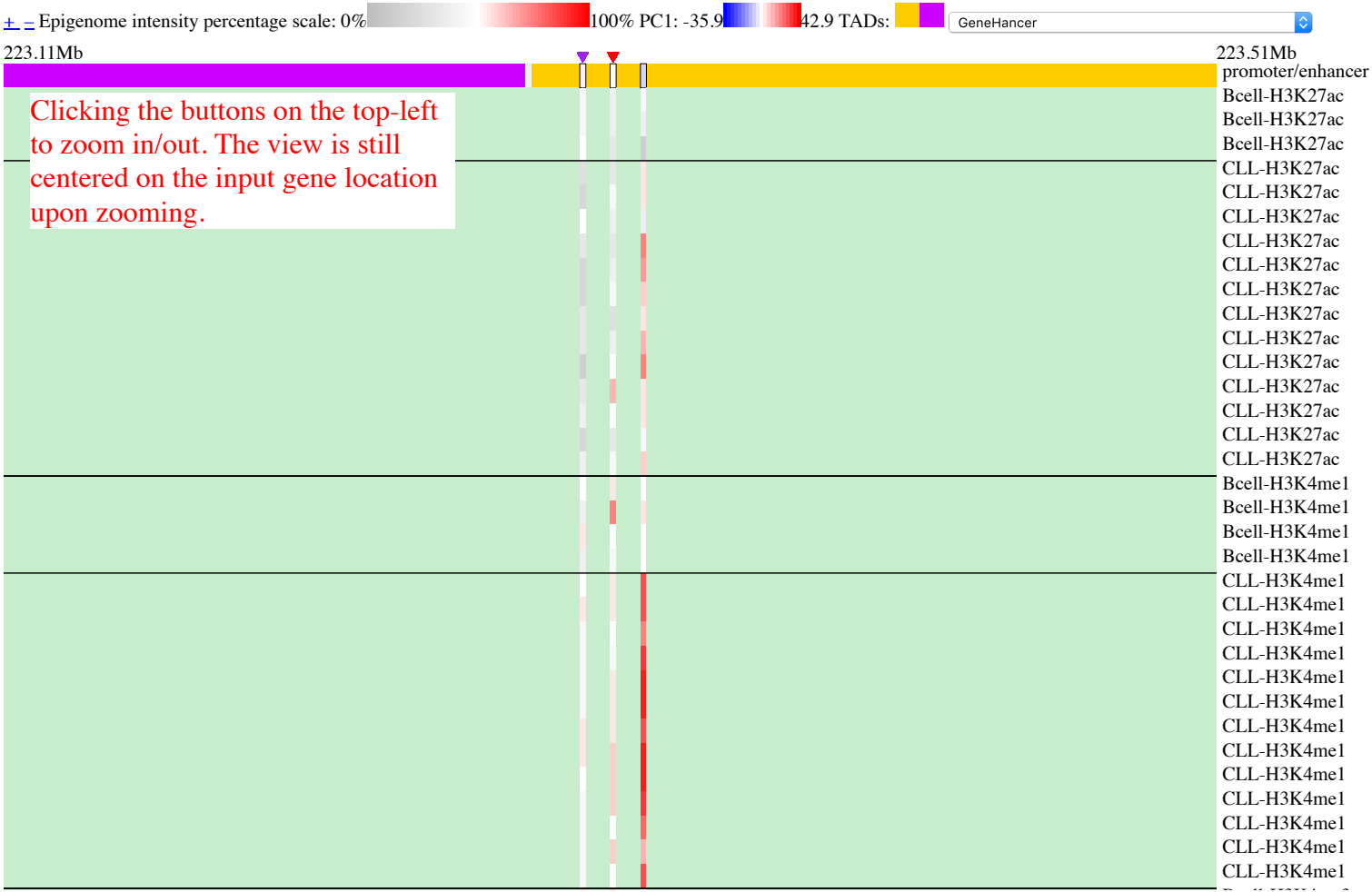

Clicking the buttons on the top-left to zoom in/out. The view is still centered on the input gene location upon zooming.