

Supplemental Table S1.

False-positive hits:		Fold induction				
		Standard	vRNA depletion	Standard	vRNA depletion	RT-PCR
Gene ID	Name	70-mer arrays		Affymetrix arrays		n.d*/n.t**
NM_016808	Usp2	<u>310.3</u>	<u>70.1</u>	<2	<2	1.2
NM_001001809	Olf218	<u>210.0</u>	<u>12.1</u>	<2	<2	n.d.
NM_010915	Klk1b4	<u>209.2</u>	<u>22.2</u>	<2	<2	n.d.
NM_012013	Figla	<u>162.7</u>	<u>23.1</u>	<2	<2	n.d.
NM_130886	Card14	<u>9.8</u>	<2	<2	<2	1.6
NM_029562	Cyp2d26	<u>9.9</u>	<2	<2	<2	n.d.
NM_175305	Lrrc19	<u>9.9</u>	<2	<2	<2	n.d.
NM_174854	Disc1	<u>9.7</u>	<2	<2	<2	n.d.
NM_007954	Es1	<u>9.5</u>	<2	<2	<2	n.d.
NM_010333	Edg5	<u>4.8</u>	<2	<u>5.5</u>	<2	1.0
NM_178403	Pus7	<2	<2	<u>4.4</u>	<2	1.2
NM_181400	Wdr47	<2	<2	<u>2.3</u>	<2	1.5
NM_133753	Errfi1	<2	<2	<u>2.1</u>	<2	1.5

*n.d. = not detectable, **n.t. = not tested

Supplemental Table S1. continued

Additional hits:		Fold induction				
		Standard	vRNA depletion	Standard	vRNA depletion	RT-PCR
Gene ID	Name	70-mer arrays		Affymetrix arrays		n.d*/n.t**
NM_010234	Fos	<2	<u>2.1</u>	<u>14.5</u>	<u>16.5</u>	<u>27.5</u>
NM_008390	Irf1	<2	<u>2.4</u>	<u>2.2</u>	<u>2.4</u>	<u>2.8</u>
NM_008562	Mcl1	<2	<u>2.3</u>	<u>2.1</u>	<u>2.0</u>	<u>2.1</u>
NM_013642	Dusp1	<2	<u>2.4</u>	<u>2.6</u>	<u>3.8</u>	n.t.
NM_009344	Phlda1	<2	<u>2.5</u>	<u>2.0</u>	<u>2.6</u>	n.t.
NM_178392	Snapc1	<2	<u>2.1</u>	<u>2.0</u>	<u>2.4</u>	n.t.
NM_010495	Id1	<2	<u>2.1</u>	<u>2.5</u>	<u>2.6</u>	n.t.
NM_008235	Hes1	<2	<u>2.0</u>	<u>2.7</u>	<u>2.7</u>	n.t.
NM_029092	Rg9mtd1	<2	<u>2.0</u>	<u>2.0</u>	<u>2.1</u>	n.t.
NM_010496	Id2	<2	<u>2.0</u>	<2	<u>2.7</u>	<u>2.8</u>
NM_172154	Lcor	<2	<u>2.0</u>	<2	<u>2.7</u>	<u>4.2</u>
NM_011498	Bhlhb2	<2	<u>2.1</u>	<2	<u>2.5</u>	n.t.
NM_010591	Jun	<2	<u>2.0</u>	<2	<u>2.3</u>	n.t.
NM_008871	Serpine1	<2	<u>2.3</u>	<2	<2	<u>2.2</u>
NM_010804	Mllt10	<2	<2	<2	<u>2.6</u>	<u>2.1</u>

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Supplemental Table S1. continued

Hits by both platforms and both methods:		Fold induction				
		Standard	vRNA depletion	Standard	vRNA depletion	RT-PCR
Gene ID	Name	70-mer arrays		Affymetrix arrays		n.d*/n.t**
NM_010907	Nfkbia	<u>2.6</u>	<u>2.7</u>	<u>2.9</u>	<u>3.5</u>	<u>3.4</u>
NM_022331	Herpud1	<u>2.4</u>	<u>2.6</u>	<u>4.3</u>	<u>5.7</u>	<u>5.1</u>
NM_008176	Cxcl1	<u>5.9</u>	<u>7.3</u>	<u>6.0</u>	<u>6.1</u>	<u>10.8</u>
NM_009140	Cxcl2	<u>3.8</u>	<u>6.3</u>	<u>6.3</u>	<u>7.7</u>	<u>7.2</u>
NM_013692	Klf10	<u>4.7</u>	<u>6.1</u>	<u>5.4</u>	<u>5.8</u>	n.t.
NM_013602	Mt1	<u>2.2</u>	<u>3.1</u>	<u>3.5</u>	<u>4.3</u>	n.t.
NM_011803	Klf6	<u>2.6</u>	<u>2.7</u>	<u>2.6</u>	<u>2.9</u>	n.t.
NM_153159	Zc3h12a	<u>2.1</u>	<u>3.4</u>	<u>2.4</u>	<u>2.8</u>	n.t.
NM_015786	Hist1h1c	<u>2.6</u>	<u>4.4</u>	<u>5.2</u>	<u>7.5</u>	n.t.
NM_133662	Ier3	<u>3.2</u>	<u>3.4</u>	<u>4.7</u>	<u>4.9</u>	n.t.
NM_008321	Id3	<u>2.6</u>	<u>2.8</u>	<u>2.9</u>	<u>3.2</u>	n.t.

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