



External control cohorts for the single-arm LIBRETTO-001 trial of selpercatinib in RET+ NSCLC cancer



Min-Hua Jen,
Senior Director, Real World & Access Analytics
Eli Lilly and the company

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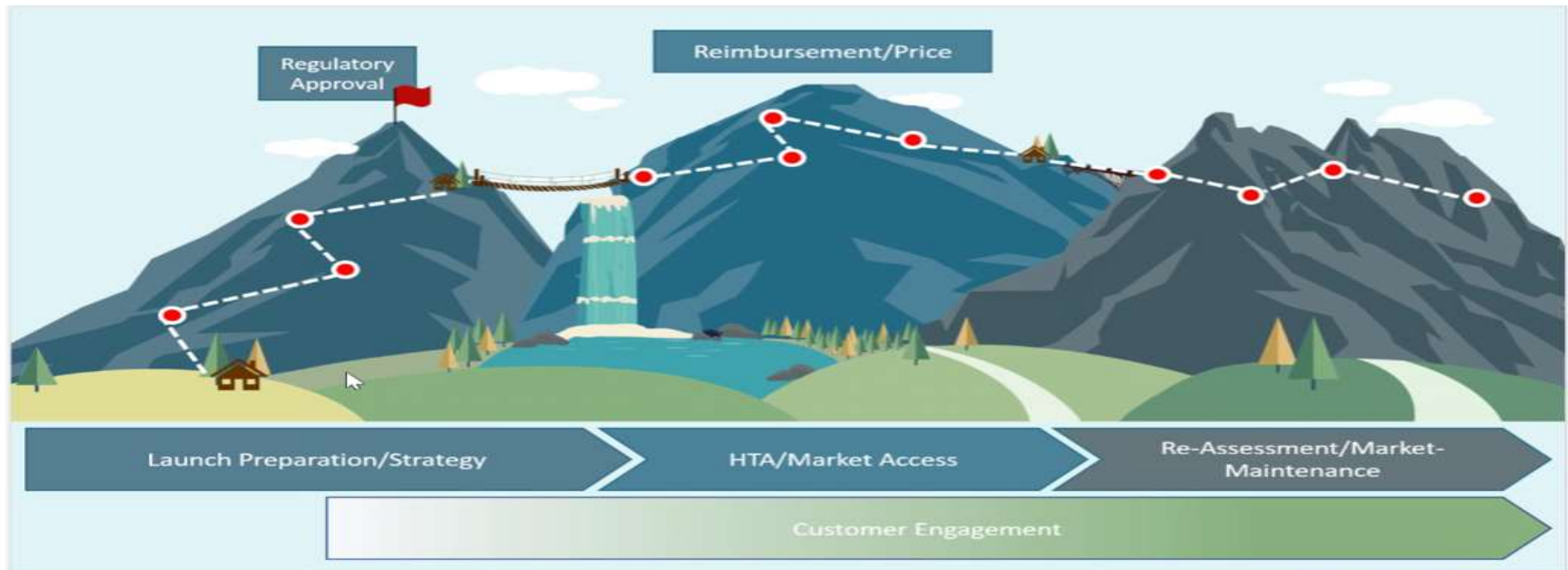
Outline

- **Setting the themes**
 - Background: An evolving paradigm to market access
 - Libretto-001: Selpercatinib in RET-altered cancers
 - Challenge: Evidence gaps inherited from single-arm trials for accelerated access
- **Optimized likelihood for accelerated access for patients**
 - Mitigated anticipated risks/uncertainties when synthesis evidence to fill the gaps inherited from single arm trial

An evolving paradigm for market access

YESTERDAY: The demonstration of a positive Benefit/Risk profile was usually enough to gain reimbursement

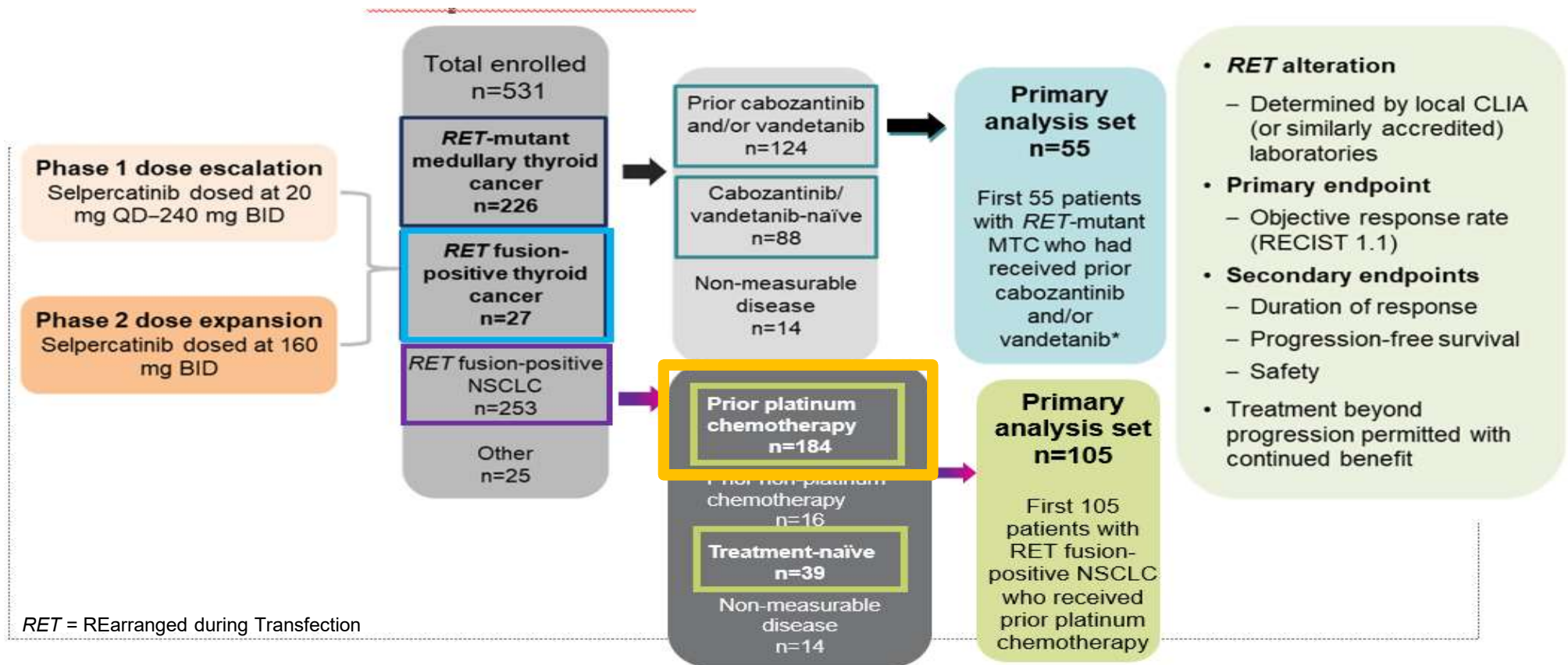
TODAY: A novel agent must demonstrate superior medical value and/or cost effectiveness benefits



Slide courtesy of the IBU stats team

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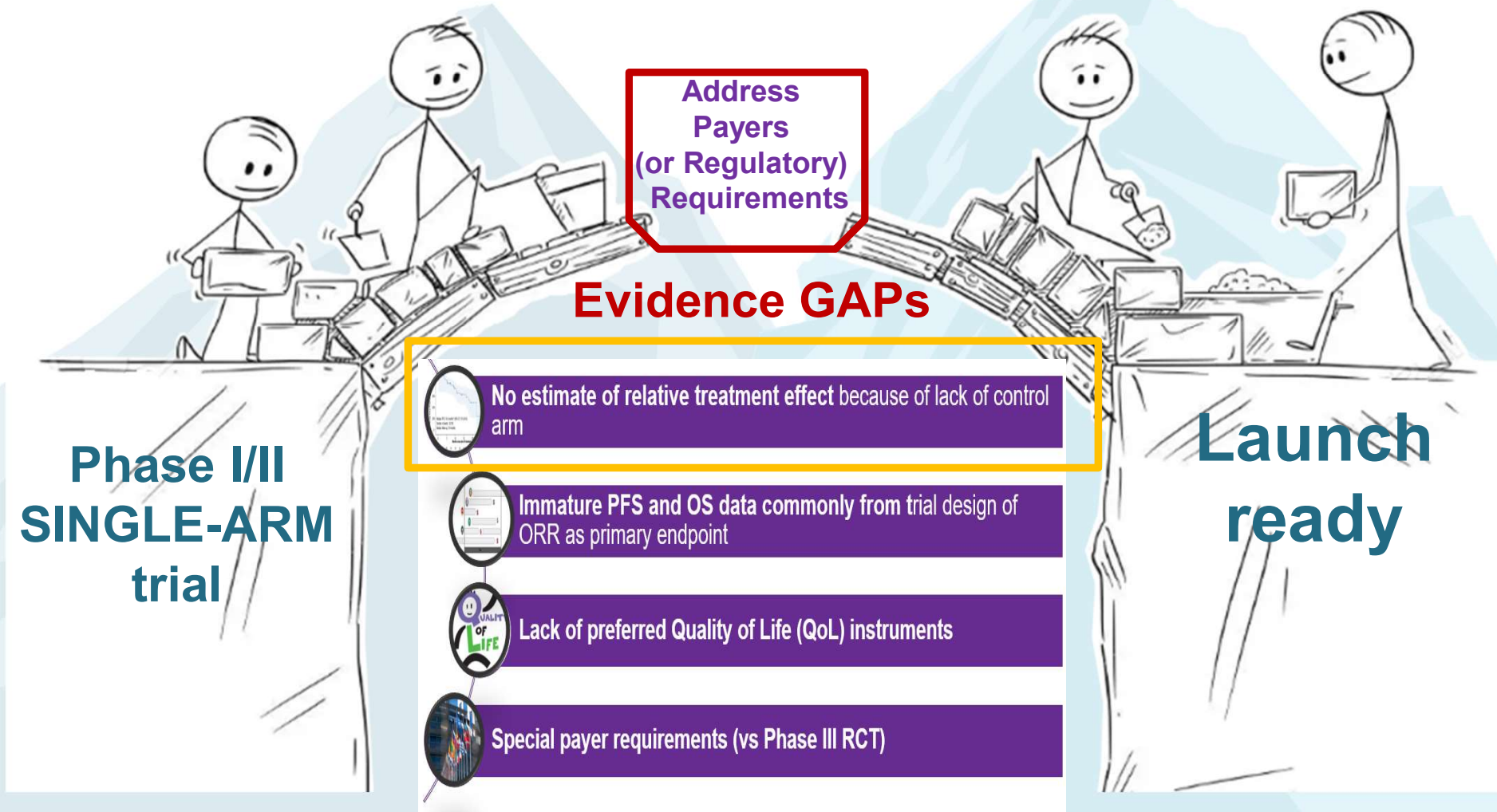
Libretto-001: Selpercatinib in RET-altered cancers



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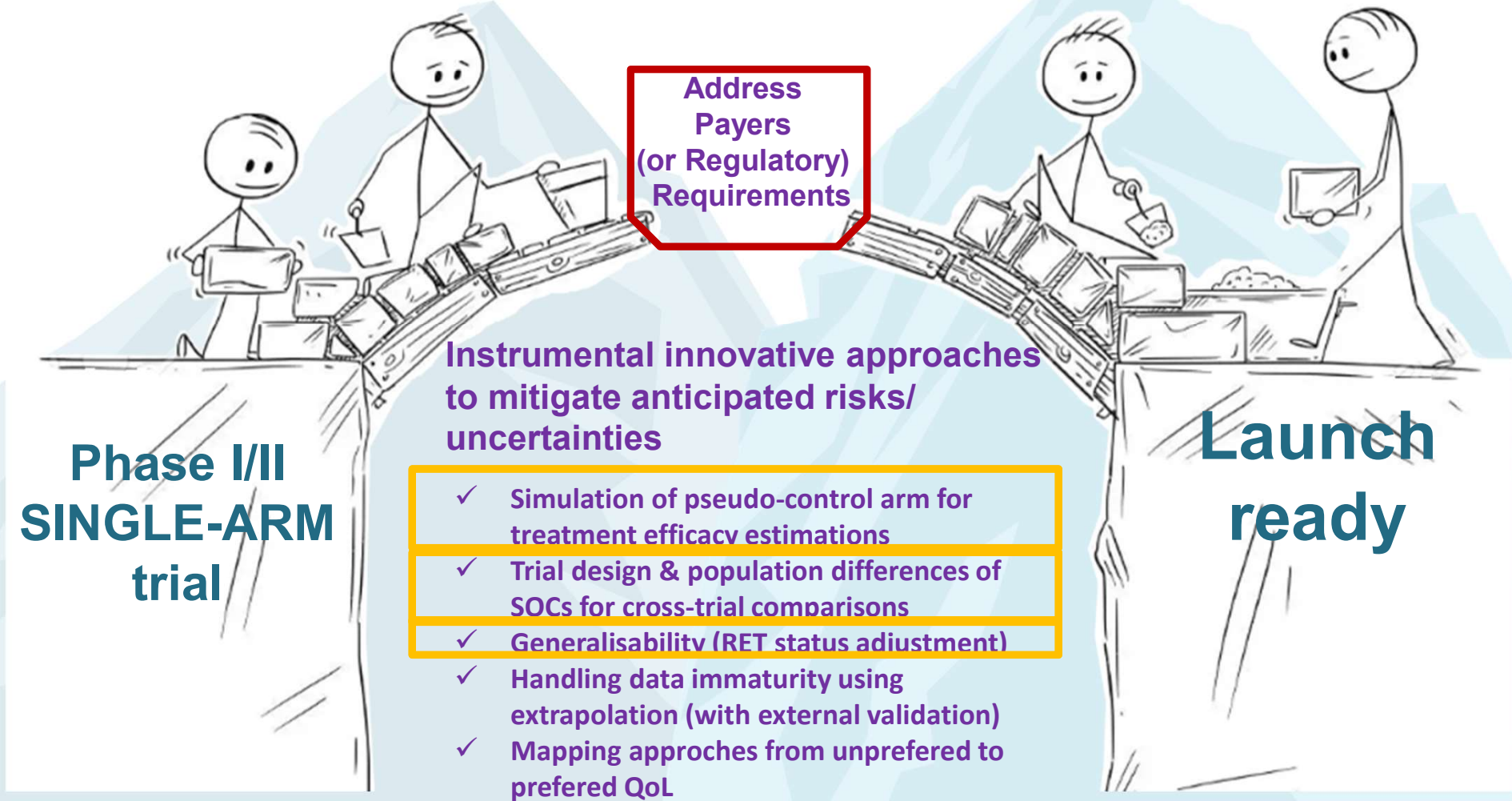


ACCELERATE market access via SINGLE-ARM trial (but not RCTs) is getting common





ACCELERATE market access via SINGLE-ARM trial (but not RCTs) is getting common





Optimized likelihood for accelerated access

- Estimating comparative efficacy of selpercatinib versus other available treatments for pre-treated RET+ lung cancer:
- Two-fold approach:
 - 1) Synthesize external control arm* using
 - ✓ Real-world data
 - ✓ Historical trial data
 - 2) Compare to standard of care(s) (SOCs) by addressing
 - ✓ Trial design and population differences of SOCs (i.e. biomarkers) cross trials

*Rolfo et al 2022 External control cohorts for the single-arm LIBRETTO-001 trial of selpercatinib in RET+ non-small-cell lung cancer

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Synthesize external control arm



Part 1: Real-world control

Objectives

- Compare efficacy (PFS, ORR and OS) of LIBRETTO-001 pre-treated NSCLC cohorts to corresponding real-world (RW) cohorts from Clinico-Genomic Database (CGDB)
- Three analytic approaches:
 - Analytic strategy 1: RET fusion positive control cohorts consistent with LIBRETTO-001
 - Analytic strategy 2: Control cohorts consistent LIBRETTO-001, other than RET status
 - Analytic strategy 3: Apply a factor to account for unknown RET status
- Primary comparison: Blended control

Eligibility of real-world cohort

- LIBRETTO-001 timeframe: Initiating index line of therapy \geq May 2017
- No clinical trial enrollment
- No history of RET inhibitor use
- No coexisting alterations (ALK, EGFR, ROS1, KRAS, BRAF)
- *RET* fusion positive

Possible sources of bias

- Favoring CGDB RW cohorts:
 - ✓ Immortal time bias in CGDB
 - ✓ Fewer patients 3rd line or later
 - ✓ More with earlier disease stage at diagnosis
- Favoring Libretto-001:
 - ✓ Any unique prognostic effect of RET fusion + status
 - ✓ More non-squamous, females and never-smokers
 - ✓ Younger, lower body weight, with better performance status
- Entropy balancing method was used to adjust for measured sources of possible bias.



Entropy Balancing (EB)

- Unlike propensity scoring methods, EB directly balances the distribution of variables between groups
- May be preferred when samples are small to avoid loss of population size, or when there are prespecified variables for the comparison groups to be balanced
- Results in a perfect match but may result in some extreme weights to achieve this.

Covariates for EB

- Systematic literature review:
 - ✓ Sex
 - ✓ Age
 - ✓ Body weight
 - ✓ ECOG performance status
 - ✓ Smoking status
 - ✓ NSCLC Histology
 - ✓ Presence of brain metastasis
 - ✓ Time from advanced/metastatic diagnosis to the start of index therapy
 - ✓ Disease stage at initial diagnosis

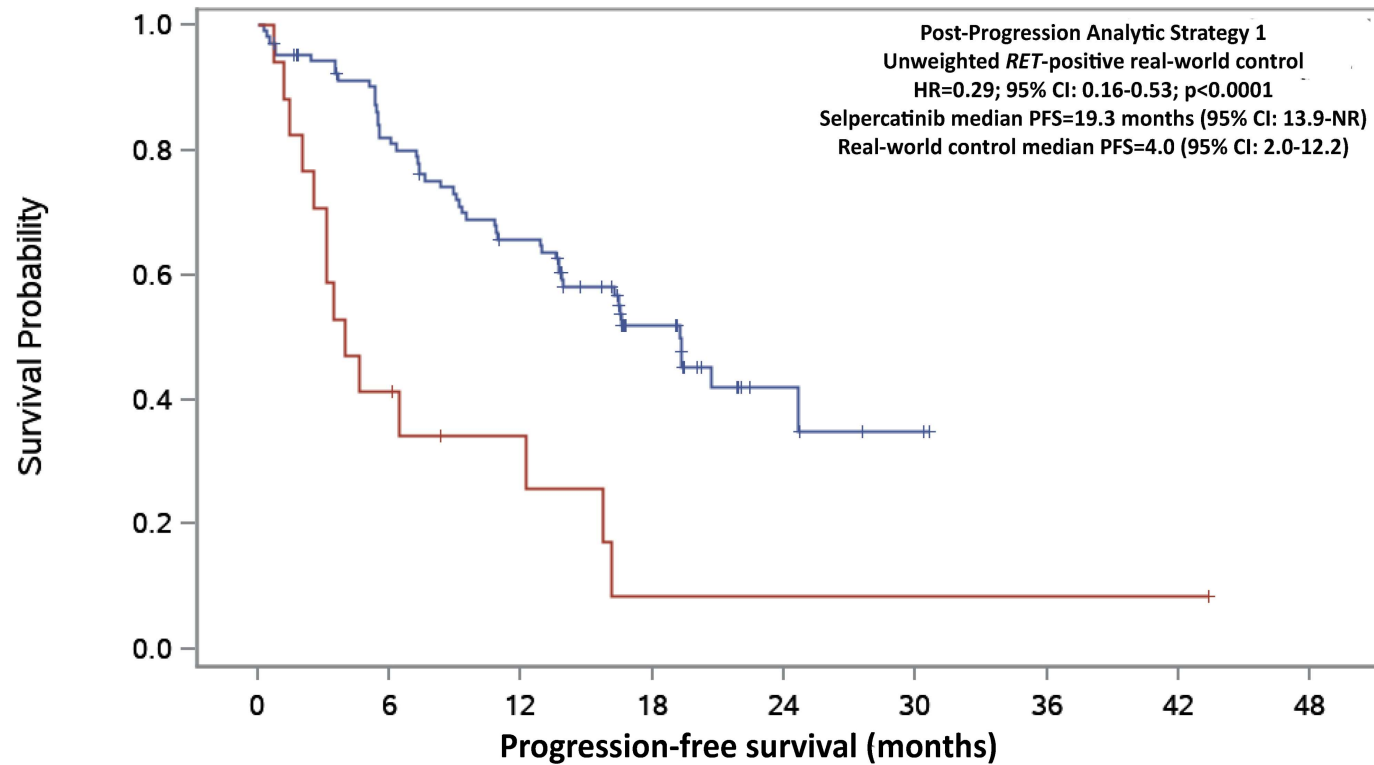


Patients included in analyses

	Real-world/ post- progression	LIBRETTO-001/ post- progression
Analytic strategy 1 (RET+)	17	105
Analytic strategy 2 & 3 (Any RET status)	1,503	

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PFS, post-progression (RET+ cohort)

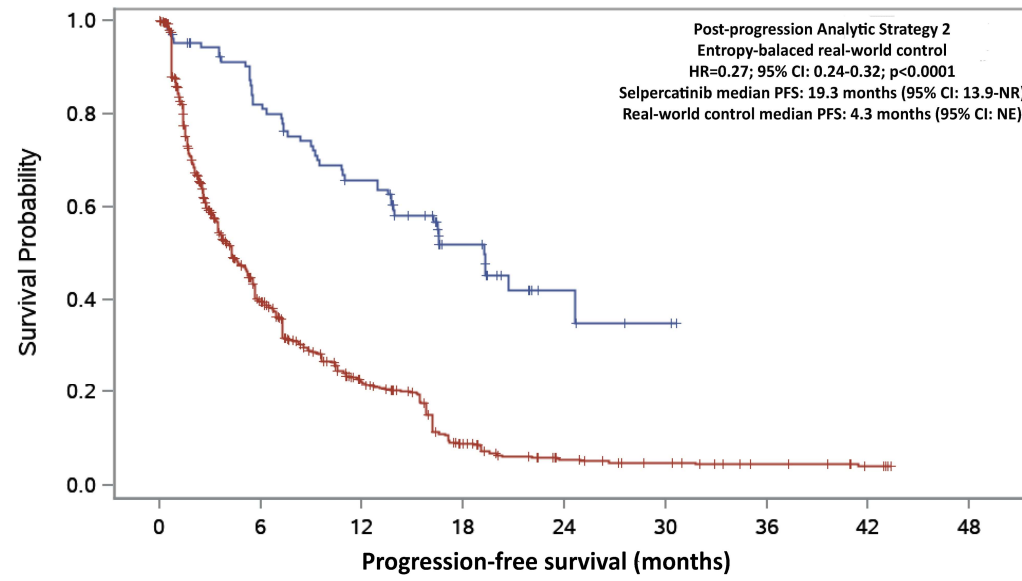


Selpercatinib (PAS)	105	81	63	27	6	3	0		
Real-world control	17	7	4	1	1	1	1	1	0

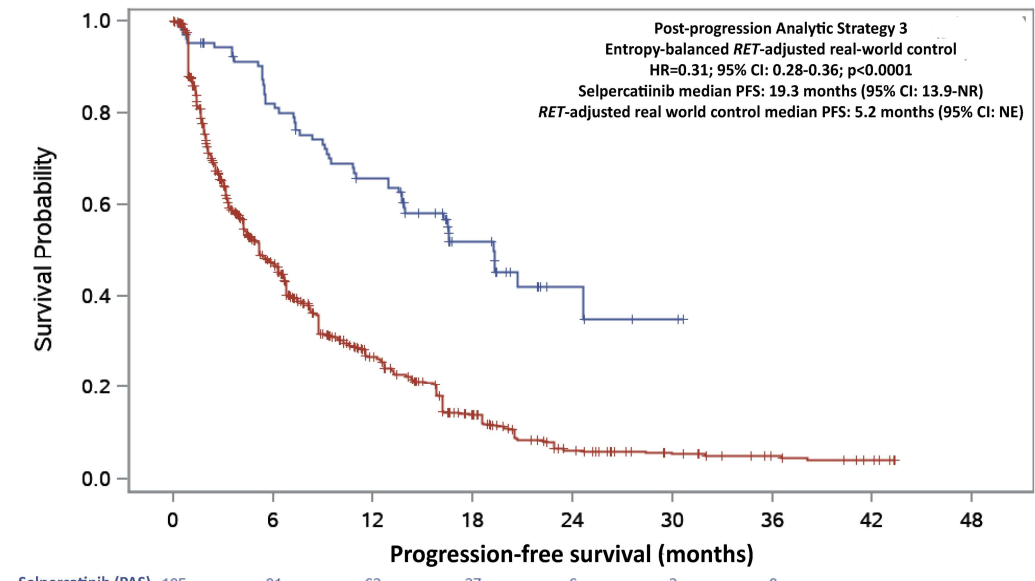
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PFS, post-progression (all patients)



Selpercatinib (PAS)	105	81	63	27	6	3	0		
Real-world control	1497	557	272	76	30	26	5	0	0



Selpercatinib (PAS)	105	81	63	27	6	3	0		
Real-world control	1497	666	333	126	35	29	6	2	0

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Synthesize external control arm

Part 2: Historical trial control

Objectives

- Compare efficacy (PFS and OS) of LIBRETTO-001 pre-treated NSCLC cohorts to relevant SOC's
 - ✓ Generalisability (i.e RET status adjustment)
 - ✓ Synthesize external control arm (used REVEL trial)
 - ✓ Compare to standard of care(s)

High level summary of the approaches

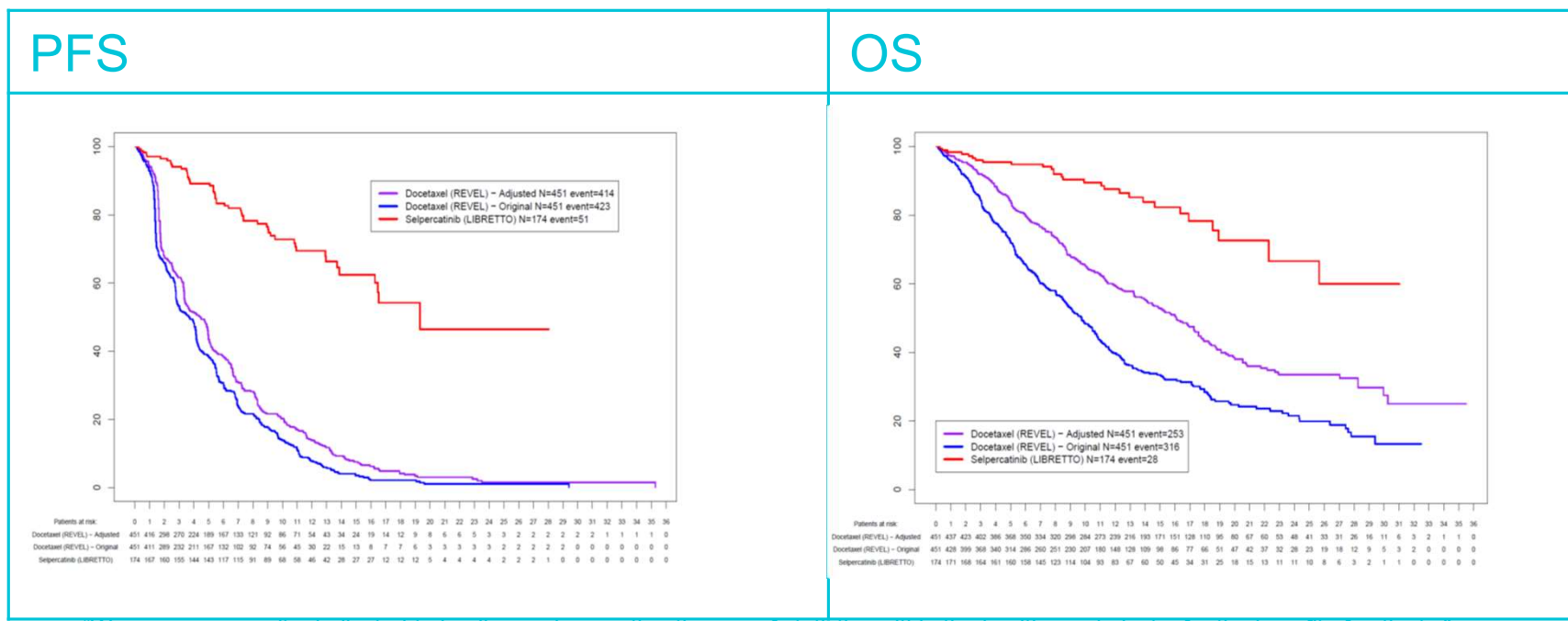
- **Generalisability**
 - ✓ Utilise Flatiron CGDB data to provide an estimate of a **time acceleration factor** for RET+, using multivariable statistical techniques.
 - ✓ Utilise the time acceleration factor to adjust the historical trial data
- Synthesize external control arm
 - ✓ Fit models to the adjusted historical trial data combined with the LIBRETTO-001 data to estimate an adjusted data set matched to the LIBRETTO-001 study
- Compare to standard of care(s)
 - ✓ Trial design and population differences of SOC(s) (i.e. biomarkers) cross trials via a hybrid Bayesian NMA approach



Adjusted historical trial data to reflect a RET+ population

- Estimated effect of RET-fusion status of RET+ vs RET- on PFS and OS from Flatiron data

Survival model	Post-progression PFS	Post-progression OS
Time acceleration factor	1.2	1.65



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High level summary of the approaches

- Generalisability
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 - ✓ Fit models to the adjusted historical trial data combined with the LIBRETTO-001 data to estimate an adjusted data set matched to the LIBRETTO-001 study
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Summary of Patient Characteristics of the REVEL and LIBRETTO Trial Populations



Characteristic	REVEL (n = 625)	LIBRETTO PAS (n = 105)	LIBRETTO IAS (n = 184)
Age (median)	62	61	62
Female	33.6%	59.0%	57.1%
Race: White	80.5%	52.4%	46.7%
Race: Asian	13.8%	38.1%	44.6%
Race: Other	5.70%	9.50%	8.70%
Never smoked	22.6%	71.4%	67.9%
Histology: Non-squamous	71.6%	100%	100%
ECOG \geq 1	67.9%	70.5%	64.2%
Prior surgery	18.1%	47.6%	45.7%
Stage IV at diagnosis	81.8%	96.2%	92.4%
Time since diagnosis to start of trial (median months)	9.2	30.1	24.2
Sum of longest diameters of tumours (mm) (median)	67	60.0	54.7
\geq 2 metastatic sites	86.4%	NR	NR
CNS metastases at baseline	3.8%	35.2%	32.6%

CNS = central nervous system; ECOG = Eastern Cooperative Oncology Group; IAS = integrated analysis set (all patients treated with platinum-based chemotherapy); NR = not reported; NSCLC = non-small cell lung cancer; PAS = Primary Analysis Set (the first consecutively enrolled patients previously treated with platinum-based chemotherapy).

Methodologies applied for comparative efficacy estimation



- **Matching methods (Multivariable regression, 1:1 matching ratio)**
 - a. Propensity score matching (on logit of the propensity score scale)
 - b. Genetic matching (weights assigned to each covariates to find the optimal balance)
- **Propensity score weighting (PSW)**
 - a. Based on a generalized boosted model
(an iterative process with multiple regression trees, $w_i=1$ for Libretto-001 cohort and $w_i = p_i/(1-p_i)$ for REVEL cohort)
 - b. Based on a logistic regression model ($w_i=p_i$ for REVEL cohort)
- **Doubly robust - targeted minimum loss-based estimation (TMLE)**

Baseline characteristics of patients treated with selpercatinib and clinical trial controls, before and after matching



Characteristic	Before adjustment or matching, post-progression setting		After matching using propensity scoring	After matching using a genetic algorithm	After PSW using a generalized boosted model	After PSW using logistic regression
	Selpercatinib cohort (LIBRETTO-001) <i>n</i> = 218 ^{a,b}	Docetaxel cohort (REVEL) <i>n</i> = 447	Docetaxel cohort (REVEL) <i>n</i> = 207	Docetaxel cohort (REVEL) <i>n</i> = 207	Docetaxel cohort (REVEL) <i>n</i> = 120	Docetaxel cohort (REVEL) <i>n</i> = 82
Age (mean, years)	58.75	59.83	59.03	59.93	59.61	59.0
Female, %	59.2	38.4	43	65.21	55.9	48.0
Race: White, %	53.4	79.1	58	49.28	53.3	52.6
Race: Asian, %	38.8	14.2	29	41.54	36.1	31.7
Race: Other, %	7.8	6.7	13	9.18	10.6	9.8
Never smoker, %	71.8	25.9	53	60.39	60.6	54.8
Stage IV, %	96.1	86	94	92.75	93.0	92.5
ECOG PS ≤ 1, %	72.8	68.3	63	61.84	66.0	64.1
Time since diagnosis to start of trial (median months)	36.63	12.04	15.61	31.08	22.61	17.6

ECOG PS, Eastern Cooperative Oncology Group performance status; PSW, propensity score weighting.

^aFour patients without non-squamous histology were excluded from further matching process.

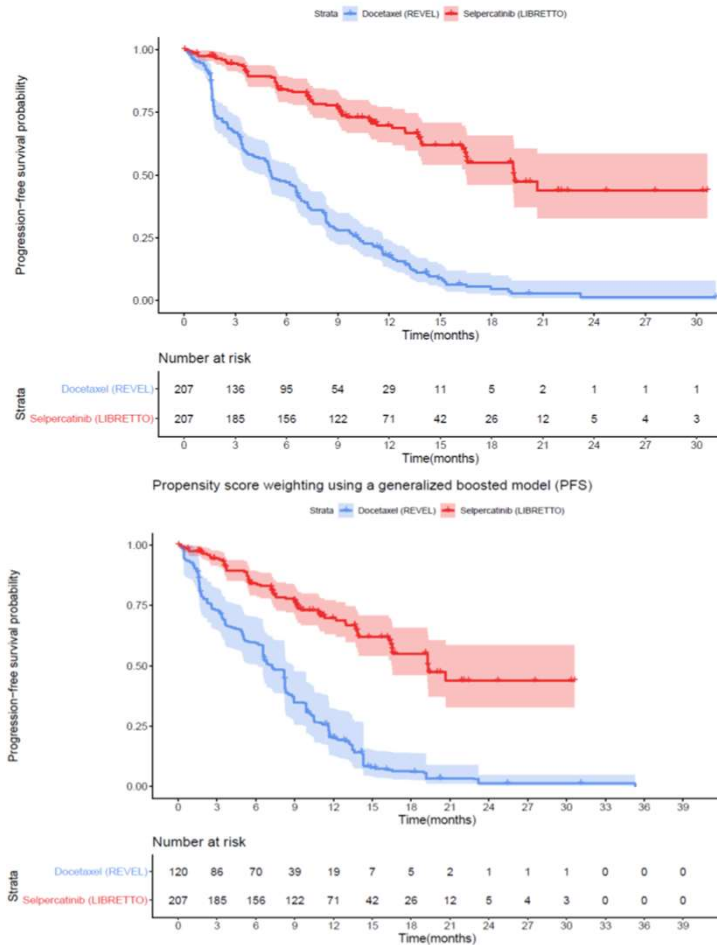
^bFive patients with ECOG PS ≥ 2 and 6 patients without non-squamous histology were excluded from further matching process.

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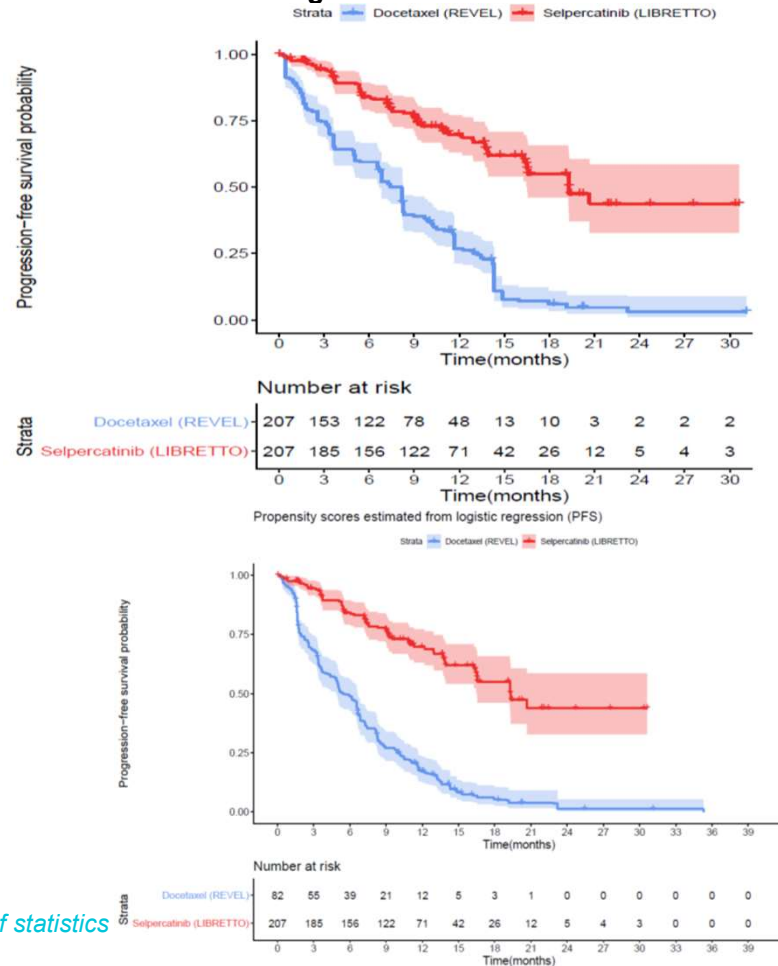
Estimation of a Control Arm (REVEL) using methodologies



- Multivariable regression: Propensity score matching

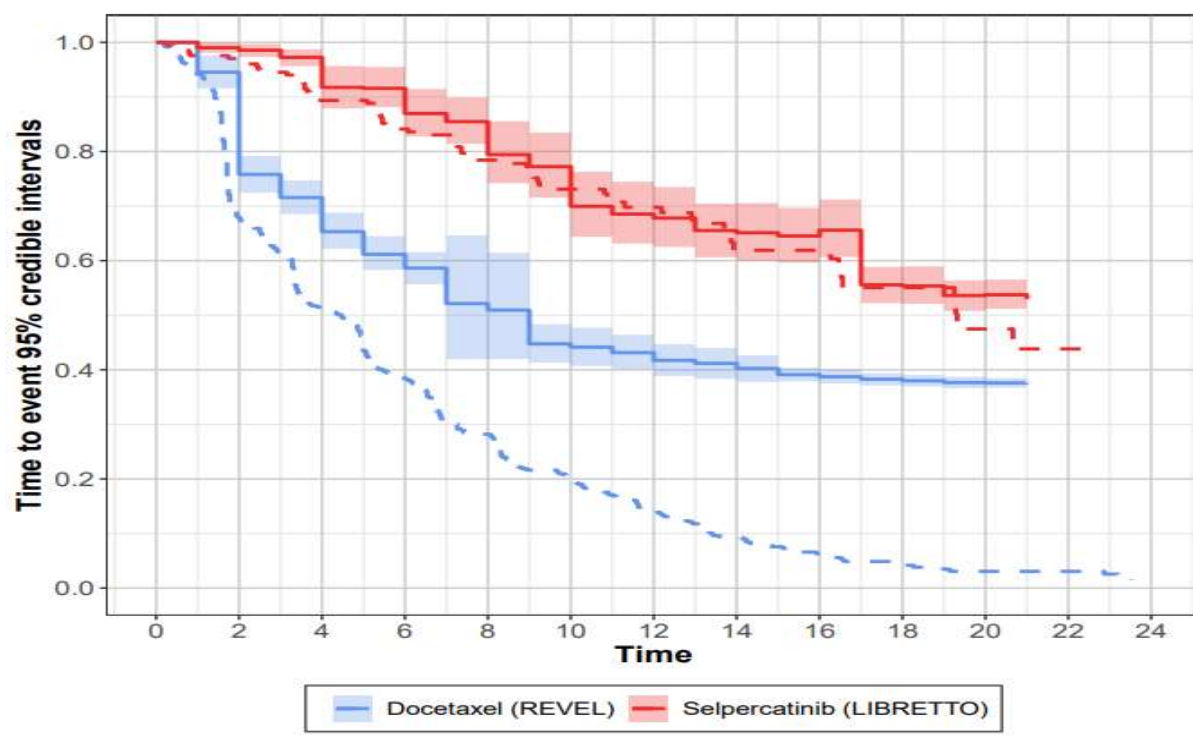


- Multivariable regression: Genetic



g the use of statistics

Estimation of a Control Arm (REVEL) using TMLE



PFS Cox HR = 0.39, 95% CI: 0.29-0.52, $p = <0.0001$

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Results - PFS

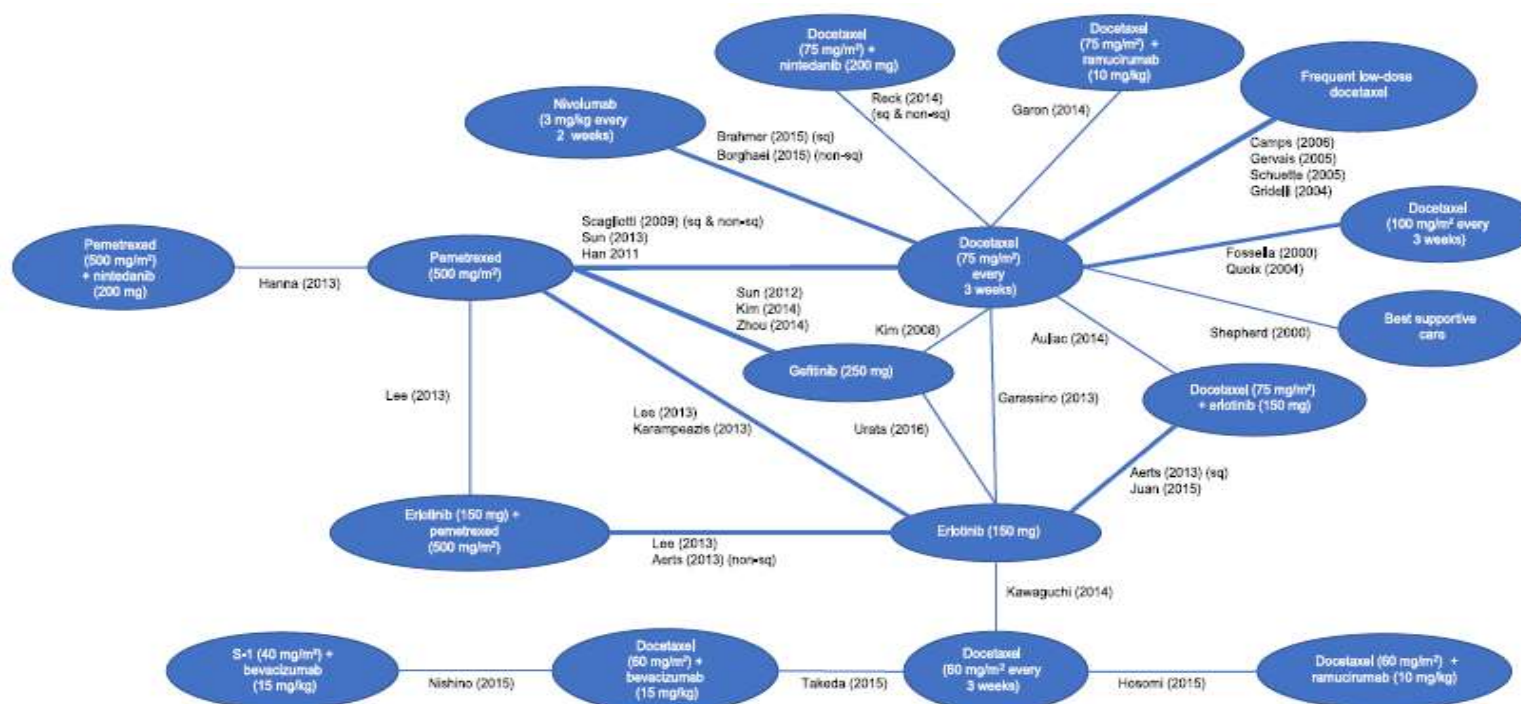
Methodologies	HR	95%CI	P-value
Targeted minimum loss-based estimation (TMLE)	0.39	0.29 - 0.52	< 0.001
Multivariable regression: Propensity score matching	0.21	0.16 - 0.27	< 0.001
Multivariable regression: Genetic matching	0.27	0.20 - 0.35	< 0.001
Propensity score weighting: a generalized boosted model	0.24	0.17 - 0.32	< 0.001
Propensity score weighting: a logistic regression	0.24	0.17 - 0.32	< 0.001

High level summary of the approaches

- Generalisability
 - ✓ Utilise Flatiron CGDB data to provide an estimate of a **time acceleration factor** for RET+, using multivariable statistical techniques.
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Clinical Data for Comparators

- Survival Network Meta-analysis (Vickers et al., 2019) has many trials investigated specifically **second-line treatment**
- An update of this network meta-analysis (NMA) had been planned, including more recent relevant trials
 - LIBRETTO-001 also included, after estimation of a control arm



Clinical Data for Comparators

The barriers for Cyramza NSCLC indication

- To incorporate differences across populations & trial designs

Ex. The following comparators were specified in the NICE (UK payer agency) scope for Cyramza NSCLC indication:

Abbreviation	Full name and dose	Patient subpopulation
RAM + DOC	Ramucirumab 10 mg/kg + Docetaxel 75 mg/m ²	All populations
NIN + DOC	Nintedanib 200 mg + Docetaxel 75 mg/m ²	Non-squamous
ERL	Erlotinib 150 mg	EGFR-negative*
DOC	Docetaxel 75 mg/m ²	All populations
NIV	Nivolumab 3mg/kg	PD-L1 expression level, Non-squamous/squamous

REVEL Direct evidence

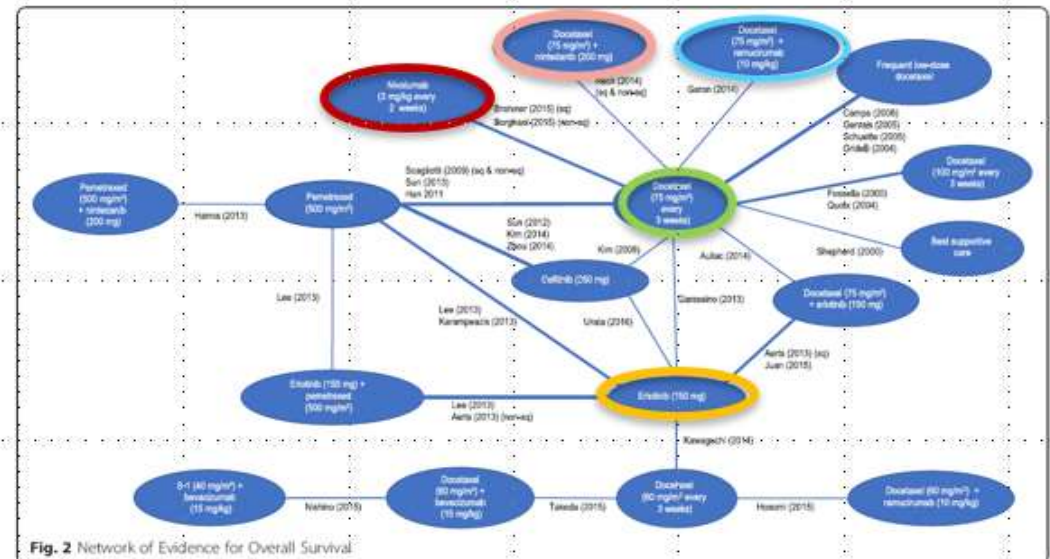


Fig. 2 Network of Evidence for Overall Survival

Clinical Data for Comparators – pre-treated *RET*+ NSCLC

Solution

- A NMA of hybrid approach of **fractional polynomials & hierarchical exchange model** was performed to allow different interaction between treatment and population type, where:
 - ERL efficacy was allowed to vary by EGFR status.
 - NIN + DOC efficacy was allowed to vary by squamous histology
 - NIV efficacy was allowed to vary by histology and PD-L1 level

A Webinar on 27th of July 2021 by NICE.....

Hierarchical Model

- Assumes exchangeability between parameters informed by RCT and NRS
- Obtain estimate of overall relative treatment effects across both types of evidence
 - Also obtain model-based estimate of relative treatment effect for RCT evidence, accounting for the NRS evidence under the model assumptions (shrunk estimate)

DSU Recommendations

- Careful consideration of **whether the observational data are sufficiently credible and how the results should be interpreted** is required.
- A **bias-adjusted base case** should be used with **other methods considered as sensitivity analyses**.
- Methods that attempt to **down-weight and adjust the observational evidence before inclusion in the synthesis are preferred (hierarchical model, design-adjusted analysis)**.
- **Naive pooling of randomised and non-randomised evidence is not recommended**, although it may be useful as a **first step analysis**, or as a **sensitivity analysis**.

A Medical Research Council (MRC) funded project is underway

NICE

Methodology evolution...



- NICE evaluation research group (ERG) considered that the company provided appropriate justification for using hierarchical models
- NICE accepted the clinical effectiveness of Cyramza submission and **grant end-of-life criteria; willingness to pay threshold increased.**

- The work has been published in BMC Cancer 2019

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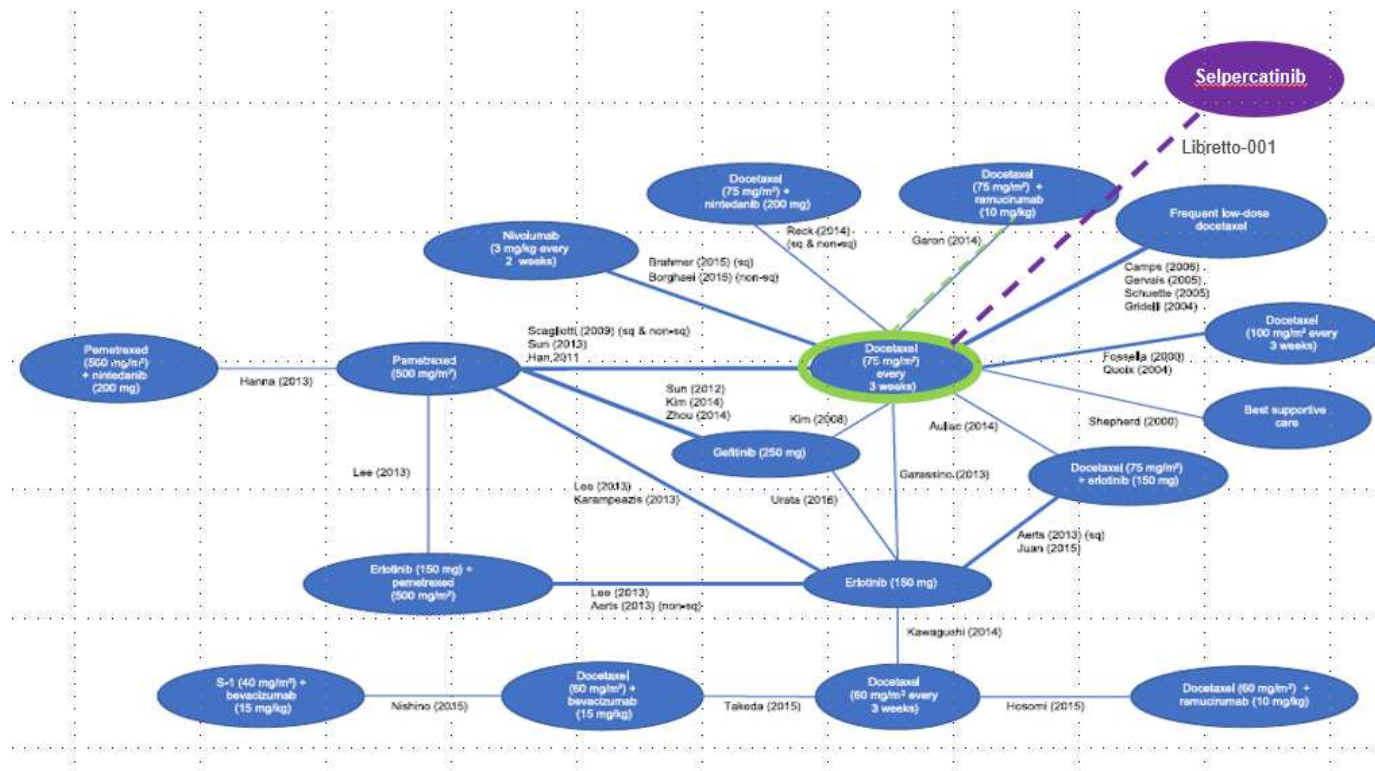
Relative efficacy of interventions in the treatment of second-line non-small cell lung cancer: a systematic review and network meta-analysis

Adrian D. Vickers¹, Katherine B. Winfree², Gebra Cuyun Carter², Urpo Kiskinen³, Min-Hua Jen⁴, Donald Stull⁵, James A. Kaye⁶ and David P. Carbone⁷



Clinical Data for Comparators

- Applied NMA of hybrid approach of **fractional polynomials & hierarchical exchange model**



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Prediction vs Actual

Lilly's Retevmo® (selpercatinib) Phase 3 Results in RET Fusion-Positive Non-Small Cell Lung Cancer and Medullary Thyroid Cancer Both Published in *The Journal of Clinical Oncology* and *Journal of Clinical Endocrinology and Metabolism* and Presented in a Presidential Session at the American Society of Clinical Oncology (ASCO) Annual Meeting in Chicago, Illinois, June 1-5, 2023.

PFS	Predicted (2019)	Actual (2023)
HR	0.424 (0.289, 0.623)	0.465 (0.309, 0.699)

October 21, 2023



- In the Phase 3 LIBRETTO-431 study, Retevmo more than doubled progression-free survival (PFS) compared to chemotherapy plus pembrolizumab in patients with advanced or metastatic rearranged during transfection (RET) fusion-positive non-small cell lung cancer (NSCLC)



- In the Phase 3 LIBRETTO-531 study, Retevmo provided a 72% improvement in PFS compared to cabozantinib or vandetanib in patients with advanced or metastatic RET-mutant medullary thyroid cancer (MTC)

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Limitations

- These are exploratory and hypothesis generating, suggesting what may be observed in a RCT, but should be confirmed in that setting
- Real-world evaluation of response does not adhere to RECIST and may measure different concepts
- Sample sizes for Analytic Strategy 1 were small, there were few events, and analyses were limited
- Weighting factors for these smaller samples could have influenced these results (inflated or decreased p values)
- Estimating a control arm in a different patient population requires several steps and each has a source of error

Conclusions

- **Clinical**
 - ✓ Despite the limitations, the consistency of results across multiple analyses of the RW and clinical trial control cohorts
 - ✓ While there may be a unique prognostic effect of RET fusion + status, it is unlikely to be strong enough to explain the large differences in observed outcomes between Libretto-001 and the alternative RW and clinical trial control cohorts

Conclusions

- **Statistical**

- ✓ With careful plannings, the proposed two-fold approaches enabled single-arm trials to estimating comparative efficacy to compare indirectly with other available treatments with good precisions.
- ✓ Results also may be used to serve as a reference for the efficacy of existing treatments for patients with a particular tumor type, where only mixed population evidence so far exists.
- ✓ The hybrid approach should help inform the decision-making process for prescribing currently available treatments and could be used to help power future trials.

Reference

- Rolfo C, Hess LM, Jen MH, Peterson P, Li X, Liu H, Lai Y, Sugihara T, Kiiskinen U, Vickers A, Summers Y. External control cohorts for the single-arm LIBRETTO-001 trial of selpercatinib in RET+ non-small-cell lung cancer. *ESMO Open*. 2022 Aug;7(4):100551. doi: 10.1016/j.esmoop.2022.100551. Epub 2022 Aug 2. PMID: 35930972; PMCID: PMC9434413.
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