

Data analysis flow:

- 1) Convert raw data files to 'compressed' files
 - a. 8 byte channel -> 1 byte
 - b. 8 byte timetag -> 8 byte timetag
 - c. 8 byte transfer number -> 2 byte integer
- 2) Create hdf5 file with sync pulse information
 - a. Use GPS signals to identify when to start correlating Alice and Bob data
 - b. Start on first sync after a common pulse-per-second GPS signal from Alice and Bob has been identified
 - c. We will use data before the common GPS signal to estimate rates to determine the stopping point in the data analysis to compute p-value
 - d. Identify bad trials due to problems with the data taking such as
 - i. Laser losing modelock
 - ii. Refrigerator warming up
 - iii. Keep trials where there is a massive time jump in timetags momentarily. This seems to be a problem with the operating system (Windows) and the timetagger driver.
 - iv. Keep the trial with the first bad sync because we didn't know the trial would be bad
- 3) Create hdf5 file with setting information
 - a. Occasionally both the "0" and "1" setting fires. Map this to "0" in the analysis
- 4) Create hdf5 file with information about clicks in the detectors, pockel cell slot number, coincidences, etc. Merge with other hdf5 files
- 5) Convert hdf5 file to text formats for further processing into hypothesis testing.

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