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April 10, 2016

Peripheral cytokine-induced CNS immune activation and effects on mesolimbic dopamine
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Abstract

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Peripheral administration of inflammatory cytokines such as interferon alpha (IFN α) has been shown to reliably induce depressive behaviors in humans and non-human primates (NHPs) that are related to reduced motivation and anhedonia. Peripheral administration of inflammatory cytokines and cytokine-inducers (e.g. endotoxin) has also been shown to reduce neural activation of the ventral striatum to hedonic reward in humans. Our previous work in NHPs suggests that the effects of peripheral inflammation on reward circuitry and behavior may be due to decreased availability and release of dopamine. However, the immunologic and neurobiologic mechanisms by which peripheral cytokines affect the dopamine system is currently unknown. Herein immune activation was assessed in relation to effects on the dopamine system in the ventral and dorsal striatum in post-mortem brain tissue from rhesus monkeys (aged 10-14 yrs) exposed to chronic IFN α (20 MIU/m² s.c. for 4 weeks, n=7) compared to saline control (n=4). Significantly less tissue content of dopamine was observed in the nucleus accumbens (p=0.004) and putamen (p=0.009) of IFN α -treated animals. Whole-gene expression analysis revealed induction of limited immune signaling pathways by IFN α in association with activation of immune cells primarily in vascular and meningeal compartments in or near putamen. Whole genome and targeted analysis also revealed alterations in genes related to dopamine neurotransmission and receptor signaling. These results support our previous findings of decreased striatal dopamine release in IFN α -treated monkeys, and indicate that effects of peripheral IFN α on brain dopamine are likely driven by peripheral immune cells trafficking to the CNS.

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Introduction

Mood disorders currently affect more than 20% of the general population in the United States (Kessler et al., 2003). One such mood disorder is major depression. While some treatment options do exist, more than 30% of those with depression fail to achieve remission of their symptoms (Fava et al., 2006, Mathew and Charney, 2009). Thus, more work is needed to better understand the pathophysiological pathways of depression.

There is evidence of a cause and effect relationship between depressive symptoms and inflammation (Felger and Lotrich, 2013, Felger and Miller, 2014). For example, elevated levels of inflammatory cytokines and other markers of innate immune activation have been consistently observed in the blood and cerebrospinal fluid of patients with major depression (Raison et al., 2006, Raison et al., 2013a, Raison et al., 2013b). Moreover, peripheral administration of inflammatory cytokines has been shown to induce depressive symptoms in both humans and laboratory animals (Haroon et al., 2012). For instance, chronic administration of interferon (IFN) α , an antiviral inflammatory cytokine that was used as an effective treatment for Hepatitis C virus (Dorr, 1993), causes clinical depression and symptoms of anhedonia in many patients (Capuron and Miller, 2004). Additionally, administration of IFN α to laboratory animals, such as non-human primates (NHPs), has been shown to induce both depressive and anhedonic-like behavior (Felger et al., 2007, Felger et al., 2013b). Anhedonia, a deficit in feelings such as pleasure and motivation, is a core symptom of depression as well as an indicator of dysfunction in the dopaminergic reward system (Hickie et al., 1999, Treadway and Zald, 2011, Tye et al., 2013).

Neuroimaging studies in humans have found that inflammatory cytokines decrease neural activation of the ventral striatum to hedonic reward, which correlated with self reported increases in depressive symptoms and reduced motivation (Eisenberger et al., 2010, Capuron et al., 2012); an effect that appears to be mediated by cytokine-induced decreases in mesolimbic dopamine (Capuron et al., 2012, Felger and Miller, 2012, Felger et al., 2013b). Indeed, decreased release of striatal dopamine in NHPs administered IFN α has been found by *in vivo* microdialysis, which correlated with decreased effort-based sucrose consumption, a measure of anhedonia in NHPs (Felger et al., 2013b). Interestingly, IFN α -induced decreases in dopamine release in NHPs were reversed by administration of the dopamine precursor levodopa (L-DOPA) via reverse microdialysis (Felger et al., 2015). Inflammatory cytokines may disrupt dopamine production by inhibition of the enzyme co-factor tetrahydrobiopterin (BH4), which is required for the conversion of phenylalanine to tyrosine by phenylalanine hydroxylase and tyrosine to L-DOPA by tyrosine hydroxylase (Neurauter et al., 2008). Additionally, inflammation-related decreases in BH4 can cause oxidative stress, which can lead to even less BH4 being available for dopamine synthesis (Cunnington and Channon, 2010). Furthermore, patients that have had chronic peripheral administration of IFN α have shown evidence of reduced BH4 activity (Neurauter et al., 2008, Zoller et al., 2012, Felger et al., 2013a), implicating the loss of BH4 as a potential factor in the loss of dopamine synthesis and release. In addition, previous studies in NHPs have also found a decrease in dopamine 2 receptor (D2R) but not dopamine transporter (DAT) binding following chronic IFN α administration, suggesting that cytokines and immune activation may also reduce dopamine receptor expression or function (Felger et al., 2013b).

Mechanistically, peripherally administered cytokines are thought to access the central nervous system (CNS) and may activate microglia and astroglia in the brain. This activation of brain immune cells may then cause local cytokine production of cytokines. For example, when IFN α is administered peripherally over several weeks, concentrations of the inflammatory cytokines interleukin (IL)-6 and tumor necrosis factor (TNF) and their soluble receptors are significantly elevated in the cerebrospinal fluid (CSF) in humans and NHPs administered IFN α (Felger et al., 2007, Raison et al., 2009). IFN α also causes the production of monocyte chemoattractant protein-1 (MCP-1), a chemokine that recruits monocytes and macrophages to the CNS (Collins et al., 1985, Smith et al., 1985, Raison et al., 2009). Previous studies in post-mortem brain tissue from depressed suicides indicate increased macrophage recruitment to perivascular spaces and selective activation or priming of microglia in the neighboring parenchyma of the prefrontal cortex (Torres-Platas et al., 2014). However, no previous study has investigated the pattern of immune cell activation in the brains of IFN α -treated subjects, or in the basal ganglia of depressed suicides. Therefore, it is unknown whether peripheral inflammatory cytokines like IFN α affect dopamine synthesis, release, and/or receptor signaling through direct effects across the blood brain barrier, or whether changes may be driven in part by accumulation of peripheral cells in perivascular and meningeal spaces or activation of local brain immune cells.

Thus, in this study, we tested the hypotheses that chronic IFN α administration was associated with increased accumulation of perivascular and meningeal macrophages, selective activation of local microglial, and increased inflammatory signaling, in association with decreased tissue content of dopamine and genes related to dopamine

synthesis, metabolism or receptor signaling. To determine the immunological and neurotransmitter mechanisms by which chronic exposure to inflammatory cytokines may affect dopamine function, post-mortem tissue from NHPs chronically administered IFN α compared to controls was examined for tissue content of dopamine by high performance liquid chromatography (HPLC), microglia and peripheral immune cell activation by immunofluorescence and confocal microscopy, and changes in gene expression related to inflammatory and neurotransmitter signaling through targeted and genome wide analyses using RNA-sequencing (RNA-seq).

Methods

Subjects and Brain Tissue Collection

Post-mortem brain tissue from rhesus monkeys (*Macaca mulatta*) aged 10-14 yrs administered pegylated IFN α 20 MIU/m² s.c. (rHu-IFN α -2b (Schering-Plough, Kenilworth, NJ)) between 0700 hours and 1000 hours, 5 days per week for 4 weeks (n=7, 3 female), similar to the treatment schedule of patients receiving IFN α monotherapy for malignant melanoma (Musselman et al., 2001), were compared to controls administered saline in equivalent volumes for 4 weeks (n=4, 2 female). All brains were collected following transcardial perfusion with cold 0.9% saline with heparin. The left hemisphere was fixed in cold 4% PFA, sunk in increasing concentrations of sucrose (10 to 30%) in 0.1 M phosphate buffer (PB), and stored at -80° C until sectioning. The right hemisphere was blocked for micro-dissections for RNA extraction and HPLC of neurotransmitters, and stored at -80° C until analysis.

HPLC for Tissue Dopamine Concentrations

Samples extract from basal ganglia tissue was outsourced to Briansonline.org for quantification of dopamine by HPLC with electrochemical detection according to established procedures (Felger and Miller, 2012, Raison et al., 2013a). Desired portions of brain tissue were dissected on dry ice and weighed. Each portion of tissue was then placed in a solution of 0.5 Molar perchloric acid and .01% (w/v) ascorbic acid for each gram of brain tissue at 4 $^{\circ}$ C. The solution was then homogenized using a tissue homogenizer (PowerGen 125, Fisher Scientific). The homogenates were then centrifuged at 4 $^{\circ}$ C for 30 mins at 13,000 rpm. The supernatant was then aliquoted and stored at -80° C

until analysis. Concentrations of dopamine were determined by using electrochemical detection using an internal standard. Samples (10 μ l) were mixed with 20 μ l internal standard (3,4-Dihydroxybenzylamine, DHBA), 20 μ l was injected onto the LC system by an automated sample injector (Gilson XL233, Gilson, France). Chromatographic separation was performed on a reversed phase Hypersil BDS C18 (150 x 2.1 mm, 3 μ m) analytical column (Thermo Scientific, keystone, USA) held at a temperature of 32.5° C. Components were separated using isocratic sodium acetate buffer, pH=4.1. Detection was performed using a potentiostat (Antec Leyden, model Intro, The Netherlands) fitted with a glassy carbon electrode set at +600 mV vs. Ag/AgCl (Antec Leyden, the Netherlands).

Gene Expression

Changes in the transcriptome induced by IFN α compared to control in the caudate and putamen were assessed by RNA-seq in collaboration with Dr. Bosinger, Yerkes NHP Genomics Core. RNA was isolated from microdissected brain tissue stored at -80 in RNAlater, and isolated with RNeasy Mini Kits (Qiagen, Hilden, Germany). RNA-seq was then run on two flowcells of an Illumina HiSeq1000. Reads were aligned to the MacaM version 7 assembly of the Indian rhesus macaque genome (Zimin et al., 2014). Alignment was performed using STAR version 2.4.0g1 (Dobin et al., 2013). Parameters were set using the annotation as a splice junction reference. Transcript abundance estimates were calculated with htseq-count version 0.6.1p1 (Anders et al., 2015). Normalization was performed using DESeq2 version 1.6.3 (Love et al., 2014), producing a both a normalized read count table and a regularized log expression table. There was an average of 19-20 million reads per sample, single end. DESeq2 was also used for differential expression estimation between treated and control samples for each brain

region. A false discovery rate (FDR) <5% was maintained to identify genes most likely to be differentially expressed while controlling the rate of false-positives. Additionally, differentially expressed gene transcripts were defined as having a $\geq 20\%$ difference (1.2 fold change) (Cole et al., 2003, Miller et al., 2008, Mehta et al., 2013, Torres et al., 2013), and were therefore highly predictive based on both statistical significance (p value) as well as biological effect size (fold change). Identified genes were examined for significantly represented pathways and transcription factor networks in Metacore. Compared with microarray, this read depth allows significantly more power to detect differentially expressed genes. A targeted analysis of genes related to the dopamine neurotransmitter system (Bibb, 2005, Bonci and Hopf, 2005) was also conducted as described below in statistical analysis.

Immunofluorescence

The fixed hemisphere was blocked and sectioned at 50-60 micron using either a freezing microtome or Vibratome 1500. Sections were then stored in cryoprotectant solution at -20°C until staining. Sections were rinsed in phosphate-buffered saline (pH 7.6; PBS) and blocked in 0.5% bovine serum albumin (BSA) with PBS at room temperature. Slices were then incubated in primary antibody in tris-buffered saline (TBS), 0.1% Triton-X, and .1% BSA overnight at $\sim 4^{\circ}\text{C}$ while on a shaker. Sections were then rinsed and incubated with secondary fluorescent antibodies in the dark for 1 hour at room temperature. For macrophage/microglia stain, rabbit anti-Iba-1 was used (Wako, 019-19741) at 1:2000. For the microglia stain, mouse anti-human B-lymphocytes was used (HLA-DR/MHC-II) (MP Biomedicals, 0869302) at 1:200. For staining of endothelial cells in the brain vasculature, anti-Von Willebrand factor was used (abcam,

ab6994) at 1:200. Sections were then counterstained with a nuclear stain, DAPI. Sections were mounted, dehydrated, and smooth cover-slipped with an antifade mount solution (Fluro-gel, Electron Microscopy Sciences). Slides were stored in a slide box at 4°C until imaging. Images were acquired using a Leica TCS SP8 confocal system with Hybrid Detector (HyD) (Emory University Winship Cancer Institute). Images were analyzed by collapsing serial Z stacks into a single image using Image J software (NIH, Bethesda, MD).

Statistics

For HPLC data and targeted analyses of dopamine-related gene expression, all statistically analyses were conducted using SPSS (IBM, Armonk, NY). To assess dopamine concentrations in the nucleus accumbens, caudate and putamen in IFN α versus control animals, we used both a one-way multivariate analysis of variance (MANOVA) with between subject contrasts, as well as two tailed independent t-tests and Bonferroni correction for multiple comparisons ($\alpha/3$). Potential group outliers were examined using two-sided Grubbs' tests ("Outlier Calculator" tool in GraphPad). To assess the effect of IFN α versus control on the D2R (gene symbol DR2D) and DAT (gene symbol SLC63A), individual t-tests were conducted separately. Pearson's product-moment correlation coefficient was used to examine relationships between genes expression and dopamine concentrations. All tests were two-tailed with α of .05.

Results

Brain tissue from all 11 NHPs (n=4 control, 7 IFN α treated) was used for HPLC and RNAseq experiments. Fixed tissue from 6 animals, 3 administered IFN α and 3 controls (1 female and 2 males per group), was used for microscopy.

Decreased Tissue Content of Dopamine in Nucleus Acumbens and Putamen

Samples were extracted from three regions of the basal ganglia (nucleus accumbens, caudate and putamen) and dopamine content was analyzed using HPLC. There was no significant main effect of treatment between control and treated animals as assessed by MANOVA ($F[3,9]=2.99$, $p=0.105$). However, independent t-tests demonstrated that there was a significant difference of means for the putamen between control ($M=53.80$, $SD=9.44$) and treated ($M=33.67$, $SD=9.44$) groups ($t=-3.089$, $df=9$, $p=0.013$; Bonferroni corrected $p=0.039$) (**Figure 1**). However, Grubbs' test indicated that one sample from each group was a significant outlier in data from the nucleus accumbens ($p > 0.05$). When these statistical outliers were removed, a significant main effect of treatment was revealed by MANOVA ($F[3,9]=17.30$, $p=0.005$) (**Figure S1**). Between-subject contrast demonstrated significantly decreased dopamine in both nucleus accumbens ($p=0.004$) and putamen ($p=0.009$). Individual t-tests also indicated significant difference in means between the control ($M=17.23$, $SD=0.92$) and treated ($M=5.72$, $SD=4.69$) animals in the nucleus accumbens ($t=-4.13$, $df=9$, $p=0.004$; Bonferroni corrected $p=0.012$).

Changes in Inflammatory and Neurotransmitter-related Gene Expression, Whole Genome Analysis

Genome wide analysis was used to determine genes that were significantly differentially regulated in the brains of IFN α -treated compared to control animals. A threshold of ± 1.20 fold change and an adjusted p-value of 0.05 was used as a cut off. Genes that were significantly differentially regulated at this threshold in putamen and caudate, respectively, are listed in **Tables S1 & S2**. In putamen, 1878 genes were differentially regulated by IFN α treatment. Of these genes, 922 were up-regulated and 956 were down-regulated by IFN α . In caudate, 187 genes were differentially regulated by IFN α treatment. Of these genes, 165 were up-regulated and only 22 were down-regulated by IFN α .

In the 922 up-regulated genes in the putamen, we observed a significant overrepresentation of genes in pathways related to development (“Thrombopoietin-regulated cell processes,” “Bone morphogenetic protein (BMP) signaling,” “Fibroblast growth factor receptor (FGFR) signaling pathway”), hormone signaling (“Ligand-independent activation of estrogen receptor 1 (ESR1) and estrogen receptor 2 (ESR2),” “Membrane-bound ESR1: Interaction with growth factors signaling”), and oncogenes (“Ligand independent activation of Androgen receptor in Prostate Cancer,” “P53 signaling pathway,” “Ovarian cancer”). Of relevance to inflammatory signaling induced by IFN α , several of these top pathways were related to nuclear factor of activated T-cells (NF-AT) signaling (“NF-AT signaling in cardiac hypertrophy”), nuclear factor-kappaB (NF-kB) (“Additional pathways of NF-kB activation in the cytoplasm”), and IFN (“IFN gamma signaling pathway”) (**Figure 2A**). Of relevance to dopamine neurotransmission

and other neurotransmitters that regulate dopamine, pathways related to melanocyte/melanoma related pathways (“Melanocyte development and pigmentation,” “Aberrant B-Raf signaling in melanoma progression”), adenosine receptors (“A2B reception: action via G-protein alpha s”), and glutamate signaling through N-methyl-D-aspartate (NMDA) receptors (“NMDA dependent postsynaptic long-term potentiation in CA1 hippocampal neurons”) (**Figure 2A**). These were the top 20 significant pathways with at least 6 genes represented per pathway ($p < 2.01e-3$). In the up-regulated genes in the putamen, there was also a significant over-representation of genes regulated by a number of transcription factors. These transcription factors were primarily involved in inflammatory and immune signaling (RELA (p65 NF-kB subunit), GATA-3, FOXP3, STAT3), oncogenes (c-Myc, p53, ZNF143, c-Jun, E2F1, IRF4, EGR1), estrogen signaling (ESR1, Sp1, Androgen receptor, RFX2), and cellular processes (Oct-3/4, YY1, NANOG, SOX2, LHX2, H1F1A, C/EBPbeta, NRSF, SP3, CREB). These were the top 24 pathways that had a p-value of $< 1.33e-141$. The up-regulated genes related to the inflammatory and IFN relevant transcription factor, STAT3, is shown in **Figure S2**.

In the 956 down-regulated genes from the putamen, there were a significant overrepresentation of genes related to oxidative phosphorylation (“Oxidative phosphorylation,” “Ubiquinone metabolism,” “ATP/ITP metabolism”), oligodendrocyte processes (“Regulation of cytoskeleton proteins in oligodendrocyte differentiation and myelination”), and cytoskeletal modifications (“Neurofilaments,” “Slit-Robo signaling,” “Regulation of actin cytoskeleton by Rho GTPases,” “Regulation of cytoskeleton proteins”) (**Figure 2B**). These were the top 8 pathways with at least 6 genes represented per pathway ($p < 1.42e-2$).

In the 189 up-regulated genes from the caudate, a significant overrepresentation of genes in pathways involving protein folding (“Posttranslational processing of neuroendocrine peptides”) and immune activation (“Lectin induced complement pathway,” “Classical complement pathway”) were observed (**Figure 3**). These were the top 3 pathways that had at least 3 genes represented per pathway ($p < 2.94e-2$). There were a variety of overrepresented genes associated with transcription factors in the up-regulated genes. These transcription factors were related to inflammatory signaling (STAT3, REIA (p65 NF-kB subunit)), oncogenes (c-Myc, p53, N-Myc, c-Jun), estrogen signaling (ESR1, Sp1, Androgen receptor, RFX2), cellular processes (Oct-3/4, H1F1A, NANOG, YY1, CREB1), and glucocorticoids (GCR, TCF7L2 (TCF4)). These were the top 15 transcription factors that had a p-value of $< 4.4e-17$. While 22 genes were significantly down-regulated in the caudate, none of the pathways identified as significant had more than one gene represented in the pathway. Thus, these pathways were not reported.

Decreased Expression of Dopamine-related Genes, Targeted Analysis

To examine changes in gene expression specific to dopamine synthesis, metabolism and neurotransmission, targeted analysis was performed (**Table 1 & 2**). A one-way multivariate analysis of variance (MANOVA) was conducted to determine if IFN α treatment significantly affected gene expression. In the putamen, there was significantly less expression of the dopamine and cAMP-regulated neuronal phosphoprotein 32 (DARP-32) ($p = 0.044$). Additionally, the gene for catechol-O-methyltransferase (COMT) was significantly decreased ($p=0.004$). Finally, two g-protein inward rectifying potassium channels were significantly decreased in the putamen

(KCNJ4, $p < 0.001$) and increased in the caudate (KCNJ16, $p = 0.030$). Gene expression values from both the caudate and putamen were examined to determine whether there was a significant correlation between dopamine availability (determined via HPLC) and gene expression (via targeted expression data). In the putamen, GNG5, a gamma subunit of a G-protein is stimulated by activation of dopamine 1 and 5 receptors, was significantly decreased (-1.31 fold, $p = 0.006$) and was also positively correlated with the concentration of dopamine ($r = 0.696$, $df = 9$, $p = 0.017$). Individual t-tests revealed that D2R was significantly decreased in the caudate (t value = 2.31, $df = 8$, $p = 0.050$), but not putamen ($p > 0.200$). DAT was not significantly differentially expressed in either caudate or putamen (both $p > 0.24$).

Increased Activated Immune Cells in and around Perivascular and Meningeal Spaces

Brain tissue from rhesus monkeys was sliced, stained and mounted for imaging. Images were taken of the vasculature within the putamen, and of the meninges and bordering tissue of the ventral surface of the brain below the putamen and globus pallidus (**Figure S3**) In one experiment, anti-Iba-1 was used to visualize microglia and peripheral monocytes, and anti-MHC-II was used to label activated microglia and peripheral macrophages. In the putamen, there was an IFN α -driven increase in accumulation of activated peripheral immune cells expressing MHC-class-II (green/yellow cells) in what appeared to be the perivascular spaces of the vasculature. In IFN α -treated animals only, MHC class-II+ activated microglia were observed in the neighboring brain parenchyma that co-localized with Iba-1 (a classic marker for brain microglia) (**Figure 4**). Furthermore, Iba-1+ microglia that did not contain MHC class-II (red cells) of IFN α -treated animals in the putamen appeared in a “primed” state with less ramification and

thicker processes. We also observed increased accumulation of activated peripheral immune MHC-class-II⁺ cells in the meningeal tissue that bordered the outside of the ventral surface of the brain below the putamen and globus pallidus, in IFN α -treated animals compared to controls (n=3 per group) (**Figure 4**). Moreover, MHC-class-II⁺ activated cells were also observed infiltrating the brain tissue parenchyma that bordered the meninges in this area in IFN α treated animals but not controls, which were found amongst the Iba-1⁺ cells (**Figure 4**). Anti-Von Willebrand and MHC-class-II staining was used to determine the location and relationship of activated monocytes or microglia with respect to the endothelial cells of the vasculature in the putamen. In the IFN α exposed animals, there was enlargement of the perivascular space, which also contained an increase in the accumulation of activated monocytes/macrophages (**Figure 5**). Furthermore, this experiment confirmed that activated microglia indeed present outside of the endothelial cells of the vasculature (**Figure 5**), indicating a selective activation of parenchymal microglia localized to tissue surrounding the vasculature. In the putamen of control animals, a few activated monocytes/macrophages were located inside the blood vessels (**Figure 5**). Only immune activation in the meningeal tissue on the edge of the caudate showed activate peripheral immune cells. Although accumulation of these cells were observed in IFN α -treated animals (similar to that observed below the putamen and globus pallidus in **Figure 4**), no parenchymal microglial cell activation was found in the caudate of either treated or control animals (data not shown).

Discussion

The findings presented support the hypotheses that chronic exposure to peripheral cytokines, such as IFN α , affects the dopamine neurotransmitter system most likely via inflammatory signaling driven by recruitment of peripheral immune cells to the CNS. We observed activation of immune cells and immune signaling pathways in the basal ganglia by immunohistochemistry and gene expression analysis, respectively, along with significantly less tissue content of dopamine in the putamen and nucleus accumbens (when statistical outliers were excluded) but not caudate. These findings are particularly interesting and important because they confirmed our previous findings that peripheral administration of cytokines can decrease dopamine synthesis and release in the striatum (Capuron et al., 2012, Felger and Miller, 2012, Felger et al., 2013b). Indeed, our previous study in NHPs found decreased dopamine release after 4 weeks of IFN α as evidenced by reduced displacement of [11 C]raclopride following amphetamine administration in putamen and nucleus accumbens but not caudate. Furthermore, findings that chronic inflammatory cytokine exposure decreased striatal D2R in the caudate without affecting DAT is consistent with that of PET studies demonstrating reduced striatal D2R but not DAT following chronic IFN α exposure (Felger et al., 2013b).

Interestingly, our findings were consistent across all three techniques HPLC, gene expression and immunohistochemistry. For instance, whereas immune cell activation and increases in inflammatory signaling pathways were observed in putamen in association with decreased dopamine by HPLC, very limited immune cell or inflammatory signaling pathway activation, as well as no change in dopamine concentration, were observed in caudate. Thus, since the only other affected pathway in the caudate was neuroendocrine

packaging and there did not appear to be a decrease in dopamine concentration or signaling, it is possible that another neurotransmitter, such as glutamate, may be affected. Indeed, glutamate has known interactions with striatal dopamine (Surmeier et al., 2007) and is thought to contribute to symptoms of cytokine-induced depression (Haroon et al., 2014, Haroon et al., 2016). In fact, our whole genome expression data indicated that exposure to IFN α caused significant up-regulation in of genes in the putamen related to increased NMDA receptor signaling. This could indicate negative downstream effects of inflammation-mediated increased glutamate receptor signaling on neuronal health and neural integrity through excitotoxicity. Indeed, other studies have found that activated microglia, which induce activation of the kynurenine pathway, are a major source of glutamate and quinolinic acid that can lead to depressive-like symptoms (Dantzer and Walker, 2014). The down regulation of oligodendrocyte gene expression in the putamen points to an issue with the glutamatergic signaling system as well, since oligodendrocytes are highly sensitive to excitotoxic environments and to increased inflammation (Domercq et al., 2005) (Matute, 2007). A targeted gene analysis of the glutamatergic signaling system, as well as further analysis on the kynurenine pathway, is necessary to determine the relationship between the glutamate system, peripheral cytokine administration, and its potential affects on dopaminergic signaling.

Also interesting was the fact that the most significantly up-regulated pathway in the putamen were melanin related pathways. It is known that melanin, specifically neuromelanin, is a major player in Parkinson's disease and the disease's hallmark decline of dopaminergic signaling in the basal ganglia (Wilms et al., 2003, Fedorow et al., 2005). It is believed that increased levels of neuromelanin outside of the dopamine cell can

cause microglial activation. Continued microglial activation exacerbates dopaminergic neurodegeneration and creates a feed forward system where increased neurodegeneration causes more neuromelanin to be present outside of a dopaminergic cell (Zhang et al., 2011). It seems likely that dopamine signaling in the putamen of IFN α -exposed animals may be affected in a similar way, except that microglia may be initially activated by chronic inflammatory stimuli to then affect neuromelanin production. Indeed, in our immunohistochemistry analysis, we have observed such activated microglia near the vasculature and meninges in the basal ganglia. Additionally, whole genome gene expression analysis demonstrated that there was down regulation of oxidative phosphorylation pathways in the putamen. This would result in a significant decrease in the amount of ATP available for metabolic processes, and thus indicative of decreased neuronal activity. To confirm if neuromelanin is accumulating in the brains of IFN α treated animals and may contribute to decreased dopamine, tissues from both groups will be compared in future assessments to see if there is an accumulation of the a black-brownish pigment in the IFN α -treated animals (Fedorow et al., 2005).

Several limitations of this work should be noted. First and foremost the small sample size, although 3-5 animals per treatment group are common in non-human primate studies (Czoty et al., 2000, Bosinger et al., 2009, Sawyer et al., 2012, Felger et al., 2013b). Additionally, the sample size of treatment was imbalanced to allow for correlation of findings between dopamine concentrations, gene expression, plasma/CSF markers and behavior in future analyses. Furthermore, additional experiments are necessary to determine whether the outliers identified in the nucleus accumbens are due to procedural issues, experimental error, or truly represent the subject's concentration of

dopamine. Also, our analysis of staining of immune cells was not quantitative. Once staining is complete in all subjects, future analysis will include quantification of immunostaining for activated immune cells using Metamorph software (Chan et al., 2013). Additional staining will be conducted to determine whether gene expression changes reflect changes in protein levels for neurotransmitter-relevant proteins (e.g. dopamine and glutamate receptors), as well as for expression of other immune-related molecules and proteins relevant to neurotransmission (e.g. quinolinic acid). In terms of other future experiments, an analysis of the behavioral symptoms for these specific subjects is also needed to determine if these neuronal changes correlate with behavioral abnormalities. Additionally, cytokines and inflammatory mediators in the plasma and CSF of these monkeys will be measured to relate to immune changes in the brain. These will help determine whether blood or CSF biomarkers of inflammation, which can easily be measured in clinical samples, are representative of inflammatory processes in the brain and their potential effects on dopamine.

This work supports the hypothesis that chronic peripheral inflammation from IFN α exposure has effects on brain dopamine. Future analyses of this data to understand better the mechanisms involved in terms of potential effects on dopamine synthesis or receptor signaling may lead to novel targets for therapeutic strategies to treat the effects of inflammation on the brain and behavior.

Tables and Figures

Table 1. Dopamine Targeted Gene transcripts in the Caudate and Putamen significantly changed in subjects administered IFN α (n=7) compared to controls (n=4). *=pvalue < 0.05

Caudate

Gene ID	Fold Change	P-value
ADCY1	4242.44	0.09
ADCY2	327.79	0.20
ADCY3	2688.54	0.38
ADCY4	67.27	0.84
ADCY5	13276.09	0.12
ADCY6	302.00	0.05*
ADCY7	239.25	0.28
ADCY8	121.84	0.23
ADCY9	875.79	0.52
ADRB1	1.14	0.54
ADRB2	-1.25	0.05*
AKT1	1.01	0.80
AKT2	1.06	0.46
AKT3	1.01	0.95
APP	1.03	0.23
ARRB1	1.03	0.58
ARRB2	1.11	0.24
CACNA1A	1.08	0.28
CACNA1G	1.48	0.36
CACNA1H	1.09	0.25
CACNA1S	1.72	0.22
CALM1	-1.07	0.09
CALM2	-1.12	0.08
CALM3	-1.10	0.19
CALML3	1.07	0.75
CALML4	1.07	0.09
CALML6	-1.08	0.60
CDK5	-1.12	0.20
CASP3	-1.10	0.15
COMT	-1.07	0.30
CREB1	1.07	0.32

CREB3	-1.13	0.07
CYP2D6	1.11	0.11
DDC	-1.06	0.85
DBH	3.20	0.37
DRD1	-1.14	0.17
DRD2	-1.22	0.10
DRD3	1.10	0.43
DRD4	1.30	0.63
DRD5	-1.08	0.70
DUSP1	-1.15	0.34
FOS	1.01	0.96
GNAI1	1.23	0.09
GNAS	1.19	0.07
GNG5	-1.09	0.32
GNG7	-1.33	0.02
GNG8	1.29	0.42
GRK5	1.07	0.47
GSK3A	-1.09	0.09
GSK3B	1.01	0.98
ITPR1	-1.15	0.20
KCNA1	-1.06	0.50
KCNA2	-1.16	0.72
KCNA3	1.13	0.78
KCNA4	1.03	0.69
KCNA5	-1.02	0.93
KCNA6	1.60	0.08
KCNJ1	1.63	0.15
KCNJ10	-1.06	0.42
KCNJ11	-1.01	0.95
KCNJ12	1.07	0.61
KCNJ13	12.16	0.14
KCNJ14	1.10	0.66
KCNJ16	2.18	0.03*
KCNJ2	1.02	0.87
KCNJ3	1.75	0.20
KCNJ4	-1.17	0.14
KCNJ5	2.60	0.22
KCNJ6	1.90	0.30
KCNJ8	-1.23	0.26
KCNJ9	-1.08	0.25

MAOA	1.09	0.32
MAOB	1.13	0.15
MAPK1	1.03	0.70
PDE10A	-1.25	0.18
PDE4A	1.12	0.11
PDE4B	-1.02	0.66
PDE4D	1.05	0.58
PHKA2	1.04	0.58
PIK3CA	1.04	0.70
PIK3CG	1.09	0.67
PLA2G5	-1.13	0.54
PLCB1	-1.15	0.25
PLCB3	1.08	0.52
PNMT	-1.22	0.23
PPP1R1B	-1.21	0.06
PPP3CA	-1.13	0.08
PPP3CB	-1.06	0.23
PPP3CC	1.00	0.96
PRKACA	-1.06	0.14
SLC18A1	2.35	0.23
SLC18A3	1.01	0.93
SNAP25	-1.13	0.11
SNCA	-1.02	0.83
SYN1	1.10	0.14
SYN2	1.15	0.20
SYN3	1.00	0.99
YWHAZ	1.03	0.46

Table 2. Dopamine Targeted Gene transcripts in the Putamen significantly changed in subjects administered IFN α (n=7) compared to controls (n=4). *=pvalue < 0.05

Putamen

Gene ID	Fold Change	P-value
ADCY1	1.23	0.48
ADCY2	1.13	0.06*
ADCY3	1.56	0.15
ADCY4	-1.30	0.64
ADCY5	-1.10	0.76

ADCY6	-1.11	0.17
ADCY7	1.58	0.80
ADCY8	-1.04	0.97
ADCY9	-1.28	0.30
ADRB1	-1.35	0.95
ADRB2	1.02	0.69
AKT1	1.16	0.33
AKT2	-1.22	0.73
AKT3	-1.19	0.01*
APP	1.10	0.31
ARRB1	-1.02	0.11
ARRB2	1.16	0.09
CACNA1A	-1.04	0.16
CACNA1G	1.01	0.13
CACNA1H	1.11	0.02*
CACNA1S	-1.12	0.83
CALM1	-1.35	0.71
CALM2	1.25	0.18
CALM3	-1.28	0.01*
CALML3	-1.41	0.85
CALML4	-1.72	0.85
CALML5	1.02	0.70
CALML6	1.16	0.52
CASP3	1.46	0.24
CDK5	1.02	0.01*
COMT	1.48	0.00*
CREB1	-1.14	0.00*
CREB3	1.53	0.01*
CYP2D6	1.37	0.00*
DBH	1.03	0.66
DDC	1.03	0.69
DRD1	1.21	0.26
DRD1_A	1.28	0.26
DRD2	-1.25	0.20
DRD3	1.17	0.36
DRD4	1.24	0.61
DRD5	1.00	0.59
DUSP1	1.01	0.50
FOS	-1.11	0.45
GNAI1	1.13	0.22

GNAS	-1.11	0.44
GNG5	1.23	0.01*
GNG7	1.36	0.74
GNG8	1.17	0.67
GRK5	1.06	0.94
GSK3A	1.17	0.03*
GSK3B	-1.15	0.01*
ITPR1	1.01	0.17
KCNA1	-1.08	0.11
KCNA10	-1.01	0.48
KCNA2	1.25	0.04*
KCNA3	1.72	0.07
KCNA4	1.17	0.16
KCNA5	1.07	0.76
KCNA6	-1.01	0.24
KCNA7	-1.10	0.29
KCNJ1	-1.18	0.10
KCNJ10	1.07	0.83
KCNJ11	-1.01	0.14
KCNJ12	-1.30	0.17
KCNJ13	-1.14	0.97
KCNJ14	2.16	0.59
KCNJ15	1.02	0.20
KCNJ16	-1.18	0.08
KCNJ18	-1.10	0.48
KCNJ2	-1.02	0.10
KCNJ3	1.09	0.07
KCNJ5	1.43	0.40
KCNJ8	1.29	0.16
KCNJ9	1.29	0.26
MAOA	-1.37	0.25
MAOB	3.06	0.50
MAPK1	-1.23	0.08
PDE10A	-1.16	0.18
PDE4A	1.08	0.71
PDE4B	1.20	0.53
PDE4D	1.24	0.12
PHKA2	1.17	0.23
PIK3CA	1.22	0.03*
PIK3CG	1.02	0.30

PLA2G5	-1.32	0.18
PLCB1	-1.04	0.12
PLCB3	1.10	1.00
PNMT	-1.05	0.10
PPP1R1B	1.03	0.04*
PPP3CA	1.06	0.53
PPP3CB	1.04	0.65
PPP3CC	-1.03	0.14
PRKACA	1.17	0.80
PRKACA_A	2.27	0.80
SLC18A1	2.20	0.43
SLC18A3	1.18	0.88
SNAP25	-1.12	0.40
SNCA	1.22	0.51
SYN1	0.00	0.90
SYN2	-1.05	0.08
SYN3	-1.30	0.00*
YWHAZ	1.23	0.60

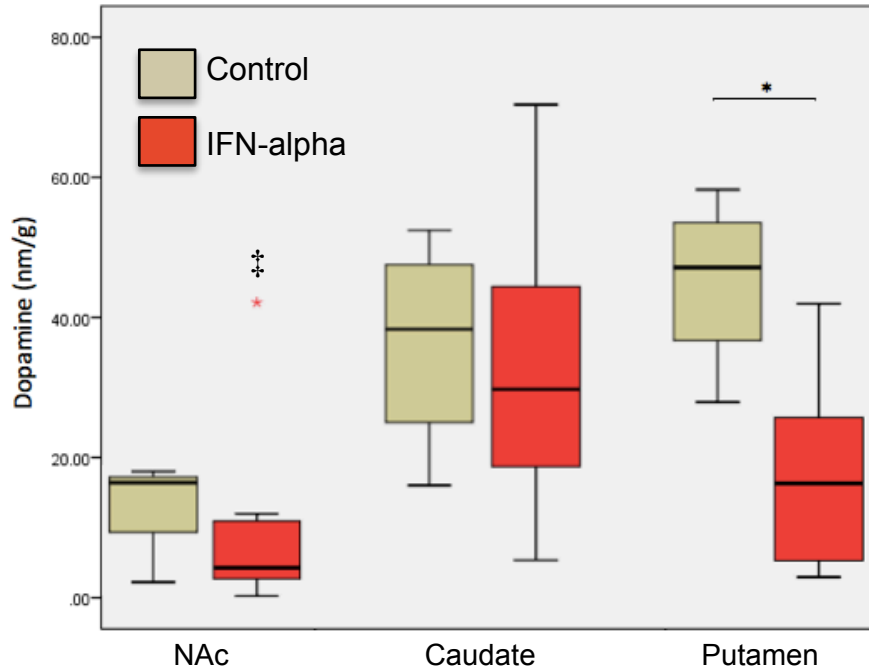


Figure 1. Dopamine Concentrations in the Basal Ganglia. Administration of interferon IFN α for 4 weeks (red boxes, n=7) compared to controls (tan boxes, n=4) significantly decrease tissue content of dopamine in the putamen, but not nucleus accumbens (NAc) or caudate. However, statistical outliers were observed in both IFN α and control groups in NAc. When these outliers were removed, a significant decreased in NAc dopamine was also observed. *p<0.01, ‡p<0.01 after removal of statistical outliers.

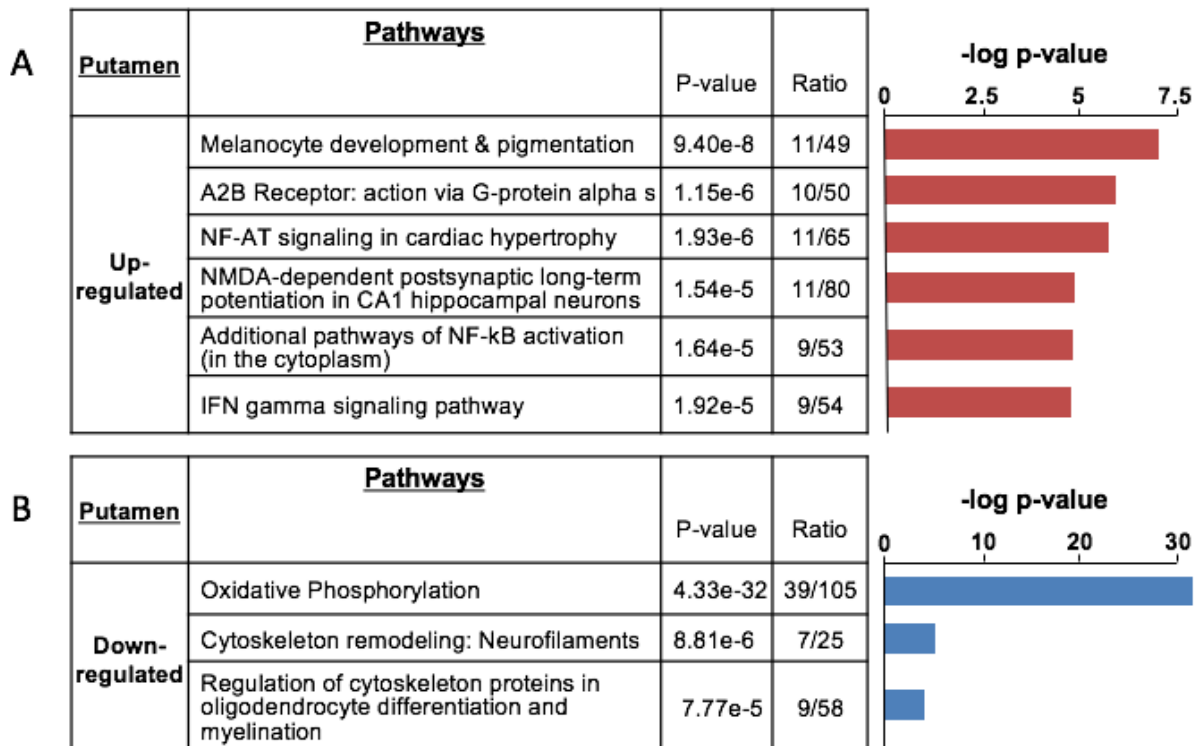


Figure 2. Pathways significantly affected by IFN α administration in treated animals compared to controls in the putamen. Pathways of interest that were significantly overrepresented in up-regulated (A) or down-regulated (B) genes in the putamen. Significant genes were defined as adjusted $p < 0.05$ and fold change of 1.20 cutoff.

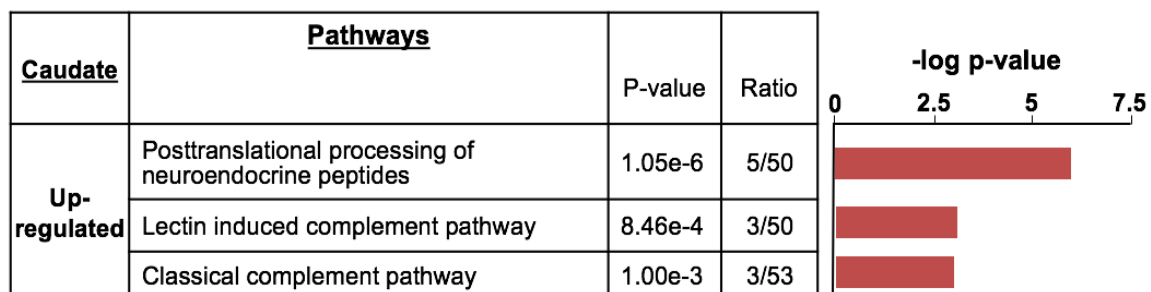


Figure 3. Pathways significantly affected by IFN α administration in Treated animals compared to Controls in the caudate. Pathways of interest that were significantly overrepresented in up-regulated genes in the caudate. Significant genes were defined as adjusted $p < 0.05$ and fold change of 1.20 cutoff. No significant pathways were observed in down-regulated genes from the caudate.

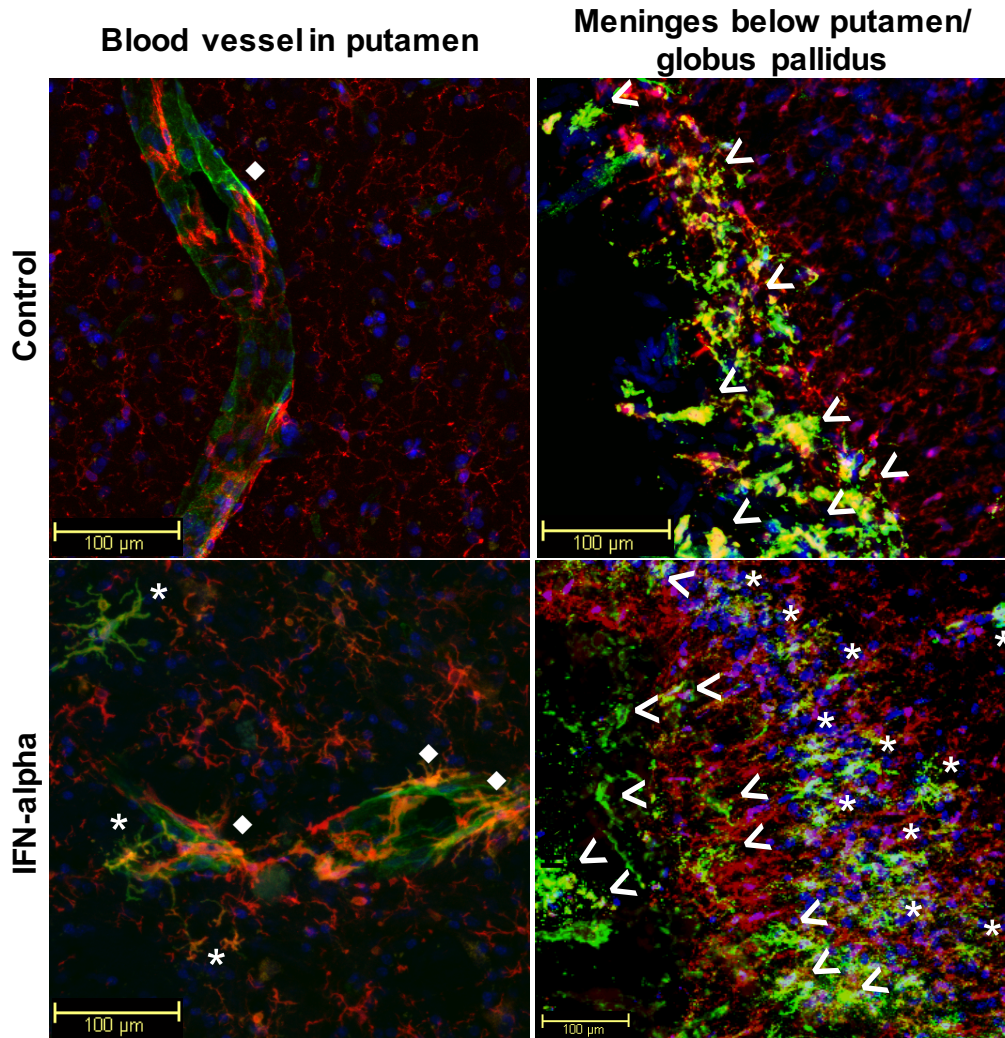


Figure 4. Interferon IFN α increased monocyte/macrophages in perivascular and meningeal spaces and activated parenchymal microglial in basal ganglia. In the putamen, IFN α increased accumulation of activated peripheral monocytes/macrophages (diamond = activated MHC class-II+ cells in what appeared to be perivascular spaces), and activated neighboring parenchymal microglial cells (asterisks = activated MHC class-II+ cells amongst red Iba-1+ microglia) in what was consistent with the neighboring parenchyma outside of the vasculature. Furthermore, in the putamen of IFN α -treated animals, Iba-1+ microglia that did not contain MHC class-II (red cells) were in a “primed” state with less ramification and thicker processes compared to that of controls. IFN α also increased accumulation of activated macrophages in meningeal spaces (arrow heads = activated MHC class-II+ peripheral cells in meninges and tissue border) just below the putamen and globus pallidus, and also activated immune cells in the neighboring tissue parenchyma (asterisks). Green=MHC class-II, red=Iba-1, blue=DAPI, scale bars = 100 μ m

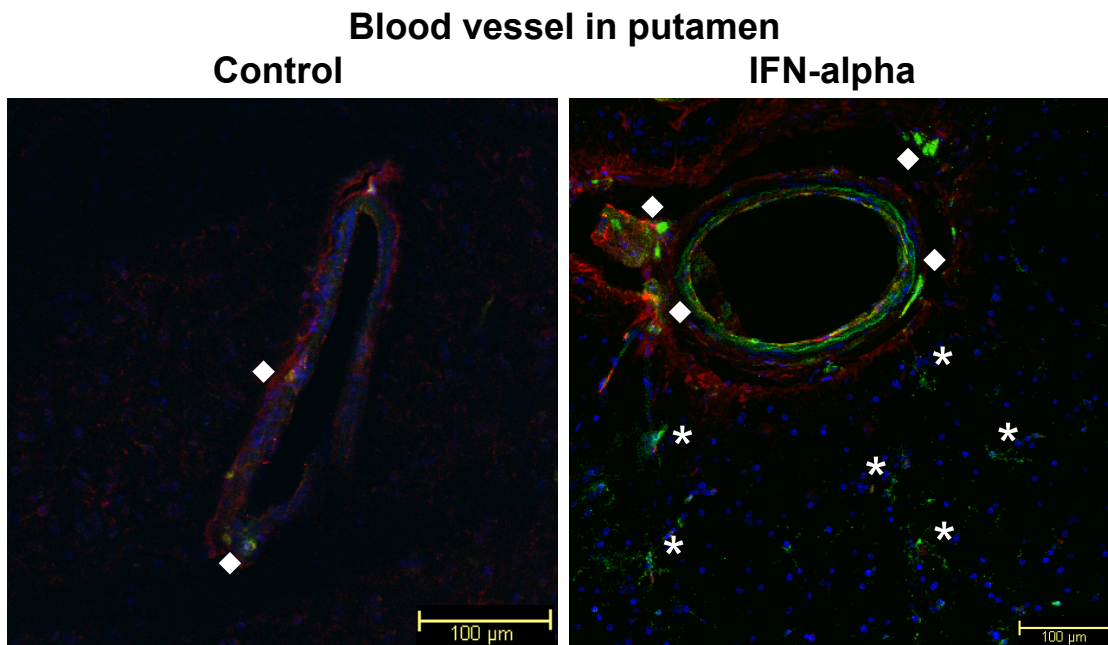


Figure 5. Interferon IFN α increased monocyte/macrophages inside of perivascular spaces and activated parenchymal microglial in putamen outside of blood vessels. In the putamen, anti-Von Willebrand factor staining for blood vessel endothelial cells (red) revealed large perivascular compartments in IFN α -treated animals and confirmed that they contained increased accumulation of perivascular cells (diamond = activated MHC class-II $^+$ inside of the blood vessel endothelial cells). This experiment also confirmed the presence of activated microglial cells (asterisks = activated MHC class-II $^+$ cells outside of the blood endothelial cells) in the neighboring parenchyma in putamen of IFN α -treated but not control animals. Green=MHC class-II, red= anti-Von Willebrand factor (endothelial cells of blood vessels), blue=DAPI, scale bars = 100 μ m

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Supplemental Materials

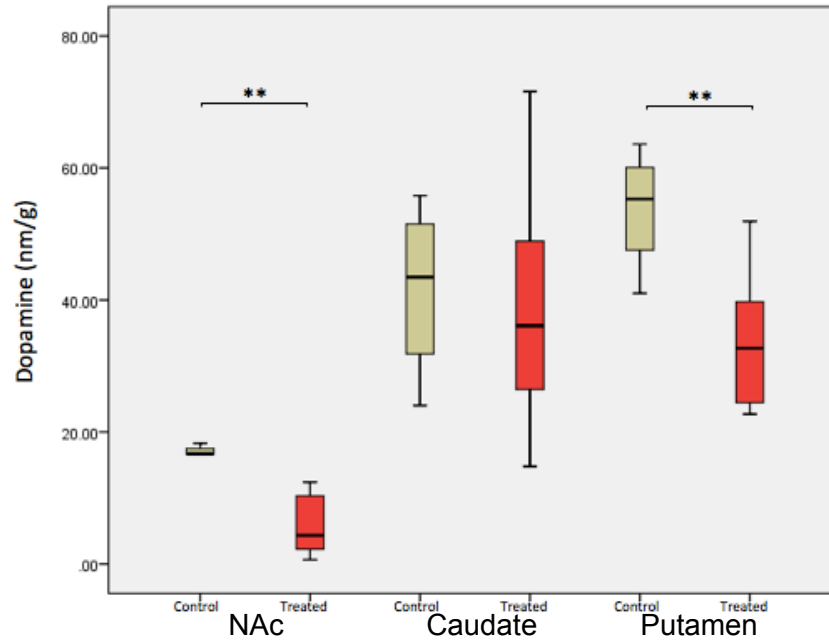


Figure S1. Dopamine Concentration in the Basal Ganglia (outliers dropped). After removal of statistical outliers in the nucleus accumbens (NAc), there was a significant difference in tissue dopamine content between interferon (IFN) α treated (red boxes) and control (tan boxes) animals in the NAc and putamen. ** p < 0.01

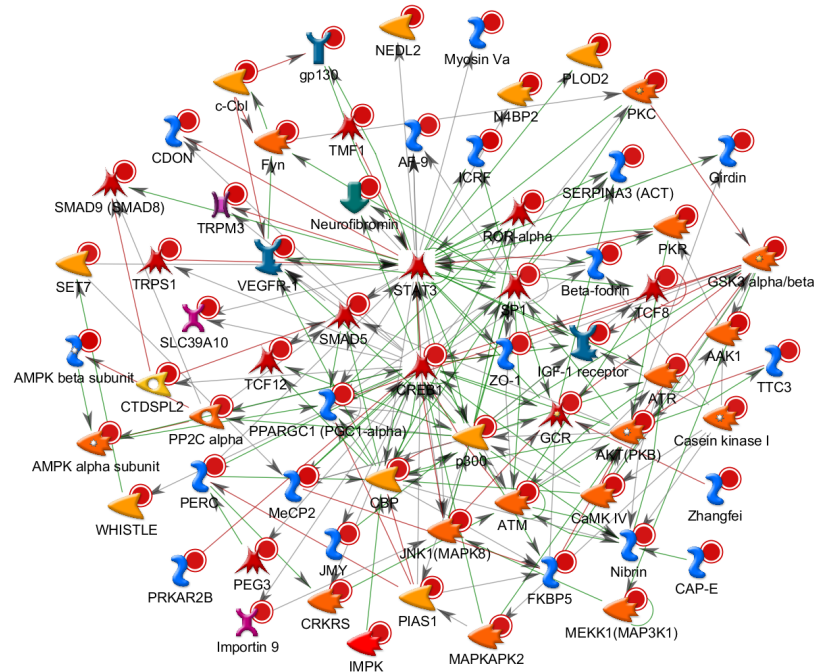


Figure S2. Up-regulated genes in putamen in a network with the IFN-related transcription factor STAT3. Effector genes and transcription factors that were up-regulated by interferon (IFN) α compared to control in the putamen that are significantly overrepresented in a network with the transcription factor STAT3. $p=1.44e-141$.

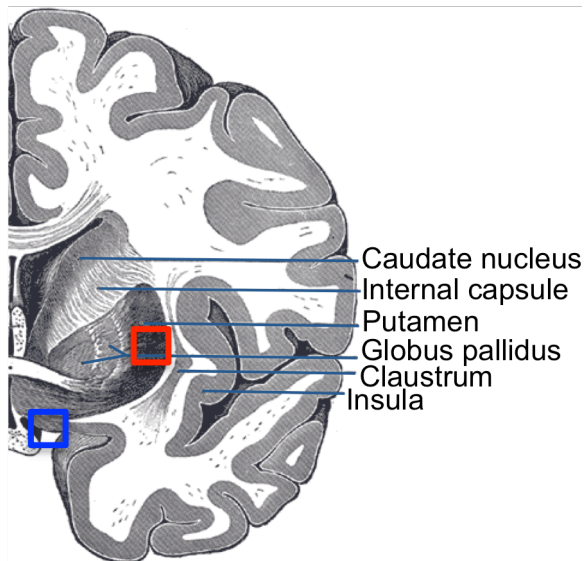


Figure S3. Location of acquisition of confocal microscopic images in basal ganglia. Activated immune cells were imaged in the perivascular spaces of putamen (red square) and meninges and neighboring parenchyma below the putamen and globus pallidus (blue square).

Table S1. Gene transcripts in the putamen significantly up- and down-regulated in subjects administered IFN α (n=7) compared to controls (n=4).

Up-Regulated Genes		
Gene-ID	Fold change	Adjusted P-value
FAM126B	1.673446159	0.001705164
SLC30A6	1.654516838	0.001227029
LCOR	1.549618326	3.82989E-05
HCN1	1.545725448	0.004815335
SLC4A7	1.54095187	0.000260465
ARHGAP26	1.53279169	0.002184831
SKIL	1.528562587	0.007104761
UNC5D	1.517132822	0.006477586
FEM1B	1.511572567	0.007642703
FRRS1L	1.511290205	0.008109344
NFAT5	1.502956998	0.008908212
GMCL1	1.502232552	0.001842061
MAN1A2	1.501493718	0.007772611
DNAJB14	1.500224637	0.007813049
DUSP18	1.49927413	0.006953418
SYN3	1.494883178	0.005030119
RRP8	1.494572104	0.008963882
ZNF770	1.493366521	0.005146403
TRIP11	1.492520735	0.003409352
FUT9	1.490785077	0.009645686
GFOD1	1.488356037	0.009892837
LONRF3	1.487830816	0.009970324
TECPR2	1.485004795	0.009858217
UHRF1BP1	1.484362405	0.006971634
HIPK3	1.483350466	0.009858067
NRIP3	1.482315826	0.004815335
PRKAB2	1.481655593	0.009484345
ZBTB24	1.48145695	0.00586721
ITSN2	1.478661728	0.004705909
TLL7	1.476879007	0.011290929
HOOK3	1.476147406	0.010725123
ZNF546	1.476138552	0.001841085
ZNF253	1.476027383	0.007513568

CELF2	1.473694261	0.007388573
TMEM106B	1.473196442	0.004705909
STOX2	1.472714718	0.011758399
DOCK5	1.472543753	0.011758399
STXBP5L	1.47209931	0.008738552
SAMD12	1.471453704	0.008863773
LMBRD2	1.469575752	0.011290929
ASXL2	1.468980497	0.010802779
XKR4	1.468750647	0.008963882
AMER1	1.466639255	0.001281582
RALGPS2	1.464664887	0.008491744
PLEKHH1	1.464269012	0.00579249
NCOA2	1.463838981	0.011904999
BRAF	1.462138738	0.012970934
RUFY2	1.460819524	0.007740931
ACER2	1.460610488	0.009483478
SOGA1	1.457280902	0.01313309
HMBOX1	1.455210339	0.012144282
KLHL11	1.454792281	0.013654591
MAP3K2	1.452979086	0.009120975
VPS13C	1.451831563	0.012750185
LRCH3	1.44950236	0.007642703
NHSL2	1.448001993	0.015447614
RGP1	1.446084354	0.011904999
MAP9	1.444404667	0.015447614
SLITRK6	1.444261666	0.011904999
EPM2AIP1	1.443133299	0.007642703
NEK10	1.443078701	0.01407053
SNX29	1.443007421	0.013057252
TOR1AIP2	1.441975486	0.012810041
GRAMD1B	1.439512512	0.008963882
KIF1B	1.437411058	0.016652026
PGM2L1	1.437311244	0.00356776
SLC24A4	1.436039231	0.011535851
COL4A5	1.435932354	0.006415636
MOB1B	1.435151535	0.0154935
LRP6	1.433391024	0.017658097
PGGT1B	1.43324365	0.017033456
GFPT1	1.432352907	0.01184407
MALT1	1.431571305	0.018185236

ZC3H8	1.431382947	0.009674447
PPP4R2	1.431348436	0.00557412
MED28	1.430797357	0.016047375
NFASC	1.429315301	0.003974097
SYT14	1.428900324	0.0154935
AGL	1.428146681	0.006283193
MDN1	1.427114095	0.004815335
SOGA3	1.427077772	0.015788454
ZNF845	1.426746449	0.009664692
TSHZ2	1.424816525	0.008738552
FNIP2	1.422738499	0.009858217
LPGAT1	1.421681998	0.003068524
EDIL3	1.421192665	0.011535851
HEATR1	1.420550788	0.008351954
BCHE	1.419883166	0.020728579
DAB1	1.41976807	0.017500413
KIAA1958	1.41954626	0.0197776
ATP7A	1.419310694	0.009484345
FAT1	1.418852108	0.008738552
C2CD3	1.418543512	0.013376019
PPM1K	1.417845008	0.006940669
CHRM3	1.417742917	0.021300163
ZFHX4	1.417648687	0.002392503
FAM169A	1.41647282	0.009790578
ZNF292	1.416036786	0.003960811
TET2	1.415895665	0.001705164
PLXNA4	1.415745034	0.020778341
LARP4	1.4152723	0.021111349
ZKSCAN8	1.414607706	0.021861218
TRUB1	1.414605517	0.019645358
RBM28	1.414536003	0.015770183
HHIP	1.413507224	0.022120754
GRM1	1.413236756	0.022120754
CLSTN2	1.412348482	0.022344274
ZNF354C	1.409904937	0.019816174
SDK2	1.409546347	0.020955744
C10orf118	1.409331179	0.009858217
ZNF462	1.409295972	0.011904999
CDH7	1.409224531	0.018845137
ERI2	1.408112388	0.007015186

VTI1A	1.40809437	0.015146862
RNF169	1.407405436	0.015788454
MYO9A	1.406123167	0.024109681
CLMN	1.405926772	0.008544808
SLC30A4	1.405807931	0.019875258
CPEB3	1.40553304	0.004815335
ABCA5	1.405304879	0.008963882
PTPRG	1.404776665	0.01863106
PLEKHM3	1.404411934	0.019816174
UBXN7	1.404364453	0.023088572
SHPRH	1.404177991	6.0804E-05
KIAA1244	1.404138565	0.018006554
MFAP3L	1.403343575	0.02473979
NHLRC2	1.403121151	0.023787082
SEMA6D	1.402880747	0.024427672
SMG1	1.402789195	0.003321961
PREPL	1.402572161	0.015507835
TTBK2	1.401991898	0.024172564
RGS7BP	1.400578126	0.022239584
C9orf72	1.400168432	0.017356917
JRK	1.400096494	0.01184407
GRIN3A	1.397601818	0.022130072
C21orf91	1.397478511	0.024630699
ALS2CR8	1.396990968	0.011514438
ADD2	1.396921978	0.023191406
UTRN	1.396192636	0.009120975
CFLAR	1.395943329	0.012464937
TMEM41B	1.395812193	0.022520735
ABL2	1.395263368	0.0260344
GPATCH2L	1.394909256	0.024672425
TRPM3	1.394778299	0.025084018
N4BP2	1.394659842	0.018922672
PLOD2	1.394560973	0.006512065
ARFGEF2	1.39400149	0.004927559
PROX1	1.393962453	0.009674447
NYAP2	1.39335688	0.026989863
ARL5B	1.39313444	0.027105868
WDR36	1.392827104	0.000892318
KITLG	1.392443103	0.005721467
CBL	1.391872572	0.022979515

ZNF45	1.391769666	0.006469316
INTU	1.391232747	0.006951912
HIP1	1.39090042	0.007643175
ZBTB16	1.390364831	0.023700158
ETAA1	1.390207642	0.00428854
HECW2	1.389944178	0.009674447
RASA2	1.389521489	0.02633489
IRAK3	1.389250731	0.011290929
UVSSA	1.389161776	0.024069776
AMER2	1.389098161	0.022040256
NIN	1.388544949	0.028788498
ATP11B	1.388440742	0.001227029
FERMT1	1.388016775	0.018963427
AAK1	1.387955946	0.010994693
CASC1	1.387773109	0.018631539
CLN8	1.386830834	0.009664692
SLC38A1	1.3865324	0.009858217
BRWD3	1.386372586	0.006361441
KCND3	1.385648822	0.029876874
FAM168A	1.385413091	0.016321343
SCN3A	1.385247875	0.007640433
TNKS	1.38419906	0.001705164
AFAP1	1.384041524	0.020797838
CYP7B1	1.383136159	0.020323404
ELAVL2	1.383004163	0.002212115
IGSF9B	1.383000293	0.030456202
MTR	1.382525376	0.030456202
PTPRD	1.382073089	0.024363791
AFF4	1.381758758	0.002318461
RNPC3	1.381668148	0.00648968
CDH9	1.381521231	0.025084018
ZNF16	1.38089655	0.003960811
CNTRL	1.380659348	0.021300163
MTMR9	1.380337192	0.009513256
CHD6	1.380011236	0.008926783
MARCH6	1.379730015	0.014468853
PTPLB	1.379197675	0.022120754
DIP2A	1.377873328	0.017056121
ATRN	1.377857463	0.01212472
C2CD2	1.377773287	0.02061029

ATG2B	1.377018366	0.001281582
SLC4A8	1.376997555	0.005528176
SMCHD1	1.376965674	0.000378597
UBN2	1.376481973	0.002919494
CEP63	1.375736518	0.00648968
BMPR2	1.37554421	0.025830266
ABCC4	1.375302506	0.033247749
TTC39C	1.37445701	0.01189794
RICTOR	1.374257181	0.004197473
RALGPS1	1.373968923	0.029461297
CNTNAP5	1.37394254	0.022979515
TCEANC2	1.37370544	0.026696033
ST8SIA1	1.373622966	0.007740931
SMAD9	1.37285168	0.009858217
GRIN2A	1.372312062	0.020142188
MAML2	1.372157286	0.020778341
TARBP1	1.371352361	0.004975442
ACER3	1.37097776	0.017017924
ZNF623	1.370347646	0.010852695
ATRX	1.370166232	0.006274202
ENTPD4	1.369995223	0.0197776
CXADR	1.369666785	0.017790469
FIGN	1.369646099	0.033389338
DGKE	1.369106579	0.031800085
GABRA2	1.368998883	0.004705909
BTA1F1	1.368496151	0.008906854
TTC39B	1.3682814	0.028911221
GNAQ	1.36777497	0.028764432
GEN1	1.367262816	0.031451737
ZFC3H1	1.367248601	0.002318461
VPS53	1.367225415	0.022593859
ALG11	1.36715763	0.036659577
ERBB4	1.365770424	0.025084018
ADCY10	1.365757735	0.007640433
TCP11L1	1.365177568	0.023892993
C6orf141	1.363798425	0.02140671
ATM	1.363771032	0.007553939
EXPH5	1.363537057	0.029559575
RAPH1	1.362450963	0.018185236
HAUS3	1.362329614	0.033972225

RAB23	1.362209335	0.021192371
C4orf29	1.361954871	0.022120754
PURB	1.361944805	0.039016286
RXFP1	1.361753655	0.031975336
SLC4A10	1.361581172	0.005944049
EIF2AK2	1.36021149	0.021848687
MGA	1.360069567	0.006416367
ZNF610	1.359825409	0.028534263
CEP350	1.359798424	0.009924803
KCNH8	1.359758553	0.030180473
ZBTB40	1.359319913	0.017950273
TSSK4	1.359088437	0.03274676
UGCG	1.358941646	0.025671159
RORA	1.358789764	0.004996376
ELOVL7	1.358376068	0.025084018
ST8SIA3	1.358348355	0.018569445
PHKA1	1.358335252	0.008436737
SGPP2	1.358297518	0.024843798
OPCML	1.357943128	0.010349222
MITF	1.357833302	0.036264465
TRIM44	1.357583545	0.007889473
ZNF704	1.35753147	0.031818025
EEF2K	1.356714062	0.034310397
AKT3	1.356596486	0.016855761
KIAA1549L	1.356406242	0.000297773
GPR113	1.356293033	0.028336583
FLT1	1.356219666	0.032943229
CEP85L	1.35603595	0.041240934
ZNF780A	1.355898699	0.0331433
TMEM245	1.355846474	0.002318461
CAMSAP2	1.355505259	0.001686657
KCNQ3	1.355494481	0.041374742
MBD5	1.355355304	0.005944049
DGKH	1.355240391	0.039562163
PIAS1	1.355096399	0.034765061
PBRM1	1.35497157	0.001705164
GLRA4	1.354578998	0.042404935
MSMP	1.353991402	0.028877571
PIK3CA	1.353970724	0.017500413
CCP110	1.353446325	0.002212115

ENTPD5	1.353196853	0.037369311
ZC3H13	1.353111711	0.011755497
LRIG2	1.352997324	0.015447614
XRN1	1.352892146	0.007947364
GUCY1A2	1.352519782	0.016261403
WSB1	1.35196357	0.001705164
ADAMTS6	1.351736009	0.021510168
SSH1	1.351673308	0.041821533
WWC2	1.351210796	0.029070016
KCNB1	1.351151569	0.030107377
NCOA3	1.351115492	0.005174853
TMF1	1.351083469	0.011421403
WDFY3	1.350903277	0.007513568
PAK3	1.350823142	0.037489112
SYNRG	1.350148566	0.007957759
SV2C	1.349810317	0.022499184
MAP3K9	1.349527746	0.045013782
LRP1B	1.349491192	0.028700037
PLXDC2	1.349055631	0.045066388
KIAA1586	1.348921068	0.035619775
CTDSPL2	1.347686259	0.004927559
INTS2	1.347469037	0.015800261
GPR83	1.347385404	0.028762545
PHC3	1.347040011	0.00356776
RASAL2	1.34695349	0.002315348
PCLO	1.34690658	0.007813049
DOPEY1	1.346317518	0.01046166
TRANK1	1.346159275	0.034765061
OGFRL1	1.346140288	0.037964311
SIPA1L1	1.345925984	0.009244314
FKBP5	1.345873776	0.039811942
ZMAT3	1.345526402	0.030786595
SMC2	1.345509145	0.029874027
PHIP	1.345391659	0.007196388
OTUD4	1.345171509	0.002315348
LPCAT2	1.344793502	0.040386875
AVL9	1.34473507	0.045854294
SECISBP2L	1.344455989	0.006692614
TNIK	1.344245353	0.019611503
AKAP6	1.344024105	0.013759989

TAF2	1.343949567	0.001705164
DMXL1	1.343568057	0.011775813
RC3H2	1.343562735	0.031648258
RAPGEF6	1.343311995	0.000378597
GSK3B	1.342824154	0.033288852
ANKRD52	1.342750264	0.029906945
BROX	1.342539824	0.030107377
ZNF236	1.342426936	0.010802779
PDZD8	1.342273787	0.030456202
PDS5B	1.34221681	0.005174853
MKL2	1.341878164	0.00428854
ERCC6L2	1.341754937	0.036611182
ODF2L	1.341376854	0.028214042
CENPF	1.341110589	0.038918656
CCSAP	1.340481881	0.021249213
WDFY2	1.340429331	0.042660196
BTBD7	1.339984329	0.046615139
AKAP11	1.339834457	0.013287069
TTPAL	1.339754242	0.024843798
KCNT2	1.339656692	0.040911557
CCDC80	1.339621123	0.035171404
ST18	1.339615861	0.008280206
ZNF33B	1.339496903	0.021249213
PITPNC1	1.339365108	0.028687065
PDPR	1.33922579	0.047017737
BDP1	1.339144521	0.038201547
SLC39A10	1.33890159	0.027867855
CAMK4	1.338884397	0.042548855
RAD54L2	1.338707457	0.037382551
POLK	1.338490045	0.039496042
DHX33	1.338091829	0.017950273
SLC12A6	1.337729122	0.024522575
TRMT44	1.337422331	0.035910039
PANK3	1.337055906	0.049403561
DCTN5	1.336548891	0.04050057
MED13	1.33647864	0.011498263
KLHL28	1.336379068	0.007947364
MAPK8	1.336331282	0.021300163
ZNF778	1.335910755	0.047451056
ZFP30	1.335593738	0.026199537

RGS17	1.335486624	0.043209514
CLVS2	1.335386558	0.042126687
DYNC2H1	1.334783031	0.031547632
KAT6A	1.333941296	0.013722216
RIF1	1.333891887	0.040806463
ZNF609	1.333778644	0.004197473
OAS3	1.333663576	0.044758482
KIAA1109	1.333329207	0.014461968
CSMD1	1.332787764	0.018922672
LIMS1	1.332526083	0.039409174
HELZ	1.332508093	0.003882908
ZNF354B	1.332390294	0.025299478
USP45	1.331591819	0.030111451
PRKCI	1.330996956	0.004862503
LACC1	1.330967435	0.044729241
DMXL2	1.330891311	0.012347375
NF1	1.330445014	0.007640433
TRAF5	1.330108518	0.024408017
ACAP2	1.330045044	0.000283858
DICER1	1.329675031	0.011758399
ATE1	1.329373639	0.028627382
AASDH	1.32904139	0.007740931
RGS8	1.329007793	0.032711986
RNF168	1.328744908	0.039562163
SLC46A3	1.328199198	0.009858217
PTAR1	1.328006934	0.021861218
WDR37	1.327971923	0.003585441
SUSD5	1.327544746	0.034490759
PCYOX1	1.327510177	0.009464122
PCDHA9	1.32706052	0.038114863
ZBTB44	1.326149783	0.001227029
PVR	1.326096501	0.037974114
PCDHGC4	1.325579209	0.033831034
SORT1	1.32552064	0.034707422
MYO5A	1.325275713	0.009858217
IGF1R	1.325269597	0.009419791
ZNF81	1.325250976	0.048759874
PNMA2	1.32517149	0.038132417
CYP2D6	1.325034188	0.004927559
HTR2C	1.324899052	0.009306156

CALCRL	1.324729381	0.039551961
CHD9	1.324024605	0.019816174
ARL5A	1.323699393	0.009120975
IQSEC3	1.323609991	0.00933751
SNX19	1.323559959	0.015698753
PLCXD3	1.322993421	0.033213799
PIGN	1.321605535	0.029876874
CPD	1.321502026	0.008926783
SLK	1.321303025	0.015103475
GRIP1	1.321053362	0.011775813
DHFR	1.320684047	0.010072491
SYT11	1.320611306	0.041936058
C14orf101	1.320313593	0.014284071
SMC5	1.319455935	0.0053165
YPEL2	1.319311613	0.033477246
SHROOM2	1.319174368	0.007772611
CDH19	1.319083017	0.021333769
PCDH7	1.318752058	0.011759109
PTPN4	1.318643406	0.004197473
AFF2	1.318591142	0.049869657
ZNF605	1.318402754	0.019645358
RFX7	1.318301024	0.011904999
KLHL36	1.318218859	0.045690716
ST8SIA4	1.318194242	0.028271767
FRMD5	1.318030896	0.013946966
PPIG	1.317972165	0.024550139
CREBZF	1.317628938	0.010344235
ZC3H6	1.317482694	0.006361441
SPATA13	1.317103112	0.018922672
ATAD2B	1.317008994	0.033867719
MED23	1.316531233	0.005581278
DIP2B	1.316500018	0.010444402
FNIP1	1.316481791	0.008738552
C2orf49	1.316330683	0.0260344
C5orf24	1.315975789	0.001705164
PLEKHA1	1.315856538	0.001705164
PXK	1.315812834	0.006283193
ZCCHC6	1.315531802	0.010233583
ATP8A1	1.31538828	0.006274202
ZMYM5	1.314927633	0.018610351

BOD1L1	1.314658192	0.01509225
ZDHHC21	1.314629271	0.010349222
SMAD5	1.313513337	0.007640433
FBXL20	1.313131914	0.026476768
LARP1B	1.31294737	0.009858217
STX17	1.312850129	0.005864323
MTF1	1.312666201	0.026690313
MKLN1	1.312554967	0.038543469
PRKD3	1.312310557	0.022495251
TRAK2	1.312107016	0.003321961
IL6ST	1.312005193	0.017585805
PCDHB16	1.311974277	0.03364734
BIRC6	1.311939324	0.019962663
MIB1	1.311367719	0.006628549
FCHO2	1.31043578	0.01611736
RBM12B	1.310066128	0.011775813
INADL	1.309858981	0.028627382
TRIM66	1.309544442	0.013366911
DPY19L4	1.309129955	0.039616959
WHSC1L1	1.309073821	0.020955744
GMEB1	1.308770256	0.024869825
MORC3	1.308750741	0.005357908
CEP120	1.308489015	0.002214478
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PEX5L	1.307172104	0.038605702
LIMD1	1.307137379	0.028600617
TOM1L1	1.306936499	0.034802506
QSER1	1.306892357	0.015262895
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PIKFYVE	1.306665943	0.013302586
NETO1	1.306501457	0.017356917
SCN1A	1.306356051	0.037396945
ZNF385D	1.306247663	0.04706971
IPP	1.306141854	0.039551961
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MAP3K1	1.305371155	0.010349222
EEA1	1.305188541	0.017585805
CDH22	1.304948256	0.024733332
RSF1	1.304928968	0.047561271
NKTR	1.304813751	0.017179796
HEATR5A	1.304659664	0.048712798
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ATP13A3	1.30399994	0.003960811
PGAP1	1.303899062	0.045530897
XPO4	1.303882541	0.018300826
TRIM33	1.302576984	0.007422927
CNOT6L	1.301747963	0.006512065
AGAP1	1.301639847	0.004705909
ZNF595	1.301520316	0.013302586
SYNM	1.301097805	0.024700299
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VAV3	1.300558866	0.039841286
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CHL1	1.300018498	0.024636896
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TMEM135	1.299195271	0.042837347
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WDR17	1.29728398	0.024843798
LPHN2	1.29722057	0.008929516
L3MBTL3	1.297144929	0.02061029
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GDAP1	1.296804952	0.003337376
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EFR3A	1.296172083	0.006355846
FAM135A	1.296055238	0.037627102
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PPIL4	1.294172815	0.045377024
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PTPRB	1.29294998	0.033780542
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BCAP29	1.292091357	0.022747603
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SPEN	1.291633835	0.009858217
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SPTBN1	1.291234929	0.019773717
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ASH1L	1.288357369	0.018961771
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NAV2	1.287626504	0.017790469
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CPEB4	1.287389406	0.013057252
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CCNT1	1.286067841	0.010994693
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CHM	1.28354499	0.00933751
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SPATA5	1.280046174	0.04087647
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KLHL24	1.276968863	0.004927559
TRPS1	1.276689206	0.021107666
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ADAM10	1.27542896	0.037152417
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PALM2	1.274479832	0.022979515
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DIS3	1.241029798	0.022236163
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CERS6	1.240631567	0.011068238
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MACF1	1.238977655	0.036611182

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ASCC3	1.238393038	0.025145719
USP36	1.23828859	0.049103485
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CREB1	1.235195209	0.007643175
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GALC	1.210212422	0.048090535
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PHF3	1.209128468	0.038201547
ZNF445	1.20908181	0.027351121
POGZ	1.208803487	0.0260344
TXLNG	1.208647518	0.025084018
PHF8	1.208624154	0.038019273
USP8	1.207952525	0.028700037
MICAL3	1.2078995	0.0208074
KDM3A	1.207739325	0.008481259
CDK19	1.207625529	0.038605702
AKAP10	1.207239516	0.016830547
CDK13	1.207186304	0.011758399
TRIM23	1.207090969	0.007718681
SP1	1.207018792	0.033767591
SRCIN1	1.20673612	0.038396482
FRMD4A	1.206534742	0.02036697
ZC3H11A	1.206455975	0.017500413
SETD7	1.206286751	0.026349829
LRRC58	1.206000724	0.028267614
IREB2	1.205877925	0.013528136
PRKCE	1.205503956	0.041845896
CSNK1G3	1.205256761	0.039208454
VPS37A	1.205106812	0.045399573
RAB33B	1.204945429	0.041907979
CENPC1	1.204849229	0.045690716
ZNF540	1.204803861	0.044941921
PNMAL1	1.204486923	0.031296614
BTBD3	1.204290233	0.0197776
IBTK	1.204265539	0.010478752
RPRD2	1.204062218	0.018642178

MLL5	1.203969696	0.038370049
MECP2	1.203680846	0.030585393
PRPF4B	1.203668084	0.028600617
NUPL1	1.203275152	0.016222844
SLC31A1	1.20317689	0.028895225
TTC14	1.201950767	0.025671159
ARHGAP21	1.201910619	0.02473979
RALGAPB	1.201838068	0.02600648
PRR14L	1.201336437	0.02956126
DNM3	1.225351642	0.022120754
VLDLR	1.225297732	0.015262895
ZEB1	1.225294435	0.018569445
CREBBP	1.225211739	0.024843798
KAT6B	1.225073597	0.023671884
CLIP4	1.224556671	0.025299478
SEC24A	1.224519944	0.037152417
TJP1	1.224463676	0.026579975
SPRY3	1.224135697	0.0372105
UBR5	1.224045884	0.040633654
RAP1GAP2	1.223999496	0.040386875
ZNF655	1.223795495	0.023809367
PPP1R2	1.223711086	0.009494836
GABRG2	1.223688216	0.038583507
YLPM1	1.222272414	0.01509225
NR3C1	1.22204784	0.038576747
PRRC2B	1.22157409	0.015082084
FAM178A	1.22125426	0.013057252
SPECC1	1.221199196	0.042993582
CDK12	1.220972135	0.043045425
SLC3A1	1.220421452	0.047986903
CHIC1	1.220190388	0.00499119
CHAMP1	1.220050466	0.015056659
NPHP3	1.219901081	0.040806463
JPH1	1.21965894	0.032133256
PCF11	1.219575472	0.022294975
CEP104	1.219441101	0.022120754
GABRB2	1.219412595	0.034765061
MAPK10	1.218253966	0.02061029
ANK3	1.218188728	0.008738552
KLHDC5	1.218067465	0.015160482

FAM76B	1.217661087	0.048937224
UPRT	1.217410939	0.045712194
WAPAL	1.217187164	0.012918559
PKN2	1.217066693	0.029842595
UBR1	1.217025927	0.034310397
SPAG9	1.216894873	0.043312737
ZFYVE26	1.216832306	0.020384098
MLLT3	1.216735075	0.030033624
ZFP91	1.216530517	0.005143666
UHMK1	1.216337346	0.029909873
TAF1	1.216241611	0.02473979
ATF6	1.215913209	0.032496254
OCRL	1.215804007	0.024295157
USP32	1.215579413	0.042112092
CCDC66	1.215546949	0.021510168
GRIA2	1.215396164	0.037880394
EPG5	1.21510216	0.040069905
CHD2	1.214744098	0.016830547
SCN8A	1.214244469	0.031975336
TFDP2	1.21414274	0.022120754
ATP6V1A	1.213964065	0.036385322
CACNB4	1.213901198	0.043335822
GPRASP1	1.213820312	0.040806463
ZNF398	1.213746115	0.025084018
PCYT1B	1.213743448	0.019816174
ANKFY1	1.213718728	0.012840137
ATR	1.213697548	0.028911221
FAM115A	1.213351499	0.039562163
ZNF644	1.213230589	0.020986884
CCSER2	1.213181575	0.02038572
CIT	1.212896157	0.007697271
LATS1	1.212610583	0.024427672
CLIP1	1.212497047	0.021000091
TSNAX	1.212370117	0.039616959
ATIC	1.212309523	0.032051314
TRIP12	1.212273811	0.018642178
RBM25	1.212243175	0.035618859
SPATS2	1.211355402	0.023034374
CPSF6	1.210924564	0.028956163
LBR	1.210274323	0.027748588

GALC	1.210212422	0.048090535
RLIM	1.209983505	0.022344274
FBXO28	1.209743639	0.005528176
LPIN2	1.209633739	0.001705164
IDS	1.209200018	0.028914709
PHF3	1.209128468	0.038201547
ZNF445	1.20908181	0.027351121
POGZ	1.208803487	0.0260344
TXLNG	1.208647518	0.025084018
PHF8	1.208624154	0.038019273
USP8	1.207952525	0.028700037
MICAL3	1.2078995	0.0208074
KDM3A	1.207739325	0.008481259
CDK19	1.207625529	0.038605702
AKAP10	1.207239516	0.016830547
CDK13	1.207186304	0.011758399
TRIM23	1.207090969	0.007718681
SP1	1.207018792	0.033767591
SRCIN1	1.20673612	0.038396482
FRMD4A	1.206534742	0.02036697
ZC3H11A	1.206455975	0.017500413
SETD7	1.206286751	0.026349829
LRRC58	1.206000724	0.028267614
IREB2	1.205877925	0.013528136
PRKCE	1.205503956	0.041845896
CSNK1G3	1.205256761	0.039208454
VPS37A	1.205106812	0.045399573
RAB33B	1.204945429	0.041907979
CENPC1	1.204849229	0.045690716
ZNF540	1.204803861	0.044941921
PNMAL1	1.204486923	0.031296614
BTBD3	1.204290233	0.0197776
IBTK	1.204265539	0.010478752
RPRD2	1.204062218	0.018642178
MLL5	1.203969696	0.038370049
MECP2	1.203680846	0.030585393
PRPF4B	1.203668084	0.028600617
NUPL1	1.203275152	0.016222844
SLC31A1	1.20317689	0.028895225
TTC14	1.201950767	0.025671159

ARHGAP21	1.201910619	0.02473979
RALGAPB	1.201838068	0.02600648
PRR14L	1.201336437	0.02956126
GPRASP1	1.213820312	0.040806463
ZNF398	1.213746115	0.025084018
PCYT1B	1.213743448	0.019816174
ANKFY1	1.213718728	0.012840137
ATR	1.213697548	0.028911221
FAM115A	1.213351499	0.039562163
ZNF644	1.213230589	0.020986884
CCSER2	1.213181575	0.02038572
CIT	1.212896157	0.007697271
LATS1	1.212610583	0.024427672
CLIP1	1.212497047	0.021000091
TSNAX	1.212370117	0.039616959
ATIC	1.212309523	0.032051314
TRIP12	1.212273811	0.018642178
RBM25	1.212243175	0.035618859
SPATS2	1.211355402	0.023034374
CPSF6	1.210924564	0.028956163
LBR	1.210274323	0.027748588
GALC	1.210212422	0.048090535
RLIM	1.209983505	0.022344274
FBXO28	1.209743639	0.005528176
LPIN2	1.209633739	0.001705164
IDS	1.209200018	0.028914709
PHF3	1.209128468	0.038201547
ZNF445	1.20908181	0.027351121
POGZ	1.208803487	0.0260344
TXLNG	1.208647518	0.025084018
PHF8	1.208624154	0.038019273
USP8	1.207952525	0.028700037
MICAL3	1.2078995	0.0208074
KDM3A	1.207739325	0.008481259
CDK19	1.207625529	0.038605702
AKAP10	1.207239516	0.016830547
CDK13	1.207186304	0.011758399
TRIM23	1.207090969	0.007718681
SP1	1.207018792	0.033767591
SRCIN1	1.20673612	0.038396482

FRMD4A	1.206534742	0.02036697
ZC3H11A	1.206455975	0.017500413
SETD7	1.206286751	0.026349829
LRRC58	1.206000724	0.028267614
IREB2	1.205877925	0.013528136
PRKCE	1.205503956	0.041845896
CSNK1G3	1.205256761	0.039208454
VPS37A	1.205106812	0.045399573
RAB33B	1.204945429	0.041907979
CENPC1	1.204849229	0.045690716
ZNF540	1.204803861	0.044941921
PNMAL1	1.204486923	0.031296614
BTBD3	1.204290233	0.0197776
IBTK	1.204265539	0.010478752
RPRD2	1.204062218	0.018642178
MLL5	1.203969696	0.038370049
MECP2	1.203680846	0.030585393
PRPF4B	1.203668084	0.028600617
NUPL1	1.203275152	0.016222844
SLC31A1	1.20317689	0.028895225
TTC14	1.201950767	0.025671159
ARHGAP21	1.201910619	0.02473979
RALGAPB	1.201838068	0.02600648
PRR14L	1.201336437	0.02956126
GPRASP1	1.213820312	0.040806463
ZNF398	1.213746115	0.025084018
PCYT1B	1.213743448	0.019816174
ANKFY1	1.213718728	0.012840137
ATR	1.213697548	0.028911221
FAM115A	1.213351499	0.039562163
ZNF644	1.213230589	0.020986884
CCSER2	1.213181575	0.02038572
CIT	1.212896157	0.007697271
LATS1	1.212610583	0.024427672
CLIP1	1.212497047	0.021000091
TSNAX	1.212370117	0.039616959
ATIC	1.212309523	0.032051314
TRIP12	1.212273811	0.018642178
RBM25	1.212243175	0.035618859
SPATS2	1.211355402	0.023034374

CPSF6	1.210924564	0.028956163
LBR	1.210274323	0.027748588
GALC	1.210212422	0.048090535
RLIM	1.209983505	0.022344274
FBXO28	1.209743639	0.005528176
LPIN2	1.209633739	0.001705164
IDS	1.209200018	0.028914709
PHF3	1.209128468	0.038201547
ZNF445	1.20908181	0.027351121
POGZ	1.208803487	0.0260344
TXLNG	1.208647518	0.025084018
PHF8	1.208624154	0.038019273
USP8	1.207952525	0.028700037
MICAL3	1.2078995	0.0208074
KDM3A	1.207739325	0.008481259
CDK19	1.207625529	0.038605702
AKAP10	1.207239516	0.016830547
CDK13	1.207186304	0.011758399
TRIM23	1.207090969	0.007718681
SP1	1.207018792	0.033767591
SRCIN1	1.20673612	0.038396482
FRMD4A	1.206534742	0.02036697
ZC3H11A	1.206455975	0.017500413
SETD7	1.206286751	0.026349829
LRRC58	1.206000724	0.028267614
IREB2	1.205877925	0.013528136
PRKCE	1.205503956	0.041845896
CSNK1G3	1.205256761	0.039208454
VPS37A	1.205106812	0.045399573
RAB33B	1.204945429	0.041907979
CENPC1	1.204849229	0.045690716
ZNF540	1.204803861	0.044941921
PNMAL1	1.204486923	0.031296614
BTBD3	1.204290233	0.0197776
IBTK	1.204265539	0.010478752
RPRD2	1.204062218	0.018642178
MLL5	1.203969696	0.038370049
MECP2	1.203680846	0.030585393
PRPF4B	1.203668084	0.028600617
NUPL1	1.203275152	0.016222844

SLC31A1	1.20317689	0.028895225
TTC14	1.201950767	0.025671159
ARHGAP21	1.201910619	0.02473979
RALGAPB	1.201838068	0.02600648
PRR14L	1.201336437	0.02956126
GPRASP1	1.213820312	0.040806463
ZNF398	1.213746115	0.025084018
PCYT1B	1.213743448	0.019816174
ANKFY1	1.213718728	0.012840137
ATR	1.213697548	0.028911221
FAM115A	1.213351499	0.039562163
ZNF644	1.213230589	0.020986884
CCSER2	1.213181575	0.02038572
CIT	1.212896157	0.007697271
LATS1	1.212610583	0.024427672
CLIP1	1.212497047	0.021000091
TSNAX	1.212370117	0.039616959
ATIC	1.212309523	0.032051314
TRIP12	1.212273811	0.018642178
RBM25	1.212243175	0.035618859
SPATS2	1.211355402	0.023034374
CPSF6	1.210924564	0.028956163
LBR	1.210274323	0.027748588
GALC	1.210212422	0.048090535
RLIM	1.209983505	0.022344274
FBXO28	1.209743639	0.005528176
LPIN2	1.209633739	0.001705164
IDS	1.209200018	0.028914709
PHF3	1.209128468	0.038201547
ZNF445	1.20908181	0.027351121
POGZ	1.208803487	0.0260344
TXLNG	1.208647518	0.025084018
PHF8	1.208624154	0.038019273
USP8	1.207952525	0.028700037
MICAL3	1.2078995	0.0208074
KDM3A	1.207739325	0.008481259
CDK19	1.207625529	0.038605702
AKAP10	1.207239516	0.016830547
CDK13	1.207186304	0.011758399
TRIM23	1.207090969	0.007718681

SP1	1.207018792	0.033767591
SRCIN1	1.20673612	0.038396482
FRMD4A	1.206534742	0.02036697
ZC3H11A	1.206455975	0.017500413
SETD7	1.206286751	0.026349829
LRRC58	1.206000724	0.028267614
IREB2	1.205877925	0.013528136
PRKCE	1.205503956	0.041845896
CSNK1G3	1.205256761	0.039208454
VPS37A	1.205106812	0.045399573
RAB33B	1.204945429	0.041907979
CENPC1	1.204849229	0.045690716
ZNF540	1.204803861	0.044941921
PNMAL1	1.204486923	0.031296614
BTBD3	1.204290233	0.0197776
IBTK	1.204265539	0.010478752
RPRD2	1.204062218	0.018642178
MLL5	1.203969696	0.038370049
MECP2	1.203680846	0.030585393
PRPF4B	1.203668084	0.028600617
NUPL1	1.203275152	0.016222844
SLC31A1	1.20317689	0.028895225
TTC14	1.201950767	0.025671159
ARHGAP21	1.201910619	0.02473979
RALGAPB	1.201838068	0.02600648
PRR14L	1.201336437	0.02956126

Gene ID	Down-Regulated Genes Fold Change	Adjusted P-value
CST3	-1.579323303	0.000332706
PHLDB3	-1.552853957	0.00579249
MZT2B	-1.536438182	0.001008709
EGR2	-1.529422034	0.005528176
C7orf50	-1.512690694	0.001307559
ROMO1	-1.50989593	0.001686657
CFD	-1.503180345	0.007367696
SLC2A8	-1.496112221	0.001281582
PARD6A	-1.485261304	0.000939807
REEP6	-1.480062655	0.003338795
RPL35	-1.47099459	0.001281582

PCSK1	-1.466128483	0.009858217
TMEM45A	-1.464570208	0.012464937
HSD17B6	-1.462273616	1.20504E-07
ORAI1	-1.459694013	0.00648968
ASGR1	-1.457283189	0.006820981
SHD	-1.456420438	0.012935763
MAMU-DRA	-1.451049191	0.013528136
C8orf82	-1.446412213	0.004815335
FAM83D	-1.442107076	0.011392688
PTRHD1	-1.44089265	0.00716266
EXOSC6	-1.440328555	0.009674847
POLR2L	-1.436581348	0.004705909
CCDC85B	-1.433238859	0.005528176
RPS19BP1	-1.430779623	0.006600468
TRAPPC5	-1.430131101	0.003170335
TMED1	-1.428991266	0.001987043
VMO1	-1.427190413	0.011498263
HRAS	-1.424488778	0.004815335
HSPE1	-1.424430657	0.000283858
CRADD	-1.419969736	0.001705164
CBR3	-1.418657213	0.011904999
FXYD1	-1.414282618	0.008544808
MYEOV2	-1.413767577	0.016009534
TMEM256	-1.413102363	0.012810041
DOHH	-1.412040003	0.002318461
C1QL1	-1.409538365	0.005379835
C16orf13	-1.406931128	0.005071953
IQCA1	-1.405564396	0.020920438
LAMTOR4	-1.405378349	0.004610624
TMEM141	-1.404633968	0.005764462
DCXR	-1.403557606	0.004790873
PARD6G	-1.403359235	0.017594157
CBY3	-1.402624224	0.017697646
RPS2	-1.402062675	0.00499119
SCAND1	-1.397957936	0.004790873
APRT	-1.395008228	0.006371191
MACROD1	-1.394788601	0.011068238
ABT1	-1.394319266	0.005286173
PRADC1	-1.393632666	0.004705909
ARL2	-1.392393176	0.001686657

MRPS12	-1.392238644	0.004815335
SDS	-1.391797274	0.025084018
TMEM219	-1.391014642	0.005864323
TXNRD2	-1.39087805	0.006274202
IFT43	-1.390248093	0.002318461
IGFLR1	-1.389094429	0.022120754
GRHPR	-1.389020585	0.002212115
TST	-1.388657376	0.010994693
AURKAIP1	-1.388290453	0.005101244
TMEM160	-1.388194475	0.007367696
POLR2F	-1.388120338	0.000283858
GDF1	-1.386445535	0.003960811
RPS27L	-1.385035365	0.013104295
CD74	-1.385025644	0.028956163
C1orf122	-1.383568005	0.004948341
CTXN1	-1.382610081	0.004790873
MPST	-1.382607405	0.008908212
JUNB	-1.382476411	0.030778793
ALKBH7	-1.382454944	0.006771791
ATP5D	-1.381858339	0.00579249
COX7A1	-1.380147454	0.011904999
RPS15	-1.379182908	0.007553939
C2orf80	-1.37787918	0.007388573
LYRM4	-1.377518956	0.009484345
SSNA1	-1.375908703	0.009484345
MAD2L2	-1.375863002	0.001705164
AEBP1	-1.375193871	0.026741357
PIIF	-1.374916592	0.011904999
RAMP1	-1.374723333	0.003338795
PRR7	-1.37469023	0.007374591
CENPM	-1.374517378	0.004927559
SIVA1	-1.3744774	0.006940669
NDUFB7	-1.374127911	0.004705909
SPATA2L	-1.373763894	0.004815335
EGR4	-1.373261599	0.028222741
FAM195A	-1.37313051	0.005864323
FKBP2	-1.373063801	0.006283193
GSTT1	-1.372635492	0.018601129
SIRT3	-1.372486196	0.002082672
SLC26A11	-1.372392625	0.022991663

JOSD2	-1.371965946	0.0154935
GP1BB	-1.371938325	0.017356917
PET100	-1.371063214	0.003451947
MRPL24	-1.370748079	0.007207603
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CHCHD5	-1.367040361	0.009120975
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TIMM13	-1.364066131	0.006283193
SMKR1	-1.363577522	0.004705909
LRRC24	-1.362692381	0.005864323
GRASP	-1.36188401	0.005143666
RND1	-1.361243826	0.00716266
GCK	-1.361040892	0.021643751
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PHPT1	-1.358548344	0.010368881
RGS10	-1.357214682	0.017500413
HSD17B10	-1.357039736	0.002919494
UBE2S	-1.356938052	0.004705909
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BRI3	-1.356442418	0.011290929
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COX4I1	-1.354220239	0.008738552
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C1QTNF4	-1.352939105	0.008065485
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MED16	-1.346033152	6.1474E-05
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ITPA	-1.343792489	0.010802779
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ENHO	-1.341518238	0.008738552
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UFSP1	-1.338807242	0.041335063
SLC7A10	-1.338295456	0.002212115
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C17orf62	-1.337461047	0.003321961
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MNF1	-1.337160159	0.007553939
TOMM22	-1.33691796	0.003338795
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DEXI	-1.336320753	0.009674447
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MANF	-1.336128383	0.005864323
SSR4	-1.335751498	0.007367696
MATK	-1.335322901	0.008431285
COX6B1	-1.335168877	0.012661285
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DUT	-1.334928621	0.004497921
C20orf27	-1.334762554	0.008738552
SNRPC	-1.334716701	0.048615487
TIMM17B	-1.334535578	0.01238385
COMMD4	-1.333937875	0.00933751
SPA17	-1.333911043	0.036110367
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C19orf10	-1.333294102	0.001705164
COX8A	-1.332263862	0.0104728
PPDPF	-1.332089907	0.010320849
TYROBP	-1.332019071	0.019442042
PCDH8	-1.331829094	0.025566529
GUK1	-1.331685182	0.009645686
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MYL6	-1.331375046	0.00408181
BAD	-1.331208351	0.013722216
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RELB	-1.330574974	0.029926471
PRR5	-1.330154095	0.028762545
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HSPBP1	-1.326751058	0.005528176
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DUSP23	-1.323843099	0.018031969
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B9D2	-1.323731894	0.02037913
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PCYT2	-1.318246892	0.001705164
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C9orf116	-1.317414464	0.045884167
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ANAPC11	-1.31549719	0.012918559
SERP2	-1.315466963	0.009120975
PSMC3	-1.315115388	0.012145374
EXOSC7	-1.315096787	0.02157716
PHB	-1.314949519	0.007642703
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SF3B5	-1.313369269	0.014234648
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GAL3ST3	-1.303028312	0.018845137
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DMAPI1	-1.290448776	0.008544808
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LAMTOR5	-1.282779624	0.011711084
CHCHD10	-1.282760721	0.024843798
MINOS1	-1.282535359	0.019560867
ALG14	-1.282451661	0.007697271
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COPE	-1.281729576	0.015996482
C12orf57	-1.281688067	0.035619775
YDJC	-1.281650239	0.020004919
PRKCDBP	-1.28143087	0.044490343
DNAJB11	-1.281398515	0.013528136
ACTB	-1.280871582	0.006685562
CYSTM1	-1.280854956	0.027451534
C19orf60	-1.280791967	0.047971764
ATP6V1G2	-1.280780782	0.005581278
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C14orf2	-1.280372395	0.027681777
ZNF205	-1.280334077	0.024172564
CTDNEP1	-1.280235322	0.004815335
GNPTG	-1.279818939	0.019244817
NDUFA4	-1.279497084	0.023787082
LRPAP1	-1.279486394	0.049307138
PFN1	-1.279422937	0.006940669
PDZD4	-1.279332791	0.008929516
ATOX1	-1.279185657	0.026690313
C12orf10	-1.278955364	0.022787837
RPL18	-1.278862757	0.04408273
COMT	-1.278748146	0.004705909
ATP6V0B	-1.278367763	0.006951912
CD63	-1.278354636	0.0197776
NDUFS5	-1.278349786	0.015160482
MED11	-1.278104962	0.011775813
SAT2	-1.278023573	0.038908626
ARMC5	-1.277630736	0.024642063
KCTD13	-1.277353989	0.008926783
PLGRKT	-1.277273771	0.033247749
RPUSD2	-1.277220242	0.026709583
EIF1AD	-1.277161123	0.024718761
LAMTOR2	-1.277096717	0.009941367
C21orf33	-1.27704303	0.021300163
RPL17	-1.27703211	0.034765061
CDKN2D	-1.276896701	0.016830547

DOLK	-1.276680946	0.012881891
TOE1	-1.276595947	0.023460099
TMEM186	-1.276571905	0.008908212
COQ3	-1.276186085	0.03660959
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C11orf68	-1.275938348	0.019776058
ZNF414	-1.275678221	0.019179881
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DBNL	-1.275576733	0.001307559
ATP5J	-1.275557644	0.012004181
BAG1	-1.275459016	0.017571215
MFSD12	-1.275297011	0.024284592
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FLYWCH2	-1.248096227	0.035445269

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PRAF2	-1.209640205	0.0108343
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CSPG5	-1.208296771	0.011498263
VDAC2	-1.208255575	0.017790469
TUBA1A	-1.208196614	0.04786864
PRMT1	-1.208066693	0.029874027
PTOV1	-1.208000952	0.029598147
HABP4	-1.207902126	0.042255696
DRG1	-1.207899147	0.025090749
LYPLA2	-1.207820942	0.011498263
CDK2AP1	-1.207772196	0.028764432
CCDC115	-1.207750637	0.04953422
RAD23A	-1.207742279	0.029653885
DCTN3	-1.207447354	0.043971583
CYP46A1	-1.206990611	0.037943714
DNAJC19	-1.206479866	0.009858217
ALG3	-1.206079898	0.031296614
C1orf85	-1.205926061	0.042404935
AUP1	-1.205570143	0.016970415
EIF4A1	-1.205424727	0.018935808
PEX14	-1.205368184	0.028700037

ARFRP1	-1.205059577	0.031827418
APEH	-1.204896054	0.005528176
POLDIP2	-1.204224984	0.023088572
GNB2L1	-1.204165464	0.040810595
SLC25A11	-1.203902195	0.030572645
EGLN2	-1.203880931	0.023818221
CFL1	-1.20383833	0.02600648
PC	-1.203824702	0.020608997
KIAA2013	-1.2038115	0.025090749
UBE2M	-1.203805776	0.029722165
FN3K	-1.203664172	0.042970021
ARFIP2	-1.203319233	0.025145719
LRRC4B	-1.203252389	0.019573236
EAPP	-1.203084915	0.033129238
BRMS1	-1.203021367	0.039000999
SENP3	-1.202978662	0.008302472
C14orf166	-1.202850834	0.018876175
TUSC2	-1.202678488	0.034390387
C9orf69	-1.202633557	0.025084018
CD3EAP	-1.202433783	0.032133256
SDF4	-1.202420489	0.030763804
RBM38	-1.201821326	0.02888021
BCAS2	-1.201033049	0.024408017
PEBP1	-1.200967661	0.028267614
RPL4	-1.200159071	0.034446707

Table S2. Gene transcripts in the caudate significantly up- and down-regulated in subjects administered IFN α (n=7) compared to controls (n=4).

Gene	Up-Regulated Genes Fold Change	Adjusted P-value
VSNL1	1.817348661	0.000109002
ALS2CR12	1.752978942	0.000152217
DCDC5	1.740066295	0.000126565
CCDC37	1.717120587	5.86E-06
LRRC46	1.691770182	0.000109843
ST6GALNAC2	1.688414149	0.000215684
DNALI1	1.686386	0.000302366
WDR16	1.665673437	9.46E-05

DPYSL3	1.662207252	0.002013253
SPAG8	1.636663296	0.000678462
FAM227A	1.632907648	0.003652052
DNAH6	1.629293999	0.002266506
LRRC48	1.628440094	0.000208213
CCDC114	1.625264444	9.46E-05
ANKMY1	1.617215095	0.004956514
DNAH9	1.615783556	0.00015828
C9orf117	1.612700694	0.000507494
AGR3	1.610635062	0.000736153
CCDC108	1.60922459	0.000317384
VWA3B	1.609066082	0.00261747
RSPH4A	1.605543874	0.000402848
TCTEX1D1	1.604257683	0.001953592
OTX2	1.599134128	0.002503812
CERKL	1.595901578	0.000302366
IQCD	1.592433439	0.000109002
HYDIN	1.590743058	0.000280333
DLEC1	1.590365788	9.46E-05
WDR63	1.58745926	2.18E-05
FXYD6	1.586216135	0.005991948
TLL9	1.584680336	0.006821024
LRRC23	1.583977391	0.001373221
CCDC80	1.580500299	0.003013556
C4A	1.578597828	0.000780953
TP73	1.576966973	0.007464268
CD24	1.576466835	0.000678462
DNAH1	1.572516691	0.000105693
NTS	1.570791617	0.000280333
KCNE1	1.568599641	0.000217715
PIFO	1.566899694	0.00387821
CCDC40	1.563471706	0.011617142
WDR52	1.558658568	0.000714425
ABI3BP	1.55490701	0.00075594
KCNJ16	1.554001503	0.014255665
FOXJ1	1.544237675	9.46E-05
MNS1	1.538566473	0.01673483
GABRQ	1.530195283	0.003298681
C20orf26	1.527426404	0.019279208
WDR96	1.526686212	0.02379065

CLHC1	1.526126284	0.020162027
C5orf49	1.521595221	0.001874562
RBM43	1.518747852	0.018886606
FHOD3	1.518258223	0.018787808
LRRC36	1.514630744	0.002265264
C9orf24	1.510544382	0.00015828
TGFBI	1.506396209	0.006272074
TMEM52	1.504121648	0.015668772
CAPS	1.500198131	0.026007145
ARHGAP6	1.499315816	0.032482838
SORD	1.498503081	0.018787808
KRT18	1.498128059	0.001723522
CCDC33	1.497774346	0.001122055
C22orf15	1.497611929	0.00015828
GJD3	1.497319804	0.032069939
SLC9A3	1.495593403	0.037705616
C6orf165	1.494219396	0.006803712
C7orf57	1.492845343	0.000736153
GLIS3	1.487463069	0.043296938
CCDC42B	1.487326303	0.039358868
CCDC19	1.485162812	0.000302366
DNAI1	1.484009292	0.026007145
CROCC	1.483721108	0.043296938
AK1	1.483251511	0.033099259
TTC18	1.481027628	0.047300226
MORN5	1.48080167	0.000280333
BTBD11	1.480517733	0.032532062
C20orf96	1.479251004	0.034531248
CD36	1.477987041	0.000507494
CELSR1	1.475397218	0.049979951
EFCAB1	1.475121591	0.002995558
ZCCHC12	1.474884616	0.02720269
HPCAL1	1.474134007	0.03676603
FAM83H	1.47273554	0.039621889
IL5RA	1.470641114	0.000105993
PI15	1.468597589	0.00015828
IQGAP2	1.467987777	0.017858261
GLI1	1.464180388	0.034973098
C2orf62	1.464095447	0.000287624
WDR65	1.462716739	0.004401297

RGS6	1.462378479	0.049655786
UBXN10	1.462112703	0.02862895
ZNF16	1.460983129	0.000507494
DZIP1L	1.459412153	0.002013253
DNAI2	1.457367698	0.012139025
TTC16	1.456589492	0.02720269
DNAH2	1.455936659	0.002333771
SLC6A7	1.450780576	0.048569724
TNFRSF11B	1.44940426	0.000714425
DCDC2	1.447315306	0.02720269
GMNC	1.443504304	0.036122479
GRAMD2	1.442703758	0.022724373
MAPK15	1.442265595	0.000217715
DTHD1	1.44160173	0.010189024
SPATA18	1.437976464	0.000678462
C11orf88	1.436669305	0.000620043
SLC28A1	1.435901678	0.007561131
WEE1	1.428876143	0.039667143
PNMA5	1.426731821	0.001122055
SLC47A2	1.419267371	0.00015828
SPAG6	1.41899184	0.015817686
CCDC81	1.415620271	0.006323771
TRPC4	1.414681579	0.0293486
RORC	1.412342785	0.011561484
TYMP	1.411567374	0.007272768
FAM92B	1.410285663	0.000881796
FAM81B	1.407121351	0.000280333
TMEM61	1.400253372	0.017306195
RSPH10B	1.39953624	0.039358868
IRX3	1.393089418	0.02720269
PLOD2	1.381606657	0.024894528
KNCN	1.376869478	0.000678462
C11orf70	1.376763442	0.013759611
ARMC3	1.376037694	0.000678462
VSIG8	1.374075957	0.02720269
NPY2R	1.37034313	0.039494564
UNC5CL	1.370342381	0.001535813
ZNF608	1.367507122	0.041495694
SMAD9	1.364200777	0.048569724
C2orf61	1.361164171	0.013299518

CLEC3A	1.360801264	0.010596677
L3MBTL3	1.356635583	0.021835843
C9orf135	1.354722057	0.00801321
ZBBX	1.349017395	0.002333771
FAM9B	1.346760557	0.041495694
ADRA2B	1.343829021	0.032958188
VWA5B1	1.342948132	0.004401297
GUCA1A	1.336789835	0.005579948
MORN3	1.328191353	0.023289956
CP	1.327784502	0.018936725
LDLRAD2	1.321612836	0.008124339
BARX2	1.321000756	0.017939739
SPATA17	1.31124825	0.039358868
ODF3B	1.310720178	0.008115443
FNDC1	1.308431115	0.044690543
HDC	1.307845163	0.009146593
TRPA1	1.298660736	0.004401297
SHROOM2	1.295118025	0.045493982
ANKRD10	1.284604386	0.036122479
SLC27A2	1.280415646	0.015817686
ADCYAP1	1.278402992	0.037705616
GSTA3	1.278212127	0.005103224
RASSF9	1.276080976	0.012769485
FHAD1	1.274464186	0.005674735
FAM183A	1.273327427	0.003471947
ATIC	1.272768359	0.036122479
LY6G6F	1.254058324	0.012119471
LY6G6D	1.252532846	0.049655786
RIBC2	1.251614863	0.014493287
CCDC60	1.243278493	0.011562918
C1orf158	1.241029566	0.008711314
PIH1D3	1.237076824	0.011429917
FAM47E	1.227523262	0.036122479
WDR38	1.213168236	0.023289956
ZNF474	1.206665454	0.035259397
CAPSL	1.204273277	0.02720269

Down-Regulated Genes

Gene ID	Fold change	Adjusted P-value
MYBPH	-1.530821916	0.021835843
EGR4	-1.490780184	0.02720269
CD93	-1.453695425	0.046759403
HTR6	-1.422228522	0.020305711
FOSL2	-1.414460501	0.046200674
REEP6	-1.391601221	0.038254861
MPC2	-1.353338283	0.000604138
IFRD1	-1.327556162	0.02720269
HAS3	-1.315952335	0.019256589
ATP6V1G2	-1.286008899	0.012119471
NLE1	-1.285049947	0.021835843
CAMKK2	-1.273395448	0.019943908
PNO1	-1.271528021	0.01278501
PRKAG2	-1.235356449	0.009146593
PEX5	-1.23277581	0.007631107
MYBPH	-1.530821916	0.021835843
EGR4	-1.490780184	0.02720269
CD93	-1.453695425	0.046759403
HTR6	-1.422228522	0.020305711
FOSL2	-1.414460501	0.046200674
REEP6	-1.391601221	0.038254861
MPC2	-1.353338283	0.000604138
IFRD1	-1.327556162	0.02720269
HAS3	-1.315952335	0.019256589
ATP6V1G2	-1.286008899	0.012119471
NLE1	-1.285049947	0.021835843
CAMKK2	-1.273395448	0.019943908
PNO1	-1.271528021	0.01278501
PRKAG2	-1.235356449	0.009146593
PEX5	-1.23277581	0.007631107