

Enhancing Interpretation of PSM through Advanced Visualization

Umayal Annamalai, KiranKumar Vyas

VeraMed



The views and opinions expressed in this presentation are solely those of the presenter and do not necessarily reflect the official policy, position, or endorsement of the company or any affiliated organization. This content is intended for educational and informational purposes only.

Agenda

01 Why PSM?

02 Challenges

03 Advanced Data Visualization

04 Case Study

05 Conclusion & Future Directions

Why PSM?

Balance Groups

Align treatment & control; adjust baseline differences

Reduce Bias

Control confounders; improve causal inference

Real-World Data

Simulate RCT-like comparability; enhance credibility

Confounding

Address systematic differences

Match by Score

Pair subjects using treatment probability

Challenges

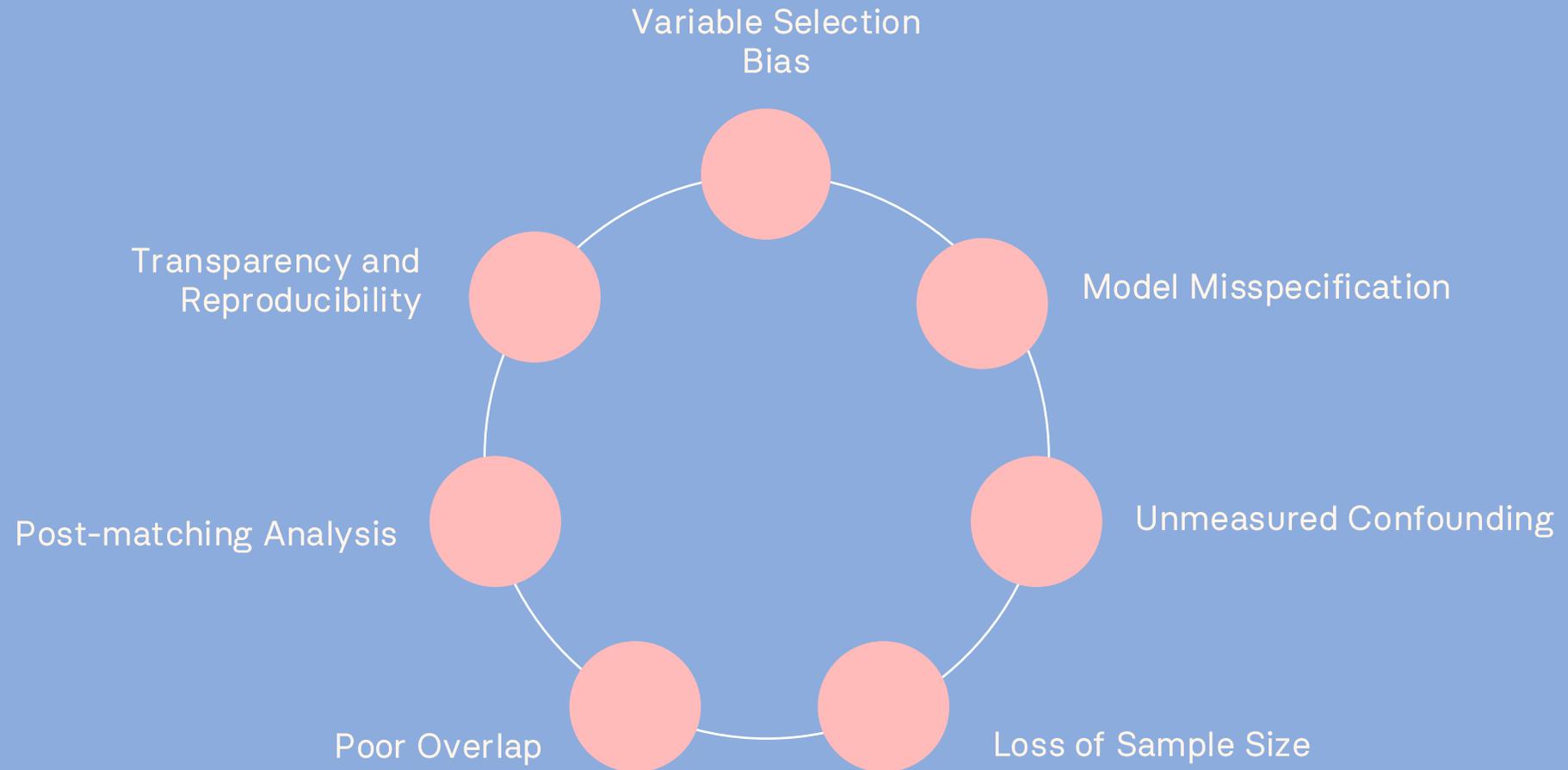


Table versus DV

“

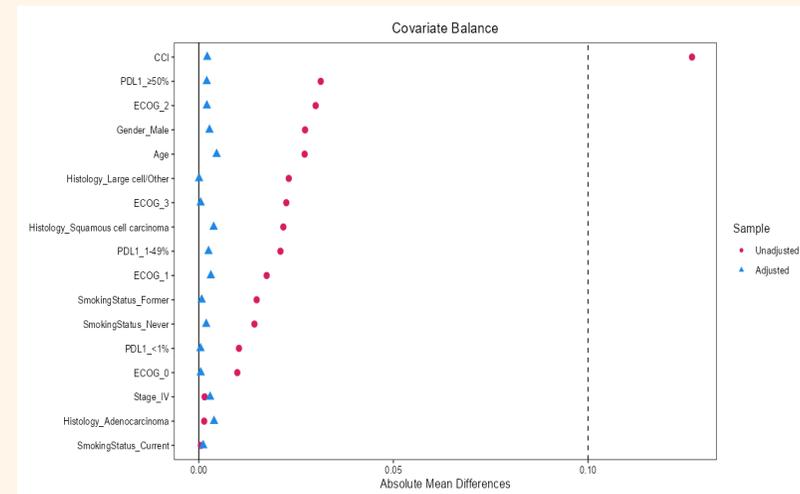
The greatest value of a picture is when it forces us to notice what we never expected to see.

— John W. Tukey

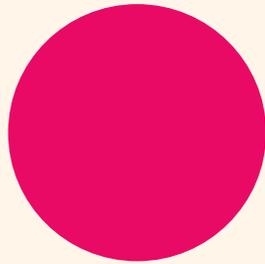
Table

Covariate	Type	Diff.Un	Diff.Adj	M.Threshold
Age	Contin.	-0.02720993422813928	-0.006538569547937281	Balanced, <0.1
Gender_Male	Binary	0.02731315734799511	-0.00206431143886121	Balanced, <0.1
SmokingStatus_Current	Binary	-0.0005720978829367174	-0.001653912697153925	Balanced, <0.1
SmokingStatus_Former	Binary	0.01488056703933316	0.00009258407667633861	Balanced, <0.1
SmokingStatus_Never	Binary	-0.01430846915639651	0.001561328620477545	Balanced, <0.1
Histology_Adenocarcinoma	Binary	-0.00140013429245045	0.003256958117260123	Balanced, <0.1
Histology_Large cell/Other	Binary	0.0231308207195787	0.0005240034172288785	Balanced, <0.1
Histology_Squamous cell carcinoma	Binary	-0.02173068642712828	-0.003780961534489036	Balanced, <0.1
Stage_IV	Binary	-0.001517564910526903	-0.002915147561267095	Balanced, <0.1
ECOG_0	Binary	-0.00991234858225111	-0.0004669896477470514	Balanced, <0.1

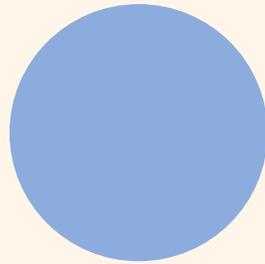
DV



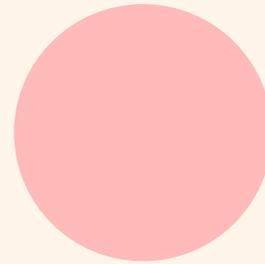
Addressing Limitations of PSM Through Visual Diagnostics



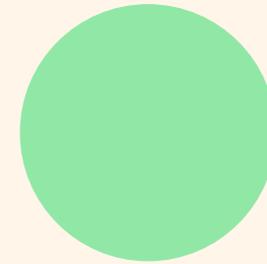
Reveals imbalanced covariates post-matching that may require re-specification.



Highlights poor propensity score overlap, reducing the risk of extrapolation bias.



Detects extreme weights in PS weighting that can distort the sample or inflate variance.



Supports sensitivity analyses with visual evidence.

The Role of Advanced Visualization techniques in Enhancing PSM Validity



Visualization bridges the gap between statistical outputs and clinical understanding.



Offers a transparent way to evaluate matching quality and model assumptions.

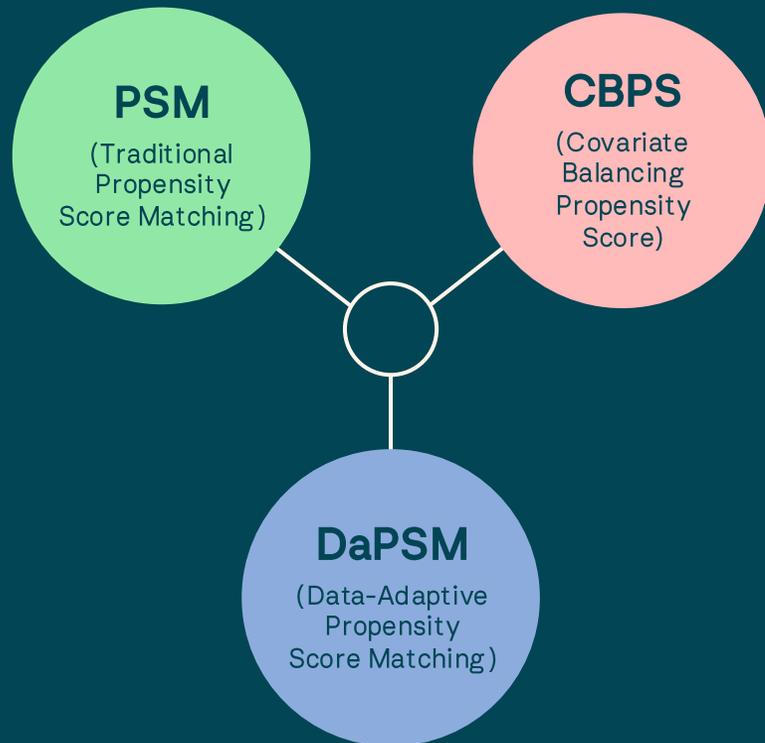


Helps uncover hidden flaws such as poor overlap, imbalance, or overfitting.



Clear communication of complex PSM results to diverse stakeholders through intuitive visual summaries.

PSM Methods and Techniques



Techniques

Nearest Neighbour matching

Each treated unit is matched with a control unit having the closest propensity score. Focuses on similarity in treatment probability based on characteristics

Caliper Matching

A stricter version of NNM. Matches are accepted only if the propensity score difference is within a predefined threshold (caliper).

Inverse probability of treatment weightage

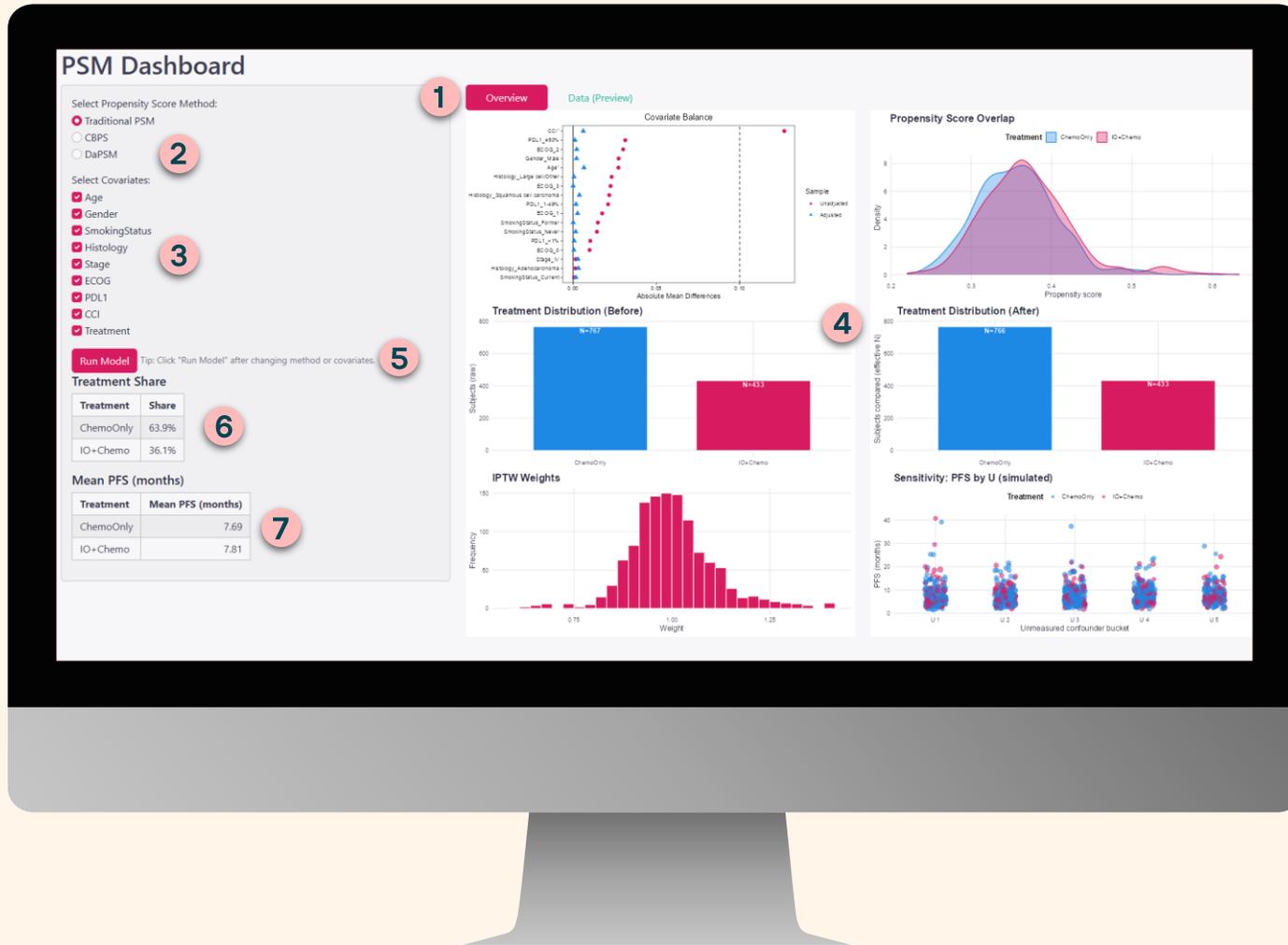
Uses weights instead of matching. Treated units are weighted by $1 / \text{propensity score (PS)}$. Control units are weighted by $1 / (1 - \text{PS})$.

The background features a dark teal gradient on the right side, transitioning to a lighter teal and then a soft pink on the left. Overlaid on this are several large, abstract, rounded shapes in shades of teal and light blue, creating a modern, layered effect.

Case study: PSM Advanced Visualizations

Dashboard Walkthrough

PSM Dashboard



1

Overview and Data (Preview) tabs showing all the plots that we have produced and original data collected.

2

PSM method selection area with radio buttons, which lets us to select the method using which we want to run our model.

3

These are list of all the possible covariates; we can include all or exclude and run the model.

4

Plot area, where we can see the plots generated.

5

Run button to run the selected model.

6

Table showing treatment share percentages of each group

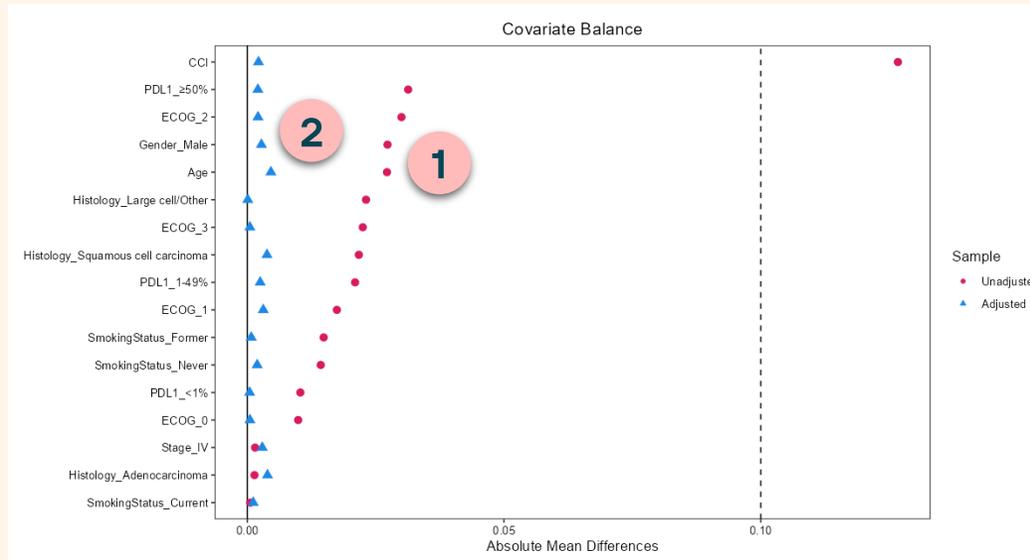
7

Calculated Mean PFS (in Months) for each treatment post weighting

PSM Interpretation with Advance Visualizations



Covariate Balance Plot



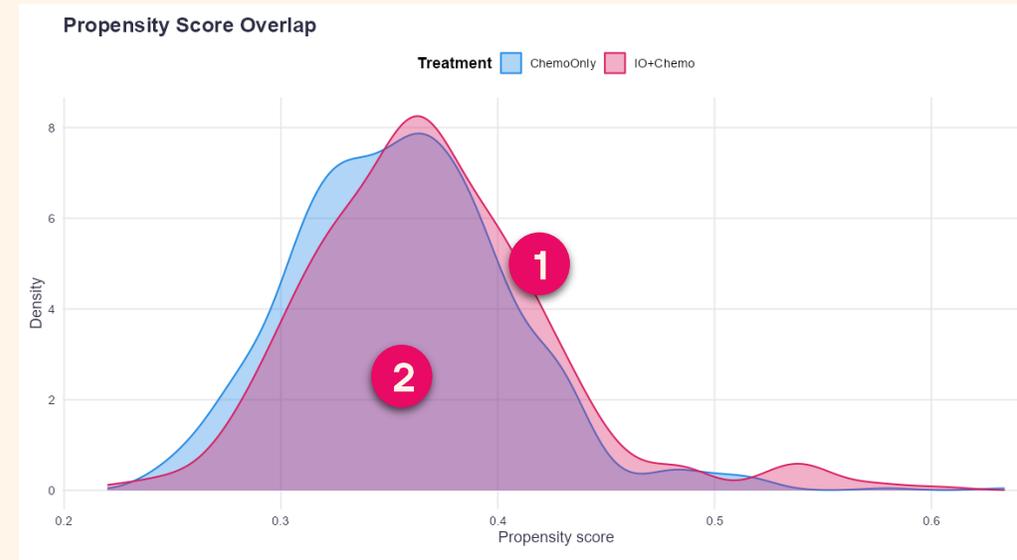
1

Large absolute mean differences before weighting

2

Adjusted mean differences after weighting

Propensity Score Overlap



1

Distribution of propensity scores.

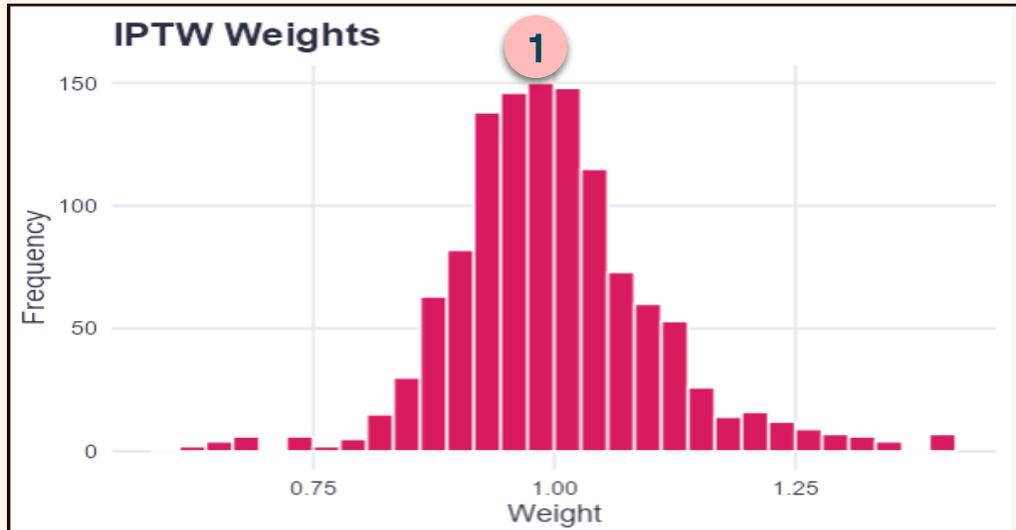
2

Good overlap = Effective & Comparable mapping

PSM Interpretation with Advance Visualizations



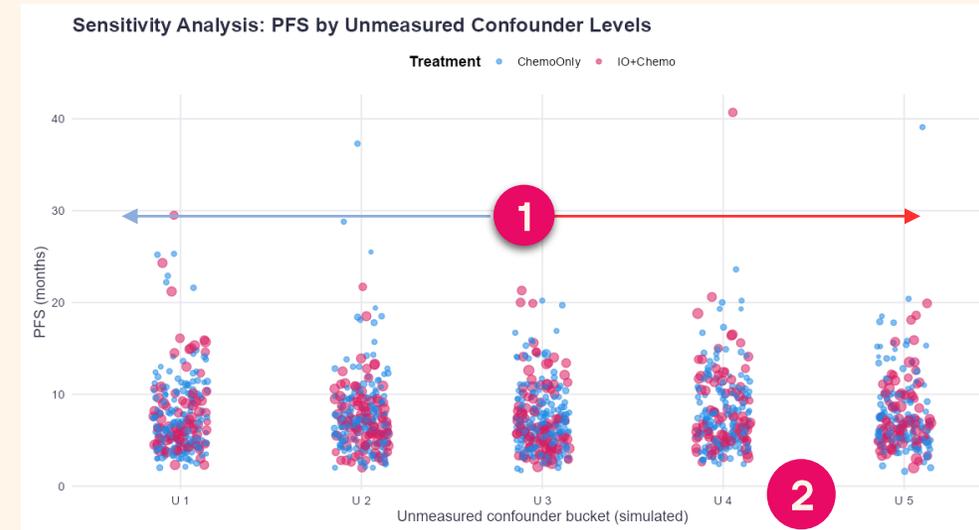
Weight Diagnostic Plot



1

A narrow, centered distribution of weights is always desired and ideal.

Sensitivity Analysis Plot (Rosenbaum Bounds plot)



1

Stable treatment effect/outcome (PFS) with changes in the strength/level of simulated unmeasured confounder.

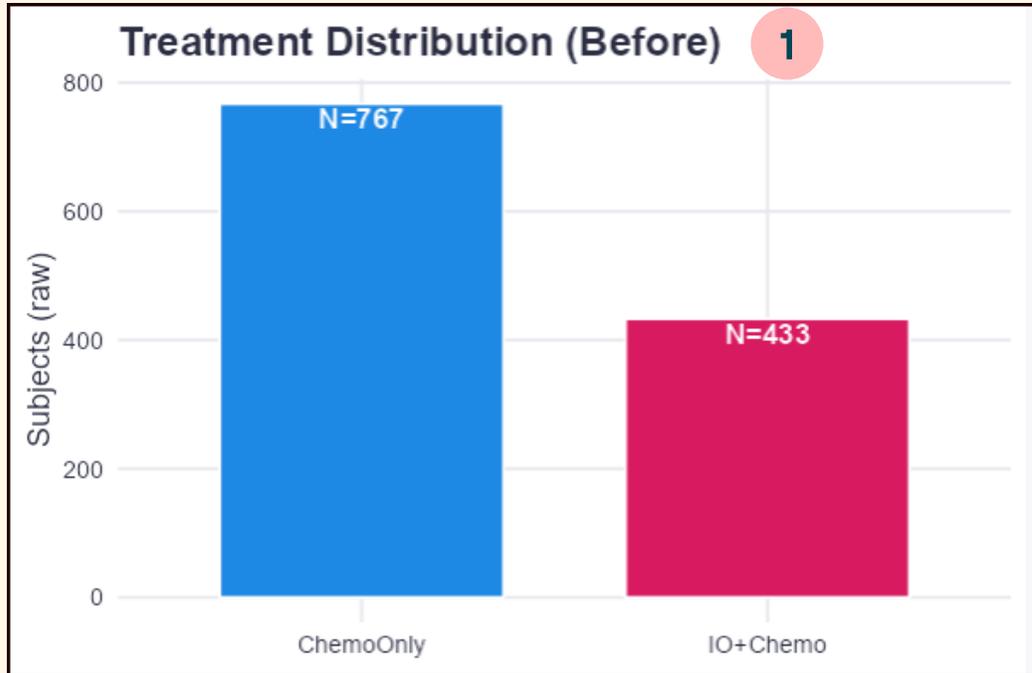
2

Levels of simulated unmeasured confounders on the Y axis.

PSM Interpretation with Advance Visualizations

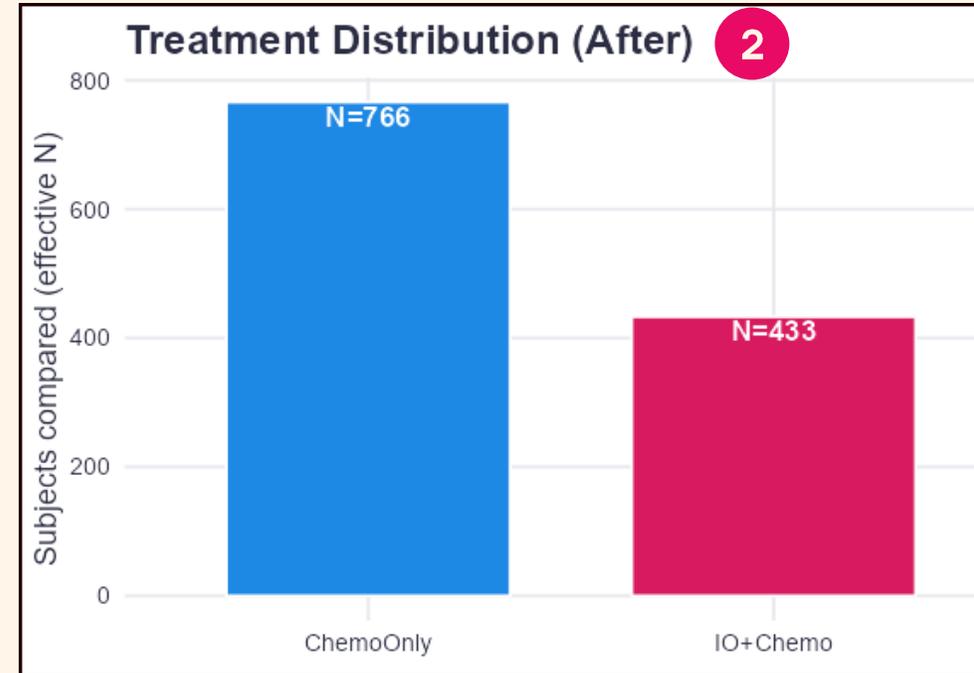


Treatment Group Distribution



1

Before Weighting: 767 ChemoOnly and 433 IO+Chemo subjects, unequal group sizes, common in real-world data.



2

After Weighting: We nearly retained sample size as per expectation. This is because weighting does not discard subjects; it re-weights them.

Outcomes



Diagnostic accuracy



Early detection of misinterpretation or overlooked bias



Streamlined process of model refinement and validation.



Simplified and Accessible to non-statisticians



Stakeholder Confidence and Communication



Transparency & Confidence in Evidence Quality



Proactive decision-making

Conclusion and Future Directions



Simplified black-box
statistical procedure



Regulatory-grade
Evidence quality



Stake holder
Collaboration



Machine learning–
enhanced PSM models



Integrated Regulatory-
focused visual
diagnostics into PSM
workflows



Tools for identifying
unmeasured confounding
analysis



Thank you

Reach out to us
umayal.annamalai@veramed.com
kirankumar.vyas@veramed.com