

New Methods to Reduce Variation in Bare Wafer Particle Inspection Results

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Agenda

Background

tpM2M (Tighter Pre-scan Map-2-Map) Method

DIOO (Dual Incidence Oblique Oblique) Method

Applying tpM2M + DIOO to Filter Evaluations

Background



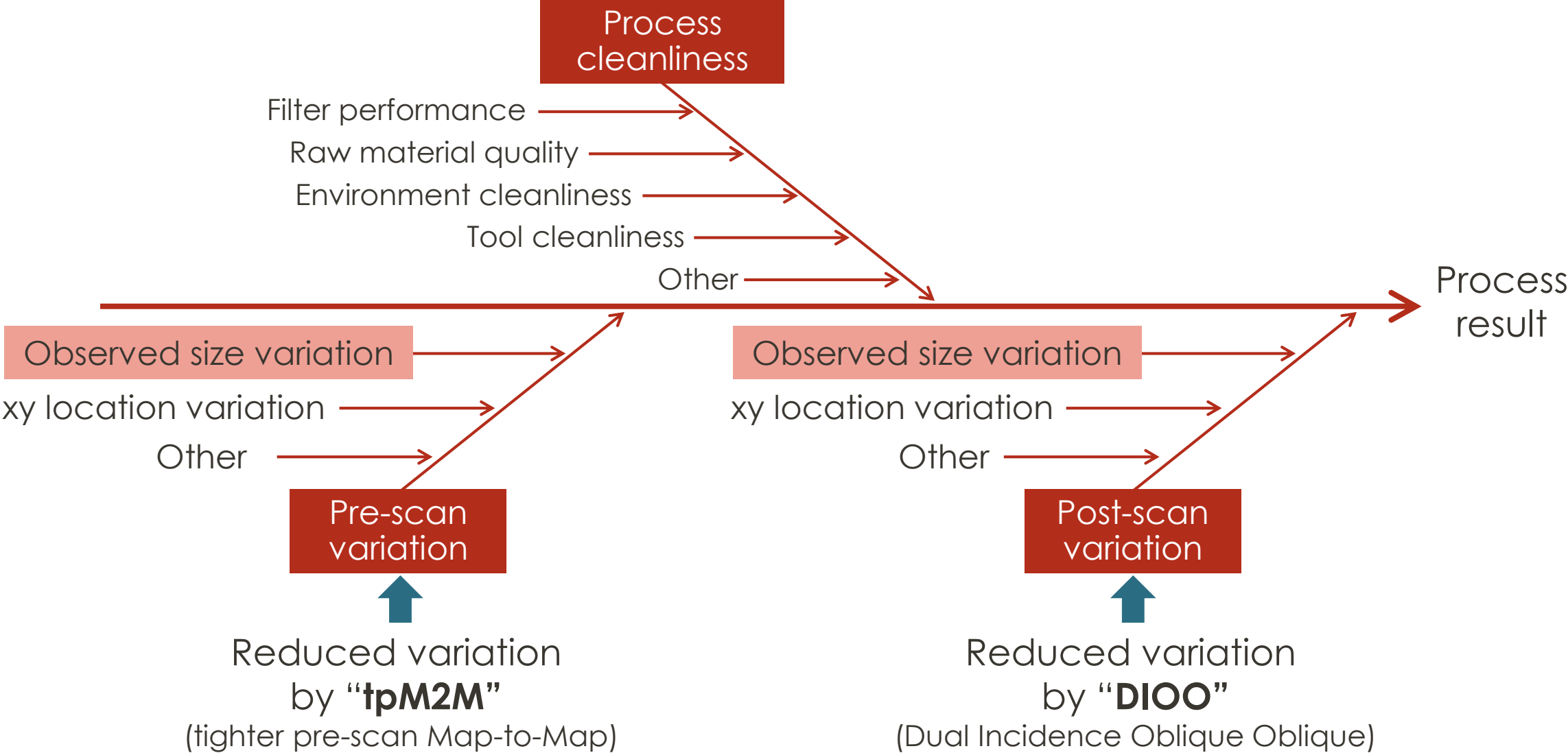
Many factors influence filter choice and defectivity performance



$$\begin{aligned} & \text{Compatibility} \\ & + \\ & \text{Morphology and flow path} \\ & + \\ & \text{Retention mechanisms} \\ & = \\ & \text{Total performance} \end{aligned}$$

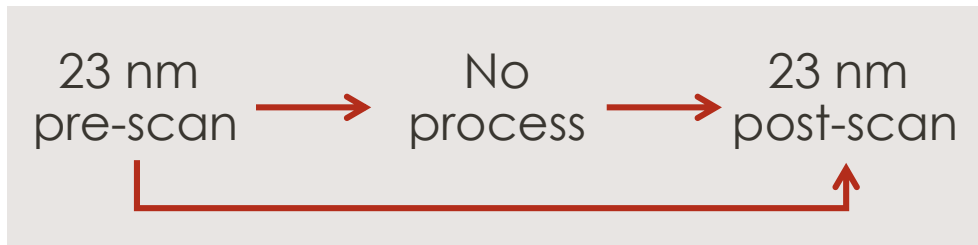
In order to best characterize filter performance in an application, metrology results need to provide meaningful, actionable results.

Fishbone diagram for process result



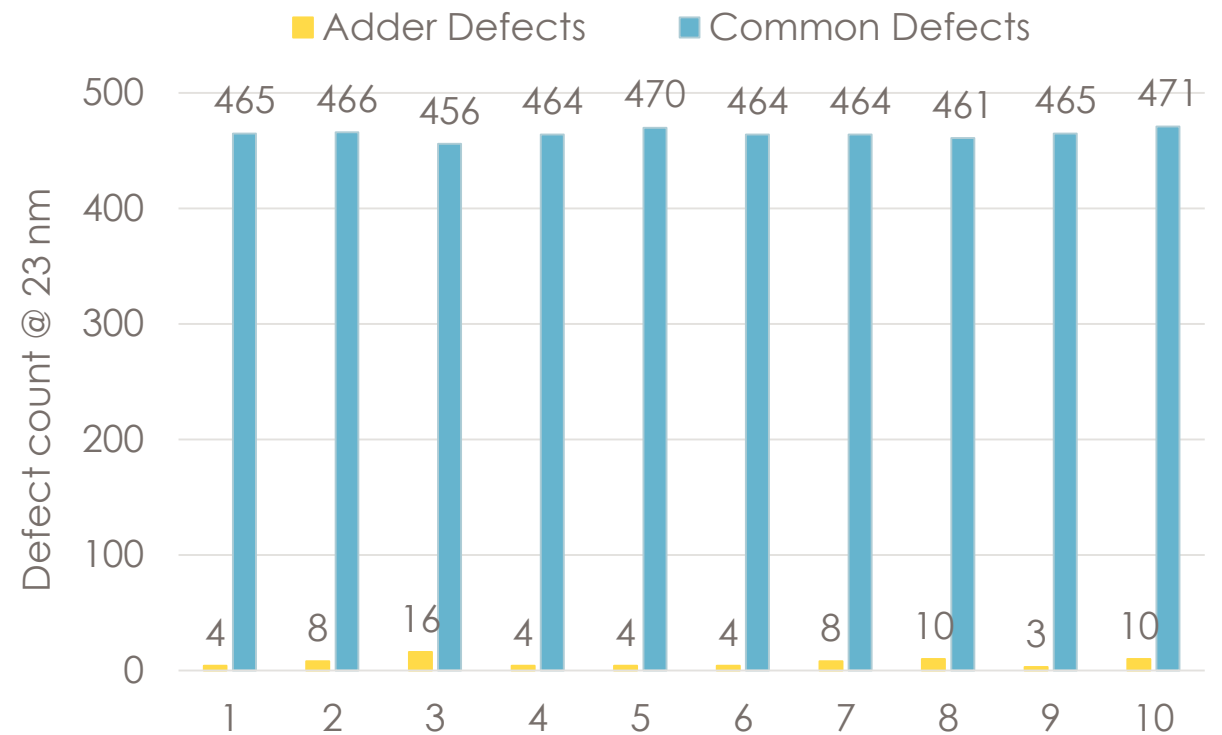
Metrology variation (pre-scan and post-scan) of a Surfscan SP3

#1 Wafer
(processed DIW before)



Overlap 2 maps and calculate adder
Ideally, there should be no adder

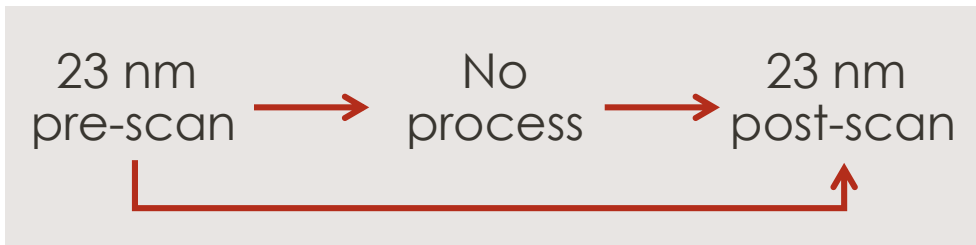
Standard Map-2-Map Adder
w/o wafer processing @ 23 nm
(10 repeat scan on same wafer)



Ideally, we would expect 0 adders in result, but some adders showed at sensitivity threshold

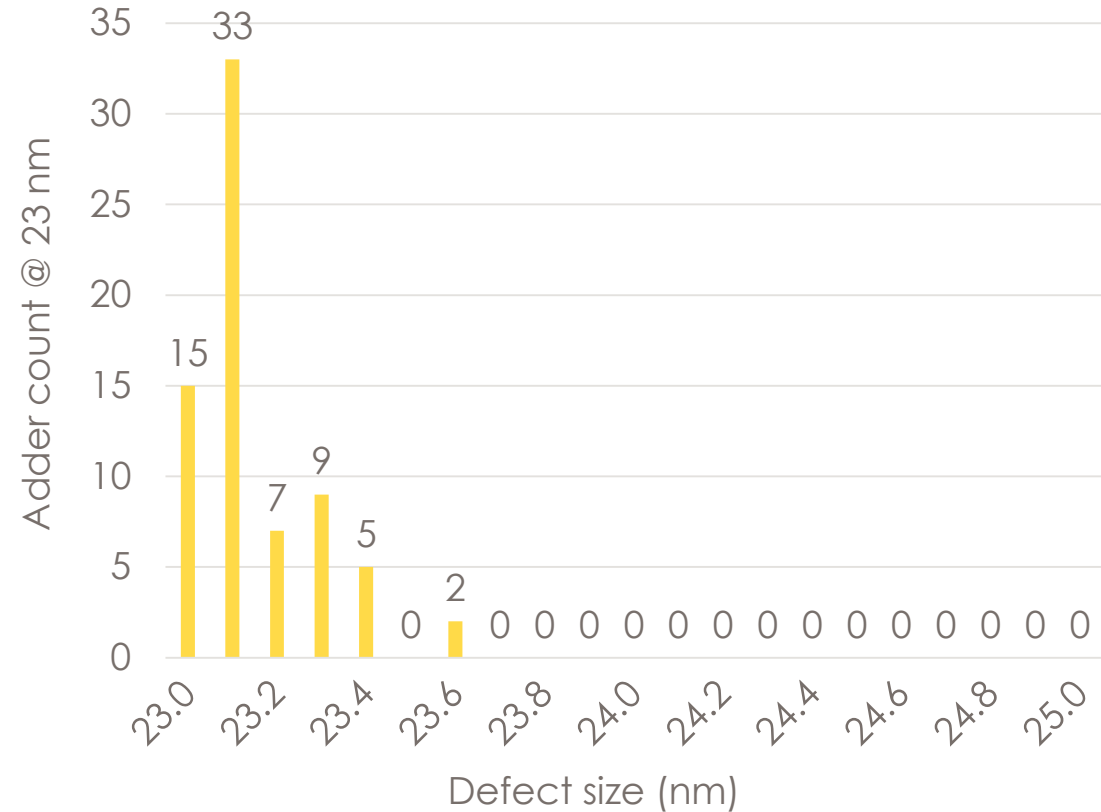
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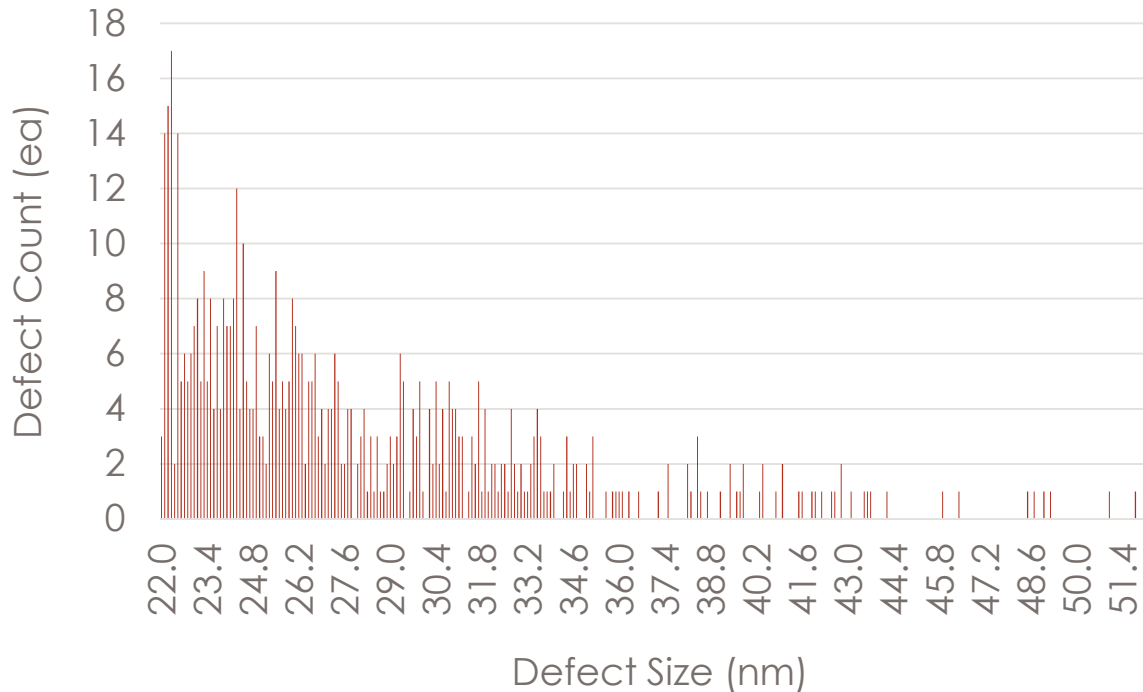
Adder Defects Size Distribution



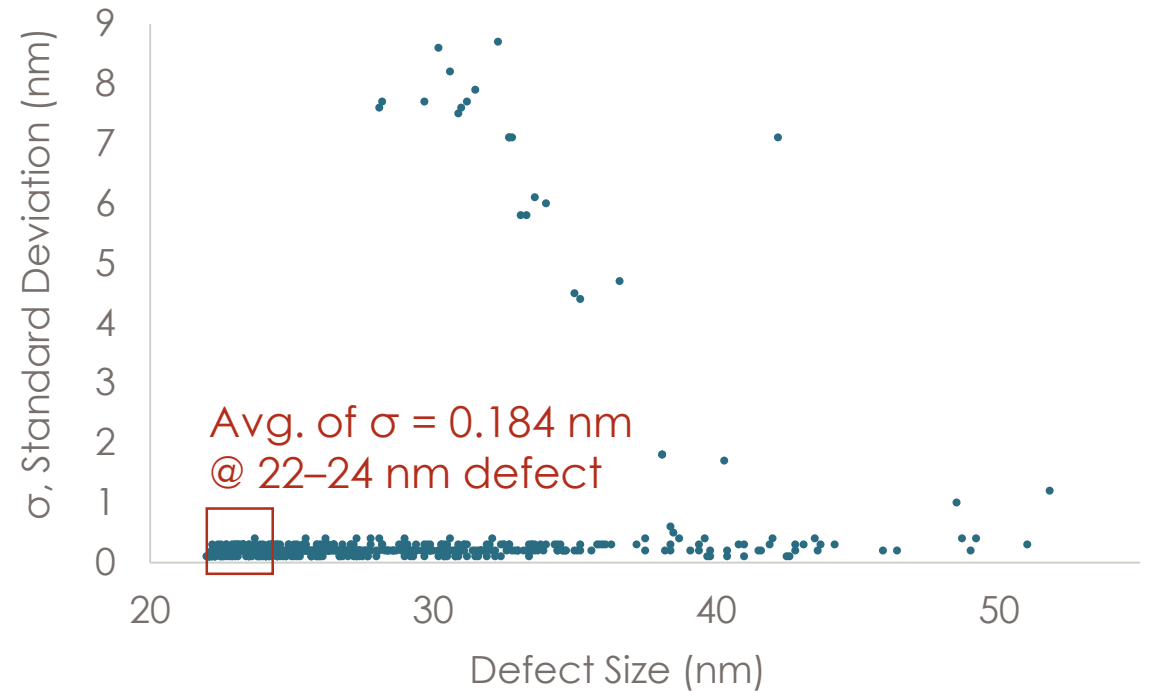
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Size deviation in a bare wafer inspection system

Defect Size Distribution >22 nm
(Average of 10 Repeat Scan at 22 nm)



Size Deviation of each Defect (Standard Deviation of 10 Repeat Scan at 22 nm)

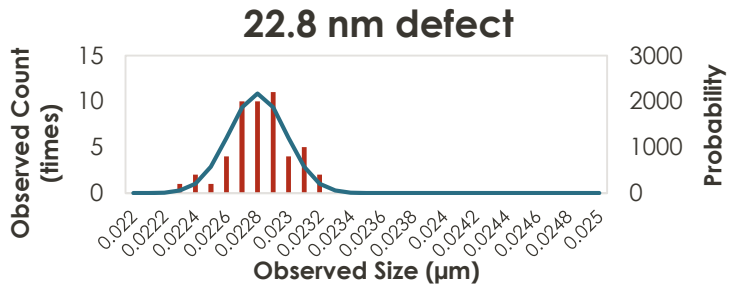
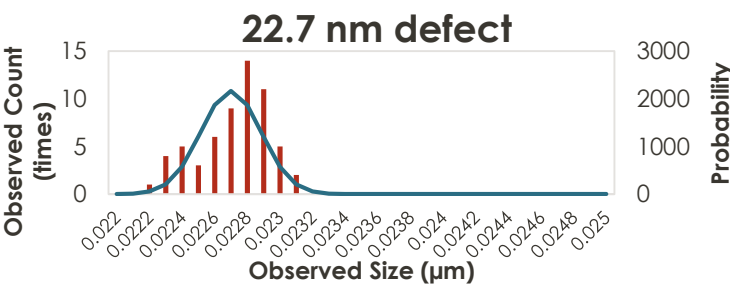
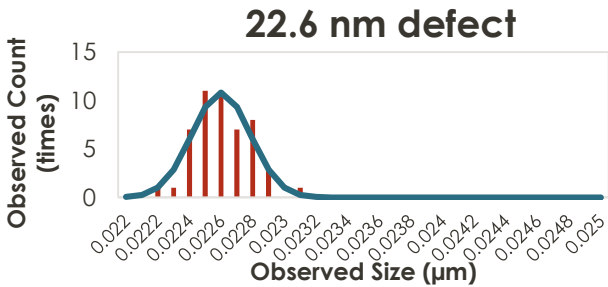
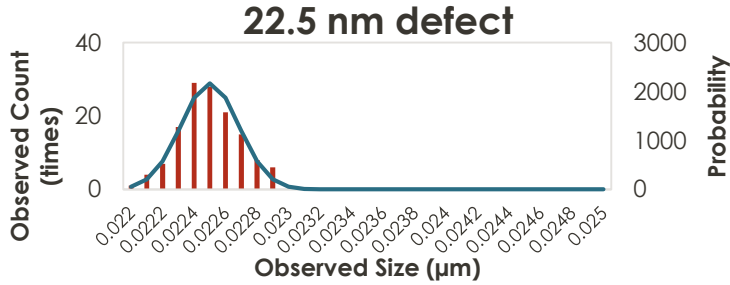
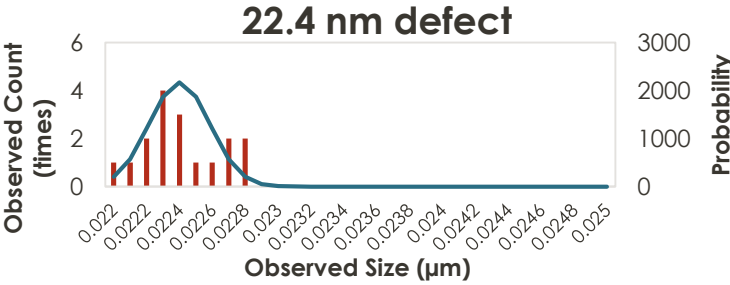
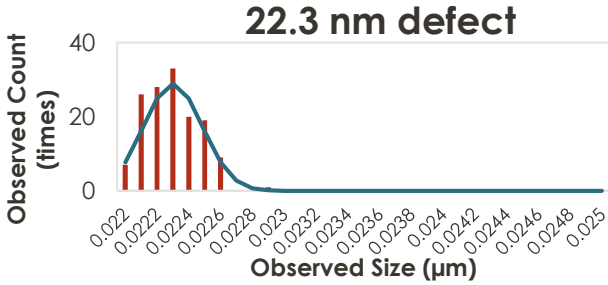
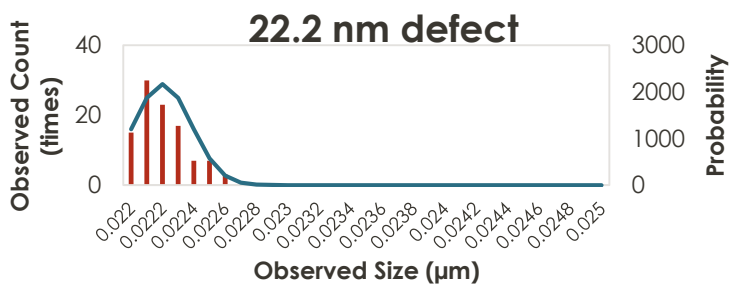
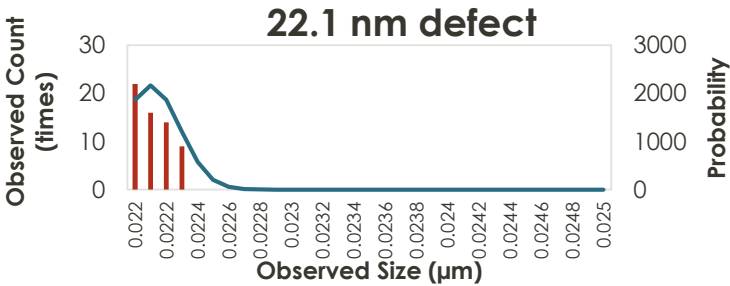
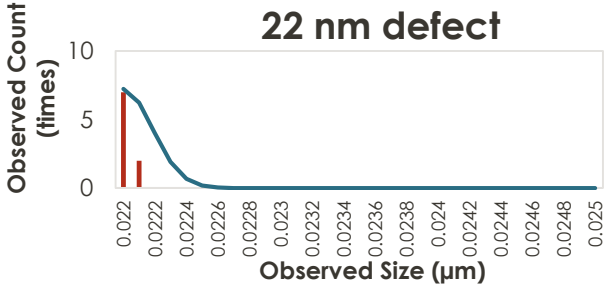


Standard deviation of the observed size is a constant, and equals 0.184nm

Observed size follows Gaussian distribution



— Actual result — Gaussian distribution



The observed size in the 10 scans followed Gaussian distribution

*Defects uncaptured in certain scans are not included into the plot
10 | SPCC 2019

Probability for a defect to be captured

After showing the result from a bare wafer inspection system follow Gaussian distribution, the probability for a defect to be captured can be calculated.

$$\text{Probability} = 1 - 0.5 * (1 + \text{erf}[(\text{Threshold Size} - \text{Actual Size}) / (\sigma * \sqrt{2})])$$

For example, a recipe threshold of 23nm is used to capture a 23nm defect.

$$\text{Probability} = 1 - 0.5 * (1 + \text{erf}[(\mathbf{23} - \mathbf{23}) / (\mathbf{0.184} * \sqrt{2})]) = 50\%$$

Probability of seeing this defect is 50%

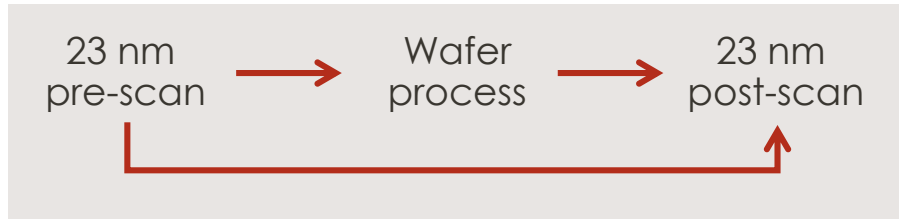
tpM2M Method and Model

(Tighter Pre-scan Map-2-Map)



tpM2M (tighter pre-scan Map-2-Map) method and model

tpM2M method



**sM2M Adder
(Standard Map-2-Map)**



tpM2M Adder

Fewer defects were missed in pre-scans by tpM2M method (fewer false adders)

tpM2M modeling

$$f(x) = \sum_{D=22.0}^{24.0} \left(\frac{1}{2} + \frac{1}{2} \operatorname{erf}\left(\frac{T_{pre} - D}{\sigma\sqrt{2}}\right) \right) * \left(\frac{1}{2} - \frac{1}{2} \operatorname{erf}\left(\frac{T_{post} - D}{\sigma\sqrt{2}}\right) \right) * N_D$$

Probability that a size=D defect CAN be seen post-scan
Count of size=D defect

Probability that a size=D defect CANNOT be seen in pre-scan

T_{pre} = Recipe Threshold at pre-scan (22.0–23.0)

D = Diameter of the defect

σ = Standard deviation in size (Assume = 0.184 = constant)

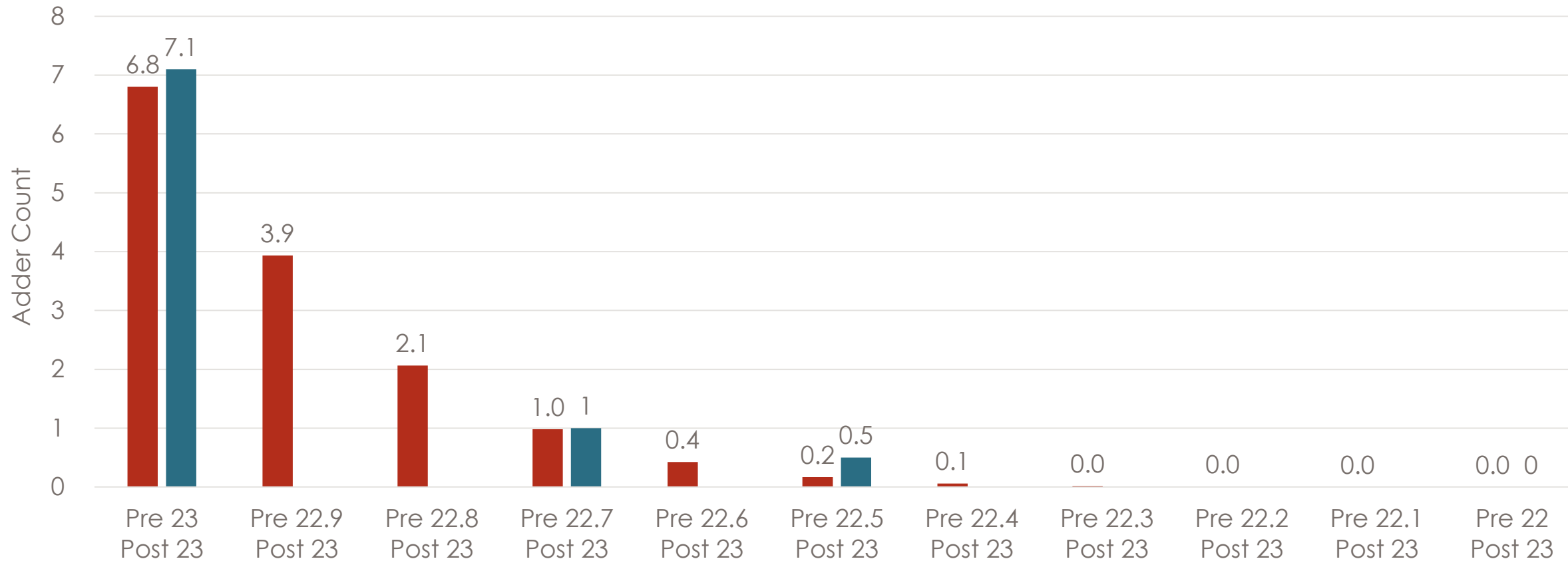
N_D = Count of size = D defect

Assumptions made:

1. All false adder defects came from observed size deviation (e.g. adder caused by x-y deviation is neglected)
2. Observed size follows Gaussian distribution
3. Standard deviation of the observed size is constant and equals 0.184 nm

Particle Adder at different Prescan threshold

■ Modeling Result ■ Actual Result (Avg of 10 Scans)



sM2M

tpM2M

DIOO Method and Model

(Dual Incidence Oblique Oblique)

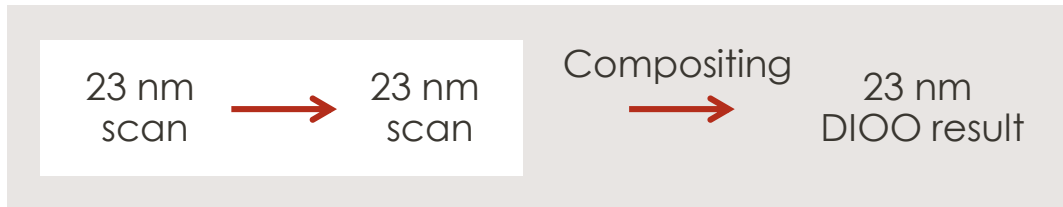


DIOO method and model

DIOO method



Standard



DIOO Method

More true defects were captured in post-scans by DIOO method

Probability for a defect to be seen at different recipe threshold

		Recipe Threshold	
		23	
		Standard	DIOO
Actual Defect Size	22.0	0.0%	0.0%
	22.1	0.0%	0.0%
	22.2	0.0%	0.0%
	22.3	0.0%	0.0%
	22.4	0.1%	0.20%
	22.5	0.3%	0.60%
	22.6	1.5%	2.98%
	22.7	5.2%	10.13%
	22.8	13.9%	25.87%
	22.9	29.3%	50.02%
	23.0	50.0%	75.00%
	23.1	70.7%	91.42%
	23.2	86.1%	98.07%
	23.3	94.8%	99.73%
	23.4	98.5%	99.98%
	23.5	99.7%	100.0%
	23.6	99.9%	100.0%
	23.7	100.0%	100.0%
	23.8	100.0%	100.0%
	23.9	100.0%	100.0%
24.0	100.0%	100.0%	

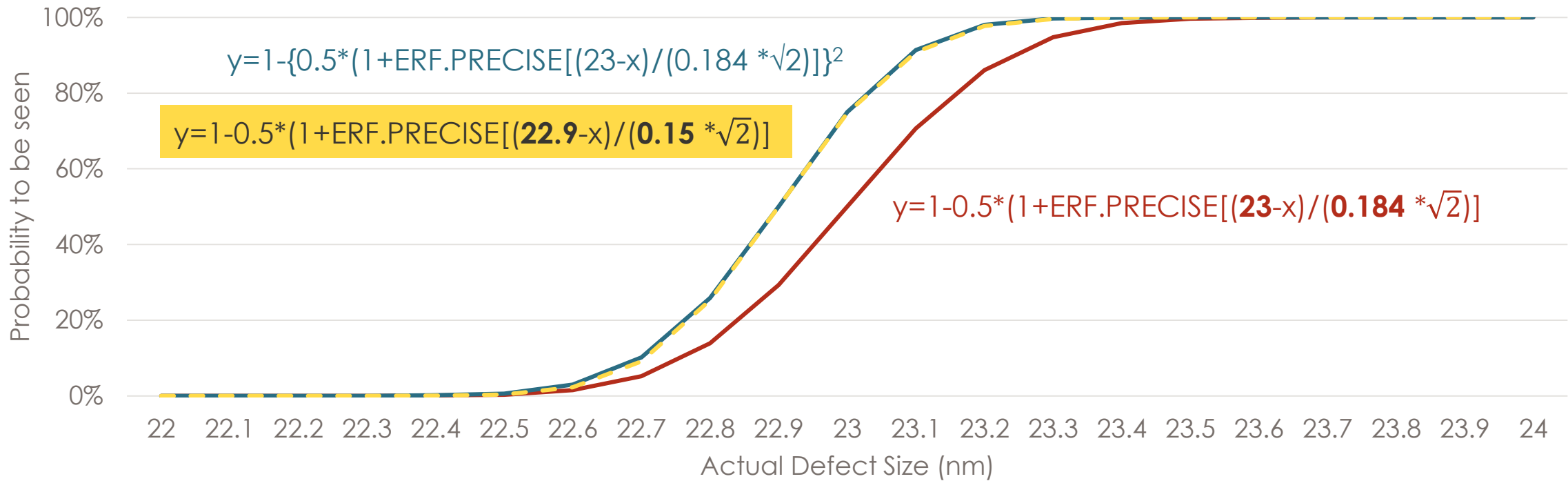
$$\text{Standard} = 1 - 0.5 * (1 + \text{ERF.PRECISE}[(\text{Threshold Size} - \text{Actual Size}) / (\sigma * \sqrt{2})])$$

$$\text{DIOO} = 1 - \{0.5 * (1 + \text{ERF.PRECISE}[(\text{Threshold Size} - \text{Actual Size}) / (\sigma * \sqrt{2})])\}^2$$

DIOO method and model

Probability for a defect to be seen at 23 nm recipe threshold

Standard DIOO Model



DIOO probability curve fit the normal distribution curve with $\mu = 22.9$ and $\sigma = 0.15$

We can consider the standard deviation of observed size is reduced from 0.184 nm to 0.15 nm

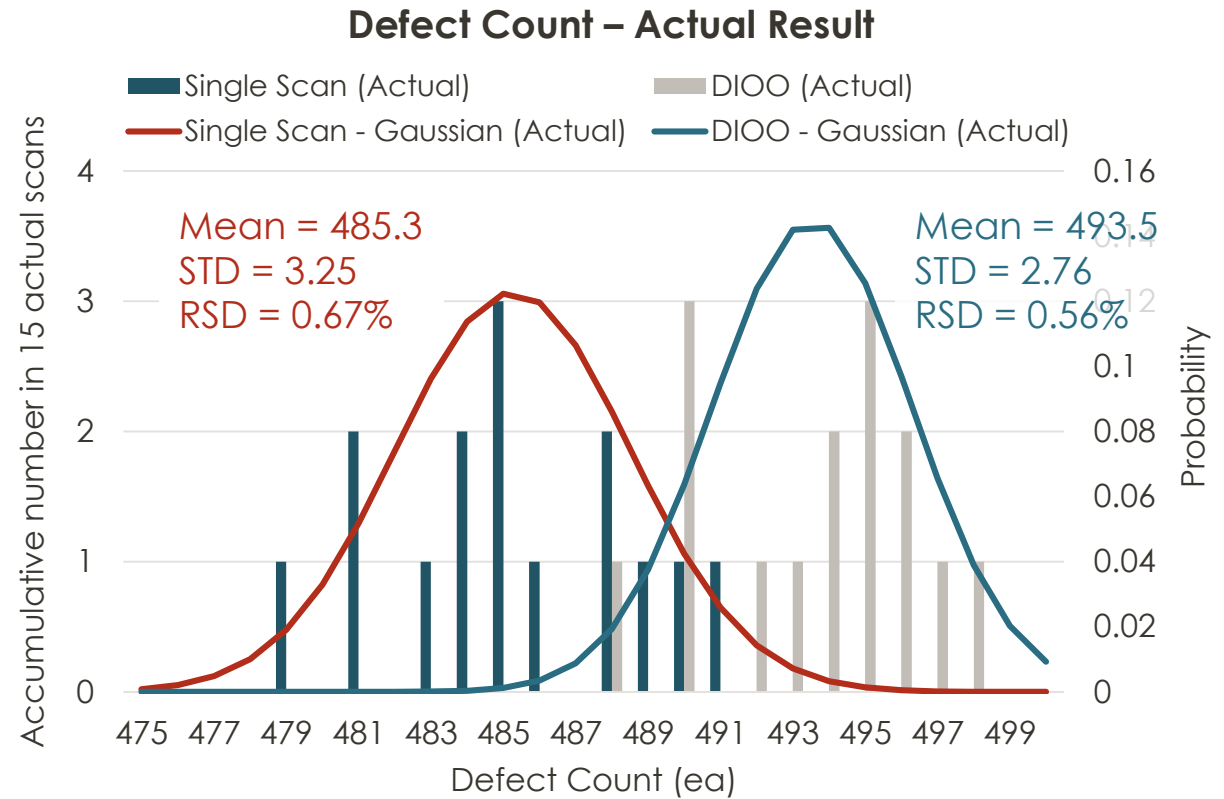
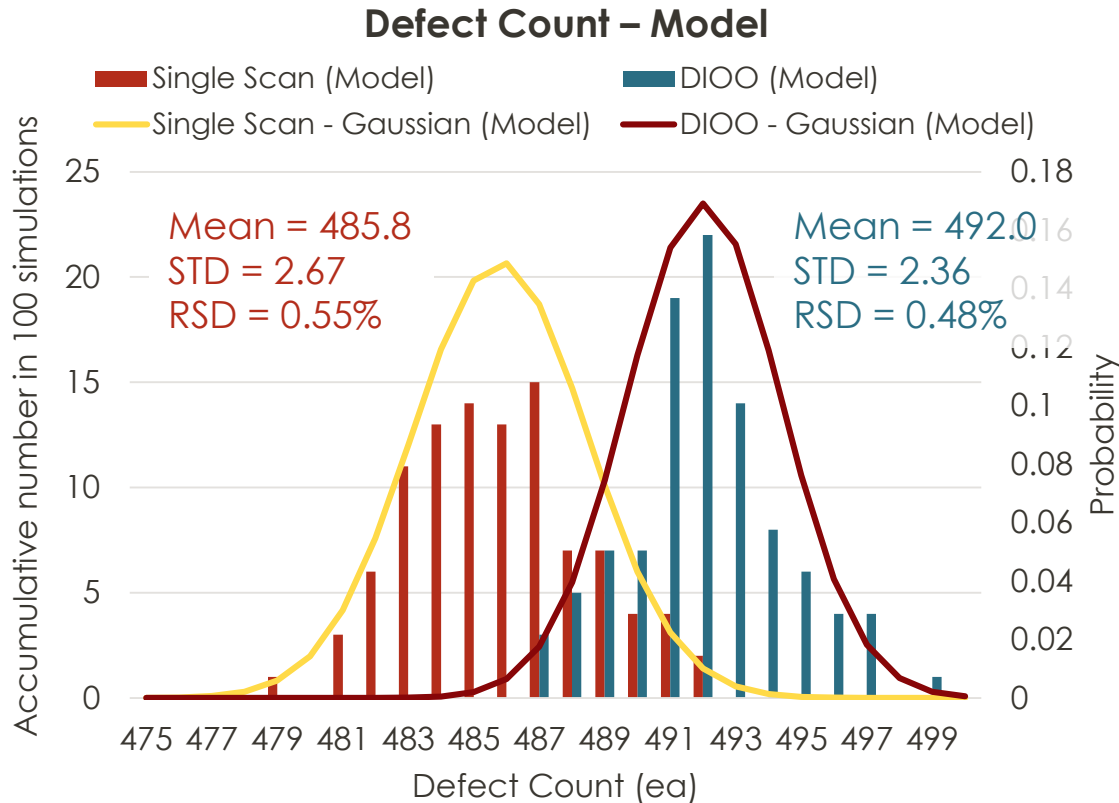
DIOO method and model



Wafer #2 was scanned 10 times at 22 nm recipe threshold

The average and standard deviation in size of each defect were used to calculate the theoretical count of 100 scans (left) at 23 nm recipe threshold

The actual wafer result of 15 scans at 23 nm (right) pretty much aligned with the prediction



Example of applying
tpM2M plus DIOO method
in Wet Process Monitor



Applying tpM2M + DIOO on filter evaluation

Filter installed onto single wafer tool



Flushed >1500 L DIW



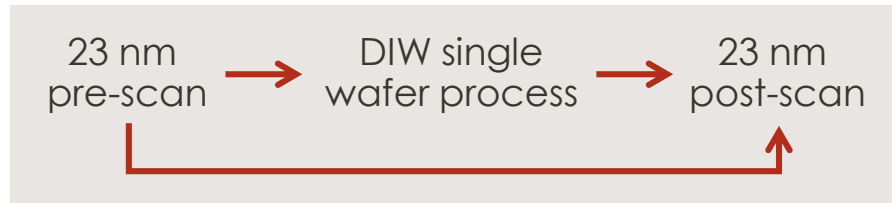
Standard



tpM2M only



DIOO only



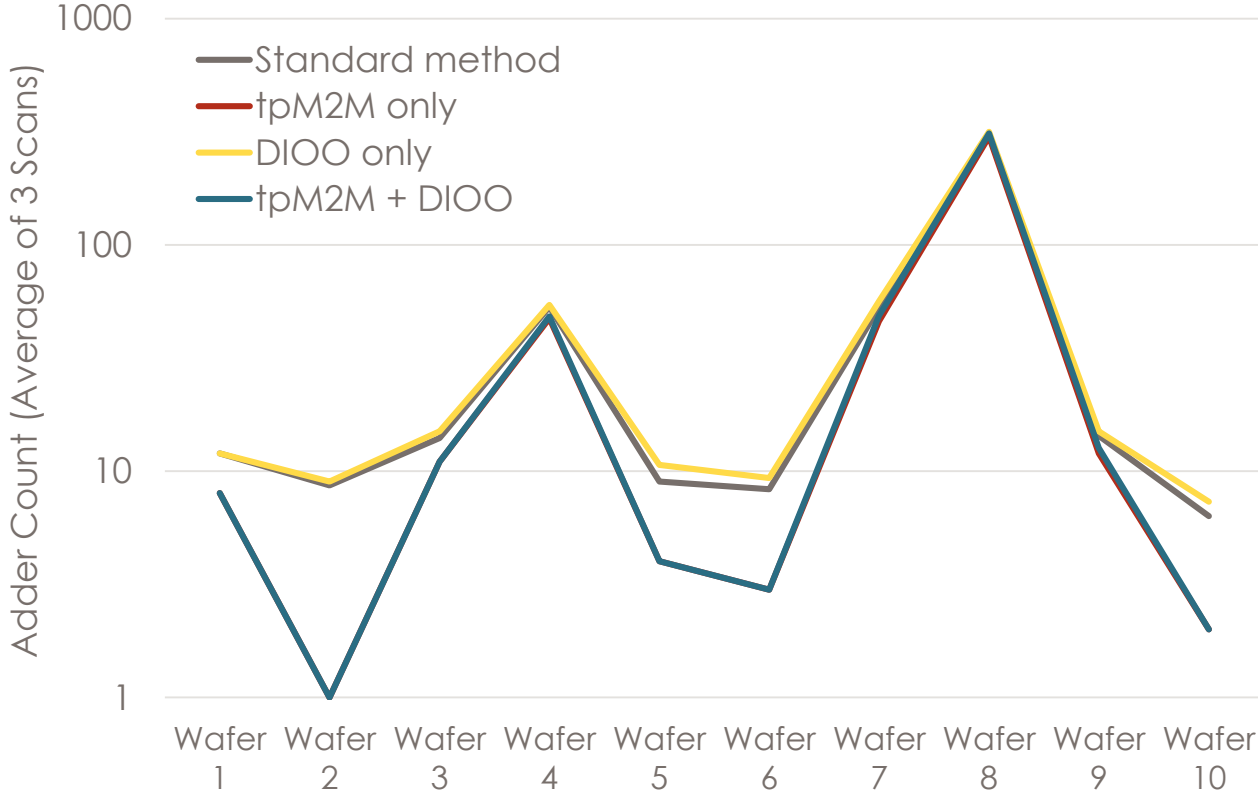
tpM2M + DIOO



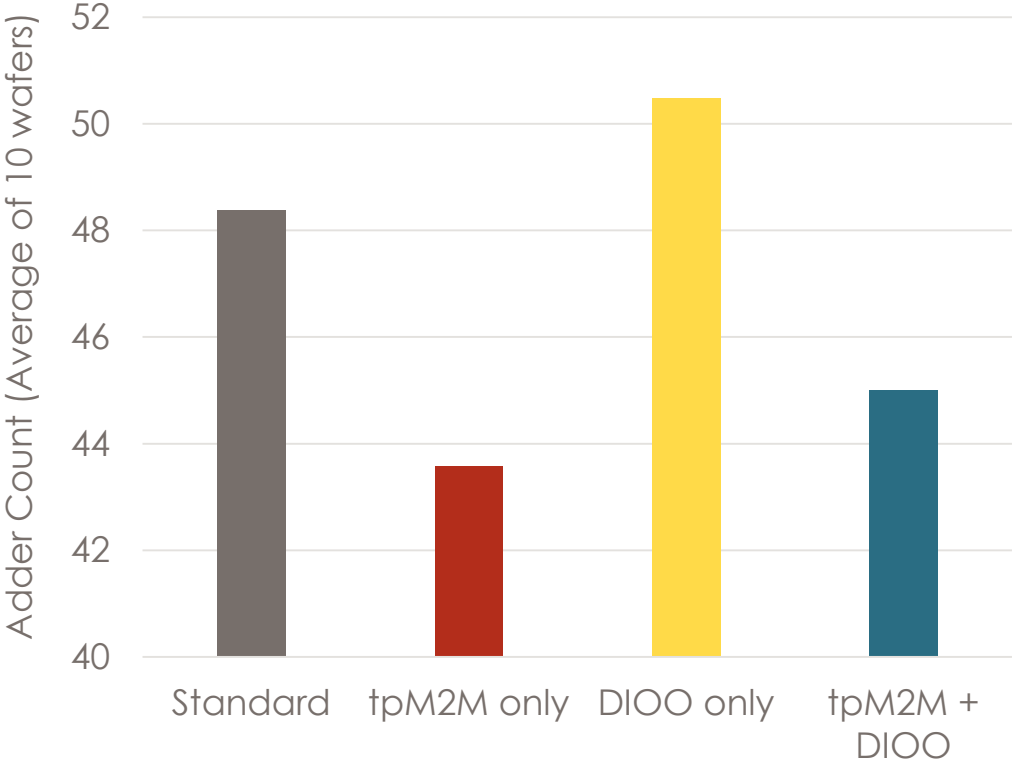
10 wafers/method
3 repeats in post-scan

Results of applying tpM2M and DIOO – average

WPA – X bar Chart of 3 Repeats



WPA – X bar bar Chart of 10 Wafers

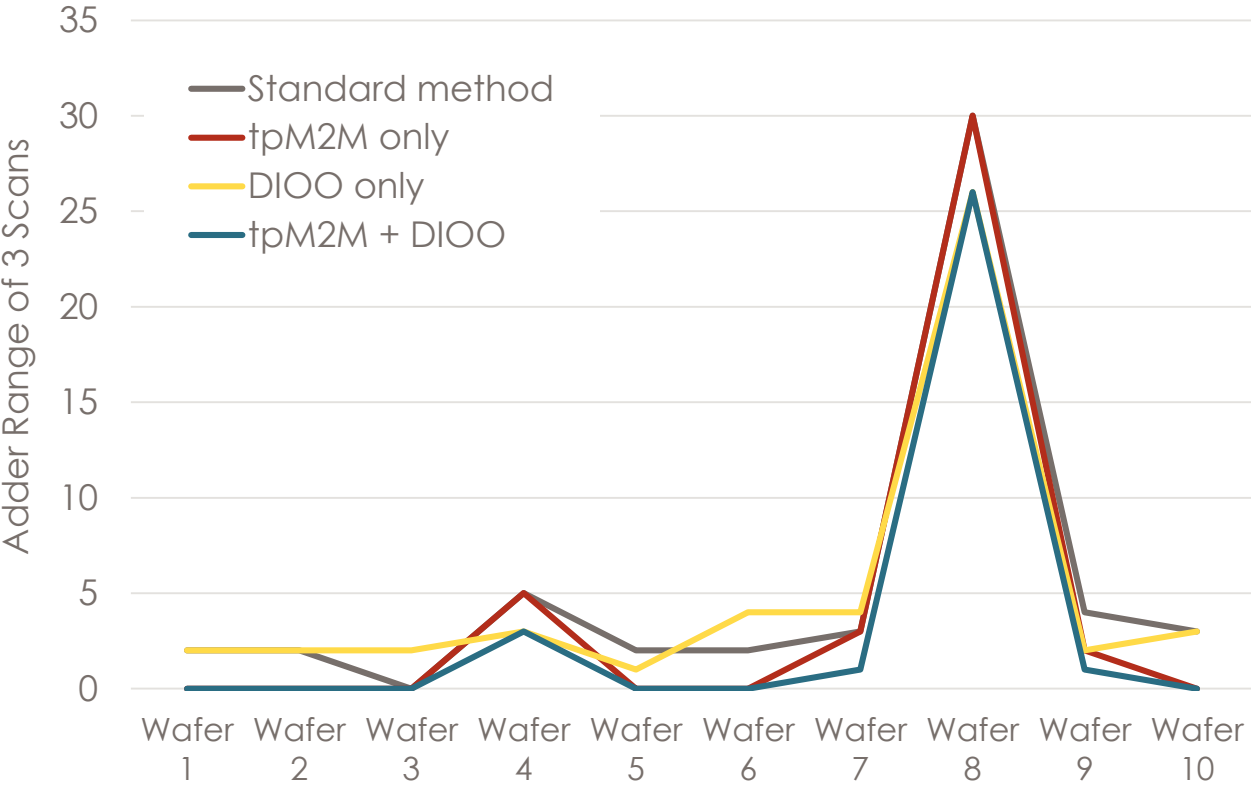


tpM2M reduced false adder and gives a lower count
DIOO slightly increased sensitivity and gives a higher count

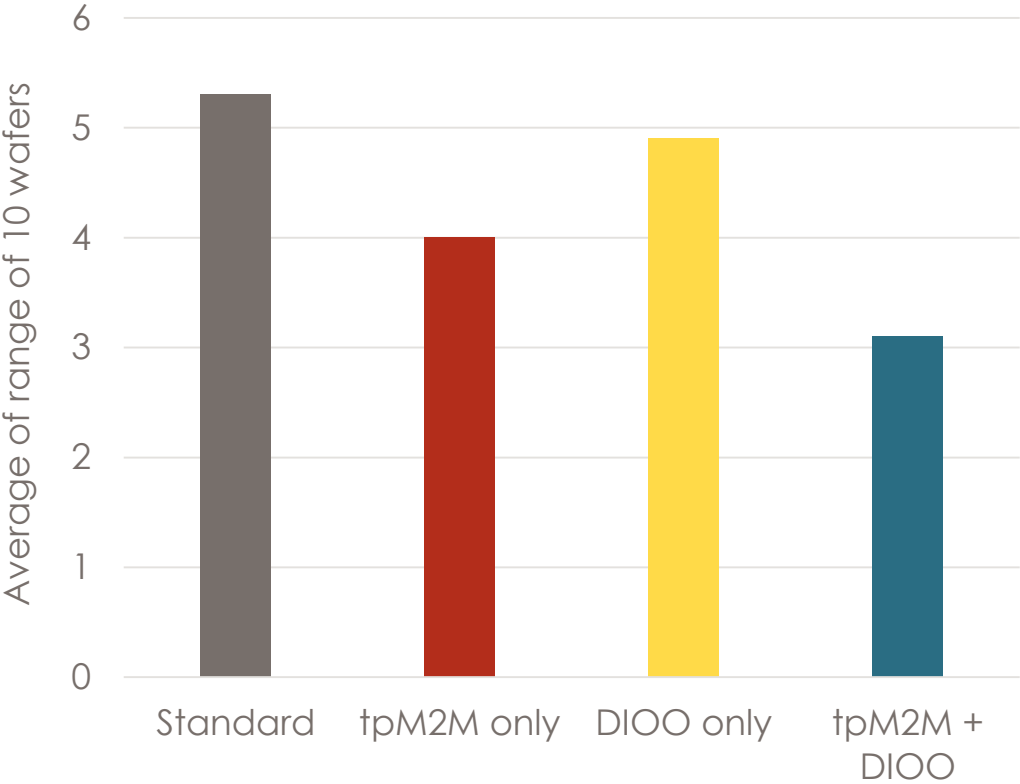


Results of applying tpM2M and DIOO – range

WPA – R Chart of 3 Scans



WPA – R bar Chart of 10 Wafers



tpM2M and DIOO method both reduced the WPA count variation of this DIW process (tpM2M more than DIOO)

A combination of applying both tpM2M and DIOO gives the best result



Summary

Reducing variation generated by on-wafer inspections is essential, so we can make better decisions regarding equipment/material

Results from an unpatterned wafer defect inspection system (e.g. Surfscan) follows Gaussian distribution

tpM2M and DIOO methods were introduced with a model to reduce the variation

A combination of tpM2M and DIOO methods is suggested to give the best result when evaluating a process/equipment/material

Thank you!

A white speech bubble with a tail pointing down and to the left, containing the text "Q&A". The bubble is centered over a background of silhouettes of people in a meeting, with a city skyline and digital data lines visible in the background.

Q&A