Data Sharing Wizard

Is your study:

- A clinical trial as defined by NIH (https://grants.nih.gov/policy/clinical-trials/definition.htm)?
- Or are you planning to publish your results in a journal that is a member of the ICMJE?
- Or are you planning an FDA-regulated study of a drug or device that falls under the FDAAA's Final Rule regarding trial registration?

If you answered "Yes" to any of these, you are required to: 1. register your study at www.ClinicalTrials.gov, through the University account; 2. periodically update the information; and 3. for some studies, publicly report results.

You can obtain assistance regarding trial registration and updating from the Clinical Research Specialist in the Clinical Research Support Office (x8409).

1. Will you generate data using microarrays or RNAseq?
If yes, you may need to deposit your data in Gene Expression Omnibus (GEO).
Yupu Liang and Thomas Carroll can assist you in submitting your data
2. Will you generate raw sequences from next generation high throughput sequencing?
If yes, you may need to deposit your data in Sequence Rea Archive (SRA).
Yupu Liang can help you submit your data.
3. Will you generate raw nucleotide sequences using gel/capillary technology?
If yes, you may need to deposit your data in Trace Archive.
Yupu Liang can help you submit your data.
4. Will you generate GWAS genotype-phenotype data?
If yes, you may need to deposit your data in the Database of Genotypes and Phenotypes (dbGAP).
Yupu Liang can assist you in depositing your data.
5. Will you generate array CGH data on structural genomic variations?
If yes, you may need to deposit your data in the Database of Genomic Structural Variation (dbVar).
Yupu Liang can assist you in depositing your data.
6. Will you generate data on short genomic variations using SNP array?
If yes, you may need to deposit your data in the Database of Short Genomic Variations (dbSNP).
Yupu Liang can assist you in submitting your data.
7. Will you generate cancer genomic data?
If yes, you may need to deposit your data in the Cancer Genomic Hub (cGHub).
Yupu Liang can assist you in submitting your data.
8. Will you generate 3-dimensional structural coordinates based data from NMR or x-ray crystallography?
If yes, you may need to deposit your data in the Protein Data Bank PDB).
Yupu Liang can assist you in submitting your data.
9. Will you generate 3-dimensional structural coordinates based data from NMR?
If yes, you may need to deposit your data in the Electron Microscopy Data Bank (EMDB).
Yupu Liang can assist you in submitting your data.