ISPOR EU 2021 Methods Workshop:

Policy and Statistical Issues in HTA Review of Histology-Independent Technologies (HIT) in Oncology





Improving healthcare decisions

Jeremy Snider - Senior Quantitative Scientist, Flatiron Health Dr Jacoline Bouvy - Technical Director, NICE Scientific Advice Joshua Ray - Head Global Evidence, Global Access, Roche

Workshop Agenda

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0	Jeremy Snider	Introduction of speakers / provide workshop overview
5	Dr Jacoline Bouvy	 Provide background for previous HIT decisions Discuss evidence and analyses that can provide the best insight for future decisions
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Introduction to Histology-Independent Analyses Using Real-World Data

Jeremy Snider, PhD MPH

Senior Quantitative Scientist Flatiron Health jsnider@flatiron.com

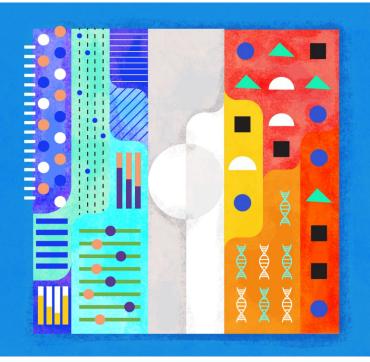


Glossary of Terms

Biomarker-defined cohort: A biological molecule found in blood, other body fluids, or tissues that is a sign of a normal or abnormal process, or of a condition or disease (US NCI Dictionary of Cancer, 2021)

Histology-Independent *(aka Tumor agnostic)* **technology**: A technology that targets a known biomarker, regardless of where in the body the tumor originates

Real world data (RWD): Data relating to patient health status and/or the delivery of health care from routinely-collected information (EMA, 2021)





Workshop goals

- Describe the current landscape of Histology-Independent Technologies (HIT) in precision oncology
- Provide a Health Technology Authority (HTA) and Pharmaceutical industry perspective around research questions and evidence gaps in HIT decision-making
- Discuss the value of longitudinal real-world data (RWD) for improving decision-making
- Discuss statistical issues and approaches that are introduced in HIT analyses





Common challenges in biomarker-defined studies

- Drugs targeting biomarkers are of growing importance in oncology research, presenting challenges to researchers:
 - Uncertainty in natural history / standard of care in newly-defined populations
 - Biomarker-defined **trial data** tend to be:
- **Immature:** capture **response** to trial drug, but often not longer-term survival outcomes



Scarce: highly-specific, (often) **rare** biomarkers where enrollment is not powered to assess longitudinal outcomes

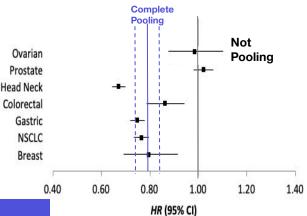


Heterogeneous: basket trials may included parallel development in adult and pediatric populations, and inclusion of patients regardless of tumour type

Heterogeneity in histology-independent data adds complexity to analyses

- Not pooling is an <u>inefficient use of data</u>, often leading to inadequate sample size for precise stratum-level inference
- Complete pooling can <u>inappropriately mask</u> <u>important differences</u> in biomarker / drug effect across histologies

These traditional analytic methods (using complete or no pooling of data) may not address current HTA decision problems





Poll: What is your experience with HITs?

- No clue hearing it about it for the first time
- Interested in the topic but limited experience
- Designing clinical trials
- Designing RWD studies
- Engagement in the HTA / Reimbursement process



Poll: Please tell us about your background!

- Academic
- Industry (Pharma / Biotech Manufacturer)
- Consultancy
- Research institution
- Government Agency / HTA
- Other



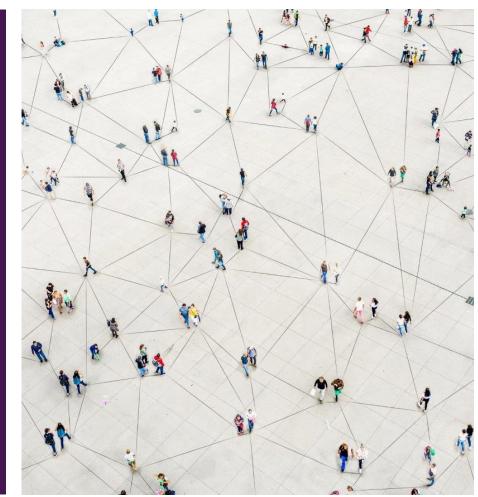
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Histology-independent cancer drugs

Dr Jacoline Bouvy Technical Director, NICE Scientific Advice





What is a histology-independent cancer drug?



19 July 2019 EMA/CHMP/391684/2019 Media and Public Relations

Press release

First 'histology-independent' treatment for solid tumours with a specific gene mutation

EMA's human medicines committee (CHMP) has recommended granting a marketing authorisation in the European Union for Vitrakvi (larotrectinib) for the treatment of adult and paediatric patients with solid tumours that display a Neurotrophic Tyrosine Receptor Kinase (NTRK) gene fusion. Treatment with Vitrakvi is recommended for patients whose disease has spread or cannot be surgically removed, and who have no other satisfactory treatment options.

Vitrakvi is the first so-called 'histology-independent' cancer treatment recommended for approval in the EU. This means that it can be used to treat non-haematological (i.e. that do not begin in the blood or bone marrow) tumours with this specific mutation, regardless of where in the body the tumour originated. Before patients can be started on the medicine, the presence of the mutation in the tumour should be confirmed by a validated test.

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diate May 23, 2017

The U.S. Food and Drug Administration today granted accelerated approval to a treatment for patients whose cancers have a specific genetic feature (biomarker). This is the first time the agency has approved a cancer treatment based on a common biomarker rather than the location in the body where the tumor originated. Why are these drugs different?

Traditional therapeutic indication First-line monotherapy of metastatic colorectal cancer (chemotherapy) RCT in patients with metastatic colorectal cancer

Histologyindependent indication

NTRK-positive solid tumours (larotrectinib, entrectinib)

Basket trial of patients with biomarker-positive solid tumours

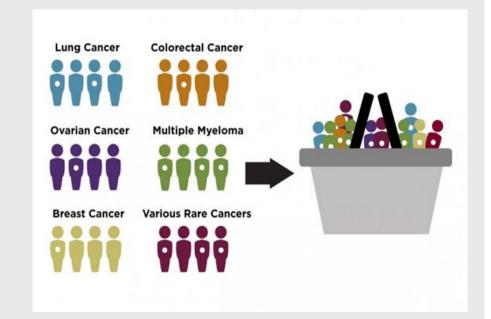
What is a basket trial?

Includes patients with lots of different tumour types whose cancer shares a genomic mutation

No comparator arm

Primary endpoints:

- Objective response rate
- Duration of response



NICE asks 2 key questions about new drugs:

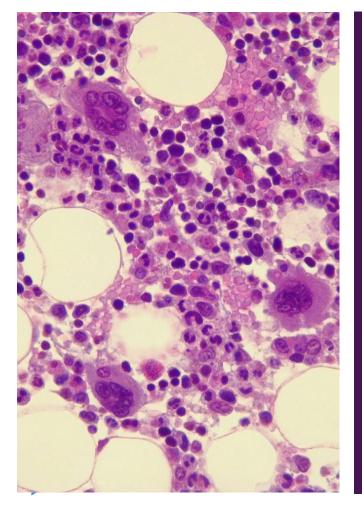


How well does the technology work compared to standard practice in the National Health service (NHS)?

How much does this course of action cost compared to standard practice in the NHS?

Cost

NICE



Key challenges for HTA

NICE

Generalisability of trial evidence

How to include testing costs

To subgroup or not to subgroup?

Developing the counterfactual

Dealing with response-based endpoints

Generalisability of trial evidence and testing costs

- Testing is required to identify patients eligible for treatment
- Point of testing in trial likely to differ from testing strategy in the NHS (no companion diagnostic)
- Extrapolation to histologies who were not included in the trial at all?



To subgroup or not to subgroup

Calculating the cost effectiveness of a new drug:

Costs of new drug – costs standard of care

Effects new drug – Effects standard of care

• Does a single ICER convey enough meaningful information on whether the drug provides value for money to the NHS across all indications?



Developing the counterfactual

- What would have happened to these patients had they received the standard of care?
- No comparator arm in basket trial
- Usual approach for single-arm studies: indirect comparison to historical control data (data from a different trial)

Trial population: NTRK positive tumours, treatment with histology independent drug **Comparator:** NTRK positive and NTRK negative tumours, treatment with standard of care

Dealing with responsebased endpoints

NICE

Traditional endpoints in cancer

Progression-free survival (PFS) and overall survival (OS)

Partitioned survival models

Can use PFS and OS survival curves to extrapolate long-term outcomes

Drug approved based on response

- Surrogate for PFS and OS or not?
- PFS and OS measured in trial, but data was immature at time of HTA

NICE National Institute for Health and Care Excellence

Thank you.

Contact details:



Jacoline.Bouvy@nice.org.uk



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Histology-independent cancer technologies Joshua Ray, Head Global Evidence, Global Access, Roche

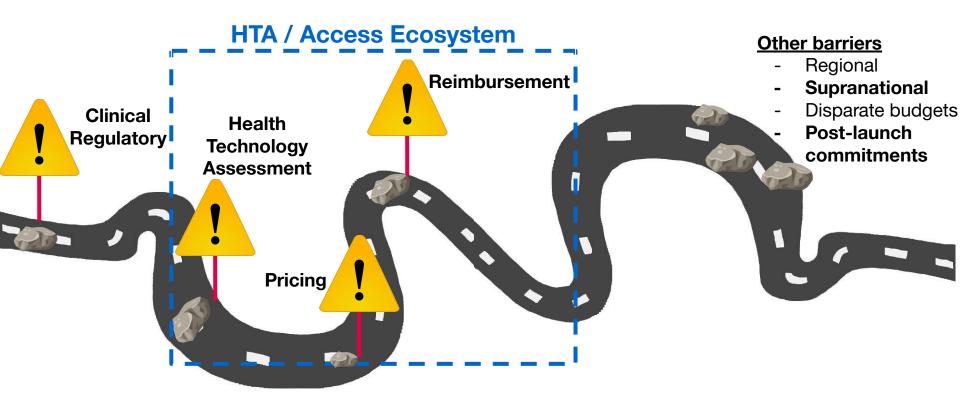




Joshua Ray is an employee of F. Hoffmann-La Roche Ltd. The views expressed in this presentation & panel discussion are his personal opinions, and may not reflect those of his employer.

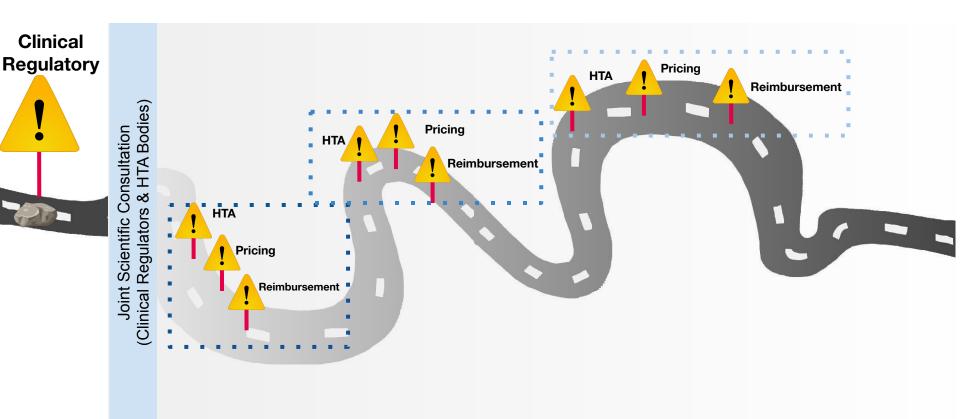


Along the road to patient access





The road to patient access <u>is likely to change</u> in the presence of histology independent technologies





What's the situation?: Limited evidence base at the time of an initial access decision

- Uncertainty around prevalence
- Partial understanding of natural history and prognostic impact of the genomic alteration
- Heterogeneity
 - Tumor origin
 - \circ Line of therapy
- Lack of a clearly defined comparator (in trials & within the HTA evaluation)

Environmental Concerns

Technology's Evidence Package

- Established reimbursement / HTA pathways
- Genomic testing
 - Clinical Guidance
 - Infrastructure
 - Reimbursement pathways



Why does this matter?



Sequential and separate processes for regulatory and reimbursement decisions creates challenges and uncertainty for HTA-bodies, Payers & Manufacturers

The environment...

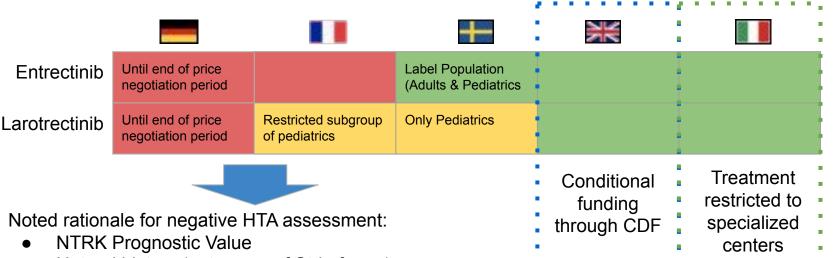
- Select Clinical Regulatory (e.g. FDA, EMA, MHLW Japan) bodies have evolved approval pathways
 - Expedite the development & assessment of technologies intended to treatment serious conditions with a high unmet need
 - Allows for the use of preliminary clinical evidence that demonstrates improvement over available therapy
- The paradigm for treating cancer continues to evolve ²
 - As of 2019, 21 biomarker targeted therapies in development assessing a tumor agnostic label ²
 - For NTRK fusions, previous estimates have estimated a tumor-specific RCT would take 17-105 years to complete ¹

1) Lozano-Ortega G. et al. Tumor-specific RCTs in rare oncogene-driven cancers: asking for the impossible? ISPOR 2019. 2) Mosele F. Recommendations for the use of NGS for patients with metastatic cancers. Annals of Oncology. 2020. 2) IQVIA Pipeline Intelligence; clinicaltrials.gov; trialtrove; Pharma Intelligence, Mar 2020





What does this mean? A Snapshot at Europe



- Natural history (outcomes of Std of care)
- Surrogate endpoint validation
- Single arm trial / no comparative evidence (or randomization)
- Trial cohort size
- Limited acceptance of intra-patient comparison & RWD

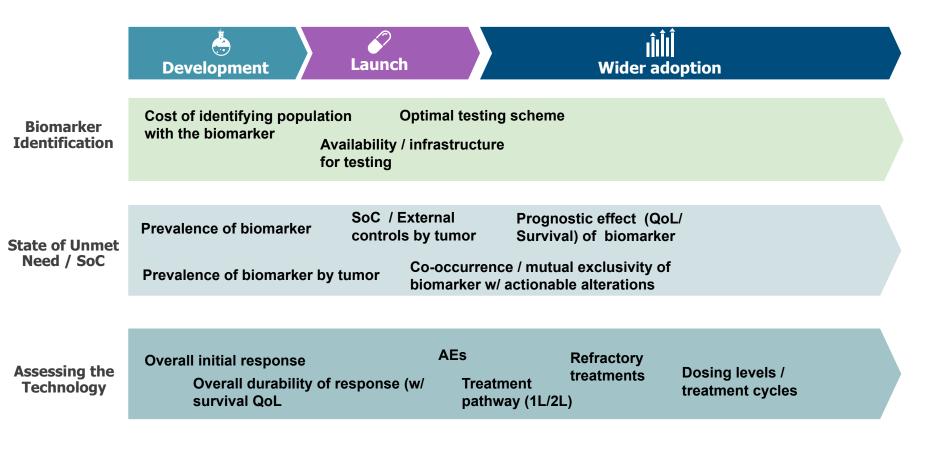
Where do we go from here?

- Utilize Joint Early Scientific Advice (Regulators & HTA) to align evidence generation plans
- Earlier evidence planning
 - Utilizing RWD
 - Intrapatient analyses
- Pivot towards continual evidence generation
 - Conditional approval & managed entry agreements
- Build upon evolving frameworks ¹:
 - Lessons from the existing histology independent technologies



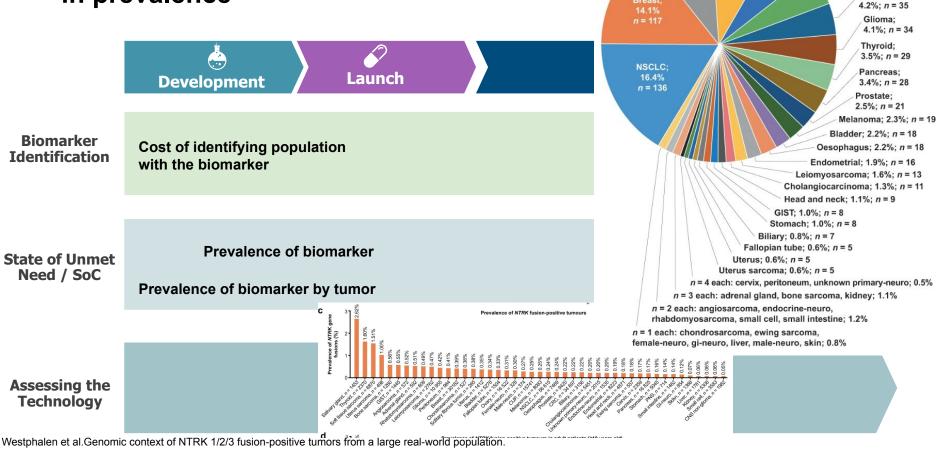


Where RWD may help narrow the gap



1001

RWD can help address uncertainty in prevalence



CRC;

9.3%

n = 77

Soft tissue

sarcoma:

9.6%

Breast:

Ovary: 6.2% n = 51

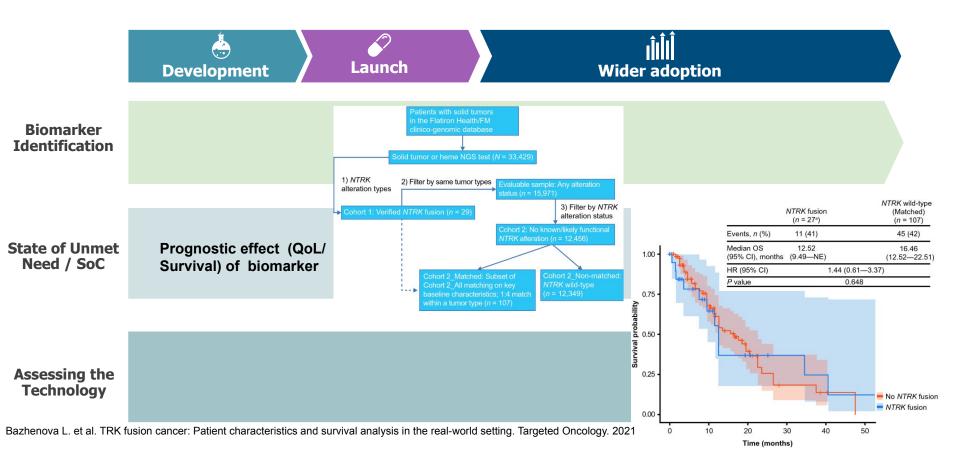
CUP; 4.8% n = 40

Salivary gland;

npj Precision Oncology. 2021

Example where RWD can address uncertainty in baseline prognosis







Summary

- Histology Independent Technologies present unique challenges and will become more commonplace
- Patient benefit may be foregone in the absence of early & multi-stakeholder dialog
 - This can be minimized by co-creating immediate and long-term solutions addressing patient-relevant outcomes and budgetary concerns
 - Focusing on both the implementation of acute treatments and their accompanying infrastructure requirements





Doing now what patients need next

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Statistical Approaches for Histology-Independent Analyses Using Real-World Data

Jeremy Snider, PhD MPH

Senior Quantitative Scientist Flatiron Health



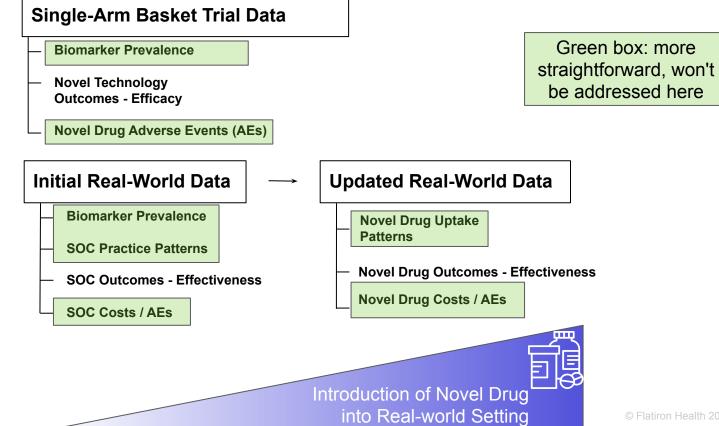
Goals for this presentation:

- Motivate using Real-World Data (RWD) in a biomarker-defined and/or histology-agnostic study
- Understand benefits and challenges of incorporating longitudinal data into histology-independent studies
- Understand how to contend with histology heterogeneity in an HIT analysis

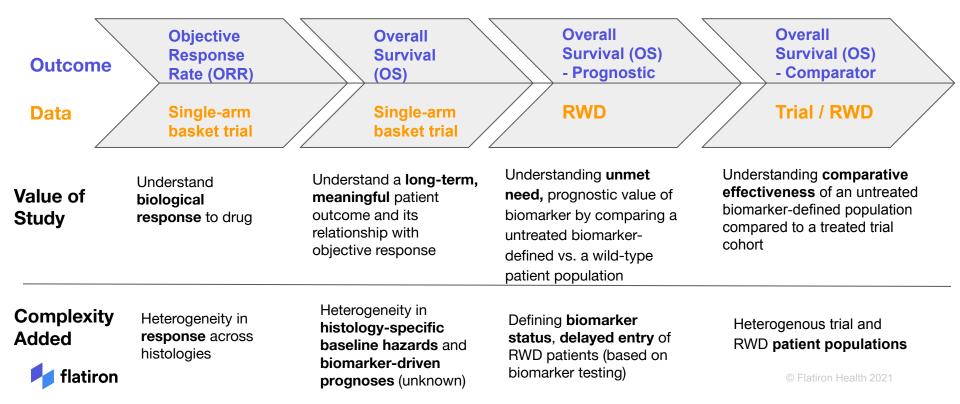




RWD can enhance biomarker-defined HEOR studies



Building evidence in histology-independent efficacy/effectiveness analyses



The first step: Objective Response Rate (ORR) from Single-Arm Trial

Histology	Observed Response	Estimated Mean Response Based on BHM (%)	95% CrI	
Fixed effects Pooled Random effects Soft-tissue sarcoma Salivary gland IFS Thyroid Lung Melanoma Colon GIST Cholangiocarcinoma Appendix Breast Pancreas Unrepresented	41/55 = 74.5% $10/11 = 90.9%$ $10/12 = 83.3%$ $7/7 = 100%$ $5/5 = 100%$ $2/4 = 50.0%$ $1/4 = 25.0%$ $3/3 = 100%$ $0/2 = 0%$ $0/1 = 0%$ $0/1 = 0%$ $0/1 = 0%$	74.20% Complete pooling suggests efficacious technology 88.10% efficacious technology 93.30% 91.60% 92.60% 52.50% 32.00% 88.30% 21.00% 30.00% 30.00% 30.00% 29.80% 56.90%	$\begin{array}{c} 62.0\% - 84.7\% \\ \hline \\ 66.0\% - 99.1\% \\ 58.0\% - 96.8\% \\ 70.5\% - 100\% \\ 63.0\% - 100\% \\ 30.4\% - 97.8\% \\ 12.4\% - 89.4\% \\ 2.6\% - 75.5\% \\ 49.3\% - 100\% \\ 0.0\% - 75.7\% \\ 0.1\% - 89.7\% \\ 0.1\% - 89.7\% \\ 0.1\% - 89.7\% \\ 0.2\% - 99.9\% \end{array}$	However, histology-specifi estimates are highly variable (11-93%)

Table 2 Probabilities of response for all histologies

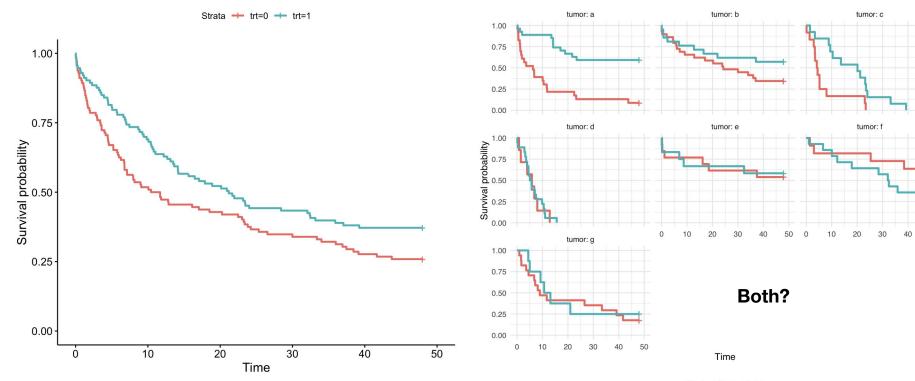


Murphy et al. (2021) Exploring Heterogeneity in Histology-Independent Technologies and the Implications for Cost-Effectiveness, Medical Decision Making

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How can do we contend with tumor heterogeneity in an Overall Survival (OS) analysis?

Complete Pooling?



No Pooling (Stratified by histology?)

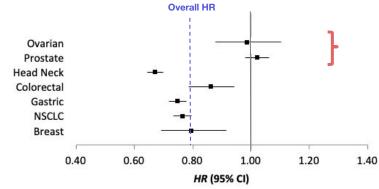
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50

Complete pooling: Can mask heterogeneity

Model: coxph(OS ~ received_drugX, data = all_mutationY_patients)

	Beta-coefficient	P-value
received_drugX	-0.23	0.002**



Should Drug X be approved for histology independent use?

HR = 0.79 Looks promising!

Tumor-specific HRs raise indicate non-effectiveness

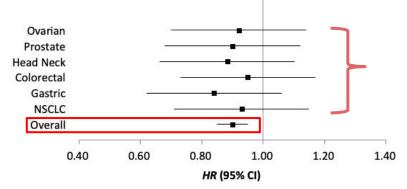
Tumor-level effects are important!



No Pooling: Stratum-level analyses do not make use of all available data

Model: coxph(OS ~ received_drugY,

data = all_mutationZ_patients_with_{cancer})



Overall effect is promising!

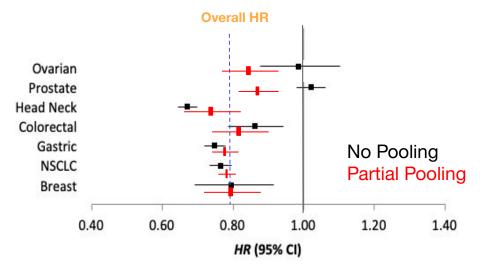
Tumor-specific effects are inconclusive

Both the **overall effect** and **tumor-level effects** are important!



Partial Pooling: A happy medium?

- Single source hierarchical models are a common approach to partial pooling
- They assume that strata-level estimates come from a single parent prior distribution
- These prior distributions for can be manually prespecified (e.g. Bayesian analysis), or calculated empirically (e.g. random effects/empirical Bayes)
- Strata-level estimates are pulled towards the global mean based on strata size





Balancing Clinical and Statistical Considerations when Interpreting Heterogeneity

Expert opinion on drug and tumor biology can inform a scientific rationale for pooling histologies. However, heterogeneity can also be interrogated in the data:

- Analysis of Variance (ANOVA): to test interaction effect between histology and biomarker/drug
- Multi-source Exchangeability Models (MEM): considers all possible pairwise exchangeability relationships, estimates the probability that subset of histologies should be considered statistically exchangeable (or poolable) -(Kane, 2019)

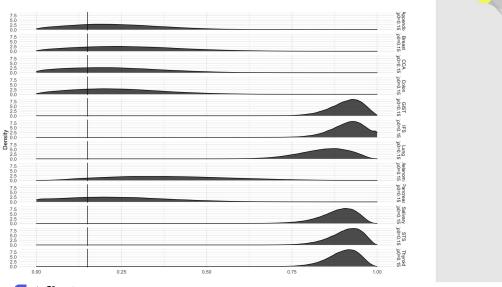
Quantifying Heterogeneity: ANOVA Interaction Test (in R)

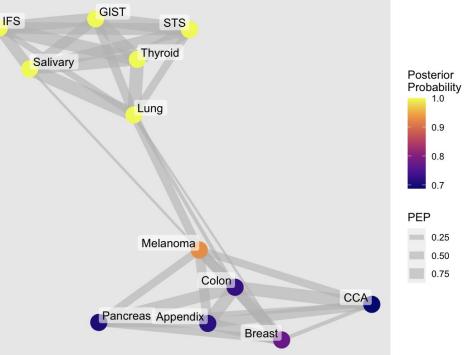
- An unstructured interaction test can help identify if effect modification by tumor histology is present in the data
- However, it won't indicate which subgroups are the source of the interaction, nor characterize the magnitude of the subgroup treatment effect differences



Quantifying Heterogeneity: Multi-source Exchangeability Models (MEM)

Analyzing Basket Trials under Multisource Exchangeability Assumptions (Kane, 2019)







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Conclusions

- RWD can enhance HEOR analyses that accompany HTA submissions:
 - Improving the breadth and depth of studied histologies
 - Providing meaningful longitudinal effectiveness data
- Longitudinal / survival data presents unique challenges in HIT analyses
- Multilevel modeling and measuring heterogeneity can improve analyses and guide methodological decisions





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Interactive case study

We'll go through one exercise together to understand how to answer a key question

1: Using Longitudinal RWD to understand the prognostic value of a biomarker across histologies

There are three additional exercises provided for your reference:

2: Using Longitudinal RWD to understand novel drug effectiveness across histologies

3: (Appendix): Combining Trial and RWD to improve precision and understand response heterogeneity in a histology-independent analysis

4 (Appendix 2): Prevalence across cohorts

Follow along at: http://bit.ly/ispor_hit



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