

How to use the diagnostic page

Load slides

Load your slides by logging into the PMA.core. You do this by clicking on the *PMA.core* node in the tree view on the left-hand side (Figure 1). This will prompt a login window where you'll enter your PMA.core credentials to connect to the PMA.core server and load the slides in the tree view (Figure 2).

Figure 1: Load slides

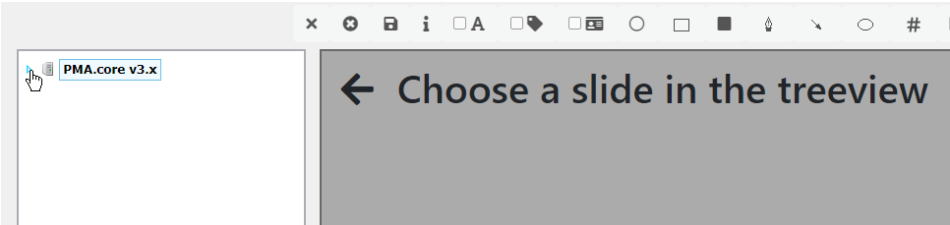
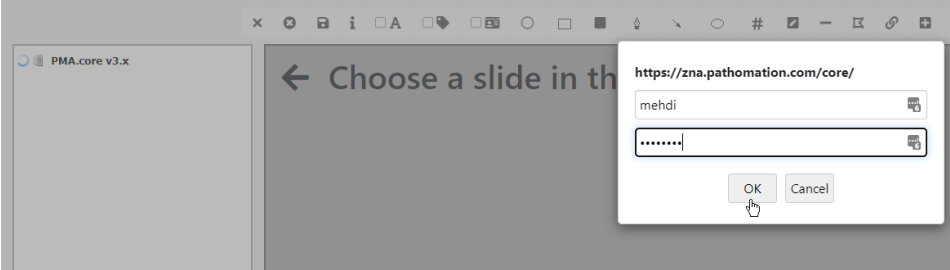


Figure 2: Load slides



Load slides on the viewer

After you logged into the PMA.core, you'll see the root directory appear. Expand this root directory which contains all your slides/cases with slides. Load your slides on the viewer by expanding one of the case folders and clicking on the slide you want to load or by clicking on the slide you want to load if the slides are not ordered by cases (Figure 3).

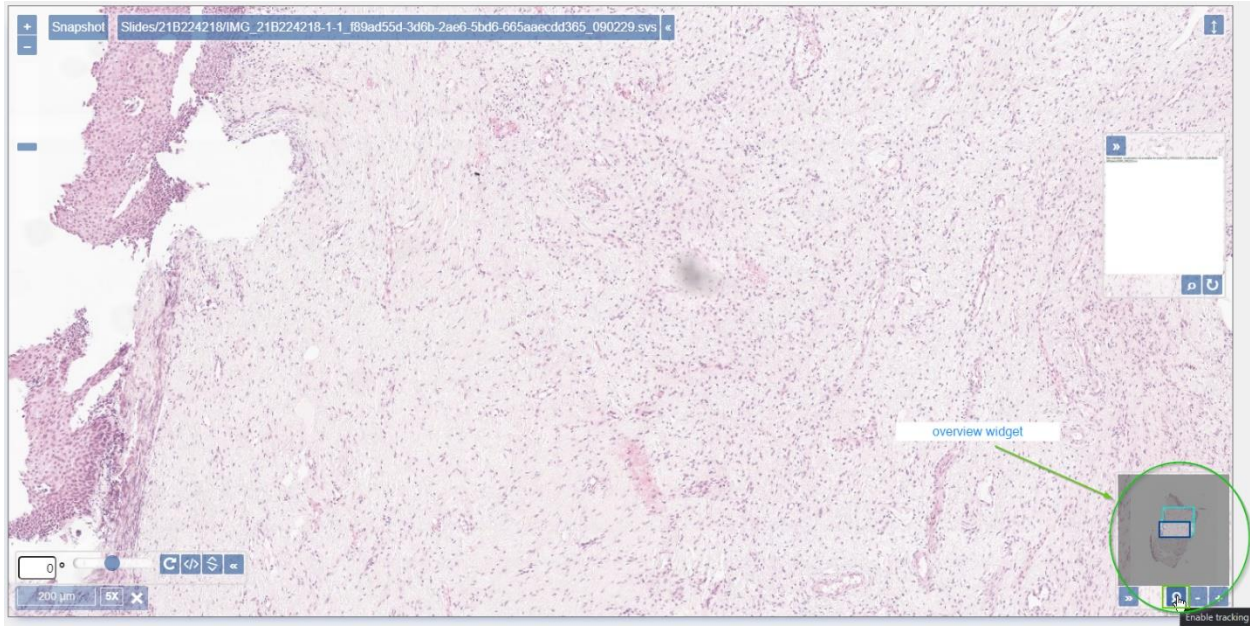
Figure 3: Load slides on the viewer



Enable tracking/heatmap

You enable tracking or heatmap in the overview widget by clicking on the magnifying glass on the overview widget. This will turn your overview widget into a heatmap where you can track where you already diagnosed the slide (Figure 4).

Figure 4: Enable tracking/heatmap



Annotations

Once you loaded the slide on the viewer, the annotations bar above gets enabled. The main annotations are the save annotations button, circle annotation, rectangle annotation, line annotation, and the measurement controls. The circle and rectangle annotations are set to 2 mm^2 . You can use the line annotation to calculate the distance. The difference between the line annotation and the measurement controls is that the measurement controls are not annotations. So, they can't be saved. Finally, you can use the save annotations button to save the annotations you made on the slide (Figure 5).

Figure 5: Annotations

