

**Table S1.** Bacterial strains, plasmids and oligonucleotides used in this study.

Strain or plasmid	Relevant characteristics <sup>a</sup>	Reference or source
<b><i>Xanthomonas campestris</i> pv. <i>vesicatoria</i></b>		
85-10	Pepper-race 2; wild type; Rif <sup>R</sup>	(1)
85-10Δ <i>hrpX</i>	85-10 derivative deleted in <i>hrpX</i> ; Rif <sup>R</sup>	(2)
85-10Δ <i>sX12</i>	85-10 derivative deleted in <i>sX12</i> ; Rif <sup>R</sup>	This study
85*	85-10 derivative containing the <i>hrpG</i> * mutation; Rif <sup>R</sup>	(3)
<b><i>Escherichia coli</i></b>		
DH5α λpir	F <sup>-</sup> <i>recA hsdR17(r<sub>k</sub><sup>+</sup>, m<sub>k</sub><sup>+</sup>) φ80dlacZ ΔM15 [λpir]</i>	(4)
TOP10	F <sup>-</sup> <i>mcrA Δ(mrr-hsdRMS-mcrBC) φ80lacZΔM15 ΔlacX74 recA1 araD139 Δ(ara-leu) 7697 galJ galK rpsL (Str<sup>R</sup>) endA1 nupG</i>	Invitrogen
TOP10F <sup>-</sup>	F <sup>-</sup> { <i>lacI<sup>q</sup> Tn10 (Tet<sup>R</sup>-)</i> } <i>mcrA Δ(mrr-hsdRMS-mcrBC) φ80lacZΔM15ΔlacX74 recA1 araD139 Δ(ara-leu) 7697 galJ galK rpsL endA1 nupG</i>	Invitrogen
<b>Plasmids</b>		
pBRM-P	Derivative of pBBR1MCS-5 without promoter; Gm <sup>R</sup>	(5)
pBRM-sX6	pBRM-P derivative expressing c-Myc-epitope tagged sX6; Gm <sup>R</sup>	This study
pCR 2.1-TOPO	General-purpose cloning vector; Ap <sup>R</sup> Km <sup>R</sup>	Invitrogen
pFG72-1	Derivative of pUFR043 containing <i>hrpG</i> *, Km <sup>R</sup>	(3)
pLAFR6	RK2 replicon, Mob <sup>+</sup> Tra <sup>+</sup> ; without promoter; multicloning site flanked by transcription terminators; Tc <sup>R</sup>	(6)
psX12	pLAFR6 derivative containing <i>sX12</i> ; Tc <sup>R</sup>	This study
pOK1	Suicide vector; <i>sacB sacQ mobRK2 oriR6K</i> ; Sm <sup>R</sup>	(7)
pOKΔ <i>sX12</i>	pOK1 derivative containing flanking regions of <i>sX12</i> ; Sm <sup>R</sup>	This study
pRK2013	ColE1 replicon, TraRK <sup>+</sup> Mob <sup>+</sup> ; Km <sup>R</sup>	(8)
<b>Oligonucleotide</b>	<b>Sequence (Purpose)</b>	<b>Gene</b>
NB76	CTGCGTGGAGTTTCTAGGCT (Northern blot probe)	<i>sX1</i>
NB2	GTCGCCTAGATGCTCTAGGG (Northern blot probe)	<i>sX2</i>
NB12	GTGGCGAGTAAGGCCAAAAAG (Northern blot probe)	<i>sX3</i>
NB13	CAAATTGTTTCAGGAACCTACGC (Northern blot probe)	<i>sX4</i>
NB81	GGACTTCGTAACGCAGGACT (Northern blot probe)	<i>sX5</i>
NB16	TCAGGATGTGCTTGAACCTTCAT (Northern blot probe)	<i>sX6</i>
NB17	CTCCTGGCCTTCGATAGATCTA (Northern blot probe)	<i>sX7</i>
NB23	ACTAGGCGCATAAGTCGTTGTT (Northern blot probe)	<i>sX8</i>
NB25	ACATCACACCGCGACCAG (Northern blot probe)	<i>sX9</i>
NB28	CCGAAGGAGTGTGTATTAGC (Northern blot probe)	<i>sX10</i>
NB31	AGTAGTAGGCTTGCTCAGAGCC (Northern blot probe)	<i>sX11</i>
NB42	TACCTTTCGACGAGGATGTG (Northern blot probe)	<i>sX12</i>
NB100	GAGAGATCTACCGACAGGCGT (Northern blot probe)	<i>sX13</i>
NB101	CACAGCTCCGACTGACATC (Northern blot probe)	<i>sX14</i>
NB67	TTACCGATCGTCGTGTAGCTG (Northern blot probe)	<i>sX15</i>
NB30	TGGTCGTAATTAAGGGACAAG (Northern blot probe)	6S
NB55	CTACCGACCCCTTACGCTACC (Northern blot probe)	<i>asX1</i>
NB89	CAGACCACCCAACAACCTTC (Northern blot probe)	<i>asX2</i>
NB45	GCCGAACCTACGTCCTTGTC (Northern blot probe)	<i>asX3</i>
NB98	CCAAAAACACGACTCAGTCG (Northern blot probe)	<i>asX4</i>
NB37	GATCAAACGAGTGGCTACTGTG (Northern blot probe)	<i>asX5</i>
NB66	AAACAGACCATGTGGCACATC (Northern blot probe)	<i>asX6</i>
NB72	CAGCTACACGACGATCGGTA (Northern blot probe)	<i>asX7</i>
NB-5S	CCTGGCGATGACCTACTCTC (Northern blot probe)	5S rRNA
5' RNA-adapter	AUAUGCGCGAAUUCUGUAGAACGAACACUAGAAGAAA (5' RACE; RNA-adapter)	
3' RNA-adapter*	P-UUCACUGUUCUAGCGCCGCAUGCUC-idT* (3' RACE; RNA-adapter)	
5'-RACE-adapter	GCGCGAATTCCTGTAGA (5' RACE; adapter-specific primer)	
3'-RACE-adapter	CGGCCGTAAGAACAG (3' RACE; adapter-specific primer)	
sX12-5' RACE1	ACCTTTCGACGAGGATG (5' RACE; reverse transcription)	<i>sX12</i>
sX12-5' RACE2	CGACGAGGATGTGCAG (5' RACE; PCR)	<i>sX12</i>
sX12-3' RACE	TGCACATCCTCGTCTGA (3' RACE; PCR)	<i>sX12</i>

Oligonucleotide	Sequence (Purpose)	Gene
asX4-5'RACE1	GAATCTGGTCACGACG (5' RACE; reverse transcription)	asX4
asX4-5'RACE2	TCTGGTCACGACGGTG (5' RACE; PCR)	asX4
asX4-3'RACE	GTCGTGACCAGATTCCTGTCT (3' RACE; PCR)	asX4
pCR2.1_colo_fw	ACGACGTTGTAACGACGG (colony PCR)	
pCR2.1_colo_rev	TTCACACAGGAAACAGCTATGAC (colony PCR)	
pCR2.1_seq	CACAGGAAACAGCTATGAC (sequencing)	
$\Delta$ sX12-left-fwd	<u>CAGGATCC</u> ATGATCAGGGTGTGAGGGTG (sX12 deletion; left fragment)	sX12
$\Delta$ sX12-left-rev	AAC <u>AAGCTT</u> ATGACCTCTGCCGGTTGTCT (sX12 deletion; left fragment)	sX12
$\Delta$ sX12-right-fwd	AAC <u>AAGCTT</u> AATCGATTTCCCGGCTTG (sX12 deletion; right fragment)	sX12
$\Delta$ sX12-right-rev	<u>GCTCTAGAGTGGCGACCTGGTACTTCAG</u> (sX12 deletion; right fragment)	sX12
sX12-comp-fwd	<u>CAGGATCCAACGGAGTTCAGCACTACGGT</u> (complementation of $\Delta$ sX12)	sX12
sX12-comp-rev	AAC <u>AAGCTT</u> CGAATTCTTGGAAATGGTCA (complementation of $\Delta$ sX12)	sX12
sX6-fwd	TTT <u>GGTCTCT</u> ATTTCGCTGAGCCGATTTTTGCAC (sX6-c-Myc tagging)	sX6
sX6-rev	TTT <u>GGTCTCT</u> CACCGCTGCAGGCTTTTCTTCTCC (sX6-c-Myc tagging)	sX6

<sup>a</sup>, Ap, ampicillin; Gm, gentamycin; Km, kanamycin; Rif, rifampicin; Sm, spectinomycin; Tc, tetracycline. <sup>R</sup>, resistance. Recognition sites of restriction enzymes are underlined.

\*, RNA oligonucleotide carrying a phosphate modification at the 5' and an inverted deoxythymidine (idT) at the 3' end.

## References

- Canteros, B.I. (1990) Ph.D. thesis. *University of Florida, Gainesville, FL.*
- Koebnik, R., Krüger, A., Thieme, F., Urban, A. and Bonas, U. (2006) Specific binding of the *Xanthomonas campestris* pv. *vesicatoria* AraC-type transcriptional activator HrpX to plant-inducible promoter boxes. *J. Bacteriol.*, **188**, 7652-7660.
- Wengelnik, K., Rossier, O. and Bonas, U. (1999) Mutations in the regulatory gene *hrpG* of *Xanthomonas campestris* pv. *vesicatoria* result in constitutive expression of all *hrp* genes. *J. Bacteriol.*, **181**, 6828-6831.
- Menard, R., Sansonetti, P.J. and Parsot, C. (1993) Nonpolar mutagenesis of the *ipa* genes defines IpaB, IpaC, and IpaD as effectors of *Shigella flexneri* entry into epithelial cells. *J. Bacteriol.*, **175**, 5899-5906.
- Szczesny, R., Jordan, M., Schramm, C., Schulz, S., Coge, V., Bonas, U. and Büttner, D. (2010) Functional characterization of the Xcs and Xps type II secretion systems from the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria*. *New Phytol.*, **187**, 983-1002.
- Bonas, U., Stall, R.E. and Staskawicz, B. (1989) Genetic and structural characterization of the avirulence gene *avrBs3* from *Xanthomonas campestris* pv. *vesicatoria*. *Mol. Gen. Genet.*, **218**, 127-136.
- Huguët, E., Hahn, K., Wengelnik, K. and Bonas, U. (1998) *hpaA* mutants of *Xanthomonas campestris* pv. *vesicatoria* are affected in pathogenicity but retain the ability to induce host-specific hypersensitive reaction. *Mol. Microbiol.*, **29**, 1379-1390.
- Figurski, D.H. and Helinski, D.R. (1979) Replication of an origin-containing derivative of plasmid RK2 dependent on a plasmid function provided *in trans*. *Proc. Natl. Acad. Sci. U. S. A.*, **76**, 1648-1652.

**Table S2.** Classification of putative TSSs which were automatically identified in the *Xcv* genome [see Figure 1B and METHODS; (1)].

TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
22092#	+	0.0001	6	0	XCVd0018	29	-	-	-	-	-
22614#	-	0.0034	7	2	-	-	-	-	XCVd0019	28/478	-
24098#	-	0.0191	3	0	-	-	-	-	-	-	+
31729#	-	0.0071	7	0	-	-	-	-	-	-	+
54664#	-	0.0344	3	0	-	-	-	-	XCVd0049	81/1463	-
56391#	-	0.0000	72	14	-	-	-	-	-	-	+
63600#	+	0.0003	6	0	-	-	-	-	XCVd0054	572/930	-
76432#	-	0.0007	7	0	XCVd0069	24	-	-	-	-	-
81111#	-	0.0032	9	2	-	-	-	-	XCVd0073	701/681	-
81749#	-	0.0014	9	2	-	-	-	-	XCVd0073	1339/43	-
82121#	+	0.0018	5	1	XCVd0074	21	-	-	-	-	-
88062#	+	0.0024	4	0	-	-	XCVd0079	757/2551	-	-	-
90670#	+	0.0341	9	0	XCVd0080	50	-	-	-	-	-
94989#	+	0.0081	4	0	-	-	XCVd0086	82/1066	-	-	-
95432#	-	0.0000	21	1	-	-	-	-	XCVd0086	525/623	-
97457#	-	0.0064	6	1	-	-	-	-	XCVd0087	1099/1000	-
99301#	-	0.0172	4	0	-	-	XCVd0089	238/370	-	-	-
100770#	+	0.0000	20	2	-	-	XCVd0091	817/13	-	-	-
101600#	+	0.0145	5	0	XCVd0093	287	-	-	XCVd0092	52/706	-
101864#	+	0.0000	156	25	XCVd0093	23	-	-	-	-	-
103085#	+	0.0369	4	3	-	-	-	-	XCVd0094	894/170	-
104001#	-	0.0000	57	11	XCVd0094	22	-	-	-	-	-
109851#	+	0.0000	35	6	-	-	-	-	XCVd0099	1283/99	-
112115#	-	0.0104	5	0	XCVd0101 XCVd0102	285 28	-	-	-	-	-
112250#	+	0.0000	41	11	XCVd0103	27	-	-	-	-	-
113697#	+	0.0159	3	1	-	-	-	-	XCVd0104	755/579	-
114827#	-	0.0000	11	0	-	-	XCVd0105	786/281	-	-	-
115639#	-	0.0000	33	4	XCVd0105	26	-	-	-	-	-
116378#	+	0.0000	80	8	XCVd0107	277	-	-	XCVd0106	31/486	-
117900#	+	0.0000	18	2	-	-	-	-	XCVd0109	740/180	-
117969#	+	0.0018	13	2	-	-	-	-	XCVd0109	671/249	-
118654#	+	0.0156	8	1	XCVd0110	201	-	-	XCVd0109	14/934	-
118665#	-	0.0000	54	9	XCVd0109	25	-	-	-	-	-
119469#	+	0.0018	5	1	XCVd0112	49	XCVd0111	375/41	-	-	-
120806#	+	0.0083	6	1	-	-	XCVd0112	1288/1684	-	-	-
120867#	-	0.0229	5	1	-	-	-	-	XCVd0112	1349/1623	-
123136#	+	0.0004	5	0	XCVd0114	218	XCVd0113	624/71	-	-	-
127227#	-	0.0439	4	2	-	-	-	-	XCVd0115	3019/62	-
127749#	-	0.0000	28	3	-	-	-	-	XCVd0116	275/1032	-
136150#	+	0.0027	3	0	-	-	XCVd0125	512/210	-	-	-
137519#	-	0.0000	15	0	XCVd0126	23	-	-	-	-	-
138153#	+	0.0020	5	0	-	-	-	-	XCVd0127	74/355	-
151691#	+	0.0026	4	0	-	-	-	-	XCVd0144	337/304	-
159251#	+	0.0000	49	5	XCVd0153	36	-	-	XCVd0152	71/2254	-
159633#	+	0.0158	3	0	-	-	XCVd0153	346/49	XCVd0154	409/74	-
164247#	-	0.0168	3	0	-	-	XCVd0155	1062/4193	-	-	-
169348#	+	0.0024	3	0	-	-	-	-	XCVd0160 XCVd0161	98/667 249/104	-
180140#	+	0.0049	3	0	-	-	-	-	XCVd0171	604/86	-
182433#	-	0.0050	3	0	XCVd0172	62	-	-	-	-	-

<sup>a</sup>, TSS position on the *Xcv* chromosome and the plasmid (pXCV183; marked with #), respectively. <sup>b</sup>, p-value indicates the significance of the annotated TSS (see METHODS). <sup>c</sup>, number of read starts at TSS position. <sup>d</sup>, TSS classification and the corresponding CDS. <sup>e</sup>, distance of the TSS to annotated start and stop codon of the corresponding CDS. <sup>f</sup>, a TSS is classified as orphan (+) if it is neither classified as primary, internal nor antisense TSS.

TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
1404	+	0.0000	15	1	XCV0002	202	-	-	-	-	-
9520	+	0.0078	5	2	XCV0007	68	-	-	-	-	-
10034	+	0.0036	4	0	-	-	XCV0007	446/744	-	-	-
10857	+	0.0001	10	1	XCV0008	78	-	-	-	-	-
11675	-	0.0001	6	0	-	-	-	-	XCV0009 XCV0008	17/775 740/72	-
15312	-	0.0000	12	0	XCV0013 XCV0014	297 20	-	-	-	-	-
17027	+	0.0000	42	18	-	-	XCV0016	23/900	-	-	-
17192	-	0.0001	12	0	-	-	-	-	XCV0016	188/735	-
18468	+	0.0022	3	0	-	-	XCV0017	538/49	-	-	-
18664	-	0.0147	3	0	-	-	-	-	-	-	+
23398	+	0.0000	13	1	-	-	XCV0022	45/317	-	-	-
38760	-	0.0000	13	0	-	-	XCV0035	12/425	-	-	-
44970	-	0.0000	14	3	XCV0037	79	-	-	-	-	-
45473	+	0.0246	5	0	XCV0039	58	-	-	XCV0038	1/376	-
45497	-	0.0170	3	0	XCV0038	23	-	-	XCV0039	34/1308	-
46397	-	0.0101	3	0	-	-	-	-	XCV0039	866/408	-
47391	-	0.0063	6	0	XCV0040	22	-	-	-	-	-
52187	+	0.0170	3	1	XCV0045	22	-	-	-	-	-
60944	-	0.0066	4	0	XCV0050	66	-	-	XCV0051	98/1191	-
62304	+	0.0206	3	0	XCV0052	23	-	-	-	-	-
72269	+	0.0213	3	0	XCV0061	116	-	-	XCV0060	27/761	-
72322	-	0.0002	9	1	XCV0060	80	-	-	XCV0061	63/404	-
72626	+	0.0001	13	2	-	-	XCV0061	241/100	-	-	-
75816	-	0.0016	4	0	-	-	-	-	-	-	+
77193	-	0.0006	6	1	XCV0064	16	-	-	-	-	-
78522	+	0.0000	23	2	-	-	XCV0066	817/13	XCV0067	231/46	-
78978	-	0.0187	8	5	XCV0067	225	-	-	-	-	-
81973	-	0.0123	6	2	XCV0069	270	-	-	-	-	-
84253	-	0.0047	3	0	XCV0072	20	-	-	-	-	-
84514	+	0.0058	3	0	-	-	XCV0073	121/460	-	-	-
85087	+	0.0000	11	0	-	-	-	-	-	-	+
87652	+	0.0469	7	0	XCV0077	22	-	-	-	-	-
89289	+	0.0065	4	0	-	-	XCV0078	1122/2840	-	-	-
103766	+	0.0206	3	0	-	-	-	-	XCV0088	747/149	-
109502	-	0.0009	6	1	XCV0092	72	-	-	-	-	-
117077	-	0.0026	4	0	XCV0099	274	XCV0100	573/266	-	-	-
118296	-	0.0401	3	1	-	-	XCV0101	675/524	-	-	-
133989	+	0.0164	3	1	-	-	XCV0117	8/588	-	-	-
138711	+	0.0011	6	1	XCV0120	26	-	-	-	-	-
139008	+	0.0333	3	0	-	-	XCV0120	271/934	-	-	-
149036	+	0.0002	7	0	-	-	-	-	-	-	+
149533	-	0.0000	13	1	-	-	XCV0125	817/13	-	-	-
152067	-	0.0152	5	0	-	-	-	-	XCV0128	155/543	-
155750	+	0.0005	11	2	XCV0131	227	-	-	-	-	-
156627	-	0.0000	15	3	-	-	-	-	XCV0132	106/295	-
192866	-	0.0118	3	0	XCV0160	31	-	-	-	-	-
193010	+	0.0008	9	2	XCV0161	0	-	-	-	-	-
193630	+	0.0149	3	1	-	-	XCV0161	620/750	-	-	-
195442	-	0.0000	22	0	XCV0162	23	-	-	-	-	-
196578	+	0.0000	37	10	XCV0164	32	-	-	-	-	-
201827	-	0.0015	7	2	XCV0168	0	-	-	XCV0169	12/1352	-
201899	+	0.0000	16	1	-	-	XCV0169	84/1280	XCV0168	72/1250	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
204493	+	0.0000	12	2	-	-	-	-	XCV0172	3775/71	-
208848	+	0.0338	4	0	-	-	XCV0173	420/248	-	-	-
212756	+	0.0000	8	0	XCV0178	116	XCV0177	396/116	-	-	-
217013	+	0.0021	4	0	-	-	XCV0181	424/562	-	-	-
217960	+	0.0416	7	4	XCV0183	22	-	-	-	-	-
220222	+	0.0000	11	0	XCV0184	23	-	-	-	-	-
222838	-	0.0000	17	0	XCV0186	21	-	-	-	-	-
224042	+	0.0000	15	2	XCV0188	35	-	-	-	-	-
225642	+	0.0161	4	2	XCV0189	118	-	-	-	-	-
237714	-	0.0365	4	1	XCV0200	47	-	-	-	-	-
238550	-	0.0323	7	0	XCV0201	0	-	-	XCV0202	57/1010	-
240862	+	0.0050	6	2	XCV0205	23	-	-	-	-	-
241374	+	0.0011	4	0	XCV0206	55	-	-	-	-	-
244105	-	0.0000	194	14	XCV0208	62	-	-	-	-	-
244587	+	0.0000	15	2	-	-	XCV0209	4/1204	-	-	-
250048	+	0.0221	3	0	-	-	-	-	XCV0213	484/535	-
252996	+	0.0000	19	1	-	-	-	-	XCV0214	72/2471	-
253127	-	0.0000	25	10	XCV0214	203	-	-	-	-	-
261157	+	0.0000	17	2	XCV0221	81	-	-	-	-	-
261741	-	0.0000	15	1	-	-	XCV0222	817/13	XCV0221	503/3	-
265771	-	0.0000	43	8	-	-	XCV0226	216/569	-	-	-
266587	-	0.0000	66	8	XCV0227	90	-	-	-	-	-
267156	+	0.0000	17	2	-	-	-	-	-	-	+
269292	+	0.0043	3	0	-	-	-	-	-	-	+
278015	-	0.0012	6	0	XCV0236	0	-	-	-	-	-
278867	+	0.0100	5	0	XCV0238	23	-	-	-	-	-
280390	+	0.0019	3	0	XCV0240	22	-	-	-	-	-
283462	+	0.0037	4	0	XCV0243	0	-	-	-	-	-
285329	+	0.0140	4	1	XCV0244	22	-	-	-	-	-
287897	-	0.0105	4	1	XCV0246	21	-	-	-	-	-
288111	+	0.0128	10	5	XCV0247	0	-	-	-	-	-
292756	+	0.0016	8	0	XCV0252	122	-	-	XCV0251	108/629	-
295006	+	0.0046	6	1	XCV0255	119	-	-	-	-	-
304274	+	0.0276	5	0	XCV0264	150	-	-	XCV0263	49/904	-
307776	+	0.0002	11	2	XCV0266	37	-	-	-	-	-
311632	-	0.0178	3	0	-	-	XCV0268	98/213	-	-	-
315387	-	0.0003	5	0	XCV0270	36	-	-	-	-	-
320398	+	0.0021	6	1	XCV0274	234	-	-	XCV0273	105/1055	-
321787	-	0.0437	7	0	-	-	-	-	XCV0275 XCV0276	448/77 77/444	-
325673	+	0.0000	24	2	XCV0281	36	-	-	-	-	-
329286	-	0.0271	4	0	XCV0283	97	-	-	XCV0284	35/1287	-
331021	-	0.0263	6	1	-	-	XCV0285	933/278	-	-	-
331070	-	0.0003	10	1	-	-	XCV0285	884/327	-	-	-
332993	+	0.0000	9	1	XCV0287	0	-	-	XCV0286	67/939	-
337456	+	0.0001	13	3	XCV0293	22	-	-	-	-	-
338466	-	0.0057	5	2	-	-	XCV0294	940/259	-	-	-
341742	-	0.0051	5	2	XCV0296	23	-	-	-	-	-
341927	+	0.0000	12	0	XCV0297	26	-	-	-	-	-
344260	-	0.0000	27	4	XCV0299	26	-	-	-	-	-
347202	-	0.0014	4	0	-	-	XCV0300	99/2498	-	-	-
347842	-	0.0030	5	0	-	-	-	-	-	-	+
348352	+	0.0081	12	5	-	-	-	-	XCV0302	16/454	-
358119	-	0.0000	13	3	XCV0313	70	-	-	-	-	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
369663	+	0.0028	3	0	-	-	XCV0322	30/833	-	-	-
375494	-	0.0081	3	0	-	-	XCV0327	24/425	XCV0328	90/1007	-
377721	+	0.0023	4	0	XCV0330	3	-	-	-	-	-
404292	-	0.0135	3	1	XCV0352	127	XCV0353	512/0	-	-	-
405223	-	0.0008	6	1	-	-	XCV0354	769/88	-	-	-
405567	+	0.0000	162	23	-	-	-	-	XCV0354	425/432	-
406317	-	0.0006	19	10	XCV0355	29	-	-	-	-	-
411941	+	0.0150	3	1	XCV0359	6	-	-	-	-	-
428338	-	0.0000	112	12	XCV0374	48	-	-	-	-	-
429323	-	0.0000	14	1	-	-	XCV0375	153/806	-	-	-
436669	+	0.0180	4	0	-	-	XCV0383	11/1368	-	-	-
441623	-	0.0130	4	0	-	-	XCV0387	26/879	-	-	-
441720	+	0.0000	11	0	XCV0388	30	-	-	XCV0387	71/976	-
447445	-	0.0001	6	2	XCV0392	52	-	-	-	-	-
448031	-	0.0000	12	2	XCV0393	23	-	-	-	-	-
448950	+	0.0098	3	0	XCV0395	226	XCV0394	784/172	-	-	-
449748	+	0.0205	5	1	XCV0396	23	-	-	-	-	-
453984	-	0.0000	24	7	-	-	XCV0400	168/758	-	-	-
455880	-	0.0012	5	1	XCV0402	24	-	-	-	-	-
458086	-	0.0001	6	0	-	-	-	-	-	-	+
462619	+	0.0014	4	0	-	-	-	-	XCV0411	2250/167	-
475337	-	0.0022	3	0	-	-	XCV0425	899/1020	-	-	-
477343	-	0.0002	8	0	XCV0426	25	-	-	-	-	-
477496	+	0.0000	22	3	XCV0427	37	-	-	-	-	-
485143	-	0.0000	13	0	-	-	-	-	XCV0435	1914/94	-
486264	-	0.0059	4	0	XCV0436	140	-	-	-	-	-
486511	+	0.0005	5	0	-	-	-	-	-	-	+
487914	-	0.0323	3	1	-	-	-	-	XCV0437	725/909	-
488081	+	0.0001	7	0	-	-	XCV0437	892/742	-	-	-
522402	-	0.0002	10	0	-	-	-	-	XCV0462	245/183	-
530343	+	0.0000	9	1	-	-	XCV0468	153/356	-	-	-
532431	-	0.0120	6	2	-	-	XCV0470	69/1289	-	-	-
532651	-	0.0025	5	0	XCV0470	151	-	-	-	-	-
532770	+	0.0009	8	1	XCV0471	0	-	-	-	-	-
533893	+	0.0033	6	0	-	-	-	-	XCV0472	302/36	-
537353	-	0.0066	4	0	XCV0475	32	XCV0476	1049/36	-	-	-
548017	+	0.0016	3	0	XCV0483	96	-	-	-	-	-
557964	+	0.0057	3	0	-	-	XCV0493	25/922	-	-	-
558968	-	0.0002	8	1	-	-	XCV0494	432/50	XCV0493	1029/82	-
568795	+	0.0038	5	1	XCV0504	27	-	-	-	-	-
573662	+	0.0040	3	0	XCV0509	53	XCV0508	156/50	-	-	-
581801	+	0.0005	4	0	XCV0514	94	-	-	-	-	-
585332	+	0.0311	8	5	XCV0518	0	XCV0517	790/4	-	-	-
586733	-	0.0003	8	0	-	-	XCV0519	327/362	-	-	-
587078	-	0.0000	62	8	XCV0519	18	-	-	-	-	-
587168	+	0.0000	39	0	XCV0520	30	-	-	-	-	-
587395	-	0.0439	3	0	-	-	-	-	XCV0520	197/594	-
589427	-	0.0000	10	0	-	-	XCV0522	42/650	-	-	-
589525	+	0.0000	73	23	XCV0523	85	-	-	XCV0522	56/748	-
590048	-	0.0012	10	0	-	-	-	-	XCV0523 XCV0524	438/13 7/382	-
590956	+	0.0000	24	1	XCV0525	24	-	-	-	-	-
592178	+	0.0000	18	5	XCV0527	24	-	-	-	-	-
598012	-	0.0331	4	2	XCV0531	22	-	-	XCV0532	86/853	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
599405	+	0.0000	18	5	XCV0534	0	-	-	-	-	-
602321	+	0.0000	16	1	-	-	XCV0537	36/887	-	-	-
605417	-	0.0000	11	1	-	-	-	-	-	-	+
616013	-	0.0182	4	0	XCV0547	25	-	-	XCV0548	49/663	-
616018	-	0.0369	3	0	XCV0547	30	-	-	XCV0548	44/658	-
624601	-	0.0081	3	1	XCV0556	46	-	-	-	-	-
629187	+	0.0177	3	1	XCV0561	39	-	-	-	-	-
633751	-	0.0002	8	2	XCV0566	21	-	-	-	-	-
636699	-	0.0000	7	0	-	-	XCV0567	194/2121	-	-	-
647837	+	0.0000	18	0	XCV0574	192	-	-	-	-	-
654736	+	0.0074	4	0	-	-	-	-	-	-	+
656637	+	0.0079	6	2	-	-	XCV0579	42/2822	-	-	-
660302	+	0.0000	12	2	-	-	-	-	-	-	+
660601	-	0.0000	8	0	-	-	-	-	XCV0581	42/1880	-
668117	-	0.0002	6	0	XCV0586	52	-	-	XCV0587	83/670	-
687581	+	0.0000	24	2	-	-	XCV0608	268/1051	-	-	-
689806	-	0.0185	11	0	-	-	-	-	XCV0610	760/772	-
690527	+	0.0000	15	4	XCV0611	54	XCV0610	1481/51	-	-	-
692255	+	0.0469	3	0	-	-	XCV0611	1674/1010	-	-	-
693668	+	0.0006	4	0	XCV0612	26	-	-	-	-	-
694560	-	0.0169	3	0	-	-	-	-	XCV0612	866/111	-
698328	+	0.0000	14	2	XCV0616	34	-	-	-	-	-
703418	-	0.0086	3	0	-	-	-	-	-	-	+
707695	+	0.0343	4	1	-	-	-	-	XCV0625 XCV0624	838/8 79/258	-
707716	-	0.0000	13	1	XCV0624	100	XCV0625	817/13	-	-	-
713610	-	0.0193	3	0	-	-	XCV0632	9/1523	-	-	-
720541	-	0.0024	3	0	XCV0639	65	-	-	-	-	-
725332	+	0.0013	10	3	-	-	-	-	XCV0642	585/1994	-
732826	+	0.0424	4	0	-	-	XCV0648	43/334	-	-	-
736976	-	0.0027	3	0	-	-	XCV0653	8/789	-	-	-
738232	+	0.0005	12	4	XCV0656	25	-	-	-	-	-
739844	-	0.0122	3	0	-	-	XCV0657	862/121	-	-	-
740729	-	0.0000	12	2	XCV0657	23	-	-	-	-	-
749385	-	0.0005	6	1	-	-	XCV0665	168/956	-	-	-
752821	-	0.0000	16	1	XCV0666	23	-	-	-	-	-
758448	-	0.0056	5	0	XCV0669	34	-	-	-	-	-
758878	+	0.0469	7	0	-	-	-	-	XCV0670	1215/212	-
763156	+	0.0000	24	1	XCV0673	23	-	-	-	-	-
765474	+	0.0141	3	0	XCV0675	282	-	-	-	-	-
765583	+	0.0143	3	0	XCV0675	173	-	-	-	-	-
770365	+	0.0023	4	0	XCV0679	0	-	-	-	-	-
772840	+	0.0000	14	0	-	-	XCV0681	177/962	-	-	-
775297	-	0.0033	4	0	XCV0683	25	-	-	-	-	-
775992	+	0.0000	13	2	XCV0685	22	-	-	-	-	-
779535	-	0.0044	8	3	XCV0686	0	-	-	-	-	-
779651	+	0.0041	4	1	XCV0687	29	-	-	-	-	-
781259	-	0.0026	4	0	-	-	-	-	XCV0687	1579/490	-
782120	+	0.0002	7	0	XCV0688	0	-	-	-	-	-
782575	+	0.0000	13	2	XCV0689	244	-	-	-	-	-
785619	+	0.0000	18	2	XCV0692	17	-	-	-	-	-
793698	+	0.0000	11	1	XCV0700	23	-	-	-	-	-
799920	-	0.0039	6	2	-	-	XCV0705	769/88	XCV0704	942/49	-
800264	+	0.0000	162	23	-	-	-	-	XCV0705	425/432	-

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801014	-	0.0004	19	10	XCV0706	29	-	-	-	-	-
815890	+	0.0058	3	0	-	-	XCV0717	436/607	-	-	-
823961	+	0.0108	4	0	XCV0723	0	-	-	-	-	-
826922	+	0.0008	13	6	XCV0725	38	-	-	-	-	-
829714	+	0.0000	24	4	XCV0728	266	XCV0727	219/275	-	-	-
837085	-	0.0214	3	0	XCV0733	26	-	-	-	-	-
842923	+	0.0015	4	0	XCV0739	38	-	-	-	-	-
844848	+	0.0000	25	2	XCV0741	21	-	-	-	-	-
847069	+	0.0177	3	1	-	-	XCV0743	91/391	-	-	-
847628	+	0.0000	30	1	XCV0744	117	-	-	-	-	-
851592	+	0.0000	8	0	XCV0747	0	-	-	-	-	-
853751	+	0.0008	4	0	-	-	XCV0749	670/511	-	-	-
873841	+	0.0010	4	0	XCV0767	31	-	-	-	-	-
884284	-	0.0299	4	0	XCV0773	0	-	-	-	-	-
893392	+	0.0005	6	0	-	-	XCV0781	42/620	-	-	-
898261	-	0.0417	6	0	XCV0784	24	-	-	XCV0785	54/1151	-
900510	+	0.0380	4	0	-	-	XCV0787	97/292	-	-	-
905920	-	0.0007	5	0	XCV0792	60	-	-	-	-	-
906061	+	0.0000	16	3	XCV0793	24	-	-	-	-	-
906749	+	0.0000	10	0	XCV0794	31	-	-	-	-	-
907111	+	0.0010	5	0	-	-	XCV0794	331/919	-	-	-
907734	+	0.0007	6	0	-	-	XCV0794	954/296	-	-	-
931121	-	0.0030	3	0	-	-	XCV0816	41/324	-	-	-
933943	+	0.0000	86	27	-	-	-	-	-	-	+
935355	-	0.0000	11	1	XCV0819 XCV0820	264 16	-	-	-	-	-
936154	-	0.0000	12	1	-	-	XCV0821	15/764	-	-	-
936452	+	0.0048	11	1	XCV0822	18	-	-	-	-	-
936455	+	0.0028	23	4	XCV0822	15	-	-	-	-	-
946406	+	0.0206	3	0	-	-	XCV0830	571/721	-	-	-
951830	+	0.0001	10	2	XCV0835	181	-	-	-	-	-
953433	+	0.0005	12	1	XCV0836	51	-	-	-	-	-
956025	+	0.0000	14	0	XCV0839	101	-	-	-	-	-
956388	+	0.0055	5	0	-	-	XCV0839	262/2473	-	-	-
959187	+	0.0001	9	2	XCV0840	0	-	-	-	-	-
963192	-	0.0156	3	1	XCV0843	22	-	-	-	-	-
963547	+	0.0026	4	0	XCV0844	20	-	-	-	-	-
964753	+	0.0000	15	1	XCV0845	31	-	-	-	-	-
973152	+	0.0000	80	8	-	-	-	-	-	-	+
973732	+	0.0349	3	0	-	-	XCV0854	279/650	-	-	-
976232	+	0.0206	3	0	-	-	-	-	XCV0856	647/792	-
977026	-	0.0014	5	0	XCV0856	147	-	-	-	-	-
994684	-	0.0007	4	0	XCV0865	33	-	-	-	-	-
997844	-	0.0000	21	0	XCV0868	0	-	-	-	-	-
1003187	-	0.0005	5	0	XCV0871	27	-	-	-	-	-
1017977	+	0.0000	67	8	XCV0887	22	-	-	-	-	-
1021354	+	0.0006	11	3	XCV0890	73	-	-	-	-	-
1024293	+	0.0000	11	2	XCV0894	27	-	-	-	-	-
1028253	-	0.0039	3	0	XCV0899	0	-	-	-	-	-
1029078	-	0.0008	5	0	XCV0900	0	-	-	-	-	-
1031658	-	0.0108	3	1	XCV0902	80	XCV0903	2346/80	-	-	-
1034027	-	0.0212	4	2	XCV0903	23	-	-	-	-	-
1036242	-	0.0129	5	0	XCV0906	23	-	-	-	-	-
1036560	+	0.0006	6	0	XCV0907	0	-	-	-	-	-

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1041488	-	0.0166	3	1	-	-	-	-	XCV0911	1007/21	-
1042407	+	0.0004	7	1	XCV0913	180	-	-	XCV0912	54/725	-
1050250	-	0.0003	6	0	-	-	XCV0920	152/429	-	-	-
1052158	-	0.0193	3	0	XCV0922	21	-	-	-	-	-
1052229	+	0.0095	4	1	XCV0923	32	-	-	XCV0922	92/994	-
1055953	-	0.0005	5	0	XCV0926	26	-	-	-	-	-
1059633	-	0.0003	12	3	XCV0930	23	-	-	-	-	-
1064780	+	0.0131	3	1	XCV0937	26	-	-	-	-	-
1067662	-	0.0119	3	0	-	-	XCV0939	104/861	-	-	-
1073410	+	0.0000	17	0	XCV0945	19	-	-	-	-	-
1086476	+	0.0025	4	0	-	-	XCV0957	74/585	-	-	-
1095646	-	0.0012	4	0	-	-	-	-	XCV0965 XCV0964	12/1205 334/86	-
1102907	-	0.0272	3	1	XCV0973	0	-	-	-	-	-
1107405	+	0.0024	4	0	XCV0977	35	-	-	XCV0976	31/1326	-
1107410	-	0.0000	8	0	XCV0976	36	-	-	XCV0977	30/1715	-
1110383	+	0.0004	16	5	-	-	XCV0979	623/261	-	-	-
1114497	+	0.0001	8	0	-	-	-	-	-	-	+
1116481	+	0.0001	7	2	XCV0986	259	XCV0985	162/242	-	-	-
1117349	+	0.0000	9	1	XCV0987	147	-	-	-	-	-
1117882	+	0.0012	7	2	XCV0988	46	XCV0987	386/39	-	-	-
1118456	+	0.0303	5	0	-	-	XCV0988	528/167	-	-	-
1118678	+	0.0000	46	10	-	-	-	-	-	-	+
1120586	+	0.0007	4	1	-	-	XCV0991	172/3976	-	-	-
1122235	+	0.0042	3	1	-	-	XCV0991	1821/2327	-	-	-
1122779	+	0.0000	26	1	-	-	XCV0991	2365/1783	-	-	-
1128623	+	0.0074	5	1	-	-	XCV0992	3957/254	-	-	-
1129024	+	0.0000	10	1	XCV0993	75	-	-	-	-	-
1130406	+	0.0005	5	0	-	-	XCV0995	315/1799	-	-	-
1134713	-	0.0133	3	0	-	-	-	-	-	-	+
1135113	+	0.0030	15	12	-	-	-	-	-	-	+
1137159	+	0.0000	79	14	XCV1002	180	XCV1001	130/166	-	-	-
1137964	+	0.0008	9	2	XCV1003	209	XCV1002	625/199	-	-	-
1142119	+	0.0000	38	13	XCV1013	108	-	-	-	-	-
1146342	+	0.0000	37	12	-	-	-	-	-	-	+
1149059	+	0.0004	4	1	XCV1024	233	XCV1023	946/49	-	-	-
1149619	+	0.0026	3	0	XCV1025	180	XCV1024	327/53	-	-	-
1150093	+	0.0000	44	3	XCV1026	23	XCV1025	294/17	-	-	-
1153493	+	0.0000	10	1	XCV1028	17	-	-	-	-	-
1160474	-	0.0000	13	2	XCV1032	43	-	-	-	-	-
1160640	+	0.0000	20	1	XCV1033	33	-	-	-	-	-
1162992	-	0.0000	35	12	XCV1035	22	-	-	-	-	-
1164003	-	0.0006	5	0	XCV1036	163	-	-	-	-	-
1168793	+	0.0000	17	8	XCV1042	80	-	-	-	-	-
1169205	+	0.0000	19	3	-	-	XCV1042	332/762	-	-	-
1170131	+	0.0000	16	0	XCV1043	32	-	-	-	-	-
1173134	-	0.0006	5	0	XCV1046	22	-	-	-	-	-
1177286	+	0.0126	3	0	XCV1051	145	-	-	-	-	-
1179872	-	0.0114	5	2	XCV1052	40	-	-	-	-	-
1184957	+	0.0000	12	1	XCV1055	25	-	-	-	-	-
1185123	-	0.0387	7	2	-	-	-	-	XCV1055	141/491	-
1189664	+	0.0048	5	0	XCV1060	99	-	-	-	-	-
1190312	+	0.0006	8	2	XCV1061	228	XCV1060	549/194	-	-	-
1206581	-	0.0144	3	0	-	-	XCV1073	42/1661	XCV1074	99/437	-

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1206687	+	0.0000	11	1	-	-	XCV1074	7/331	XCV1073	64/1767	-
1207568	+	0.0000	8	0	XCV1076	72	-	-	-	-	-
1209562	-	0.0027	5	0	XCV1077	20	-	-	-	-	-
1209662	+	0.0017	10	0	-	-	-	-	-	-	+
1217572	+	0.0134	3	1	-	-	XCV1083	1641/827	-	-	-
1218498	+	0.0000	46	8	XCV1084	118	-	-	-	-	-
1219137	-	0.0112	3	0	-	-	-	-	-	-	+
1219503	+	0.0007	6	1	XCV1085	25	-	-	-	-	-
1223677	-	0.0046	3	0	-	-	XCV1087	426/338	-	-	-
1226768	-	0.0027	5	1	XCV1091	27	-	-	-	-	-
1227290	+	0.0000	10	0	-	-	XCV1092	172/1306	-	-	-
1233373	+	0.0000	19	2	-	-	XCV1098	817/13	-	-	-
1233578	+	0.0000	28	5	-	-	-	-	-	-	+
1235373	+	0.0020	5	1	-	-	-	-	-	-	+
1240320	+	0.0038	6	1	XCV1107	20	-	-	-	-	-
1242175	-	0.0000	91	11	-	-	-	-	-	-	+
1244521	-	0.0000	16	1	XCV1110	22	-	-	-	-	-
1249118	+	0.0002	20	9	XCV1118	29	XCV1117	910/16	-	-	-
1249868	-	0.0000	170	21	-	-	-	-	XCV1119	425/432	-
1250212	+	0.0041	5	1	XCV1120	153	XCV1119	769/88	-	-	-
1257067	-	0.0053	3	0	-	-	XCV1126	180/1334	-	-	-
1257541	+	0.0001	14	4	XCV1127	21	-	-	-	-	-
1261798	-	0.0002	16	6	XCV1131	65	XCV1132	1884/65	-	-	-
1265655	-	0.0171	3	1	XCV1134	0	-	-	XCV1135	40/1807	-
1270936	+	0.0000	47	4	-	-	XCV1141	37/517	-	-	-
1272952	-	0.0053	3	0	-	-	-	-	-	-	+
1273770	+	0.0000	36	10	XCV1145	23	-	-	-	-	-
1278951	+	0.0001	8	0	XCV1150	29	-	-	-	-	-
1284606	-	0.0088	4	1	-	-	XCV1155	27/1619	-	-	-
1287704	+	0.0032	3	0	-	-	XCV1158	694/1897	-	-	-
1295774	+	0.0131	4	1	XCV1164	42	XCV1163	308/12	-	-	-
1298129	-	0.0224	3	0	-	-	-	-	XCV1164	2313/2585	-
1309475	+	0.0005	8	1	XCV1172	65	-	-	XCV1171	98/196	-
1312832	+	0.0002	9	0	XCV1175	61	-	-	-	-	-
1321929	-	0.0414	5	0	XCV1181	22	-	-	XCV1182	50/637	-
1340207	+	0.0008	6	1	XCV1198	233	XCV1197	282/212	-	-	-
1343656	+	0.0005	5	0	XCV1202	63	-	-	-	-	-
1364483	+	0.0191	5	2	XCV1217	20	-	-	-	-	-
1368267	-	0.0345	6	3	XCV1221 XCV1222	249 19	-	-	-	-	-
1376702	+	0.0016	8	2	-	-	XCV1228	90/857	-	-	-
1381773	+	0.0121	3	0	XCV1232	23	-	-	-	-	-
1384583	-	0.0028	7	2	XCV1233	20	-	-	-	-	-
1384820	+	0.0000	17	3	XCV1234	46	-	-	-	-	-
1386923	+	0.0068	3	0	-	-	XCV1236	66/1907	-	-	-
1390988	-	0.0193	3	0	-	-	XCV1239	35/150	-	-	-
1399557	-	0.0435	4	2	XCV1243	291	-	-	-	-	-
1404809	+	0.0275	3	0	XCV1247	89	-	-	-	-	-
1411639	-	0.0000	18	4	XCV1251	22	-	-	-	-	-
1419087	+	0.0005	8	1	XCV1260	0	-	-	XCV1259	78/833	-
1421227	+	0.0000	11	0	XCV1262	48	-	-	-	-	-
1424474	-	0.0060	4	0	XCV1265	171	-	-	-	-	-
1426111	-	0.0236	10	0	XCV1268	21	-	-	-	-	-
1429267	-	0.0016	4	0	XCV1271	24	-	-	-	-	-

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1433010	-	0.0000	18	5	XCV1275	29	-	-	-	-	-
1436219	-	0.0000	8	0	XCV1279	24	-	-	-	-	-
1437031	+	0.0023	6	0	XCV1281	28	-	-	XCV1280	4/763	-
1437682	-	0.0023	3	0	-	-	XCV1282	365/36	-	-	-
1440046	-	0.0054	3	0	-	-	XCV1283	964/1996	-	-	-
1441077	-	0.0028	5	0	XCV1283	67	-	-	-	-	-
1443612	-	0.0000	49	7	XCV1287	41	-	-	-	-	-
1443614	-	0.0007	48	7	XCV1287	43	-	-	-	-	-
1445416	+	0.0000	17	4	XCV1289	51	-	-	-	-	-
1450021	+	0.0001	8	1	XCV1292	0	-	-	-	-	-
1451420	+	0.0001	14	1	-	-	-	-	-	-	+
1451553	+	0.0055	10	2	-	-	-	-	-	-	+
1454260	-	0.0125	3	1	-	-	-	-	XCV1295	821/15	-
1456955	+	0.0000	37	0	XCV1299	25	-	-	-	-	-
1457771	+	0.0000	12	2	-	-	XCV1299	791/84	-	-	-
1458095	-	0.0000	15	1	-	-	XCV1300	819/11	-	-	-
1460678	+	0.0000	9	0	-	-	-	-	XCV1303	554/672	-
1461842	-	0.0113	3	1	XCV1304	23	-	-	-	-	-
1463947	-	0.0002	7	1	XCV1306	112	-	-	-	-	-
1464022	+	0.0151	9	0	-	-	-	-	-	-	+
1464279	+	0.0000	107	6	XCV1307	118	-	-	-	-	-
1464610	+	0.0122	3	0	-	-	XCV1307	213/719	-	-	-
1468435	+	0.0000	36	10	XCV1311	30	-	-	-	-	-
1470573	+	0.0005	6	0	XCV1312	22	-	-	-	-	-
1498473	-	0.0002	8	0	-	-	XCV1332	8/3195	-	-	-
1498586	+	0.0000	18	3	XCV1333	21	-	-	-	-	-
1502160	+	0.0000	15	3	XCV1336	21	-	-	-	-	-
1504286	-	0.0000	15	0	XCV1337	0	-	-	-	-	-
1504669	+	0.0019	5	1	XCV1338	28	-	-	-	-	-
1506240	+	0.0023	7	2	XCV1339	21	-	-	-	-	-
1508171	+	0.0000	62	7	XCV1342	45	-	-	-	-	-
1508253	-	0.0275	8	0	-	-	-	-	XCV1342	37/217	-
1508464	+	0.0113	6	1	XCV1343	23	XCV1342	248/6	-	-	-
1509902	+	0.0000	34	3	XCV1345	97	-	-	-	-	-
1529221	+	0.0200	8	0	-	-	XCV1357	388/1945	-	-	-
1543610	+	0.0094	6	0	-	-	XCV1365	301/493	-	-	-
1546532	+	0.0000	41	11	XCV1368	22	-	-	-	-	-
1549846	-	0.0000	19	0	XCV1369	30	-	-	-	-	-
1549864	+	0.0081	6	1	XCV1370	189	-	-	XCV1369	48/2126	-
1550020	+	0.0162	5	1	XCV1370	33	-	-	-	-	-
1553394	+	0.0000	25	3	XCV1373	50	-	-	-	-	-
1555991	+	0.0086	7	2	XCV1375	180	XCV1374	651/146	-	-	-
1567734	+	0.0048	3	0	XCV1387	38	XCV1386	1047/35	-	-	-
1573503	-	0.0021	3	0	-	-	XCV1393	482/2016	-	-	-
1580759	-	0.0050	6	1	XCV1399	248	XCV1400	322/16	-	-	-
1581048	-	0.0441	5	1	-	-	XCV1400	33/305	-	-	-
1582399	-	0.0000	14	2	XCV1401	20	-	-	-	-	-
1582406	+	0.0046	5	1	XCV1402	128	-	-	XCV1401	27/1199	-
1589016	+	0.0002	9	1	XCV1406	69	-	-	-	-	-
1589822	+	0.0000	10	1	XCV1408	22	-	-	-	-	-
1590522	-	0.0020	3	0	-	-	XCV1409	338/225	-	-	-
1591652	-	0.0054	5	1	XCV1410	154	-	-	-	-	-
1593751	-	0.0000	7	0	XCV1413	68	-	-	-	-	-
1595169	+	0.0009	7	0	XCV1415	21	-	-	XCV1414	76/1140	-

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1601206	+	0.0360	5	2	XCV1423 XCV1424	0 264	-	-	-	-	-
1601450	+	0.0022	5	0	XCV1424	20	-	-	-	-	-
1609450	+	0.0000	17	0	-	-	-	-	XCV1430	151/391	-
1614026	-	0.0069	3	0	-	-	XCV1434	128/1002	-	-	-
1618771	-	0.0010	5	1	XCV1440	52	-	-	-	-	-
1618915	+	0.0038	4	0	-	-	-	-	XCV1441	2116/35	-
1624149	-	0.0192	5	0	-	-	-	-	XCV1446	1/678	-
1629856	-	0.0009	5	0	XCV1450	22	-	-	-	-	-
1631636	-	0.0147	3	0	XCV1452	8	-	-	-	-	-
1631992	+	0.0137	3	1	XCV1453	10	-	-	-	-	-
1639397	+	0.0060	4	1	XCV1459	28	-	-	XCV1458	46/645	-
1641414	-	0.0122	3	0	XCV1461	44	-	-	-	-	-
1642409	-	0.0000	20	5	XCV1462	22	-	-	-	-	-
1644256	-	0.0088	3	0	-	-	XCV1463	1905/1682	-	-	-
1648431	-	0.0007	5	0	-	-	XCV1465	8/1308	-	-	-
1653692	-	0.0000	18	6	XCV1470	209	XCV1471	1221/122	-	-	-
1658315	-	0.0000	49	5	XCV1475	25	-	-	-	-	-
1659232	-	0.0166	3	1	XCV1476	32	-	-	-	-	-
1659403	+	0.0034	4	0	-	-	-	-	XCV1477	1427/93	-
1660847	-	0.0038	6	2	XCV1477	17	-	-	-	-	-
1663194	-	0.0000	15	4	XCV1479	242	-	-	-	-	-
1669473	+	0.0000	12	1	XCV1485	0	-	-	-	-	-
1676072	+	0.0012	7	0	XCV1490	220	-	-	-	-	-
1684643	+	0.0000	70	11	XCV1495	21	-	-	-	-	-
1690056	+	0.0113	4	0	XCV1499	63	-	-	XCV1498	51/899	-
1690128	+	0.0290	3	0	-	-	XCV1499	9/437	-	-	-
1700939	-	0.0019	3	0	XCV1506	26	-	-	XCV1507	39/1895	-
1702829	+	0.0009	5	0	XCV1508	16	XCV1507	1929/5	-	-	-
1704019	-	0.0037	4	0	XCV1509	69	-	-	-	-	-
1704140	+	0.0005	8	1	XCV1510	38	-	-	-	-	-
1708364	-	0.0001	18	7	XCV1513	70	-	-	-	-	-
1710153	-	0.0031	3	0	XCV1515	25	-	-	-	-	-
1712207	-	0.0034	4	0	XCV1516	0	-	-	-	-	-
1712476	+	0.0277	4	2	XCV1517	30	-	-	-	-	-
1719682	+	0.0000	20	2	XCV1525	0	-	-	XCV1524	75/500	-
1721663	+	0.0000	20	5	XCV1528	34	-	-	-	-	-
1724343	-	0.0193	3	0	-	-	-	-	XCV1530	504/1025	-
1726530	+	0.0027	3	0	XCV1532	29	-	-	-	-	-
1733457	+	0.0000	12	2	XCV1537	39	-	-	-	-	-
1736086	+	0.0264	3	0	XCV1539	131	-	-	XCV1538	1/1030	-
1745530	+	0.0024	3	0	-	-	XCV1545	189/41	-	-	-
1750856	+	0.0000	21	2	-	-	XCV1551	817/13	XCV1552	537/46	-
1753470	-	0.0000	31	1	-	-	-	-	-	-	+
1756674	+	0.0000	18	6	XCV1557	73	-	-	XCV1556	12/488	-
1756686	-	0.0000	27	2	XCV1556	0	-	-	XCV1557	61/489	-
1758176	-	0.0000	17	1	XCV1559	24	-	-	XCV1560	81/488	-
1758200	+	0.0493	3	0	XCV1560	57	-	-	XCV1559	48/440	-
1762031	+	0.0060	8	4	XCV1563	36	-	-	-	-	-
1762679	+	0.0002	7	1	XCV1564	48	-	-	-	-	-
1768866	-	0.0152	5	0	-	-	XCV1568	81/197	-	-	-
1778677	-	0.0061	3	0	-	-	XCV1576	903/536	-	-	-
1779063	-	0.0195	3	0	-	-	XCV1576	517/922	-	-	-
1783856	-	0.0000	18	5	-	-	XCV1578	65/2904	-	-	-

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1783909	+	0.0004	5	0	-	-	-	-	XCV1578 XCV1579	12/2957 978/52	-
1788756	-	0.0008	4	0	XCV1582	28	-	-	XCV1583	7/1117	-
1791356	-	0.0000	13	0	XCV1584	147	-	-	-	-	-
1793470	-	0.0004	7	0	-	-	XCV1586	5/441	-	-	-
1797706	-	0.0000	9	0	XCV1591	280	-	-	-	-	-
1798406	+	0.0000	19	7	XCV1593	47	-	-	-	-	-
1803085	+	0.0019	5	1	XCV1598	11	-	-	-	-	-
1814647	+	0.0433	9	0	-	-	XCV1606	149/1542	-	-	-
1816736	+	0.0000	17	0	XCV1607	1	-	-	-	-	-
1819516	+	0.0183	3	0	XCV1608	23	-	-	-	-	-
1820299	+	0.0000	12	0	XCV1609	22	-	-	-	-	-
1821313	+	0.0001	9	1	-	-	XCV1610	140/3054	-	-	-
1825238	-	0.0001	8	1	-	-	XCV1611	6/632	-	-	-
1830728	-	0.0186	3	0	-	-	XCV1617	37/1051	-	-	-
1833891	+	0.0000	87	4	-	-	XCV1620	63/338	-	-	-
1838797	+	0.0000	40	12	XCV1626	51	-	-	-	-	-
1843191	+	0.0014	5	0	XCV1631	0	-	-	XCV1630	24/596	-
1859472	-	0.0028	7	2	XCV1646	0	-	-	-	-	-
1859602	+	0.0008	6	1	XCV1647	26	-	-	-	-	-
1864572	+	0.0002	7	0	-	-	XCV1654	8/699	-	-	-
1868556	-	0.0000	40	4	XCV1659	30	-	-	-	-	-
1868712	+	0.0002	7	1	XCV1660	19	-	-	-	-	-
1874518	+	0.0001	15	4	XCV1665	38	-	-	-	-	-
1875752	-	0.0005	6	0	-	-	-	-	XCV1666	458/78	-
1880820	-	0.0193	3	0	-	-	-	-	XCV1669 XCV1670	1001/42 70/753	-
1882951	+	0.0000	12	3	XCV1672	98	-	-	-	-	-
1895289	-	0.0047	5	1	XCV1679	25	-	-	XCV1680	36/1403	-
1895299	+	0.0072	4	0	XCV1680	26	-	-	XCV1679	35/1237	-
1900120	-	0.0065	5	1	XCV1684	90	-	-	-	-	-
1903361	+	0.0000	26	5	XCV1689	21	-	-	-	-	-
1908861	-	0.0008	5	0	XCV1693	19	-	-	XCV1694	93/1370	-
1908930	+	0.0000	31	1	XCV1694	24	-	-	XCV1693	88/756	-
1911229	+	0.0005	7	1	XCV1696	12	-	-	XCV1695	89/670	-
1912983	+	0.0148	3	1	-	-	XCV1697	180/1178	-	-	-
1914145	+	0.0003	19	9	XCV1698	113	XCV1697	1342/16	-	-	-
1914895	-	0.0000	170	21	-	-	-	-	XCV1699	425/432	-
1915239	+	0.0039	5	1	-	-	XCV1699	769/88	-	-	-
1921894	-	0.0000	44	2	-	-	XCV1704	297/491	-	-	-
1925949	+	0.0000	11	1	XCV1707	18	-	-	-	-	-
1938634	-	0.0204	3	1	XCV1719	95	-	-	-	-	-
1961959	+	0.0022	3	0	XCV1741	21	-	-	-	-	-
1964107	-	0.0151	3	0	-	-	XCV1743	400/937	-	-	-
1964739	+	0.0021	3	0	XCV1744	41	-	-	-	-	-
1966625	+	0.0056	3	0	XCV1746	28	-	-	-	-	-
1971387	-	0.0000	8	0	-	-	XCV1748	108/1910	-	-	-
1971505	+	0.0000	15	1	-	-	-	-	XCV1748	10/2028	-
1972062	+	0.0000	13	1	XCV1749	22	-	-	-	-	-
1974295	-	0.0437	7	0	-	-	-	-	XCV1750	326/501	-
1975236	+	0.0034	4	0	XCV1752	47	-	-	-	-	-
1976107	+	0.0055	5	0	-	-	XCV1752	824/465	-	-	-
1980337	+	0.0000	22	0	XCV1758	17	-	-	XCV1757	83/622	-
1984126	-	0.0005	4	0	XCV1763	24	-	-	XCV1764	42/671	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
1984810	+	0.0051	8	3	XCV1765	45	-	-	-	-	-
1988882	+	0.0042	9	2	XCV1769 XCV1768	299 9	-	-	-	-	-
1989065	+	0.0275	4	0	XCV1769	116	XCV1768	174/101	-	-	-
1993650	+	0.0050	4	1	XCV1772	0	-	-	-	-	-
1996311	+	0.0230	7	3	XCV1775	74	-	-	-	-	-
1996655	+	0.0419	5	1	-	-	XCV1775	270/2375	-	-	-
1998924	+	0.0002	18	1	XCV1776	249	XCV1775	2539/106	-	-	-
1999090	+	0.0000	30	4	XCV1776	83	-	-	-	-	-
2011844	-	0.0138	4	0	XCV1789	264	XCV1790	374/249	-	-	-
2012054	-	0.0424	4	0	-	-	XCV1790	164/459	-	-	-
2014218	-	0.0000	15	3	XCV1793	19	-	-	-	-	-
2018225	+	0.0012	6	1	XCV1798	30	-	-	-	-	-
2041528	+	0.0004	8	2	XCV1810	30	-	-	-	-	-
2046892	-	0.0002	8	1	-	-	-	-	XCV1814	28/1392	-
2046910	+	0.0095	3	0	XCV1814	10	-	-	-	-	-
2048788	+	0.0014	5	1	XCV1816	24	-	-	-	-	-
2050617	+	0.0000	11	0	XCV1818	26	-	-	-	-	-
2056210	+	0.0105	3	0	XCV1823	3	-	-	-	-	-
2061162	+	0.0026	5	1	-	-	-	-	XCV1825	1738/259	-
2063383	+	0.0003	5	0	-	-	XCV1826	165/809	-	-	-
2064776	-	0.0056	3	0	-	-	XCV1827	166/397	-	-	-
2066165	-	0.0265	6	3	XCV1828	0	-	-	-	-	-
2066489	+	0.0029	7	0	XCV1829	0	-	-	-	-	-
2076935	-	0.0010	4	0	XCV1838	55	-	-	-	-	-
2077009	+	0.0344	6	0	-	-	-	-	-	-	+
2077238	+	0.0116	7	1	XCV1839	103	-	-	-	-	-
2078061	+	0.0225	4	1	-	-	-	-	XCV1840	838/8	-
2078080	-	0.0000	14	1	-	-	XCV1840	819/11	-	-	-
2080165	+	0.0001	20	9	XCV1843	29	-	-	XCV1842	17/930	-
2080915	-	0.0000	170	21	-	-	-	-	XCV1844	425/432	-
2081259	+	0.0038	5	1	XCV1845	156	XCV1844	769/88	-	-	-
2081955	+	0.0103	3	0	-	-	XCV1845	540/11	XCV1846	816/76	-
2088259	+	0.0013	5	0	XCV1852	36	-	-	-	-	-
2094165	+	0.0004	6	0	XCV1859	103	-	-	-	-	-
2100728	+	0.0004	5	0	-	-	XCV1861	3048/8078	-	-	-
2108278	-	0.0259	3	0	-	-	-	-	XCV1861	10598/528	-
2109973	+	0.0000	20	2	XCV1864	151	XCV1863	817/13	-	-	-
2111307	+	0.0000	17	4	XCV1866	226	-	-	-	-	-
2118507	-	0.0193	3	0	-	-	XCV1871	90/668	-	-	-
2118674	+	0.0000	9	0	XCV1872	31	-	-	XCV1871	77/835	-
2133276	-	0.0001	11	2	XCV1886	0	-	-	-	-	-
2134171	-	0.0080	5	0	-	-	XCV1887	625/802	-	-	-
2136428	-	0.0036	5	0	XCV1888	83	-	-	XCV1889	100/648	-
2136615	-	0.0356	4	0	XCV1888	270	-	-	XCV1889	87/461	-
2138655	+	0.0022	5	0	-	-	XCV1891	50/1437	-	-	-
2140105	+	0.0393	3	1	XCV1892	25	-	-	-	-	-
2149915	+	0.0034	11	0	-	-	XCV1904	174/1199	-	-	-
2153351	+	0.0000	33	2	XCV1907	111	-	-	-	-	-
2163985	+	0.0000	45	9	XCV1915	22	-	-	-	-	-
2171527	+	0.0078	4	1	-	-	-	-	XCV1920	303/563	-
2173745	-	0.0019	6	0	XCV1921	56	-	-	XCV1922	21/488	-
2173764	+	0.0177	3	1	-	-	XCV1922	40/469	XCV1921	75/1754	-
2177450	-	0.0000	7	0	XCV1924	34	-	-	-	-	-

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2177544	+	0.0048	6	1	XCV1926 XCV1925	265 20	-	-	-	-	-
2181033	+	0.0009	4	0	XCV1928	17	-	-	-	-	-
2185929	+	0.0206	3	0	-	-	-	-	XCV1931	407/186	-
2190346	-	0.0000	10	0	XCV1934	56	-	-	-	-	-
2191418	+	0.0014	4	0	-	-	-	-	XCV1936	728/54	-
2192469	+	0.0054	3	0	-	-	-	-	XCV1937	282/20	-
2201842	+	0.0206	3	0	-	-	-	-	XCV1941	2237/234	-
2221729	-	0.0018	3	0	-	-	XCV1952	1758/416	-	-	-
2226060	+	0.0031	3	0	-	-	-	-	XCV1954	1097/1173	-
2236742	-	0.0198	5	0	XCV1962	157	-	-	-	-	-
2239894	+	0.0087	3	0	-	-	XCV1963	2441/657	-	-	-
2245024	+	0.0000	19	2	-	-	XCV1969	817/13	-	-	-
2246333	+	0.0376	3	1	XCV1972	25	-	-	-	-	-
2247573	-	0.0193	3	0	-	-	-	-	XCV1973	582/893	-
2283523	-	0.0087	3	0	XCV2000	53	-	-	-	-	-
2298800	-	0.0002	6	0	XCV2016	37	-	-	-	-	-
2299637	-	0.0000	29	3	-	-	XCV2017	8/654	-	-	-
2303997	-	0.0000	15	1	XCV2022	88	-	-	-	-	-
2316851	-	0.0000	7	0	XCV2034	0	-	-	-	-	-
2317891	+	0.0138	4	1	XCV2036	22	-	-	-	-	-
2319528	+	0.0009	3	0	-	-	XCV2038	789/434	-	-	-
2331979	+	0.0000	28	2	-	-	XCV2048	60/323	-	-	-
2332427	+	0.0011	9	3	XCV2049	21	-	-	-	-	-
2332843	-	0.0400	4	0	-	-	-	-	XCV2049	395/1884	-
2332879	+	0.0100	5	0	-	-	XCV2049	431/1848	-	-	-
2335263	-	0.0000	35	5	XCV2051	40	-	-	-	-	-
2339189	-	0.0139	3	1	-	-	XCV2055	457/508	-	-	-
2339660	+	0.0000	10	1	XCV2056	270	-	-	XCV2055	14/979	-
2345698	+	0.0167	5	2	-	-	-	-	XCV2059	3686/477	-
2349615	+	0.0006	5	0	XCV2060	0	-	-	-	-	-
2350997	+	0.0135	5	2	XCV2061	171	-	-	-	-	-
2356401	+	0.0000	27	1	XCV2066	90	-	-	-	-	-
2358828	+	0.0000	30	1	XCV2069	108	XCV2068	677/108	-	-	-
2359054	+	0.0064	4	0	-	-	XCV2069	118/739	-	-	-
2359959	+	0.0145	11	6	-	-	XCV2070	92/543	-	-	-
2363433	+	0.0023	8	0	XCV2073	0	-	-	-	-	-
2363454	+	0.0023	8	0	-	-	XCV2073	21/1196	-	-	-
2368751	+	0.0309	4	0	-	-	XCV2075	2367/293	-	-	-
2373126	-	0.0028	4	0	XCV2080	22	-	-	-	-	-
2374746	+	0.0005	6	0	XCV2083	29	-	-	-	-	-
2382135	+	0.0134	3	0	XCV2090	23	-	-	XCV2089	58/1032	-
2384696	-	0.0077	4	0	-	-	-	-	XCV2092	15/1220	-
2393149	+	0.0000	29	5	XCV2097	35	-	-	-	-	-
2395930	+	0.0076	4	1	XCV2101	25	-	-	-	-	-
2396500	+	0.0018	5	0	-	-	XCV2101	545/150	-	-	-
2398280	-	0.0085	3	0	-	-	XCV2102	315/1493	-	-	-
2399163	+	0.0000	17	2	XCV2103	185	-	-	-	-	-
2400264	-	0.0193	3	0	-	-	-	-	XCV2103	916/9274	-
2411526	-	0.0190	3	0	XCV2105	0	-	-	-	-	-
2414657	-	0.0000	19	2	-	-	XCV2107	1507/1582	-	-	-
2476734	-	0.0043	4	0	-	-	-	-	XCV2155	1024/1990	-
2480236	-	0.0000	183	10	XCV2156	22	-	-	-	-	-
2480585	-	0.0122	5	1	-	-	XCV2157	769/88	-	-	-

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2480929	+	0.0000	162	23	-	-	-	-	XCV2157	425/432	-
2481679	-	0.0003	19	10	XCV2158	29	-	-	XCV2159	48/686	-
2483366	+	0.0000	12	0	-	-	XCV2161	708/938	-	-	-
2484379	-	0.0000	210	29	-	-	-	-	XCV2161 XCV2162	1721/75 5/222	-
2485206	-	0.0008	5	0	XCV2163	45	-	-	-	-	-
2485799	+	0.0003	6	0	-	-	XCV2165	73/2704	-	-	-
2490307	-	0.0101	3	0	-	-	-	-	XCV2166	1804/247	-
2495693	-	0.0096	4	0	-	-	-	-	-	-	+
2499035	-	0.0056	8	2	XCV2173	21	-	-	XCV2174	76/978	-
2511922	-	0.0000	15	0	XCV2183	98	-	-	XCV2184	18/938	-
2523234	+	0.0469	7	0	-	-	XCV2193	308/1317	-	-	-
2529317	+	0.0002	6	0	XCV2200	43	-	-	-	-	-
2531282	-	0.0031	5	1	XCV2201	124	XCV2202	2254/121	-	-	-
2532845	-	0.0013	7	2	-	-	XCV2202	691/1684	-	-	-
2533596	-	0.0000	83	7	XCV2202	60	-	-	XCV2203	83/901	-
2534689	+	0.0009	11	4	XCV2204	0	-	-	-	-	-
2536986	+	0.0000	11	2	-	-	XCV2207	27/632	-	-	-
2537657	+	0.0000	11	0	XCV2208	19	-	-	-	-	-
2540067	+	0.0068	3	0	XCV2210 XCV2211	22 228	-	-	-	-	-
2542431	+	0.0000	15	2	XCV2214	32	-	-	-	-	-
2548437	-	0.0272	6	0	XCV2215	25	-	-	-	-	-
2549014	-	0.0000	26	5	XCV2216	19	-	-	-	-	-
2552155	+	0.0018	5	0	XCV2222	125	XCV2221	185/54	-	-	-
2557002	-	0.0193	3	0	-	-	-	-	XCV2230	276/854	-
2570503	-	0.0055	3	0	-	-	XCV2237	457/1099	-	-	-
2573369	+	0.0000	65	12	XCV2240 XCV2241	14 241	-	-	XCV2239	26/895	-
2573375	-	0.0408	4	2	XCV2239	32	-	-	XCV2240	8/175	-
2580164	+	0.0023	3	0	-	-	-	-	XCV2250	1090/29	-
2580190	-	0.0020	3	0	-	-	-	-	-	-	+
2583951	+	0.0000	19	5	XCV2252	149	XCV2251	2402/144	-	-	-
2589488	+	0.0004	7	1	-	-	XCV2258	81/1415	-	-	-
2590477	+	0.0005	5	0	-	-	XCV2258	1070/426	-	-	-
2595113	-	0.0028	5	1	-	-	XCV2262	769/88	XCV2261	1384/68	-
2595457	+	0.0000	162	23	-	-	-	-	XCV2262	425/432	-
2596207	-	0.0004	19	10	XCV2263	23	XCV2264	759/146	-	-	-
2608263	+	0.0000	52	5	XCV2275	22	-	-	-	-	-
2609179	-	0.0441	4	3	-	-	-	-	XCV2275	894/170	-
2610400	-	0.0000	108	15	XCV2277	23	-	-	-	-	-
2613410	+	0.0000	33	2	XCV2280	23	-	-	-	-	-
2613918	-	0.0087	4	0	-	-	-	-	XCV2280	485/588	-
2616298	-	0.0486	4	4	-	-	XCV2283	1577/546	-	-	-
2616321	+	0.0280	7	4	-	-	-	-	XCV2283	1554/569	-
2625423	+	0.0094	3	0	-	-	-	-	XCV2290	44/796	-
2629491	+	0.0001	20	9	XCV2295	29	-	-	XCV2294	88/345	-
2630241	-	0.0000	170	21	-	-	-	-	XCV2296	425/432	-
2630585	+	0.0061	5	1	XCV2297	207	XCV2296	769/88	-	-	-
2631345	+	0.0001	8	0	-	-	-	-	-	-	+
2631738	+	0.0076	5	1	XCV2298	211	-	-	-	-	-
2634620	-	0.0125	3	0	XCV2299	37	-	-	XCV2300	68/358	-
2642688	+	0.0005	12	4	XCV2309	26	-	-	-	-	-
2645525	+	0.0293	6	0	-	-	-	-	XCV2311	235/721	-

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2646914	+	0.0085	3	0	XCV2314	1	-	-	-	-	-
2651289	-	0.0067	3	0	XCV2319	31	-	-	-	-	-
2651497	+	0.0000	151	12	-	-	XCV2320	11/780	-	-	-
2651654	+	0.0232	8	1	-	-	XCV2320	168/623	-	-	-
2652046	-	0.0009	6	1	-	-	-	-	XCV2320	560/231	-
2656549	+	0.0112	8	4	-	-	-	-	XCV2325	877/35	-
2656965	+	0.0000	12	0	-	-	-	-	XCV2325	461/381	-
2659679	+	0.0004	7	1	-	-	-	-	XCV2329	559/232	-
2660227	-	0.0000	151	14	-	-	XCV2329	11/780	XCV2330	10/501	-
2671874	+	0.0004	7	1	-	-	-	-	XCV2340	559/232	-
2672422	-	0.0000	151	14	-	-	XCV2340 XCV2341	11/780 831/50	-	-	-
2676073	+	0.0000	17	4	XCV2345	205	XCV2344	9/209	-	-	-
2680420	+	0.0140	4	1	-	-	-	-	XCV2350	539/474	-
2680631	+	0.0092	3	0	-	-	-	-	XCV2350	328/685	-
2682681	-	0.0000	1040	80	XCV2352	34	-	-	XCV2353	69/860	-
2682761	+	0.0000	151	12	-	-	XCV2353	11/780	-	-	-
2683310	-	0.0014	6	1	-	-	-	-	XCV2353	560/231	-
2684765	+	0.0021	3	0	XCV2355	119	-	-	-	-	-
2694331	+	0.0395	7	0	-	-	-	-	XCV2365	346/145	-
2740409	+	0.0004	9	1	-	-	-	-	-	-	+
2740875	+	0.0037	7	3	-	-	-	-	-	-	+
2744537	-	0.0193	3	0	XCV2419	9	-	-	XCV2420	39/401	-
2744576	+	0.0318	7	4	XCV2420	0	-	-	XCV2419	48/488	-
2745041	+	0.0026	4	0	XCV2421	0	-	-	-	-	-
2751220	-	0.0000	33	2	-	-	-	-	XCV2424	30/1013	-
2756332	+	0.0000	14	3	XCV2429	0	-	-	XCV2428	89/442	-
2764488	+	0.0011	5	0	XCV2434	22	-	-	-	-	-
2767760	-	0.0000	8	1	XCV2435	187	-	-	-	-	-
2768957	+	0.0122	8	1	-	-	-	-	-	-	+
2768989	-	0.0017	10	3	-	-	-	-	XCV2436	1026/67	-
2770178	-	0.0242	5	0	XCV2437	22	-	-	-	-	-
2779019	+	0.0351	4	0	-	-	-	-	XCV2445	298/565	-
2780755	+	0.0206	3	0	-	-	-	-	XCV2448	701/321	-
2788996	+	0.0193	3	1	XCV2461	27	-	-	-	-	-
2798003	+	0.0002	20	9	XCV2477	29	-	-	XCV2476	49/789	-
2798753	-	0.0000	170	21	-	-	-	-	XCV2478	425/432	-
2799097	+	0.0038	5	1	-	-	XCV2478	769/88	XCV2479	274/68	-
2801025	+	0.0293	5	0	-	-	XCV2482	18/1007	-	-	-
2803434	-	0.0110	4	1	-	-	XCV2484	242/957	-	-	-
2803838	-	0.0000	24	7	XCV2484	162	-	-	-	-	-
2805975	-	0.0102	3	0	-	-	XCV2486	1154/300	-	-	-
2806891	-	0.0465	3	0	-	-	XCV2486	238/1216	-	-	-
2807153	-	0.0000	12	1	XCV2486	24	-	-	-	-	-
2811009	-	0.0111	3	1	XCV2489	296	XCV2490	771/119	-	-	-
2811397	-	0.0049	3	0	-	-	XCV2490	383/507	-	-	-
2811810	-	0.0000	28	1	XCV2490	30	XCV2491	1877/27	-	-	-
2813614	+	0.0206	3	0	-	-	-	-	XCV2491 XCV2492	73/1831 1055/90	-
2817226	-	0.0044	9	9	-	-	XCV2496	1085/597	-	-	-
2818655	-	0.0000	20	5	-	-	XCV2497	522/158	-	-	-
2819280	+	0.0000	54	6	XCV2498	70	-	-	-	-	-
2819686	+	0.0000	11	0	XCV2499	31	-	-	-	-	-
2833387	-	0.0215	5	1	XCV2509	242	XCV2510	2840/135	-	-	-

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2837417	-	0.0039	7	2	XCV2511	34	-	-	-	-	-
2838360	-	0.0015	4	0	XCV2514	31	-	-	-	-	-
2838423	+	0.0000	16	1	-	-	-	-	XCV2514	94/390	-
2843234	-	0.0000	29	2	XCV2518	24	-	-	-	-	-
2843374	+	0.0051	8	0	XCV2519	0	-	-	-	-	-
2853818	+	0.0003	8	1	XCV2531	0	-	-	-	-	-
2859395	+	0.0011	5	1	-	-	XCV2535	11/1653	-	-	-
2865835	-	0.0152	5	0	XCV2539	37	-	-	-	-	-
2874392	-	0.0011	4	0	-	-	XCV2546	54/554	-	-	-
2880224	-	0.0000	36	5	XCV2551	264	-	-	-	-	-
2880816	-	0.0000	17	3	-	-	XCV2552	80/540	-	-	-
2880924	+	0.0000	9	0	XCV2553	15	-	-	XCV2552	28/648	-
2885613	-	0.0125	6	1	-	-	-	-	XCV2557 XCV2558	622/43 46/336	-
2894891	+	0.0000	45	4	XCV2568	229	-	-	XCV2567	73/690	-
2894892	+	0.0006	30	0	XCV2568	228	-	-	XCV2567	74/691	-
2897112	-	0.0028	3	0	-	-	-	-	XCV2569	681/332	-
2897115	+	0.0019	3	0	-	-	XCV2569	684/329	-	-	-
2899257	+	0.0068	3	0	-	-	XCV2571	1189/478	-	-	-
2901871	-	0.0041	5	0	-	-	XCV2574	299/417	-	-	-
2905203	-	0.0074	4	0	XCV2577	95	-	-	XCV2578	8/1033	-
2910245	+	0.0043	4	0	XCV2582	71	-	-	XCV2581	47/808	-
2915414	-	0.0000	21	2	XCV2586	0	-	-	-	-	-
2915525	+	0.0037	5	1	XCV2587	0	-	-	-	-	-
2917592	-	0.0001	6	0	XCV2588	30	-	-	-	-	-
2919213	-	0.0000	11	1	-	-	XCV2589	117/1553	XCV2590	36/1154	-
2921819	-	0.0000	349	6	XCV2592	157	-	-	-	-	-
2924141	+	0.0383	3	1	XCV2595	16	-	-	-	-	-
2927298	+	0.0210	3	0	XCV2597	4	-	-	XCV2596	94/969	-
2929160	+	0.0000	24	4	XCV2599	24	-	-	-	-	-
2931916	-	0.0004	5	0	XCV2601	29	XCV2602	1848/26	-	-	-
2933879	-	0.0041	4	0	XCV2602	115	-	-	-	-	-
2935603	-	0.0000	17	4	XCV2603	0	-	-	-	-	-
2937657	-	0.0056	4	1	XCV2605	0	-	-	XCV2606	25/1089	-
2941785	-	0.0000	30	0	XCV2608	20	-	-	-	-	-
2944333	+	0.0008	5	0	XCV2611	33	-	-	XCV2610	86/2068	-
2945265	+	0.0005	5	0	-	-	XCV2611	899/189	-	-	-
2947490	+	0.0206	3	0	-	-	-	-	XCV2612	790/1825	-
2952716	+	0.0000	22	2	-	-	XCV2618	817/13	-	-	-
2953172	-	0.0002	6	0	-	-	-	-	XCV2619	75/347	-
2953268	+	0.0000	16	2	XCV2620	279	XCV2619	21/251	-	-	-
2958333	+	0.0424	4	0	-	-	-	-	XCV2625	769/886	-
2959110	+	0.0295	3	0	XCV2626	290	-	-	XCV2625	8/1663	-
2966234	-	0.0001	8	1	XCV2630	0	-	-	-	-	-
2967021	-	0.0000	29	3	XCV2631	73	-	-	-	-	-
2970083	-	0.0000	34	8	XCV2634	71	-	-	-	-	-
2971502	-	0.0000	31	4	XCV2635	0	-	-	-	-	-
2972659	-	0.0031	4	0	-	-	XCV2636	105/962	-	-	-
2976928	+	0.0008	7	0	XCV2642	28	-	-	-	-	-
3000010	+	0.0000	45	7	XCV2661	23	-	-	-	-	-
3005175	-	0.0036	5	0	-	-	XCV2665	479/363	-	-	-
3011678	+	0.0040	3	0	XCV2671	212	XCV2670	1236/65	-	-	-
3024175	-	0.0047	6	0	-	-	XCV2674	97/5035	-	-	-
3024425	+	0.0317	3	0	XCV2675	0	-	-	-	-	-

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3027493	+	0.0024	5	1	-	-	XCV2677	1074/2093	-	-	-
3038824	-	0.0029	3	0	XCV2683	25	-	-	-	-	-
3045902	-	0.0048	5	1	XCV2688	242	XCV2689	769/88	-	-	-
3046246	+	0.0000	162	23	-	-	-	-	XCV2689	425/432	-
3046996	-	0.0002	19	10	XCV2690	29	-	-	-	-	-
3047619	-	0.0002	16	6	-	-	-	-	-	-	+
3062769	+	0.0020	3	0	XCV2703	21	-	-	-	-	-
3065332	-	0.0000	14	0	XCV2704	0	XCV2705	110/0	-	-	-
3068986	-	0.0043	6	2	XCV2709	24	XCV2710	596/24	-	-	-
3069778	+	0.0000	10	1	XCV2711	43	-	-	-	-	-
3075487	+	0.0184	4	0	-	-	XCV2715	2113/2521	-	-	-
3078655	+	0.0001	7	0	XCV2716	22	-	-	-	-	-
3087900	+	0.0000	24	5	-	-	XCV2723	817/13	-	-	-
3093405	+	0.0044	4	1	-	-	-	-	XCV2725	892/1999	-
3094186	-	0.0004	6	0	-	-	XCV2725	111/2780	-	-	-
3094539	-	0.0000	8	0	XCV2725	242	-	-	-	-	-
3095807	-	0.0025	3	0	-	-	XCV2726	769/1048	-	-	-
3103362	-	0.0033	3	0	-	-	XCV2729	22/4333	-	-	-
3106814	-	0.0000	16	1	XCV2730	214	XCV2731	164/84	-	-	-
3108818	-	0.0114	3	0	-	-	XCV2732	620/1734	-	-	-
3112840	+	0.0171	4	1	-	-	-	-	XCV2736	959/672	-
3113638	+	0.0206	3	0	-	-	-	-	XCV2736	161/1470	-
3129258	-	0.0000	71	9	XCV2748	0	-	-	-	-	-
3130321	+	0.0015	5	0	-	-	XCV2750	16/1636	-	-	-
3142239	+	0.0017	3	0	XCV2763	4	-	-	-	-	-
3143314	+	0.0000	47	7	XCV2764	19	-	-	-	-	-
3151089	-	0.0019	4	0	-	-	XCV2770	46/1018	-	-	-
3151886	+	0.0278	4	1	-	-	-	-	XCV2771 XCV2772	11/616 838/8	-
3151905	-	0.0000	17	2	XCV2771	30	XCV2772	819/11	-	-	-
3166096	-	0.0024	3	0	-	-	XCV2786	1363/58	-	-	-
3168287	-	0.0000	20	1	XCV2787	133	-	-	-	-	-
3174731	-	0.0001	7	1	XCV2794	263	XCV2795	1753/148	-	-	-
3184488	-	0.0001	10	2	-	-	-	-	XCV2800	857/1221	-
3193300	+	0.0000	20	2	-	-	XCV2806	817/13	-	-	-
3196040	-	0.0001	12	1	XCV2809	257	-	-	-	-	-
3198093	-	0.0001	8	0	XCV2811	25	-	-	-	-	-
3201040	-	0.0005	7	1	XCV2812	90	-	-	XCV2813	21/518	-
3201949	+	0.0156	4	1	-	-	-	-	XCV2814	203/201	-
3202518	+	0.0117	3	0	-	-	-	-	XCV2815	3413/330	-
3205497	-	0.0050	3	0	-	-	XCV2815	434/3309	-	-	-
3206053	+	0.0226	4	1	-	-	-	-	XCV2816	838/8	-
3206074	-	0.0000	15	2	XCV2815	143	XCV2816	817/13	-	-	-
3207206	-	0.0292	6	2	XCV2817	16	-	-	-	-	-
3209078	+	0.0000	12	0	-	-	-	-	XCV2820	266/210	-
3209434	-	0.0000	28	2	XCV2820	90	XCV2821	425/90	-	-	-
3209880	-	0.0000	14	0	XCV2821	21	-	-	-	-	-
3211387	-	0.0000	69	15	XCV2822	18	-	-	-	-	-
3214823	-	0.0020	3	0	XCV2823	123	-	-	-	-	-
3217042	-	0.0146	3	1	XCV2826	57	-	-	XCV2827	64/639	-
3219491	-	0.0000	15	1	XCV2829	21	-	-	XCV2830	38/304	-
3219506	+	0.0001	7	0	XCV2831 XCV2830	289 23	-	-	XCV2829	36/536	-
3223002	-	0.0142	3	0	-	-	XCV2833	1321/790	-	-	-

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3223483	-	0.0042	4	0	-	-	XCV2833	840/1271	-	-	-
3227994	-	0.0041	6	1	-	-	XCV2837	1172/1539	-	-	-
3229766	-	0.0087	3	0	-	-	XCV2838	1007/501	-	-	-
3231114	-	0.0002	6	1	-	-	XCV2839	254/333	-	-	-
3233114	-	0.0008	3	0	-	-	XCV2840	192/1268	-	-	-
3239149	-	0.0102	4	1	-	-	XCV2846	433/655	-	-	-
3247561	-	0.0000	10	1	XCV2854	185	XCV2855	604/148	-	-	-
3248192	-	0.0005	9	0	XCV2855	27	-	-	XCV2856	86/883	-
3258473	+	0.0169	3	1	-	-	-	-	XCV2864	21/905	-
3259835	-	0.0036	5	1	XCV2865	289	XCV2866	769/88	-	-	-
3260179	+	0.0000	162	23	-	-	-	-	XCV2866	425/432	-
3260929	-	0.0004	19	10	XCV2867	29	XCV2868	266/75	-	-	-
3262486	-	0.0007	10	3	XCV2869	31	-	-	XCV2870	74/997	-
3267661	-	0.0000	16	3	-	-	XCV2874	105/1976	-	-	-
3270081	-	0.0406	4	2	-	-	XCV2876	462/647	-	-	-
3272560	-	0.0050	4	1	XCV2878	23	-	-	XCV2879	58/705	-
3272598	+	0.0103	4	1	XCV2879	20	-	-	XCV2878	61/984	-
3275968	-	0.0158	6	3	XCV2881	21	-	-	-	-	-
3276220	+	0.0041	5	1	XCV2882	20	-	-	-	-	-
3278748	+	0.0019	5	1	XCV2884	23	-	-	-	-	-
3284009	-	0.0058	3	0	XCV2887	42	-	-	-	-	-
3285246	+	0.0000	12	1	XCV2890	23	-	-	-	-	-
3285826	+	0.0001	8	1	XCV2891	21	-	-	-	-	-
3290997	-	0.0001	12	2	-	-	-	-	-	-	+
3291196	+	0.0042	4	1	XCV2895	107	-	-	-	-	-
3292629	+	0.0216	3	0	-	-	XCV2896	652/1678	-	-	-
3296402	+	0.0003	5	0	XCV2898	28	-	-	-	-	-
3298935	+	0.0001	7	0	-	-	XCV2899	236/2115	-	-	-
3301211	+	0.0002	8	1	XCV2900	27	-	-	-	-	-
3304175	-	0.0012	14	0	XCV2901	18	-	-	-	-	-
3304605	-	0.0124	3	0	XCV2902	0	-	-	-	-	-
3309674	+	0.0010	4	0	XCV2907	27	-	-	XCV2906	83/559	-
3309679	-	0.0000	34	2	XCV2906	88	-	-	XCV2907	22/561	-
3310327	+	0.0000	35	2	XCV2908	26	-	-	-	-	-
3311814	-	0.0012	7	2	XCV2909	23	-	-	-	-	-
3316609	+	0.0010	4	0	-	-	XCV2914	211/1138	-	-	-
3317782	+	0.0000	48	8	XCV2916 XCV2915	262 0	-	-	-	-	-
3320950	-	0.0001	12	3	XCV2919	23	-	-	-	-	-
3323956	-	0.0145	3	1	-	-	XCV2923	1168/274	-	-	-
3325124	-	0.0022	8	3	XCV2923	0	-	-	XCV2924	51/848	-
3332244	+	0.0000	19	1	XCV2929	17	-	-	-	-	-
3334516	-	0.0000	17	1	XCV2931	0	-	-	-	-	-
3336764	-	0.0000	16	0	XCV2933	20	-	-	-	-	-
3339362	-	0.0000	14	0	XCV2936	0	-	-	-	-	-
3341585	-	0.0005	9	2	-	-	XCV2938	1132/1663	-	-	-
3343425	+	0.0027	7	0	XCV2940	28	-	-	-	-	-
3348398	+	0.0001	12	3	XCV2944	173	-	-	-	-	-
3349667	-	0.0231	3	0	-	-	-	-	XCV2944	1096/1102	-
3355947	+	0.0211	3	0	-	-	XCV2950	704/711	-	-	-
3358746	+	0.0002	11	3	XCV2955	22	XCV2954	148/1	-	-	-
3373502	-	0.0004	6	0	XCV2966	25	-	-	-	-	-
3377499	+	0.0000	8	0	XCV2970	0	-	-	-	-	-
3379701	-	0.0001	7	1	-	-	XCV2971	37/1087	-	-	-

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3380658	-	0.0147	4	0	-	-	XCV2972	39/719	-	-	-
3381360	-	0.0035	3	0	-	-	XCV2973	1843/559	-	-	-
3385409	-	0.0000	12	1	-	-	XCV2974	46/619	XCV2975	46/619	-
3388971	-	0.0035	5	0	XCV2981	23	-	-	-	-	-
3390399	+	0.0035	3	0	XCV2983	0	-	-	-	-	-
3395819	-	0.0027	5	2	XCV2987	82	-	-	-	-	-
3402000	+	0.0000	13	3	-	-	XCV2992	179/66	-	-	-
3403164	-	0.0051	3	0	-	-	-	-	XCV2993	743/1092	-
3408906	-	0.0127	4	0	-	-	-	-	-	-	+
3410761	+	0.0000	15	3	XCV3000	6	-	-	-	-	-
3413795	-	0.0049	5	0	-	-	XCV3002	256/1246	-	-	-
3418545	+	0.0055	4	0	XCV3005	16	-	-	-	-	-
3424210	-	0.0016	5	0	-	-	XCV3010	63/599	-	-	-
3426246	+	0.0002	10	2	XCV3013	178	-	-	-	-	-
3429740	+	0.0009	4	0	XCV3016	0	-	-	-	-	-
3449971	-	0.0042	4	1	XCV3029	27	-	-	XCV3030	76/786	-
3450047	+	0.0067	4	1	XCV3030	0	-	-	-	-	-
3452981	+	0.0000	14	1	XCV3033	110	-	-	-	-	-
3455491	+	0.0025	6	0	XCV3036	21	-	-	-	-	-
3459115	-	0.0105	10	5	-	-	XCV3037	8/693	-	-	-
3473908	-	0.0000	57	0	XCV3049	55	-	-	-	-	-
3474110	+	0.0001	9	1	XCV3050	19	-	-	-	-	-
3477013	-	0.0179	3	1	-	-	XCV3053	848/555	-	-	-
3478034	-	0.0001	6	0	XCV3053	173	XCV3054	1163/189	-	-	-
3482294	-	0.0016	4	0	XCV3056	49	-	-	-	-	-
3485108	-	0.0003	5	0	-	-	XCV3059	65/621	-	-	-
3485693	-	0.0001	11	0	XCV3060	23	-	-	-	-	-
3487971	-	0.0010	7	0	XCV3063	0	-	-	-	-	-
3488858	+	0.0200	8	0	-	-	XCV3065	12/893	-	-	-
3491956	-	0.0038	3	0	-	-	XCV3067	280/847	-	-	-
3493428	-	0.0000	11	1	XCV3068	42	-	-	-	-	-
3496031	+	0.0002	13	8	-	-	XCV3071	421/346	-	-	-
3496288	+	0.0000	13	3	XCV3072	199	XCV3071	678/89	-	-	-
3497562	+	0.0024	3	0	XCV3074	22	-	-	-	-	-
3498161	+	0.0052	7	2	XCV3075	21	-	-	-	-	-
3505699	+	0.0000	18	4	XCV3083	0	-	-	-	-	-
3518387	-	0.0000	994	98	XCV3096	0	-	-	-	-	-
3525219	-	0.0000	10	0	XCV3103	14	-	-	-	-	-
3525280	+	0.0031	3	0	XCV3104	57	-	-	XCV3103	75/1184	-
3528942	+	0.0037	3	0	-	-	XCV3108	507/173	-	-	-
3531766	-	0.0000	12	1	XCV3111	281	XCV3112	8/246	XCV3113	32/1030	-
3533944	+	0.0020	4	1	XCV3116	2	XCV3115	551/21	-	-	-
3534466	+	0.0045	3	1	XCV3117	40	XCV3116	520/37	-	-	-
3536664	+	0.0000	95	12	XCV3119	129	XCV3118	1356/38	-	-	-
3539930	+	0.0284	24	15	XCV3123	0	-	-	-	-	-
3539954	+	0.0062	11	0	-	-	XCV3123	24/365	-	-	-
3544261	+	0.0001	10	1	XCV3127	23	-	-	-	-	-
3545188	-	0.0029	5	1	-	-	XCV3128	970/43	-	-	-
3551148	-	0.0015	4	0	XCV3131	23	-	-	-	-	-
3557788	-	0.0203	4	0	-	-	XCV3138	1006/385	-	-	-
3573400	-	0.0028	8	3	XCV3149	21	-	-	XCV3150	79/1449	-
3574632	-	0.0019	4	0	-	-	-	-	XCV3150	1153/217	-
3575831	-	0.0011	3	0	-	-	XCV3152	752/411	-	-	-
3581398	-	0.0058	4	1	XCV3159	23	-	-	-	-	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
3583485	+	0.0094	5	1	XCV3163 XCV3162	269 0	-	-	-	-	-
3590242	-	0.0437	7	0	XCV3165	21	-	-	-	-	-
3601774	-	0.0055	4	1	XCV3168	22	-	-	XCV3169	75/734	-
3601849	+	0.0033	8	3	XCV3169	0	-	-	XCV3168	97/3669	-
3603357	+	0.0017	3	0	-	-	XCV3171	276/290	-	-	-
3610521	-	0.0000	53	4	XCV3177	188	-	-	-	-	-
3616498	-	0.0000	26	5	XCV3182	24	XCV3183	982/61	-	-	-
3620330	+	0.0000	31	3	XCV3187	89	-	-	-	-	-
3620973	+	0.0033	9	13	-	-	XCV3187	554/2343	-	-	-
3622644	-	0.0072	3	0	-	-	-	-	XCV3187	2225/672	-
3622778	+	0.0000	6	3	-	-	XCV3187	2359/538	-	-	-
3624042	-	0.0000	14	2	-	-	XCV3188	42/605	-	-	-
3628252	-	0.0020	4	0	XCV3191	23	-	-	-	-	-
3629839	-	0.0000	9	0	XCV3192	69	-	-	-	-	-
3630079	+	0.0000	41	5	XCV3193	25	-	-	-	-	-
3630391	+	0.0204	4	0	-	-	XCV3193	287/1089	-	-	-
3634081	-	0.0009	6	1	XCV3195	36	-	-	-	-	-
3634438	-	0.0010	7	0	-	-	XCV3196	865/241	-	-	-
3635324	-	0.0001	10	0	XCV3196	21	-	-	-	-	-
3639271	+	0.0000	16	2	XCV3202	0	-	-	XCV3201	33/587	-
3639290	-	0.0003	10	2	XCV3201	52	-	-	XCV3202	19/1396	-
3643121	+	0.0000	43	3	-	-	-	-	XCV3205	79/246	-
3672452	-	0.0000	9	0	-	-	XCV3222	757/304	-	-	-
3673257	-	0.0074	4	1	XCV3222	48	-	-	-	-	-
3680238	-	0.0221	4	1	-	-	XCV3228	238/967	-	-	-
3682598	-	0.0019	6	1	-	-	XCV3229	5082/2132	-	-	-
3683097	-	0.0018	3	0	-	-	XCV3229	4583/2631	-	-	-
3684784	-	0.0103	3	1	-	-	XCV3229	2896/4318	-	-	-
3691234	-	0.0115	16	7	XCV3233	53	-	-	-	-	-
3691397	+	0.0047	4	0	XCV3234	21	-	-	-	-	-
3698089	-	0.0219	3	1	-	-	XCV3240	342/179	-	-	-
3700913	-	0.0008	7	1	XCV3241	21	-	-	-	-	-
3710992	+	0.0001	11	0	XCV3249	14	-	-	-	-	-
3713718	+	0.0000	10	1	XCV3251	45	-	-	-	-	-
3721917	-	0.0320	3	1	XCV3253	44	-	-	XCV3254	59/604	-
3725397	+	0.0001	7	0	-	-	XCV3257	7/421	-	-	-
3727405	+	0.0000	21	1	-	-	-	-	-	-	+
3728571	+	0.0000	32	1	XCV3260	18	-	-	-	-	-
3732545	+	0.0081	3	0	XCV3264	21	-	-	XCV3263	81/704	-
3735352	+	0.0000	13	1	XCV3268	26	-	-	XCV3267	23/1255	-
3738534	-	0.0000	177	26	XCV3271	42	-	-	-	-	-
3739927	-	0.0263	4	1	XCV3272	58	-	-	-	-	-
3744037	-	0.0006	6	1	XCV3277	0	-	-	-	-	-
3746306	-	0.0008	5	0	-	-	XCV3279	796/1105	-	-	-
3748830	-	0.0020	3	0	XCV3281	265	XCV3282	594/131	-	-	-
3749453	-	0.0000	22	4	XCV3282	29	-	-	-	-	-
3750836	-	0.0003	6	0	XCV3284	22	-	-	XCV3285	23/435	-
3763733	-	0.0193	3	0	XCV3293	20	-	-	XCV3294	81/953	-
3765356	-	0.0062	4	0	XCV3295	19	-	-	-	-	-
3767403	+	0.0000	14	2	XCV3297	91	-	-	XCV3296	41/2002	-
3793853	-	0.0018	4	0	XCV3315	51	XCV3316	466/55	-	-	-
3804274	-	0.0017	3	0	-	-	XCV3324	604/1978	-	-	-
3819837	+	0.0000	21	0	XCV3337	152	-	-	-	-	-

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3827398	-	0.0129	3	1	-	-	XCV3342	613/379	-	-	-
3828101	+	0.0000	14	0	XCV3343	32	-	-	XCV3342	90/1082	-
3832104	-	0.0006	5	0	XCV3345	170	XCV3346	132/155	-	-	-
3834377	+	0.0206	3	0	-	-	-	-	XCV3349	459/707	-
3837008	+	0.0000	18	0	-	-	XCV3351	60/1391	-	-	-
3840372	-	0.0000	59	3	XCV3352	22	-	-	-	-	-
3840688	+	0.0000	27	0	-	-	-	-	XCV3353	224/201	-
3841040	-	0.0000	41	9	XCV3353	128	-	-	-	-	-
3841288	-	0.0375	6	0	-	-	-	-	XCV3354	34/1225	-
3841563	+	0.0014	3	0	-	-	XCV3354	309/950	-	-	-
3842377	+	0.0037	4	1	XCV3355	146	XCV3354	1123/136	-	-	-
3845146	-	0.0280	3	1	-	-	-	-	XCV3357	953/84	-
3850198	+	0.0000	23	2	-	-	XCV3360	817/13	-	-	-
3854811	-	0.0000	12	1	XCV3366	29	-	-	-	-	-
3859554	-	0.0002	6	0	XCV3370	25	-	-	-	-	-
3861697	-	0.0004	6	0	-	-	-	-	-	-	+
3862413	+	0.0057	7	3	XCV3376	0	-	-	-	-	-
3866613	-	0.0029	3	0	XCV3380	25	-	-	-	-	-
3868645	-	0.0063	3	0	-	-	XCV3381	1945/2014	-	-	-
3871397	-	0.0028	5	1	XCV3382	115	XCV3383	769/88	-	-	-
3871741	+	0.0000	162	23	-	-	-	-	XCV3383	425/432	-
3872491	-	0.0072	19	10	XCV3384	29	-	-	XCV3385	17/252	-
3872641	-	0.0000	20	0	XCV3384	179	-	-	XCV3385	167/102	-
3873924	+	0.0000	20	2	-	-	XCV3387	817/13	XCV3388	285/46	-
3877920	-	0.0000	26	2	XCV3391	247	-	-	-	-	-
3880139	-	0.0000	52	3	-	-	XCV3392	46/1567	XCV3393	35/703	-
3880635	-	0.0105	3	0	-	-	-	-	XCV3393	461/207	-
3884056	+	0.0001	20	9	XCV3397	29	XCV3396	1586/75	-	-	-
3884806	-	0.0000	170	21	-	-	-	-	XCV3398	425/432	-
3885150	+	0.0048	5	1	-	-	XCV3398	769/88	-	-	-
3886210	+	0.0000	17	6	XCV3399	173	-	-	-	-	-
3886493	-	0.0037	4	0	-	-	-	-	XCV3399	110/1965	-
3897296	-	0.0000	42	7	XCV3408	24	-	-	-	-	-
3897851	+	0.0000	162	23	-	-	-	-	XCV3409	425/432	-
3898601	-	0.0002	19	10	XCV3410	29	-	-	-	-	-
3899941	-	0.0104	5	0	-	-	XCV3413	1081/61	XCV3412	888/64	-
3903440	+	0.0000	13	1	XCV3416 XCV3417	0 250	-	-	-	-	-
3908082	+	0.0002	7	0	XCV3421	27	-	-	-	-	-
3909587	+	0.0055	3	0	XCV3424	36	-	-	-	-	-
3910153	+	0.0074	4	1	XCV3425	25	-	-	-	-	-
3920060	-	0.0008	8	1	XCV3430	142	-	-	XCV3431	77/2038	-
3927580	+	0.0000	15	1	-	-	-	-	-	-	+
3931813	-	0.0000	14	1	XCV3437	22	-	-	-	-	-
3935639	+	0.0044	4	0	-	-	XCV3440	1960/1993	-	-	-
3939554	-	0.0018	5	1	-	-	-	-	XCV3442	1004/306	-
3944243	-	0.0036	4	0	XCV3444	21	-	-	-	-	-
3944604	-	0.0033	3	0	-	-	-	-	-	-	+
3956443	-	0.0040	8	3	-	-	XCV3452	111/434	-	-	-
3956726	+	0.0022	5	0	-	-	XCV3453	144/1214	-	-	-
3962678	+	0.0000	9	0	XCV3460	19	-	-	-	-	-
3963400	+	0.0000	26	4	XCV3461	0	-	-	-	-	-
3964399	-	0.0202	3	0	-	-	XCV3462	760/241	-	-	-
3965186	-	0.0002	5	0	XCV3462	27	-	-	-	-	-

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3969443	-	0.0020	3	0	XCV3465	258	-	-	-	-	-
3974418	-	0.0003	6	0	XCV3468	65	-	-	-	-	-
3975609	-	0.0000	36	5	XCV3469	45	-	-	-	-	-
3976811	+	0.0112	5	2	XCV3471	26	-	-	-	-	-
3977787	-	0.0001	9	1	-	-	-	-	XCV3472	209/402	-
3978409	+	0.0002	8	0	-	-	XCV3473	141/812	-	-	-
3980115	+	0.0007	5	0	XCV3475	23	-	-	-	-	-
3984582	-	0.0403	3	1	-	-	XCV3481	71/381	-	-	-
3991134	+	0.0046	3	0	XCV3486	0	-	-	-	-	-
3999749	-	0.0225	3	0	-	-	XCV3491	21/1304	-	-	-
4001676	+	0.0155	3	1	-	-	-	-	XCV3492	620/1125	-
4007577	-	0.0000	28	2	XCV3496	0	-	-	-	-	-
4008425	-	0.0006	6	3	-	-	XCV3497	1539/407	-	-	-
4012132	-	0.0017	6	0	XCV3500	174	XCV3501	884/171	-	-	-
4013200	+	0.0010	6	1	XCV3502	29	-	-	-	-	-
4018650	-	0.0000	242	26	XCV3505 XCV3506	272 27	-	-	-	-	-
4018651	-	0.0025	76	7	XCV3505 XCV3506	273 28	-	-	-	-	-
4019632	+	0.0011	8	2	-	-	-	-	XCV3507	1/934	-
4020256	-	0.0157	3	0	-	-	XCV3508	1653/497	-	-	-
4022331	-	0.0000	11	1	XCV3509	21	-	-	-	-	-
4024561	-	0.0001	9	1	XCV3510	0	-	-	-	-	-
4025035	-	0.0000	25	2	XCV3511	47	-	-	-	-	-
4026894	-	0.0018	7	2	XCV3513	22	-	-	XCV3514	88/810	-
4027759	-	0.0224	3	0	-	-	-	-	XCV3514 XCV3515	777/55 32/334	-
4037253	+	0.0022	14	7	XCV3525	218	XCV3524	207/218	-	-	-
4037865	+	0.0000	2627	442	-	-	-	-	-	-	+
4037913	-	0.0000	36	17	-	-	-	-	-	-	+
4041002	+	0.0024	3	0	-	-	XCV3530	147/842	-	-	-
4042199	+	0.0120	3	0	-	-	XCV3532	28/691	XCV3531	5/204	-
4046356	+	0.0469	7	0	-	-	-	-	XCV3534	558/2621	-
4061390	-	0.0014	4	0	XCV3544	28	-	-	XCV3545	45/914	-
4064047	-	0.0329	5	0	-	-	XCV3547	69/656	-	-	-
4064199	+	0.0071	5	1	XCV3548	0	-	-	XCV3547	83/808	-
4064434	+	0.0046	4	0	-	-	XCV3548	235/1849	-	-	-
4068348	+	0.0000	18	2	-	-	-	-	XCV3551	110/483	-
4069816	-	0.0013	7	1	XCV3552	0	-	-	-	-	-
4069950	+	0.0000	35	10	-	-	-	-	-	-	+
4070819	-	0.0003	7	0	XCV3553	78	-	-	XCV3554	82/1335	-
4072489	+	0.0000	27	2	XCV3556 XCV3555	240 94	-	-	-	-	-
4084692	-	0.0083	3	0	-	-	-	-	XCV3566	176/1173	-
4086583	-	0.0069	6	2	XCV3567	29	-	-	-	-	-
4089769	+	0.0000	44	3	XCV3570	32	-	-	-	-	-
4090439	+	0.0085	4	1	XCV3571	45	-	-	-	-	-
4098408	+	0.0006	5	0	-	-	-	-	XCV3574	31/2493	-
4100738	-	0.0000	25	1	XCV3575	184	-	-	-	-	-
4107864	-	0.0154	3	0	XCV3578	29	-	-	-	-	-
4108094	+	0.0000	30	8	-	-	-	-	-	-	+
4109000	+	0.0025	4	0	-	-	XCV3579	513/485	-	-	-
4112499	+	0.0105	3	0	XCV3583	186	XCV3582	929/186	-	-	-
4117962	-	0.0038	4	0	XCV3586	43	-	-	XCV3587	78/959	-

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TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
4122387	+	0.0009	8	2	XCV3589	0	-	-	-	-	-
4123145	+	0.0016	5	0	XCV3590	24	-	-	-	-	-
4130785	-	0.0035	9	3	XCV3596 XCV3595	28 236	-	-	-	-	-
4137131	-	0.0000	10	0	XCV3598	28	-	-	-	-	-
4160234	+	0.0144	3	0	-	-	XCV3617	377/2082	-	-	-
4165173	-	0.0193	3	0	XCV3619	34	-	-	-	-	-
4167914	+	0.0000	10	1	XCV3622	24	-	-	-	-	-
4171902	-	0.0026	5	1	-	-	XCV3625	15/773	XCV3626	61/984	-
4171936	+	0.0001	12	3	XCV3626	27	-	-	XCV3625	19/807	-
4176511	+	0.0483	3	0	-	-	XCV3630	293/984	-	-	-
4183458	-	0.0153	5	0	XCV3636	21	-	-	-	-	-
4199423	-	0.0272	6	0	XCV3647	2	-	-	-	-	-
4205166	-	0.0363	4	0	XCV3651	104	-	-	-	-	-
4213401	-	0.0000	14	3	-	-	XCV3658	2181/95	-	-	-
4214082	-	0.0152	3	1	-	-	XCV3658	1500/776	-	-	-
4223074	-	0.0022	3	0	-	-	XCV3668	1292/432	-	-	-
4224385	-	0.0043	3	0	XCV3668	19	-	-	-	-	-
4245230	-	0.0001	8	1	XCV3675	24	-	-	-	-	-
4247726	-	0.0193	3	0	-	-	XCV3678	165/131	-	-	-
4258733	+	0.0037	4	0	XCV3686	283	-	-	XCV3685	43/2833	-
4259551	+	0.0000	45	4	-	-	XCV3686	535/1456	-	-	-
4264986	-	0.0000	16	5	XCV3690	0	-	-	-	-	-
4265843	+	0.0000	23	1	XCV3692	21	-	-	-	-	-
4267656	+	0.0000	13	1	XCV3693	46	-	-	-	-	-
4269632	+	0.0001	7	0	-	-	XCV3695	102/1124	-	-	-
4270597	+	0.0017	4	0	XCV3696	248	XCV3695	1067/159	-	-	-
4274499	+	0.0003	8	1	XCV3700	27	-	-	XCV3699	89/685	-
4277595	-	0.0000	23	6	XCV3702	26	-	-	-	-	-
4279066	+	0.0042	5	2	XCV3704	31	-	-	-	-	-
4282216	-	0.0002	7	0	XCV3705	260	-	-	-	-	-
4283564	-	0.0041	3	1	-	-	XCV3707	189/365	-	-	-
4283579	+	0.0018	5	0	-	-	-	-	XCV3707	174/380	-
4284080	-	0.0011	5	0	-	-	XCV3708	557/327	-	-	-
4284757	-	0.0000	13	2	XCV3708	120	XCV3709	991/61	-	-	-
4285797	-	0.0017	5	0	XCV3709	49	-	-	-	-	-
4287776	+	0.0000	20	2	-	-	-	-	-	-	+
4292574	-	0.0066	3	0	-	-	-	-	XCV3716	871/1378	-
4293883	+	0.0001	7	0	-	-	XCV3716	2180/69	-	-	-
4296181	-	0.0047	6	2	XCV3718	82	-	-	-	-	-
4309263	-	0.0000	12	1	XCV3726	32	-	-	-	-	-
4309871	-	0.0055	9	3	XCV3727	186	-	-	-	-	-
4311306	+	0.0001	9	2	XCV3730	100	-	-	XCV3729	1/379	-
4311336	-	0.0000	8	0	XCV3729	29	-	-	XCV3730	70/1200	-
4314643	-	0.0000	12	0	XCV3732	23	-	-	-	-	-
4314757	+	0.0008	7	2	XCV3733	241	-	-	-	-	-
4316512	-	0.0299	3	0	-	-	-	-	XCV3734	10/360	-
4317032	+	0.0000	12	2	-	-	XCV3735	82/367	-	-	-
4318226	-	0.0018	5	1	XCV3736	37	-	-	-	-	-
4318420	+	0.0010	7	1	XCV3737	0	-	-	-	-	-
4321777	+	0.0001	8	1	XCV3741	33	-	-	-	-	-
4323846	-	0.0166	3	1	-	-	XCV3743	672/32	-	-	-
4326863	-	0.0448	3	1	XCV3744	182	-	-	-	-	-
4328091	+	0.0001	13	3	-	-	XCV3746	16/439	-	-	-

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4339048	+	0.0200	4	0	-	-	-	-	XCV3754	365/1461	-
4339435	-	0.0006	9	3	XCV3754	22	-	-	-	-	-
4349332	-	0.0151	3	0	XCV3762	23	-	-	XCV3763	92/502	-
4350995	-	0.0003	6	0	XCV3765	72	-	-	-	-	-
4358104	-	0.0422	3	1	-	-	XCV3773	121/676	-	-	-
4358669	-	0.0000	13	2	XCV3774	86	-	-	-	-	-
4364685	-	0.0000	45	0	XCV3779	31	-	-	-	-	-
4366433	+	0.0296	3	0	XCV3782	40	-	-	-	-	-
4367233	-	0.0359	11	0	-	-	-	-	XCV3782 XCV3783	760/77 89/694	-
4367263	+	0.0001	6	0	XCV3783	59	-	-	-	-	-
4367880	-	0.0082	8	1	-	-	-	-	XCV3783	558/47	-
4369125	-	0.0193	3	0	XCV3784	23	-	-	-	-	-
4369290	+	0.0000	12	2	-	-	-	-	-	-	+
4371772	-	0.0000	68	5	-	-	-	-	-	-	+
4376389	-	0.0017	4	0	XCV3789	20	-	-	-	-	-
4377581	+	0.0009	11	3	XCV3791	25	-	-	XCV3790	83/976	-
4385950	+	0.0030	4	0	-	-	-	-	XCV3800	30/1145	-
4386005	-	0.0017	5	0	XCV3800	25	-	-	-	-	-
4386948	-	0.0021	4	0	-	-	-	-	XCV3801	10/270	-
4388305	-	0.0077	4	0	-	-	XCV3802	292/1042	-	-	-
4392400	+	0.0000	11	1	XCV3807	41	-	-	XCV3806	13/652	-
4393809	-	0.0194	4	2	-	-	XCV3808	538/556	-	-	-
4394950	-	0.0017	4	0	-	-	XCV3809	686/600	-	-	-
4395515	-	0.0030	4	0	-	-	XCV3809	121/1165	-	-	-
4398395	+	0.0001	8	0	XCV3814	26	-	-	-	-	-
4400663	+	0.0000	23	2	-	-	XCV3816	151/604	-	-	-
4402676	+	0.0018	9	3	-	-	XCV3820	49/301	-	-	-
4403069	+	0.0130	3	0	-	-	-	-	-	-	+
4407078	-	0.0020	3	0	XCV3823	68	-	-	-	-	-
4409801	-	0.0009	4	0	-	-	XCV3825	422/1011	-	-	-
4411740	-	0.0008	7	1	XCV3826	0	-	-	-	-	-
4413048	-	0.0400	4	0	XCV3827	252	-	-	XCV3828	85/978	-
4420847	+	0.0009	5	0	XCV3836	104	-	-	-	-	-
4423078	-	0.0000	12	0	-	-	XCV3837	245/1347	-	-	-
4423362	-	0.0192	6	1	XCV3837	39	-	-	-	-	-
4454565	+	0.0000	13	1	XCV3871	226	-	-	-	-	-
4480327	-	0.0021	6	0	XCV3894	262	XCV3895	199/232	-	-	-
4485330	-	0.0000	11	0	XCV3903	218	-	-	-	-	-
4488575	+	0.0019	3	0	-	-	-	-	-	-	+
4492811	-	0.0000	41	3	XCV3912	37	-	-	-	-	-
4492830	+	0.0125	3	0	-	-	-	-	XCV3913 XCV3912	493/56 56/1930	-
4493424	+	0.0109	4	1	-	-	XCV3914	102/911	-	-	-
4498709	-	0.0005	7	0	XCV3919	22	-	-	-	-	-
4498825	+	0.0000	16	0	-	-	-	-	-	-	+
4499967	+	0.0085	3	0	XCV3921	43	-	-	XCV3920	39/1013	-
4500051	-	0.0006	6	1	XCV3920	45	-	-	XCV3921	41/780	-
4507411	-	0.0004	7	1	XCV3926	42	-	-	XCV3927	90/1223	-
4507470	+	0.0000	19	5	XCV3927	31	-	-	-	-	-
4509798	+	0.0000	54	10	XCV3929	99	XCV3928	1089/53	-	-	-
4515928	+	0.0000	10	1	-	-	XCV3934	156/773	-	-	-
4518620	-	0.0000	168	23	XCV3936	297	-	-	XCV3937	38/250	-
4521105	-	0.0000	18	0	XCV3938	37	-	-	-	-	-

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4522609	+	0.0016	6	0	XCV3940	114	-	-	XCV3939	80/1453	-
4526218	+	0.0424	4	0	XCV3943	0	-	-	-	-	-
4533862	-	0.0000	86	11	-	-	XCV3949	9/872	-	-	-
4537266	-	0.0001	8	0	XCV3953	0	-	-	-	-	-
4539141	+	0.0000	16	2	XCV3955	121	-	-	XCV3954	78/1796	-
4539754	-	0.0167	4	0	-	-	-	-	-	-	+
4542837	-	0.0000	25	4	XCV3957	22	-	-	-	-	-
4548120	-	0.0002	8	0	-	-	XCV3960	101/2127	-	-	-
4550453	-	0.0002	7	1	XCV3963	0	-	-	-	-	-
4551938	-	0.0012	6	1	XCV3964	21	-	-	-	-	-
4562352	-	0.0000	10	0	-	-	XCV3969	75/596	-	-	-
4562806	+	0.0469	7	0	-	-	-	-	XCV3970	425/138	-
4563705	+	0.0177	3	1	-	-	-	-	XCV3971	422/327	-
4564127	-	0.0008	4	0	XCV3971	0	-	-	-	-	-
4566249	-	0.0006	6	0	XCV3976	71	-	-	-	-	-
4578759	-	0.0060	3	0	XCV3988	27	-	-	-	-	-
4580486	-	0.0047	3	0	XCV3990	23	-	-	-	-	-
4580654	+	0.0015	22	14	XCV3991	88	-	-	-	-	-
4583277	+	0.0041	6	2	XCV3994	0	-	-	-	-	-
4585144	+	0.0165	3	0	-	-	XCV3995	114/1034	-	-	-
4587148	+	0.0089	4	1	-	-	XCV3996	796/781	-	-	-
4589368	-	0.0000	8	0	-	-	XCV3998	862/298	-	-	-
4590816	-	0.0271	9	5	XCV3999	0	-	-	-	-	-
4596722	-	0.0000	22	4	-	-	XCV4007	147/500	-	-	-
4596940	+	0.0002	14	5	XCV4008	21	-	-	XCV4007	71/718	-
4606228	+	0.0261	5	0	-	-	-	-	-	-	+
4606630	-	0.0003	7	1	-	-	-	-	-	-	+
4608071	-	0.0000	11	2	XCV4010	33	-	-	-	-	-
4608171	+	0.0000	21	4	XCV4011	38	-	-	-	-	-
4615446	+	0.0000	12	0	XCV4018	30	-	-	-	-	-
4626132	+	0.0272	3	1	XCV4029	1	-	-	-	-	-
4627723	+	0.0051	9	4	XCV4030	299	XCV4029	1590/116	-	-	-
4628082	+	0.0415	5	2	-	-	XCV4030	60/821	-	-	-
4634680	+	0.0206	3	0	-	-	-	-	XCV4035	178/1606	-
4636053	+	0.0000	17	3	-	-	-	-	XCV4036	1018/1186	-
4641920	-	0.0001	8	0	XCV4040	31	-	-	-	-	-
4646308	-	0.0070	4	1	XCV4043	27	-	-	-	-	-
4647214	+	0.0010	6	0	XCV4046	73	-	-	-	-	-
4649419	+	0.0057	3	0	-	-	XCV4048	26/528	-	-	-
4652571	-	0.0001	4	0	-	-	XCV4052	907/211	-	-	-
4654564	-	0.0343	9	6	-	-	XCV4053	18/1073	XCV4054	93/737	-
4654612	+	0.0110	3	0	XCV4054	45	-	-	XCV4053	30/1121	-
4664567	+	0.0141	3	0	XCV4067	173	-	-	-	-	-
4665917	+	0.0024	3	0	-	-	XCV4068	143/519	-	-	-
4670222	-	0.0002	6	0	XCV4073	78	-	-	-	-	-
4674977	+	0.0001	13	3	XCV4079	0	-	-	-	-	-
4676672	+	0.0078	3	0	-	-	XCV4080	916/451	-	-	-
4691793	-	0.0012	4	0	XCV4094	27	-	-	XCV4095	20/523	-
4694340	-	0.0000	28	4	XCV4096	32	-	-	-	-	-
4694851	+	0.0043	3	0	-	-	-	-	XCV4097	880/139	-
4697095	-	0.0028	3	0	XCV4098	24	-	-	-	-	-
4698077	-	0.0000	105	8	XCV4100 XCV4101	268 41	-	-	-	-	-
4699089	-	0.0000	14	1	XCV4102	23	-	-	-	-	-

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4701994	-	0.0000	23	7	-	-	-	-	XCV4106	81/257	-
4706354	+	0.0070	4	1	-	-	XCV4111	10/301	-	-	-
4711085	+	0.0206	3	0	-	-	-	-	XCV4115	902/510	-
4714461	-	0.0015	4	0	XCV4118	44	-	-	-	-	-
4715131	-	0.0000	43	10	XCV4119	29	-	-	XCV4120	31/573	-
4720514	-	0.0088	4	0	XCV4123	1	-	-	-	-	-
4727508	-	0.0162	4	1	-	-	-	-	-	-	+
4728553	-	0.0000	15	1	XCV4131	20	-	-	-	-	-
4729028	-	0.0051	6	2	XCV4132	22	-	-	-	-	-
4731606	-	0.0104	3	0	-	-	XCV4134	17/1809	-	-	-
4735763	-	0.0011	7	2	XCV4137	177	-	-	-	-	-
4735903	+	0.0000	23	2	XCV4138	67	-	-	-	-	-
4736549	+	0.0008	6	0	-	-	XCV4138	579/2432	-	-	-
4737286	+	0.0422	3	0	-	-	XCV4138	1316/1695	-	-	-
4738388	+	0.0401	5	4	-	-	XCV4138	2418/593	-	-	-
4738712	+	0.0402	4	0	-	-	XCV4138	2742/269	-	-	-
4740424	+	0.0070	3	0	-	-	XCV4140	560/450	-	-	-
4748322	-	0.0132	5	1	XCV4147	160	-	-	-	-	-
4748727	-	0.0177	3	0	-	-	-	-	XCV4148	153/317	-
4751394	-	0.0007	4	0	-	-	XCV4151	23/801	XCV4152	60/1052	-
4751412	+	0.0006	7	1	XCV4152	42	-	-	XCV4151	5/819	-
4757260	+	0.0000	21	2	-	-	-	-	-	-	+
4757357	-	0.0000	12	3	-	-	-	-	-	-	+
4761503	-	0.0156	4	1	-	-	-	-	-	-	+
4768772	+	0.0000	14	1	XCV4166	17	-	-	-	-	-
4775972	-	0.0100	3	0	-	-	XCV4167	36/4274	-	-	-
4776441	-	0.0000	37	12	-	-	-	-	-	-	+
4784392	-	0.0001	13	3	XCV4170 XCV4171	287 34	-	-	-	-	-
4788659	-	0.0193	3	1	-	-	-	-	XCV4175	1294/325	-
4793007	-	0.0025	3	0	-	-	XCV4181	2017/346	-	-	-
4796313	+	0.0079	3	0	-	-	-	-	XCV4184 XCV4185	69/713 829/142	-
4798298	-	0.0023	7	2	XCV4187	111	XCV4188	689/54	-	-	-
4799322	-	0.0160	3	0	XCV4189	13	-	-	-	-	-
4804093	+	0.0039	3	0	XCV4194	26	-	-	-	-	-
4808258	-	0.0073	5	1	-	-	-	-	XCV4197	626/36	-
4808628	+	0.0000	17	1	XCV4198	0	-	-	-	-	-
4813172	-	0.0110	6	3	XCV4200	0	-	-	XCV4201	77/364	-
4857728	-	0.0087	3	0	XCV4227	23	-	-	-	-	-
4863179	+	0.0000	19	2	-	-	XCV4235	817/13	-	-	-
4882281	-	0.0001	7	0	-	-	XCV4251	86/1434	-	-	-
4887536	-	0.0000	15	2	XCV4257	162	-	-	-	-	-
4891644	+	0.0008	8	2	-	-	-	-	-	-	+
4895203	-	0.0000	14	2	XCV4262	0	-	-	-	-	-
4898418	-	0.0007	8	2	XCV4264	0	-	-	-	-	-
4902712	+	0.0151	3	0	XCV4268	41	-	-	XCV4267	53/679	-
4904300	-	0.0041	5	1	XCV4270	9	-	-	-	-	-
4906974	+	0.0031	4	0	-	-	-	-	XCV4273	40/274	-
4913268	+	0.0000	10	1	XCV4279	23	-	-	-	-	-
4914278	-	0.0017	6	1	-	-	-	-	-	-	+
4923729	-	0.0000	10	0	XCV4289	42	-	-	-	-	-
4946992	-	0.0000	9	0	XCV4306	0	-	-	-	-	-
4955269	-	0.0002	7	1	XCV4311	25	-	-	XCV4312	65/1876	-

<sup>a</sup>, TSS position on the *Xcv* chromosome and the plasmid (pXCV183; marked with #), respectively. <sup>b</sup>, p-value indicates the significance of the annotated TSS (see METHODS). <sup>c</sup>, number of read starts at TSS position. <sup>d</sup>, TSS classification and the corresponding CDS. <sup>e</sup>, distance of the TSS to annotated start and stop codon of the corresponding CDS. <sup>f</sup>, a TSS is classified as orphan (+) if it is neither classified as primary, internal nor antisense TSS.

TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
4966194	-	0.0004	8	2	-	-	XCV4315	42/1799	-	-	-
4968288	-	0.0126	3	0	-	-	-	-	XCV4318	845/84	-
4970190	-	0.0000	21	2	XCV4320 XCV4321	270 27	-	-	-	-	-
4971181	-	0.0000	20	0	XCV4322	37	-	-	-	-	-
4971307	+	0.0006	6	1	-	-	XCV4323	9/953	-	-	-
4973545	-	0.0041	4	0	-	-	-	-	-	-	+
4980145	-	0.0129	5	0	XCV4332	86	-	-	XCV4333	50/2283	-
4994114	-	0.0156	3	0	XCV4342	22	-	-	-	-	-
4994366	+	0.0134	3	0	XCV4343	22	-	-	-	-	-
5006692	-	0.0001	6	0	XCV4353	22	-	-	-	-	-
5010792	+	0.0013	13	0	-	-	-	-	XCV4356	93/2675	-
5012026	+	0.0135	5	0	-	-	-	-	XCV4357	362/1053	-
5019614	+	0.0206	3	0	XCV4363	39	-	-	-	-	-
5024943	+	0.0000	9	0	-	-	XCV4369	15/200	-	-	-
5031007	+	0.0000	17	4	XCV4373	137	-	-	-	-	-
5032259	+	0.0000	24	1	XCV4374	105	-	-	-	-	-
5036212	-	0.0257	3	0	-	-	-	-	XCV4375	493/637	-
5042497	-	0.0288	3	1	-	-	XCV4379	311/978	-	-	-
5045604	-	0.0003	5	0	XCV4382	48	-	-	-	-	-
5047832	+	0.0013	11	4	XCV4384	0	-	-	-	-	-
5047838	-	0.0096	3	0	XCV4383	212	-	-	XCV4384	6/620	-
5051823	-	0.0000	15	0	-	-	XCV4387	28/1621	-	-	-
5060407	+	0.0261	5	0	-	-	-	-	-	-	+
5060810	-	0.0086	5	1	-	-	-	-	-	-	+
5068657	-	0.0001	8	0	XCV4395	63	-	-	-	-	-
5070952	-	0.0002	6	0	-	-	XCV4399	3606/446	-	-	-
5084851	-	0.0027	3	0	XCV4408	27	-	-	-	-	-
5086838	+	0.0032	3	0	-	-	XCV4411	726/1118	-	-	-
5087567	-	0.0028	3	0	-	-	-	-	XCV4411	1455/389	-
5088118	+	0.0132	4	0	XCV4412	38	-	-	-	-	-
5094547	-	0.0008	6	0	-	-	-	-	-	-	+
5102472	+	0.0003	4	0	-	-	XCV4425	136/127	-	-	-
5103051	+	0.0028	3	0	XCV4426	18	-	-	-	-	-
5105859	+	0.0000	10	1	-	-	-	-	XCV4430	391/94	-
5106753	-	0.0061	4	1	-	-	XCV4431	457/16	-	-	-
5108588	+	0.0068	4	0	XCV4434	28	-	-	XCV4433	4/625	-
5110351	-	0.0466	4	0	-	-	XCV4435	195/626	-	-	-
5110576	-	0.0061	5	0	XCV4435	30	-	-	-	-	-
5110677	+	0.0008	4	0	XCV4436	130	-	-	-	-	-
5116206	-	0.0185	11	0	XCV4440	222	XCV4441	2272/100	-	-	-
5126147	-	0.0050	4	1	-	-	XCV4443	137/3414	-	-	-
5131412	-	0.0017	7	0	XCV4444	107	-	-	-	-	-
5134201	-	0.0000	10	1	XCV4449	25	-	-	-	-	-
5134535	+	0.0000	17	5	XCV4450	24	-	-	-	-	-
5139920	-	0.0007	5	0	XCV4457	21	-	-	-	-	-
5142226	-	0.0414	10	0	XCV4459	138	-	-	-	-	-
5146436	+	0.0002	6	0	XCV4466	26	-	-	-	-	-
5153717	-	0.0193	3	0	XCV4468	112	-	-	-	-	-
5154683	-	0.0047	3	0	XCV4469	22	-	-	-	-	-
5156544	-	0.0002	8	0	-	-	XCV4470	25/1768	-	-	-
5164171	-	0.0166	3	1	XCV4476	0	-	-	-	-	-
5164230	+	0.0072	3	0	XCV4477	206	-	-	XCV4476	59/1222	-
5165820	+	0.0130	4	1	XCV4478	0	-	-	-	-	-

<sup>a</sup>, TSS position on the *Xcv* chromosome and the plasmid (pXCV183; marked with #), respectively. <sup>b</sup>, p-value indicates the significance of the annotated TSS (see METHODS). <sup>c</sup>, number of read starts at TSS position. <sup>d</sup>, TSS classification and the corresponding CDS. <sup>e</sup>, distance of the TSS to annotated start and stop codon of the corresponding CDS. <sup>f</sup>, a TSS is classified as orphan (+) if it is neither classified as primary, internal nor antisense TSS.

TSS position <sup>a</sup>	Strand	p-value <sup>b</sup>	Library 2 <sup>c</sup>	Library 1 <sup>c</sup>	Primary <sup>d</sup> to CDS	5' UTR (bp)	Internal <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Antisense <sup>d</sup> to CDS	Distance to start/stop <sup>e</sup> (bp)	Orphan <sup>f</sup>
5170524	+	0.0000	14	0	XCV4482	28	-	-	-	-	-
5178255	-	0.0000	293	20	XCV4486 XCV4487	210 31	-	-	-	-	-
5178392	+	0.0000	23	2	-	-	-	-	-	-	+

## Reference

1. Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria* revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.

<sup>a</sup>, TSS position on the *Xcv* chromosome and the plasmid (pXCV183; marked with #), respectively. <sup>b</sup>, p-value indicates the significance of the annotated TSS (see METHODS). <sup>c</sup>, number of read starts at TSS position. <sup>d</sup>, TSS classification and the corresponding CDS. <sup>e</sup>, distance of the TSS to annotated start and stop codon of the corresponding CDS. <sup>f</sup>, a TSS is classified as orphan (+) if it is neither classified as primary, internal nor antisense TSS.

**Table S3.** Putative chromosomal TSSs within the first 50 bp of annotated CDSs (1).

CDS	Strand	Annotated translation start	TSS position	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	Distance of annotated start codon and TSS (bp)
XCV0016	+	17004	17027	42	18	24
XCV0022	+	23353	23398	13	1	46
XCV0035	-	38772	38760	13	0	13
XCV0117	+	133981	133989	3	1	9
XCV0209	+	244583	244587	15	2	5
XCV0322	+	369633	369663	3	0	31
XCV0327	-	375518	375494	3	0	25
XCV0383	+	436658	436669	4	0	12
XCV0387	-	441649	441623	4	0	27
XCV0493	+	557939	557964	3	0	26
XCV0522	-	589469	589427	10	0	43
XCV0537	+	602285	602321	16	1	37
XCV0579	+	656595	656637	6	2	43
XCV0632	-	713619	713610	3	0	10
XCV0648	+	732783	732826	4	0	44
XCV0653	-	736984	736976	3	0	9
XCV0781	+	893350	893392	6	0	43
XCV0816	-	931162	931121	3	0	42
XCV0821	-	936169	936154	12	1	16
XCV1073	-	1206623	1206581	3	0	43
XCV1074	+	1206680	1206687	11	1	8
XCV1141	+	1270899	1270936	47	4	38
XCV1155	-	1284633	1284606	4	1	28
XCV1239	-	1391023	1390988	3	0	36
XCV1332	-	1498481	1498473	8	0	9
XCV1400	-	1581081	1581048	5	1	34
XCV1465	-	1648439	1648431	5	0	9
XCV1499	+	1690119	1690128	3	0	10
XCV1586	-	1793475	1793470	7	0	6
XCV1611	-	1825244	1825238	8	1	7
XCV1617	-	1830765	1830728	3	0	38
XCV1654	+	1864564	1864572	7	0	9
XCV1922	+	2173724	2173764	3	1	41
XCV2017	-	2299645	2299637	29	3	9
XCV2073	+	2363433	2363454	8	0	22
XCV2207	+	2536959	2536986	11	2	28
XCV2320	+	2651486	2651497	151	12	12
XCV2329	-	2660238	2660227	151	14	12
XCV2340	-	2672433	2672422	151	14	12
XCV2341	-	2673253	2672422	151	14	12
XCV2353	+	2682750	2682761	151	12	12
XCV2482	+	2801007	2801025	5	0	19
XCV2535	+	2859384	2859395	5	1	12
XCV2729	-	3103384	3103362	3	0	23
XCV2750	+	3130305	3130321	5	0	17
XCV2770	-	3151135	3151089	4	0	47
XCV2971	-	3379738	3379701	7	1	38
XCV2972	-	3380697	3380658	4	0	40
XCV2974	-	3385412	3385409	12	1	4
XCV3037	-	3459123	3459115	10	5	9
XCV3065	+	3488846	3488858	8	0	13
XCV3123	+	3539930	3539954	11	0	25
XCV3188	-	3624084	3624042	14	2	43
XCV3257	+	3725390	3725397	7	0	8
XCV3392	-	3880185	3880139	52	3	47
XCV3491	-	3999770	3999749	3	0	22
XCV3532	+	4042171	4042199	3	0	29
XCV3625	-	4171917	4171902	5	1	16

<sup>a</sup>, number of read starts at TSS position.

CDS	Strand	Annotated translation start	TSS position	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	Distance of annotated start codon and TSS (bp)
XCV3746	+	4328075	4328091	13	3	17
XCV3949	-	4533871	4533862	86	11	10
XCV4048	+	4649393	4649419	3	0	27
XCV4053	-	4654582	4654564	9	6	19
XCV4111	+	4706344	4706354	4	1	11
XCV4134	-	4731623	4731606	3	0	18
XCV4151	-	4751417	4751394	4	0	24
XCV4167	-	4776008	4775972	3	0	37
XCV4315	-	4966236	4966194	8	2	43
XCV4323	+	4971298	4971307	6	1	10
XCV4369	+	5024928	5024943	9	0	16
XCV4387	-	5051851	5051823	15	0	29
XCV4470	-	5156569	5156544	8	0	26

### Reference

1. Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria* revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.

<sup>a</sup>, number of read starts at TSS position.



**Table S4.** Predicted antisense TSSs located close to the 3' end of annotated CDSs (+/- 100 bp) (1).

TSS position <sup>a</sup>	Strand	Library 2 <sup>b</sup>	Library 1 <sup>b</sup>	Distance of CDS end to TSS <sup>c</sup> (bp)	Gene product
81749#	-	9	2	43	XCVd0073: hypothetical protein
109851#	+	35	6	-99	XCVd0099: putative zeta toxin of the postsegregational killing system
127227#	-	4	2	-62	XCVd0115: Tn5044 transposase
159633#	+	3	0	-74	XCVd0154: hypothetical protein
180140#	+	3	0	-86	XCVd0171: hypothetical protein
11675	-	6	0	-72	XCV0008: energy transducer TonB protein
78522	+	23	2	-46	XCV0067: hypothetical protein
204493	+	12	2	-71	XCV0172: indolepyruvate ferredoxin oxidoreductase
261741	-	15	1	3	XCV0221: hypothetical protein
321787	-	7	0	-77	XCV0275: putative DNA-binding protein
485143	-	13	0	-94	XCV0435: HrcC protein
533893	+	6	0	-36	XCV0472: hypothetical protein
558968	-	8	1	-82	XCV0493: hypothetical protein
590048	-	10	0	-13	XCV0523: 50S ribosomal protein L13
707695	+	4	1	-8	XCV0625: IS1477 transposase
799920	-	6	2	-49	XCV0704: hypothetical protein
1041488	-	3	1	21	XCV0911: putative secreted protein
1095646	-	4	0	-86	XCV0964: hypothetical protein
1454260	-	3	1	-15	XCV1295: putative phage replication protein
1618915	+	4	0	-35	XCV1441: methionyl-tRNA synthetase
1659403	+	4	0	93	XCV1477: putative signal transduction protein
1750856	+	21	2	-46	XCV1552: hypothetical protein
1783909	+	5	0	-52	XCV1579: hypothetical protein
1875752	-	6	0	-78	XCV1666: hypothetical protein
1880820	-	3	0	-42	XCV1669: lysyl-tRNA synthetase
2078061	+	4	1	-8	XCV1840: IS1477 transposase
2081955	+	3	0	-76	XCV1846: ISxcC1 transposase
2191418	+	4	0	54	XCV1936: hypothetical protein
2192469	+	3	0	20	XCV1937: hypothetical protein
2484379	-	210	29	-75	XCV2161: conjugal transfer protein TrbL
2580164	+	3	0	-29	XCV2250: putative secreted protein
2595113	-	5	1	-68	XCV2261: phage-related integrase
2656549	+	8	4	-35	XCV2325: streptomycin 6-kinase
2768989	-	10	3	-67	XCV2436: IS1595 transposase
2799097	+	5	1	-68	XCV2479: putative replication protein C (fragment)
2813614	+	3	0	-90	XCV2492: glycosyltransferase
2885613	-	6	1	43	XCV2557: hypothetical protein
3151886	+	4	1	-8	XCV2772: IS1477 transposase
3206053	+	4	1	-8	XCV2816: IS1477 transposase
3845146	-	3	1	84	XCV3357: putative secreted protein
3873924	+	20	2	-46	XCV3388: hypothetical protein
3899941	-	5	0	-64	XCV3412: ISxac2 transposase
4027759	-	3	0	-55	XCV3514: ribonuclease PH
4367233	-	11	0	-77	XCV3782: OmpW family outer membrane protein
4367880	-	8	1	47	XCV3783: putative secreted protein
4492830	+	3	0	-56	XCV3913: D-tyrosyl-tRNA(Tyr) deacylase
4808258	-	5	1	-36	XCV4197: hypothetical protein
4968288	-	3	0	84	XCV4318: hypothetical protein
5105859	+	10	1	94	XCV4430: hypothetical protein

<sup>a</sup>, TSS position on the *Xcv* chromosome and the plasmid (pXCV183; marked with #), respectively. <sup>b</sup>, number of read starts at TSS position.

<sup>c</sup>, positive and negative sign indicates TSS within and downstream the CDS, respectively.

## Reference

1. Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria* revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.

**Table S5.** 5' UTRs of type III effector genes, leaderless genes with TSSs  $\leq 10$  bp upstream of the annotated start codon and genes containing long 5' UTRs (150 to 300 bp) referring to the annotated genome sequence of *Xcv* strain 85-10 (1). TSSs marked with # are located on pXCV183. TSSs of type III effector genes marked in bold exhibit a PIP box and -10 T/A-rich element as previously described (2). The PIP box of the type III effector gene *XCV2280* was identified in this study and is shown in brackets.

TSS position	Strand	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	5' UTR (bp)	Gene product
<b>5' UTRs of Type III Effectors</b>					
114827#	-	11	0	375	XCVd0104: avirulence protein AvrBs1
<b>62304</b>	+	3	0	23	XCV0052: avirulence protein AvrBs2
<b>486511</b>	+	5	0	678	XCV0437: xanthomonas outer protein D (XopD)
<b>532770</b>	+	8	1	0	XCV0471: avirulence protein AvrRxv
<b>660302</b>	+	12	2	341	XCV0581: xanthomonas outer protein B (XopB)
<b>1184957</b>	+	12	1	25	XCV1055: xanthomonas outer protein O (XopO)
<b>2480236</b>	-	183	10	22	XCV2156: xanthomonas outer protein J1 (XopJ1)
2613410 (2613343-2613403)	+	33	2	23	XCV2280: xanthomonas outer protein E2 (XopE2)
<b>2767760</b>	-	8	1	187	XCV2435: xanthomonas outer protein C (XopC)
3348398	+	12	3	173	XCV2944: xanthomonas outer protein N (XopN)
<b>4371772</b>	-	68	5	477	XCV3785: xanthomonas outer protein AA (XopAA)
<b>Leaderless Transcripts</b>					
193010	+	9	2	0	XCV0161: putative amidase/aminoacylase/peptidase family protein
201827	-	7	2	0	XCV0168: putative secreted protein
238550	-	7	0	0	XCV0201: uroporphyrinogen-III synthase
278015	-	6	0	0	XCV0236: predicted rRNA methylase
283462	+	4	0	0	XCV0243: acyl-CoA synthetase
288111	+	10	5	0	XCV0247: hypothetical protein
332993	+	9	1	0	XCV0287: predicted hydrolase of the alpha/beta superfamily
377721	+	4	0	3	XCV0330: putative NADH flavin oxidoreductase
411941	+	3	1	6	XCV0359: predicted Fe-S-cluster oxidoreductase
<b>532770</b>	+	8	1	0	XCV0471: avirulence protein AvrRxv
585332	+	8	5	0	XCV0518: phosphoserine phosphatase
599405	+	18	5	0	XCV0534: putative Iron-sulfur cluster assembly accessory protein
770365	+	4	0	0	XCV0679: porphobilinogen deaminase
779535	-	8	3	0	XCV0686: aspartate/tyrosine/aromatic aminotransferase
782120	+	7	0	0	XCV0688: hypothetical protein
823961	+	4	0	0	XCV0723: membrane-bound lytic transglycosylase
851592	+	8	0	0	XCV0747: 2-amino-4-hydroxy-6-hydroxymethylidihydropteridine pyrophosphokinase
884284	-	4	0	0	XCV0773: tRNA nucleotidyltransferase/poly(A) polymerase
959187	+	9	2	0	XCV0840: thiamine monophosphate synthase
997844	-	21	0	0	XCV0868: tRNA-dihydrouridine synthase
1028253	-	3	0	0	XCV0899: hypothetical protein
1029078	-	5	0	0	XCV0900: dimethyladenosine transferase (rRNA methylation)
1036560	+	6	0	0	XCV0907: 2-octaprenyl-6-methoxyphenyl hydroxylase
1102907	-	3	1	0	XCV0973: molybdenum cofactor biosynthesis protein B
1265655	-	3	1	0	XCV1134: MoxR-like ATPase
1419087	+	8	1	0	XCV1260: two-component system sensor protein
1450021	+	8	1	0	XCV1292: 4-hydroxy-3-methylbut-2-enyl diphosphate reductase
1504286	-	15	0	0	XCV1337: putative cytochrome C assembly protein
1601206	+	5	2	0	XCV1423: hypothetical protein
1631636	-	3	0	8	XCV1452: hypothetical protein
1631992	+	3	1	10	XCV1453: hypothetical protein
1669473	+	12	1	0	XCV1485: methionine aminopeptidase
1712207	-	4	0	0	XCV1516: ABC transporter permease and ATP-binding protein
1719682	+	20	2	0	XCV1525: NAD/FAD-binding protein
1756686	-	27	2	0	XCV1556: tmRNA-binding protein SmpB
1816736	+	17	0	1	XCV1607: RNase R

<sup>a</sup>, number of read starts at the TSS position.

TSS position	Strand	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	5' UTR (bp)	Gene product
1843191	+	5	0	0	XCV1631: hypothetical protein
1859472	-	7	2	0	XCV1646: carbonic anhydrase
1988882	+	9	2	9	XCV1768: RNA-binding protein Hfq
1993650	+	4	1	0	XCV1772: LexA repressor
2046910	+	3	0	10	XCV1814: tRNA nucleotidyltransferase/poly(A) polymerase
2056210	+	3	0	3	XCV1823: Glucan 1,4-beta-glucosidase precursor
2066165	-	6	3	0	XCV1828: two-component system sensor protein
2066489	+	7	0	0	XCV1829: 4-hydroxy-3-methylbut-2-en-1-yl diphosphate synthase
2133276	-	11	2	0	XCV1886: methylthioribulose-1-phosphate dehydratase
2316851	-	7	0	0	XCV2034: chemotaxis signal transduction protein
2349615	+	5	0	0	XCV2060: putative ATPase
2363433	+	8	0	0	XCV2073: molybdopterin biosynthesis protein
2411526	-	3	0	0	XCV2105: two-component system sensor kinase
2534689	+	11	4	0	XCV2204: hypothetical protein
2646914	+	3	0	1	XCV2314: ArsR family transcriptional regulator
2744537	-	3	0	9	XCV2419: FKBP-type peptidyl-prolyl cis-trans isomerase
2744576	+	7	4	0	XCV2420: putative membrane protein
2745041	+	4	0	0	XCV2421: predicted sulfurtransferase
2756332	+	14	3	0	XCV2429: thioredoxin-like protein
2843374	+	8	0	0	XCV2519: glutamine cyclotransferase
2853818	+	8	1	0	XCV2531: homoserine O-acetyltransferase
2915414	-	21	2	0	XCV2586: ABC transporter ATPase
2915525	+	5	1	0	XCV2587: ATP-dependent RNA helicase
2927298	+	3	0	4	XCV2597: glutamyl-tRNA synthetase
2935603	-	17	4	0	XCV2603: N-acetylmuramoyl-L-alanine amidase precursor
2937657	-	4	1	0	XCV2605: hypothetical protein
2966234	-	8	1	0	XCV2630: stringent starvation protein B
2971502	-	31	4	0	XCV2635: putative soluble lytic murein transglycosylase precursor
3024425	+	3	0	0	XCV2675: TetR family transcriptional regulator
3065332	-	14	0	0	XCV2704: cellulase precursor
3129258	-	71	9	0	XCV2748: proline racemase
3142239	+	3	0	4	XCV2763: HNH endonuclease family protein
3304605	-	3	0	0	XCV2902: putative 6-O-methylguanine-DNA methyltransferase
3317782	+	48	8	0	XCV2915: exodeoxyribonuclease VII small subunit
3325124	-	8	3	0	XCV2923: putative modulator of DNA gyrase
3334516	-	17	1	0	XCV2931: rRNA large subunit methyltransferase
3339362	-	14	0	0	XCV2936: rare lipoprotein B
3377499	+	8	0	0	XCV2970: hypothetical protein
3390399	+	3	0	0	XCV2983: 3-methyladenine DNA glycosylase
3410761	+	15	3	6	XCV3000: LysR family transcriptional regulator
3429740	+	4	0	0	XCV3016: hypothetical protein
3450047	+	4	1	0	XCV3030: deoxyribonuclease V
3487971	-	7	0	0	XCV3063: hypothetical protein
3505699	+	18	4	0	XCV3083: acetyltransferase (GNAT) family protein
3518387	-	994	98	0	XCV3096: ComEA-related DNA uptake protein
3533944	+	4	1	2	XCV3116: putative secreted protein
3539930	+	24	15	0	XCV3123: putative enzyme II of the phosphotransferase system
3583485	+	5	1	0	XCV3162: hypothetical protein
3601849	+	8	3	0	XCV3169: hypothetical protein
3639271	+	16	2	0	XCV3202: adenosylmethionine--8-amino-7-oxononoate transaminase
3744037	-	6	1	0	XCV3277: Holliday junction DNA helicase
3862413	+	7	3	0	XCV3376: hypothetical protein
3903440	+	13	1	0	XCV3416: thiamin S sulphur transfer protein
3963400	+	26	4	0	XCV3461: hypothetical protein
3991134	+	3	0	0	XCV3486: putative secreted protein
4007577	-	28	2	0	XCV3496: MoxR-like ATPase
4024561	-	9	1	0	XCV3510: guanosine-3',5'-bis(diphosphate) 3'-pyrophosphohydrolase

<sup>a</sup>, number of read starts at the TSS position.

TSS position	Strand	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	5' UTR (bp)	Gene product
4064199	+	5	1	0	XCV3548: putative membrane-associated phospholipid phosphatase
4069816	-	7	1	0	XCV3552: UDP-N-acetylmuramate--L-alanine ligase
4122387	+	8	2	0	XCV3589: TetR family transcriptional regulator
4199423	-	6	0	2	XCV3647: nicotinate phosphoribosyltransferase
4264986	-	16	5	0	XCV3690: CDP-diacylglycerol--serine O-phosphatidyltransferase
4318420	+	7	1	0	XCV3737: putative signal transduction protein
4411740	-	7	1	0	XCV3826: hypothetical protein
4526218	+	4	0	0	XCV3943: primosomal protein N
4537266	-	8	0	0	XCV3953: ABC transporter ATP-binding protein involved in cell division
4550453	-	7	1	0	XCV3963: hypothetical protein
4564127	-	4	0	0	XCV3971: HAD superfamily hydrolase
4583277	+	6	2	0	XCV3994: DNA primase
4590816	-	9	5	0	XCV3999: hypothetical protein
4626132	+	3	1	1	XCV4029: arginyl-tRNA synthetase
4674977	+	13	3	0	XCV4079: predicted hydrolase (HAD superfamily)
4720514	-	4	0	1	XCV4123: ATP-dependent DNA helicase
4808628	+	17	1	0	XCV4198: coproporphyrinogen III oxidase
4813172	-	6	3	0	XCV4200: DNA polymerase I
4895203	-	14	2	0	XCV4262: putative secreted protein
4898418	-	8	2	0	XCV4264: methyltransferase
4904300	-	5	1	9	XCV4270: exodeoxyribonuclease III
4946992	-	9	0	0	XCV4306: putative outer membrane protein
5047832	+	11	4	0	XCV4384: thymidine kinase
5164171	-	3	1	0	XCV4476: hypothetical protein
5165820	+	4	1	0	XCV4478: putative membrane protein

#### long 5' UTRs

101600#	+	5	0	287	XCVd0093: putative secreted protein
116378#	+	80	8	277	XCVd0107: ISxac2 transposase (fragment)
118654#	+	8	1	201	XCVd0110: hypothetical protein
123136#	+	5	0	218	XCVd0114: putative cointegrate resolution protein S
1404	+	15	1	202	XCV0002: DNA polymerase III subunit beta chain
78978	-	8	5	225	XCV0067: hypothetical protein
81973	-	6	2	270	XCV0069: putative secreted protein
117077	-	4	0	274	XCV0099: putative secreted protein
155750	+	11	2	227	XCV0131: hypothetical protein
253127	-	25	10	203	XCV0214: putative transposase
304274	+	5	0	150	XCV0264: malate synthase
320398	+	6	1	234	XCV0274: TetR family transcriptional regulator
448950	+	3	0	226	XCV0395: hypothetical protein
532651	-	5	0	151	XCV0470: ATP-dependent RNA helicase
647837	+	18	0	192	XCV0574: phospho-2-dehydro-3-deoxyheptonate aldolase
765474	+	3	0	282	XCV0675: glucans biosynthesis glucosyltransferase H
765583	+	3	0	173	XCV0675: glucans biosynthesis glucosyltransferase H
782575	+	13	2	244	XCV0689: diaminopimelate epimerase
829714	+	24	4	266	XCV0728: lipoate-protein ligase B
951830	+	10	2	181	XCV0835: cell division protein FtsZ
1042407	+	7	1	180	XCV0913: protocatechuate 4,5-dioxygenase beta chain
1116481	+	7	2	259	XCV0986: transcription antitermination factor
1137159	+	79	14	180	XCV1002: 50S ribosomal protein L2
1137964	+	9	2	209	XCV1003: 30S ribosomal protein S19
1149059	+	4	1	233	XCV1024: 50S ribosomal protein L17
1149619	+	3	0	180	XCV1025: hypothetical protein
1164003	-	5	0	163	XCV1036: hypothetical protein
1190312	+	8	2	228	XCV1061: Glutamine phosphoribosylpyrophosphate amidotransferase
1250212	+	5	1	153	XCV1120: hypothetical protein
1340207	+	6	1	233	XCV1198: hypothetical protein
1399557	-	4	2	291	XCV1243: glycine cleavage system P-protein

<sup>a</sup>, number of read starts at the TSS position.

TSS position	Strand	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	5' UTR (bp)	Gene product
1424474	-	4	0	171	XCV1265: putative D-alanyl-D-alanine carboxypeptidase
1549864	+	6	1	189	XCV1370: RNA polymerase ECF-type sigma factor RpoE2
1555991	+	7	2	180	XCV1375: putative membrane protein
1580759	-	6	1	248	XCV1399: putative secreted protein
1591652	-	5	1	154	XCV1410: hypothetical protein
1653692	-	18	6	209	XCV1470: outer membrane antigen
1663194	-	15	4	242	XCV1479: 30S ribosomal protein S2
1676072	+	7	0	220	XCV1490: asparagine synthetase B
1797706	-	9	0	280	XCV1591: GntR family transcriptional regulator
1998924	+	18	1	249	XCV1776: carbon storage regulator
2011844	-	4	0	264	XCV1789: hypothetical protein
2081259	+	5	1	156	XCV1845: putative membrane protein
2109973	+	20	2	151	XCV1864: hypothetical protein
2111307	+	17	4	226	XCV1866: bifunctional aspartokinase/homoserine dehydrogenase I
2136615	-	4	0	270	XCV1888: amino acid-polyamine-organocation superfamily protein
2236742	-	5	0	157	XCV1962: Flagellar motor component MotA
2339660	+	10	1	270	XCV2056: cell division protein FtsK
2350997	+	5	2	171	XCV2061: Putative membrane-associated chromosome condensation protein
2399163	+	17	2	185	XCV2103: putative filamentous hemagglutinin-like protein
2630585	+	5	1	207	XCV2297: ISxac2 transposase
2631738	+	5	1	211	XCV2298: phage-related integrase
2676073	+	17	4	205	XCV2345: putative cytosine-specific DNA methylase
<b>2767760</b>	-	8	1	187	XCV2435: xanthomonas outer protein C (XopC)
2803838	-	24	7	162	XCV2484: phage-related integrase
2811009	-	3	1	296	XCV2489: glutathione-regulated potassium efflux protein B
2833387	-	5	1	242	XCV2509: putative secreted protein
2880224	-	36	5	264	XCV2551: N-acetylornithine carbamoyltransferase
2894891	+	45	4	229	XCV2568: putative secreted protein
2894892	+	30	0	228	XCV2568: putative secreted protein
2921819	-	349	6	157	XCV2592: major cold shock protein
2953268	+	16	2	279	XCV2620: putative DNA-binding protein
2959110	+	3	0	290	XCV2626: 3-methyladenine DNA glycosylase
3011678	+	3	0	212	XCV2671: two-component system sensor histidine kinase-response regulator hybridprotein
3045902	-	5	1	242	XCV2688: hypothetical protein
3094539	-	8	0	242	XCV2725: TonB-dependent outer membrane receptor
3106814	-	16	1	214	XCV2730: TonB-dependent outer membrane receptor
3174731	-	7	1	263	XCV2794: translation initiation factor IF-3
3196040	-	12	1	257	XCV2809: putative secreted protein
3247561	-	10	1	185	XCV2854: preprotein translocase subunit SecG
3259835	-	5	1	289	XCV2865: tryptophan synthase subunit alpha
3348398	+	12	3	173	XCV2944: xanthomonas outer protein N (XopN)
3426246	+	10	2	178	XCV3013: putative cysteine protease precursor
3478034	-	6	0	173	XCV3053: UDP-N-acetylmuramoyl-L-alanyl-D-glutamate synthetase
3496288	+	13	3	199	XCV3072: membrane-bound metalloendopeptidase
3531766	-	12	1	281	XCV3111: hypothetical protein
3610521	-	53	4	188	XCV3177: homoserine O-acetyltransferase
3748830	-	3	0	265	XCV3281: Holliday junction resolvase
3819837	+	21	0	152	XCV3337: glucose dehydrogenase
3832104	-	5	0	170	XCV3345: NAD synthetase
3872641	-	20	0	179	XCV3384: ISxac3 transposase
3877920	-	26	2	247	XCV3391: hypothetical protein
3886210	+	17	6	173	XCV3399: hypothetical protein
3969443	-	3	0	258	XCV3465: phosphoglycerate kinase
4012132	-	6	0	174	XCV3500: Tfp pilus assembly protein PilN
4037253	+	14	7	218	XCV3525: hypothetical protein
4100738	-	25	1	184	XCV3575: thiamine biosynthesis protein ThiC

<sup>a</sup>, number of read starts at the TSS position.

TSS position	Strand	Library 2 <sup>a</sup>	Library 1 <sup>a</sup>	5' UTR (bp)	Gene product
4112499	+	3	0	186	XCV3583: 2-isopropylmalate synthase
4258733	+	4	0	283	XCV3686: soluble lytic murein transglycosylase precursor
4270597	+	4	0	248	XCV3696: ABC transporter ATP-binding protein
4282216	-	7	0	260	XCV3705: UDP-glucose dehydrogenase
4309871	-	9	3	186	XCV3727: hypothetical protein
4314757	+	7	2	241	XCV3733: ATP-dependent RNA helicase
4326863	-	3	1	182	XCV3744: oligopeptidase A
4413048	-	4	0	252	XCV3827: hypothetical protein
4454565	+	13	1	226	XCV3871: hypothetical protein
4480327	-	6	0	262	XCV3894: methionine aminopeptidase
4485330	-	11	0	218	XCV3903: hypothetical protein
4518620	-	168	23	297	XCV3936: tropinone reductase
4627723	+	9	4	299	XCV4030: putative secreted protein
4664567	+	3	0	173	XCV4067: hypothetical protein
4735763	-	7	2	177	XCV4137: putative glutathione transferase
4748322	-	5	1	160	XCV4147: putative membrane protein
4887536	-	15	2	162	XCV4257: 50S ribosomal protein L28
5047838	-	3	0	212	XCV4383: ATP-dependent DNA helicase
5116206	-	11	0	222	XCV4440: hypothetical protein
5164230	+	3	0	206	XCV4477: tellurium resistance protein

## References

1. Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria* revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.
2. Koebnik, R., Krüger, A., Thieme, F., Urban, A. and Bonas, U. (2006) Specific binding of the *Xanthomonas campestris* pv. *vesicatoria* AraC-type transcriptional activator HrpX to plant-inducible promoter boxes. *J. Bacteriol.*, **188**, 7652-7660.

<sup>a</sup>, number of read starts at the TSS position.

**Table S6.** Candidate riboswitches and widely conserved RNAs in *Xcv*.

Riboswitch (Rfam entry) <sup>a</sup>	Chromosomal position (strand) <sup>b</sup>	TSS position <sup>c</sup>	Downstream CDS <sup>d</sup>
FMN (RF00050)	911608-911787 (+)	-	Riboflavin synthase subunit alpha (XCV0800)
SAH (RF01057)	976910-977023 (-)	977026 (5/0)	S-adenosyl-L-homocysteine hydrolase (XCV0856)
Glycine (RF00504)	1399365-1399480 (-)	1399557 (4/2)	Glycine dehydrogenase (XCV1243)
	1399482-1399550 (-)	-	Glycine dehydrogenase (XCV1243)
SAM (RF00162)	3610402-3610523 (-)	3610521 (53/4)	Homoserine O-acetyltransferase (XCV3177)
Cobalamin (RF00174)	3801922-3802147 (-)	-	TonB-dependent outer membrane receptor (XCV3323)
TPP (RF00059)	4100645-4100743 (-)	4100738 (25/1)	Thiamine biosynthesis protein ThiC (XCV3575)
YybP-ykoY candidate (RF00080)	4924876-4924995 (+)	-	Hypothetical/putative membrane protein (XCV4291)
YybP-ykoY candidate (RF00080)	5164221-5164342 (+)	5164230 (3/0)	Tellurium resistance export protein (XCV4477)
sRNA (Rfam entry) <sup>a</sup>	Chromosomal position (strand) <sup>b</sup>	TSS position <sup>c</sup>	Chromosomal context <sup>e</sup>
RNaseP bacterial (RF00010)	933945-934295 (+)	933943 (86/27)	--> XCV0818 --> RNaseP --> XCV819
RtT (RF00391)	1235501-1235623 (+)	-	--> XCV1100 --> RtT --> XCV1101
SRP bacterial (RF00169)	1241912-1242175 (-)	1242175 (91/11)	--> XCV1109 <-- SRP <-- XCV1110
tmRNA (RF00023)	1753051-1753447 (-)	1753470 (31/1)	<-- XCV1553 <-- tmRNA --> XCV1554
6S RNA (RF00013)	4037865-4038049 (+)	4037865 (2627/442)	--> XCV3525 --> 6S RNA --> XCV3526

<sup>a</sup>, Putative riboswitches and widely conserved RNAs identified in *Rfam* database [see SI; (1)]. <sup>b</sup>, chromosomal position in *Xcv* corresponding to *Rfam* entry. <sup>c</sup>, TSS position in the *Xcv* chromosome (2). Numbers in brackets indicate read starts at TSS position (library 2/library 1). <sup>d</sup>, gene product of CDS annotated downstream of putative riboswitch (2). <sup>e</sup>, chromosomal orientation of sRNA gene and flanking CDSs indicated by arrows.

## References

- Gardner, P.P., Daub, J., Tate, J., Moore, B.L., Osuch, I.H., Griffiths-Jones, S., Finn, R.D., Nawrocki, E.P., Kolbe, D.L., Eddy, S.R. *et al.* (2010) Rfam: Wikipedia, clans and the "decimal" release. *Nucleic Acids Res.*, **39**, D141-145.
- Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. *vesicatoria* revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.

**Table S7.** RNAcode prediction of novel chromosomal ORFs in *Xcv* (1, 2). Alternative start codons are listed under “chromosomal position”. RNAcode6 is highlighted in bold and corresponds to sX6 which was experimentally verified (see Figure 3).

RNAcode prediction	Chromosomal position (Strand)	Transcript Support <sup>a</sup>	TSS position <sup>b</sup>
RNAcode1	285154-285243 (+)	0/0	-
RNAcode2	465113-465232 (+)	0/0	-
RNAcode3	722692-722736 (+)	1/0	-
RNAcode4	1281664-1281846 (+)	1/1	-
	1281757-1281846 (+)		
RNAcode5	1435086-1435148 (+)	0/0	-
	1435113-1435148 (+)		
<b>RNAcode6 (sX6)</b>	<b>1971528-1971770 (+)</b>	<b>15/1</b>	<b>1971505</b>
	1971591-1971770 (+)		
RNAcode7	3457704-3457808 (+)	0/0	-
	3457734-3457808 (+)		
RNAcode8	3463701-3463796 (+)	3/4	-
	3463743-3463796 (+)		
RNAcode9	3727337-3727555 (+)	21/1	3727405
	3727361-3727555 (+)		
	3727445-3727555 (+)		
RNAcode10	3848998-3849048 (+)	1/0	-
RNAcode11	4083126-4083464 (+)	0/0	-
	4083348-4083464 (+)		
RNAcode12	4368388-4368489 (+)	0/4	-
RNAcode13	4545859-4545918 (+)	0/0	-
RNAcode14	5098317-5098562 (+)	0/0	-
	5098473-5098562 (+)		
RNAcode15	644457-644513 (-)	0/0	-
RNAcode16	1219379-1219462 (-)	0/0	-
	1219379-1219450 (-)		
RNAcode17	1360037-1360129 (-)	0/0	-
	1360037-1360120 (-)		
RNAcode18	2429470-2429610 (-)	0/0	-
	2429470-2429568 (-)		
RNAcode19	2602554-2602655 (-)	0/0	-
RNAcode20	3215182-3215274 (-)	1/0	-
	3215182-3215223 (-)		
RNAcode21	3408680-3408883 (-)	4/0	3408906
RNAcode22	3881142-3881246 (-)	2/0	-
RNAcode23	4104001-4104081 (-)	1/1	-
	4104001-4104060 (-)		
RNAcode24	5054413-5054448 (-)	1/0	-

<sup>a</sup>, number of reads (library 2/library 1) mapped to the respective loci. <sup>b</sup>, automatically annotated TSS (see Table S2); “-“ no TSS identified.

## References

1. Thieme, F., Koebnik, R., Bekel, T., Berger, C., Boch, J., Büttner, D., Caldana, C., Gaigalat, L., Goesmann, A., Kay, S. *et al.* (2005) Insights into genome plasticity and pathogenicity of the plant pathogenic bacterium *Xanthomonas campestris* pv. vesicatoria revealed by the complete genome sequence. *J. Bacteriol.*, **187**, 7254-7266.
2. Washietl, S., Findeiß, S., Müller, S.A., Kalkhof, S., von Bergen, M., Hofacker, I.L., Stadler, P.F. and Goldman, N. (2011) RNAcode: Robust discrimination of coding and noncoding regions in comparative sequence data. *RNA*, **17**, 578-594.



**Table S8.** Genome sequences used for the calculation of multiple sequence alignments.

NCBI ID	Species name
NC_008752.1	<i>Acidovorax avenae</i> subsp. <i>citrulli</i> AAC00-1, complete genome
NC_004547.2	<i>Erwinia carotovora</i> subsp. <i>atroseptica</i> SCRI1043, complete genome
NC_010943.1	<i>Stenotrophomonas maltophilia</i> K279a, complete genome
NC_011071.1	<i>Stenotrophomonas maltophilia</i> R551-3, complete genome
NC_002516.2	<i>Pseudomonas aeruginosa</i> PAO1, complete genome
NC_009656.1	<i>Pseudomonas aeruginosa</i> PA7, complete genome
NC_008463.1	<i>Pseudomonas aeruginosa</i> UCBPP-PA14, complete genome
NC_005773.3	<i>Pseudomonas syringae</i> pv. <i>phaseolicola</i> 1448A, complete genome
NC_007005.1	<i>Pseudomonas syringae</i> pv. <i>syringae</i> B728a, complete genome
NC_004578.1	<i>Pseudomonas syringae</i> pv. <i>tomato</i> DC3000, complete genome
NC_003295.1	<i>Ralstonia solanacearum</i> GMI1000, complete genome
NC_003919.1	<i>Xanthomonas axonopodis</i> pv. <i>citri</i> 306, complete genome
NC_003902.1	<i>Xanthomonas campestris</i> pv. <i>campestris</i> ATCC 33913, complete genome
NC_007086.1	<i>Xanthomonas campestris</i> pv. <i>campestris</i> 8004, complete genome
NC_007508.1	<i>Xanthomonas campestris</i> pv. <i>vesicatoria</i> 85-10, complete genome
NC_002488.3	<i>Xylella fastidiosa</i> 9a5c, complete genome
NC_004556.1	<i>Xylella fastidiosa</i> Temecula1, complete genome
NC_006834.1	<i>Xanthomonas oryzae</i> pv. <i>oryzae</i> KACC10331, complete genome
NC_007705.1	<i>Xanthomonas oryzae</i> pv. <i>oryzae</i> MAFF 311018, complete genome

**Table S9.** The extended two-by-two confusion matrix summarizes the predictive power of the automated TSS annotation approach. A subsample of the manually curated TSS map of *H. pylori* (1) was used as reference data set and evaluated with our method. The analyzed data-set contained 392 manually annotated TSSs. In total 566 genomic positions fulfilled the criteria (accumulation of at least three read starts) to be analyzed by our automated TSS annotation approach. According to Fawcett (2006), true positives (classified as TSS and also manually annotated) and false positives (classified as TSS but not manually annotated) as well as true negatives (neither classified as TSS nor manually annotated) and false negatives (not classified as TSS but manually annotated) were evaluated (2). Based on these values the calculated values for sensitivity, specificity as well as positive and negative predictive values are listed.

	Manually annotated TSSs (392)		
Automatic classification (566)	True Positives: 321	False Positives: 124	Positive predictive value: 72%
	False Negatives: 71	True Negatives: 50	Negative predictive value: 59%
	Sensitivity: 82%	Specificity: 29%	

## References

1. Sharma, C.M., Hoffmann, S., Darfeuille, F., Reignier, J., Findeiss, S., Sittka, A., Chabas, S., Reiche, K., Hackermüller, J., Reinhardt, R. *et al.* (2010) The primary transcriptome of the major human pathogen *Helicobacter pylori*. *Nature*, **464**, 250-255.
2. Fawcett, T. (2006) An introduction to ROC analysis. *Pattern recognition letters*, **27**, 861-874.