

Society for Immunotherapy of Cancer

# Mutation Drivers of Immunological Responses to Cancer

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La Caixa Junior Leader Fellow

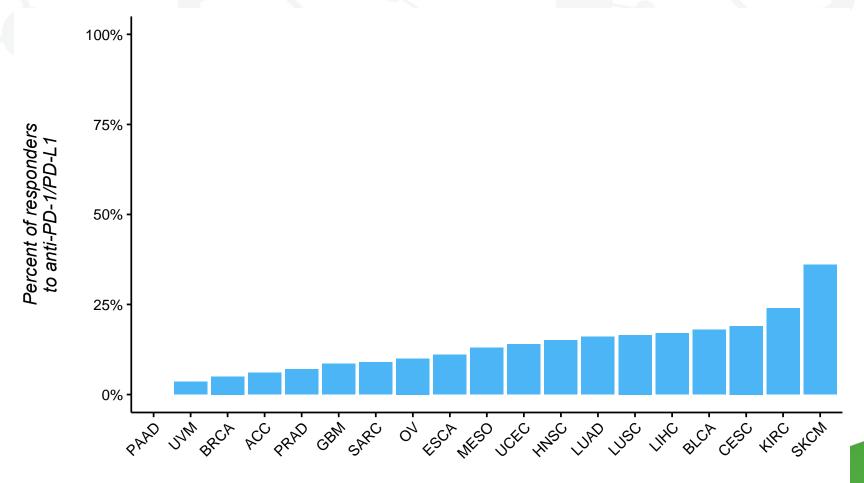
Barcelona Supercomputing Center (BSC)

# Financial conflicts of interest

None

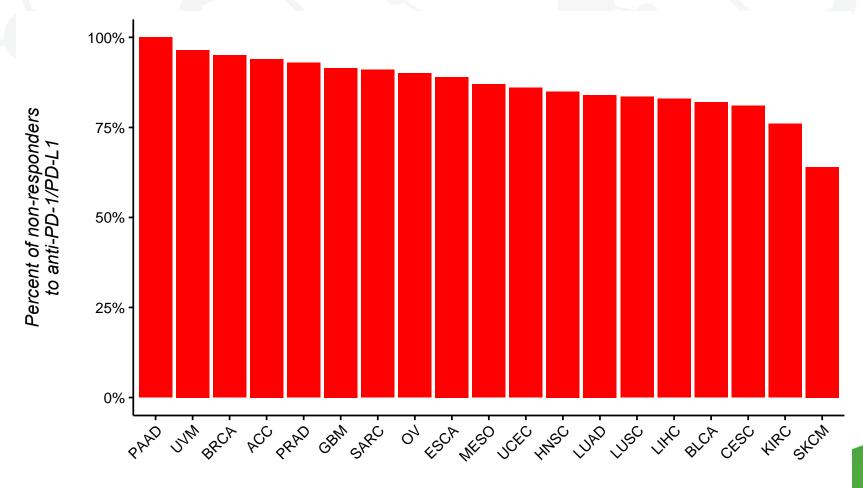


# **Exciting times in cancer immunology**



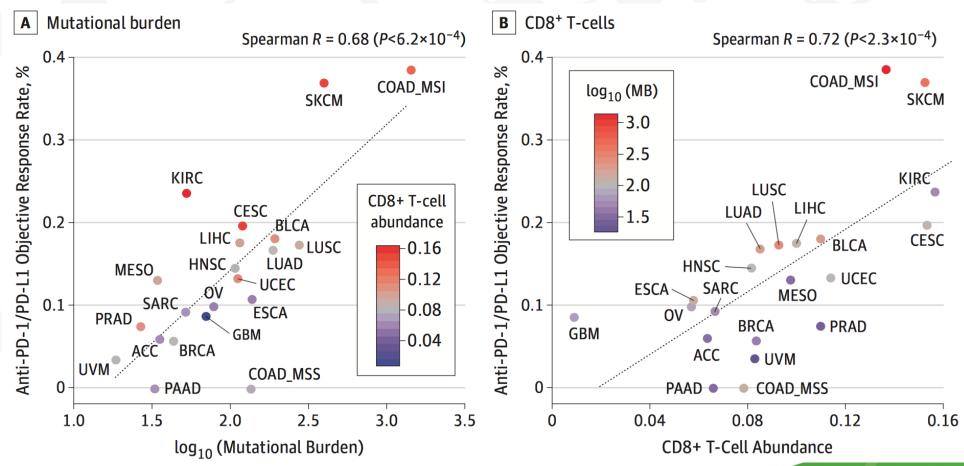


#### ... but there's still a lot to do!



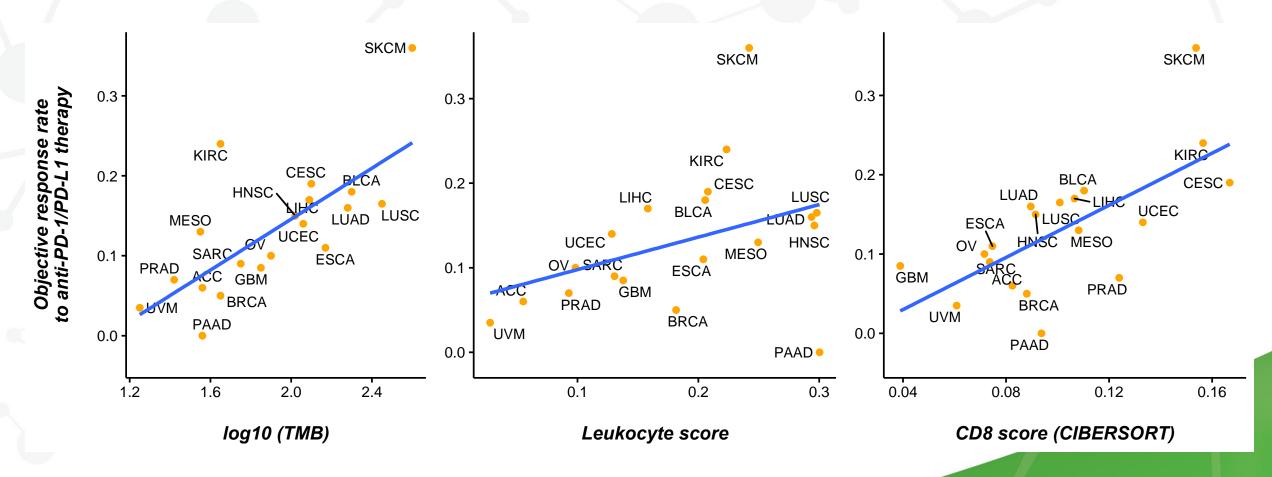


#### Predicting success of immune-based therapies





#### This is also true with other immune scores

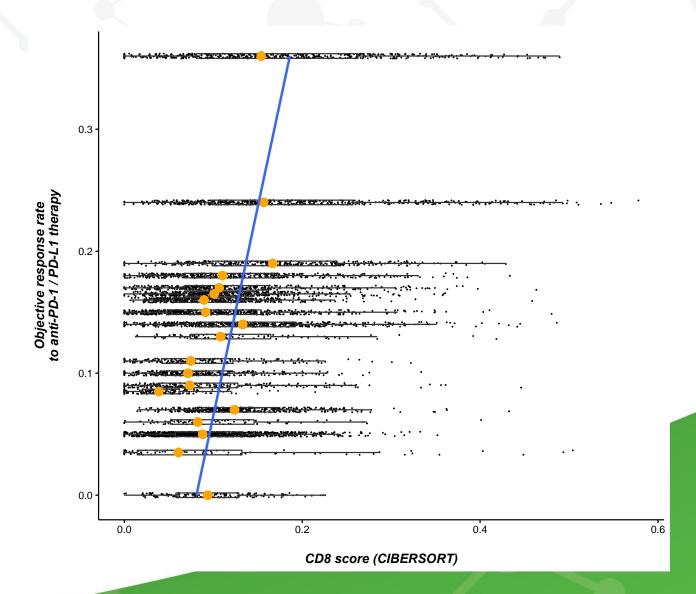




# Medians hide an immense variability

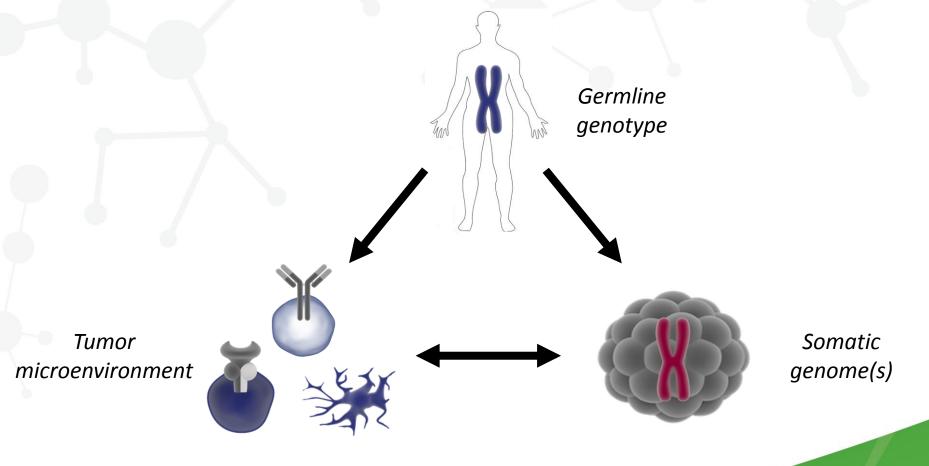
Each black dot is a TCGA sample.

Orange dots represent the median of CD8 score for each cancer type



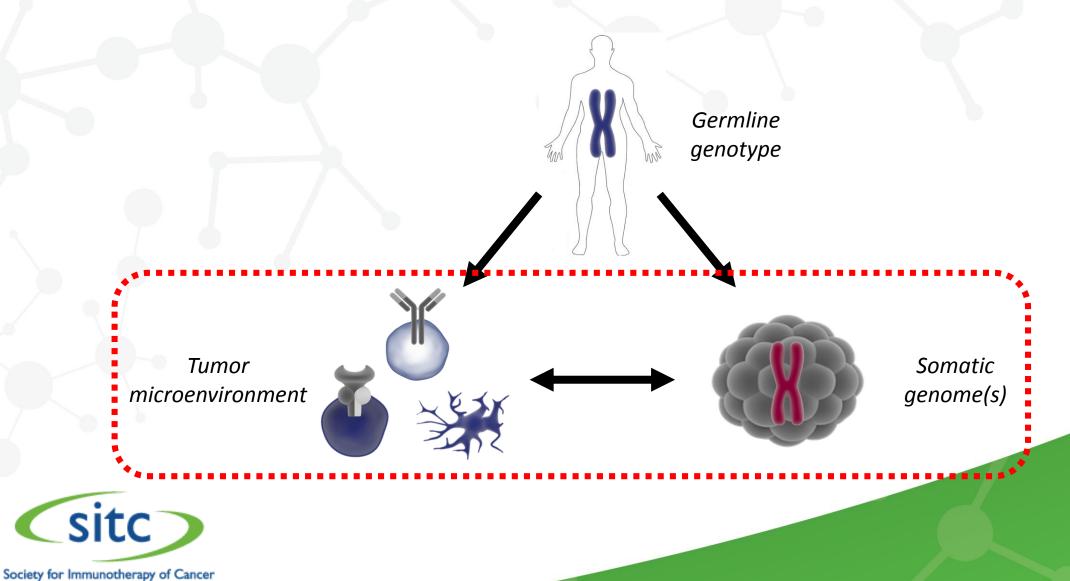


# This variability stems from the "trialogue"





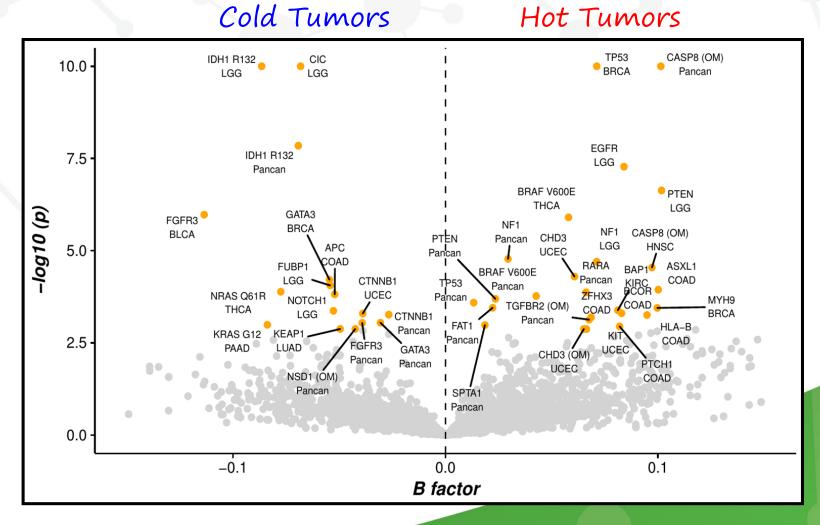
# This variability stems from the "trialogue"



#### **Known driver – TME interactions**

Correlation between the presence of driver mutations in TCGA samples and the leukocyte score (methylation)

Positive B values indicate more leukocytes and viceversa





Rooney et al., Cell, 2015 Tamborero et. al., Clin Can Res, 2018 Thorsson et. al., Immunity, 2018

# **Open questions**



Victoria Ruiz-Serra

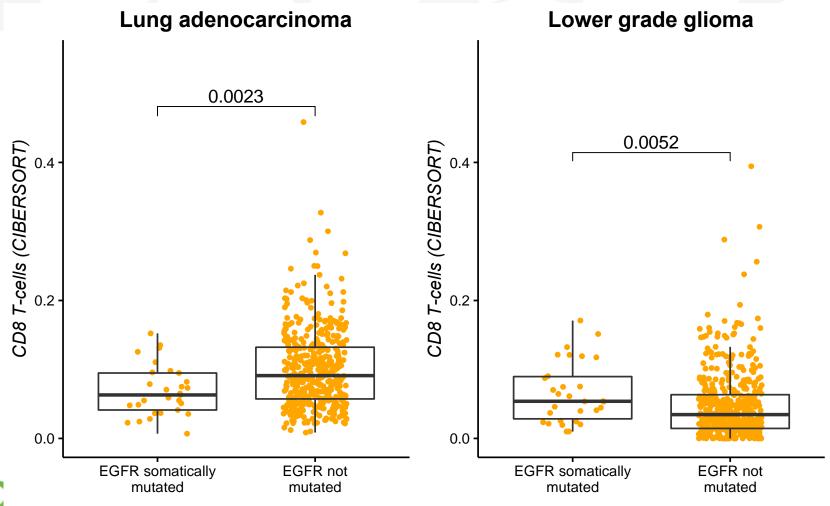
- 1 Do mutations in cancer driver genes always correlate in the same direction?
- 2 Do all mutations in cancer driver genes correlate similarly with the immune component of the TME?
- 3 Do different mutations interact with each other?



### Correlations might depend on the tissue



Victoria Ruiz-Serra





# **Open questions**

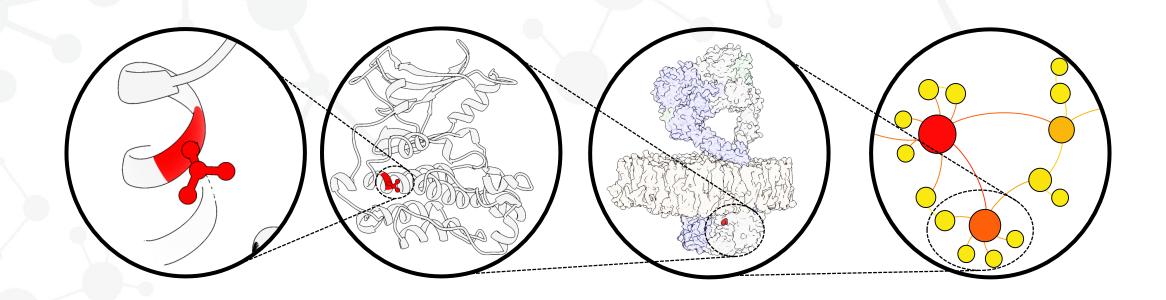


Victoria Ruiz-Serra

- 1 Do mutations in cancer driver genes always correlate in the same direction? **They might!**
- 2 Do all mutations in cancer driver genes correlate similarly with the immune component of the TME?
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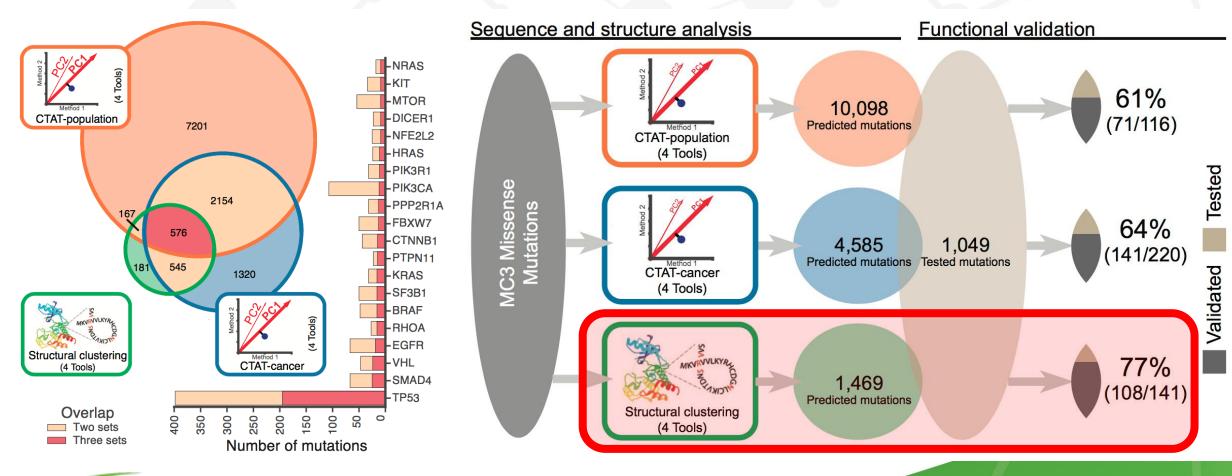


# Perspective is important





# Not all driver mutations are equal





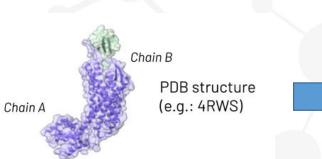
# Structure – guided analysis

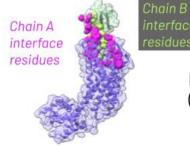
>200.000 PDB files analyzed

~100.000 have >50% sequence identity with human protein AND potential interface

Catalogue with ~150.000 interfaces (protein, ligands or nucleic acids)





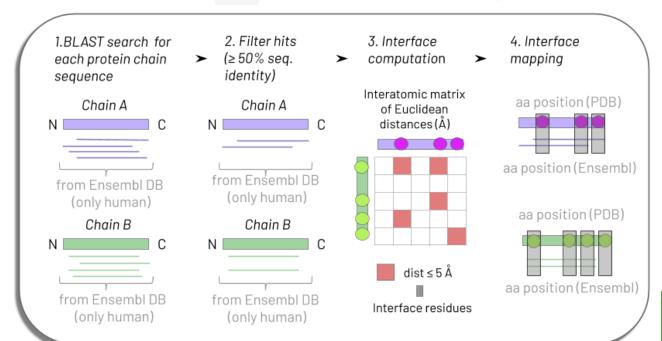


interface residues

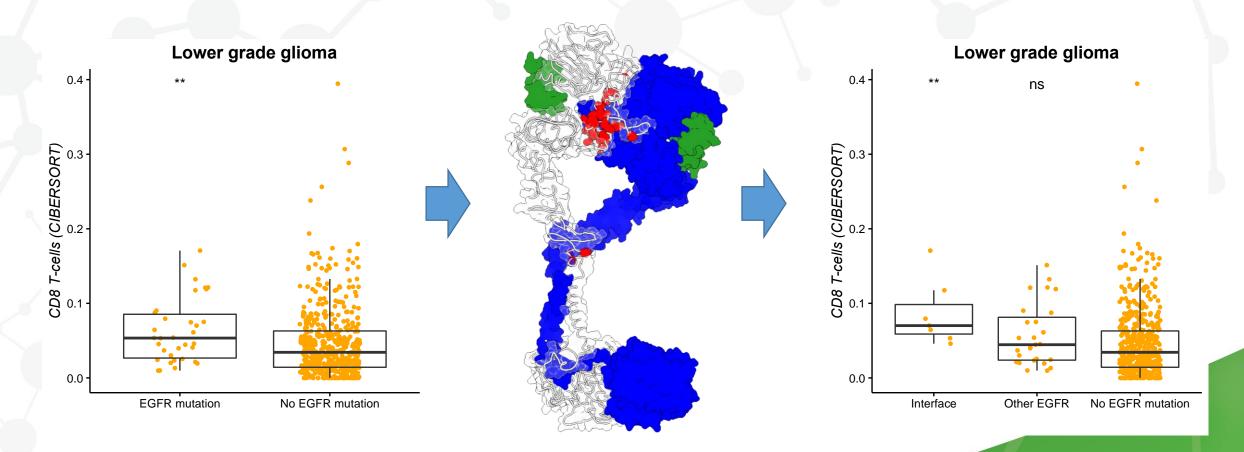
PDB structure (e.g.: 4RWS)



Victoria Ruiz-Serra



# Structure analysis of EGFR - LGG



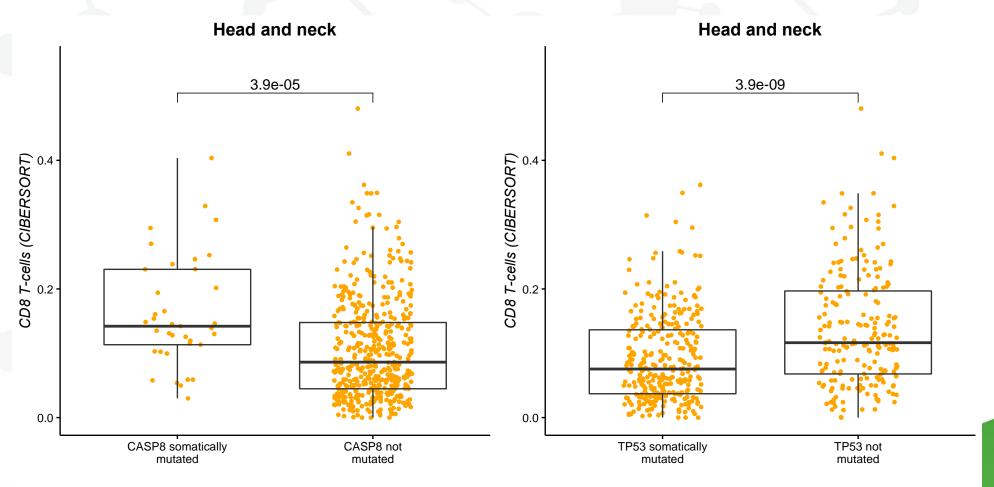


### **Open questions**

- 1 Do mutations in cancer driver genes always correlate in the same direction? **They might!**
- 2 Do all mutations in cancer driver genes correlate similarly with the immune component of the TME? **No!**
- 3 How do different mutations interact with each other?

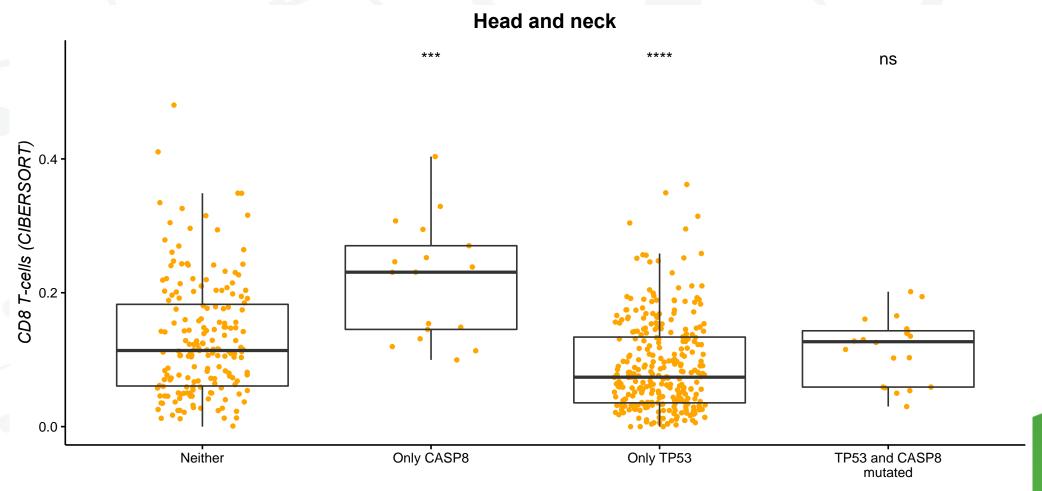


#### Different drivers can have different effects...





# ...and they can interact with each other!





### **Open questions**



Victoria Ruiz-Serra

- 1 Do mutations in cancer driver genes always correlate in the same direction? **They might!**
- 2 Do all mutations in cancer driver genes correlate similarly with the immune component of the TME? **No!**
- 3 Do different mutations interact with each other? They can!



# Thank you!

#### **Collaborators**



Victoria Ruiz-Serra
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Valencia's lab

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Germline Immune working group

#### **Funding Agencies**





