

Figure S7. Screenshot of IGV visualization on *L. monocytogenes* RO15 for condition-specific prophage TSSs. A) TSS:679897_f, B) TSS:2178605_r , and C) TSS:2221338_f. Nine rows of data, in descending order: 0 min control sample for 200 MPa cap-seq reads, 0 min 200 MPa treated samples cap-seq reads, 0 min control sample for 400 MPa cap-seq reads, 0 min 400 MPa treated samples cap-seq reads, 24 h control sample for 400 MPa cap-seq reads, 24 h 400 MPa treated sample cap-seq reads, 24 h control sample for 400 MPa direct RNA-seq reads, 24 h 400 MPa treated sample direct RNA-seq reads, and Illumina short RNA-seq reads. Red lines are forward reads, and blue lines are reverse reads. Grey bars are coverage bar plots.

A)

CADEHJ010000001.1:679,254-680,832

1,579 bp

679,400 bp 679,600 bp 679,800 bp 680,000 bp 680,200 bp 680,400 bp 680,600 bp 680,800 bp

T0 Control for 200 MPa Cap-seq

[0 - 27]

T0 200 MPa Treated Cap-seq

[0 - 10.00]

T0 Control for 400 MPa Cap-seq

[0 - 13]

T0 400 MPa Treated Cap-seq

[0 - 10.00]

T8 Control for 400 MPa Cap-seq

[0 - 19]

T8 400 MPa Treated Cap-seq

[0 - 10.00]

Nanopore Direct RNA-seq T8 Control

[0 - 10.00]

Nanopore Direct RNA-seq T8 400 MPa Treated

[0 - 10.00]

Illumina RNA-seq combine

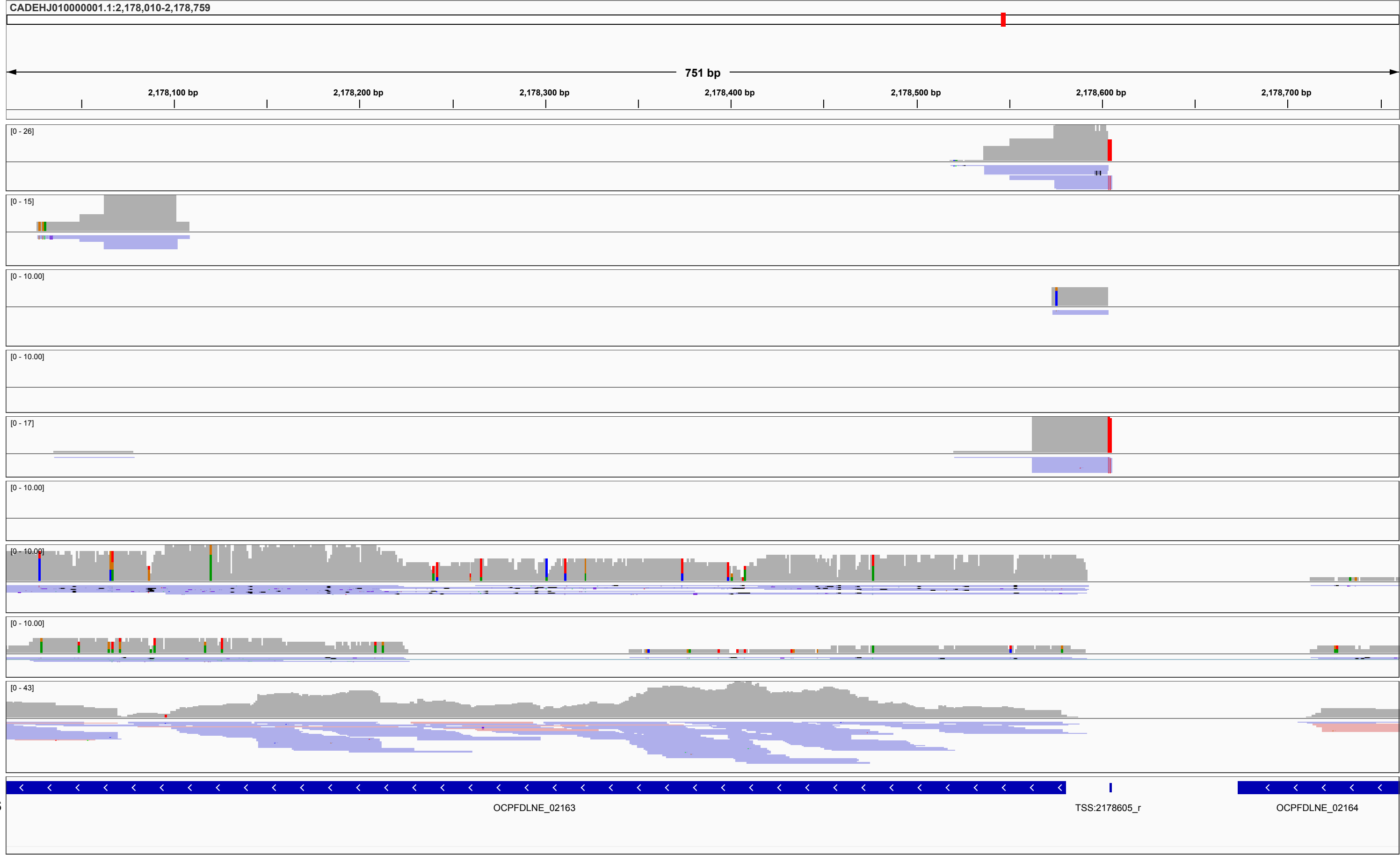
[0 - 10.00]

Gene and TSS Annotation

hhaIM

CPFDLNE_00656 TSS:679897_f OCPFDLNE_00658

Gene and TSS Annotation



C)

T0 Control for 200 MPa Cap-seq

**T0 200 MPa
Treated
Cap-seq**

T0 Control for 400 MPa Cap-seq

**T0 400 MPa
Treated
Cap-seq**

T8 Control for 400 MPa Cap-seq

**T8 400 MPa
Treated
Cap-seq**

**Nanopore
Direct
RNA-seq
T8 Control**

**Nanopore
Direct
RNA-seq
T8 400 MPa
Treated**

Illumina RNA-seq combine

Gene and TSS Annotation

