Continuing Conversations are informal discussions designed for meeting attendees to discuss key issues relevant to the lymphoma community. The discussions during these sessions are meant to be interactive so attendees should come prepared to actively participate. One of the Continuing Conversations at the 2016 ASH Meeting on Lymphoma Biology will be on comprehensive genomic analysis of lymphoid malignancies.

The purpose of this discussion is to identify gaps in existing/ongoing efforts in genomic profiling of lymphoid malignancies and to gain better understanding of information contained in existing datasets. Additionally, the discussion will touch upon effective data-sharing strategies and opportunities (e.g., NCI’s Genomic Data Commons).

Discussion Questions:

1. What datasets already exist for lymphoid malignancies?
   - What kind of data are there?
   - Where are the datasets?
   - Are they easily accessible/open?
   - How are these data shared? Is there a need for one comprehensive database?

2. Is there a need for additional comprehensive sequencing efforts in lymphoma?
   - What diseases/subtypes?
   - What would it take to organize these efforts?
   - How can the lymphoma community help to move these efforts forward?
     - Scientific community
     - Funding organizations
     - Professional societies
     - Patient advocacy groups