

# Multi-cancer mutual exclusivity analysis of genomic alterations

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**Giovanni Ciriello**  
Computational Biology - MSKCC  
*TCGA Annual Symposium*  
Washington DC, 2011

# Mutually exclusive alterations in Cancer

## Recurrent genomic alterations target specific pathways

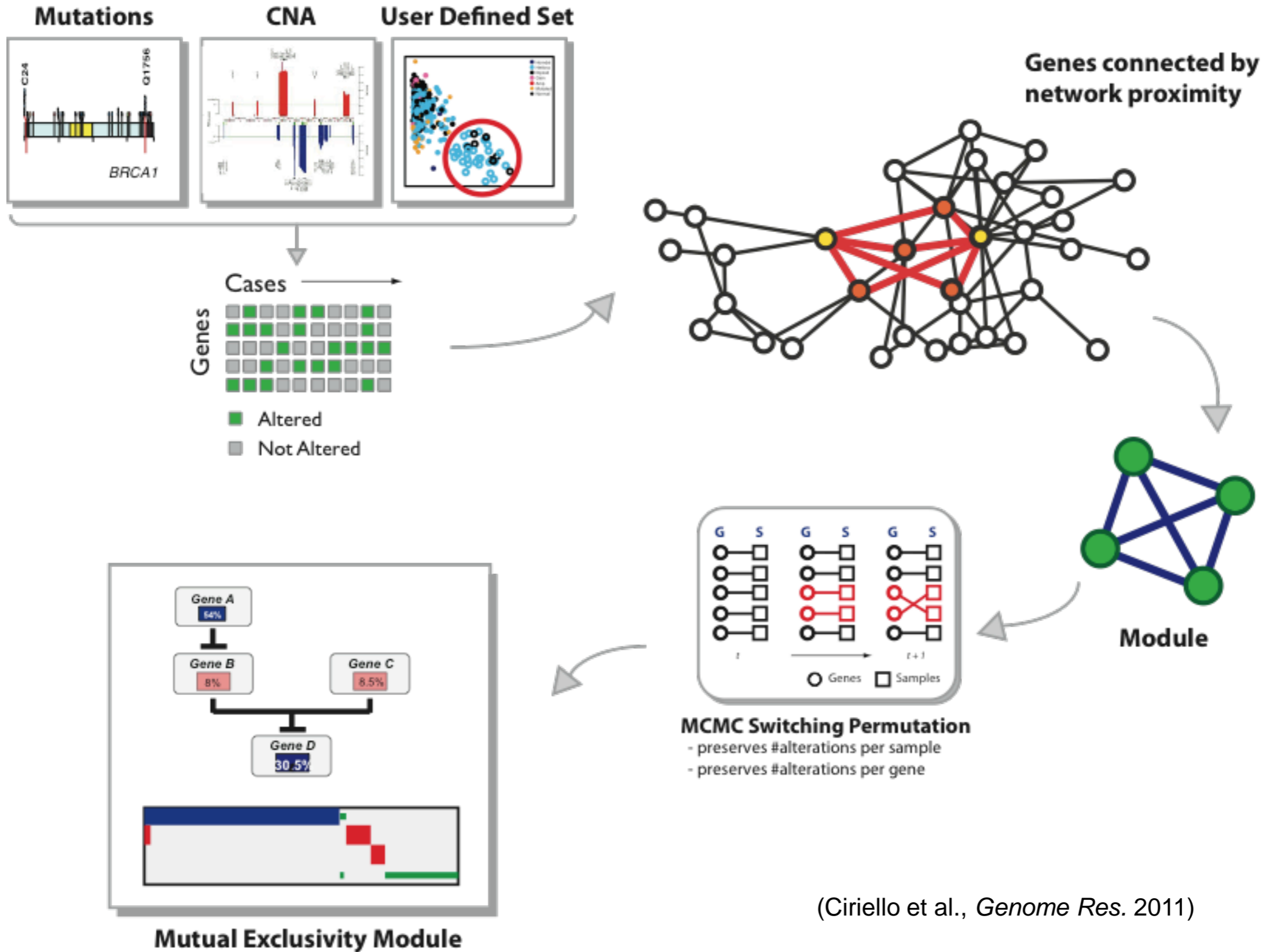


**Functional alterations targeting the same pathway frequently occur in a mutually exclusive manner**



(TCGA, Nature, 2011)

# MEMo: Mutual Exclusivity Modules



(Ciriello et al., *Genome Res.* 2011)



# MEMo results on TCGA Datasets

MEMo has been applied to the following TCGA projects:

- **Glioblastoma Multiforme (GBM)**
  - Phase 2 338 samples
- **Serous Ovarian Cancer (OVCA)**
  - Updated dataset 384 samples
- **Colon and Rectum Adenocarcinoma (COAD)**
  - Non hyper-mutators 151 samples
- **Uterine Corpus Endometriod Carcinoma (UCEC)**
  - Non serous / Non hyper-mutators 144 samples
- **Invasive Breast Cancer (BRCA)**
  - 463 samples

Mutually exclusive patterns of alteration identified in several oncogenic pathways:

- Rb - signaling
- p53 - signaling
- DNA repair
- PI(3)K/Akt signaling



# MEMo results on TCGA Datasets

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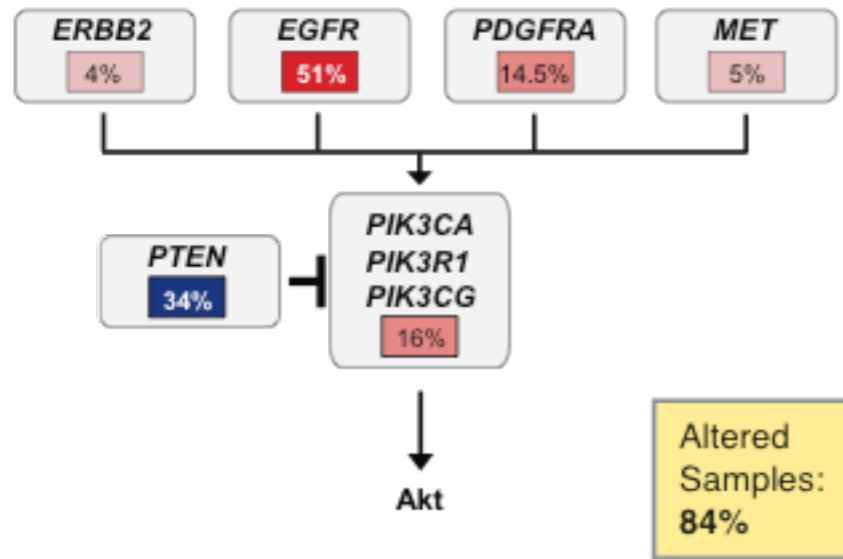
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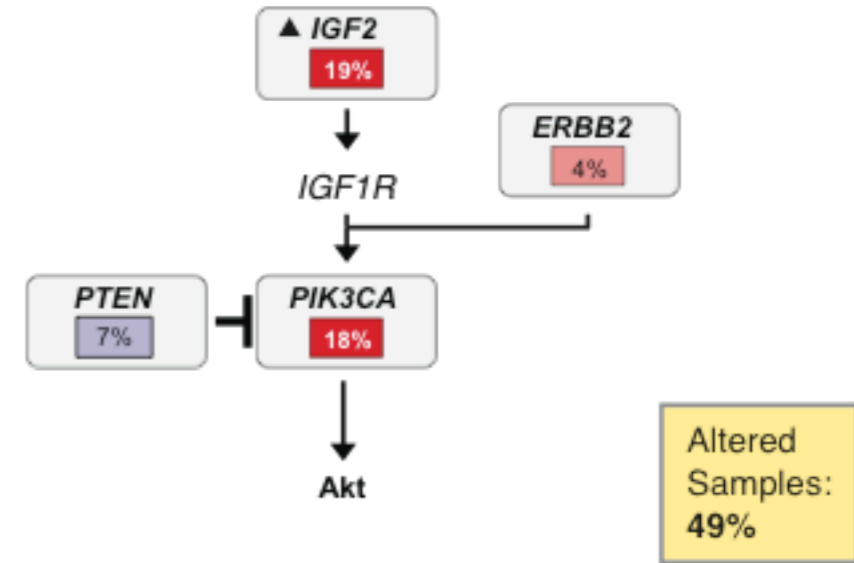
- Rb - signaling
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- DNA repair
- **PI(3)K/Akt signaling**

# Mutual exclusivity in PI(3)K/Akt

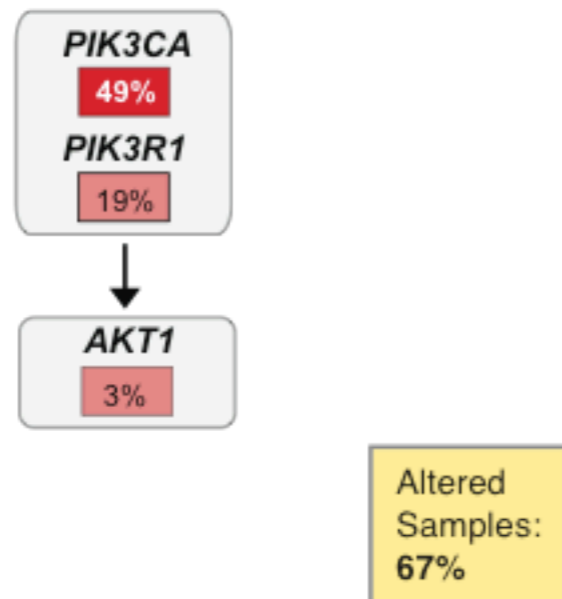
## GBM (338 samples)



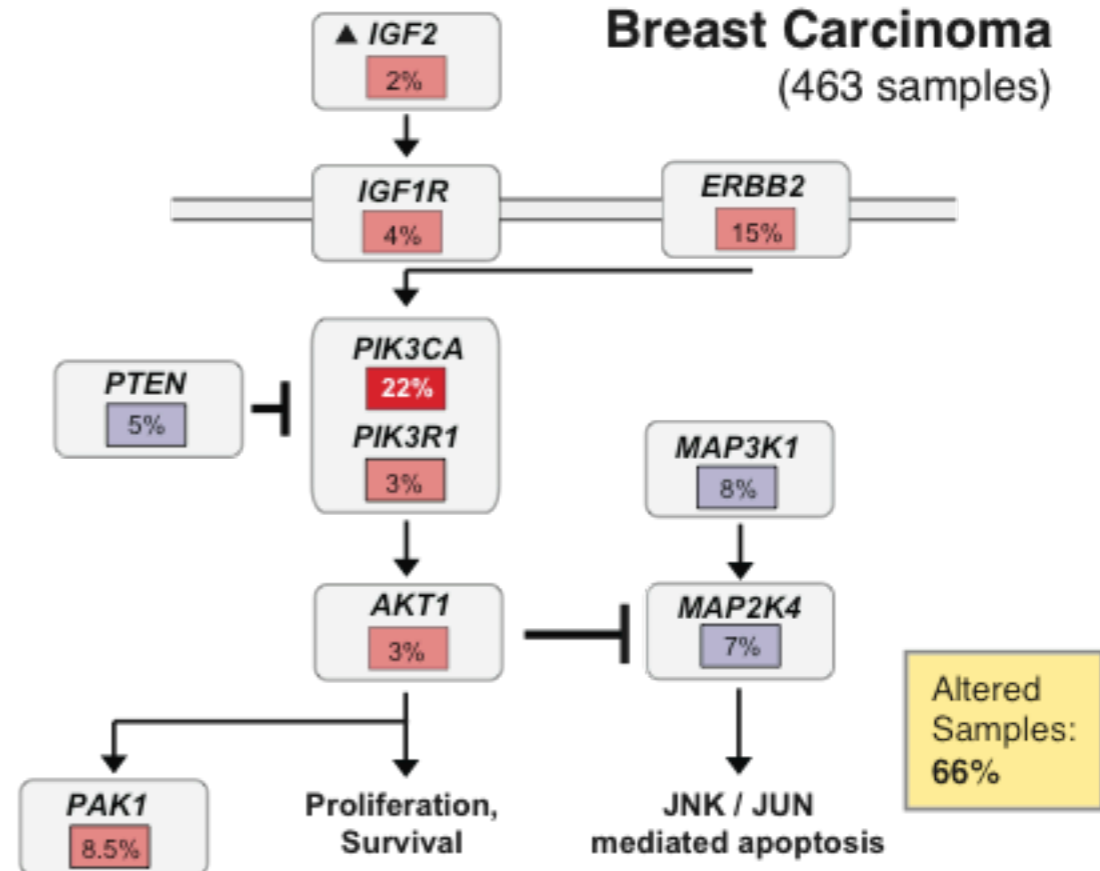
## Colon and Rectum Adenocarcinoma (151 samples)



## Endometriod Carcinoma (144 samples)

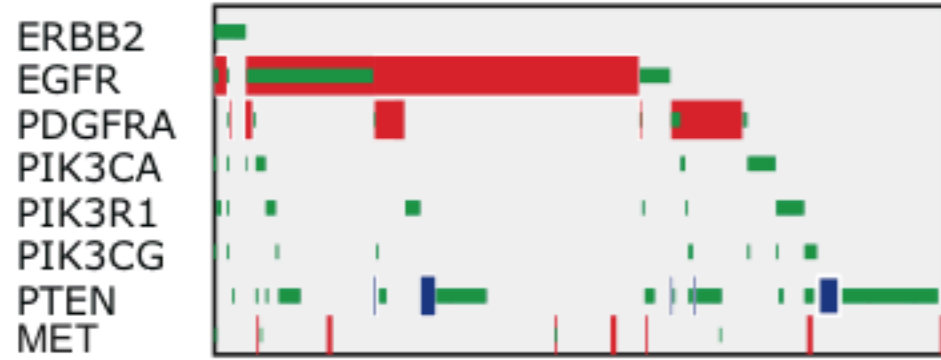


## Breast Carcinoma (463 samples)



# Mutual exclusivity in PI(3)K/Akt

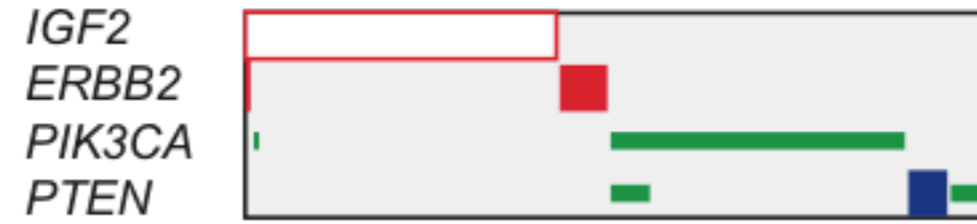
**GBM** (338 samples)



Altered Samples:  
**84%**

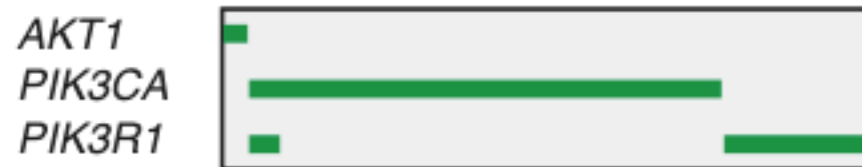
**Colon and Rectum Adenocarcinoma**

(151 samples)



Altered Samples:  
**49%**

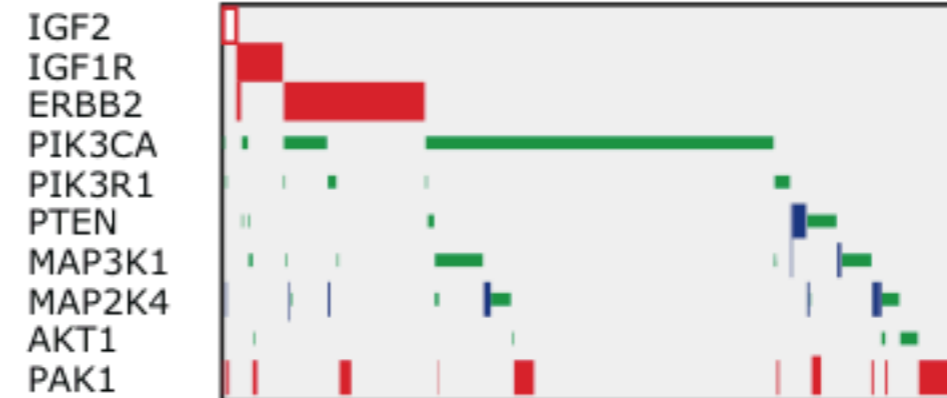
**Endometriod Carcinoma** (144 samples)



Altered Samples:  
**67%**

**Breast Carcinoma**

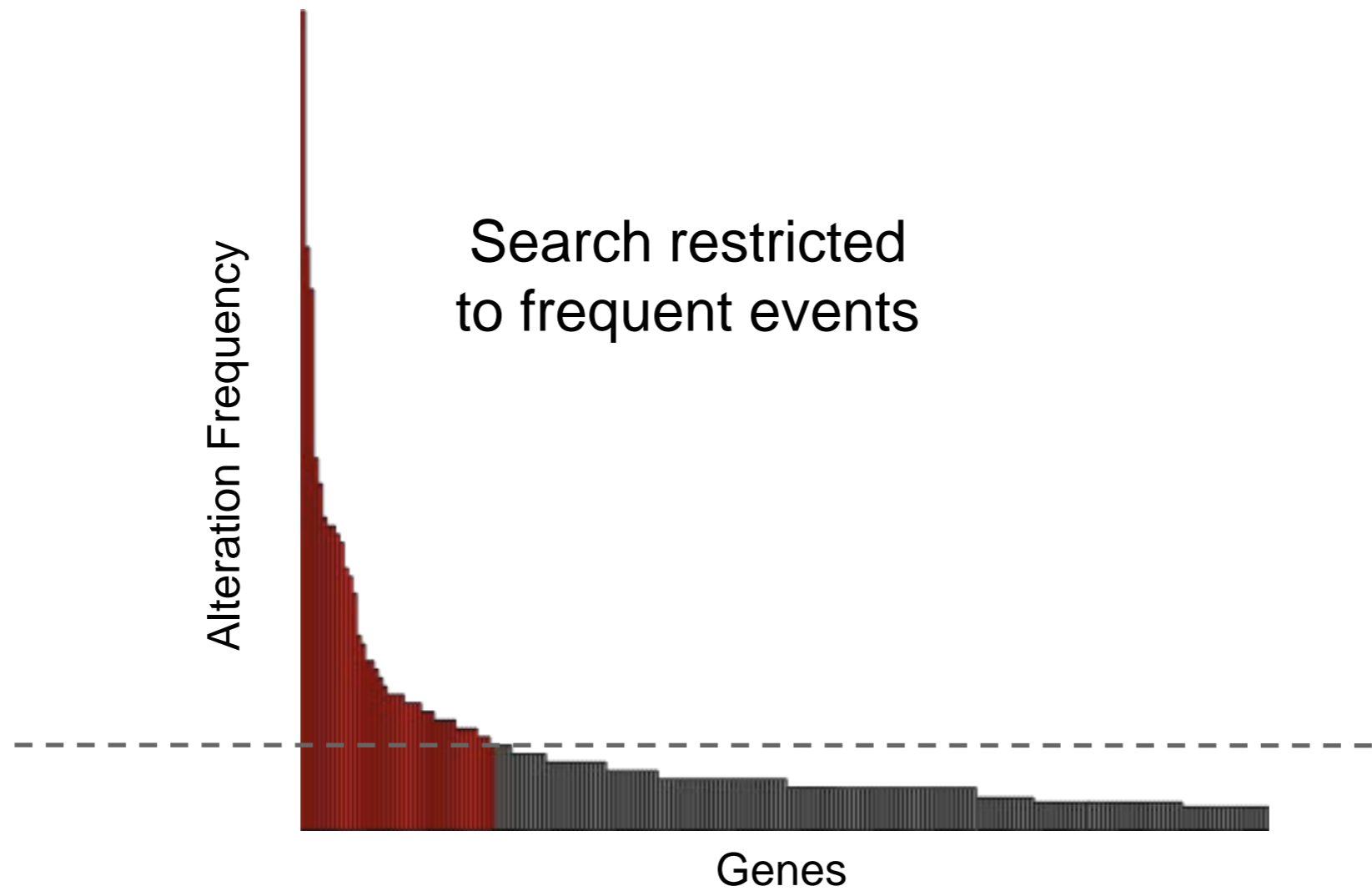
(463 samples)



Altered Samples:  
**66%**

# PI(3)K/Akt alterations in Ovarian Carcinoma

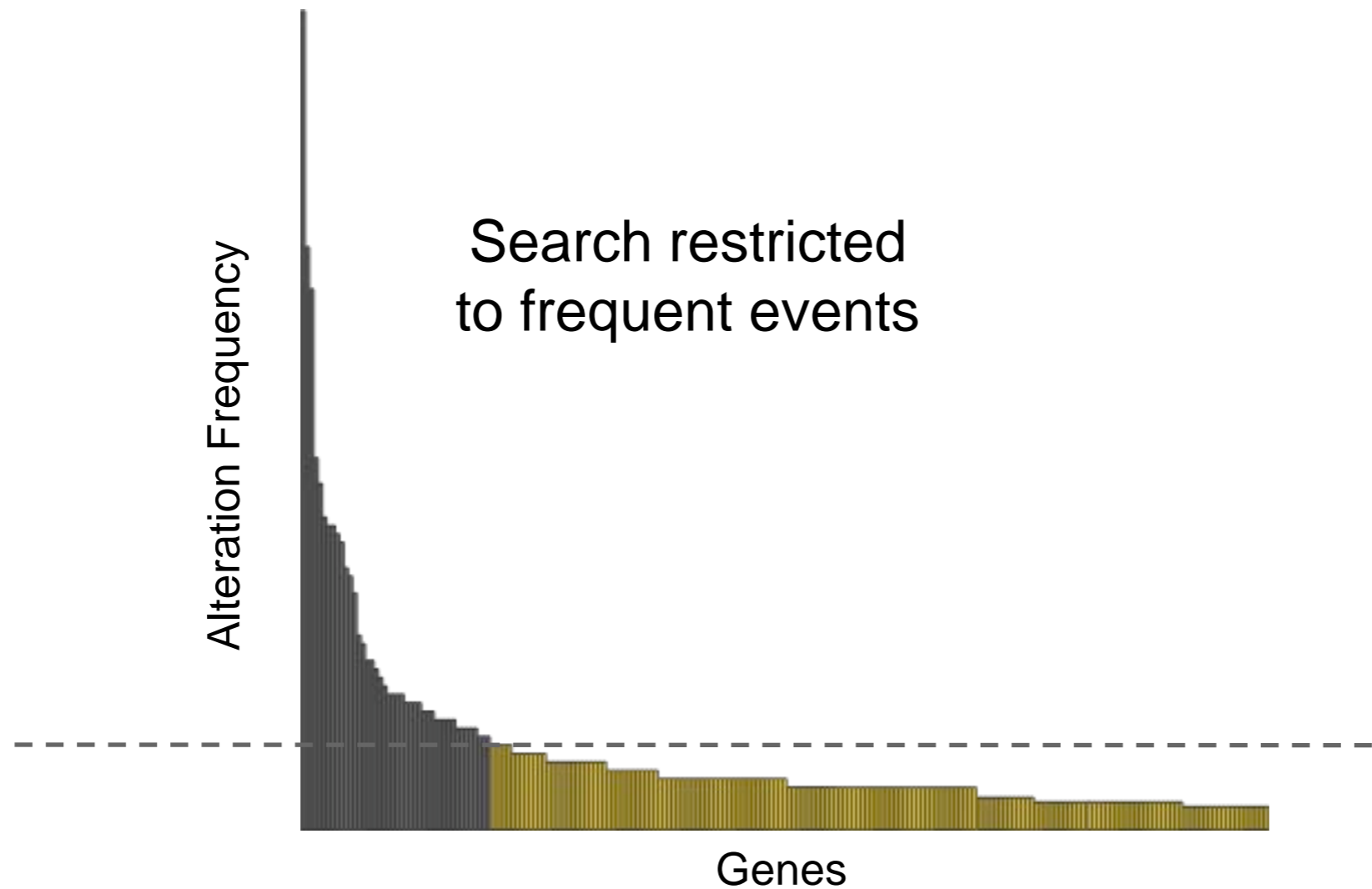
MEMo does not find PI(3)K/Akt modules





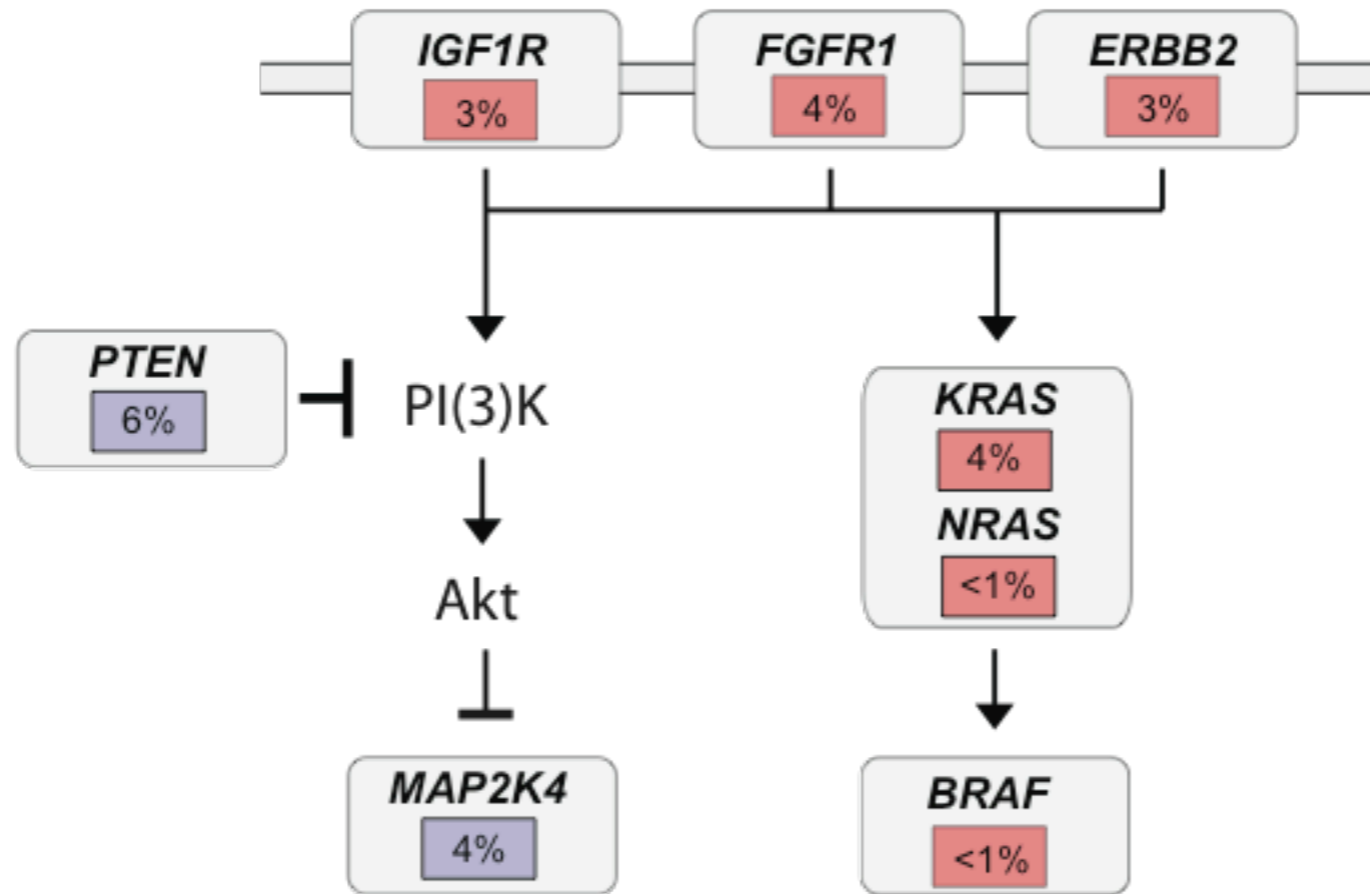
# PI(3)K/Akt alterations in Ovarian Carcinoma

MEMo does not find PI(3)K/Akt modules



Are there low-frequency but functional events affecting this pathway?

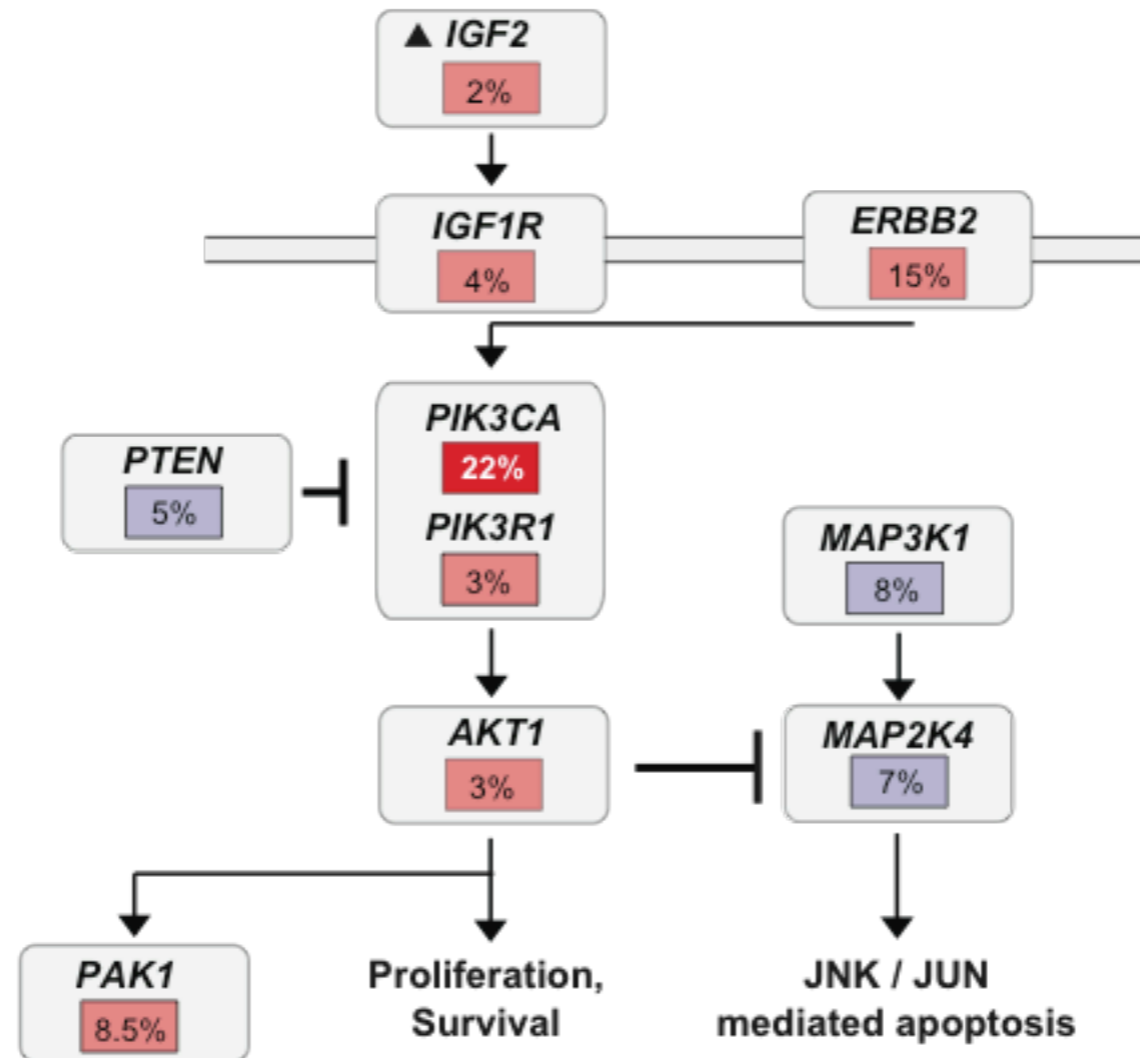
# Multiple Low-frequency events target PI(3)K pathway



24% Altered Samples

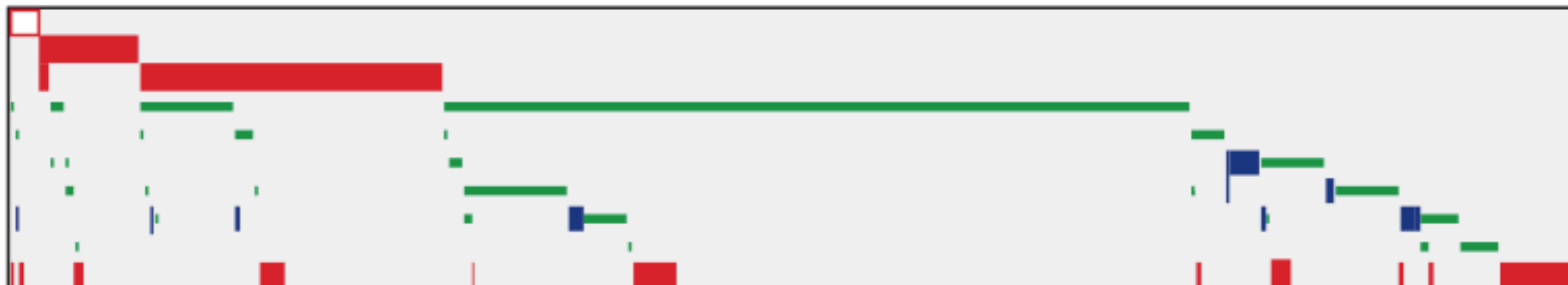


# Breast Cancer (463 samples)



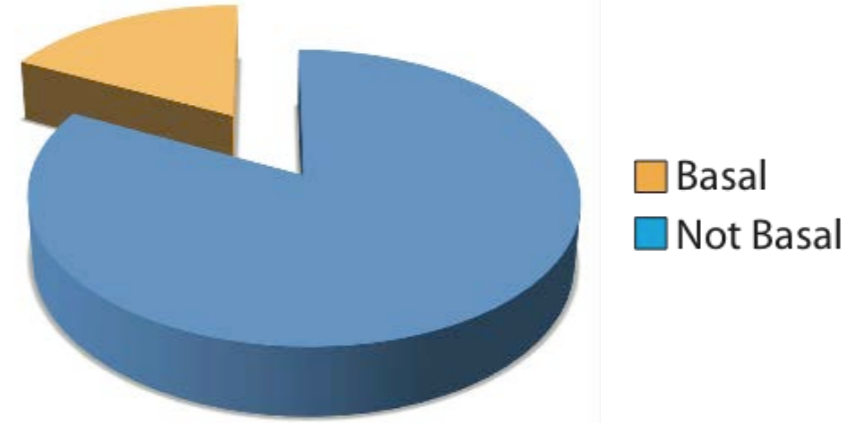
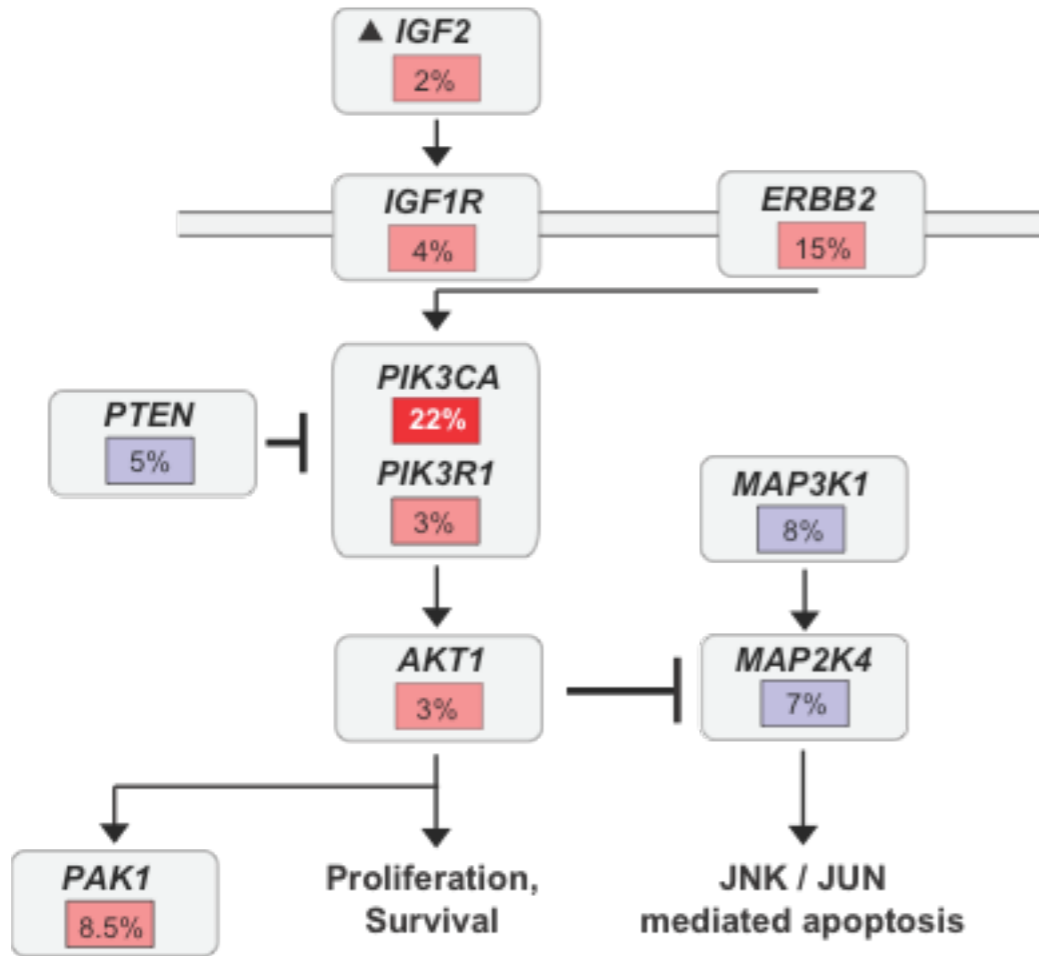
Altered Samples 66%

IGF2  
IGF1R  
ERBB2  
PIK3CA  
PIK3R1  
PTEN  
MAP3K1  
MAP2K4  
AKT1  
PAK1



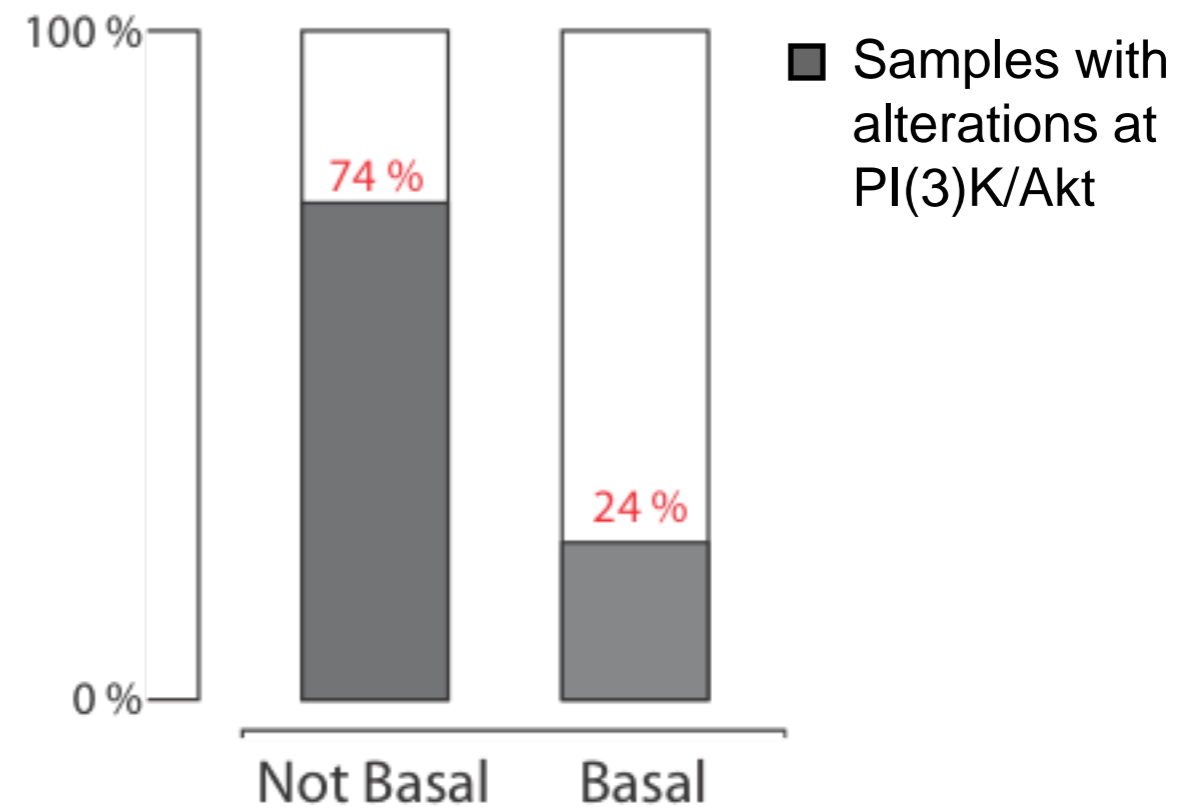
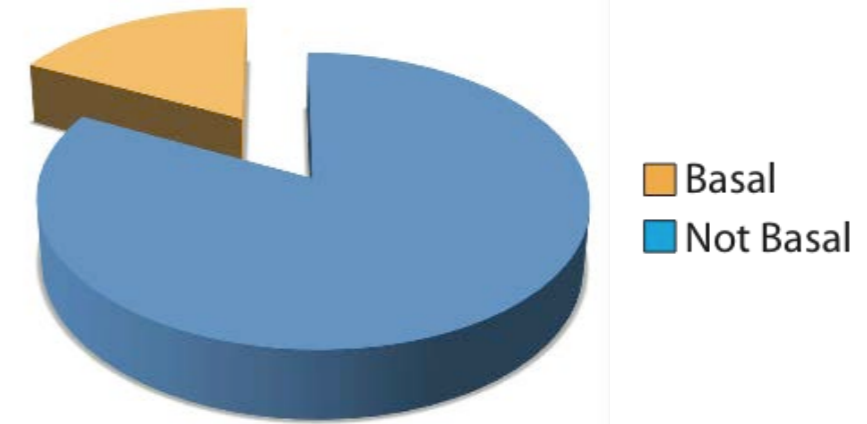
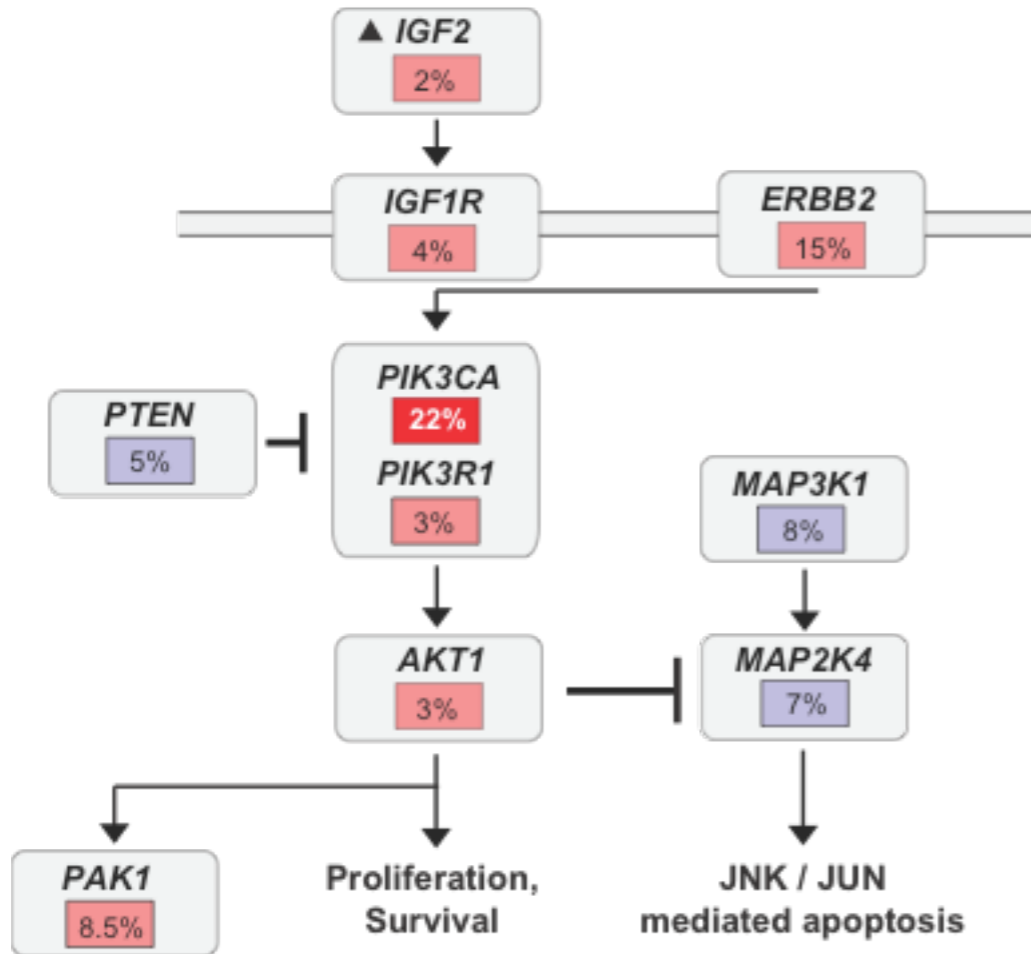
Over-expressed Amplified Hom. Del. Mutated

# Basal vs. Not Basal



Basal  
Not Basal

# Basal vs. Not Basal

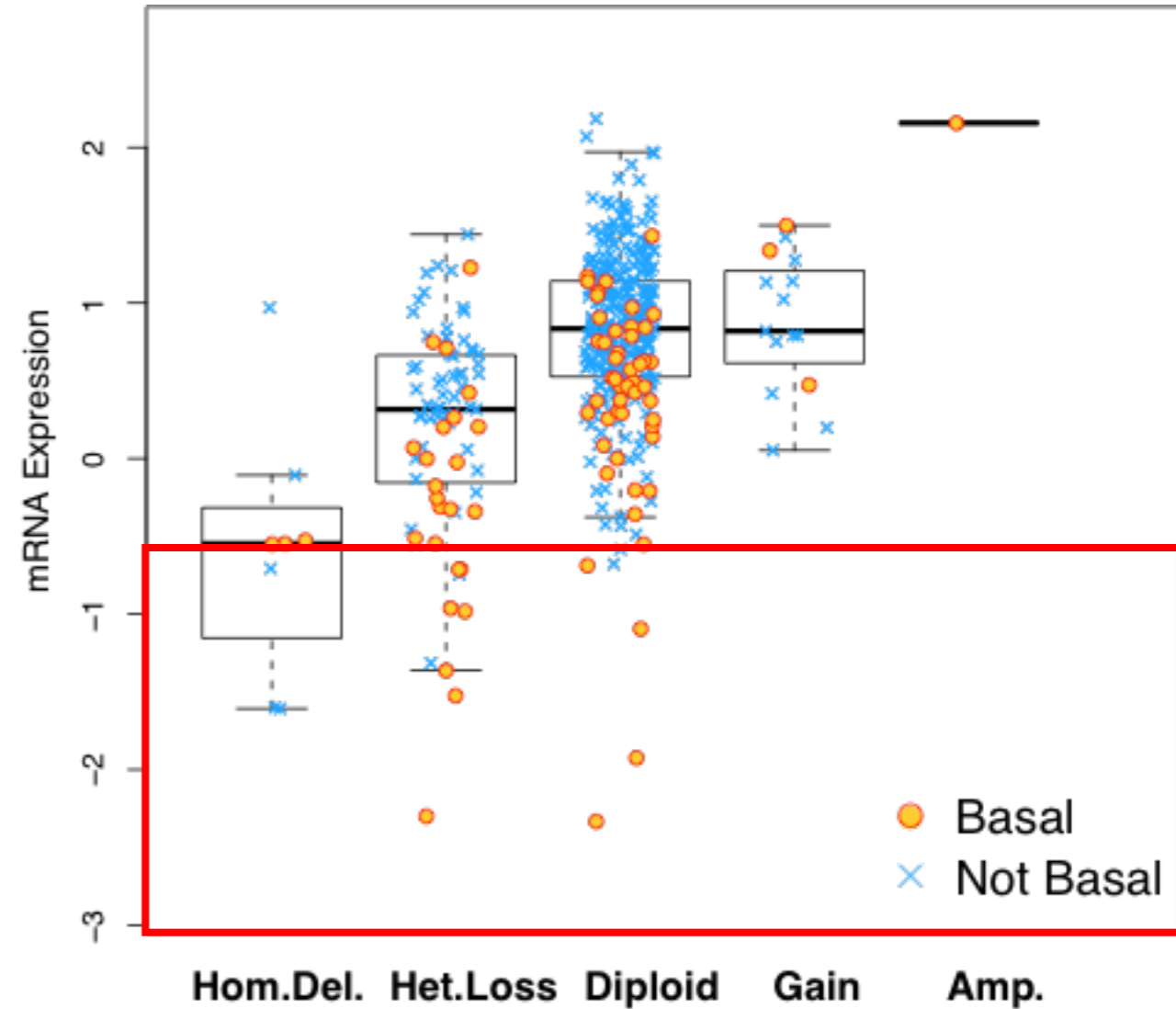


Is the PI(3)K pathway altered by other means in Basal tumors?

# PTEN is down-regulated in Basal Breast Cancer

- PTEN is down-regulated in Basal tumors

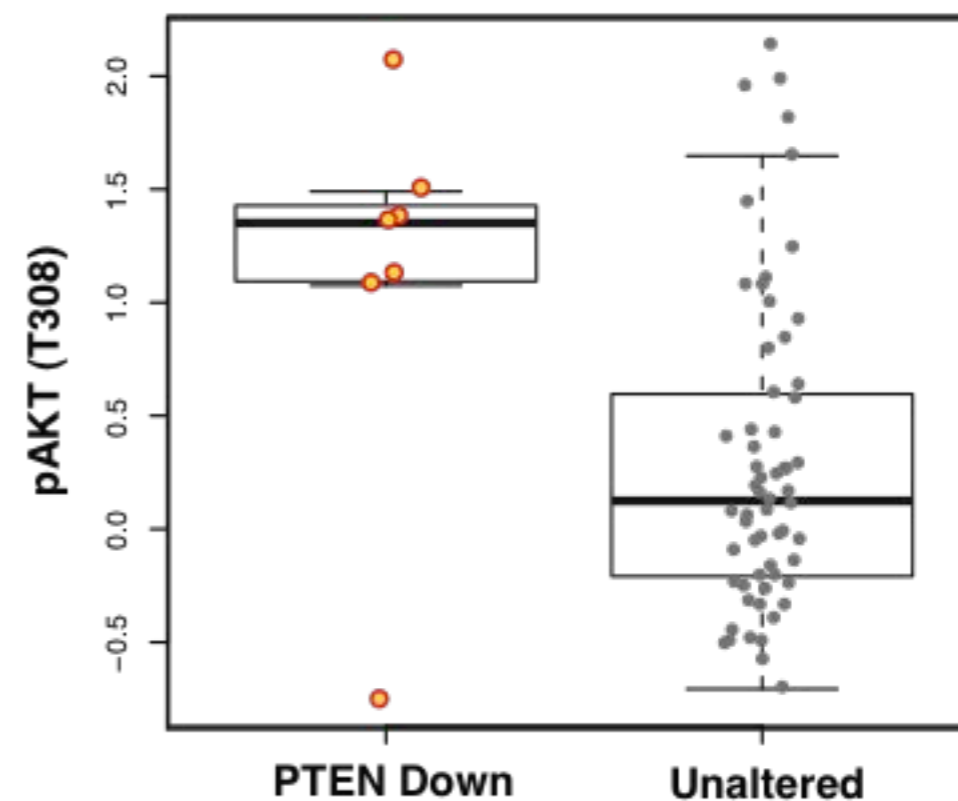
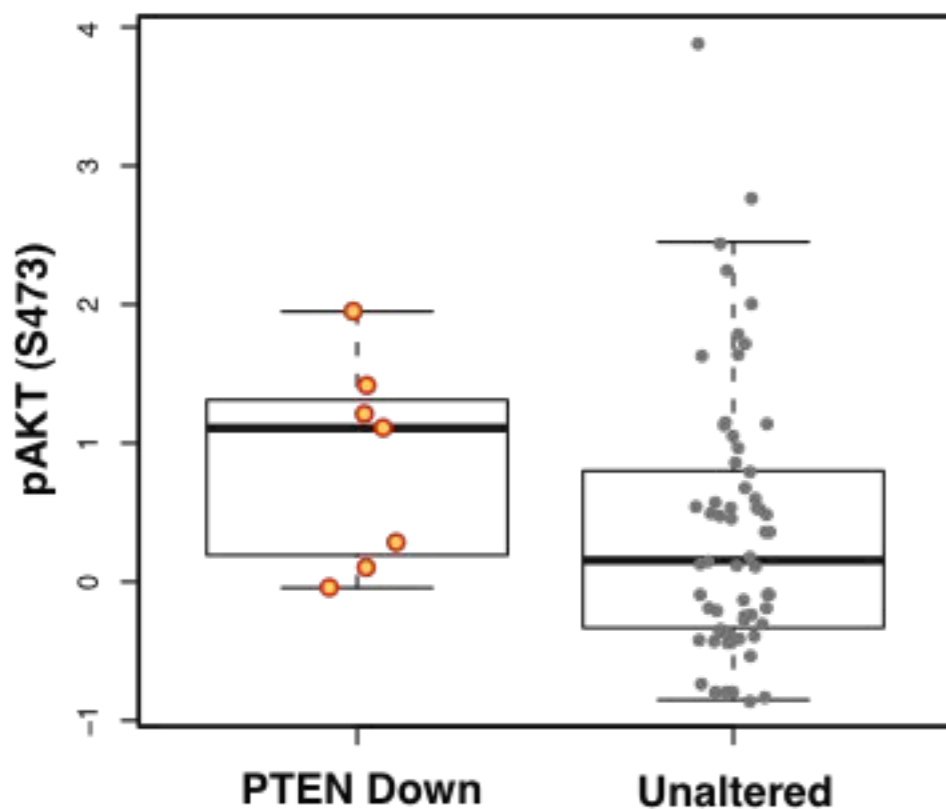
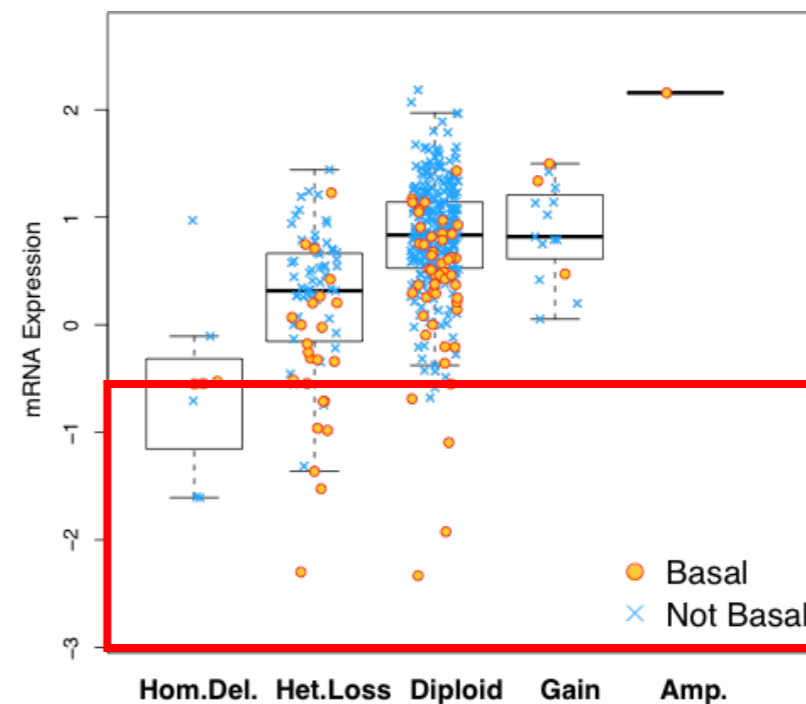
**10% of Basal Tumors**



# PTEN down-regulation activates Akt

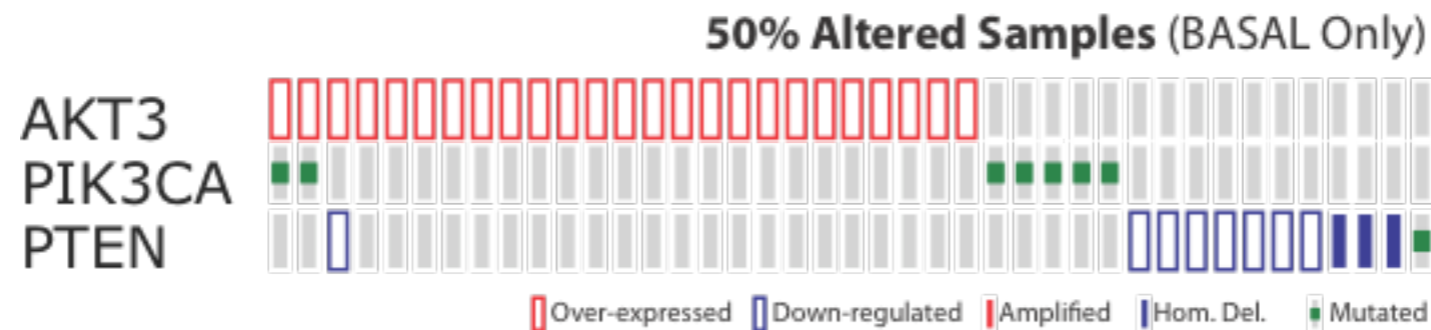
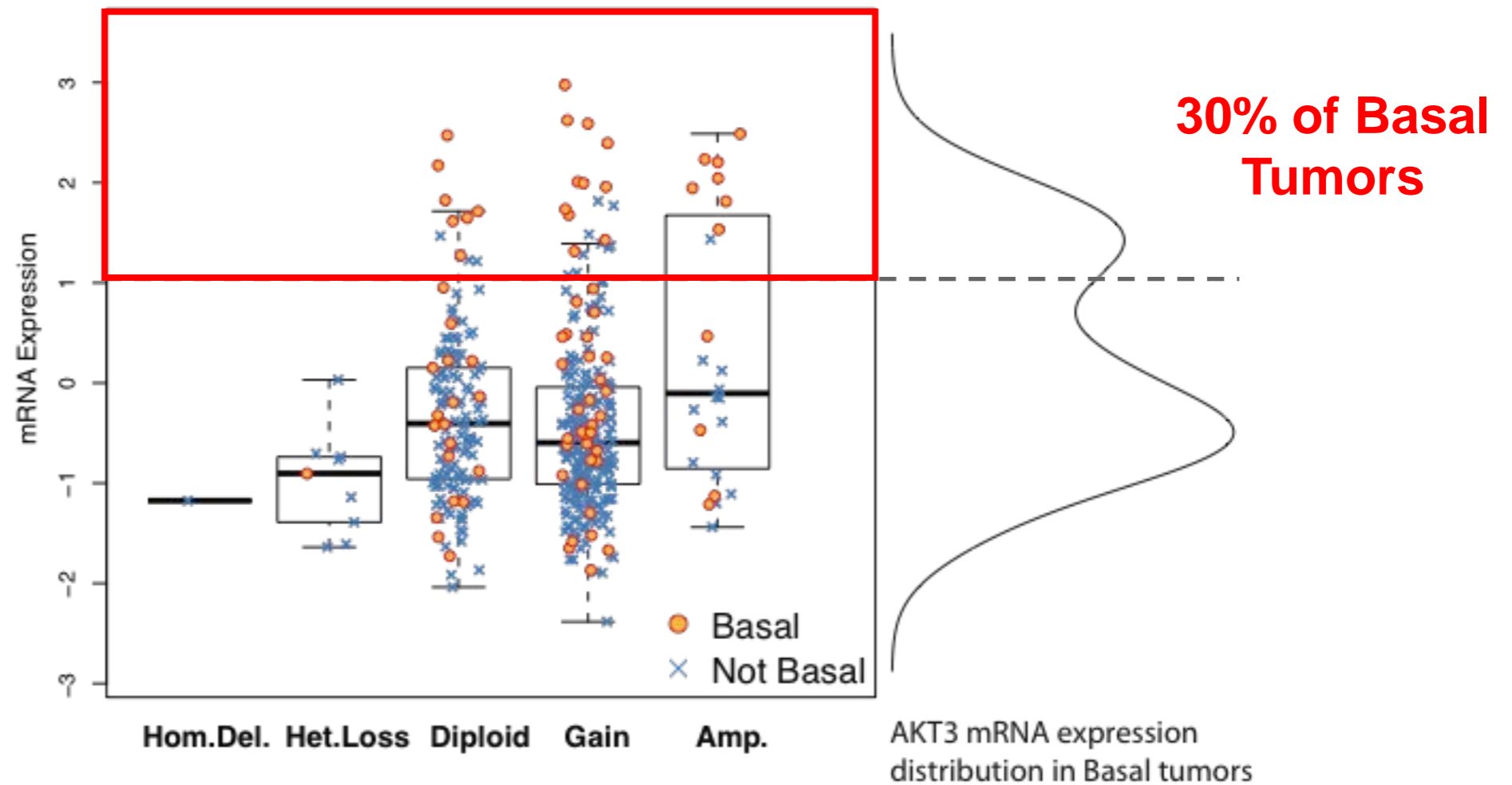
- PTEN is down-regulated in Basal tumors
- Down-regulated samples show higher Akt phosphorylation

10% of Basal Tumors



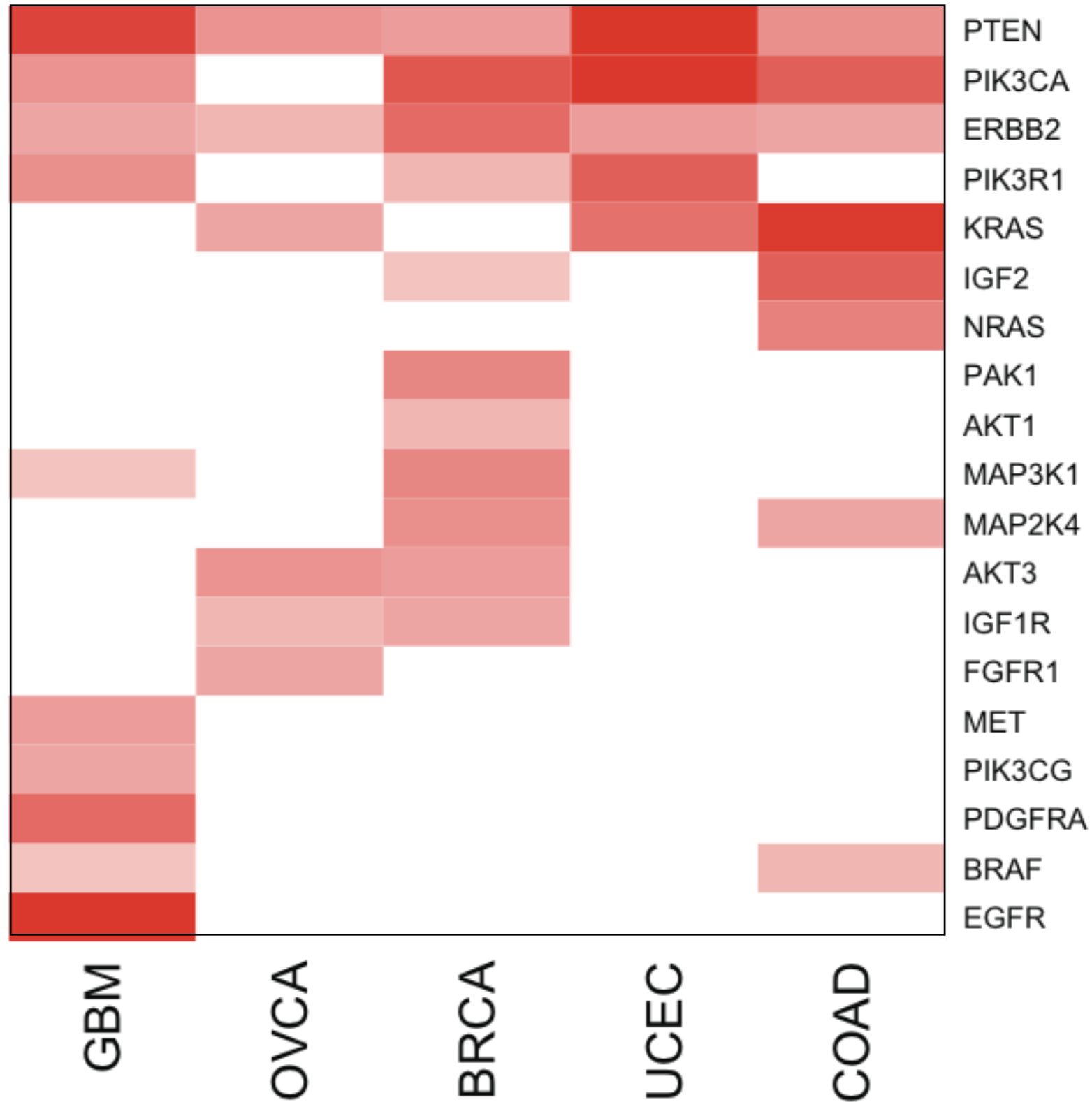
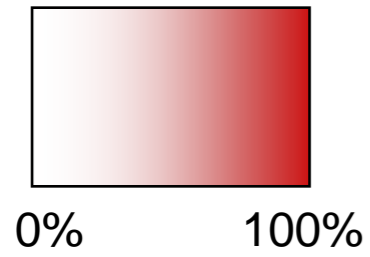
# AKT3 is over-expressed in Basal Breast Cancer

## AKT3 in Basal Breast Cancer





# Overall Extent of Alteration





# Conclusions

- **MEMo** systematically identifies **mutually exclusive** alterations targeting oncogenic pathways across multiple cancer types
- **PI(3)K /Akt** signaling is consistently altered in cancers, with different **extents of alteration**, and by **different mechanisms**
- Mutual exclusivity analysis **across multiple cancers** unveils the underlying heterogeneity of the disease, thus suggesting candidate **therapeutic targets in different subtypes**



# Thanks!

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... and everyone at cBio!

