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## SMALL MOLECULE BIOMARKER ANALYSES BY LC-MS

SMALL MOLECULE BIOMARKER LC-MS ASSAYS DESIGNED TO QUICKLY DELIVER FIT-FOR-PURPOSE STUDY DATA THAT MEETS OR EXCEEDS DATA QUALITY AND EXPERIMENTAL OBJECTIVES FOR NON-REGULATED STUDIES. Bioanalysis of biomarkers presents multiple, inherent challenges including, separation of isobaric interferences, instability/interconversions of analytes in biological matrices, limited sample volume for use across several assays, and quantification of concentrations below naturally occurring endogenous levels. The scientists at Meadowhawk Biolabs are well versed in applying flexible and innovative experimental approaches to effectively address these challenges for both simple and complex assays across a wide range of matrices including plasma and tissue homogenates (Refer to Table below).

| LC-MS Biomarker Assay Design Considerations |                          |  |  |   |  |  |  |
|---|--------------------------|--|--|---|--|--|--|
| Biomarker<br>Modulation                     | Experimental<br>Approach | Blank Matrix   | Calibration<br>Standards & QCs                                     | Assessments   |  |  |  |
| Up or Down<br>Regulation                    | Stripped Matrix          | Authentic biological matrix modified to<br>eliminate target endogenous compound(s)<br>(e.g. activated charcoal)                              | Actual Biomarker<br>Analytical Reference<br>Standard(s)            | <ul> <li>Confirm absence of analyte across multiple preparations/lots</li> <li>Create matrix pool for consistency between batches and studies</li> </ul>  |  |  |  |
|   | Surrogate<br>Matrix      | Target compound free substitute matrix<br>mimicking the behavior of the biological<br>sample matrix (e.g. Bovine Serum Albumin<br>in buffer) | Actual Biomarker<br>Analytical Reference<br>Standard(s)            | <ul> <li>Establish surrogate and authentic matrices equivalency through<br/>dilutional linearity and spike recovery experiments</li> <li>Prepare QCs in surrogate, authentic or combination of matrices<br/>based on endogenous analyte levels</li> </ul> |  |  |  |
|   | Surrogate<br>Analyte     | Study sample matched, authentic<br>biological matrix   | Targeted Biomarker<br>Stable Isotope Labeled<br>Reference Standard | <ul> <li>Evaluate and balance LC-MS responses of native biomarker and<br/>stable isotope reference standard</li> <li>Demonstrate parallelism of calibrants prepared by authentic and<br/>surrogate analytes</li> </ul>                                    |  |  |  |
| Up Regulation<br>(≥ 3-fold<br>change)       | Standard<br>Addition     | Study sample matched, authentic<br>biological matrix   | Actual Biomarker<br>Analytical Reference<br>Standard(s)            | <ul> <li>Demonstrate acceptable levels of endogenous background<br/>concentrations versus quantification range</li> <li>Create matrix pool for consistency between batches and studies</li> </ul>   |  |  |  |

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To meet study objectives at points along the drug discovery continuum, Meadowhawk offers a tiered, fit-for-purpose approach to best manage time, costs and complexity. Meadowhawk's facilities feature state-of-the-art instrumentation, including Sciex API 7500 triple quadrupole mass spectrometers paired with Shimadzu UPLC systems for ultimate sensitivity and maximum method versatility. Additionally, for our clients in both the Boston and San Francisco Biohubs, Meadowhawk provides free, same day courier services to simplify logistics and conserve time.

Count on the team at Meadowhawk to flexibly and creatively support your small molecule biomarker bioanalytical needs supported with exceptional customer service.



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|  |                       | Biomarker Assay Tier   | ring (LC-MS Platform)  |  |  |
|--|-----------------------|--|--|--|--|
| Attribute                              |                       | Exploratory  | Optimization   | Pre-IND  |  |
| Research Objective                     |                       | To qualitatively monitor fold-changes in up or down regulation of potential biomarkers                             | To provide quantitative data supporting<br>efficacy studies, PK/PD correlations and<br>other assessments                                 | To provide definitive quantitative data to<br>support the design and execution of<br>IND-enabling and clinical studies |  |
| Turnaround Time                        |                       | +  | ++   | +++  |  |
| Experimental Rigor                     |                       | ++   | +++  | ++++   |  |
| Cost                                   |                       | +  | ++   | +++  |  |
|  |                       | Biomarker Ass  | say Parameters   |  |  |
| Precision/Accuracy Criteria            |                       | Relative Quantification  | ±20% (±25% LLOQ)   | ±15% (±20% LLOQ)   |  |
| Analytical<br>Reference<br>Standard(s) | Reference Material    | Preferred if available (used for confirmation of peak identification)  | Required   | Required   |  |
|  | Salt Form             | Required   | Required   | Required   |  |
|  | Purity Information    | Not required, assume 100% in absence   | Not required, assume 100% in absence   | Required   |  |
|  | Quantity Required     | Pre-weighed or $\geq$ 2 mg (if applicable)   | Pre-weighed or $\geq$ 2 mg   | Pre-weighed or $\geq$ 10 mg  |  |
|  | Weighings             | Single (if applicable)   | Single   | Triplicate   |  |
|  | Stock Solution Checks | No   | No   | Yes (equivalency within 5%)  |  |
| Method Qualification                   |                       | No   | No   | Required   |  |
| Calibration Standards                  |                       | Relative quantification  | Predefined linear range  | Customized range for biomarker/study   |  |
| Quality Control Samples                |                       | No   | Independently prepared at 3-4 concentration levels (extracted in triplicate)   | Independently prepared at 3-4<br>concentration levels (extracted in triplica   |  |
| Linearity of Dilution                  |                       | Not Applicable   | Independent dilution QC (2-3X ULOQ)<br>processed and analyzed at highest dilution<br>factor used in batch                                | Independent dilution QC (≥ 3X ULOQ<br>qualified at highest dilution factor utilize<br>during sample processing         |  |
| Internal Standard                      |                       | Cocktail Internal Standard or<br>Structural Dialog   | Structural Analog or Stable Isotope Label  | Structural Analog or Stable Isotope Lab  |  |
| Sample Dilutions                       |                       | Not Applicable   | Based on historical data (if available) and/or dilution into predefined matrices   |  |  |
| Samples Above Quantitation Limit       |                       | Not Applicable   | Fit-for-purpose data extrapolation or post-extraction dilution with reanalysis   | Sample reprocessing with required diluti for quantification  |  |
| Parameters Evaluated                   |                       | Optional estimation of concentrations by<br>single point calibration standard<br>(if reference material available) | Parallelism; Dilution Linearity; Endogenous Concentrations; Spike Recovery;<br>Short-Term Matrix Stability; Freeze/Thaw Matrix Stability |  |  |
| Data Deliverable                       |                       | Standard Excel Tabular Summary<br>Additional options: Bioanalytical Word Report                                    |  |  |  |

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