

PCR hyperplexed without compromise

# Democratizing high-performance multipathogen surveillance with next-gen Hyperplex PCR™

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**Chief Scientific Officer** 

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- Founded in 2020
- Headquartered in Karolinska Institutet Campus, Stockholm
- Formed by industry and academic leaders in Molecular Diagnostics behind companies like Olink, Cartana, ParAllele Bioscience, Qlinea, and Halo Genomics



Mission

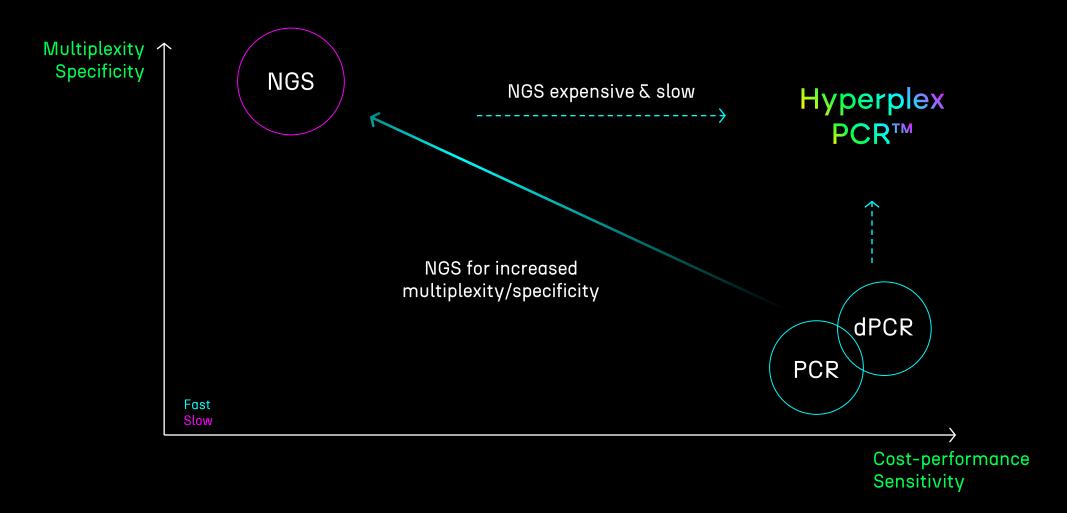
To push the boundaries of molecular analysis enabling unprecedented multiplexing, sensitivity and specificity without compromise

Vision

To be a globally recognized leader in molecular analysis technology, providing multiple, gold-standard hyperplexing solutions for healthcare and sustainability

#### Unmet Need In Efficient Molecular Diagnostics





#### Introducing Hyperplex PCR™





Multiplexity of 100+

No splitting, no sequencing







New probes on-demand

With NGS-grade specificity, no optimization



With results within 1 day





High & low abundance targets

Simultaneous detection in one tube

Single-molecule sensitivity

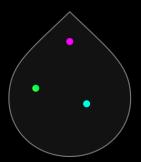
Ability to count PCR amplicons



#### Hyperplex PCR™ - Multiplexing Unleashed

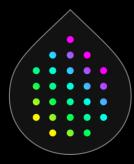


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Conventional PCR 3-5 biomarkers

Limited information split sample or rely on sequencing



Hyperplex PCR 100+ biomarkers

Complete picture unparalleled simple solution

#### With Cutting-Edge Nanoprobes and Molecular Biology

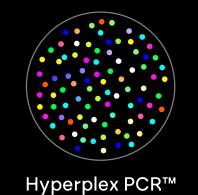
Based on Over 20 Years of Research

#### We Have Taken PCR to a Whole New Level



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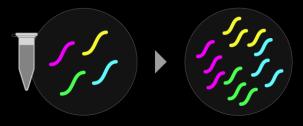


Multiplex/well 100+ LoD ~5 (copies/reaction) Single-nucleotide specificity X Simple probe design \$ \$\$ \$ Cost Counts 10,000 1,000,000

## Hyperplex PCR™ Assay



**01** PCR Amplification



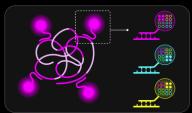
02 Padlock probing & Ligation

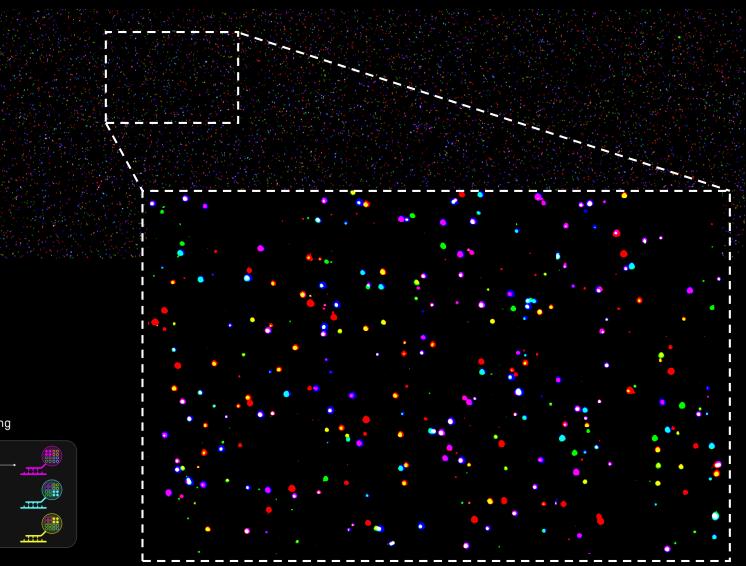


03 Rolling-Circle Amplification



04 Nanopixel Labelling



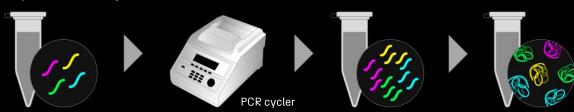


## Simple Workflow, Common Instruments



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hpPCR assay - PCR/Ligation/RCA



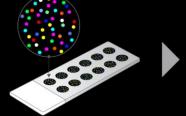
Mixed sample >100 markers

Compatible with any off-the-shelf extraction kit

Sample capture - slide or plate

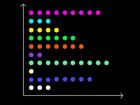


Fluorescence imaging and automated image analysis









>100-plex output in a single step

Turnover time < 1 day

Fluorescence microscope or reader (4-5 common filters)

#### hpPCR used for COVID WW monitoring in Sweden Aplex<sup>BIO</sup>



on behalf of Swedish Environmental Epidemiology Center (SEEC)

Powered by Nanopixels™

#### Targets Monitored

N3	S:T547I
S:R346T	S:Q613H
S:K444T	Orf9b:I5T
S:N460K	Orf9b:N55S
S:F490S	S:483del
S:F456L	S:F157S/R1580
S:Q52H	orf1:A7842G
S:F486P	S:69-70del
S:T478R	PMMoV

Application note released on our website

https://www.aplex.bio/application-note-wbe



Sites Monitored

On weekly basis.

Results are delivered within 3 days.





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## Hyperplexed Variant Surveillance



COVID Monitoring Oct '22-Nov '23

700+

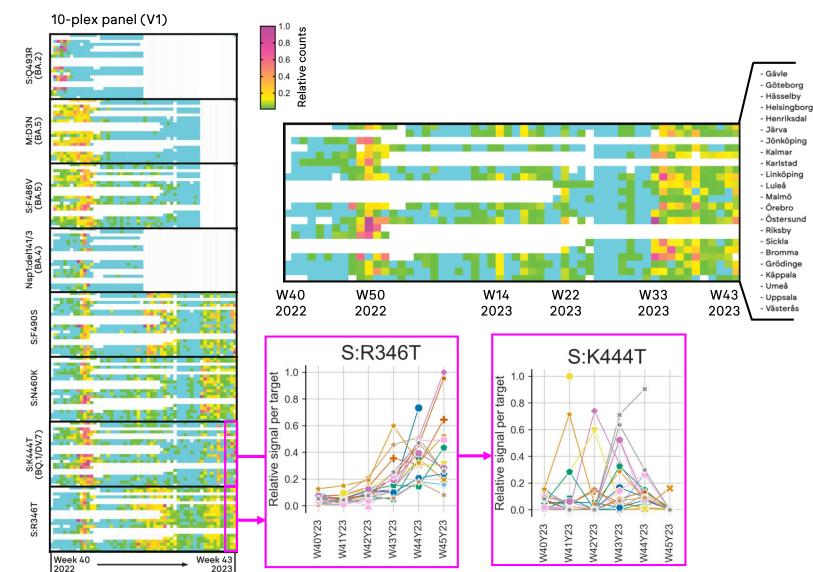
Samples analysed (duplicates)

10,000+

Data points

18-plex+

Per sample, modified on demand



### Hyperplexed Variant Surveillance



## COVID Monitoring Oct '22-Nov '23

700+

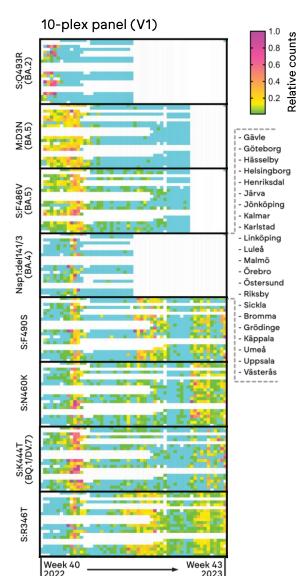
Samples analysed (duplicates)

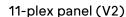
10,000+

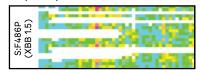
Data points

18-plex+

Per sample, modified on demand

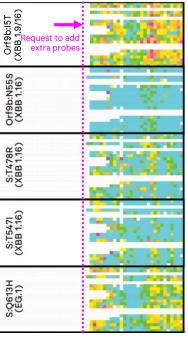






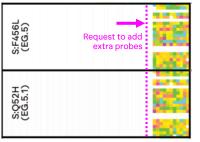
Additional probe for S:F486P (XBB 1.5) requested W49/2022

#### 14-plex panel (V3)



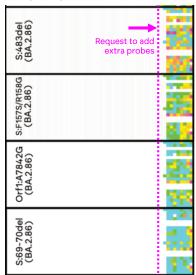
Additional probes for mutations characteristic of XBB 1.9/16 and EG.1 BA.2+BA.4 probes excluded

#### 16-plex panel (V4)



Probes for BA.5 excluded

#### 18-plex panel (V5)

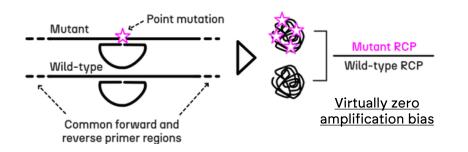


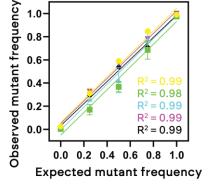
All panel modifications carried out within 2-weeks

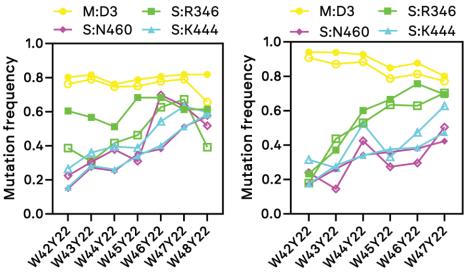
#### Mutation Frequency and Earlier Detection

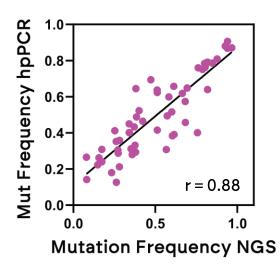


hpPCR allows precise and sensitive quantification of mutation frequency in wastewater samples

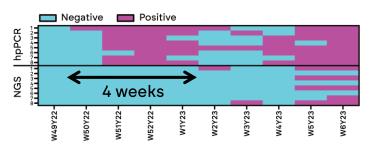








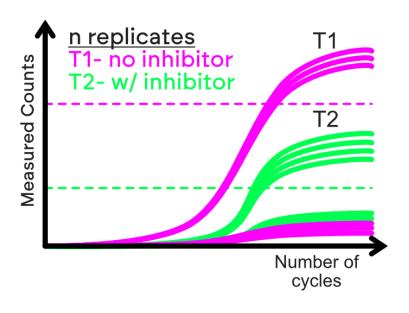
hpPCR allows early detection of emerging variants vs NGS at least 4 weeks earlier



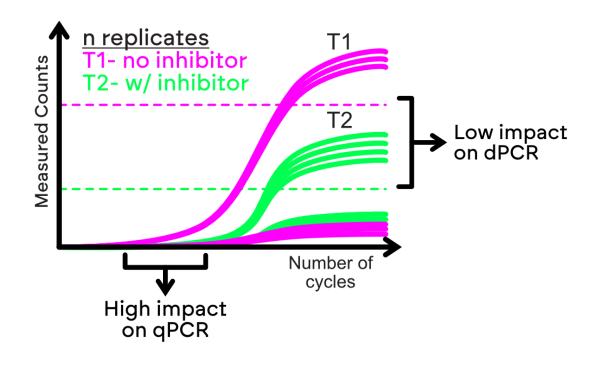
Weekly measurement of S:F486P, characteristic of XBB 1.5 using hpPCR and NGS (Ion torrent). 8 wastewater collection sites were measured at each time point.

Both measurements were performed on the same extracted nucleic acid sample.



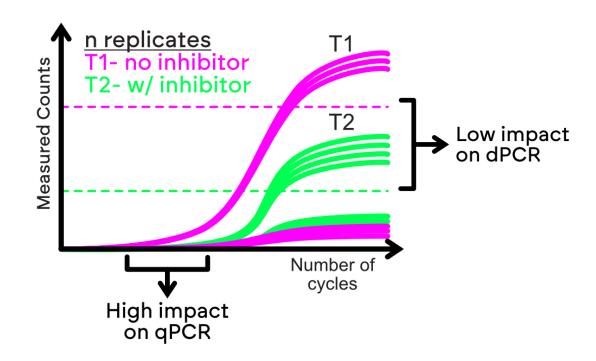




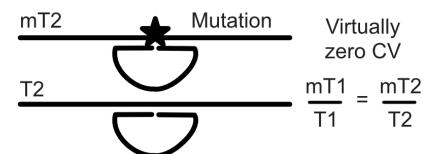




Mutation frequency readout and internal controls neutralize PCR bias



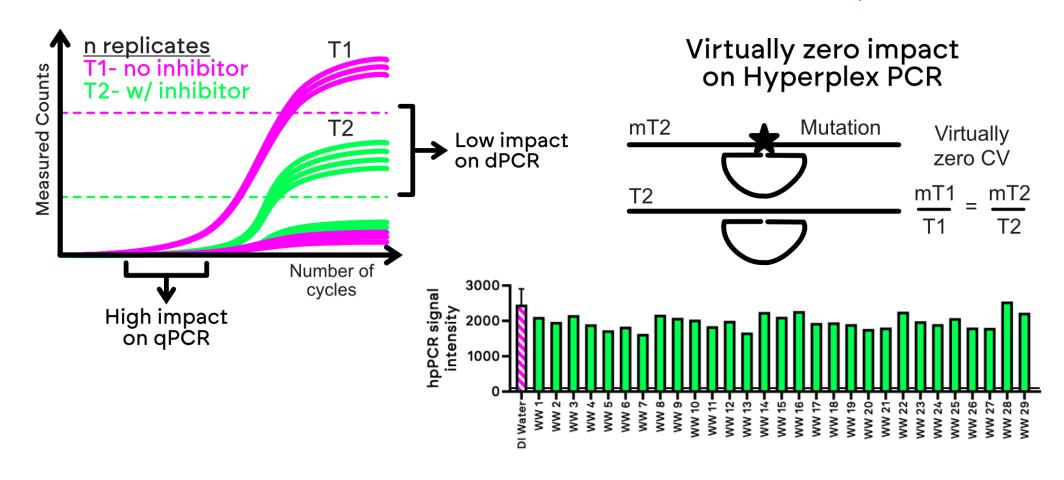
## Virtually zero impact on Hyperplex PCR





Mutation frequency readout and internal controls neutralize PCR bias

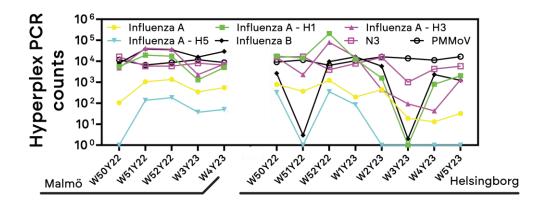
+ Intensity of RCA signals independent of inhibition in wastewater samples



### Quantitative Multi-Pathogen Analysis



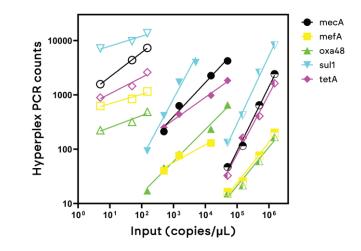
- Versatile multi-pathogen panels
  - Influenza
  - AMR
  - SARS-CoV-2



Monitoring of wastewater in two sites using a 7-plex multi-pathogen / multi-subtype panel.

All plotted signals are significantly above the blank baseline by at least 3 standard deviations. 10<sup>o</sup> means non-detection.

- Absolute quantification
- Dynamic range >6 orders
- High & low abundance in same sample



Site	mecA (cop/µL)	mefA (cop/µL)	oxa48 (cop/µL)	sul1 (cop/µL)	tetA (cop/µL)
1	< 10	1.1x10 <sup>4</sup>	< 10	8.9x10 <sup>4</sup>	1,5x10 <sup>3</sup>
2	< 10	1.2x10 <sup>4</sup>	< 10	1.3x10⁵	1,7x10 <sup>3</sup>
3	< 10	1.2x10 <sup>4</sup>	< 10	1.2x10⁵	5,3x10 <sup>3</sup>
4	1.2x10 <sup>2</sup>	2.0x10 <sup>4</sup>	< 10	4.7x10⁵	4,9x10 <sup>3</sup>
5	48	3.4x10⁵	< 10	2.8x10⁵	1,4x10 <sup>4</sup>
6	2,2x10 <sup>2</sup>	7.1x10 <sup>4</sup>	< 10	6.7x10 <sup>4</sup>	4,2x10 <sup>3</sup>
7	1,2x10 <sup>3</sup>	1.0x10 <sup>4</sup>	< 10	2.3x10⁵	3,3x10 <sup>4</sup>
8	9,8x10 <sup>2</sup>	9.8x10 <sup>3</sup>	< 10	8.3x10 <sup>4</sup>	2,6x10 <sup>4</sup>
9	5.8x10 <sup>3</sup>	9.8x10 <sup>2</sup>	< 10	2.4x10⁵	7,8x10 <sup>3</sup>
10	< 10	5.0x10 <sup>2</sup>	N.D.	2.6x10⁵	6,1x10 <sup>2</sup>

## Summary

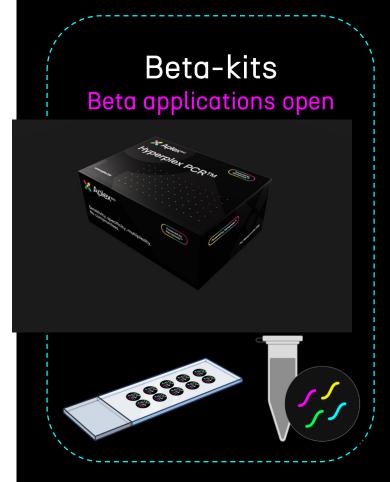


- ✓ 4-weeks+ earlier detection of variants of concern compared to NGS
- ✓ Dynamic panel modification with new probes within 2 weeks
- ✓ Multiplex capability of 100+ targets per sample no sample splitting
- ✓ High and low abundance targets in one reaction
- ✓ NGS-grade mutation frequency quantification
- ✓ No need to rely on NGS for monitoring variants/mutations.

#### Time for Action — Method Standardization is Key

- ✓ Comparability across European labs is essential
- ✓ Hyperplexed multi-pathogen panels are now available
- ✓ Beta kits are now ready to be delivered to your labs
- ✓ Apply today and get ahead with Hyperplex PCR™





## Hyperplex PCR™

Powered by Nanopixels™

#### Revolutionizing Molecular Diagnostics

Contact:
Danai Nikou
Business Developer
danai@aplex.bio



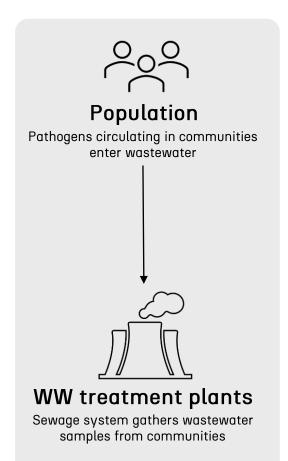


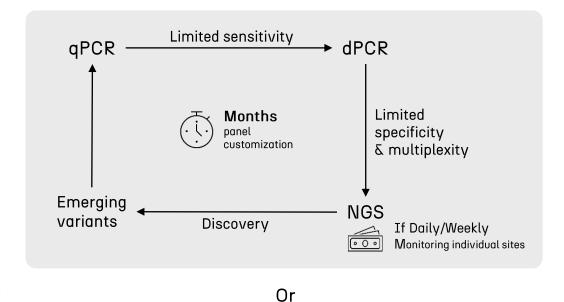
#### Extra slides

#### Sampling

#### **Discovery & Monitoring**



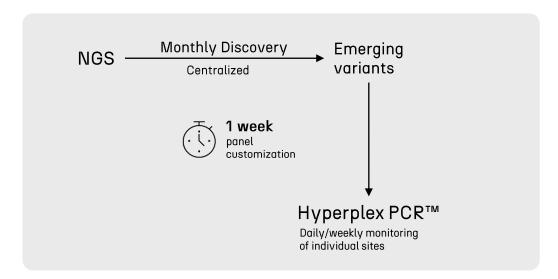




Time for Action

Comparable and
Quantitative Results
Between sites

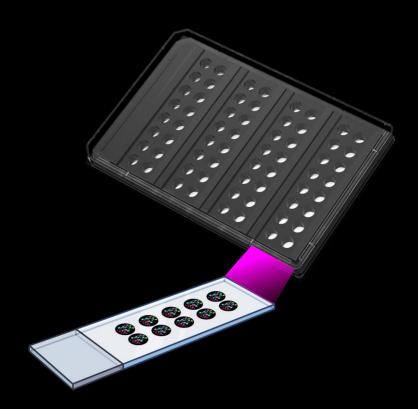
 $\sum$ 1 day



## Throughput and read-out



SBS format
Up to 384 samples at once
Automated off the shelf



Feature	hpPCR					
Multiplex/well	100+					
Counts/well	Up to 1 000 000					
Throughput/day	User/automation defined >128 samples					
Sample type	Extracted DNA/RNA					
Workflow	4-8 hr / <30 min hands-on					
Software	Dedicated image analysis pipeline included					
Instrumentation	PCR cycler Fluorescence microscope					
Microscope	Objective: 20x/0.8 Channels: 4-6					





			Current: Bet	а		SCALED			HIG	H THROUG	HPUT		
					1 2 3	4 5 6	7 8 9		1 2	3 4 5 6 7 8	9 10 11 12		
				A B C D				•	B C C C C C C C C C C C C C C C C C C C			Þ	
				F					J OC				
	Samples		18			54 (3x18)	•			144 (3x48	5)		
	Hyperplexing (per spot)		20			• 100 •				100			
	Or max multi (using all spots)	plex	360			5 400				14 400			

Consumables are designed to be compatible with standard plate formats as well as to facilitate flexibility with microscope slides