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# Driver Kinase Fusions in Cancer

TCGA 4<sup>th</sup> Annual Scientific Symposium – May 12<sup>th</sup>, 2015  
Nicolas Stransky, PhD

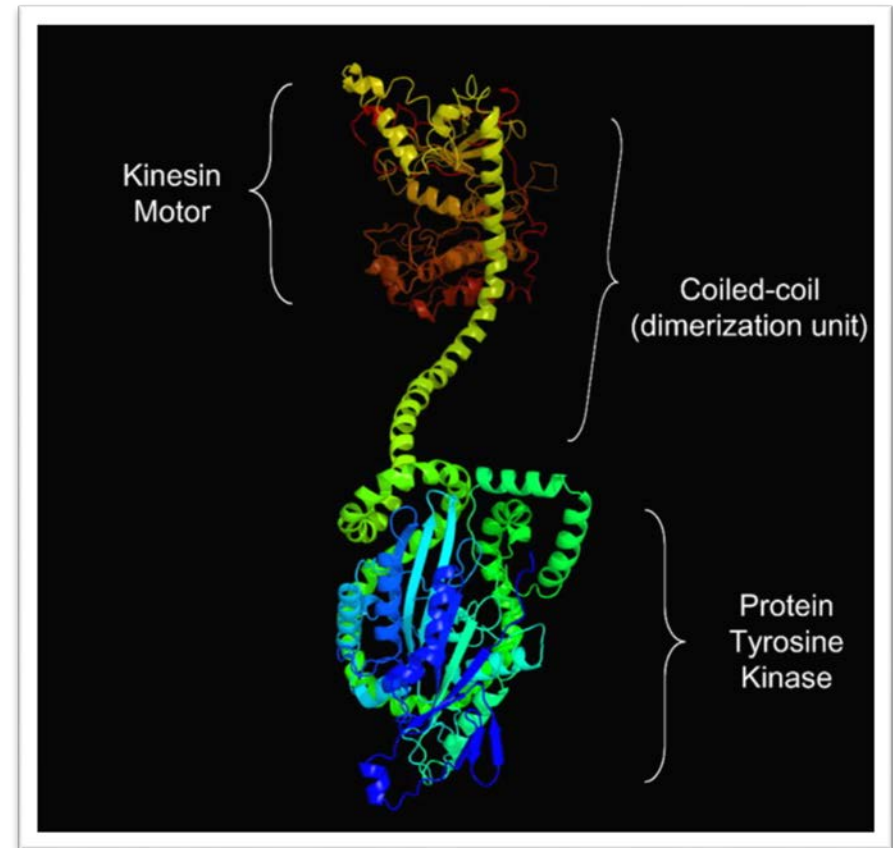


# What are Kinase Fusions?



- Genomic instability, a hallmark of cancer, can result in chromosomal translocations or other complex rearrangements
- These events can produce chimeric genes called “fusions”
- Known driver kinase events include *BCR-ABL1* in CML, *EML4-ALK* in Lung adenocarcinoma

## *KIF5B-RET* Fusion



Ju Y S *et al.* Genome Res. 2012

# May 2015:

## >10,000 RNAseq Samples in TCGA, 33 Tumor Types



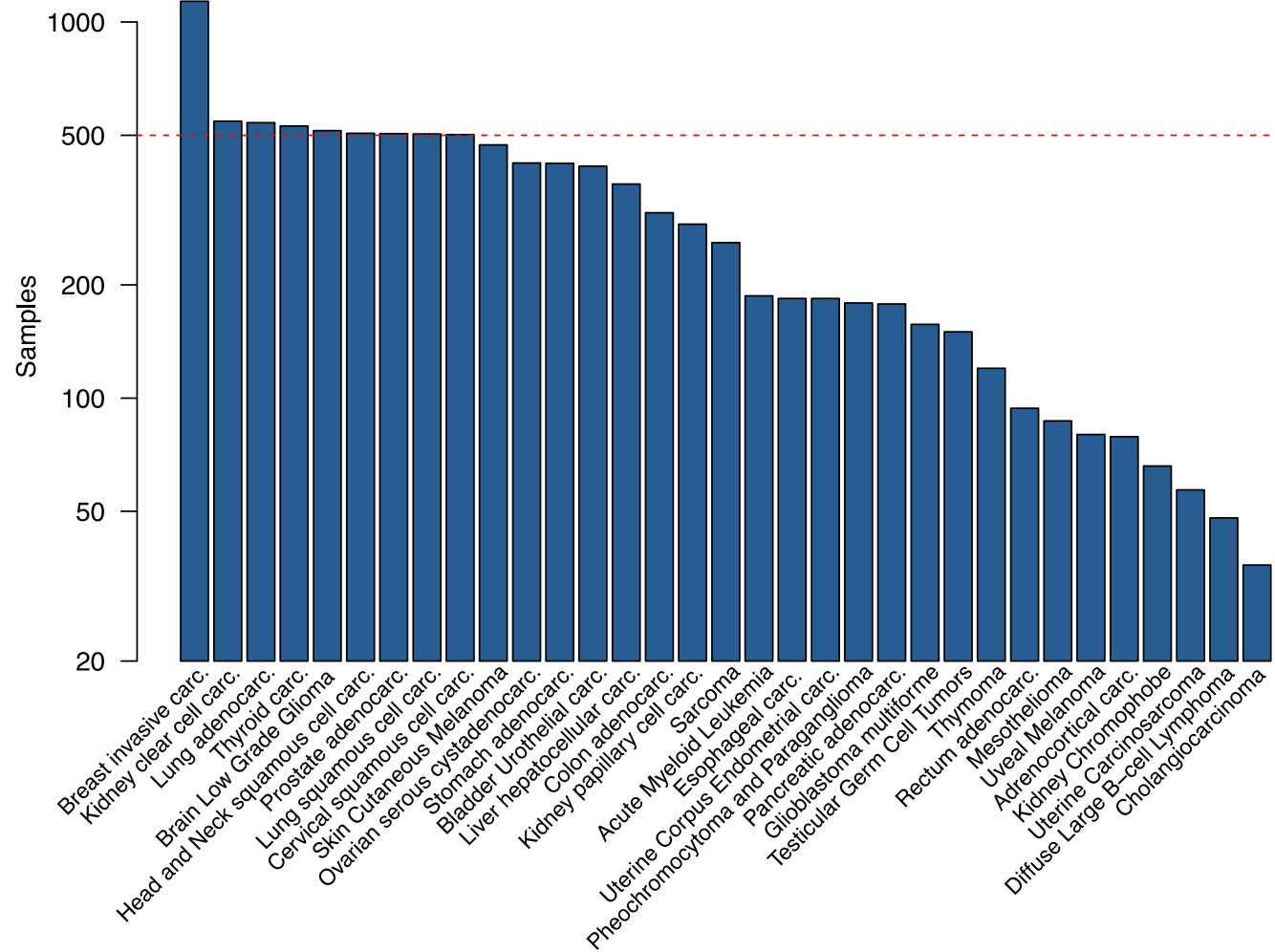
TCGA RNA-seq data  
for ~10,000 tumors



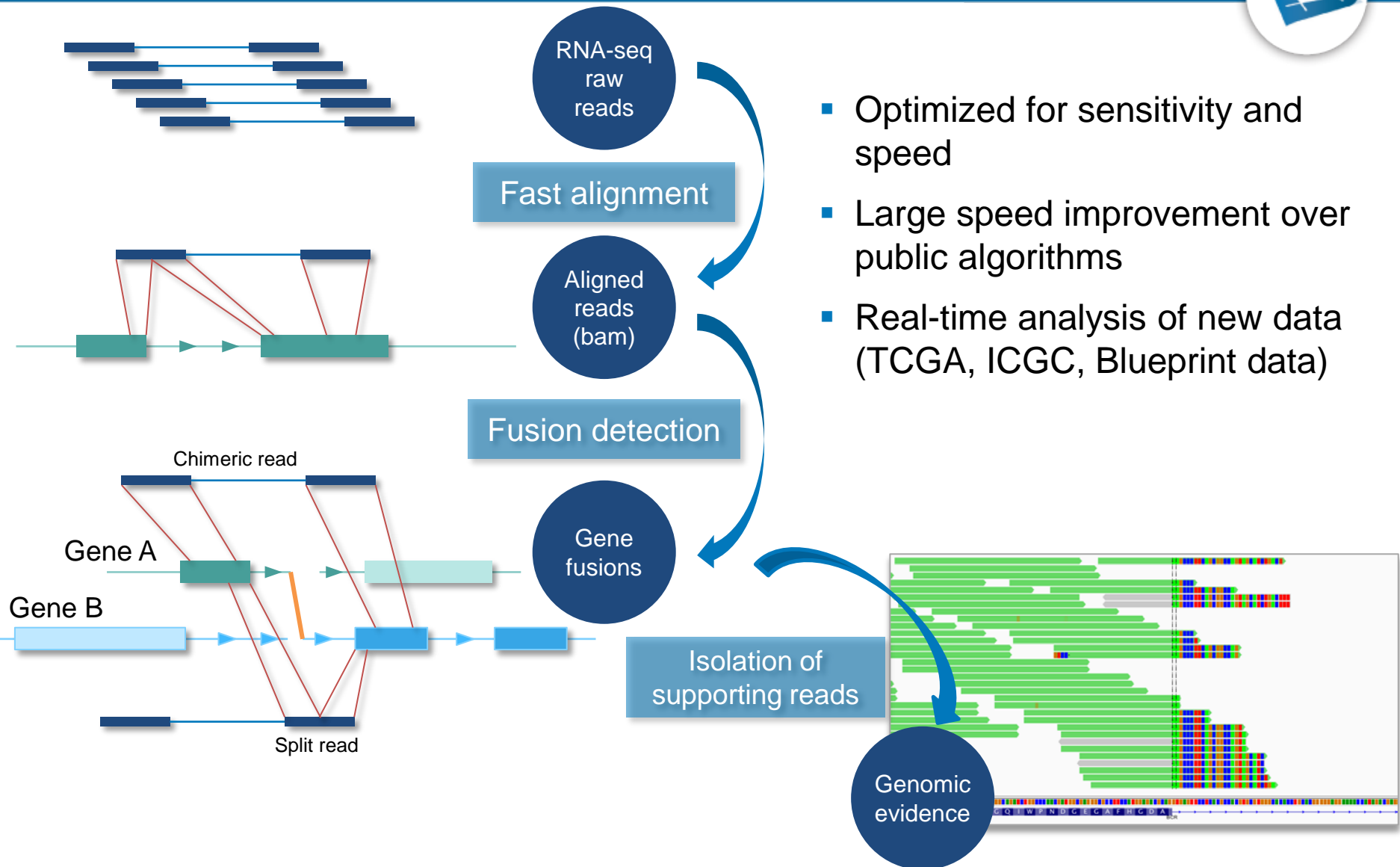
Fusion finding  
algorithm



First pan-cancer  
evaluation of fusions



# Novel Algorithm for Rapid Kinase Gene Fusion Detection



# Computational Pipeline for Fusion Detection



Fusion Detection

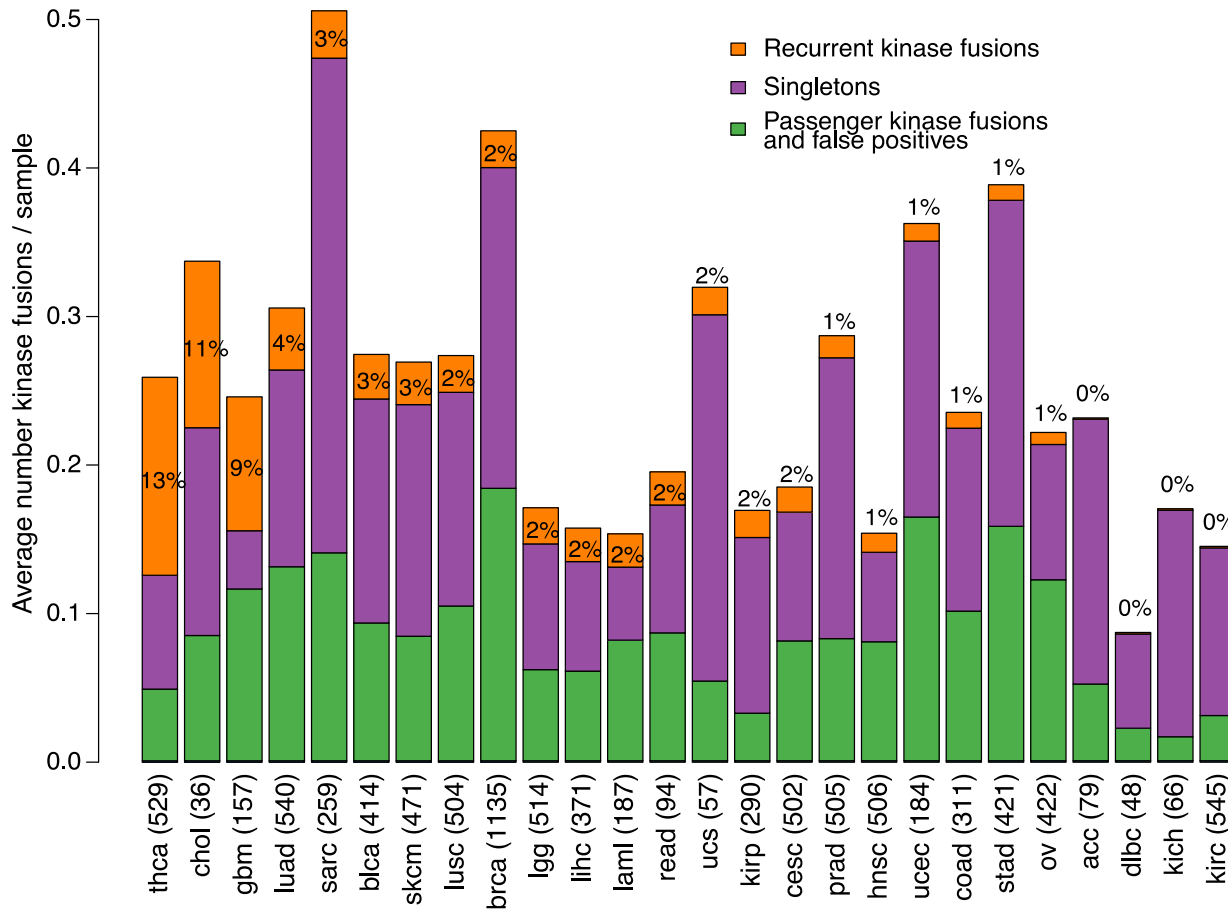
Post-processing

 Report  
&  
Annotate

- Core algorithm
  - Identifies gene-gene fusions in RNA-seq data
  - Reports supporting evidence for each fusion
- Post-processing
  - Heuristics to filter out passenger events
    - Intergenic junctions (between two exons)
    - Coding sequence in frame
    - **Presence of kinase catalytic domain**
  - Heuristics to filter out false-positives
    - Fusions present in normal
    - Alignment artifacts (repetitive sequences)
    - High expression level of one partner
- Reporting tools
  - Reporting of pipeline outputs, fusion frequencies
  - Manually review and annotate fusions

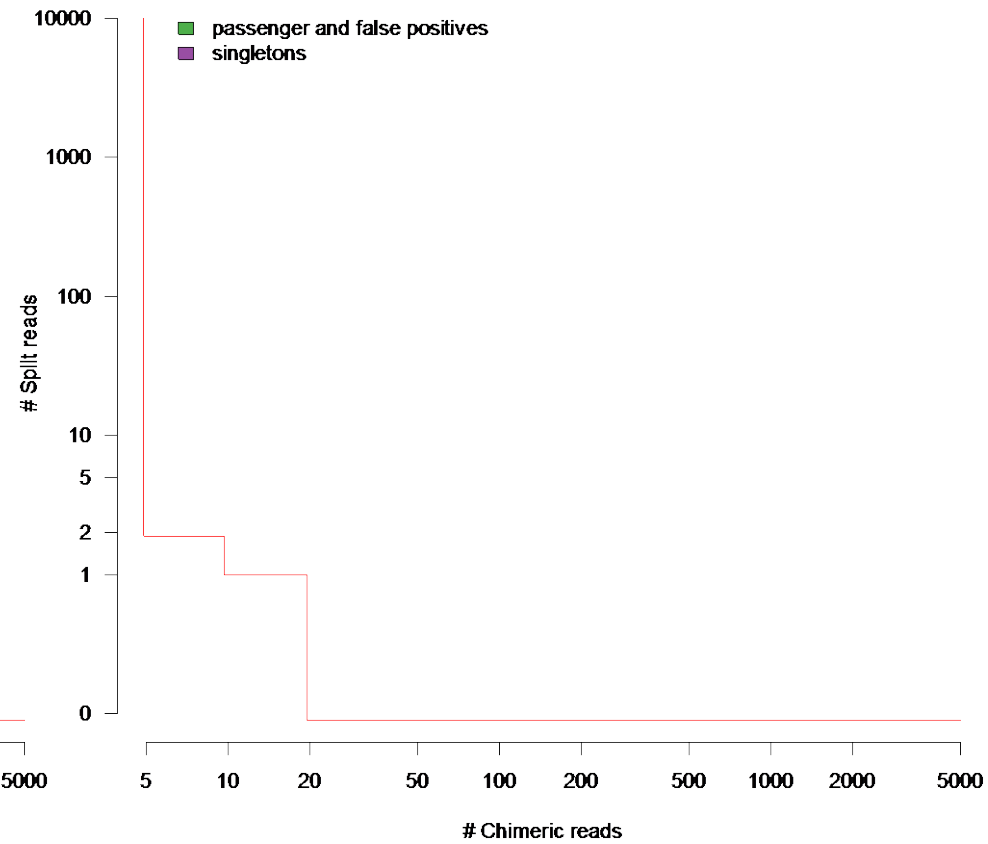
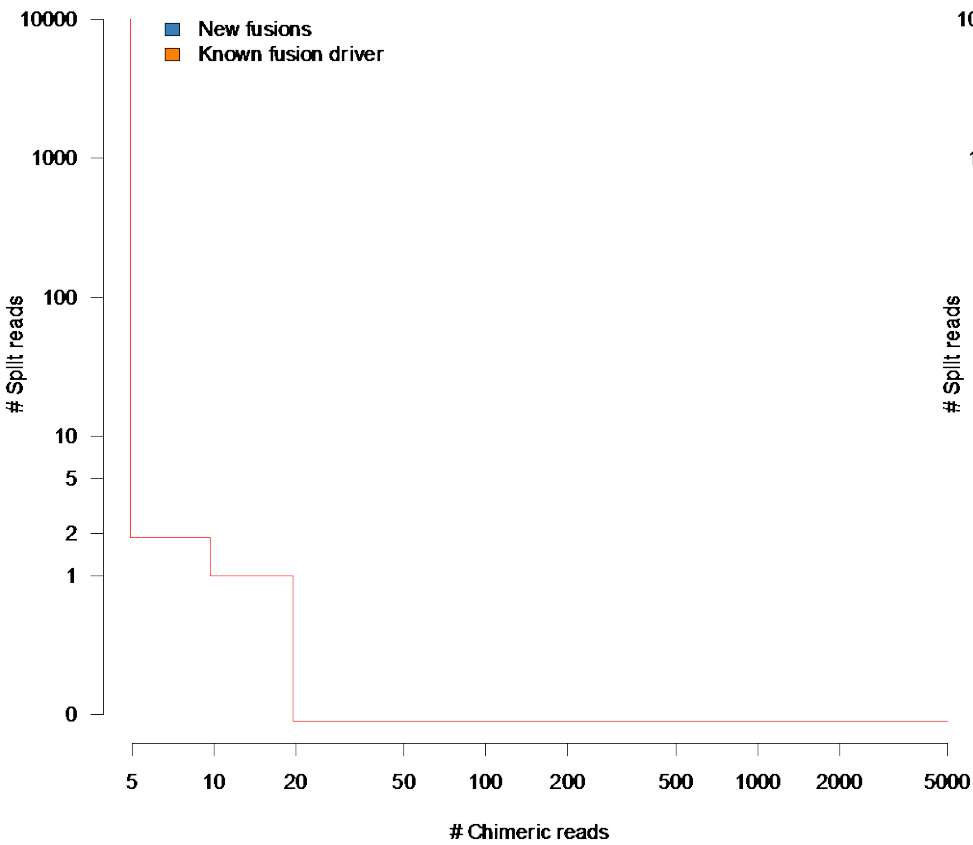
Therapeutic  
relevance

# Pipeline output: kinase fusions after manual review

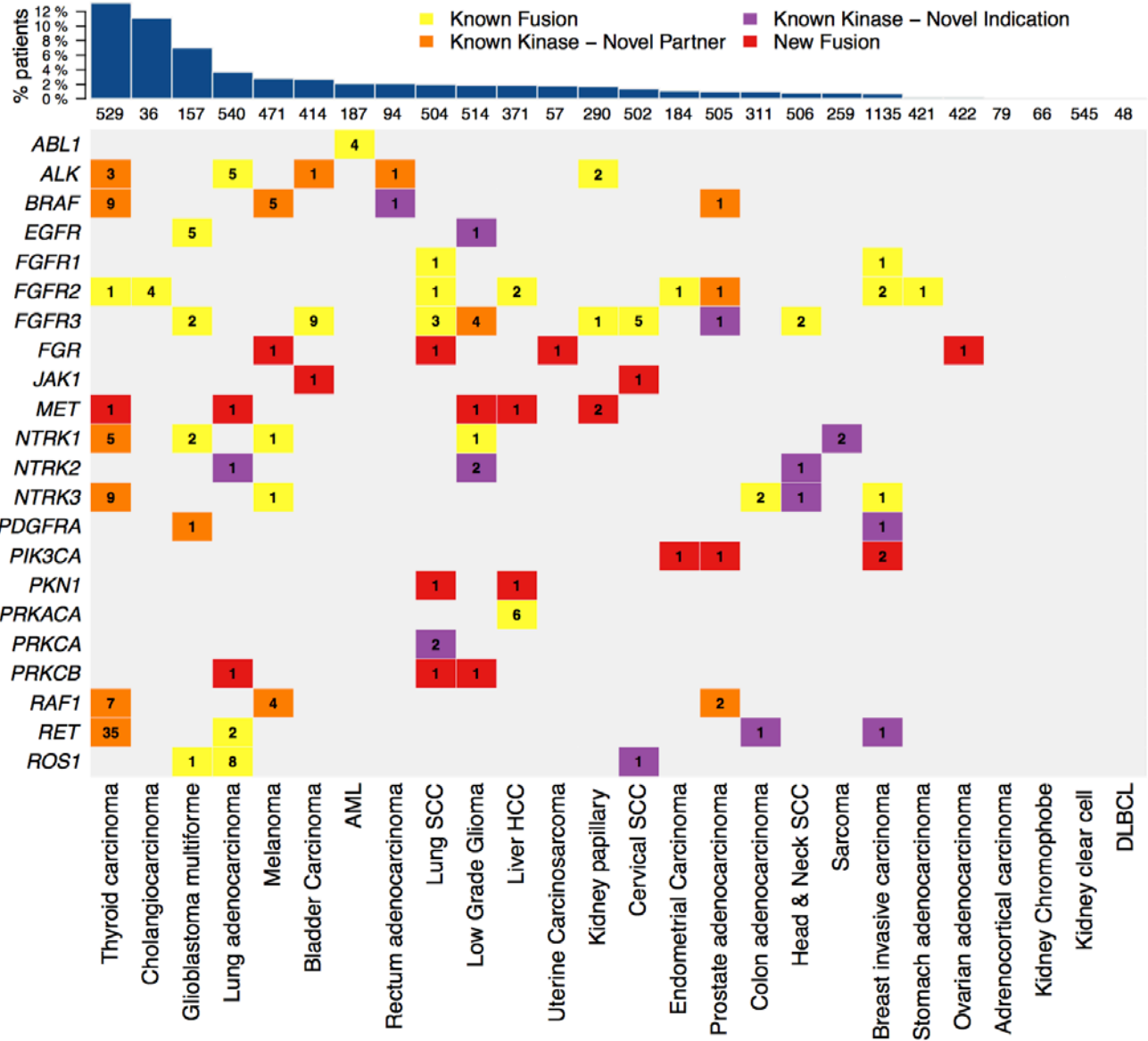


- 2.8 % of tumor samples contain a likely oncogenic kinase fusion (2.0 % excluding thyroid cancer)
- Thyroid cancer, sarcoma and glioblastoma have the highest proportion of recurrent kinase fusions
- Kidney clear cell and kidney chromophobe have almost no kinase fusions

# Genomic evidence for novel kinase fusion events



# The Landscape of Kinase Fusions in Cancer



New Indications  
and New Gene  
Partners

Novel Recurrent  
Kinase Fusions

Adapted from Stransky *et al.*  
Nature Communications, 2014

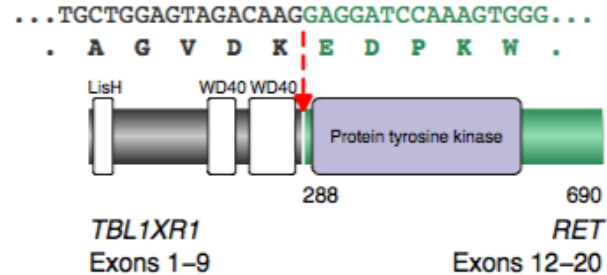


# Novel partners and novel indications for kinase fusions

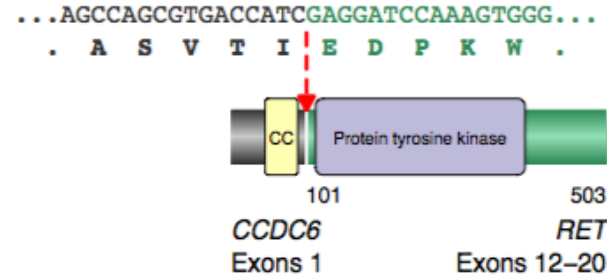


## RET

participant ID: 47ad3d2e-9096-4e6d-884f-83507fd8e970  
cancer type: thyroid carcinoma

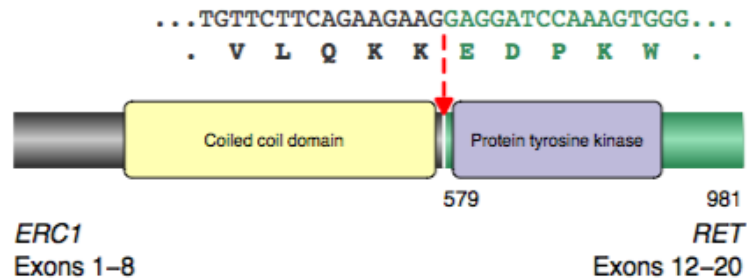


participant ID: c8d7eb0f-2502-4277-a8ef-435b08f3c863  
cancer type: colon adenocarcinoma



Known partners

participant ID: 8a209c68-bef9-4191-b6ec-7a1a9258b3bd  
cancer type: breast invasive carcinoma

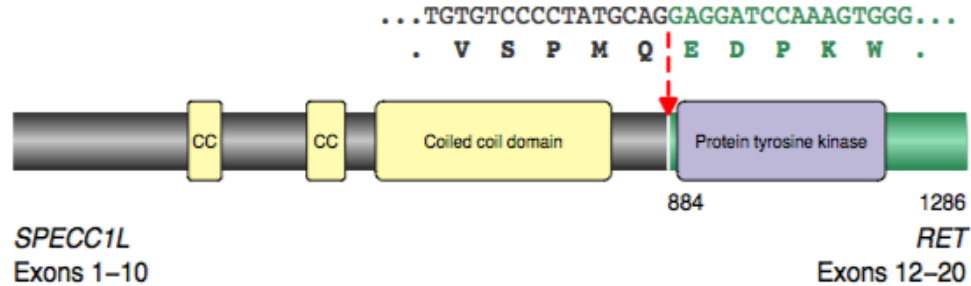


# Novel partners and novel indications for kinase fusions

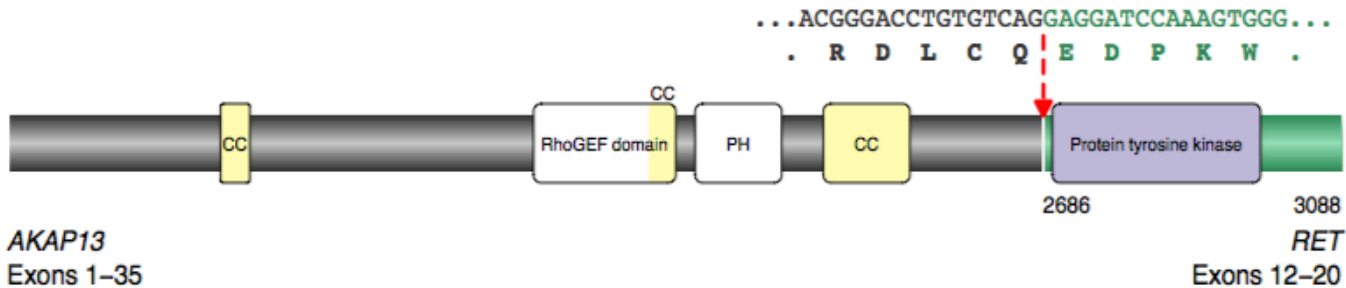


## RET

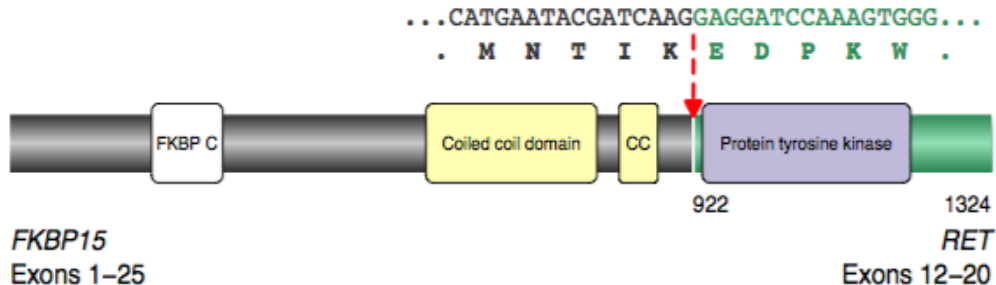
participant ID: 3d114411-0215-4870-87d6-7ff08ca8f771  
cancer type: thyroid carcinoma



participant ID: 494fec2d-a438-4b21-903c-6ef07fb08fe4  
cancer type: thyroid carcinoma



participant ID: 449ee008-36eb-4cb8-87f7-be76a568481f  
cancer type: thyroid carcinoma

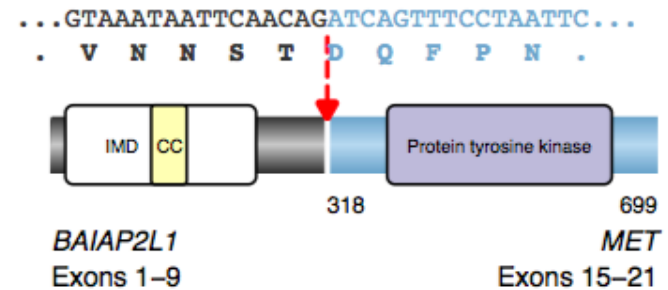
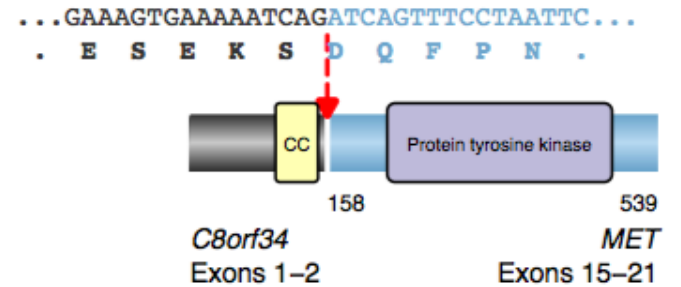
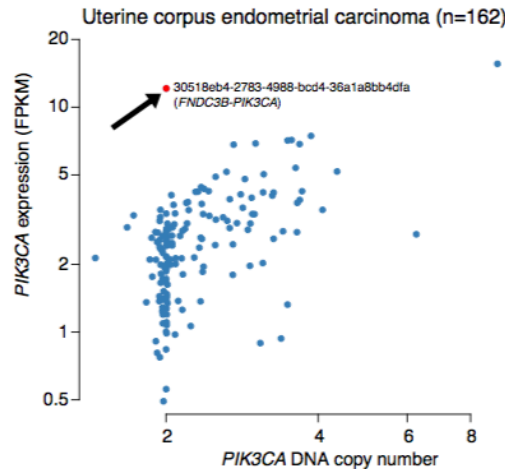
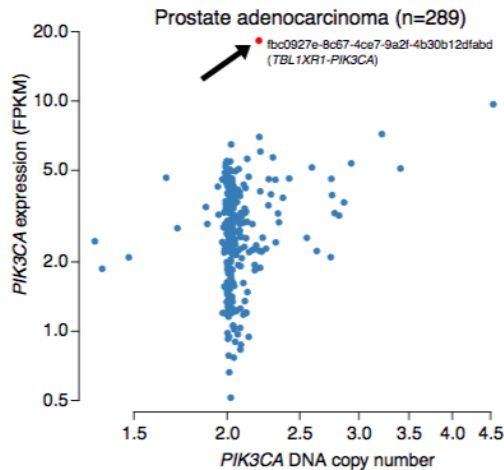
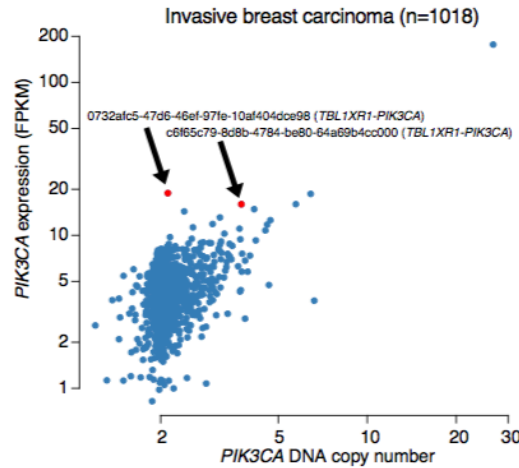
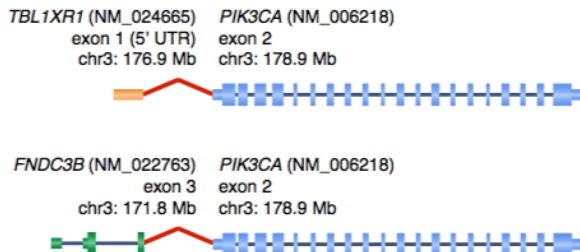


Novel Partners, all with dimerization motifs

# Novel *MET* and *PIK3CA* Fusions



- MET* and *PIK3CA* fusions occur in solid tumors where mutations and amplifications are already driver events

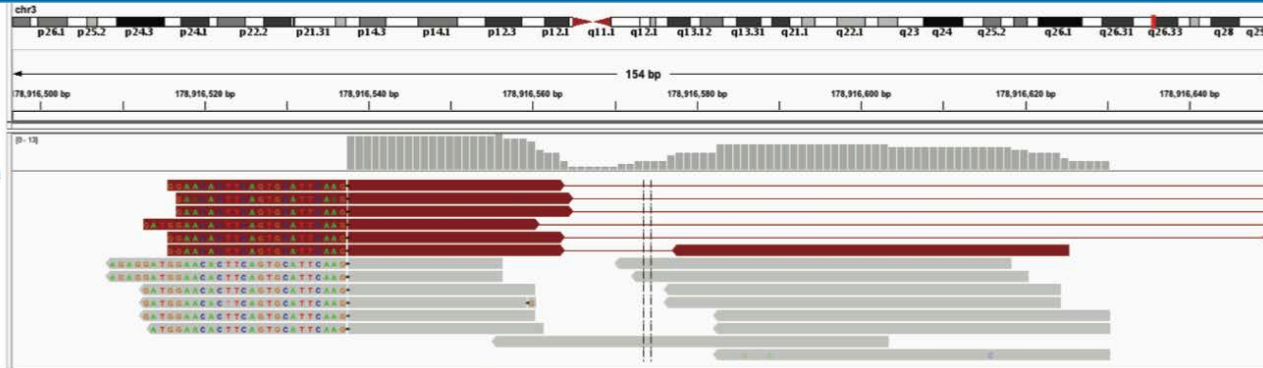


*MET* fusions in kidney papillary cell carcinoma

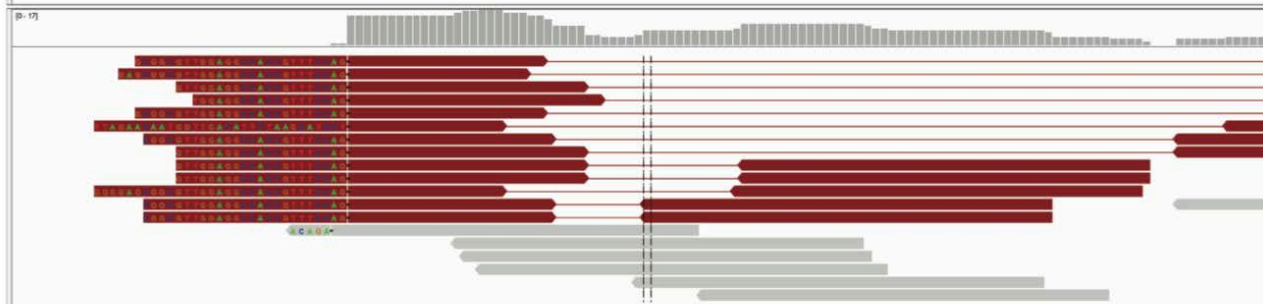
# Novel *PIK3CA* fusions – supporting reads



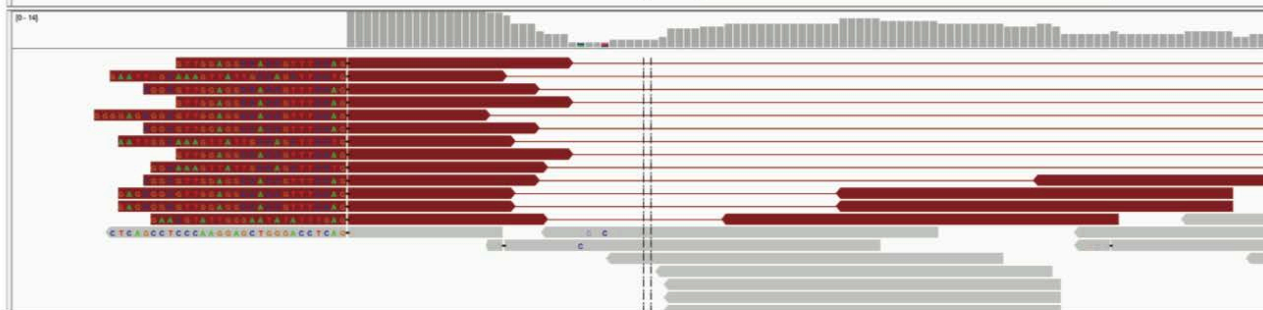
FNDC3B : *PIK3CA*  
sample: 30518eb4-2783-4988-bcd4-36a1a8bb4dfa  
Cancer type: Uterine Corpus Endometrial Carcinoma



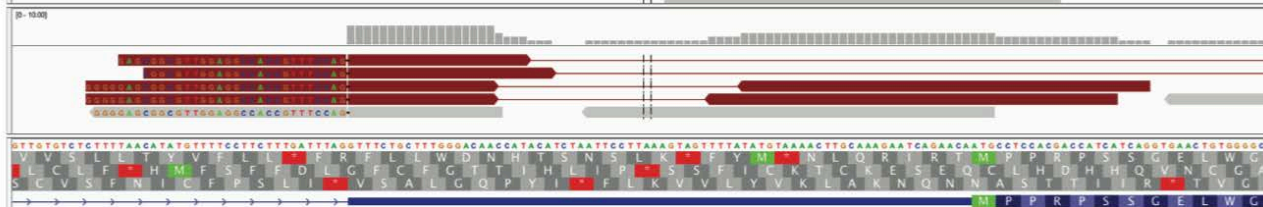
TBL1XR1 : *PIK3CA*  
sample: 0732afc5-47d6-46ef-97fe-10af404dce98  
Cancer type: Breast invasive carcinoma



TBL1XR1 : *PIK3CA*  
sample: fbc927e-8c67-4ce7-9a2f-4b30b12dfabd  
Cancer type: Prostate adenocarcinoma



TBL1XR1 : *PIK3CA*  
sample: c6f65c79-8d8b-4784-be80-64a69b4cc000  
Cancer type: Breast invasive carcinoma



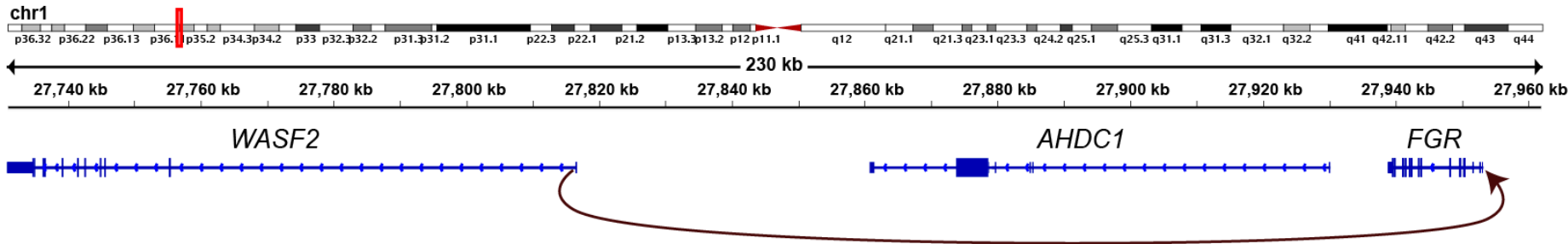
**PIK3CA**

# New *WASF2-FGR* 5'-UTR Fusions



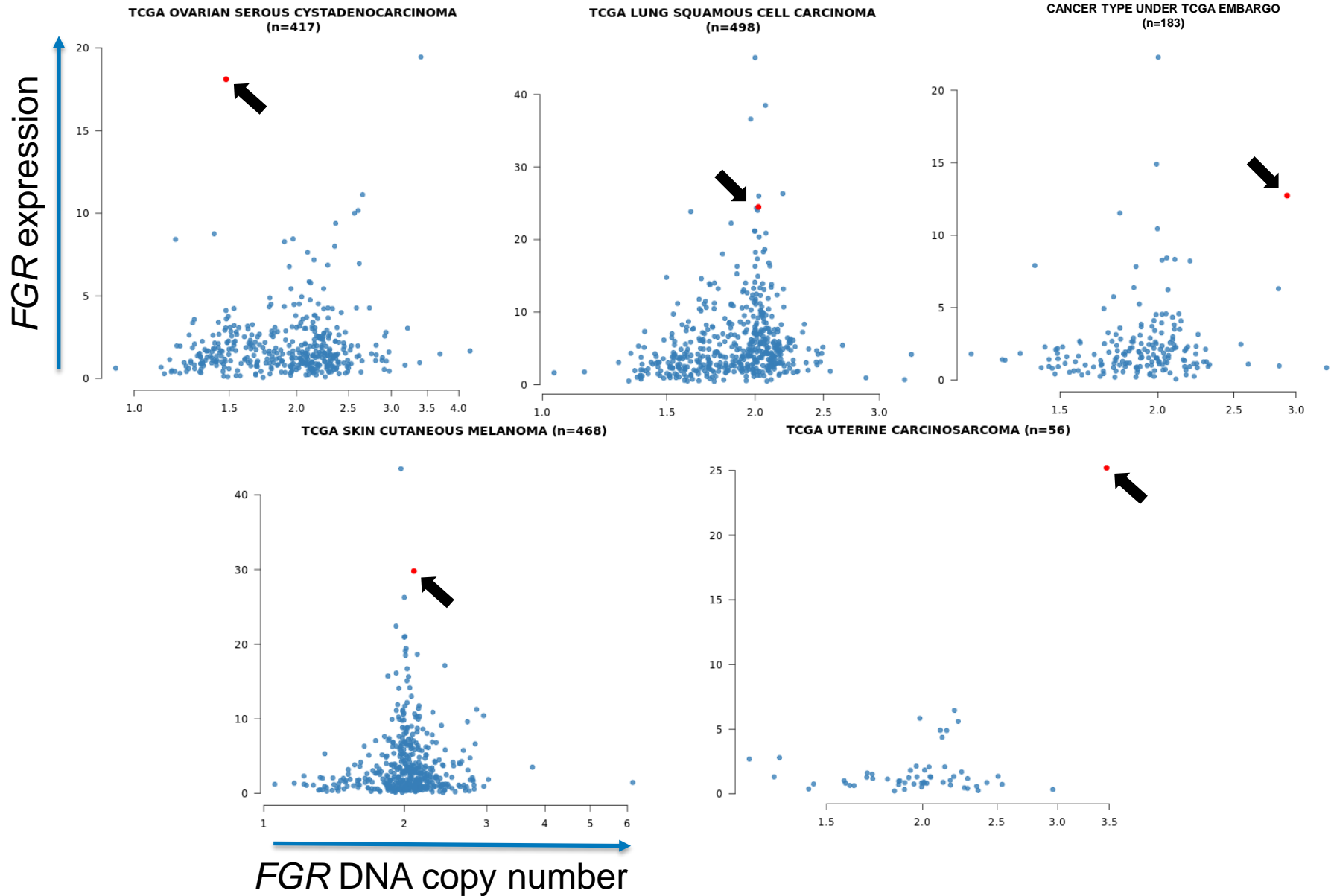
*WASF2* (NM\_006990)  
exon 1 (5' UTR)  
chr1: 27.8 Mb

*FGR* (NM\_005248)  
exon 2  
chr1: 27.9 Mb



- Src family kinase
- Highly expressed in some hematopoietic cells and malignancies
- Oncogenic potential - viral oncogene homolog
- A new promoter fusion not previously implicated in cancer

# New *WASF2-FGR* 5'-UTR Fusions



# NTRK1/2/3 Fusions



- Certain fusions are very recurrent across tumors
  - 9/26 tumor types with *NTRK1/2/3* fusions for a total of 29 fusions
  - Additional recurrent fusions exist in other cancers under embargo

■ Known Fusion      ■ Known Kinase – Novel Indication  
■ Known Kinase – Novel Partner      ■ New Fusion





## New insights into the kinase fusion “landscape”

- 6 additional TCGA cancer types surveyed
- 10% of *FGFR2* fusions in cholangiocarcinoma
- Novel *ALG14-JAK1* fusions
- 2 new *FGR* fusions in solid tumors
- New pan-cancer *NTRK1/2/3* fusions
- *PRKACA* fusions in Liver cancer (FL-HCC)

## Key Takeaways

- First pan-cancer fusion analysis
- New fusion analysis framework, designed with speed and sensitivity in mind
- Focus on kinase fusions as driver events
- Profound implications for diagnosis, patient treatment and drug discovery





- **The Cancer Genome Atlas**
  
- **Blueprint Fusions team**
  - Andy Garner
  - Christoph Lengauer
  - Ethan Cerami
  - Joseph Kim
  - Klaus Hoeflich
  - Nicolas Stransky
  - Stefanie Schalm
  
- **Blueprint Informatics**
  - Adam Whelan
  - Tat Chu
  - Will Oemler