

# Unanswered questions from MedDream technical training (2022-03-22)

## Hanging Protocols configuration:

1. When I set a predefined W/W level for a specific series in a hanging protocol, it sets it but only for the displayed image of the series. As soon as I scroll the series, the predefined W/W in the hanging protocol is not applied to all this series images.
2. Could you demonstrate how to display two separate studies for the same patient side-by-side?

# 1 QUESTION:

**When I set a predefined W/W level for a specific series in a hanging protocol, it sets it but only for the displayed image of the series. As soon as I scroll the series, the predefined W/W in the hanging protocol is not applied to all this series images.**

If the series contains non single image files like multi-frame images, pdf and so on W/W level set does not apply. Otherwise, it should apply. If your study does not contain non single image files and W/W level from the hanging protocol is not applied, please send it to our support team for valuation:  
[support@softneta.com](mailto:support@softneta.com).

## 2 QUESTION:

**Could you demonstrate how to display two separate studies for the same patient side-by-side?**

**Step 1:** Create a Hanging protocol group for the study modality, that will be used:

The screenshot shows a 'CREATE GROUP' dialog box overlaid on a 'Hanging protocol' configuration screen. The dialog has a title bar with 'CREATE GROUP' and a close button. It contains several fields and options:

- Name \***: A text input field.
- Description**: A text input field.
- Devices**: A set of radio buttons with options: Mobile, Tablet, Desktop, 1 display (selected), and Desktop, 2 displays.
- Group conditions**: A section with an 'Add condition' link.
- Active**: A set of radio buttons with options: Enable and Disable (selected).

At the bottom of the dialog are 'Create' and 'Close' buttons. In the background, the 'Hanging protocol' screen shows a list of groups, with 'Add new group' highlighted by a red box.

# 2 QUESTION:

**Step 2:** Click the “Add new protocol” button to create a new Hanging protocol:

**CREATE PROTOCOL** [X]

Hanging protocol

Metadata fetching

Do not fetch metadata for these modalities

**Add new protocol**

Group for desktop MG

Group for CT (all devices)

Group for desktop US

Group for desktop XA

Group for desktop OT

Prior

Description

Layout columns: 1

Layout rows: 1

Study conditions: Add condition

Views: 1x1

Prior conditions: Add condition

Windowing: Width, Center

Rotation: 0

Instance conditions: Add condition

Flip horizontally: Enable, **Disable**

Flip vertically: Enable, **Disable**

**Create** Close

## 2 QUESTION:

**Step 3:** Set the layout rows and columns. In the preferred layout section add a precondition for the previous study.



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**Step 3:** Set the layout rows and columns. In the preferred layout section add a precondition for the previous study.



**Step 4:** In the precondition section select “no NumberOfPriorStudy” and enter the number of the study in the Patient History list. The operand is always set to “=”.

**NOTE:** Please note, that the number of the prior study is taken from the patient history list. If 3 is set as the number, then the third study will be taken from the list that is ordered from oldest to newest.