

Supplementary Material

Pyoverdine Binding Aptamers and Label-Free Electrochemical Detection of Pseudomonads

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TABLES

Table S1. Comparison of methods for detecting pyoverdine

Method	Detection method	sensitivity	detection limits
Colorimetric	Binding of pyoverdine and other siderophores to iron results in a color change (Louden et al., 2011)	Positive for all siderophores	Visual indicator, and quantification could be challenging
Fluorescence Assays	Intrinsic fluorescence of chromogenic metabolite (Cueva et al., 2020)	Linearity range of 10- 1000 ng/ml (7.3 nM – 732 nM)	LOD 0.9 ng/ml (0.6 nM)
Electrochemical Sensing	PVD oxidation on carbon electrodes coated with electrochemically generated graphene functionalized with gold nanoparticles (Gandouzi et al., 2018)	A linear range of 1- 100 μM pyoverdine in PBS. Detected 25 μM pyoverdine in serum, saliva and tap water matrices.	LOD: 333.33 nM
	Pyoverdine sensing with aptamers (this report)	A dynamic range over three log units from 5 - 1000 nM. Specific for pyoverdine over other siderophores - enterobactin and pseudobactin	LOD: 1.3 nM

Legend: A summary is given of the current methods for detecting pyoverdine. Some (colorimetric, fluorescence) require first extracting the pyoverdine to separate it from many other compounds that contribute to background noise.

Table S2. Sequences of aptamers

15THR1A	GGTTGGTGTGG
21COD1A	TGGGTCGGGAGGGAATGCGGG
32THR4A	GGTAGGGCAGGTTGGGTGTTTTCACTTTTGGG
55PYO2A	CTAGGAATCAGGAGCGAATGTTAGAGTATTTGGATAGTATGTTTATT
	GTATCGGC
55PYO4A	GATCGGAATGGGAGCGA
	ATGAG GAGGGAGCGCGGACATGGTGCAGCAGCCCGACG
BRC	GAGACGGUCGGGUCCAGAUAUUCGUAUCUGUCGAGUAGAGUGUGGGC
	UC

Table S3. Sequences of other nucleic acids used in DNA SELEX

F _	
Oligo 5615	GCCTGTTGTGAGCCTCCTGTCGAA
5' primer	
Oligo 5617	GGGAGACAAGAATAAACGCTC
3' primer	
Oligo 5625	TTCGACAGGAGGCTCACAACAGGC
5' primer	
complement	
Oligo 5636	GCCTGTTGTGAGCCTCCTGTCGAACTAGGAATCAGGAGC
55PYO2A with	GAATGTTAGAGTATTTGGATAGTATGTTTATTGTATCGGCG
primer binding	AGCGTTTATTCTTGTCTCCC
region	
Oligo 5637	GCCTGTTGTGAGCCTCCTGTCGAAGTATGCGATAGGAGC
Control	GAATGATATATCAAGTTACCACACGGATGTACTTGCTTG
oligonucleotide	AGCGTTTATTCTTGTCTCCC
Oligo 5685	GCCTGTTGTGAGCCTCCTGTCGAAGATCGGAATGGGAGC
55PYO4A with	GAATGAGGAGGGAGCGCGGACATGGTGCAGCAGCCCGA
primer binding	CGGAGCGTTTATT
region	

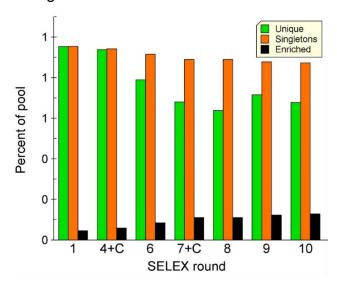
Table S4: Selection scheme for DNA aptamers that bind PVD-Pf5.

Round	ssDNA: PVD-Pf5	Target	Length of capture oligo (nt)	Incubation temperature
1	1:1	Ferric PVD-Pf5	6	4
2	2:1	Ferric PVD-Pf5	6	4
3	3:1	Ferric PVD-Pf5	7	4
Counter selection	1:1	Ferric enterobactin and Ferric ornibactin	7	4
4	5:1	Ferric PVD-Pf5	8	4
5	6:1	Ferric PVD-Pf5	8	23
6	7:1	Ferric PVD-Pf5	8	23
Counter selection	1:1	Ferric enterobactin and Ferric ornibactin	8	23
7	8:1	Ferric PVD-Pf5	8	23
8	10:1	Ferric PVD-Pf5	9	23
9	10:1	Ferric PVD-Pf5	9	23
10	10:1	Ferric PVD-Pf5	9	23

Legend: Ferric-enterobactin and ferric-ornibactin were used as counter selection targets. Selection was performed in SSMA2.

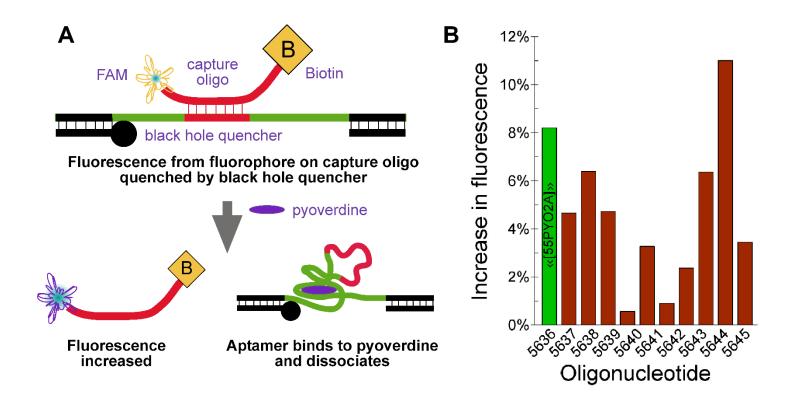
FIGURES

Figure \$1. NextGEN sequencing data



Legend: Next gen sequencing data from DNA SELEX showed downward trends of the unique fraction (number of species in pool divided by pool size) and singletons (species present only once in the database) and a rise in the enriched species (species present more than once in the database).

Figure S2: Screening for aptamers



LEGEND: Aptamer screening by fluorescence quenching assay **A)** ssDNA aptamer candidates were hybridized with FAM (6-Carboxyfluorescein) and biotin-labeled capture oligo and black hole quencher labeled blocking oligo on 5' end and blocking oligo on 3' end. FAM fluorescence was quenched by the black hole quencher in the hybridized complex (capture oligo-aptamer-blocking oligos). After binding with pyoverdine the aptamer dissociates from complex and fluorescence increases from the fluorophore attached to the capture oligo. This assay was performed in streptavidin coated microwells by which the biotinylated capture oligo was held to the surface. The depiction in A is based on Fig. 1 for aptamer screening in (Zhao et al., 2019). **B)** Binding of 40nM candidate aptamers with 800nM pyoverdine. Candidate 5636 contains the sequence of 55PYO2A

Figure S3: The structure of PVD-Pf5

$$H_2N$$
 H_2N
 H_2N

Legend: The structure of PVD-Pf5 is shown (Hartney et al., 2013)

Citations

- Cueva, A. R., Pham, O., Diaby, A., Fleming, D., Rumbaugh, K. P. & Fernandes, G. E. 2020. Pyoverdine Assay for Rapid and Early Detection of Pseudomonas aeruginosa in Burn Wounds. *ACS Applied Bio Materials*, 3, 5350-5356.
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