

# Using artificial intelligence to distinguish subjects with prostate cancer (PCa) from benign prostate hyperplasia (BPH) through immunophenotyping of MDSCs and lymphocyte cell populations

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Society for Immunotherapy of Cancer

# Disclosure Information

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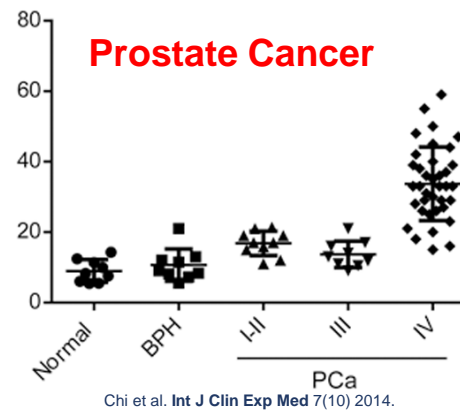
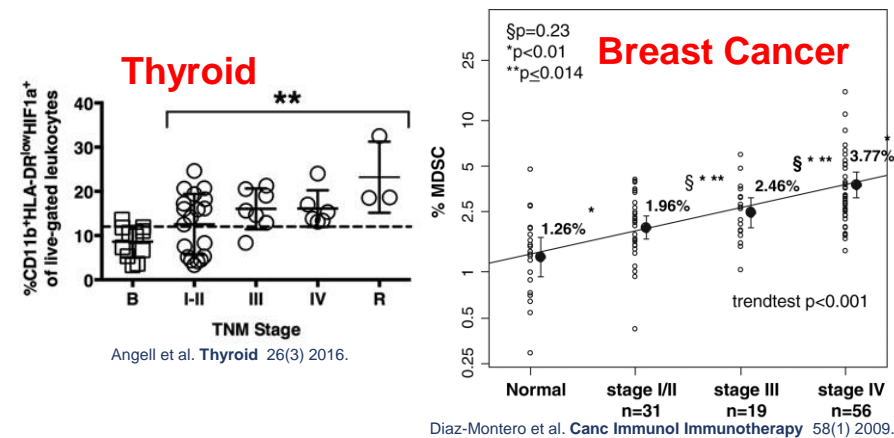
George Dominguez

Employee - Anixa Biosciences, Inc.

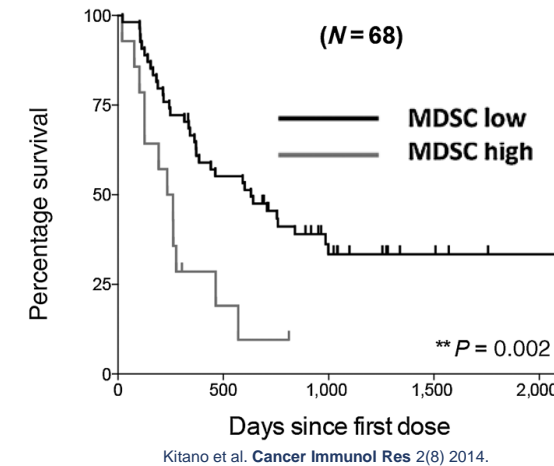
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# What can MDSCs tell us?

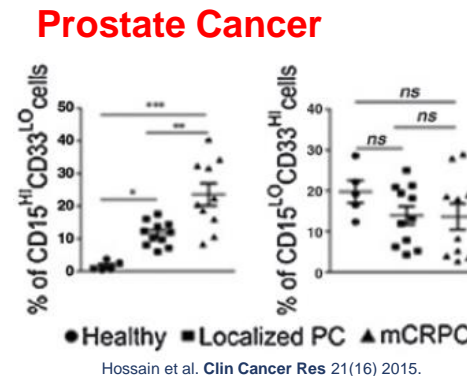
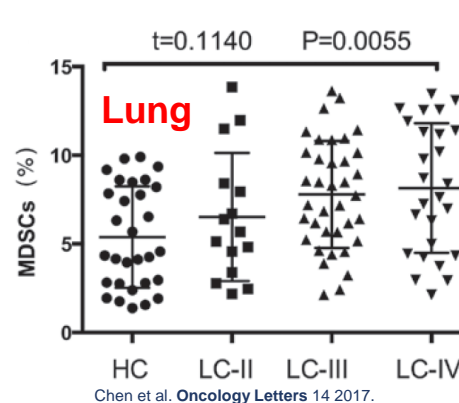
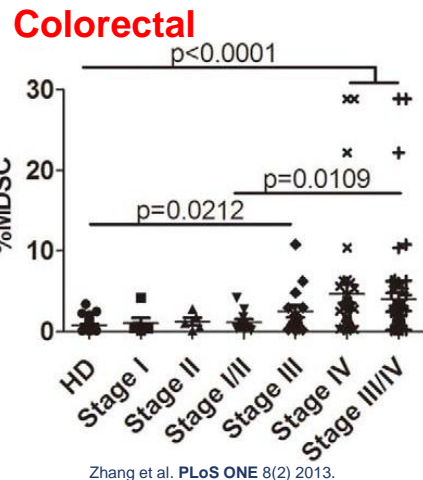
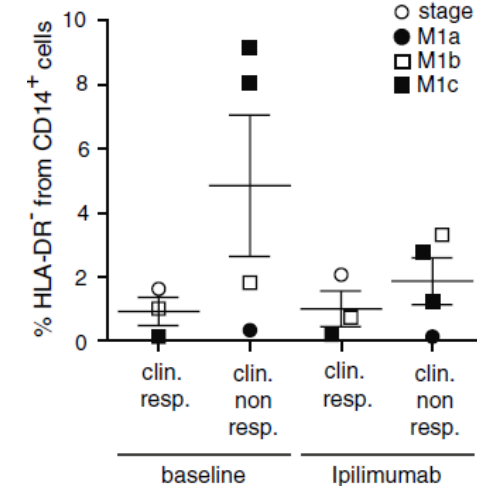
## Indicative of Solid Tumors and Severity



## Predictors of Immunotherapy Response?



## CTLA-4 Responses in Late Stage Melanoma



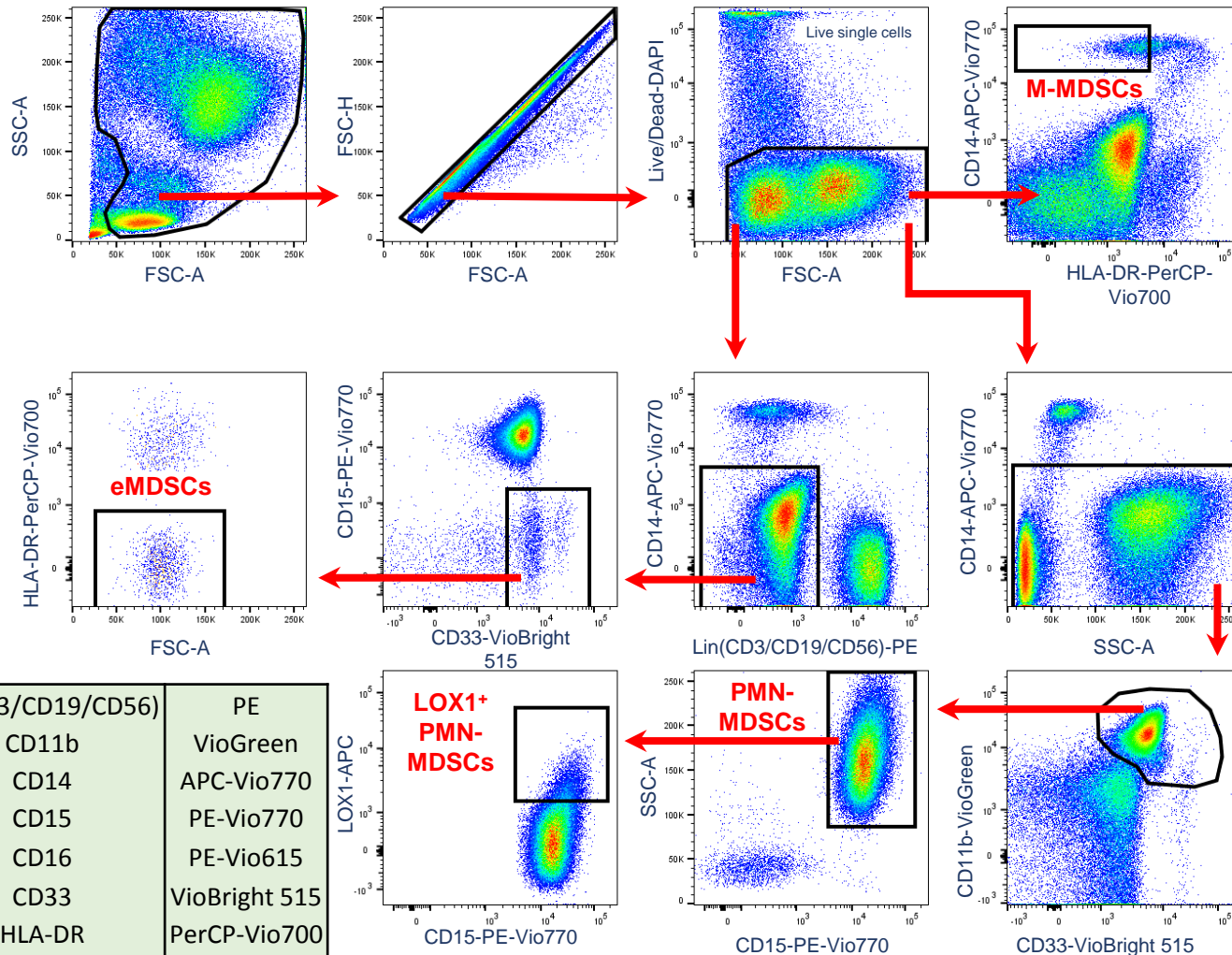
# Question

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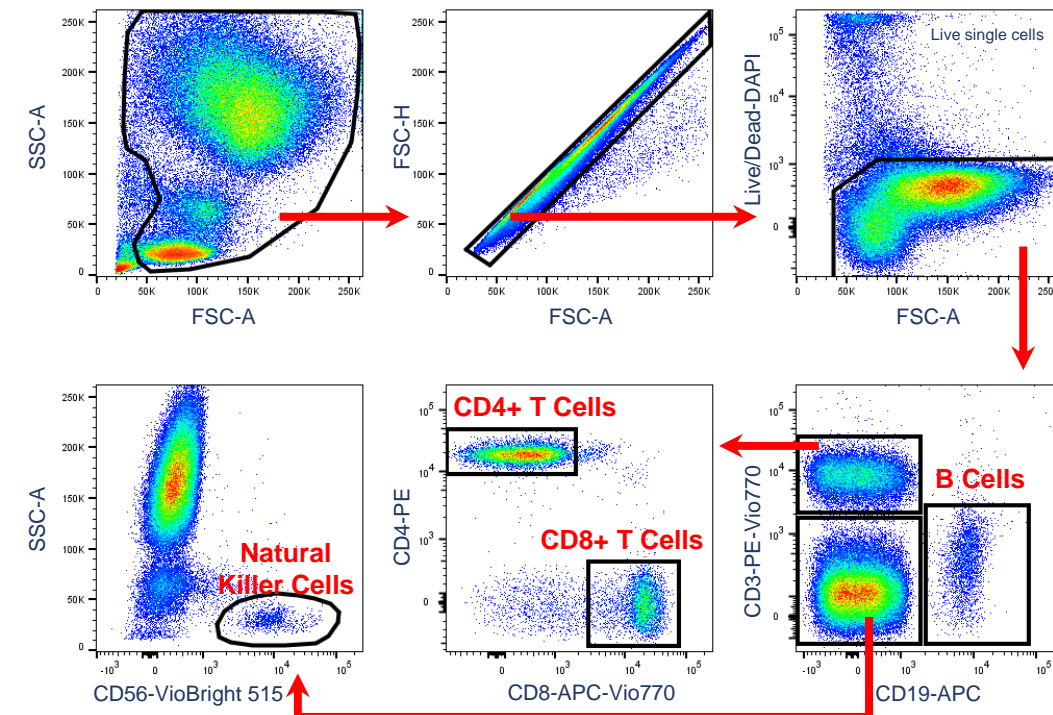
Can we use MDSCs as an indicator for higher risk prostate cancer (PCa) and distinguish from benign prostatic hyperplasia (BPH)/lower risk PCa?

# What are we measuring?

## Myeloid Panel (8 Markers)



## Lymphoid Panel (5 Markers)



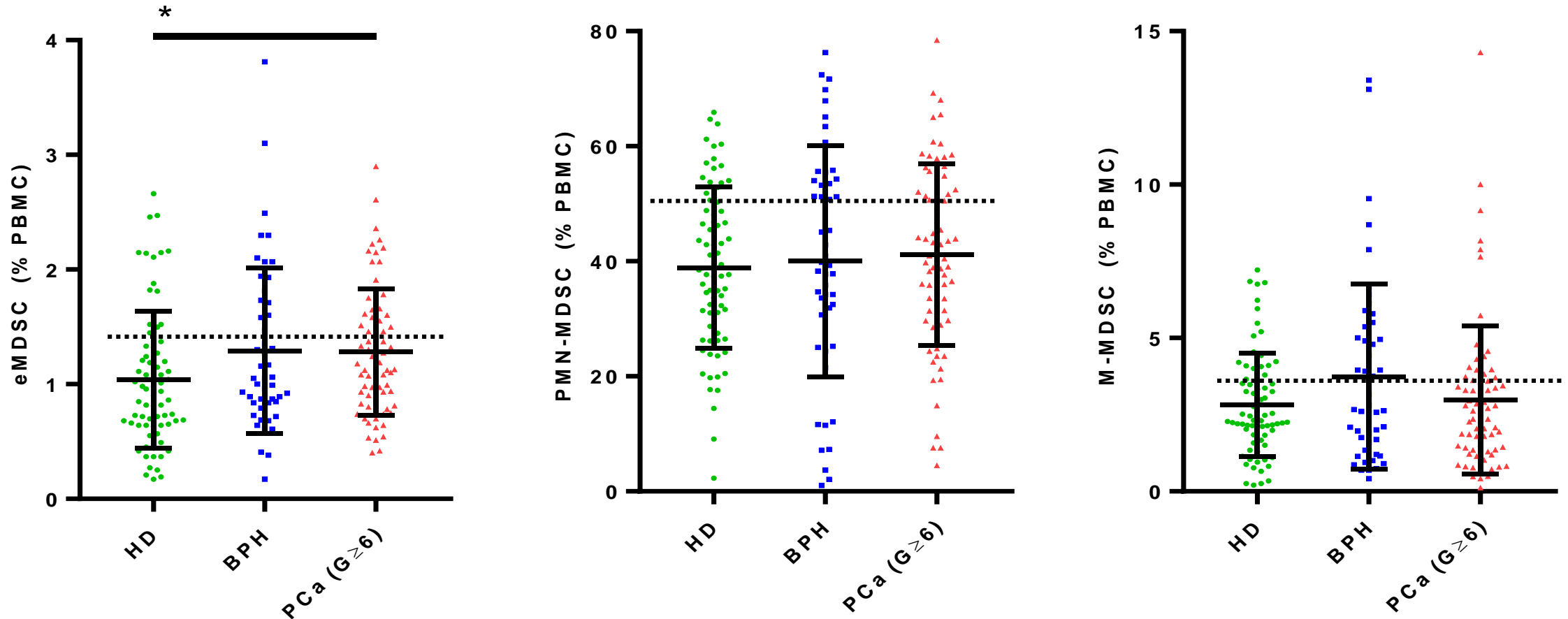
CD3	PE-Vio770
CD4	PE
CD8	APC-Vio770
CD19	APC
CD56	VioBright 515
Live/Dead	DAPI

# Clinical Characteristics and Categorization

- Prospective blood collection – processed within 20 to 30 hours
- All subjects were already scheduled to undergo a transrectal ultrasound guided prostate (TRUSP) biopsy
- Subjects not included if they had:
  - previous history of cancer (excluding active surveillance)
  - any previous medical intervention for PCa
  - on active treatment for BPH

Characteristic	PCa	BPH	HD
Total	73	48	73
Median Age	65	62	53
Age Range	44 - 86	40 - 81	22 - 79
Gleason Score			
6	26		
7 (3+4)	14		
7 (4+3)	15		
>8	18		
Tumor Stage			
T1c	43		
T2a	2		
Unknown	28		

# Traditional Gating: Manual Counting - MDSCs



***Simple cell counts can provide information about trends, but can only categorize some subjects***

Healthy Donor (HD) n = 73

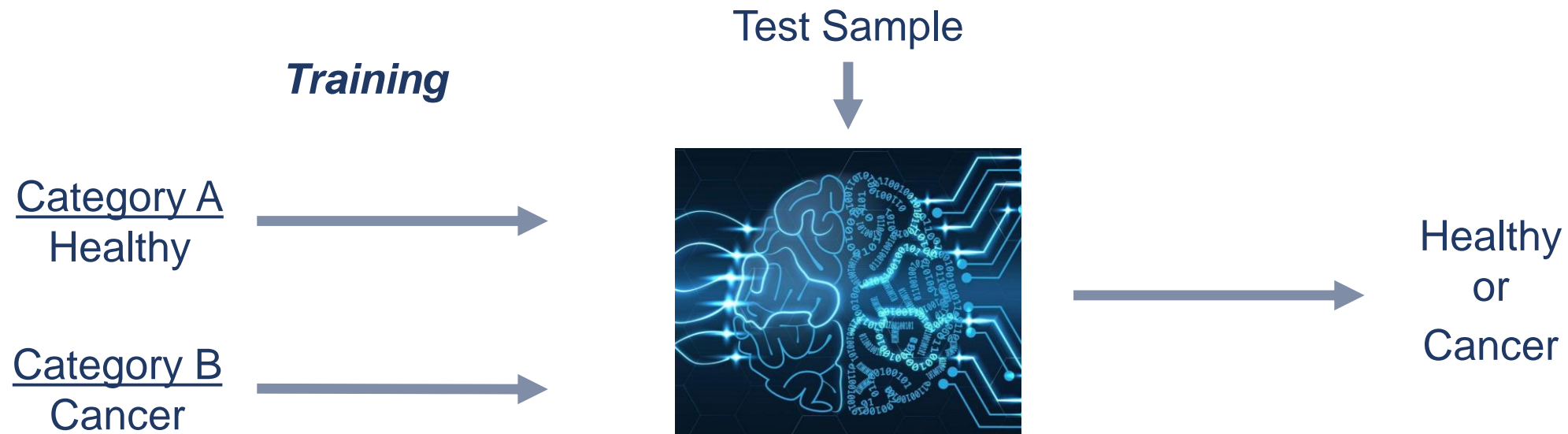
Benign Prostatic Hyperplasia (BPH) n = 48

Prostate Cancer (PCa) n = 73

\*The dotted line represents the 75<sup>th</sup> percentile for HD values (75% specificity)

# Our Question

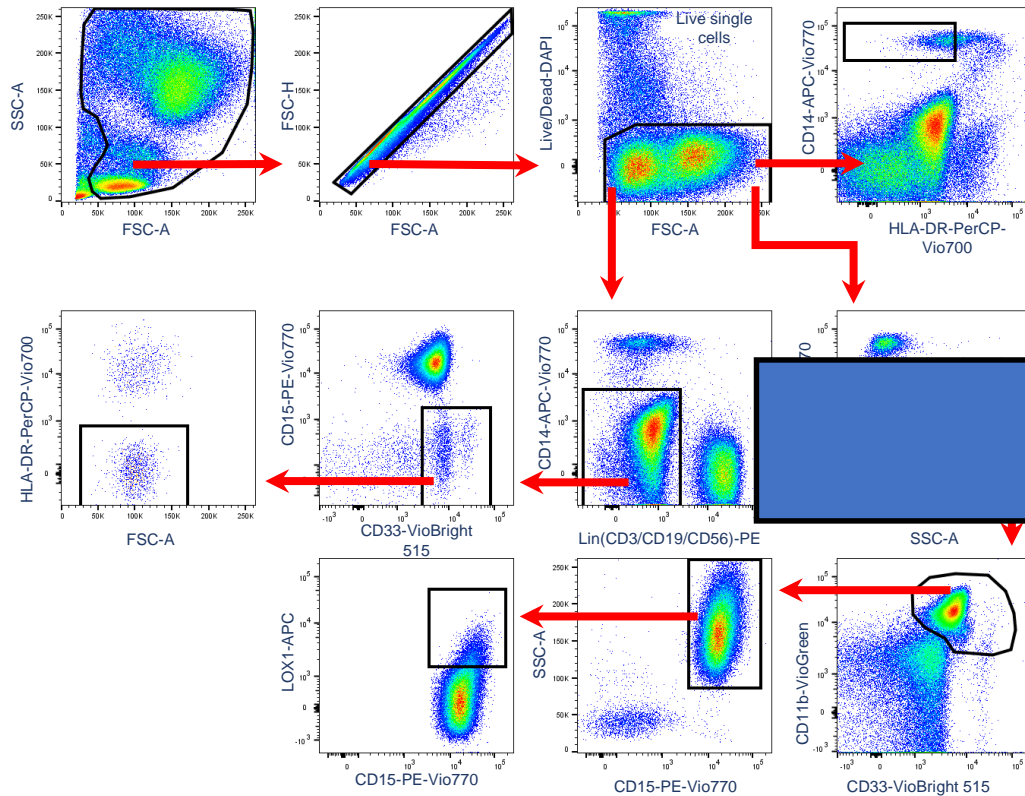
Can we use machine learning (neural networks) to analyze the flow cytometry data to categorize patients?



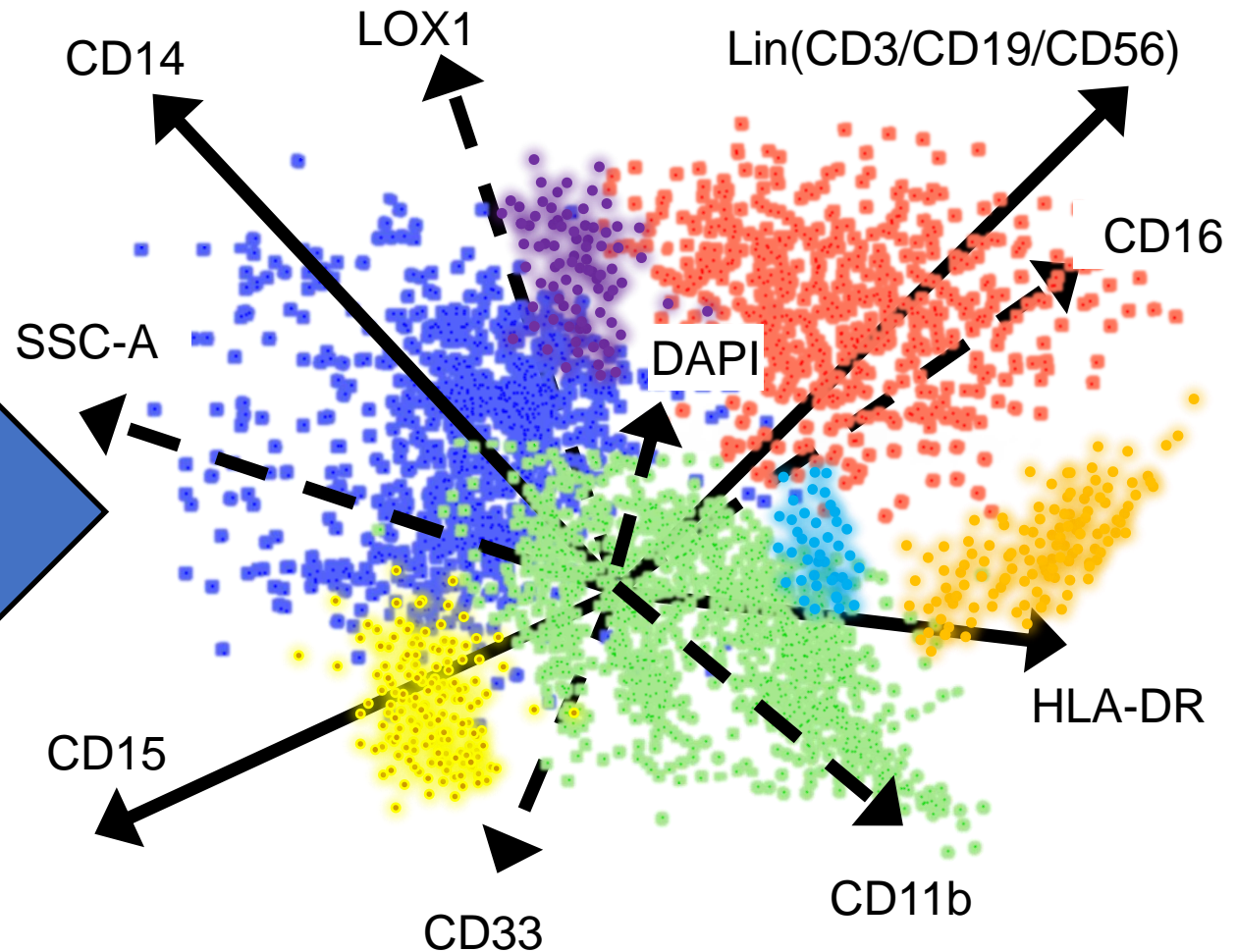


# The Inputs – Event Counts

FCS File → CSV File (Event counts for each channel)



## Multidimensional Space

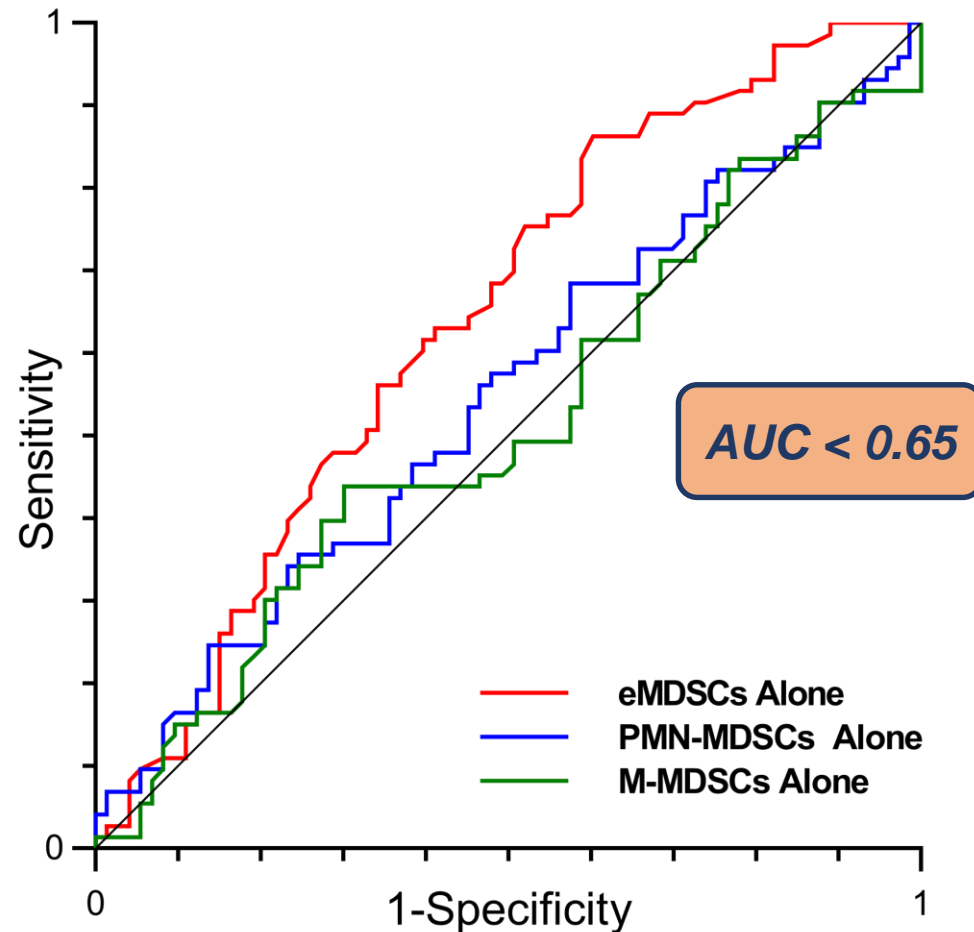


The Neural Network does not see gates, markers, or fluorophores – it just sees numbers

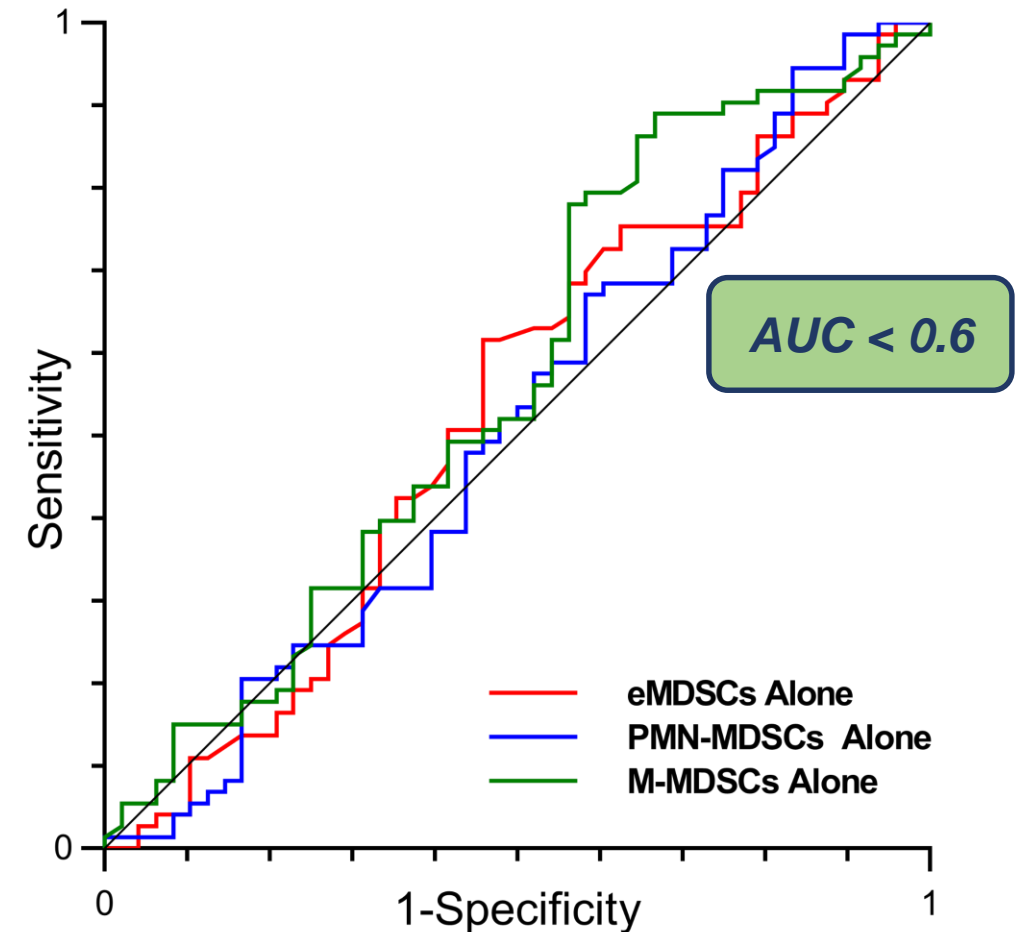
# Manual Gating – not enough...

\*PCa (Adenocarcinoma) = Gleason  $\geq 6$

## Healthy Donor vs Prostate Cancer



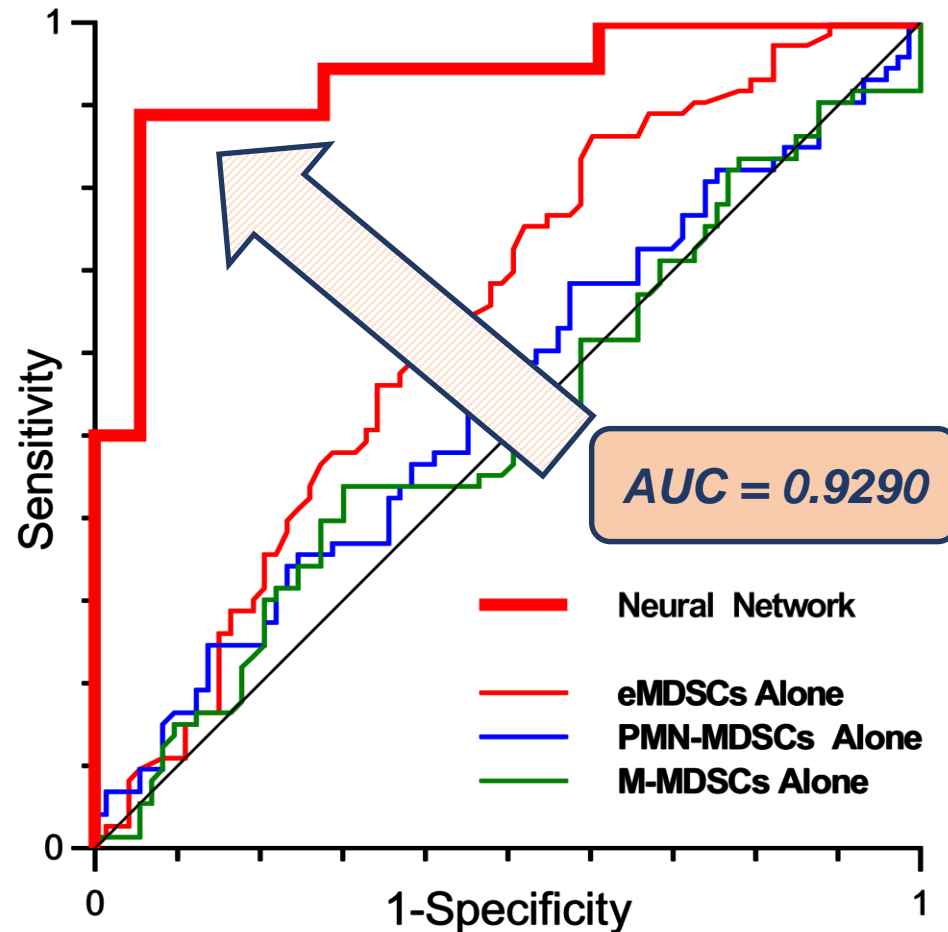
## BPH vs Prostate Cancer



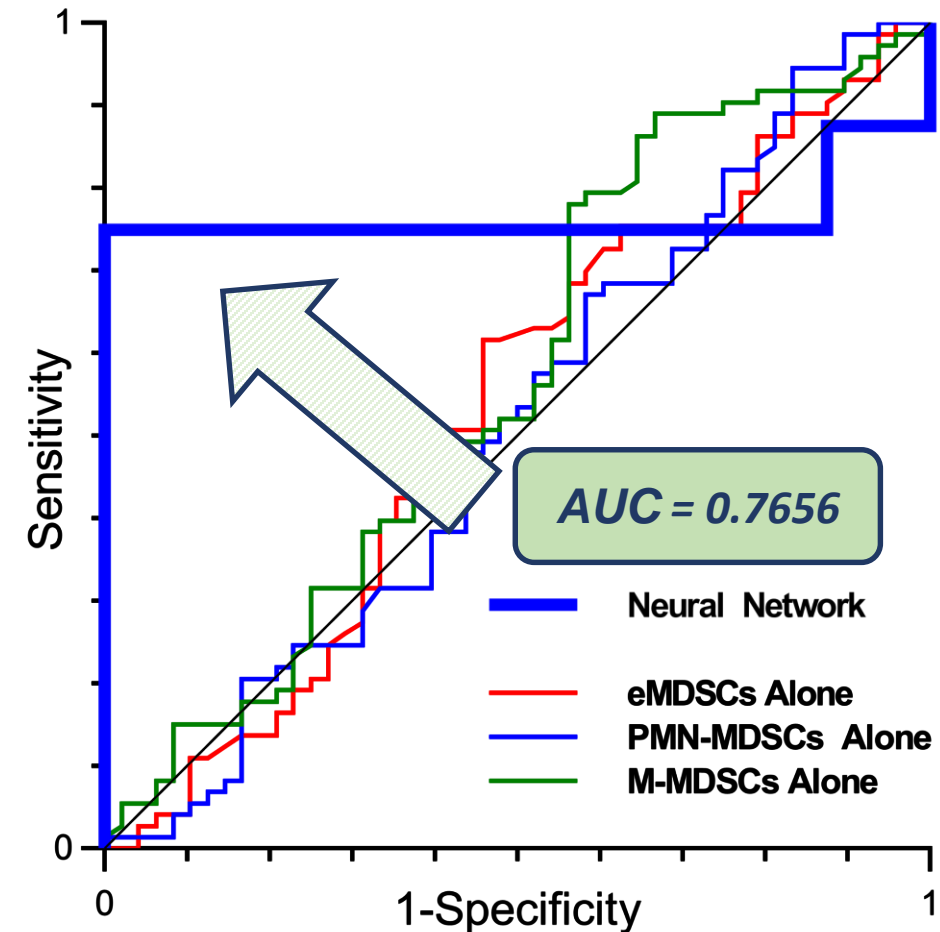
# Manual Gating – not enough...

\*PCa (Adenocarcinoma) = Gleason  $\geq 6$

## Healthy Donor vs Prostate Cancer



## BPH vs Prostate Cancer



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What is a clinical application of this technology?

# Clinical Application: Confirmatory Testing

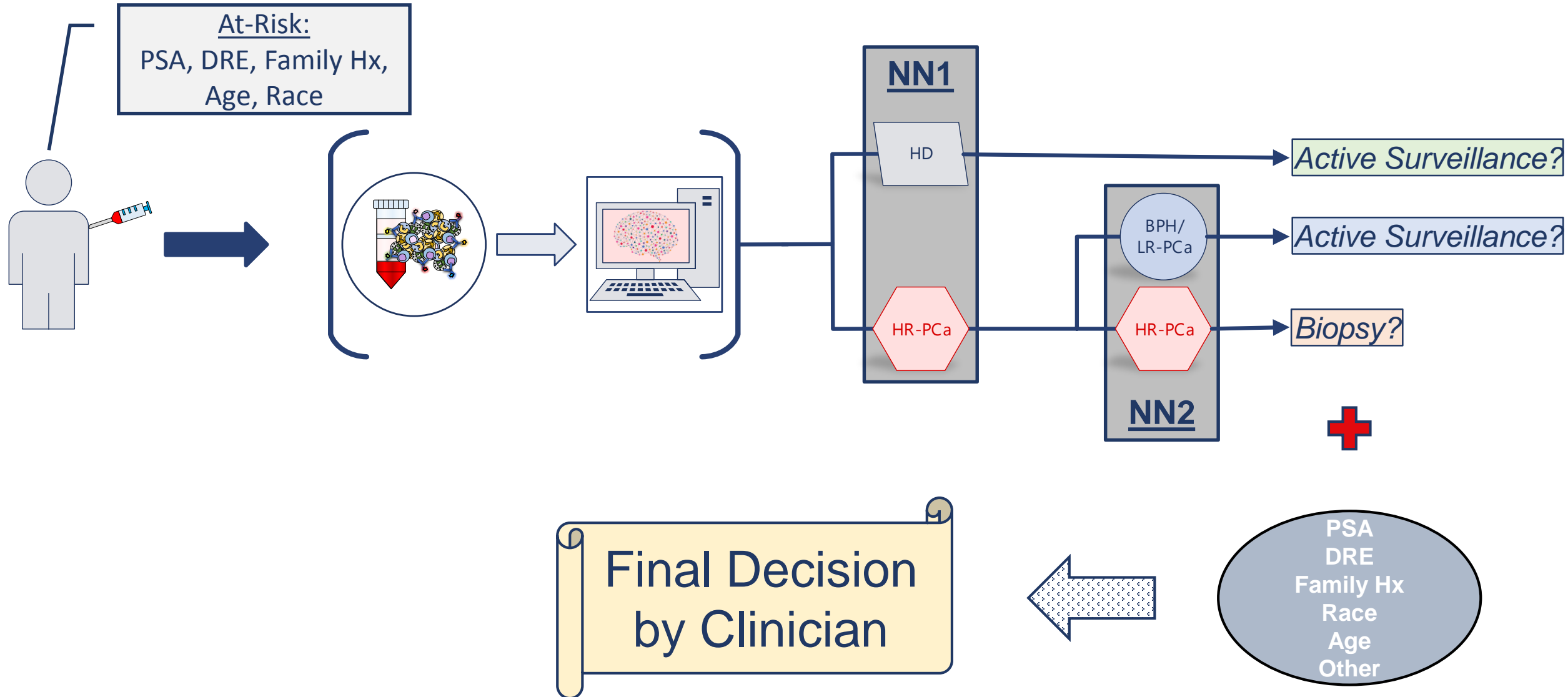
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- PSA is not reliable (large numbers of false positives)
- Majority of biopsies are negative
- 20% to 50% of men diagnosed through screening may be over diagnosed
- Gold Standard for Confirming → Prostate Biopsy (invasive/stressful)

## Risks of Screening and Overdiagnosis/Overtreatment

- 1% of prostate biopsies result in hospitalization
- 1 in 5 men who undergo prostatectomy may develop long-term urinary incontinence
- 2 in 3 men may experience long-term erectile dysfunction
- 1 in 6 men may experience long-term bothersome bowel symptoms

# Clinical Application: Confirmatory Testing for PCa Bx

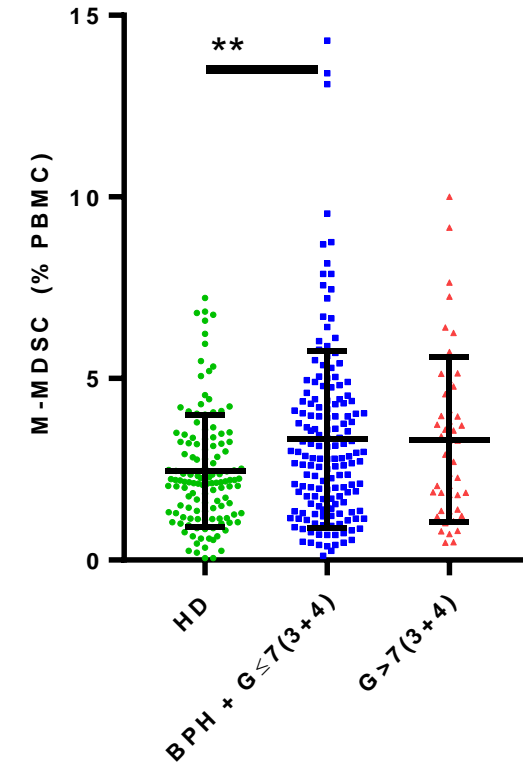
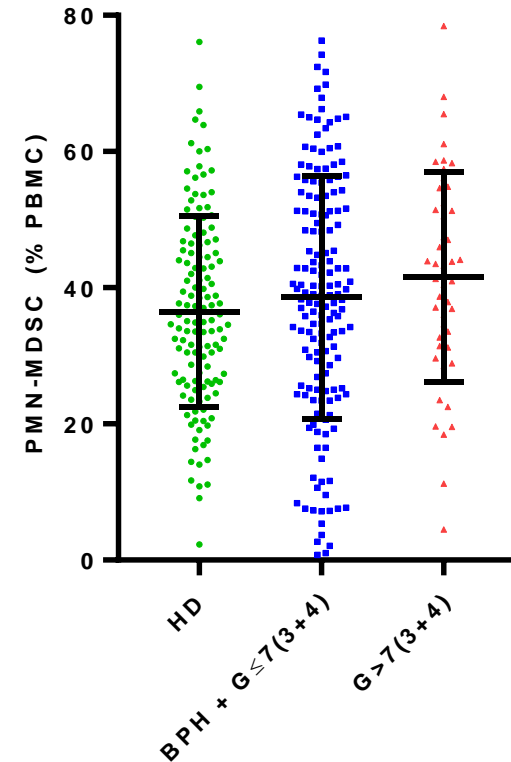
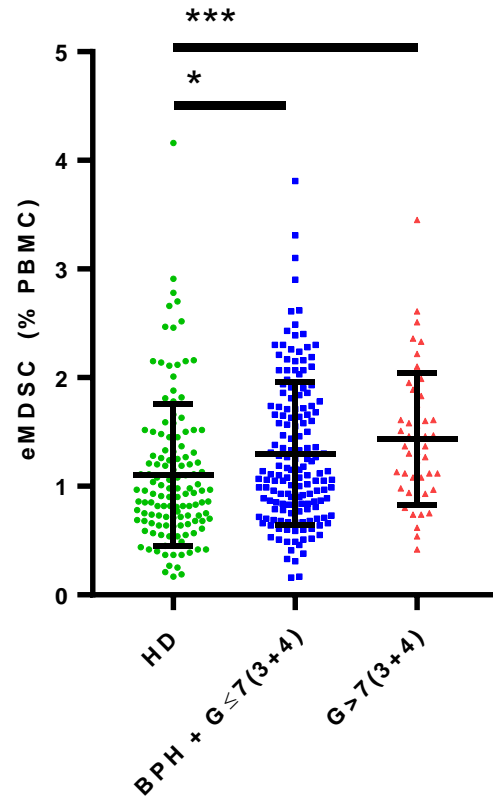


# Clinical Characteristics and Manual Counting

➤ Additional samples were collected

- + 41 PCa
- + 41 BPH
- + 43 Male HD

Characteristic	PCa	BPH	HD
Total	114	89	116
Median Age	67	62	52
Age Range	42 – 86	40 – 81	18 – 79
Gleason Score			
6	44		
7 (3+4)	26		
7 (4+3)	22		
>8	22		
Tumor Stage			
T1c	75		
T2a	5		
T2c	2		
Unknown	32		



***Still...simple cell counts can provide information about trends, but not really categorize subjects***

# Clinical Application: Confirmatory Testing

		Classified	
		Biopsy Recommended	Biopsy Not Recommended
Measured	<b>Gleason <math>\geq 7(4+3)</math></b>	9	1
	Needs Biopsy		
	<b>Gleason <math>\leq 7(3+4)</math> + BPH</b>	24	26
	Does Not Need Biopsy		
	<i>Sens. (%)</i>	90	
	<i>Spec. (%)</i>	52	
	<i>Prec. (%)</i>	27.27	
	<i>Acc. (%)</i>	58.33	

- Classified 26 BPH/LR-PCa samples as “Biopsy Not Recommended” → potentially reduce the number of unnecessary biopsies
- Mis-classified 1 out of the 10 HR-PCa samples → other factors may still suggest biopsy
  - subject had an abnormal DRE and a PSA > 20 ng/ml



# Conclusions

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- We demonstrated that machine learning can be used to analyze flow cytometry data of MDSC and lymphocytes
- We have applied this technique to distinguish between HD/PCa and BPH/PCa in a small number of samples
- We also demonstrated that this has the potential to reduce the number of unnecessary prostate biopsies (confirmatory testing)
  - PSA results have high false positive rate
  - Over 1 million prostate biopsies performed annually - overwhelmingly negative

# Future Work

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- Incorporate DRE results? PSA? Age? Race?
- Identify the critical relationships between cell populations that are used to make the classifications → unexpected relationships?
- Can this technique be applied to other flow cytometry data sets with different cancers? (retrospective analysis)
- Can this be used for predicting tumor recurrence, treatment and/or immunotherapy responses?
  - Collaborative projects

# Thank You!

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## **Anixa Biosciences**

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## **The Wistar Institute**

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Flow Cytometry Core Facility

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*MD Anderson Cancer Center at Cooper*

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# Questions?

Visit our poster (O2) tonight if you have more questions or interested in more details.