

Supplementary Table 1. Primers used for RT-qPCR analysis of striatal and nigral tissue.

Gene	Forward Primer (5'-3')	Reverse Primer (5'-3')
Dopaminergic Markers		
TH	CTG GCC ATT GAT GTA CTG GA	ACA CAC ATG GGA AAG CCT CT
DAT	TGA ACT TTG GCC TAG CCT GT	ACA TGG AGC CAC TCA GCT CT
Cellular Markers		
GFAP	CGA GTC CCT AGA GCG GCA AAT G	CGG ATC TGG AGG TTG GAG AAA GTC
MAC-1	TTC TCA TGG TCA CCT CCT GC	GGT CTG ACC ATC TGA ACC TG
Inflammatory Enzymes		
gp91 phox	CAG GAG TTC CAA GAT GCC TG	GAT TGG CCT GAG ATT CAT CC
p67phox	CAG CCA GCT TCG GAA CAT G	GAC AGT ACC AGG ATT ACA TC
Cytokines		
NFkB	GCG TAC ACA TTC TGG GGA GT	ACC GAA GCA GGA GCT ATC AA
Antioxidant Markers		
NQO1	GCG AGA AGA GCC CTG ATT GTA CTG	TCT CAA ACC AGC CTT TCA GAA TGG
HO1	CAA GCC GAG AAT GCT GAG TTC ATG	GCA AGG GAT GAT TTC CTG CCA G
GCLM	GCC ACC AGA TTT GAC TGC CTT TG	TGC TCT TCA CGA TGA CCG AGT ACC
GCLC	ACA TCT ACC ACGC AGT CAA GG ACC	CTC AAG AAC ATC GCC TCC ATT CAG
Peroxiredoxin 3	CAC CCA GCA CGC GCC CTA TT	GGC CCA CCG GAA GGT CGT TG
Peroxiredoxin 5	GTG TCT CTC TTT GGG AAT CGT C	CTC AGA GTT GAG AGA GGA TGT TGG
SOD2	GAC TGG CAG CAT TTA GGG TT	GGA CCC AAT GAG CCA AAG AA
Thioredoxin 2	TCG CCA AGC AGC ACG GGA AG	TGA CCC CCA CAC CAG TGG CA
Protein Kinases		
AMPK	GGG GAC ACG CTT GGT GTC GG	GCT GGA ACA GAC GGC GGC TT
MAPK	CCT CAA GCC TGA AAA TCT GC	GTC TCT GCG CTT CCA GTA CC
PPAR-γ markers		
PPAR-γ	CCA GAG TCT GCT GAT CTG CG	GCC ACC TCT TTG CTC TGA TC
PGC1-α	TGA TGT GAA TGA CTT GGC TAC AGA CA	GCT CAT TGT TGT ACT GGT TGG ATA TG
Housekeeping gene		
Beta-actin	CAT GAA GAT CCT GAC CGA GCG TG	TCT GCT GGA AGG TGG ACA GTG AGG

Supplementary Table 2. Effect of LSN862 (LSN) on striatal mRNA expression. Striatal tissue was collected at 24 hrs, 72 hrs, and 21 days after the last dose of MPTP and 24 hrs after LSN862. The groups were vehicle + PBS (24 hrs: n = 5; 72 hrs: n = 4; 21 days: n = 4); vehicle + MPTP (24 hrs: n = 5; 72 hrs: n = 5; 21 days: n = 6); 30 mg/kg LSN + PBS (24 hrs: n = 4; 72 hrs: n = 3; 21 days n = 4); 30 mg/kg LSN + MPTP (24 hrs: n = 5; 72 hrs n = 4; 21 days n = 5). All data are standardized to β -actin presented as mean \pm SEM compared to vehicle + PBS controls. Two-Way ANOVA with Repeated Measures: ^aVehicle + PBS vs. Vehicle + MPTP, $P < 0.05$; ^bVehicle + MPTP vs. 30mg/kg LSN + MPTP, $P < 0.05$; ^cVehicle + PBS vs. 30mg/kg LSN + PBS, $P < 0.05$.

Gene	Vehicle + PBS			Vehicle + MPTP			30 mg/kg LSN + PBS			30 mg/kg LSN + MPTP		
	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days
Dopaminergic markers												
TH	1.00 \pm 0.34	1.00 \pm 0.58	1.00 \pm 0.09	0.37 \pm 0.19	0.09 \pm 0.06	0.39 \pm 0.08	0.98 \pm 0.57	0.98 \pm 0.57	1.30 \pm 0.36	0.20 \pm 0.06	0.52 \pm 0.01	1.15 \pm 0.51
DAT	1.00 \pm 0.49	1.00 \pm 0.65	1.00 \pm 0.39	0.49 \pm 0.21	0.004 \pm 0.003	0.27 \pm 0.10	0.97 \pm 0.36	0.88 \pm 0.01	1.41 \pm 0.73	0.45 \pm 0.15	0.30 \pm 0.11	1.20 \pm 0.59
Cellular Markers												
GFAP	1.00 \pm 0.40	1.00 \pm 0.35	1.00 \pm 0.21	7.38 \pm 0.98^{ab}	2.49 \pm 1.16^a	0.95 \pm 0.26	0.43 \pm 0.03	0.71 \pm 0.67	0.45 \pm 0.15	0.92 \pm 0.22^b	0.77 \pm 0.46	0.51 \pm 0.14
MAC-1	1.00 \pm 0.42	1.00 \pm 0.42	1.00 \pm 0.17	4.13 \pm 1.26^{ab}	4.92 \pm 2.83	0.83 \pm 0.23	0.85 \pm 0.21	0.70 \pm 0.04	0.94 \pm 0.35	1.55 \pm 0.56^b	0.23 \pm 0.18	0.99 \pm 0.15
Inflammatory Enzymes												
gp91phox	1.00 \pm 0.58	1.00 \pm 0.48	1.00 \pm 0.44	2.5 \pm 0.39^{ab}	0.31 \pm 0.17	0.10 \pm 0.03	0.95 \pm 0.62	0.10 \pm 0.08	0.07 \pm 0.01	1.14 \pm 0.38^b	0.15 \pm 0.08	0.10 \pm 0.03
p67phox	1.00 \pm 0.37	1.00 \pm 0.32	1.00 \pm 0.36	1.40 \pm 0.58	0.02 \pm 0.01	37.48 \pm 11.47^{ab}	0.86 \pm 0.36	0.05 \pm 0.003	2.46 \pm 0.47	0.02 \pm 0.01	0.47 \pm 0.44	2.74 \pm 1.31^b
NFkB	1.00 \pm 0.16	1.00 \pm 0.59	1.00 \pm 0.15	5.80 \pm 3.45^a	0.03 \pm 0.02	0.18 \pm 0.04	2.02 \pm 1.33	0.71 \pm 0.25	0.19 \pm 0.05	3.43 \pm 1.01	0.33 \pm 0.16	0.14 \pm 0.04
Antioxidant Markers												
NQO1	1.00 \pm 0.37	1.00 \pm 0.43	1.00 \pm 0.18	5.00 \pm 1.94^{ab}	0.51 \pm 0.25	0.12 \pm 0.02	0.91 \pm 0.27	0.85 \pm 0.79	0.34 \pm 0.12	2.31 \pm 0.72^b	1.46 \pm 0.63	0.11 \pm 0.02
HO-1	1.00 \pm 0.11	1.00 \pm 0.49	1.00 \pm 0.48	2.27 \pm 0.20	0.18 \pm 0.09	2.57 \pm 0.84	1.11 \pm 0.25	0.61 \pm 0.43	1.81 \pm 1.49	0.92 \pm 0.35	1.23 \pm 0.44	2.69 \pm 0.92
GCLM	1.00 \pm 0.34	1.00 \pm 0.45	1.00 \pm 0.46	1.55 \pm 0.46	5.80 \pm 2.82^a	1.02 \pm 0.31	0.56 \pm 0.21	1.32 \pm 0.66	1.16 \pm 0.32	0.96 \pm 0.30	4.07 \pm 1.79	1.24 \pm 0.40
GCLC	1.00 \pm 0.38	1.00 \pm 0.41	1.00 \pm 0.09	3.75 \pm 1.23	12.92 \pm 3.65^a	0.15 \pm 0.07	2.01 \pm 0.92	0.87 \pm 0.26	0.08 \pm 0.02	1.69 \pm 0.81	15.47 \pm 3.32	0.03 \pm 0.01
Peroxiredoxin 5	1.00 \pm 0.47	1.00 \pm 0.35	1.00 \pm 0.36	0.31 \pm 0.12	1.24 \pm 0.62	0.58 \pm 0.15	0.58 \pm 0.16	0.83 \pm 0.17	0.70 \pm 0.23	0.45 \pm 0.39	0.74 \pm 0.40	0.41 \pm 0.08
SOD2	1.00 \pm 0.51	1.00 \pm 0.17	1.00 \pm 0.32	0.45 \pm 0.14	0.05 \pm 0.02	0.44 \pm 0.14	0.16 \pm 0.05	1.34 \pm 0.91	0.28 \pm 0.06	0.88 \pm 0.38	0.51 \pm 0.20	0.35 \pm 0.09
Thioredoxin 2	1.00 \pm 0.54	1.00 \pm 0.51	1.00 \pm 0.40	2.55 \pm 0.56	0.78 \pm 0.30	0.62 \pm 0.21	0.72 \pm 0.21	0.06 \pm 0.01	0.51 \pm 0.18	1.14 \pm 0.49	0.42 \pm 0.41	0.22 \pm 0.05
Protein Kinases												
AMPK	1.00 \pm 0.16	1.00 \pm 0.47	1.00 \pm 0.36	3.41 \pm 0.56^{ab}	0.68 \pm 0.20^b	0.06 \pm 0.02	2.79 \pm 1.03^c	1.18 \pm 0.69	0.62 \pm 0.27	1.82 \pm 0.66^b	3.73 \pm 0.85^b	0.13 \pm 0.03
MAPK	1.00 \pm 0.58	1.00 \pm 0.08	1.00 \pm 0.35	0.05 \pm 0.02	0.03 \pm 0.01	0.41 \pm 0.09	13.31 \pm 8.74^c	1.69 \pm 1.19	5.36 \pm 4.92	4.34 \pm 2.64	3.15 \pm 0.90	4.93 \pm 2.23
PPAR-γ markers												
PPAR- γ	1.00 \pm 0.35	1.00 \pm 0.17	1.00 \pm 0.31	2.04 \pm 0.25^{ab}	0.30 \pm 0.12^a	0.39 \pm 0.08	1.39 \pm 0.33	7.01 \pm 1.5^c	0.91 \pm 0.31	0.63 \pm 0.25^b	3.56 \pm 0.64	1.09 \pm 0.32
PGC1- α	1.00 \pm 0.42	1.00 \pm 0.57	1.00 \pm 0.49	22.65 \pm 0.57^{ab}	0.10 \pm 0.42	0.31 \pm 0.07	0.35 \pm 0.23	0.37 \pm 0.23	0.58 \pm 0.07	0.44 \pm 0.19^b	0.61 \pm 0.20	0.43 \pm 0.13

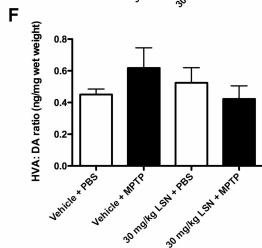
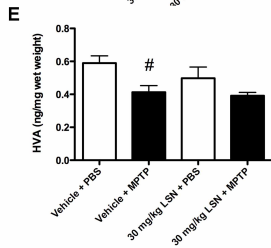
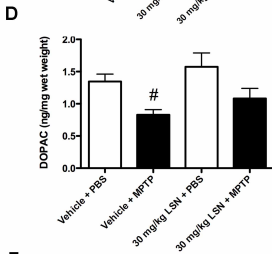
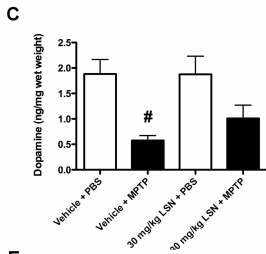
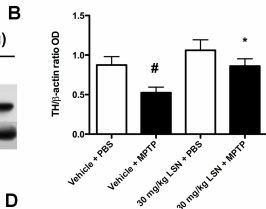
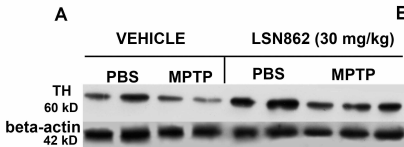
Supplementary Table 3. Effect of LSN862 (LSN) on nigral mRNA expression. Substantia nigra tissue was collected at 24 hrs, 72 hrs, and 21 days after the last dose of MPTP and 24 hrs after LSN862. The groups were vehicle + PBS (24 hrs: n = 5; 72 hrs: n = 4; 21 days: n = 4); vehicle + MPTP (24 hrs: n = 5; 72 hrs: n = 5; 21 days: n = 6); 30 mg/kg LSN + PBS (24 hrs: n = 4; 72 hrs: n = 3; 21 days: n = 4); 30 mg/kg LSN + MPTP (24 hrs: n = 5; 72 hrs: n = 4; 21 days: n = 5). All data are standardized to β -actin presented as mean \pm SEM compared to vehicle + PBS controls. Two-Way ANOVA with Repeated Measures: ^aVehicle + PBS vs. Vehicle + MPTP, $P < 0.05$; ^bVehicle + MPTP vs. 30mg/kg LSN + MPTP, $P < 0.05$; ^cVehicle + PBS vs. 30mg/kg LSN + PBS, $P < 0.05$.

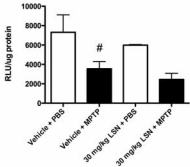
Gene	Vehicle + PBS			Vehicle + MPTP			30 mg/kg LSN + PBS			30 mg/kg LSN + MPTP		
	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days	24 hrs	72 hrs	21 days
Dopaminergic markers												
TH	1.00 \pm 0.53	1.00 \pm 0.35	1.00 \pm 0.45	0.06 \pm 0.04	0.02 \pm 0.01	0.17 \pm 0.04	1.27 \pm 0.75	0.86 \pm 0.33	1.33 \pm 0.53	0.58 \pm 0.32	0.15 \pm 0.04	0.47 \pm 0.34
DAT	1.00 \pm 0.66	1.00 \pm 0.62	1.00 \pm 0.41	0.06 \pm 0.05	0.62 \pm 0.26	0.12 \pm 0.04	1.45 \pm 0.77	1.15 \pm 0.32	1.33 \pm 0.74	0.18 \pm 0.09	0.58 \pm 0.22	0.32 \pm 0.05
Cellular Markers												
GFAP	1.00 \pm 0.29	1.00 \pm 0.47	1.00 \pm 0.40	262.7 \pm 76.92^{a,b}	3.84 \pm 1.39	1.39 \pm 0.65	32.0 \pm 6.96	0.04 \pm 0.02	0.90 \pm 0.72	18.96 \pm 10.82^b	0.15 \pm 0.09	1.58 \pm 0.63
MAC-1	1.00 \pm 0.35	1.00 \pm 0.33	1.00 \pm 0.10	5.16 \pm 2.43^{a,b}	3.24 \pm 0.46^{a,b}	2.32 \pm 0.72	1.47 \pm 0.62	0.03 \pm 0.01	1.34 \pm 0.32	1.38 \pm 0.91^b	0.80 \pm 0.53^b	2.06 \pm 0.80
Inflammatory Enzymes												
gp91phox	1.00 \pm 0.46	1.00 \pm 0.30	1.00 \pm 0.62	7.12 \pm 3.11^{a,b}	0.74 \pm 0.32	1.18 \pm 0.52	1.03 \pm 0.56	0.20 \pm 0.18	3.15 \pm 0.81	1.63 \pm 0.92^b	0.89 \pm 0.44	1.02 \pm 0.24
p67phox	1.00 \pm 0.32	1.00 \pm 0.29	N/A	4.47 \pm 1.99^b	2.75 \pm 1.20^{a,b}	N/A	0.70 \pm 0.39	0.30 \pm 0.12	N/A	0.09 \pm 0.03^b	0.94 \pm 0.79^b	N/A
NFkB	1.00 \pm 0.35	1.00 \pm 0.14	1.00 \pm 0.47	3.29 \pm 1.20	0.54 \pm 0.24	1.05 \pm 0.39	0.37 \pm 0.17	0.15 \pm 0.13	0.94 \pm 0.22	2.40 \pm 0.73	0.02 \pm 0.01	0.62 \pm 0.22
Antioxidant Markers												
NQO1	1.00 \pm 0.38	1.00 \pm 0.22	1.00 \pm 0.34	5.66 \pm 1.86^a	0.88 \pm 0.32	0.68 \pm 0.10	0.98 \pm 0.46	0.08 \pm 0.05	1.13 \pm 0.38	4.29 \pm 1.81	0.34 \pm 0.14	0.33 \pm 0.10
HO-1	1.00 \pm 0.24	1.00 \pm 0.10	1.00 \pm 0.62	1.01 \pm 0.20	0.35 \pm 0.13	1.63 \pm 0.47	0.51 \pm 0.12	0.67 \pm 0.25	8.35 \pm 4.99	2.47 \pm 1.15	0.38 \pm 0.06	0.82 \pm 0.29
GCLM	1.00 \pm 0.21	1.00 \pm 0.40	1.00 \pm 0.04	1.72 \pm 0.25	3.06 \pm 0.53^a	0.01 \pm 0.002	0.78 \pm 0.38	0.99 \pm 0.42	0.001 \pm 0.0005	1.43 \pm 0.50	2.47 \pm 0.76	0.003 \pm 0.0009
GCLC	1.00 \pm 0.28	1.00 \pm 0.30	1.00 \pm 0.07	3.42 \pm 0.96^a	3.09 \pm 1.10^a	0.06 \pm 0.004	0.27 \pm 0.16	0.31 \pm 0.14	0.02 \pm 0.01	1.94 \pm 0.85	6.11 \pm 1.85	0.02 \pm 0.006
Peroxioredoxin 5	1.00 \pm 0.29	1.00 \pm 0.36	1.00 \pm 0.19	2.23 \pm 0.61	0.80 \pm 0.19	0.96 \pm 0.26	6.31 \pm 3.76	0.24 \pm 0.21	0.72 \pm 0.06	3.24 \pm 1.31	0.64 \pm 0.45	0.89 \pm 0.31
SOD2	1.00 \pm 0.35	1.00 \pm 0.39	1.00 \pm 0.42	0.74 \pm 0.21	0.23 \pm 0.15	2.53 \pm 0.38^{a,b}	0.09 \pm 0.04	0.03 \pm 0.01	1.29 \pm 0.38	1.04 \pm 0.38	0.07 \pm 0.04	1.14 \pm 0.31^b
Thioredoxin 2	1.00 \pm 0.43	1.00 \pm 0.15	1.00 \pm 0.40	0.62 \pm 0.09	3.09 \pm 1.49	0.62 \pm 0.22	0.40 \pm 0.21	0.29 \pm 0.15	0.51 \pm 0.19	0.72 \pm 0.29	0.13 \pm 0.04	0.22 \pm 0.05
Protein Kinases												
AMPK	1.00 \pm 0.41	1.00 \pm 0.39	1.00 \pm 0.27	1.93 \pm 0.02	3.40 \pm 2.04	0.37 \pm 0.13	4.40 \pm 2.53	0.91 \pm 0.33	0.37 \pm 0.09	13.12 \pm 8.55	0.76 \pm 0.15	0.22 \pm 0.05
MAPK	1.00 \pm 0.27	1.00 \pm 0.66	1.00 \pm 0.19	0.04 \pm 0.01	0.17 \pm 0.08	1.01 \pm 0.18	0.51 \pm 0.15	2.98 \pm 1.81	0.62 \pm 0.05	2.67 \pm 1.39	1.92 \pm 0.76	0.37 \pm 0.05
PPAR-γ markers												
PPAR- γ	1.00 \pm 0.15	1.00 \pm 0.43	1.00 \pm 0.47	0.87 \pm 0.28	0.02 \pm 0.01	1.90 \pm 1.14	0.37 \pm 0.18	4.95 \pm 1.67	0.71 \pm 0.17	0.90 \pm 0.54	1.89 \pm 0.5	0.99 \pm 0.30
PGC1-a	1.00 \pm 0.14	1.00 \pm 0.33	1.00 \pm 0.25	7.25 \pm 4.12	0.19 \pm 0.06	0.57 \pm 0.12	0.79 \pm 0.23	0.12 \pm 0.06	0.85 \pm 0.22	2.93 \pm 1.32	0.11 \pm 0.01	0.43 \pm 0.15

Supplementary Figure Legends

Supplementary Figure 1. LSN862 confers neuroprotection against MPTP toxicity in ARE-hPAP mice. Striata from ARE-hPAP mice were harvested 7 days after the last dose of MPTP and 24 hrs after treatment with LSN. A) Western blot of TH protein levels. B) Densitometric analysis of TH western blot relative to β -actin (means \pm SEM). Groups: vehicle + PBS (n = 7), vehicle + MPTP (n = 7), 30 mg/kg LSN + PBS (n = 8), 30 mg/kg LSN + MPTP (n = 6). # $P = 0.029$ (vehicle + PBS vs. vehicle + MPTP), * $P = 0.05$, (vehicle + MPTP vs. 30 mg/kg LSN + MPTP). HPLC analysis of striatal catecholamine levels vehicle + PBS (n = 8), vehicle + MPTP (n = 9), 30 mg/kg LSN + PBS (n = 9), 30 mg/kg LSN + MPTP (n = 7), C) Dopamine: # $P = 0.003$ (vehicle + PBS vs. vehicle + MPTP). D) DOPAC: # $P = 0.022$ (vehicle + PBS vs. vehicle + MPTP). E) HVA: # $P = 0.014$ (vehicle + PBS vs. vehicle + MPTP), F) HVA:DA ratio.

Supplementary Figure 2. LSN862 does not affect striatal and nigral hPAP activity in ARE-hPAP mice. Groups: vehicle + PBS (n = 5), vehicle + MPTP (n = 5), 30 mg/kg LSN + PBS (n = 3), 30 mg/kg LSN + MPTP (n = 5). A) Striatal hPAP activity levels (means \pm SEM). # $P = 0.040$ (vehicle + PBS vs. vehicle + MPTP). B) Nigral hPAP activity levels; # $P = 0.031$ (vehicle + PBS vs. vehicle + MPTP).



A**Striatal hPAP activity****B****Nigral hPAP activity**