

*Supplemental Materials*

***In Silico* Analysis Reveals Overlapping Molecular Mechanisms  
Between COVID-19 and Attention-Deficit/Hyperactivity Disorder**

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**Supplementary Tables 1-4**

**Supplementary Table 1. List of genes for each enrichment term in GO Molecular Function pathways.**

<b>Enrichment terms</b>	<b>List of genes</b>
Growth factor activity	<i>AMH, HGF, IL2, IL5, REG1A, PSPN, FGF2, EGF, CSF2, AGT, NGF, IGF1, IL6, IL10, BDNF</i>
Hormone activity	<i>SCT, AMH, GAST, PMCH, AGT, AVP, IGF1, INS, POMC, ADIPOQ, PTH, ADM, HAMP</i>
Growth factor receptor binding	<i>IL2, IL5, PSPN, IL1RN, IL1A, IL1B, FGF2, EGF, APP, CSF2, IL6, IL10</i>
1-phosphatidylinositol-3-kinase activity	<i>PIK3CA, ATM, PIK3CD, PIK3CG, PIK3CB</i>
Monooxygenase activity	<i>CYP4F3, CYP27B1, CYP2B6, DBH, TBXAS1, PAH, NOS1, CYP2D6, CYP3A4</i>
1-phosphatidylinositol-4-phosphate 3-kinase activity	<i>PIK3CA, PIK3CD, PIK3CG, PIK3CB</i>
Phosphatidylinositol-4,5-bisphosphate 3-kinase activity	<i>PIK3CA, PIK3CD, PIK3CG, PIK3CB</i>
Copper ion binding	<i>IL1A, SOD1, ALB, MT-CO2, DBH, HAMP, SNCA</i>
Phosphatidylinositol-3,4-bisphosphate 5-kinase activity	<i>PIK3CA, PIK3CD, PIK3CG, PIK3CB</i>
Anandamide 8,9 epoxidase activity	<i>CYP2B6, CYP2D6, CYP3A4</i>
Anandamide 11,12 epoxidase activity	<i>CYP2B6, CYP2D6, CYP3A4</i>
Antioxidant activity	<i>CAT, APOE, SOD1, ALB, TPO, HP, TXN</i>
Anandamide 14,15 epoxidase activity	<i>CYP2B6, CYP2D6, CYP3A4</i>
Oxidoreductase activity, acting on paired donors, with incorporation or reduction of molecular oxygen, NAD(P)H as one donor, and incorporation of one atom of oxygen	<i>CYP27B1, CYP2B6, NOS1, CYP2D6, CYP3A4</i>
Death receptor binding	<i>CASP3, NGF, EDA, BDNF</i>
Tau protein binding	<i>STUB1, APOE, S100B, GSK3B, SNCA</i>
Low-density lipoprotein particle receptor binding	<i>APOE, CRP, APP, HSPG2</i>
Insulin receptor substrate binding	<i>PIK3CA, INSR, PIK3CB</i>
Neurotrophin receptor binding	<i>NGF, BDNF, NTRK1</i>
Death receptor agonist activity	<i>NGF, EDA</i>

**Supplementary Table 2. List of genes for each enrichment term in GO Cellular Component pathways.**

<b>Enrichment terms</b>	<b>List of genes</b>
Secretory granule lumen	<i>MAPK1, HGF, NFKB1, CAT, CFP, EGF, APP, ALB, SERPINF2, FN1, HP, IGF1, DBH, INS, POMC, SERPINA1, PPIA</i>
Axon	<i>SNAP25, NRPI, SOD1, SLC6A3, APP, BCR, USP9X, GSK3B, SCN8A, COMT, CNTNAP2, DMD, INSR, CNR1, NGF, OPRM1, DDC, GABRG2, BDNF, NTRK1, ADORA2A, SLC6A1, PRKCB, SNCA</i>
Endoplasmic reticulum lumen	<i>MAPK1, STS, PROC, CFP, APOE, HRC, BCHE, MFGE8, APP, ALB, FN1, COL18A1, COL11A2, INS, IL6, SERPINA1</i>
Platelet alpha granule lumen	<i>HGF, EGF, APP, ALB, SERPINF2, FN1, IGF1, SERPINA1</i>
Membrane raft	<i>MAPK1, SLC6A3, APP, CASP3, F2R, ADCY2, DMD, CNR1, ADRA1A, OPRM1, PTPRC, SLC2A1, NOS1, HK1, FAS</i>
Phosphatidylinositol 3-kinase complex, class IA	<i>PIK3CA, PIK3CD, PIK3CG, PIK3CB</i>
Blood microparticle	<i>PON1, APOE, BCHE, ALB, SERPINF2, FN1, HP, AGT, SLC2A1</i>
Synaptic membrane	<i>CLTA, SNAP25, NRPI, SLC6A3, F2R, HTR3A, DMD, CNR1, SRPX2, ADRA1A, NLGN1, OPRM1, GABRG2, ADORA2A, SLC6A1</i>
Integral component of postsynaptic membrane	<i>NRPI, SLC6A3, ADRA1A, NLGN1, OPRM1, GABRG2, ADORA2A, SLC6A1</i>
Integral component of synaptic membrane	<i>NRPI, SLC6A3, CNR1, ADRA1A, NLGN1, OPRM1, GABRG2, ADORA2A, SLC6A1</i>
Organelle envelope lumen	<i>CAT, HTRA2, BCHE, SOD1, APP, PANK2, COA7</i>
Postsynaptic membrane	<i>NRPI, SLC6A3, F2R, HTR3A, DMD, ADRA1A, NLGN1, OPRM1, GABRG2, ADORA2A, SLC6A1</i>
Integral component of presynaptic membrane	<i>SLC6A3, CNR1, ADRA1A, OPRM1, ADORA2A, SLC6A1</i>
Presynaptic membrane	<i>CLTA, SNAP25, SLC6A3, CNR1, ADRA1A, OPRM1, ADORA2A, SLC6A1</i>
Exocytic vesicle	<i>CLTA, SNAP25, APP, DMD, NGF, IGF1, DDC, BDNF, RAB40B, SNCA</i>

Synaptic vesicle	<i>CLTA, SNAP25, APP, DMD, NGF, DDC, BDNF, RAB40B, SNCA</i>
Phosphatidylinositol 3-kinase complex, class IB	<i>PIK3CA, PIK3CG</i>
GABA-ergic synapse	<i>CNR1, ADRA1A, NLGN1, GABRG2, SLC6A1</i>
Caveola	<i>MAPK1, SLC6A3, F2R, ADRA1A, SLC2A1</i>
Cytoplasmic side of membrane	<i>EEF1A2, SNAP25, KRAS, HTRA2, GNAS, PTEN, PTPRC, AKT1</i>

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**Supplementary Table 3. List of genes for each enrichment term in GO Biological Processes pathways.**

<b>Enrichment terms</b>	<b>List of genes</b>
Dopamine catabolic process	<i>SLC6A3, MAOA, COMT, MAOB, DBH</i>
Regulation of calcidiol 1-monooxygenase activity	<i>NFKB1, CYP27B1, IFNG, IL1B</i>
Alkaloid metabolic process	<i>BCHE, DDC, CYP2D6, CYP3A4</i>
Maintenance of blood vessel diameter homeostasis by renin-angiotensin	<i>SERPINF2, AGT, AGTR2</i>
Brain renin-angiotensin system	<i>TACR1, AGT, AGTR2</i>
G protein-coupled receptor signaling pathway coupled to cGMP nucleotide second messenger	<i>AGT, AGTR2</i>
Superior temporal gyrus development	<i>GSK3B, CNTNAP2</i>
Regulation of Golgi lumen acidification	<i>SLC9A7, UBE3A</i>
Fever generation	<i>IL1A, IL1B</i>
Angiotensin-mediated drinking behavior	<i>TACR1, AGT</i>
Alkaloid catabolic process	<i>CYP2D6, CYP3A4</i>
Negative regulation of calcidiol 1-monooxygenase activity	<i>NFKB1, CYP27B1</i>
Isoquinoline alkaloid metabolic process	<i>DDC, CYP2D6</i>
Natural killer cell chemotaxis	<i>PIK3CD, PIK3CG</i>
Negative regulation of serotonin uptake	<i>NOS1, SNCA</i>
Positive regulation of calcidiol 1-monooxygenase activity	<i>IFNG, IL1B</i>
Trans-synaptic signaling by endocannabinoid, modulating synaptic transmission	<i>F2R, CNR1</i>
Positive regulation of plasma cell differentiation	<i>IL2, IL10</i>
Response to melanocyte-stimulating hormone	<i>MC4R, POMC</i>
Regulation of hydrogen peroxide catabolic process	<i>HP, SNCA</i>

**Supplementary Table 4. List of genes for each enrichment term in KEGG pathway.**

<b>Enrichment terms</b>	<b>List of genes</b>
EGFR tyrosine kinase inhibitor resistance	<i>MAPK1, HGF, KRAS, PIK3CA, FGF2, EGF, ERBB2, GSK3B, EIF4EBP1, PTEN, PIK3CD, IGF1, IL6, AKT1, PRKCB, PIK3CB</i>
Prostate cancer	<i>MAPK1, NFKB1, KRAS, PIK3CA, EGF, ERBB2, KLK3, GSK3B, INSRR, PTEN, PIK3CD, IGF1, INS, AKT1, IKBKG, PIK3CB</i>
HIF-1 signaling pathway	<i>MAPK1, NFKB1, IFNG, PIK3CA, EGF, ERBB2, EIF4EBP1, PIK3CD, IGF1, INS, IL6, SLC2A1, AKT1, HK1, PRKCB, PIK3CB</i>
T cell receptor signaling pathway	<i>MAPK1, NFKB1, IL2, IFNG, IL5, KRAS, PIK3CA, CSF2, GSK3B, PIK3CD, PTPRC, IL10, AKT1, IKBKG, PIK3CB</i>
Acute myeloid leukemia	<i>MAPK1, NFKB1, RARA, KRAS, PIK3CA, CSF2, EIF4EBP1, PIK3CD, AKT1, IKBKG, PIK3CB</i>
Central carbon metabolism in cancer	<i>MAPK1, KRAS, PIK3CA, ERBB2, PTEN, PIK3CD, SLC2A1, NTRK1, AKT1, HK1, PIK3CB</i>
Pancreatic cancer	<i>MAPK1, NFKB1, KRAS, PIK3CA, EGF, ERBB2, PIK3CD, BRCA2, AKT1, IKBKG, PIK3CB</i>
Aldosterone-regulated sodium reabsorption	<i>MAPK1, KRAS, PIK3CA, NR3C2, PIK3CD, IGF1, INS, PRKCB, PIK3CB</i>
Melanoma	<i>MAPK1, HGF, KRAS, PIK3CA, FGF2, EGF, PTEN, PIK3CD, IGF1, AKT1, PIK3CB</i>
Endometrial cancer	<i>MAPK1, KRAS, PIK3CA, EGF, ERBB2, GSK3B, PTEN, PIK3CD, AKT1, PIK3CB</i>
Longevity regulating pathway – multiple species	<i>CAT, KRAS, PIK3CA, SOD1, ADCY2, PIK3CD, IGF1, INS, AKT1, PIK3CB</i>
Allograft rejection	<i>IL2, IFNG, IL5, CD40, IL10, LOC102723407, FAS</i>
African trypanosomiasis	<i>IFNG, IL1B, IL6, IL10, LOC102723407, PRKCB, FAS</i>
Type II diabetes mellitus	<i>MAPK1, PIK3CA, PIK3CD, INS, ADIPOQ, HK1, PIK3CB</i>
Malaria	<i>HGF, IFNG, IL1B, GYPA, CD40, IL6, IL10</i>
Tyrosine metabolism	<i>TPO, MAOA, COMT, MAOB, DBH, DDC</i>
Graft-versus-host disease	<i>IL2, IFNG, IL1A, IL1B, IL6, FAS</i>
Type I diabetes mellitus	<i>IL2, IFNG, IL1A, IL1B, INS, FAS</i>
Asthma	<i>IL5, CCL11, CD40, IL10, LOC102723407</i>
Phenylalanine metabolism	<i>MAOA, MAOB, DDC, PAH</i>