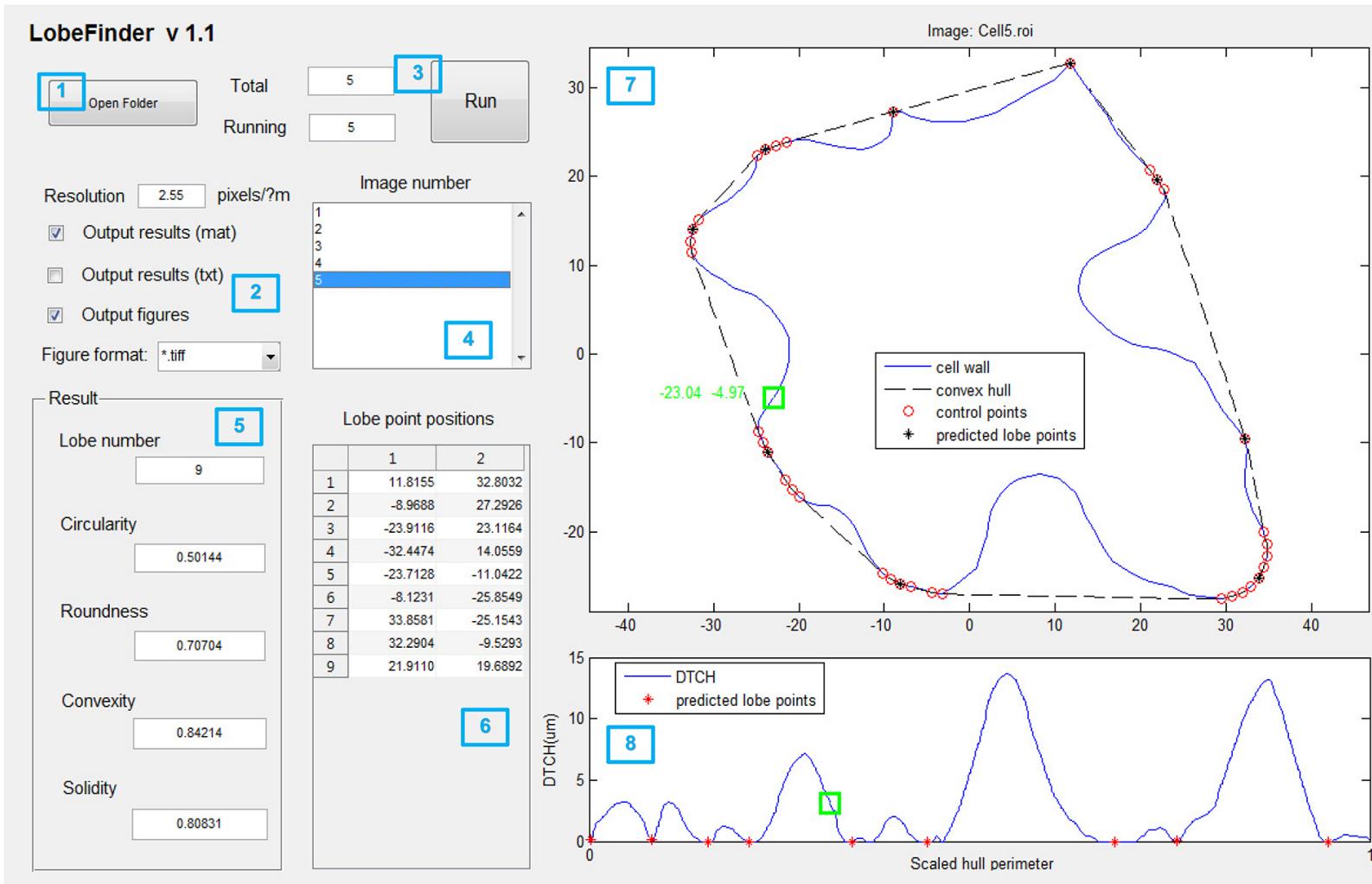


Starting the LobeFinder_GUI

1. Click the “Open Folder” button to select folder where the ROI files are located.
2. Select checkbox(es) for output data type and/or figure format options.
3. Click “Run” button to start LobeFinder.



Results of LobeFinder_GUI

4. Select image ROI number in the box to display the LobeFinder result.
5. Predicted lobe numbers and Cell-shape descriptors.
6. Predicted lobe x-y coordinates.
7. Cell perimeter, convex hull, and predicted lobe point positions.
8. Corresponding distance to the convex hull with predicted lobe point locations on it.

Supplemental Figure S3. Snapshot of the graphical user interface of LobeFinder.

Operation steps and navigation of the results are outlined.