

**ISMARA****Project****Final-1****Navigation**

[All motifs sorted by activity significance](#)

[All samples sorted alphabetically](#)

[Mean activities](#)

**Downloads**

[Activities table](#)

[Activity deltas table](#)

[Regulatory interactions](#)

[Motifs sorted by significance \(text\)](#)

[Download the whole report](#)

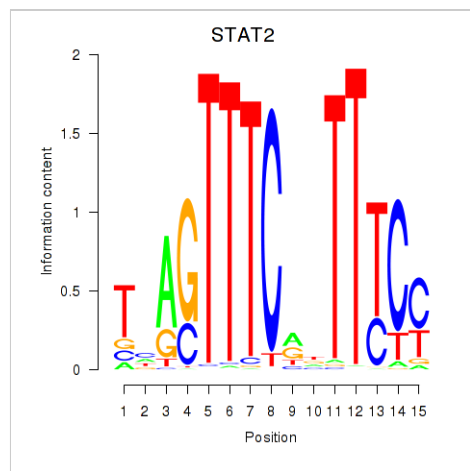
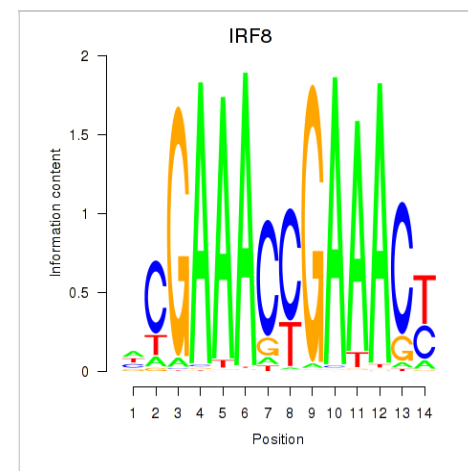
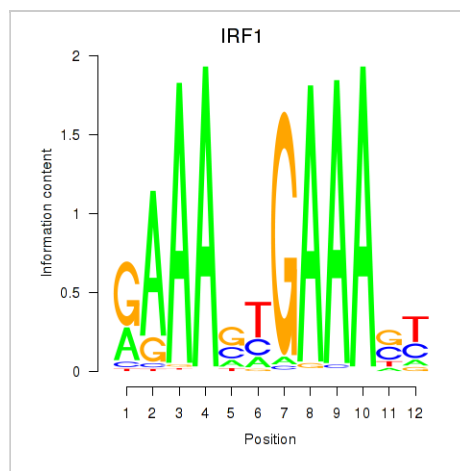
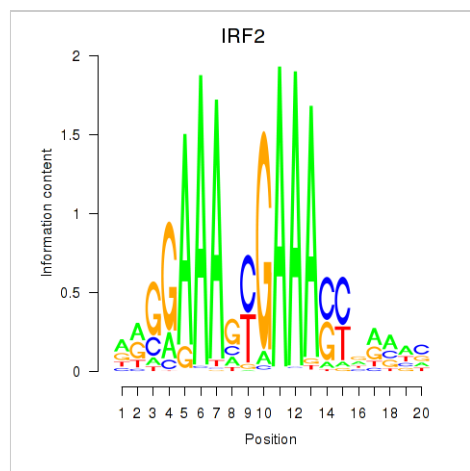
[Activity graph](#)

[Z-values graph](#)

[Top targets](#)

[GO analysis](#)

[Back to top](#)

**Motif ID: IRF2\_STAT2\_IRF8\_IRF1****Z-value: 5.971****Transcription factors associated with IRF2\_STAT2\_IRF8\_IRF1:**

Gene Symbol	Entrez ID	Gene Name
<a href="#">IRF1</a>	<a href="#">ENSG00000125347.9</a>	<a href="#">IRF1</a>

<a href="#">IRF2</a>	<a href="#">ENSG00000168310.6</a>	<a href="#">IRF2</a>
<a href="#">IRF8</a>	<a href="#">ENSG00000140968.6</a>	<a href="#">IRF8</a>
<a href="#">STAT2</a>	<a href="#">ENSG00000170581.9</a>	<a href="#">STAT2</a>

**ISMARA****Project**

Final-1

**Navigation**

[All motifs sorted by activity significance](#)

[All samples sorted alphabetically](#)

[Mean activities](#)

**Downloads**

[Activities table](#)

[Activity deltas table](#)

[Regulatory interactions](#)

[Motifs sorted by significance \(text\)](#)

[Download the whole report](#)

[Activity graph](#)

[Z-values graph](#)

[Top targets](#)

[GO analysis](#)

[Back to top](#)

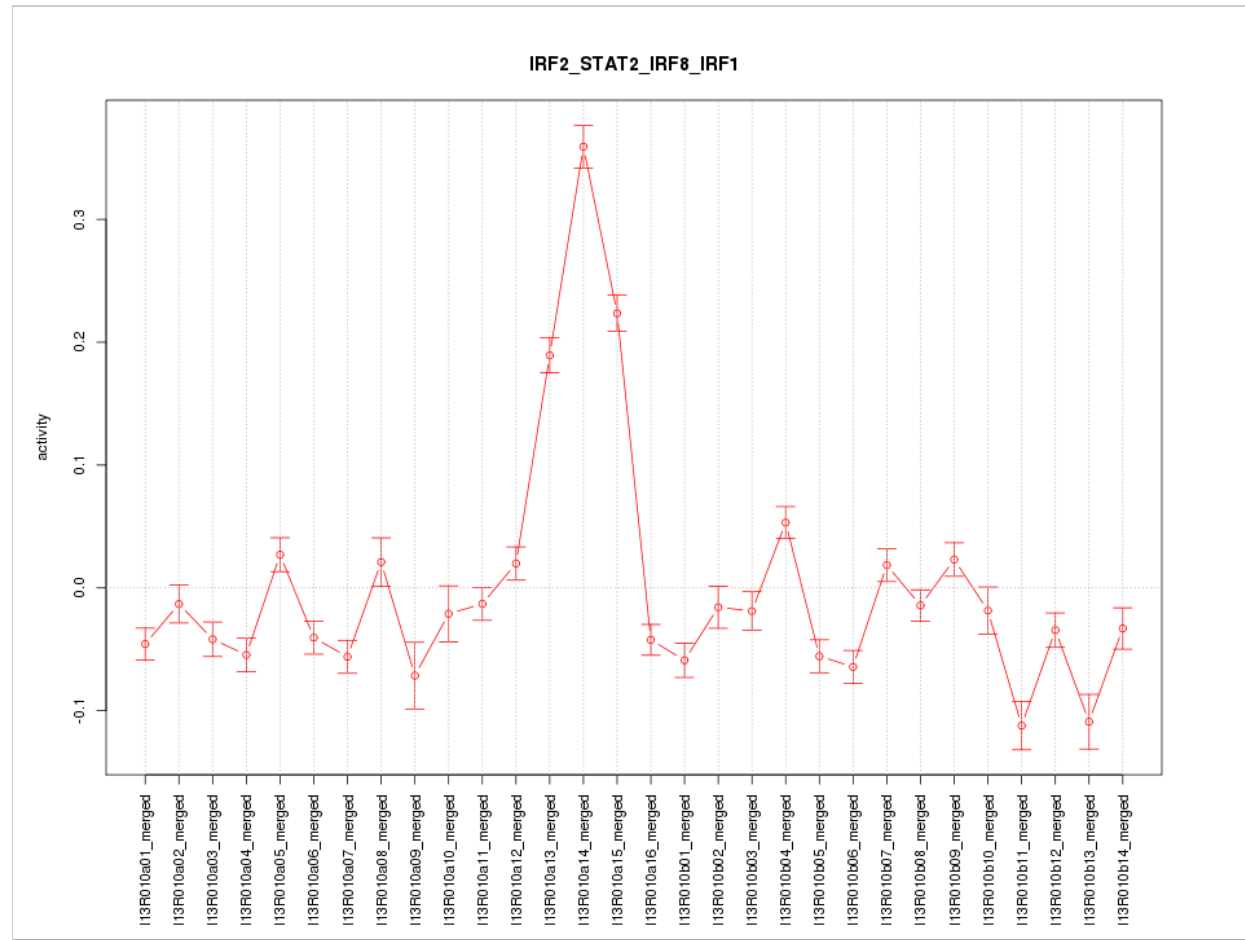
**Activity-expression correlation:**

Gene Symbol	Promoter	Pearson corr. coef.	P-value	Plot
STAT2	<a href="#">hg19_v2_chr12 - 56753858_56753930</a>	0.82	2.1e-08	<a href="#">Click!</a>
IRF1	<a href="#">hg19_v2_chr5 - 131826457_131826514</a>	0.70	1.9e-05	<a href="#">Click!</a>
IRF2	<a href="#">hg19_v2_chr4 - 185395672_185395734</a>	0.62	2.8e-04	<a href="#">Click!</a>
IRF8	<a href="#">hg19_v2_chr16 + 85936295_85936444</a>	0.29	1.3e-01	<a href="#">Click!</a>

**Activity profile for motif IRF2\_STAT2\_IRF8\_IRF1.**

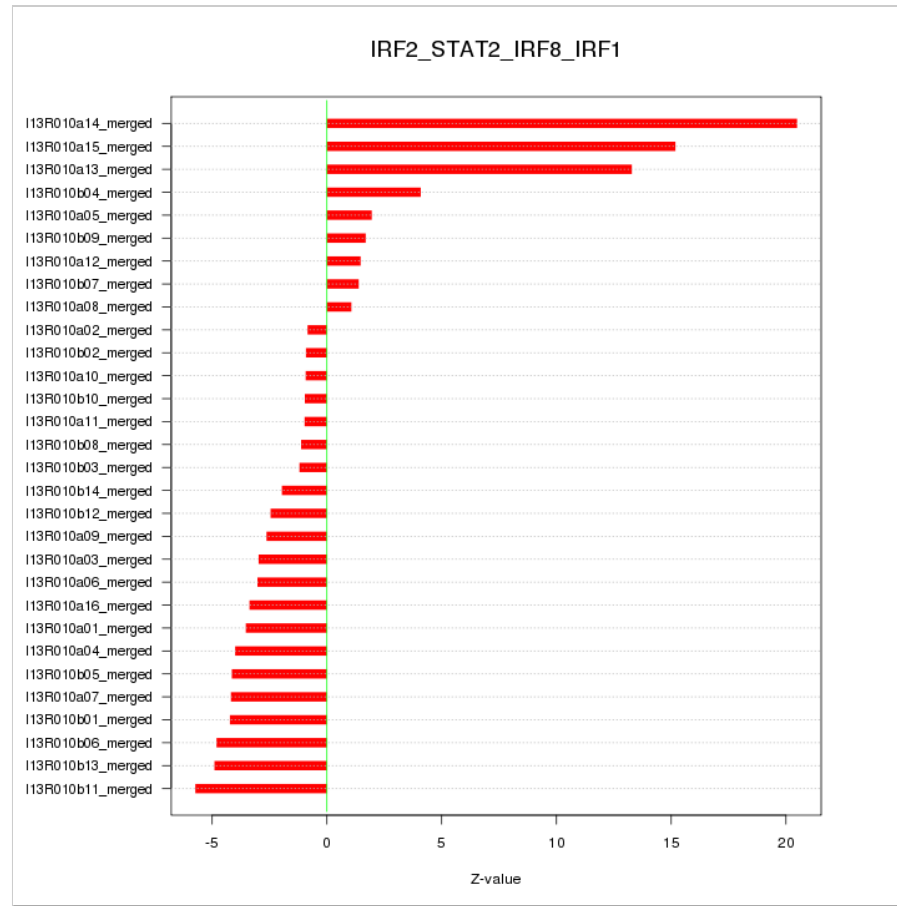
**ISMARA****Project**

Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)**Sorted Z-values histogram for motif IRF2\_STAT2\_IRF8\_IRF1**[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

**ISMARA****Project**

Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)

**Network of associations between targets according to the [STRING database](#).**

[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity  
significance

All samples sorted  
alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by  
significance (text)

Download the whole report

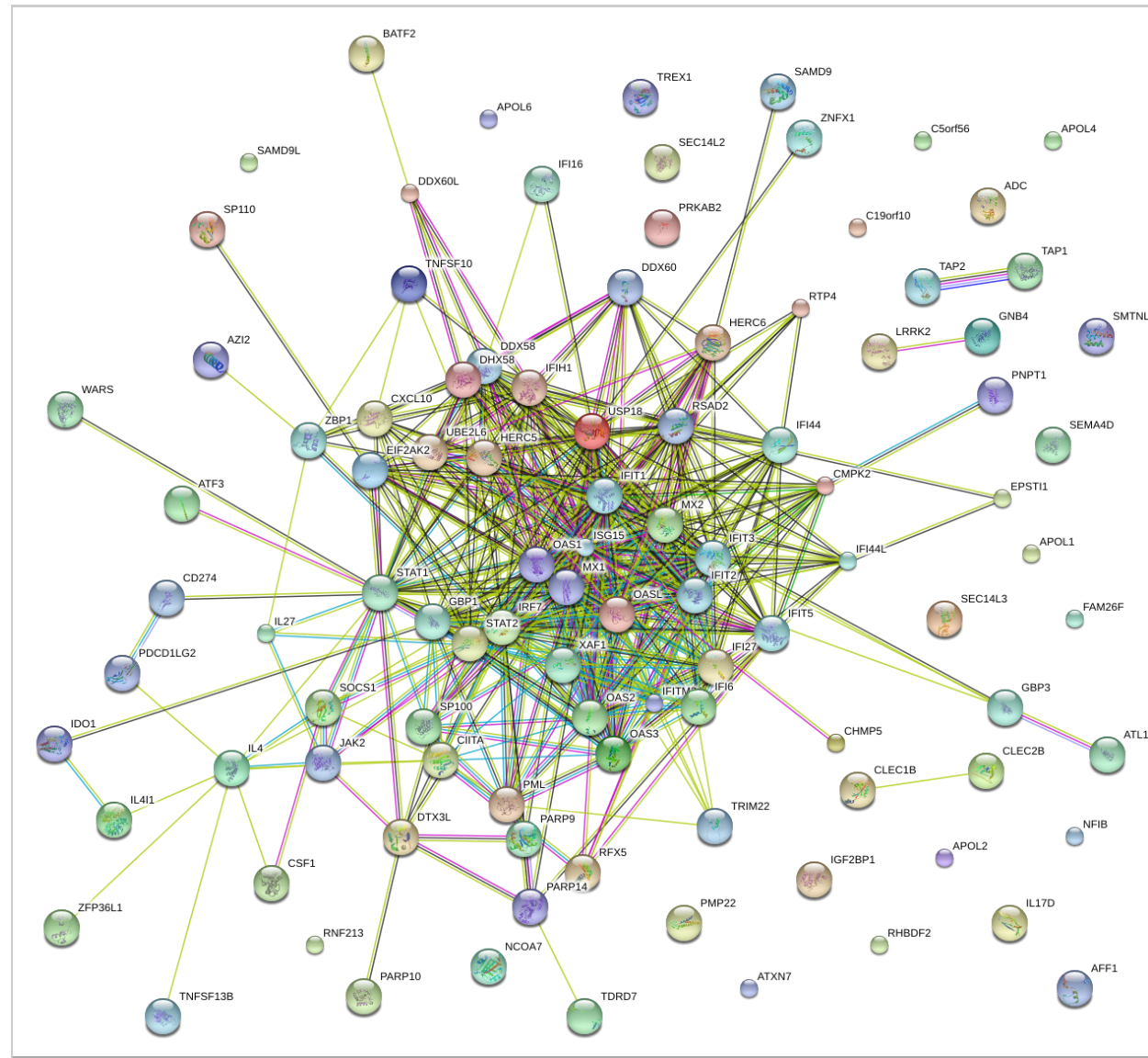
Activity graph

Z-values graph

Top targets

GO analysis

Back to top

**First level regulatory network of IRF2\_STAT2\_IRF8\_IRF1**

**ISMARA**

**Project**

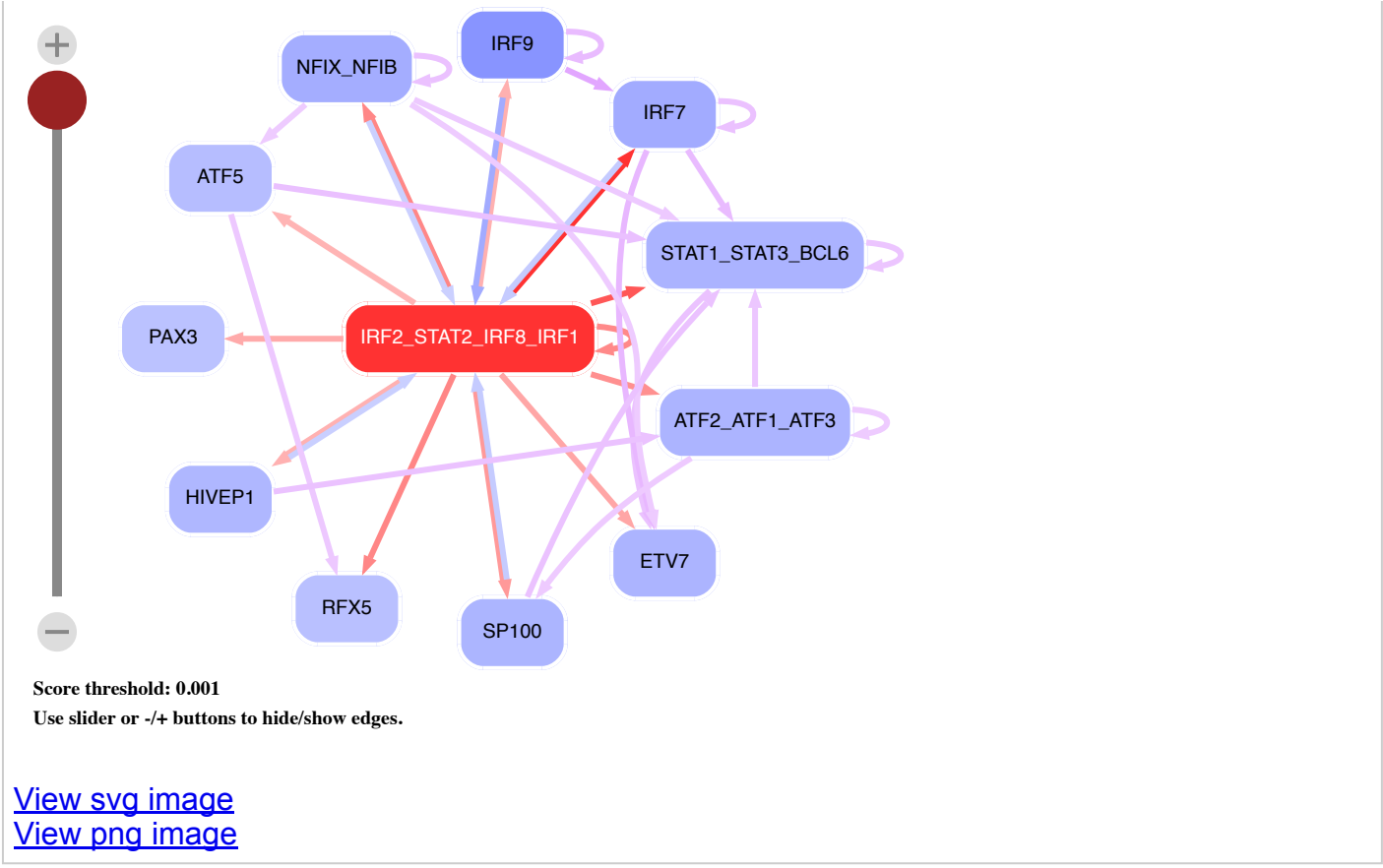
Final-1

**Navigation**

- All motifs sorted by activity significance
- All samples sorted alphabetically
- Mean activities

**Downloads**

- Activities table
- Activity deltas table
- Regulatory interactions
- Motifs sorted by significance (text)
- Download the whole report



**Top targets:**

Search:

Show  entries

Showing 1 to 200 of 200 entries

- Activity graph
- Z-values graph
- Top targets
- GO analysis
- Back to top

Promoter	Score	Refseq	Gene Symbol	Gene Name
<a href="#">chr2_+_7005959</a>	69.597	<a href="#">ENST00000442639.1</a>	<a href="#">RSAD2</a>	<a href="#">radical S-adenosyl methionine domain containing 2</a>
<a href="#">chr2_+_7017796</a>	56.421	<a href="#">ENST00000382040.3</a>	<a href="#">RSAD2</a>	<a href="#">radical S-adenosyl methionine domain containing 2</a>
				<a href="#">interferon-induced protein 4</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr1_+_79086088</a>	56.379	<a href="#">ENST00000370751.5</a> <a href="#">ENST00000342282.3</a>	<a href="#">IFI44L</a> <a href="#">IFI44L</a>	<a href="#">4-like interferon-induced protein 4 4-like</a>
<a href="#">chr10_+_91152303</a>	44.091	<a href="#">ENST00000371804.3</a>	<a href="#">IFIT1</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 1</a>
<a href="#">chr1_+_948803</a>	40.058	<a href="#">ENST00000379389.4</a>	<a href="#">ISG15</a>	<a href="#">ISG15 ubiquitin-like modifier</a>
<a href="#">chr10_+_91087651</a>	38.585	<a href="#">ENST00000371818.4</a>	<a href="#">IFIT3</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 3</a>
<a href="#">chr2_-_7005785</a>	38.074	<a href="#">ENST00000458098.1</a> <a href="#">ENST00000256722.5</a> <a href="#">ENST00000404168.1</a>	<a href="#">CMPK2</a> <a href="#">CMPK2</a>	<a href="#">cytidine monophosphate (UMP-CMP) kinase 2, mitochondrial</a> <a href="#">cytidine monophosphate (UMP-CMP) kinase 2, mitochondrial</a>
<a href="#">chr10_+_91092241</a>	37.841	<a href="#">ENST00000371811.4</a>	<a href="#">IFIT3</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 3</a>
<a href="#">chr12_+_113376157</a>	32.420	<a href="#">ENST00000228928.7</a>	<a href="#">OAS3</a>	<a href="#">2'-5'-oligoadenylate synthetase 3, 100kDa</a>
<a href="#">chr17_+_6659354</a>	31.302	<a href="#">ENST00000574907.1</a>	<a href="#">XAF1</a>	<a href="#">XIAP associated factor 1</a>
<a href="#">chr22_+_18632666</a>	30.767	<a href="#">ENST00000215794.7</a>	<a href="#">USP18</a>	<a href="#">ubiquitin specific peptidase 18</a>
<a href="#">chr4_+_89378261</a>	26.528	<a href="#">ENST00000264350.3</a>	<a href="#">HERC5</a>	<a href="#">HECT and RLD domain containing E3 ubiquitin protein ligase 5</a>
<a href="#">chr17_+_6658878</a>	26.450	<a href="#">ENST00000574394.1</a>	<a href="#">XAF1</a>	<a href="#">XIAP associated factor 1</a>
<a href="#">chr1_+_79115503</a>	25.933	<a href="#">ENST00000370747.4</a> <a href="#">ENST00000438486.1</a> <a href="#">ENST00000545124.1</a>	<a href="#">IFI44</a> <a href="#">IFI44</a>	<a href="#">interferon-induced protein 4 4</a> <a href="#">interferon-induced protein 4 4</a>
<a href="#">chr12_+_113416340</a>	25.685	<a href="#">ENST00000552756.1</a>	<a href="#">OAS2</a>	<a href="#">2'-5'-oligoadenylate synthetase 2, 100kDa</a>

**ISMARA**

**Project**

**Final-1**

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

				<a href="#">ase 2. 69/71kDa</a>
<a href="#">chr17_-40264321</a>	25.529	<a href="#">ENST00000430773.1</a> <a href="#">ENST00000413196.2</a>	<a href="#">DHX58</a> <a href="#">DHX58</a>	<a href="#">DEXH (Asp-Glu-X-His) box polypeptide 58</a> <a href="#">DEXH (Asp-Glu-X-His) box polypeptide 58</a>
<a href="#">chr12_+113376249</a>	25.524	<a href="#">ENST00000551007.1</a> <a href="#">ENST00000548514.1</a>	<a href="#">OAS3</a> <a href="#">OAS3</a>	<a href="#">2'-5'-oligoadenylate synthetase 3. 100kDa</a> <a href="#">2'-5'-oligoadenylate synthetase 3. 100kDa</a>
<a href="#">chr4_-169239921</a>	24.929	<a href="#">ENST00000514995.1</a> <a href="#">ENST00000393743.3</a>	<a href="#">DDX60</a> <a href="#">DDX60</a>	<a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 60</a> <a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 60</a>
<a href="#">chr13_-43566301</a>	24.729	<a href="#">ENST00000313624.7</a> <a href="#">ENST00000398762.3</a> <a href="#">ENST00000313640.7</a>	<a href="#">EPST11</a> <a href="#">EPST11</a>	<a href="#">epithelial stromal interaction 1 (breast)</a> <a href="#">epithelial stromal interaction 1 (breast)</a>
<a href="#">chr11_-64764435</a>	24.560	<a href="#">ENST00000301887.4</a> <a href="#">ENST00000534177.1</a>	<a href="#">BATF2</a> <a href="#">BATF2</a>	<a href="#">basic leucine zipper transcription factor, ATF-like 2</a> <a href="#">basic leucine zipper transcription factor, ATF-like 2</a>
<a href="#">chr14_+94577074</a>	24.433	<a href="#">ENST00000555744.1</a> <a href="#">ENST00000444961.1</a> <a href="#">ENST00000448882.1</a> <a href="#">ENST00000557098.1</a> <a href="#">ENST00000554800.1</a> <a href="#">ENST00000556544.1</a> <a href="#">ENST00000298902.5</a> <a href="#">ENST00000555819.1</a> <a href="#">ENST00000557634.1</a>	<a href="#">IFI27</a> <a href="#">IFI27</a>	<a href="#">interferon, alpha-inducible protein 27</a> <a href="#">interferon, alpha-inducible protein 27</a>
<a href="#">chr10_+91174486</a>	22.921	<a href="#">ENST00000416601.1</a>	<a href="#">IFIT5</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 5</a>
<a href="#">chr4_+89300158</a>	22.724	<a href="#">ENST00000502870.1</a>	<a href="#">HERC6</a>	<a href="#">HECT and RLD domain containing E3 ubiquitin protei</a>



**ISMARA**

**Project**

**Final-1**

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

				<a href="#">n ligase family member 6</a>
<a href="#">chr4_-76944621</a>	22.397	<a href="#">ENST00000306602.1</a>	<a href="#">CXCL10</a>	<a href="#">chemokine (C-X-C motif) ligand 10</a>
<a href="#">chr12_+113416265</a>	21.771	<a href="#">ENST00000449768.2</a>	<a href="#">OAS2</a>	<a href="#">2'-5'-oligoadenylate synthetase 2, 69/71kDa</a>
<a href="#">chr12_+113416191</a>	21.351	<a href="#">ENST00000342315.4</a> <a href="#">ENST00000392583.2</a>	<a href="#">OAS2</a> <a href="#">OAS2</a>	<a href="#">2'-5'-oligoadenylate synthetase 2, 69/71kDa</a> <a href="#">2'-5'-oligoadenylate synthetase 2, 69/71kDa</a>
<a href="#">chr9_+5450503</a>	19.893	<a href="#">ENST00000381577.3</a> <a href="#">ENST00000381573.4</a>	<a href="#">CD274</a> <a href="#">CD274</a>	<a href="#">CD274 molecule</a> <a href="#">CD274 molecule</a>
<a href="#">chr21_+42792442</a>	19.836	<a href="#">ENST00000398600.2</a>	<a href="#">MX1</a>	<a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a>
<a href="#">chr10_+91061712</a>	19.794	<a href="#">ENST00000371826.3</a>	<a href="#">IFIT2</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 2</a>
<a href="#">chr3_+122399697</a>	19.502	<a href="#">ENST00000494811.1</a>	<a href="#">PARP14</a>	<a href="#">poly (ADP-ribose) polymerase family, member 14</a>
<a href="#">chr10_+91174314</a>	19.292	<a href="#">ENST00000371795.4</a>	<a href="#">IFIT5</a>	<a href="#">interferon-induced protein with tetratricopeptide repeats 5</a>
<a href="#">chr1_-27998689</a>	17.923	<a href="#">ENST00000339145.4</a> <a href="#">ENST00000362020.4</a> <a href="#">ENST00000361157.6</a>	<a href="#">IFI6</a> <a href="#">IFI6</a>	<a href="#">interferon, alpha-inducible protein 6</a> <a href="#">interferon, alpha-inducible protein 6</a>
<a href="#">chr17_+6659153</a>	17.782	<a href="#">ENST00000361842.3</a> <a href="#">ENST00000441631.1</a> <a href="#">ENST00000438512.1</a> <a href="#">ENST00000346752.4</a>	<a href="#">XAF1</a> <a href="#">XAF1</a>	<a href="#">XIAP associated factor 1</a> <a href="#">XIAP associated factor 1</a>
<a href="#">chr3_+187086120</a>	17.094	<a href="#">ENST00000259030.2</a>	<a href="#">RTP4</a>	<a href="#">receptor (chemosensory) transporter protein 4</a>
<a href="#">chr16_-28518153</a>	16.848	<a href="#">ENST00000356897.1</a>	<a href="#">IL27</a>	<a href="#">interleukin 27</a>
<a href="#">chr9_+5510558</a>	16.841	<a href="#">ENST00000397747.3</a>	<a href="#">PDCD1LG2</a>	<a href="#">programmed cell death 1 ligand 2</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr9_+_5510492</a>	15.654	<a href="#">ENST00000397745.2</a>	<a href="#">PDCD1LG2</a>	<a href="#">programmed cell death 1 ligand 2</a>
<a href="#">chr17_-_40264692</a>	15.241	<a href="#">ENST00000591220.1</a> <a href="#">ENST00000251642.3</a>	<a href="#">DHX58</a> <a href="#">DHX58</a>	<a href="#">DEXH (Asp-Glu-X-His) box polypeptide 58</a> <a href="#">DEXH (Asp-Glu-X-His) box polypeptide 58</a>
<a href="#">chr7_-_92777606</a>	14.970	<a href="#">ENST00000318238.4</a> <a href="#">ENST00000437805.1</a> <a href="#">ENST00000446959.1</a> <a href="#">ENST00000439952.1</a> <a href="#">ENST00000414791.1</a> <a href="#">ENST00000446033.1</a> <a href="#">ENST00000411955.1</a>	<a href="#">SAMD9L</a> <a href="#">SAMD9L</a>	<a href="#">sterile alpha motif domain containing 9-like</a> <a href="#">sterile alpha motif domain containing 9-like</a>
<a href="#">chr11_-_615942</a>	14.786	<a href="#">ENST00000397574.2</a> <a href="#">ENST00000397562.3</a> <a href="#">ENST00000330243.5</a> <a href="#">ENST00000397570.1</a>	<a href="#">IRF7</a> <a href="#">IRF7</a>	<a href="#">interferon regulatory factor 7</a> <a href="#">interferon regulatory factor 7</a>
<a href="#">chr1_-_89488510</a>	14.490	<a href="#">ENST00000564665.1</a> <a href="#">ENST00000370481.4</a>	<a href="#">GBP3</a> <a href="#">GBP3</a>	<a href="#">guanylate binding protein 3</a> <a href="#">guanylate binding protein 3</a>
<a href="#">chr20_-_56195449</a>	14.359	<a href="#">ENST00000541799.1</a>	<a href="#">ZBP1</a>	<a href="#">Z-DNA binding protein 1</a>
<a href="#">chr9_-_32526184</a>	13.721	<a href="#">ENST00000545044.1</a> <a href="#">ENST00000379868.1</a>	<a href="#">DDX58</a> <a href="#">DDX58</a>	<a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 58</a> <a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 58</a>
<a href="#">chr1_-_89531041</a>	13.642	<a href="#">ENST00000370473.4</a>	<a href="#">GBP1</a>	<a href="#">guanylate binding protein 1, interferon-inducible</a>
<a href="#">chr9_-_32526299</a>	12.917	<a href="#">ENST00000379883.2</a> <a href="#">ENST00000379882.1</a>	<a href="#">DDX58</a> <a href="#">DDX58</a>	<a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 58</a> <a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 58</a>
<a href="#">chr2_-_163175133</a>	12.832	<a href="#">ENST00000263642.2</a> <a href="#">ENST00000421365.2</a>	<a href="#">IFIH1</a> <a href="#">IFIH1</a>	<a href="#">interferon induced with helicase C domain 1</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

				<a href="#">interferon induced with heli case C domain 1</a>
<a href="#">chr12_+_113344811</a>	12.371	<a href="#">ENST00000551241.1</a> <a href="#">ENST00000553185.1</a> <a href="#">ENST00000550689.1</a>	<a href="#">OAS1</a> <a href="#">OAS1</a>	<a href="#">2'-5'-oligoadenylate synthetase 1, 40/46kDa</a> <a href="#">2'-5'-oligoadenylate synthetase 1, 40/46kDa</a>
<a href="#">chr2_-37384175</a>	12.008	<a href="#">ENST00000390013.3</a> <a href="#">ENST00000411537.2</a> <a href="#">ENST00000233057.4</a> <a href="#">ENST00000395127.2</a>	<a href="#">EIF2AK2</a> <a href="#">EIF2AK2</a>	<a href="#">eukaryotic translation initiation factor 2-alpha kinase 2</a> <a href="#">eukaryotic translation initiation factor 2-alpha kinase 2</a>
<a href="#">chr11_-615570</a>	12.006	<a href="#">ENST00000525445.1</a> <a href="#">ENST00000348655.6</a> <a href="#">ENST00000397566.1</a>	<a href="#">IRF7</a> <a href="#">IRF7</a>	<a href="#">interferon regulatory factor 7</a> <a href="#">interferon regulatory factor 7</a>
<a href="#">chr3_+_122283175</a>	11.651	<a href="#">ENST00000383661.3</a>	<a href="#">DTX3L</a>	<a href="#">deltex 3-like (Drosophila)</a>
<a href="#">chr20_-56195525</a>	11.602	<a href="#">ENST00000371173.3</a> <a href="#">ENST00000395822.3</a> <a href="#">ENST00000340462.4</a> <a href="#">ENST00000343535.4</a>	<a href="#">ZBP1</a> <a href="#">ZBP1</a>	<a href="#">Z-DNA binding protein 1</a> <a href="#">Z-DNA binding protein 1</a>
<a href="#">chr11_+_5711010</a>	11.426	<a href="#">ENST00000454828.1</a>	<a href="#">TRIM22</a>	<a href="#">tripartite motif containing 22</a>
<a href="#">chr5_+_131746575</a>	11.406	<a href="#">ENST00000337752.2</a> <a href="#">ENST00000378947.3</a> <a href="#">ENST00000407797.1</a>	<a href="#">C5orf56</a> <a href="#">C5orf56</a>	<a href="#">chromosome 5 open reading frame 56</a> <a href="#">chromosome 5 open reading frame 56</a>
<a href="#">chr2_-191878681</a>	11.226	<a href="#">ENST00000409465.1</a>	<a href="#">STAT1</a>	<a href="#">signal transducer and activator of transcription 1, 91kDa</a>
<a href="#">chr9_+_100174232</a>	11.106	<a href="#">ENST00000355295.4</a>	<a href="#">TDRD7</a>	<a href="#">tudor domain containing 7</a>
<a href="#">chr14_-100841794</a>	10.771	<a href="#">ENST00000556295.1</a> <a href="#">ENST00000554820.1</a>	<a href="#">WARS</a> <a href="#">WARS</a>	<a href="#">tryptophanyl-tRNA synthetase</a> <a href="#">tryptophanyl-tRNA synthetase</a>

**ISMARA**

**Project**

**Final-1**

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr4_-169401628</a>	10.759	<a href="#">ENST00000514748.1</a> <a href="#">ENST00000512371.1</a> <a href="#">ENST00000260184.7</a> <a href="#">ENST00000505890.1</a> <a href="#">ENST00000511577.1</a>	<a href="#">DDX60L</a>    <a href="#">DDX60L</a>	<a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 60-like</a>    <a href="#">DEAD (Asp-Glu-Ala-Asp) box polypeptide 60-like</a>
<a href="#">chr3_+122399444</a>	10.698	<a href="#">ENST00000474629.2</a>	<a href="#">PARP14</a>	<a href="#">poly (ADP-ribose) polymerase family, member 14</a>
<a href="#">chr9_+100174344</a>	10.510	<a href="#">ENST00000422139.2</a>	<a href="#">TDRD7</a>	<a href="#">tudor domain containing 7</a>
<a href="#">chr3_-122283079</a>	10.248	<a href="#">ENST00000471785.1</a> <a href="#">ENST00000466126.1</a>	<a href="#">PARP9</a> <a href="#">PARP9</a>	<a href="#">poly (ADP-ribose) polymerase family, member 9</a> <a href="#">poly (ADP-ribose) polymerase family, member 9</a>
<a href="#">chr21_+42797958</a>	10.197	<a href="#">ENST00000419044.1</a>	<a href="#">MX1</a>	<a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a>
<a href="#">chr21_+42798158</a>	10.118	<a href="#">ENST00000441677.1</a>	<a href="#">MX1</a>	<a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a>
<a href="#">chr21_+42798124</a>	10.063	<a href="#">ENST00000417963.1</a>	<a href="#">MX1</a>	<a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a>
<a href="#">chr11_-57334732</a>	9.683	<a href="#">ENST00000527022.1</a> <a href="#">ENST00000526659.1</a>	<a href="#">UBE2L6</a> <a href="#">UBE2L6</a>	<a href="#">ubiquitin-conjugating enzyme E2L 6</a> <a href="#">ubiquitin-conjugating enzyme E2L 6</a>
<a href="#">chr2_-191878874</a>	9.548	<a href="#">ENST00000392322.3</a> <a href="#">ENST00000392323.2</a> <a href="#">ENST00000424722.1</a> <a href="#">ENST00000361099.3</a>	<a href="#">STAT1</a>   <a href="#">STAT1</a>	<a href="#">signal transducer and activator of transcription 1, 91kDa</a>   <a href="#">signal transducer and activator of transcription 1, 91kDa</a>
<a href="#">chr17_+78234625</a>	9.508	<a href="#">ENST00000508628.2</a> <a href="#">ENST00000582970.1</a>	<a href="#">RNF213</a>	<a href="#">ring finger protein 213</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

		<a href="#">ENST00000456466.1</a> <a href="#">ENST00000319921.4</a>	<a href="#">RNF213</a>	<a href="#">ring finger protein 213</a>
<a href="#">chr3 - 122283424</a>	9.493	<a href="#">ENST00000360356.2</a> <a href="#">ENST00000477522.2</a>	<a href="#">PARP9</a> <a href="#">PARP9</a>	<a href="#">poly (ADP-ribose) polymerase family, member 9</a> <a href="#">poly (ADP-ribose) polymerase family, member 9</a>
<a href="#">chr11 + 5710919</a>	9.441	<a href="#">ENST00000425490.1</a> <a href="#">ENST00000379965.3</a>	<a href="#">TRIM22</a> <a href="#">TRIM22</a>	<a href="#">tripartite motif containing 22</a> <a href="#">tripartite motif containing 22</a>
<a href="#">chr22 + 36649170</a>	9.417	<a href="#">ENST00000440669.2</a> <a href="#">ENST00000438034.1</a> <a href="#">ENST00000427990.1</a> <a href="#">ENST00000347595.7</a> <a href="#">ENST00000397279.4</a> <a href="#">ENST00000433768.1</a>	<a href="#">APOL1</a> <a href="#">APOL1</a>	<a href="#">apolipoprotein L, 1</a> <a href="#">apolipoprotein L, 1</a>
<a href="#">chr3 + 63897605</a>	8.963	<a href="#">ENST00000487717.1</a>	<a href="#">ATXN7</a>	<a href="#">ataxin 7</a>
<a href="#">chr22 - 36635598</a>	8.802	<a href="#">ENST00000454728.1</a>	<a href="#">APOL2</a>	<a href="#">apolipoprotein L, 2</a>
<a href="#">chr3 - 179169330</a>	8.471	<a href="#">ENST00000232564.3</a>	<a href="#">GNB4</a>	<a href="#">guanine nucleotide binding protein (G protein), beta polypeptide 4</a>
<a href="#">chr22 - 36600727</a>	8.436	<a href="#">ENST00000328429.4</a> <a href="#">ENST00000397275.2</a>	<a href="#">APOL4</a> <a href="#">APOL4</a>	<a href="#">apolipoprotein L, 4</a> <a href="#">apolipoprotein L, 4</a>
<a href="#">chr17 - 74489215</a>	8.428	<a href="#">ENST00000585701.1</a> <a href="#">ENST00000591192.1</a> <a href="#">ENST00000589526.1</a>	<a href="#">RHBDF2</a> <a href="#">RHBDF2</a>	<a href="#">rhomboid 5 homolog 2 (Drosophila)</a> <a href="#">rhomboid 5 homolog 2 (Drosophila)</a>
<a href="#">chr9 + 33264861</a>	8.355	<a href="#">ENST00000223500.8</a>	<a href="#">CHMP5</a>	<a href="#">charged multivesicular body protein 5</a>
<a href="#">chr22 + 36044411</a>	8.329	<a href="#">ENST00000409652.4</a>	<a href="#">APOL6</a>	<a href="#">apolipoprotein L, 6</a>
<a href="#">chr12 + 113344755</a>	8.308	<a href="#">ENST00000550883.1</a>	<a href="#">OAS1</a>	<a href="#">2'-5'-oligoadenylate synthetase 1, 40/46kDa</a>
<a href="#">chr3 - 122283100</a>	8.227	<a href="#">ENST00000462315.1</a> <a href="#">ENST00000492382.1</a>	<a href="#">PARP9</a> <a href="#">PARP9</a>	<a href="#">poly (ADP-ribose) polymerase family, member 9</a> <a href="#">poly (ADP-ribose) polymerase family, member 9</a>
				<a href="#">charged multivesicular bod</a>

**ISMARA****Project**

Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

<a href="#">chr9 + 33265011</a>	8.208	<a href="#">ENST00000419016.2</a>	<a href="#">CHMP5</a>	<a href="#">y protein 5</a>
<a href="#">chr8 - 145060593</a>	8.144	<a href="#">ENST00000313059.5</a> <a href="#">ENST00000524918.1</a> <a href="#">ENST00000313028.7</a> <a href="#">ENST00000525773.1</a>	<a href="#">PARP10</a>  <a href="#">PARP10</a>	<a href="#">poly (ADP-ribose) polymerase family, member 10</a>  <a href="#">poly (ADP-ribose) polymerase family, member 10</a>
<a href="#">chr21 + 42733870</a>	8.122	<a href="#">ENST00000330714.3</a> <a href="#">ENST00000436410.1</a> <a href="#">ENST00000435611.1</a>	<a href="#">MX2</a>  <a href="#">MX2</a>	<a href="#">myxovirus (influenza virus) resistance 2 (mouse)</a>  <a href="#">myxovirus (influenza virus) resistance 2 (mouse)</a>
<a href="#">chr21 + 42798094</a>	8.073	<a href="#">ENST00000424365.1</a> <a href="#">ENST00000398598.3</a> <a href="#">ENST00000455164.2</a>	<a href="#">MX1</a> <a href="#">MX1</a>	<a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a> <a href="#">myxovirus (influenza virus) resistance 1, interferon-inducible protein p78 (mouse)</a>
<a href="#">chr6 - 32821599</a>	7.992	<a href="#">ENST00000354258.4</a>	<a href="#">TAP1</a>	<a href="#">transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)</a>
<a href="#">chr2 - 166702601</a>	7.951	<a href="#">ENST00000428888.1</a>	<a href="#">AC009495.4</a>	<a href="#">AC009495.4</a>
<a href="#">chr6 + 37400974</a>	7.883	<a href="#">ENST00000373451.4</a> <a href="#">ENST00000455891.1</a>	<a href="#">CMTR1</a> <a href="#">CMTR1</a>	<a href="#">cap methyltransferase 1</a> <a href="#">cap methyltransferase 1</a>
<a href="#">chr14 - 100841670</a>	7.821	<a href="#">ENST00000555410.1</a> <a href="#">ENST00000557297.1</a> <a href="#">ENST00000555813.1</a> <a href="#">ENST00000557135.1</a> <a href="#">ENST00000556698.1</a> <a href="#">ENST00000554509.1</a>	<a href="#">WARS</a> <a href="#">WARS</a>	<a href="#">tryptophanyl-tRNA synthetase</a> <a href="#">tryptophanyl-tRNA synthetase</a>
<a href="#">chr15 + 74287118</a>	7.658	<a href="#">ENST00000563500.1</a>	<a href="#">PML</a>	<a href="#">promyelocytic leukemia</a>
<a href="#">chr12 + 40618764</a>	7.654	<a href="#">ENST00000343742.2</a>	<a href="#">LRRK2</a>	<a href="#">leucine-rich repeat kinase 2</a>
				<a href="#">family with sequence simila</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr6 + 116782527</a>	7.551	<a href="#">ENST00000368606.3</a> <a href="#">ENST00000368605.1</a>	<a href="#">FAM26F</a> <a href="#">FAM26F</a>	<a href="#">rity 26, member F</a> <a href="#">family with sequence simila</a> <a href="#">rity 26, member F</a>
<a href="#">chr9 + 4985016</a>	7.498	<a href="#">ENST00000539801.1</a>	<a href="#">JAK2</a>	<a href="#">Janus kinase 2</a>
<a href="#">chr15 + 74287035</a>	7.444	<a href="#">ENST00000359928.4</a> <a href="#">ENST00000395132.2</a> <a href="#">ENST00000268059.6</a> <a href="#">ENST00000354026.6</a> <a href="#">ENST00000268058.3</a> <a href="#">ENST00000565898.1</a> <a href="#">ENST00000569477.1</a> <a href="#">ENST00000569965.1</a> <a href="#">ENST00000567543.1</a> <a href="#">ENST00000436891.3</a> <a href="#">ENST00000435786.2</a> <a href="#">ENST00000564428.1</a>	<a href="#">PML</a> <a href="#">PML</a>	<a href="#">promyelocytic leukemia</a> <a href="#">promyelocytic leukemia</a>
<a href="#">chr1 - 146644036</a>	7.404	<a href="#">ENST00000425272.2</a>	<a href="#">PRKAB2</a>	<a href="#">protein kinase, AMP-activat</a> <a href="#">ed, beta 2 non-catalytic sub</a> <a href="#">unit</a>
<a href="#">chr11 - 57335280</a>	7.139	<a href="#">ENST00000287156.4</a>	<a href="#">UBE2L6</a>	<a href="#">ubiquitin-conjugating enzy</a> <a href="#">me E2L 6</a>
<a href="#">chr19 + 17516624</a>	7.107	<a href="#">ENST00000596322.1</a> <a href="#">ENST00000600008.1</a> <a href="#">ENST00000601885.1</a>	<a href="#">CTD-2521M24.9</a>	<a href="#">CTD-2521M24.9</a>
<a href="#">chr12 + 113344582</a>	7.068	<a href="#">ENST00000452357.2</a> <a href="#">ENST00000202917.5</a> <a href="#">ENST00000445409.2</a>	<a href="#">OAS1</a> <a href="#">OAS1</a>	<a href="#">2'-5'-oligoadenylate synthet</a> <a href="#">ase 1, 40/46kDa</a> <a href="#">2'-5'-oligoadenylate synthet</a> <a href="#">ase 1, 40/46kDa</a>
<a href="#">chr15 + 74287009</a>	6.858	<a href="#">ENST00000395135.3</a>	<a href="#">PML</a>	<a href="#">promyelocytic leukemia</a>
<a href="#">chr13 + 108922228</a>	6.837	<a href="#">ENST00000542136.1</a>	<a href="#">TNFSF13B</a>	<a href="#">tumor necrosis factor (ligan</a> <a href="#">d) superfamily, member 13</a> <a href="#">b</a>
<a href="#">chr1 - 151319654</a>	6.746	<a href="#">ENST00000412774.1</a> <a href="#">ENST00000430227.1</a>	<a href="#">RFX5</a> <a href="#">RFX5</a>	<a href="#">regulatory factor X, 5 (influe</a> <a href="#">nces HLA class II expressio</a> <a href="#">n)</a> <a href="#">regulatory factor X, 5 (influe</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

				<a href="#">nces HLA class II expression)</a>
<a href="#">chr16 - 11350036</a>	6.717	<a href="#">ENST00000332029.2</a>	<a href="#">SOCS1</a>	<a href="#">suppressor of cytokine signaling 1</a>
<a href="#">chr9 - 92020841</a>	6.714	<a href="#">ENST00000433650.1</a>	<a href="#">SEMA4D</a>	<a href="#">sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D</a>
<a href="#">chr22 - 36635225</a>	6.691	<a href="#">ENST00000529194.1</a>	<a href="#">APOL2</a>	<a href="#">apolipoprotein L, 2</a>
<a href="#">chr5 + 132009675</a>	6.681	<a href="#">ENST00000350025.2</a> <a href="#">ENST00000231449.2</a>	<a href="#">IL4</a> <a href="#">IL4</a>	<a href="#">interleukin 4</a> <a href="#">interleukin 4</a>
<a href="#">chr9 - 14322319</a>	6.643	<a href="#">ENST00000606230.1</a>	<a href="#">NFIB</a>	<a href="#">nuclear factor I/B</a>
<a href="#">chr3 - 28389922</a>	6.587	<a href="#">ENST00000415852.1</a>	<a href="#">AZI2</a>	<a href="#">5-azacytidine induced 2</a>
<a href="#">chr3 - 172241250</a>	6.485	<a href="#">ENST00000241261.2</a> <a href="#">ENST00000420541.2</a>	<a href="#">TNFSF10</a> <a href="#">TNFSF10</a>	<a href="#">tumor necrosis factor (ligand) superfamily, member 10</a> <a href="#">tumor necrosis factor (ligand) superfamily, member 10</a>
<a href="#">chr12 - 56753858</a>	6.449	<a href="#">ENST00000314128.4</a> <a href="#">ENST00000557235.1</a> <a href="#">ENST00000418572.2</a>	<a href="#">STAT2</a> <a href="#">STAT2</a>	<a href="#">signal transducer and activator of transcription 2, 113kDa</a> <a href="#">signal transducer and activator of transcription 2, 113kDa</a>
<a href="#">chr6 + 126240442</a>	6.422	<a href="#">ENST00000444128.1</a> <a href="#">ENST00000448104.1</a> <a href="#">ENST00000438495.1</a>	<a href="#">NCOA7</a> <a href="#">NCOA7</a>	<a href="#">nuclear receptor coactivator 7</a> <a href="#">nuclear receptor coactivator 7</a>
<a href="#">chr2 - 55920952</a>	6.390	<a href="#">ENST00000447944.2</a>	<a href="#">PNPT1</a>	<a href="#">polyribonucleotide nucleotidyltransferase 1</a>
<a href="#">chr3 - 28390415</a>	6.265	<a href="#">ENST00000414162.1</a> <a href="#">ENST00000420543.2</a>	<a href="#">AZI2</a> <a href="#">AZI2</a>	<a href="#">5-azacytidine induced 2</a> <a href="#">5-azacytidine induced 2</a>
<a href="#">chr11 + 57308979</a>	6.187	<a href="#">ENST00000457912.1</a>	<a href="#">SMTNL1</a>	<a href="#">smoothelin-like 1</a>
				<a href="#">transporter 2, ATP-binding</a>



**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr6 - 32806483</a>	6.166	<a href="#">ENST00000374899.4</a>	<a href="#">TAP2</a>	<a href="#">cassette, sub-family B (MD R/TAP)</a>
<a href="#">chr7 - 92747269</a>	6.074	<a href="#">ENST00000379958.2</a> <a href="#">ENST00000446617.1</a>	<a href="#">SAMD9</a> <a href="#">SAMD9</a>	<a href="#">sterile alpha motif domain containing 9</a> <a href="#">sterile alpha motif domain containing 9</a>
<a href="#">chr22 + 36649056</a>	5.885	<a href="#">ENST00000397278.3</a> <a href="#">ENST00000422706.1</a> <a href="#">ENST00000426053.1</a> <a href="#">ENST00000319136.4</a>	<a href="#">APOL1</a>  <a href="#">APOL1</a>	<a href="#">apolipoprotein L 1</a>  <a href="#">apolipoprotein L 1</a>
<a href="#">chr11 - 321050</a>	5.869	<a href="#">ENST00000399808.4</a>	<a href="#">IFITM3</a>	<a href="#">interferon induced transmembrane protein 3</a>
<a href="#">chr1 + 212782012</a>	5.775	<a href="#">ENST00000341491.4</a> <a href="#">ENST00000366985.1</a>	<a href="#">ATF3</a> <a href="#">ATF3</a>	<a href="#">activating transcription factor 3</a> <a href="#">activating transcription factor 3</a>
<a href="#">chr14 - 100841930</a>	5.724	<a href="#">ENST00000553934.1</a> <a href="#">ENST00000555031.1</a> <a href="#">ENST00000553395.1</a> <a href="#">ENST00000553545.1</a> <a href="#">ENST00000344102.5</a> <a href="#">ENST00000556338.1</a> <a href="#">ENST00000392882.2</a>	<a href="#">WARS</a> <a href="#">WARS</a>	<a href="#">tryptophanyl-tRNA synthetase</a> <a href="#">tryptophanyl-tRNA synthetase</a>
<a href="#">chr12 - 121476750</a>	5.708	<a href="#">ENST00000543677.1</a>	<a href="#">OASL</a>	<a href="#">2'-5'-oligoadenylate synthetase-like</a>
<a href="#">chr14 - 69262789</a>	5.658	<a href="#">ENST00000557022.1</a>	<a href="#">ZFP36L1</a>	<a href="#">ZFP36 ring finger protein-like 1</a>
<a href="#">chr12 - 10022735</a>	5.643	<a href="#">ENST00000228438.2</a>	<a href="#">CLEC2B</a>	<a href="#">C-type lectin domain family 2, member B</a>
<a href="#">chr12 - 121476959</a>	5.633	<a href="#">ENST00000339275.5</a>	<a href="#">OASL</a>	<a href="#">2'-5'-oligoadenylate synthetase-like</a>
<a href="#">chr14 - 69262947</a>	5.607	<a href="#">ENST00000557086.1</a>	<a href="#">ZFP36L1</a>	<a href="#">ZFP36 ring finger protein-like 1</a>
				<a href="#">indoleamine 2,3-dioxygenase</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr8 + 39770803</a>	5.606	<a href="#">ENST00000518237.1</a>	<a href="#">IDO1</a>	<a href="#">se 1</a>
<a href="#">chr2 - 231084617</a>	5.601	<a href="#">ENST00000409815.2</a>	<a href="#">SP110</a>	<a href="#">SP110 nuclear body protein</a>
<a href="#">chr3 - 28390298</a>	5.538	<a href="#">ENST00000457172.1</a>	<a href="#">AZI2</a>	<a href="#">5-azacytidine induced 2</a>
<a href="#">chr22 - 36635563</a>	5.496	<a href="#">ENST00000451256.2</a>	<a href="#">APOL2</a>	<a href="#">apolipoprotein L 2</a>
<a href="#">chr1 + 158979686</a>	5.448	<a href="#">ENST00000368132.3</a> <a href="#">ENST00000295809.7</a>	<a href="#">IFI16</a> <a href="#">IFI16</a>	<a href="#">interferon, gamma-inducible protein 16</a> <a href="#">interferon, gamma-inducible protein 16</a>
<a href="#">chr1 + 158979792</a>	5.367	<a href="#">ENST00000430894.2</a> <a href="#">ENST00000359709.3</a>	<a href="#">IFI16</a> <a href="#">IFI16</a>	<a href="#">interferon, gamma-inducible protein 16</a> <a href="#">interferon, gamma-inducible protein 16</a>
<a href="#">chr12 - 121477039</a>	5.319	<a href="#">ENST00000257570.5</a>	<a href="#">OASL</a>	<a href="#">2'-5'-oligoadenylate synthetase-like</a>
<a href="#">chr13 + 108921977</a>	5.304	<a href="#">ENST00000375887.4</a> <a href="#">ENST00000430559.1</a>	<a href="#">TNFSF13B</a> <a href="#">TNFSF13B</a>	<a href="#">tumor necrosis factor (ligand) superfamily, member 13 b</a> <a href="#">tumor necrosis factor (ligand) superfamily, member 13 b</a>
<a href="#">chr19 + 17516494</a>	5.298	<a href="#">ENST00000534306.1</a>	<a href="#">CTD-2521M24.9</a>	<a href="#">CTD-2521M24.9</a>
<a href="#">chr1 + 158979680</a>	5.282	<a href="#">ENST00000340979.6</a> <a href="#">ENST00000368131.4</a>	<a href="#">IFI16</a> <a href="#">IFI16</a>	<a href="#">interferon, gamma-inducible protein 16</a> <a href="#">interferon, gamma-inducible protein 16</a>
<a href="#">chr14 - 69262916</a>	5.257	<a href="#">ENST00000553375.1</a>	<a href="#">ZFP36L1</a>	<a href="#">ZFP36 ring finger protein-like 1</a>
<a href="#">chr19 - 50400125</a>	5.245	<a href="#">ENST00000593956.1</a>	<a href="#">IL411</a>	<a href="#">interleukin 4 induced 1</a>
<a href="#">chr1 + 110453608</a>	5.239	<a href="#">ENST00000369801.1</a>	<a href="#">CSF1</a>	<a href="#">colony stimulating factor 1 (macrophage)</a>
<a href="#">chr16 + 10972818</a>	5.164	<a href="#">ENST00000576601.1</a>	<a href="#">CIITA</a>	<a href="#">class II, major histocompatibility complex, transactivator</a>
<a href="#">chr20 - 47894936</a>	5.099	<a href="#">ENST00000371754.4</a>	<a href="#">ZNF1</a>	<a href="#">zinc finger, NFX1-type containing 1</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr3_+_48507210</a>	5.028	<a href="#">ENST00000444177.1</a> <a href="#">ENST00000433541.1</a> <a href="#">ENST00000422277.2</a> <a href="#">ENST00000436480.2</a>	<a href="#">TREX1</a> <a href="#">TREX1</a>	<a href="#">three prime repair exonuclease 1</a> <a href="#">three prime repair exonuclease 1</a>
<a href="#">chr2_+_231280954</a>	4.876	<a href="#">ENST00000341950.4</a> <a href="#">ENST00000409824.1</a> <a href="#">ENST00000409341.1</a> <a href="#">ENST00000409112.1</a> <a href="#">ENST00000340126.4</a>	<a href="#">SP100</a> <a href="#">SP100</a>	<a href="#">SP100 nuclear antigen</a> <a href="#">SP100 nuclear antigen</a>
<a href="#">chr4_+_87928140</a>	4.852	<a href="#">ENST00000307808.6</a>	<a href="#">AFF1</a>	<a href="#">AF4/FMR2 family member 1</a>
<a href="#">chr1_+_113217073</a>	4.838	<a href="#">ENST00000369645.1</a>	<a href="#">MOV10</a>	<a href="#">Mov10, Moloney leukemia virus 10, homolog (mouse)</a>
<a href="#">chr2_+_231280908</a>	4.828	<a href="#">ENST00000432979.1</a> <a href="#">ENST00000427101.2</a>	<a href="#">SP100</a> <a href="#">SP100</a>	<a href="#">SP100 nuclear antigen</a> <a href="#">SP100 nuclear antigen</a>
<a href="#">chr14_-_67955426</a>	4.828	<a href="#">ENST00000554480.1</a>	<a href="#">TMEM229B</a>	<a href="#">transmembrane protein 229 B</a>
<a href="#">chr17_-_54991369</a>	4.825	<a href="#">ENST00000537230.1</a>	<a href="#">TRIM25</a>	<a href="#">tripartite motif containing 25</a>
<a href="#">chr11_-_104916034</a>	4.811	<a href="#">ENST00000375704.3</a> <a href="#">ENST00000528513.1</a> <a href="#">ENST00000375706.2</a>	<a href="#">CARD16</a> <a href="#">CARD16</a>	<a href="#">caspase recruitment domain family, member 16</a> <a href="#">caspase recruitment domain family, member 16</a>
<a href="#">chr1_+_113217345</a>	4.802	<a href="#">ENST00000357443.2</a>	<a href="#">MOV10</a>	<a href="#">Mov10, Moloney leukemia virus 10, homolog (mouse)</a>
<a href="#">chr6_-_32806506</a>	4.799	<a href="#">ENST00000374897.2</a> <a href="#">ENST00000452392.2</a>	<a href="#">TAP2</a> <a href="#">TAP2</a>	<a href="#">transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)</a> <a href="#">Uncharacterized protein</a>
<a href="#">chr1_+_113217043</a>	4.774	<a href="#">ENST00000413052.2</a>	<a href="#">MOV10</a>	<a href="#">Mov10, Moloney leukemia virus 10, homolog (mouse)</a>
<a href="#">chr14_+_35451880</a>	4.773	<a href="#">ENST00000555746.1</a> <a href="#">ENST00000554803.1</a>	<a href="#">SRP54</a> <a href="#">SRP54</a>	<a href="#">signal recognition particle 54kDa</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

				<a href="#">signal recognition particle 5 4kDa</a>
<a href="#">chr7_+_18535321</a>	4.741	<a href="#">ENST00000413380.1</a> <a href="#">ENST00000430454.1</a>	<a href="#">HDAC9</a> <a href="#">HDAC9</a>	<a href="#">histone deacetylase 9</a> <a href="#">histone deacetylase 9</a>
<a href="#">chr1_-_150693318</a>	4.665	<a href="#">ENST00000442853.1</a> <a href="#">ENST00000368995.4</a> <a href="#">ENST00000368993.2</a> <a href="#">ENST00000361824.2</a> <a href="#">ENST00000322343.7</a>	<a href="#">HORMAD1</a>  <a href="#">HORMAD1</a>	<a href="#">HORMA domain containing 1</a>  <a href="#">HORMA domain containing 1</a>
<a href="#">chr2_+_85811525</a>	4.651	<a href="#">ENST00000306384.4</a>	<a href="#">VAMP5</a>	<a href="#">vesicle-associated membrane protein 5</a>
<a href="#">chr15_+_89182156</a>	4.649	<a href="#">ENST00000379224.5</a>	<a href="#">ISG20</a>	<a href="#">interferon stimulated exonuclease gene 20kDa</a>
<a href="#">chr6_+_32821924</a>	4.619	<a href="#">ENST00000374859.2</a> <a href="#">ENST00000453265.2</a>	<a href="#">PSMB9</a> <a href="#">PSMB9</a>	<a href="#">proteasome (prosome, macrocatalytic) subunit, beta type, 9</a> <a href="#">proteasome (prosome, macrocatalytic) subunit, beta type, 9</a>
<a href="#">chr3_-_146262352</a>	4.556	<a href="#">ENST00000462666.1</a>	<a href="#">PLSCR1</a>	<a href="#">phospholipid scramblase 1</a>
<a href="#">chr6_+_25963020</a>	4.518	<a href="#">ENST00000357085.3</a>	<a href="#">TRIM38</a>	<a href="#">tripartite motif containing 38</a>
<a href="#">chrX_+_37639302</a>	4.490	<a href="#">ENST00000545017.1</a> <a href="#">ENST00000536160.1</a>	<a href="#">CYBB</a> <a href="#">CYBB</a>	<a href="#">cytochrome b-245, beta polypeptide</a> <a href="#">cytochrome b-245, beta polypeptide</a>
<a href="#">chr3_+_122283064</a>	4.466	<a href="#">ENST00000296161.4</a>	<a href="#">DTX3L</a>	<a href="#">deltex 3-like (Drosophila)</a>
<a href="#">chr3_-_146262428</a>	4.422	<a href="#">ENST00000486631.1</a>	<a href="#">PLSCR1</a>	<a href="#">phospholipid scramblase 1</a>
<a href="#">chr19_-_17516449</a>	4.420	<a href="#">ENST00000252593.6</a>	<a href="#">BST2</a>	<a href="#">bone marrow stromal cell antigen 2</a>
<a href="#">chr9_+_26746951</a>	4.411	<a href="#">ENST00000523363.1</a>	<a href="#">RP11-18A15.1</a>	<a href="#">RP11-18A15.1</a>
<a href="#">chr1_+_113217309</a>	4.367	<a href="#">ENST00000369644.1</a> <a href="#">ENST00000544796.1</a>	<a href="#">MOV10</a> <a href="#">MOV10</a>	<a href="#">Mov10, Moloney leukemia virus 10, homolog (mouse)</a> <a href="#">Mov10, Moloney leukemia virus 10, homolog (mouse)</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr12_-_118498911</a>	4.337	<a href="#">ENST00000544233.1</a>	<a href="#">WSB2</a>	<a href="#">WD repeat and SOCS box containing 2</a>
<a href="#">chr4_+_100737954</a>	4.286	<a href="#">ENST00000296414.7</a> <a href="#">ENST00000512369.1</a>	<a href="#">DAPP1</a> <a href="#">DAPP1</a>	<a href="#">dual adaptor of phosphotyrosine and 3-phosphoinositides</a> <a href="#">dual adaptor of phosphotyrosine and 3-phosphoinositides</a>
<a href="#">chr3_-_28390581</a>	4.271	<a href="#">ENST00000479665.1</a>	<a href="#">AZI2</a>	<a href="#">5-azacytidine induced 2</a>
<a href="#">chr11_-_104905840</a>	4.230	<a href="#">ENST00000436863.3</a> <a href="#">ENST00000526568.1</a> <a href="#">ENST00000393136.4</a> <a href="#">ENST00000531166.1</a> <a href="#">ENST00000534497.1</a> <a href="#">ENST00000527979.1</a> <a href="#">ENST00000446369.1</a> <a href="#">ENST00000353247.5</a> <a href="#">ENST00000528974.1</a> <a href="#">ENST00000533400.1</a> <a href="#">ENST00000525825.1</a>	<a href="#">CASP1</a> <a href="#">CASP1</a>	<a href="#">caspase 1, apoptosis-related cysteine peptidase</a> <a href="#">caspase 1, apoptosis-related cysteine peptidase</a>
<a href="#">chr8_+_27184320</a>	4.218	<a href="#">ENST00000522517.1</a>	<a href="#">PTK2B</a>	<a href="#">protein tyrosine kinase 2 beta</a>
<a href="#">chr2_-_166651152</a>	4.198	<a href="#">ENST00000412248.1</a> <a href="#">ENST00000431484.1</a>	<a href="#">GALNT3</a> <a href="#">GALNT3</a>	<a href="#">UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 3 (GalNAc-T3)</a> <a href="#">UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 3 (GalNAc-T3)</a>
<a href="#">chr11_-_104972158</a>	4.141	<a href="#">ENST00000598974.1</a> <a href="#">ENST00000593315.1</a> <a href="#">ENST00000594519.1</a> <a href="#">ENST00000415981.2</a>	<a href="#">CASP1</a>	<a href="#">caspase 1, apoptosis-related cysteine peptidase</a>  <a href="#">caspase recruitment domain</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

		<a href="#">ENST00000525374.1</a> <a href="#">ENST00000375707.1</a>	<a href="#">CARD16</a> <a href="#">CARD17</a>	<a href="#">n family, member 16</a> <a href="#">caspase recruitment domain family, member 17</a>
<a href="#">chr8_+_144640499</a>	4.128	<a href="#">ENST00000534018.1</a> <a href="#">ENST00000525721.1</a>	<a href="#">GSDMD</a> <a href="#">GSDMD</a>	<a href="#">gasdermin D</a> <a href="#">gasdermin D</a>
<a href="#">chr3_-_28390120</a>	4.128	<a href="#">ENST00000334100.6</a>	<a href="#">AZI2</a>	<a href="#">5-azacytidine induced 2</a>
<a href="#">chr19_+_17516909</a>	4.108	<a href="#">ENST00000600514.1</a> <a href="#">ENST00000601007.1</a> <a href="#">ENST00000594913.1</a> <a href="#">ENST00000599975.1</a>	<a href="#">MVB12A</a> <a href="#">CTD-2521M24.9</a>	<a href="#">multivesicular body subunit 12A</a> <a href="#">CTD-2521M24.9</a>
<a href="#">chr15_+_89182178</a>	4.101	<a href="#">ENST00000559876.1</a>	<a href="#">ISG20</a>	<a href="#">interferon stimulated exonuclease gene 20kDa</a>
<a href="#">chr10_+_92980517</a>	4.077	<a href="#">ENST00000336126.5</a>	<a href="#">PCGF5</a>	<a href="#">polycomb group ring finger 5</a>
<a href="#">chr1_+_110453462</a>	4.041	<a href="#">ENST00000488198.1</a>	<a href="#">CSF1</a>	<a href="#">colony stimulating factor 1 (macrophage)</a>
<a href="#">chr10_-_9801179</a>	4.023	<a href="#">ENST00000419836.1</a>	<a href="#">RP5-1051H14.2</a>	<a href="#">RP5-1051H14.2</a>
<a href="#">chr9_-_80437915</a>	4.005	<a href="#">ENST00000397476.3</a>	<a href="#">GNAQ</a>	<a href="#">guanine nucleotide binding protein (G protein), q polypeptide</a>
<a href="#">chr2_-_231084659</a>	3.978	<a href="#">ENST00000392048.3</a> <a href="#">ENST00000258381.6</a> <a href="#">ENST00000358662.4</a> <a href="#">ENST00000455674.1</a>	<a href="#">SP110</a> <a href="#">SP110</a>	<a href="#">SP110 nuclear body protein</a> <a href="#">SP110 nuclear body protein</a>
<a href="#">chr11_-_4629367</a>	3.972	<a href="#">ENST00000533021.1</a>	<a href="#">TRIM68</a>	<a href="#">tripartite motif containing 68</a>
<a href="#">chr3_-_146262637</a>	3.966	<a href="#">ENST00000472349.1</a> <a href="#">ENST00000342435.4</a>	<a href="#">PLSCR1</a> <a href="#">PLSCR1</a>	<a href="#">phospholipid scramblase 1</a> <a href="#">phospholipid scramblase 1</a>
<a href="#">chr5_-_39203093</a>	3.946	<a href="#">ENST00000515010.1</a>	<a href="#">FYB</a>	<a href="#">FYN binding protein</a>
<a href="#">chr13_+_50070491</a>	3.943	<a href="#">ENST00000442195.1</a> <a href="#">ENST00000496612.1</a> <a href="#">ENST00000357596.3</a> <a href="#">ENST00000485919.1</a>	<a href="#">PHF11</a> <a href="#">PHF11</a>	<a href="#">PHD finger protein 11</a> <a href="#">PHD finger protein 11</a>
<a href="#">chr3_-_146262488</a>	3.918	<a href="#">ENST00000487389.1</a>	<a href="#">PLSCR1</a>	<a href="#">phospholipid scramblase 1</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

<a href="#">chr7 + 134832808</a>	3.895	<a href="#">ENST00000275767.3</a>	<a href="#">TMEM140</a>	<a href="#">transmembrane protein 140</a>
<a href="#">chr6 - 109702885</a>	3.878	<a href="#">ENST00000504373.1</a>	<a href="#">CD164</a>	<a href="#">CD164 molecule, sialomucin</a>
<a href="#">chr2 - 145275828</a>	3.859	<a href="#">ENST00000409211.1</a> <a href="#">ENST00000392861.2</a>	<a href="#">ZEB2</a> <a href="#">ZEB2</a>	<a href="#">zinc finger E-box binding homeobox 2</a> <a href="#">zinc finger E-box binding homeobox 2</a>
<a href="#">chrX - 30595959</a>	3.859	<a href="#">ENST00000378962.3</a>	<a href="#">CXorf21</a>	<a href="#">chromosome X open reading frame 21</a>
<a href="#">chr2 - 231084820</a>	3.846	<a href="#">ENST00000338556.3</a> <a href="#">ENST00000258382.5</a>	<a href="#">SP110</a> <a href="#">SP110</a>	<a href="#">SP110 nuclear body protein</a> <a href="#">SP110 nuclear body protein</a>
<a href="#">chr3 - 146262365</a>	3.845	<a href="#">ENST00000448787.2</a>	<a href="#">PLSCR1</a>	<a href="#">phospholipid scramblase 1</a>
<a href="#">chr6 + 32811885</a>	3.788	<a href="#">ENST00000458296.1</a> <a href="#">ENST00000413039.1</a> <a href="#">ENST00000429600.1</a> <a href="#">ENST00000412095.1</a> <a href="#">ENST00000415067.1</a> <a href="#">ENST00000395330.1</a>	<a href="#">TAPSAR1</a>  <a href="#">TAPSAR1</a> <a href="#">PSMB9</a>	<a href="#">TAP1 and PSMB8 antisense RNA 1</a>  <a href="#">TAP1 and PSMB8 antisense RNA 1</a> <a href="#">proteasome (prosome, macropain) subunit, beta type, 9</a>
<a href="#">chr5 - 95158375</a>	3.785	<a href="#">ENST00000512469.2</a> <a href="#">ENST00000379979.4</a> <a href="#">ENST00000505427.1</a> <a href="#">ENST00000508780.1</a>	<a href="#">GLRX</a>  <a href="#">GLRX</a>	<a href="#">glutaredoxin (thioltransferase)</a>  <a href="#">glutaredoxin (thioltransferase)</a>
<a href="#">chr16 + 57023406</a>	3.762	<a href="#">ENST00000436936.1</a> <a href="#">ENST00000262510.6</a> <a href="#">ENST00000308149.7</a>	<a href="#">NLRC5</a> <a href="#">NLRC5</a>	<a href="#">NLR family, CARD domain containing 5</a> <a href="#">NLR family, CARD domain containing 5</a>
<a href="#">chr2 + 33701707</a>	3.758	<a href="#">ENST00000423159.1</a> <a href="#">ENST00000425210.1</a>	<a href="#">RASGRP3</a> <a href="#">RASGRP3</a>	<a href="#">RAS guanyl releasing protein 3 (calcium and DAG-regulated)</a> <a href="#">RAS guanyl releasing prote</a>

**ISMARA****Project**

Final-1

**Navigation**

All motifs sorted by activity significance

All samples sorted alphabetically

Mean activities

**Downloads**

Activities table

Activity deltas table

Regulatory interactions

Motifs sorted by significance (text)

Download the whole report

		<a href="#">ENST00000444784.1</a>		<a href="#">in 3 (calcium and DAG-regulated)</a>
<a href="#">chr19 - 50400212</a>	3.749	<a href="#">ENST00000391826.2</a>	<a href="#">IL411</a>	<a href="#">interleukin 4 induced 1</a>
<a href="#">chr19 + 17516531</a>	3.701	<a href="#">ENST00000528911.1</a> <a href="#">ENST00000528604.1</a> <a href="#">ENST00000595892.1</a> <a href="#">ENST00000500836.2</a> <a href="#">ENST00000598546.1</a> <a href="#">ENST00000600369.1</a> <a href="#">ENST00000598356.1</a> <a href="#">ENST00000594426.1</a>	<a href="#">MVB12A</a>  <a href="#">CTD-2521M24.9</a>	<a href="#">multivesicular body subunit 12A</a>  <a href="#">CTD-2521M24.9</a>
<a href="#">chr6 - 36355486</a>	3.624	<a href="#">ENST00000538992.1</a>	<a href="#">ETV7</a>	<a href="#">ets variant 7</a>
<a href="#">chr13 + 50070077</a>	3.601	<a href="#">ENST00000426879.1</a> <a href="#">ENST00000378319.3</a>	<a href="#">PHF11</a> <a href="#">PHF11</a>	<a href="#">PHD finger protein 11</a> <a href="#">PHD finger protein 11</a>
<a href="#">chr14 - 67981870</a>	3.573	<a href="#">ENST00000555994.1</a>	<a href="#">TMEM229B</a>	<a href="#">transmembrane protein 229 B</a>
<a href="#">chr6 - 36355513</a>	3.569	<a href="#">ENST00000373737.4</a> <a href="#">ENST00000340181.4</a>	<a href="#">ETV7</a> <a href="#">ETV7</a>	<a href="#">ets variant 7</a> <a href="#">ets variant 7</a>
<a href="#">chr15 + 89181974</a>	3.557	<a href="#">ENST00000306072.5</a>	<a href="#">ISG20</a>	<a href="#">interferon stimulated exonu lease gene 20kDa</a>
<a href="#">chr7 + 150382781</a>	3.555	<a href="#">ENST00000474605.1</a> <a href="#">ENST00000223293.5</a>	<a href="#">GIMAP2</a> <a href="#">GIMAP2</a>	<a href="#">GTPase, IMAP family mem ber 2</a> <a href="#">GTPase, IMAP family mem ber 2</a>

**Gene Ontology Analysis**

Activity graph

Z-values graph

Top targets

GO analysis

Back to top

**Gene overrepresentation in biological\_process category:**

Search:

Show  entries

Showing 1 to 20 of 209 entries



**ISMARA****Project**

Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)

Log-likelihood per target	Total log-likelihood	Term	Description
5.3	628.3	<a href="#">GO:0071357</a>	type I interferon signaling pathway(GO:0071357)
1.1	131.0	<a href="#">GO:0051607</a>	defense response to virus(GO:0051607)
0.1	117.2	<a href="#">GO:0006955</a>	immune response(GO:0006955)
1.1	73.2	<a href="#">GO:0032480</a>	negative regulation of type I interferon signaling pathway(GO:0032480)
1.1	51.0	<a href="#">GO:0042130</a>	negative regulation of T cell proliferation(GO:0042130)
11.0	44.1	<a href="#">GO:0051097</a>	negative regulation of helicase activity(GO:0051097)
0.1	38.2	<a href="#">GO:0007249</a>	I-kappaB kinase/NF-kappaB signaling(GO:0007249)
9.5	38.1	<a href="#">GO:0009138</a>	pyrimidine nucleoside diphosphate metabolism(GO:0009138)
0.2	36.8	<a href="#">GO:0042787</a>	protein ubiquitination involved in ubiquitin-mediated proteolysis(GO:0042787)
2.4	29.2	<a href="#">GO:0002230</a>	positive regulation of defense response(GO:0002230)
0.2	29.0	<a href="#">GO:0009615</a>	response to virus(GO:0009615)
0.6	26.3	<a href="#">GO:0060135</a>	maternal process involved in female pronuclear migration(GO:0060135)
4.1	24.3	<a href="#">GO:0006436</a>	tryptophanyl-tRNA aminoacylation(GO:0006436)
1.9	24.2	<a href="#">GO:0017121</a>	phospholipid scrambling(GO:0017121)
7.3	22.0	<a href="#">GO:0032938</a>	negative regulation of translation in response to stress(GO:0032938)
0.3	20.8	<a href="#">GO:0060333</a>	interferon-gamma-mediated signaling pathway(GO:0060333)
0.1	20.4	<a href="#">GO:0007281</a>	germ cell development(GO:0007281)
0.5	19.1	<a href="#">GO:0070206</a>	protein trimerization(GO:0070206)
4.7	19.0	<a href="#">GO:0046967</a>	cytosol to ER transport(GO:0046967)
4.7	18.8	<a href="#">GO:0035279</a>	mRNA cleavage involved in gene silencing by microRNAs(GO:0035279)

**Gene overrepresentation in cellular\_component category:**Search: Show  entries

Showing 1 to 20 of 86 entries

Log-likelihood per target	Total log-likelihood	Term	Description
0.1	289.0	<a href="#">GO:0005829</a>	cytosol(GO:0005829)
0.0	239.2	<a href="#">GO:0005575</a>	cellular_component(GO:0005575)
0.0	168.0	<a href="#">GO:0005737</a>	cytoplasm(GO:0005737)

[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

**ISMARA****Project**

Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)

0.4	162.1	<a href="#">GO:0005792</a>	obsolete microsome(GO:0005792)
0.1	152.8	<a href="#">GO:0005739</a>	mitochondrion(GO:0005739)
0.2	135.8	<a href="#">GO:0005789</a>	endoplasmic reticulum membrane(GO:
0.1	58.5	<a href="#">GO:0010008</a>	endosome membrane(GO:0010008)
0.1	53.0	<a href="#">GO:0048471</a>	perinuclear region of cytoplasm(GO:00
0.0	52.7	<a href="#">GO:0005730</a>	nucleolus(GO:0005730)
0.0	27.8	<a href="#">GO:0005615</a>	extracellular space(GO:0005615)
2.6	23.8	<a href="#">GO:0042825</a>	TAP complex(GO:0042825)
0.5	22.3	<a href="#">GO:0012505</a>	endomembrane system(GO:0012505)
7.3	22.0	<a href="#">GO:0042406</a>	extrinsic component of endoplasmic re
0.2	21.7	<a href="#">GO:0016605</a>	PML body(GO:0016605)
1.7	21.5	<a href="#">GO:0033391</a>	chromatoid body(GO:0033391)
0.3	19.8	<a href="#">GO:0015030</a>	Cajal body(GO:0015030)
0.3	19.2	<a href="#">GO:0000932</a>	cytoplasmic mRNA processing body(G
0.1	16.6	<a href="#">GO:0016363</a>	nuclear matrix(GO:0016363)
0.1	16.5	<a href="#">GO:0009897</a>	external side of plasma membrane(GO
1.6	14.3	<a href="#">GO:0008074</a>	guanylate cyclase complex, soluble(GC

**Gene overrepresentation in molecular\_function category:**Search: Show  entries

Showing 1 to 20 of 182 entries

Log-likelihood per target	Total log-likelihood	Term	Description
0.9	163.3	<a href="#">GO:0016779</a>	nucleotidyltransferase activity(GO:0016779)
0.6	134.1	<a href="#">GO:0004386</a>	helicase activity(GO:0004386)
1.5	122.4	<a href="#">GO:0051536</a>	iron-sulfur cluster binding(GO:0051536)
0.2	97.4	<a href="#">GO:0003924</a>	GTPase activity(GO:0003924)
0.0	92.1	<a href="#">GO:0008270</a>	zinc ion binding(GO:0008270)
0.4	79.5	<a href="#">GO:0016881</a>	acid-amino acid ligase activity(GO:0016881)
1.6	73.3	<a href="#">GO:0003950</a>	NAD+ ADP-ribosyltransferase activity(GO:0003950)

[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

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Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)

0.6	40.8	<a href="#">GO:0036459</a>	thiol-dependent ubiquitin-specific prote
13.4	40.1	<a href="#">GO:0031386</a>	protein tag(GO:0031386)
5.4	38.1	<a href="#">GO:0004127</a>	cytidylate kinase activity(GO:0004127)
0.3	31.2	<a href="#">GO:0003704</a>	obsolete specific RNA polymerase II tra
0.1	29.6	<a href="#">GO:0043565</a>	sequence-specific DNA binding(GO:00
5.7	28.6	<a href="#">GO:0003726</a>	double-stranded RNA adenosine deam
1.4	25.6	<a href="#">GO:0008035</a>	high-density lipoprotein particle binding
0.1	24.7	<a href="#">GO:0003714</a>	transcription corepressor activity(GO:00
4.1	24.3	<a href="#">GO:0004830</a>	tryptophan-tRNA ligase activity(GO:000
1.9	24.2	<a href="#">GO:0017128</a>	phospholipid scramblase activity(GO:00
0.0	23.5	<a href="#">GO:0004872</a>	receptor activity(GO:0004872) molecu
0.5	22.5	<a href="#">GO:0008603</a>	cAMP-dependent protein kinase regula
5.5	22.0	<a href="#">GO:0032183</a>	SUMO binding(GO:0032183)

**Gene overrepresentation in C2:CP category:**Search: Show  entries

Showing 1 to 18 of 18 entries

Log-likelihood per target	Total log-likelihood	Term
2.4	28.3	<a href="#">ST_INTERFERON_GAMMA_PATHWAY</a>
0.1	23.3	<a href="#">SIG_PIP3_SIGNALING_IN_CARDIAC_MYOCYTES</a>
0.6	21.6	<a href="#">SA_MMP_CYTOKINE_CONNECTION</a>
0.4	11.8	<a href="#">SIG_PIP3_SIGNALING_IN_B_LYMPHOCYTES</a>
0.3	10.5	<a href="#">SA_FAS_SIGNALING</a>
0.4	7.0	<a href="#">SIG_IL4RECEPTOR_IN_B_LYMPHOCYTES</a>
0.3	6.7	<a href="#">ST_INTERLEUKIN_4_PATHWAY</a>
0.2	5.7	<a href="#">SA_CASPASE_CASCADE</a>
0.7	5.2	<a href="#">ST_TYPE_I_INTERFERON_PATHWAY</a>
0.3	4.4	<a href="#">SA_PTEN_PATHWAY</a>
0.3	3.4	<a href="#">ST_PAC1_RECEPTOR_PATHWAY</a>

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Final-1

**Navigation**[All motifs sorted by activity significance](#)[All samples sorted alphabetically](#)[Mean activities](#)**Downloads**[Activities table](#)[Activity deltas table](#)[Regulatory interactions](#)[Motifs sorted by significance \(text\)](#)[Download the whole report](#)

0.1	2.5	<a href="#">ST_ADRENERGIC</a>
0.0	1.8	<a href="#">SIG_REGULATION_OF_THE_ACTIN_CYTOSKELET</a>
0.0	1.6	<a href="#">ST_WNT_BETA_CATENIN_PATHWAY</a>
0.0	0.8	<a href="#">ST_P38_MAPK_PATHWAY</a>
0.1	0.7	<a href="#">ST_JAK_STAT_PATHWAY</a>
0.0	0.6	<a href="#">SIG_INSULIN_RECEPTOR_PATHWAY_IN_CARDIAC</a>
0.0	0.3	<a href="#">SA_PROGRAMMED_CELL_DEATH</a>



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[Activity graph](#)[Z-values graph](#)[Top targets](#)[GO analysis](#)[Back to top](#)