202203_nrg_comparison

March 12, 2022

[1]: from typing	g import List, Union, Optional
import lmf	it as lm
from dat_ar	nalysis import get_dat, get_dats
from dat_ar	nalysis.analysis_tools import nrg
from dat_ar	nalysis.analysis_tools.general_fitting import calculate_fit, FitInfo
from dat_ar	nalysis.useful_functions import mean_data, get_data_index
from dat_ar	nalysis.plotting.mpl.util import make_axes, ax_setup
from dat_ar	nalysis.plotting.mpl.plots import display_2d, waterfall_plot
import matp	plotlib.pyplot as plt
import matp	plotlib as mpl
from datacl	lasses import dataclass
import nump	by as np
Import nump	yy as up

[2]: %%latex \tableofcontents

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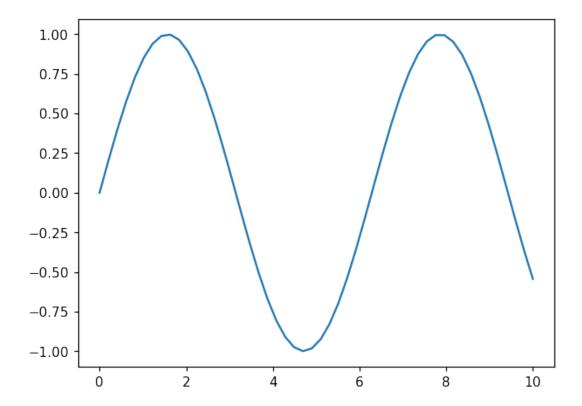
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$1\quad \text{Quick setup to make figures look OK}$

Adjust dpi and figsize to make figure below look like a full size figure with reasonable text size

```
[3]: %matplotlib inline
mpl.rcParams.update({
        'figure.figsize': (6.4,4.8),
        'figure.dpi': 110, # 27in 1440p = 110
})
plt.plot(np.linspace(0,10), np.sin(np.linspace(0,10)))
```

[3]: [<matplotlib.lines.Line2D at 0x1bbd8a8f640>]



2 Loading Data

Experimental data presented in 'Entropy measurement of a strongly correlated quantum dot' are from measurements Feb/March 2021. Measured on what we refer to as Owen's DD (the first two chamber electron heating device that had 2 additional QDs near the entropy sensing QD).

I'll first get the information we are going to use later from each of the HDF files now, and put that into a few containers to keep track of it.

I'm using my functions to load the HDF file only so that I don't have to worry about annoying differences between how data is recorded in different cooldowns (i.e. my library standardizes the interface with measurement data)

```
[4]: @dataclass
     class Data:
         datnum: int
         # Recorded data
         original_x: np.ndarray
         y: np.ndarray # This won't change with processing
         original_i_sense2d: np.ndarray
         # Other recorded attributes
         x label: str
         y label: str
         coupling_gate_val: float # What potential on the coupling gate (i.e._
      →useful for identifying weak/strong coupled data)
         num_samples_per_setpoint: int # How many datapoints per heating setpoint
         num_steps: int # Number of full heating cycles (DACs set at the beginning ...
      →of each cycle)
         measurement_frequency_hz: float # Rate of datapoint measurement
         # Variables required for processing
         center_before_averaging: bool # Whether to center the data with_
      → individual transition fits to unheated data before averaging
         fitting_width_mv: float # How much data on either side of zero to use for_
      → fitting (in units of sweep gate)
         setpoint_averaging_delay: float = 0.01 # how many seconds of data to throw_
      →out after each setpoint change
         forced_theta: Optional[float] = None # Forced theta value for fitting_
      → (necessary for gamma broadened)
         forced gamma: Optional[float] = 0.001 # Forced gamma value for fitting
      → (necessary for thermally broadened)
         # Calculated things
         centers: List[float] = None # Center positions of each row of data (only_
      \rightarrow for weakly coupled data)
         cold fit: FitInfo = None # Result from fitting averaged unheated i sense
         hot_fit: FitInfo = None # Result from fitting averaged heated i_sense
         # Calculated data
        x: np.ndarray = None # x array with num_steps shape (i.e. one x val per_
      \rightarrow DAC step)
         i sense separated: np.ndarray = None # I sense data separated into each
      →part of heating square wave (after averaging setpoints)
         i_sense_cold: np.ndarray = None # Average together the unheated parts of ___
      \hookrightarrow data
         i_sense_hot: np.ndarray = None  # Average together the heated parts of data
         entropy_signal: np.ndarray = None # Average heated minus average unheated_
      \rightarrow i_sense (2D entropy signal)
```

```
avg_x: np.ndarray = None # x_array centered around 0 after averaging data_
 \rightarrow (i.e. if centering data anyway, might as well define 0 as center)
    average_i_sense: np.ndarray = None # Averaged i_sense using center values_
\rightarrow to align first
    average_i_sense_std: np.ndarray = None # stdev of i_sense data averaged
    average_i_sense_cold: np.ndarray = None # Average together the unheated_
 \rightarrow parts of data
    average_i_sense_hot: np.ndarray = None # Average together the heated parts_
 \rightarrow of data
    average_entropy_signal: np.ndarray = None # Averaged entropy using center_
 \rightarrow values to align first
    average_entropy_signal_std: np.ndarray = None # stdev of entropy data_
\rightarrow averaged
dats = get_dats([2164, 2121, 2167, 2133], exp2hdf='febmar21tim')
fit_widths = [50, 100, 200, 400]
datas = []
for dat, fit_width in zip(dats, fit_widths):
    data = Data(
        datnum=dat.datnum,
        original_x=dat.Data.x_array,
        y=dat.Data.y_array,
        original_i_sense2d=dat.Data.i_sense,
        x_label = dat.Logs.xlabel,
        y_label = dat.Logs.ylabel,
        coupling_gate_val=dat.Logs.dacs['ESC'],
        num_samples_per_setpoint=int(dat.Logs.awg.wave_len/4),
        num_steps=dat.Logs.awg.num_steps,
        measurement_frequency_hz=dat.Logs.measure_freq,
        center_before_averaging=True if dat.Logs.dacs['ESC'] < -225 else False,
        fitting_width_mv=fit_width,
    )
    datas.append(data)
```

```
[5]: # Temporary cell for quickly checking values
dat = dats[0]
dat.Logs.awg
for data in datas:
    print(data.center_before_averaging, data.coupling_gate_val)
```

```
True -349.43
True -239.87
False -219.73
False -189.51
```

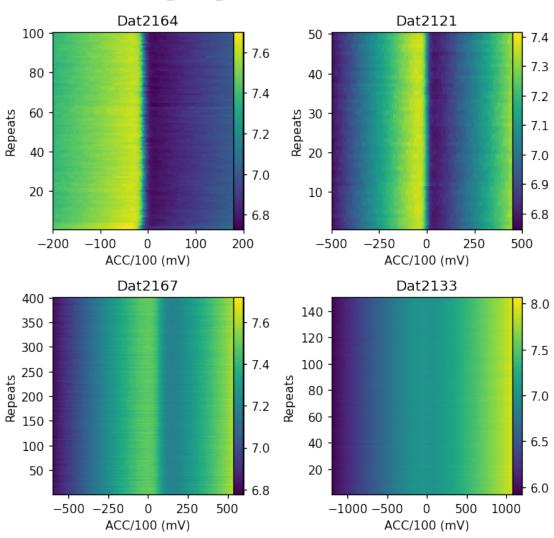
I'll just plot the recorded data in a few different ways first to give a reasonable idea of what the measurements look like

```
fig, axs = make_axes(len(datas))

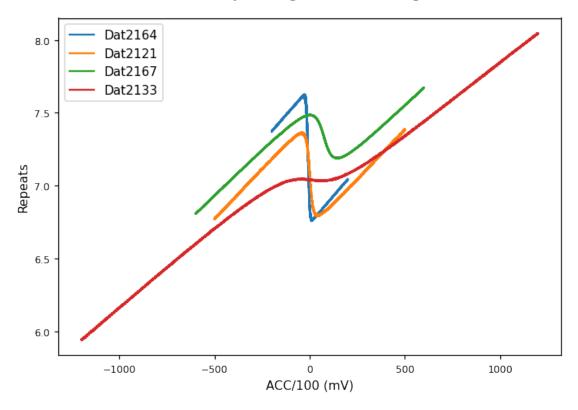
for data, ax in zip(datas, axs):
    ax = display_2d(
        data.original_x, data.y, data.original_i_sense2d,
        ax=ax, colorscale=True,
        x_label=data.x_label, y_label=data.y_label,
        auto_bin=True
        )
    ax.set_title(f'Dat{data.datnum}')

fig.suptitle('Recorded i_sense_2d Binned to 1000 points per row')
fig.tight_layout()
```

Recorded i_sense_2d Binned to 1000 points per row



Data blindly averaged (no centering)



2.1 Standard processing of square wave heated data

Need to: - separate out the recorded data into the different heating sections - average the data from each setpoint (excluding a settling time) - average together the heated/unheated parts and generate entropy signal - fit the unheated parts (to obtain center values. Only necessary for not very gamma broadened data) - average 2D to 1D data using centers (where calculated, otherwise just blindly average)

2.1.1 Separate into heating sections AND average data from each setpoint

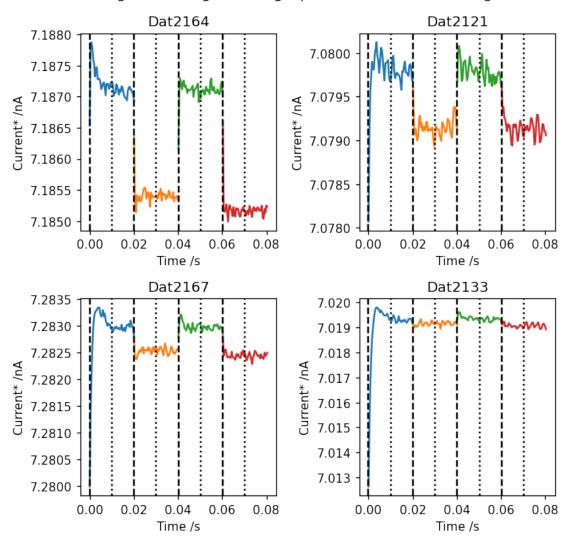
```
[8]: fig1, axs1 = make_axes(len(datas))
     fig1.suptitle(f'Data averaged to a single heating square wave (to see settling_
     →time)')
     for ax in axs1:
         ax.set_xlabel('Time /s')
         ax.set_ylabel('Current* /nA')
     fig2, axs2 = make_axes(len(datas))
     fig2.suptitle(f'Single row of separated and setpoint averaged data')
     for ax in axs2:
         ax.set_ylabel('Current /nA')
     for data, ax1, ax2 in zip(datas, axs1, axs2):
         ax1.set_title(f'Dat{data.datnum}')
         ax2.set title(f'Dat{data.datnum}')
         ax2.set_xlabel(data.x_label)
         # Separate by setpoint
         delay_index = round(data.setpoint_averaging_delay*data.
      →measurement_frequency_hz)
         data_by_setpoint = data.original_i_sense2d.reshape((data.original_i_sense2d.
      →shape[0], -1, 4, data.num_samples_per_setpoint))
                                                                          # (repeats, ...
      \rightarrowsteps, 4 sections, samples)
         # Plot the square waves averaged on top of each other
         setpoint_duration = data.num_samples_per_setpoint/data.
      →measurement frequency hz
         for i in range(4):
             startx = 0 + i*setpoint_duration
             endx = startx + setpoint_duration
             averaged_to_single_setpoint = np.mean(data_by_setpoint[:,:,i,:],_
      \rightarrowaxis=(0,1))
             ax1.plot(np.linspace(startx, endx, data.num_samples_per_setpoint),_
      →averaged_to_single_setpoint, label=i)
             ax1.axvline(startx+data.setpoint_averaging_delay, linestyle=':',__
      ax1.axvline(startx, linestyle='--', color='k')
         # Average parts of setpoint after delay
         data.i_sense_separated = np.mean(data_by_setpoint[:, :, :, delay_index:],_
      \rightarrowaxis=-1)
         data.x = np.linspace(data.original_x[0], data.original_x[-1], data.
      →num_steps)
```

```
for i in range(4):
    ax2.plot(data.x, data.i_sense_separated[0, :, i], label=i) # Plotting
    row 0 of each data
    ax2.legend()

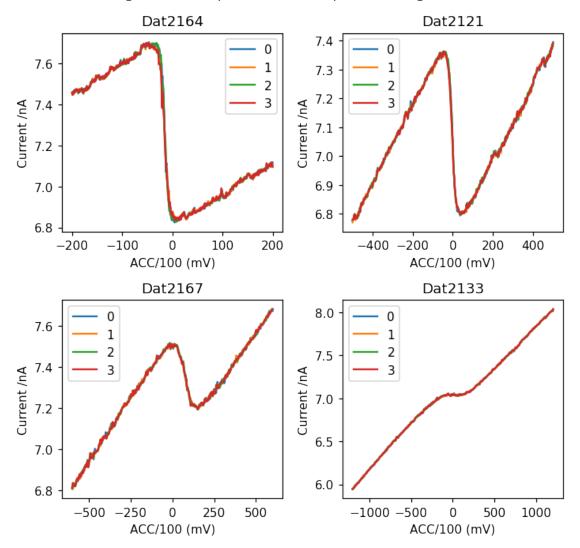
# Calculate E

for fig in fig1, fig2:
    fig.tight_layout()
```

Data averaged to a single heating square wave (to see settling time)



Single row of separated and setpoint averaged data



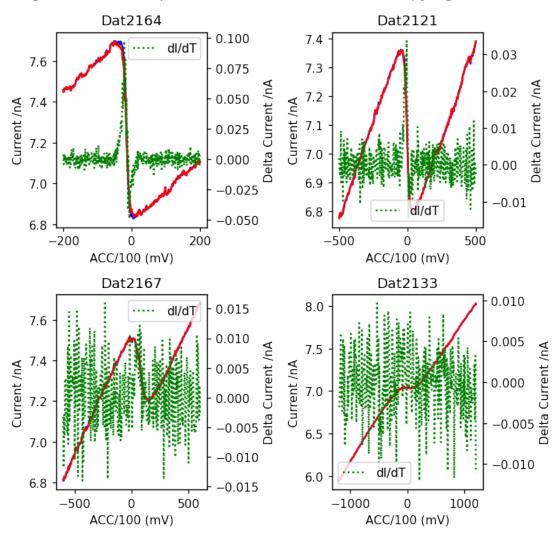
The first set of figures here show the averaged heating square waves (i.e. every heating square wave from the entired dataset averaged on top of each other), where the dashed line indicates heating changes, and the dotted line indicates the point at which we will start averaging the data (i.e. throwing out data before the dotted lines). This is to determine how much data has to be ignored before averaging the setpoints (mostly determined by the first section which occurs after DAC steps)

The second set of graphs shows a single sweep of the data separated into the 4 parts of the square wave. In the weakly coupled regime you can see the difference between heated and unheated

2.1.2 Average together heated/unheated parts and generate entropy signal

```
[9]: fig, axs = make_axes(len(datas))
     fig.suptitle(f'Single row of data separated into hot/cold and the entropy_
     ⇔signal from them')
     for ax in axs:
         ax.set_ylabel('Current /nA')
     for data, ax in zip(datas, axs):
         ax.set_title(f'Dat{data.datnum}')
         ax.set_xlabel(data.x_label)
         data.i_sense_cold = np.mean(data.i_sense_separated[:,:,(0,2)], axis=-1)
         data.i_sense_hot = np.mean(data.i_sense_separated[:,:,(1,3)], axis=-1)
         data.entropy_signal = data.i_sense_cold - data.i_sense_hot
         ax.plot(data.x, data.i_sense_cold[0], 'b-')
         ax.plot(data.x, data.i_sense_hot[0], 'r-')
         ax2 = ax.twinx()
         ax2.plot(data.x, data.entropy_signal[0], 'g:', label='dI/dT')
         ax2.set_ylabel('Delta Current /nA')
         ax2.legend()
     fig.tight_layout()
```





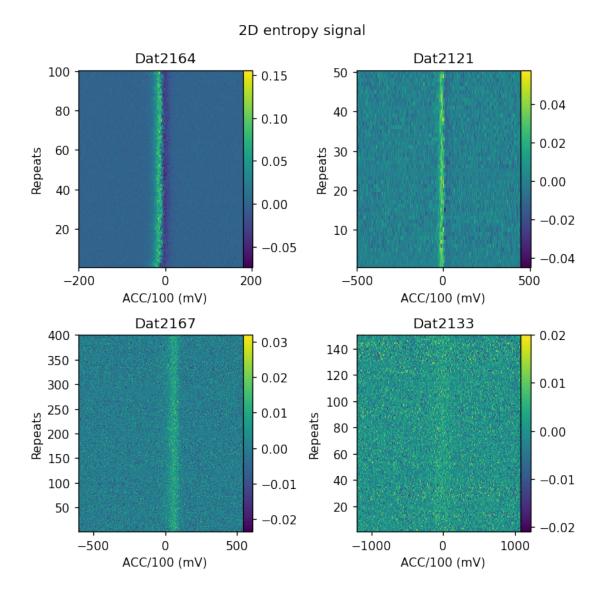
Since this is only a single row of data, for the high Gamma measurements, the signal to noise is extremely weak. After averaging we'll see more.

I'll quickly plot the 2D entropy signal which does show something

```
fig, axs = make_axes(len(datas))
fig.suptitle(f'2D entropy signal')

for data, ax in zip(datas, axs):
    display_2d(data.x, data.y, data.entropy_signal, ax, x_label=data.x_label,
    y_label=data.y_label, colorscale=True)
    ax.set_title(f'Dat{data.datnum}')

fig.tight_layout()
```



Can see that there is a dI/dT signal even at higher Gamma

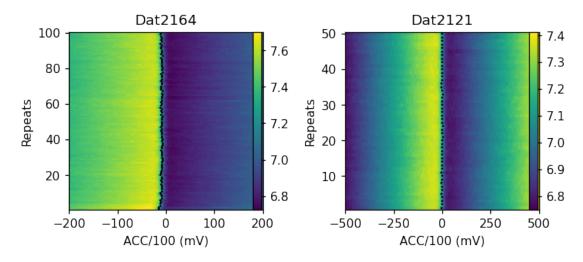
2.1.3 Fit unheated parts to obtain centers for averaging

```
[11]: def weak_transition_func(x, mid, theta, amp, lin, const):
    """Charge transition shape for weak coupling only"""
    arg = (x - mid) / (2 * theta)
    return -amp / 2 * np.tanh(arg) + lin * (x - mid) + const

fig, axs = make_axes(sum([data.center_before_averaging for data in datas]))
    fig.suptitle('I_sense data with center values plotted')
    i = 0
    for data in datas:
```

```
if data.center_before_averaging:
        ax = axs[i]
        i+=1
        ax.set_title(f'Dat{data.datnum}')
        d = data.i_sense_cold
        x = data.x
        params = lm.Parameters()
        params.add_many(
            # param, value, vary, min, max
            lm.Parameter('mid', np.mean(x), True, np.min(x), np.max(x)),
            lm.Parameter('amp', np.max(d)-np.min(d), True, 0, 2),
            lm.Parameter('const', np.mean(d), True),
            lm.Parameter('lin', 0, True, 0, 0.01),
            lm.Parameter('theta', 10, True, 0, 100),
        fits = [calculate_fit(x, d1d, params, weak_transition_func,_
                                     # basically wraps model.fit from lmfit
 →method='leastsq') for d1d in d]
        data.centers = [fit.best_values.mid for fit in fits]
        display_2d(x, data.y, d, ax, colorscale=True, x_label=data.x_label,_
 →y label=data.y label)
        ax.scatter(data.centers, data.y, s=2, c='k', marker='+')
    else:
        data.centers = np.zeros(data.y.shape[0])
fig.tight_layout()
```

I_sense data with center values plotted



Graphs above are to check that the fitting of each individual row of unheated data has done a good job of finding a reasonable center point that will be used for averaging

2.1.4 Average 2D to 1D using centers calculated

```
[12]: for data in datas:
          # Average Transition Data
          averaged datas = []
          for i in range(4):
              averaged_datas.append(
                  mean_data(data.x, data.i_sense_separated[:,:,i],
                             centers=data.centers, return_x=True, return_std=True,__
       →nan_policy='omit')
              # Note: mean_data returns [avg_data, centered_x_array, stdev of data_
       \rightarrow averaged]
          data.average_i_sense = np.array([d[0] for d in averaged_datas])
          assert np.all(np.all([averaged_datas[0][1] == averaged_datas[i][1] for i inu
       \rightarrowrange(3)])) # Ensure the x_arrays returned are all identical to each other
          data.avg_x = averaged_datas[0][1]
          data.average_i_sense = np.array([d[0] for d in averaged_datas])
          data.average_i_sense_std = np.array([d[2] for d in averaged_datas])
          data.average_i_sense_cold = np.mean(data.average_i_sense[(0,2), :], axis=0)
          data.average_i_sense_hot = np.mean(data.average_i_sense[(1,3), :], axis=0)
          # Average Entropy Data
          data.average_entropy_signal, x_, data.average_entropy_signal_std =__
       →mean_data(data.x, data.entropy_signal,
                             centers=data.centers, return_x=True, return_std=True,_
       →nan_policy='omit')
          # Note: mean data returns [avg_data, centered x_array, stdev of data_
       \rightarrow averaged]
          assert np.all(data.avg x == x_) # Ensure same x axis for entropy data
```

I'm using my own mean_data function here, effectively what it is doing is centering the data first using the center values provided, then averaging along axis 0. To do the centering requires some interpolation to end up with the same x_axis for each row of data before averaging, so that's why I'm not showing it all in the cell here. mean_data returns three things: - the averaged data - the new centered x_array for that data - the standard deviation of all the values averaged together for each datapoint. (Note that it is the standard deviation, NOT the standard error. To convert to standard error it should be divided by \sqrt{N} where N is the number of repeats (shape of data.y)

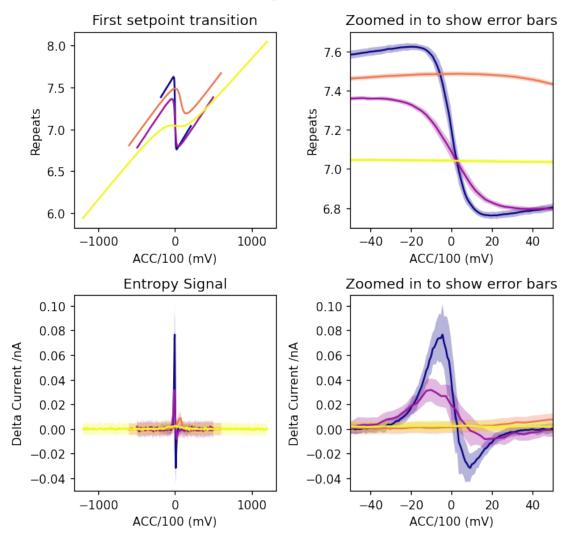
```
[13]: fig, axs = make_axes(len(datas))
fig.suptitle(f'Averaged Data')
transition_axs = axs[0:2]
entropy_axs = axs[2:]

colors = plt.get_cmap('plasma')(np.linspace(0, 1, len(datas)))

for ax in transition_axs: # Plot exactly the same thing twice
```

```
for data, c in zip(datas, colors):
        d, std = data.average_i_sense[0], data.average_i_sense_std[0] # /np.
\rightarrow sqrt(data.y.shape[0])
        ax.plot(data.avg_x, d, c=c, label=f'Dat{data.datnum}')
        ax.fill_between(data.avg_x, d-std, d+std, facecolor=c, alpha=0.3)
    ax.set ylabel(datas[0].y label)
# Zoom in on one of the plots
transition_axs[0].set_title(f'First setpoint transition')
transition_axs[1].set_title(f'Zoomed in to show error bars')
transition_axs[1].set_xlim(-50, 50)
transition_axs[1].set_ylim(6.7, 7.7)
for ax in entropy_axs: # Plot exactly the same thing twice
    for data, c in zip(datas, colors):
        d, std = data.average_entropy_signal, data.average_entropy_signal_std #_
\rightarrow/np.sqrt(data.y.shape[0])
        ax.plot(data.avg_x, d, c=c, label=f'Dat{data.datnum}')
        ax.fill_between(data.avg_x, d-std, d+std, facecolor=c, alpha=0.3)
    ax.set ylabel('Delta Current /nA')
# Zoom in on one of the plots
entropy_axs[0].set_title(f'Entropy Signal')
entropy_axs[1].set_title(f'Zoomed in to show error bars')
entropy_axs[1].set_xlim(-50, 50)
# entropy_axs[1].set_ylim(6.7, 7.7)
for ax in axs:
    ax.set xlabel(datas[0].x label)
fig.tight_layout()
```

Averaged Data



This is the data averaged after centering (for the more weakly coupled data), where the standard deviation, σ , of all data averaged together is plotted. Note that this is **NOT** the standard error (σ/\sqrt{N}) . Although it should make more sense to plot standard error, in practice the error bars end up invisibly small, and realistically, there is a significant uncertainty in these measurements. At the same time, the pure standard deviation is too large of an error bar. So this is really just for guidance as to how the data looks, and the **error bars should not be taken seriously**.

3 Fitting to NRG

For details on how I work with NRG data, see the 202203_nrg_fitting_explained.pdf.

When fitting data to NRG, I can only vary one of theta or gamma at a time as they are too closely correlated otherwise, and I end up getting very unreliable results. Because of that, it is necessary to first figure out what theta should be for the gamma broadened measurements.

As a first approximation, we can just use the theta we find for the weakly coupled data where gamma = 0. That is what I will plot initially, as it may be insightful enough.

The assumption that theta remains constant does not take into account the change in lever arm strength of the plunger gate as a function of the coupling gate. I.e. as the coupling gate is made more positive to open up the QD to the reservoir, this also has the effect of making the plunger gate (ACC) less effective at changing dot occupation.

Additionally, the charge sensor does not quite have a linear response, and so for the very broadened measurements, this plays a larger role. Again, we will first ignore this and only include it later if necessary

3.1 Fitting weakly coupled

```
[14]: weak_datas = datas[0:1] # Currently, only first dataset is weakly coupled
     fig, axs = make_axes(len(weak_datas)*2)
     nrg_helper = nrg.NrgUtil()
     for i, data in enumerate(weak_datas):
         # initial guess for params
         params = lm.Parameters()
         params.add many(
             # name, value, vary, min, max
             lm.Parameter('mid', 0, True),
             lm.Parameter('amp', 1, True, 0, 2),
             lm.Parameter('const', 7, True),
             lm.Parameter('lin', 0.0001, True, 0, 0.01),
             lm.Parameter('theta', 10, True, 0, 50),
             lm.Parameter('g', 0.001, False), # NOT varying g while fitting
             lm.Parameter('occ_lin', 0, False), # Not usually necessary to vary this
         )
         x, d_c, d_h = data.avg_x, data.average_i_sense_cold, data.
      →average_i_sense_hot
         indexes = get_data_index(x, [-data.fitting_width_mv, data.fitting_width_mv])
         s = np.s [indexes[0]:indexes[1]]
         x, d_c, d_h = x[s], d_c[s], d_h[s]
         data.cold_fit = nrg_helper.get_fit(x, d_c, initial_params=params,_
      →which_data='i_sense')
         data.hot_fit = nrg_helper.get_fit(x, d_h, initial_params=params,__
      ⇔which_data='i_sense')
         print(f'Dat{data.datnum} Fit values:\n'
               f'\tUnheated Theta: {data.cold_fit.best_values.theta:.3f}\u00b1{data.
      f'\tHeated Theta: {data.hot_fit.best_values.theta:.3f}\u00b1{data.
      →hot_fit.params["theta"].stderr:.3f} mV\n'
               f'\tDelta Theta: {data.hot_fit.best_values.theta - data.cold_fit.
       ⇒best values.theta:.3f} mV\n'
```

```
for ax, d, fit in zip(axs[i*2:i*2+2], [data.average_i_sense_cold, data.

→average_i_sense_hot], [data.cold_fit, data.hot_fit]):

ax.scatter(data.avg_x, d, label='Data', s=30, marker='o', u

→facecolors='none', edgecolors='b')

ax.plot(data.avg_x, fit.eval_fit(x=data.avg_x), label='Fit', c='r')

ax.set_xlabel(data.x_label)

ax.set_ylabel('Current /nA')

ax.set_xlim(-50, 50)

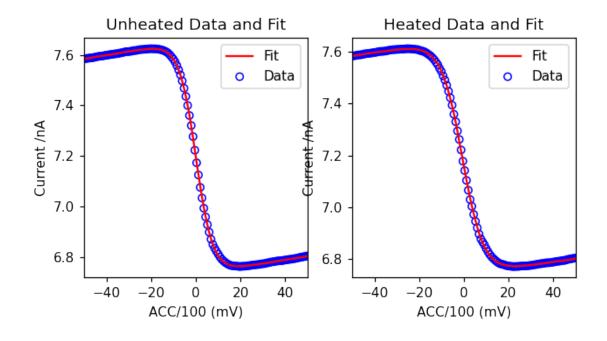
ax.legend()

axs[2*i].set_title('Unheated Data and Fit')

axs[2*i+1].set_title('Heated Data and Fit')
```

Dat2164 Fit values:

Unheated Theta: 3.925±0.006 mV Heated Theta: 4.910±0.007 mV Delta Theta: 0.985 mV



Calculated fits to both heated and unheated data.

```
[15]: cold_theta = datas[0].cold_fit.best_values.theta
hot_theta = datas[0].hot_fit.best_values.theta
delta_theta = hot_theta-cold_theta
```

3.2 Fitting Strongly Coupled

For now we will assume the thetas from weak coupling can be used in strong coupling without any adjustment (an OK approximation)

```
[16]: strong_datas = datas[1:]
      fig, axs = plt.subplots(len(strong datas), 2, figsize=(7, 3.
      →5*len(strong datas)))
      axs = axs.flatten()
      nrg_helper = nrg.NrgUtil()
      for i, data in enumerate(strong_datas):
          # Only use data near transition
          x, d_c, d_h = data.avg_x, data.average_i_sense_cold, data.
       →average_i_sense_hot
          indexes = get_data_index(x, [-data.fitting_width_mv, data.fitting_width_mv])
          s = np.s_[indexes[0]:indexes[1]]
          x, d_c, d_h = x[s], d_c[s], d_h[s]
          # initial quess for params
          params = lm.Parameters()
          params.add_many(
              # name, value, vary, min, max
              lm.Parameter('mid', np.mean(x), True, np.min(x), np.max(x)),
              lm.Parameter('amp', np.nanmax(d)-np.nanmin(d), True, 0.1, 1),
              lm.Parameter('const', np.nanmean(d_c), True),
              lm.Parameter('lin', 0.001, True, 0, 0.005),
              lm.Parameter('theta', cold_theta, False), # Don't vary theta for
       →Strong coupling fitting
              lm.Parameter('g', 40, True, cold_theta/1000, cold_theta*50),
              lm.Parameter('occ_lin', 0, False), # Not usually necessary to vary this
          )
          # Fit Cold
          data.cold_fit = nrg_helper.get_fit(x, d_c, initial_params=params,__
       ⇔which_data='i_sense')
          # Force hot theta then fit hot
          params['theta'].value = hot_theta
          data.hot_fit = nrg_helper.get_fit(x, d_h, initial_params=params,__
       →which_data='i_sense')
          cold_g, hot_g = data.cold_fit.best_values.g, data.hot_fit.best_values.g
          print(f'Dat{data.datnum} Fit values:\n'
```

```
f'\tUnheated Gamma: {cold_g:.3f}\u00b1{data.cold_fit.params["g"].
 \rightarrowstderr:.3f} mV\n'
          f'\tHeated Gamma: {hot_g:.3f}\u00b1{data.hot_fit.params["g"].stderr:.
 \rightarrow3f} mV\n'
           f'\tGamma Error: {abs(hot_g-cold_g)/(np.mean([hot_g, cold_g]))*100:.
 →3f} % error\n'
           )
    for ax, d, fit in zip(axs[i*2:i*2+2], [data.average_i_sense_cold, data.
 →average_i_sense_hot], [data.cold_fit, data.hot_fit]):
         ax.scatter(data.avg_x, d, label='Data', s=30, marker='o',_

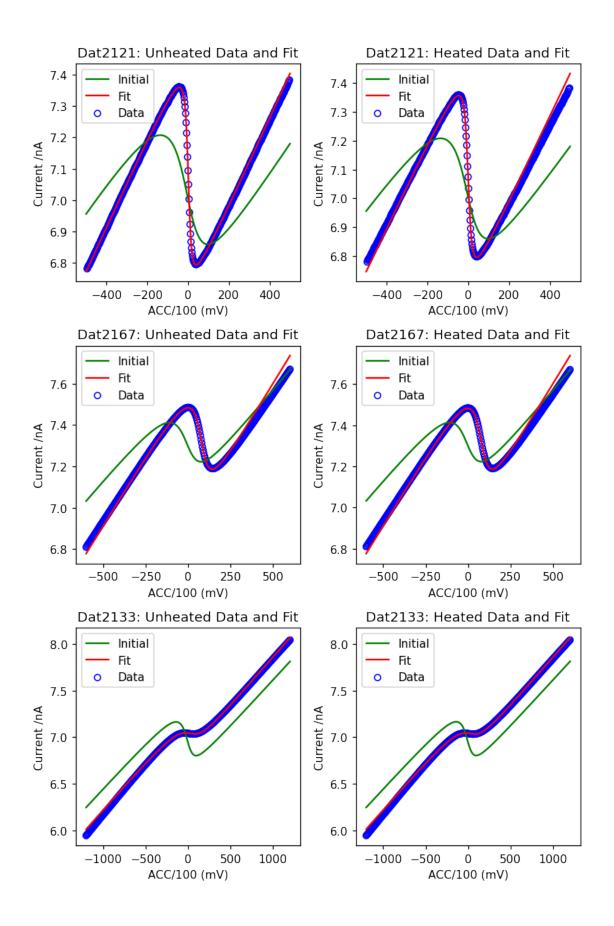
→facecolors='none', edgecolors='b')
         ax.plot(data.avg_x, fit.eval_init(x=data.avg_x), label='Initial', c='g')
         ax.plot(data.avg_x, fit.eval_fit(x=data.avg_x), label='Fit', c='r')
        ax.set_xlabel(data.x_label)
        ax.set_ylabel('Current /nA')
        ax.legend()
    axs[2*i].set title(f'Dat{data.datnum}: Unheated Data and Fit')
    axs[2*i+1].set_title(f'Dat{data.datnum}: Heated Data and Fit')
fig.tight_layout()
Dat2121 Fit values:
        Unheated Gamma: 6.173±0.091 mV
        Heated Gamma: 6.971±0.097 mV
        Gamma Error: 12.142 % error
```

Dat2167 Fit values:

Unheated Gamma: 24.448±0.195 mV Heated Gamma: 25.124±0.179 mV Gamma Error: 2.729 % error

Dat2133 Fit values:

Unheated Gamma: 77.132±1.263 mV Heated Gamma: 77.920±0.916 mV Gamma Error: 1.016 % error



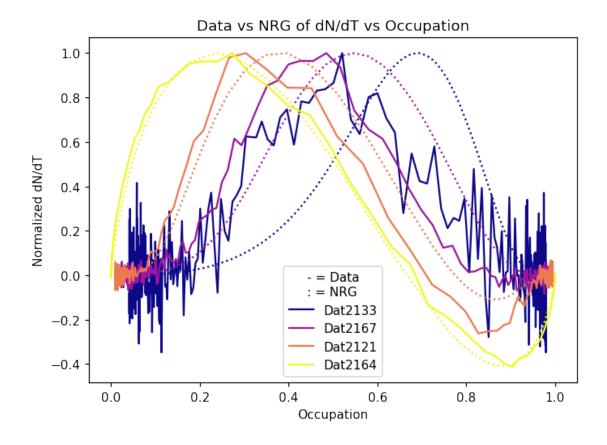
Getting NRG data to fit is a bit more finicky, so it's helpful to see what the initial guesses are. They need to be quite close in order for fitting to work.

Overall, we see that we have quite good fits (i.e. gamma calculated from unheated/heated are pretty similar, line seems to follow well near the middle of the transition - where we are fitting).

We can now use these fit parameters to compare the dNdT to NRG

4 Comparison of Data to NRG

```
[17]: fig, ax = plt.subplots(1)
      colors = plt.get_cmap('plasma')(np.linspace(0, 1, len(datas)))
      nrg_helper = nrg.NrgUtil()
      for data, c in (zip(reversed(datas), colors)):
          cold_values, hot_values = data.cold_fit.best_values, data.hot_fit.
       →best_values
          nrg_helper.init_params(
              mid = np.mean([hot_values.mid, cold_values.mid]),
              amp = np.mean([hot_values.amp, cold_values.amp]),
              const = np.mean([hot_values.const, cold_values.const]),
              lin = np.mean([hot_values.lin, cold_values.lin]),
              theta = np.mean([hot_values.theta, cold_values.theta]),
              g = np.mean([hot_values.g, cold_values.g]),
              occ_lin = np.mean([hot_values.occ_lin, cold_values.occ_lin]),
          nrg_dndt = nrg_helper.data_from_params(x=data.avg_x, which_data='dndt',__
       →which_x='occupation')
          # Note: This returns both data and x in a Data1D object
          # Normalize both NRG and Data entropy signal to have max of 1
          data_dndt = data.average_entropy_signal/np.nanmax(data.
       →average_entropy_signal)
          nrg_dndt.data = nrg_dndt.data/np.nanmax(nrg_dndt.data)
          ax.plot(nrg_dndt.x, data_dndt, c=c, linestyle='-', label=f'Dat{data.
       →datnum}')
          ax.plot(nrg_dndt.x, nrg_dndt.data, c=c, linestyle=':')
      ax.legend(title='- = Data\n: = NRG')
      ax.set_xlabel('Occupation')
      ax.set_ylabel('Normalized dN/dT')
      ax.set_title(f'Data vs NRG of dN/dT vs Occupation')
      fig tight_layout()
```



Using the average fitting parameters of the heated/unheated data to generate the NRG data. Plotting both against occupation.

We see good agreement in the weakly coupled data, but not for any of the more strongly coupled data.