Weekly Report June 21-June 27

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Monte-Carlo Simulation

Some improvements:

- 1. Preprocess function: added rotate_pads=True for the simulated data only
 - a. The real & simulated data are now on the same coordinates
- 2. Commented out the "process shifted z by 38mm due to trigger delay" command in the preprocess function
 - a. Rather nice fitting for event_305
 - b. However, some events who do go through trigger delays now experience a shift in z-direction

Gradient Descent - a local optimization method

Gradient Descent Principles

- 1. We choose a starting point within a function, and then take steps towards the direction with negative (approximate) gradient
- 2. Conjugate gradient: used to approximate the answer x to the equation Qx = b, where Q is a symmetrical matrix -> transformed to this form from our objective equation
 - a. Let $\{d0,...,dn-1\}$ be a set of nonzero Q-orthogonal vectors, the sequence $x_{k+1} = x_k + a_k * d_k$ (where a is step size) converges to a unique solution $Q\mathbf{x} = \mathbf{b}$ that minimizes the function.

Chi-square function

- 1. The function we used to evaluate the chi-square value is minimizer.run_tracks
 - a. Takes in our initial guess (in the form of a 2D array), the experimental data points (x y z) and the experimental hit pattern (pad amplitudes indexed by pad number).
 - Returns an array consisting of the three chi-square values (position, energy, vertex components)

Scipy function construction

1. We define a function f(y) that can be accepted by Scipy.optimize functions

Scipy function construction

- We define the jacobian with a numerical derivative function scipy.optimize.approx_fprime so we can implement it into our optimization function to get a more accurate result
- 3. We then create a callback function (callbackF) which prints out the intermediate result of each iteration when we call the optimization function
- 4. Run the optimization function
 - results = scipy.optimize.minimize(f #object function, ctr0 #initial guess, method="CG", jac=jacobian, callback=callbackF, options={'eps':1e-13} #step size)

Differential evolution - a global optimization method

Differential Evolution Principles

- 1. The method iteratively try to improve an existing candidate solution -> stochastic method, the best solution is not guaranteed to be found
- 2. Starts with a "population" of candidate solutions points with random positions in the search space.
- 3. Pick two distinct candidates (a, b) from the population
- 4. Compute the candidate's potentially new position (aka trial candidate):
 - a. We want to "mutate" the best candidate c0
 - b. C' = c0 + mutation*(a-b)
 - c. For 'best1bin' strategy, we take a random number in [0,1) and if it is less than the recombination constant (explained later) the parameters is loaded from c' (else it's loaded from c0).
 - d. If the trial candidate c' is better than the original candidate c0, it replaces c0.

Scipy function parameters

- 1. Bounds: (min, max) pairs for each element in our input vector; defines search space.
- Maxitr: # of iterations allowed
- 3. Popsize: population size. Default is 15; larger size improves chance of convergence
- 4. Mutation: a float or a range larger mutation value increases the search radius ("step size") but will slow down convergence. If mutation is given a range, the constant is randomly changed per iteration to increase speed.
- 5. Recombination constant: float in [0,1]. Increased RC will allow more mutants to become trial candidates, but will decrease population stability

Results

Event #305

1. Conjugate Gradient chi = 24.713

```
results = scipy.optimize.minimize(f, ctr0, method="CG", jac=jacobian, callback=callbackF, options={'gtol': 30.0, 'eps':le-1
cq = results.x
302 0.000360-0.000141 0.734463 2.614517-3.002859 1.951079 24.713554
303 0.000360-0.000141 0.734463 2.614517-3.002859 1.951079 24.713554
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308 0.000360-0.000141 0.734463 2.614517-3.002859 1.951079 24.713554
309 0.000360-0.000141 0.734462 2.614516-3.002859 1.951079 24.713481
310 0.000360-0.000141 0.734462 2.614516-3.002858 1.951079 24.713390
311 0.000360-0.000141 0.734462 2.614515-3.002858 1.951079 24.713326
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319 0.000360-0.000141 0.734462 2.614515-3.002858 1.951079 24.712220
320 0.000360-0.000141 0.734462 2.614515-3.002858 1.951079 24.713326
```

2. Monte-Carlo

```
uvw, (cu, cv) = mcfitter.preprocess(xyzs[:,0:5], center=(cx, cy), rotate_pads=False) # get calibrated set of data
uvw_values = uvw.values #transform pd file to arrays
print(cu,cv)
mcres, minChis, all_params, good_param_idx = mcfitter.process_event(uvw, cu, cv, return_details=True)
|-43.91625428724363 118.39379347393248
{'x0': 0.004871498439906168, 'y0': -0.011977943314897467, 'z0': 0.7347184090083536, 'enu0': 2.8186679462143145, 'azi0': -2.920989043120735, 'pol0': 1.9263176741477495, 'posChi2': 3.97700712247824, 'enChi2': 7.1981754856024605, 'vertChi2': 3.3
44052462098103, 'lin_scat_ang': 1.2420273973461806, 'lin_beam_int': 720.0566082022592, 'lin_chi2': 36.34021928440234, 'rad curv': 117.8081237563669, 'brho': 0.20911788771969209, 'curv_en': 2.0944258199546852, 'curv_ctr_x': -43.91625428724363,
```

Chi = 3.977+7.198+3.344 = 14.519

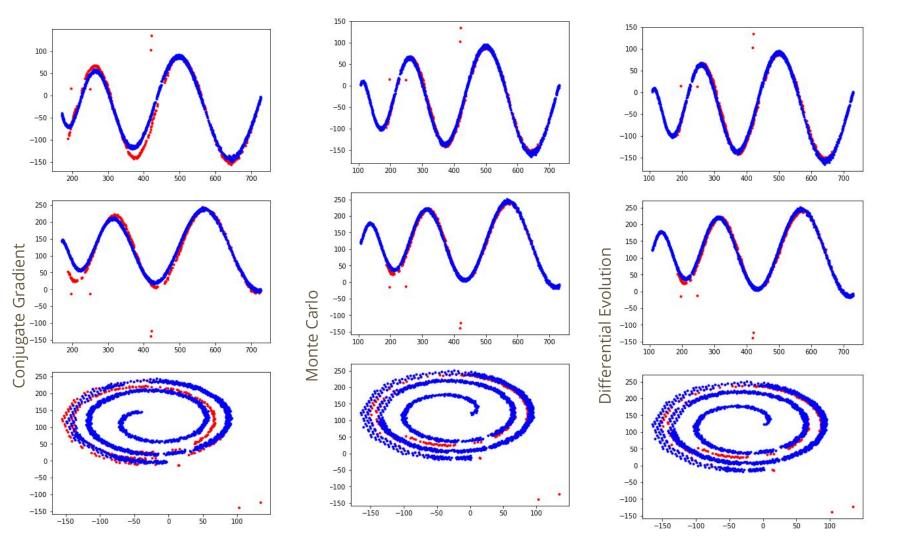
'curv ctr y': 118.39379347393248}

3. Differential evolution

chi=14.208

```
i = 1
def callbackF(x, convergence=10):
    global i
    print('{0:4d}{1: 3.6f}{2: 3.6f}{4: 3.6f}{4: 3.6f}{5: 3.6f}{6: 3.6f}{7: 3.6f}\'.format(i,x[0],x[1],x[2],x[3],x[4],x[5],
    i += 1
bounds = [(-1,1), (-1,1), (0,1), (0,5), (-2 * pi, 2 * pi), (-2 * pi, 2 * pi)]
results = scipy.optimize.differential evolution(f, bounds, callback=callbackF, maxiter=10000, strategy='best1bin',\
                                                 recombination=0.8, popsize=15, mutation=(0.5, 1.0))
       The differential evolution strategy should be one of:
            'bestlbin' - shorter time but usually needs to run multiple times to get the best result
            'bestlexp'- shorter time but usually needs to run multiple times to get the best result
            'randlexp' - takes a long time but converges
            'randtobestlexp' - takes a long time but converges
            'currenttobestlexp' - takes a long time but converges
            'best2exp'
            'rand2exp'
             'randtobest1bin'
            'currenttobest1bin'
             'best2bin'
            'rand2bin'
            'rand1bin'
    The default is 'best1bin'.
  95 0.003659-0.012275 0.733378 2.809433 0.211693-1.926856 14.577127
  96 0.003659-0.012275 0.733378 2.809433 0.211693-1.926856 14.577127
  97 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398
```

98 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 99 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 100 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 101 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 102 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 103 0.002564-0.011759 0.733055 2.809754 0.207638-1.926092 14.260398 104 0.002306-0.011989 0.732534 2.808862 0.204325-1.926166 14.207878 105 0.002306-0.011989 0.732534 2.808862 0.204325-1.926166 14.207878 106 0.002306-0.011989 0.732534 2.808862 0.204325-1.926166 14.207878



Event #456

Monte Carlo: chi = 12.365+7.088+0.033=19.486

Gradient: 110.8 (fail)

Differential evolution: with population size=30, recombination constant = 0.6, chi= 27.32

Monte-Carlo: chi = 94.97 + 6.71 + 0.03 = 101.71

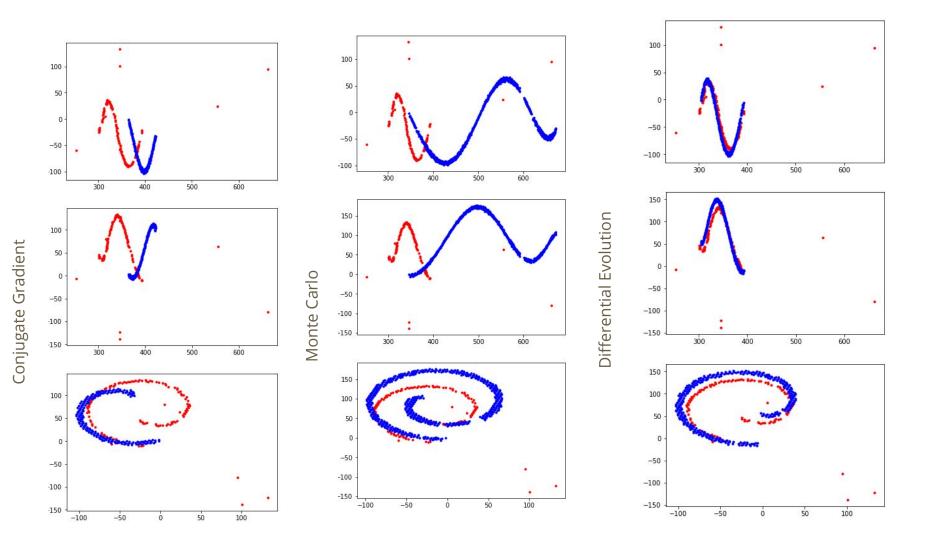
Conjugate Gradient: chi=104.62 (failed)

Differential evolution: chi=37.7

Monte-Carlo: chi = 88.75+11.09+0.21 = 100.05

Conjugate Gradient: chi=106.98 (failed)

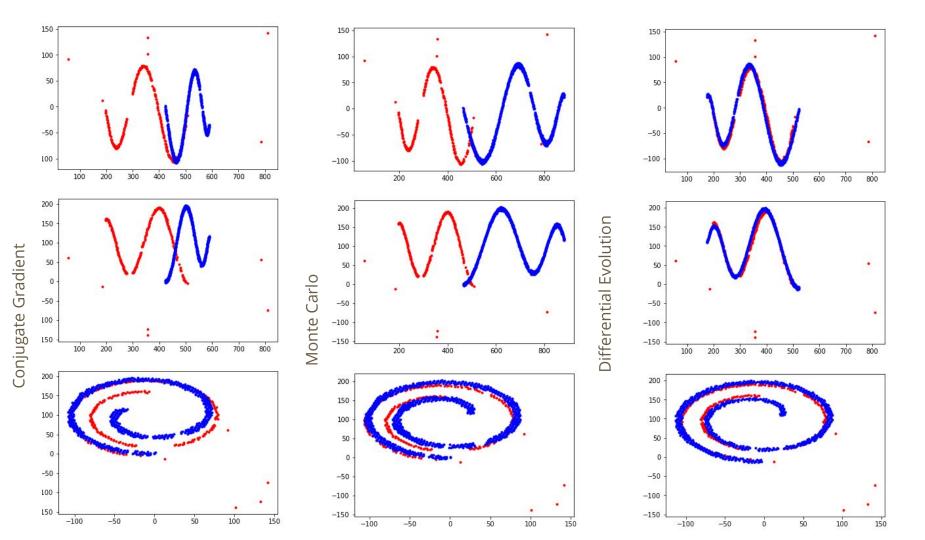
Differential evolution: chi=35.06



Monte-Carlo: chi = 95.92 + 3.626 + 0.04 = 99.59

Conjugate Gradient: chi=102.09

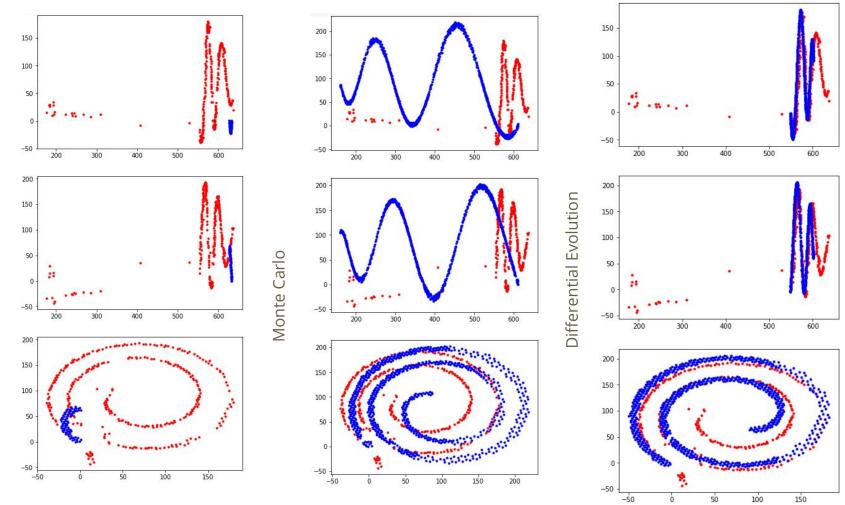
Differential evolution: chi=15.58



Monte-Carlo: chi = 88.79+1.211+0.04 = 90.041

Conjugate Gradient: chi=101.06

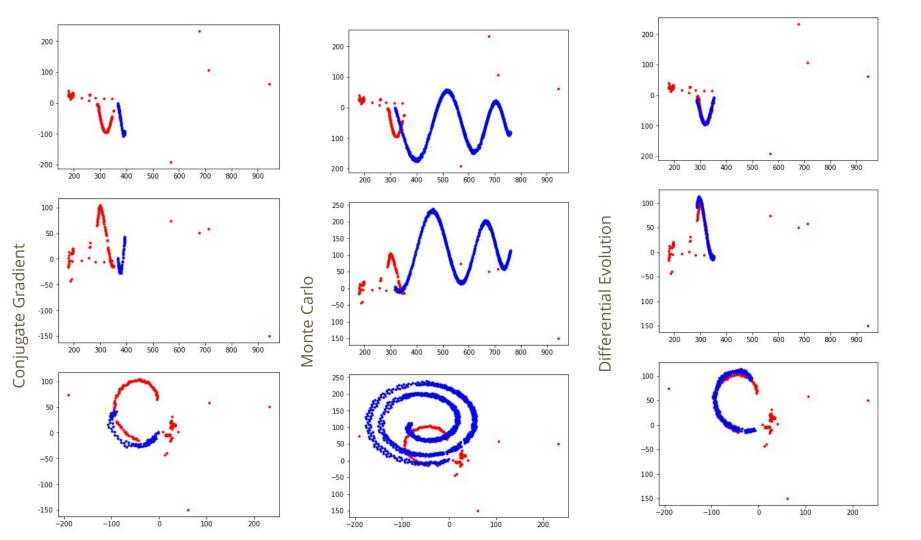
Differential evolution: chi=53.08



Monte-Carlo: chi = 89.9+5.044+0.037 = 94.98

Conjugate Gradient: chi=105.539 (failed)

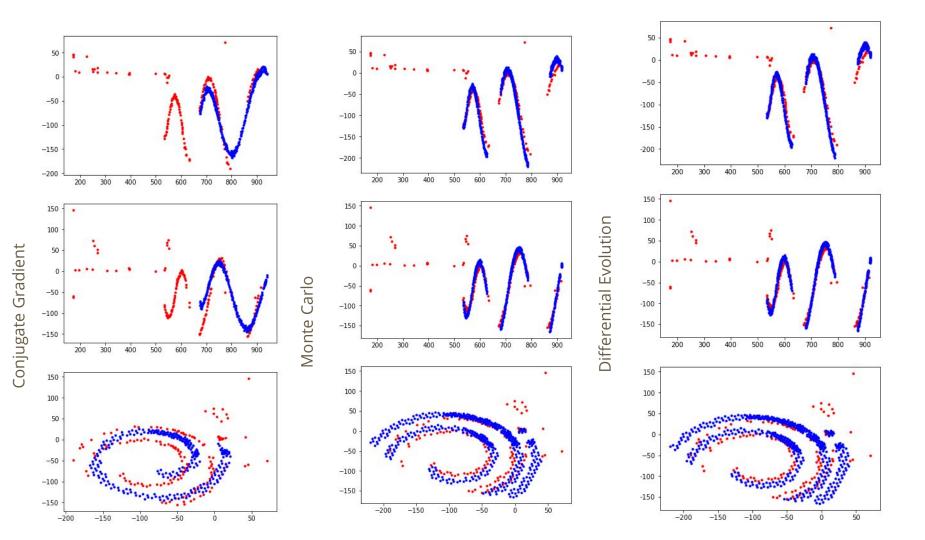
Differential evolution: chi=40.36



Monte-Carlo: chi = 28.9+5.96+1.68 = 36.54

Conjugate Gradient: chi=86.62

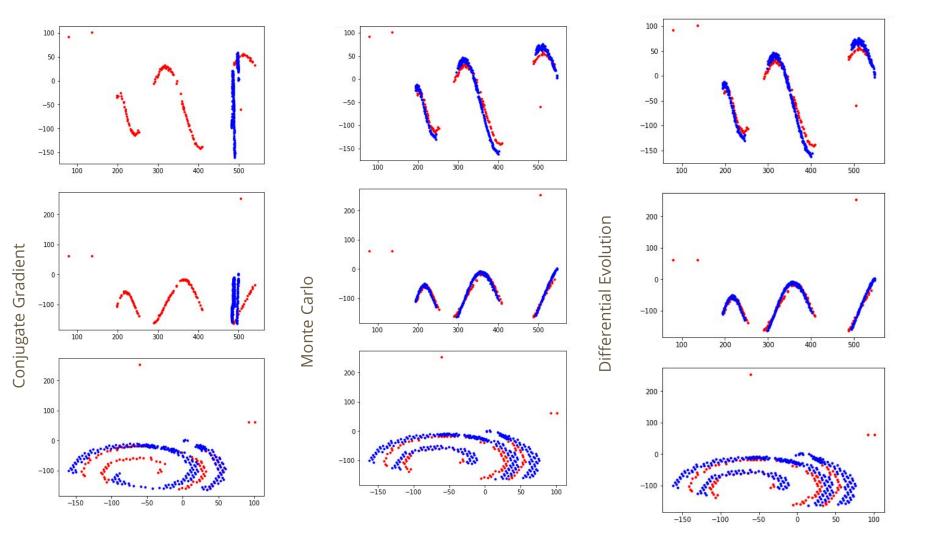
Differential evolution: chi=36.427



Monte-Carlo: chi = 12.22+7.26+0.09 = 19.57

Conjugate Gradient: chi=110.822 (failed)

Differential evolution: chi=18.97



Monte-Carlo: chi = 14.68 + 5.23 + 1.15 = 21.06

Conjugate Gradient: chi=39.18

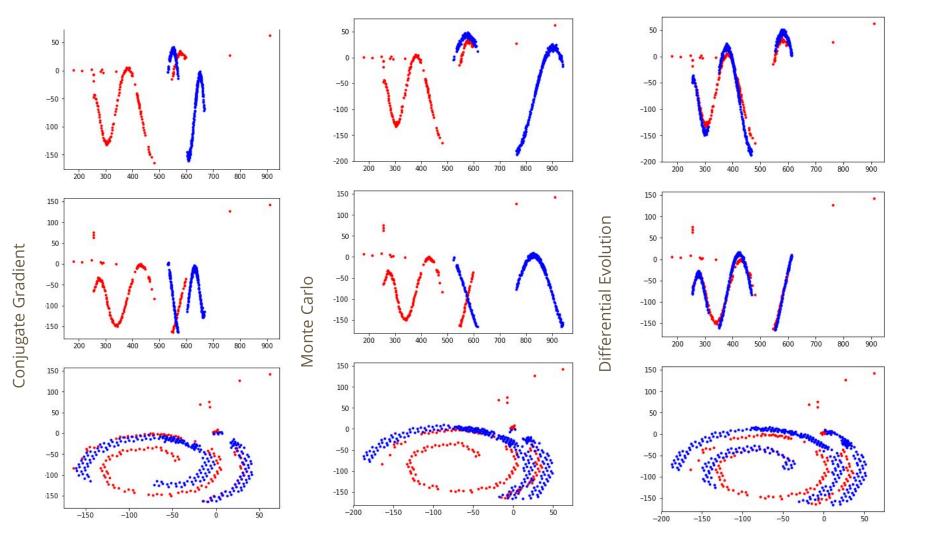
Differential evolution: chi=19.84

150 --50 -50 -100-100 Differential Evolution Conjugate Gradient -50 -50 -50 -75 **Monte Carlo** .100 -100 -100 -150-150 -25 -50 -50 -50 -75 -100-100-150 -100 -50 -150 -100 -50 -100 -50

Monte-Carlo: chi = 94.47 + 5.04 + 0.015 = 99.53

Conjugate Gradient: chi=106.07 (failed)

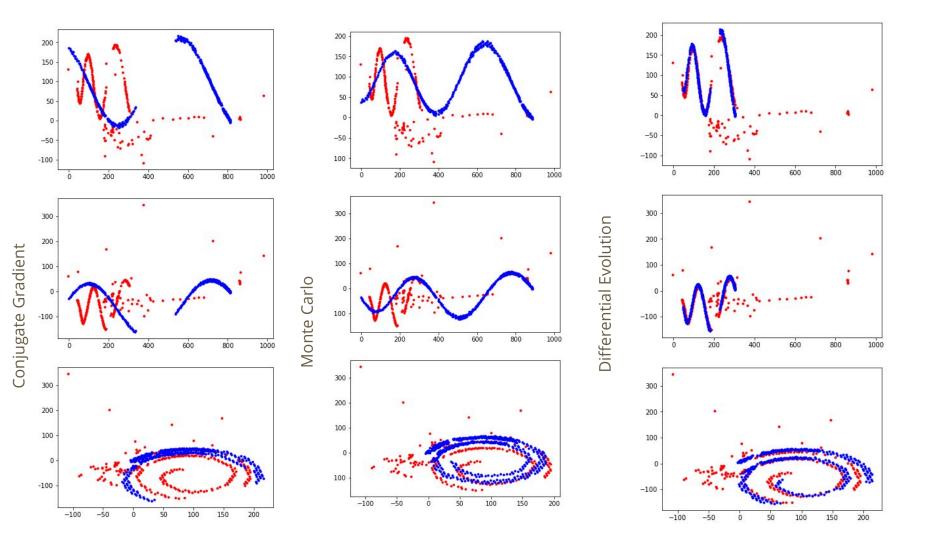
Differential evolution: chi=30.29



Monte-Carlo: chi = 80.5 + 6.45 + 0.05 = 96.45

Conjugate Gradient: chi=92.45

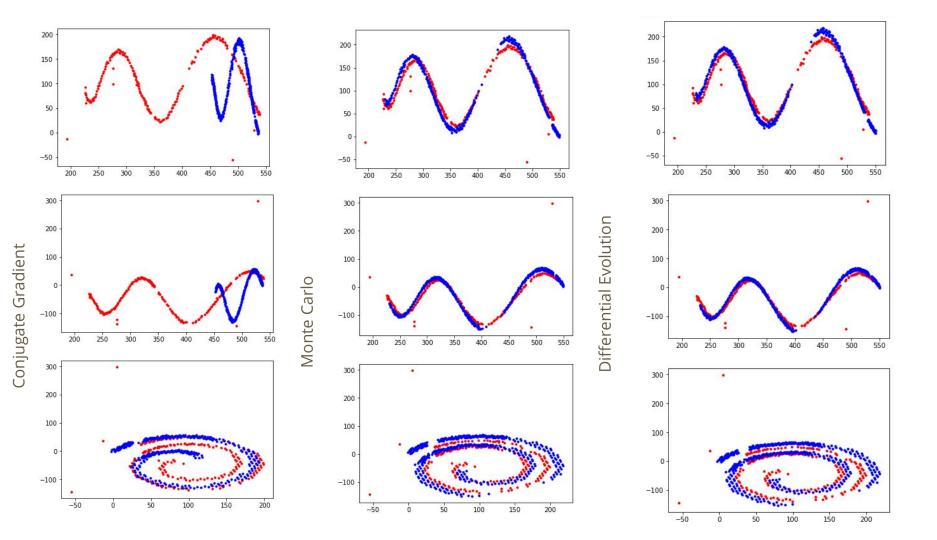
Differential evolution: chi=42.15



Monte-Carlo: chi = 13.67+7.23+0.11 = 30.01

Conjugate Gradient: chi=101.44

Differential evolution: chi=20.76



Other Methods

The Basin-Hopping Algorithm

Similar to the Simulated Annealing method (but more performant according to Scipy)

- 1. Chooses a starting point
- 2. Computes the local minimum (s) using one of the local minimum functions in scipy.optimize.minimize
- 3. Applies a random perturbation (candidate state) to the local minimum (s')
- 4. When $T \sim 0$, if energy at s' is larger than energy at s, the probability of transition from s to s' is approximately zero
- 5. However when T is large, the system is more likely to update to s' in order to have a broader region of search space
- 6. T becomes smaller per iteration so that the system converges

However it gave very unstable & inefficient performance on our data - could be that our function is not smooth enough; we are not sure which T or step size to choose; the method usually fails before giving a significantly lower chi-square than our initial guess