

# Supporting Information

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**Table S1.** Genes differentially expressed in WT and TNFRI KO nonoccluded brain tissue

Function	Accession no.	Fold change
Cell death		
Survival motor neuron	U77714	-8.66
BH3 interacting (with BCL2 family) domain, apoptosis agonist	D83698	5.96
Signaling		
v-abl Abelson murine leukemia oncogene 1	L10656	-6.93
Chemokine-like receptor 1	U79525	-6.48
Estrogen related receptor, $\alpha$	U85259	5.73
5-Hydroxytryptamine (serotonin) receptor 1B	Z11597	6.19
Androgen receptor	X53779	6.22
Cytoskeleton		
Sequestosome 1	U40930	-6.36
Cortactin	U03184	5.68
Actin, $\beta$ , cytoplasmic	M12481	11.69
Laminin, $\alpha$ 5	U37501	15.47
Ion transporters		
Potassium large conductance calcium-activated channel, subfamily M, $\beta$ member 1	AF020711	-12.06
Solute carrier family 16 (monocarboxylic acid transporters), member 1	AF058055	-6.14
ATPase, Ca $^{++}$ transporting, plasma membrane 2	AF053471	-6.07
Potassium voltage-gated channel, subfamily H (eag-related), member 1	U04294	-6.06
Solute carrier family 6 (neurotransmitter transporter, serotonin), member 4	AF013604	-5.45
ATP-binding cassette, sub-family G (WHITE), member 1	U34920	-5.22
Friedreich ataxia	U95736	5.79
Metabolism		
Prostaglandin I2 (prostacyclin) synthase	AB001607	-6.81
Tryptophan hydroxylase	J04758	-6.43
Fatty acid amide hydrolase	U82536	-5.84
Protein convertase subtilisin/kexin type 1	M58589	-5.47
Serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	M33960	5.16
Hypoxanthine guanine phosphoribosyl transferase	J00423	5.81
Plasminogen	J04766	6.82
Ubiquitin B	X51703	7.47
Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	D78647	7.64
Serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2	X16490	7.87
Stress response		
Heat shock protein, 84 kDa 1	M36829	-5.78
Transcription/translation		
Nuclear DNA-binding protein	AF031426	-16.77
NK2 transcription factor related, locus 2 (Drosophila)	U31566	-5.63
Iroquois related homeobox 3 (Drosophila)	Y15001	-5.42
fos-like antigen 2	X83971	-5.34
Ribosomal protein S29	L31609	12.42

This list of genes shows those with transcripts that were different by  $\geq 2$ -fold in TNFRI KO compared with WT. The genes have been grouped according to known functions (i.e., Entrez definition).

**Table S2. Genes differentially expressed in WT and TNFRI KO lesions 3 h after pMCAO**

Function	Accession no.	Fold change
<b>Signaling</b>		
Adenomatosis polyposis coli	M88127	-2.06
raf-related oncogene	M13071	-2.13
abl proto-oncogene	L10656	-2.18
DnaJ-like protein 1	L16953	-2.65
Multidrug resistance protein 1	M14757	-3.20
Serine/arginine-rich protein specific kinase 2	U92456	-2.64
Protein tyrosine kinase 6	U16805	-2.05
Protein kinase C inhibitor protein 1, zeta	D78647	-2.52
Protein kinase C inhibitor protein 1, eta	U57311	2.19
TRAF family member-associated Nf-kappa B activator	U51907	2.03
Presenilin 2	U57324	3.11
Tumor necrosis factor receptor superfamily, member 1b	M59378	2.24
<b>Transcription/translation</b>		
ELK1, member of ETS oncogene family	X87257	-3.14
fos-like antigen 2 (Fra-2)	X83971	-3.35
DNA polymerase delta 1, catalytic domain	Z21848	2.55
Thyroid autoantigen 70 kDa	M38700	3.56
CCCTC-binding factor	U51037	2.12
Est2 repressor factor	U58533	2.02
<b>Cell cycle</b>		
Cyclin-dependent kinase inhibitor 2C	U19596	-2.31
<b>Development/differentiation</b>		
Myeloblastosis oncogene-like 1	X82327	-2.58
Gli oncogene; zinc finger transcription factor	S65038	-2.09
Eph receptor B4	Z49085	-2.47
Eph receptor B2	L25890	-2.93
Transferrin receptor	X57349	2.07
Clusterin	L08235	2.40
<b>Growth factors</b>		
Thrombopoietin receptor	Z22649	-2.02
Angiotensin converting enzyme	J04946	-2.13
Insulin-like growth factor binding protein 1	X81579	-2.41
Neuroleukin	M14220	2.21
MAD homolog 2 (Drosophila) (SMAD 2)	U60530	2.39
<b>Cytoskeleton</b>		
Heat shock protein, 25 kDa	U03560	-2.09
Keratin complex 1, acidic, gene 18	M11686	3.12
<b>Metabolism</b>		
Calcium-binding protein, intestinal	J05186	-2.27
Glutathione S-transferase, mu 2	J04696	2.38
Glutathione S-transferase, theta 1	X98055	-2.20
Ubiquitin B	X51703	-2.18
Glutathione peroxidase 3	U13705	2.22
Glutathione S-transferase, $\alpha$ 2 (Yc2)	J03958	2.14
<b>Cell death</b>		
Neuronal death protein DP5	D83698	2.64
Bcl-associated death promoter (BAD)	L37296	2.10
Defender against cell death 1	U83628	3.03
Adenosine A1 receptor	U05671	3.29
Fas antigen	U39643	2.09
Programmed cell death 1	X67914	2.15
CASP8 and FADD-like apoptosis regulator	U97076	2.48
Adenosine A2a receptor	U05672	-3.19
BH3 interacting domain death agonist	U75506	-3.08
<b>DNA repair</b>		
Excision repair 1	X07414	2.33
DNA excision repair protein ERCC-5	D16306	3.30
RAD23b homolog (S. cerevisiae)	X92411	2.44
Translin	X81464	2.26

This is a list of genes with transcripts that were different by at least 2-fold between WT and TNFRI KO pMCAO. The genes have been grouped accordingly to known functions (i.e., Entrez definition).

**Table S3. Genes differentially expressed in WT and TNFRI KO lesions 6 h after pMCAO**

Function	Accession no.	Fold change
Signaling		
Adenomatosis polyposis coli	M88127	2.32
Lymphocyte protein tyrosine kinase	M12056	2.60
Harvey rat sarcoma virus oncogene	Z50013	2.60
Neurofibromatosis 2	L27105	3.70
Jun oncogene	J04115	3.30
Map3k8	D13759	6.06
Transcription factor RelB	M83380	3.00
GTP-binding protein	U10551	16.73
Promyelocytic leukemia	U33626	3.13
Hemopoietic cell kinase	Y00487	2.69
DnaJ-like protein 1	U28423	2.24
Zinc finger protein 36	M57422	2.17
Protein tyrosine phosphatase, receptor type, G	L09562	2.72
Heat shock 70-kD protein 5	D78645	3.35
NF-κ B inhibitor β	U19799	2.47
Plasminogen activator inhibitor, type I	M33960	2.18
IL-12b	M86671	-2.53
Tumor susceptibility gene 101	U52945	-2.29
T-cell lymphoma invasion and metastasis 1	U05245	-2.18
MAPK kinase 4	U18310	-2.38
Lysosomal-associated protein transmembrane 4A	U34259	-2.52
MAPK kinase 6	X97052	-2.01
PKC, α	M25811	-2.11
PKC, β	X53532	-2.18
I-TRAF	U51907	-2.03
Tnf receptor-associated factor 3	U21050	-5.23
CD22 antigen	L16928	-2.56
CD152 antigen	X05719	-2.06
Selectin, platelet (p-selectin) ligand	X91144	-2.02
Growth factors		
Colony-stimulating factor 1 receptor	X68932	-2.37
Erythropoietin receptor	J04843	-2.75
Insulin receptor substrate 1	L24563	-2.93
Fibroblast growth factor receptor 1	M28998	-2.71
GM-CSF-R-α	M85078	-2.41
Corticotropin-releasing hormone receptor	X72305	-4.66
Insulin-like growth factor 1	X04480	-2.46
Myosin light chain, alkali, nonmuscle	U04443	-2.14
Small inducible cytokine A4	M35590	-2.67
Epidermal growth factor	L21671	-4.48
Vascular endothelial growth factor	M95200	-5.06
Nerve growth factor, α	M11434	-3.32
Tyrosine kinase receptor 1	X80764	5.20
Plasminogen activator, urokinase	X02389	2.56
Small inducible cytokine subfamily, member 2	X53798	3.04
Neuroleukin	M14220	2.06
Glial cell line derived neurotrophic factor	D49921	2.35
Cell cycle		
Cell division cycle 2 homolog (S. pombe)-like 1	M58633	3.82
Cyclin-dependent kinase inhibitor 1A (P21)	U09507	2.31
Cell division cycle 25 homolog A (S. cerevisiae)	U27323	2.72
Cyclin G	Z37110	2.66
Retinoblastoma 1	M26391	2.76
Wee 1 homolog (S. pombe)	D30743	-2.13
Development/differentiation		
Fyn proto-oncogene	U70324	2.14
Engrailed 1	L12703	2.07

**Table S3. Continued**

Function	Accession no.	Fold change
Leukemia inhibitory factor	D26177	2.79
POU domain, class 3, transcription factor 1	X56959	-2.03
Paired box gene 6 (Pax6)	X63963	-2.12
Stromal cell-derived factor 1 receptor	D87747	-3.68
Homeo box B5	M26283	-2.57
Kruppel-like factor 3 (basic)	U36340	-2.23
Growth differentiation factor 9 (BMP15)	X77113	-2.18
Tissue inhibitor of metalloproteinase 3	L19622	-3.11
Thrombopoietin	L34169	-2.34
Cathepsin B	M14222	-2.38
Cathepsin D	X53337	3.12
Follistatin	Z29532	2.72
Cell death		
Defender against cell death 1	U83628	32.22
BCL2-binding athanogene 1 (BAG1)	U17162	-2.30
DNA repair		
Ligase I, DNA, ATP-dependent	U04674	-2.83
Translin	X81464	-2.02
Protein tyrosine kinase 6	U16805	-2.05
Transcription/translation		
Transcription factor MTF-1	X71327	2.34
POU domain, class 2, associating factor 1	U43788	2.27
Kruppel-like factor 4 (gut)	U20344	9.99
Zinc finger protein, subfamily 1A, 1 (Ikarus)	L03547	2.75
Nuclease sensitive element binding protein 1	X57621	2.71
Activating transcription factor 3	U19118	2.04
Transcription factor UBF	X60831	-2.26
Neuronal function		
5-hydroxytryptamine (serotonin) receptor 2C	X72230	-3.66
5-hydroxytryptamine (serotonin) receptor 1E $\beta$	Z14224	-2.15
Glutamate receptor, ionotropic, AMPA1 ( $\alpha$ 1)	X57497	-2.14
Cytoskeleton		
Keratin complex 2, basic, gene 1	M10937	-2.09
Myosin Ib	L00923	-2.99
Myosin light chain, alkali, cardiac atria	M19436	2.20

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