TABLE S1 Regulatory factors/conditions affecting expression of SPI1

Regulator	Description	Mechanism of action/comments	References
HilA	Transcriptional activator, OmpR/ToxR family	Direct activation of prg/org and inv/spa operons	(8; 23; 27; 56; 57)
HilD	AraC-like transcriptional activator	Direct activation of hilA, hilD, hilC, and rtsA	(29; 88)
HilC	AraC-like transcriptional activator	Direct activation of hilA, hilD, hilC, and rtsA	(28; 29; 46; 88)
RtsA	AraC-like transcriptional activator	Direct activation of hilA, hilD, hilC, and rtsA	(29; 30)
HilE		Repression of <i>hilA</i> by binding to and preventing HilD function	(11); Chubiz JE (unpublished)
FliZ	Enhancer of class II flagellar genes expression	Activation of <i>hilA</i> via post- translational regulation of HilD	(20; 45; 54; 59; 85; 102)
EnvZ/OmpR	Two-component regulatory system; regulation of outer membrane porin genes, and virulence	Activation of <i>hilA</i> via HilD	(29; 58)
FadD	Acyl-CoA synthetase; degradation of long-chain fatty acids	Activation of <i>hilA</i> via an unknown mechanism	(59);Ellermeier JR (unpublished)
SirA	Transcriptional regulator; two-component regulatory system BarA/SirA; regulation of carbohydrate metabolism, motility, biofilm formation, and invasion	Activation of <i>hilA</i> via activation of <i>csrB/csrC</i> to block CsrA repression of <i>hilD</i>	(1; 29; 46; 52; 62; 99; 102);Ellermeier JR (unpublished)
Dam	DNA methylase	Activation of <i>hilA</i> via post- transcriptional regulation of <i>hilD</i>	(55)
Ack Pta	acetate kinase and phosphotransacetylase	Activation of <i>hilA</i> and <i>hilD</i> by formate via an unknown mechanism	(43)

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Regulator	Description	Mechanism of action/comments	References
YfgL	Outer membrane lipoprotein; assembly of the outer membrane β- barrel proteins in complex with YaeT, YfiO, and NIpB	Activation of SPI1genes via an unknown mechanism	(2; 37)
Trk (potassium)	Potassium transporter	Activation of SPI1 genes via an unknown mechanism	(94)
Fnr	Transcriptional regulator; cytoplasmic oxygen sensor	Repression of hilA via an unknown mechanism	(102);Ellermeier JR (unpublished)
PhoPQ (PhoQ24)	Two-component regulatory system; response to divalent cation limitation, pH and antimicrobial peptides	Repression of <i>hilA</i> , most likely direct	(9; 10; 14; 78; 111);Ellermeier JR (unpublished)
H-NS	Nucleoid protein	Direct repression of hilA, hilC, hilD and rtsA	(73; 74; 89)
Hha	Nucleoid protein	Direct repression of hilA, hilC, hilD and rtsA	(36; 73; 74; 102)
Fis	Nucleoid protein	Activation of SPI1 genes	(22; 48; 89; 110)}
HU	Nucleoid protein	Activation of SPI1 genes	(60; 89)
RfaH	Transcriptional anti- terminator; long operons for LPS core and O-antigen biosynthesis	Activation of SPI1genes via an unknown mechanism	(60; 67)
Fur	Transcriptional regulator; response to iron	Activation of hilA via an unknown regulation of HilD; repression of H-NS; direct binding of Fur to hilD promoter	(32; 98; 101)
TdcA	Transcriptional regulator of <i>tdc</i> operon; transport and metabolism of L-	Activation of <i>fliZ</i> and SPI1 genes expression	(50)

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Regulator	Description	Mechanism of action/comments	References
	threonine and L-serine		
FlhDC	Transcriptional regulator; activation of class II flagellar genes	Activation of <i>hilA</i> via activation of <i>fliZ</i>	(20; 54)
DsbA	Periplasmic disulfide bond oxidase	Activation of <i>hilA</i> via activation of FliZ, including repression of RcsCDB	(31; 54)
RcsCDB	phosphorelay system: sensor RcsC, response regulator RcsB, and phosphotransfer protein RcsD; regulation of capsule synthesis and biofilm formation	Repression of <i>hilA</i> via repression of FliZ and an independent unknown regulation of HilD	(54)
Lon	ATP-dependent protease	Repression of <i>hilA</i> via degradation of HilD, HilC, and FliZ	(16; 20; 95; 97)
ClpXP	ATP-dependent protease	Repression of SPI1 genes via FliZ	(47)
Formate	Short chain fatty acid	Activation of <i>hilA</i> and <i>hilD</i> ( see Ack Pta)	(43)
Dimethyl sulfide/ DMSO		Repression of SPI1 genes via an unknown mechanism	(4)
Temperature		Activation of SPI1 genes when shifted from 25° to 37°C in H-NS- dependent manner	(75)
Butyrate	Short chain fatty acid	Repression of SPI1 genes via an unknown mechanism	(41)
ррGрр	Small signaling molecule; stringent response during starvation	Activation of <i>hilA</i> via an unknown mechanism	(79; 92; 93; 100);Ellermeier JR (unpublished)

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Regulator	Description	Mechanism of action/comments	References
FimZY	Transcriptional regulators; control of type 1 fimbriae gene expression	Repression of <i>hilA</i> via activation of <i>hilE</i>	(12; 87)
FimW	negative regulator of type 1 fimbriae	Activation of SPI1 genes via an unknown mechanism	(38)
Mlc	Transcriptional regulator; regulation of sugar uptake and metabolism	Repression of <i>hilA</i> via activation of <i>hilE</i>	(53)
CRP	cAMP-receptor protein	Activation of invasion via an unknown mechanism	(17)
СрхА	Sensor kinase, two- component regulatory system CpxRA; periplasmic stress responce	Activation of <i>hilA</i> via an unknown mechanism, apparently independent of CpxR	(68)
Lrp	Transcriptional regulator	Overproduction of Lrp represses SPI1 genes via an unknown mechanism	(6)
PmrM	Part of the <i>pmrHFIJKLM</i> operon	Activation of <i>hilA</i> via an unknown mechanism	(61; 65; 102)
ApaH/YgdP	Dinucleoside polyphosphate hydrolases	Activation of invasion via an unknown mechanism	(44)
PreAB (QseBC)	Two-component regulatory system; regulation of motility and virulence in response to quorum-sensing and hormonal signals	Activation of SPI1 genes via an unknown mechanism	(64; 66)
LuxS	Autoinducer 2 synthase	Activation of <i>invF</i>	(19)
PhoBR	Two-component regulatory system; phosphate limitation	Repression of <i>hilA</i> via PhoBR	(59)

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Regulator	Description	Mechanism of action/comments	References
SprB	Transcriptional regulator	Repression of <i>hilA</i> via repression of <i>hilD</i> transcription	(86)
RamA	AraC/XyIS family transcriptional activator; regulation of multidrug resistance	Repression of SPI1 genes via an unknown mechanism	(7)
РРК	Polyphosphate kinase	Activation of invasion via an unknown mechanism	(49)
PNPase	Polynucleotide phosphorylase	Repression of SPI1 genes via an unknown mechanism	(21)
Hfq	RNA shaperone	Activation of SPI1 genes	(3; 90; 91)
SmpB	RNA-binding protein	Activation of SPI1 genes	(3)
IHF	Nucleoid protein	Activation of <i>hilA</i> ; counteraction of H-NS mediated silencing	(35; 83)
RNAseE	5 -end-dependent endoribonuclease; part of degradosome complex	Repression of SPI1genes; exact mechanism unclear	(35)
Pag		Repression of SPI1 genes via an unknown mechanism	(35)
SirB		Activation of <i>hilA</i> ; exact mechanism unclear	(84)
Sig32	Sigma factor; heat shock responce	Repression of <i>hilA</i> via degradation of HilD by Lon protease	(63)
TolC/AcrAB	Multidrug efflux pump	Activation of SPI1 genes via an unknown mechanism	(15; 103; 107)
AsmA	Outer membrane protein	Required for invasion; unknown mechanism	(80)
CorA	Mg <sup>2+</sup> channel	Activation of SPI1 genes via an unknown mechanism	(76; 77)

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Regulator	Description	Mechanism of action/comments	References
PoxA	paralog of lysyl tRNA- synthetase	Repression of SPI1 genes via an unknown mechanism	(70)
YjeK	putative 2,3-β-lysine aminomutase	Repression of SPI1 genes via an unknown mechanism	(70)
Antimicrobial peptides		Repression of SPI1 genes via PhoPQ	(5)
Macrophages		Repression of SPI1 genes	(33; 96)
Epithelial cells		Activation of SPI1 genes	(42)
Bile	Role in lipid digestion	Repression of <i>hilA</i> via SirA	(81; 82)
Propanediol	Product of decomposition of rhamnose and fucose	Repression of <i>hilA</i> via an unknown mechanism	(69)
Microgravity	low-shear modeled microgravity	Repression of SPI1 genes	(109)
Lactobacillus supernatant; probiotics		Repression of SPI1 genes via an unknown mechanism	(25); (13)
Tetracycline		Activation of SPI1 genes	(108)
Nalidixic acid		Repression of SPI1 genes	(26)
Salicylidene acylhydrasides		Repression of SPI1 genes	(71)
Fluoroquinolone resistance		Repression of SPI1 genes	(34)