A rough road map for the MOSE project

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Chapter 1

K wall networks

1.1 Intersection algorithm - done

This should be a function called find_intersections(trajectory_1,trajectory_2) returning the intersection(s) of the two given trajectories. It shold be placed in the numerics.py file If the trajectories don't intersect, it should return [] otherwise it should return a list [[point, index_1, index_2], ...] where

- point is the exact point of intersection, given as a complex number
- index_1 is the index of the element of trajectory_1.coordinates that corresponds to the exact intersection point. (Same for index_2). This is important because it will be used to determine the pertinent charge to use in computing progenie by means of the KSWCF. We are taking the shortcut of not adding the actual point to the trajectory, so we introduce a tiny approximation. Keep in mind if trouble arises.
- The exact point of intersection should be added to the lists trajectory_1.coordinates and trajectory_2.coordinates, at the appropriate point warning: coordinates attributes in the Trajectory class are given as a 2-vector, not as a complex numer.
- Moreover, the corresponding value of the period of η for that trajectory should also be added to trajectory_1.periods (and same for trajectory_2), this goes as a complex number.

A very primitive intersection algorithm is already present in numerics.py.

1.2 KSWCF - done (maybe can improve a bit more)

This should be a function called progeny_2(data) placed in the file kswcf.py, that takes data = [γ_1 , $\Omega(\gamma_1)$], $[\gamma_2$, $\Omega(\gamma_2)$]] and returns a list of new charges and corresponding degeneracies, excluding the parents: [. . . [$n\gamma_1 + m\gamma_2$, $\Omega(n\gamma_1 + m\gamma_2)$] . . .]. A primitive example is provided in the file kswcf.py.

1.3 Rotation of the phase - done

A function called phase_scan(theta_range) in structure.py, that repeats the whole computation of primary k-walls, their intersections, generation of descendants, and iterations to get new intersections, new descendats etc.. for different phases.

The argument theta_range is of the form theta_range = [theta_in, theta_fin, steps].

At each phase, it stores the information about intersection points and their genealogy. This information will then be used to construct full MS walls.

As of now, it keeps track of all intersections and all k-walls at every phase, and returns a big array containing them: this has the shape [all_ints, all_trajs] where all_ints = [... [int1, int2, ...] contains arrays of intersections of given phase and similarly all_trajs = [... [traj1, traj2, ...] ...] contains arrays of trajectories of given phase.

1.4 Genealogy - done

Not sure how to structure this yet. But it should be a function called build_genealogy_tree(intersection_point that takes as argument an object of the class IntersectionPoint. It should use the attribute intersection_point.parents to recursively determine the trajectories from whom the point descends. It should eventually return a structure (type of it is to be determined) that contains the genealogy of such point, meaning which branch-points are the farthest ancestors of the intersection, and in which associative order, eg: [[BP1,BP2],BP3]. Not sure if keeping track of branch-point ancestors will be enough, maybe not. Maybe should keep track of trajectories in the middle as well.

The purpose of genealogy is to identify which intersections (occurring at different phases) belong to the same wall of marginal stability.

1.5 Construction of MS walls - done

Once the phase-rotation algorithm is working and the genealogy is ready, we should be able to group together intersection points coming from different phase-snapshots, and build analytic MS walls out of them. This will need some thinking

1.6 Branch cuts

In the function prepare_branch_locus in file structure.py (warning: the whole structure of the code has undergone a major change, need to look for this function), should create an array of branch cuts called bcts that is to be returned (see the function there). Each branch point will generate a branch cut. Cuts should be straight lines going to infinity, at a certain angle THETA_CUTS to be specified in the file config.py. A tentative implementation of this kind of cuts is already provided by the init method of the BranchCut class. It remains to

- tune the parameter branch_cut_cutoff in file parameters.py (length of the branch cut)
- write the method check_cuts in the class Trajectory (Warning: now this class has been replaced by the KWall class and its daughter classes), which is supposed to detect intersection of trajectories with cuts, and determine the Trajectory.splittings (find exact

point, add it to Trajectory.coordinates, then the splitting contains the indices of these exact points. The corresponding kind of addition must be done to Trajectory.periods) and determine the values of Trajectory.local_charge on segments of the trajectory between the various splittings. See the dummy method in the Trajectory class.

• Update the intersection-searching procedure: when two K-walls intersect, the intersection pairing should be evaluated based on the charge of the K-wall at the intersection, this of course can only happen after K-walls have evolved and intersections with cuts have been taken into account.

1.7 Charges of branch points

This should be an algorithm that, at the very beginning —while creating the branch locus—determines automatically which charges to assign to each branch point (hence to each branch cut). It should also specify the DSZ matrix. All this information should replace DSZ_MATRIX and FIXED_CHARGES which are now given by hand in the config.py file.

1.8 Storing Data - done (see instructions)

When running a long computation, involving probing several (such as O(100)) phases, it would be good to store all data on a separate file. In particular, we should

- Create an external file(s) whose name contains date and time
- Store data in such a way that it is retrievable: just need to store the output of the function phase_scan
- Write a module for retrieving (reloading in the RAM) the above data ready for analysis
- Bonus: write a module that (using the external data) will create a separate folder, containing snapshots of the k-walls on the moduli space for all scanned phases
- Extra bonus: write a module that uses the above screenshots to provide a window with a slider tool and lets you browse the k-wall evolution. This is quite useful for debugging and checking the numerics (esp. the undesired bending of k-wall at critical phases)

Instructions To save data, either a single Kwall network, or an entire phase-scan and the related MS walls, use the option -w in the command line.

To read the data, from the python shell, import the function f_recover from the file save_to_file.py, i.e. use:

```
>>> from save_to_file.py import f_recover
```

Then use this function to load the data from a *.mose file:

```
>>> f_recover('some_file_of_type.mose')
```

Now there are two possibilities:

- If the filename begins with single_network then the above command return an instance of the class KWallNetwork.
- If the filename begins with phase_scan then the above command will return a list of the form

[[KWN₁, KWN₂, ..., KWN_m], [MS₁, ...MS_{$$\ell$$}]]

with KWN, MS being instances respectively of the classes KWallNetwork, MarginalStabilityWall

1.9 Refined numerical issues with evolution - done

Trajectories evolve according to the PF equations, these are governed by a matrix which becomes singular at the singular locus of the moduli space (since they contain the discriminant). The odeint algorithm will get into trouble integrating PF in the neighborhood of these points. In particular, the problems seem to consist in trajectories bending abruptly when they run close to singularities.

Possible solutions:

- Once a trajectory was evolved, search backwards to see if it passed near a singularity. If it did, cut it in a neighborhood of the singular point (where it enters a disk of radius ϵ centered there), then invoke a new evolution algorithm (such as that of primay k-walls), then revert back to PF.
- Once a trajectory was evolved, search backwards to see if it passed near a singularity. If it did, erase the trajectory before next iteration, so it cannot create bogus offsprings due to bad numerics (e.g. the bending will make it intersect k-walls it wasn't supposed to meet).

The first possibility seems better, however the evolution method of primary walls will likely run into trouble too, recall the behavior of that method near a MS wall (which certainly would be met by the evolving kwall, when crossing the disc centered around the singular point).

The second possibility might be numerically heavy, but it could be greatly improved by tracking e.g. the determinant of the PF matrix, and using that as a red flag, instead of computing distances from singularities. It would clearly leave an unexplored area nearby a singularity, however this can be fixed by the idea of section 1.10

Note: solved with the second option. Introduced a parameter called TRAJECTORY_SINGULARITY_THRESHOLD that controls the value of the **determinant of the picard-fuchs matrix** along the trajectory's evolution, above which a trajectory is deemed too close to a singularity. To start with, this threshold is set to be 10^6 .

1.10 Enhancing MS walls

1.10.1 Adding endpoints - done

A very simple thing to do: from the genealogy data of an MS wall it should be possible to tell whether the wall starts/ends at certain singularities. Then, make a module that creates new intersection-like objects from data of those singularities, and add these to the MS wall.

1.10.2 Interpolating function

Replace numerical data points with an approximate interpolating function to have a nice smooth MS-wall. A simpler alternative is just to draw a line matching all the dots, although it won't be as nice.

1.11 Fibration-handling module

1.11.1 Storing and calling fibrations

We have a code that handles a class of fibrations. The separate data specifying each fibration should be stored in a tidy way, and called by the core part of the program with a simple function. A possibility is to have a different text file containing the data of each fibration, formatted in a standard way, and to load such data by giving the name of the file (e.g. SU2_Nf=0.xml).

1.11.2 Creating arbitrary fibrations

A nice addition would be a function which allows the user to add a custom-made fibration. For example, one could be prompted to specify the number of UV parameters m, Λ, \ldots , then be prompted for $g_2(u), g_3(u)$. Then the code would store this data in one of the fibration files described above, permanently, with a user-defined name. Then this fibration would be available to the user for future study.

1.12 Visualization module

As we will be analyzing a big array of data, and storing it for later analysis, it is desirable to have a separate suite that handles visualization of this formatted data. This can be tackled once we agree on how stored data will be formatted.

1.13 GUI

Write a graphical user interface, be creative.

Chapter 2

Extracting Refined Topological Invariants

For the distant future: figure out how to handle "finite networks", and automatically recover topological information on the fibration.

Chapter 3

Hyperkahler metrics

For the distant future: figure alorithms for solving TBA equations and getting the explicit metrics.