

TABLE S6 Summary of mapped *C. perfringens* and murine RNA-seq reads.

<i>C. perfringens</i>	Mapped reads	
	<i>in vivo</i> ^a	<i>in vitro</i> ^a
Replicate 1	2,164,911	1,819,981
Replicate 2	1,823,412	1,389,633
Replicate 3	1,759,987	1,374,988
Total reads	5,748,310	4,584,602
Host	<i>C. perfringens</i> infected host	Mock-infected host
Replicate 1	21,241,035	27,713,718
Replicate 2	22,033,729	25,463,014
Replicate 3	19,942,403	29,983,040
Total reads	63,217,167	83,159,772

^a RNA-seq analysis of the enriched RNA samples from the *in vivo*- and *in vitro*-derived *C. perfringens* cells was carried out on Illumina Hi-seq and Mi-seq instruments, respectively, which accounts for higher number of reads in the *in vivo* samples. .