

CoU implementation at Roche

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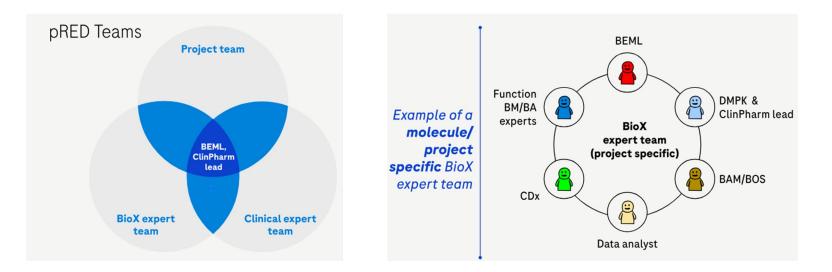
Roche Innovation Center Basel

16.11.23 | EBF meeting

One BioX (Biosamples, Biomarkers and Bioanalytics)



- BioX expert team operates at a Project level including all relevant stakeholders and integrating feedback from Biomarker, DMPK, Clinical Science and Clinical Pharmacology strategies
- BioX Expert Teams develop and discuss biomarker strategies; while integrating other topics like PK/PD modeling, immunogenicity, PoM, patient enrichment and disease understanding.



Biomarker Request Form to understand the CoU



Biomarker Request Form (BRF)

Document Location	Project Folder
Electronic version *	<add folder="" link="" relevant="" to=""></add>
Document guidance	 This document serves as entry point for all biomarker requests The document is to be filled out and signed by the requester (BEML, TBL, etc) in consultation with the supporting PS scientists Completion of this document is a prerequisite to initiate any biomarker activities Any changes to the initial biomarker request should be captured in this document, agreed upon, versioned and appended

Name	Department	Signature and date
BioAM Scientist(s) (responsible for the activity)		
BEML / TBL / BEMS / <u>ClinPharm</u> (activity requestor)		
Biosample Operations Specialist		
BAM (if applicable, delete line if not)		

Document History	Version	Reason for Change	Date
	Version 1.0	Initial Request	



FBE (Functional Biomarker Expert) BEML (Biomarker Experimental Medicine Leader)

BAM (Bioanalytical Manager) BOS (Biosample Operation Specialist)

- Setup per study/Project
- Consolidated information on all Biomarker assessments
- Entry point for all biomarker requests
- Evolving document that is versioned
- All relevant functions are involved

Biomarker Request Questionnaire



Scientific Information and Rationale

- Scientific Background biological questions, disease indication
- Rationale to measure the biomarkers
- Are different proteoforms described for any of the biomarkers? If yes, which proteoform is of interest? Do PTMs, splice variants or disease specific SNPs play a role?
- Are there any known binding partners? Is the aim to measure bound or unbound protein complexes?
- What is the expected range of biomarker concentrations in the relevant sample type and study populations?
- What is the expected level of change? Are the levels expected to increase or decrease? Please specify respectively for total vs free (target engagement) analyte levels
- Are change in levels expected due to therapeutic intervention or general biology (eg, disease progression, protein shedding from cell surface)

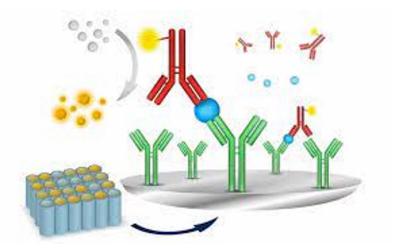


Biomarker Request Questionnaire



Technical Aspects

- Sample type and matrix/tube of choice (eg, anticoagulants).
 Consider artificial release during sample processing (eg, platelet activation, temperature instability, aggregation, tricky stability such as complement proteins)
- Technologies in consideration
- Is there an existing prototype assay or internal/external evidence from specific technology?
- Any known or expected drug interference?
- Are there any pre-meds or co-meds that could interfere with the measurements?
- What is the minimal change the assay needs to demonstrate?
- Is a custom assay generation needed? Are reagents available?



Biomarker Request Questionnaire

Operational and Protocol related

- Is there access to relevant diseased samples for development/validation work?
- Does the ICF wording allow samples to be used for method development (eg, in-study clinical verification/validation extension)?
- Estimated total sample volume available
- How will the data be utilized? (For eg, fold changes in levels, relative/absolute concentration reporting)
- What does this entail in terms of decision making? (For eg, internal decision making, dose selection, patient safety monitoring, subject inclusion/exclusion, label claim, hypothesis testing, efficacy readouts, others)
- Is there potential for the data to support a regulatory claim or for submission to HA?
- How are the biomarkers placed in terms of clinical endpoints?
- What is the expected analysis frequency (eg, batch analysis, cohort driven analysis, end of the study analysis, etc)







Capturing the CoU



Table-1

Definition and agreement on the Context of Use

To be filled by BEML and BioAM scientist. This table captures a summary of biomarkers needed and their intended Coll in the study. For cellular biomarkers, please only list critical populations of interest. Define Coll per biomarker in a clear and concise way

(for eg. The purpose of the assay is to measure [free/unbound] biomarker [A] in [matrix X] on [platform X] to evaluate levels in [population Y] to enable a change of [Y%] from baseline to allow decision [Z])

Biomarker	Туре	Intended Context of Use
	Cellular	
	soluble	
	soluble	
	Cellular	
	Cellular	

Technical CoU statement:

- Intended CoU captured in the Biomarker Request Form (tabular format) - what is needed
- Signed and agreed with stakeholders (BioA Scientist, BPL, BEML, FBE)
- Final technical CoU statement (what the method can deliver) to be captured in the validation report

The purpose of the assay is to measure [free/unbound] biomarker [A] in [matrix_X] on [platform_X] to evaluate levels in [population_Y] to enable a change of [Y%] from baseline to allow decision [Z]

Doing now what patients need next