

Development of Biomarkers to Assist New Drug Development for CNS Disorders

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Outline



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 - Biomarkers in new drug development
 - Biomarkers for new drug development in patients with CNS disorder
- Case Examples
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Biomarkers in New Drug Development



- Biomarkers are a key <u>medical product development tool</u> capable of facilitating development of medical products and spurring innovation.
 - Lab values, medical imagines, pharmacodynamic assessment, clinical outcomes, etc.
- Examples of usage in drug development:
 - Patient enrichment:
 - Patient selection:
 - Bridging biomarkers:
 - Surrogate endpoint:
 - Reasonably likely surrogate endpoint
 - Validated surrogate endpoint
- Quantitative/data science tools for biomarker selection

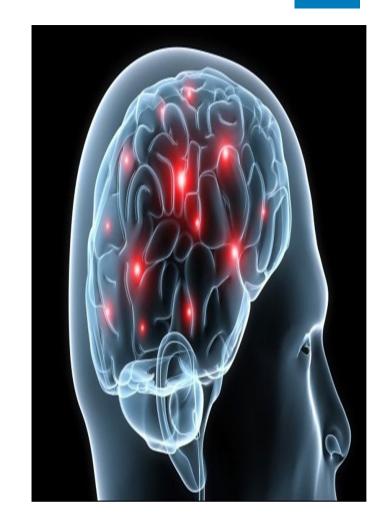
PK/PD Exposure-Response In Silico Biomarker- Clinical Trial • Clinica **Simulations** Respons **MIDD** Disease Models AI / ML Clinical Trial Models Systems **Biology** • QSP CiPA

Source: Biomarkers at FDA | FDA

Biomarkers for Drug Development in Patients with CNS Disorder



- Various biomarkers are evaluated in drug development programs for patients with CNS disorders.
- Most biomarkers have been used to inform target engagement, upstream/downstream changes, proof of efficacy, dose selection, and patient selection.
- Limited examples for CNS biomarkers being considered as surrogate endpoints that can be used to support accelerated approval/approval.
- Given the complexity of CNS disorders, use of multimodal biomarkers may provide better assistance for future new drug development.



Review Example: Drug B



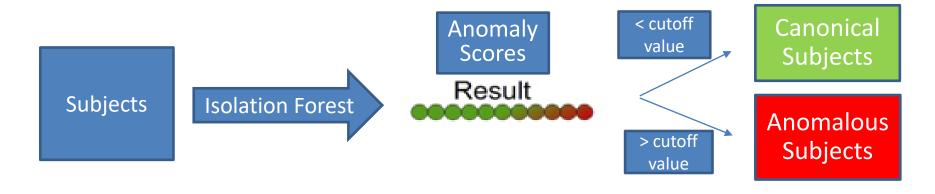
Context of Use:

- Drug B failed to reach statistical significance in prespecified primary efficacy analysis for the Phase 2 trial in patients with bipolar depression.
 - Unexpectedly large improvement in the primary efficacy endpoint from baseline for the placebotreated cohort.
 - This primary efficacy endpoint was a composite endpoint with many items.
 - Motivation to explore ML method to identify patients with abnormal patterns.
- The sponsor is proposing an ML-based enrichment strategy for two planned Phase 3 studies
 - Decrease inter-subject variability prior to randomization:
 - Select canonical subjects whose symptoms are characterized by greater similarity to the typical patient in the target population.

Review Example: Drug B (continued)



- The dataset for machine learning (ML) development includes:
 - The Phase 2 trial of Drug B
 - Multiple trials of Drug C for the same indication
- The anomaly score is calculated by applying a trained isolation forest model on the components of the efficacy endpoint at screening and baseline.
 - Patient characteristics include
 - The anomaly score for a given subject indicates how easily this subject can be isolated from the rest of the subjects.



Review Example: Drug B (continued)



- Encouraging trend was observed when ML-based inclusion criterion was applied retrospectively to the phase II data of drug B
 - Larger effect size
- In general, similar trends were observed when it was applied to trials of drug C
- Conclusion:

FDA:

Suggest using the selected criterion in only one of the two planned adequate and well-controlled studies.

Due to the potential challenges related to the **explainability** of the model and the **generalizability** of the results, it would be helpful to include the "anomalous" patients in at least one of the adequate and well-controlled studies.

Research Example: Enrichment Strategy







Identify
biomarkers
associated with
"placebo
responders"



Remove patients with biomarkers consistent with "placebo responders"



Compare treatment effect after removing patient subgroups Placebo responder:
patients receiving
placebo showing
improvement in clinical
symptoms at different
defined range (%
improvement)



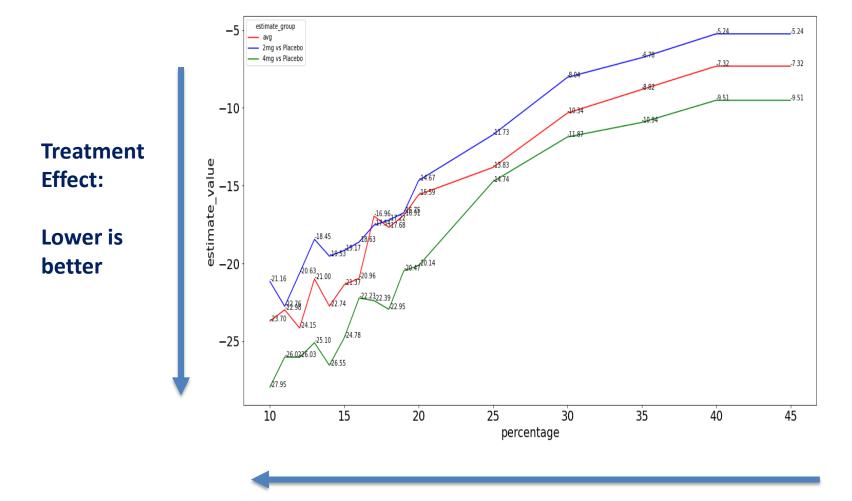




To improve future clinical trials

Research Example: Improved treatment effect





This trend appears to be consistent for all pivotal clinical trials (Phase 3 trials) from NDAs in schizophrenia patients.

General Considerations (Personal Opinion, Still Evolving)



- Data Source:
 - Meta analysis: (e.g., endpoints, lab values, patients, biomarker measurement, clinical outcome assessment, standard of care, etc)
 - Data standard (e.g., variables, format, etc), data transformation, handling of missing data and outliers...
- Model Development:
 - model or covariate (patient feature) selection
 - Model validation
 - Assumption check
- Model Interpretation:
 - Generalizability (critical for AI/ML Models)
 - Training data should be unbiased and diverse/inclusive
 - Methods need to be developed for performance guarantee
 - Transparency/interpretability/explainability (Important for empirical models)
 - Methods are being developed to improve interpretability/explainability
- We need to develop best practice

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