Adipose	
FUNCTION	ADJ_P_VALUE
vasculature development	2×10^{-15}
blood vessel development	8×10^{-15}
blood vessel morphogenesis	2×10^{-14}
angiogenesis	5×10^{-13}
cell chemotaxis	2×10^{-8}
regulation of angiogenesis	1×10^{-7}
regulation of vasculature development	2×10^{-7}
positive regulation of cell motility	2×10^{-7}
positive regulation of cell migration	2×10^{-7}
leukocyte chemotaxis	2×10^{-7}

Liver

FUNCTION	ADJ_P_VALUE
monocarboxylic acid metabolic process	7×10^{-51}
small molecule catabolic process	7×10^{-39}
organic acid catabolic process	4×10^{-36}
carboxylic acid catabolic process	2×10^{-35}
steroid metabolic process	7×10^{-33}
small molecule biosynthetic process	4×10^{-31}
cellular amino acid metabolic process	1 × 10 ⁻²⁸
fatty acid metabolic process	6×10^{-28}
organic hydroxy compound metabolic process	9 × 10 ⁻²⁸
alpha-amino acid metabolic process	3 × 10 ⁻²⁷

Cervix

FUNCTION	ADJ_P_VALUE
axoneme assembly	1 × 10 ⁻⁴
cilium assembly	1×10^{-4}
cilium organization	3×10^{-4}
plasma membrane bounded cell projection assembly	4×10^{-4}
animal organ formation	4×10^{-4}
microtubule bundle formation	5×10^{-4}
cell projection assembly	5 × 10 ⁻⁴
cilium movement	6 × 10 ⁻⁴
morphogenesis of a branching structure	1 × 10 ⁻³
skeletal system development	1 × 10 ⁻³

Adrenal Gland

FUNCTION	ADJ_P_VALUE
sulfur compound metabolic process	3×10^{-3}
lipid biosynthetic process	3×10^{-3}
mitochondrial respiratory chain complex assembly	3×10^{-3}
steroid metabolic process	2×10^{-2}
secondary alcohol metabolic process	2×10^{-2}
sterol metabolic process	2×10^{-2}
regulation of lipid biosynthetic process	2×10^{-2}
cholesterol metabolic process	3×10^{-2}
steroid biosynthetic process	3×10^{-2}
mitochondrial transmembrane transport	4 × 10 ⁻²

Lung

FUNCTION	ADJ_P_VALUE
immune effector process	1 × 10 ⁻³⁸
regulation of cell activation	1×10^{-38}
regulation of leukocyte activation	1×10^{-35}
positive regulation of immune response	3×10^{-35}
adaptive immune response	1×10^{-34}
leukocyte mediated immunity	3×10^{-34}
lymphocyte activation	5×10^{-34}
cytokine production	4×10^{-33}
regulation of cytokine production	1×10^{-32}
T cell activation	1 × 10 ⁻³²

Skin

FUNCTION	ADJ_P_VALUE
epidermis development	2 × 10 ⁻²¹
skin development	9×10^{-21}
keratinization	6×10^{-17}
keratinocyte differentiation	2×10^{-14}
epidermal cell differentiation	1×10^{-10}
intermediate filament cytoskeleton organization	9×10^{-10}
intermediate filament-based process	1×10^{-9}
intermediate filament organization	6 × 10 ⁻⁹
regulation of water loss via skin	2 × 10 ⁻⁸
establishment of skin barrier	1×10^{-7}

Minor Salivary Gland

FUNCTION	ADJ_P_VALUE
epidermis development	2 × 10 ⁻⁵
skin development	9×10^{-5}
muscle cell development	1 × 10 ⁻⁴
multicellular organismal movement	2×10^{-4}
musculoskeletal movement	2×10^{-4}
myofibril assembly	2×10^{-4}
striated muscle cell development	2×10^{-4}
skeletal muscle organ development	2×10^{-4}
sarcomere organization	3×10^{-4}
skeletal muscle tissue development	3 × 10 ⁻⁴

Fallopian Tube

FUNCTION	ADJ_P_VALUE
axoneme assembly	2 × 10 ⁻⁷
cilium movement	2×10^{-6}
microtubule bundle formation	2×10^{-6}
cilium assembly	2×10^{-5}
cilium organization	2×10^{-5}
cilium or flagellum-dependent cell motility	3×10^{-4}
cilium-dependent cell motility	3×10^{-4}
cilium movement involved in cell motility	3×10^{-4}
axonemal dynein complex assembly	3×10^{-4}
microtubule-based movement	5×10^{-4}

Pituitary

FUNCTION	ADJ_P_VALUE
microtubule-based movement	7×10^{-14}
synaptic signaling	1×10^{-13}
chemical synaptic transmission	2×10^{-13}
anterograde trans-synaptic signaling	2×10^{-13}
trans-synaptic signaling	3×10^{-13}
cilium organization	2×10^{-11}
cilium assembly	3×10^{-11}
axoneme assembly	3×10^{-9}
cilium movement	3×10^{-9}
microtubule-based transport	4 × 10 ⁻⁹

Whole Blood

FUNCTION	ADJ_P_VALUE
lymphocyte activation	2×10^{-58}
immune response-regulating signaling pathway	4×10^{-49}
adaptive immune response	8×10^{-49}
immune effector process	4×10^{-47}
innate immune response	1×10^{-45}
T cell activation	4×10^{-45}
leukocyte mediated immunity	1×10^{-42}
positive regulation of immune response	7×10^{-42}
regulation of cell activation	1×10^{-41}
cytokine production	4 × 10 ⁻⁴⁰

Muscle - Skeletal

FUNCTION	ADJ_P_VALUE
muscle structure development	7×10^{-29}
muscle system process	8×10^{-27}
muscle organ development	5×10^{-25}
muscle cell development	1×10^{-21}
muscle contraction	1×10^{-21}
muscle cell differentiation	4×10^{-20}
striated muscle cell differentiation	4×10^{-19}
myofibril assembly	3×10^{-18}
striated muscle cell development	3×10^{-18}
striated muscle contraction	1×10^{-16}

Pancreas

FUNCTION	ADJ_P_VALUE
signal release	1×10^{-3}
protein secretion	1×10^{-3}
establishment of protein localization to extracellular region	1×10^{-3}
protein localization to extracellular region	2×10^{-3}
peptide secretion	2×10^{-3}
digestion	2×10^{-3}
hormone secretion	2×10^{-3}
regulation of secretion	2×10^{-3}
hormone transport	2×10^{-3}
peptide hormone secretion	2×10^{-3}

Ovary

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FUNCTION	ADJ_P_VALUE
genitalia development	3 × 10 ⁻²
extracellular structure organization	4×10^{-2}
male sex differentiation	4×10^{-2}
collagen fibril organization	4×10^{-2}
male genitalia development	4×10^{-2}
extracellular matrix organization	4×10^{-2}
regulation of smoothened signaling pathway	4×10^{-2}
external encapsulating structure organization	4×10^{-2}

Thyroid

FUNCTION	ADJ_P_VALUE
thyroid hormone generation	1 × 10 ⁻⁴
plasma membrane bounded cell projection assembly	2×10^{-4}
cell projection assembly	2×10^{-4}
thyroid hormone metabolic process	2×10^{-4}
cilium assembly	5×10^{-4}
cilium organization	3×10^{-3}
positive regulation of cell-substrate adhesion	3×10^{-2}
cilium movement	3×10^{-2}
regulation of cell-substrate adhesion	4×10^{-2}

Brain

2.5	
FUNCTION	ADJ_P_VALUE
synaptic signaling	9×10^{-43}
trans-synaptic signaling	9×10^{-43}
chemical synaptic transmission	9×10^{-43}
anterograde trans-synaptic signaling	9 × 10 ⁻⁴³
modulation of chemical synaptic transmission	9×10^{-25}
regulation of trans-synaptic signaling	9×10^{-25}
neuron projection morphogenesis	2×10^{-18}
cell projection morphogenesis	8×10^{-18}
plasma membrane bounded cell projection morphogenesis	2×10^{-17}
cell part morphogenesis	2 × 10 ⁻¹⁷

Bladder

FUNCTION	ADJ_P_VALUE
tissue morphogenesis	4×10^{-5}
muscle tissue development	5×10^{-5}
cardiac muscle tissue development	1×10^{-4}
morphogenesis of an epithelium	1×10^{-4}
striated muscle tissue development	1×10^{-4}
mesenchyme development	1×10^{-4}
epithelial to mesenchymal transition	2×10^{-4}
muscle structure development	3×10^{-4}
mesenchymal cell differentiation	6×10^{-4}
urogenital system development	9×10^{-4}

Spleen

FUNCTION	ADJ_P_VALUE
lymphocyte activation	1 × 10 ⁻⁶⁵
T cell activation	3×10^{-56}
adaptive immune response	4×10^{-53}
immune effector process	5×10^{-50}
positive regulation of immune response	8×10^{-50}
regulation of cell activation	3×10^{-49}
regulation of leukocyte activation	4×10^{-49}
regulation of lymphocyte activation	8×10^{-48}
leukocyte cell-cell adhesion	2 × 10 ⁻⁴²
immune response-regulating signaling pathway	1 × 10 ⁻⁴¹

Prostate

FUNCTION	ADJ_P_VALUE
tissue morphogenesis	4 × 10 ⁻⁴
morphogenesis of a branching structure	5×10^{-4}
morphogenesis of an epithelium	7×10^{-4}
morphogenesis of a branching epithelium	1×10^{-3}
branching morphogenesis of an epithelial tube	2×10^{-3}
regulation of morphogenesis of a branching structure	3×10^{-2}
muscle structure development	3×10^{-2}
regulation of morphogenesis of an epithelium	3×10^{-2}
striated muscle tissue development	3×10^{-2}
steroid hormone mediated signaling pathway	3×10^{-2}

Artery

FUNCTION	ADJ_P_VALUE
muscle structure development	5 × 10 ⁻¹³
muscle system process	3×10^{-11}
muscle cell development	4×10^{-10}
extracellular matrix organization	1×10^{-9}
muscle cell differentiation	1×10^{-9}
extracellular structure organization	1×10^{-9}
external encapsulating structure organization	1×10^{-9}
actin cytoskeleton organization	3×10^{-9}
muscle contraction	8×10^{-9}
muscle tissue development	3×10^{-8}

Nerve - Tibial

FUNCTION	ADJ_P_VALUE
cell morphogenesis involved in differentiation	1 × 10 ⁻¹²
ensheathment of neurons	1×10^{-12}
axon ensheathment	1×10^{-12}
cell part morphogenesis	3×10^{-12}
myelination	3×10^{-12}
cell morphogenesis involved in neuron differentiation	3×10^{-12}
cell projection morphogenesis	3×10^{-12}
axonogenesis	4×10^{-12}
neuron projection morphogenesis	4×10^{-12}
plasma membrane bounded cell projection morphogenesis	7×10^{-12}

Esophagus

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FUNCTION	ADJ_P_VALUE
keratinization	3 × 10 ⁻⁶
epidermis development	3×10^{-6}
keratinocyte differentiation	4×10^{-6}
epithelial cell differentiation	2×10^{-5}
mitotic cell cycle process	2×10^{-5}
epidermal cell differentiation	2×10^{-5}
regulation of cell cycle process	5×10^{-5}
cell division	6×10^{-5}
intermediate filament-based process	6×10^{-5}
sister chromatid segregation	1 × 10 ⁻⁴

Stomach

FUNCTION	ADJ_P_VALUE
digestion	1 × 10 ⁻⁴
regulation of hormone levels	6×10^{-4}
gastric acid secretion	2×10^{-3}
peptide hormone secretion	2×10^{-3}
peptide transport	2×10^{-3}
peptide secretion	3×10^{-3}
digestive system process	6×10^{-3}
amide transport	6×10^{-3}
hormone transport	6×10^{-3}
hormone secretion	6×10^{-3}

Testis

FUNCTION	ADJ_P_VALUE
male gamete generation	1 × 10 ^{-3!}
spermatogenesis	3 × 10 ⁻³⁴
microtubule-based movement	2 × 10 ⁻²⁰
cilium organization	2 × 10 ⁻²⁰
cilium assembly	2 × 10 ⁻²³
microtubule cytoskeleton organization	2 × 10 ⁻²³
meiotic cell cycle	2 × 10 ⁻²³
cilium movement	3×10^{-20}
nuclear division	1 × 10 ⁻¹⁸
meiotic cell cycle process	2 × 10 ⁻¹⁸

Heart

ADJ_P_VALU
3 × 10 ⁻
1 × 10-
2 × 10 ⁻
nt 2 × 10 ⁻
3 × 10 ⁻
nent 3 × 10 ⁻
3 × 10 ⁻
nent 2 × 10 ⁻
5 × 10 ⁻
1 × 10 ⁻