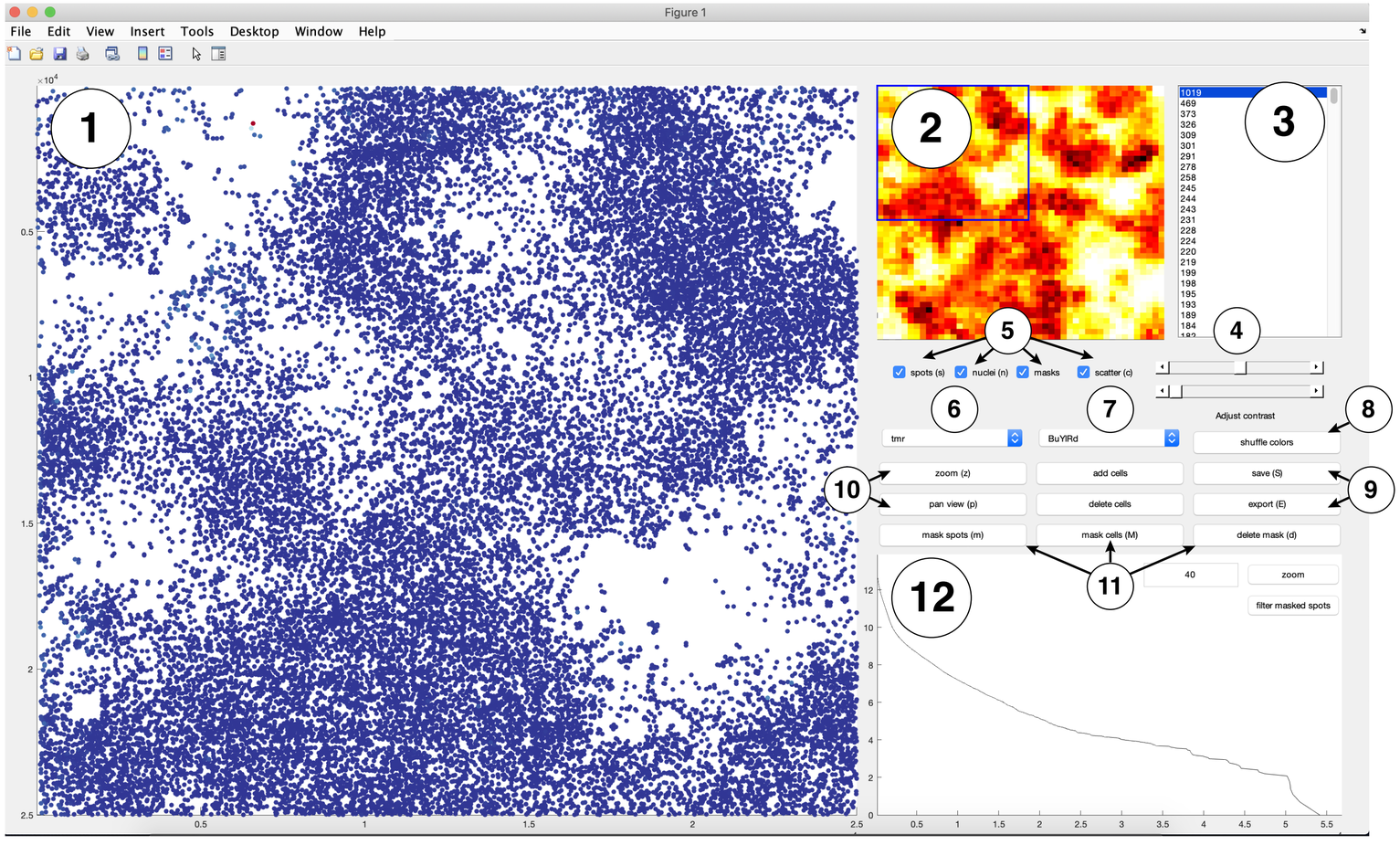

1. Main axes:
   - Click and drag to zoom/pan view
   - Scatter shows all nuclei colored by spot count
   - Use DAPI+FISH overlay to mask spots & create/delete cells
2. Thumbnail Axes:
   - Shows nuclei density across the entire scan
   - Rectangle corresponds to the position displayed in the main axes
   - Drag rectangle or double click to change positions
3. Centroid List:
   - Spot counts for all nuclei (ranked high to low)
   - Double click on value to move to corresponding cell
   - Use up/down arrow keys to move through list
4. Sliders:
   - Adjust contrast of the FISH image in main axes
5. Checkbox:
   - Toggle plot settings for main axes
6. Channel drop-down:
   - Change current FISH channels
7. Colormap drop-down:
   - Change colormap for scatter plot in main axes
8. Shuffle colors:
   - Change spot and centroid colors in main axes
9. Save and export:
   - Save spot, nuclei, and mask tables
   - Export spot summary table (spots per nucleus)
10. Zoom and pan view:
    - While in zoom mode, click and drag on main axes to zoom-in
    - While in pan view mode, click and drag to move through scan
    - Right click to zoom-out by 2X
    - Double click to return to home view
11. Add/delete masks and nuclei:
    - Mask spots in current channel or mask nuclei for all channels
    - Delete erroneous nuclei and add missing nuclei
    - If nucleus is masked/added/deleted, spots in view will be reassigned
12. Threshold axes:
    - Drag vertical blue line to set spot intensity threshold
    - Or use text box in upper right corner