

Supplementary table 1 The binding site of SPI1 and S100A8/A9

Matrix ID	TF	Score	Relative score	Sequence ID	Start	End	Strand	Predicted sequence
<u>MA0080.2</u>	SPI1	11.542796	1.00000001	S100A8	663	669	+	aggaagt
<u>MA0080.1</u>	SPI1	8.578765	0.9944638	S100A8	411	416	+	gggaag
MA0080.1	SPI1	8.578765	0.994463805	S100A9	769	774	+	gggaag

Notes: SPI, salmonella pathogenicity island; TF, transfer factor.