

Additional file 3.

Candidate genes with potential impact for virulence in pathogenic erwinias encoded in the genomes of *E. pyrifoliae*, *E. billingiae* and *E. tasmaniensis*. If no gene name could be assigned, the corresponding locus tags from the *E. pyrifoliae* are provided. Names of the genes encoding proteins with the potential to act as effectors made bold.

Secretion systems and effectors					
Functional system	Product	Gene name	Ep1/96	Et1/99	Eb661
hrp/hsv/dsp cluster (T3SS)	Type III effector	hrpK	+	-	-
	Hrp-associated systemic virulence proteins	<i>hsvA</i>	+	-	-
		<i>hsvB</i>	+	-	-
	Hrp/hrc secretion/translocation pathway proteins	<i>hsvC</i>	+	-	-
		<i>hrcU</i>	+	+	-
		<i>hrcT</i>	+	+	-
		<i>hrcS</i>	+	+	-
		<i>hrcR</i>	+	+	-
		<i>hrcQ</i>	+	+	-
		<i>hrpP</i>	+	+	-
		<i>hrpO</i>	+	+	-
		<i>hrcN</i>	+	+	-
		<i>hrpQ</i>	+	+	-
		<i>hrpI/hrcV</i>	+	+	-
		<i>hrpJ</i>	+	+	-
		<i>hrpL</i>	+	+	-
		<i>hrpX</i>	+	+	-
		<i>hrpY</i>	+	+	-
		<i>hrpS</i>	+	+	-
		<i>hrpA</i>	+	+	-
		<i>hrpB</i>	+	+	-
		<i>hrcJ</i>	+	+	-
		<i>hrpD</i>	+	+	-
		<i>hrpE</i>	+	+	-
		<i>hrpF</i>	+	+	-
		<i>hrpG</i>	+	+	-
		<i>hrcC</i>	+	+	-
		<i>hrpT</i>	+	+	-
		<i>hrpV</i>	+	+	-
	Hrp elicitor/effector region	hrpN	+	+	-
		<i>orfA</i>	+	+	-
		<i>orfB</i>	+	+	-
		<i>orfC</i>	+	+	-
	Chaperone for DspA/E	hrpW	+	+	-
		dspA/E	+	+	-
		<i>dspB/F</i>	+	+	-
Salmonella SPI-1-like Type III secretion system and related exported proteins	Cell adherence/invasion protein	<i>InvH</i>	-	-	-
	Secretion chaperone	<i>invF</i>	+	+	-
	Secretion protein	<i>invG</i>	+	+	-
	Cell invasion protein	<i>invE</i>	+	+	-
	Invasion protein	<i>invA</i>	+	+	-
	Surface presentation of antigens	<i>invB</i>	+	+	-
	ATPase	<i>invC/spaL</i>	+	+	-

	Surface presentation of antigens	<i>invI/spaM</i>	-	-	-
		<i>invJ/spaN</i>	-	-	-
		<i>spaO</i>	+	+	-
		<i>spaP</i>	+	+	-
		<i>spaQ</i>	+	+	-
		<i>spaR</i>	+	+	-
		<i>spaS</i>	+	+	-
		<i>spaT</i>	-	-	-
		<i>sicA</i>	+	+	-
	Secretion protein				
	Chaperone protein (<i>Salmonella</i> invasin chaperone)				
	Cell invasion protein	<i>sipB</i>	+	+	-
		<i>sipC</i>	-	-	-
		<i>sipD</i>	+	+	-
		<i>sipA</i>	-	-	-
		<i>sptP</i>	-	-	-
	Protein tyrosine phosphatase	<i>prgH</i>	+	+	-
	Cell invasion proteins	<i>prgI</i>	+	+	-
		<i>prgJ</i>	(+)	(+)	-
		<i>prgK</i>	+	+	-
Type V secretion system		<i>orgA</i>	+	+	-
	Oxygen-regulated invasion protein	<i>ipaB</i>	-	-	-
	Invasion plasmid antigen	<i>ipaC</i>	-	-	-
	Cell invasion protein	<i>ipaD</i>	-	-	-
	Invasion plasmid antigen	<i>ipaA</i>	-	-	-
	Invasin				
	Autotransporter/effector proteins	Porin- /Pertactin-like	-	-	+
	Lipoprotein VCA0113	EpC_06160	+	+	+
	conserved uncharacterized protein	EpC_06170	+	+	+
	OmpA/MotB-like protein	<i>vasF1</i>	+	+	+
	IcmF-like protein	EpC_06190	+	+	+
	T6SS associated protein	EpC_06200	+	+	+
	T6SS associated ImpA protein	EpC_06210	+	+	+
	conserved uncharacterized protein	EpC_06220	+	+	+
	conserved uncharacterized protein	EpC_06230	+	+	+
	conserved uncharacterized protein	EpC_06240	+	+	-
	Virulence factor for secretion apparatus	<i>hcp</i>	+	+	+
	putative exported protein	EpC_06290	+	-	+
	FHA domain-containing protein	EpC_06300	+	+	+
Type VI secretion system	Protein phosphatase 2C-like protein	EpC_06310	+	+	+
	conserved uncharacterized protein	EpC_06320	+	+	+
	conserved uncharacterized protein	EpC_06330	+	+	+
	conserved uncharacterized protein	EpC_06340	+	+	+
	T6SS family protein, VCA0110	<i>vasA</i>	+	+	+
	conserved uncharacterized protein	EpC_06360	+	+	+
	conserved uncharacterized protein	EpC_06370	+	+	-
	conserved uncharacterized protein	EpC_06380	+	+	-
	T6SS family ATPase. ClpV1	EpC_06390	+	+	+
	Serine/threonine protein kinase	EpC_06400	+	+	+
	Rhs element Vgr protein	<i>vgrG1</i>	+	+	+
	Rhs element Vgr protein	<i>vgrG2</i>	+	+	+
	Putative invasin YchP	<i>ychP</i>	+	+	+
	OmpA family protein	<i>vasF2</i>	+	+	+
	IcmF protein	EpC_19530	+	+	+
	T6SS associated protein	EpC_19540	+	+	+
	FHA domain-containing protein	EpC_19550	+	-	+
Effector and virulence proteins	SsrAB regulated protein	<i>srfA</i>	+	+	+
	Virulence protein	<i>srfB</i>	+	+	+
	Virulence effector protein	<i>srfC</i>	+	+	+

	Secreted effector protein maintenance of virulence plasmid, toxin Virulence protein Two-component response regulator of virulence determinants Virulence sensor histidine kinase mouse virulence protein Virulence-related outer membrane protein X Virulence outer membrane protein Putative virulence protein Putative tyrosine-protein phosphatase YopH (Virulence protein)	sopA <i>mvpT</i> <i>msgA</i> <i>phoP</i> <i>phoQ</i> <i>mviN</i> <i>ompX</i> <i>pagC</i> <i>virK</i> yopH	+	-	-
			+	+	-
			+	+	+
			+	+	+
			+	+	+
			+	+	+
			+	+	+
			+	-	+
			-	+	+
			+	+	-
Metabolism					
Functional system	Product	Gene name	Ep1/96	Et1/99	Eb661
Capsular polysaccharide biosynthesis	UDP-galactose-lipid carrier transferase	<i>ams/cpsG</i>	+	+	+
	Periplasmic protein involved in polysaccharide export	<i>ams/cpsH</i>	+	+	+
	Low molecular weight protein-tyrosine-phosphatase	<i>ams/cpsI</i>	+	+	+
	Tyrosine-protein kinase	<i>ams/cpsA</i>	+	+	+
	Glycosyltransferase	<i>ams/cpsB</i>	+	+	+
	Exopolysaccharide biosynthesis protein	<i>ams/cpsC</i>	+	+	+
	Glycosyltransferase	<i>ams/cpsD</i>	+	+	+
		<i>ams/cpsE</i>	+	(+)	(+)
	Exopolysaccharide biosynthesis proteins	<i>ams/cpsF</i>	+	+	+
		<i>ams/cpsJ</i>	+	+	+
		<i>ams/cpsK</i>	+	+	+
		<i>ams/cpsL</i>	+	+	+
	Glycosyltransferase	<i>galF</i>	+	+	+
	Exopolysaccharide biosynthesis protein	<i>galE</i>	+	+	+
	Possible subunit of GalU	<i>ymcA</i>	+	+	+
	UDP-glucose 4-epimerase	<i>ymcB</i>	+	+	+
	Putative capsular polysaccharide lipoproteins	<i>ymcC</i>	+	+	+
		<i>rcsA</i>	+	+	+
	Activator of capsular EPS synthesis	<i>rcsB</i>	+	+	+
	Regulator of capsular EPS synthesis	<i>rcsC</i>	+	+	+
		<i>rcsD</i>	+	+	+
Levan metabolism	Levansucrase	<i>lsc</i>	-	+	+
	Levanase	EpC_17920	+	-	-
	Regulators of levansucrase expression	<i>rlsA</i>	+	+	-
		<i>rlsB</i>	+	+	-
		<i>rlsC</i>	+	+	-
Sorbitol-Operon	Glucitol/sorbitol-specific enzyme IIC	<i>srlA</i>	+	-	+
	Enzyme IIB of PTS	<i>srlE</i>	+	-	+
	Enzyme IIA of PTS	<i>srlB</i>	+	-	+
	Sorbitol-6-phosphate 2-dehydrogenase	<i>srlD</i>	+	-	+
	Activator of <i>srl</i> operon	<i>srlM</i>	+	-	+
	Repressor of <i>srl</i> operon	<i>srlR</i>	+	-	+
Sucrose metabolism	Fructokinase	<i>srcK</i>	+	+	-
	Porin	<i>srcY</i>	+	+	-
	Enzyme II	<i>srcA</i>	+	+/+	-
	Hydrolase	<i>srcB</i>	+	+/+	-
	Repressor	<i>srcR</i>	+	+	-
NRPS	Non-ribosomal peptide synthetase	<i>eppT</i>	+	(+)	-

Adhesion and extracellular factors					
Functional system	Product	Gene name	Ep1/96	Et1/99	Eb661
Necrosis factors	Probable cytotoxic necrotizing factor 1 factor 2	<i>cnf1</i>	+	+	-
		<i>cnf2</i>	+	+	-
Proteases	Protease III	<i>ptrA</i>	+	+	+
	Protease II	<i>ptrB</i>	+	+	+
	Protease A of <i>E. amylovora</i>	<i>prrA,D,E,F</i>	-	-	-
Siderophores	Ferrioxamine receptor	<i>foxR</i>	+	+	+
	Siderophore biosynthesis enzyme, L-lysine 6-monooxygenase (NADPH)	<i>dfoA</i>	+	+	-
	Siderophore biosynthesis protein, probable alcaligin	<i>alcA</i>	+	+	-