



***RET* fusion gene detection in NSCLC: how (not) to FISH?**

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Disclosures

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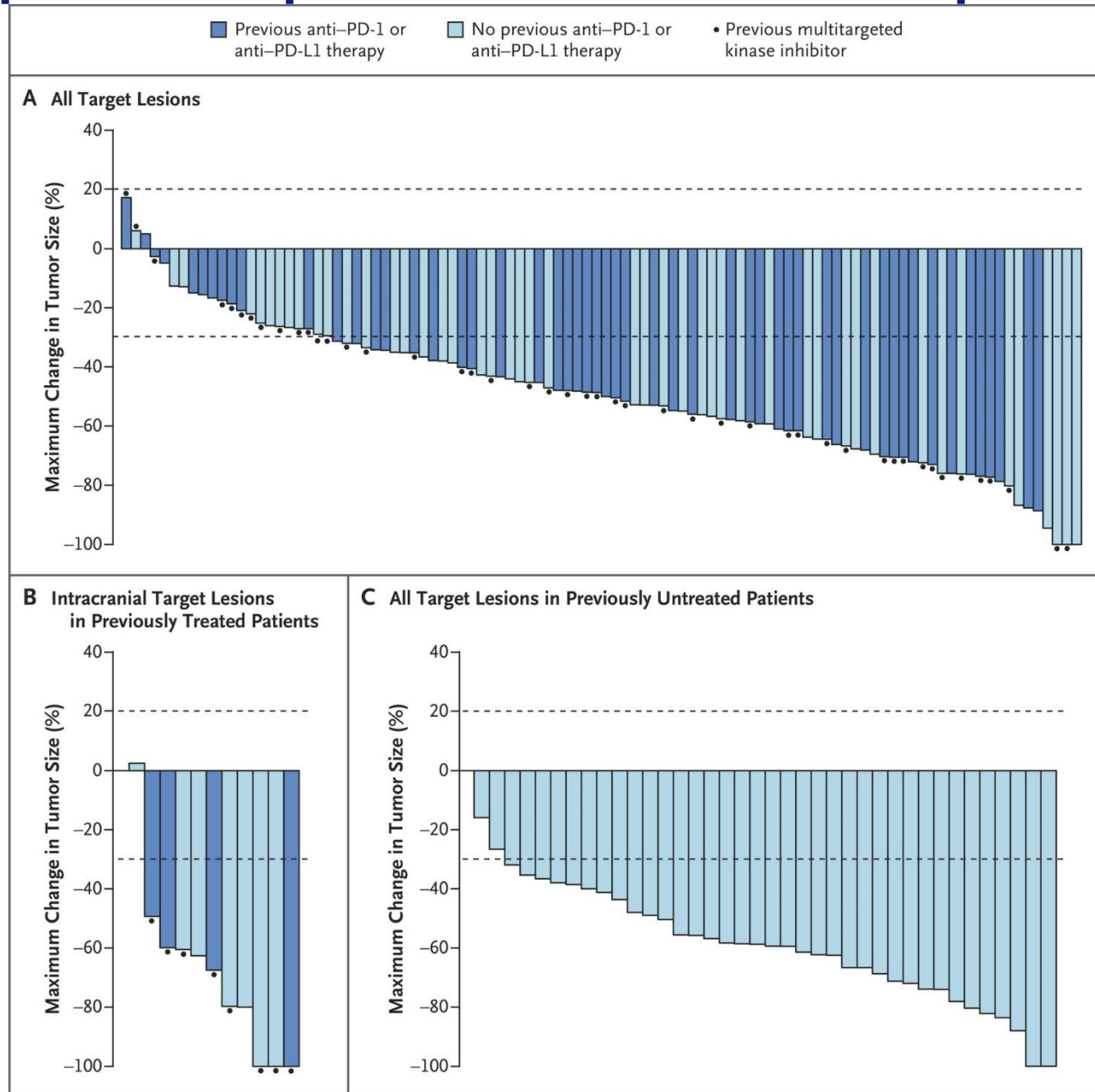
Advisory boards : Abbvie, AstraZeneca, Bayer, Janssen, Lilly, Pfizer, PGDx

Honorarium: AstraZeneca, Lilly, Pfizer

***RET*-targeted diagnostics**

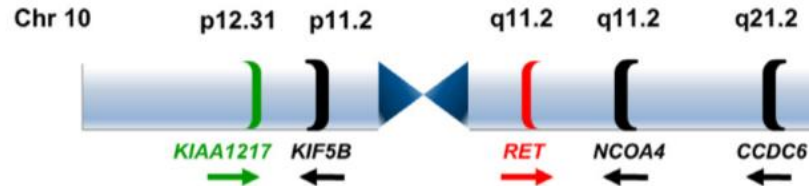
- *RET* fusion genes are oncogenic drivers in NSCLC (adenocarcinoma)
 - Predominantly found in younger NSCLC patients without smoking history
 - Prevalence of *RET* fusion genes in NSCLC: 1-2%
 - Multikinase *RET* inhibitors: objective response rates (ORR) of 30%
 - Selective *RET* kinase inhibitors, e.g pralsetinib (BLU-667) and selpercatinib (LOXO-292): ORR of 64-85%
- FDA approval for NSCLC, EMA soon expected

Waterfall plots of selpercatinib-treated NSCLC patients



RET fusion genes in cancer

RET-fusion proteins



Thyroid

- CCDC6-RET
- NCOA4-RET
- TRIM33-RET
- ERC1-RET
- GOLGA5-RET
- HOOK3-RET
- KTN1-RET
- PCM1-RET
- PRKAR1A-RET
- TRIM24-RET
- TRIM27-RET
- AKAP13-RET
- FKBP15-RET
- SPECC1L-RET
- TBL1XR1-RET

10-20% in PTC

Lung

- KIF5B-RET
- CCDC6-RET
- NCOA4-RET
- TRIM33-RET
- CUX1-RET
- KIAA1468-RET
- KIAA1217-RET

1-2% in NSCLC

Others

Colon

- CCDC6-RET
- NCOA4-RET

Skin

- GOLGA5-RET

CML

- BRC-RET
- FGFR1OP-RET

Breast

- ERC1-RET

Low rates

Kanker Instituut



From ESMO recommendations 2021 on the standard methods to detect *RET* fusions

Table 1. Summary of main features, strengths and weaknesses of all available techniques to detect *RET* rearrangements

| Method | Sensitivity | Specificity | Detection of partner | Detection of expression |
|-------------|----------------------------|----------------------------|----------------------|-------------------------|
| IHC | Moderate ^a | Moderate ^b | No | Yes |
| FISH | High | High | No/Yes ^c | No |
| RT-PCR | Moderate/high ^d | High | Yes/No ^e | Yes |
| DNA-seq NGS | Moderate ^f | High/moderate ^g | Yes | No |
| RNA-seq NGS | High | High | Yes | Yes ^h |

→ FISH is the currently used standard for the detection of *RET* fusions, with good sensitivity and specificity

BUT

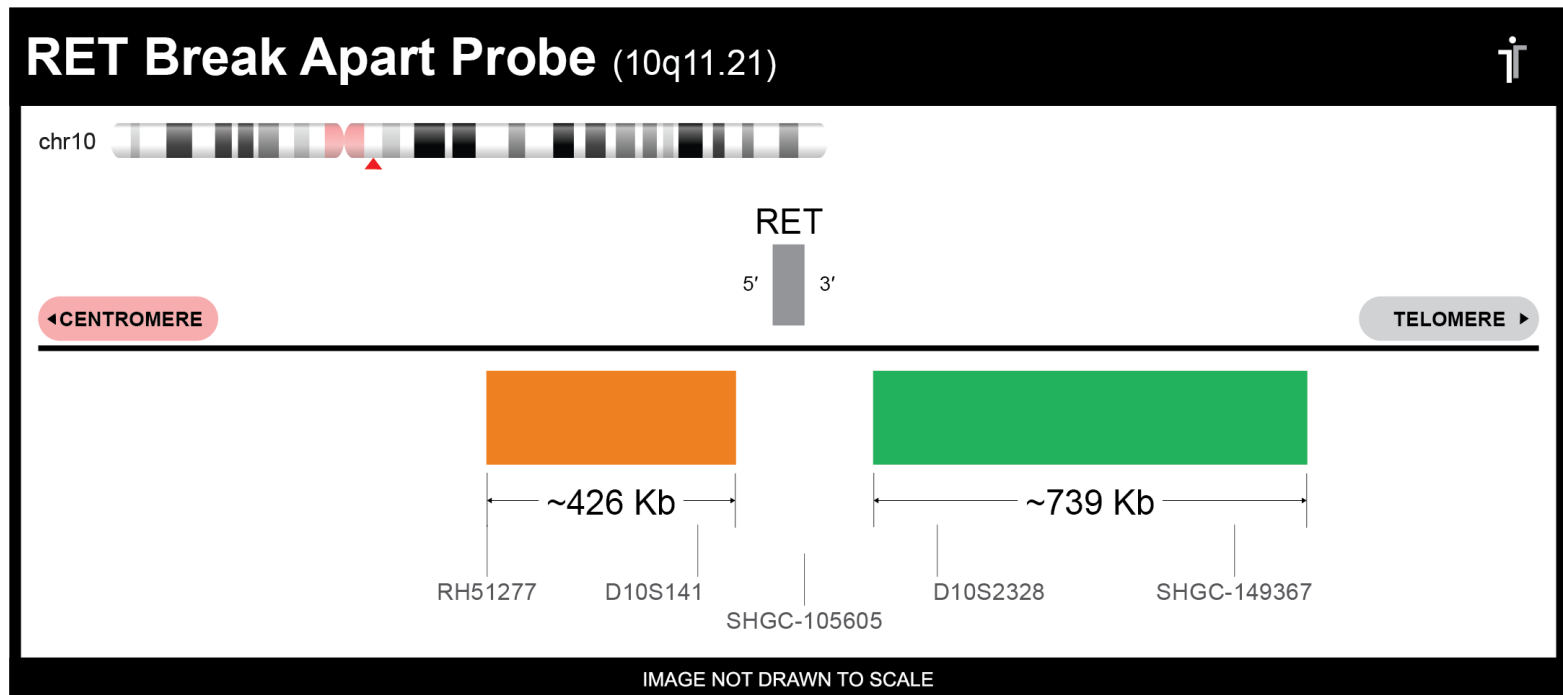
In routine practice often discordant results between FISH and RNA NGS results

Estimation of sensitivity and specificity of *RET* FISH
analysis of NSCLC

RET FISH analysis

Scoring system according to lab-specific routine practice

- 15% or more of cells with a split signal of at least more than 1 signal diameter between the 5' and 3' signals
- single 3' signal
- complex pattern: any pattern except usual split or single 3' patterns
- Single 5' signals: clinically negative, because of lost kinase domain



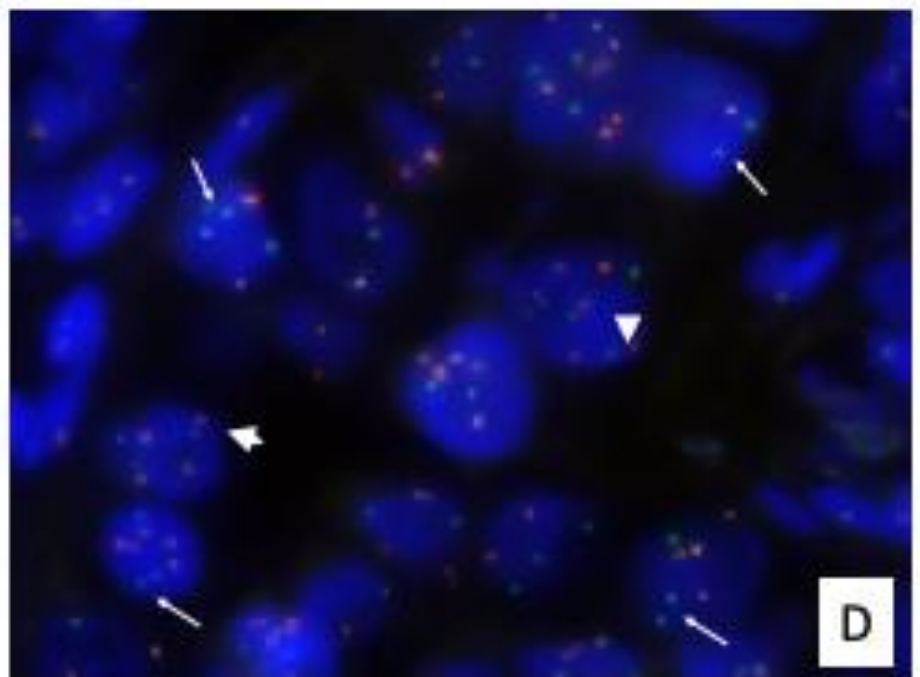
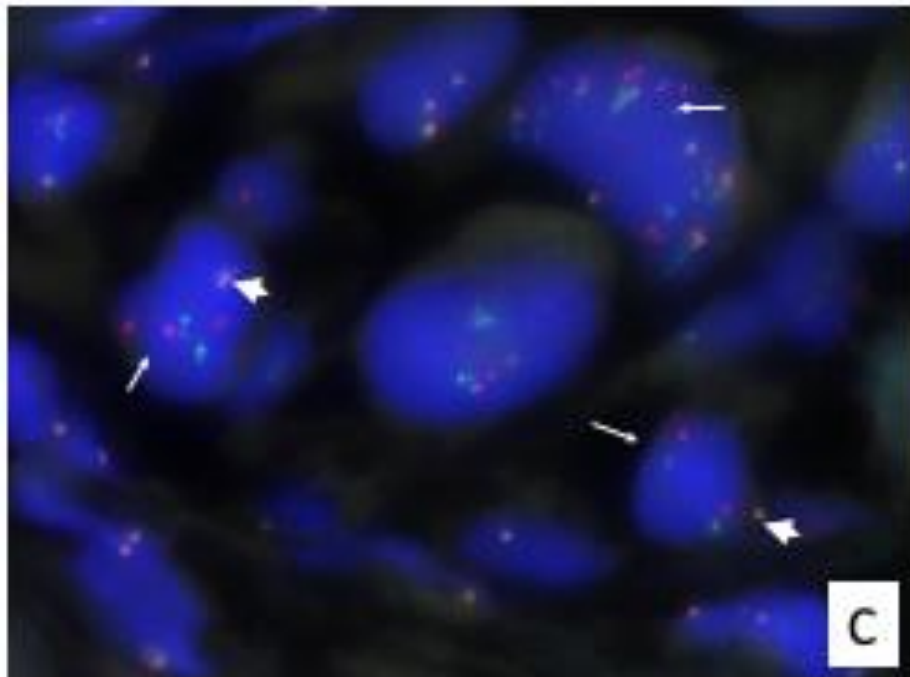
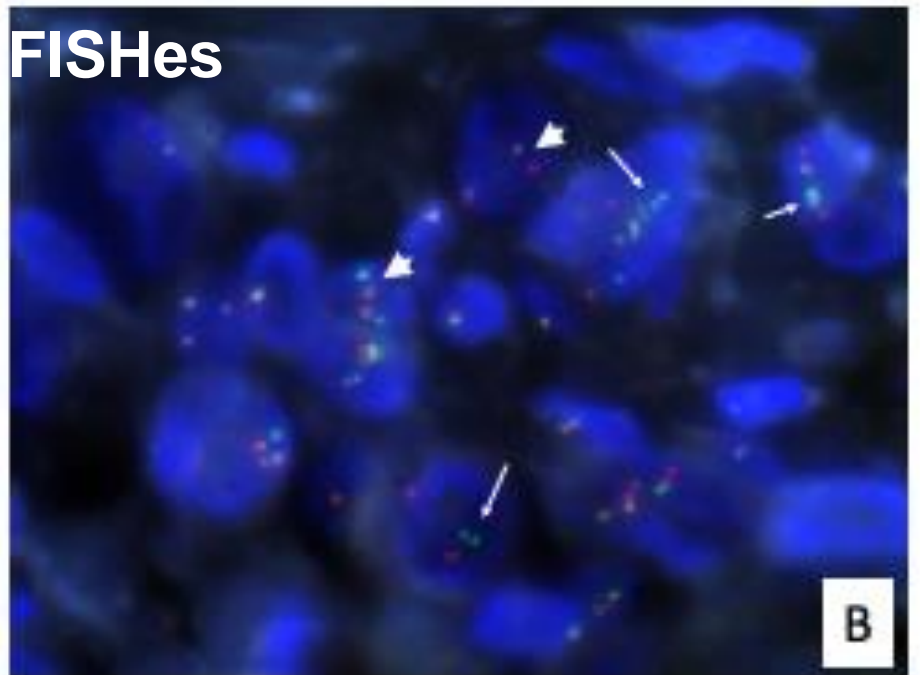
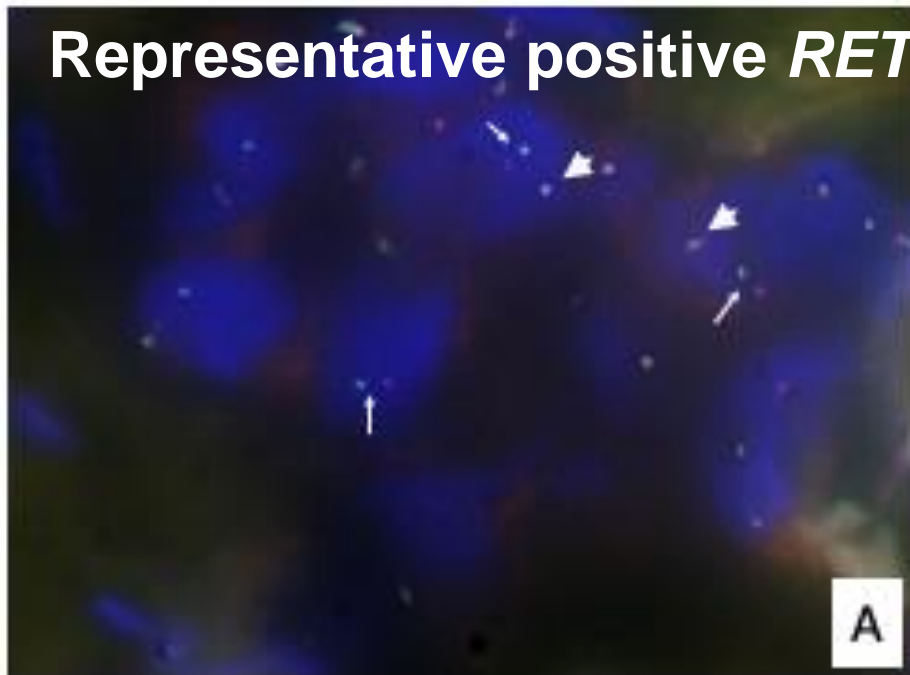
Patient cohort: 4873 NSCLC patients

- Case selection from January 2016 to November 2019
- Routine screening for *RET* fusions according to local logistics
FISH: 2858; by RNA NGS: 2015
- Collaborating institutes:
***RET* FISH:** Amsterdam UMC, Erasmus MC, Maastricht UMC+, and Netherlands Cancer Institute, the Netherlands
RNA NGS: St. James Hospital, Dublin, Ireland and Leiden University Medical Center, the Netherlands

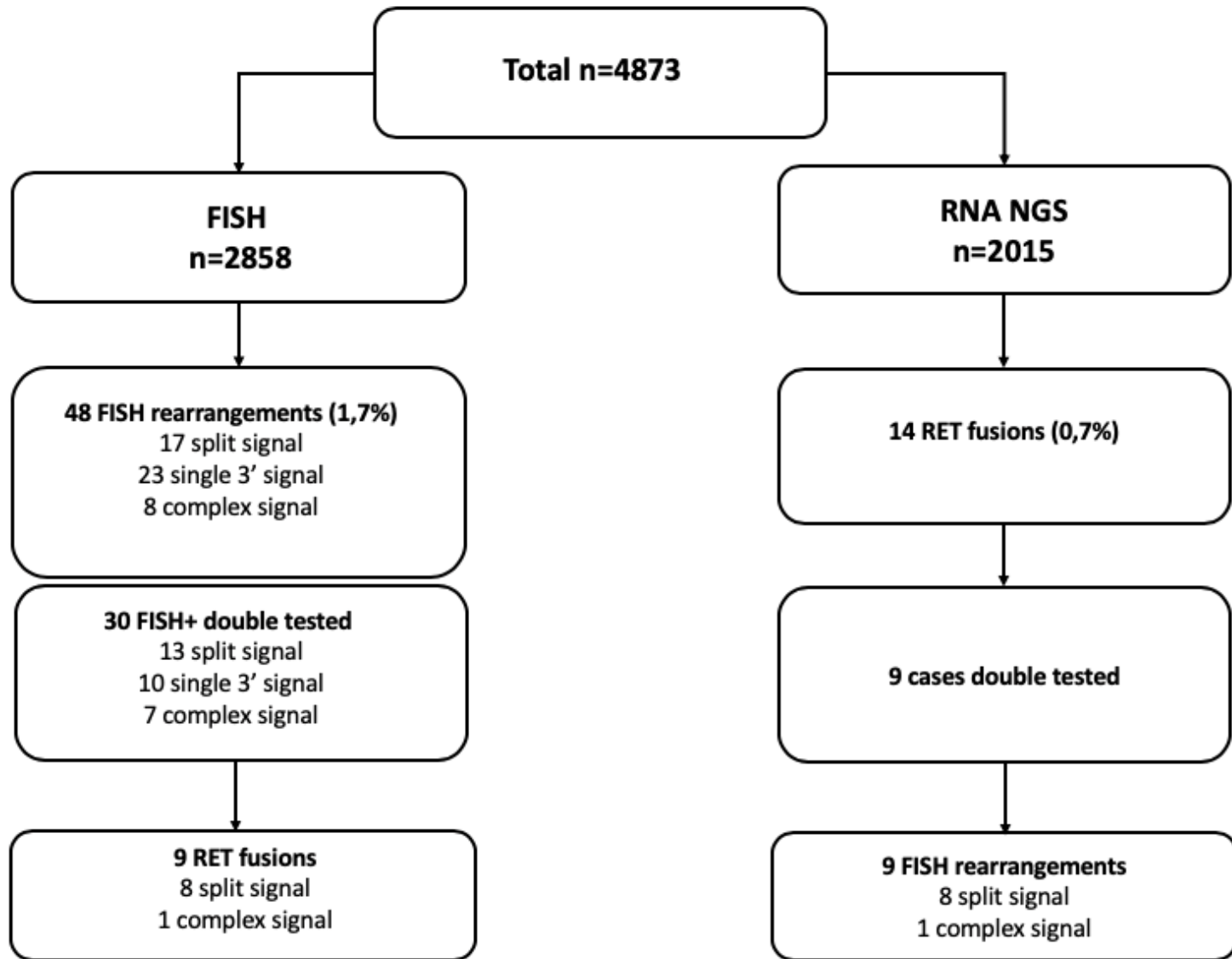
Study set up

- Comparison of data from routine FISH and RNA NGS analysis according to lab-specific procedures
- Four different *RET* break apart FISH assays: Vysis, SureFISH, Kreatech, and Cytocell Aquarius
- Three different RNA NGS methods: Archer, Oncomine, and Asuragen Quantidex
- Cross validation if possible

Representative positive *RET* FISHes



Yield



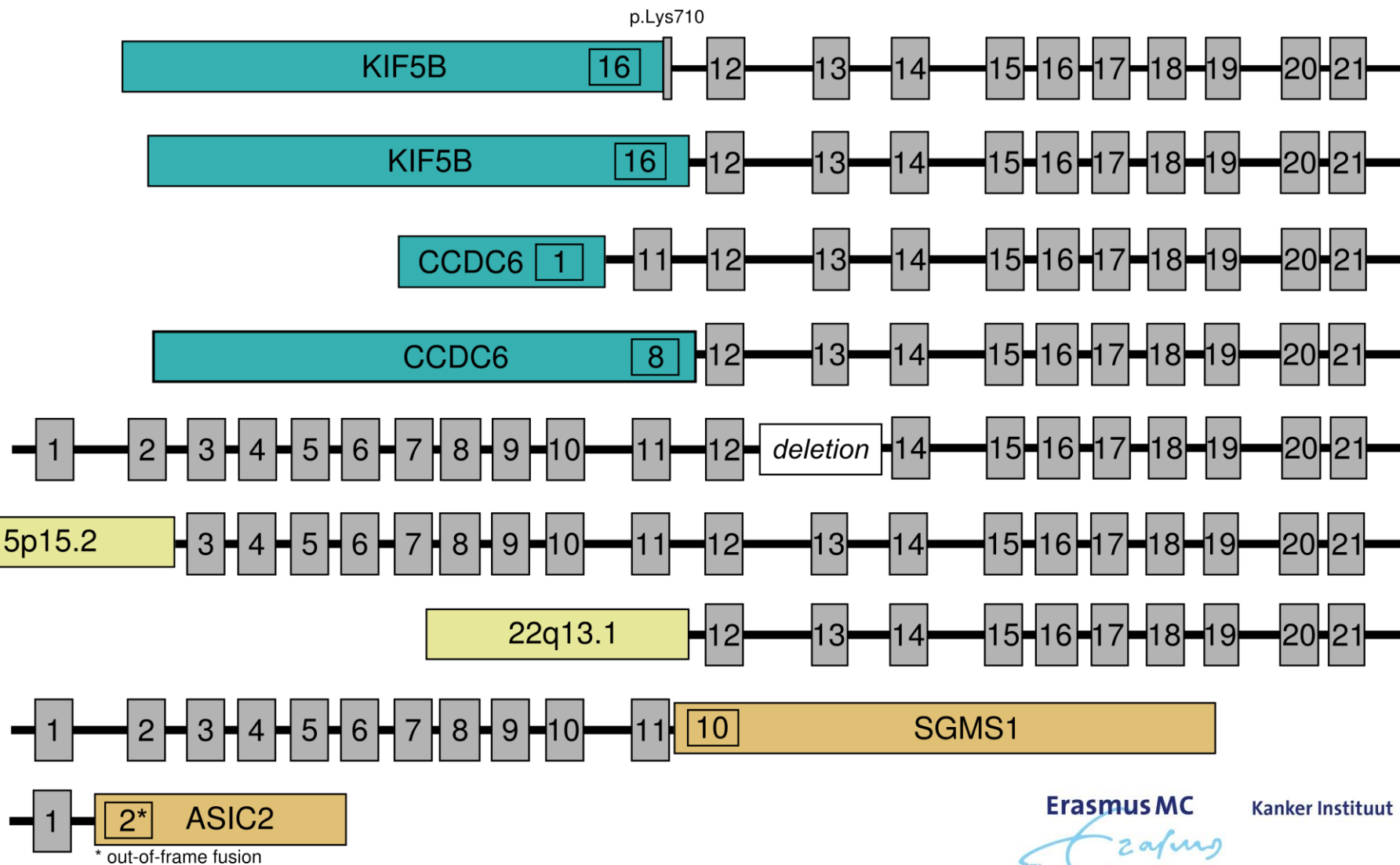
Conclusions 1

- *RET* FISH results in 70% of cases in false-positive results
- All RNA NGS positive cases could be detected by *RET* FISH

WGS cohort – Hartwig medical foundation

- WGS data from 520 NSCLC samples
- Stage IV lung cancer patients in all lines of screening and therapy
- Analysis of genomic loci of *RET*, *ALK*, *ROS1*

WGS analysis of *RET* locus for functional fusions (4) and disruptive events (5)



WGS results and conclusion

- Functional fusions compared to all disruptive events per locus:
 - *RET* (4/9, 44%)
 - *ALK* (27/34, 79%)
 - *ROS1* (9/12, 75%)
- In NSCLC the *RET* locus shows more rearrangements not resulting in functional gene fusions than shown for the *ALK* and *ROS1* loci
- WGS provides a possible explanation that aberrations observed by *RET* FISH often could not be substantiated by RNA NGS

Final conclusions

- *RET* FISH analysis is a sensitive but highly unspecific screening method for *RET* fusions in lung cancer
- *RET* screening by FISH always requires confirmation by an orthogonal technique, due to frequently occurring *RET* rearrangements not resulting in functional fusions in NSCLC
- Ranodic et al (2021), *J Thorac Oncol*, accepted for publication



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| Case | FISH | FISH pattern | RNA NGS |
|------|--|------------------|------------------------|
| 1 | Kreatech and Vysis | Split | KIF5B(ex15): RET(ex12) |
| 2 | Kreatech and Vysis | Split | KIF5B(ex15): RET(ex12) |
| 3 | Kreatech and Vysis | Split | KIF5B(ex15): RET(ex12) |
| 4 | Kreatech and Vysis | Split | CCDC6(ex1):RET(ex12) |
| 5 | Sure FISH | Split | CCDC6(ex1):RET(ex12) |
| 6 | Sure FISH | Split | KIF5B(ex15): RET(ex12) |
| 7 | Vysis | Split | KIF5B(ex15): RET(ex12) |
| 8 | Vysis | Split | KIF5B(ex15): RET(ex12) |
| 9 | Kreatech and Vysis | Split | KIF5B (ex15):RET(ex11) |
| 10 | Vysis | Split | KIF5B(ex15): RET(ex12) |
| 11 | Sure FISH | Split | KIF5B(ex15): RET(ex12) |
| 12 | Sure FISH | Split | CCDC6(ex1):RET(ex12) |
| 13 | CytoCELL Aquarius | Split | KIF5B(ex15): RET(ex12) |
| 14 | Sure FISH and CytoCELL Aquarius | Split | KIF5B(ex15): RET(ex12) |
| 15 | CytoCELL Aquarius | Split | CCDC6(ex1):RET(ex12) |
| 16 | CytoCELL Aquarius | Split | KIF5B(ex15): RET(ex12) |
| 17 | CytoCELL Aquarius | Split | No fusion |
| 18 | CytoCELL Aquarius | Split | No fusion |
| 19 | CytoCELL Aquarius | Split | No fusion |
| 20 | Kreatech | Split | No fusion |
| 21 | <i>Sure FISH and CytoCELL Aquarius</i> | <i>Split</i> | <i>No fusion</i> |
| 22 | CytoCELL Aquarius | Single 3' | No fusion |
| 23 | Kreatech | Single 3' | No fusion |
| 24 | Sure FISH | Single 3' | No fusion |
| 25 | <i>Sure FISH and CytoCELL Aquarius</i> | <i>Single 3'</i> | <i>No fusion</i> |
| 26 | <i>Sure FISH, Kreatech and Vysis</i> | <i>Single 3'</i> | <i>No fusion</i> |
| 27 | Sure FISH, Kreatech and Vysis | Single 3' | No fusion |
| 28 | Sure FISH, Kreatech and Vysis | Single 3' | No fusion |
| 29 | Vysis | Single 3' | No fusion |
| 30 | Vysis | Single 3' | No fusion |
| 31 | Vysis | Single 3' | No fusion |
| 32 | Sure FISH and Vysis | Complex | KIF5B(ex15): RET(ex12) |
| 33 | Sure FISH | Complex | KIF5B(ex15): RET(ex12) |
| 34 | Kreatech | Complex | No fusion |
| 35 | Kreatech | Complex | No fusion |
| 36 | Kreatech and Vysis | Complex | No fusion |
| 37 | <i>Sure FISH, Kreatech and Vysis</i> | <i>Complex</i> | <i>No fusion</i> |
| 38 | Vysis | Complex | No fusion |
| 39 | Vysis | Complex | No fusion |