# ATRC @ UCSF OVERVIEW



\*Antibody technology research center

# Goals of the ATRC

Robust approaches for rAbs to the proteome

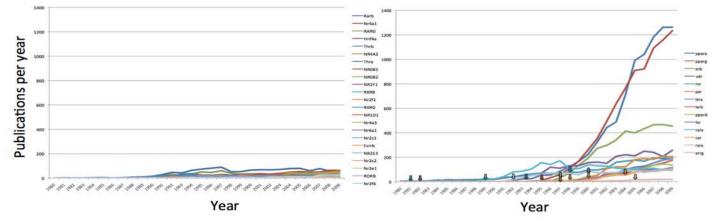
- Initially focus (largely) on secreted proteins
  - Single and multipass membrane proteins/proteases
  - PTMs (phospho-specific rAbs)

Technologies

- Phage Ab
- Automation and high throughput screening
- In vitro antigen and Ab expression
- Automatable cell selection

# **Existing antibodies**

#### Functional Abs not available for >90% of proteome



### Where available

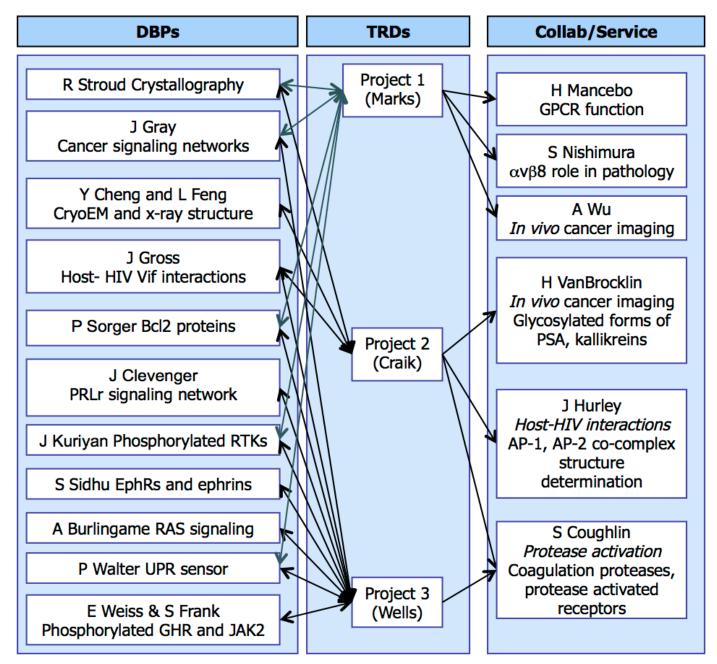
- Half or more are not specific
- Lot-to-lot and vendor-to-vendor variability
- Relatively expensive
- Not renewable (not cloned)
- Typically IgG only



# **ATRC Solution**

Phage Ab based rAb generation

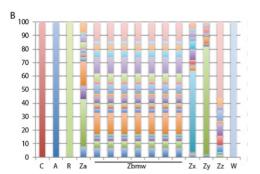
- rAbs defined by their sequence
- Forever renewable
- Can be made as fragment or with any Fc (species/isotype)
- A technology-driven extension and expansion of what we are doing at a cottage industry level
  - Generating robust phage Ab libraries
  - Automation of rAb generation
  - Generating rAbs to challenging antigens
  - Providing technology training

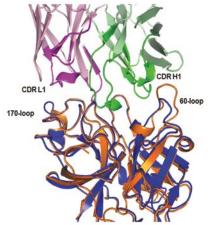


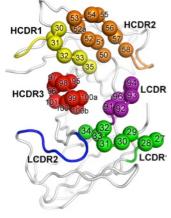
ATRC is funded by a P41 mechanism, that requires TRD, DBP, CSP

# TR&Ds

- TR&D 1 (Marks):
  - Next generation phage Ab libraries
  - Robust rAb generation to secreted and single and multipass membrane proteins
- TR&D 2 (Craik):
  - Generate secreted and type 2 TM serine proteases, receptors and intramembrane proteases
  - rAbs to protease targets including inhibitory rAbs
- TR&D 3 (Wells):
  - Phage libraries/technologies to generate
    rAbs to phosphorylated antigens,
    neo- and conformational-epitopes
  - Automation of selections

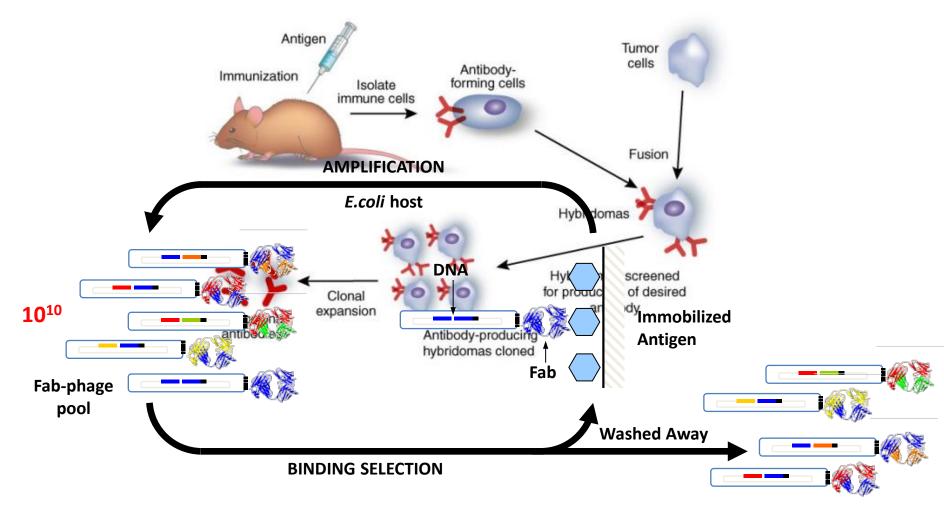






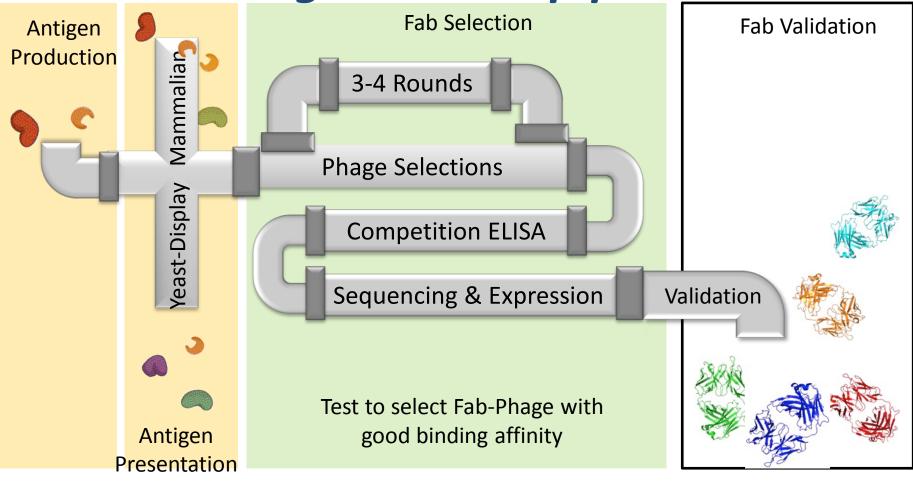
## <u>Technologies: in vivo vs in vitro Abs</u>

Adapted from www.medscape.com



Non-binding phage

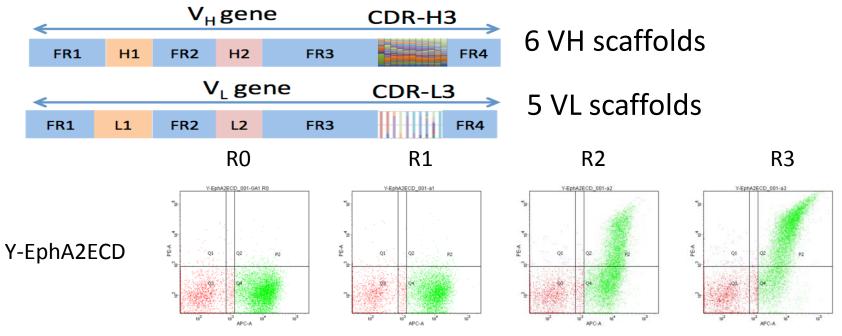
# Ab generation pipeline



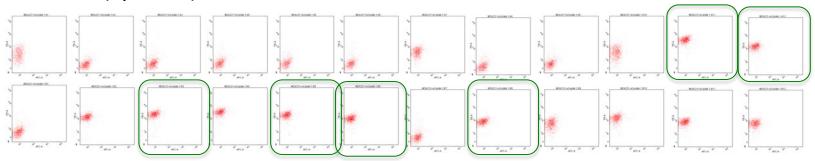
Antigen QC: SDS-PAGE; SEC

rAb QC: Sequence, affinity, specificity

## TRD 1: Nature-inspired synthetic antibody libraries

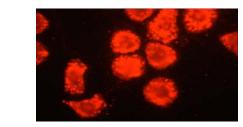


> 30 unique antibodies MDA-MB-231 (EphA2 +) cells



## Antibody fragments for nanoliposome targeting

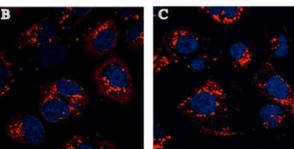
Anti-HER2 liposomal doxorubicin MM-302 Registration Phase 2 HER2+, progress on T DM-1 Ar

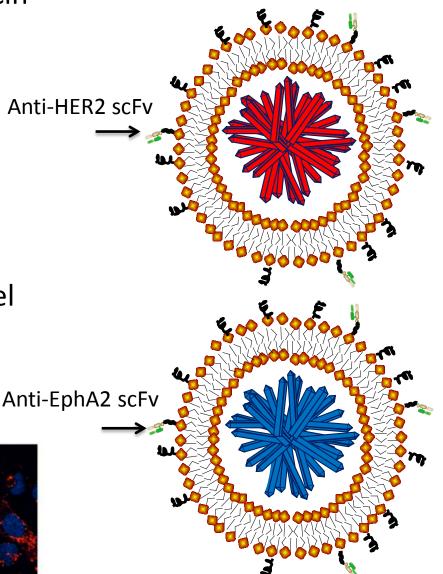


Anti-EphA2 liposomal docetaxel MM-310 Completed manufacturing Phase 1 Q2 2016

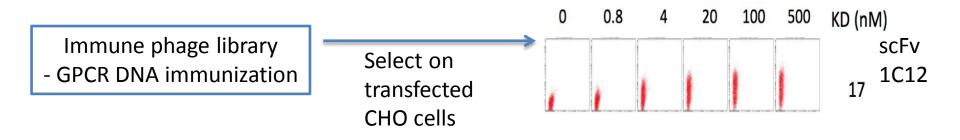
scFv D2-1A7

scFv F5

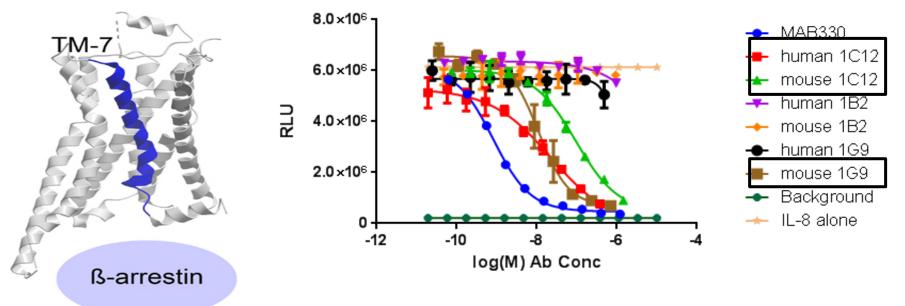




## rAbs to CXCR1 with functional activity

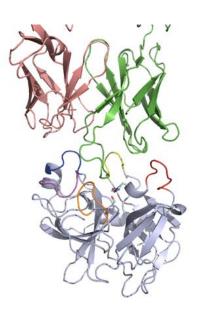


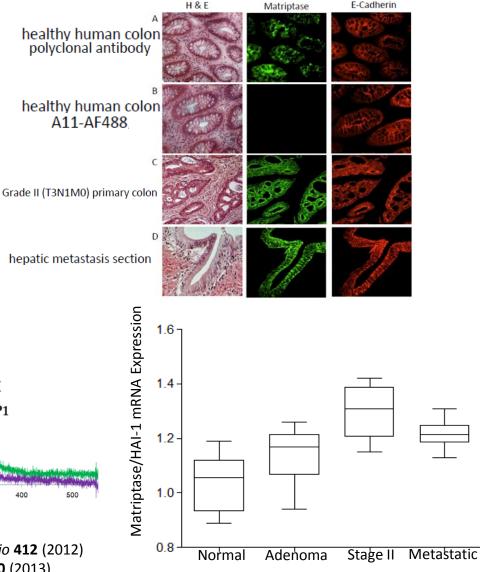
B-arrestin assay for mAb biologic activity: inhibit IL-6 CXCR1 activation



#### Marks Lab CXCR1 Antibodies

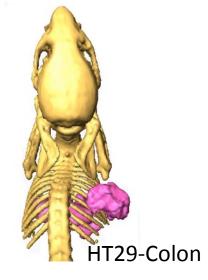
### Ab Activity Based Probe to Matriptase Provides **Diagnostic Information in Colon Cancer**

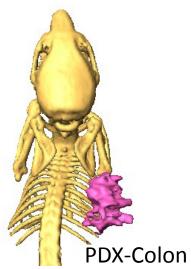


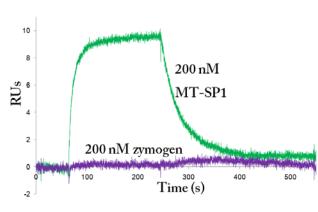


E-Cadherin





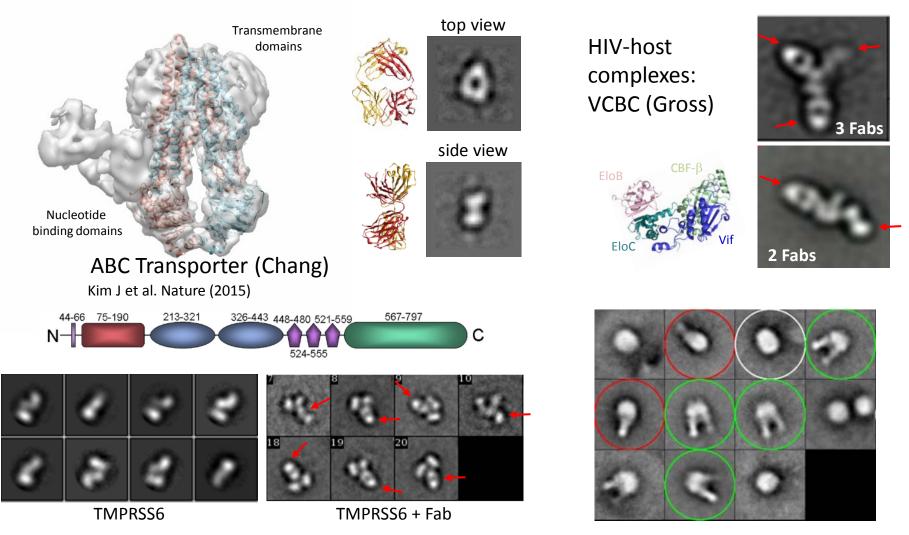




Schneider E, et al. J. Mol Bio 412 (2012) LeBeau AM, et al. PNAS 110 (2013)

### Conformationally Selective Fabs to Challenging Targets





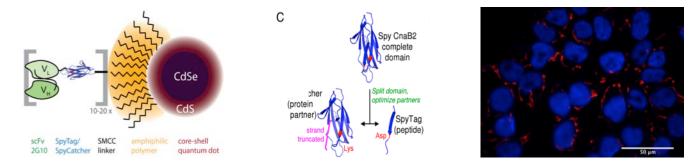
TMPRSS6 membrane anchored serine protease (Bayer)

Dimeric Chloride Channel (Jan)

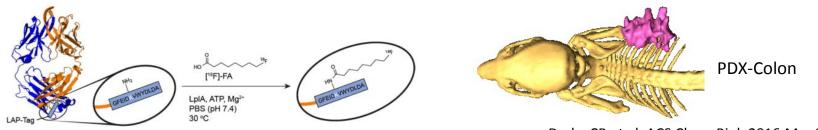
### **Engineered Antibody Projects**



#### Antibody-Qdot conjugates using the SpyTag/SpyCatcher system

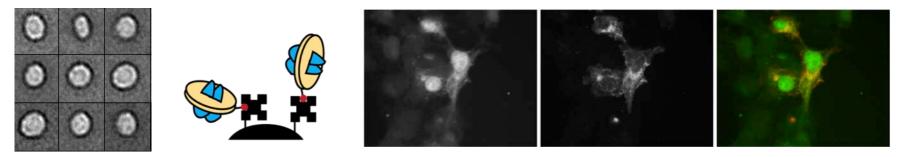


Labeling using LpIA acceptor peptide (LAP) peptide or aldehyde peptide (CxPxR)



Drake CR et al. ACS Chem Biol. 2016 Mar 31.

Nanobodies selected against VGLUT2 reconstituted in nanodiscs



# **Many Thanks**

Natalia Sevillano Kathrin Zuberbuhler Melody Lee

**Jim Marks** 



Jim Wells







Cheng Lab



#### Gross Lab

**Robert Stroud** 













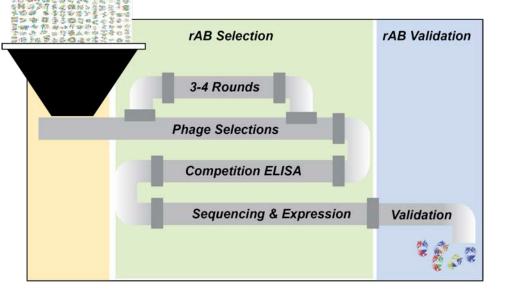


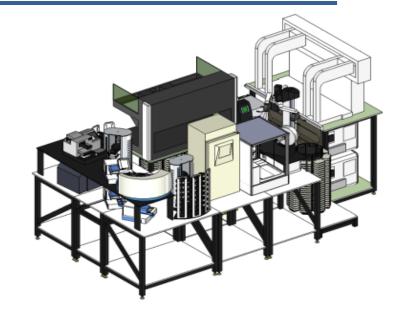




- Industrialized Selections
- Mutant and PTM specific selections
- How is the Surfaceome changed by RAS transformation?
- Multiplexed antibody detection for proteomics
  - BaNGS (Bar-coded antibodies NGS)
  - PhaNGS (Phage-antibodies NGS)

# Industrialized Antibody Pipeline





~3000 Fab 346 TFs 211 Epigenetic Factors Ave affinity: 10-20nM Expression: 1-10 mg/mL







THE UNIVERSITY OF CHICAGO

#### Hornsby et al 2015



## Selection for high resolution antibodies

### Conformations: On/Off state

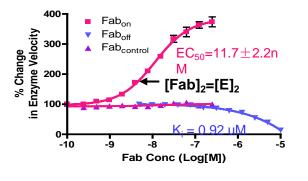
	<b>Fab</b> on			<b>Fab</b> off		
	"on" form	"off" form	аро	"on" form	"off" form	аро
K <sub>D</sub> (10⁻ ⁰M)	2.5	>1000	330	99	4.7	17
<i>k<sub>on</sub></i> (10 <sup>4</sup> M <sup>-1</sup> s <sup>-1</sup> )	66	N.D.	0.8	1.6	135	55
<i>k<sub>off</sub></i> (10 <sup>-</sup> <sup>3</sup> s <sup>-1</sup> )	1.7	N.D.	2.6	1.6	6.4	9.5

### PTMs: Phosphorylation

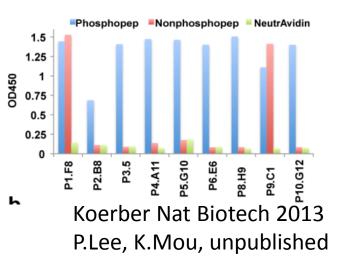
## Peptide Phosphospecific "Reader "Anchor" Domain"

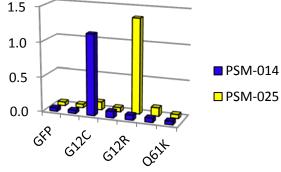
### Mutations: KRAS

Gly12	Gly13	Gln61	
Α	А	E	
С	С	Н	
D	D	K	
R	R	L	
V	V	Р	
S	S	R	



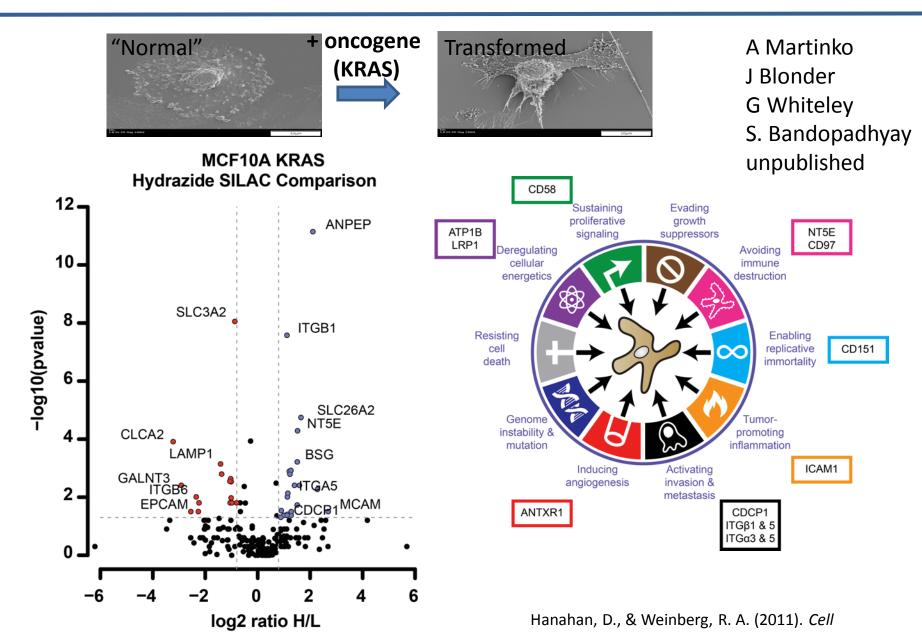
Gao PNAS 2009 Thomson PNAS 2013



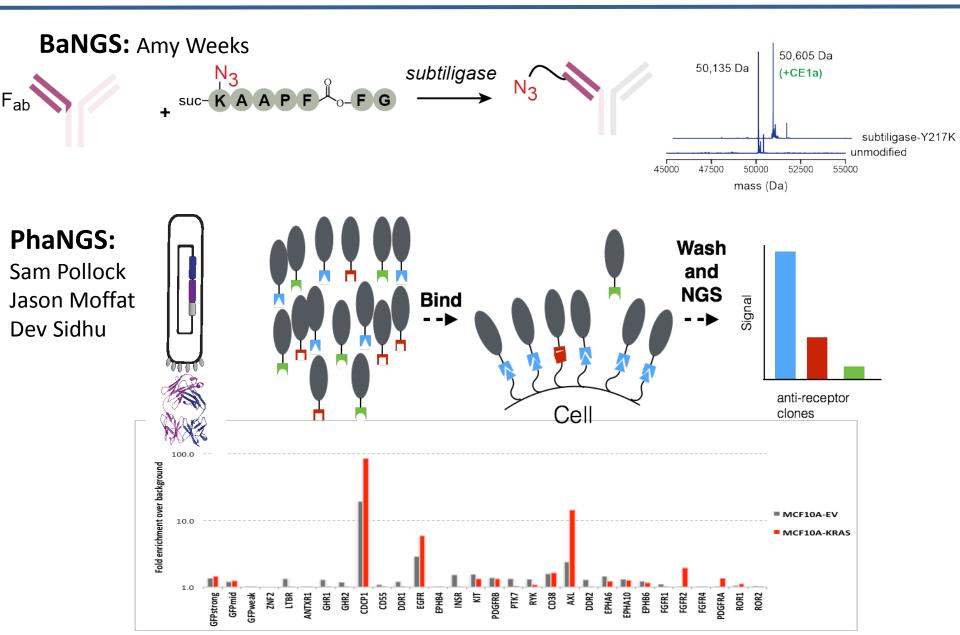


P. Marinec, Unpublished,

## How oncogenes remodel cell surfaces



## **DNA-barcoding for multiplex detection**





# How can FNLCR help?

- Protein Antigens (100/yr)
  - Fc fusions
  - Membrane proteins and complexes
- IgG conversion, expression, distribution (100/yr)
- Access to high priority targets and collaborations