ARDS, RESPIRATORY FAILURE AND BLOOD BIOMARKERS

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Friday, January 18, 2019 – 1:55 p.m. – 2:40 p.m.

Angela Rogers, MD, MPH, received her medical degree from Harvard Medical School, and her Masters in public health from the Harvard School of Public Health, and pursued post-graduate training at the Brigham and Women's Hospital and Harvard Combined fellowship. She is an Assistant Professor in Pulmonary and Critical Care Medicine at Stanford University, where her research focuses on using genetics and genomics to identify novel biology in ARDS.

Precision medicine & the role for biomarkers in ARDS

Angela Rogers Stanford University California Thoracic Society January 18, 2019

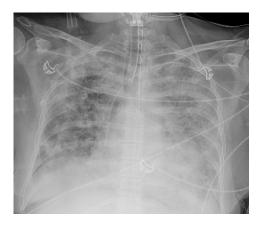
Conflicts of Interest

• I have no conflicts of interest

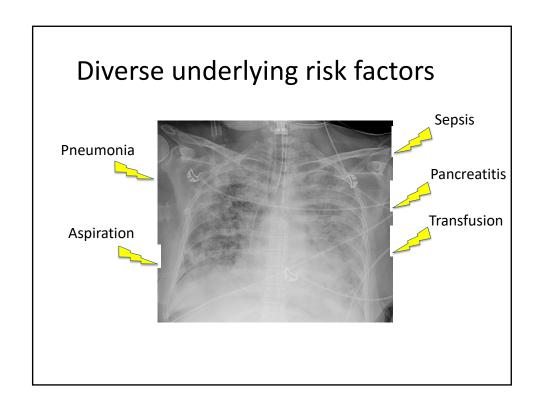
Learning objectives

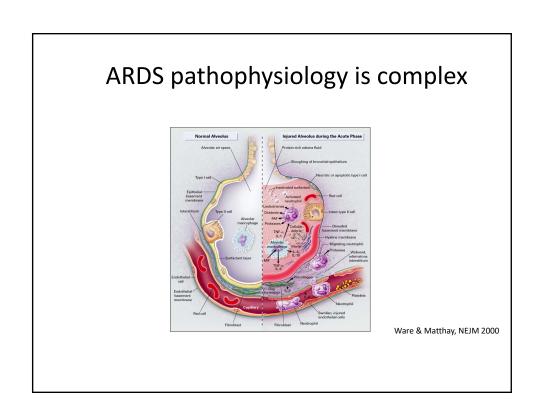
- To understand the need for biomarkers in ARDS
- PaO2:FIO2: A biomarker that works in ARDS
- Biomarkers for endotyping or "splitting" ARDS:
 - Latent class modeling of plasma
 - Molecular phenotyping of edema fluid

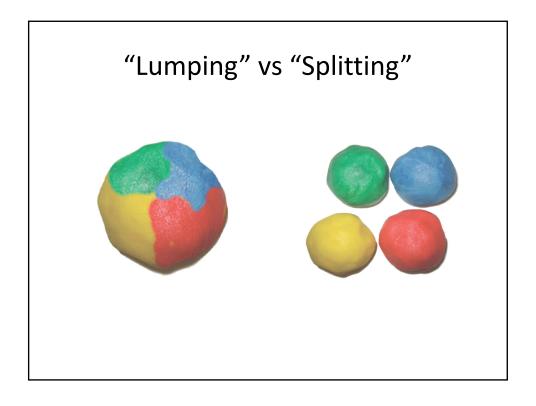
ARDS is defined very simply



- Intubated
- Acute
- •P:F ratio <300
- Bilateral opacities
- •Not explained by hydrostatic edema





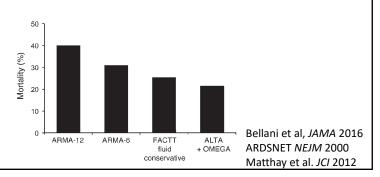


What have we learned from lumping?

- ARDS is common
 - 10% of all ICU & 23% of acute respiratory failure admissions



- In real world carries high mortality rate
- Major benefit of low tidal ventilation

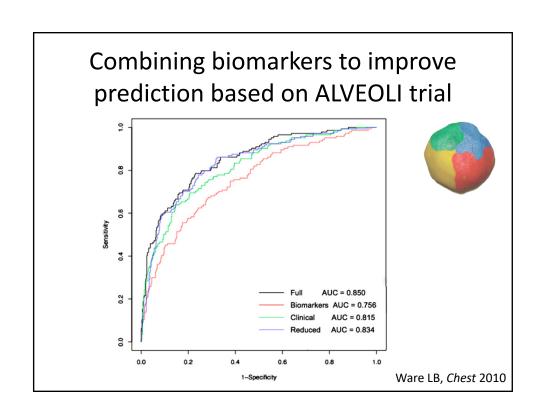


Biomarkers in all of ARDS

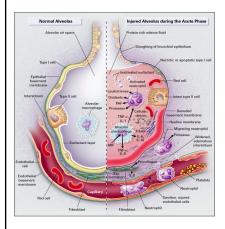


		90-day mortality		у
Pathway	Biomarker	Alive	Die	p- value
Inflammation	IL-6	209	322	0.004
	IL-8	35	64	<0.001
	TNFR	3668	6914	<0.001
Coagulation &	Protein C	82	68	.011
fibrinolysis	PAI-1	54	111	<0.001
Endothelial injury	ICAM	854	1072	<0.001
	VWF	370	477	<0.001
Epithelial injury	SP-D	92	124	.01

Ware LB, Chest 2010



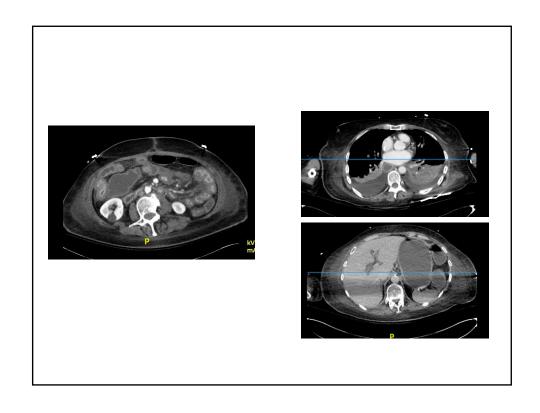
Is it possible that lumping all of ARDS together is harming ARDS clinical trials & science?

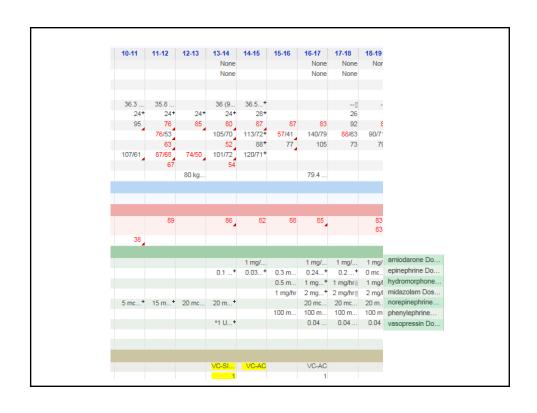


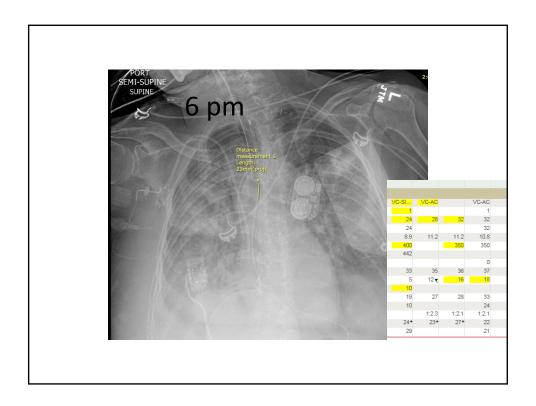


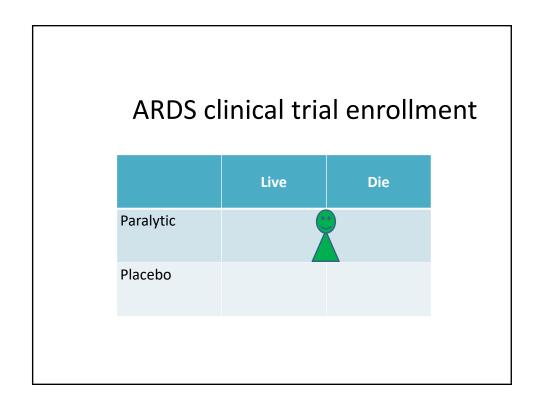
Why splitting matters: a case to classify

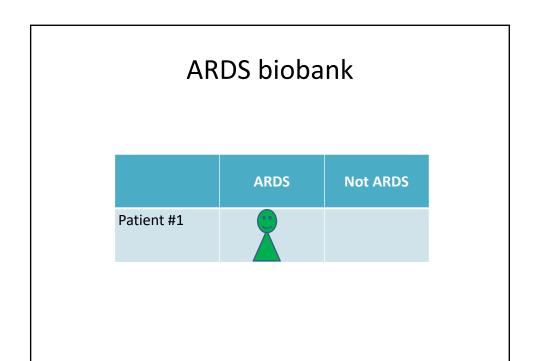
- 70 yo F with colon CA on chemo, recently discharged after 1 week admission for failure to thrive
- Per husband, was nauseated, "gurgling" all night
- Returns to ED critically ill

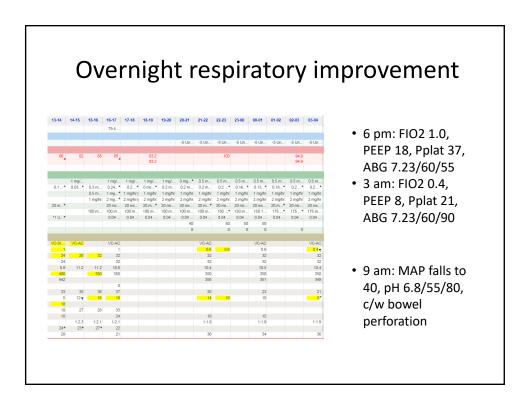






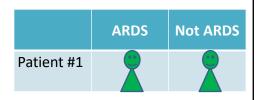






What does this case do to our clinical trial and biobank?

	Live	Die
Paralytic		
Placebo		



Misclassification in ARDS really matters for clinical trials

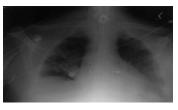
 Inter-rater CXR interpretation varies from κ ~.4-.9



	ARDS	Not ARDS
ARDS	50	
Not		50



	ARDS	Not ARDS
ARDS	40	10
Not	10	40



K = .4

		ARDS	Not ARDS
4	ARDS	40	20
	Not	10	30

Rubenfeld et al. Chest 1999

Power for clinical trials dramatically falls with misclassification

RCT power estimate when ARDS enrollment is imperfect from a patient cohort with 25% ARDS prevalence

Inter-observer		Power in 1500	Sample size for
Agreement	Kappa	patient trial	90% power
Perfect	1.00	0.92	1402
Almost perfect	0.85	0.87	1664
Substantial	0.72	0.81	1968
	0.61	0.74	2320
Moderate	0.51	0.67	2726
	0.42	0.60	3198

Sjoder et al. Annals ATS 2016



A major role for biomarkers may be in "Splitting" ARDS

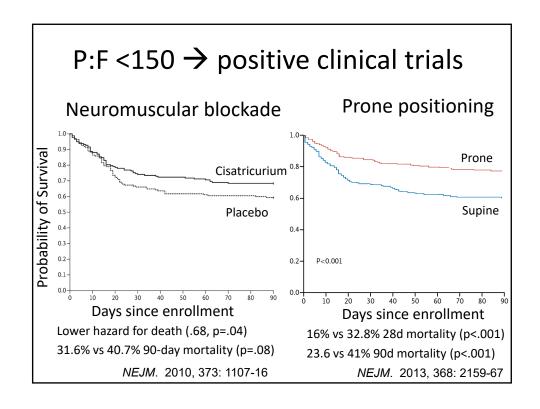
- Prognostic: Identify patients at highest risk of bad outcomes and death
- Predictive: Identify patients who would benefit most from treatment

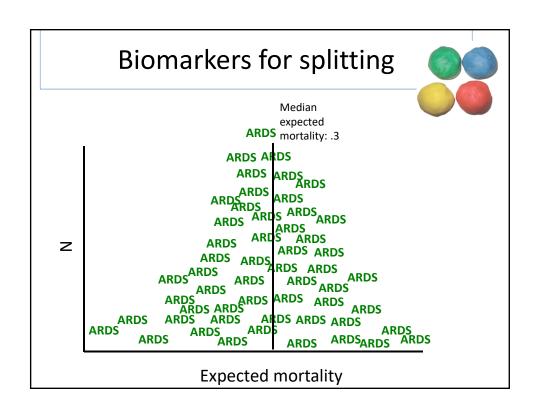


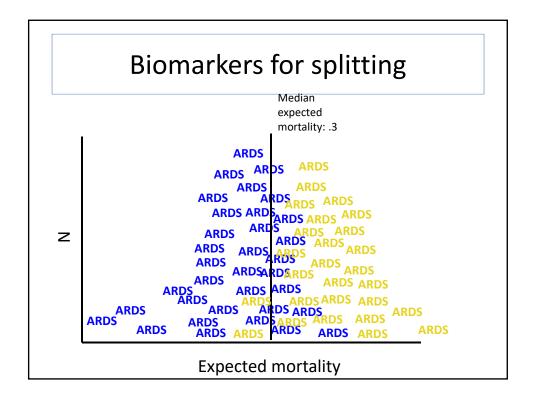
PaO2:FIO2 ratio as a critical ARDS biomarker

- PaO2:FIO2
 - P:F ratio defines disease severity
 - Prognostic, outperforms other, more complex models
 - Enriches clinical trials: recruiting based on more stringent thresholds
 - · Predictive enrichment

AECC consensus conf, AJRCCM 1994 Berlin definition, JAMA 2014







2 examples

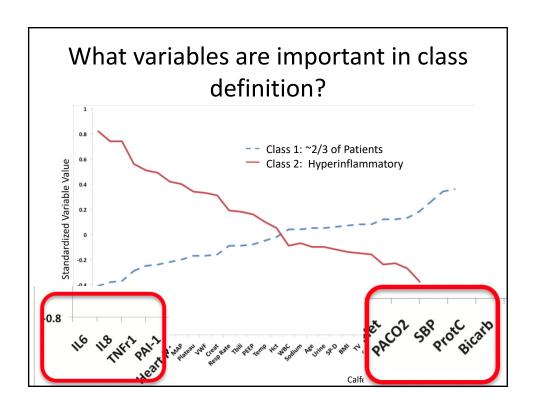


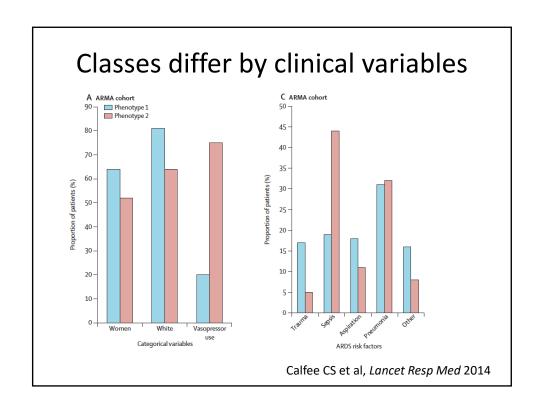
- Latent class modeling, identified plasma biomarkers
- Metabolomics of pulmonary edema fluid

Latent Class Analysis: Are There Distinct Subtypes of ARDS?

- Study population: Three ARDSnet clinical trials
 - First cohort: ARMA (low tidal volume only; n=479)
 - Second cohort: ALVEOLI (low vs. high PEEP; n=549)
 - Third cohort: FACCT (conservative vs liberal fluid; n=1000)
- Clinical and biomarker data from baseline in each study as inputs that "identify" class (endotype)
 - Analysis conducted independently in each cohort
 - Outcomes not considered in class modeling

Calfee CS et al, Lancet Resp Med 2014 Famous K et al, AJRCCM 2016





Mortality differs by class

	90-day mortality		
Study	Class 1 (~2/3) ARDS	Class 2 (~1/3) ARDS	p-value
ARMA	23%	44%	0.006
ALVEOLI	19%	51%	<0.001
FACTT	22%	45%	<0.0001

Class could be defined w/ >90% AUC with 3 factors: IL8, TNFr1, bicarbonate

Calfee CS et al, *Lancet Resp Med* 2014 Famous K et al, *AJRCCM* 2016

Response to Therapy differs by class

ALVEOLI (p_{interaction}=.049)

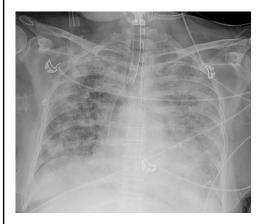
	Mortality in Class 1 ARDS (n=404)	Mortality in Class 2 ARDS (n=145)
Low PEEP	16%	51%
High PEEP	24%	40%

FACCT (p_{interaction}=.004)

	Mortality in Class 1 ARDS (n=727)	Mortality in Class 2 ARDS (n=273)
Liberal fluid	18%	50%
Conservative fluid	26%	40%

Calfee CS et al, Lancet Resp Med 2014 Famous K et al, AJRCCM 2016

Metabolomics of pulmonary edema fluid: ARDS vs CHF





Pulmonary edema fluid metabolomics

- Undiluted pulmonary edema fluid in ARDS
 - High edema: plasma protein ratio (>.65) associated with ARDS (AUC >.8)
- Pulmonary edema fluid at time of intubation
 - 16 ARDS vs 13 CHF
 - Collected at Vanderbilt and UCSF

Rogers et al. AJP Lung 2017

CHF vs ARDS Phenotyping

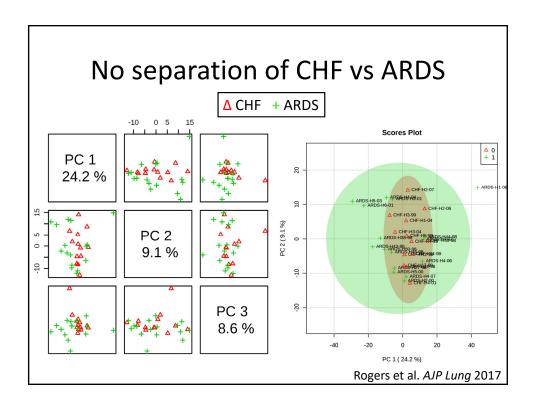
	ARDS (N=16)	CHF (N=13)	P value
Age	43.7	50.5	.3
Gender (%M)	50%	62%	.7
Sepsis	44%	0%	.008
Mortality	44%	15%	.12
Primary Diagnosis	Pneumonia (4) Sepsis (4) Anaphylaxis (2) Aspiration (1) TRALI (2) Fulm Hep Fail (1) Reperfusion edema (1) Tumor lysis (1)	Vol overload/CHF (5) MI/Ischemia (2) Cardiac arrest (1) Post-obstructive (2) Cardiogenic shock (1) TRALI (1) Neurogenic (1)	.01

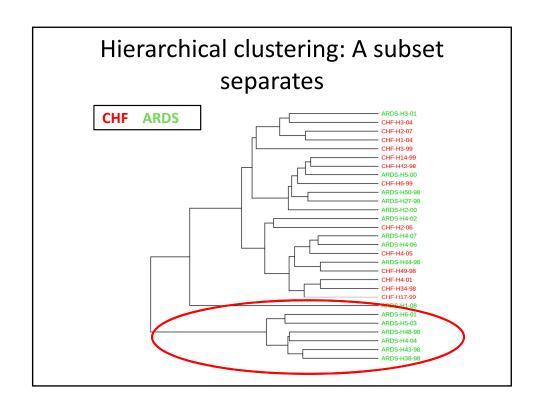




Metabolic profiling strategy

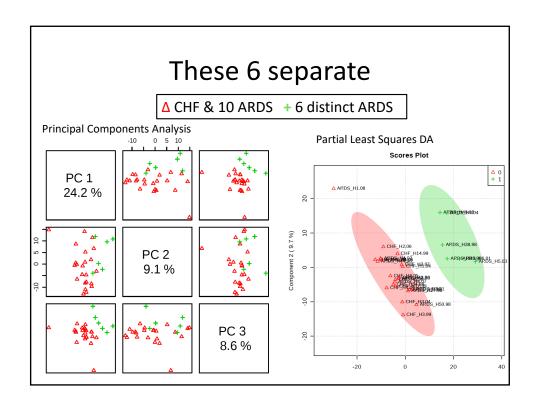
- Undiluted pulmonary edema fluid profiled by Metabolon
- Tests up to 3000 human plasma metabolites with high accuracy
- Metabolite levels log₂ normalized and auto scaled
- Differences in classes assessed using machine learning
 - Principle components analysis
 - Partial least squares-discriminant analysis
 - Hierarchical clustering

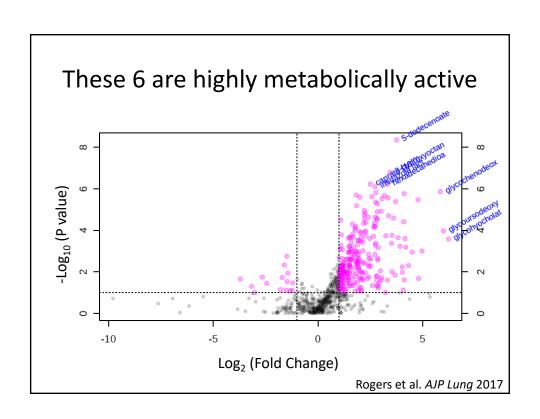


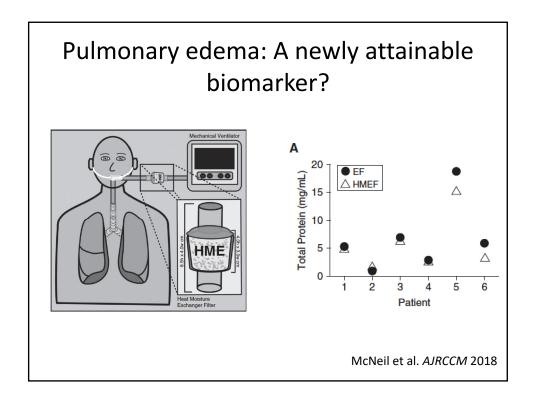


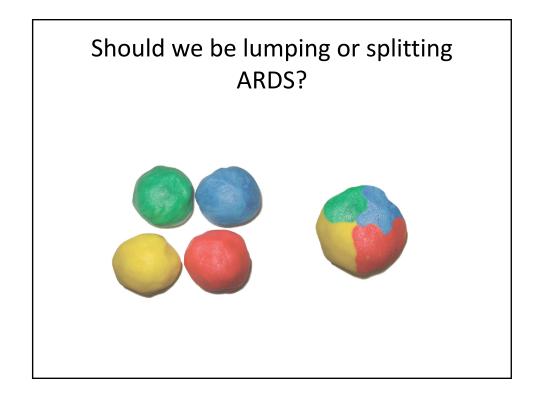
6 Separate ARDS

	6 Distinct ARDS	10 Remaining ARDS	CHF
Age	36	49	51
Gender (%M)	33	60%	62%
Sepsis	66%	30%	0%
Mortality	66%	30%	15%
Primary Diagnosis	Sepsis (3) Fulm Hep Fail (1) Anaphylaxis (1) Aspiration (1)	Pneumonia (4) Sepsis (1) Anaphylaxis (1) TRALI (2) Reperfusion edema (1) Tumor lysis (1)	Vol overload/CHF (5) MI/Ischemia (2) Cardiac arrest (1) Post-obstructive (2) Cardiogenic shock (1) TRALI (1) Neurogenic (1)









Should we be lumping AND splitting ARDS?





Should we be lumping AND splitting ARDS?

Low tidal volume, lung protective ventilation

- Clearly helps mortality in ARDS
- Little downside in some misclassification



Should we be lumping AND splitting ARDS?



ARDS clinical trials

- Genomics tells us we need to be careful with lumping
 - Endotypes
 - Frank misclassification
- Kills power of trial & puts patients who can't benefit at risk

Conclusions

- Lumping all of ARDS as a single phenotype has been very successful for lung protective ventilation and reduced mortality
- For moving toward precision medicine:
 - To date our only established biomarker in ARDS is the P:F ratio, which is prognostic and predictive
 - Biomarkers will likely be critical in endotyping ARDS & moving toward personalized medicine in practice and clinical trials

