GMes expt

What steve would like to be able to show with this is:

* Equations exist to estimate rubisco and cyt b6f amounts from Vcmax and Jmax
* Are they close?
* How important is mesophyll conductance to these equations (when this is low, does it screw up equations)
  + Estimated rubisco is based on assumption that Gmes is not limiting
  + Where Gmes is limiting, we expect estimations to be wrong, i.e. different from quant proteomics data
  + We want nice relationships between proteomics rubisco and Vcmax / Jmax
  + We can use the difference between Cc (chloroplastic CO2) and Ci to correct A/Ci curves (since licor assumes Cc and Ci are equal), and then see if we get better relationships between rubisco and adjusted Vcmax/Jmax

Data processing:

TDL didn’t use daylight savings time but licor (probably) did. DST started Sunday Oct 6th, 2013, so all dates after that will need to be corrected.

To do:

* + correct all timestamps in the TDL files and resave them
  + then use the licor log numbers for each sampleID to get the TDL time range (-30sec from first log number, +30 from last log number) and remove kruft
  + stitch everything together
  + everything has to be done by date