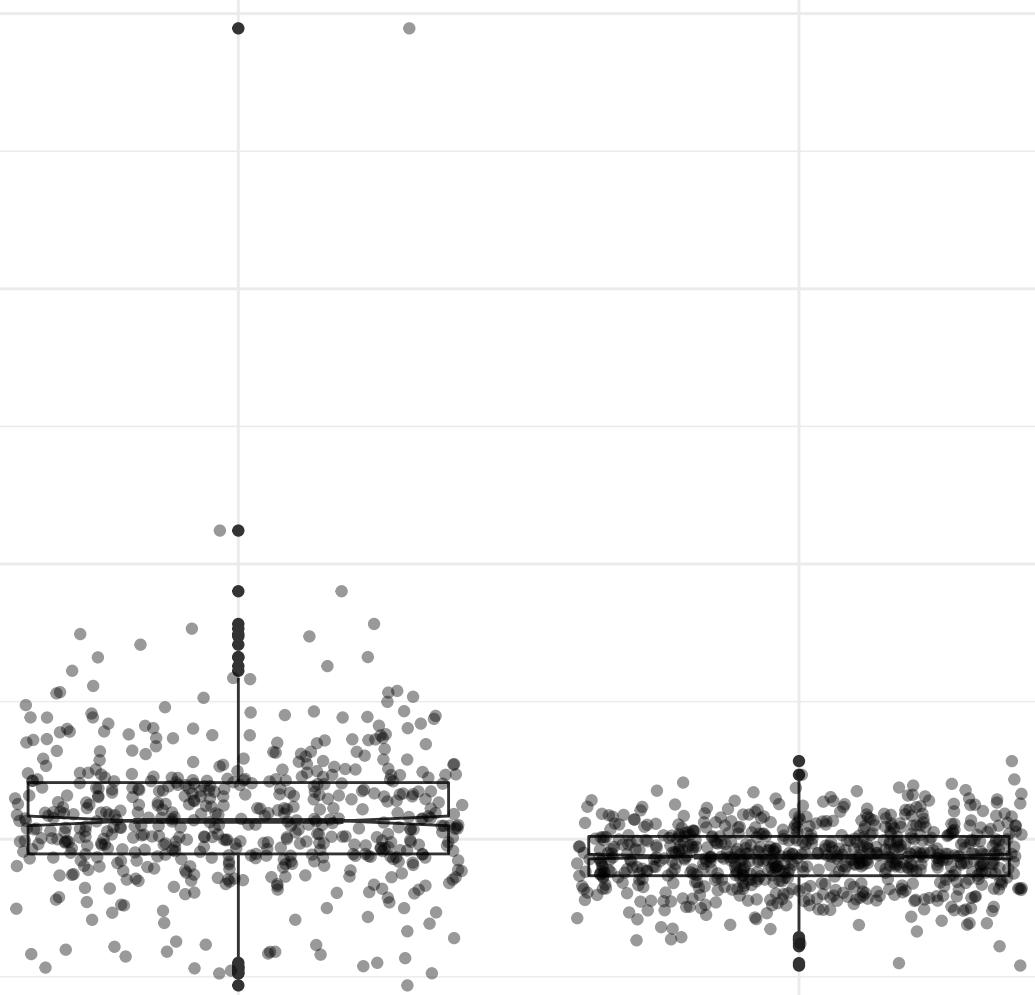
0.1

0.0

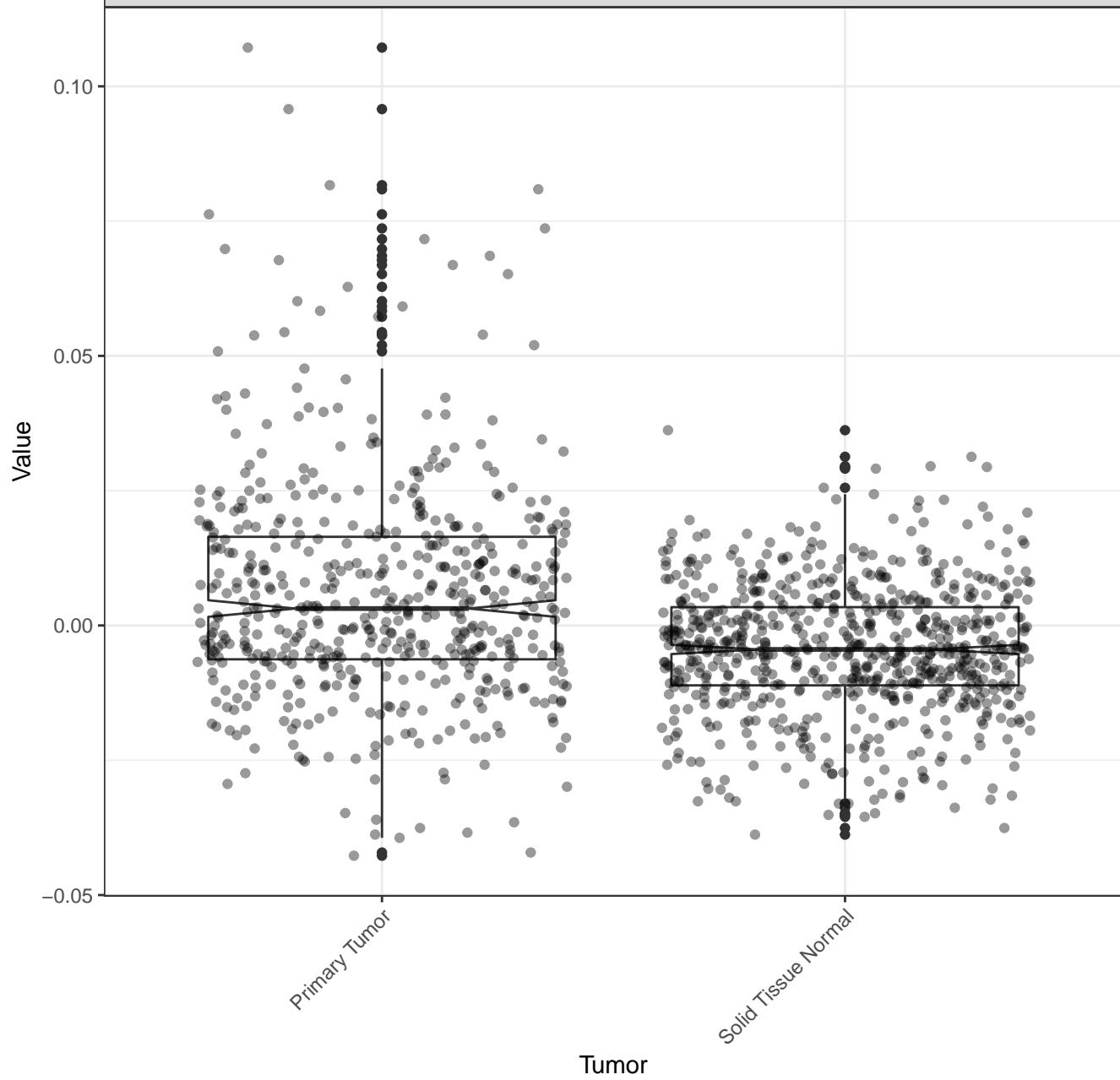
Primary Tumor

Solid Tissue Normal

Tumor



264,REACTOME_TRANSMEMBRANE_TRANSPORT_OF_SMALL_MOLECULES



Value

0.05

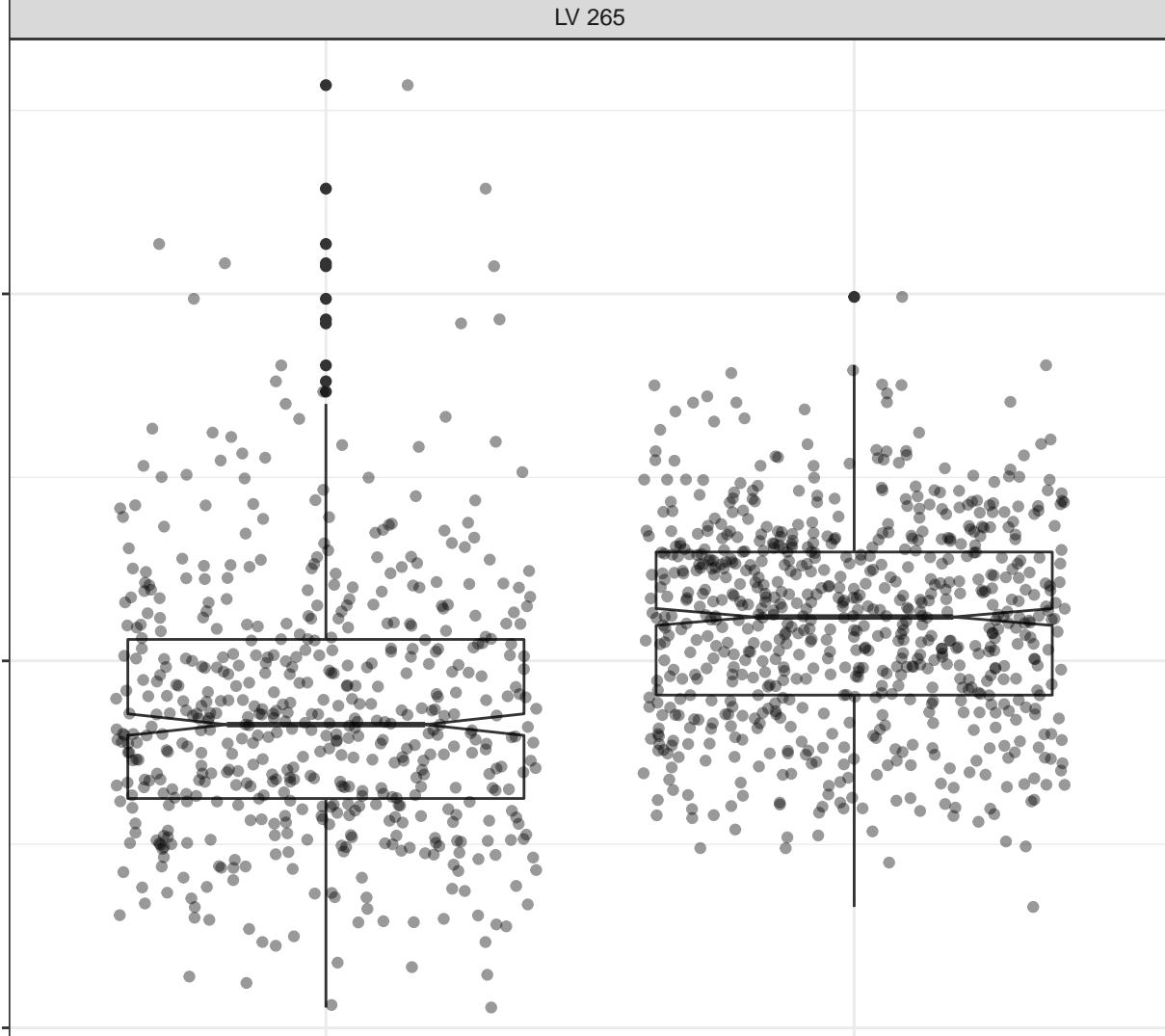
0.00

-0.05

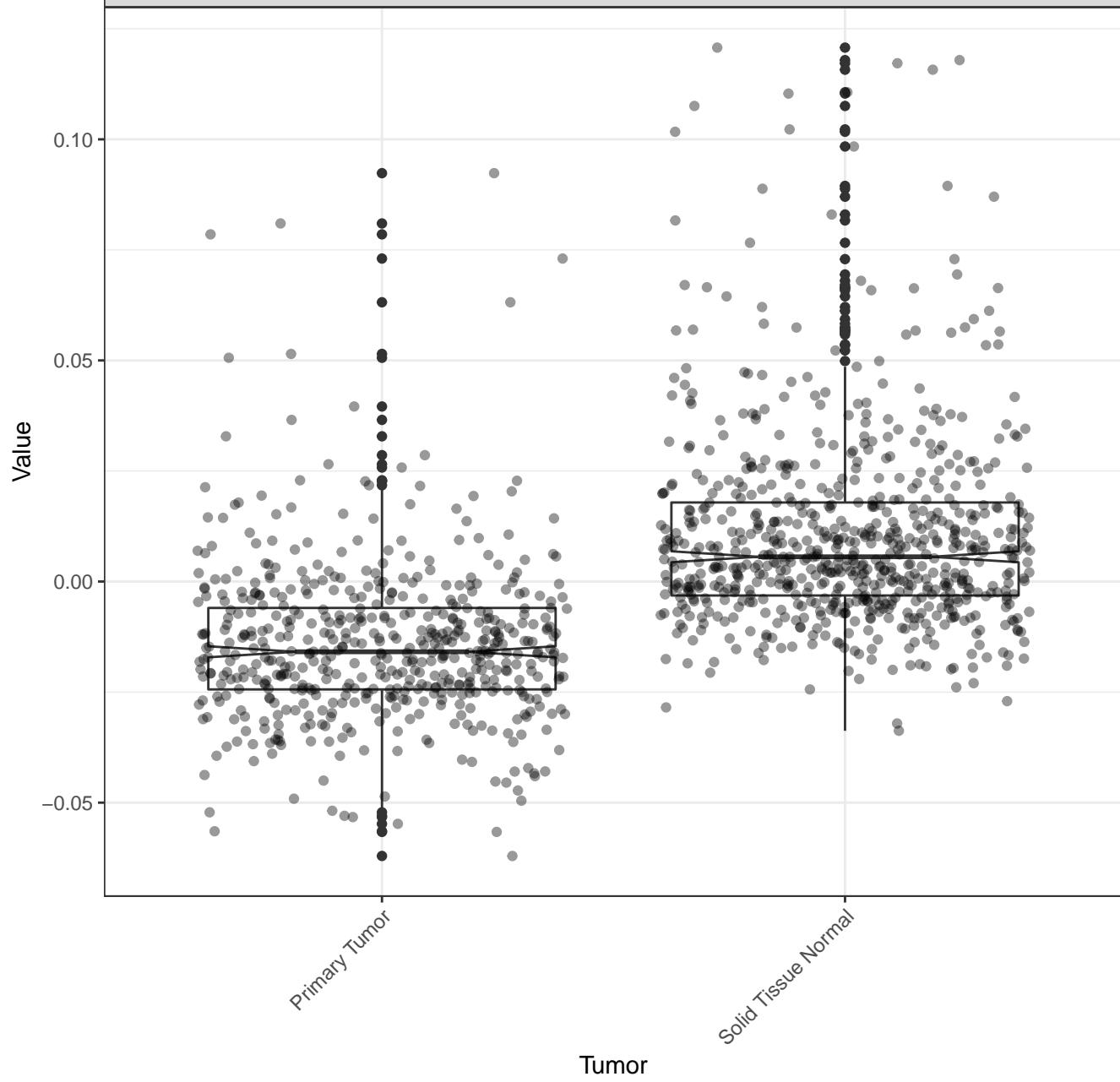
Primary Tumor

Solid Tissue Normal

Tumor



266,IRIS_Monocyte–Day1



267,KEGG_VASOPRESSIN_REGULATED_WATER_REABSORPTION

Value

0.10

0.05

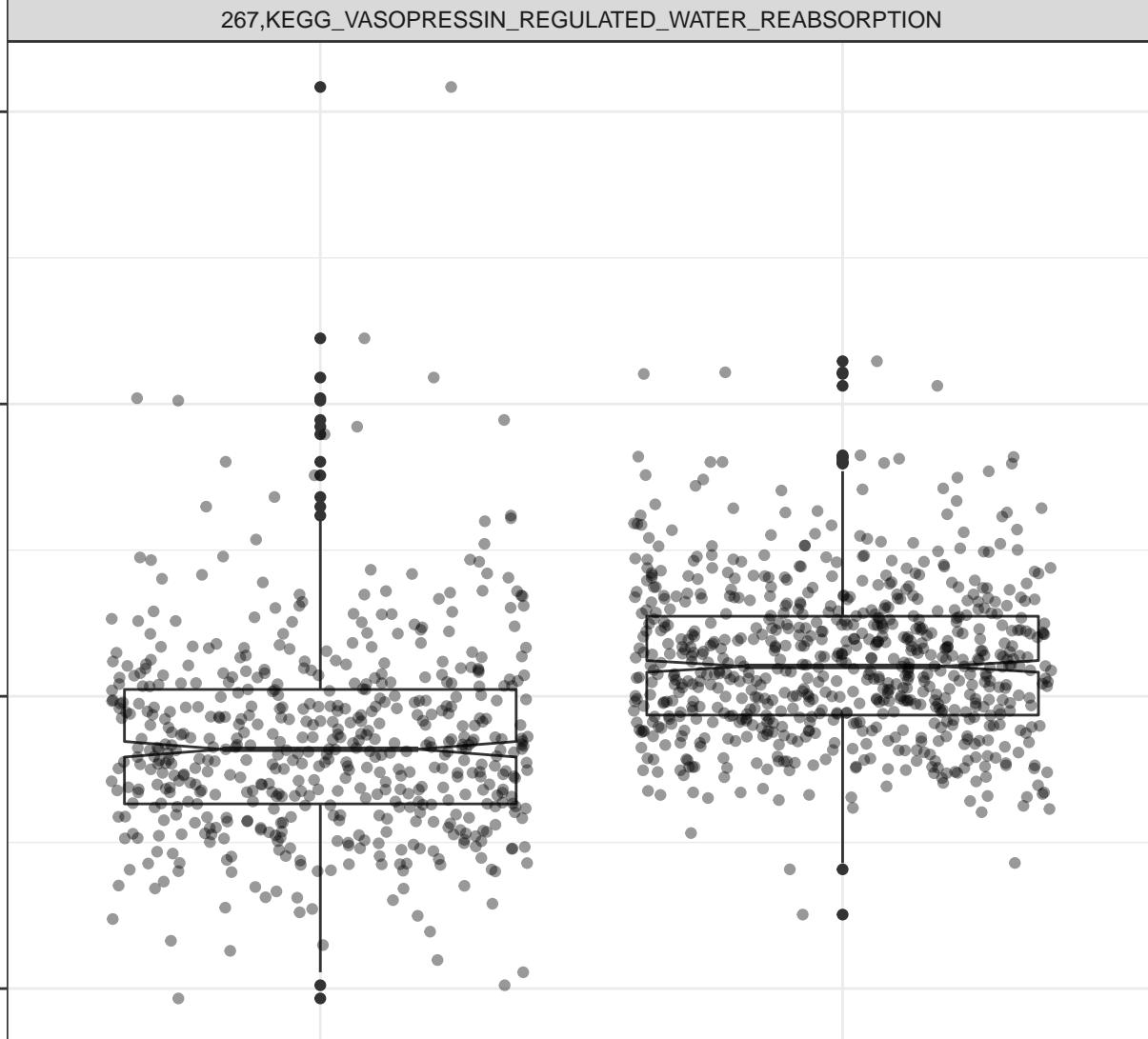
0.00

-0.05

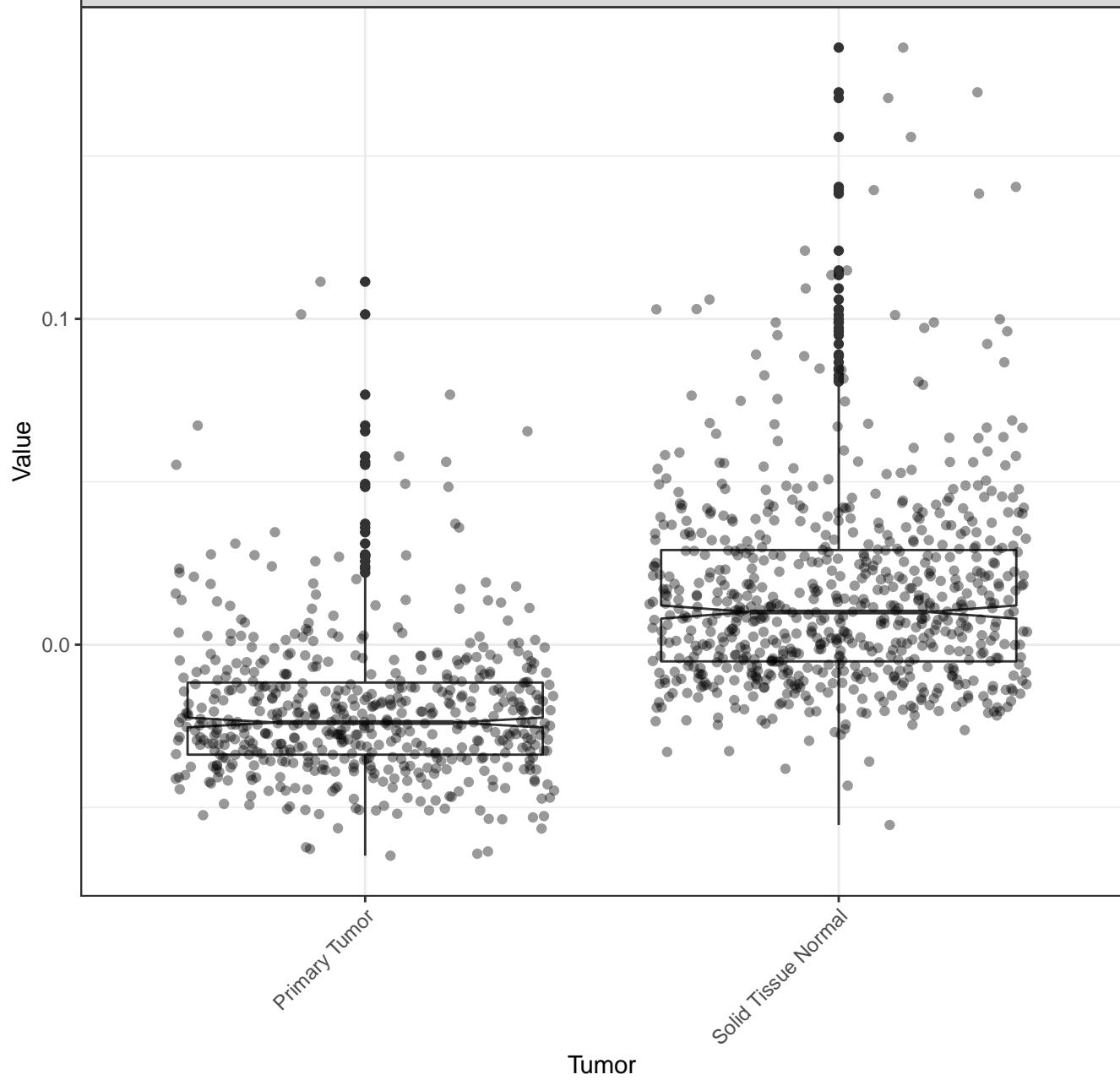
Primary Tumor

Solid Tissue Normal

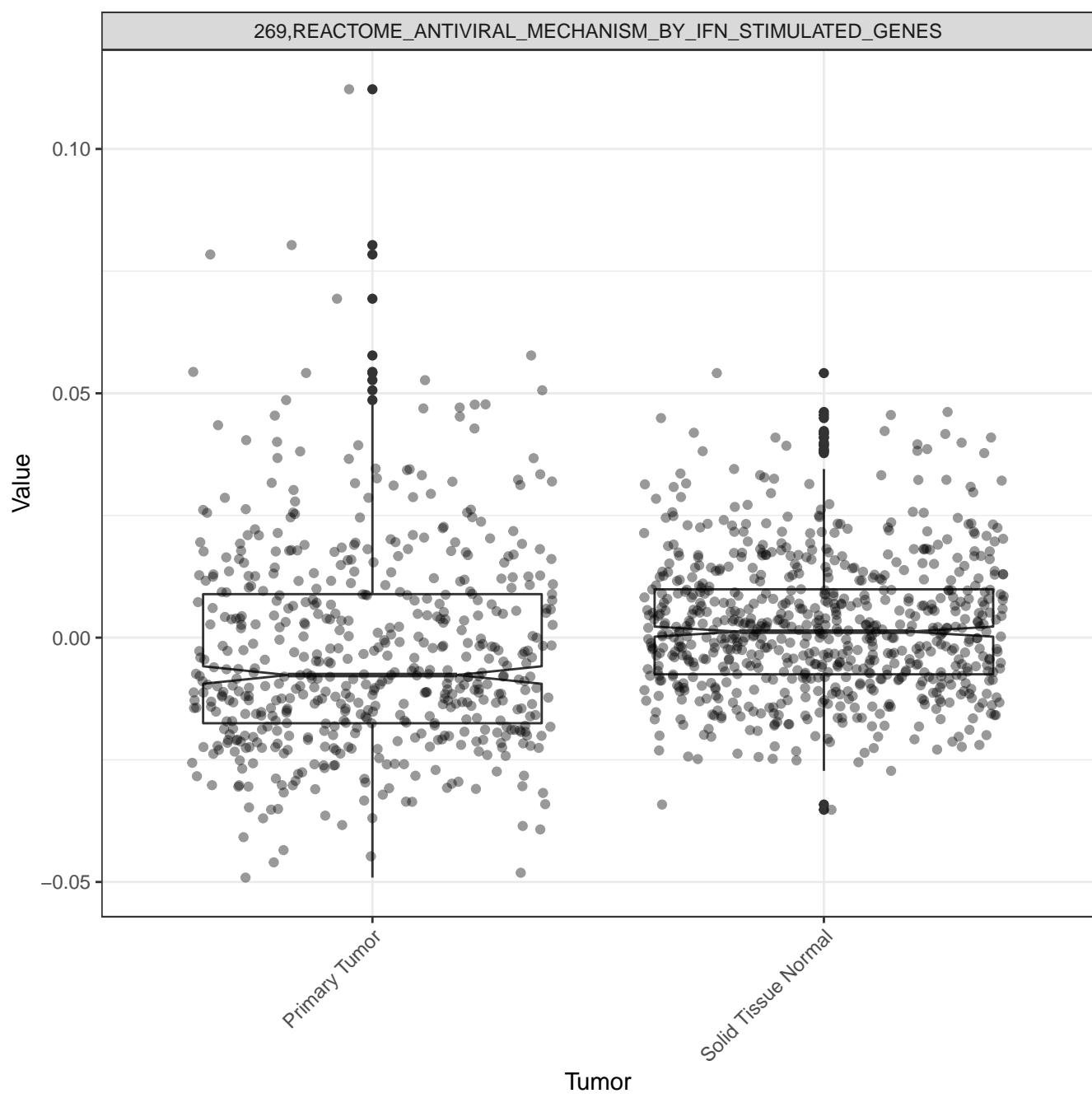
Tumor



268,SVM Mast cells activated



269,REACTOME_ANTIVIRAL_MECHANISM_BY_IFN_STIMULATED_GENES



270,KEGG_CELL_CYCLE

Value

0.10

0.05

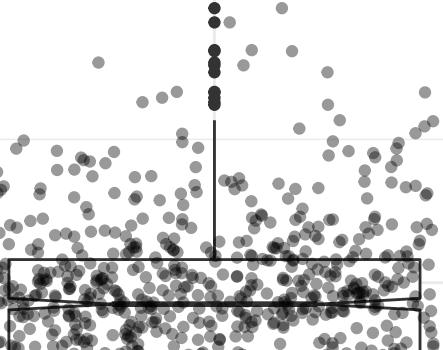
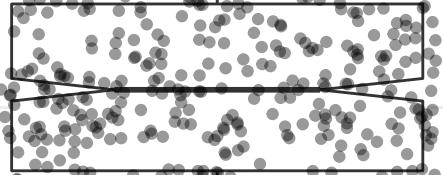
0.00

-0.05

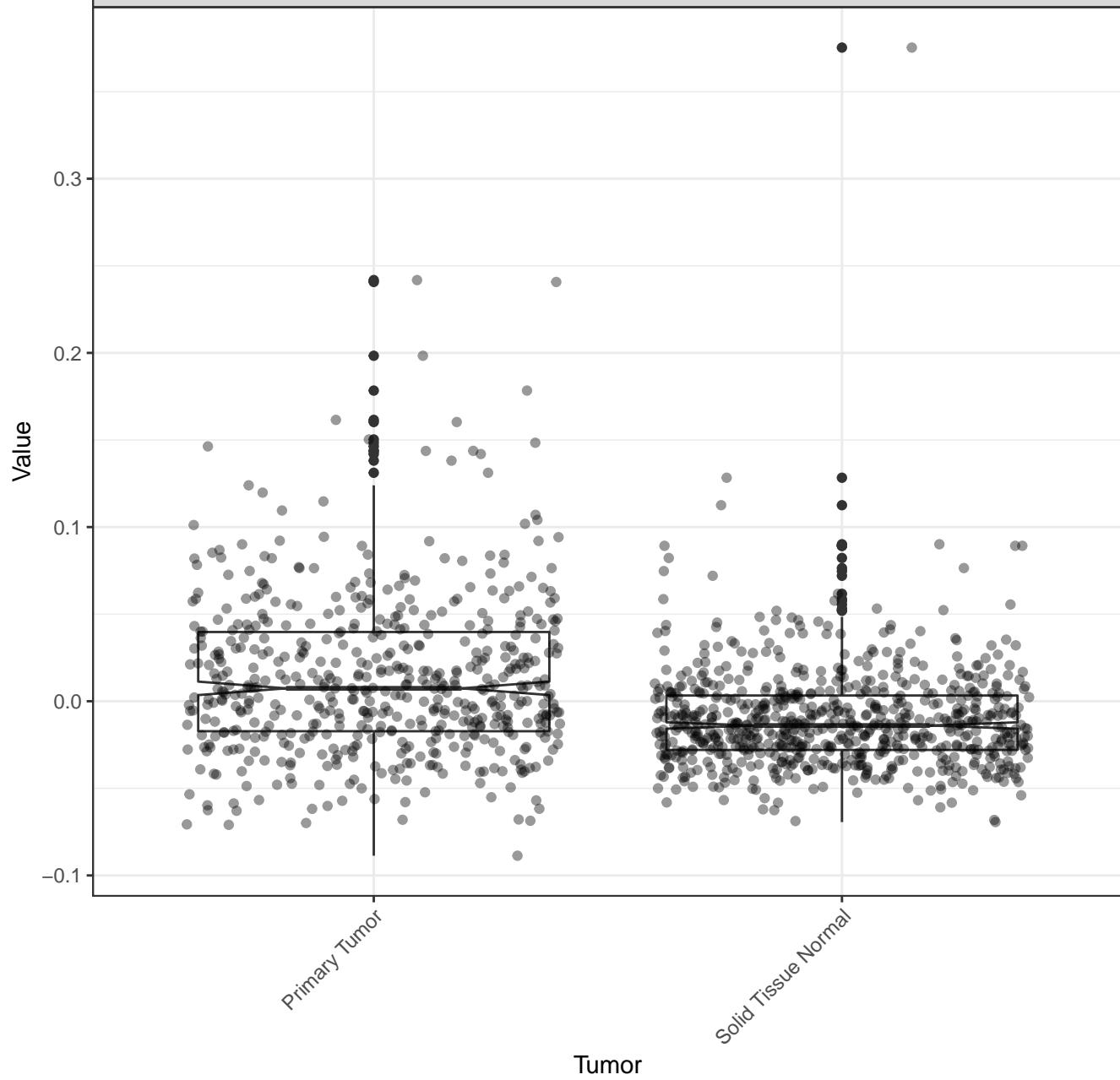
Primary Tumor

Solid Tissue Normal

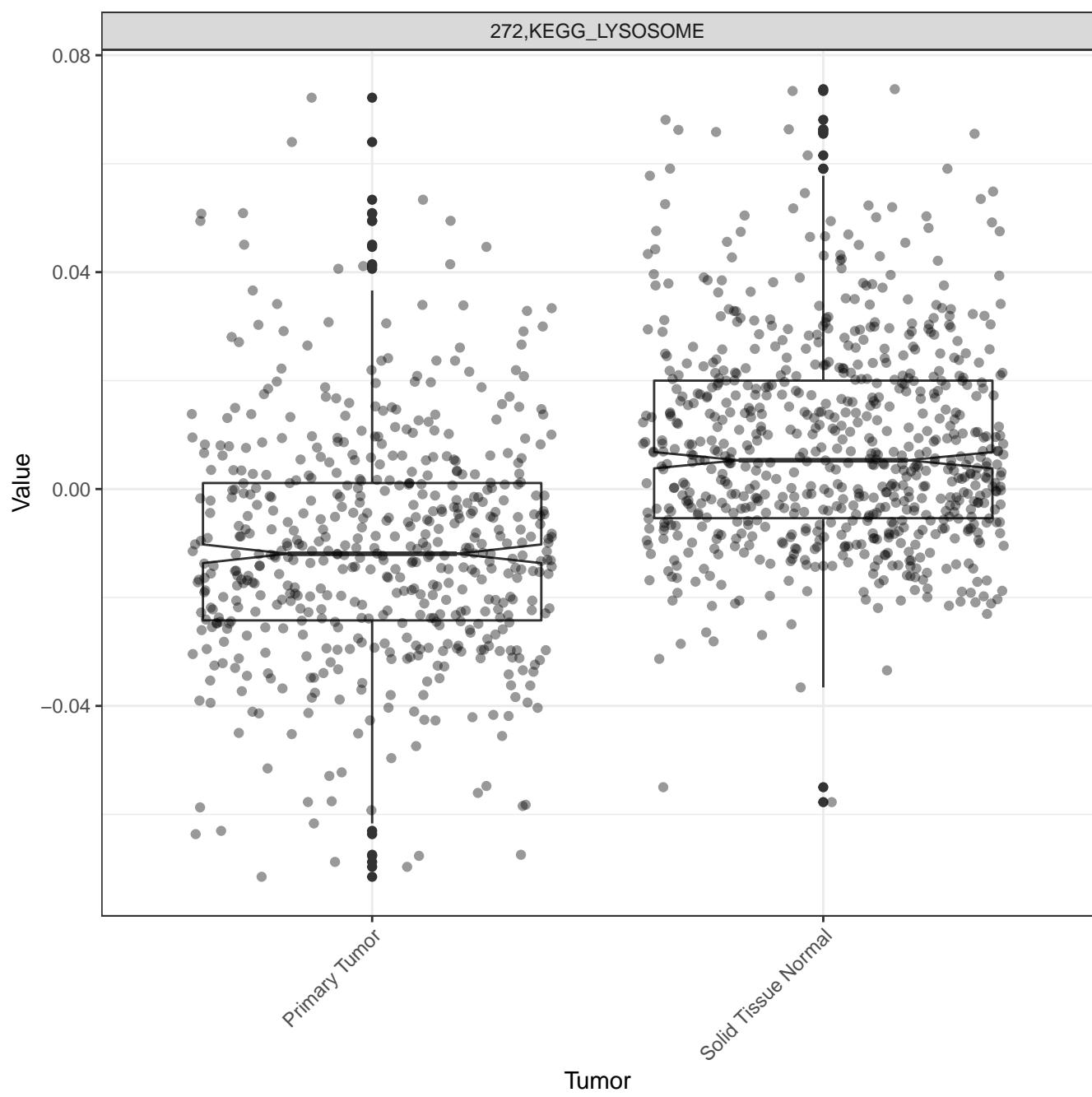
Tumor



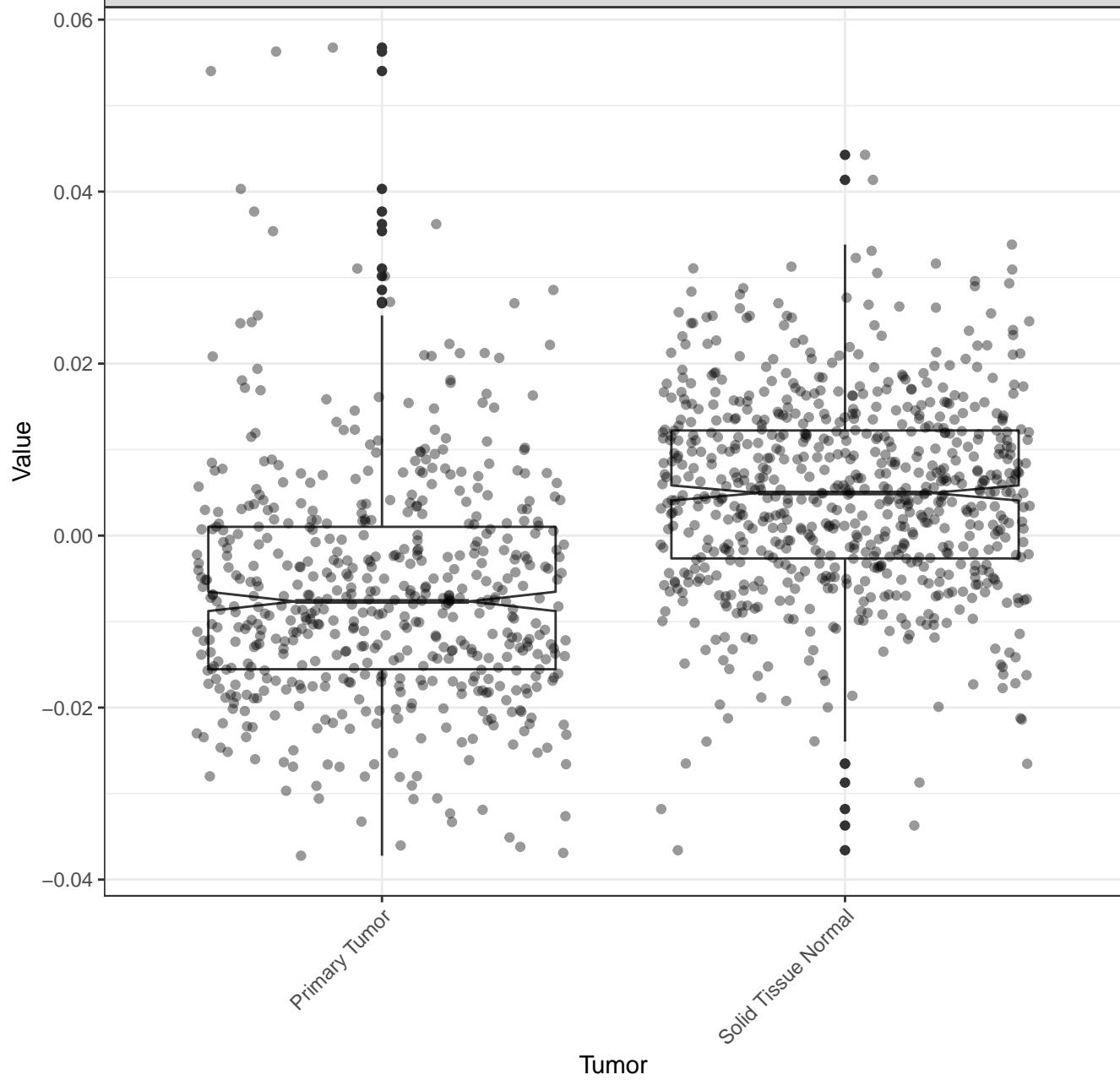
271,REACTOME_INTERFERON_ALPHA_BETA_SIGNALING



272,KEGG_LYSOSOME



273,PID_RHOA_PATHWAY



LV 274

Value

0.15

0.10

0.05

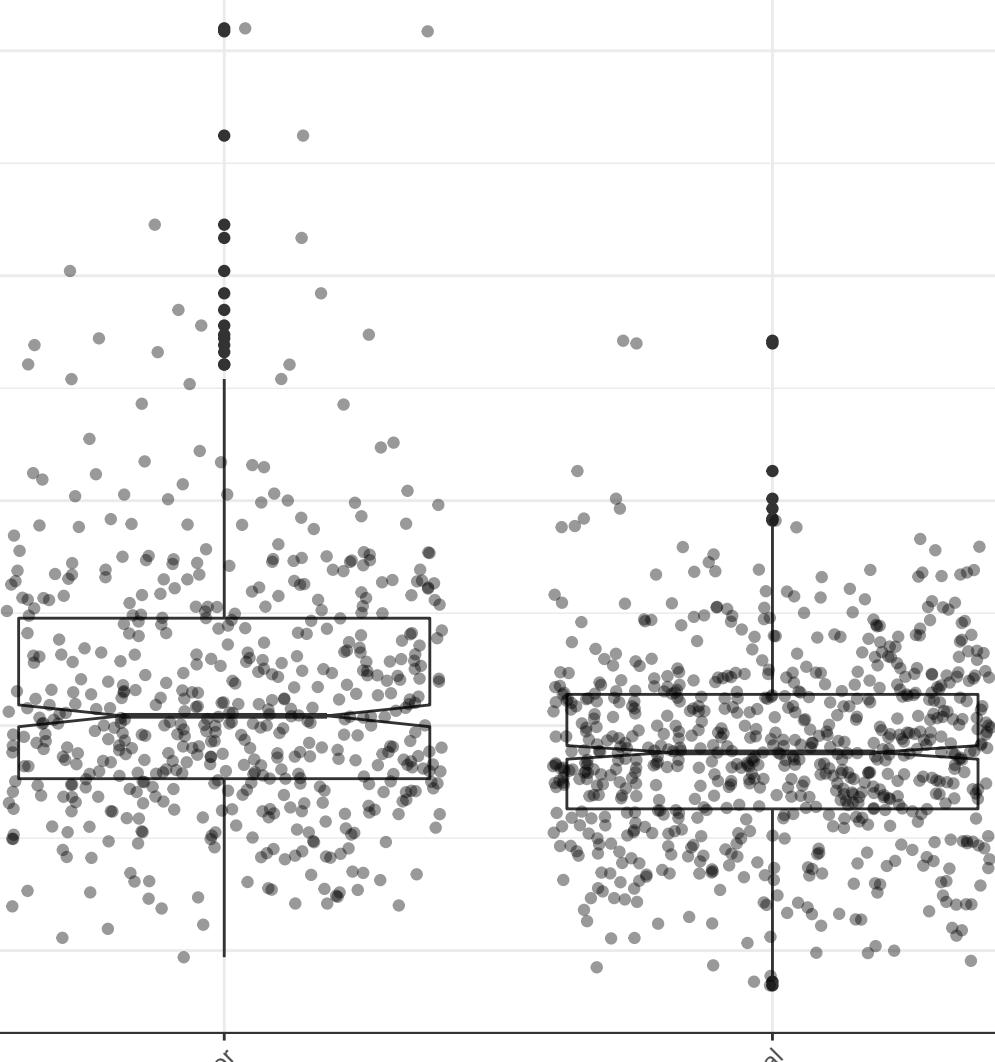
0.00

-0.05

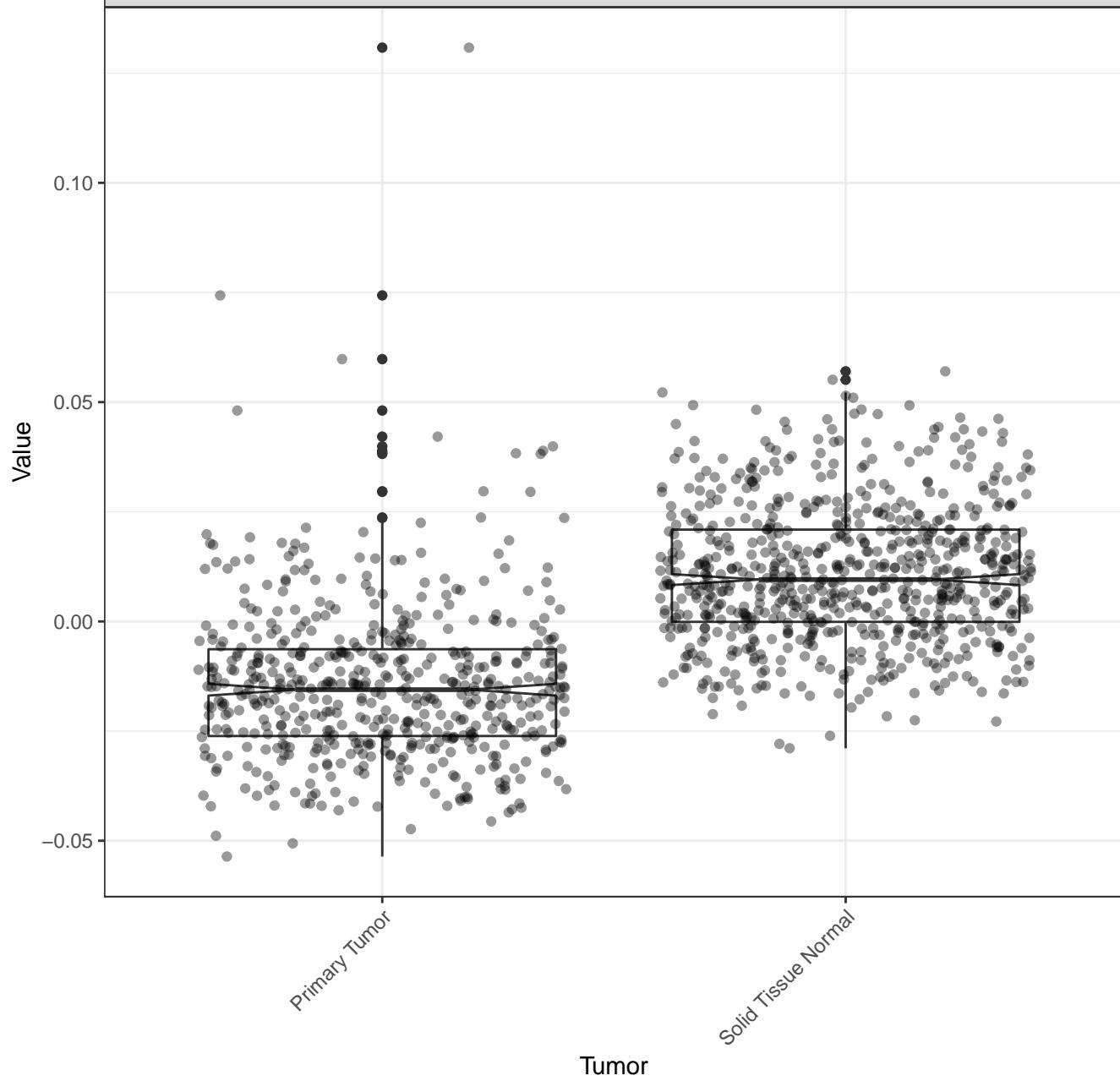
Primary Tumor

Tumor

Solid Tissue Normal



275,REACTOME_METABOLISM_OF_PROTEINS



LV 276

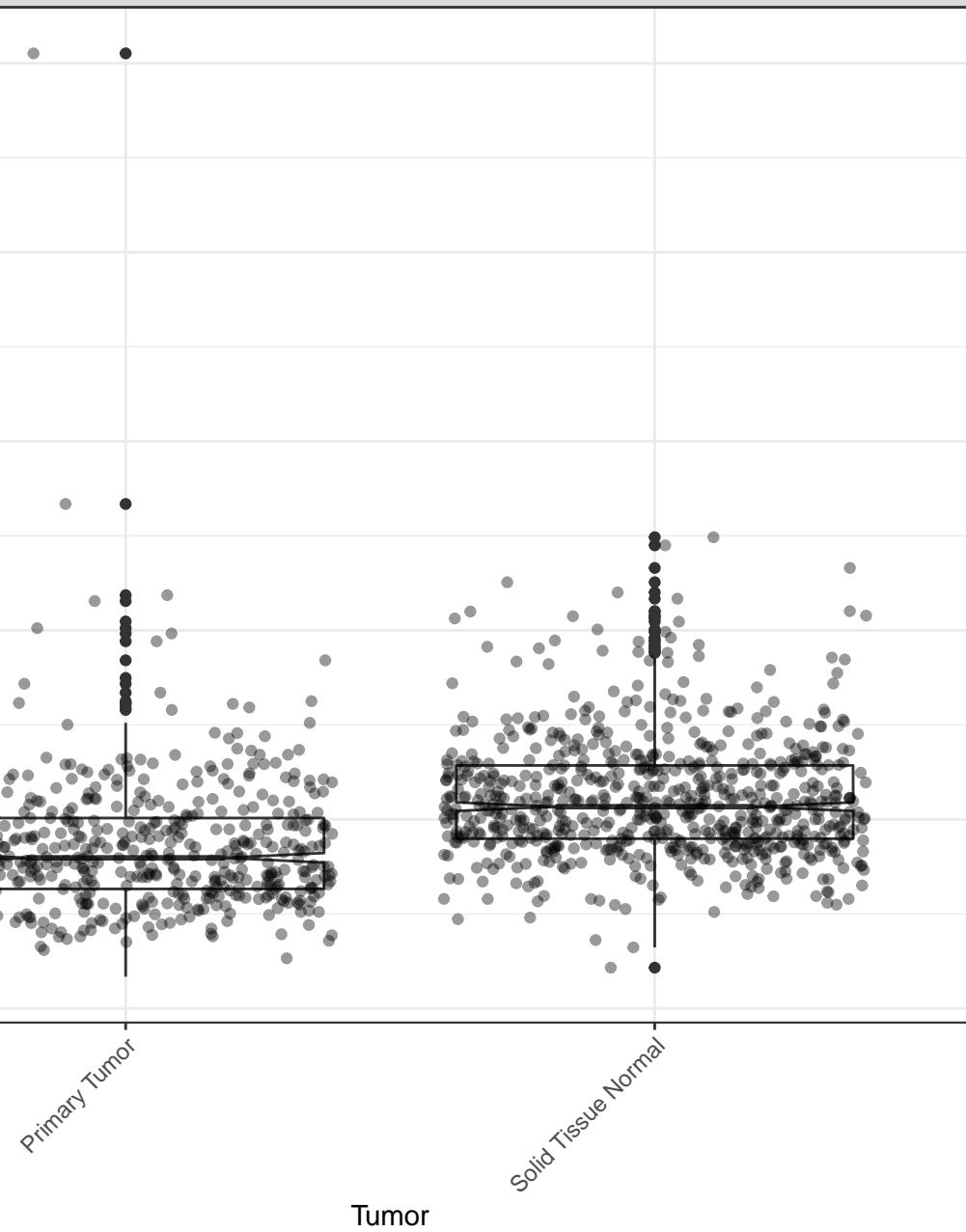
Value

0.20
0.15
0.10
0.05
0.00
-0.05

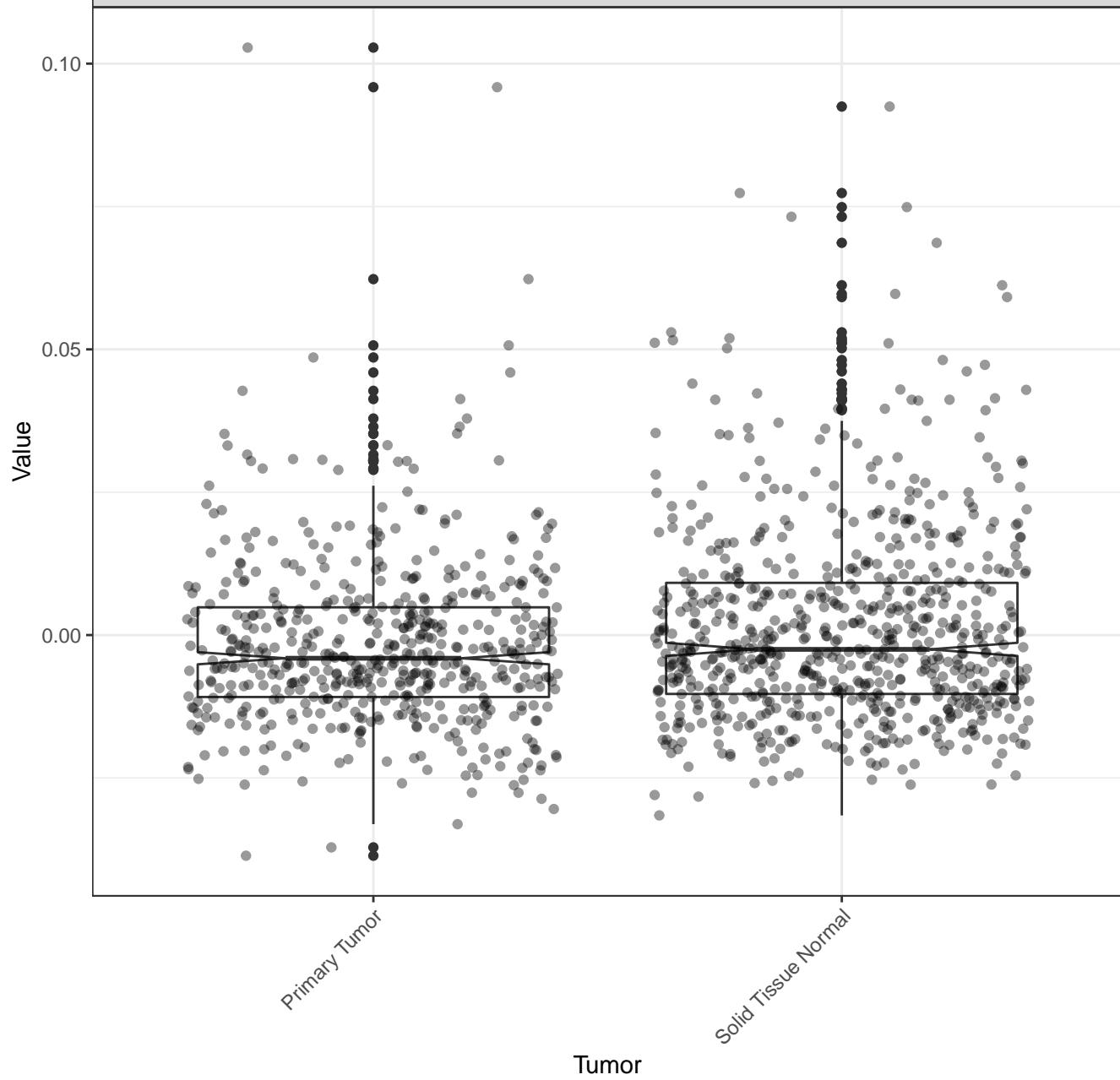
Primary Tumor

Solid Tissue Normal

Tumor



277,PID_IL4_2PATHWAY



278,REACTOME_SLC_MEDIATED_TRANSMEMBRANE_TRANSPORT

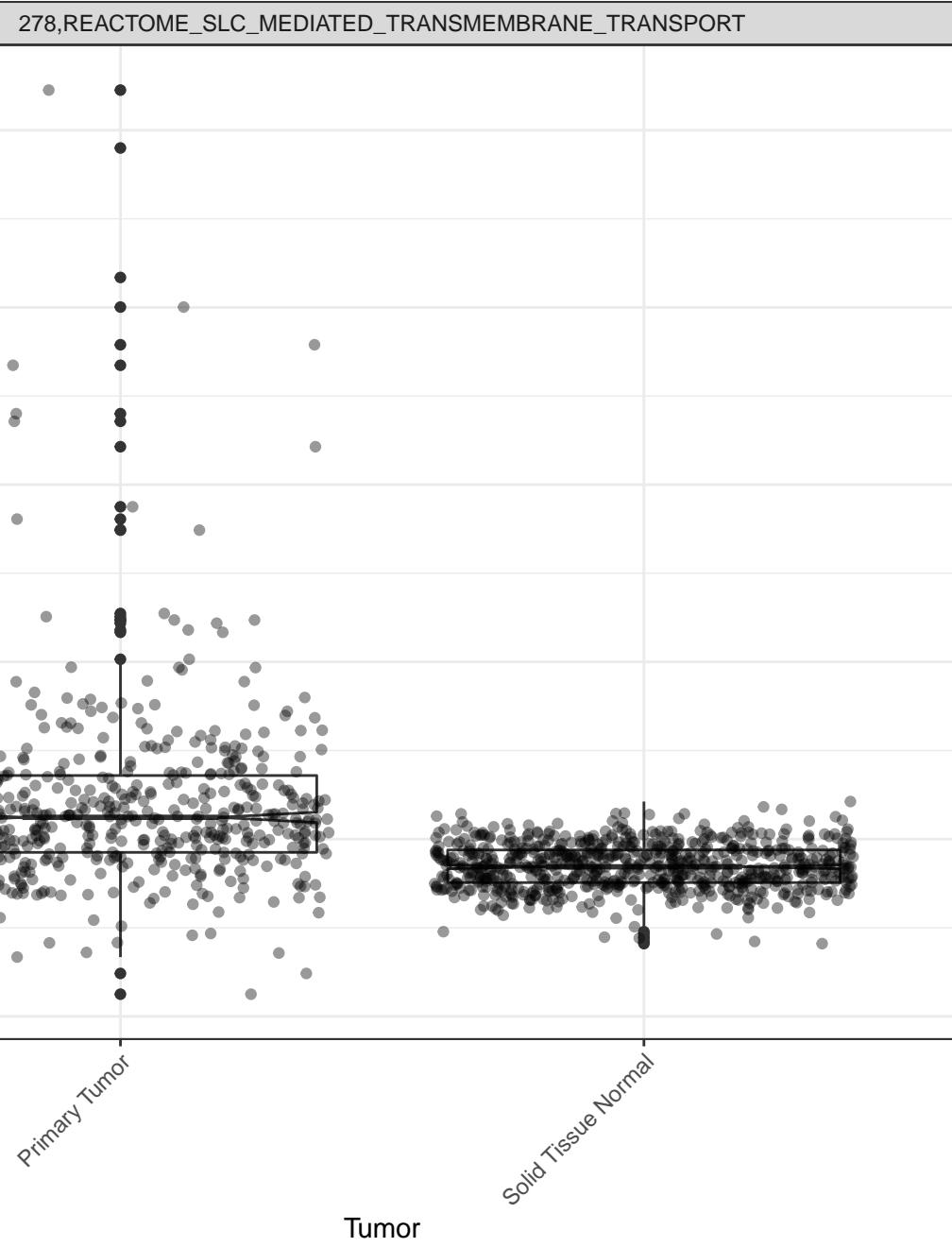
Value

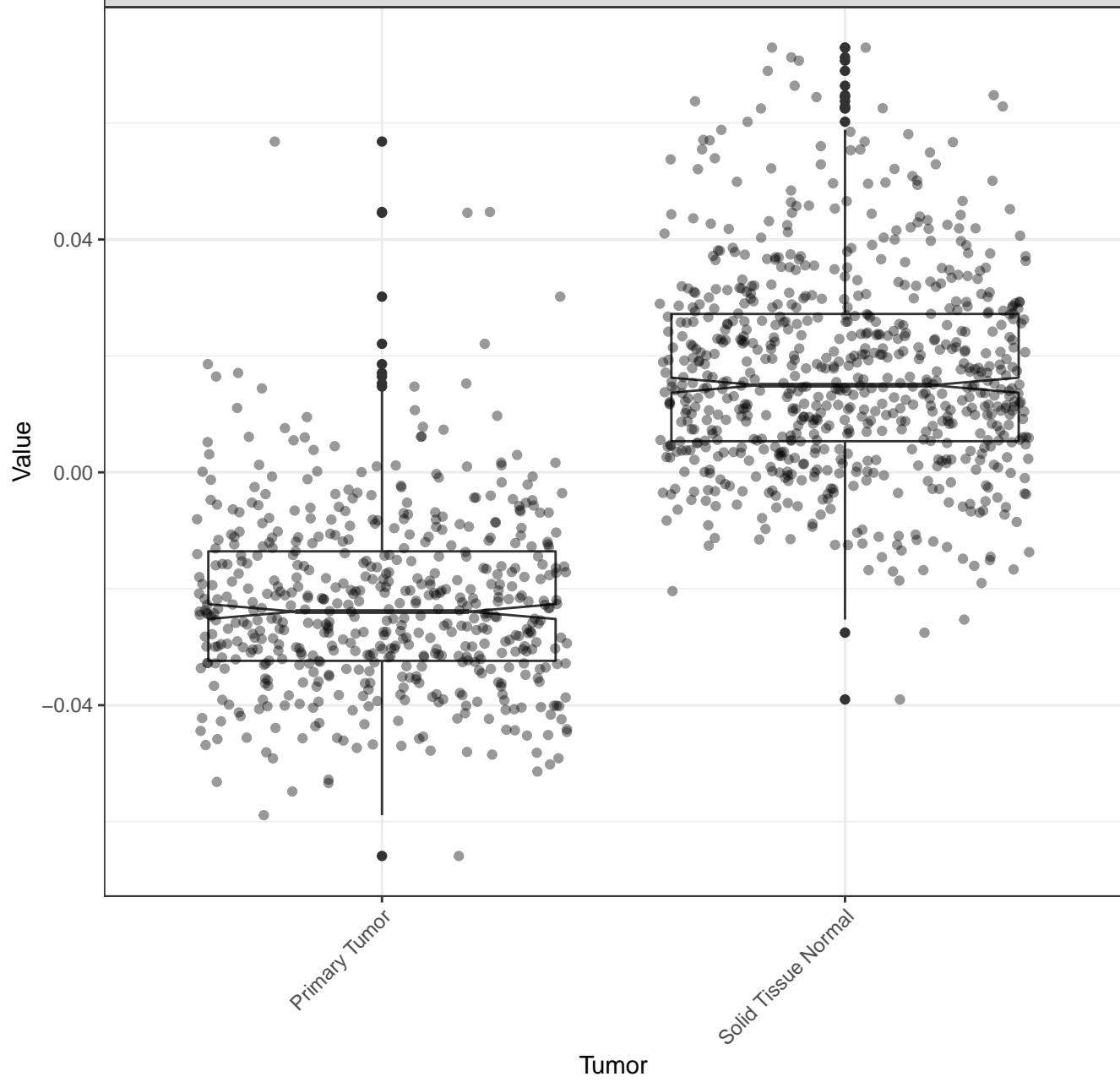
0.4
0.3
0.2
0.1
0.0
-0.1

Primary Tumor

Solid Tissue Normal

Tumor





LV 280

Value

0.10

0.05

0.00

-0.05

Primary Tumor

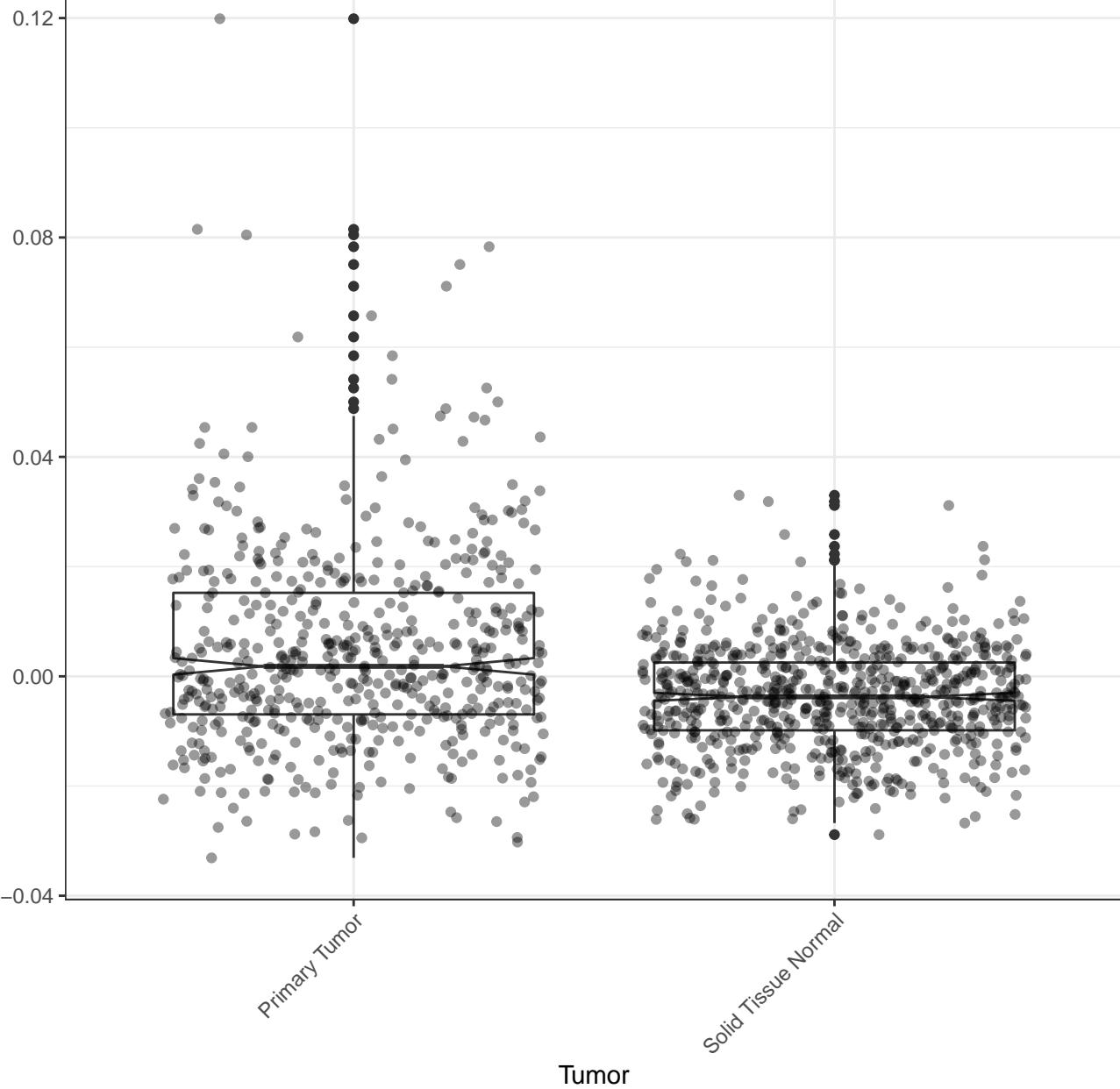
Solid Tissue Normal

Tumor



281,REACTOME_METABOLISM_OF_AMINO_ACIDS_AND_DERIVATIVES

Value



282,KEGG_RNA_DEGRADATION

Value

0.10

0.05

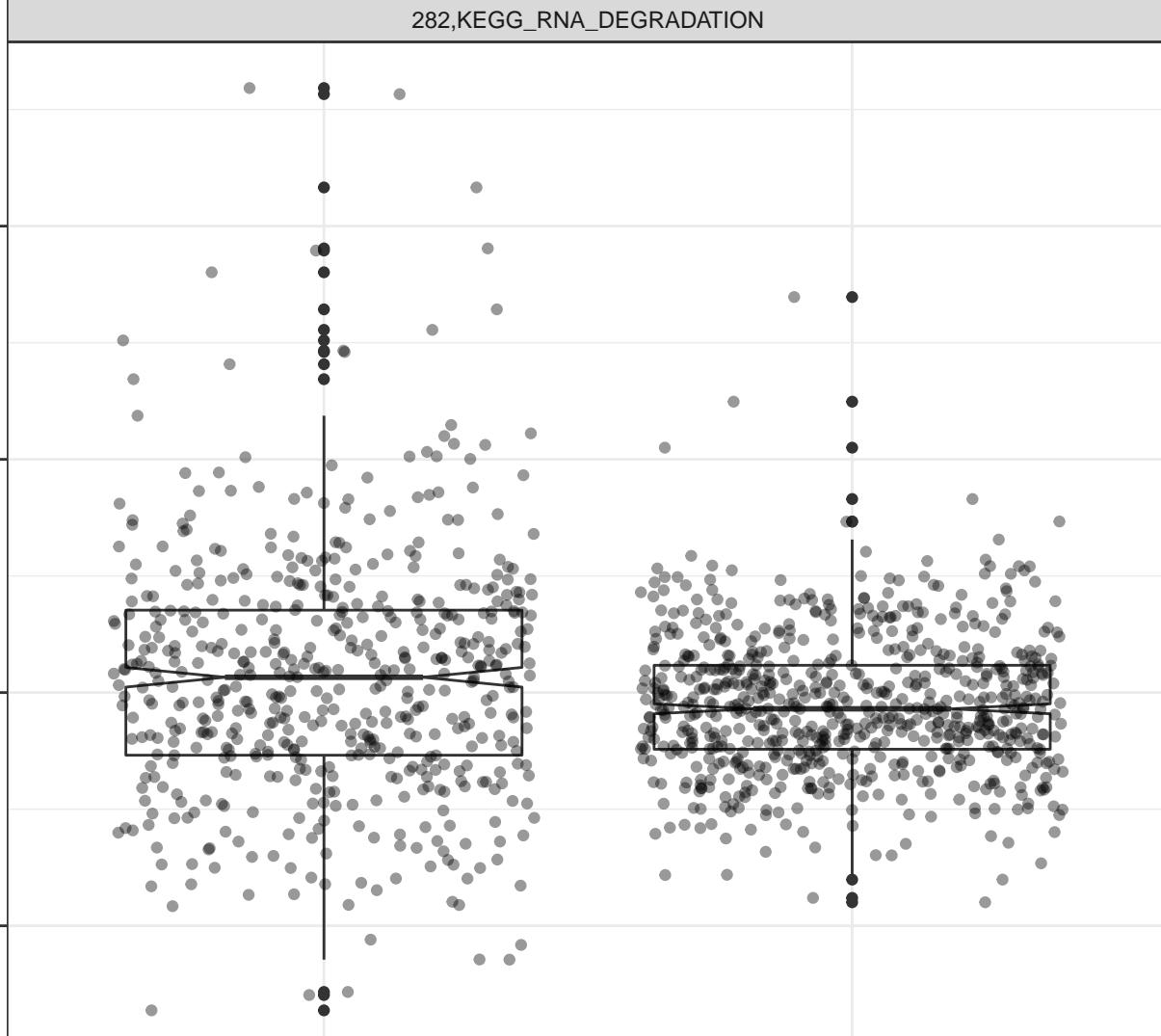
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



283,IRIS_Monocyte–Day0

Value

0.05

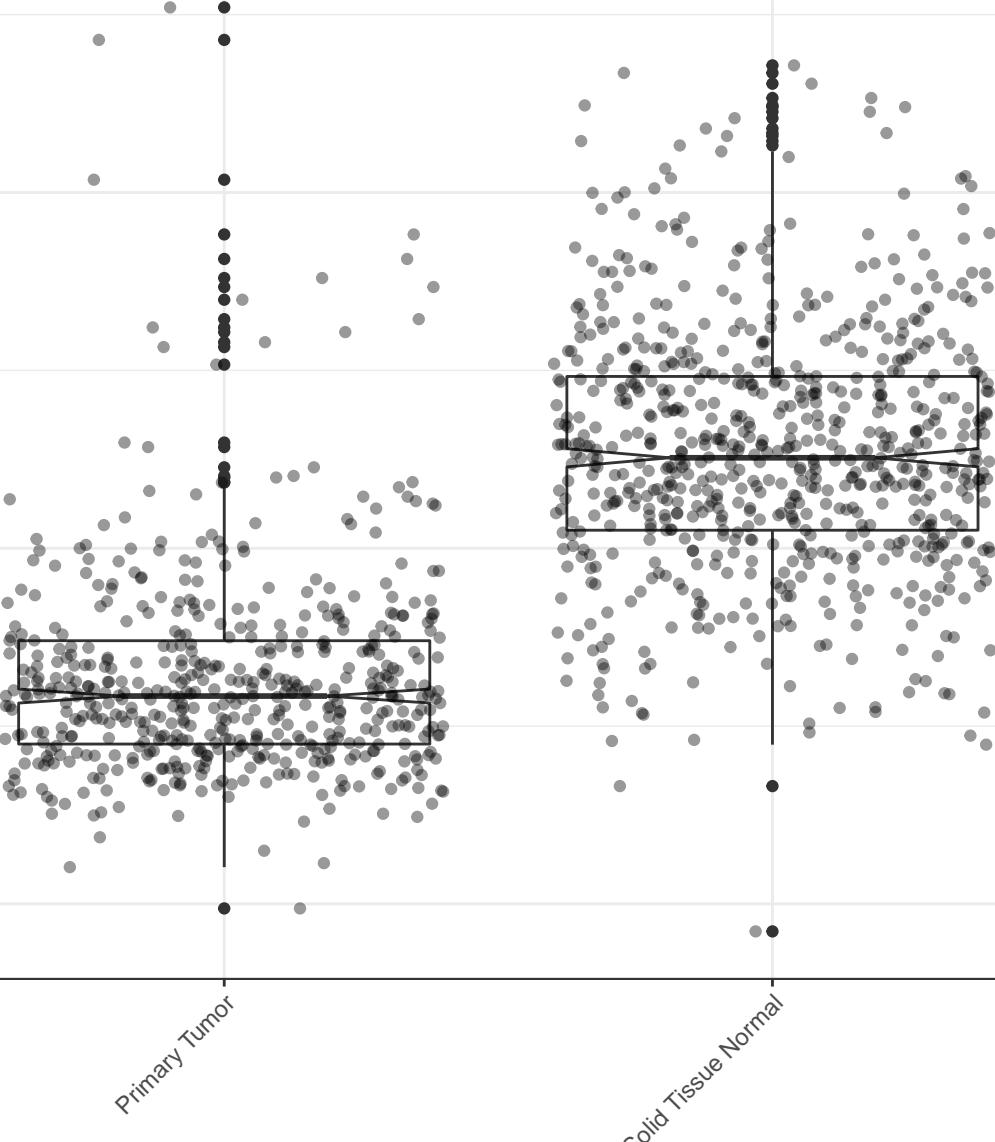
0.00

-0.05

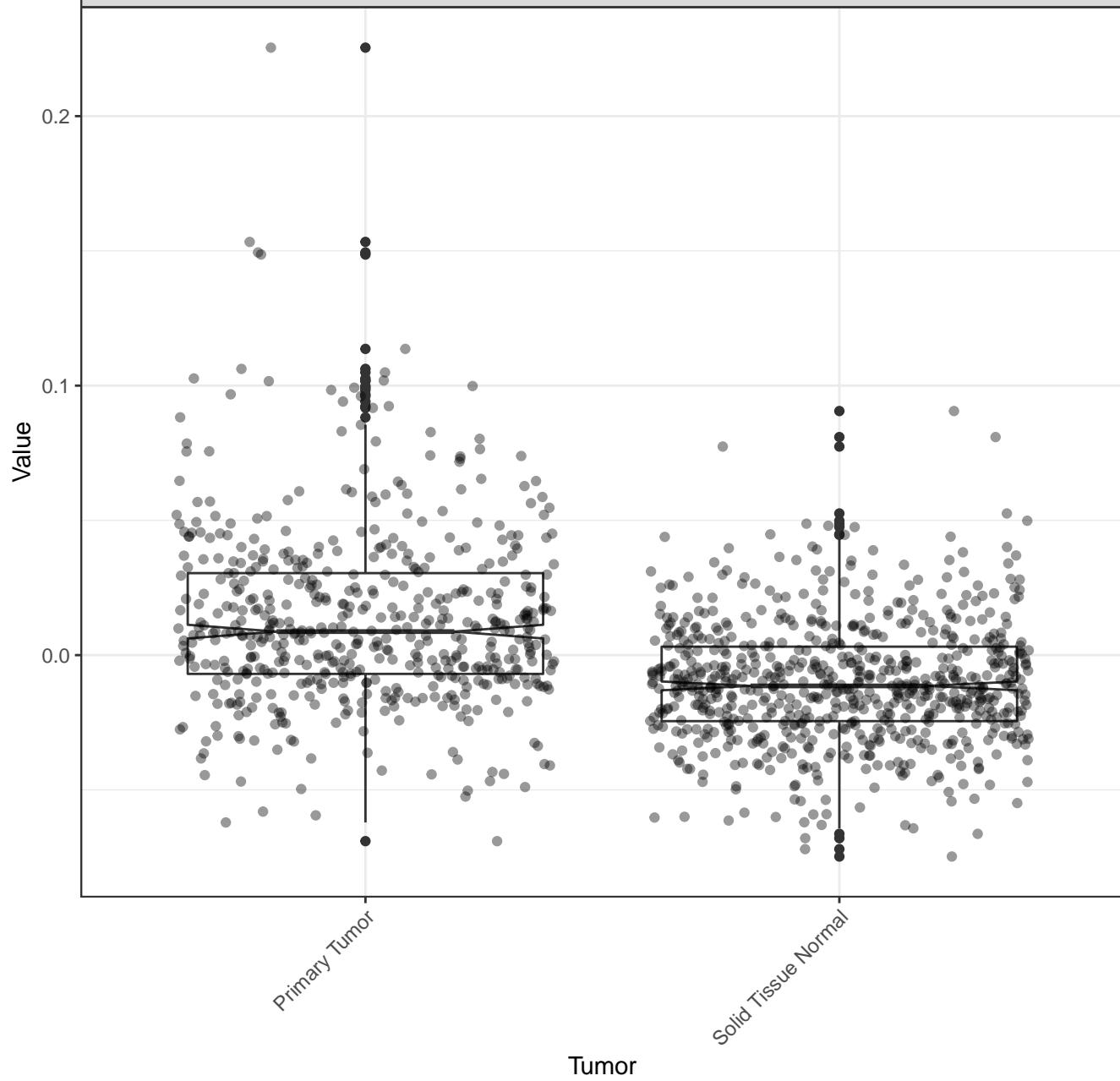
Primary Tumor

Solid Tissue Normal

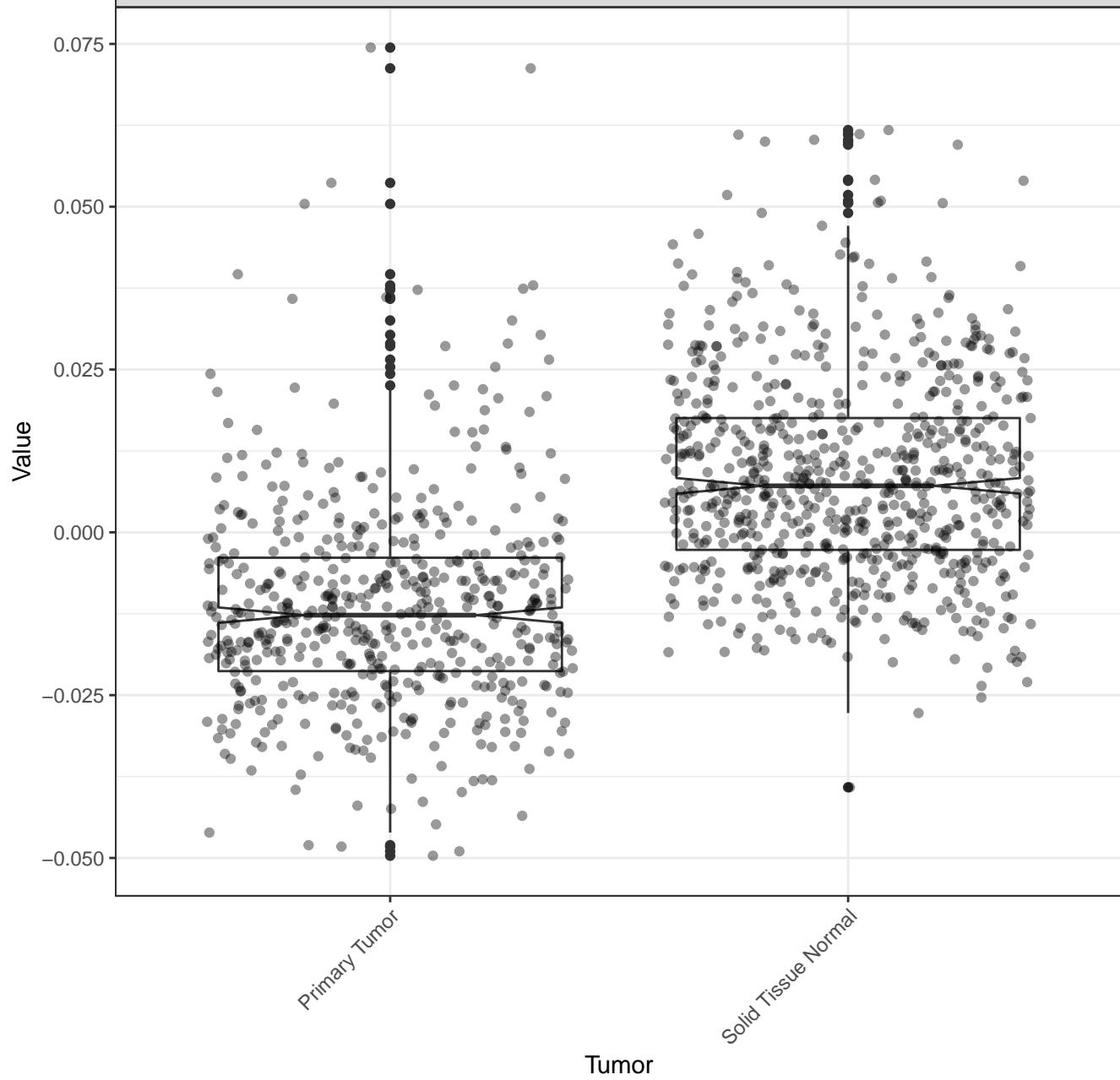
Tumor



284,KEGG_ECM_RECEPтор_INTERACTION



285,KEGG_SNARE_INTERACTIONS_IN_VESICULAR_TRANSPORT



LV 286

Value

0.12

0.08

0.04

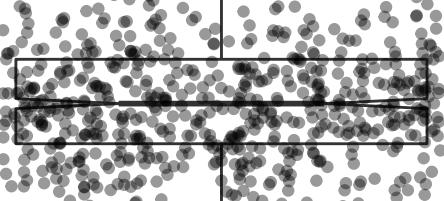
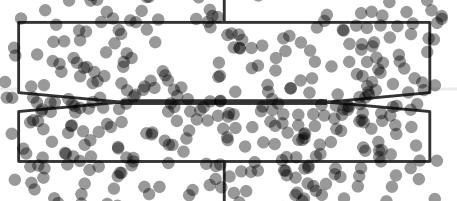
0.00

-0.04

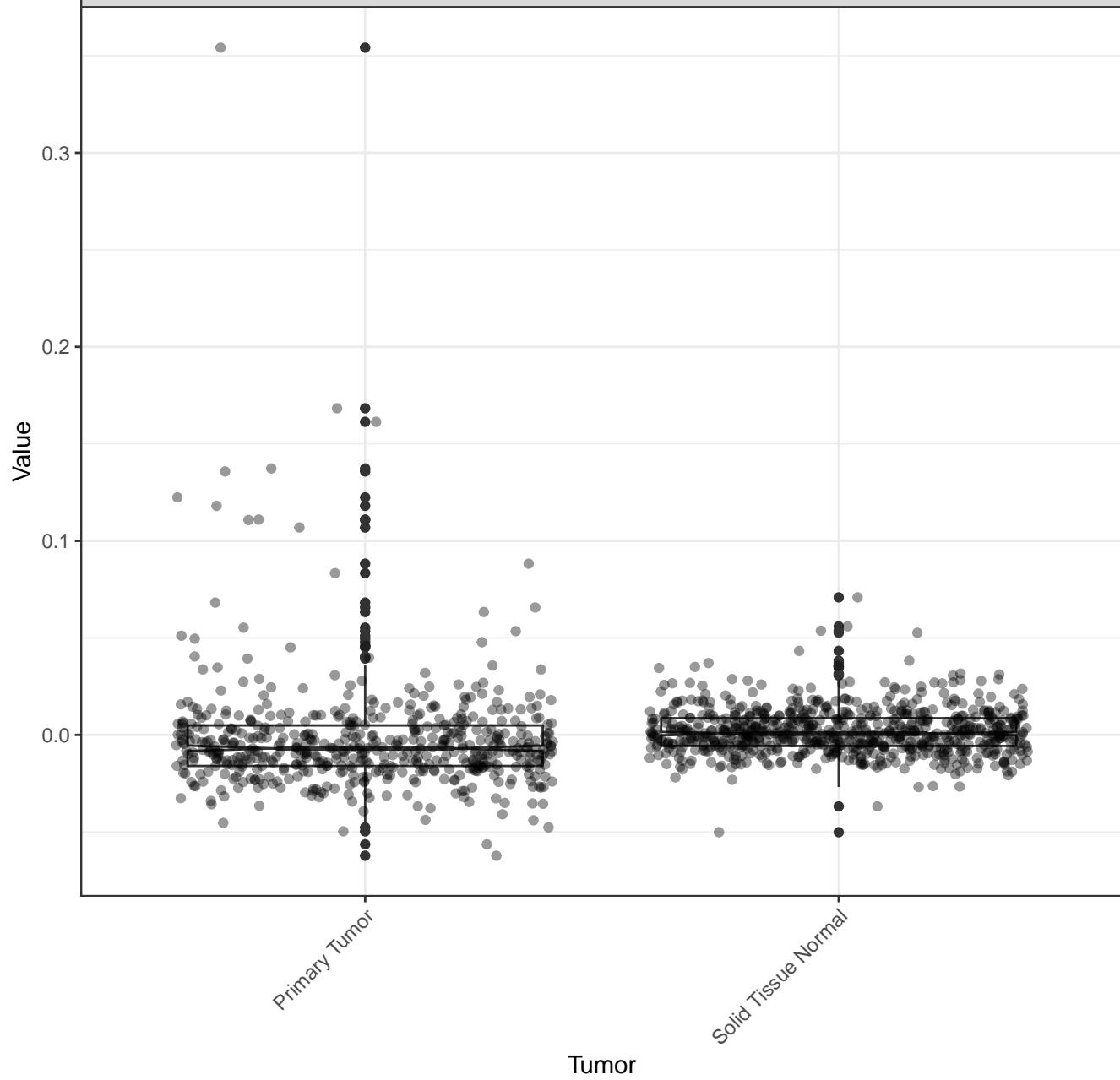
Primary Tumor

Solid Tissue Normal

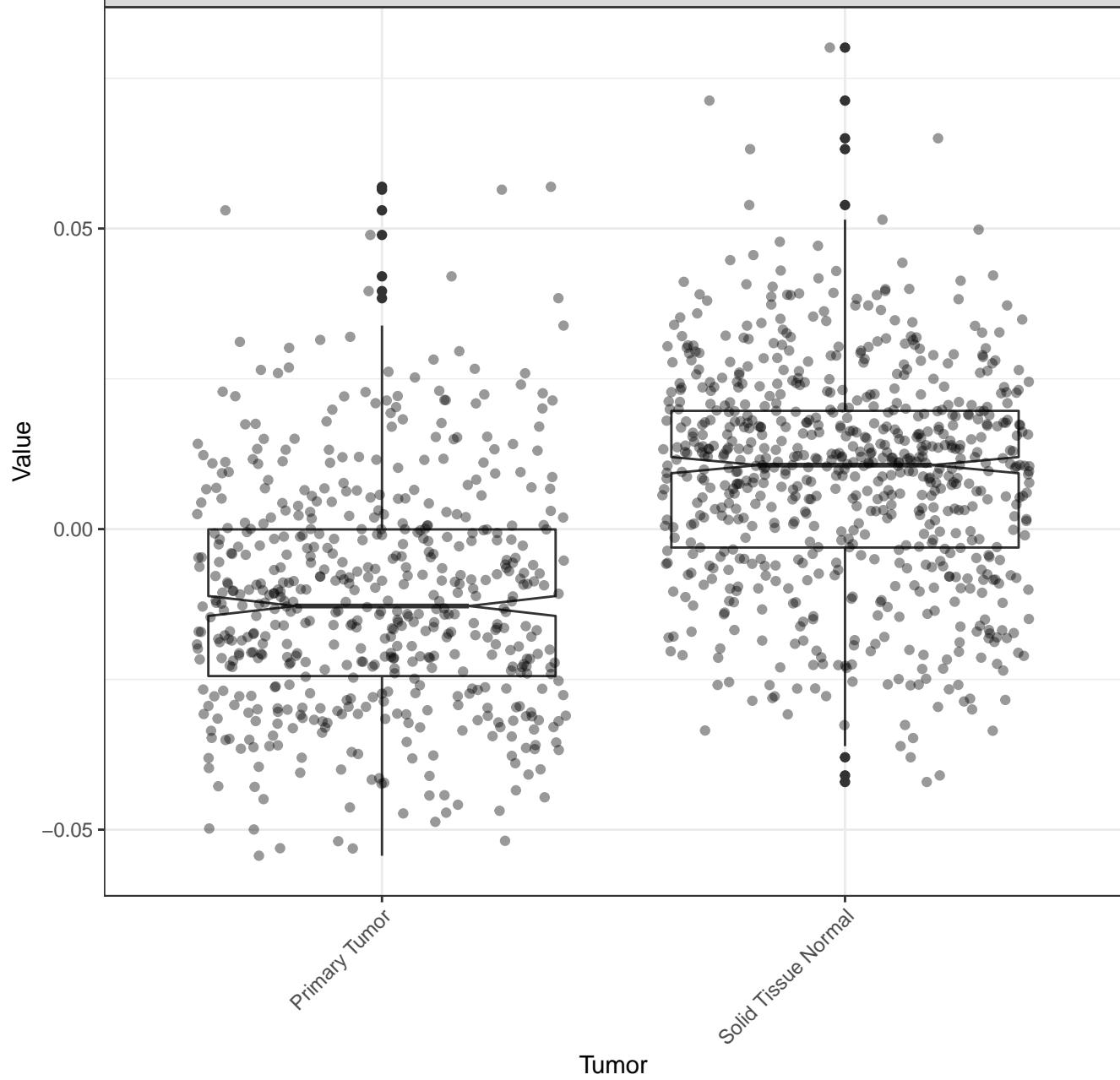
Tumor



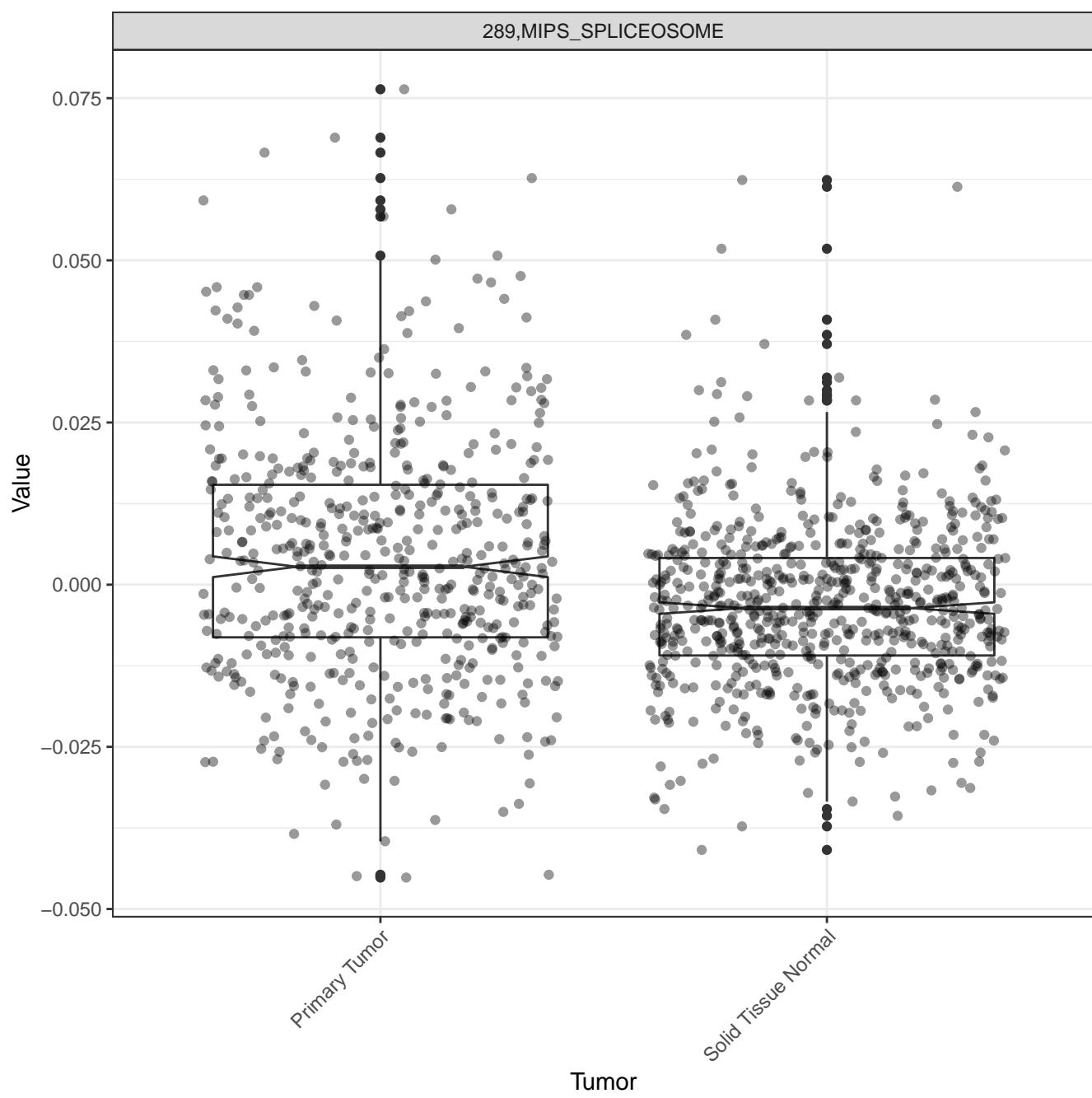
287,PID_RET_PATHWAY



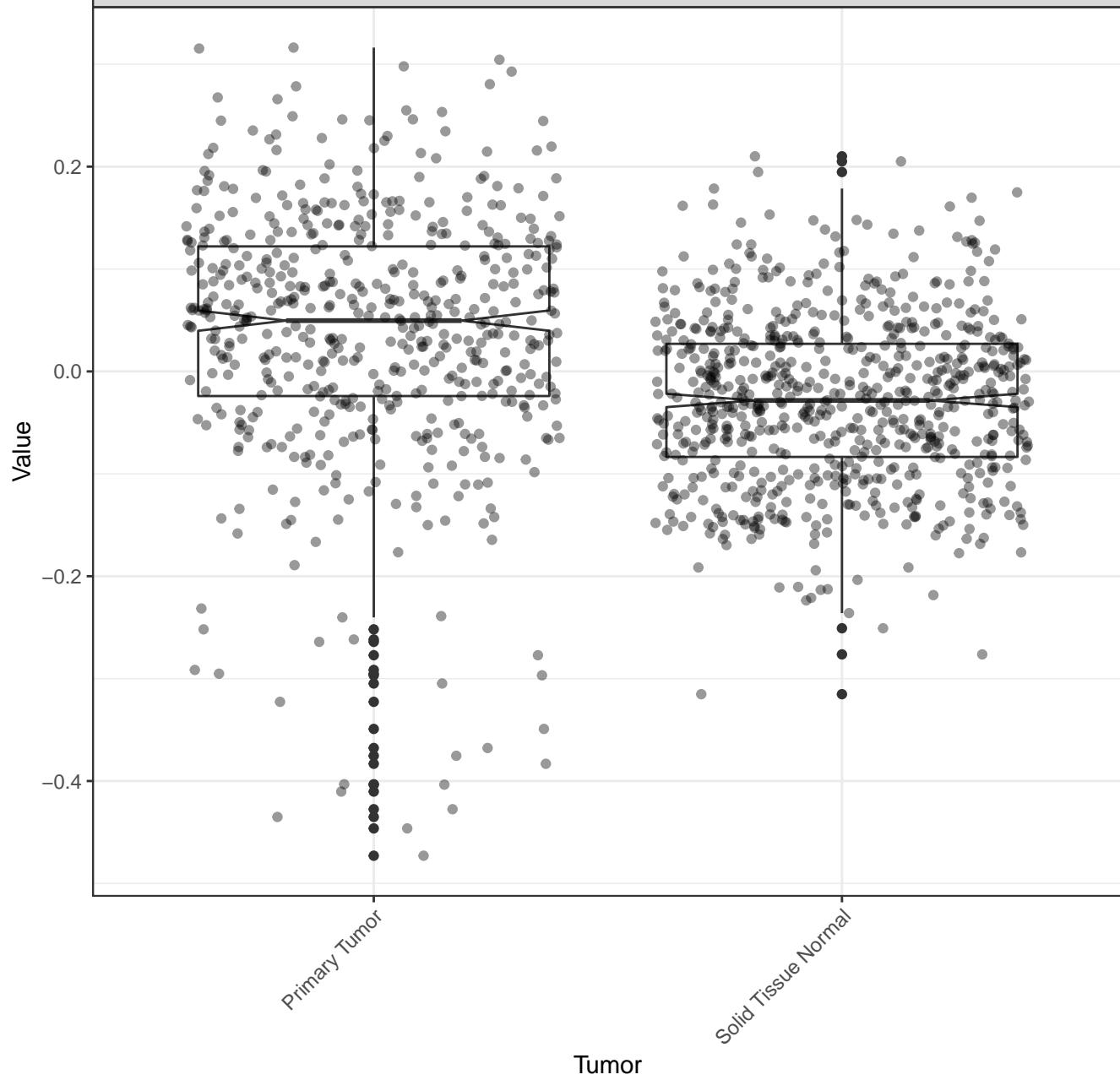
288,KEGG_PURINE_METABOLISM



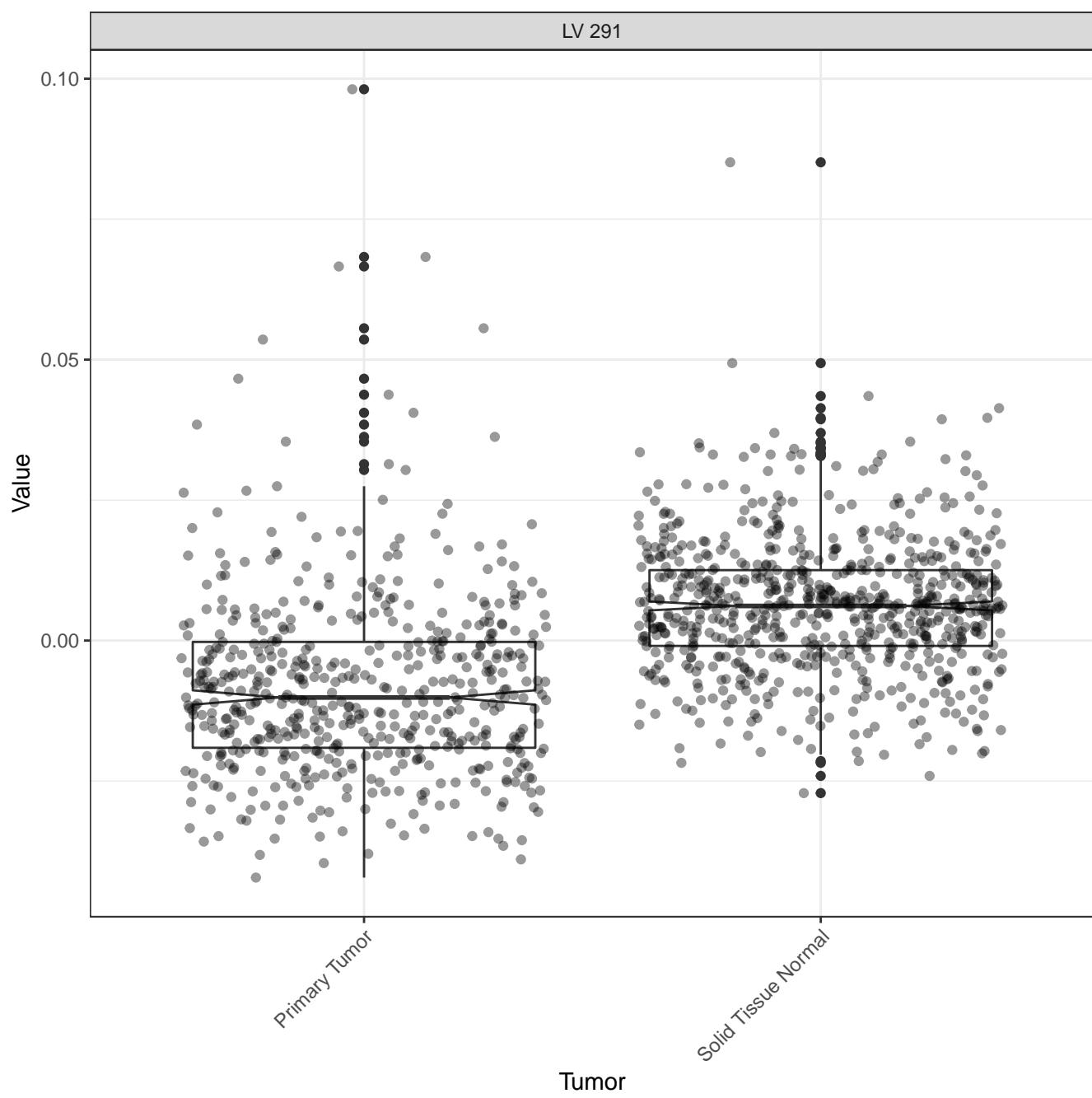
289,MIPS_SPLICEOSOME



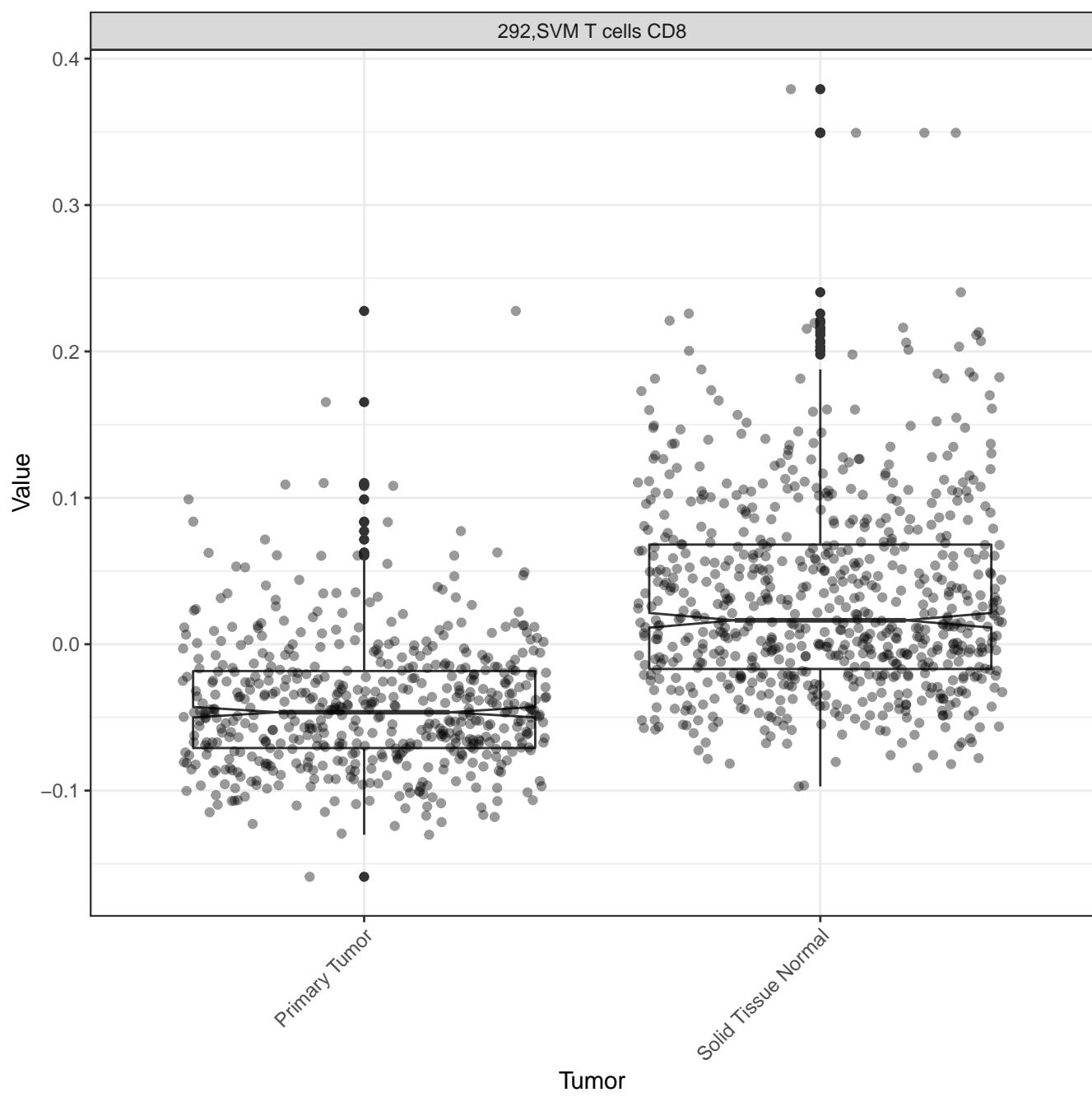
290,REACTOME_TRNA_AMINOACYLATION



LV 291



292,SVM T cells CD8



LV 293

Value

0.10

0.05

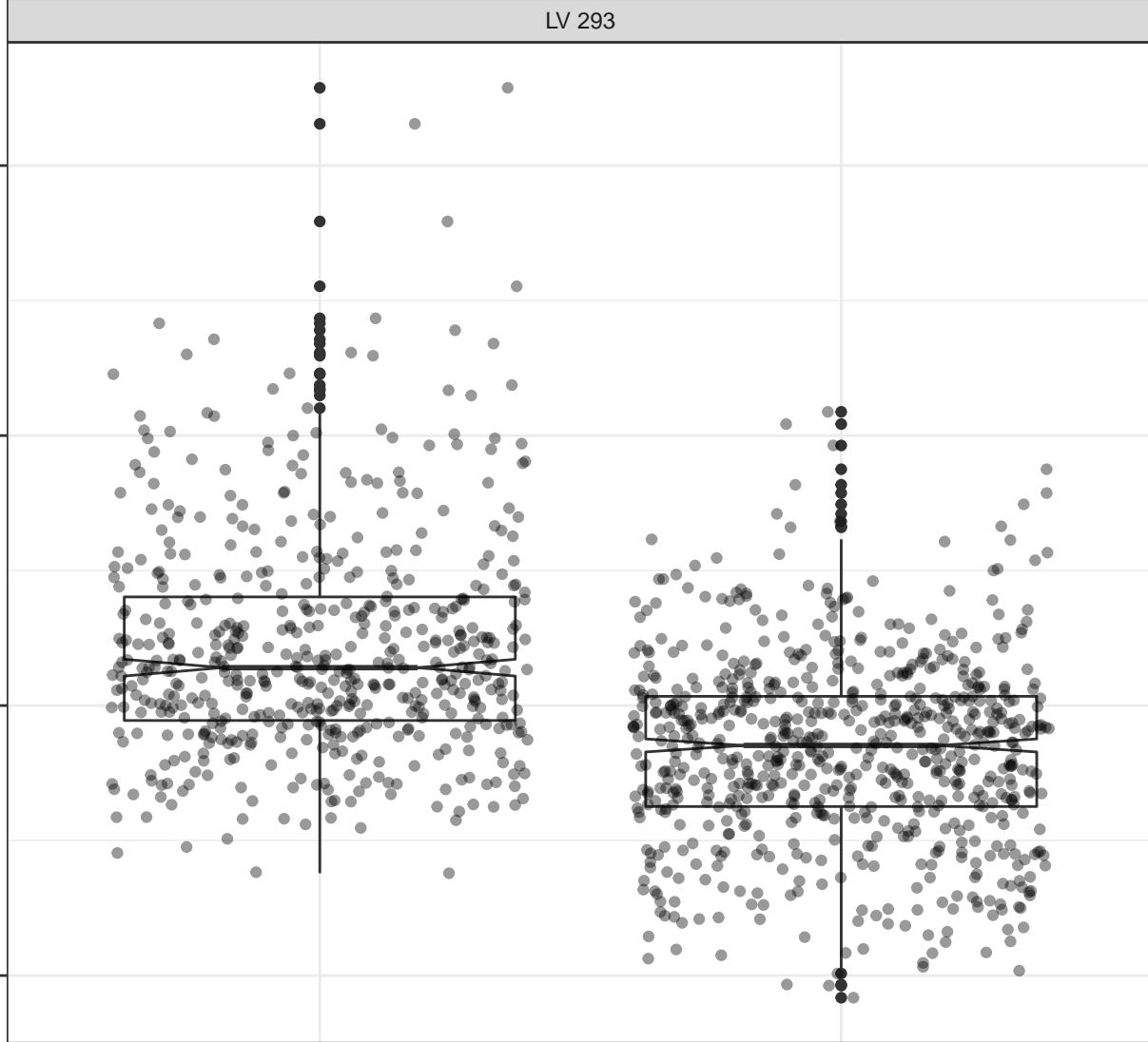
0.00

-0.05

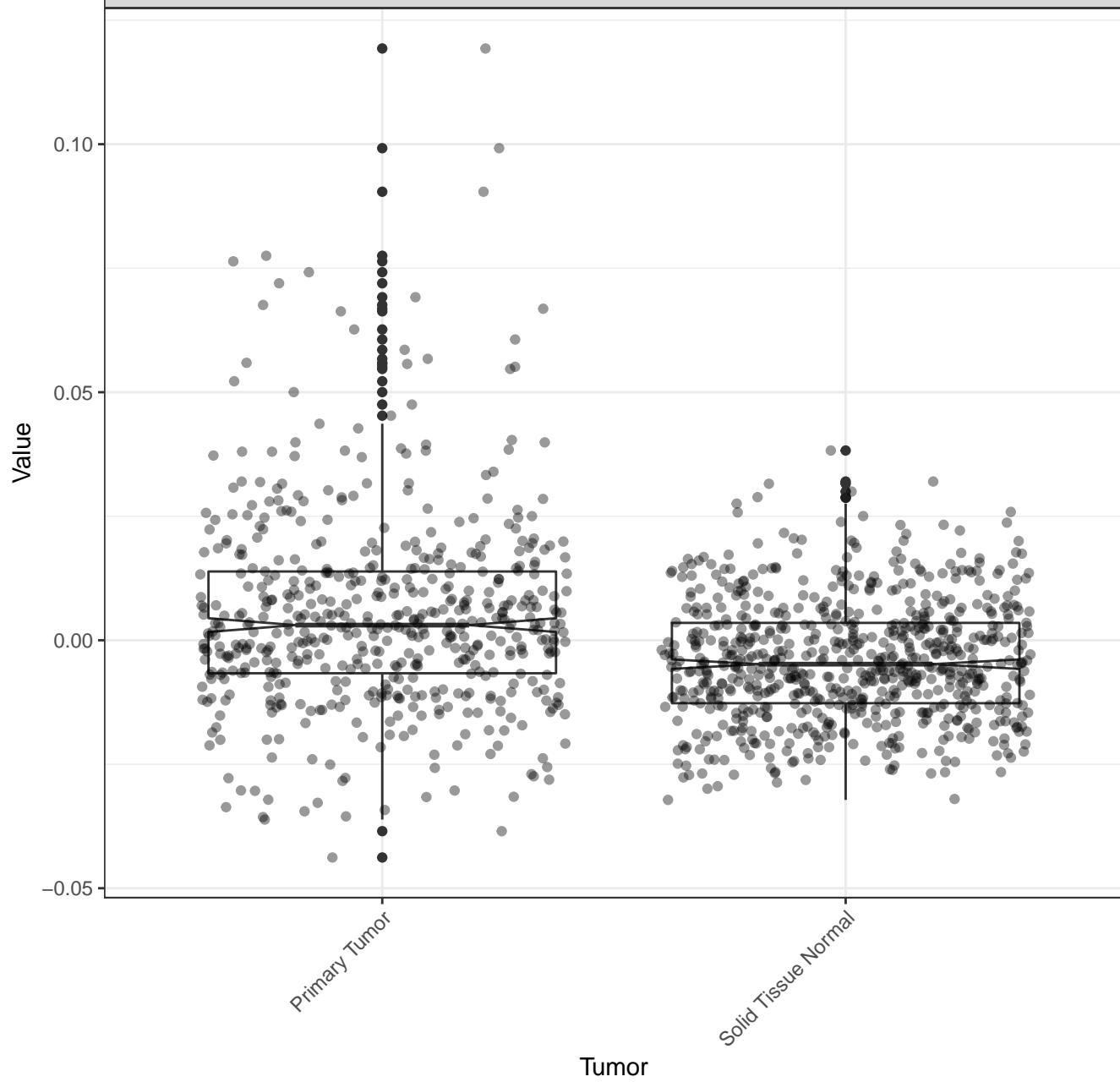
Primary Tumor

Solid Tissue Normal

Tumor



294,REACTOME_METABOLISM_OF_PROTEINS



295,PID_NFAT_TFPATHWAY

Value

0.4

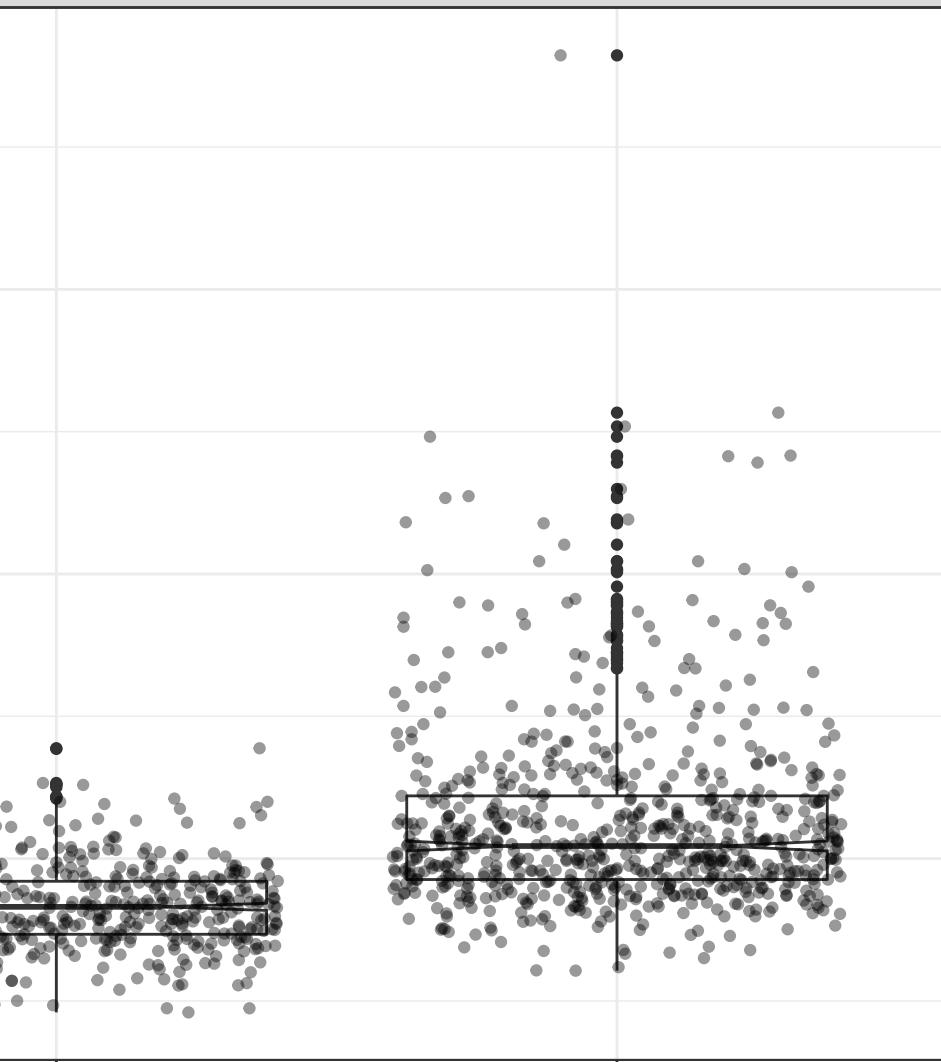
0.2

0.0

Primary Tumor

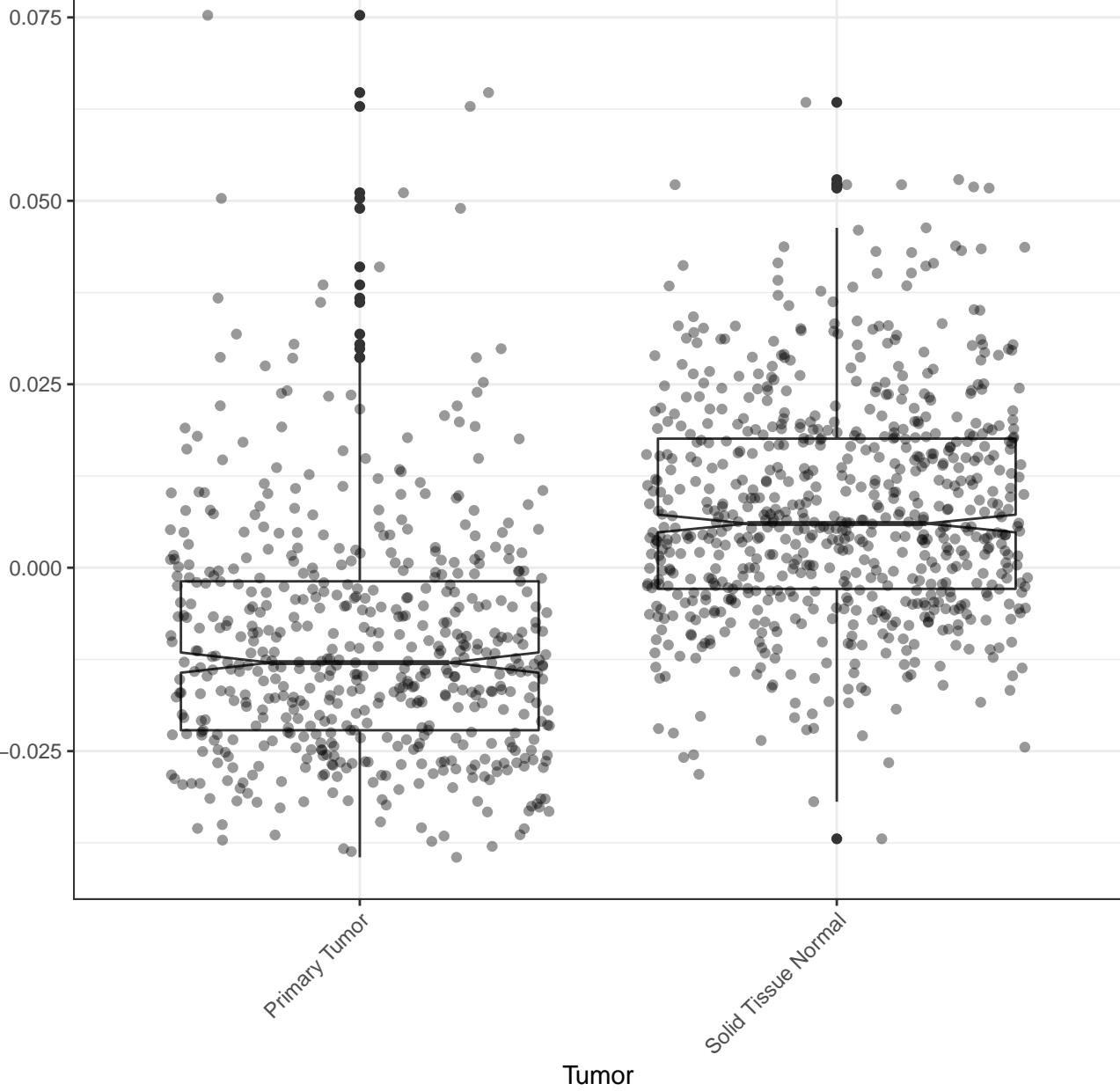
Solid Tissue Normal

Tumor



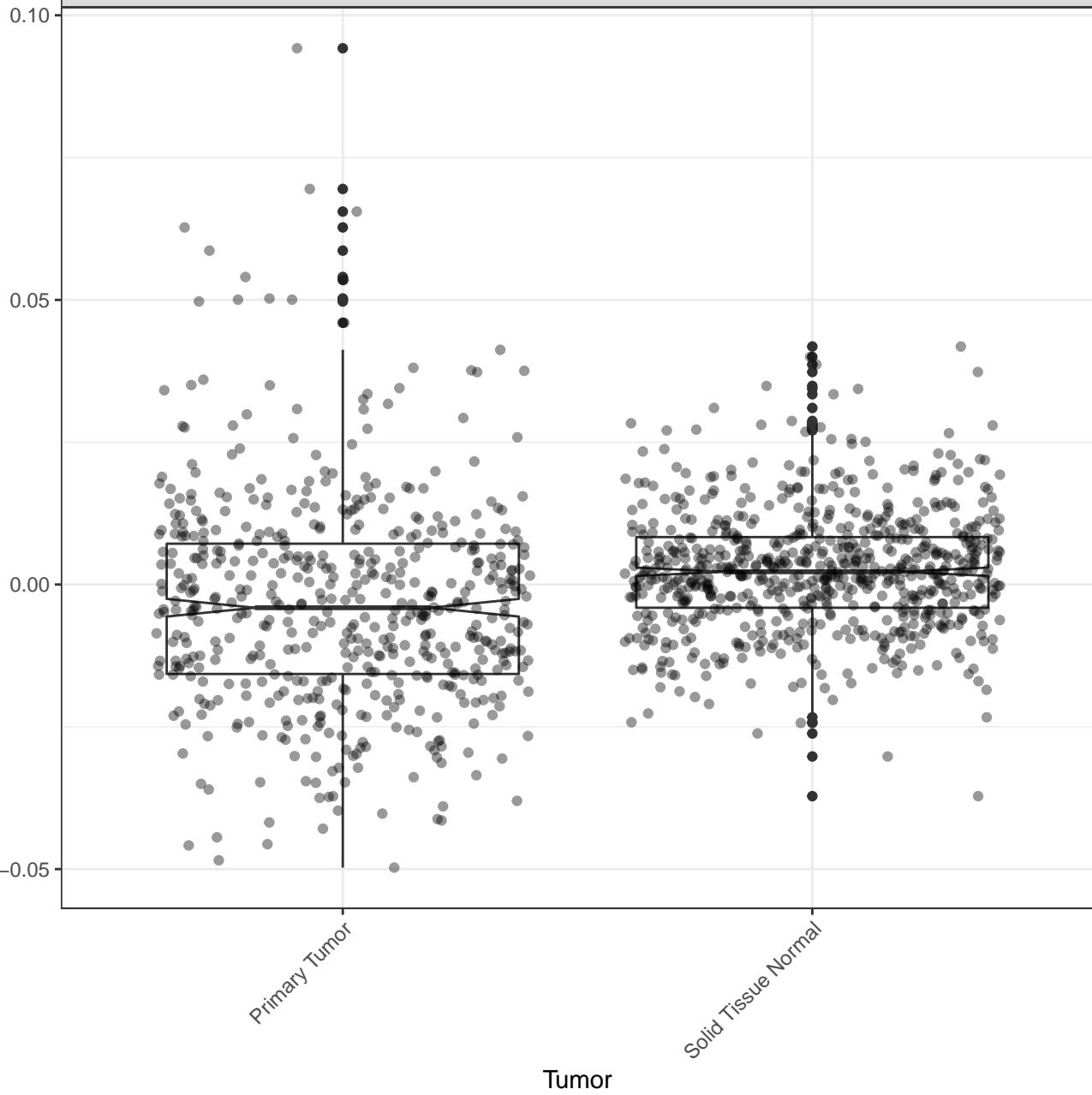
296,REACTOME_ACTIVATED_TLR4_SIGNALLING

Value

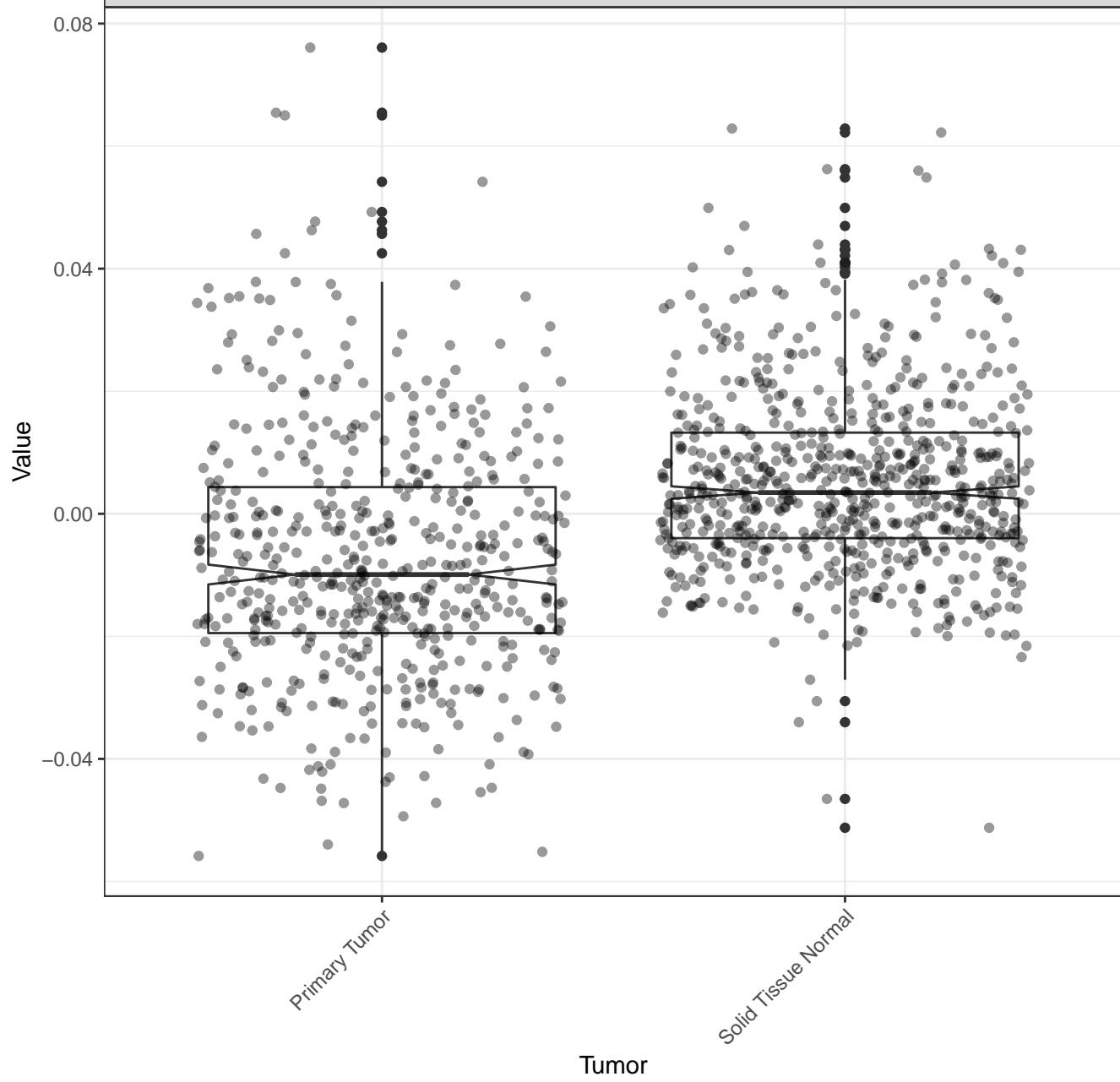


297,PID_MAPKTRKPATHWAY

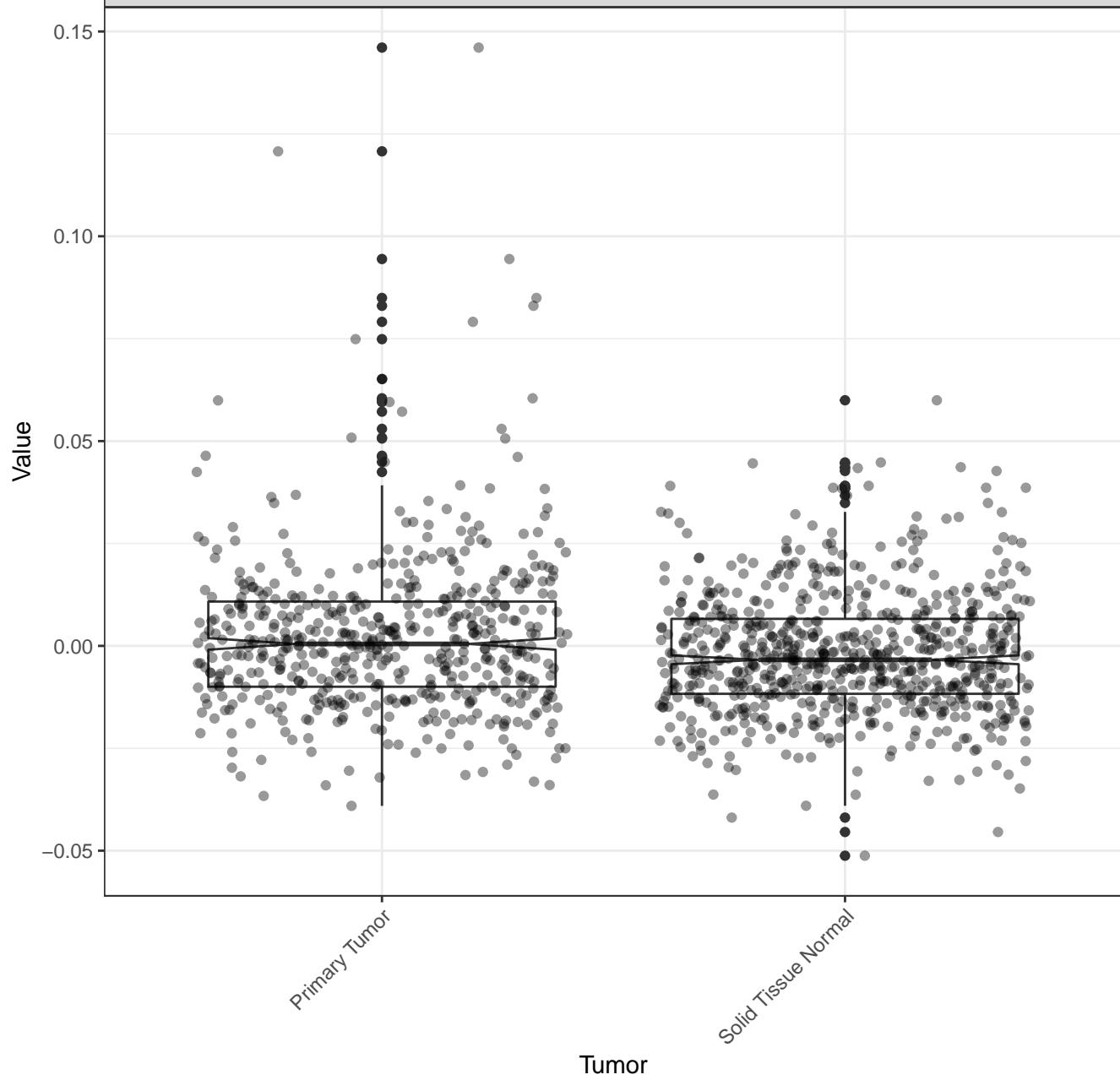
Value



LV 298



299,KEGG_MELANOGENESIS



LV 300

Value

0.10

0.05

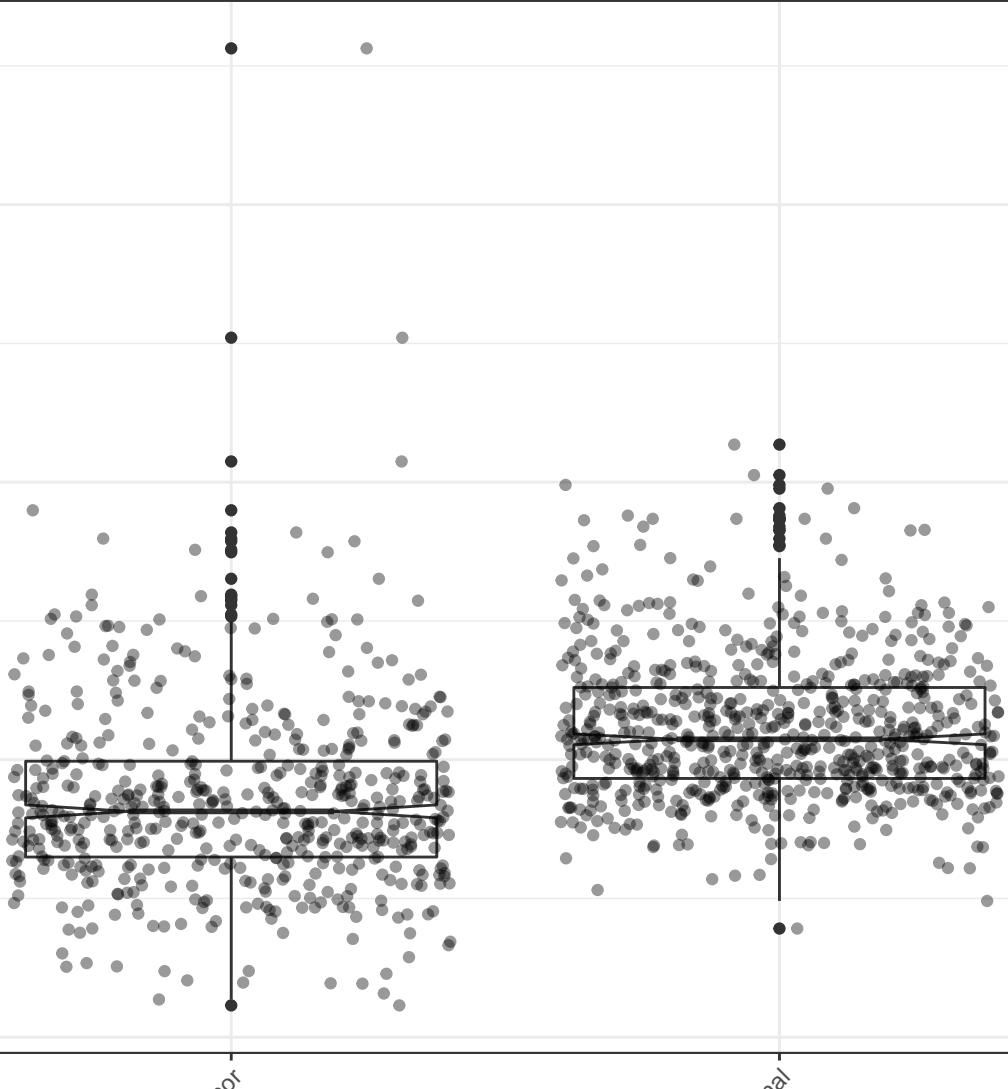
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.10

0.05

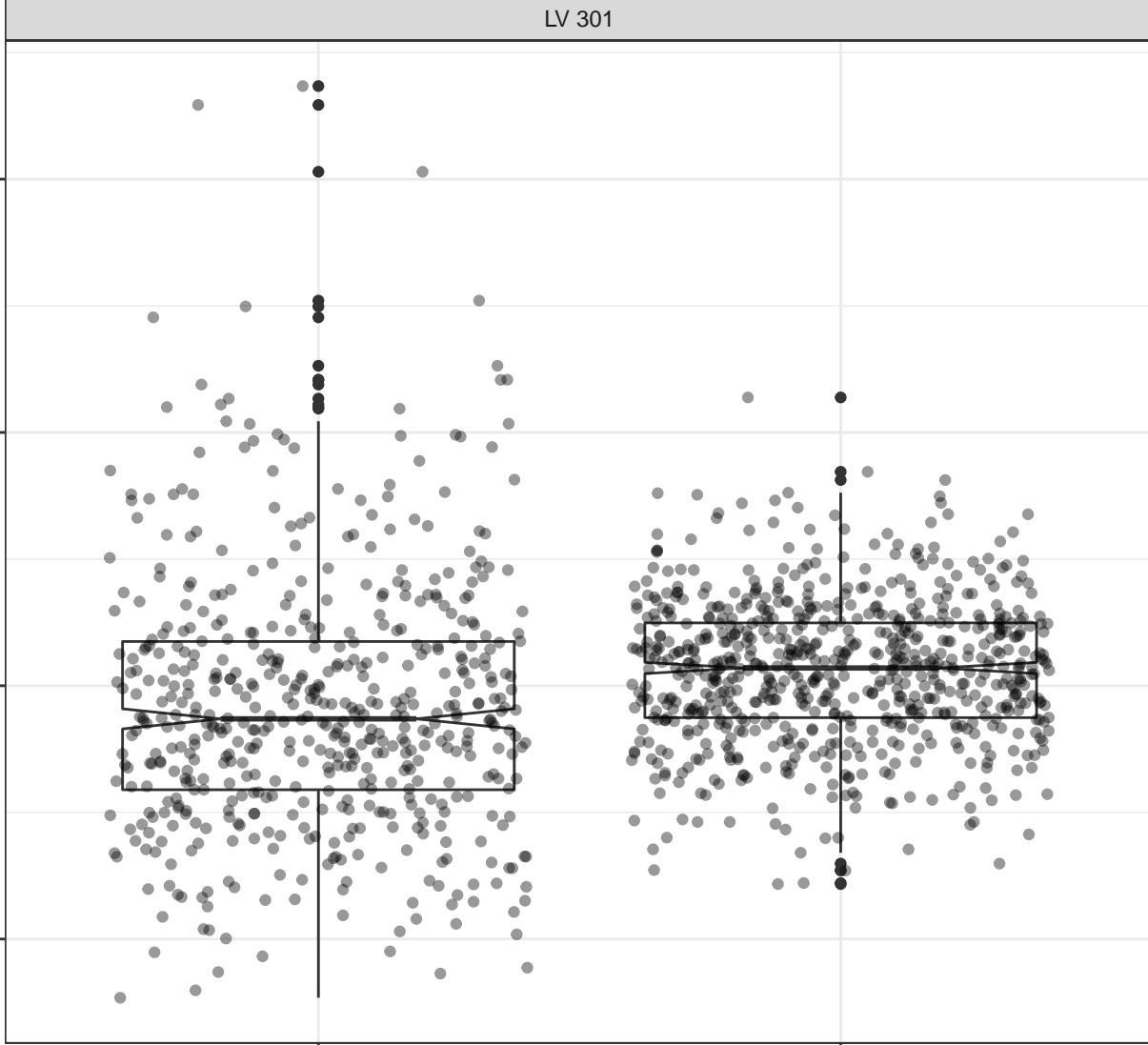
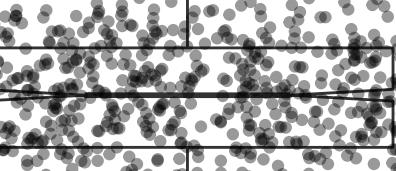
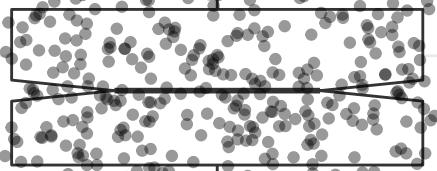
0.00

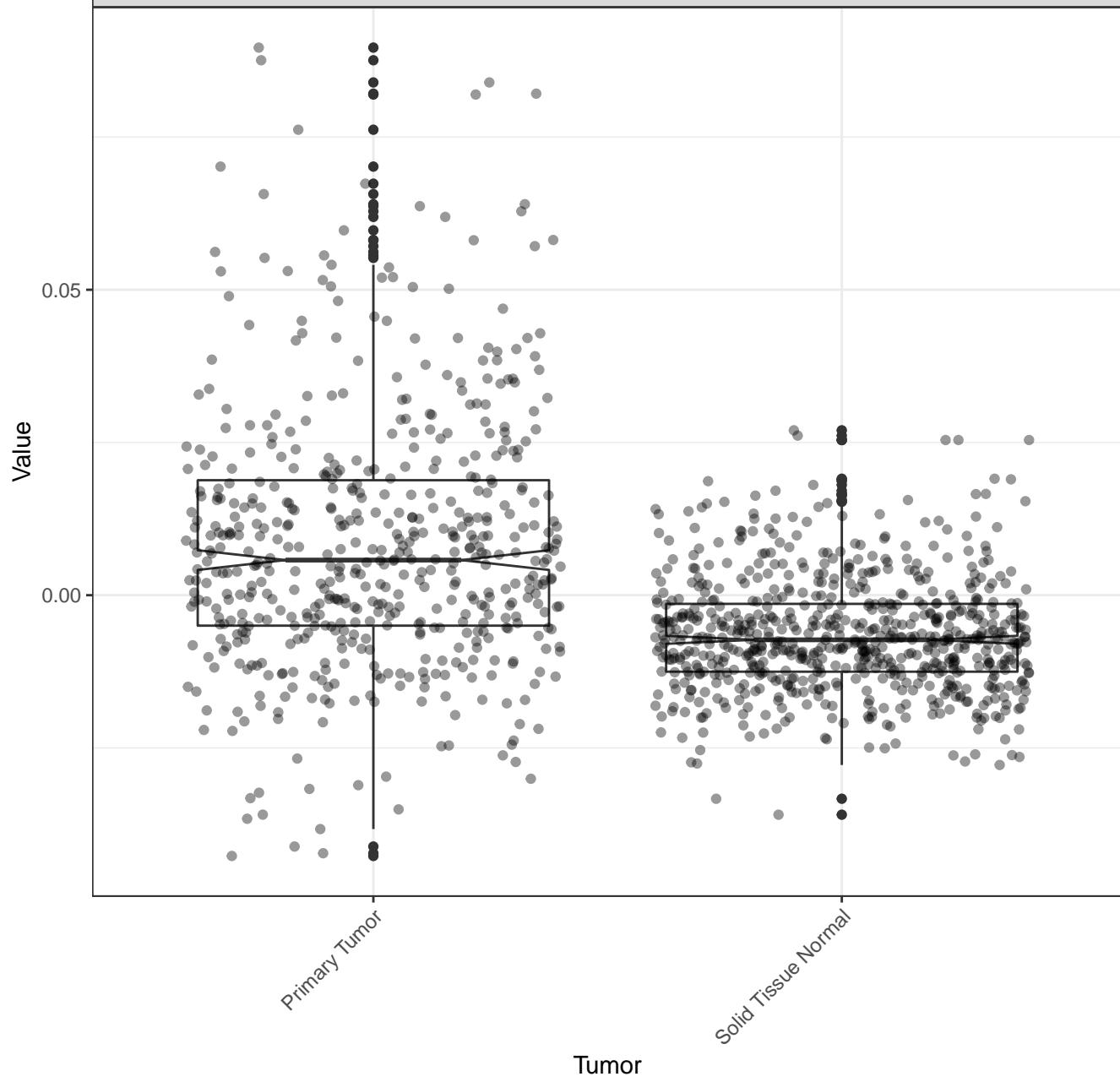
-0.05

Primary Tumor

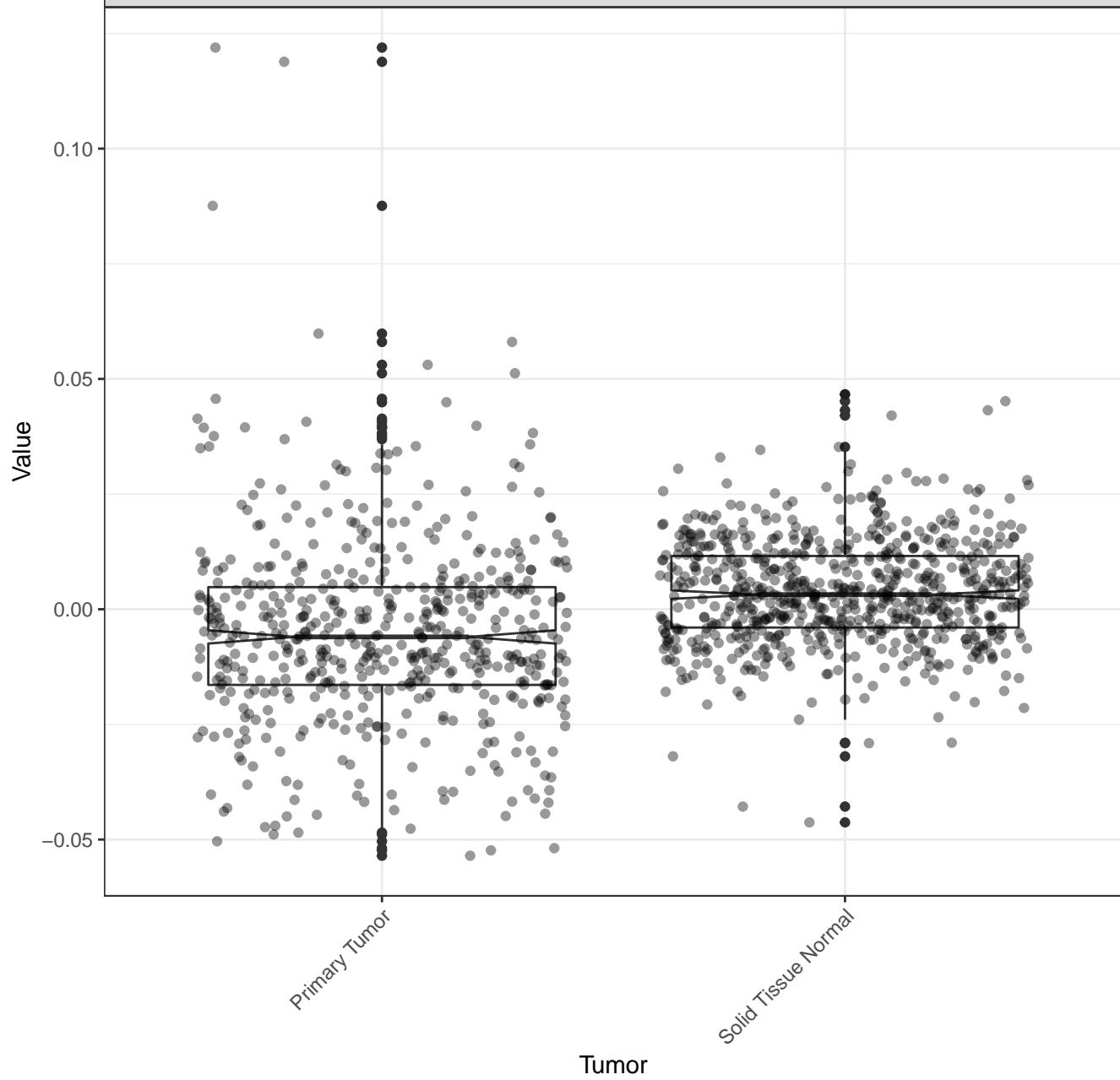
Solid Tissue Normal

Tumor

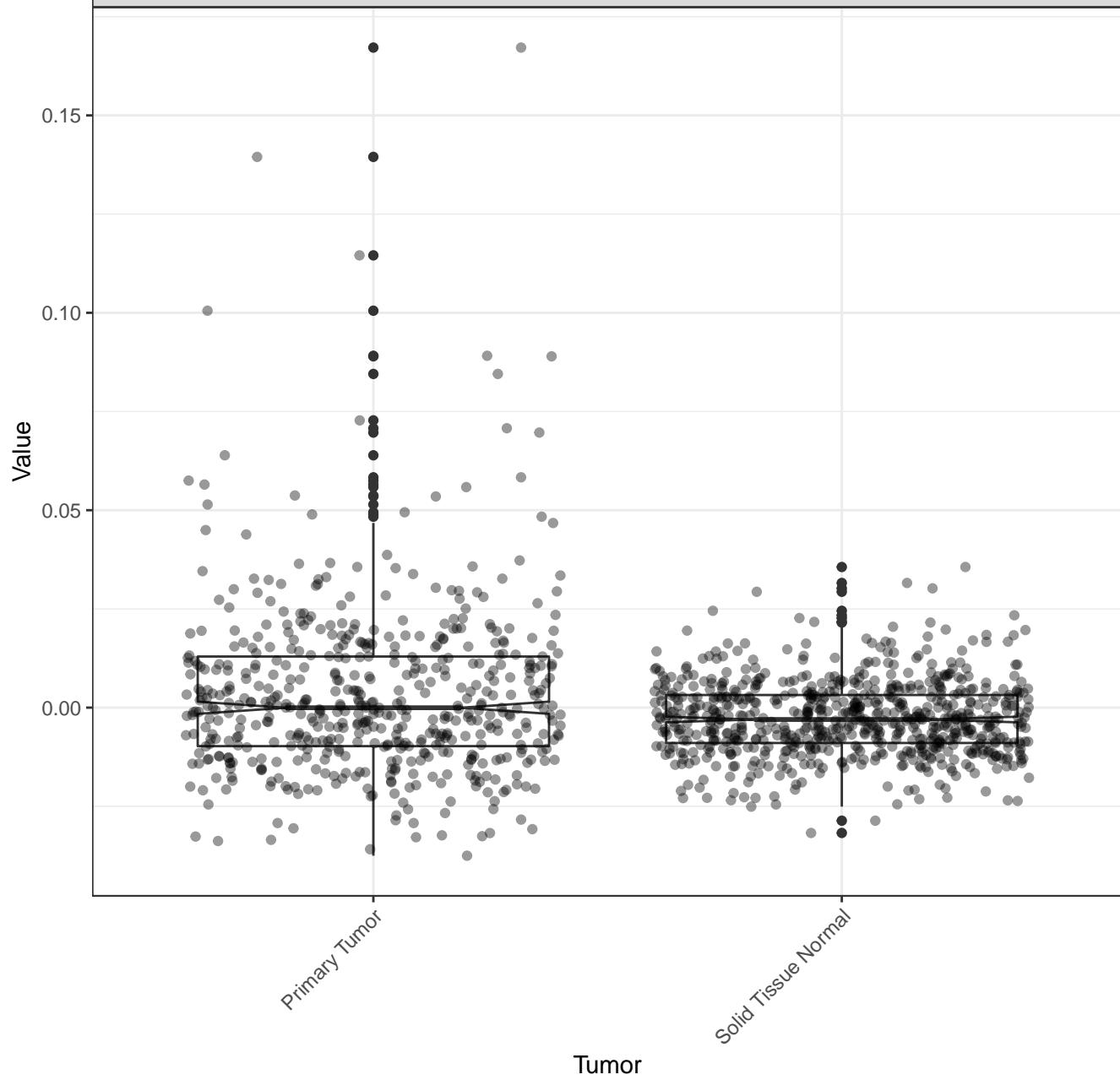




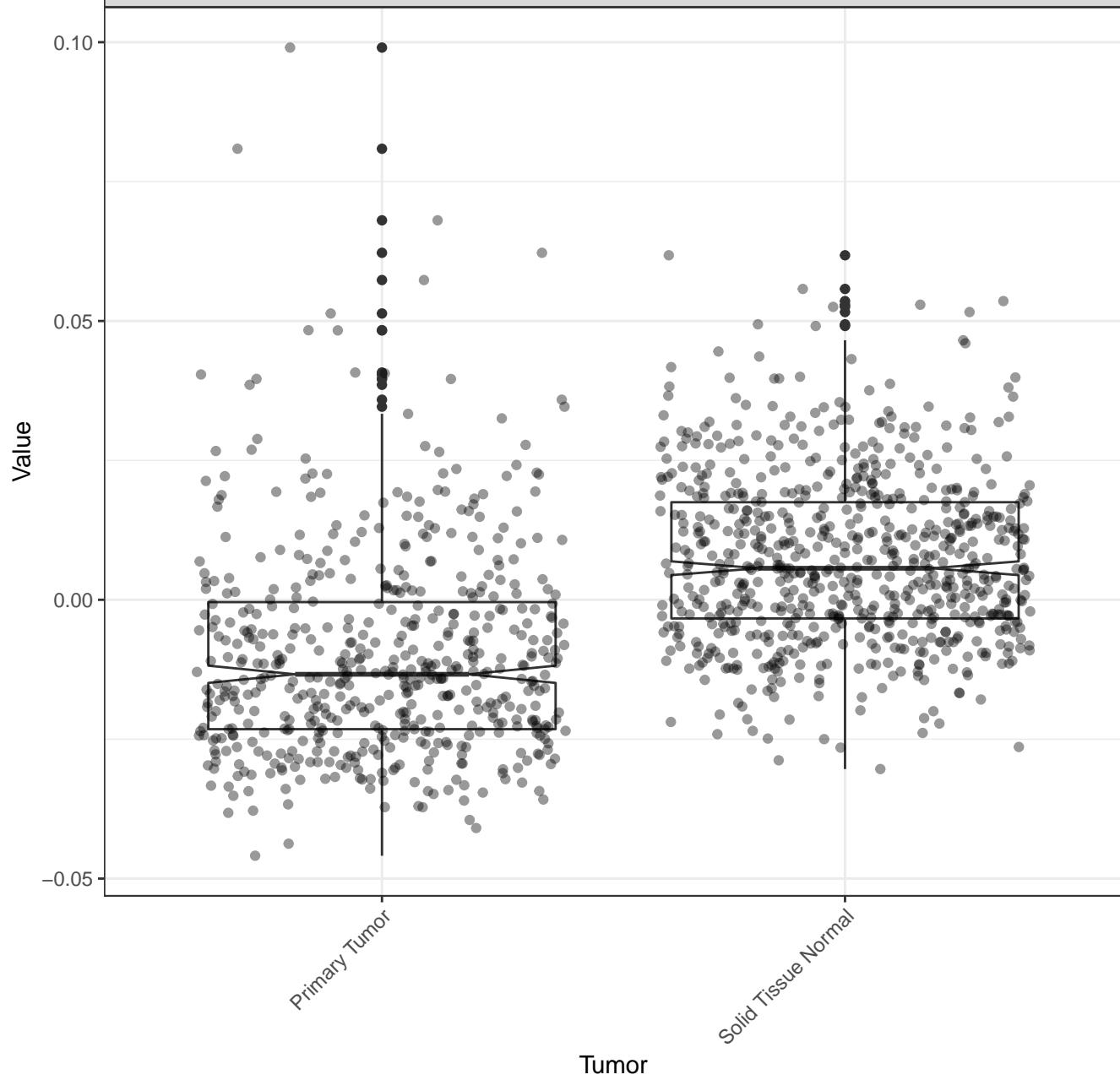
303,MIPS_NOP56P_ASSOCIATED_PRE_RRNA_COMPLEX



304,IRIS_DendriticCell-Control



305,REACTOME_SIGNALING_BY_RHO_GTPASES



Value

0.05

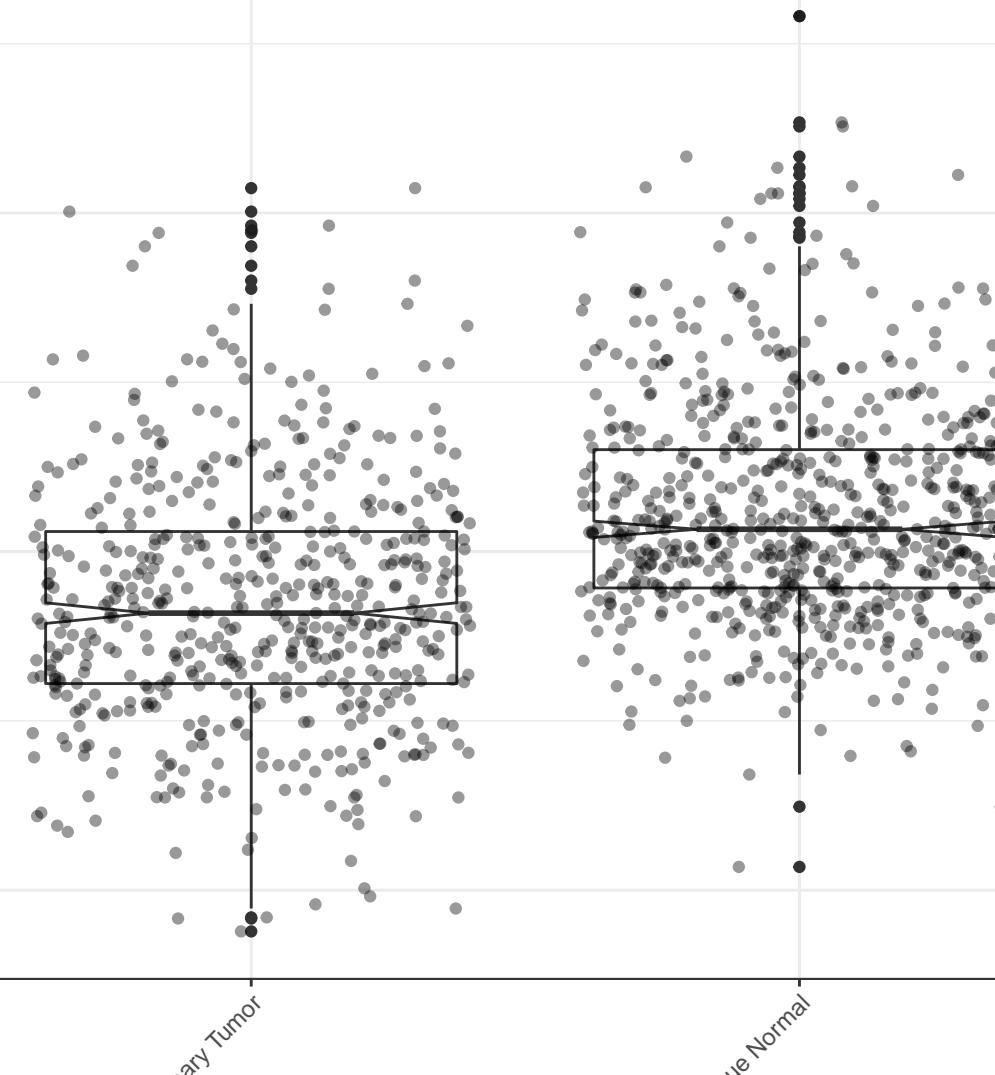
0.00

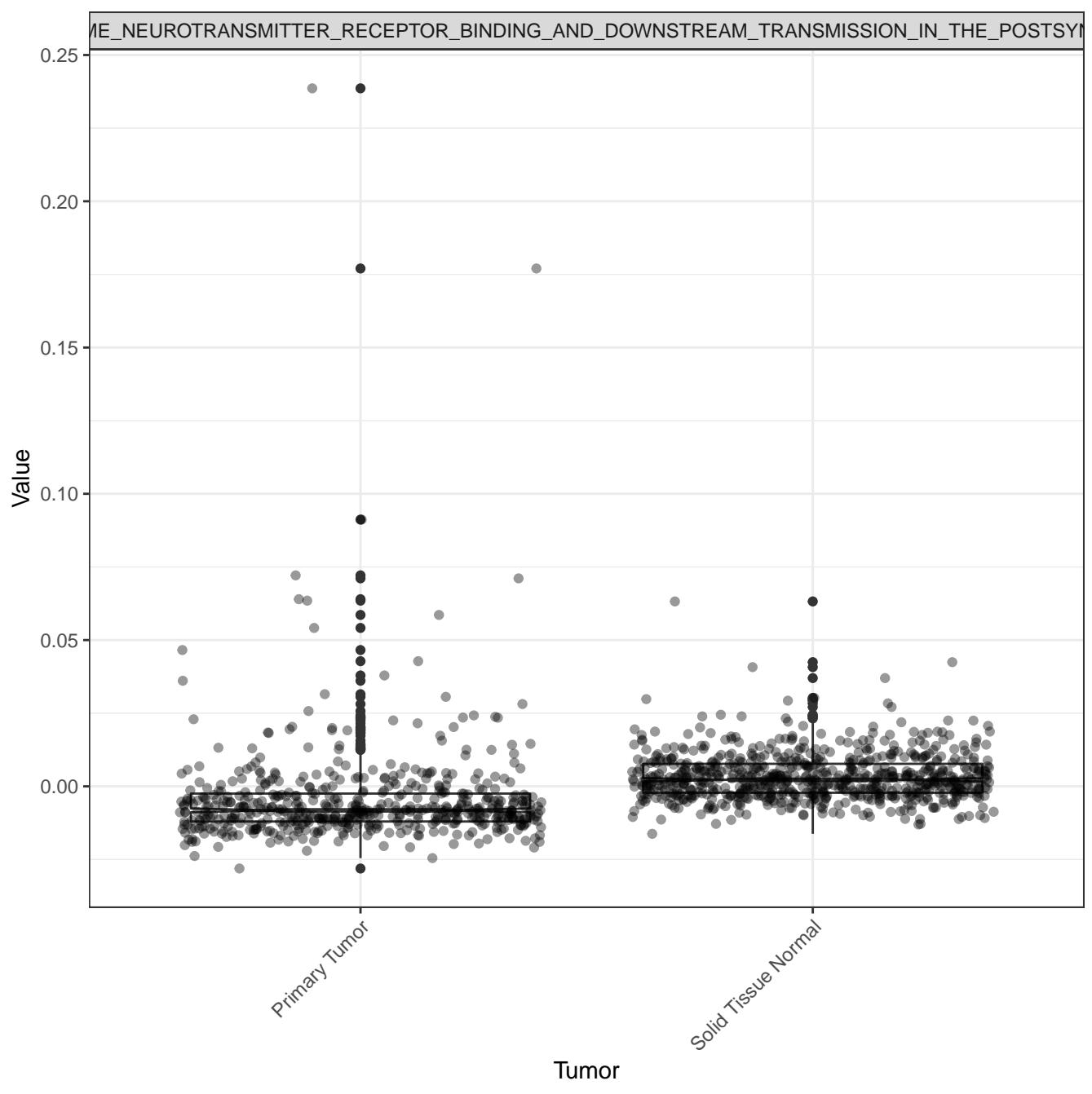
-0.05

Primary Tumor

Solid Tissue Normal

Tumor





LV 308

Value

0.10

0.05

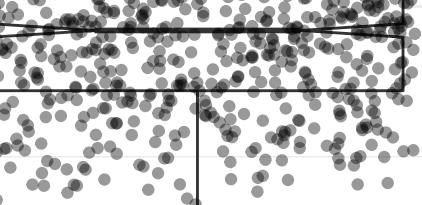
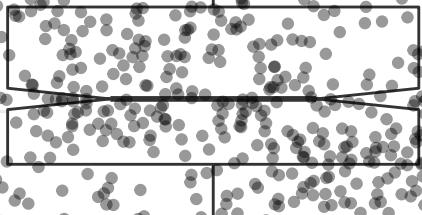
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 309

Value

0.15

0.10

0.05

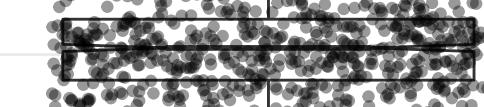
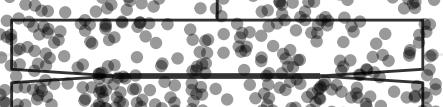
0.00

-0.05

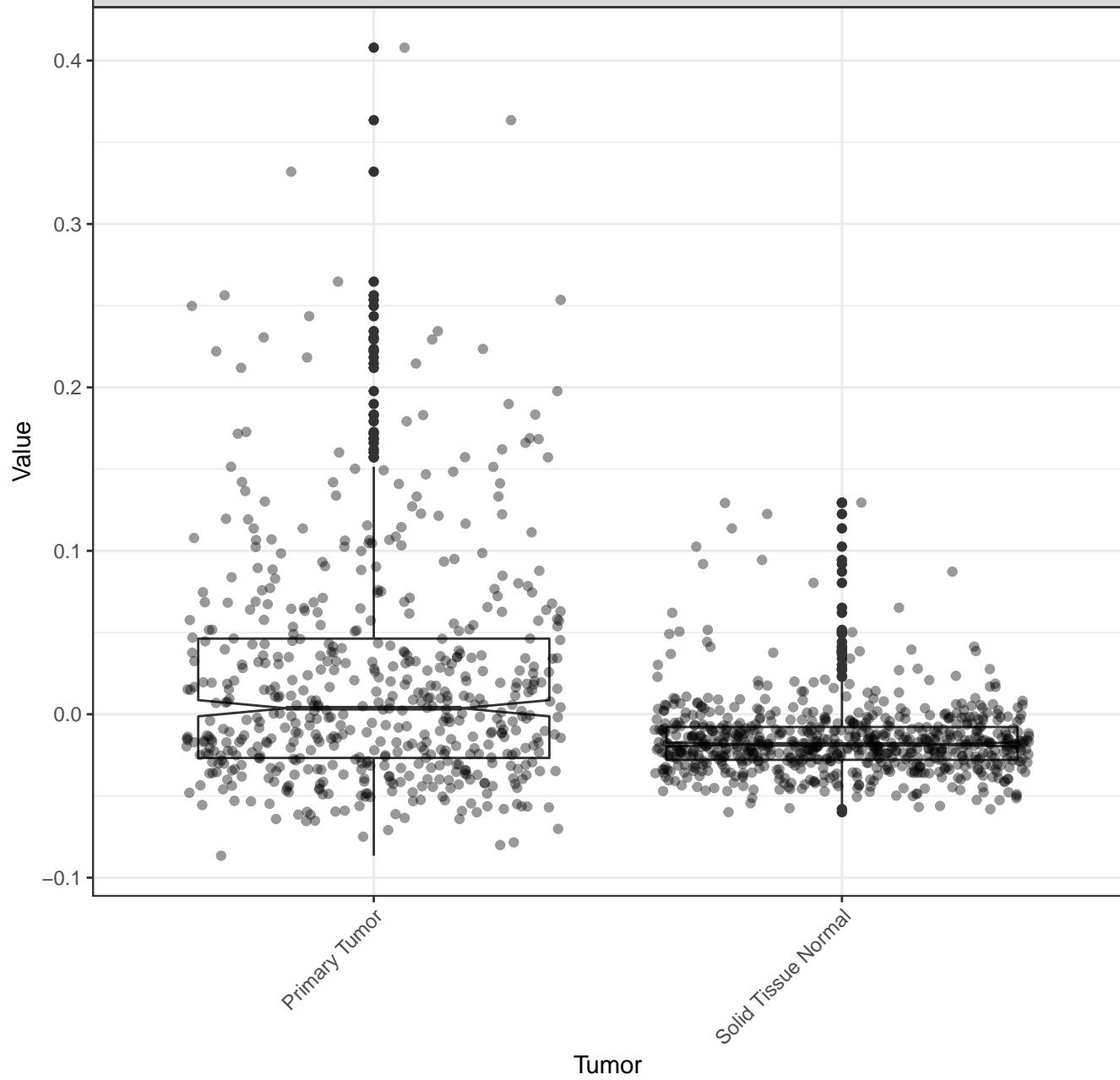
Primary Tumor

Solid Tissue Normal

Tumor



310,REACTOME_DNA_REPLICATION



311,REACTOME_METABOLISM_OF_PROTEINS

Value

0.15

0.10

0.05

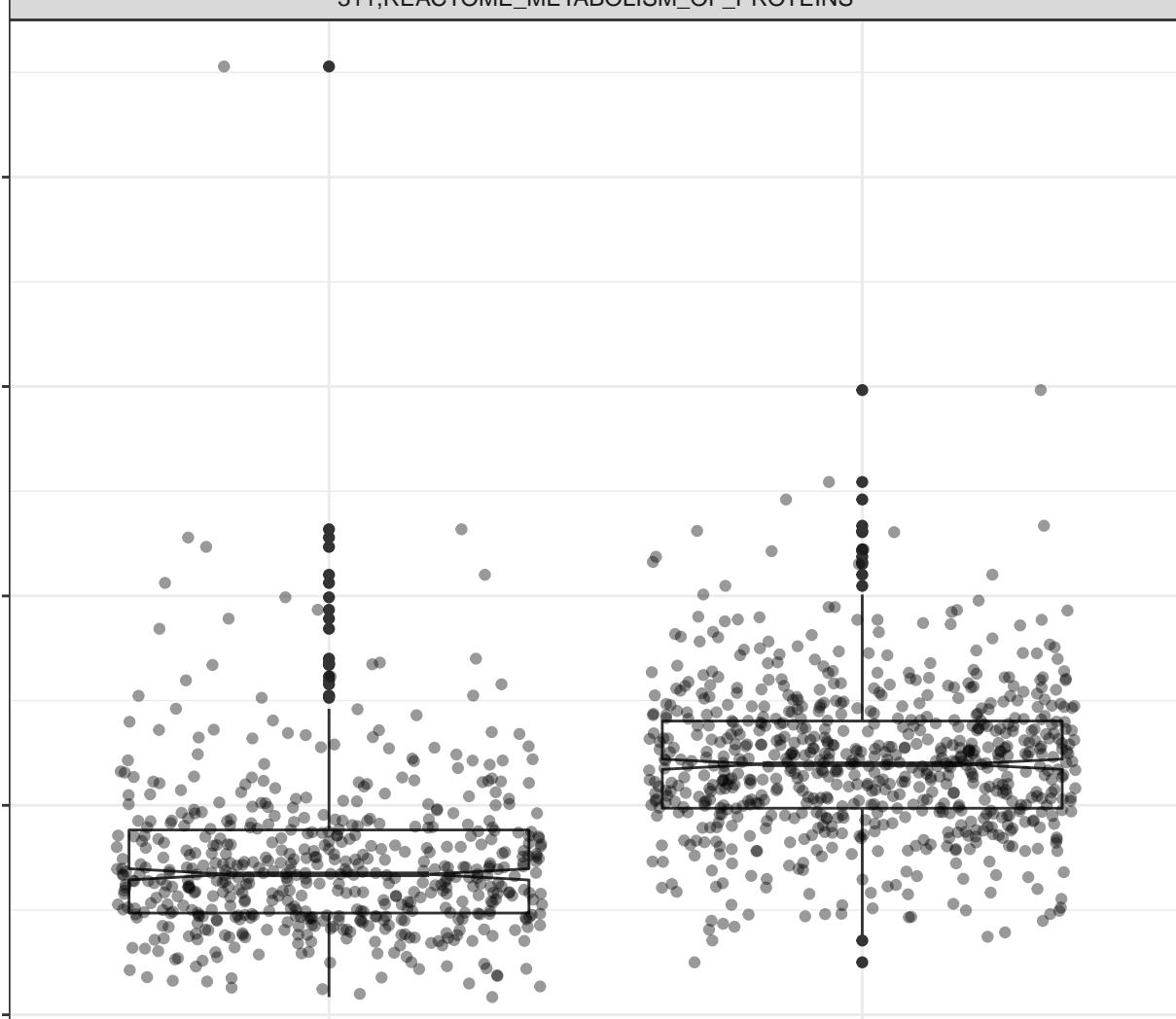
0.00

-0.05

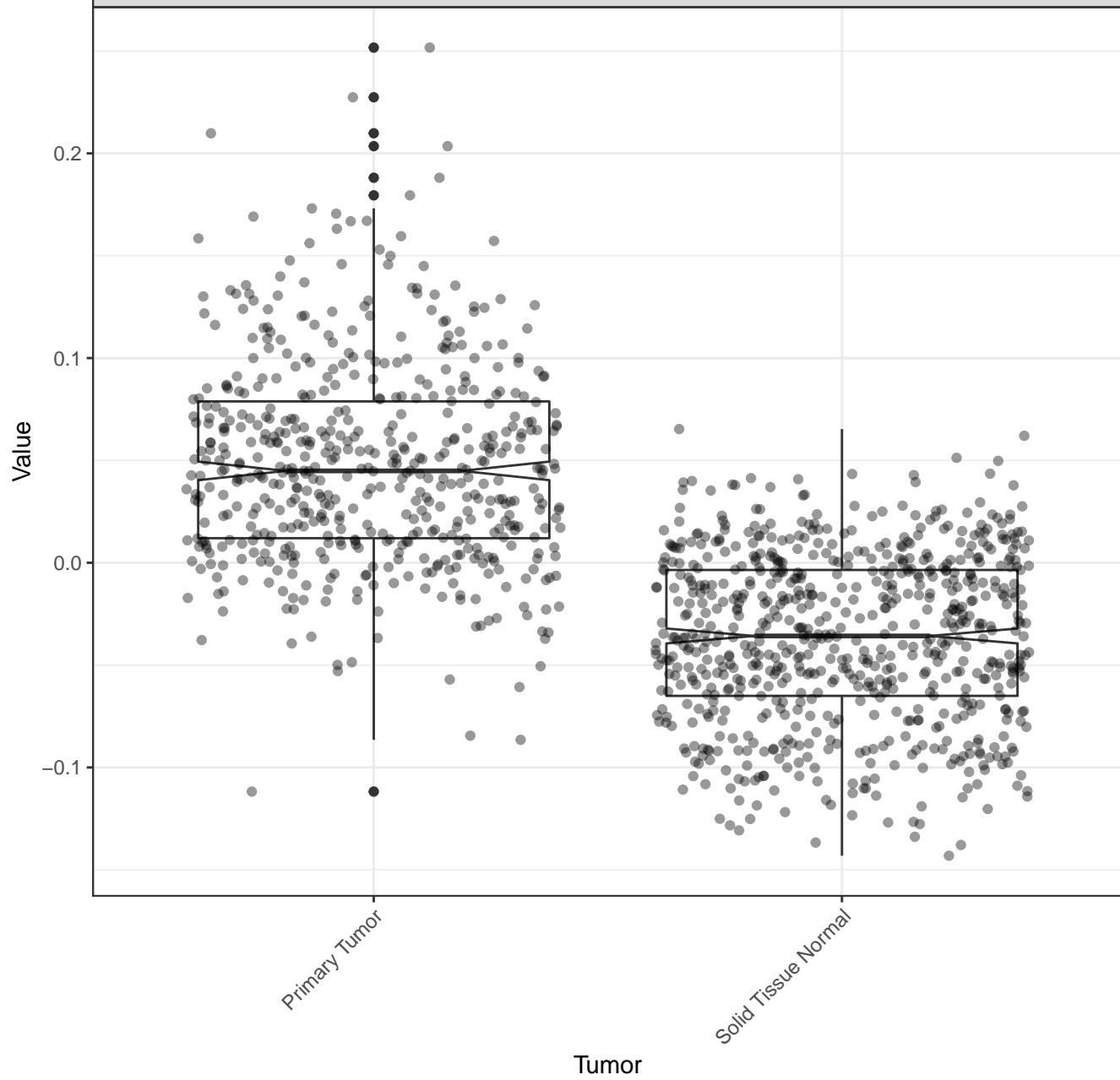
Primary Tumor

Solid Tissue Normal

Tumor



312,KEGG_PEROXISOME



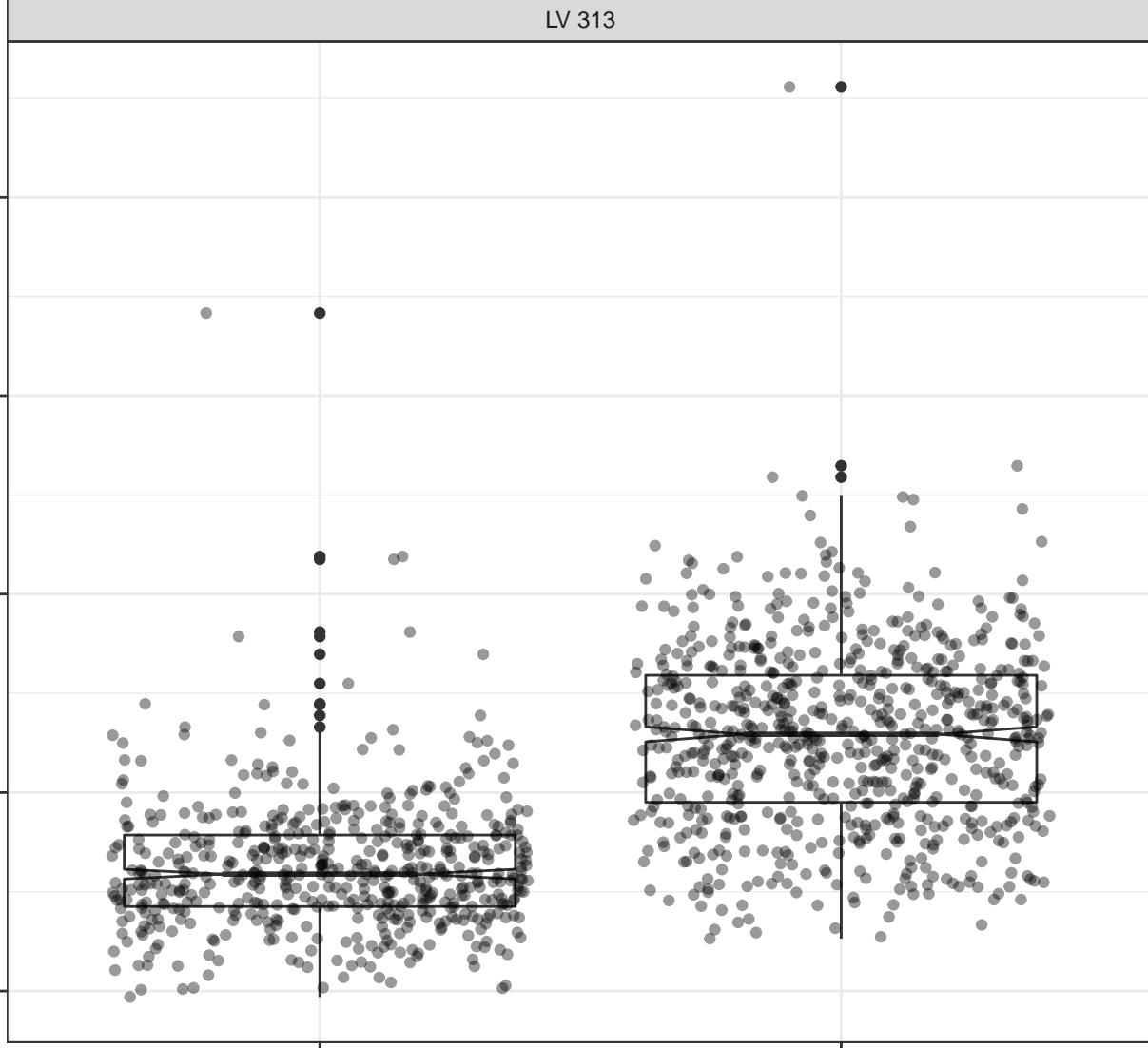
Value

0.15
0.10
0.05
0.00
-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 314

Value

0.15

0.10

0.05

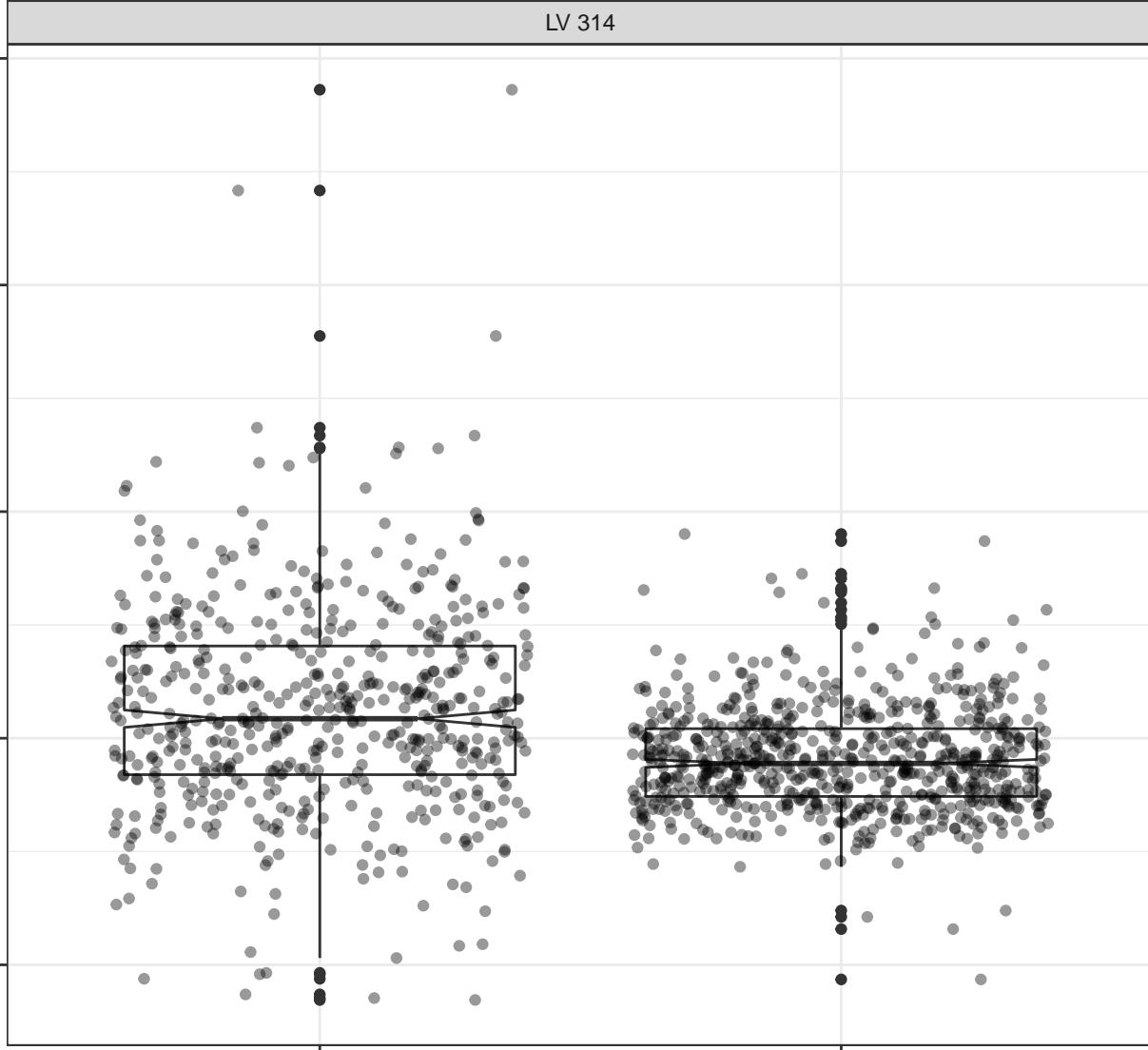
0.00

-0.05

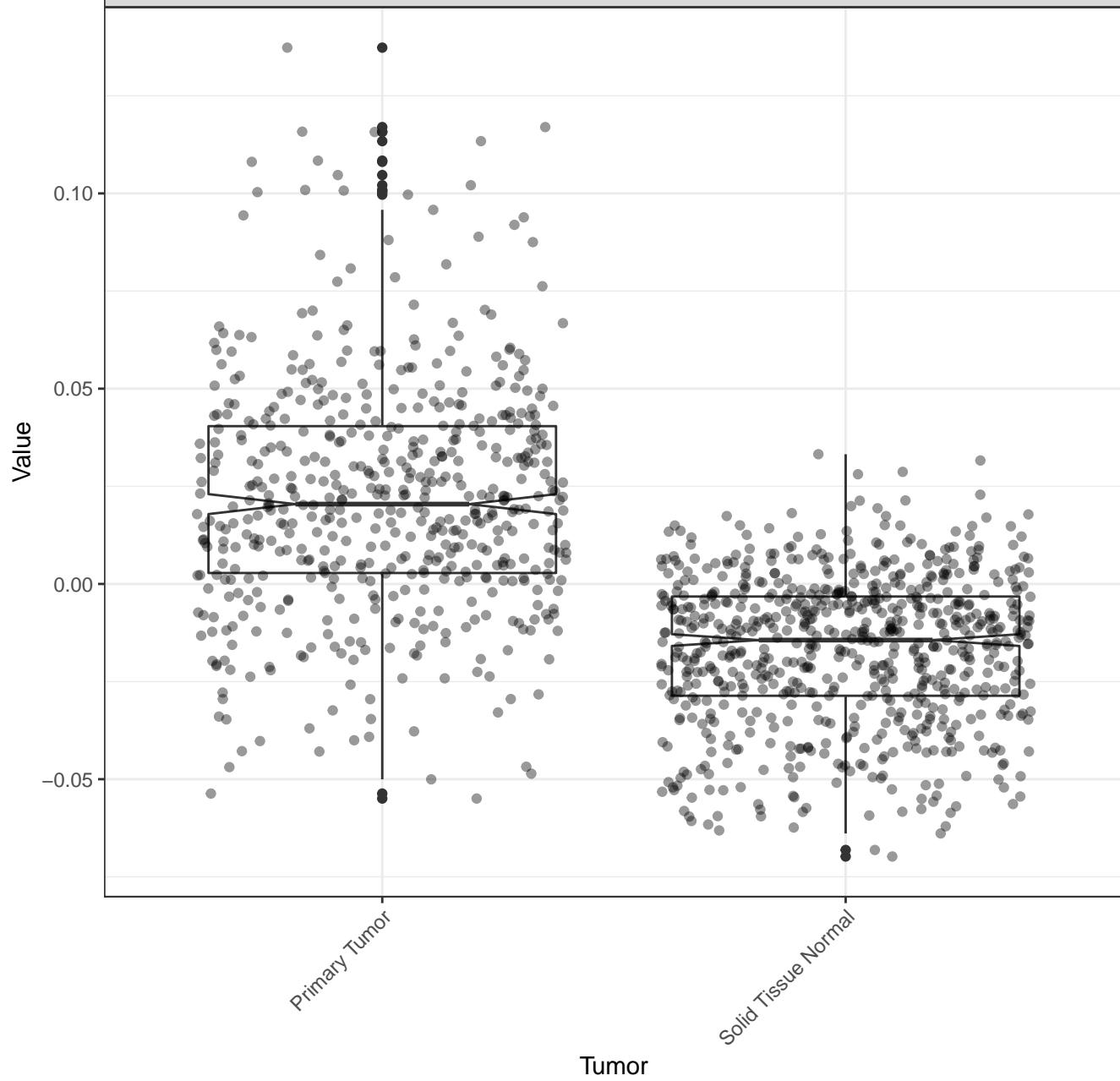
Primary Tumor

Solid Tissue Normal

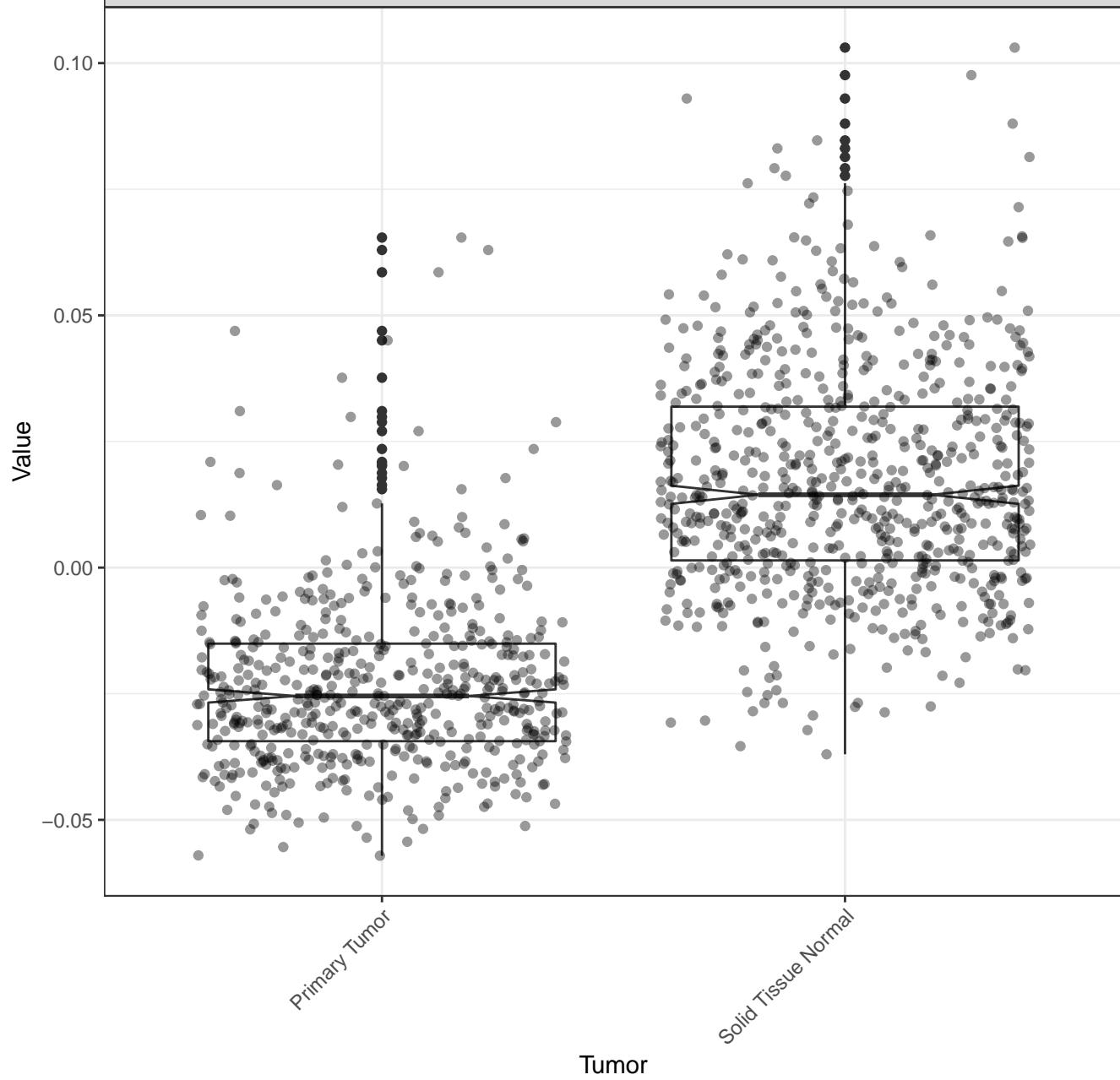
Tumor



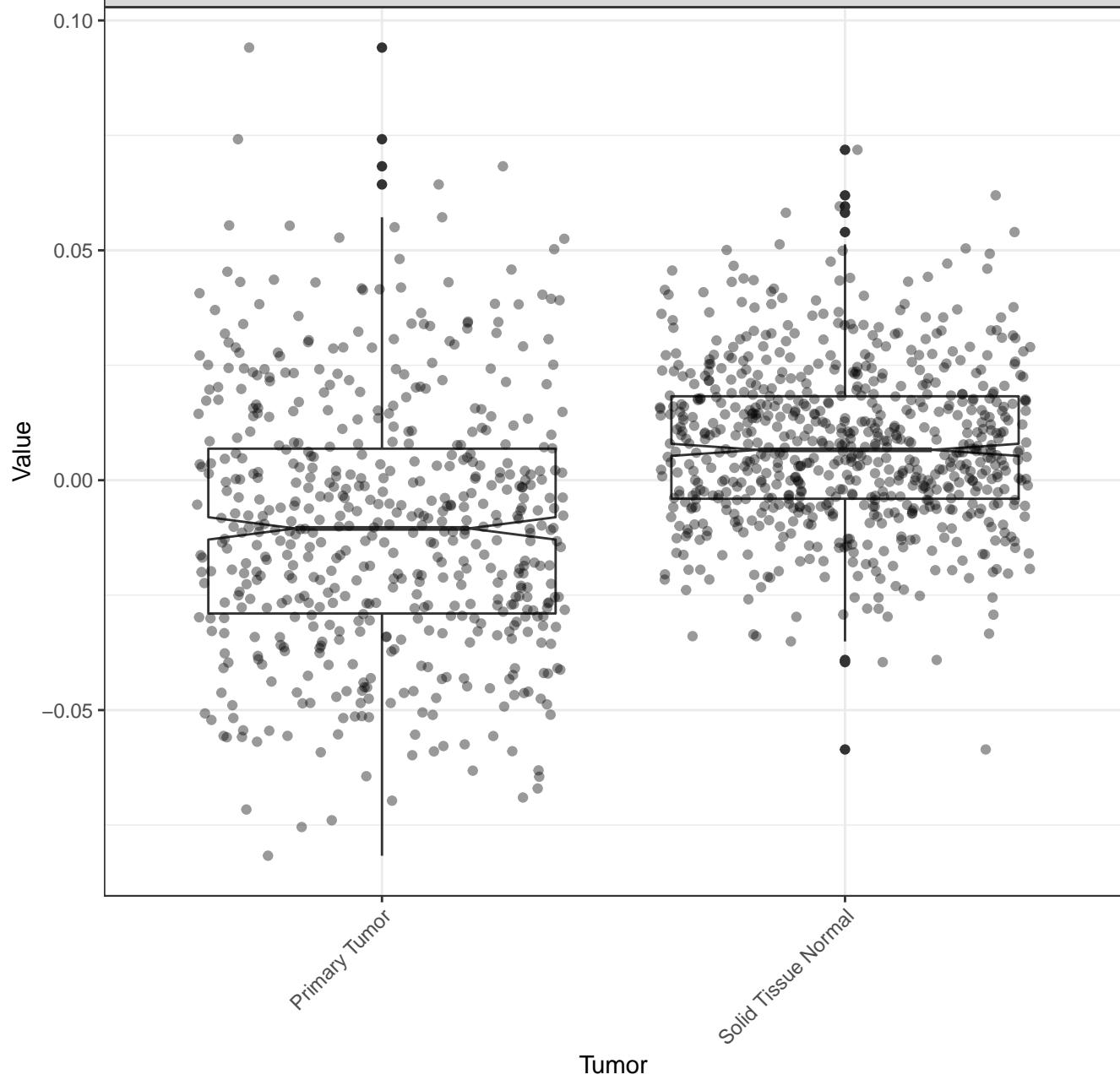
315,REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION

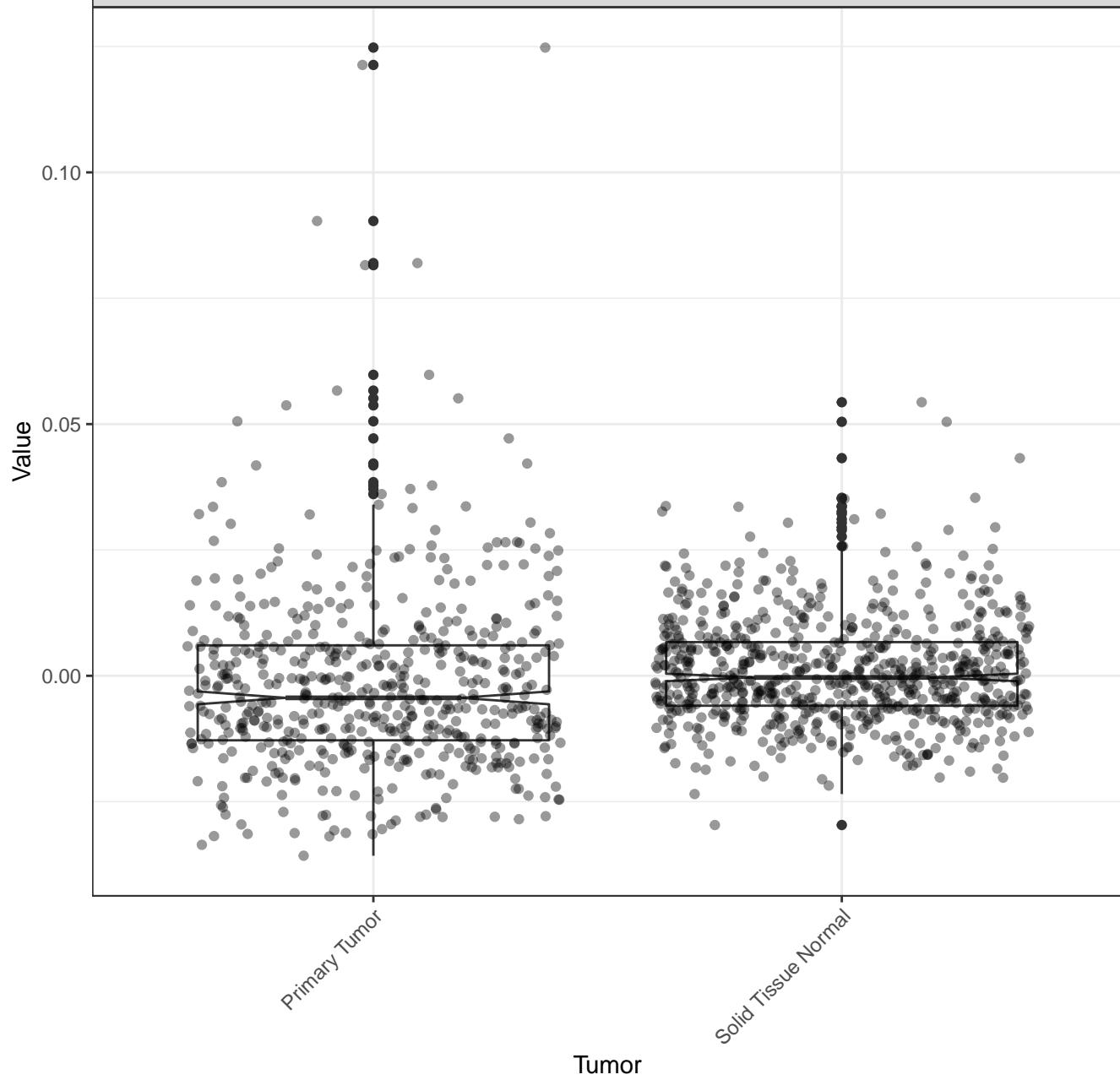


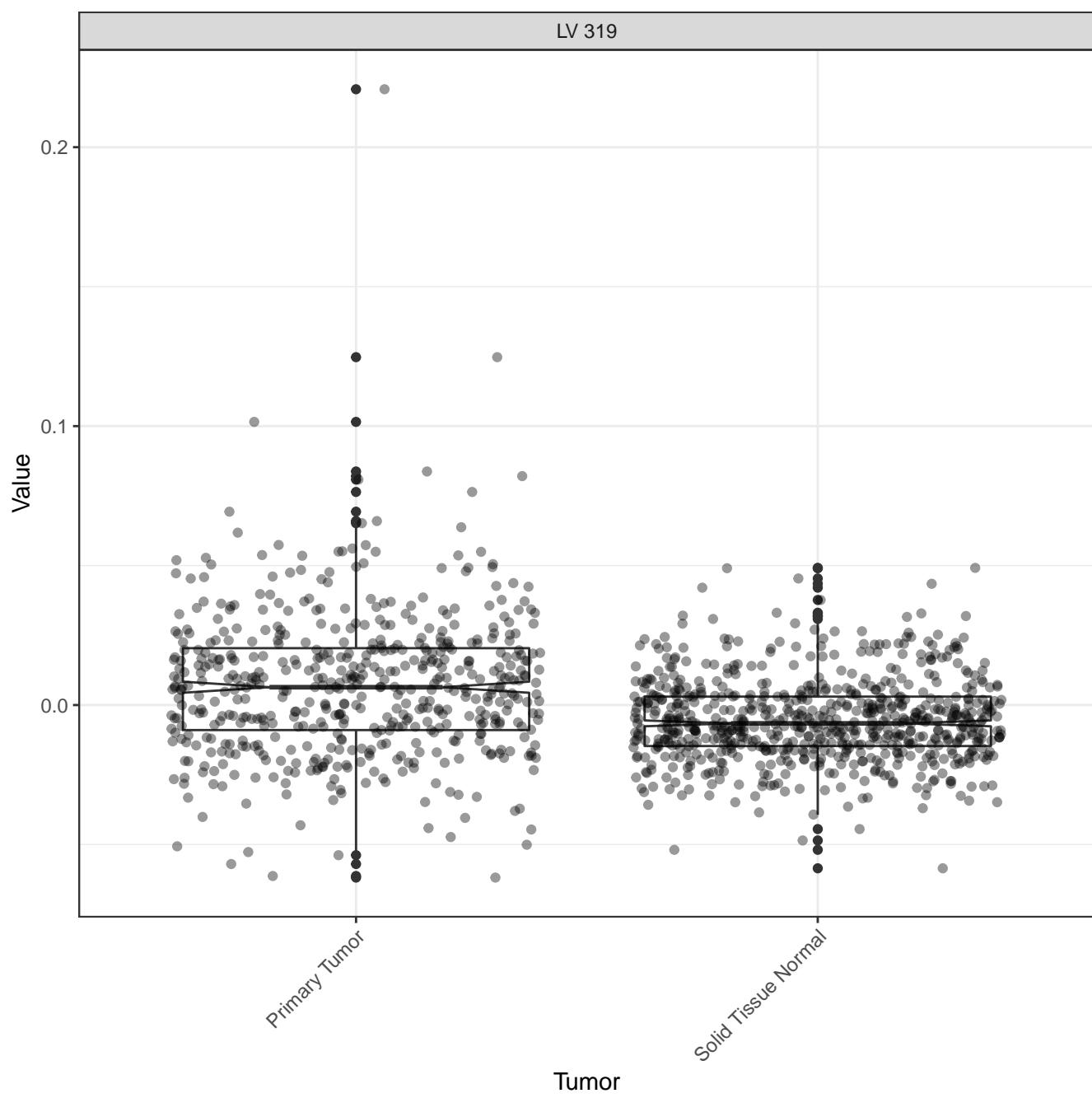
316,KEGG_MAPK_SIGNALING_PATHWAY



LV 317







320,REACTOME_METABOLISM_OF_MRNA

Value

0.10

0.05

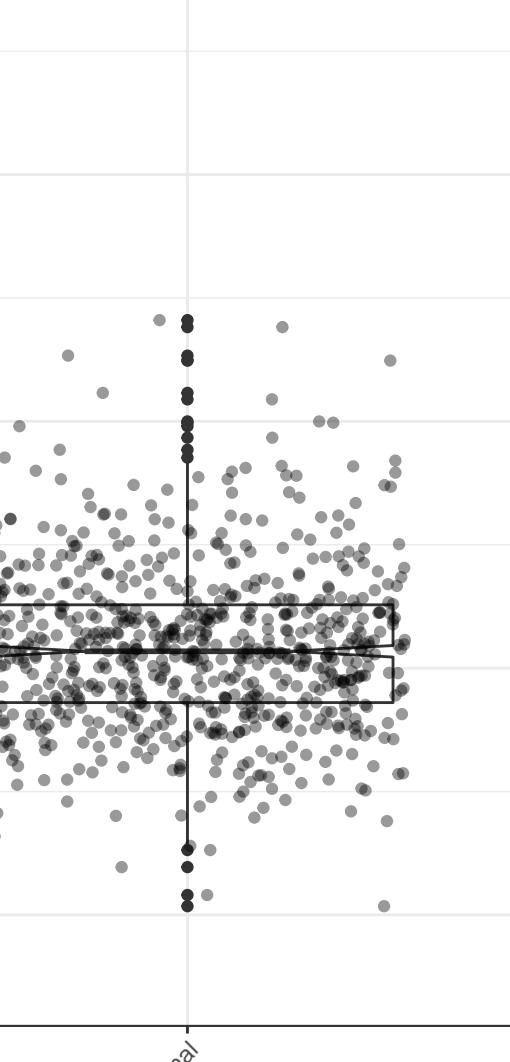
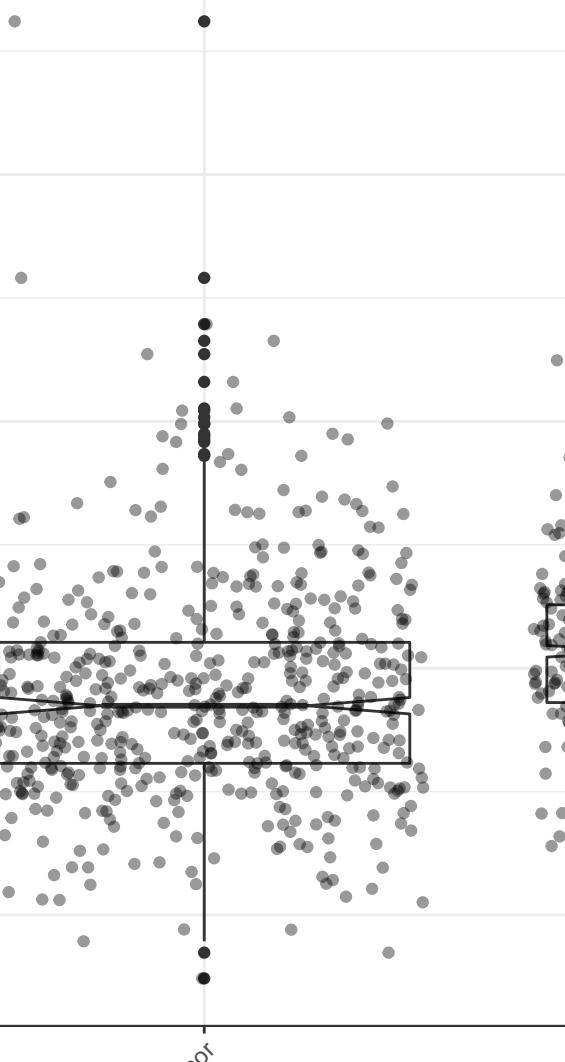
0.00

-0.05

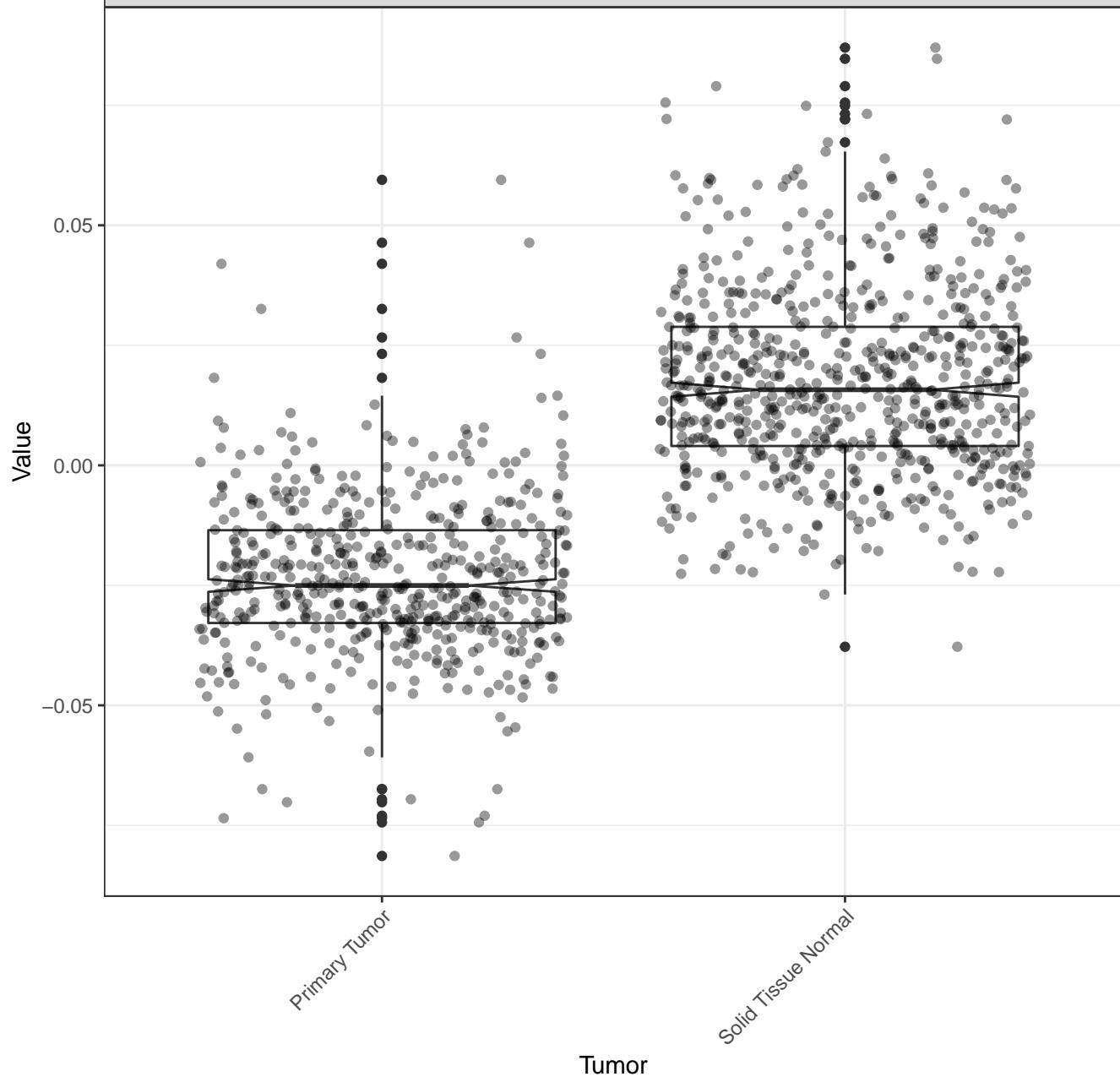
Primary Tumor

Solid Tissue Normal

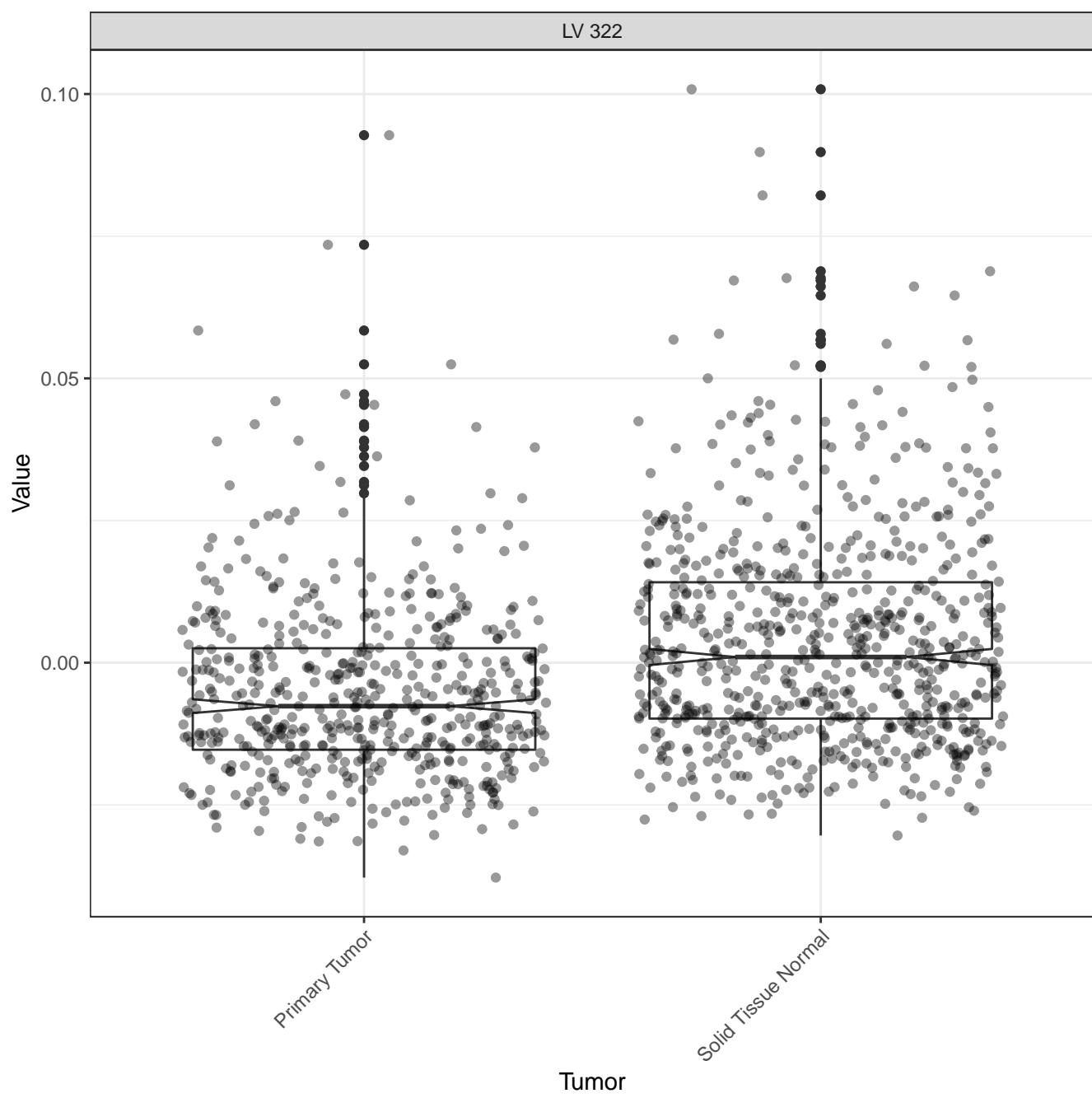
Tumor



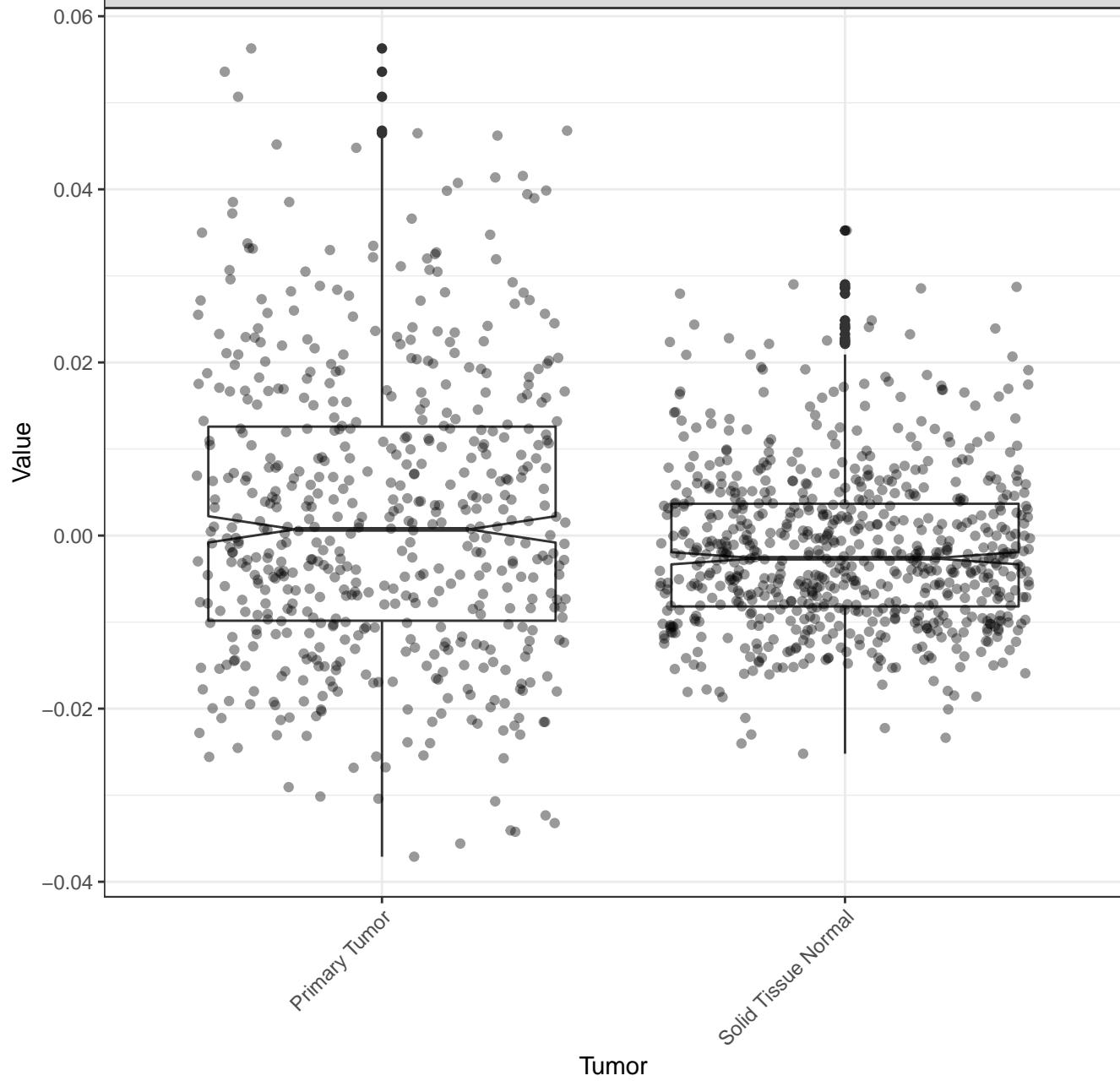
321,KEGG_VASCULAR_SMOOTH_MUSCLE_CONTRACTION



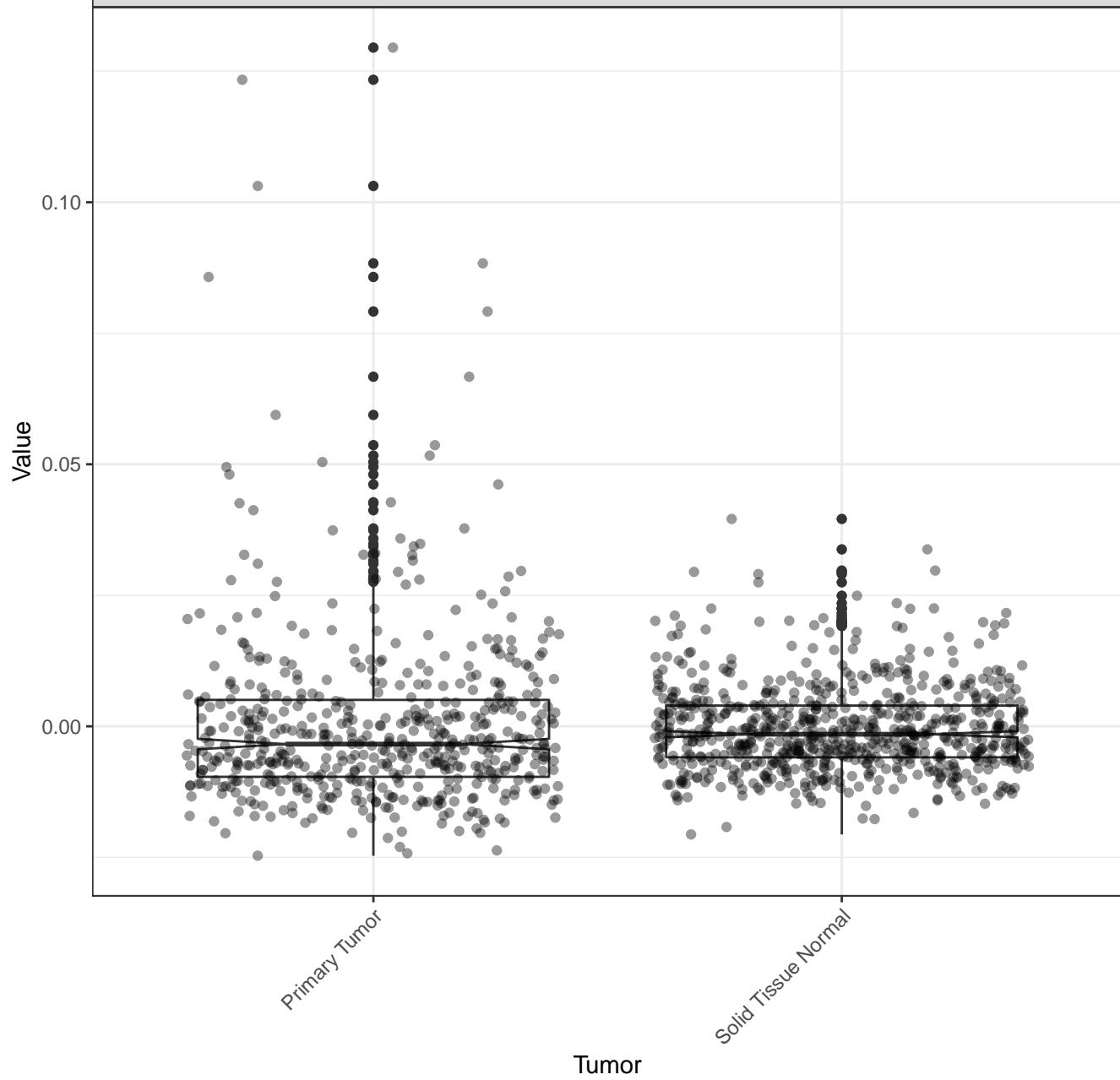
LV 322



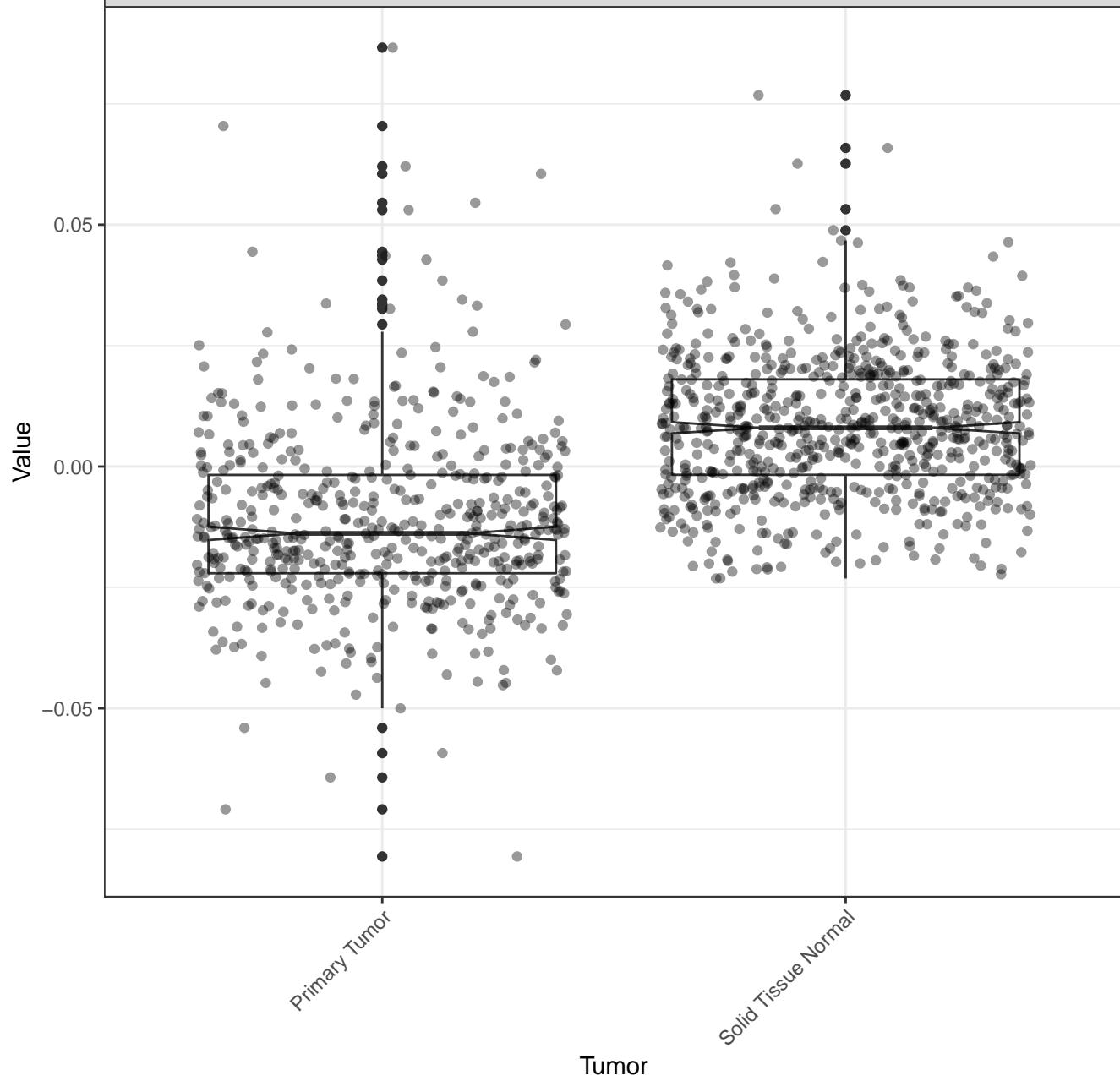
323,KEGG_LYSOSOME



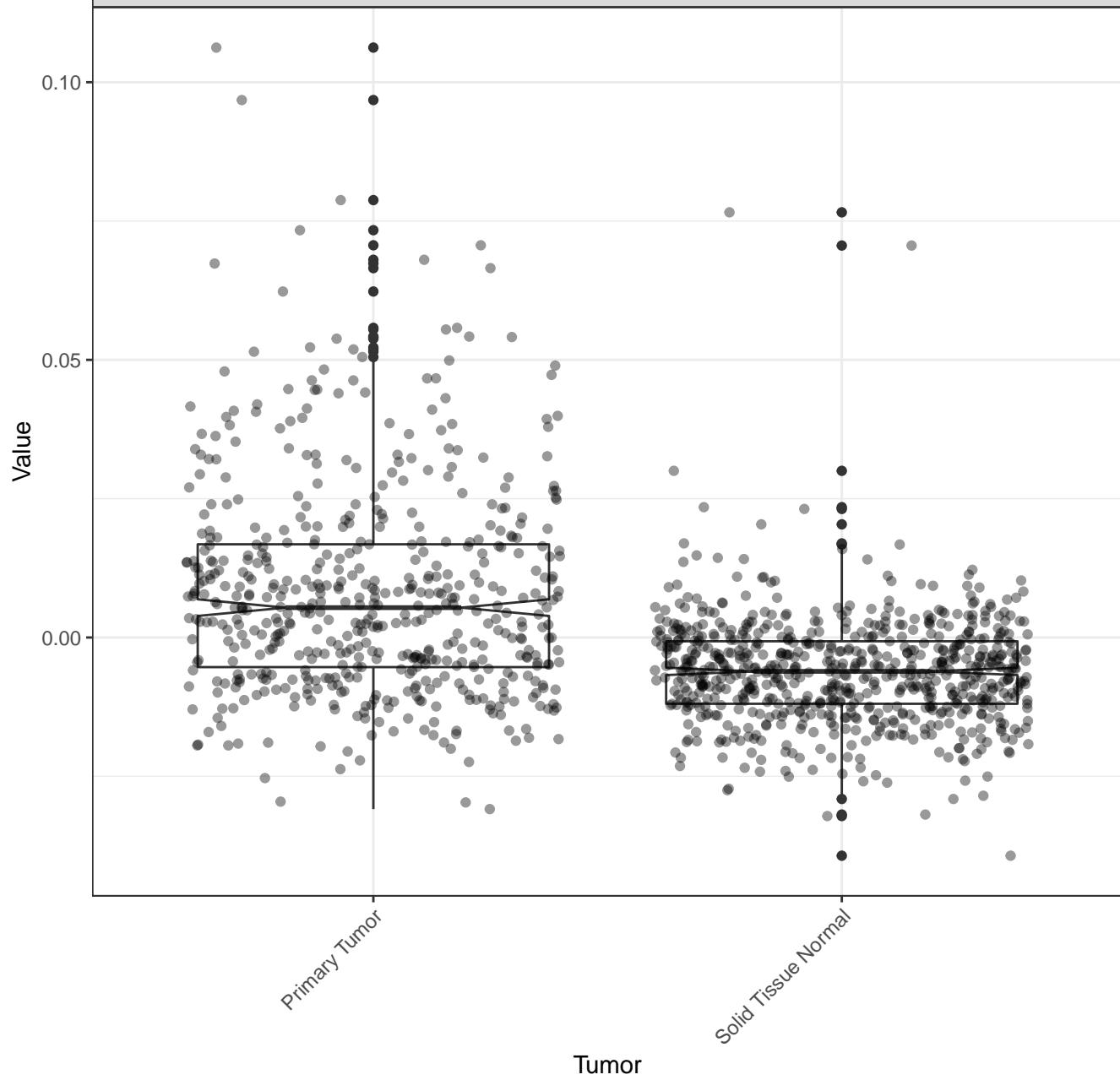
324,REACTOME_PI_METABOLISM



325,REACTOME_NEURONAL_SYSTEM



326,KEGG-ENDOCYTOSIS



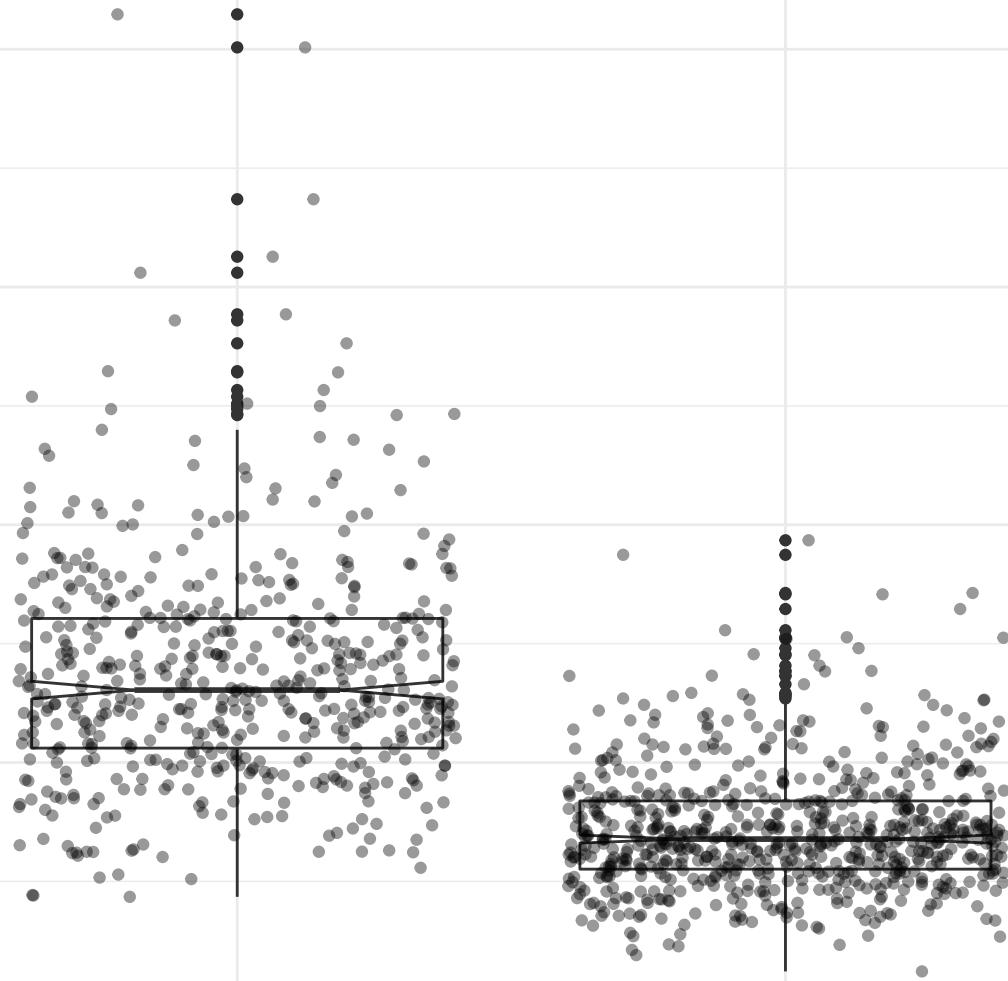
Value

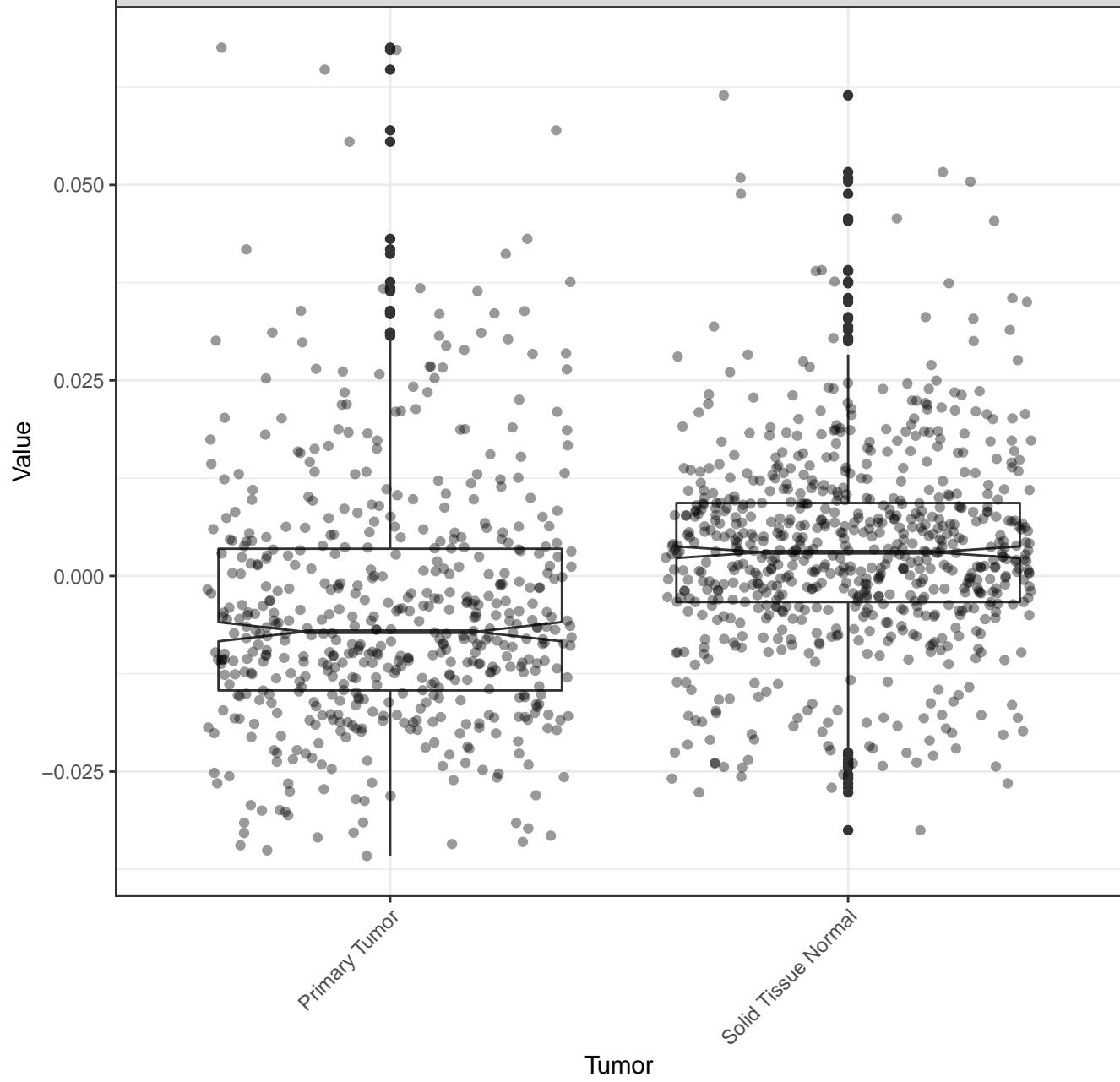
0.15
0.10
0.05
0.00
-0.05

Primary Tumor

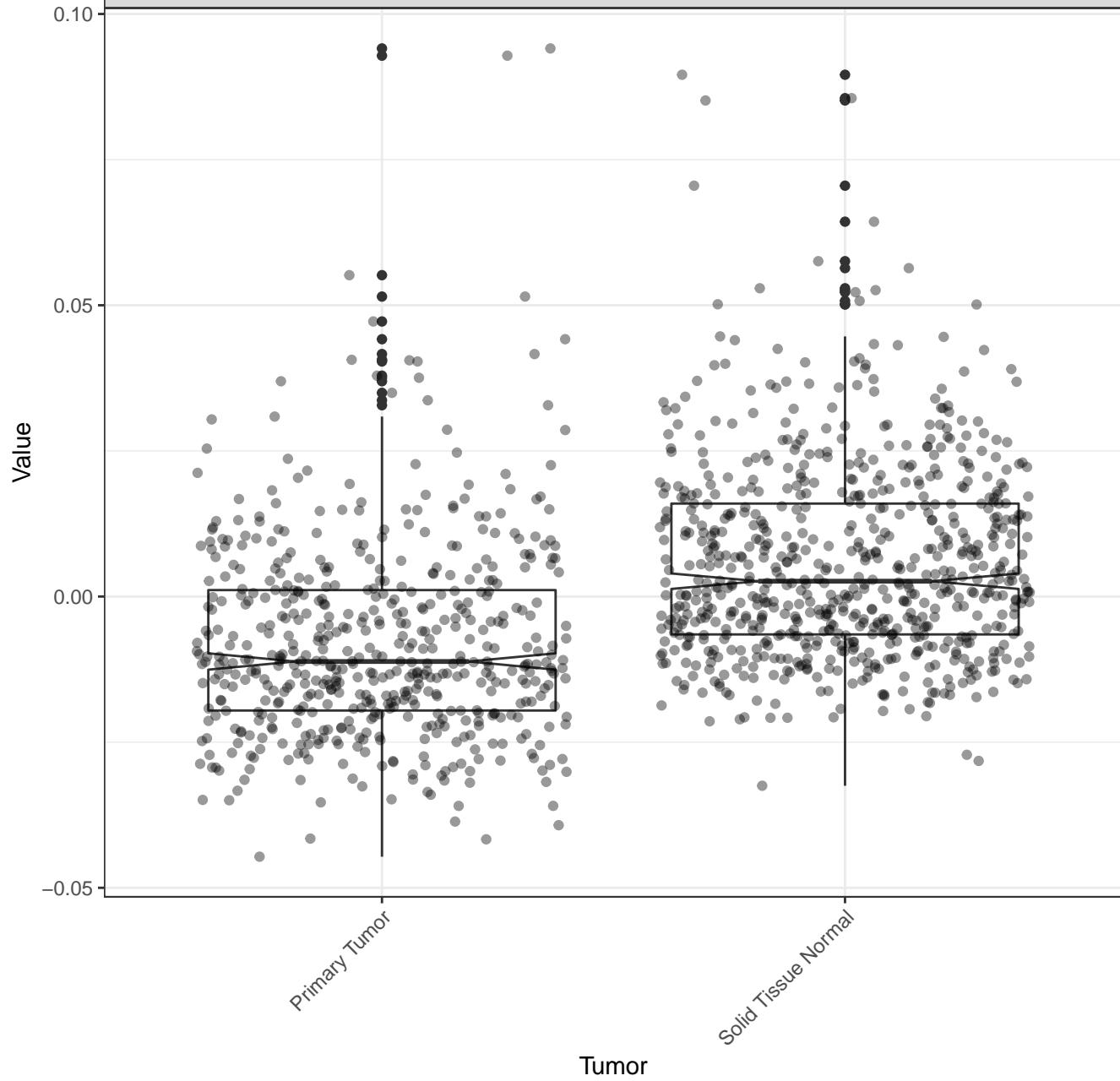
Solid Tissue Normal

Tumor





329,REACTOME_TOLL_RECEPTOR_CASCADES



330,MIPS_40S_RIBOSOMAL_SUBUNIT_CYTOPLASMIC

Value

0.10

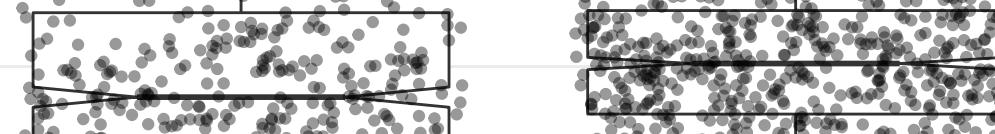
0.05

0.00

Primary Tumor

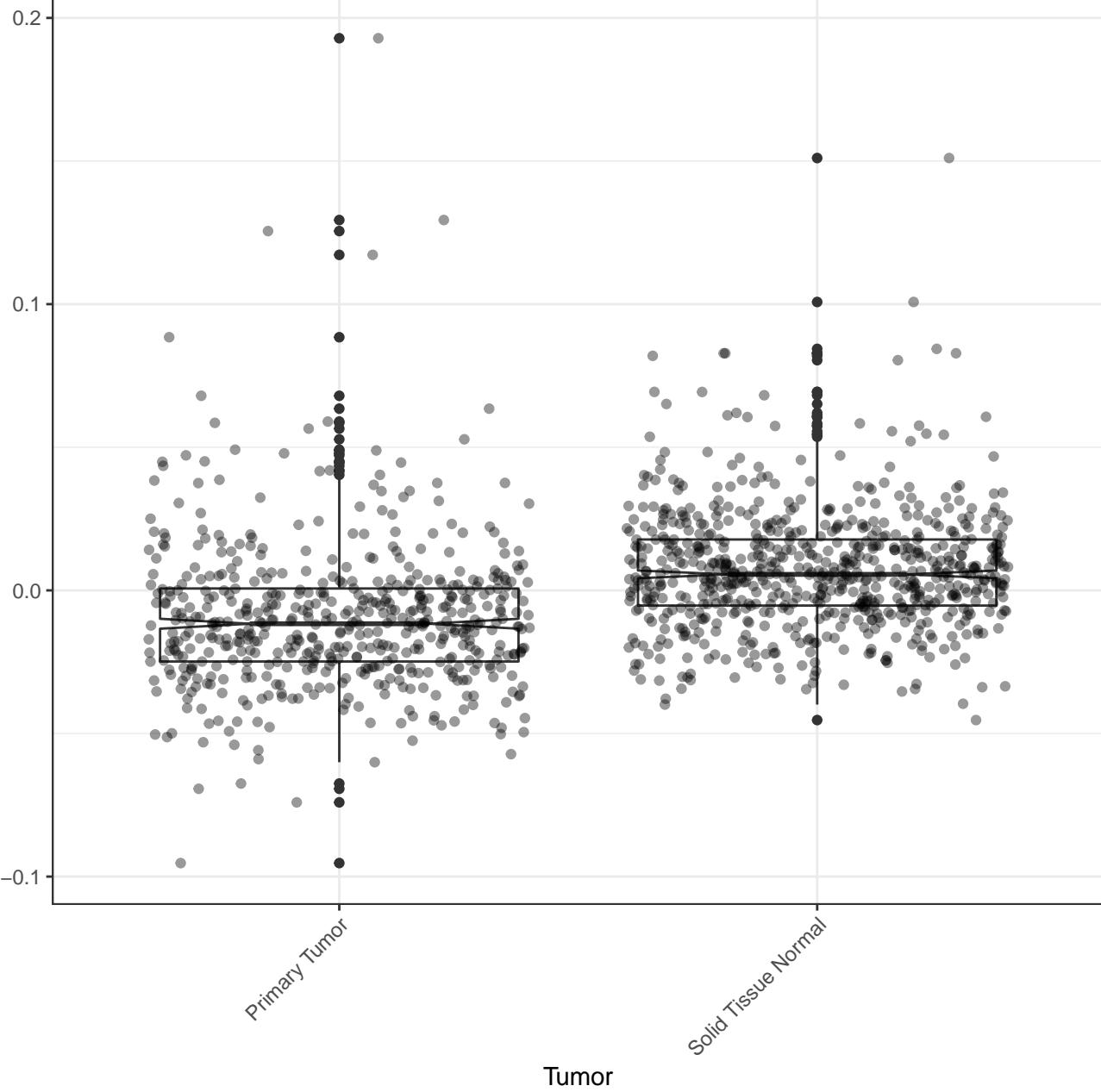
Solid Tissue Normal

Tumor



331,DMAP_ERY3

Value



332,REACTOME_DOWNSTREAM_SIGNALING_EVENTS_OF_B_CELL_RECECTOR_BCR

Value

0.05

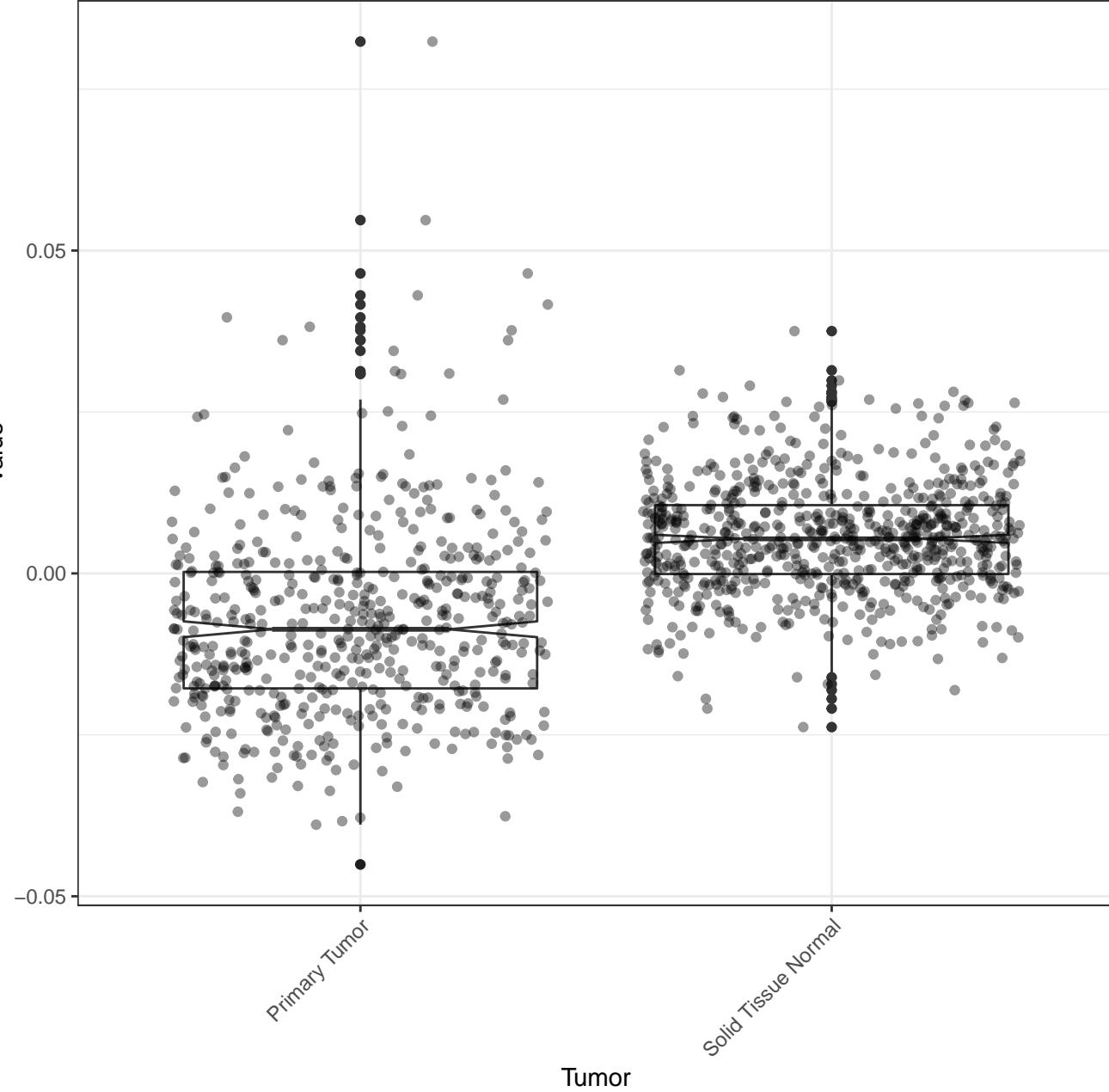
0.00

-0.05

Primary Tumor

Tumor

Solid Tissue Normal



333,KEGG_SYSTEMIC_LUPUS_ERYTHEMATOSUS

Value

0.05

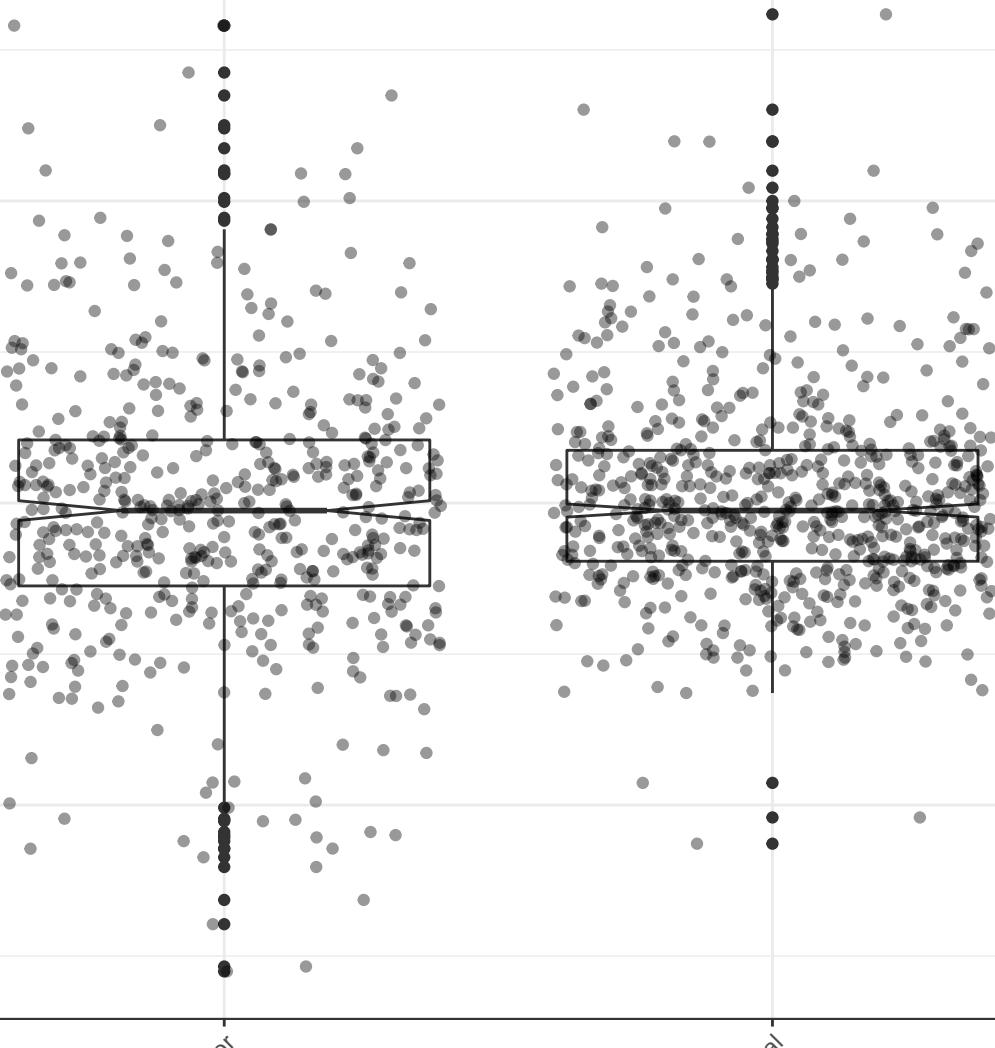
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



334,REACTOME_AXON_GUIDANCE

Value

0.05

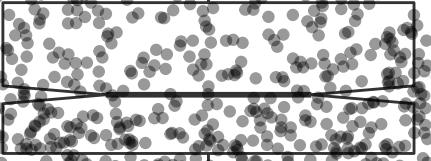
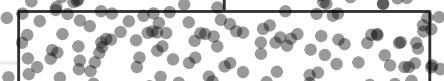
0.00

-0.05

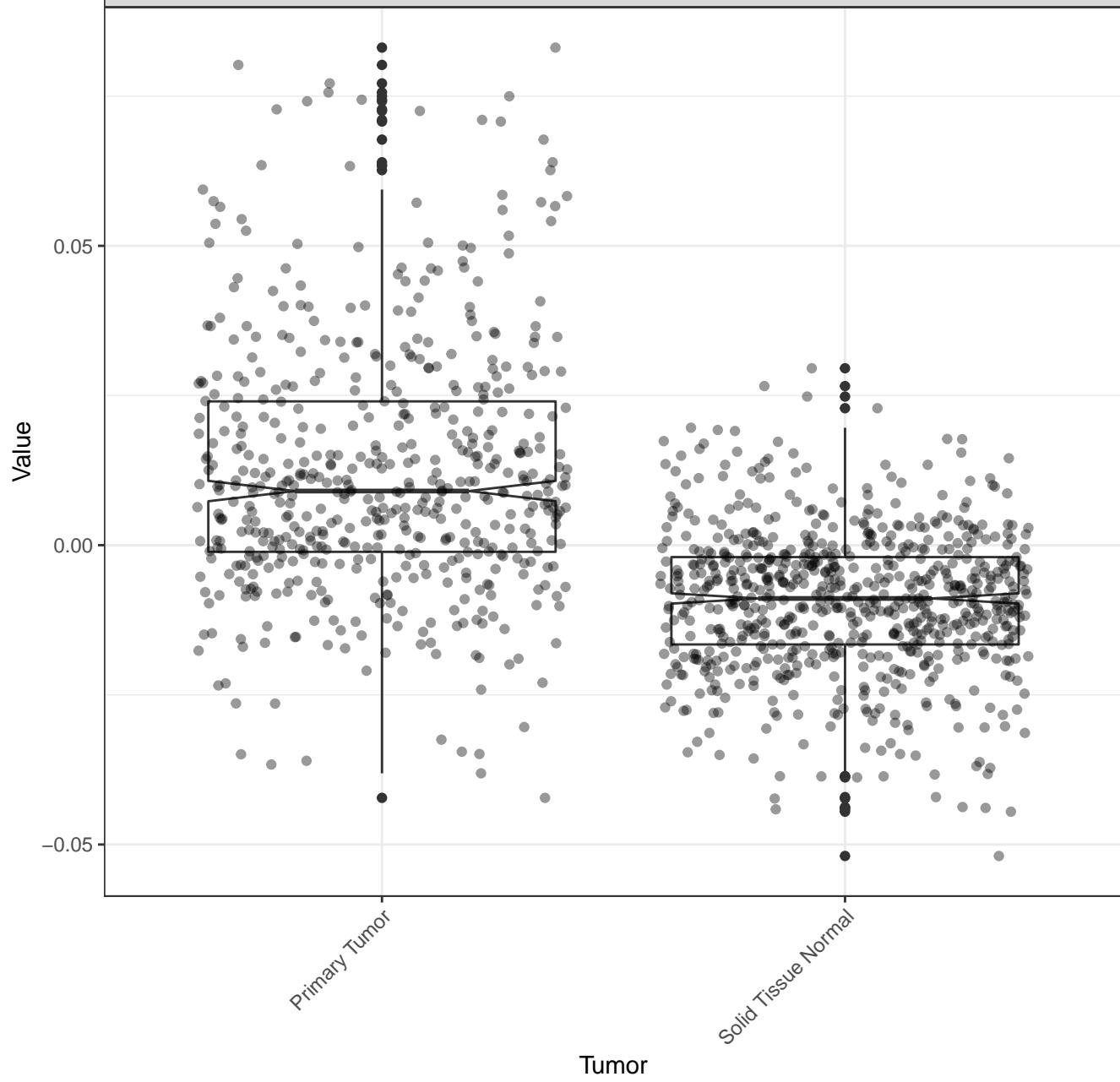
Primary Tumor

Solid Tissue Normal

Tumor



335,KEGG_ADHERENS_JUNCTION



LV 336

Value

0.10

0.05

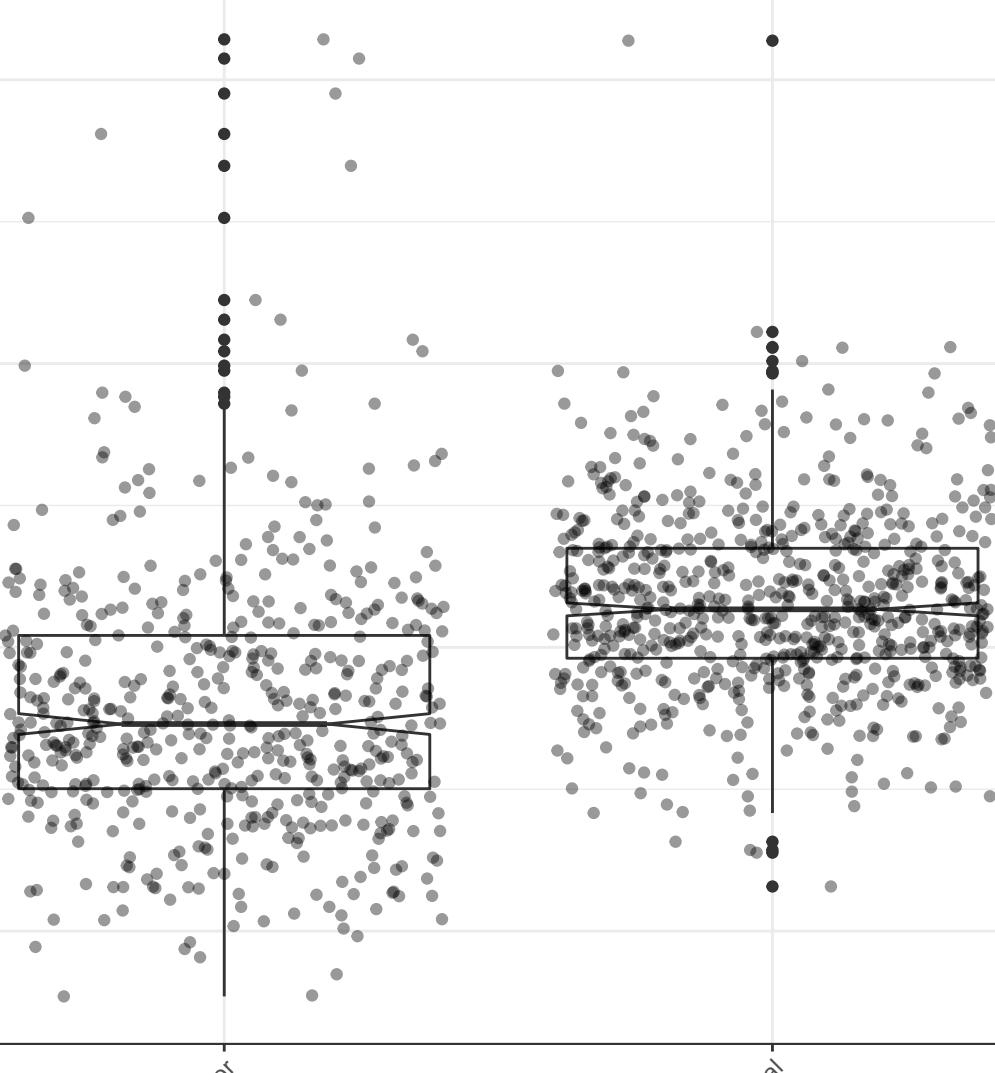
0.00

-0.05

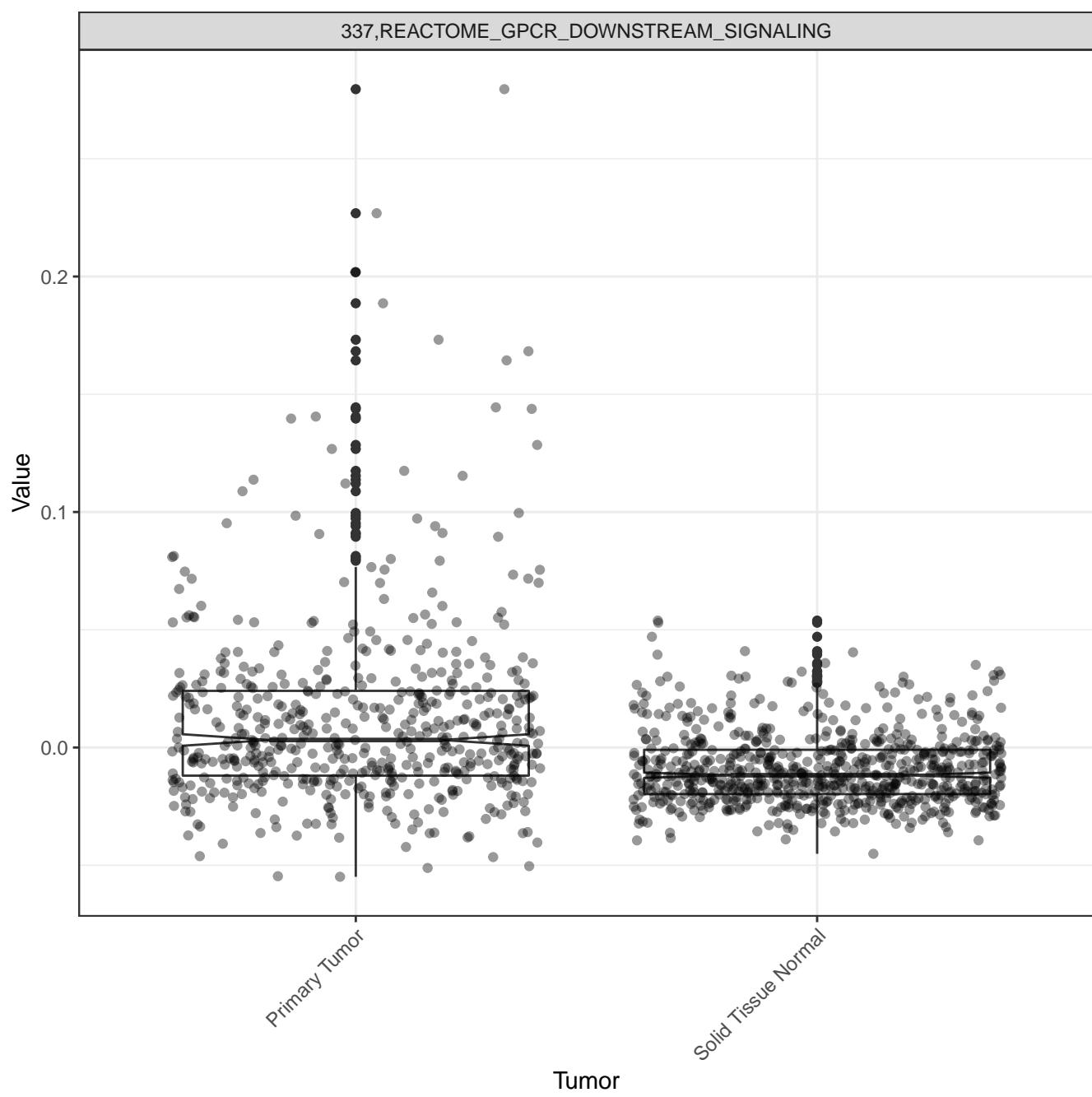
Primary Tumor

Solid Tissue Normal

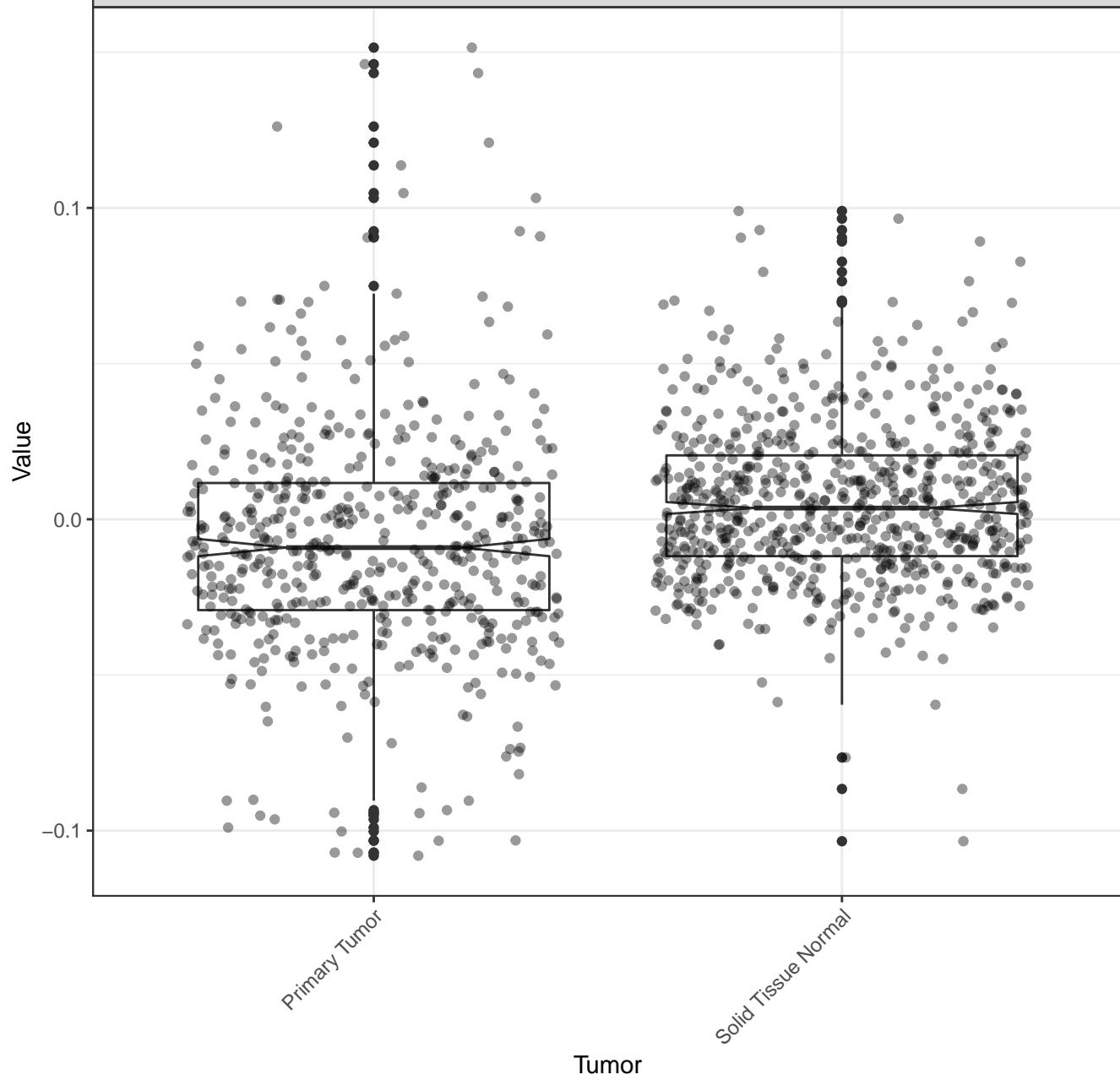
Tumor



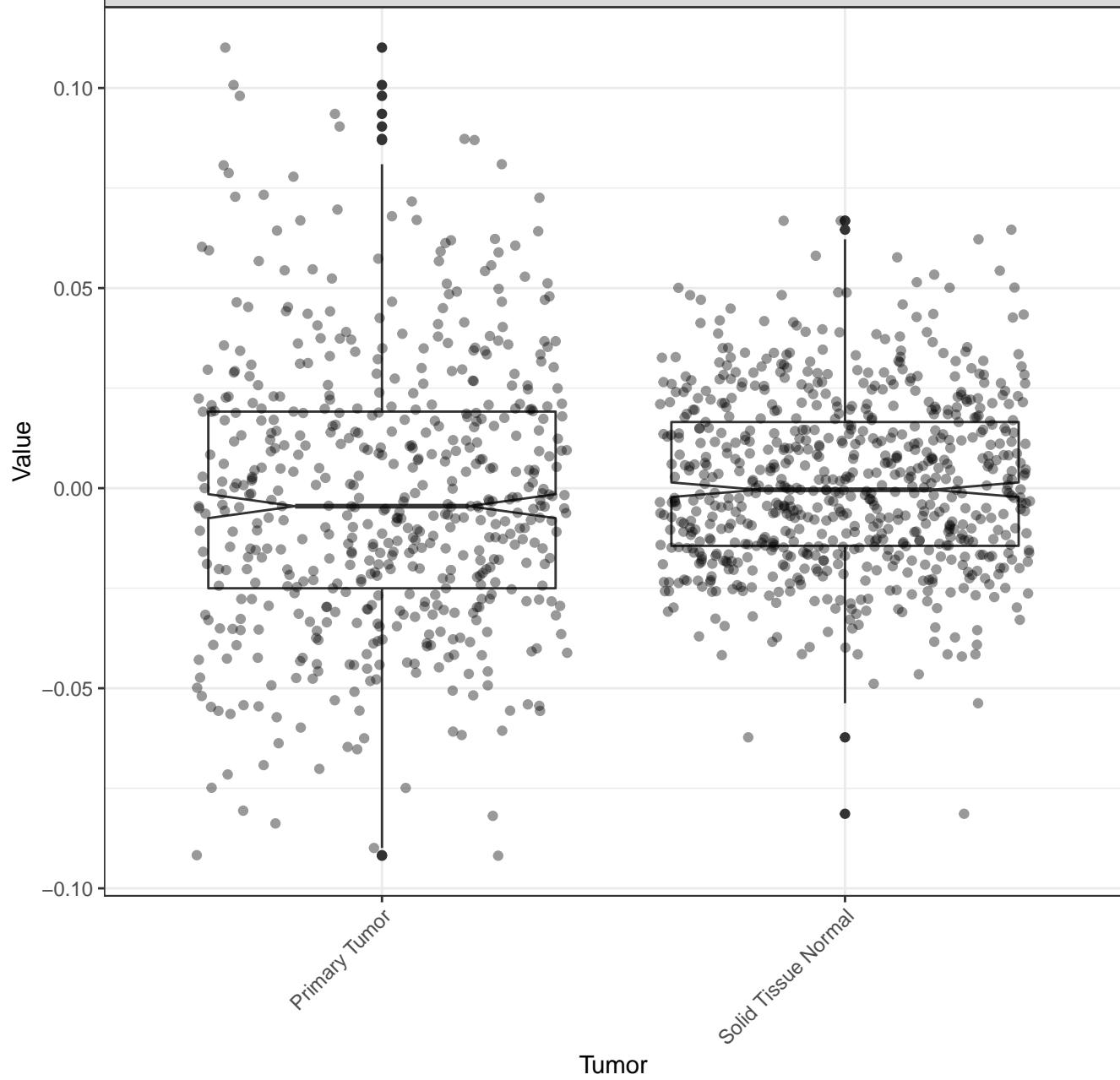
337,REACTOME_GPCR_DOWNSTREAM_SIGNALING



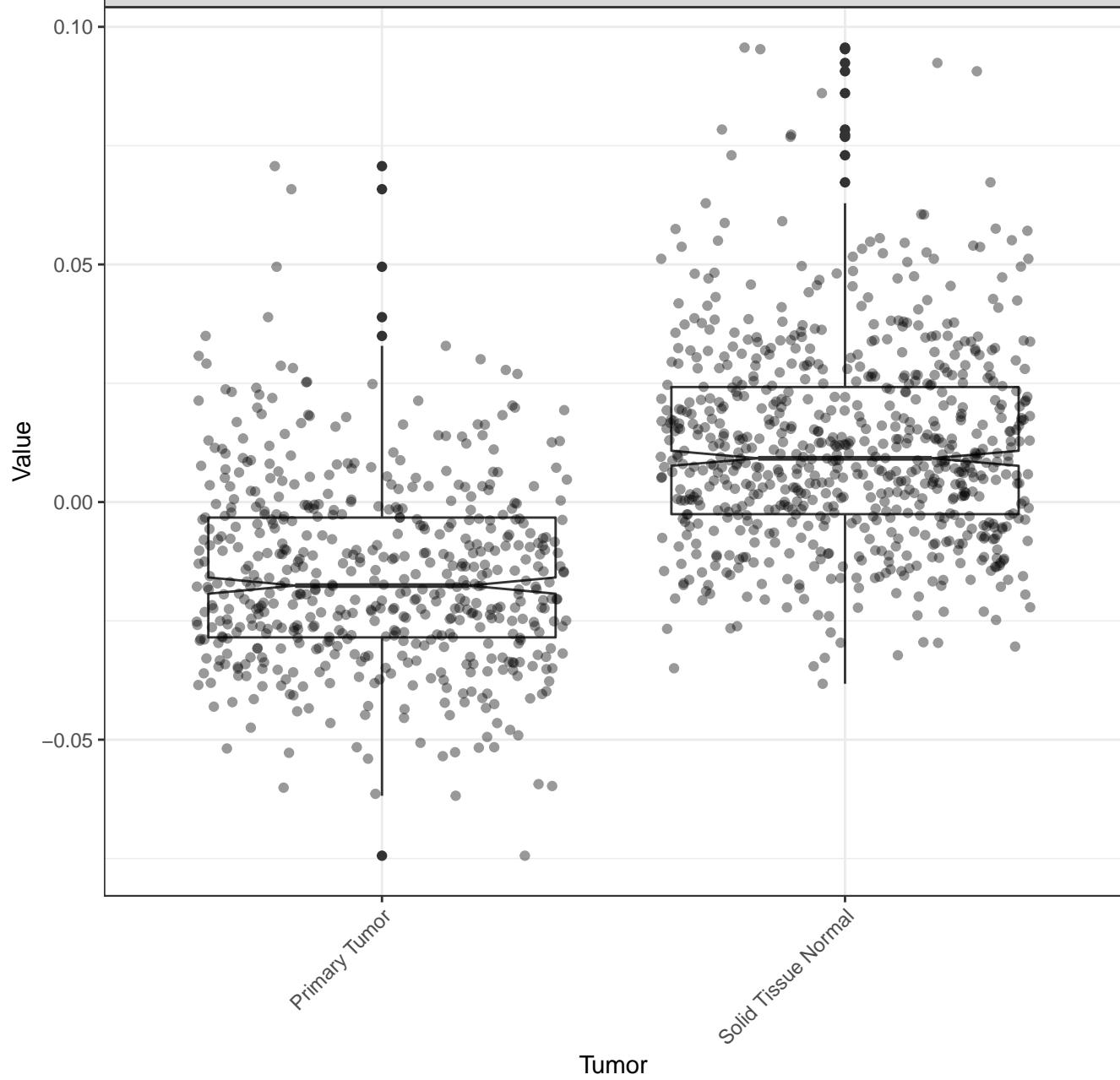
338,KEGG_SPLICEOSOME

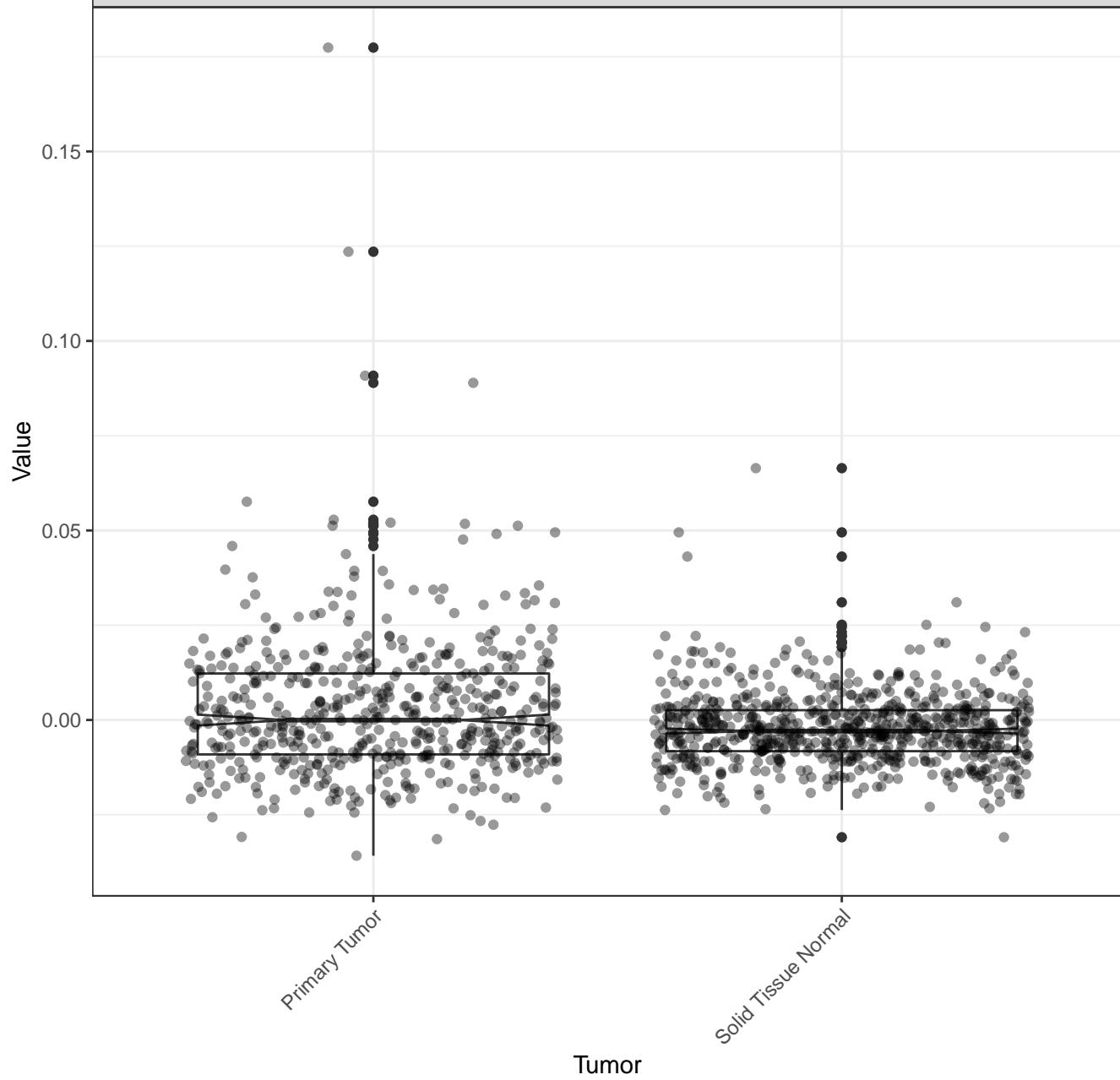


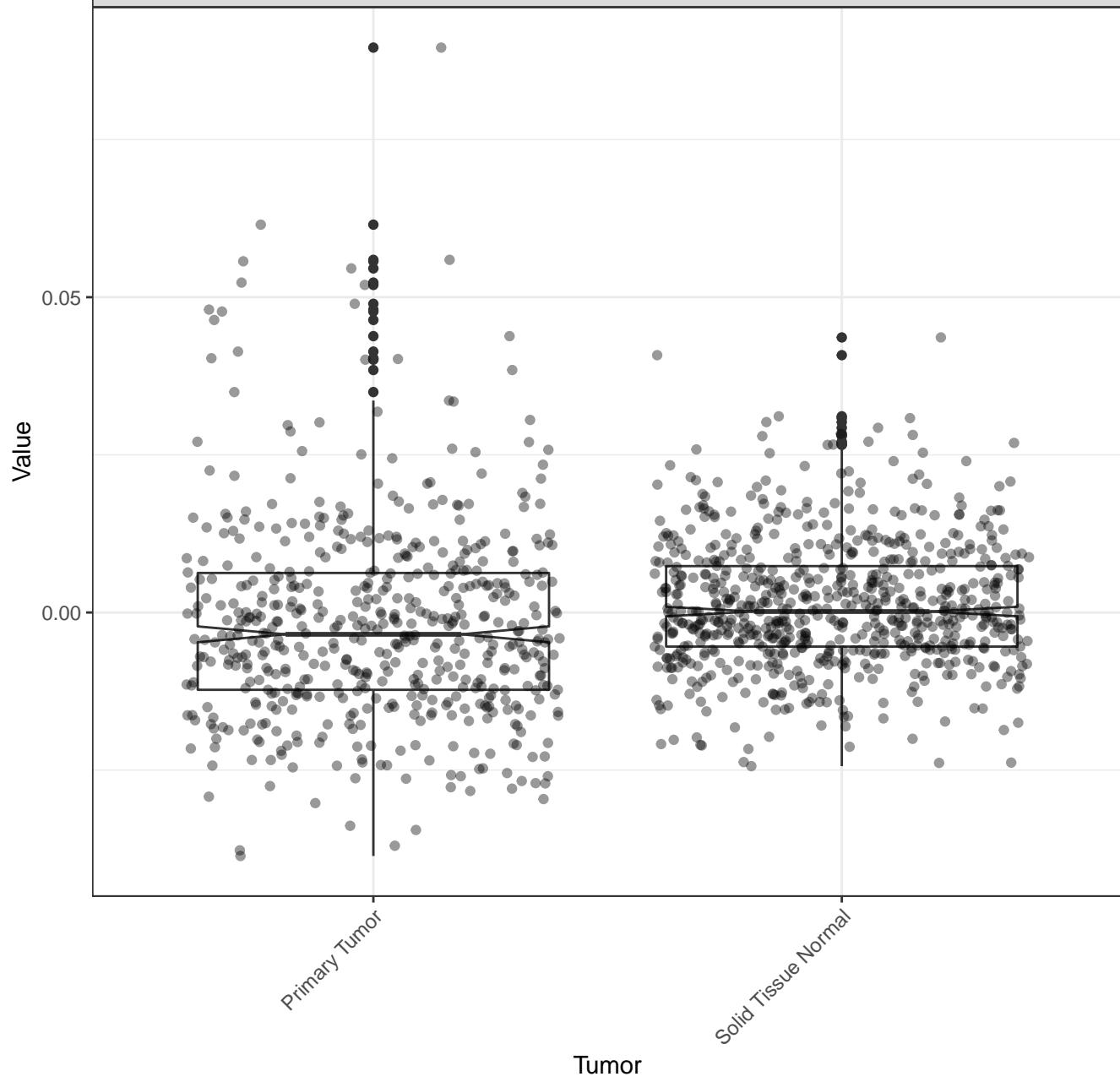
339,REACTOME_METABOLISM_OF_LIPIDS_AND_LIPOPROTEINS



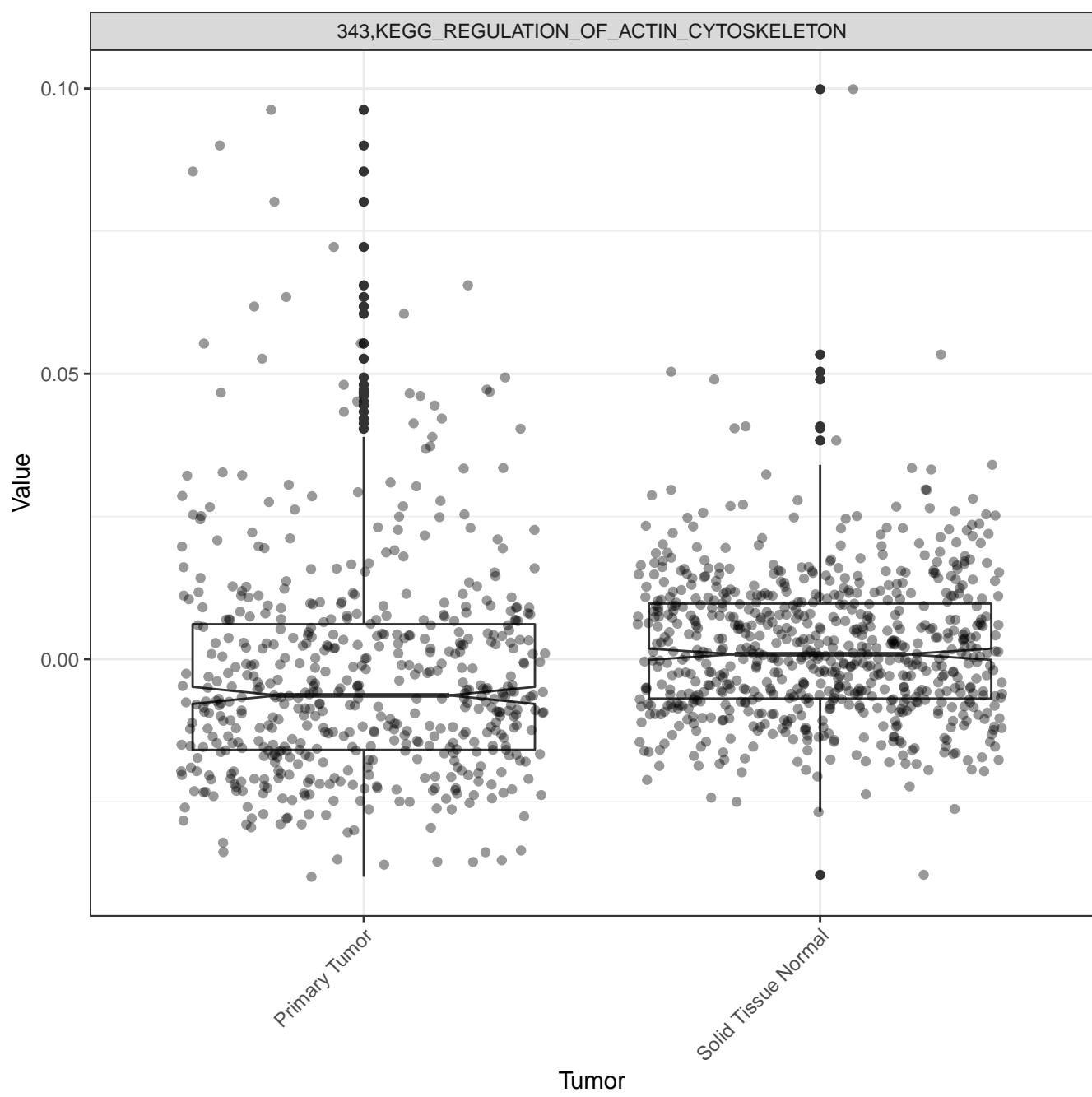
340,REACTOME_PLATELET_ACTIVATION_SIGNALING_AND_AGGREGATION



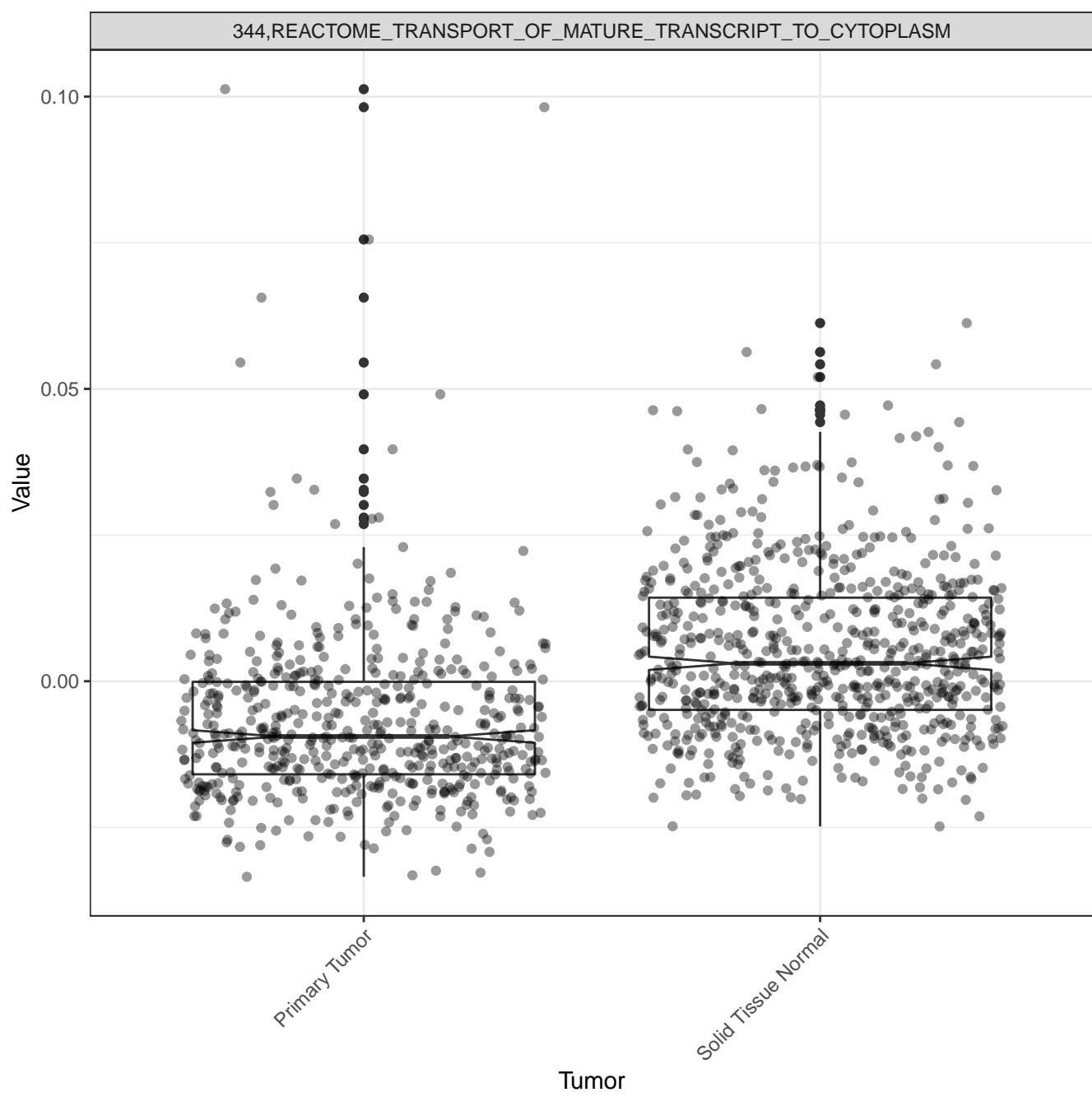




343,KEGG_REGULATION_OF_ACTIN_CYTOSKELETON

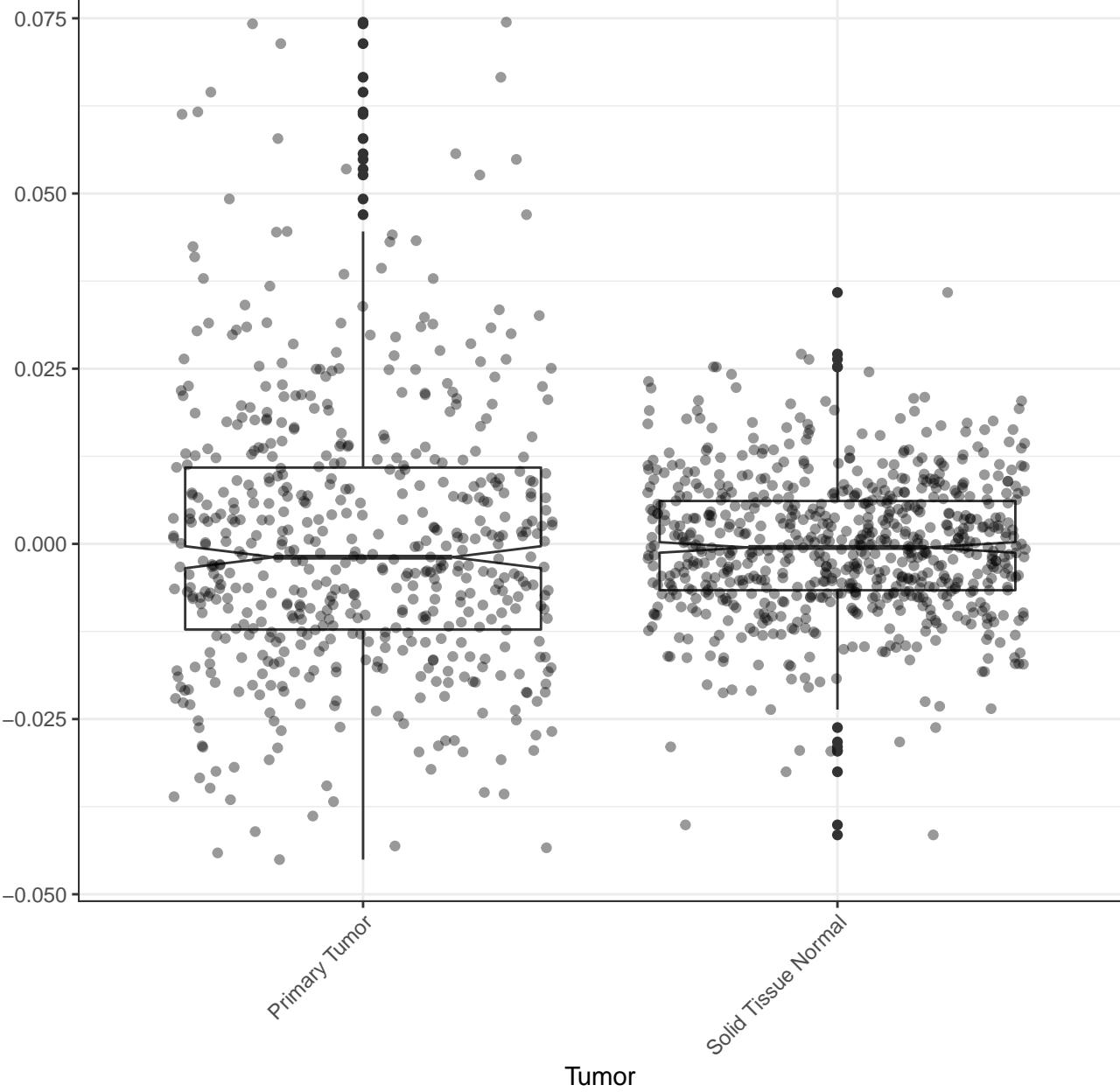


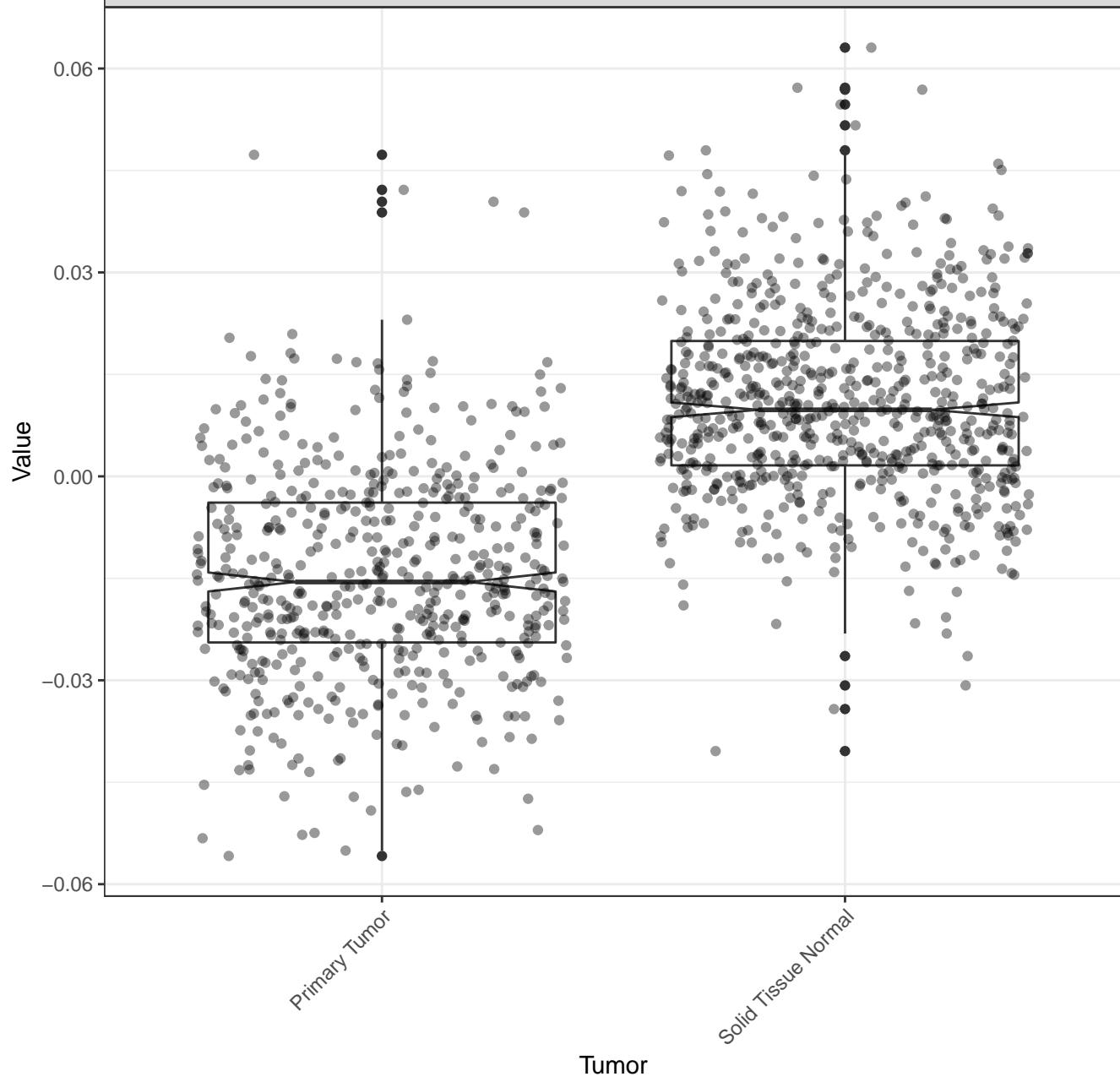
344,REACTOME_TRANSPORT_OF_MATURE_TRANSCRIPT_TO_CYTOPLASM

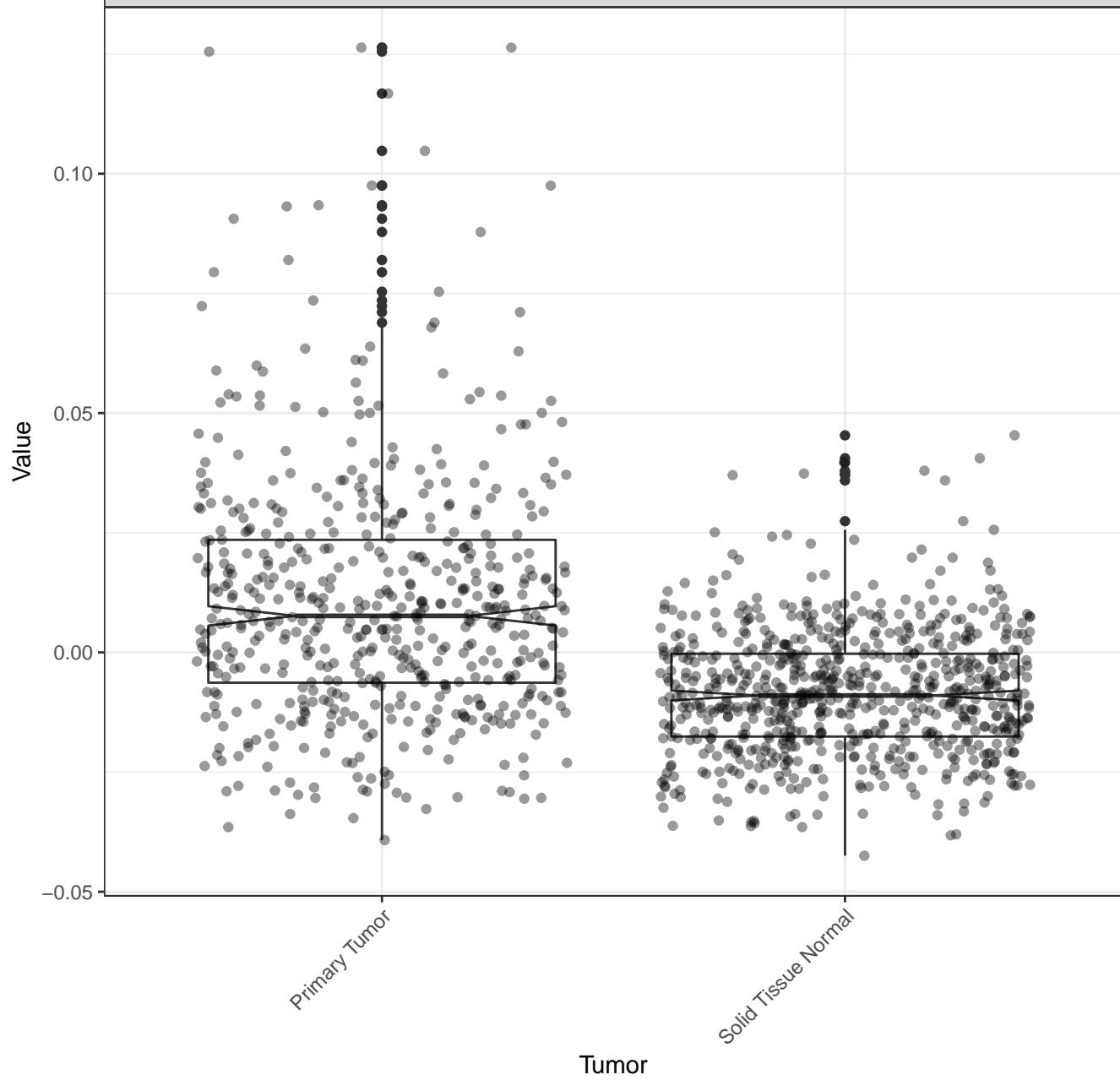


LV 345

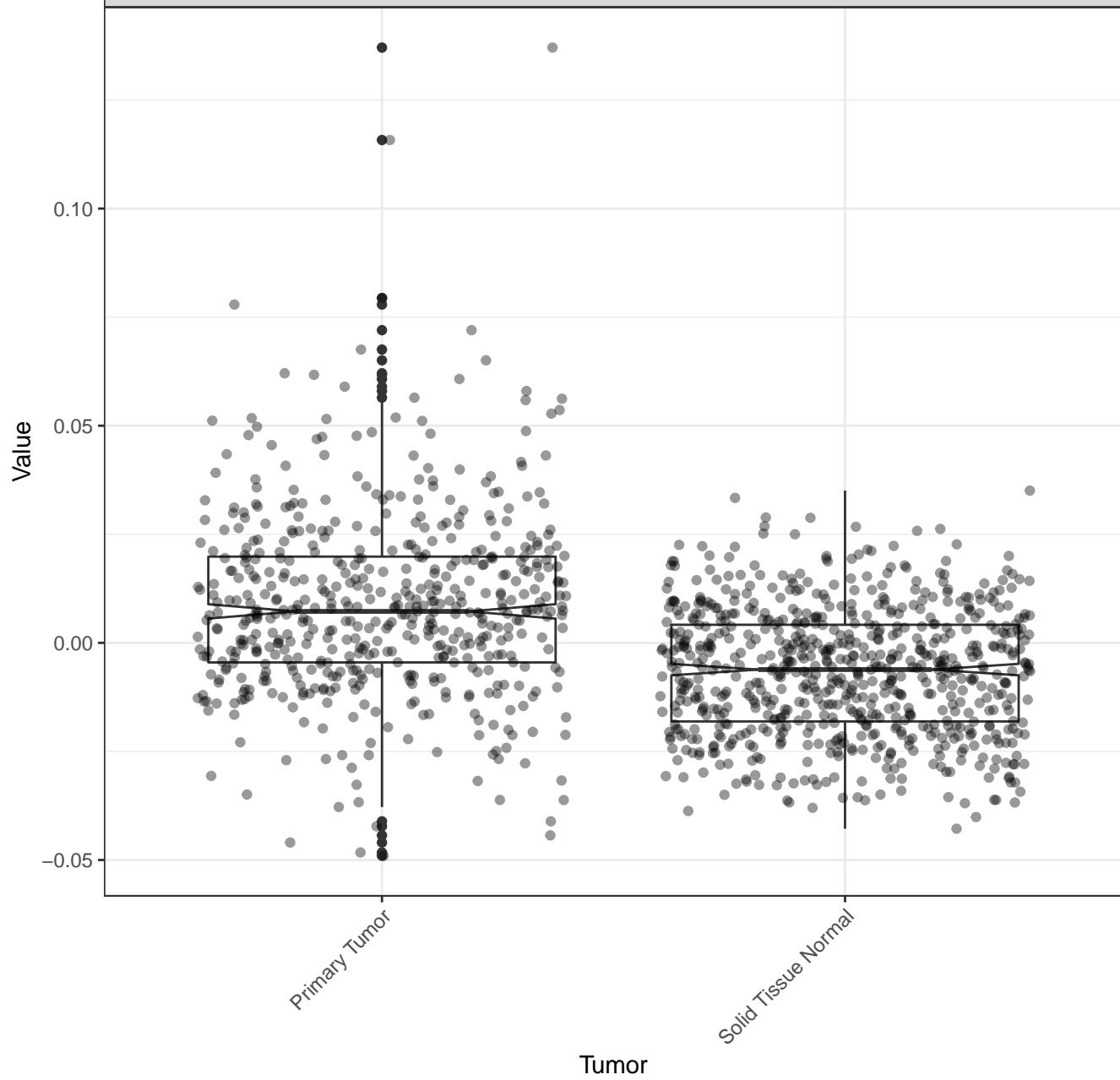
Value







348,KEGG_EPITHELIAL_CELL_SIGNALING_IN_Helicobacter_pyloRI_INFECTIOn



LV 349

Value

0.08

0.04

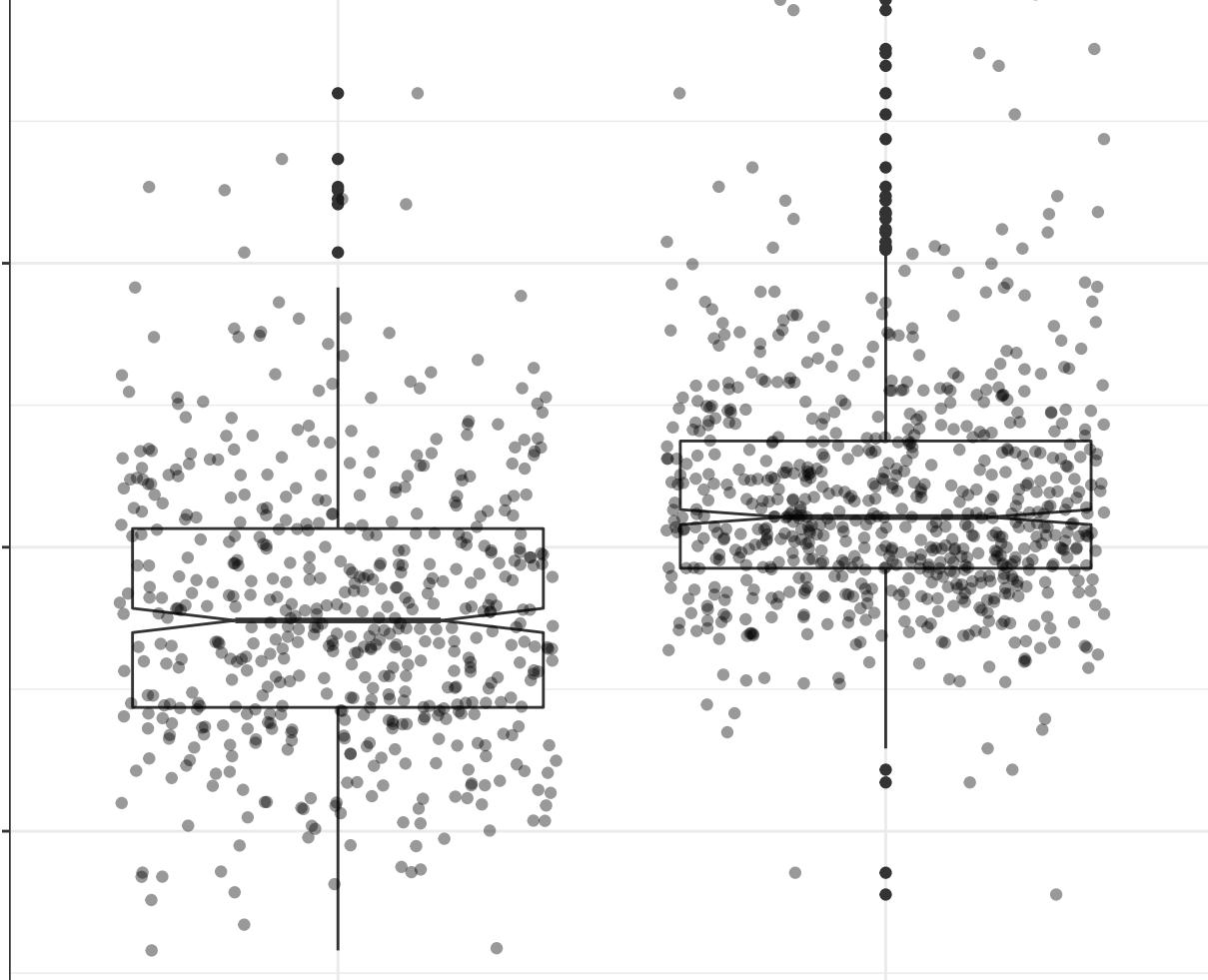
0.00

-0.04

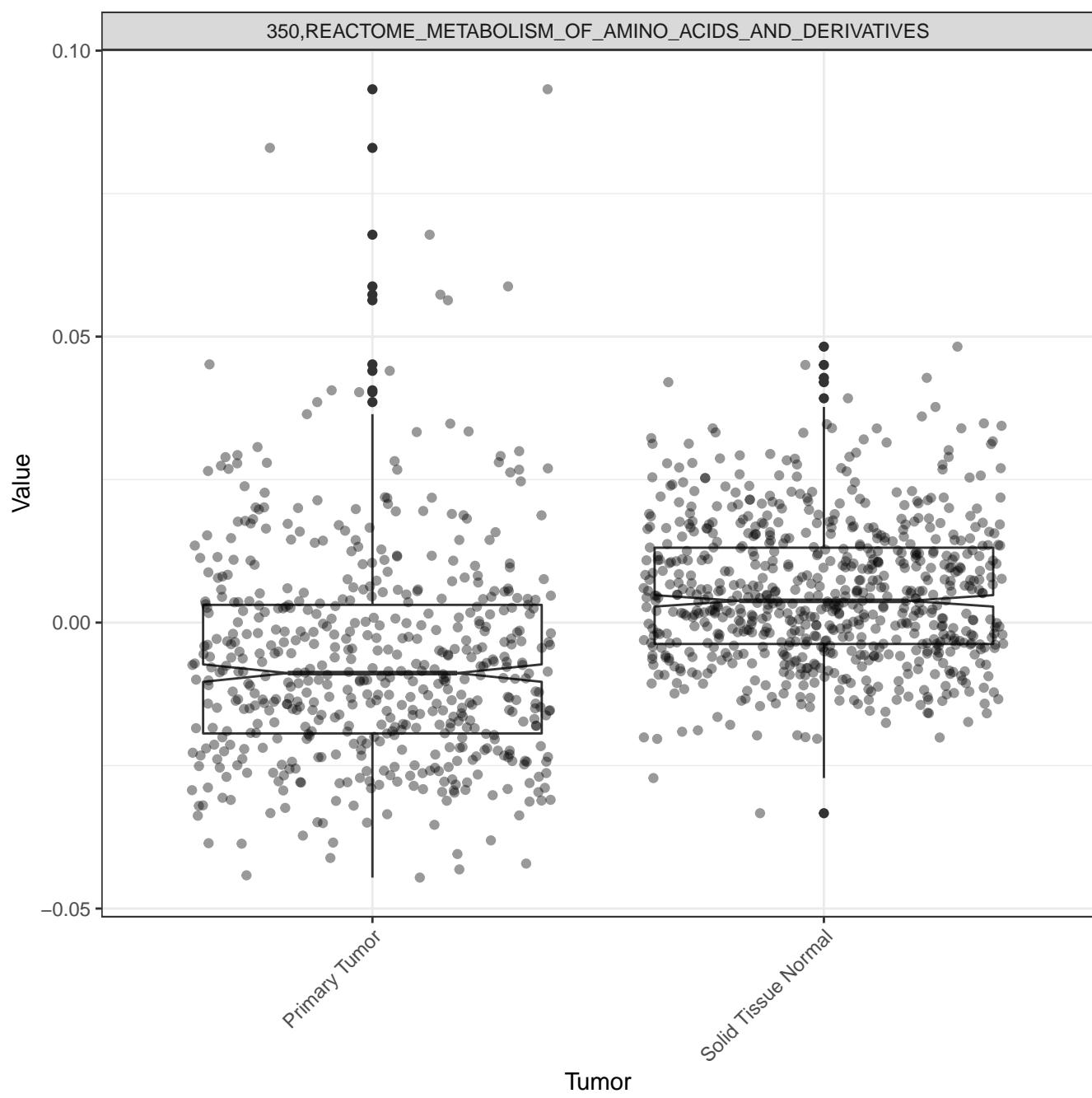
Primary Tumor

Solid Tissue Normal

Tumor



350,REACTOME_METABOLISM_OF_AMINO_ACIDS_AND_DERIVATIVES



351,REACTOME_METABOLISM_OF_MRNA

Value

0.1

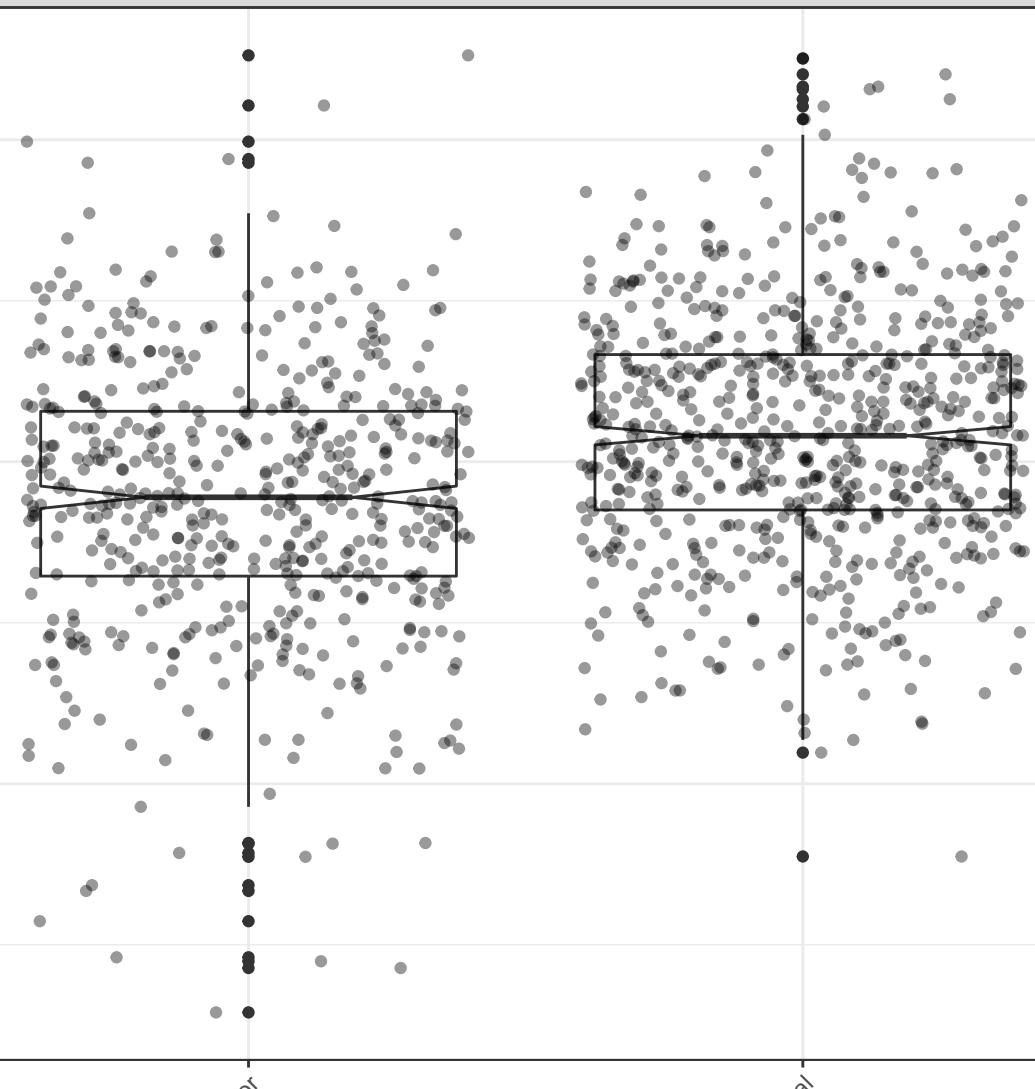
0.0

-0.1

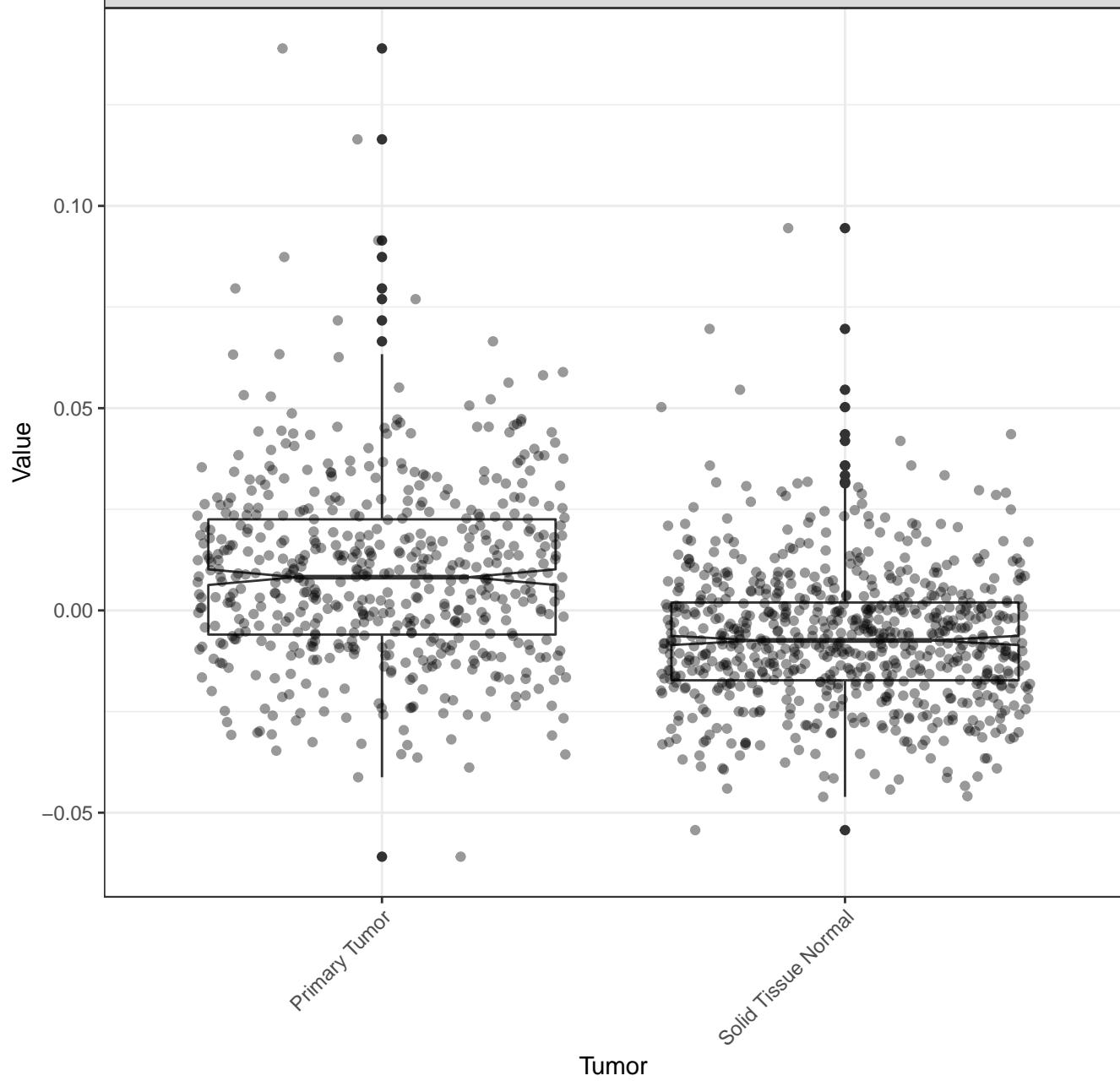
Primary Tumor

Solid Tissue Normal

Tumor



352, DMAP_DENDA1



Value

0.10

0.05

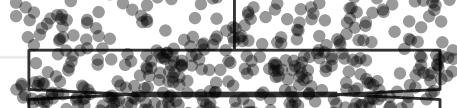
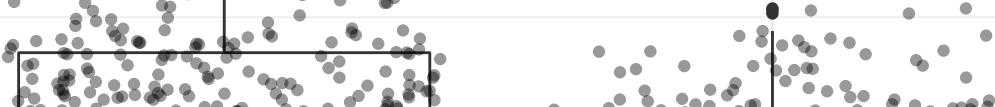
0.00

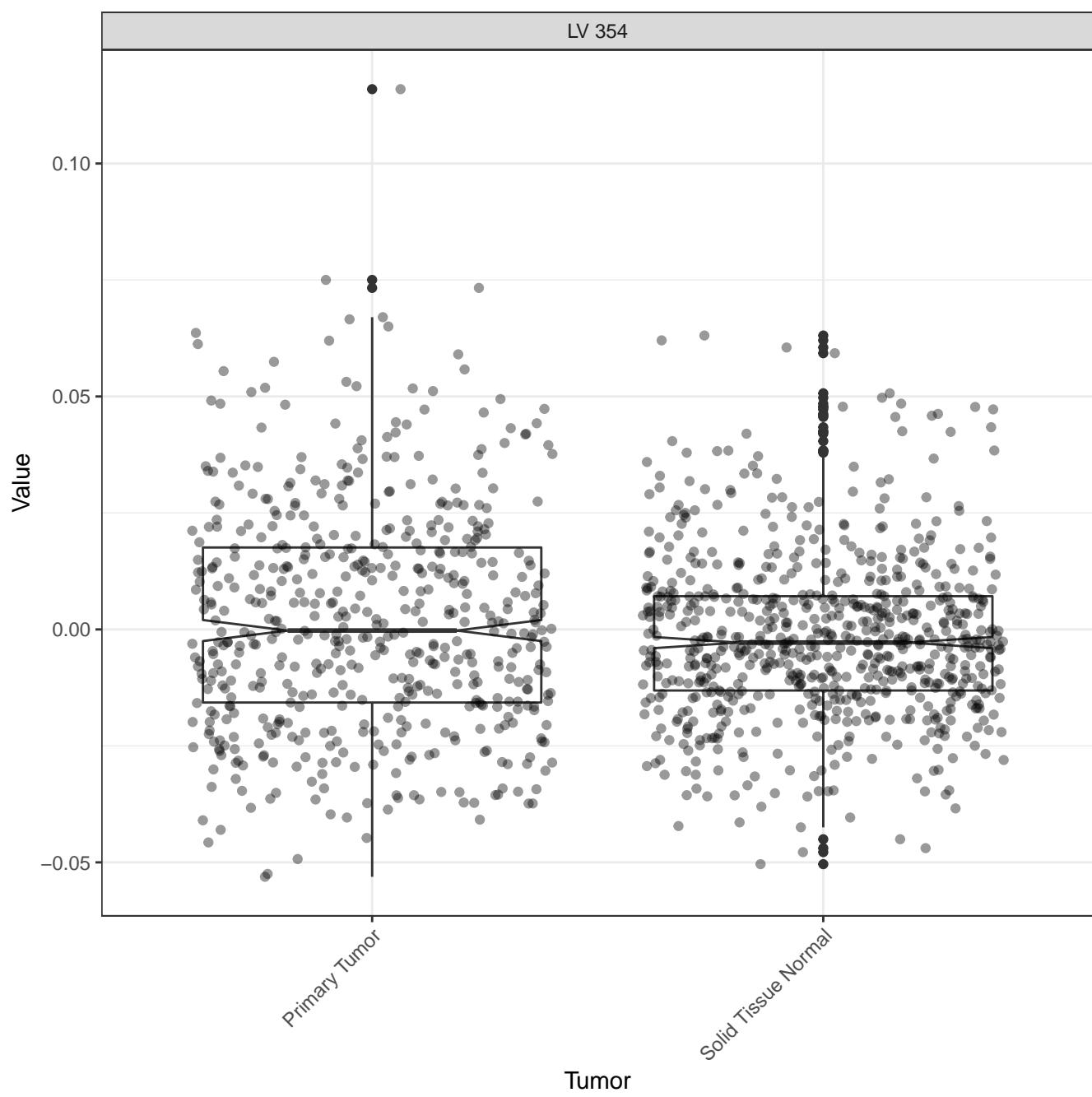
-0.05

Primary Tumor

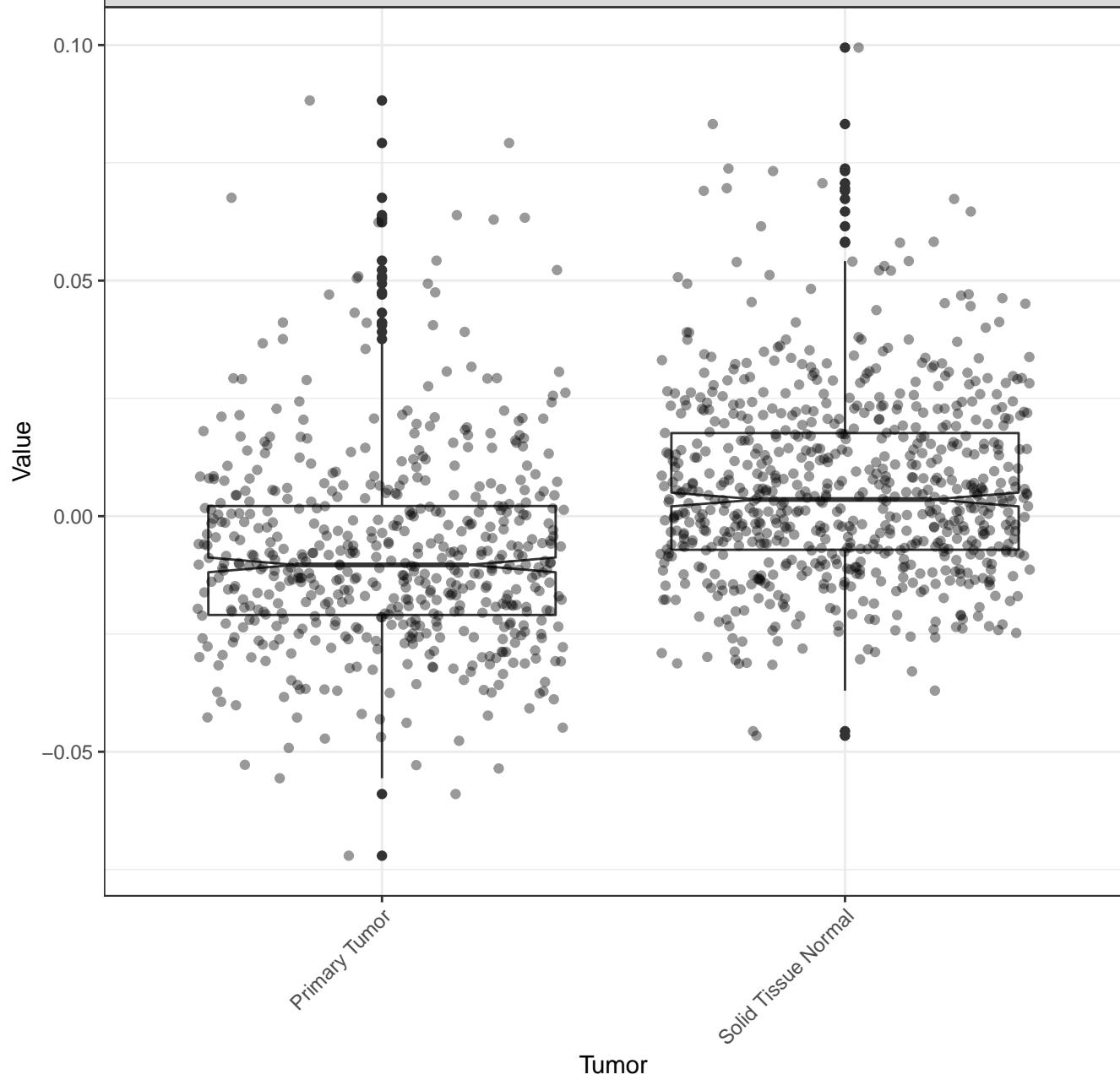
Solid Tissue Normal

Tumor

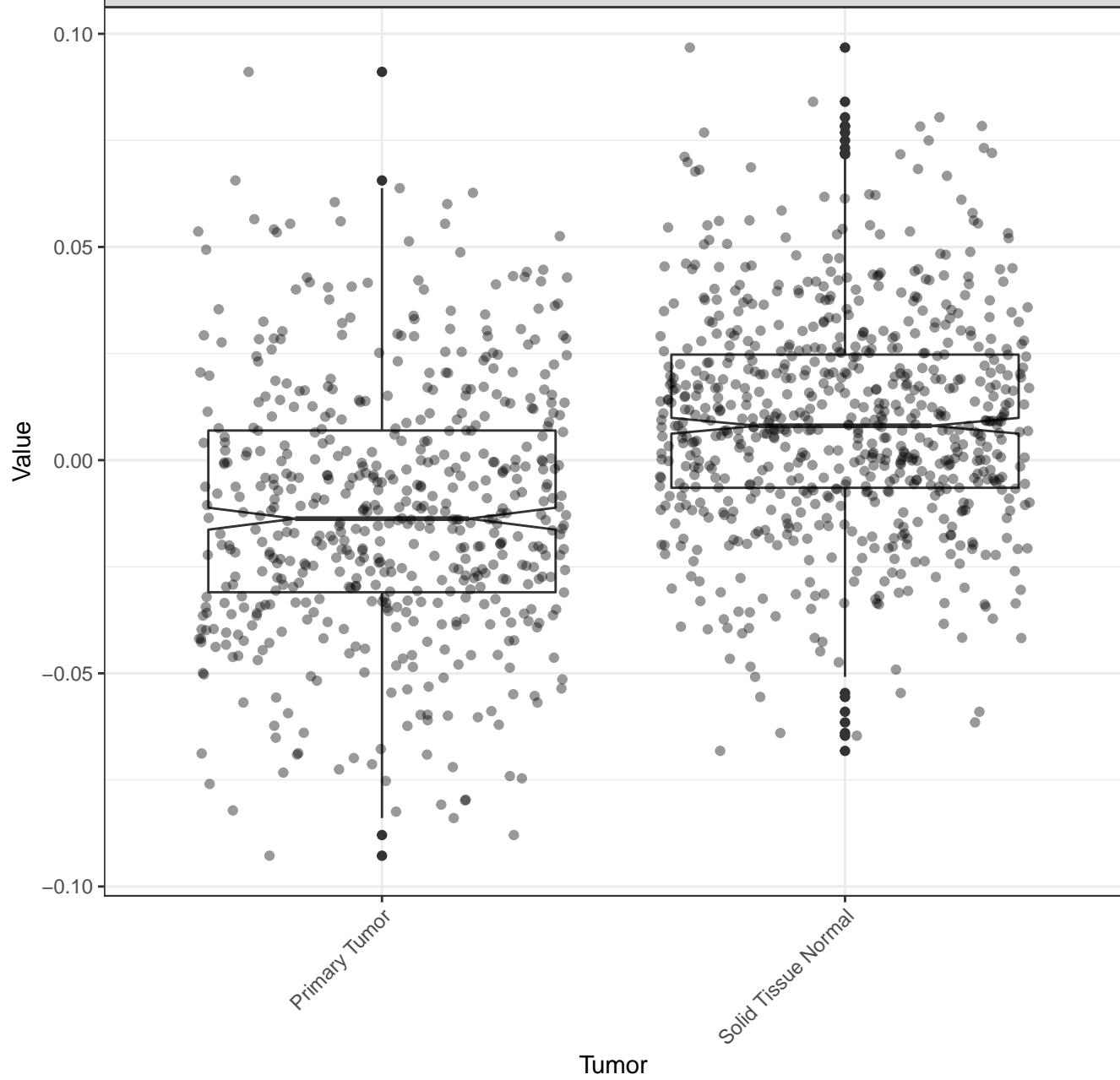




355,KEGG_HEDGEHOG_SIGNALING_PATHWAY



356,REACTOME DEVELOPMENTAL_BIOLOGY



LV 357

Value

0.10

0.05

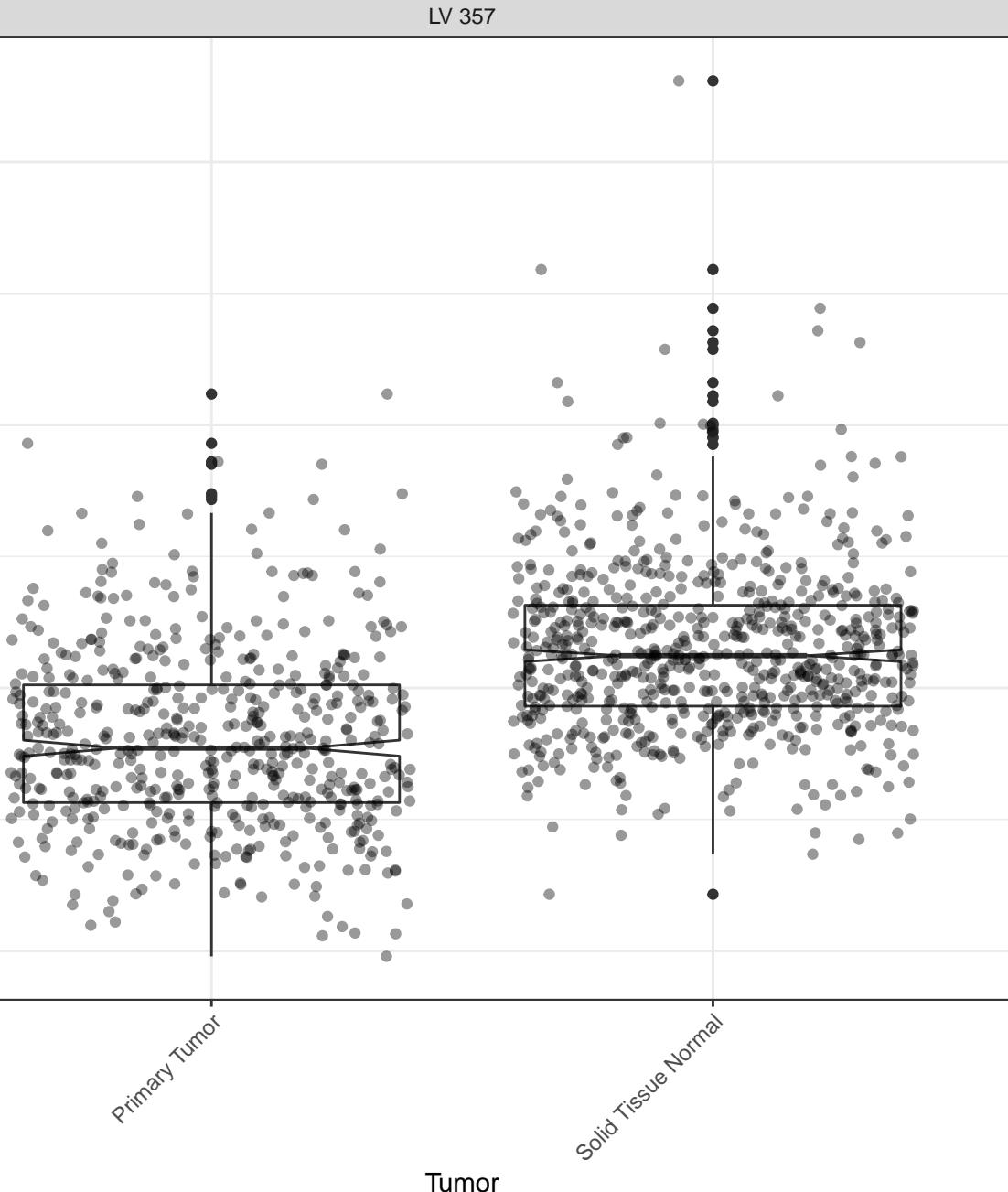
0.00

-0.05

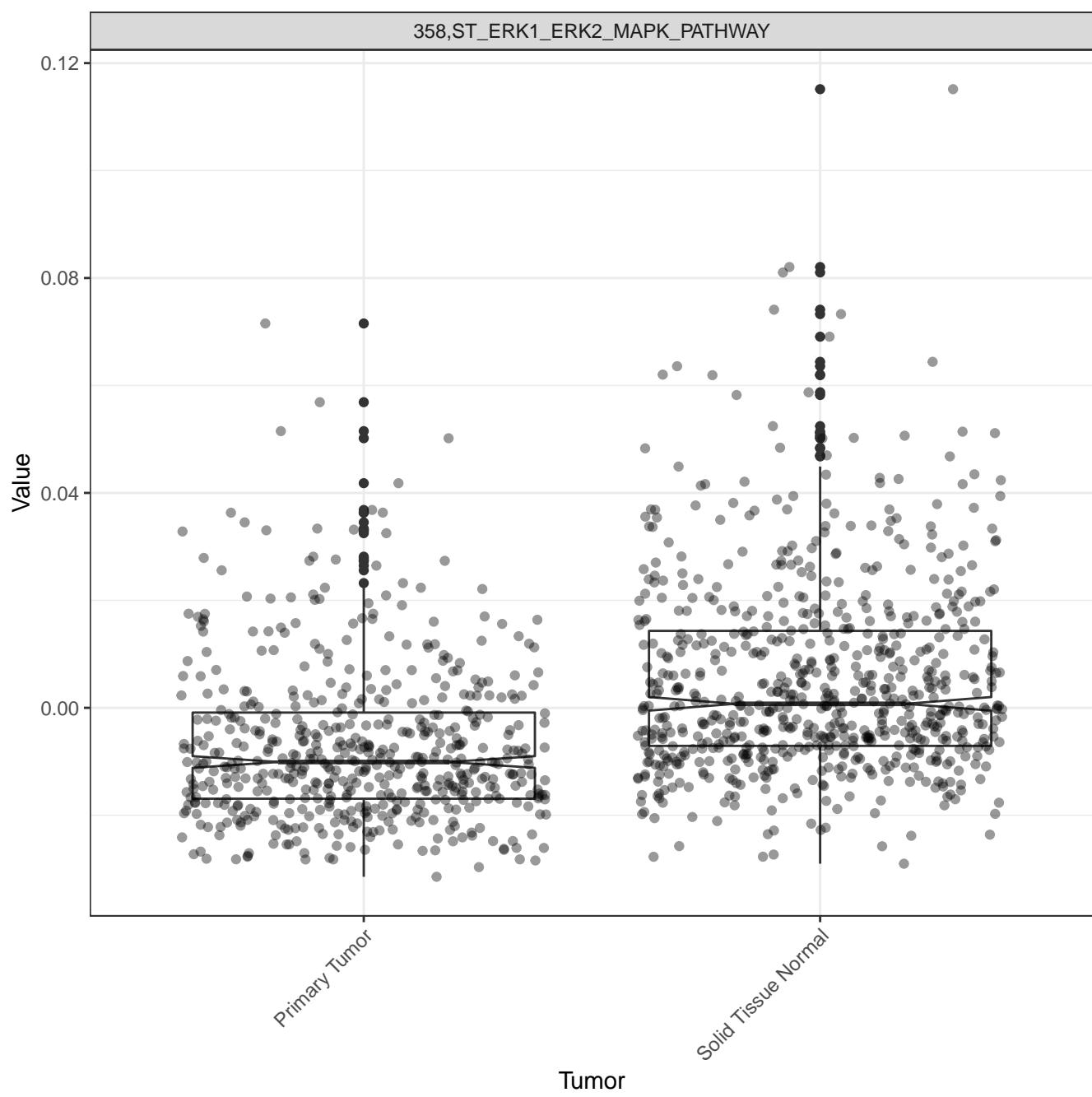
Primary Tumor

Solid Tissue Normal

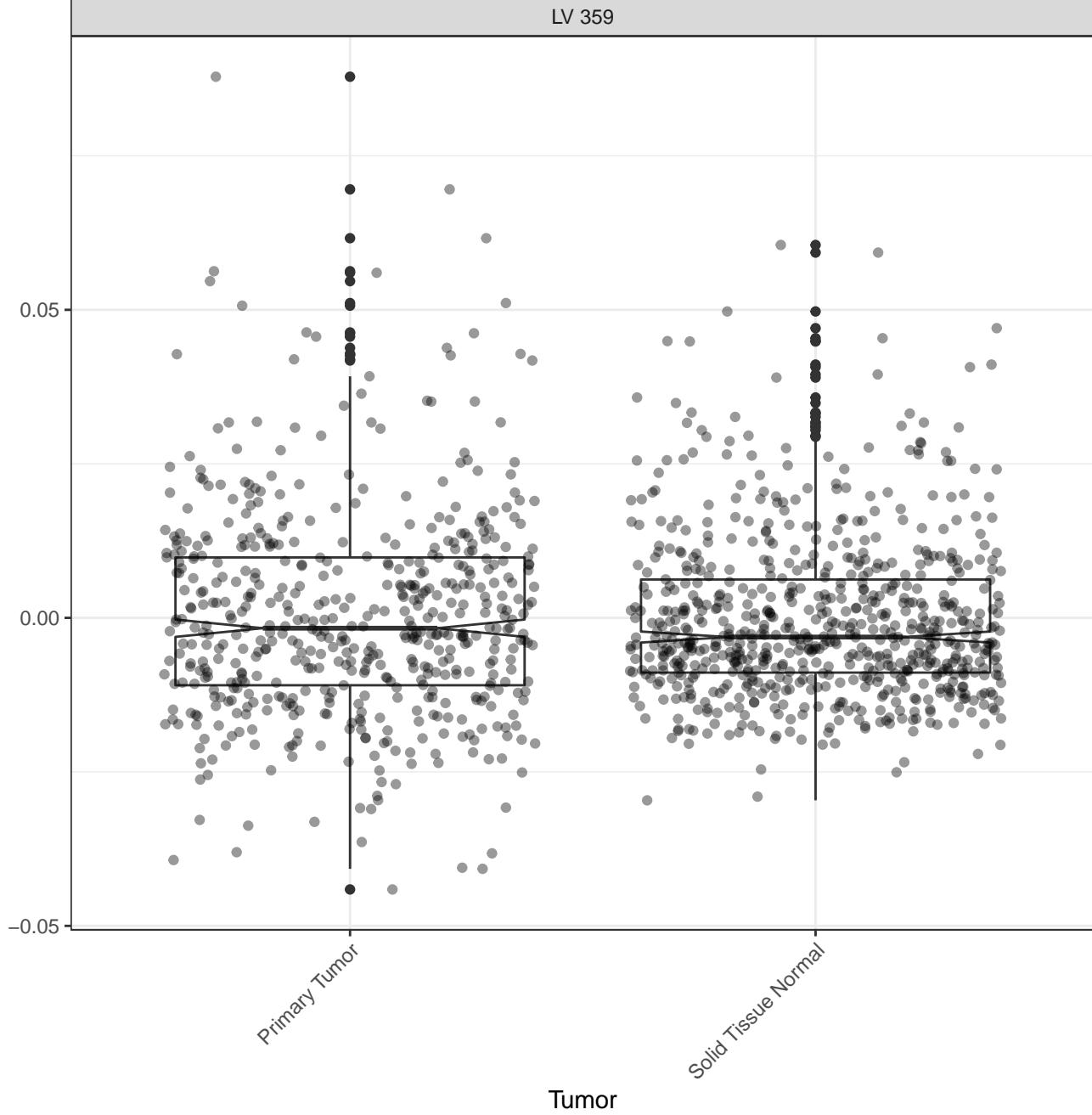
Tumor



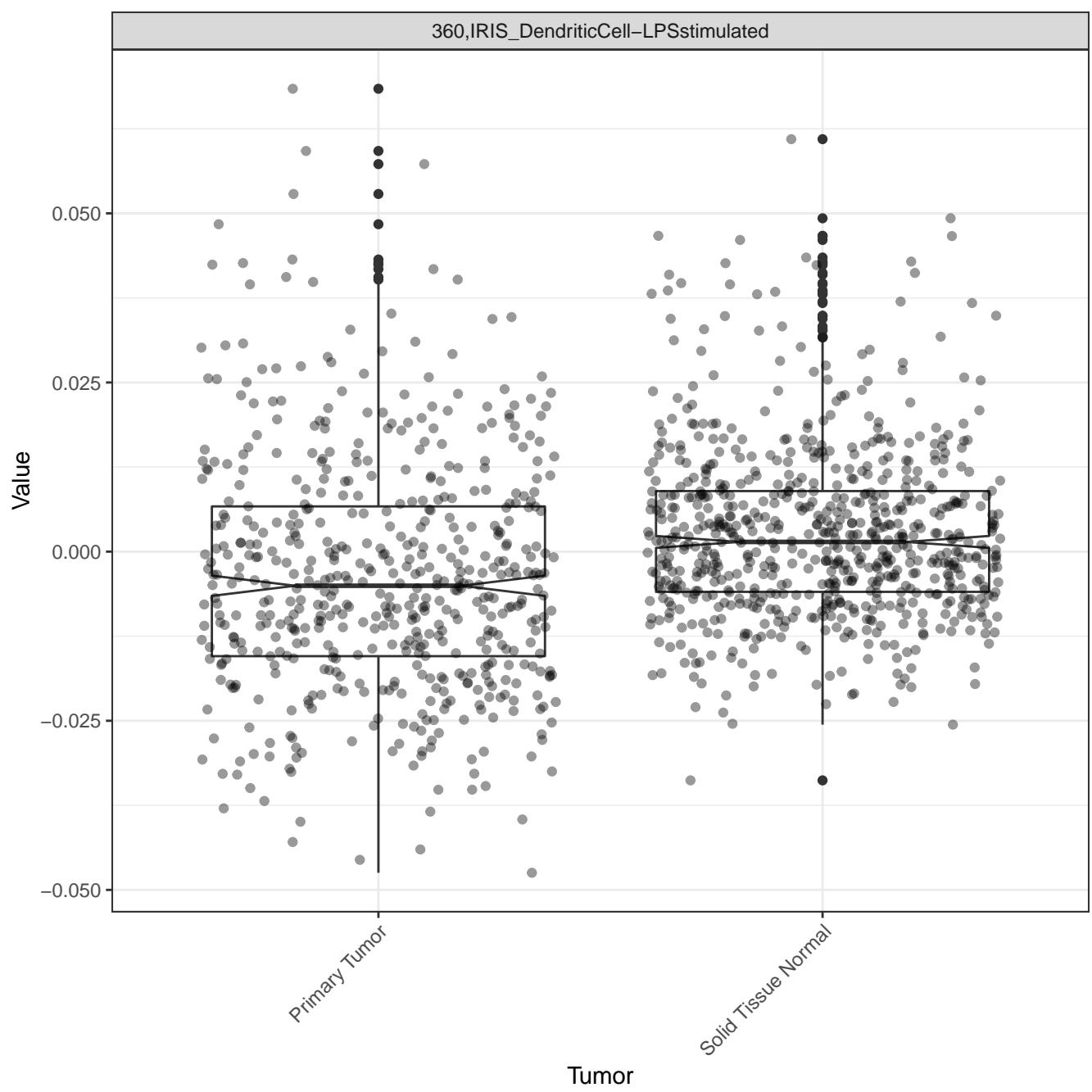
358,ST_ERK1_ERK2_MAPK_PATHWAY



Value



360,IRIS_DendriticCell-LPSstimulated



Value

0.10

0.05

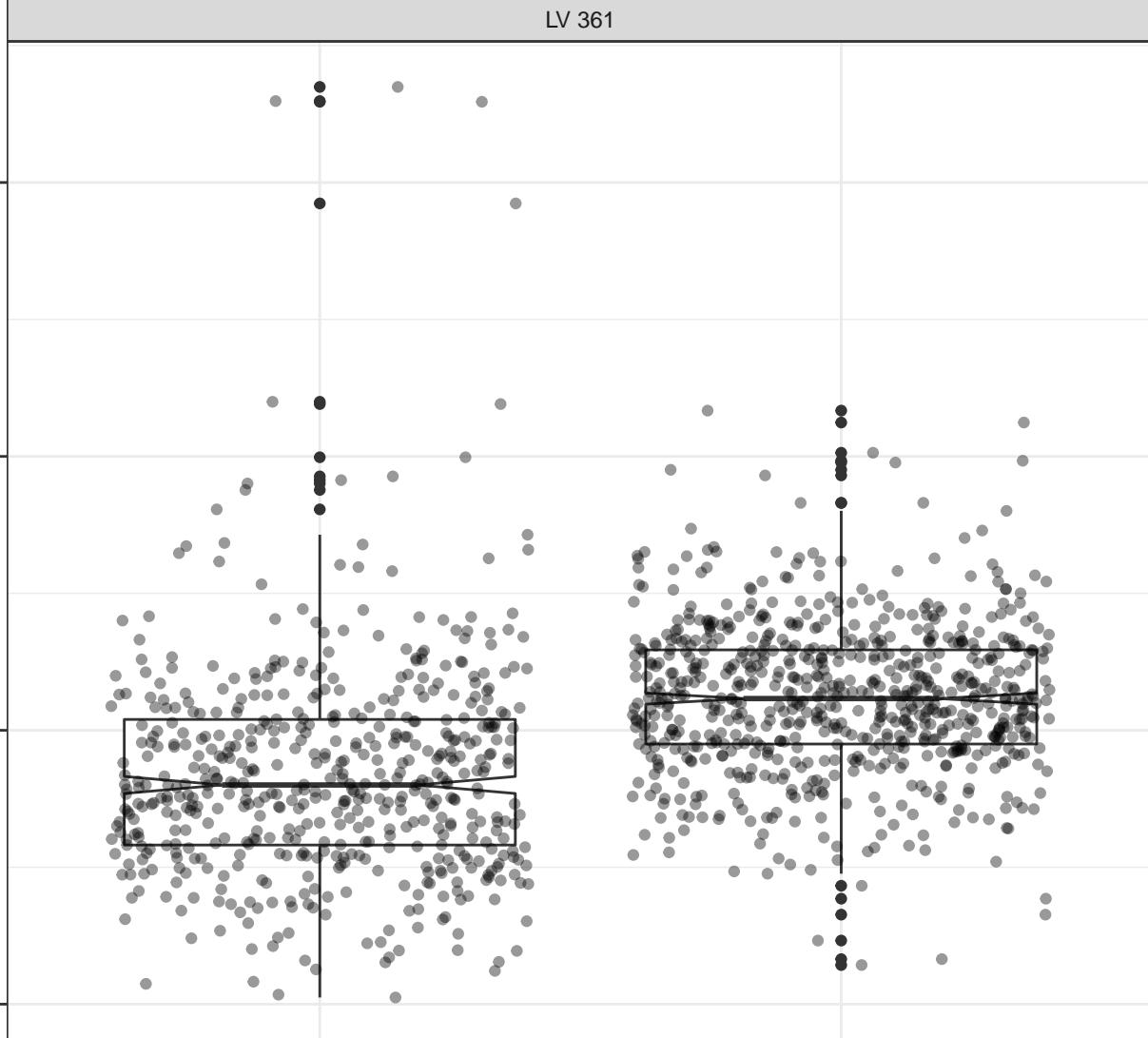
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.10

0.05

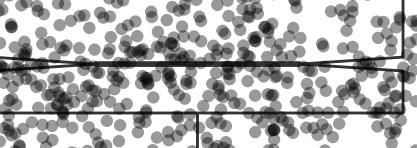
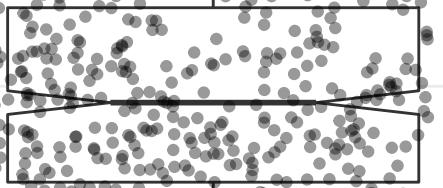
0.00

-0.05

Primary Tumor

Solid Tissue Normal

Tumor



LV 363

Value

0.20

0.15

0.10

0.05

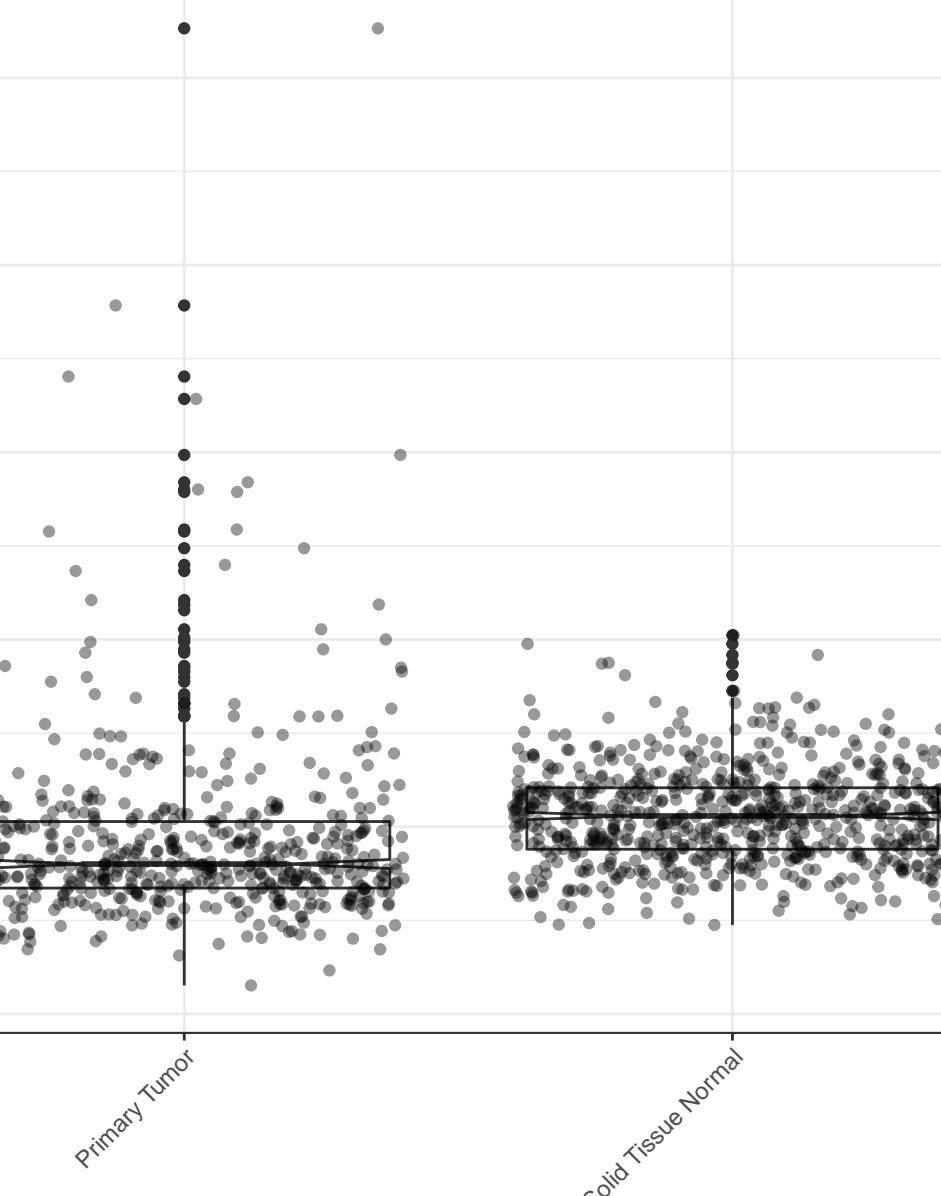
0.00

-0.05

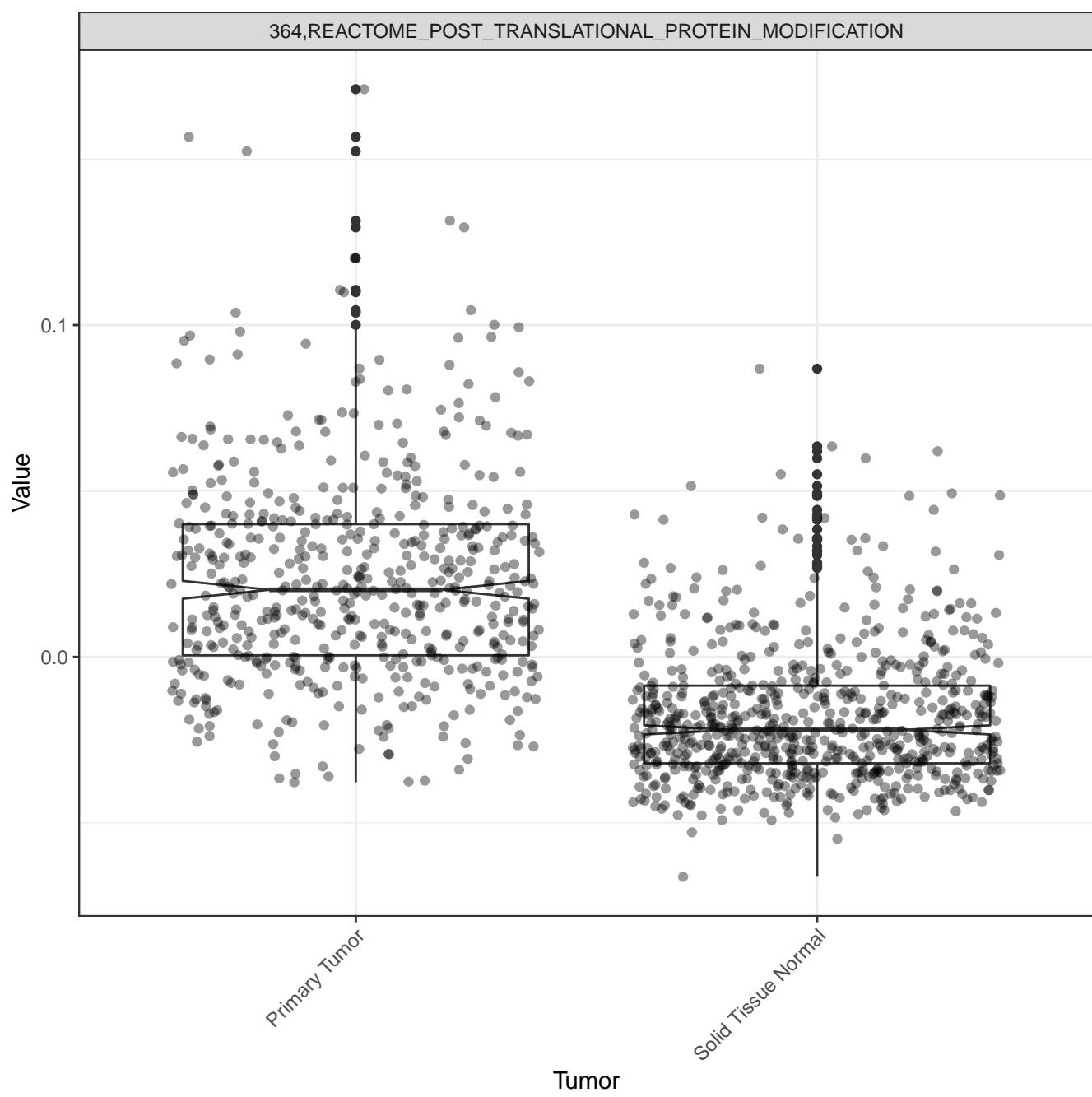
Primary Tumor

Solid Tissue Normal

Tumor



364,REACTOME_POST_TRANSLATIONAL_PROTEIN_MODIFICATION



Value

0.15

0.10

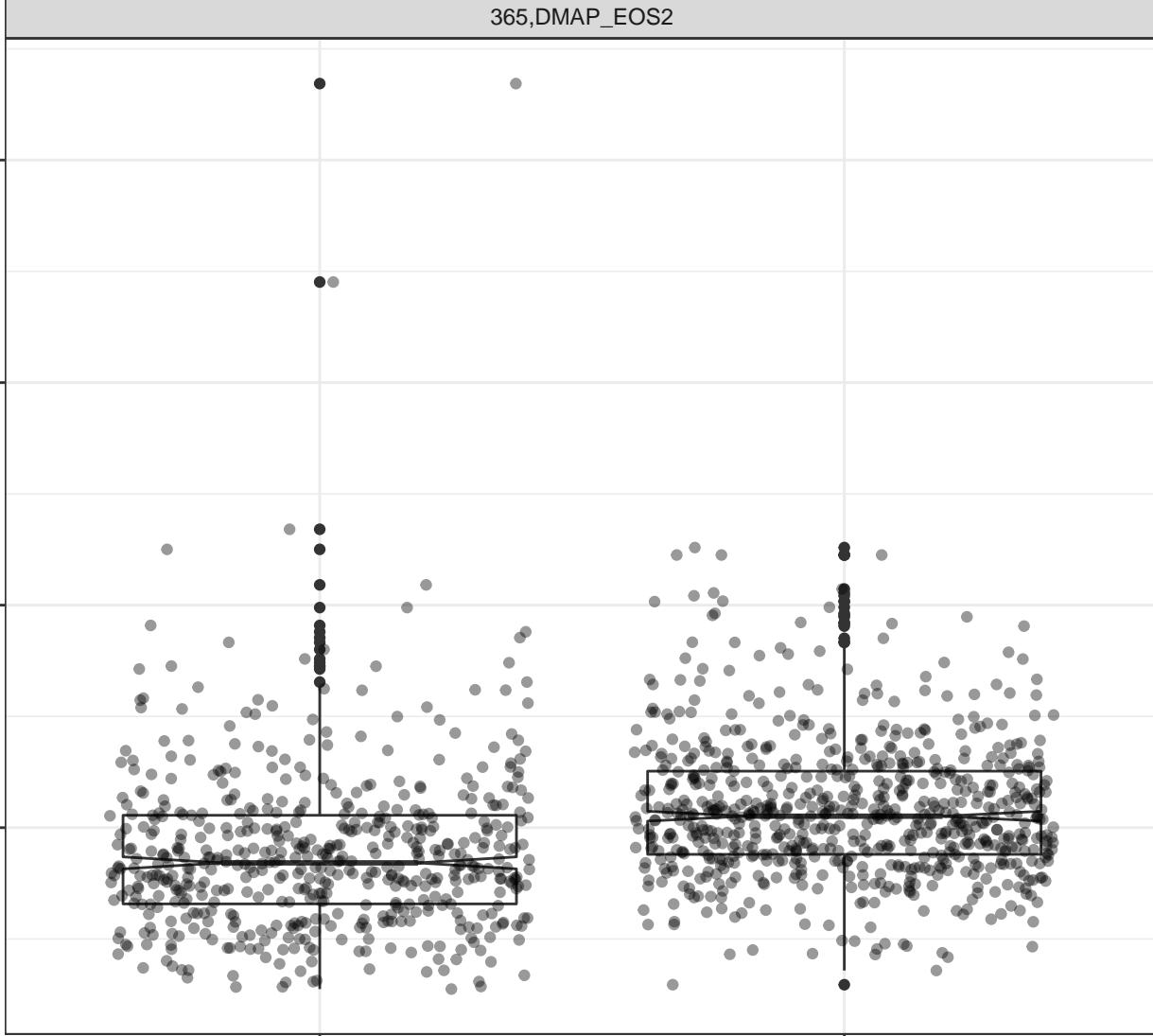
0.05

0.00

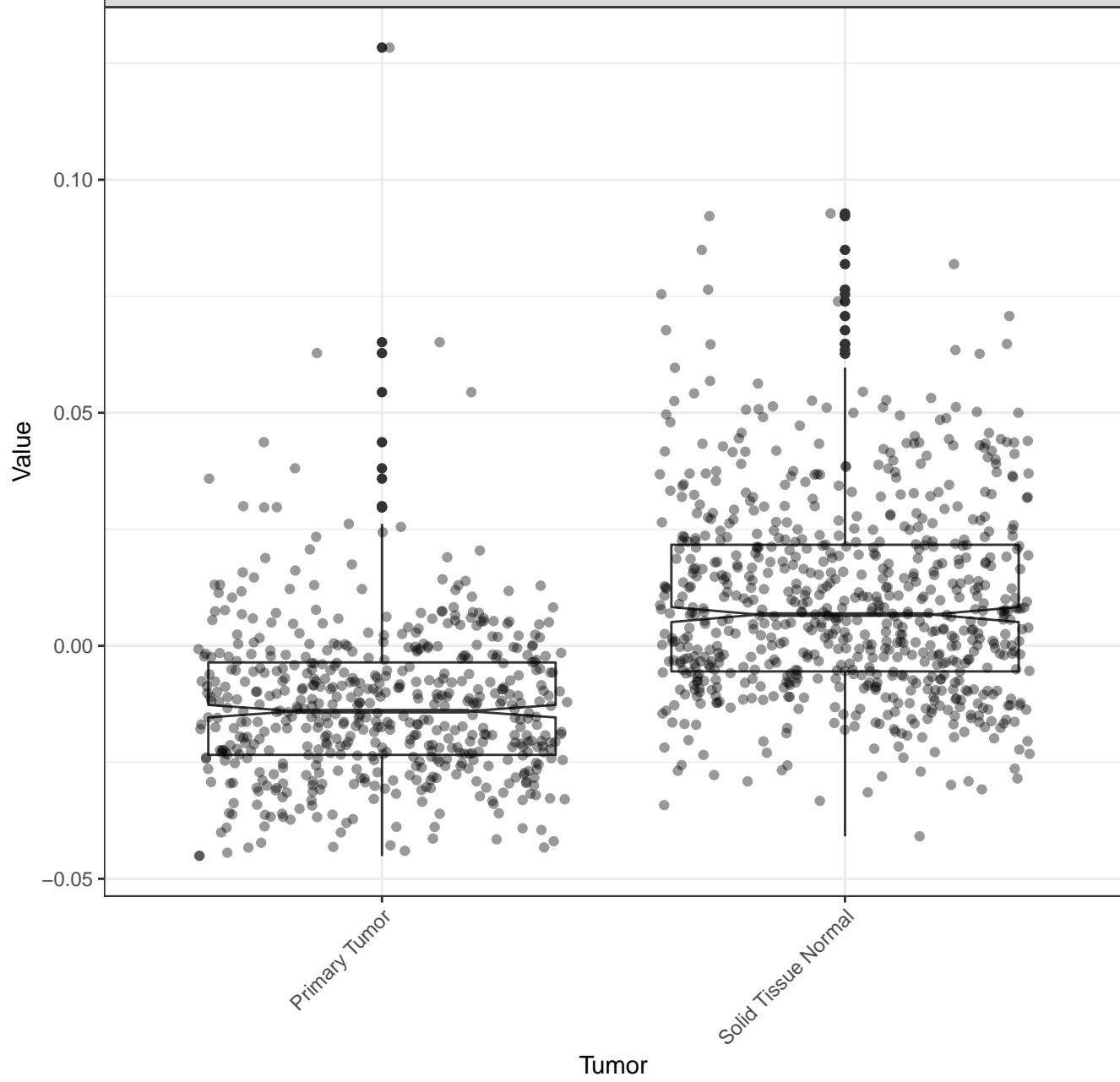
Primary Tumor

Solid Tissue Normal

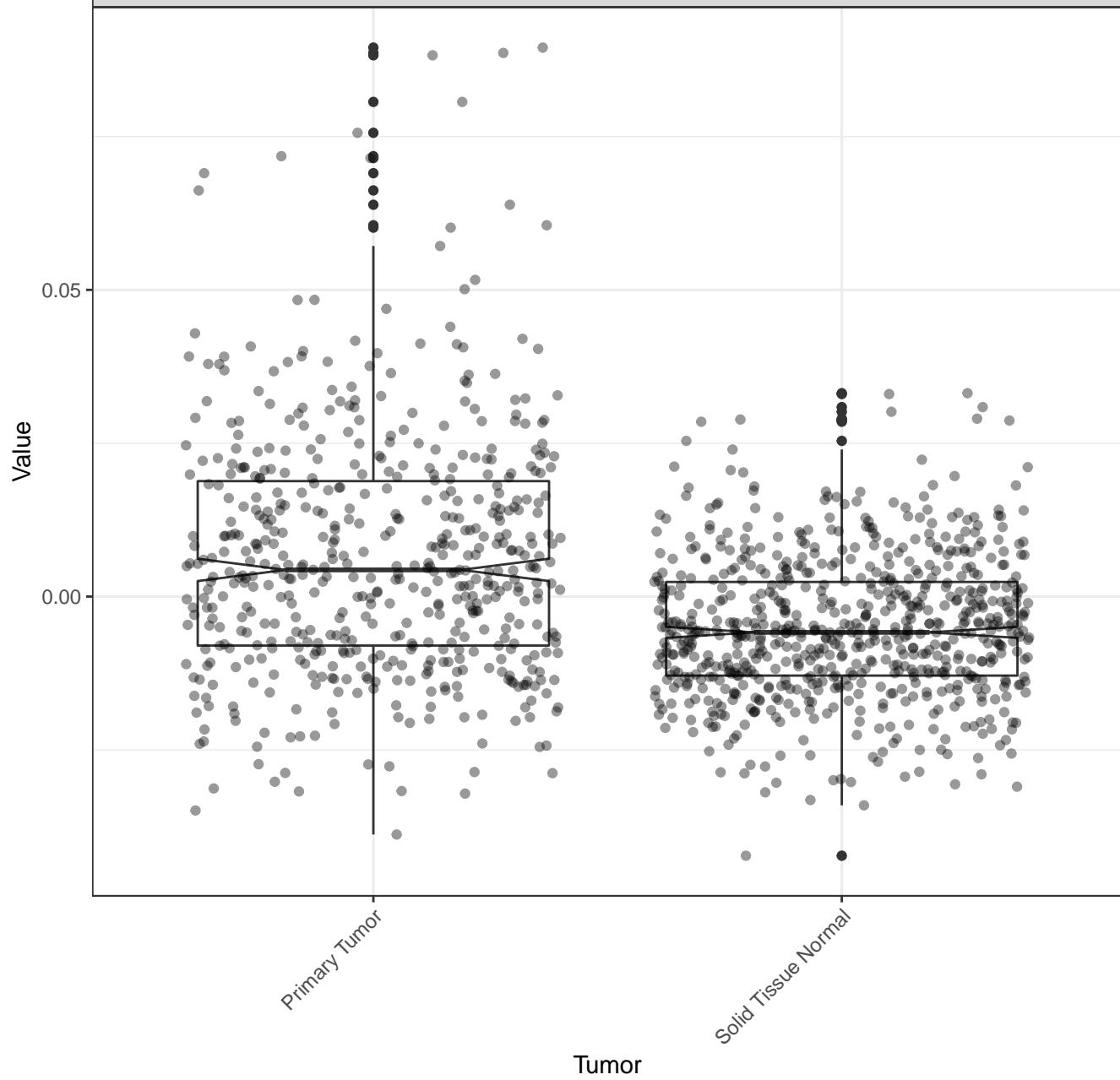
Tumor



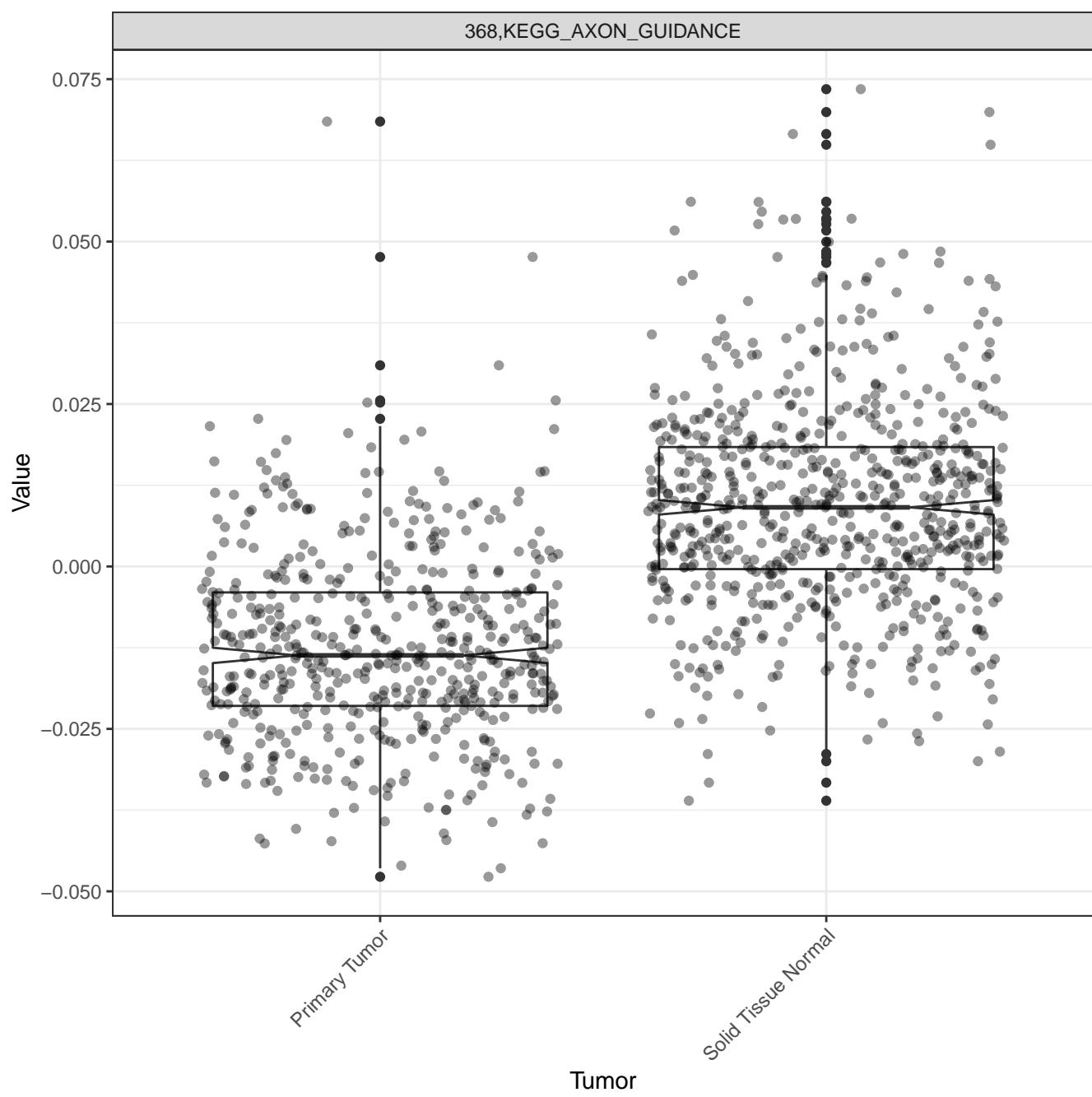
LV 366



367,PID_FANCONI_PATHWAY



368,KEGG_AXON_GUIDANCE



369,DMAP_MONO1

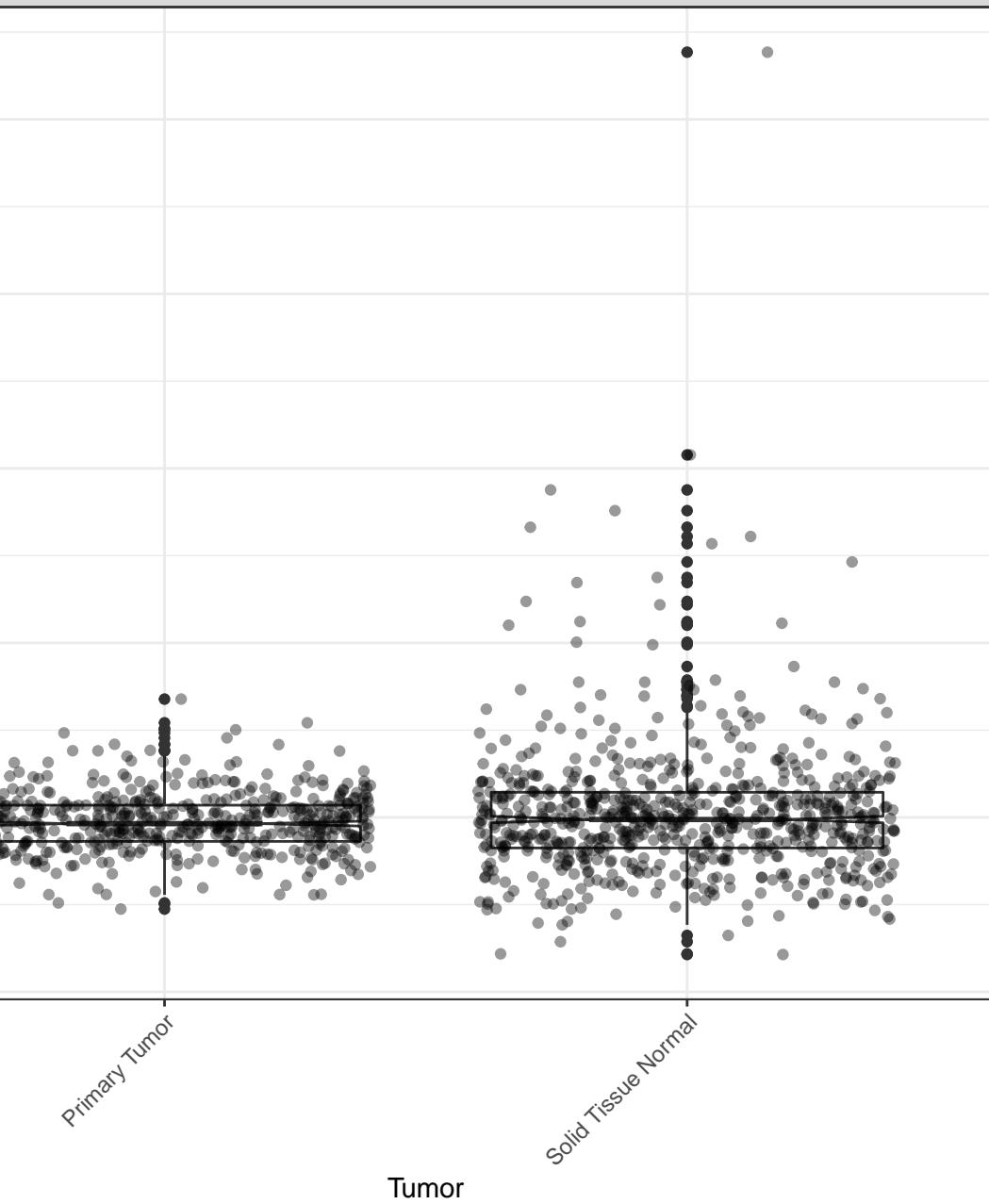
Value

0.4
0.3
0.2
0.1
0.0
-0.1

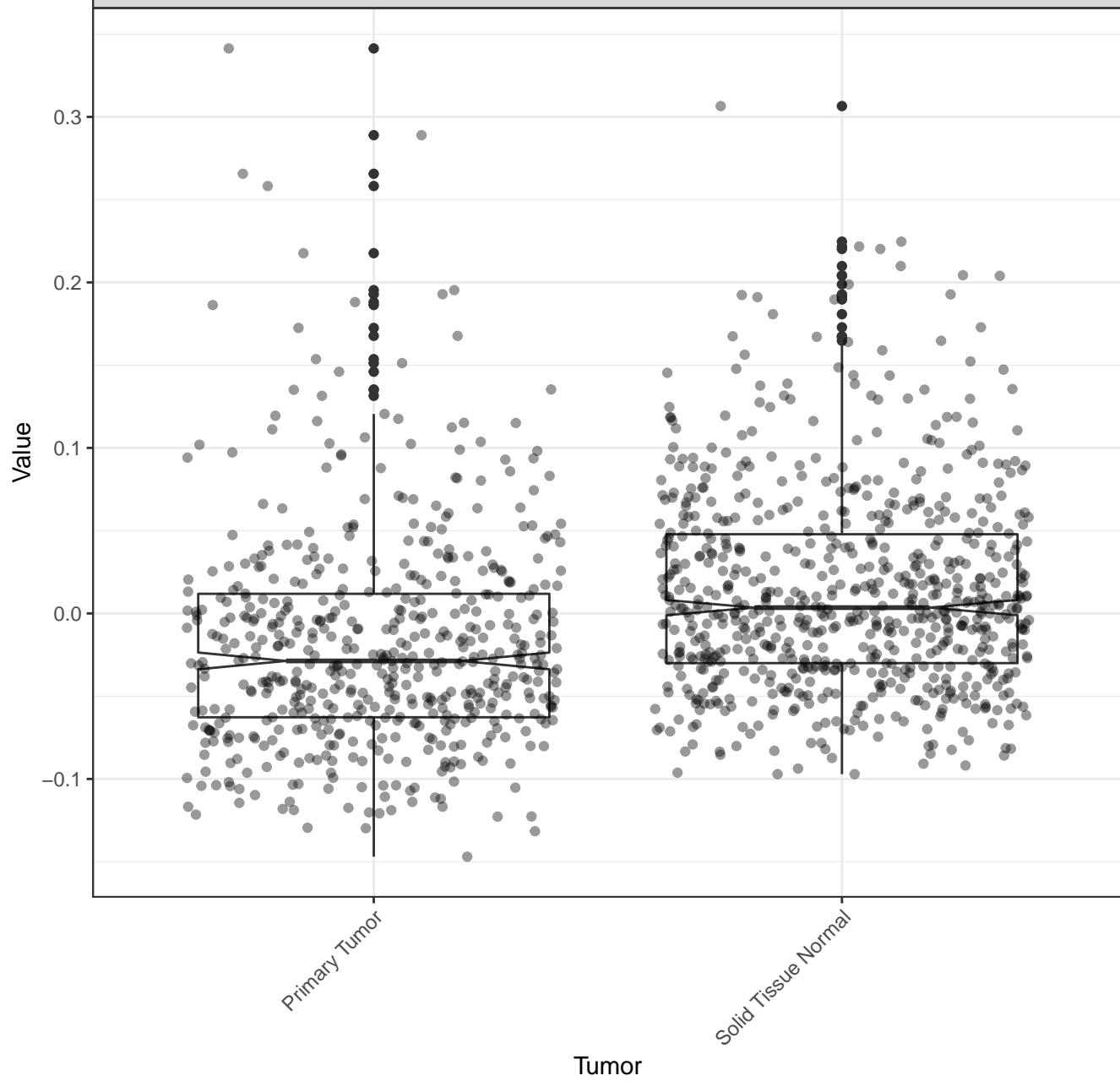
Primary Tumor

Solid Tissue Normal

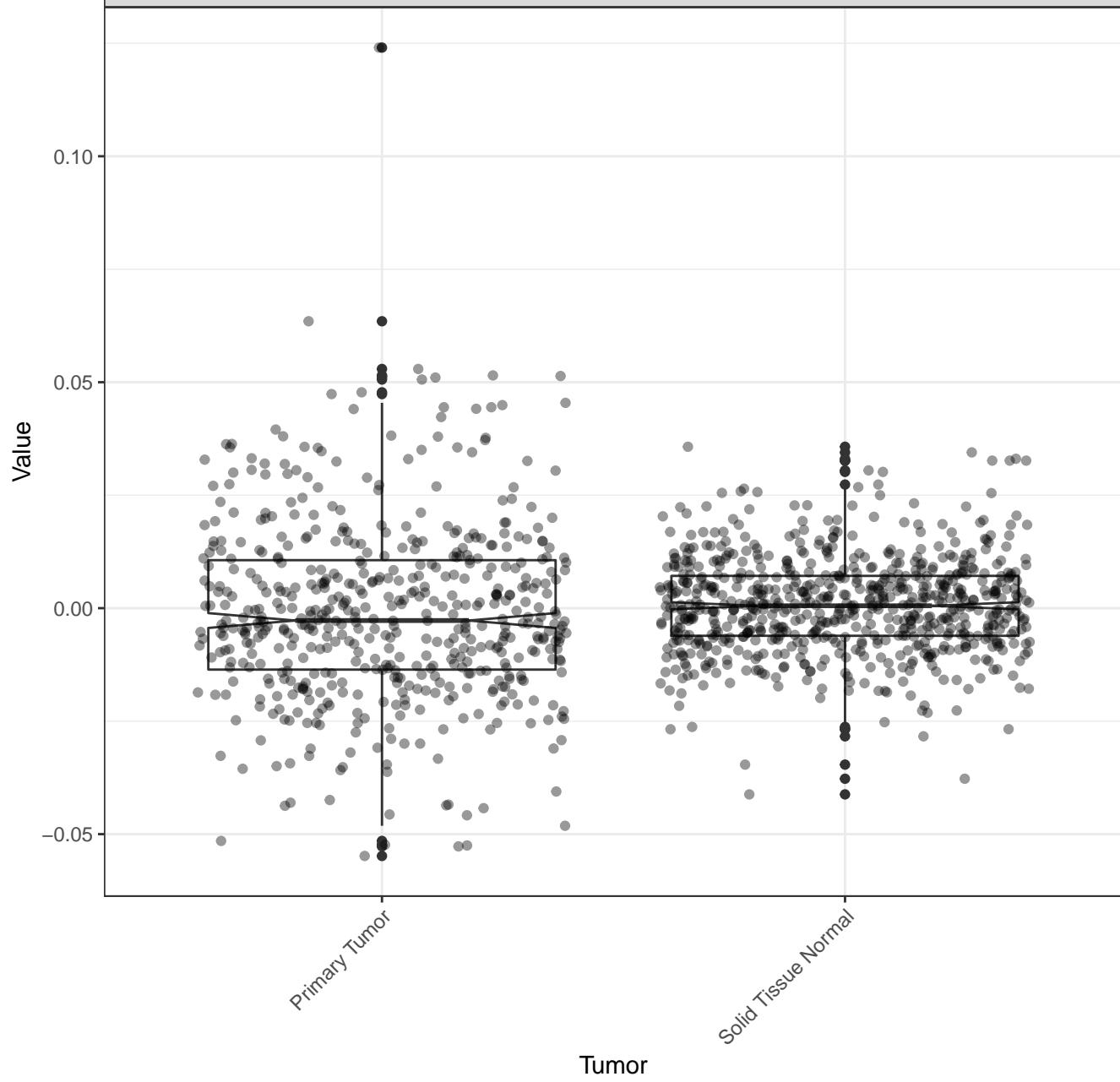
Tumor



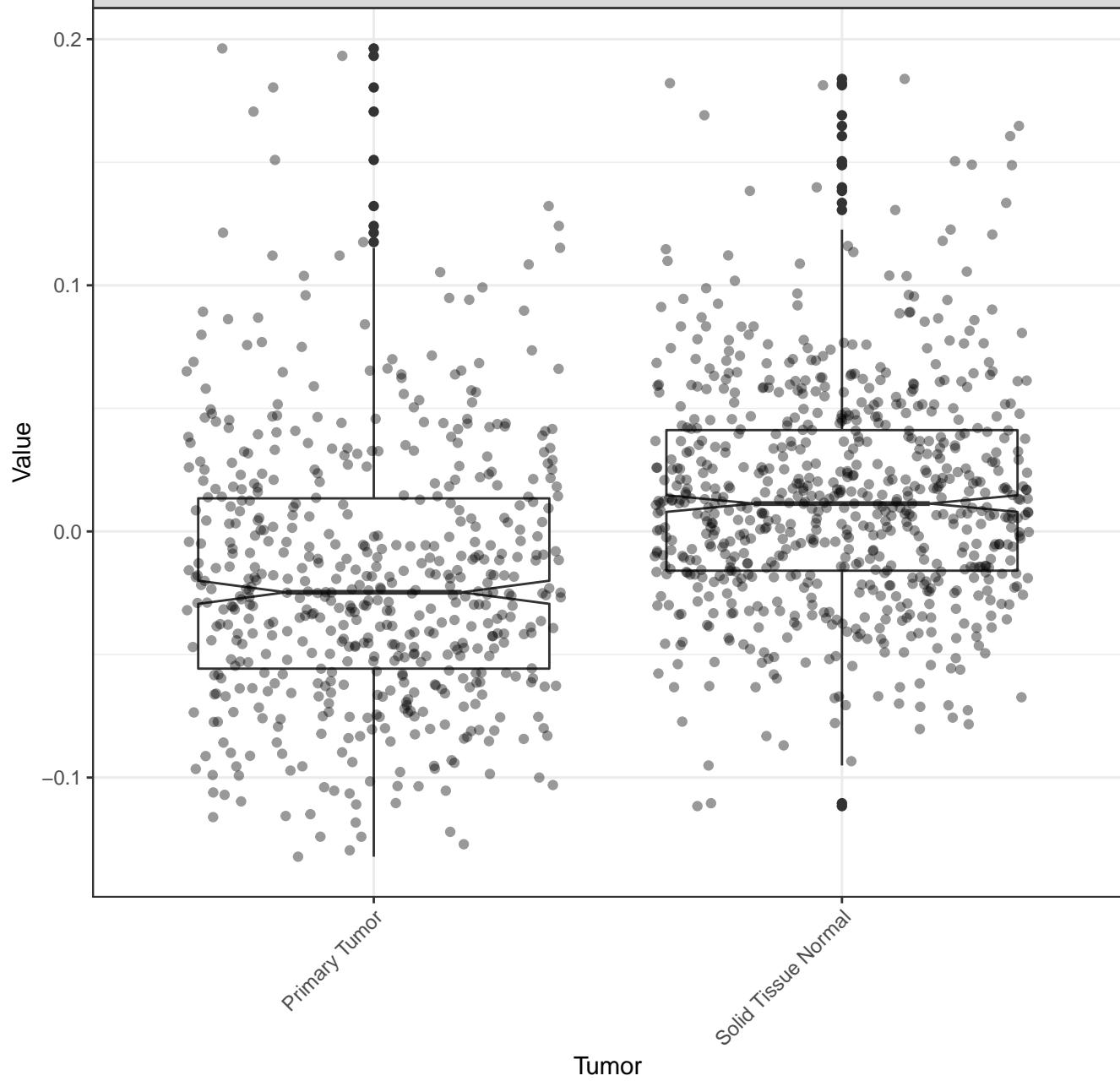
370,REACTOME_GENERIC_TRANSSCRIPTION_PATHWAY



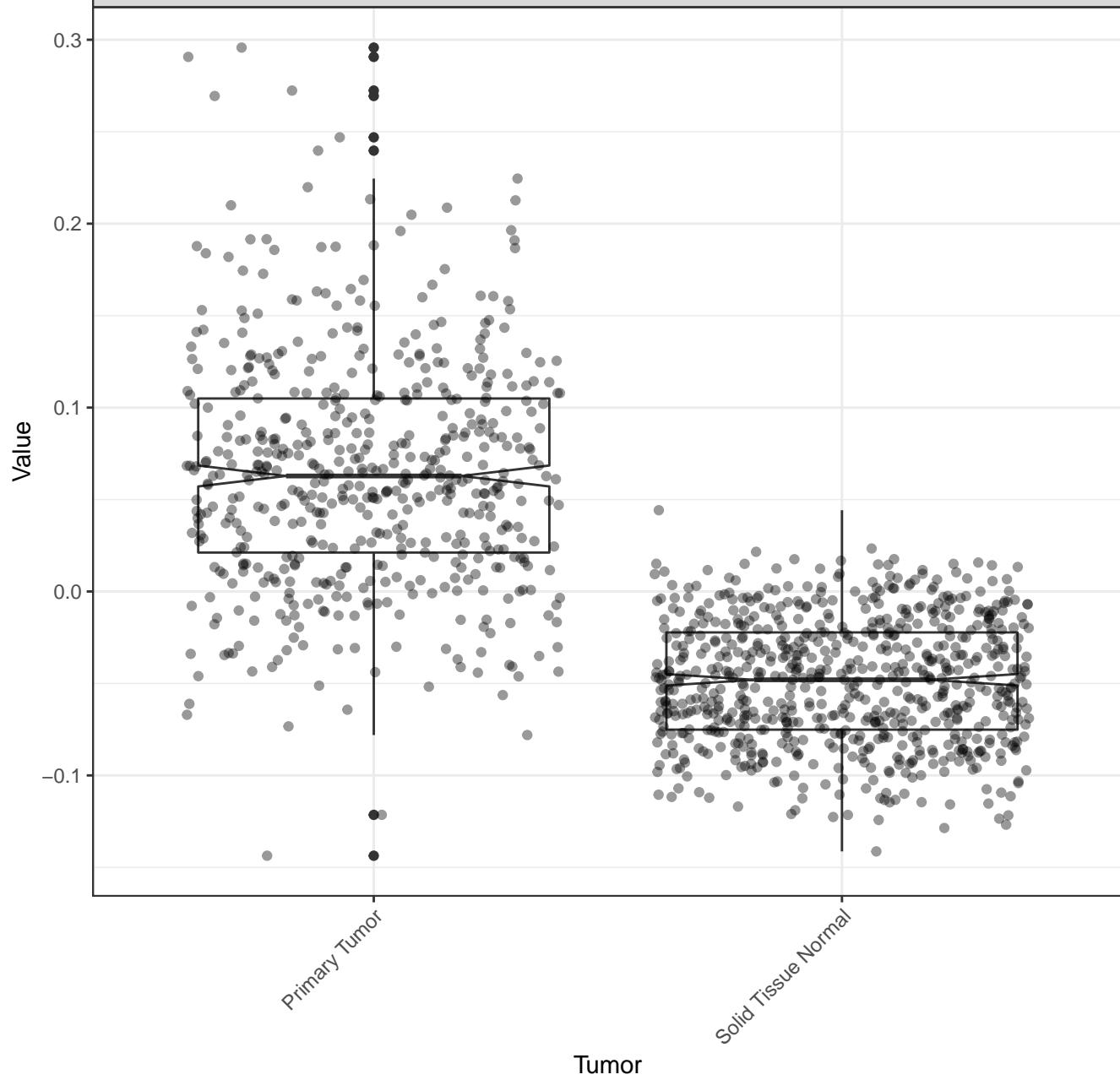
371,PID_ERBB1_DOWNSTREAM_PATHWAY



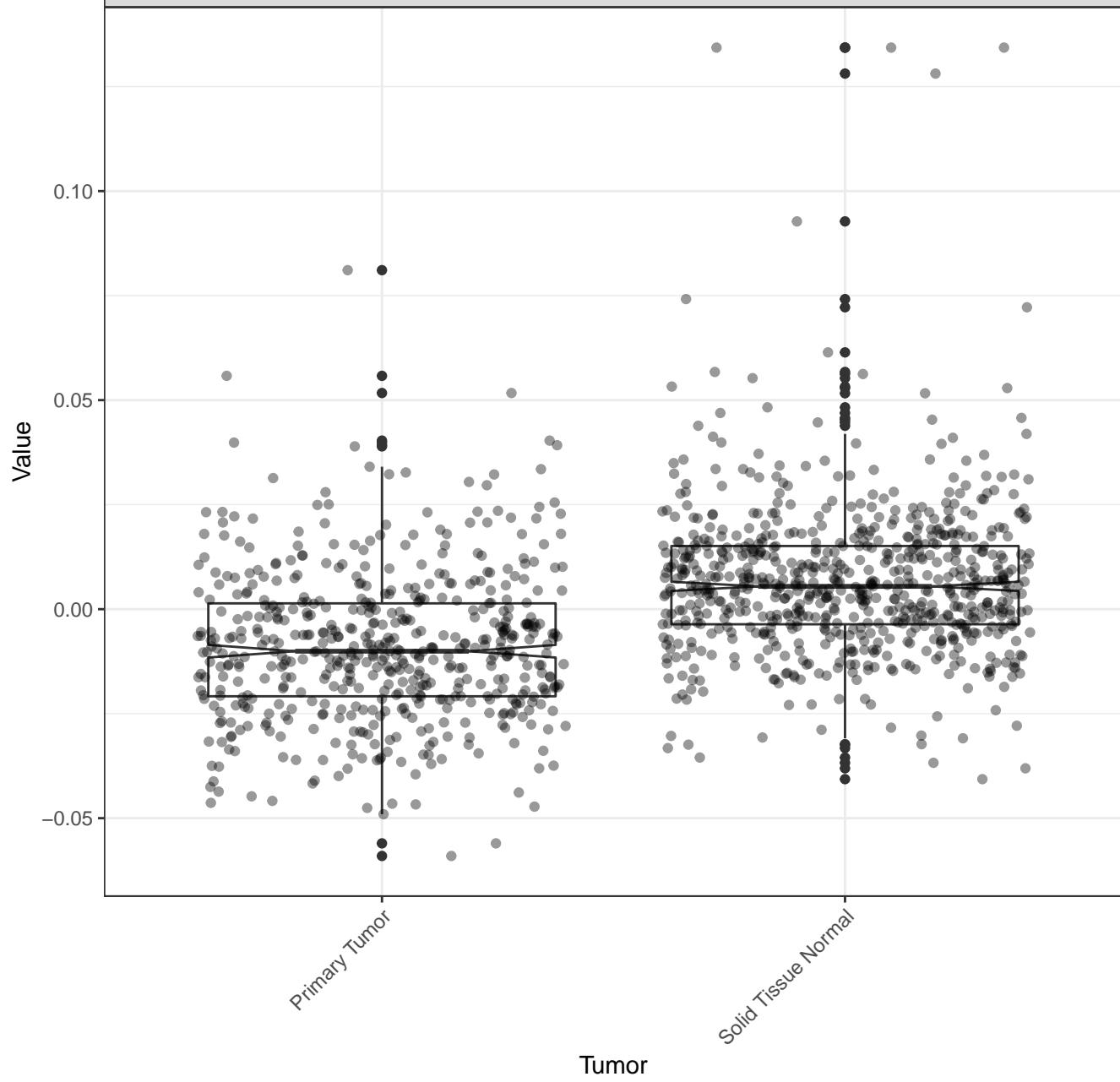
372,REACTOME_METABOLISM_OF_RNA



373,REACTOME_METABOLISM_OF_LIPIDS_AND_LIPOPROTEINS



374,DMAP_TCELLA2



375,KEGG_NUCLEOTIDE_EXCISION_REPAIR

Value

0.10

0.05

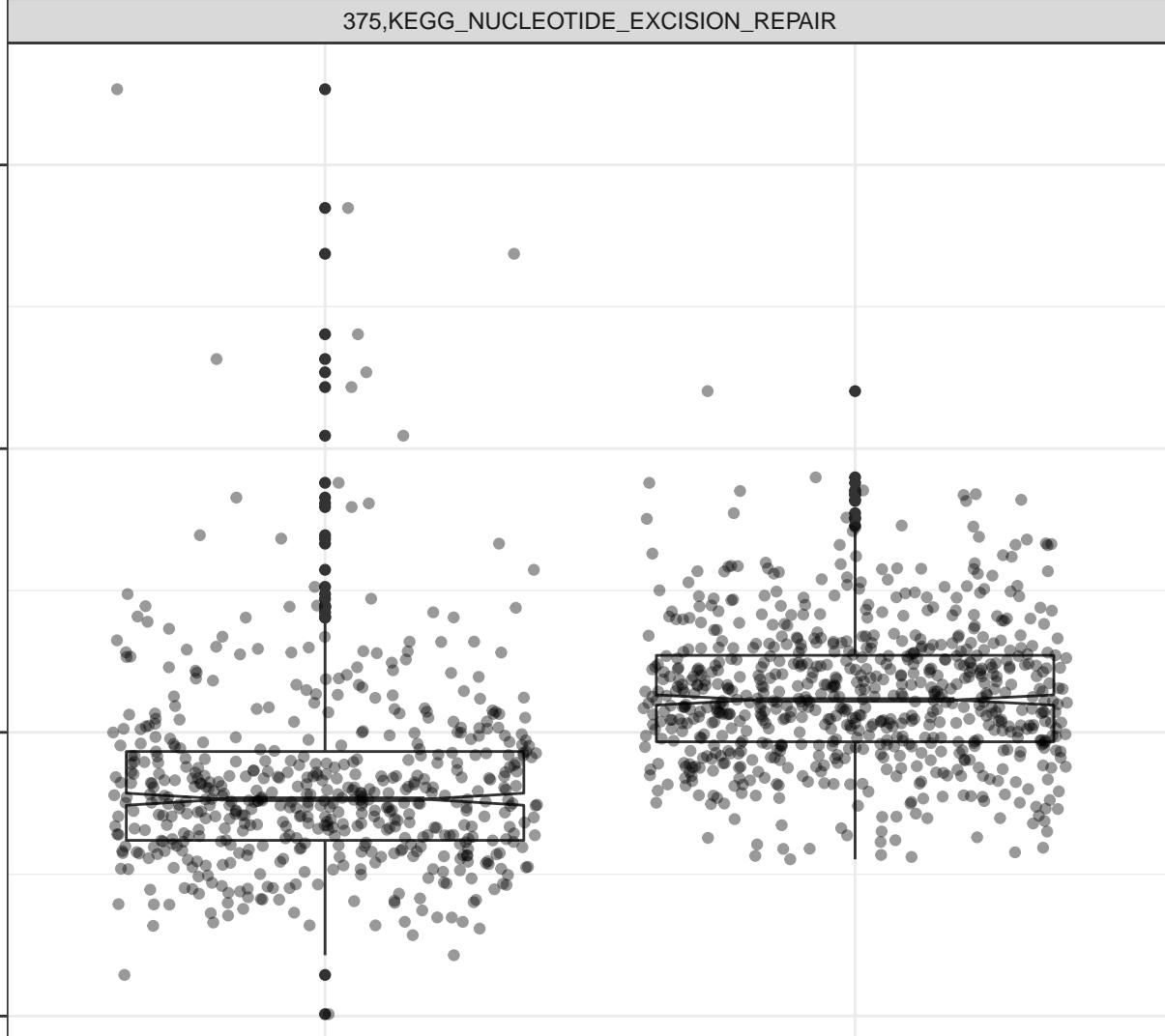
0.00

-0.05

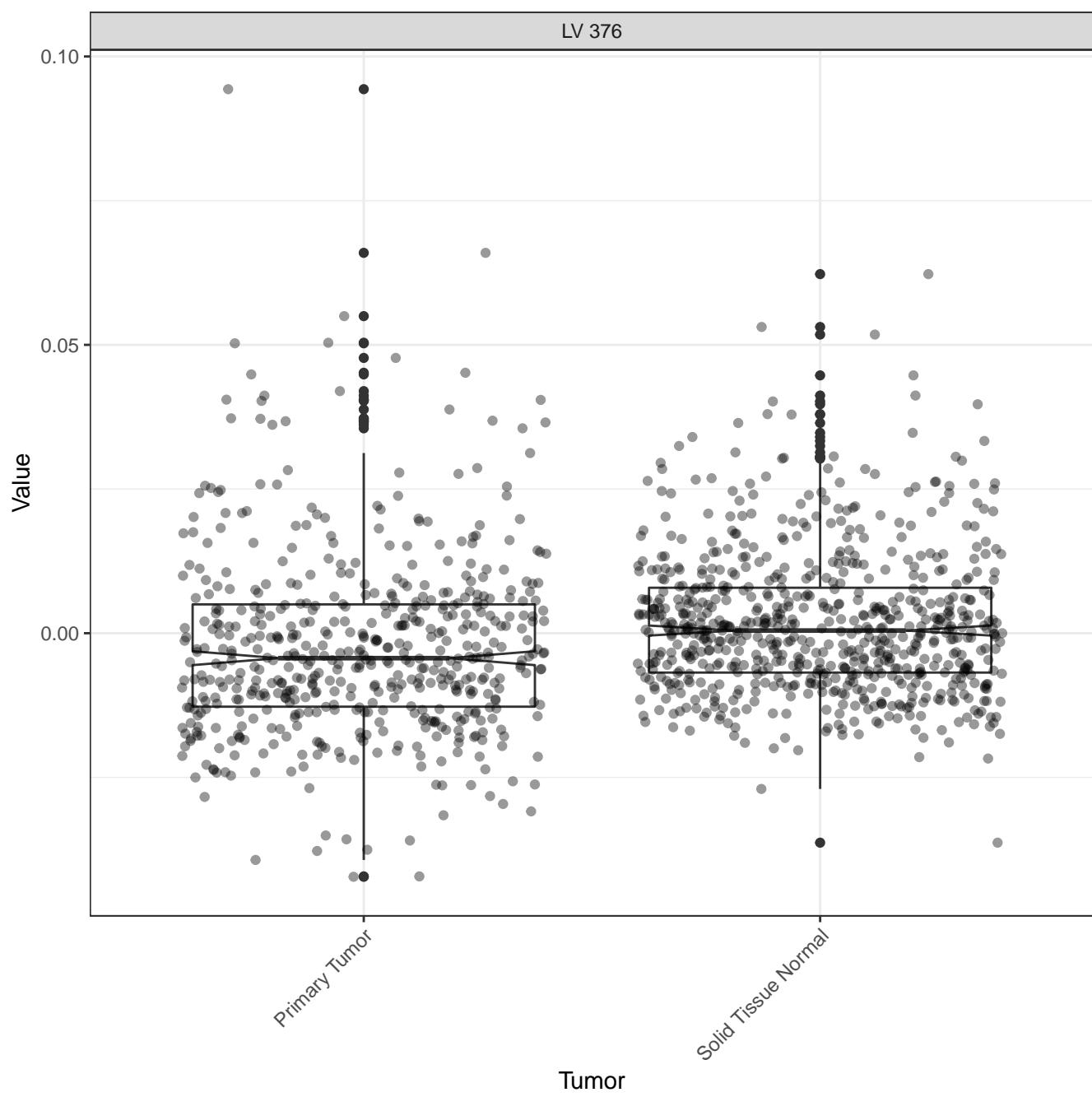
Primary Tumor

Solid Tissue Normal

Tumor



LV 376



377,DMAP_ERY3

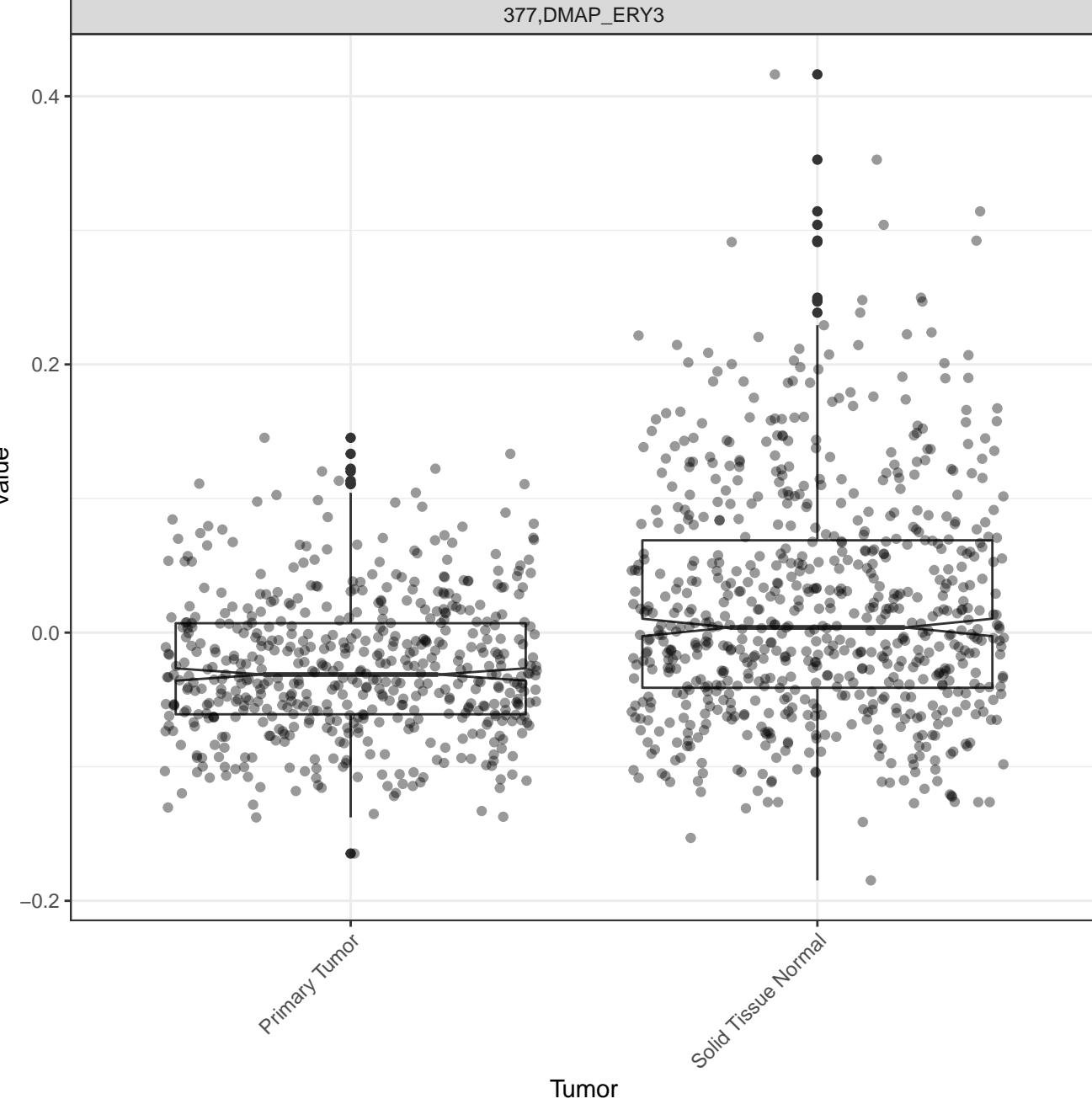
Value

0.4
0.2
0.0
-0.2

Primary Tumor

Tumor

Solid Tissue Normal



378,REACTOME_HOST_INTERACTIONS_OF_HIV_FACTORS

Value

0.05

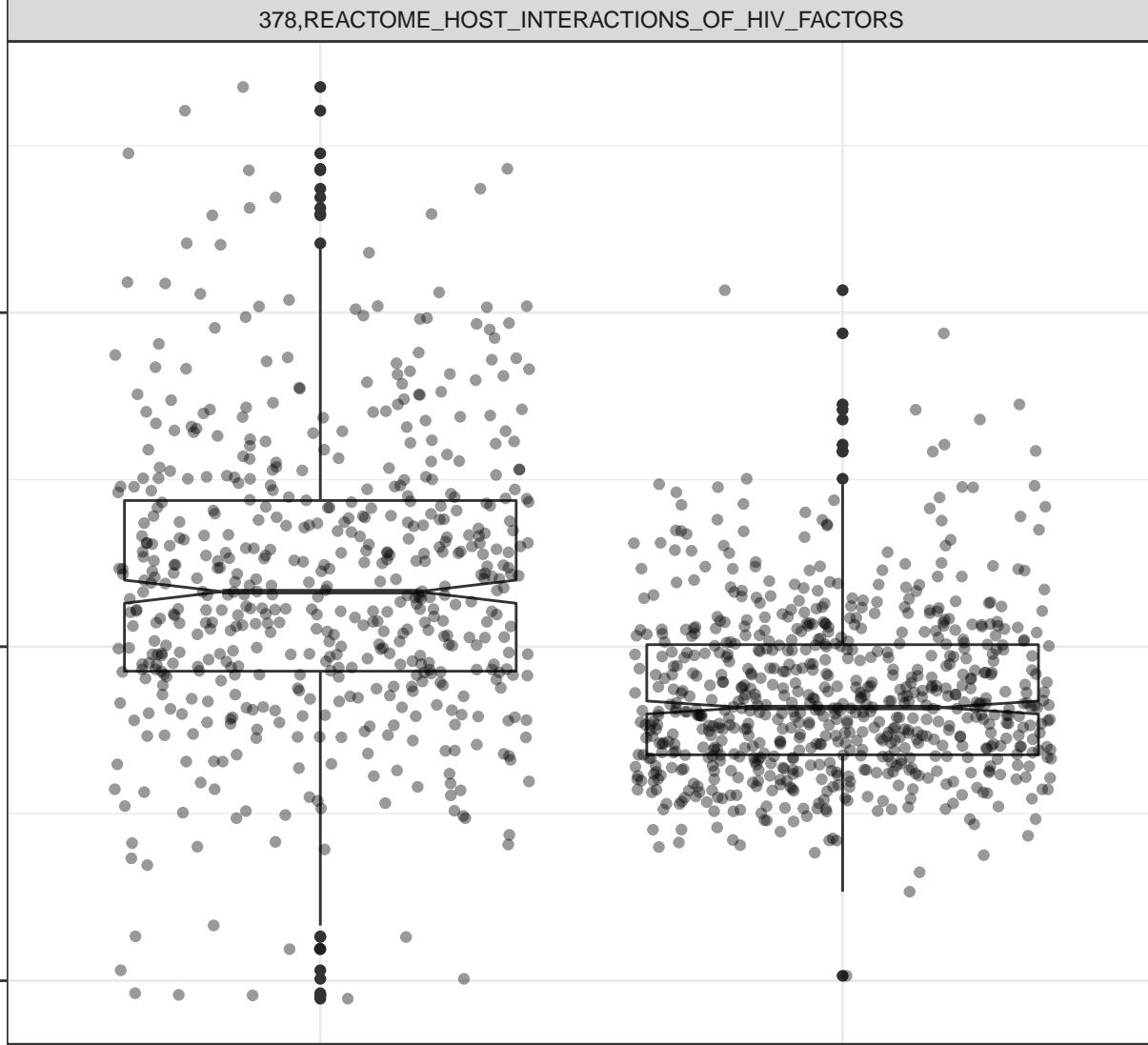
0.00

-0.05

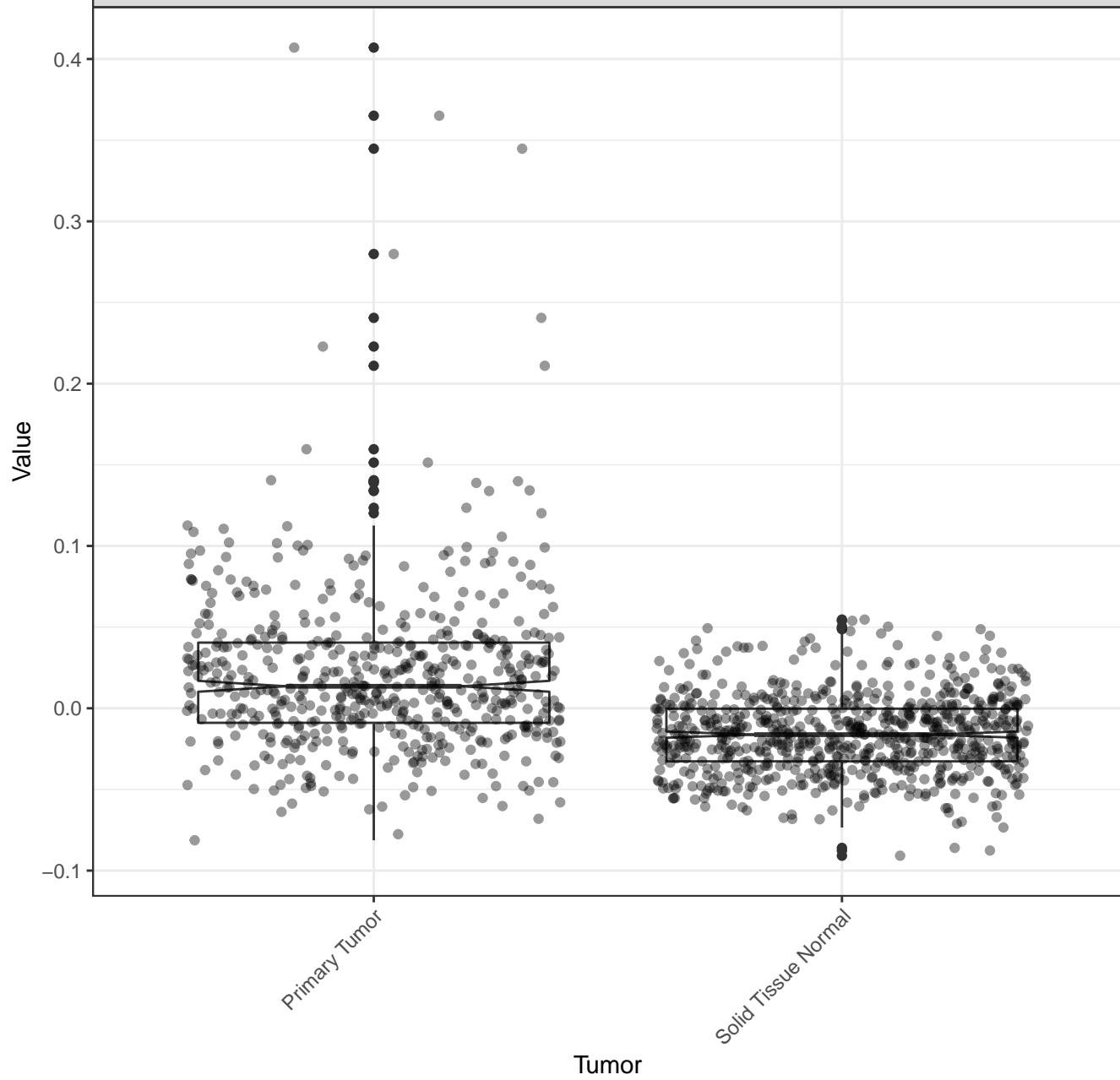
Primary Tumor

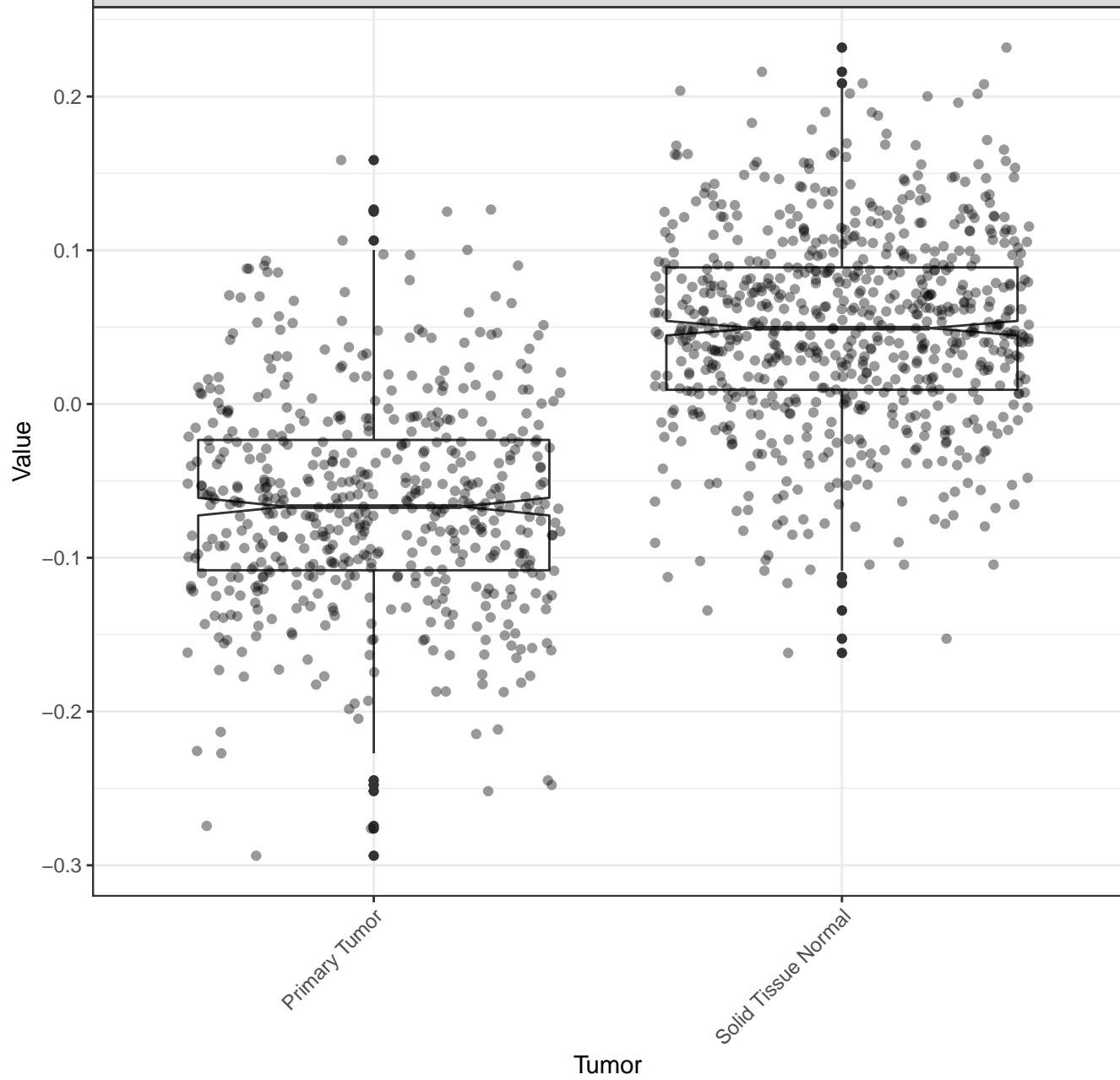
Solid Tissue Normal

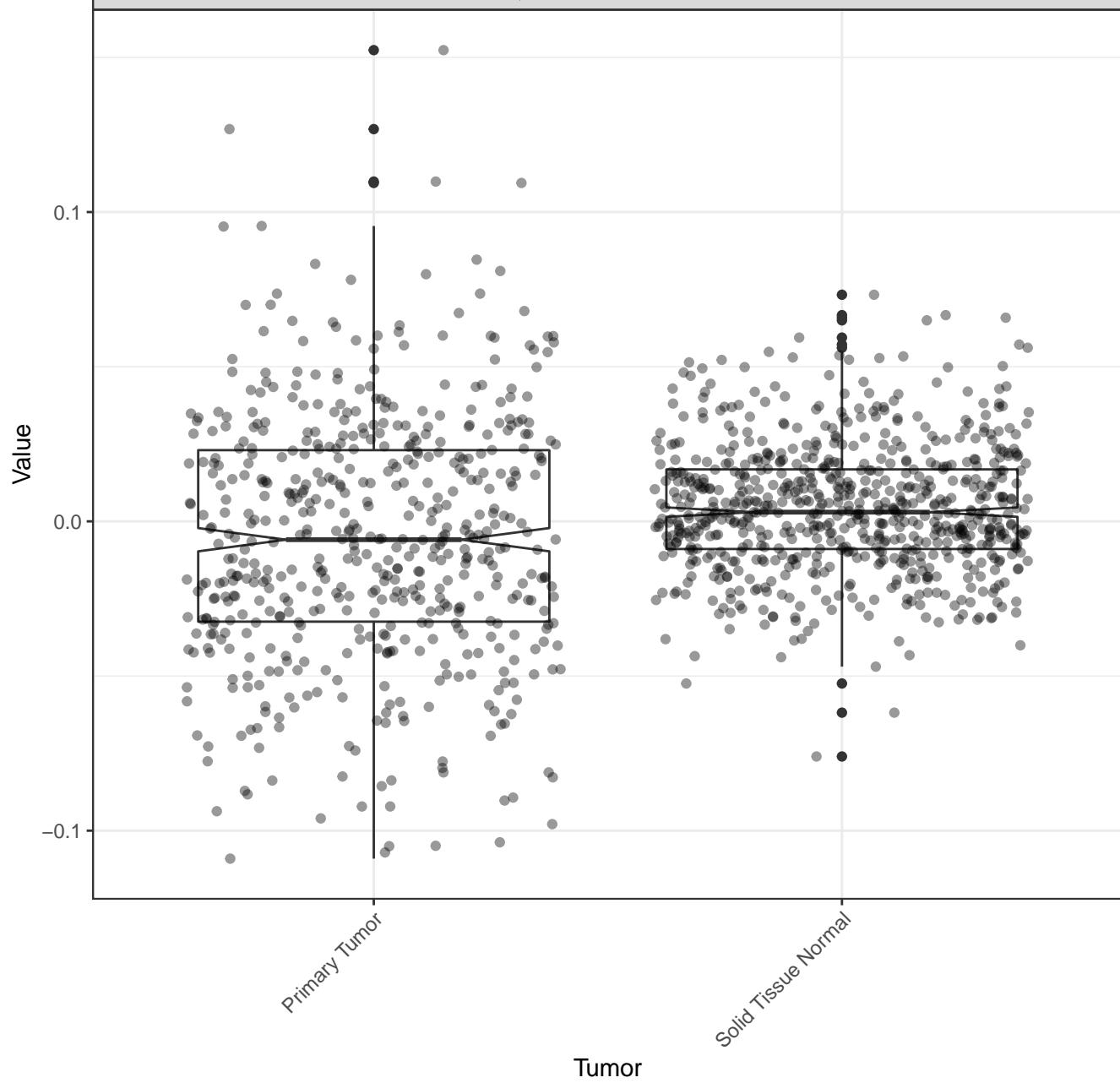
Tumor



379,KEGG_LYSOSOME







382,MIPS_NOP56P_ASSOCIATED_PRE_RRNA_COMPLEX

Value

0.75

0.50

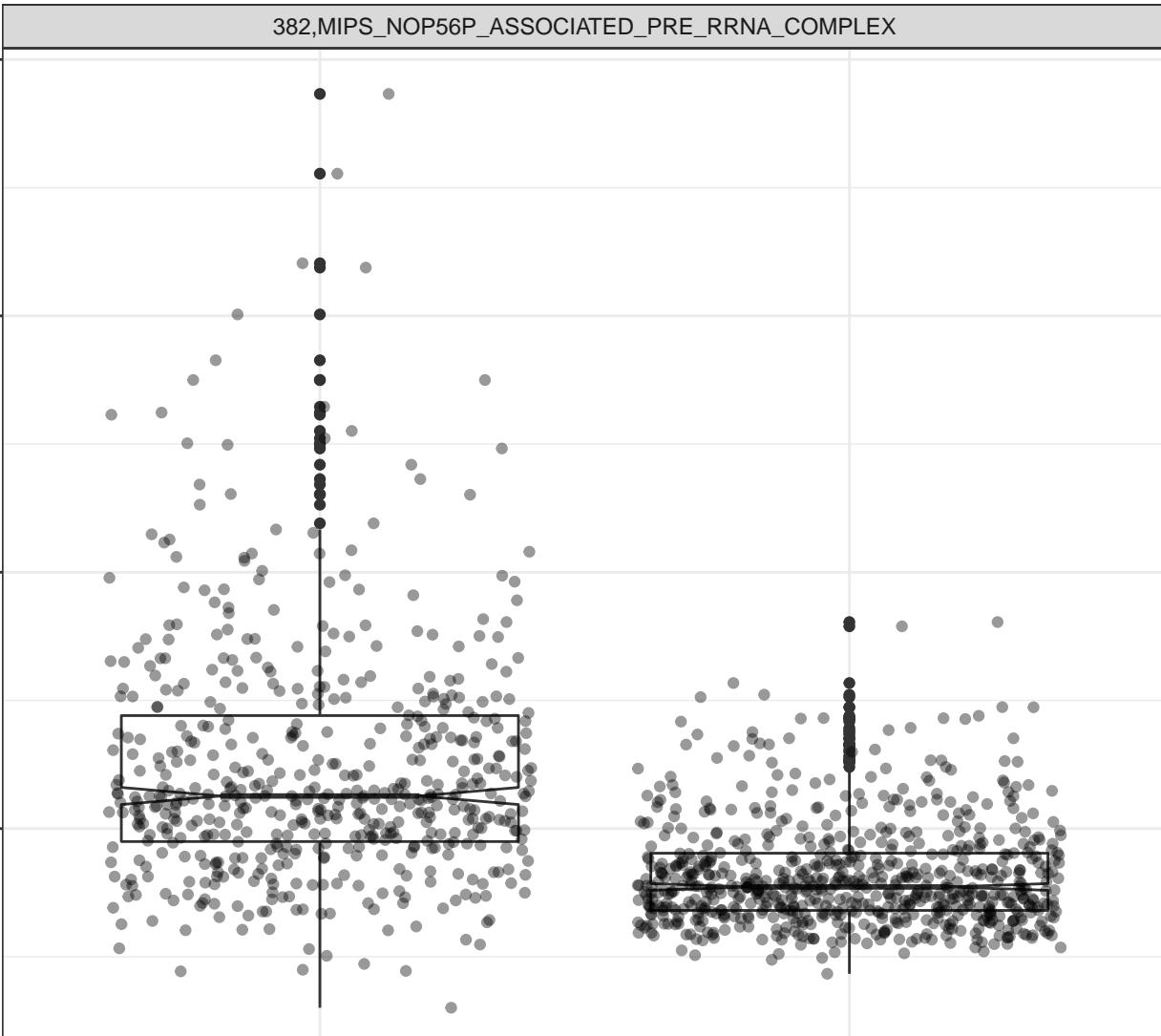
0.25

0.00

Primary Tumor

Solid Tissue Normal

Tumor



Value

0.15

0.10

0.05

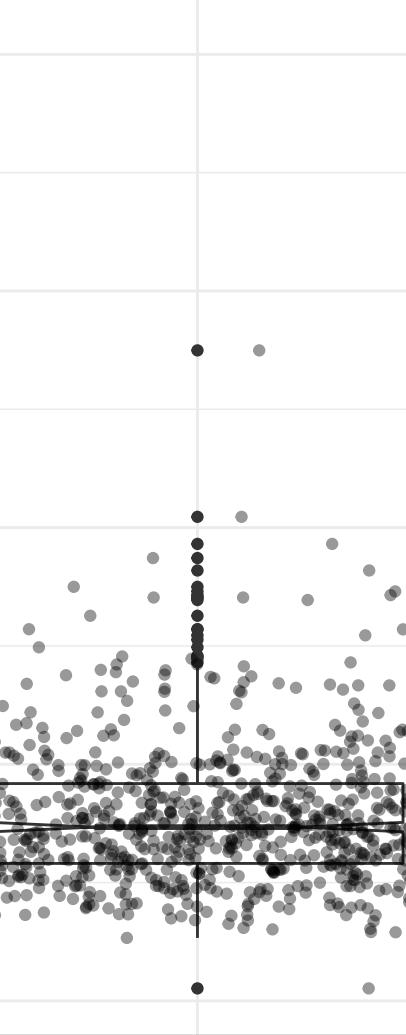
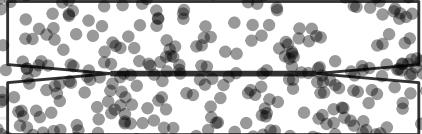
0.00

-0.05

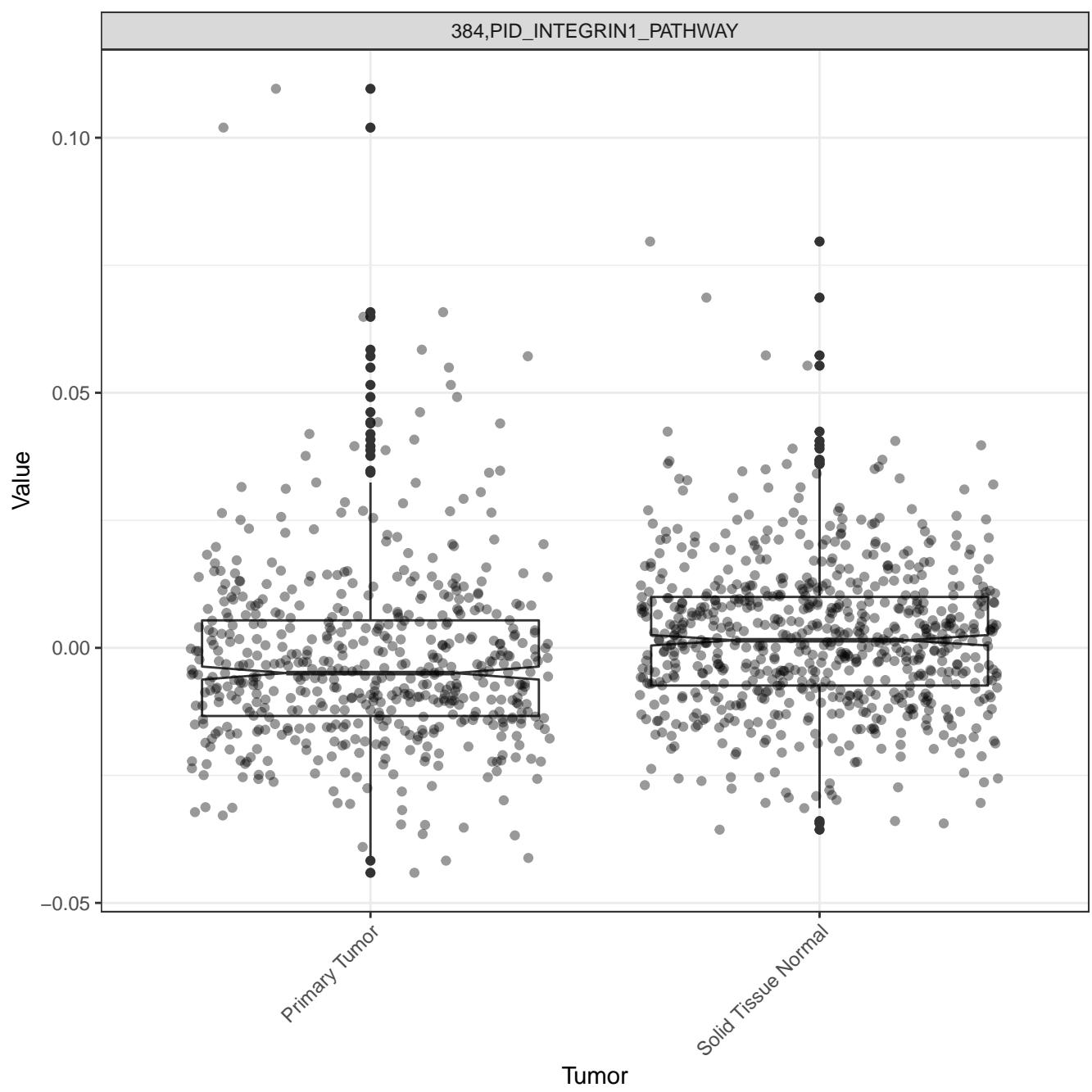
Primary Tumor

Solid Tissue Normal

Tumor



384,PID_INTEGRIN1_PATHWAY



LV 385

