**RESOURCE SHARING PLAN**

Data generated in these studies will be freely available to members of the research community with a goal to enable dissemination of data, while also protecting the privacy of the participants and the utility of the data, by de-identifying and masking potentially sensitive data elements in compliance with the NIH public data sharing policy ([http://grants.nih.gov/grants/policy/data\_sharing](http://grants.nih.gov/grants/policy/data_sharing" \t "_blank)). All other resources developed in the course of the proposed studies will be available by request to qualified academic investigators for non-commercial research.

For all studies, we will follow the National Institute of Health's Genomic Data Sharing Policy. The raw datasets corresponding to expression, genomic, and genetic data generated by these studies will be submitted to Gene Expression Omnibus (GEO, <http://www.ncbi.nlm.nih>. gov/geo/) or the Sequence Read Archive (SRA, www.ncbi.nlm.nih.gov/sra) for use by other investigators. As datasets are analyzed, then validated, we will proceed with deposition in ImmPort according to a timeline negotiated with the Program Officer. Sample data in the YCAAD biorepository are, and will continue to be, available on the internet through the YCCI research accelerator, a publicly accessible platform for scientific collaboration (ycci.researchaccelerator.org).

Yale University School of Medicine and all investigators will adhere to the NIH Grants Policy on Sharing of Unique Research Resources including the “Sharing of Biomedical Research Resources: Principles and Guidelines for Recipients of NIH Grants and Contracts” issued in December 1999. Specifically, material transfers would be made with no more restrictive terms than in the Simple Letter Agreement or the UBMTA and without reach through requirements. Should any intellectual property arise which requires a patent, we would ensure that the technology remains widely available to the research community in accordance with the NIH Principles and Guidelines document. In addition, we will provide relevant protocols and published data upon request. Accepted versions of final, peer-reviewed manuscripts emanating from this research will be deposited on-line to PubMed Central in accord with NIH Public Access.