Subject: Remove 'abs' and other chiral center labels in structure depiction Posted by zhentg on Mon, 16 Mar 2020 15:26:55 GMT View Forum Message <> Reply to Message

chiral labels are mandatory in structure depiction(pls check the attached image), it is helpful, but if I want to present the structures in a slide or manuscript, I want to remove these labels to keep the structure depiction simple and clear.

How to remove the chiral labels.

File Attachments

, downloaded 629 times

Subject: Re: Remove 'abs' and other chiral center labels in structure depiction Posted by zhentg on Wed, 25 Mar 2020 08:52:19 GMT View Forum Message <> Reply to Message

It would be nice if there is an option to switch on/off the 'abs' label display.

Subject: Re: Remove 'abs' and other chiral center labels in structure depiction Posted by thomas on Sun, 12 Jul 2020 19:08:25 GMT View Forum Message <> Reply to Message

For the structure view there are now flexible structure display modes concerning and stereo labels and usage of color. With this structure images should be more appropriate for presentation, papers or patents.

This change is available in the current development version.

Subject: Re: Remove 'abs' and other chiral center labels in structure depiction Posted by zhentg on Wed, 09 Sep 2020 00:24:30 GMT View Forum Message <> Reply to Message

Thanks for the added option. I can turn off the stereo labels in "Structure View" panel now through the "Set Structure Stereo Display Mode" option. Am I using the function correctly?

File Attachments 1) dw.png, downloaded 562 times Subject: Re: Remove 'abs' and other chiral center labels in structure depiction Posted by thomas on Sun, 13 Sep 2020 10:13:14 GMT View Forum Message <> Reply to Message

Looks OK to me. One should be aware that in case of multiple stereo centers informarmation may be lost, e.g. in case of epimers. The 'stereo display mode' can be combined with the 'color display mode' as patent layers seem to prefer black & white structures with no stereo labels.

Page 2 of 2 ---- Generated from openmolecules.org Forum