



**NTP**  
National Toxicology Program

# Lung Breakout Group





## Breakout Group Participants

- **Steven Kleeberger**, NIEHS (**Chair**)
- **Mike Jokinen**, Pathology Associates (**Rappateur**)
- **James Bonner**, Chemical Industry Institute of Toxicology
- **Rajendra Chhabra**, NIEHS
- **Daniel Costa**, U.S. Environmental Protection Agency
- **Angela King-Herbert**, NIEHS
- **George Leikauf**, University of Cincinnati
- **Robert Maronpot**, NIEHS
- **Joe Mauderly**, Lovelace Respiratory Research Institute
- **Daniel Morgan**, NIEHS
- **Brooke Mossman**, University of Vermont
- **Edward Postlethwait**, University of Alabama at Birmingham
- **Sally Tinkle**, NIEHS



## Recommendation #1: Bronchoalveolar Lavage Analysis

- Is it applicable to rodents and humans? **Yes.** If not currently applied to rodents could it be? **Yes.**
- What is the disease process(es) evaluated (e.g., tissue injury, altered function, altered metabolism)? **Especially useful for markers of injury, inflammation, and other tissue response indicators.** Does it identify early or late events? **Largely useful for early events, but may be a good indicator of selected late events as well.**
- Is it sensitive, specific, and/or predictive of the disease process? **It could be all of these, depending on the panel of indicators chosen for investigation.** Could it be used to demonstrate a NOAEL? **Probably not.**
- What type of specimen/measurement is needed? **Whole lung or partial lung lavages are doable.** Is obtaining it noninvasive and easily accessible? **It is largely invasive in small animals, but easily accessible.** What is the appropriate time for sample or measurement collection? **Likely most informative at early time points.**
- Are there other special considerations with including it in NTP studies (e.g., extra animals needed)? **Depending on the number of other biomarkers assessed, extra animals may be necessary.**
- What technology is required? **Trivial.** Is it accurate, reproducible, and cost effective? **It can be. We recommend strongly that a SOP be developed to maximize these outcomes.**



## **Bronchoalveolar Lavage Fluid Analysis – other considerations**

- Especially useful for cell counts and differentials
- Propose that panels of molecular markers (chemokines, cytokines, antioxidant, albumin, etc) should be considered instead of measuring a specific few of these.
  - marker panels could be designed to evaluate processes such as innate immunity, acquired immunity, and inflammation.
  - cost-effective methods have been developed that make this approach reasonable



## Recommendation #2: Enhanced Histopathology

- Is it applicable to rodents and humans? **Yes**. If not currently applied to rodents could it be?
- What is the disease process(es) evaluated (e.g., tissue injury, altered function, altered metabolism)? **Tissue injury, inflammation, apoptosis, repair, etc.** Does it identify early or late events? **Early and late.**
- Is it sensitive, specific, and/or predictive of the disease process? **Yes for all.** Could it be used to demonstrate a NOAEL? **Perhaps.**
- What type of specimen/measurement is needed? **Appropriately fixed lung or lung lobe(s).** Is obtaining it noninvasive and easily accessible? **Invasive and easily accessible.** What is the appropriate time for sample or measurement collection? **Investigator defined – likely appropriate at all times.**
- Are there other special considerations with including it in NTP studies (e.g., extra animals needed)? **Depending on other biomarkers assessed in the lung, extra animals may be needed.**
- What technology is required? **Good pathologists...** Is it accurate, reproducible, and cost effective?



## **Enhanced Histopathology – other considerations**

- Investigation/development of fixation techniques to allow for morphometric analysis (shrinkage of tissue)
- Immunohistochemistry for proteins, particularly those that may be found in BAL analysis. Emphasis on cross-platform confirmation
- Perform trichrome and PAS stains
- Stain for Ki67 protein for cell proliferation



## Recommendation #3: Gene Expression Analysis

- Is it applicable to rodents and humans? **Yes**. If not currently applied to rodents could it be?
- What is the disease process(es) evaluated (e.g., tissue injury, altered function, inflammation, altered metabolism)? **It may be useful for multiple disease processes. Informatics have developed or will be developed to better understand process.** Does it identify early or late events? **Early and late.**
- Is it sensitive, specific, and/or predictive of the disease process? **It may be very sensitive, specific, and predictive.** Could it be used to demonstrate a NOAEL? **Uncertain.**
- What type of specimen/measurement is needed? **High quality RNA.** Is obtaining it noninvasive and easily accessible? **It could be noninvasive, and is easily accessible.** What is the appropriate time for sample or measurement collection? **Investigation/question-defined.**
- Are there other special considerations with including it in NTP studies (e.g., extra animals needed)? **May require additional animals.**
- What technology is required? **Multiple microarray platforms.** Is it accurate, reproducible, and cost effective? **Answers to these questions are forthcoming.**



## **Gene Expression Analysis – other considerations**

- Analyses will identify some genes that are part of disease and some part of the process
- Probably not for routine use – however, save tissue for future evaluation if warranted
- Cost for large microarray chips coming down so can make evaluation of very large numbers of genes feasible in near future
- Need to have phenotypic changes (anchoring) to correlate with genotypic changes





## **Additional biomarkers for consideration (up to 5)**

- #1
  - What is the disease process(es) evaluated?
  - Why is it recommended with a lower priority?
- #2
  - What is the disease process(es) evaluated?
  - Why is it recommended with a lower priority?
- #3
  - What is the disease process(es) evaluated?
  - Why is it recommended with a lower priority?



## Biomarkers that offer promise, but need more work

- Imaging

- What is the disease process evaluated? **Many.** A particular advantage is the value-added that imaging may represent. Multiple organ systems may be evaluated simultaneously.
- Could it provide information not otherwise obtainable? **Perhaps, but real strength is that it may be non-invasive, and repeated measures could be possible in the same animal.**
- Would it be applicable to rodents and humans? **Yes**
- What type of development is needed to make it useful?
  - What is the roadblock(s) to it being developed? **None – progress is ongoing to refine techniques**
  - If resources were available, how long would it take? **Projected 1 yr before ready for prime-time.**



## Other recommendations or comments

- Investigate routinely freezing tissues for future evaluation
- Investigate use of imaging techniques
  - Need to define information desired from imaging
- Suggest that the NTP conducts internal validation work with a known chemical to determine if these biomarkers are useful.



## Conclusions

- Plethora of available biomarkers
- Recommend utilizing bronchoalveolar lavage and enhanced histopathology to develop suites or clusters of biomarkers that may be used and applied to developing systems analyses
- Gene expression may be a very useful biomarker, with caveats
- We recommend that plans should be made to appropriately archive multiple tissues for biomarker application using methods not currently ready for implementation, but will be available shortly



## Other questions, time permitting

- Gene Chip Technology: to investigate patterns of gene expression
  - How could this tool be used? Should NTP put resources into its development? Why or why not?
    - Is there evidence that this methodology is appropriate for use in preclinical rodent studies?
    - Is there a specific list of genes that should be targeted?
    - How frequently should the list of targeted genes be reviewed?
    - What is the potential for identifying new gene targets?
- Proteomic Profiling Technology: to investigate patterns of protein expression
  - How could this tool be used? Should NTP put resources into its development? Why or why not?
    - Is there evidence that this methodology is appropriate for use in preclinical rodent studies?
    - What is the potential for identifying new protein targets?