A genetically encoded probe for monitoring and detection of iron in real-time

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Fig. S1 SocA sequence alignment analysis: comparing the original sequence with obtained results.

Score		Expect	Identities	Gaps	Strand
582 bi	ts(315)	3e-171	315/315(100%)	0/315(0%)	Plus/Plus
Query	1	ATGAACGACAGTGAAT	TTCATCGCCTGGCTGATCA	ACTCTGGCTGACCATTGA	AGAACGC 60
Sbjct	1	ATGAACGACAGTGAAT	TTCATCGCCTGGCTGATCA	ACTCTGGCTGACCATTGA	AGAACGC 60
Query	61	CTGGACGACTGGGATG	GCGACAGCGATATCGACTG	CGAAATCAACGGCGGCGT	ACTGACC 120
Sbjct	61	CTGGACGACTGGGATG	GCGACAGCGATATCGACTG	GAAATCAACGGCGGCGT	ACTGACC 120
Query	121	ATTACCTTTGAGAATG	GCAGCAAAATCATTATCAA	CCGCCAGGAGCCGCTGCA	CCAGGTA 18
Sbjct	121	ATTACCTTTGAGAATG	GCAGCAAAATCATTATCAA	CCGCCAGGAGCCGCTGCA	CCAGGTA 18
Query	181	TGGCTGGCAACCAAAC	AGGGCGGCTACCATTTTGA	CCTGAAAGGCGATGAGTG	GATTTGT 24
Sbjct	181	teecteectaaccaaac	AGGGCGGCTACCATTTGA	icctgaaaggcgatgagtg	GATTTGT 24
Query	241	GATCGCAGCGGCGAAA	CCTTCTGGGATTTGCTGGA	ACAGGCGGCGACGCAGCA	GGCGGGT 30
Sbjct	241	GATCGCAGCGGCGAAA	CCTTCTGGGATTTGCTGGA	ACAGGCGGCGACGCAGCA	GGCGGGT 30
Query	301	GAAACAGTCAGTTTC	315		
sbjct	301	GAAACAGTCAGTTTC	315		

Fig. S2 Designing of nanosensor constructs in different expression vectors (pRSET-B, pYEST-DEST52, pCDNA3.1(-)) to facilitate cellular expression in *E. coli*, *S. cerevisiae*, and HEK-293T cells.



Fig. S3 Electropherograms showing the sequencing results of (A) I17C (B) A10D-I17C and (C) D76H sensor variants. The indicated mutated nucleotides lead to the expression of cysteine, aspartic acid/cysteine, and histidine at positions 17, 10/17, and 76 in the corresponding variants I17C, A10D-I17C, and H76D.



Fig. S4 Confocal microscopy image demonstrating efficient expression of the genetically encoded FeOS in bacterial cells.



Fig. S5 FRET ratio changes recorded before and after incubation of yeast cell with metal ions for 10 minutes.



Fig. S5 FRET ratio changes recorded before and after incubation of HEK cell with metal ions for 10 minutes.

