Supplementary information

Pathogen-specific DNA sensing with engineered zinc finger proteins immobilized on

polymer chip

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Target 1 Stx2_233/stx2_2685'-GAC GGC TTG ATG TCT ATC AGG CGC GTT TTG ACC ATC TTC GGG TTTT CCC GAA GAT GGTCAA AAC GCG CCT GAT AGA CATCAA GCC GTC-3'

Target 2 Stx2_525/stx2_560

5'-GĂC TGC ATC CAG AGC AGT TCT GCG TTT TGT CAC TGT CAC GGG TTTT CCC GTG ACA GTG ACA AAA CGC AGA ACT GCT CTG GAT GCA GTC-3'

Target 3 Stx2_1093/stx2_1128

5'-GĂC AAG TGG CCG GGA AAG AAT ACT GGA CCA GTC GCT GGA CCC TTTT GGG TCC AGC GAC TGG TCC AGT ATT CTT TCC CGG CCA CTT GTC-3'

Hairpin oligonucleotide of Zif268/PBSII site, used as an irrelevant control: 5'-GGC TTT CCA CAC CGC CCA CGC GGG TTTT CCC GCG TGG GCG GTG TGG AAA GCC-3'

Fig. S1 Sequences of hairpin target DNA oligonucleotides for stx2 ZFP pairs with ZFP binding sites shown in bold.

1	atgaagtgta	tattatttaa	atgggtactg	tgcctgttac	tgggtttttc	ttcggtatcc
61	tattcccggg	aatttacgat	agacttttcg	actcaacaaa	gttatgtctc	ttcgttaaat
121	agtatacgga	cagagatatc	gacccctctt	gaacatatat	ctcaggggac	cacatcggtg
181	tctgttatta	accacacccc	accgggcagt	tattttgctg	tggatatacg	ag <mark>ggcttgat</mark>
241	<mark>gtctatcagg</mark>	<mark>cgcgttttga</mark>	<mark>ccatcttc</mark> gt	ctgattattg	agcaaaataa	tttatatgtg
301	gccgggttcg	ttaatacggc	aacaaatact	ttctaccgtt	tttcagattt	tacacatata
361	tcagtgcccg	atgtgacaac	ggtttccatg	acaacggaca	gcagttatac	cactctgcaa
421	cgtgtcgcag	cgctggaacg	ttccggaatg	caaatcagtc	gtcactcact	ggtttcatca
481	tatctggcgt	taatggagtt	cagtggtaat	acaatgacca	gaga <mark>tgcatc</mark>	cagagcagtt
541	ctgcgttttg	tcactgtcac	agcagaagcc	ttacgcttca	ggcagataca	gagagaattt
601	cgtcaggcac	tgtctgaaac	tgctcctgtg	tatacgatga	cgccgggaga	tgtggacctc
661	actctgaact	gggggcgaat	cagcaatgtg	cttccggagt	atcggggaga	ggatggtgtc
721	agagtgggga	gaatatcctt	taataatata	tcagcgatac	tgagtactgt	ggccgttata
781	ctgaattgcc	atcatcaggg	ggcgcgttct	gttcgcgccg	tgaatgaaga	gagtcaacca
841	gaatgtcaga	taactggcga	caggcccgtt	ataaaataa	acaatacatt	atgggaaagt
901	aatacagcag	cagcgtttct	gaacagaaag	tcacagtttt	tatatacaac	gggtaaataa
961	aggagttaag	tatgaagaag	atgtttatgg	cggttttatt	tgcattagtt	tctgttaatg
1021	caatggcggc	ggattgcgct	aaaggtaaaa	ttgagttttc	caagtataat	gagaatgata
1081	cattcacagt	aa <mark>aagtggcc</mark>	gggaaagaat	actggaccag	tcgctggaat	ctgcaaccgt
1141	tactgcaaag	tgctcagctg	acaggaatga	ctgtcacaat	caaatccagt	acctgtgaat
1201	caggctccgg	atttgctgaa	gtgcagttta	attttgaatg	a	

Fig. S2 Location of target regions of stx2 ZFP pairs in *stx2* gene (1241 bp). Target sites are highlighted for stx2_233/stx2_268 (yellow), stx2_525/stx2_560 (cyan), and stx2_1093/stx2_1128 (light green).



Fig. S3 Illustration of EMSA (electromobility shift assay) of engineered zinc finger proteins (ZFPs) to target different sequences within stx2 gene. Exemplary data are shown for (a) Stx2_233, (b) Stx2_268, and (c) Stx2_525. The ZFP concentrations (nM) are given at the top. Top bands indicate the ZFP and DNA complex and the bottom bands show free DNA.

		Motifs	Seed	Enrichment	Protein cont.	Salt cont.
		GTGACAGTGACAAAACGC		·		
	Stx2_560		АСААААА	0.157	50 nM	50 mM
	Stx2_1093	AAGTGGCCGGGAAAGAAT				•
			CTTTCCCGCC	26.432	500 nM	100 mM
	Stx2_1128	ACTGGACCAGTCGCTGGA		·	·	
			AGTTGGTCG	4.345	50 nM	1 mM

Table. S1 Binding motifs identified by the Bind-n-Seq for ZFP stx2_560, stx2_1093,and stx2_1128.