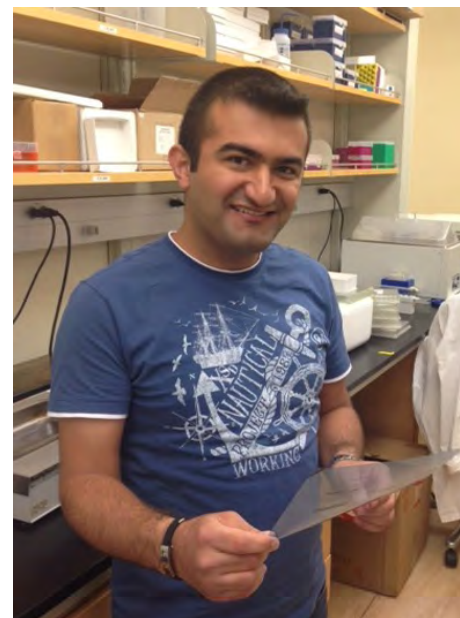


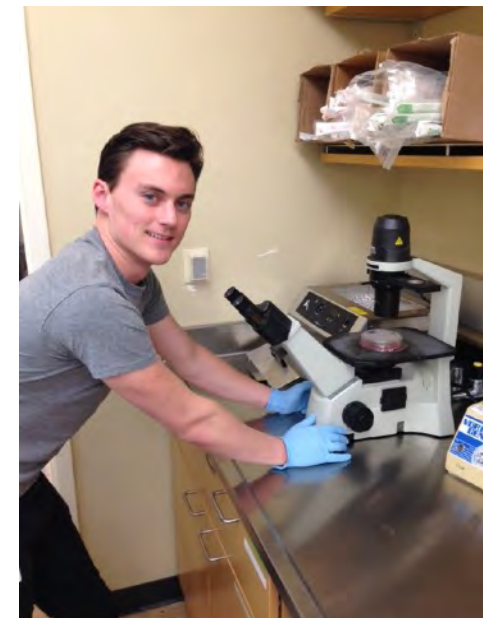
# MYC drives a unique molecular and therapeutically-relevant subset of SCLC

Trudy G. Oliver, PhD  
SCLC Consortium  
March 15, 2018

Gurkan  
Mollaoglu

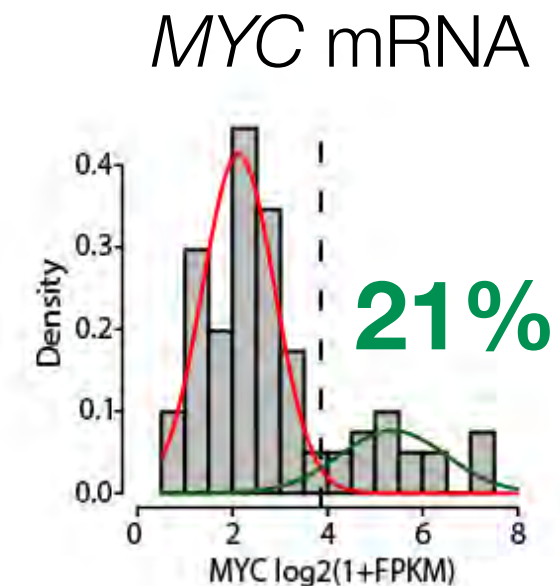
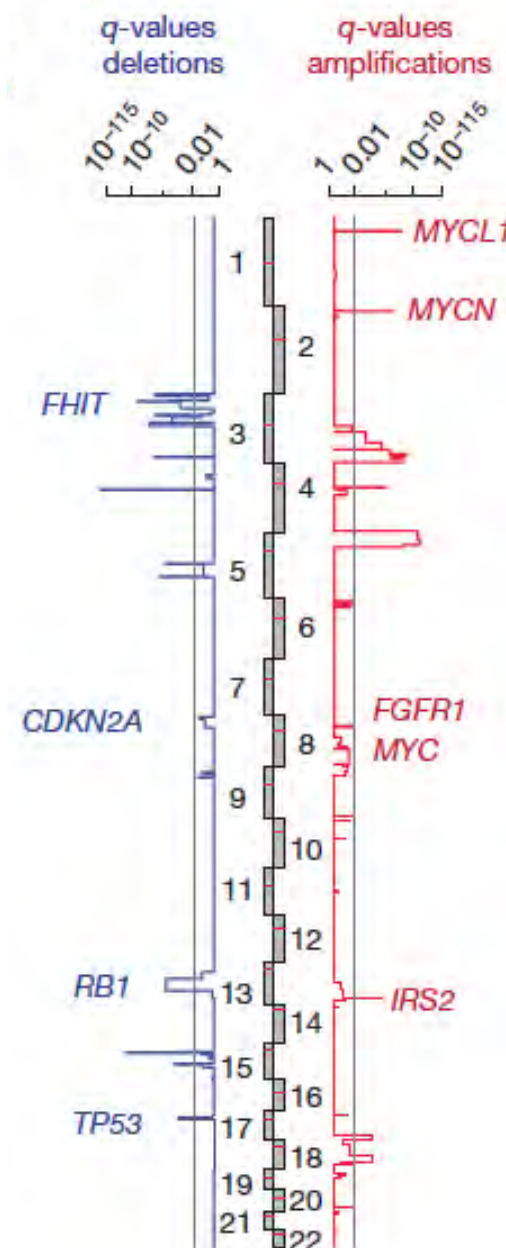
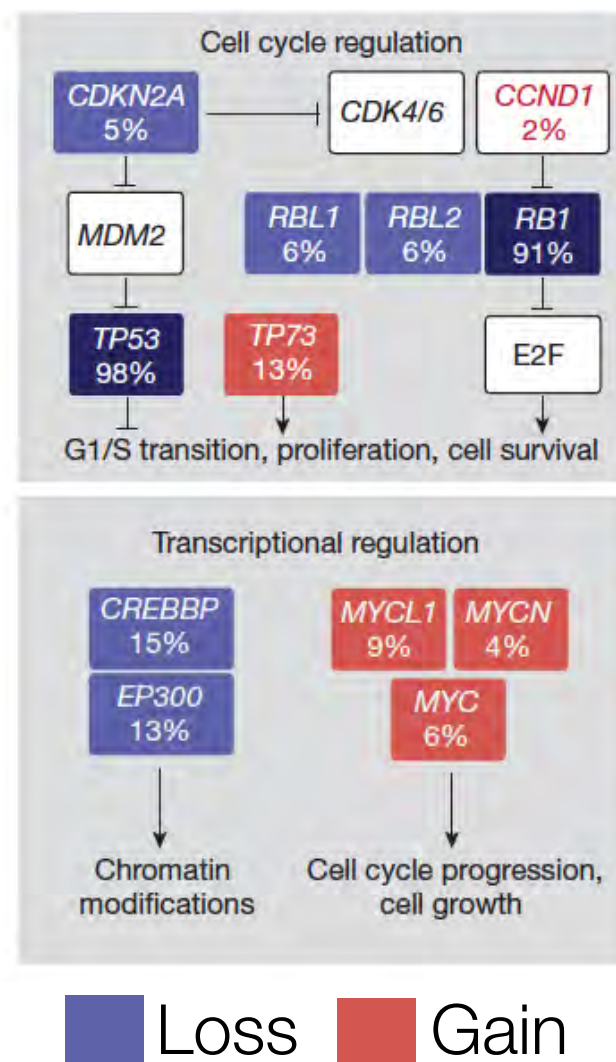


Matt  
Guthrie



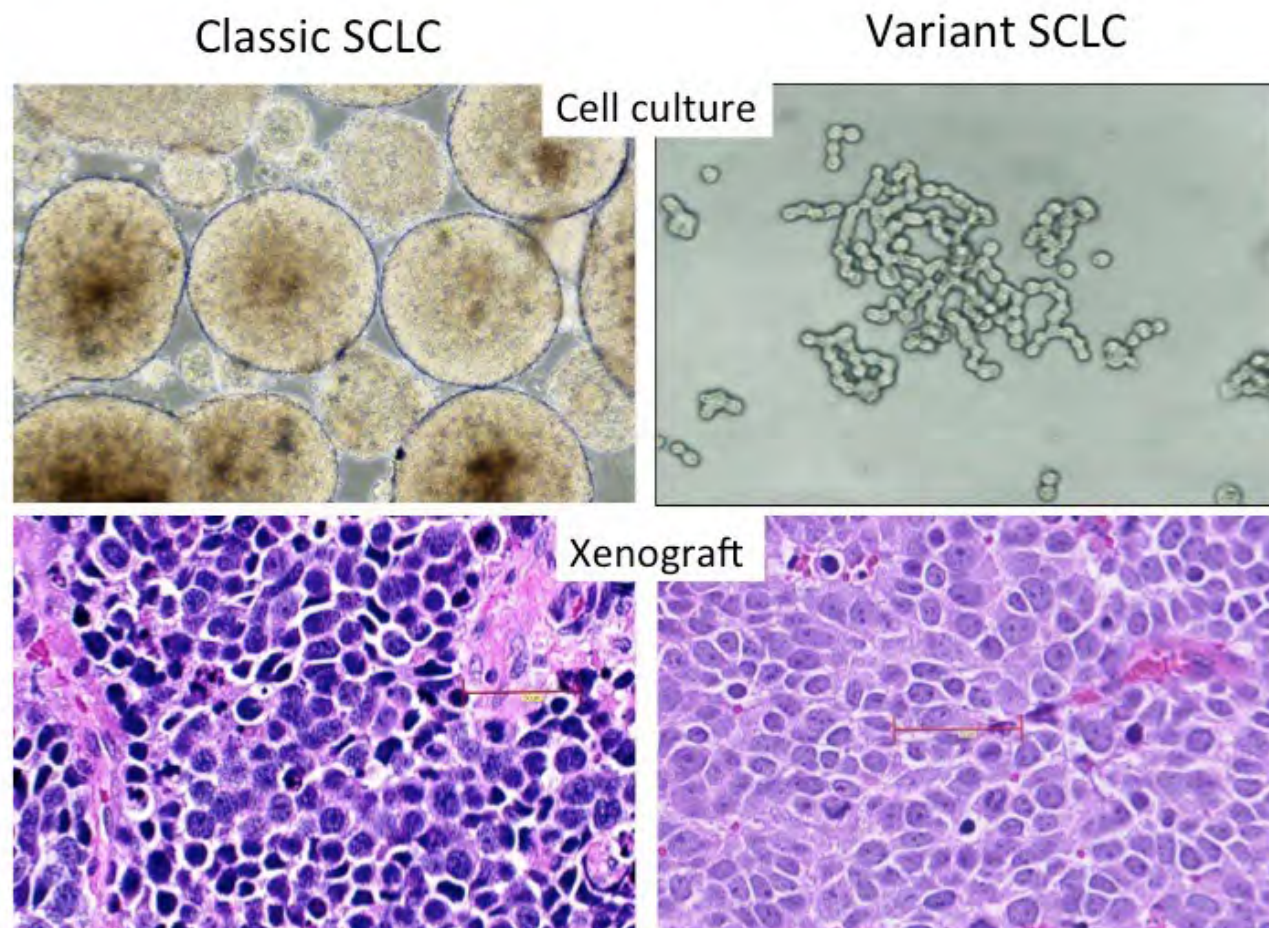
# Genomics of human SCLC

*Loss of RB1 and P53, plus mutually exclusive gain of MYC family members*



# In the 1980's.....classic and variant SCLC

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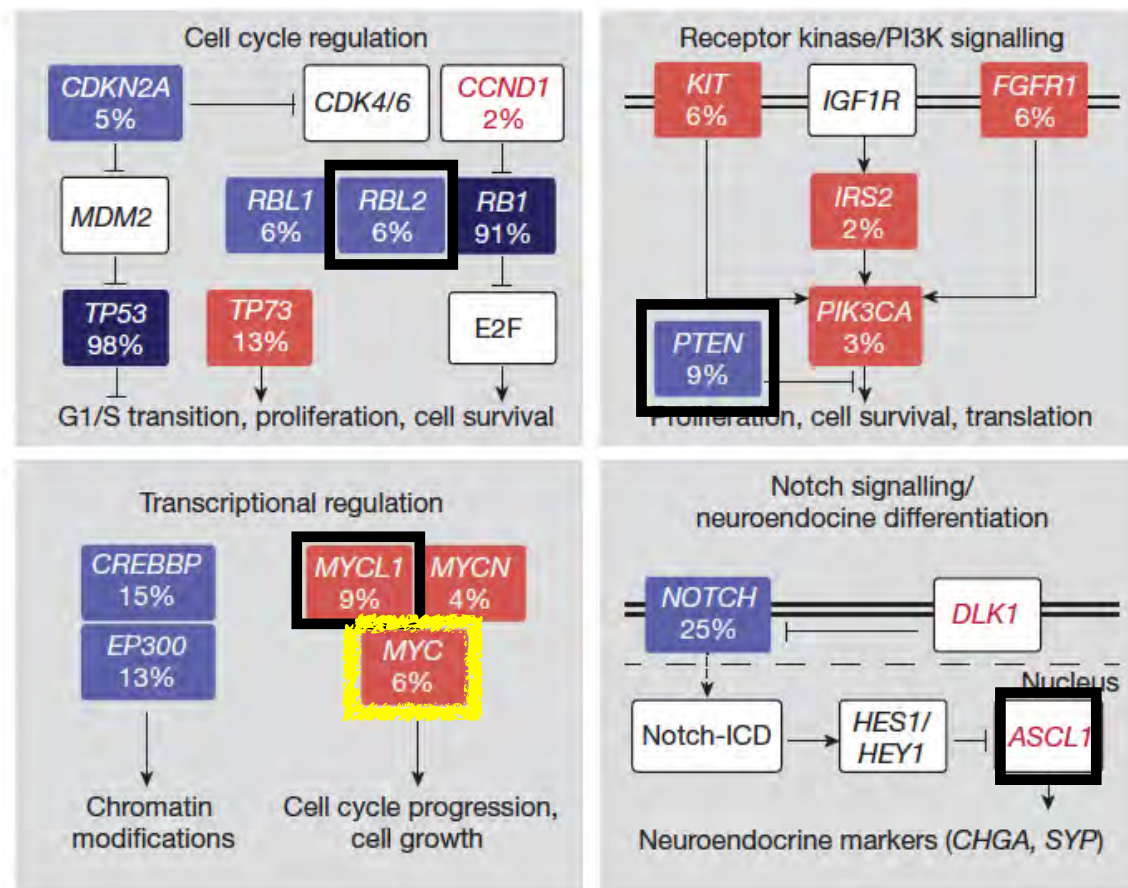
*Provided by Drs. Adi Gazdar and John Minna*

## Variant SCLC

- More commonly derived from relapsed patients / at autopsy
- Associated with shorter survival time
- Reduced expression of neuroendocrine markers
- Faster proliferation
- More radio-resistant
- Associated with *c-MYC* amplifications



# Mouse models of SCLC based on *Rb1/p53* loss

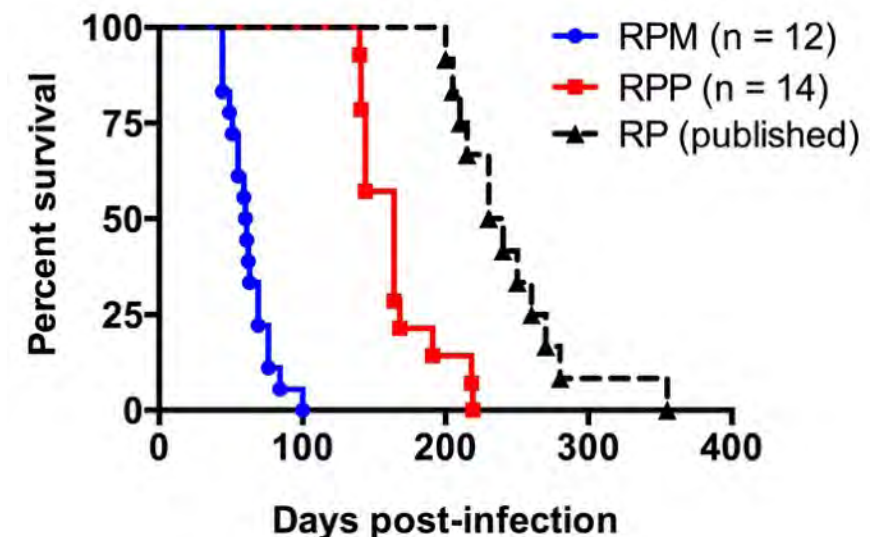


George et al Nature, 2015

■ Loss ■ Gain



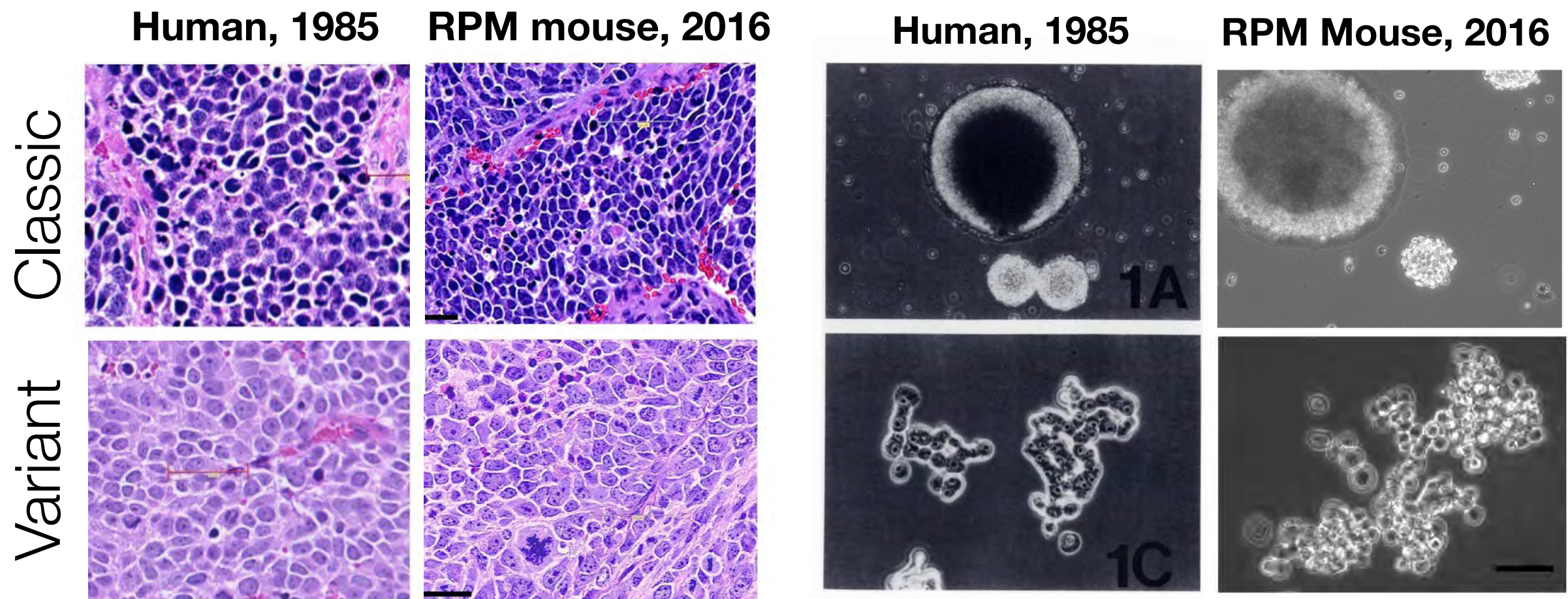
Rb1/p53 (RP, Berns)  
 Rb1/p53/Rbl2 (RPR2, Sage)  
 Rb1/p53/Pten (RPP, Jacks, MacPherson)  
 Rb1/p53/L-myc (Berns)  
 SV40 + Ascl1 (Linnoila)  
 Rb1/p53/Myc<sup>T58A</sup> (RPM, Oliver)



**Classic SCLC, LCNEC or other NSCLC phenotypes  
 often with high *L-Myc* expression and latencies of ~5-12 months  
 (Gazdar et al, JTO, 2015)**



# C-MYC-driven tumors and cell lines resemble variant SCLC



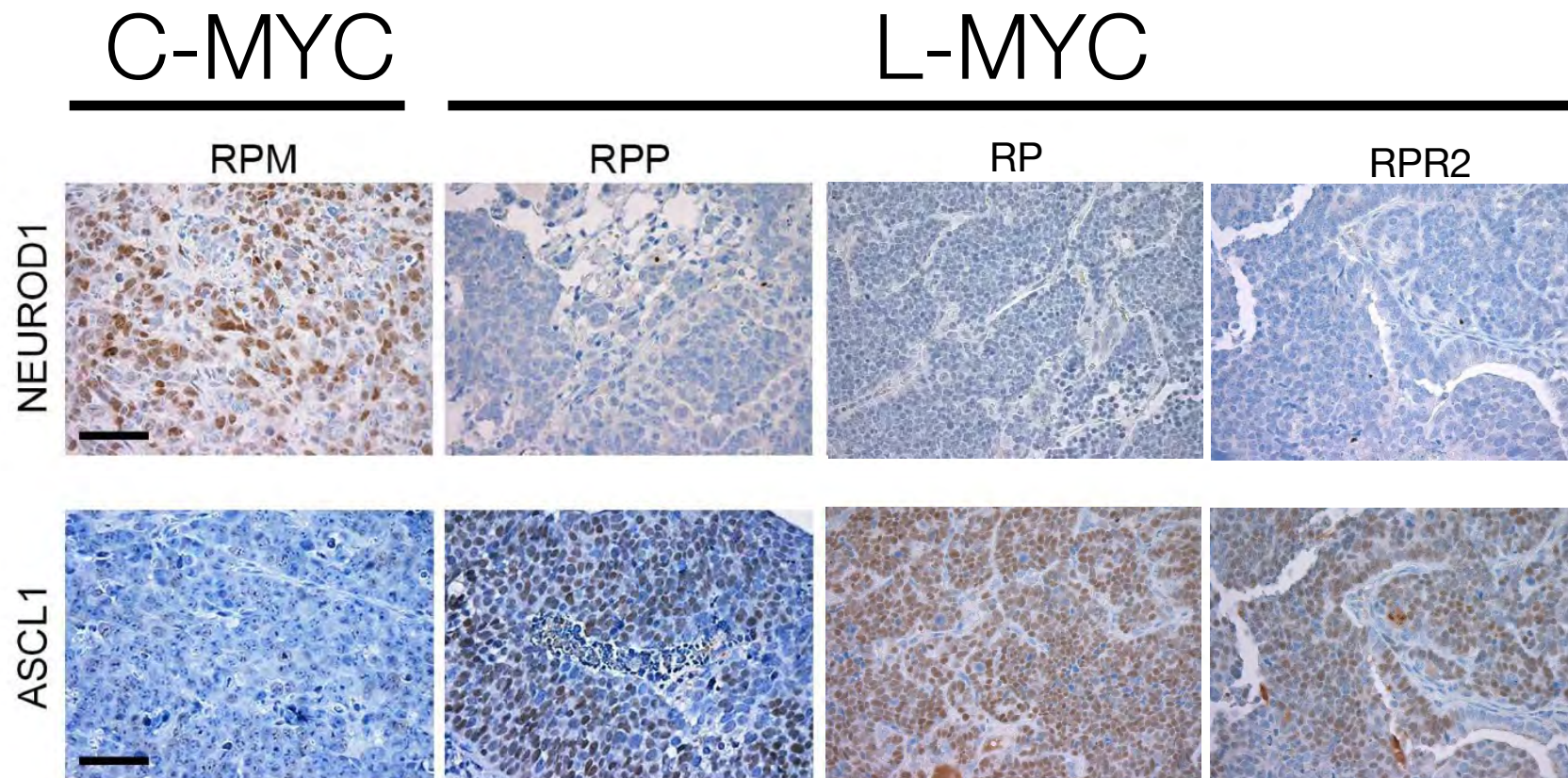
Are they neuroendocrine?



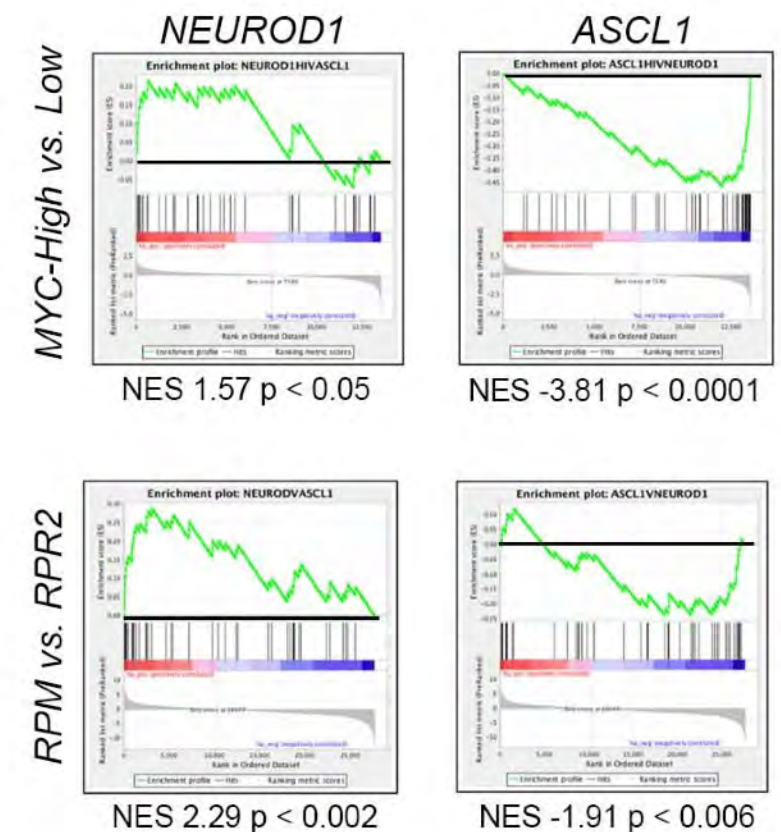




# MYC-driven mouse and human tumors express NEUROD1 and its target genes



## 81 human tumors



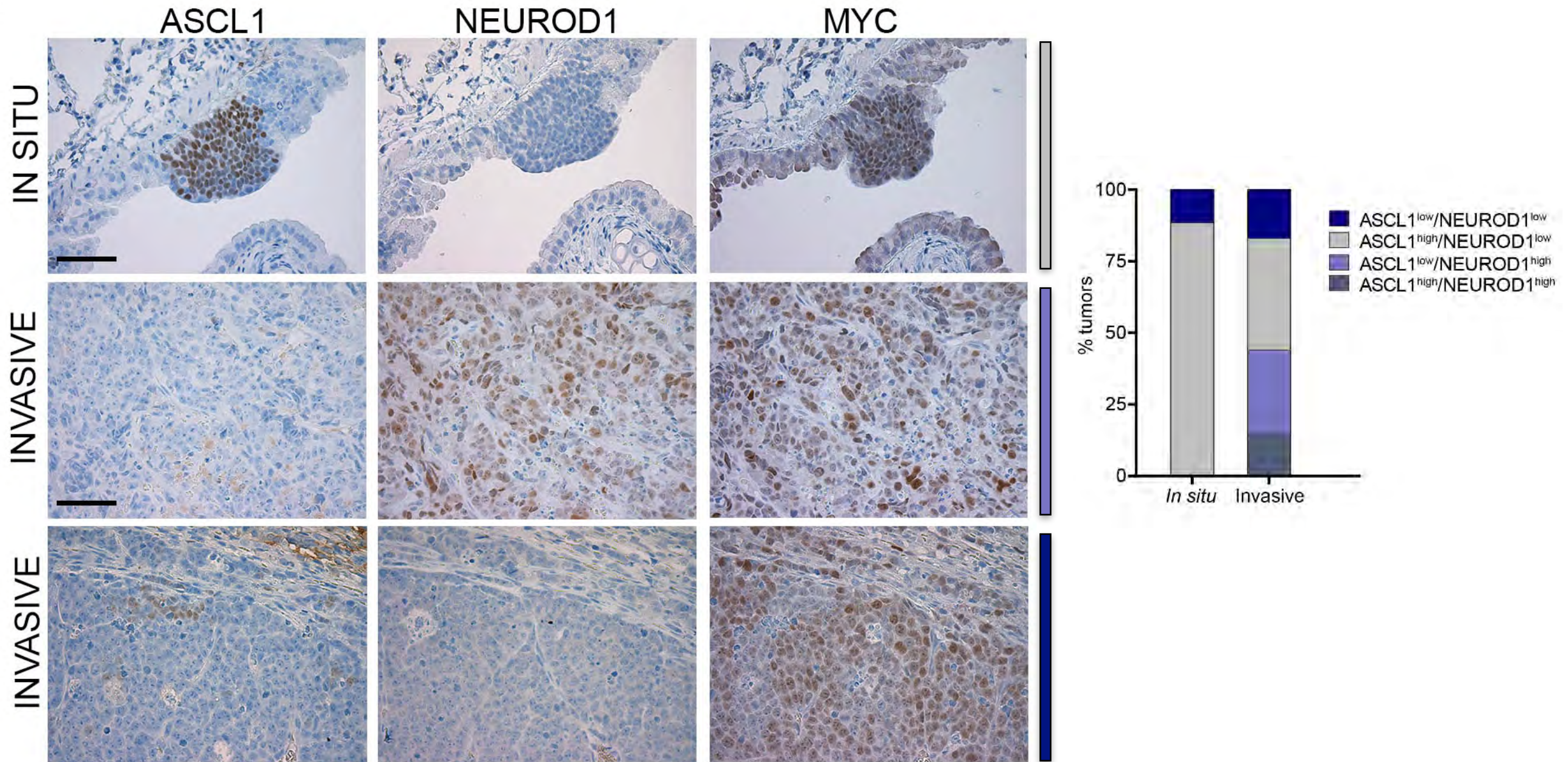
## 26 mouse tumors



Borromeo et al, *Cell Reports*, 2016



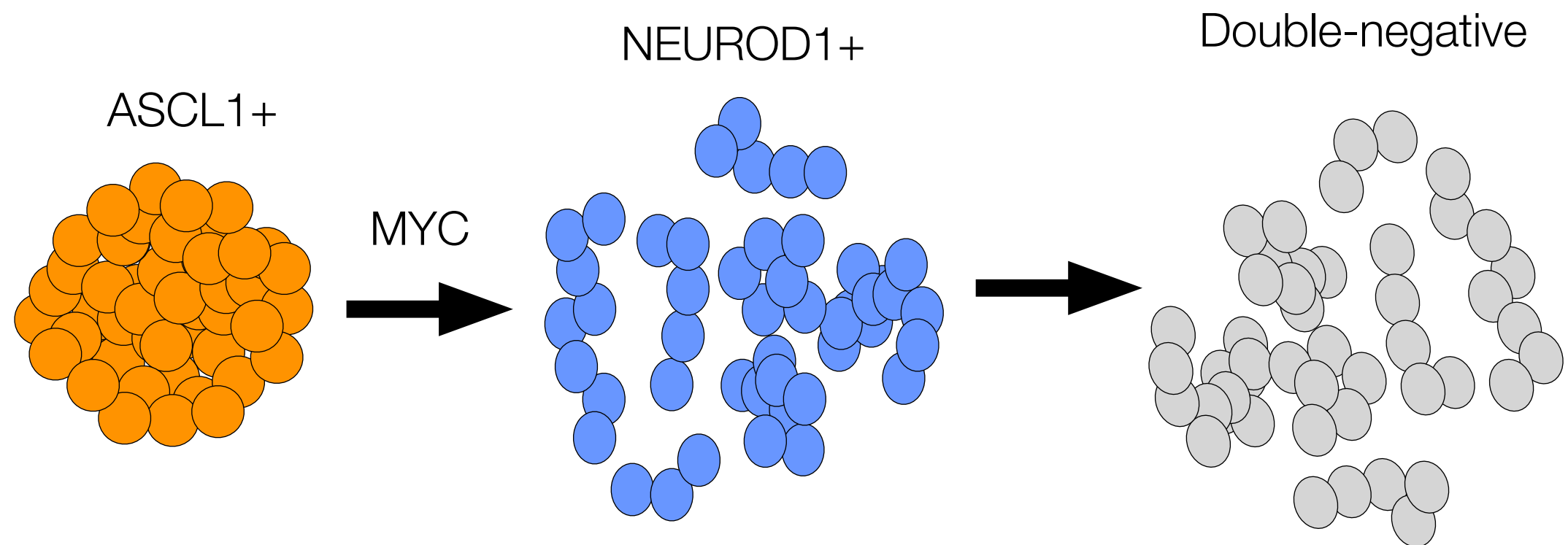
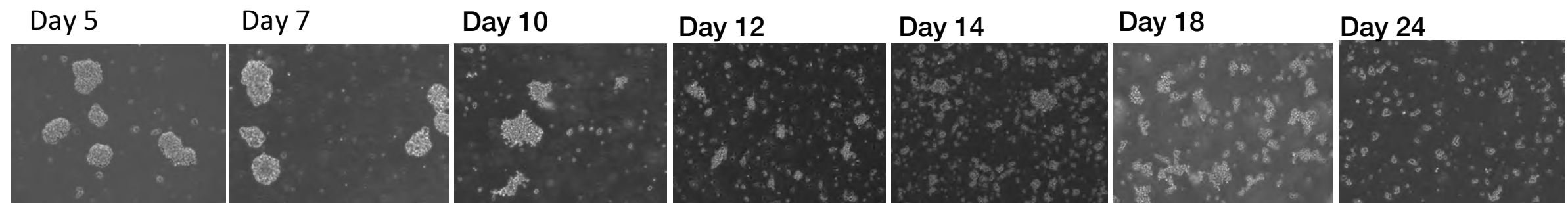
# Hypothesis: SCLC originates in ASCL1+ cells and progresses to NEUROD1+ or double-negative state





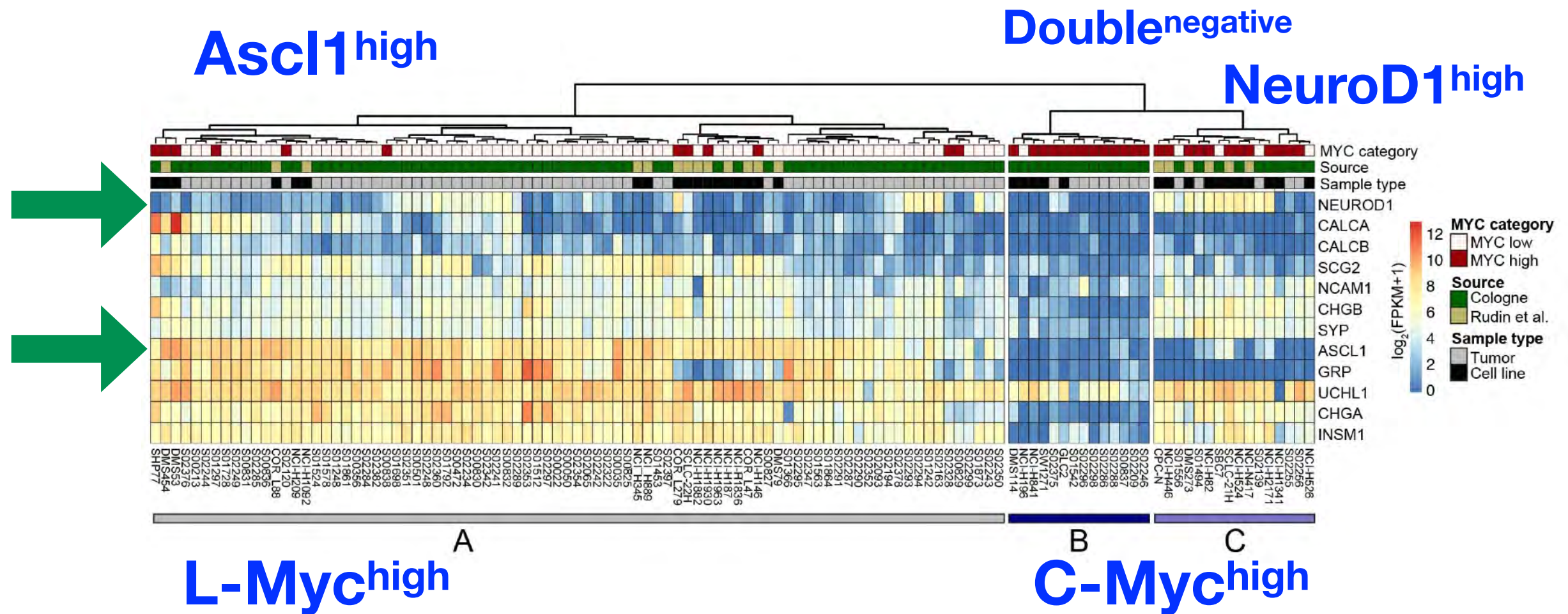
# Hypothesis: MYC promotes transition of ASCL1+ cells to NEUROD1+ to double-negative

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# MYC-high human SCLC exhibits low neuroendocrine gene expression and a switch in ASCL1/NEUROD1

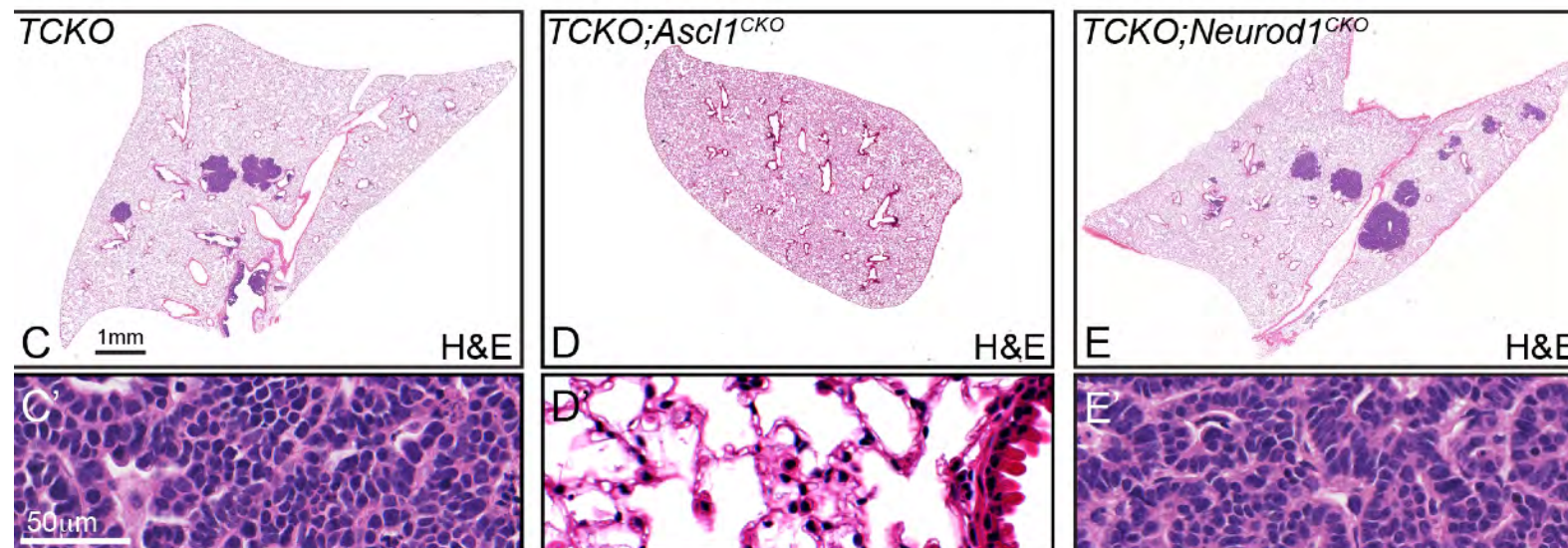
Unsupervised clustering of 81 human tumors and 34 human SCLC cell lines





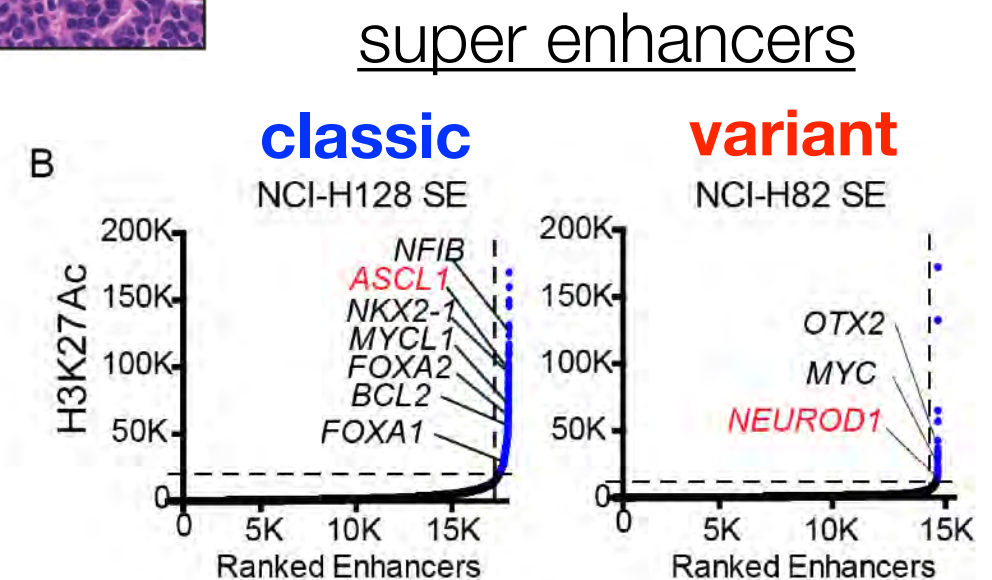
# ASCL1 is required for classic SCLC development

- ASCL1 is required, but NEUROD1 is not required for classic (L-MYC-associated) SCLC



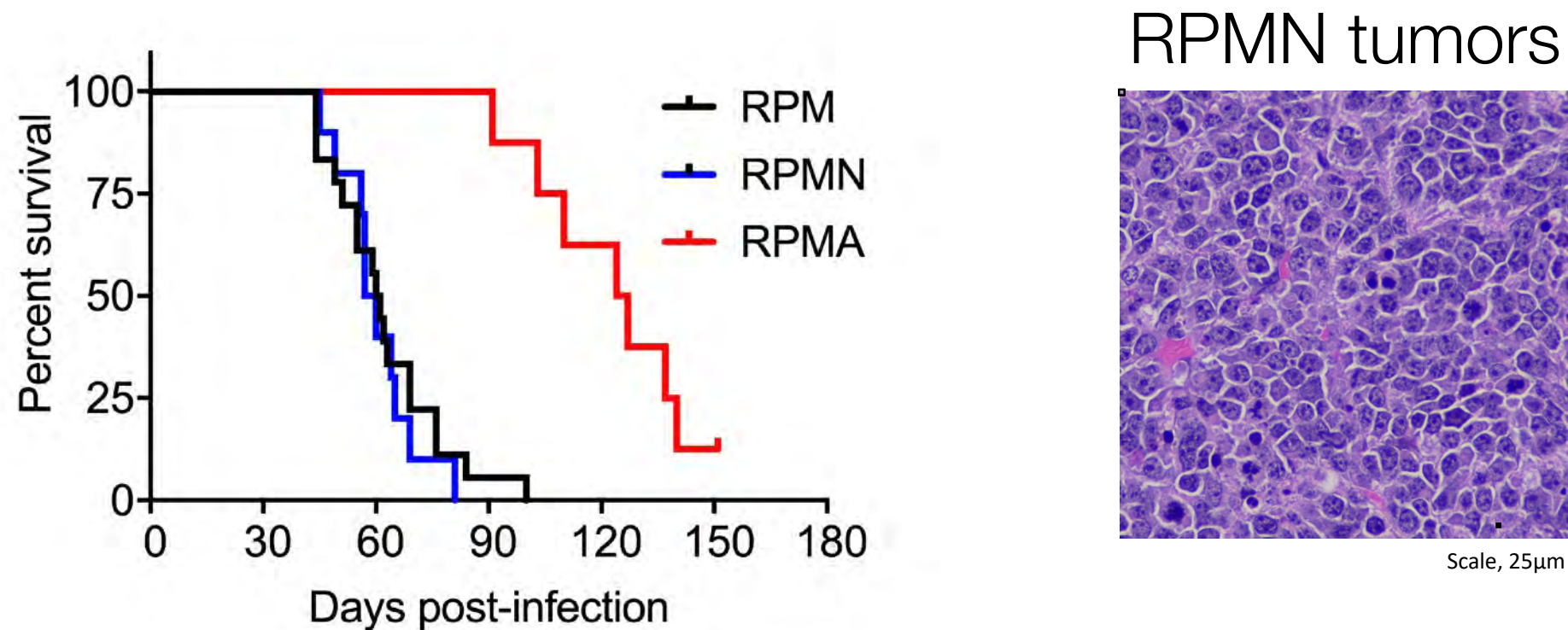
RPR2 model (Sage)

**Are ASCL1 and NEUROD1  
required for MYC-driven SCLC?**



Borromeo et al, Cell Reports, 2016  
Christensen, Cancer Cell, 2014

# NEUROD1 is not required for MYC-driven SCLC

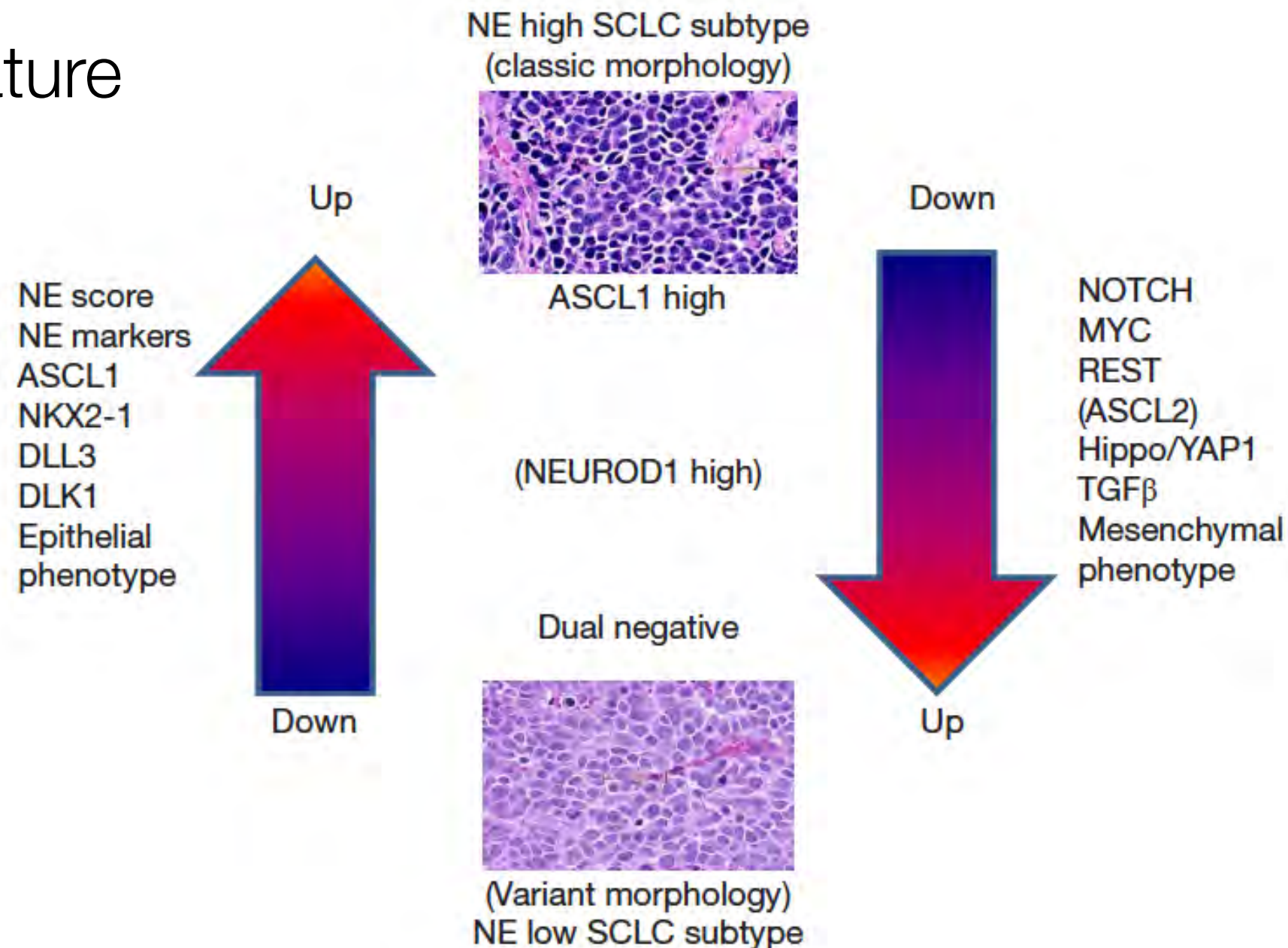


MYC-driven tumors develop in the absence of ASCL1,  
but what are they?

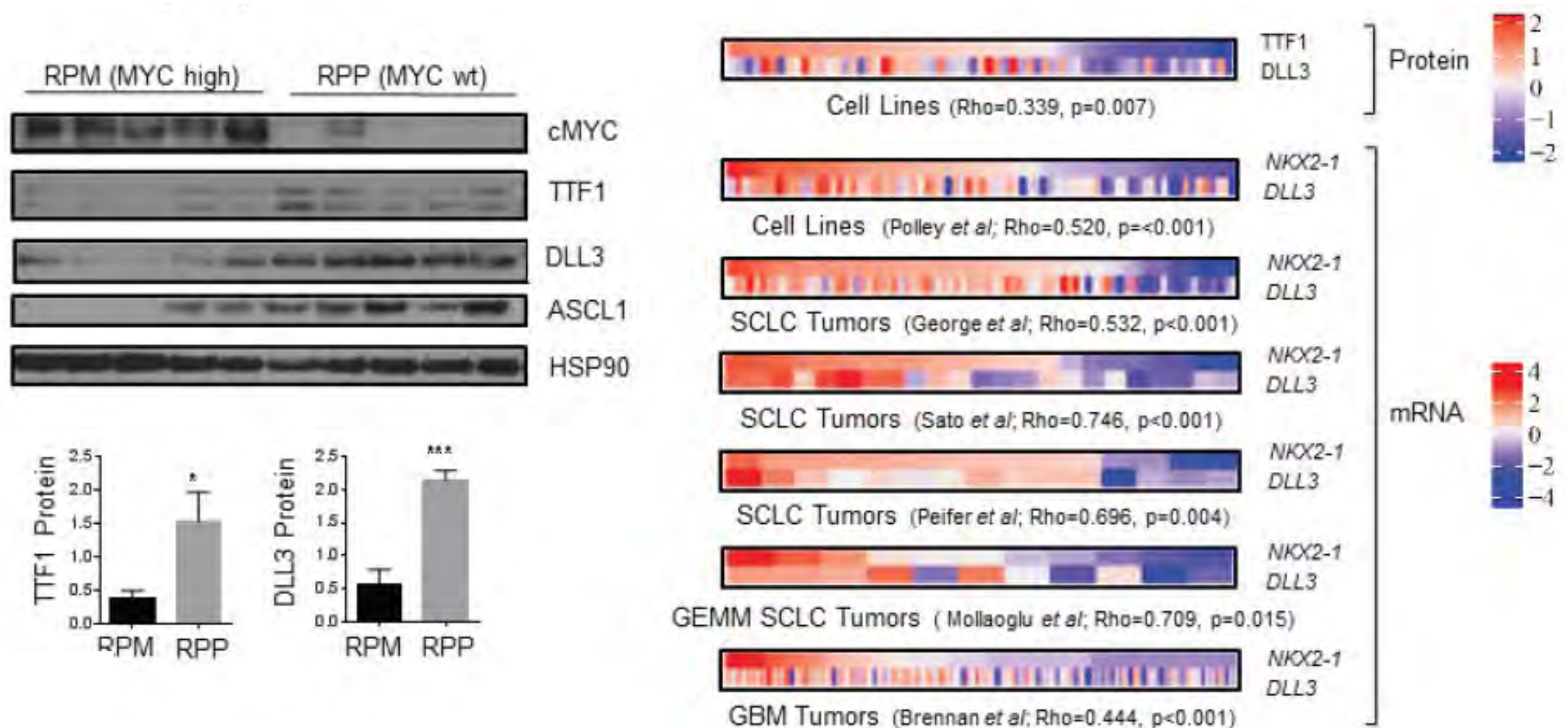


# Features associated with the double-negative “NE-low” state in mouse and human

## NE signature

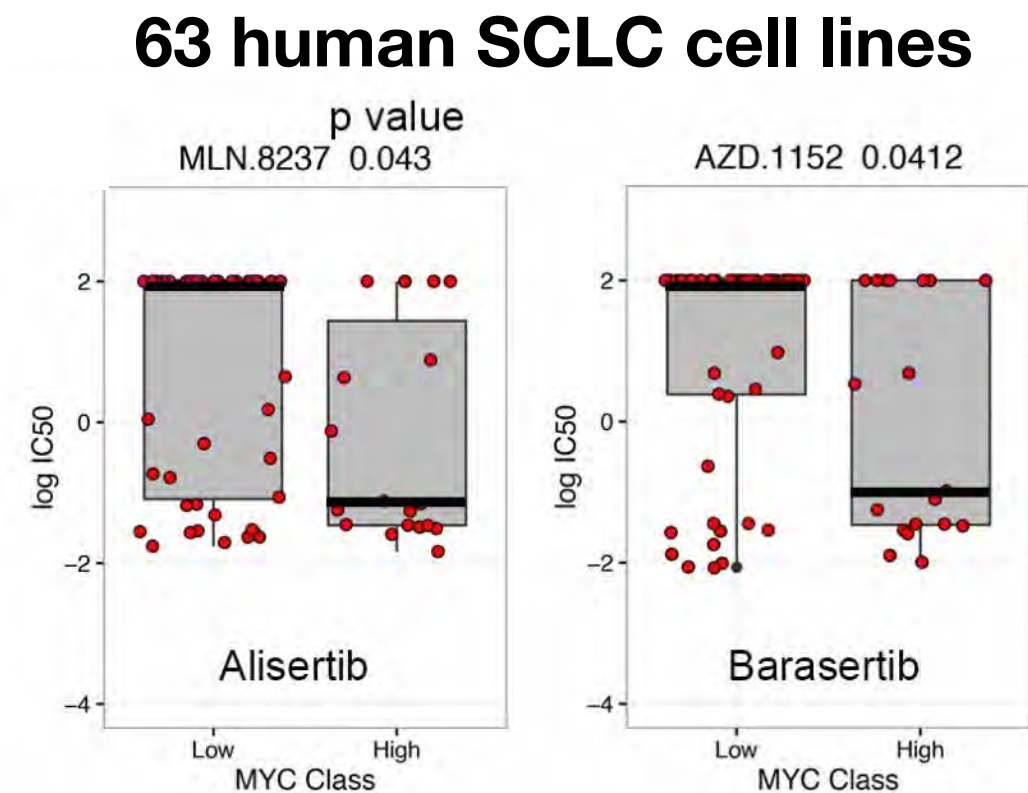
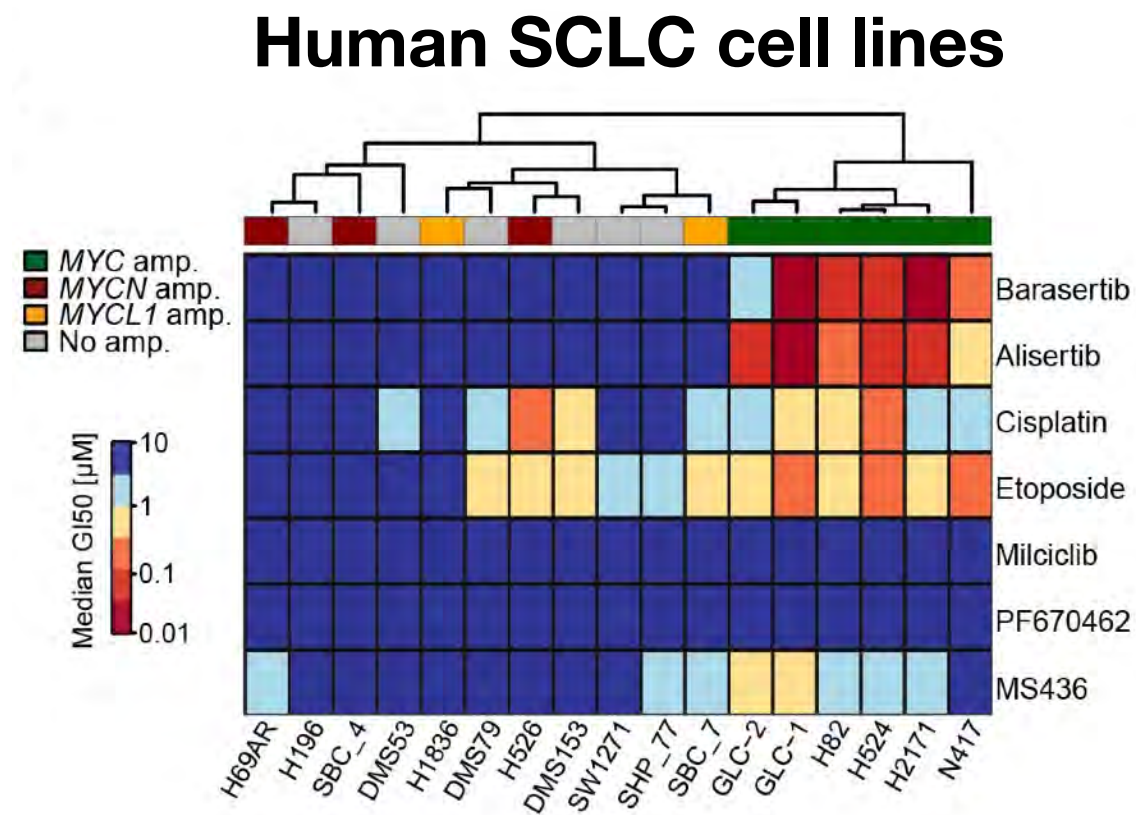


# MYC-driven tumors have reduced NKX2-1/TTF1 and DLL3 (a new SCLC drug target, “Rova-T”)





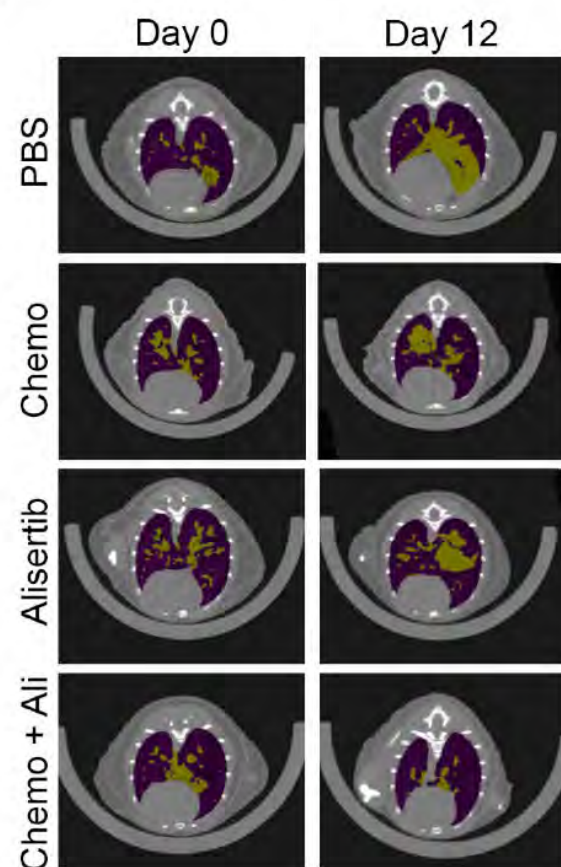
# MYC-high SCLC cell lines preferentially respond to Aurora kinase A/B inhibition



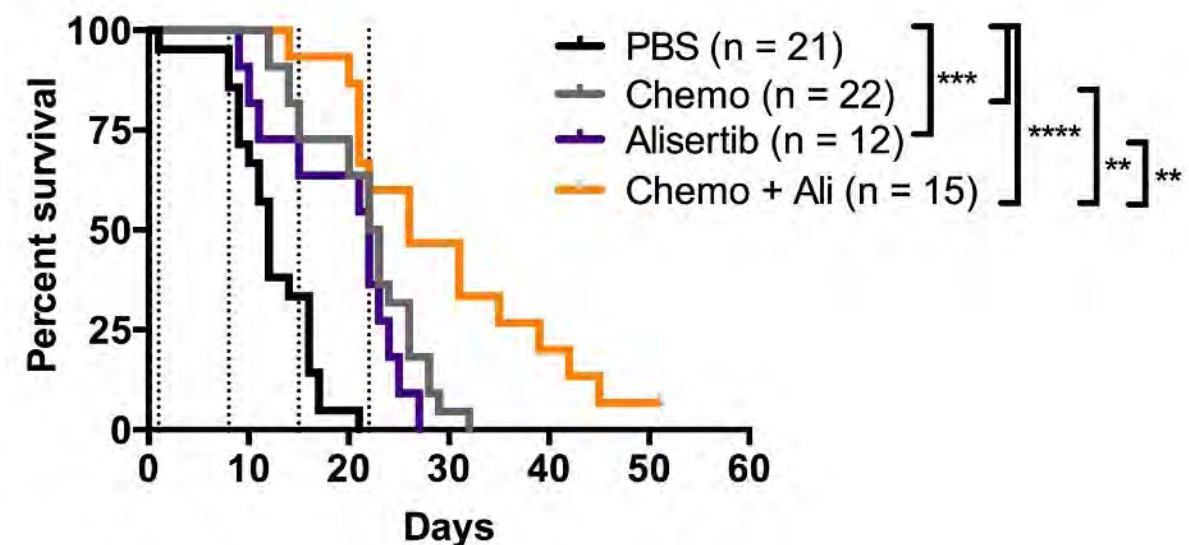
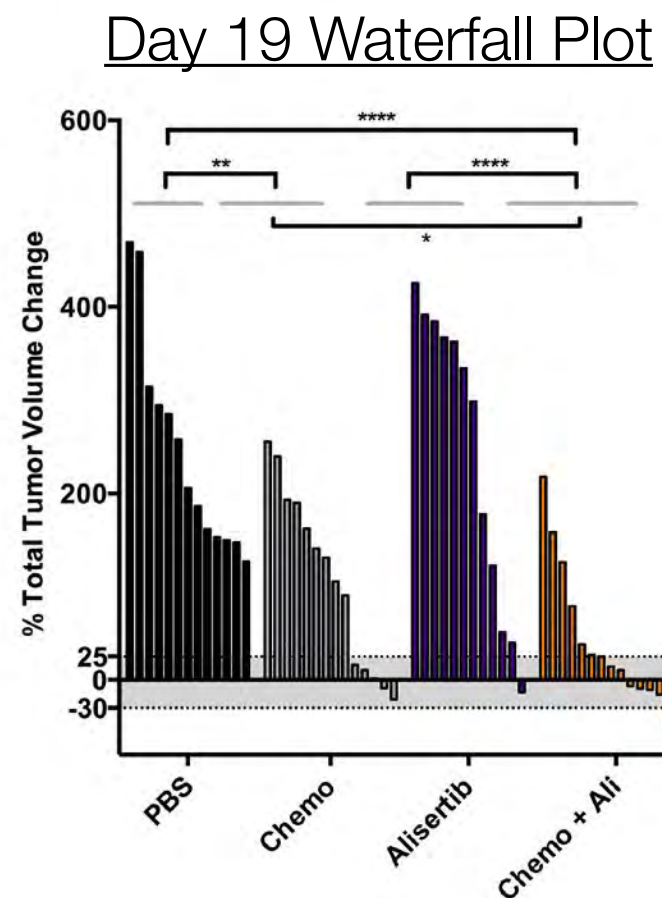
*Polley et al, JNCI, 2016*

21% of relapsed SCLC responded to Alisertib monotherapy  
(Melichar et al, *Lancet Oncology*, 2015)

# Alisertib in combination with chemotherapy significantly improves tumor control and survival



**Tumor / Lung**

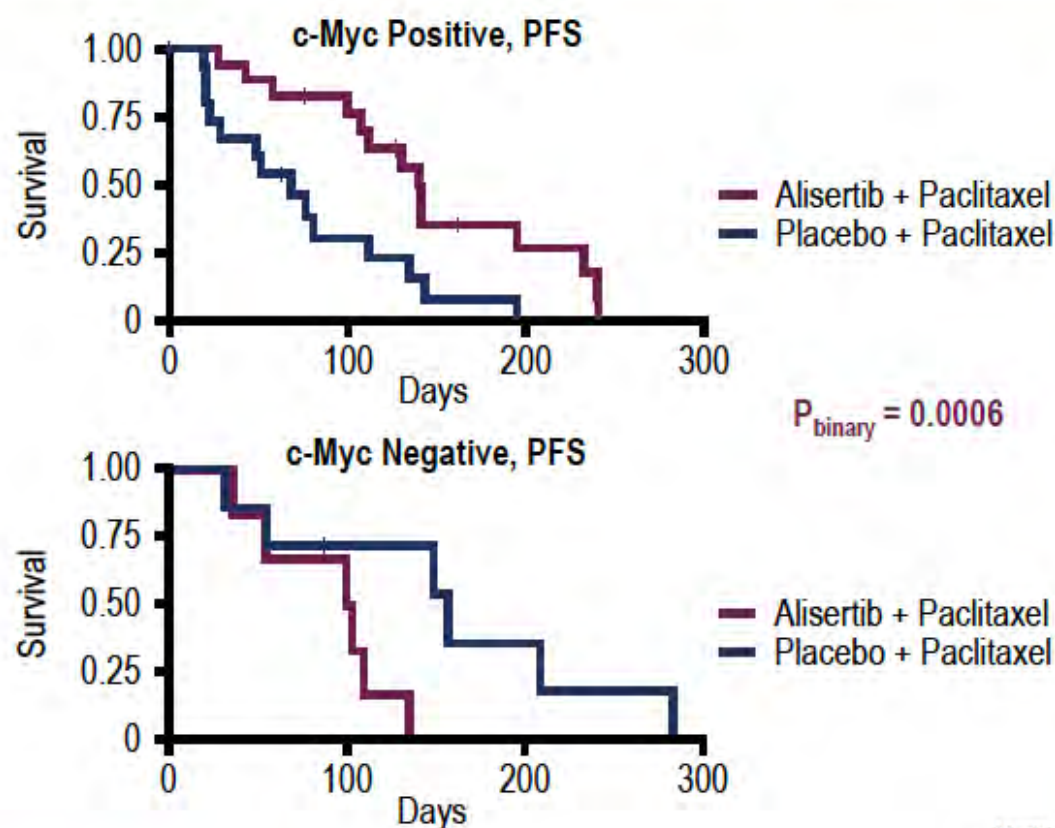


*NCI-funded R21 related to MYC & Alisertib  
RPM mice deposited into JAX #029971*



# Improved survival in patients with MYC-high tumors

n = 46 out of 89 patient samples assessed by MYC IHC



Arm	c-Myc positive	
	n	Median PFS (months)
Alisertib + Paclitaxel	17	4.64
Placebo + Paclitaxel	16	2.27
Hazard Ratio (95% CI)	0.29 (0.12–0.72)	

Arm	c-Myc negative	
	n	Median PFS (months)
Alisertib + Paclitaxel	6	3.32
Placebo + Paclitaxel	7	5.16
Hazard Ratio (95% CI)	11.8 (1.52–91.2)	

\*Archived tumor tissue available from 46 patients. Modal intensity for c-Myc positive= 1+, 2+, 3+ IHC score.  
Modal intensity for c-Myc negative = 0 IHC score.



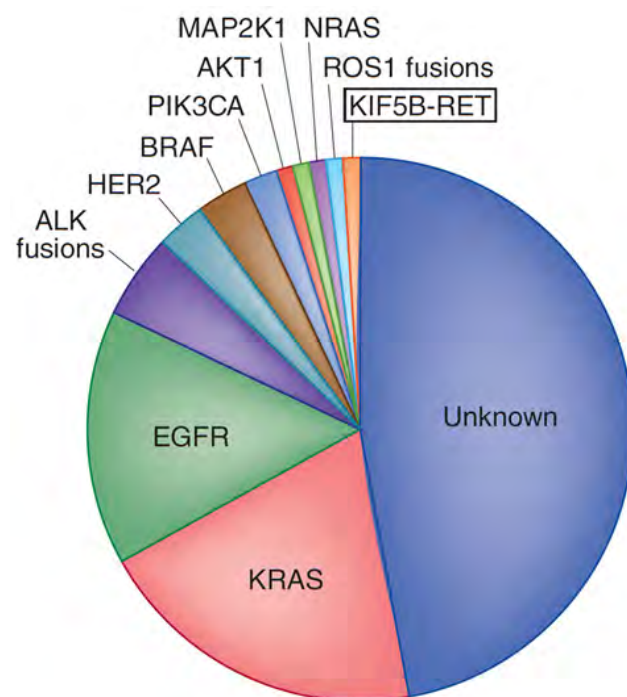
**MYC-high SCLC is preferentially sensitive to Aurora kinase inhibition**

MYC promotes aggressive, neuroendocrine-low, “variant” subtype (NEUROD1<sup>+</sup> or Double<sup>Negative</sup>) of SCLC that is vulnerable to Aurora kinase inhibition

Molecular and functional subtypes of SCLC exist (MYC vs. L/N-MYC) **with clinical implications**

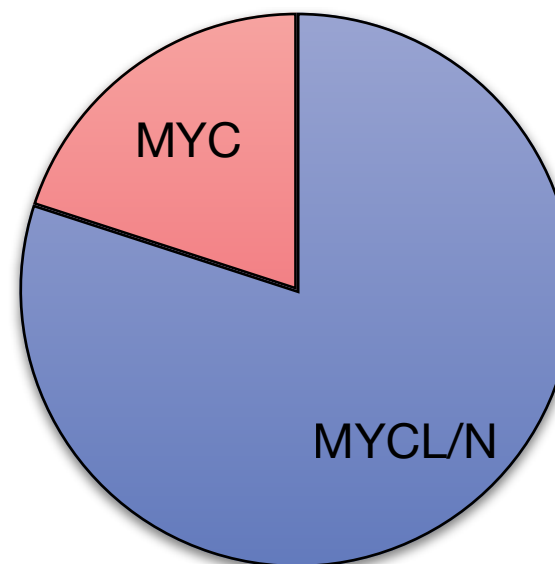
While subtype-specific therapies are not yet approved (and need to be improved), **preclinical and clinical trials** should examine these biomarkers to best determine subtype-specific therapies

NSCLC



Source : Nature Medicine 18, 349–351 (2012)

SCLC

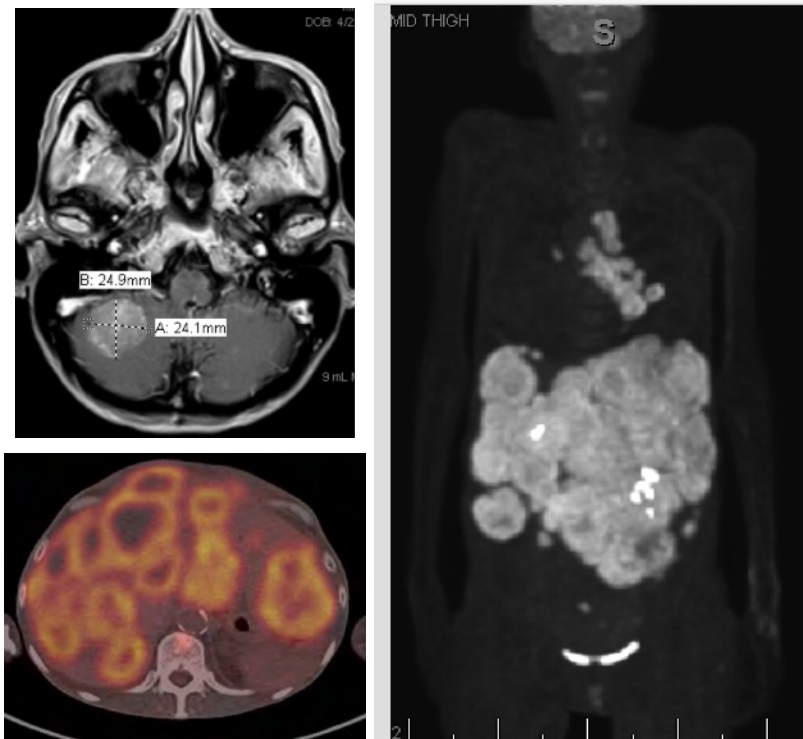


**SCLC subset differences:**

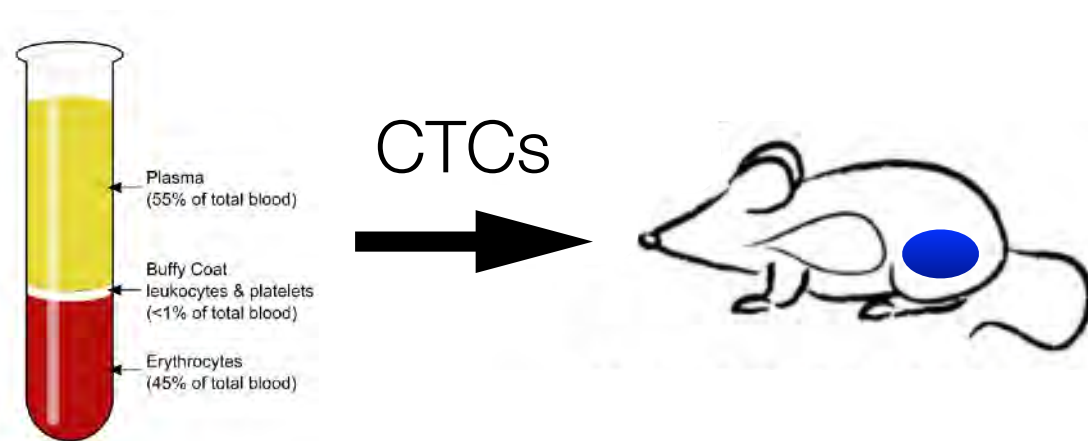
- Metabolic profile
- Apoptotic BH3 profile
- Immune profile
- Metastatic drivers



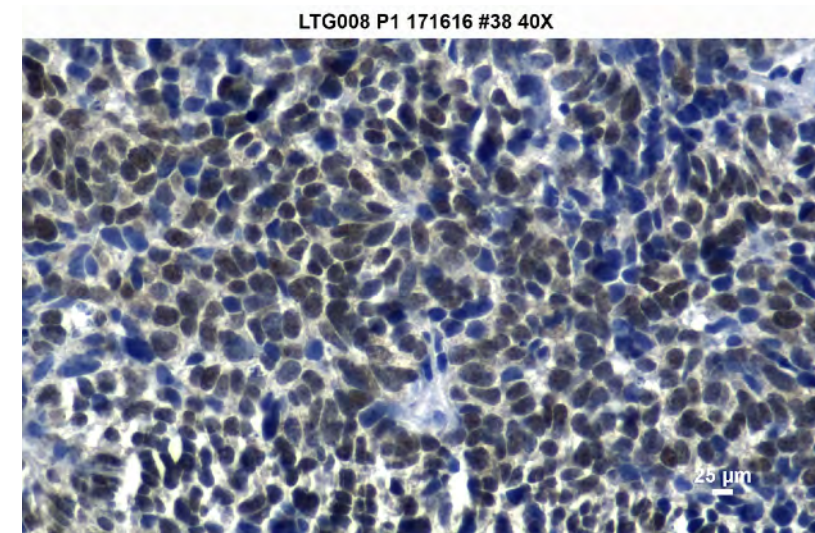
# CDX models at Huntsman Cancer Institute, n = 11



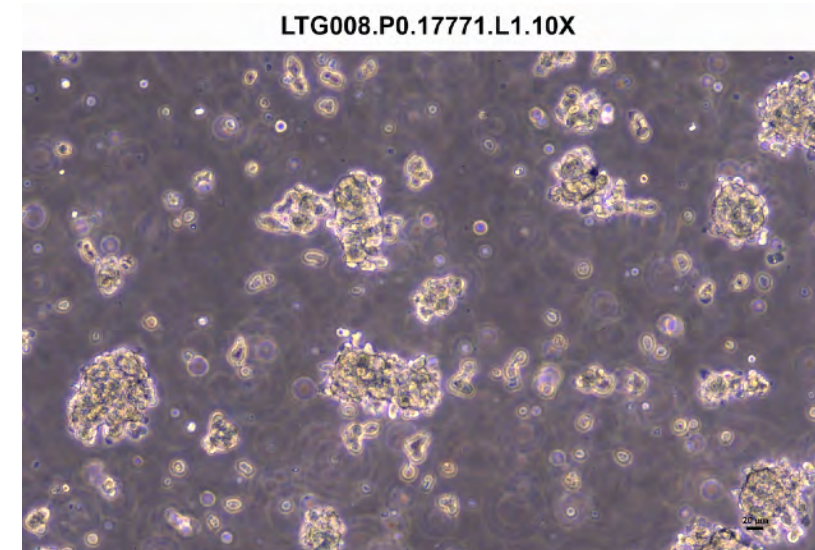
*Female, 2/6/2018*



CDX tumor, ASCL1+



CDX-derived cell line



*With planned sequencing from Charlie Rudin, JT Poirier, Roman Thomas and Julie George.....  
to be deposited into cBioportal*





## The Oliver Lab

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Matt  
Milind  
Rachelle  
Abbie  
Sarah  
Chris  
Sangmin  
Sophia



*Always hiring*



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Eric Snyder, MD/PhD  
Mohamed Salama, MD  
Ben Witt, MD  
Bioinformatics Core

**Martin Sos Lab:** Stef & Johannes



Damon Runyon  
**Cancer Research**  
Foundation



Mini-grants from Cell Response & Regulation and UADOT



