A list of 158 host factors related to influenza virus infection verified in vivo.

Gene	Entrez	Direction of genes	
symbol	ID	(lead to pathology or virus	References
		replication)	547
LCAD	33	1	[1]
ADORA1	134	<u> </u>	[2]
ANXA1	301	Indeterminable	[3,4]
SAP	325	Indeterminable	[5]
BIRC3	330	↓	[6]
ARNTL	406	↓	[7]
AXL	558	↓ • • • • • • • • • • • • • • • • • • •	[8]
BCL2A1	597	Inessential	[9]
<i>C3</i>	718	↓	[10]
C4A	720	↓	[10]
C4B	721 727	↓ • • • • • • • • • • • • • • • • • • •	[10]
C5	727	Inessential	[10]
CAMP	820	↓	[11]
CASP1	834	<u> </u>	[12]
CASP6	839	+	[13]
CD79A	973	Inessential	[14]
CCR2	1231	<u>↑</u>	[15]
CCR5	1234	\downarrow	[16]
TPL2	1326	<u> </u>	[17]
CRP	1401	1	[18]
CSF2	1437	↓	[19]
CSF2RA	1438	\downarrow	[20]
CSF2RB	1439	\downarrow	[19]
NOX2	1536	\uparrow	[21]
S1P1	1901	<u> </u>	[22]
SERPINB1	1992	.	[23]
FAP	2191	Inessential	[24]
FCER1G	2207	\downarrow	[25]
FGF2	2247	\downarrow	[26]
FHL2	2274	1	[27]
FMRP	2332	<u> </u>	[28]
ADAP	2533	\	[29]
GALNT3	2591	<u></u>	[30]
CXCR3	2833	↑	[31]
CXCL2	2920	<u> </u>	[32]
IFI35	3430	↑	[33]
IFIT1	3434	Inessential	[34]
IFNAR1	3454	<u> </u>	[35]
IFNAR2	3455	↓	[36]
IFNG	3458	Indeterminable	[37,38]
IGHM	3507	↓	[39]
IL1A	3552	↓	[40]
IL1B	3553	\	[40]
IL1R1	3554	<u> </u>	[40]
IL6	3569	\downarrow	[41]

Gene	Entrez ID	Direction of genes	
symbol		(lead to pathology or virus replication)	References
IL7	3574	Inessential	[42]
<i>IL10</i>	3586	↑	[43]
IL12A	3592	·	[44]
IL12H	3593	*	[44]
IL12B IL15	3600	↓	[45]
IL13 IL17A	3605	 ↑	[46]
IL17A IL18	3606		[47]
CXCL10	3627	↓	[31]
IRF3	3661		[48]
IRF5	3663	→	[49]
IRF7	3665		[48]
ASK1	4217	↓ 	[50]
MLK3	4296	Inessential	[51]
CD200	4345		[52]
MPO MPO	4353	*	[53]
MX1	4599		[54]
MYD88	4615	*	[55]
NFE2L2	4780	*	[56]
NOS2	4843	*	[57]
NRAS	4893		[58]
P2XR7	5027	*	[59]
SERPINE1	5054		[60]
PCBP2	5094	*	[61]
CXCL4	5196	j.	[62]
PIK3CG	5294	j	[63]
PLG	5340	<u> </u>	[64]
PKR	5610	į.	[65]
PRNP	5621	j	[66]
PTAFR	5724	<u> </u>	[67]
PTX3	5806	į.	[68]
RAG2	5897	\downarrow	[69]
RNASEL	6041	\downarrow	[70]
CCL2	6347	↑	[71]
<i>SFTPD</i>	6441		[72]
ST6GAL1	6480	Inessential	[73]
SOD1	6647	\downarrow	[74]
SOD2	6648	\downarrow	[74]
SOD3	6649	↓	[74]
STAT1	6772	\downarrow	[75]
STAT2	6773	\downarrow	[76]
TGFB1	7040	\downarrow	[77]
TGFB2	7042	\downarrow	[77]
TGFB3	7043	.	[77]
TLR2	7097	Indeterminable	[78,79]
TLR3	7098	Indeterminable	[79,80]
TLR4	7099	Indeterminable	[78,79]
TMPRSS2	7113	<u> </u>	[81]

	T	Direction of genes	
Gene	Entrez	(lead to pathology or virus	References
symbol	ID	replication)	
TNF	7124		[82]
TNFAIP3	7128	<u>†</u>	[83]
<i>TP53</i>	7157	į	[84]
UBE1L	7318	\	[85]
OASL	8638	Į.	[70,86]
MARCO	8685	↑	[87]
TNFSF10	8743	Indeterminable	[88,89]
<i>RIPK2</i>	8767	\downarrow	[90]
TNFRSF18	8784	\downarrow	[91]
NCR1	9437	\downarrow	[92]
AIM2	9447	Indeterminable	[93,94]
THEMIS2	9473	Inessential	[95]
PTGES	9536	↑	[96]
CXCL14	9547	Inessential	[97]
ISG15	9636	\downarrow	[85,98]
SOCS5	9655	\downarrow	[99]
MVP	9961	\downarrow	[100]
<i>ZMPSTE24</i>	10269	\downarrow	[101]
IFITM3	10410	\downarrow	[102]
MINDIN	10417	\downarrow	[103]
ATG7	10533	↑	[104]
RIPK3	11035	\downarrow	[105]
ADAMTS5	11096	\downarrow	[106]
IRAK3	11213	\downarrow	[107]
ATG14	22863	<u> </u>	[104]
PAD4	23569	Inessential	[108]
DDX58	23586	\downarrow	[109]
CLEC5A	23601	<u>†</u>	[110]
TRIM29	23650	<u> </u>	[111]
IL17RA	23765	<u> </u>	[46]
NOX1	27035	<u>↑</u>	[21]
DMAC2L	27109	\	[10]
ASC	29108	↓	[112]
DUOX2	50506	↓	[113]
NOX4	50507	Ť	[21]
NOX3	50508	Tu = = = = 1	[21]
IL21R	50615	Inessential	[114]
TLR7	51284	*	[79,115]
DUOX1 ATG16L1	53905 55054		[21]
	56300	↓	[116]
IL36G MAVS	50300 57506	\rightarrow	[117] [109]
EPG5	57724	↓	[104]
NLRC4	58484	Inessential	
ACE2	59272	1110550111181	[112] [118]
NOD2	64127	↓ Inessential	
NOD2 CARD9	64170	messemai	[119] [120]
NOX5	79400	I ↑	[21]
ΙνΟΛΙ	/ 5400		[41]

Gene symbol	Entrez ID	Direction of genes (lead to pathology or virus replication)	References
NLRX1	79671	Inessential	[121]
ZBP1	81030	↓	[122]
UNC93B1	81622	Inessential	[123]
TRIM56	81844	Inessential	[124]
<i>NLRC5</i>	84166	Inessential	[125]
RSAD2	91543	Inessential	[126]
<i>NLRP3</i>	114548	\downarrow	[12,112]
SOCS4	122809	\downarrow	[99]
CD200R	131450	\downarrow	[52]
PAR1	145624	↑	[127]
TRIF	148022	↑	[78]
IFNLR1	163702	\downarrow	[128]
MLKL	197259	Inessential	[129]
SPRED2	200734	↓	[130]
TREML4	285852	↓	[131]
SFTPA1	653509	↓	[132]
SFTPA2	729238	\downarrow	[132]
IRG1	730249	Inessential	[133]

The 28 genes in this list overlapped with the 784 DEGs derived from IAV-infected patients are highlighted.

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