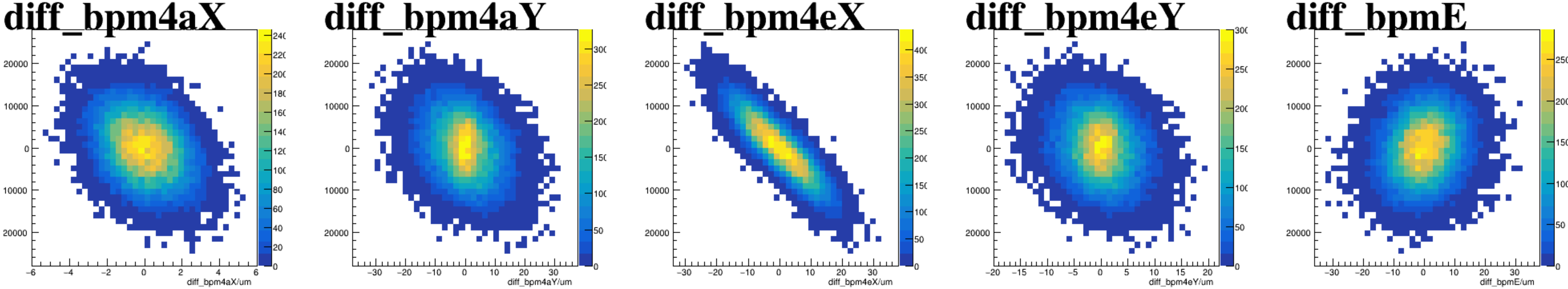
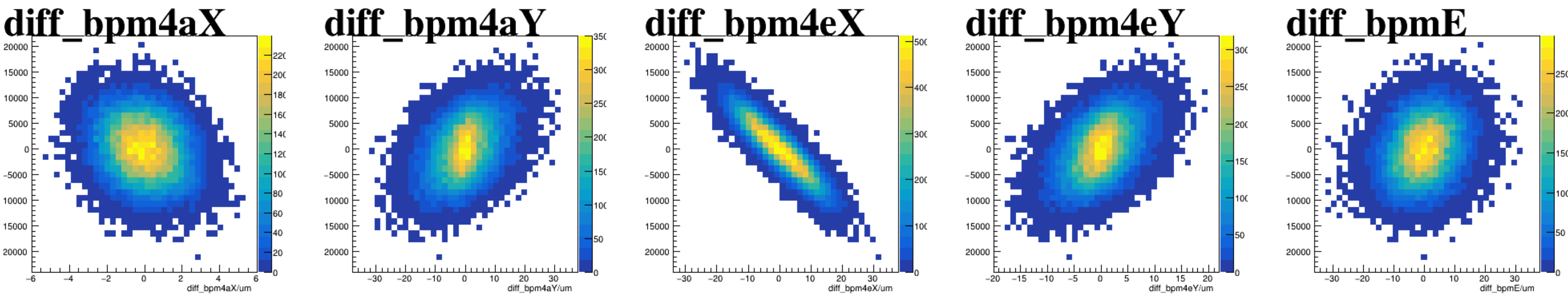


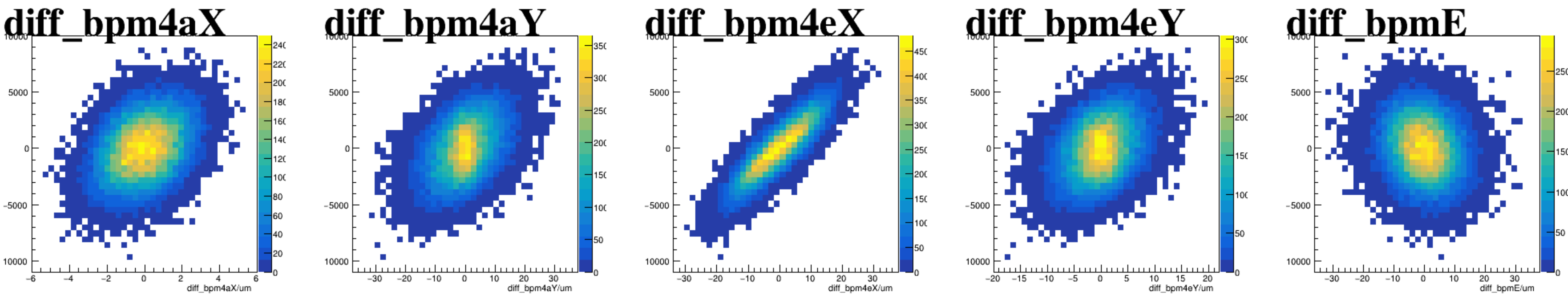
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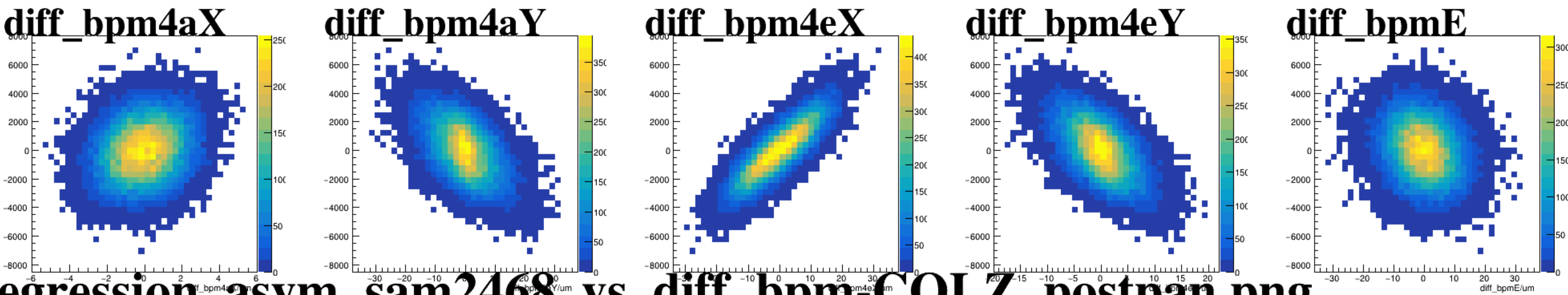
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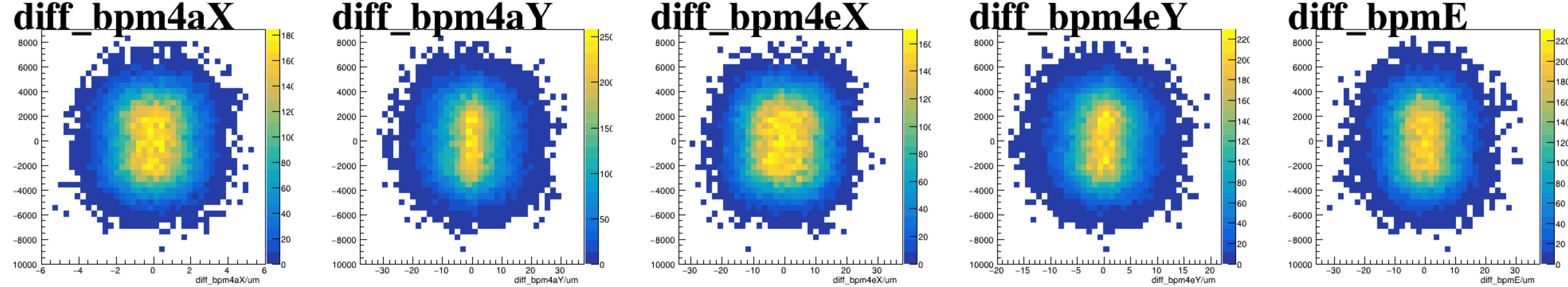
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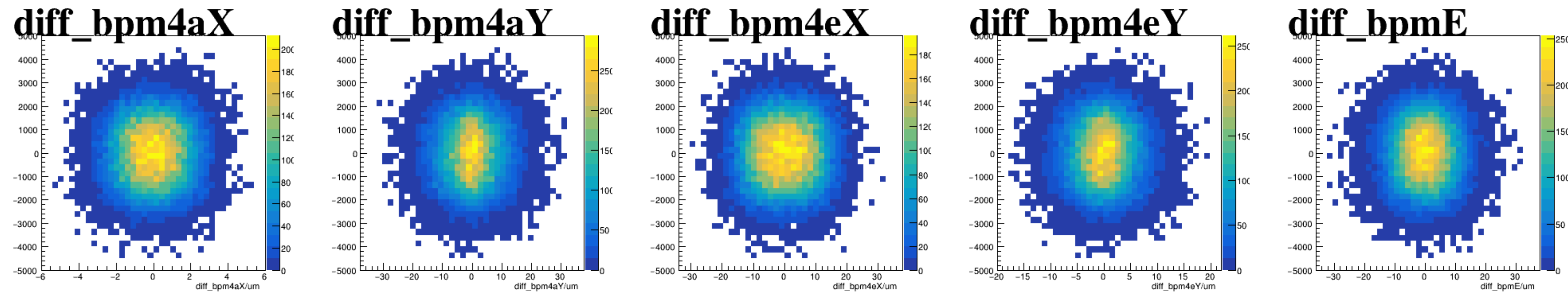
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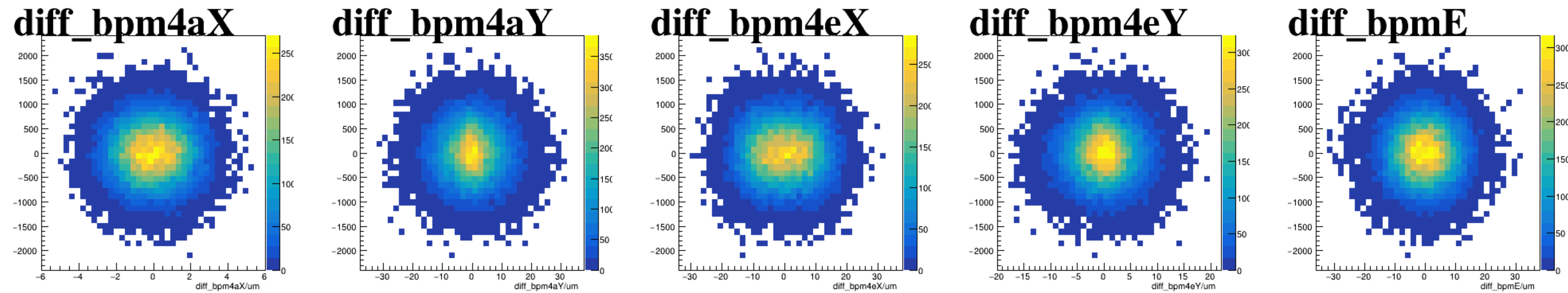
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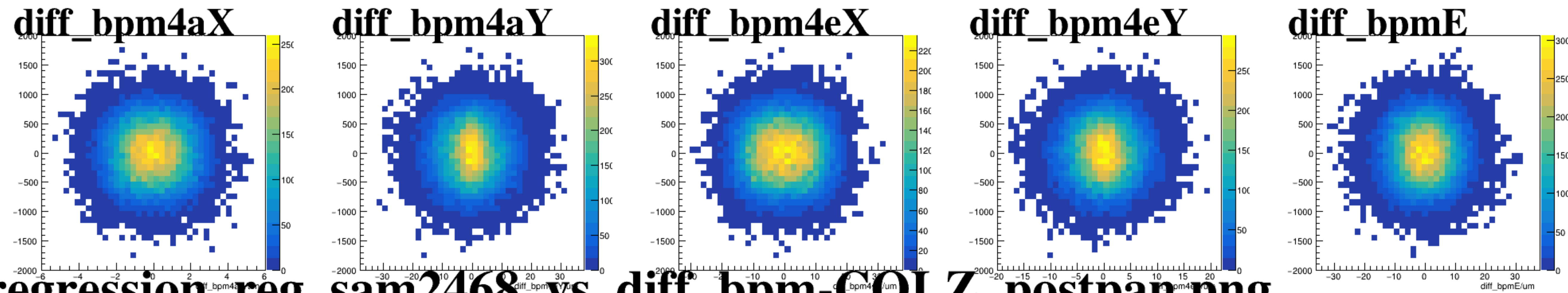
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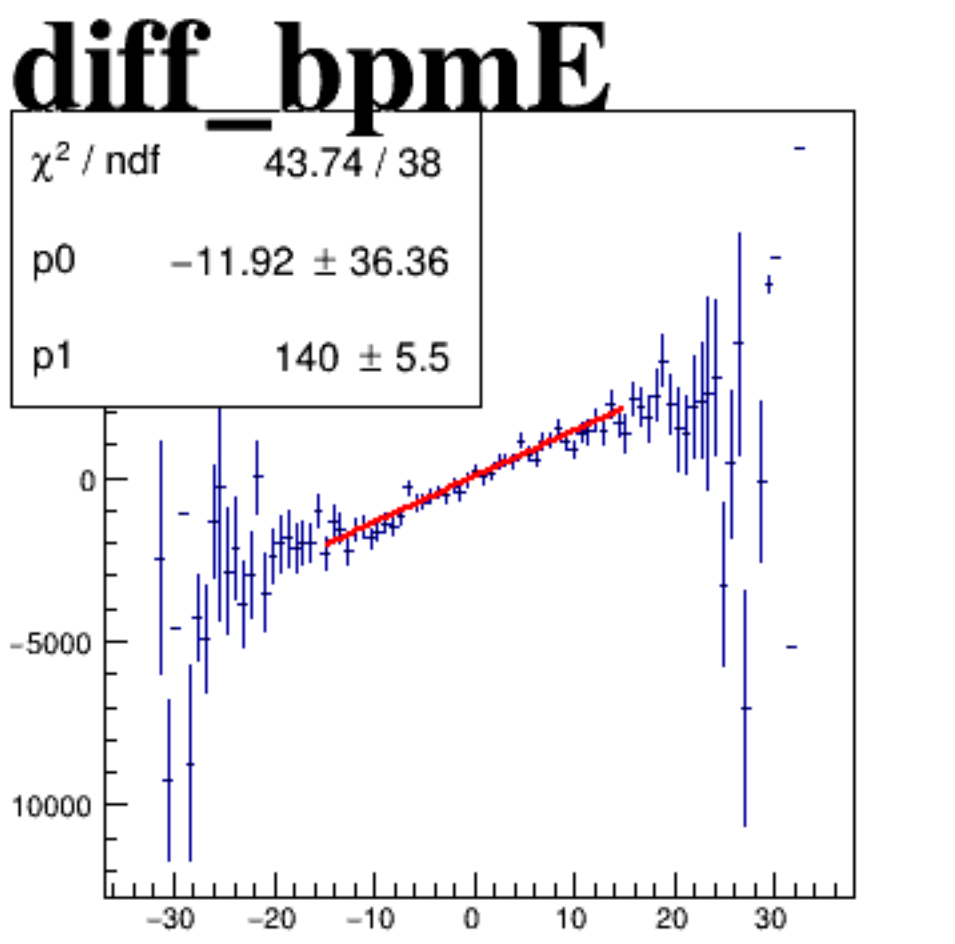
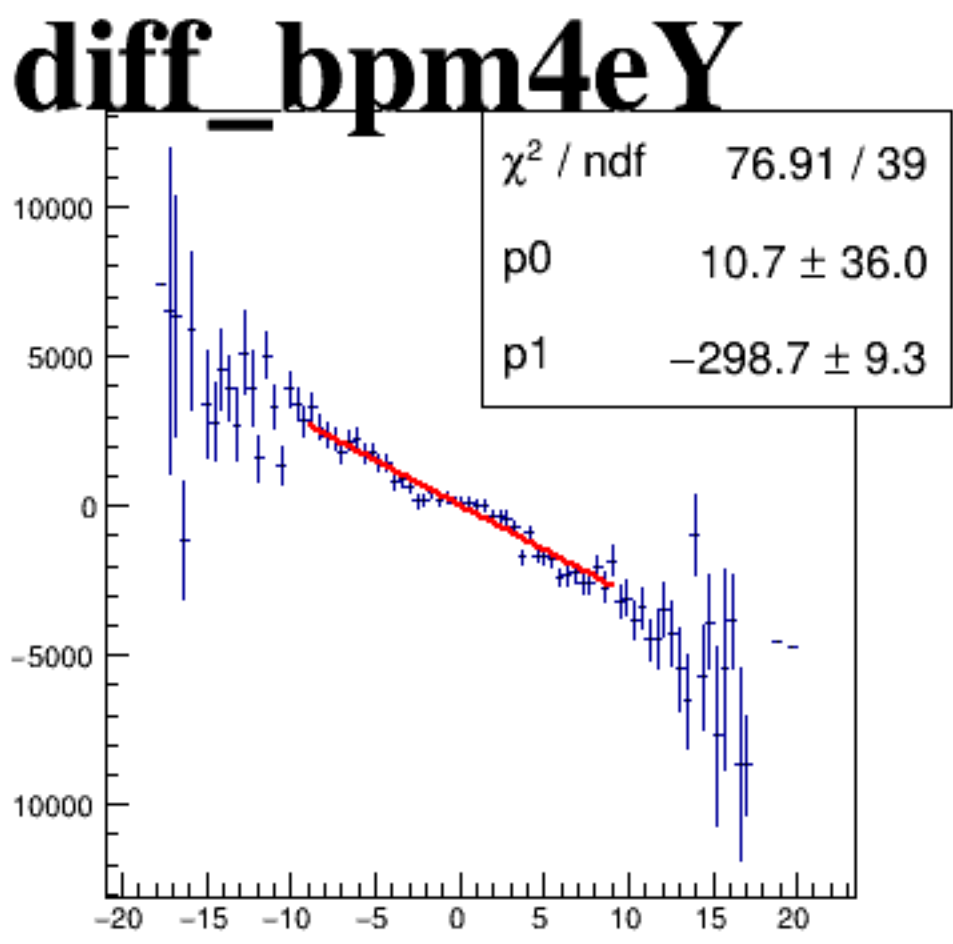
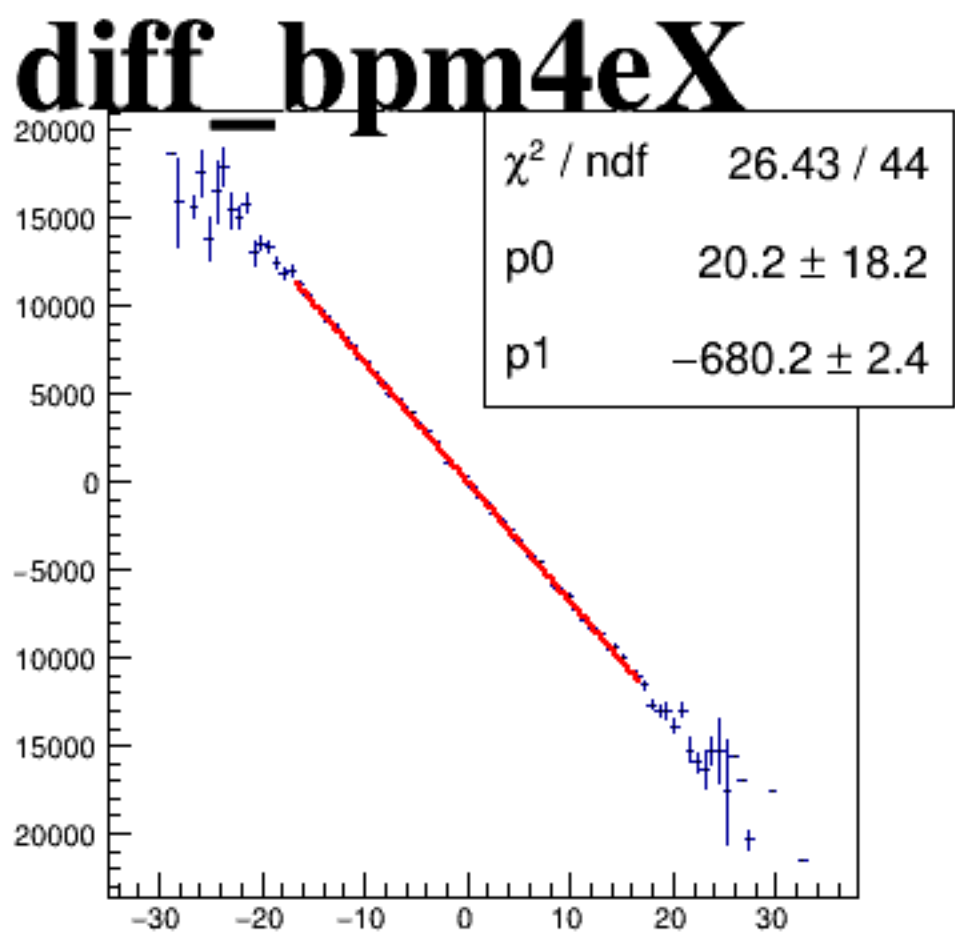
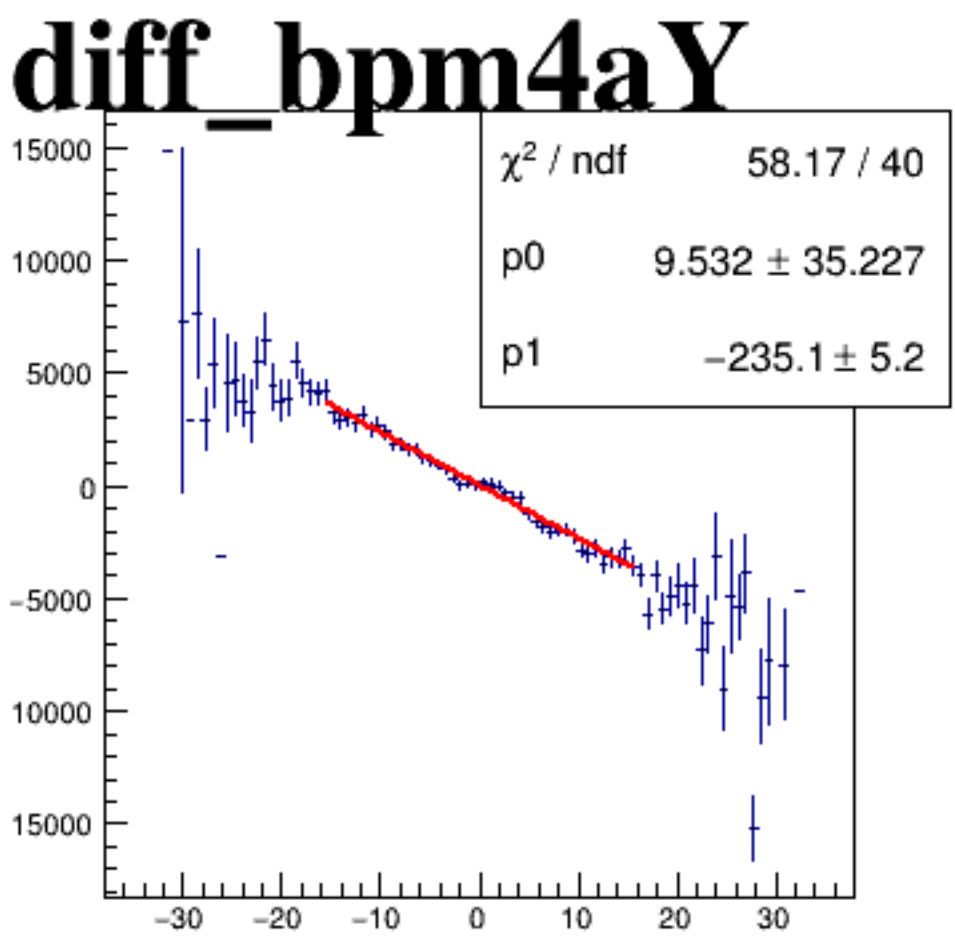
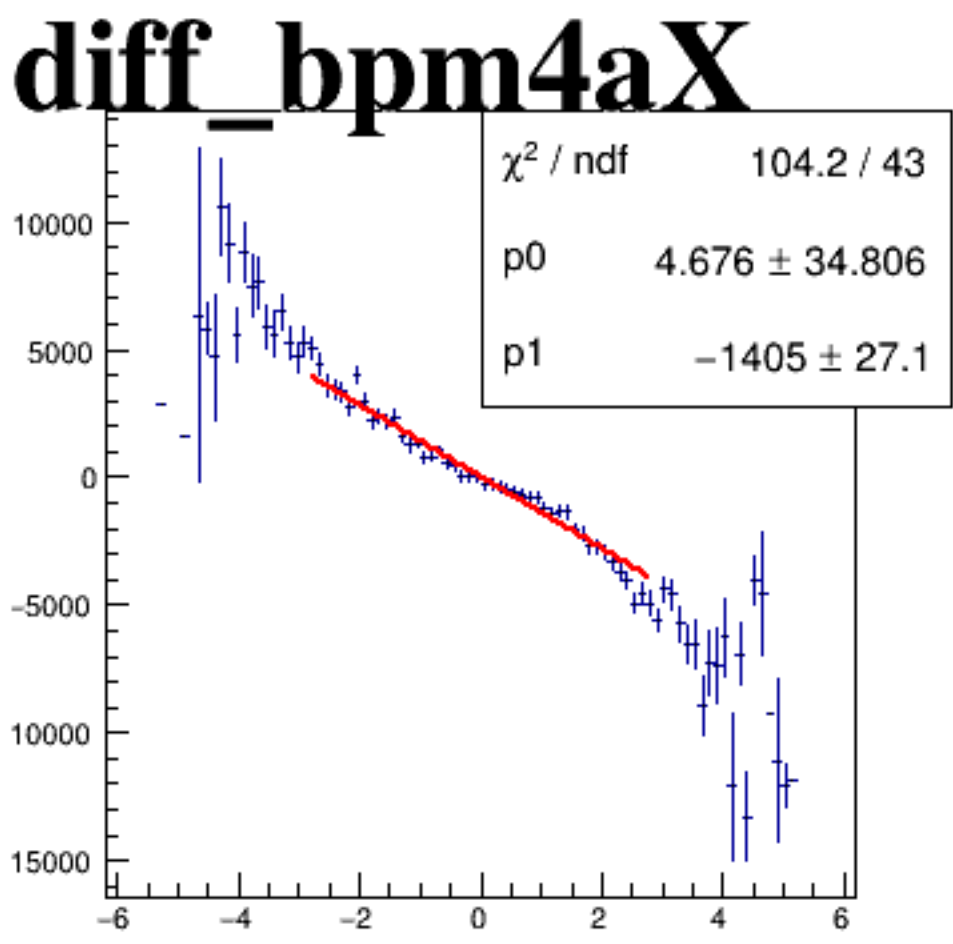
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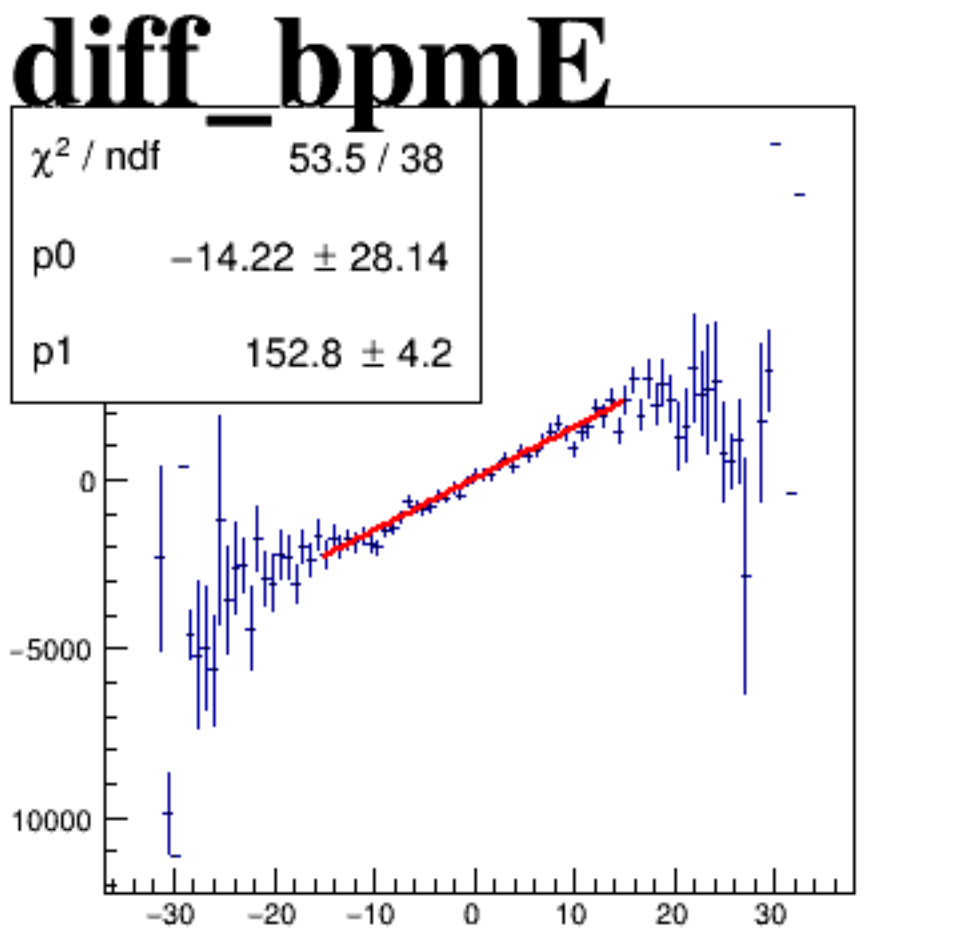
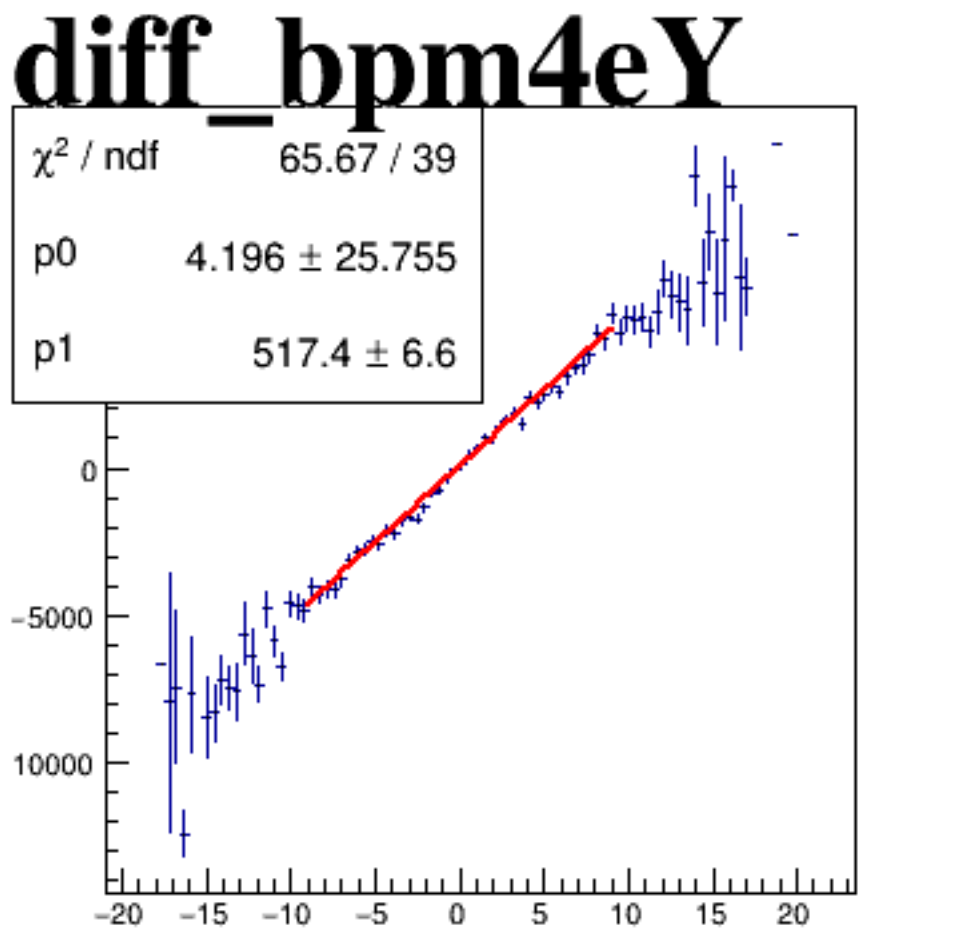
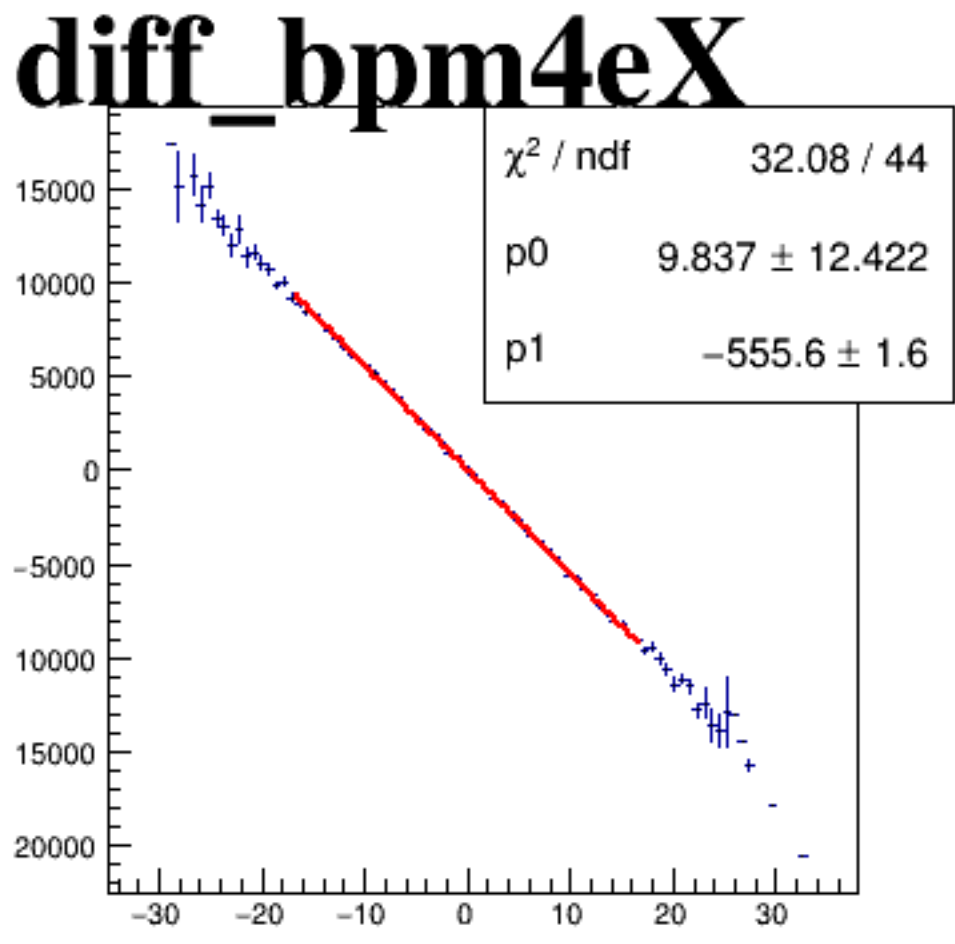
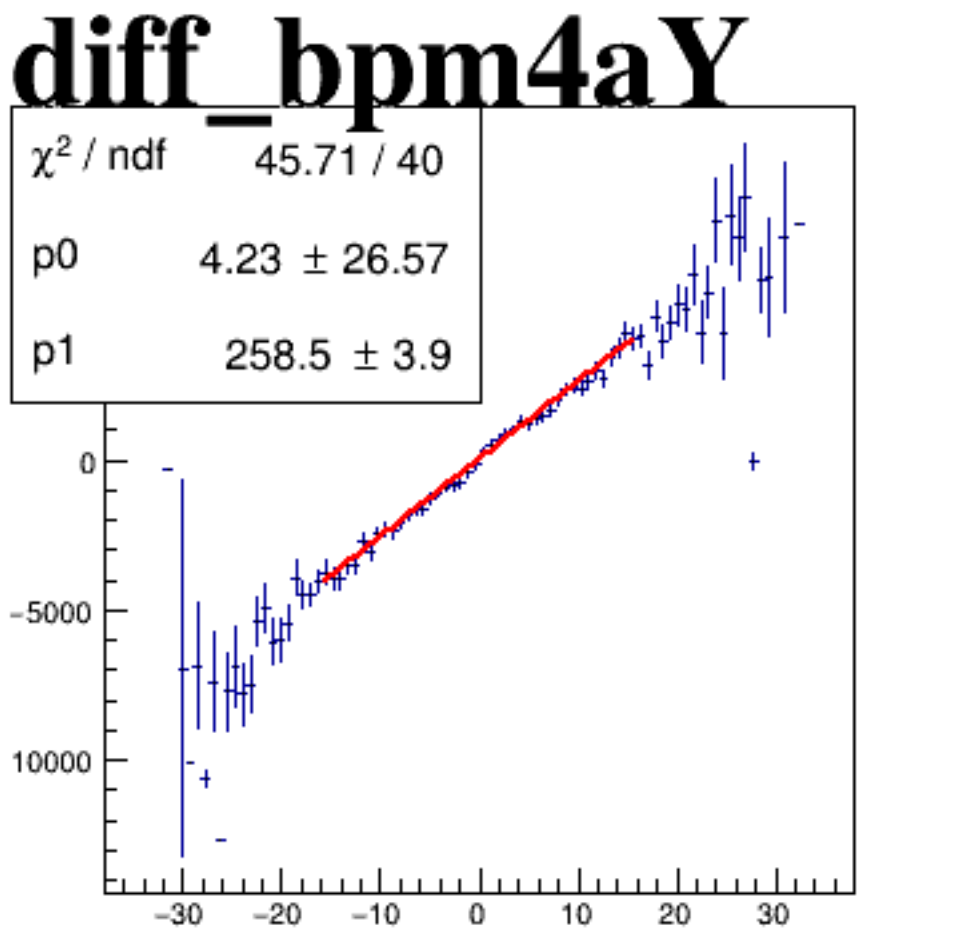
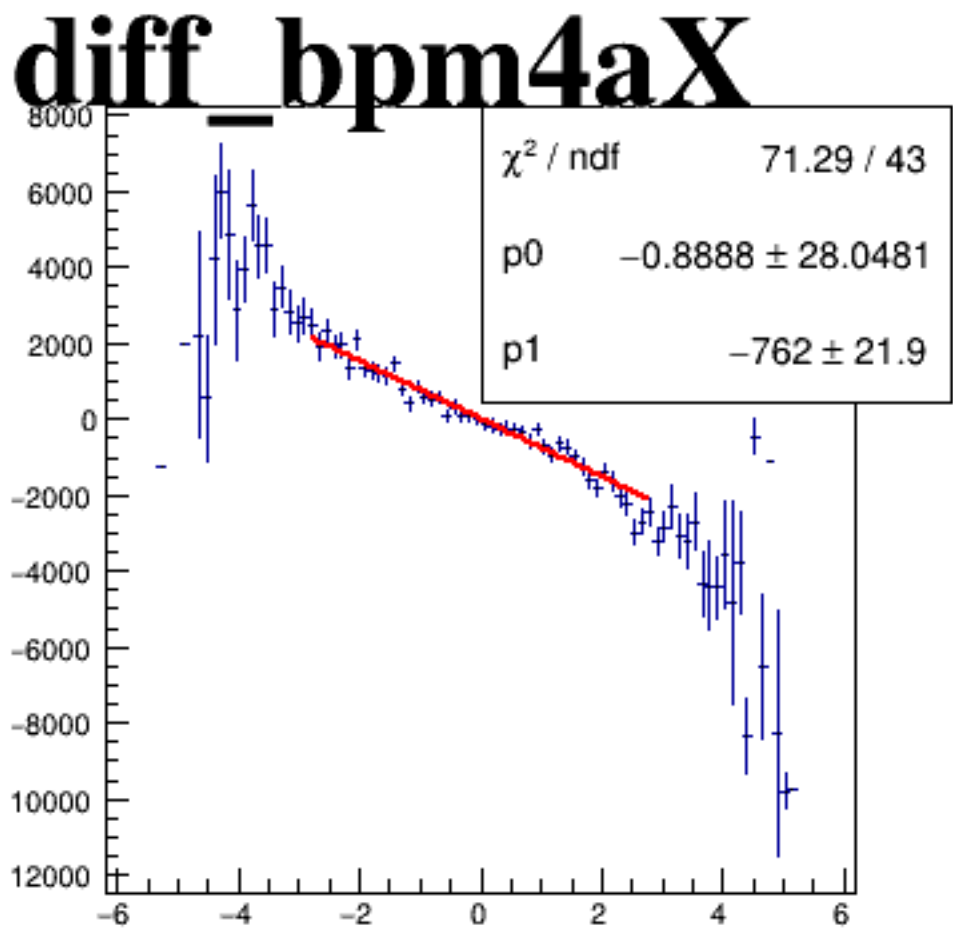
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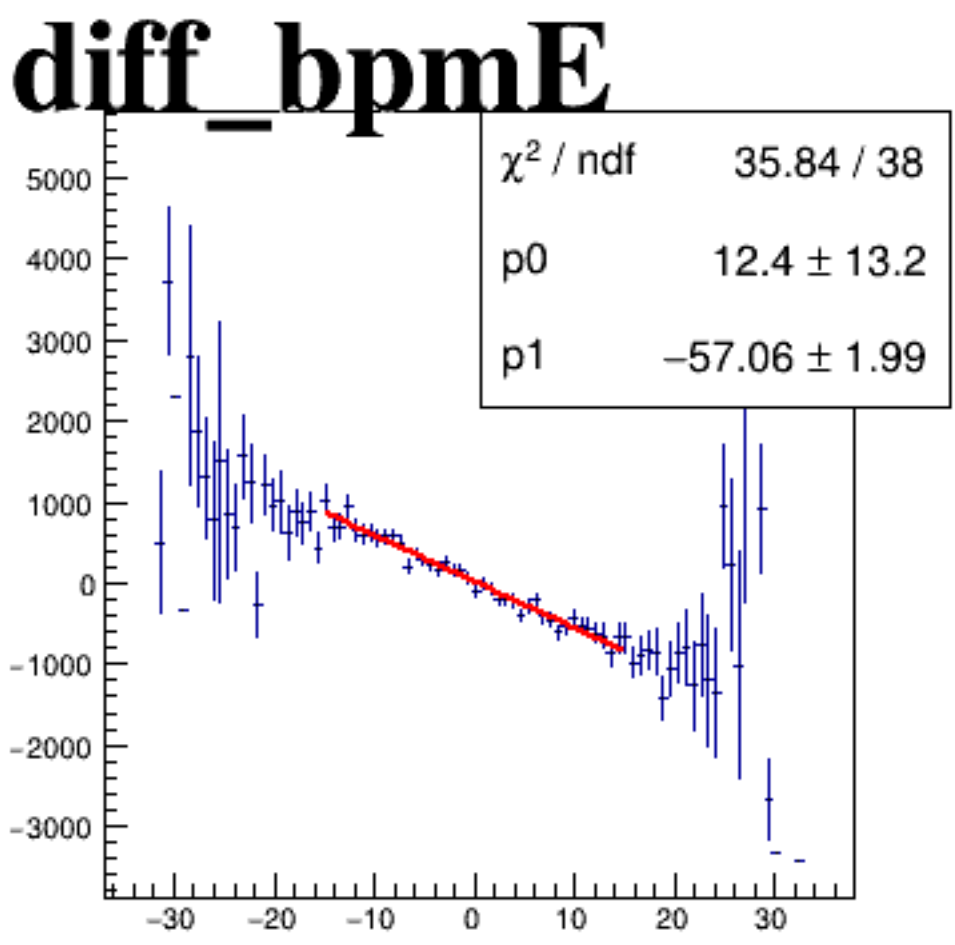
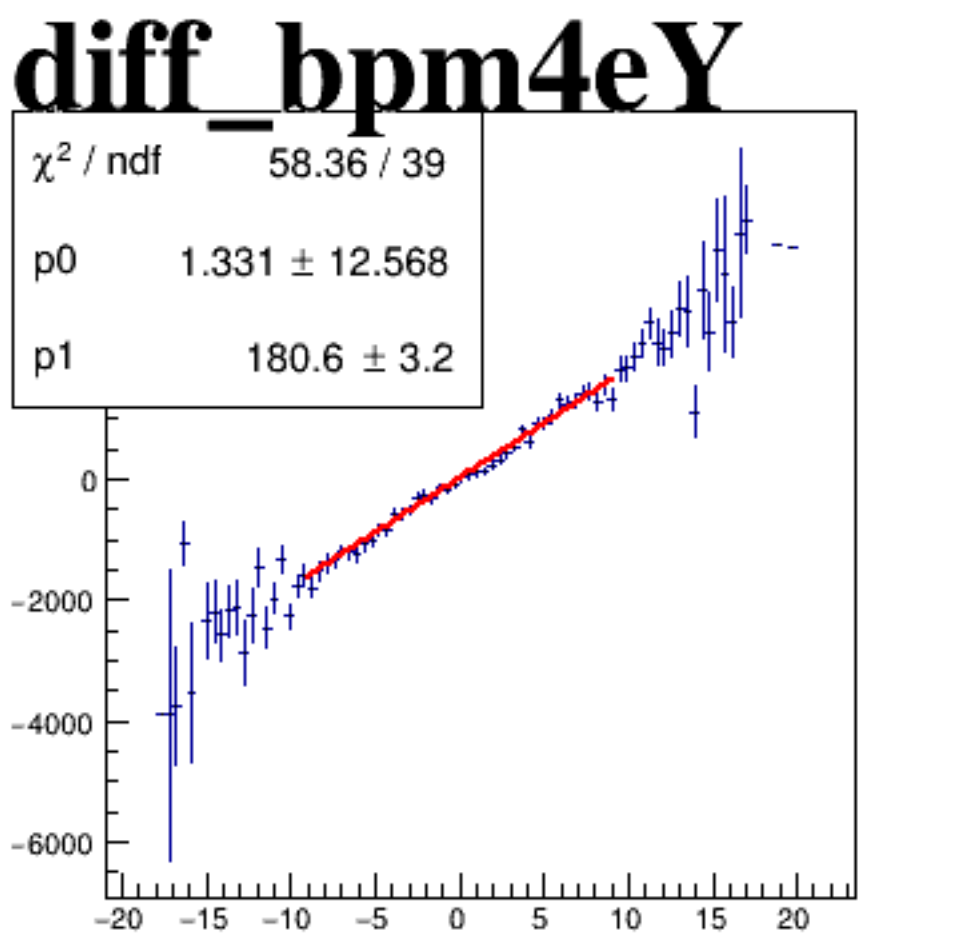
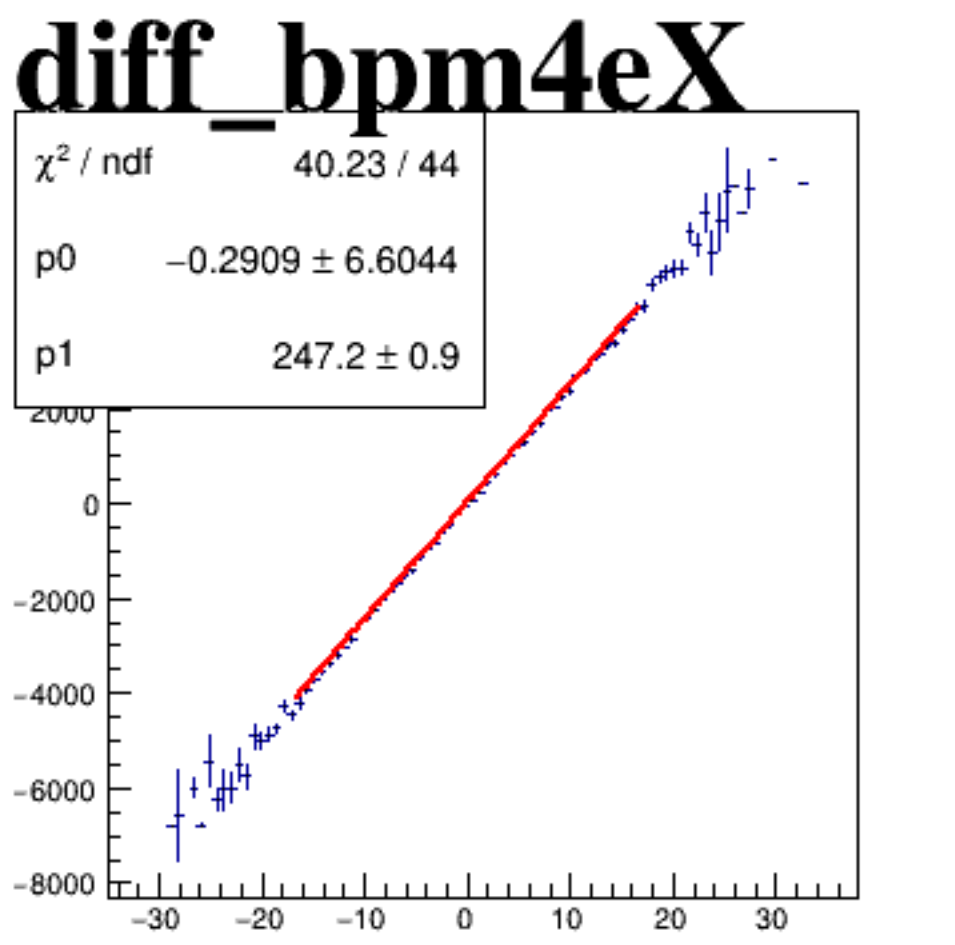
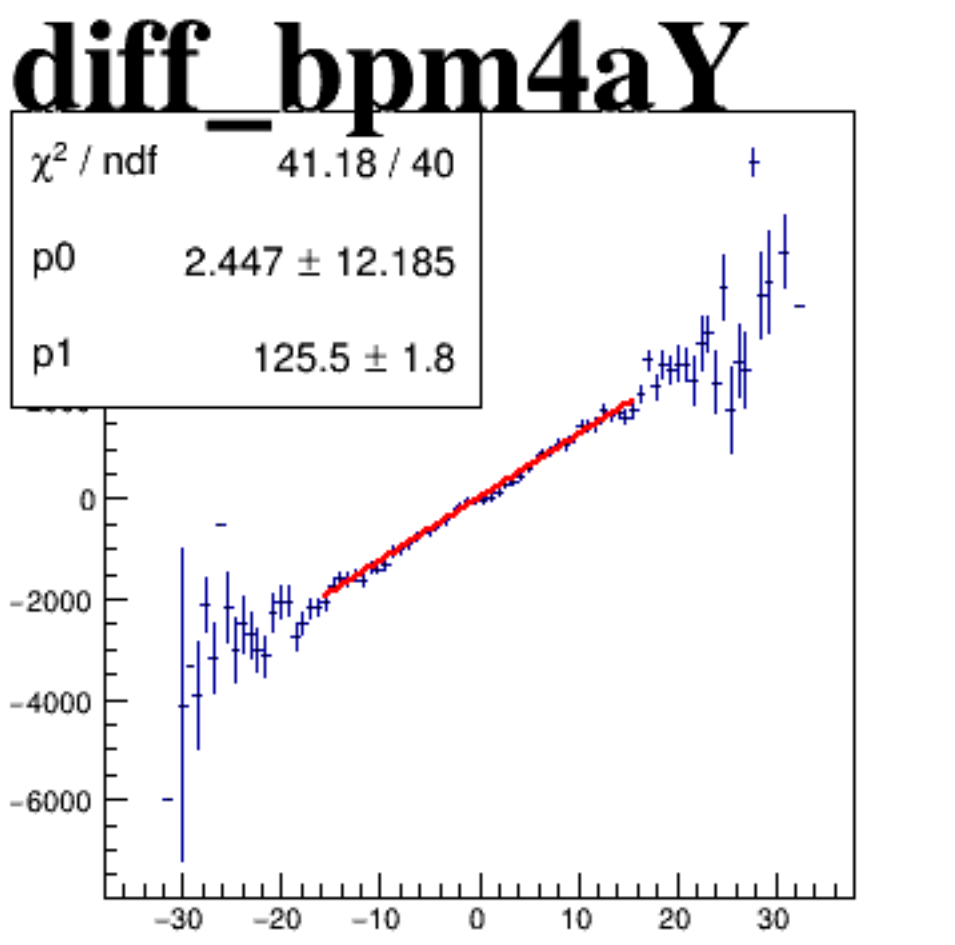
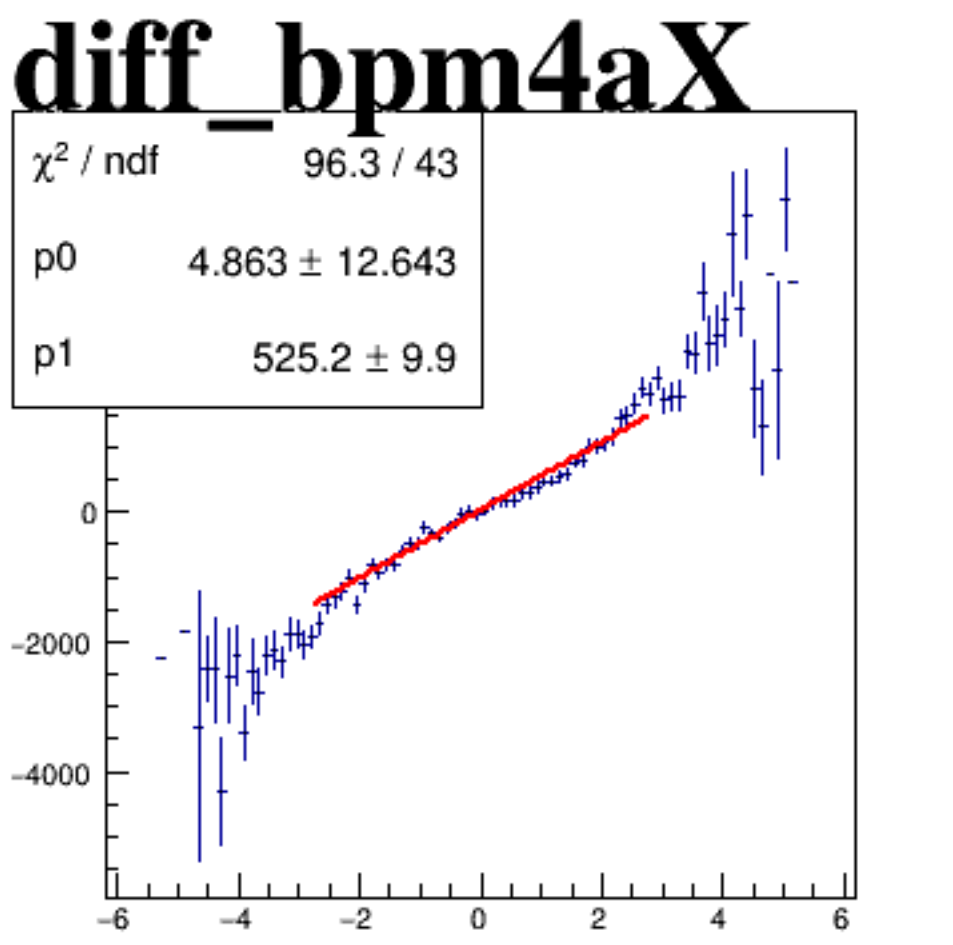
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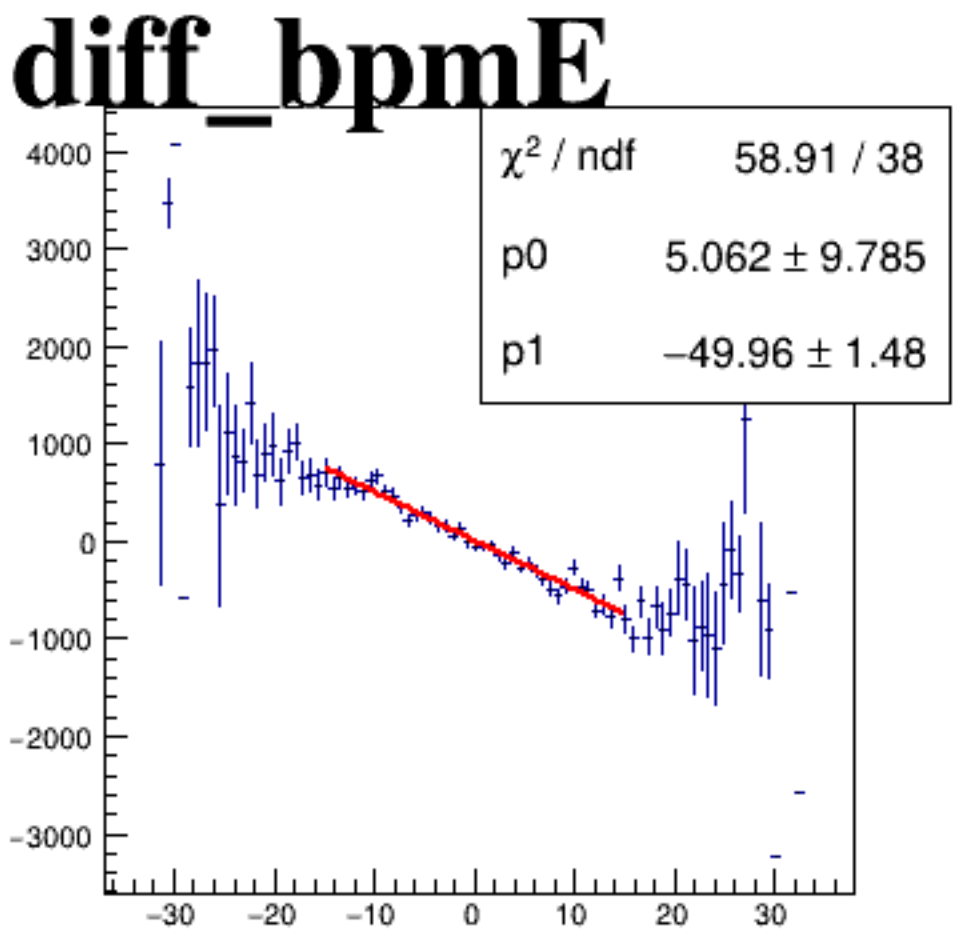
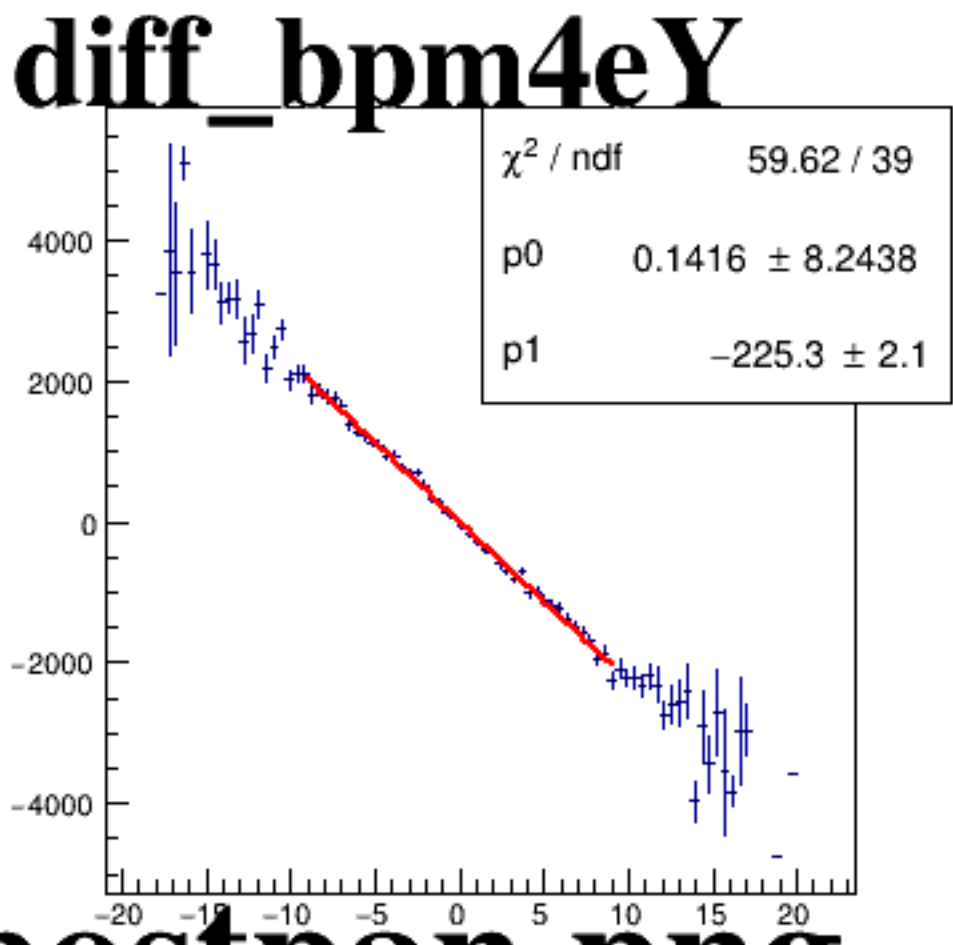
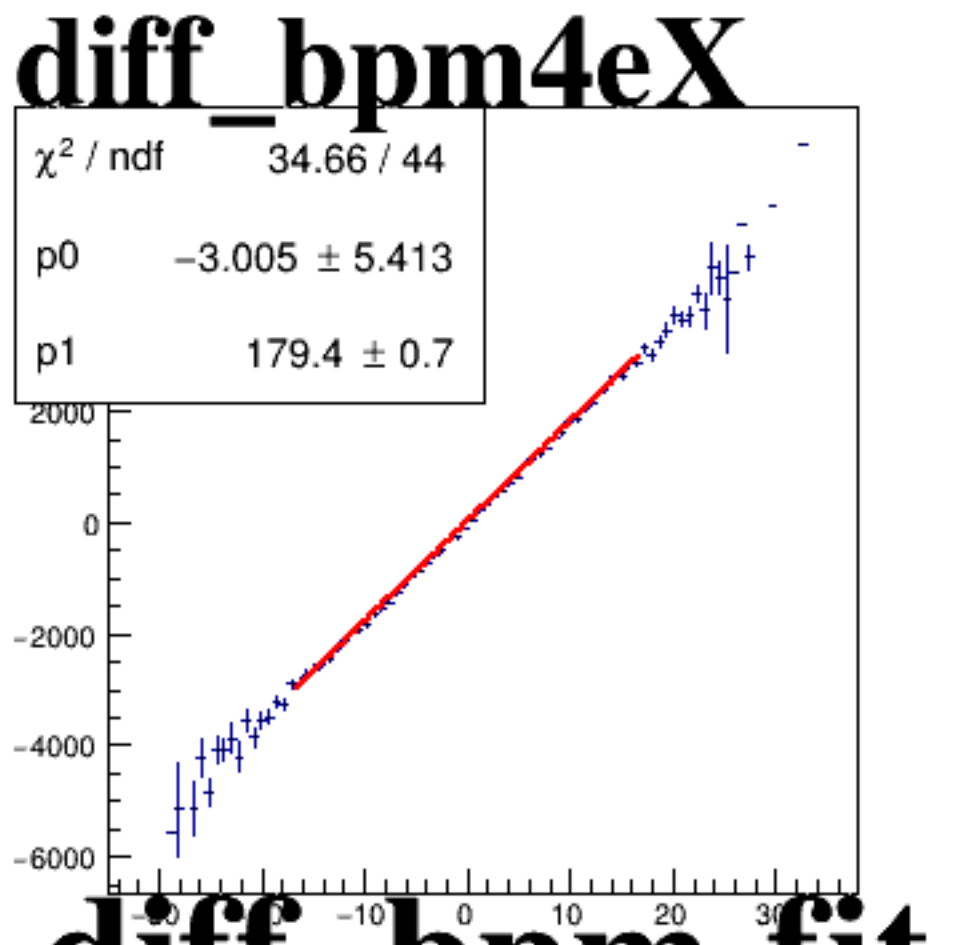
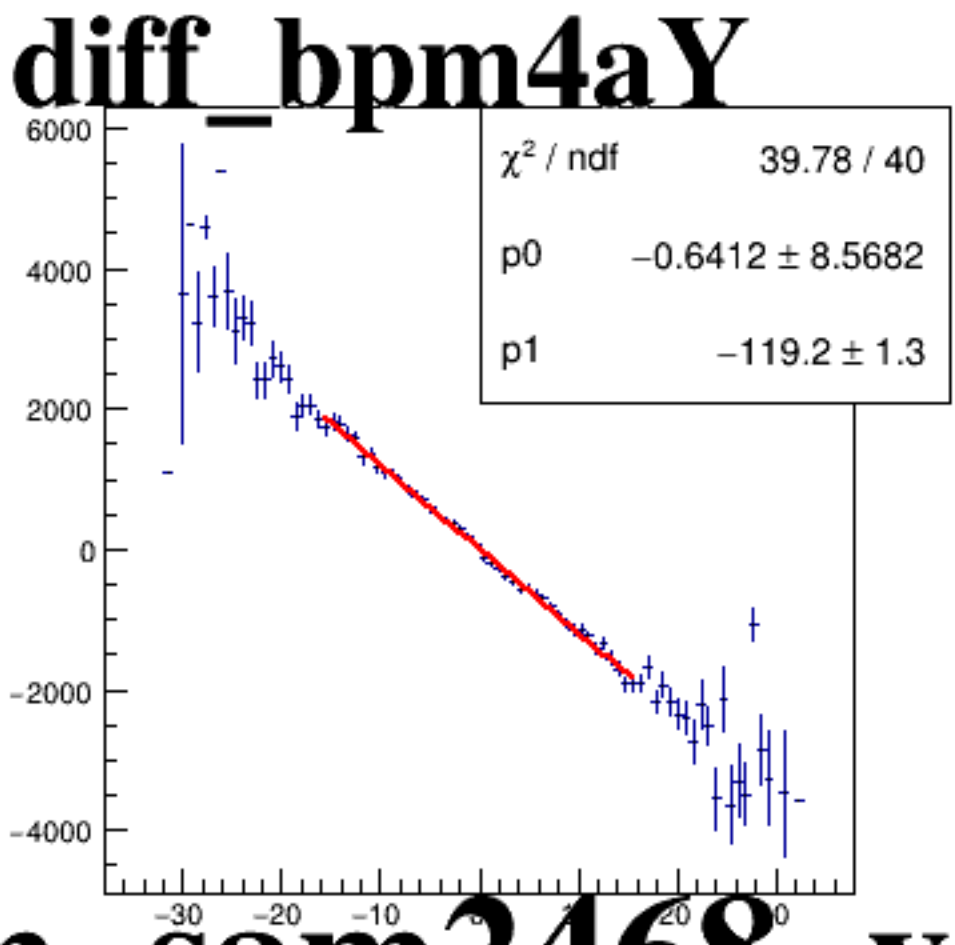
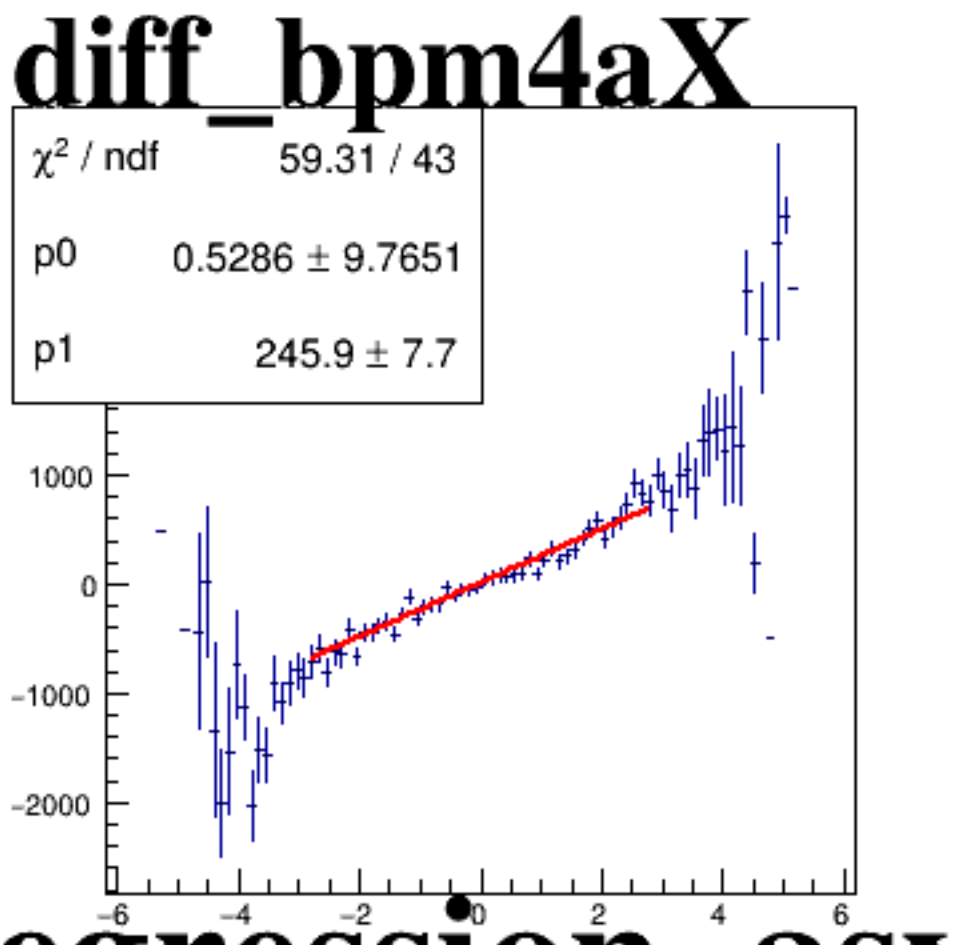
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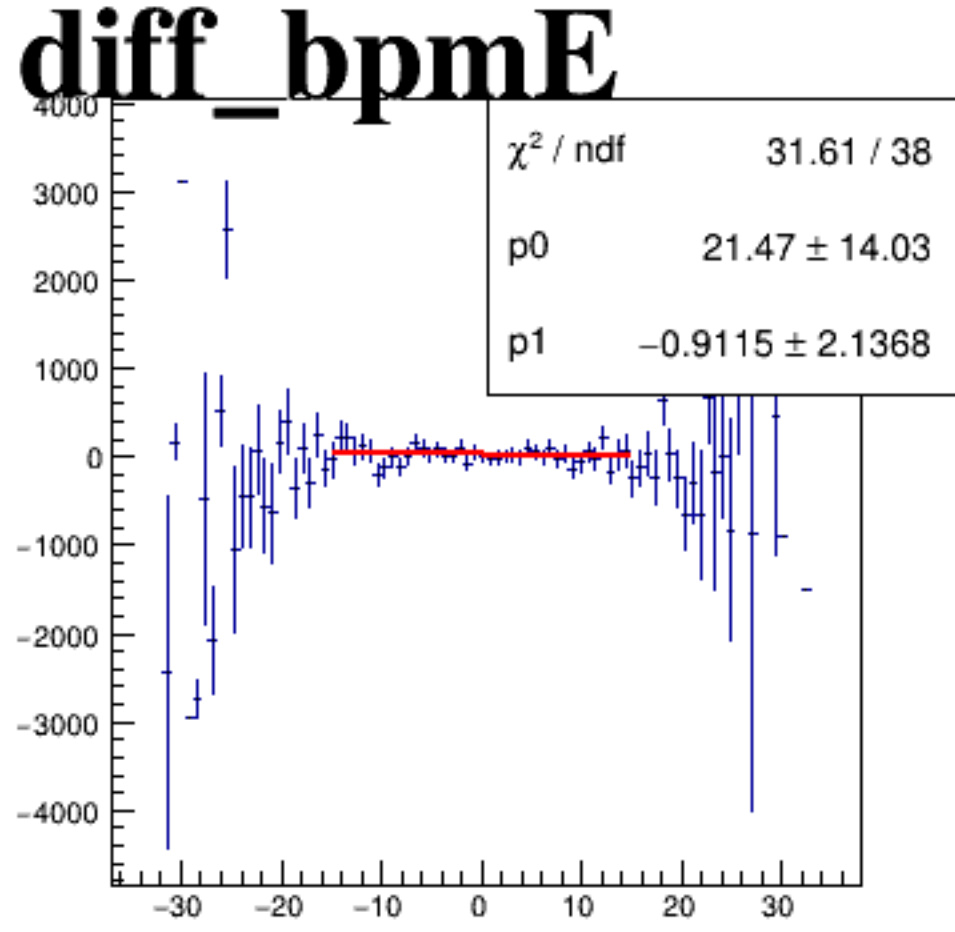
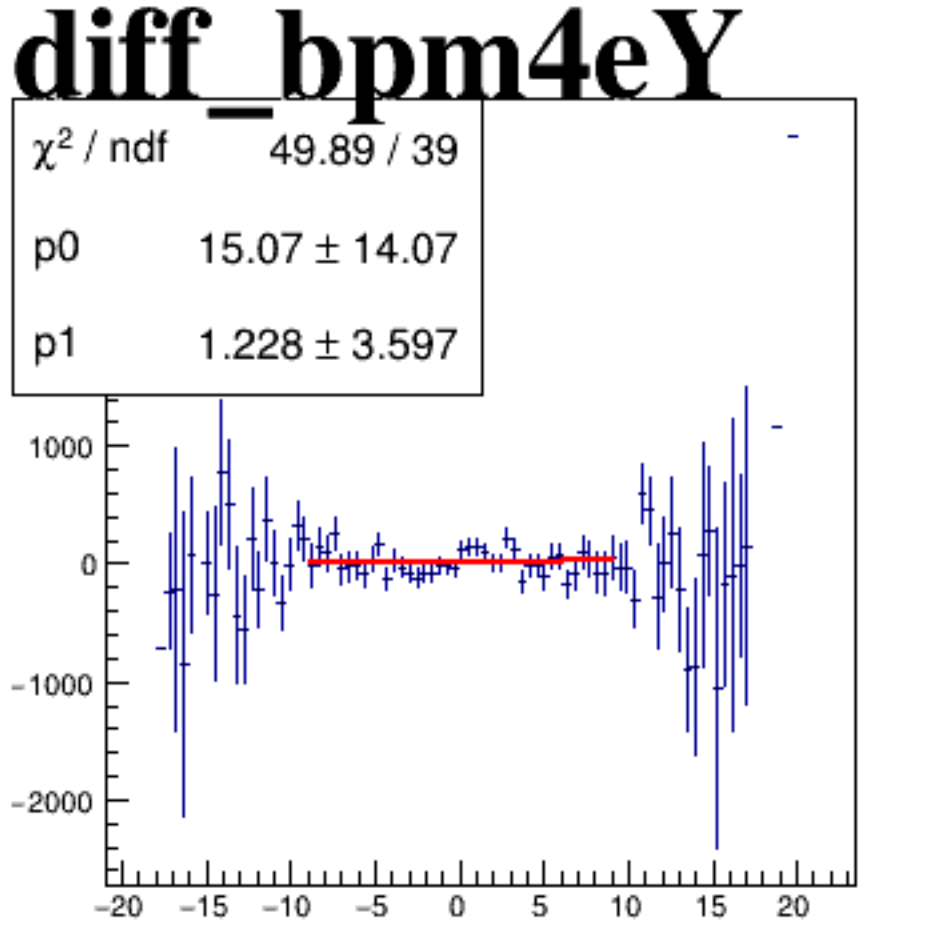
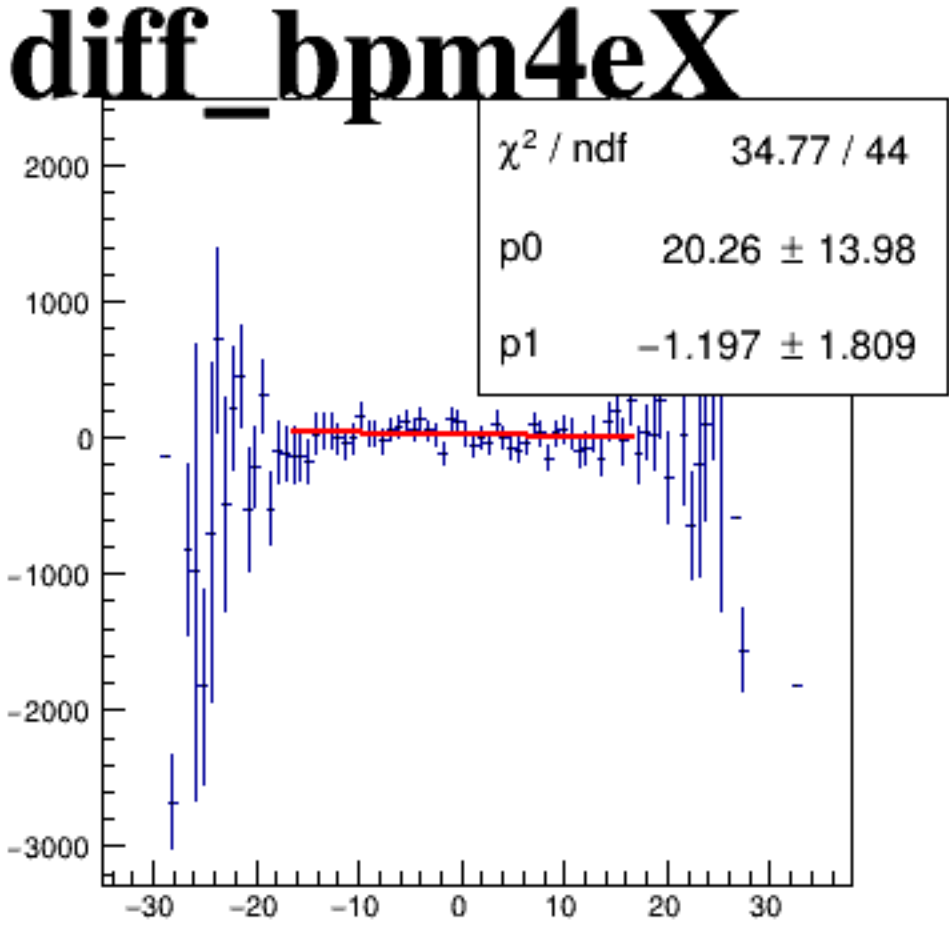
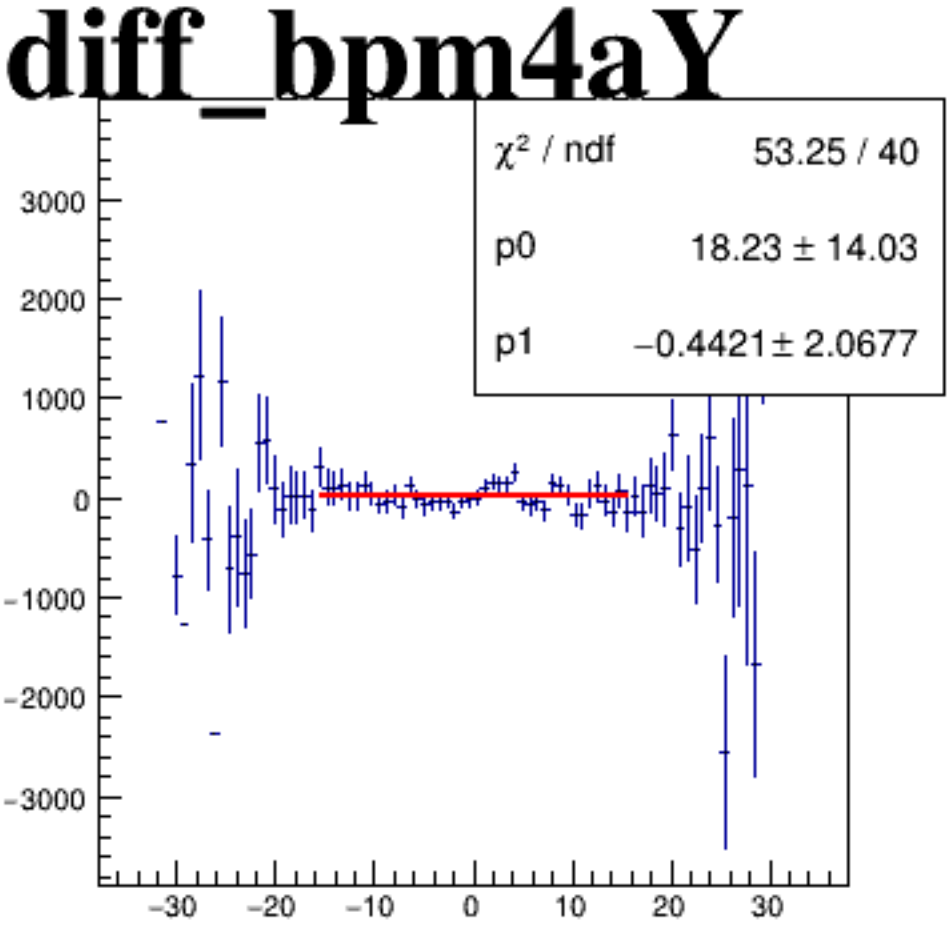
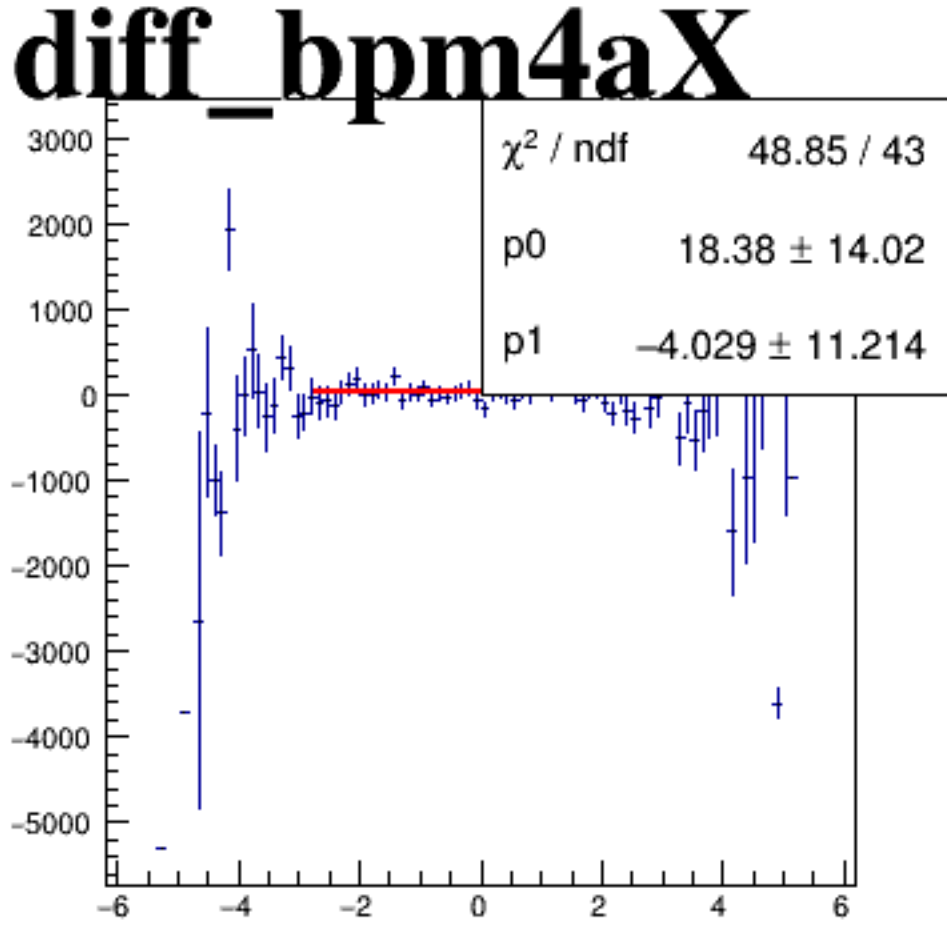
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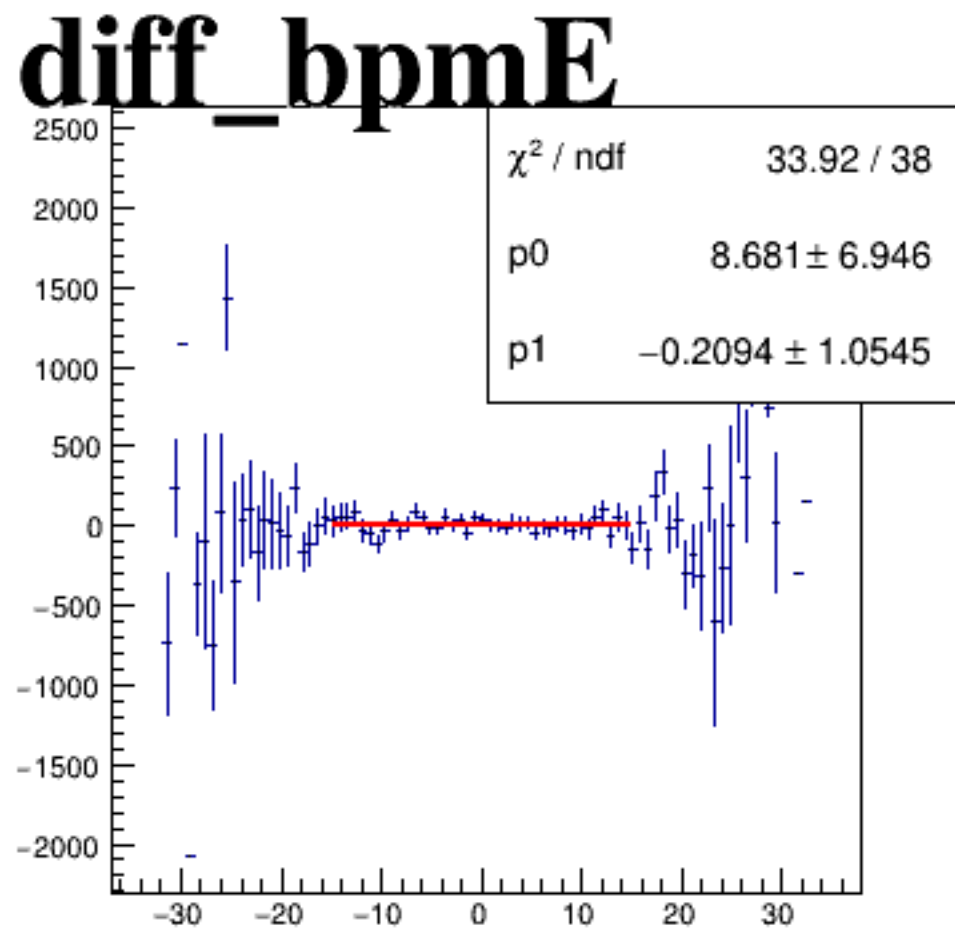
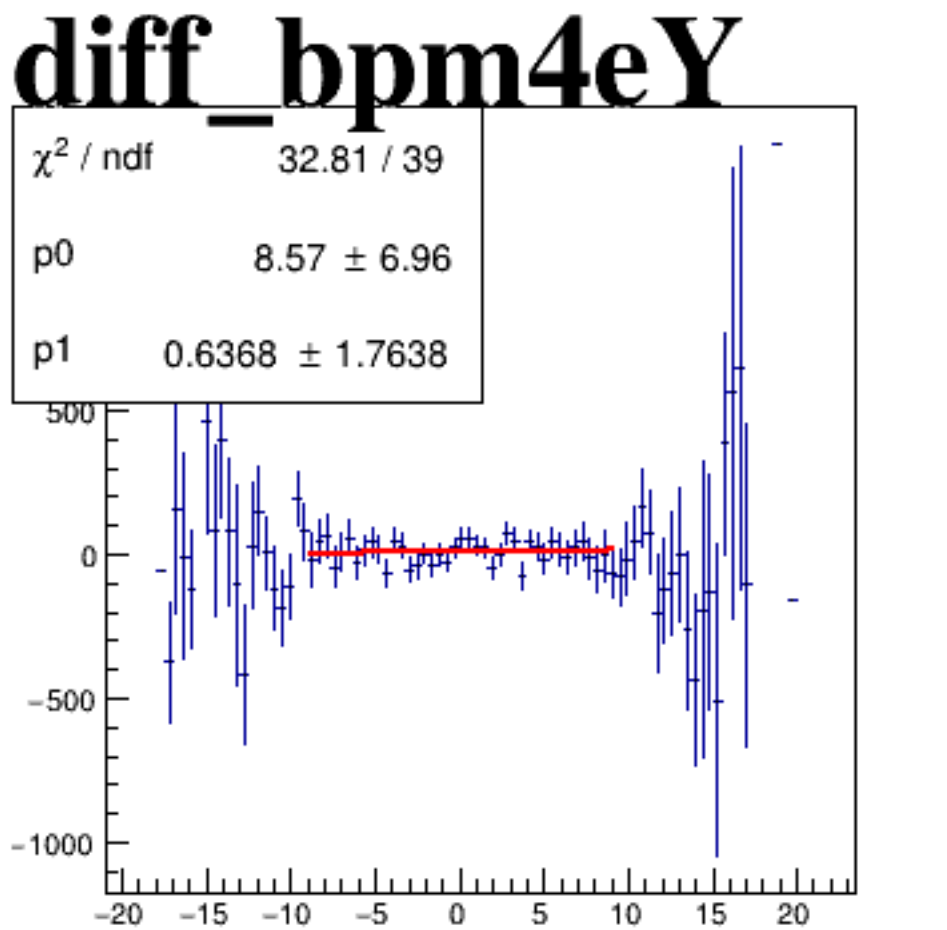
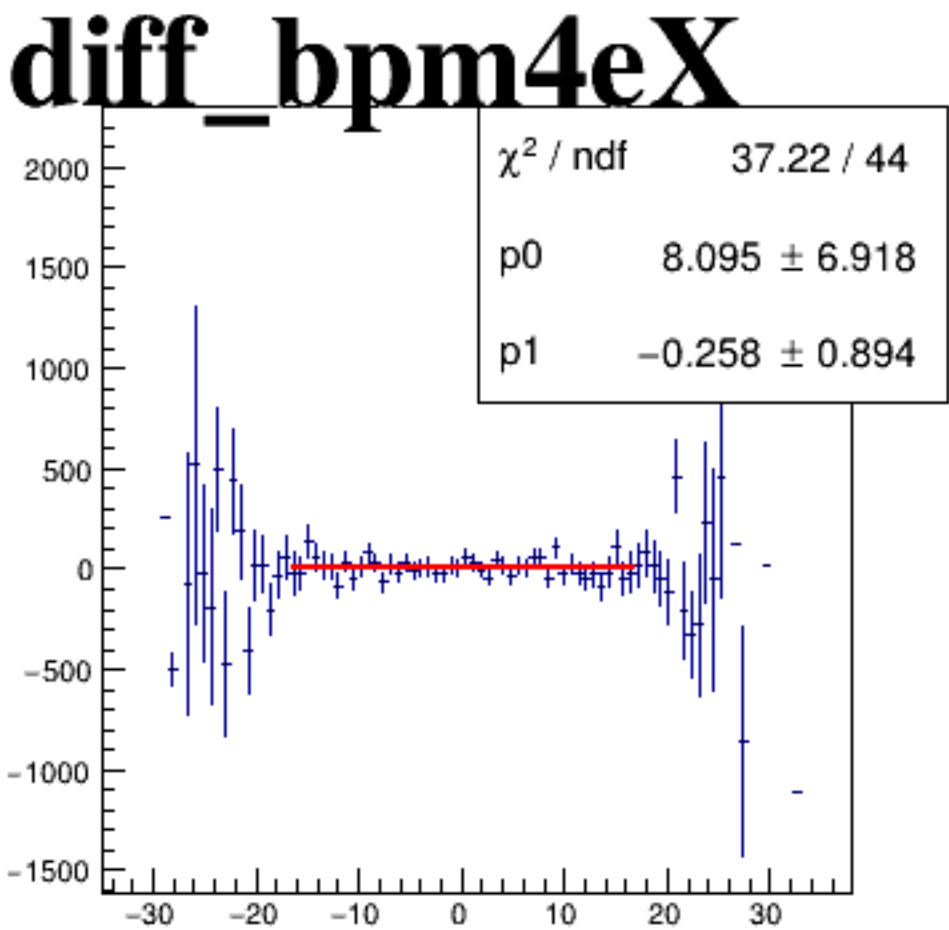
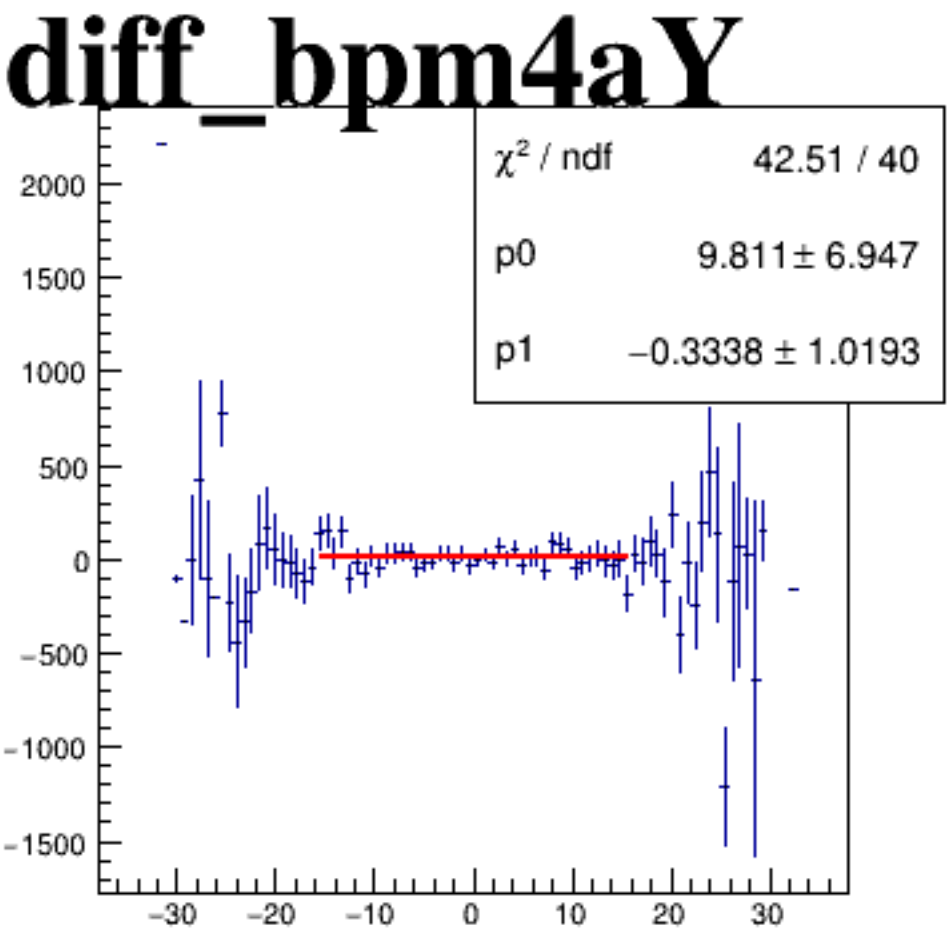
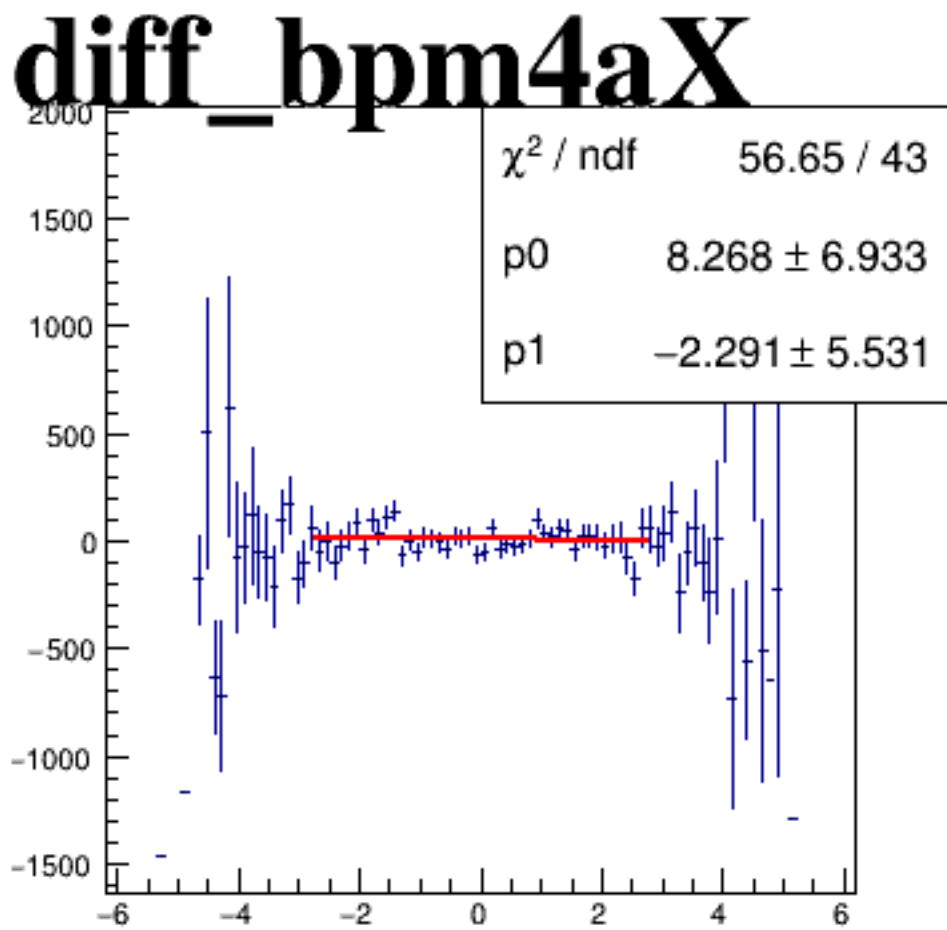
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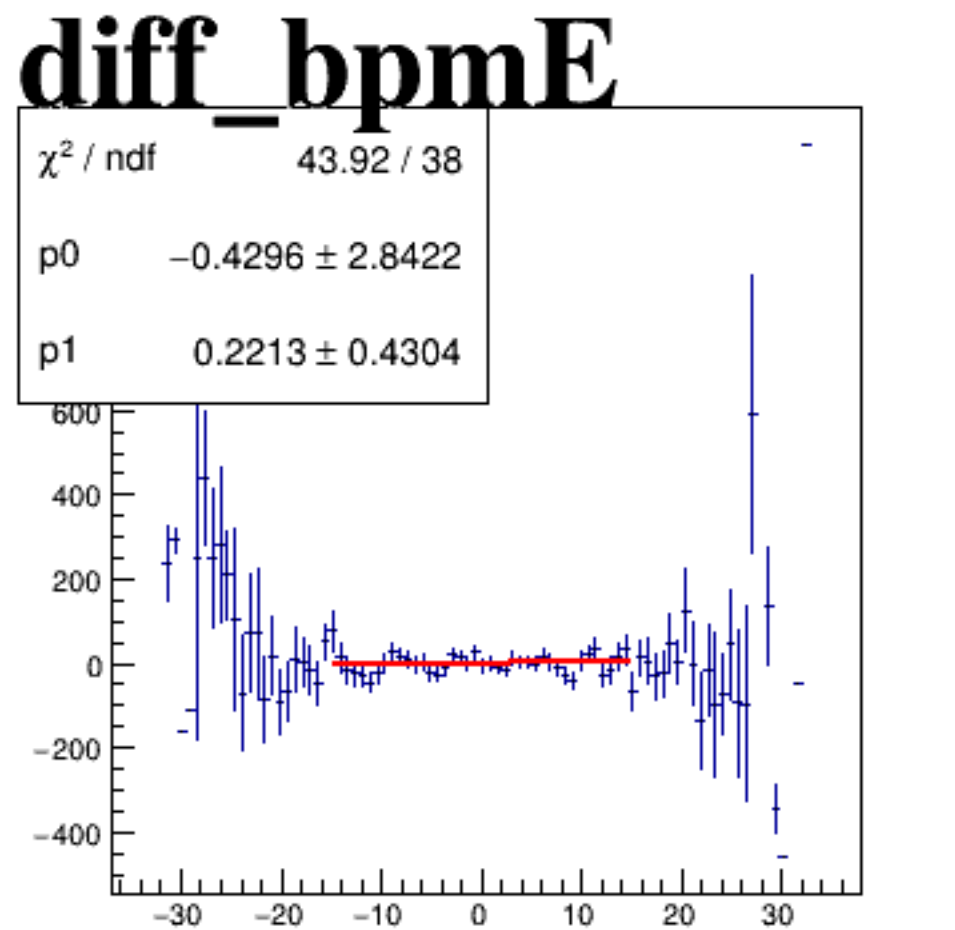
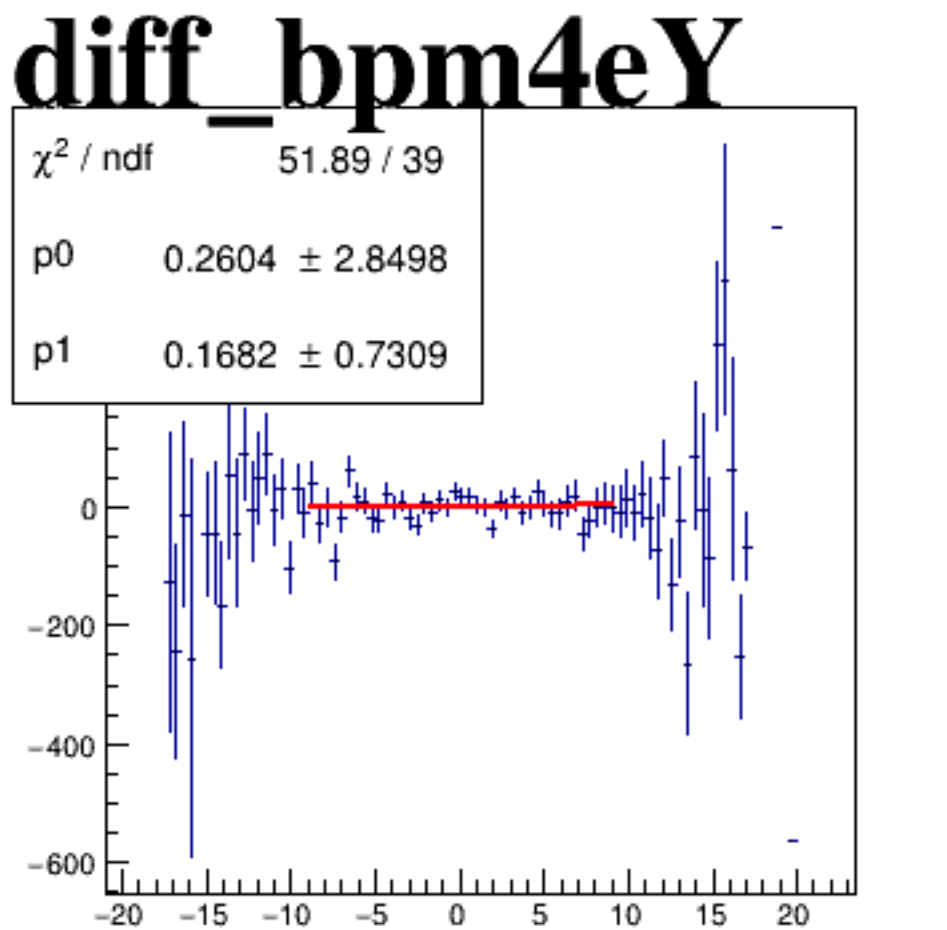
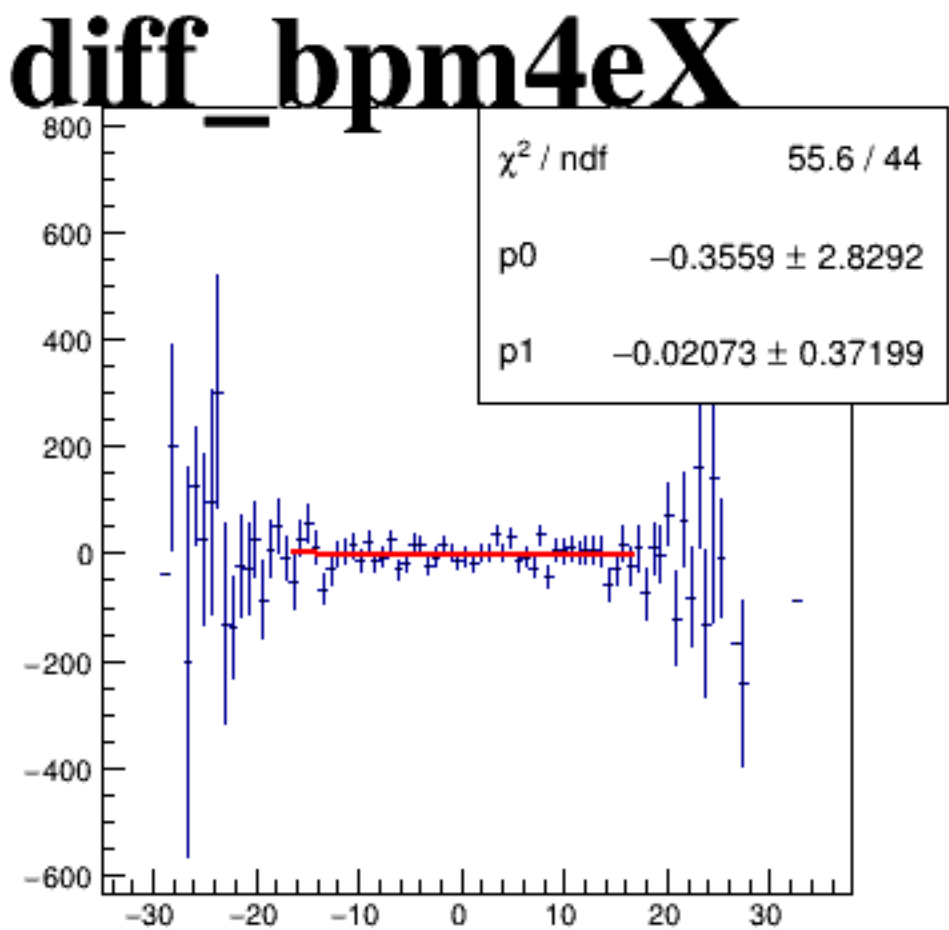
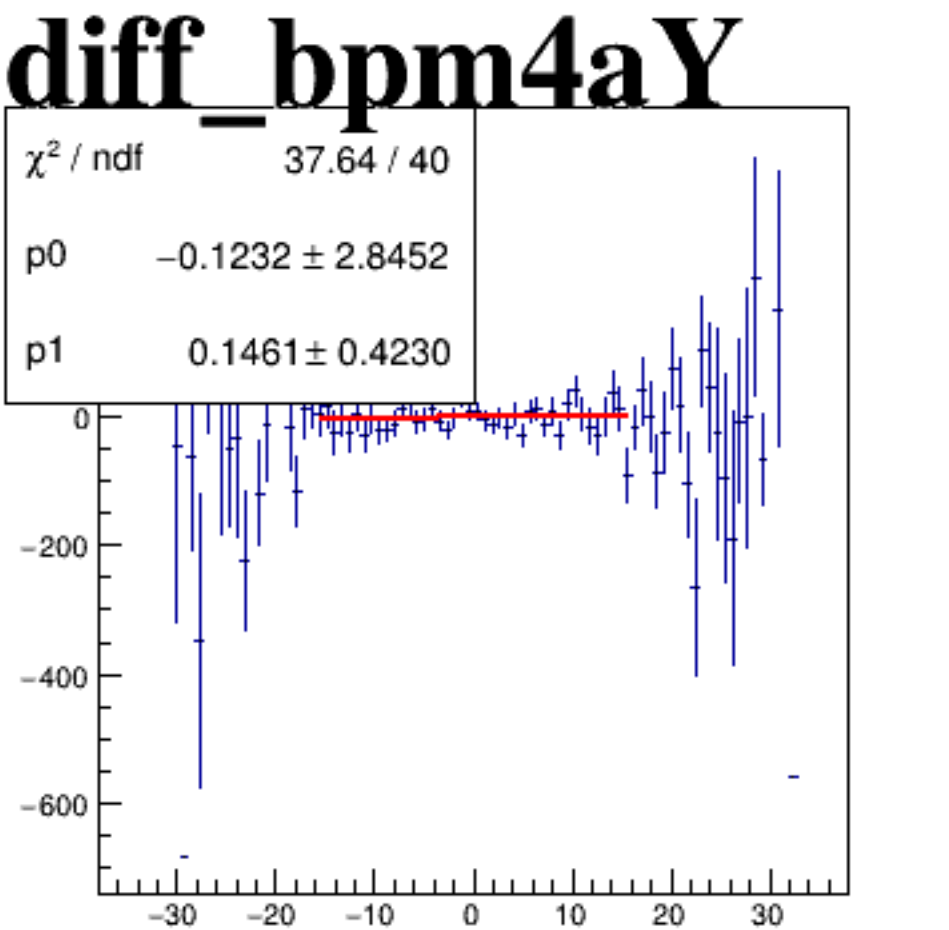
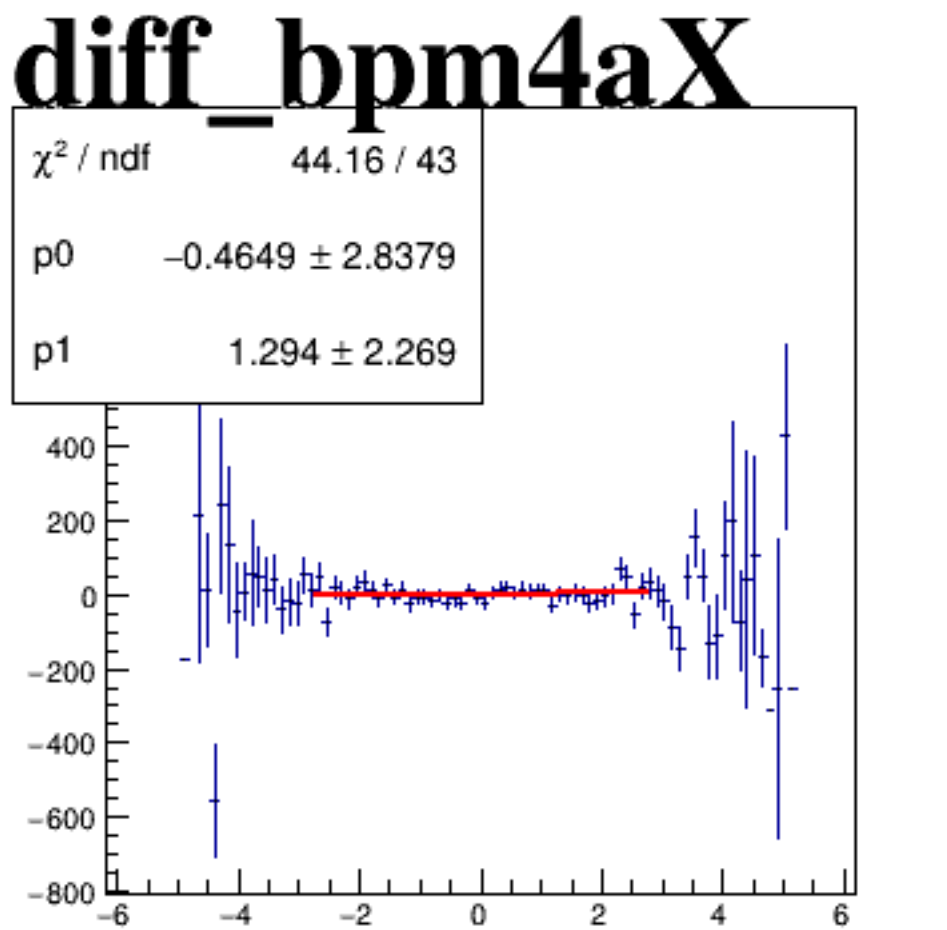
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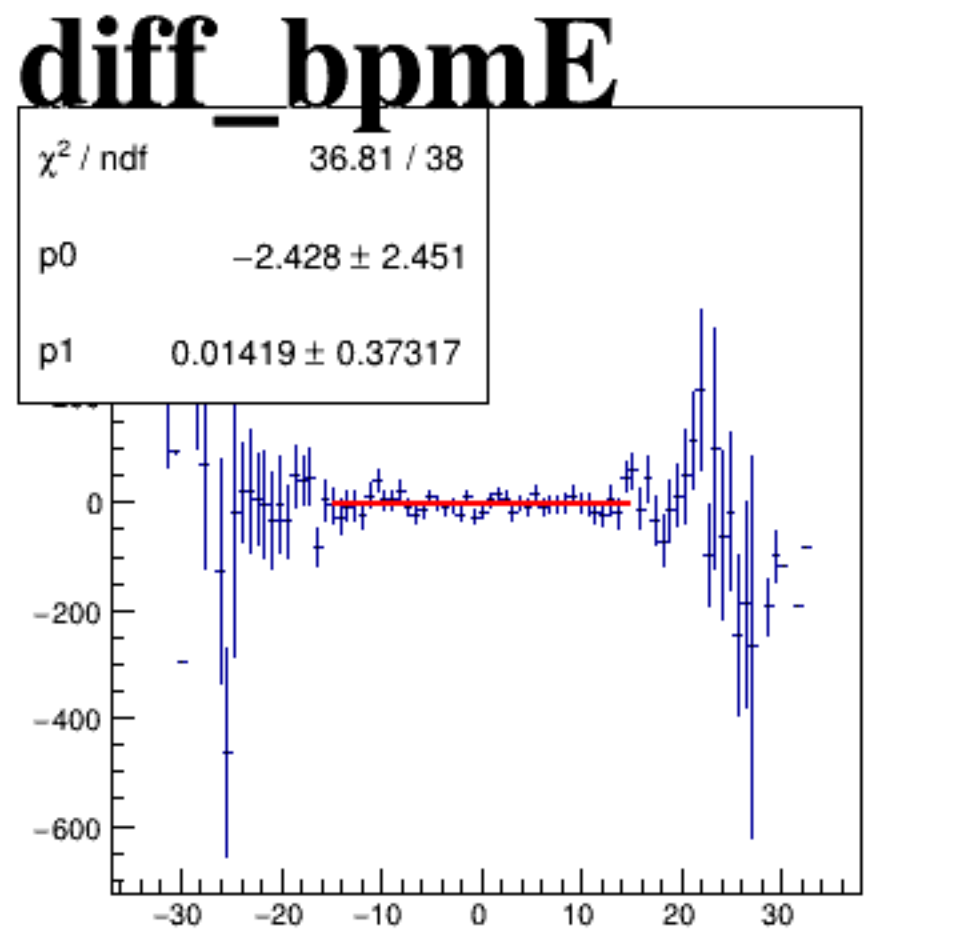
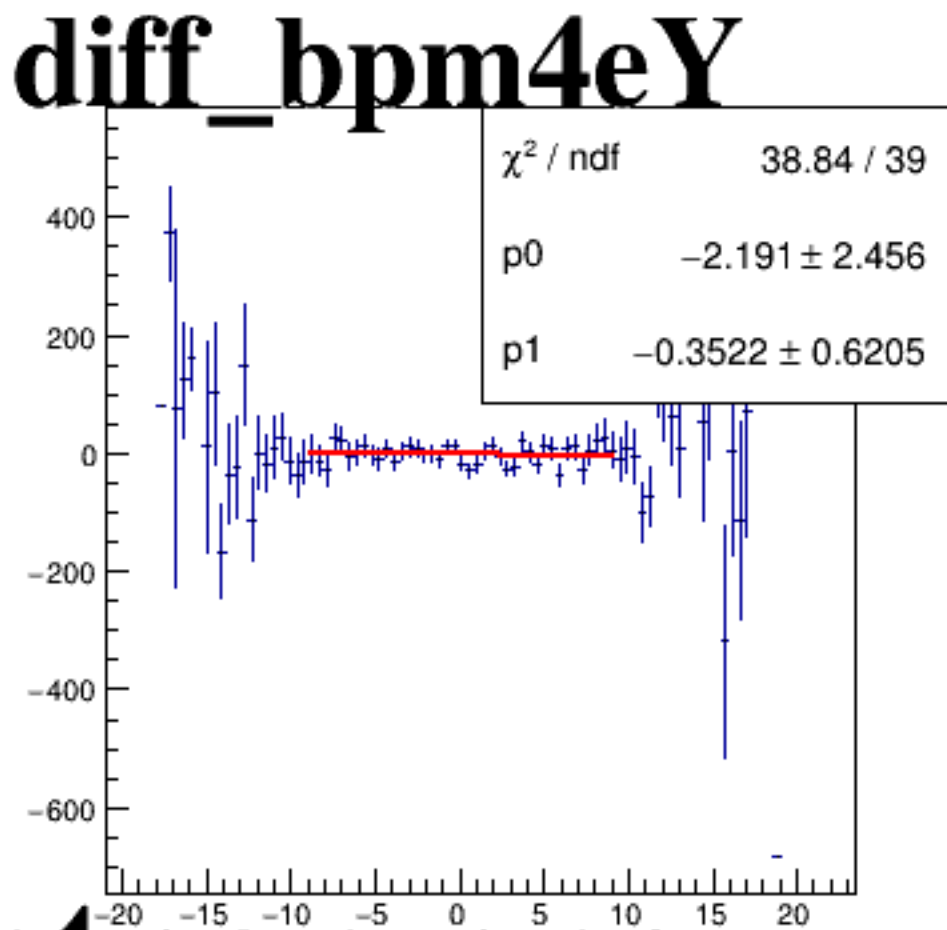
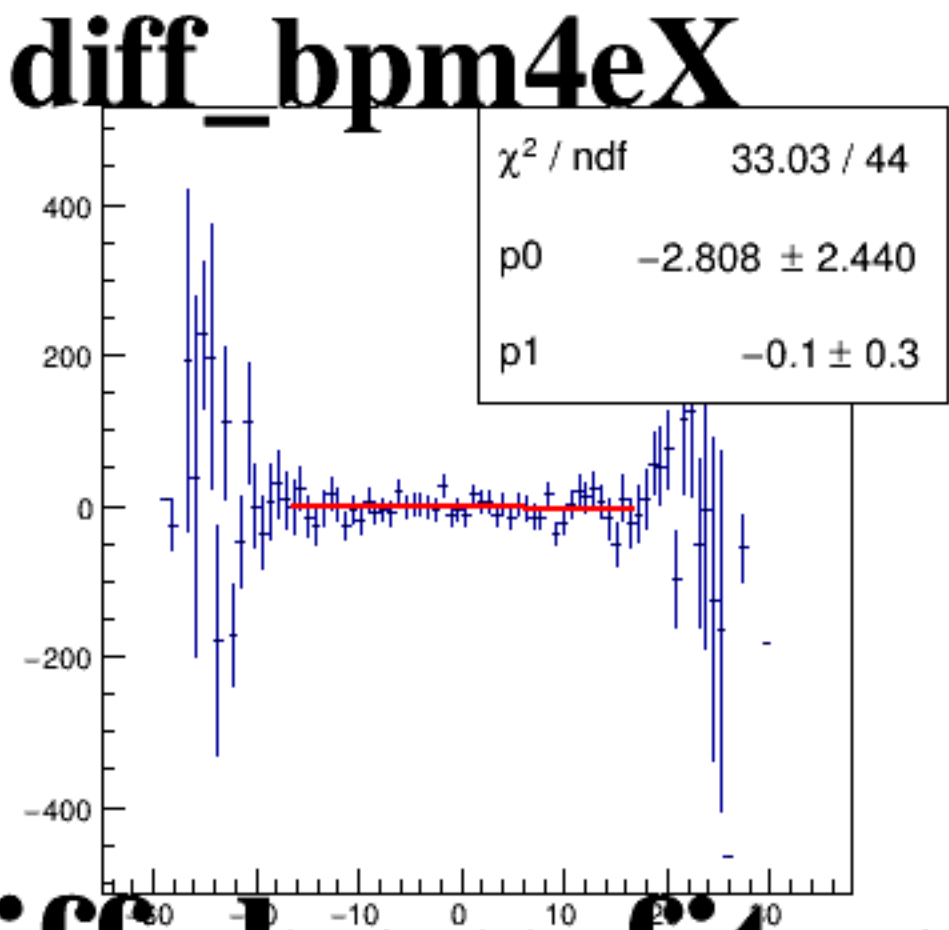
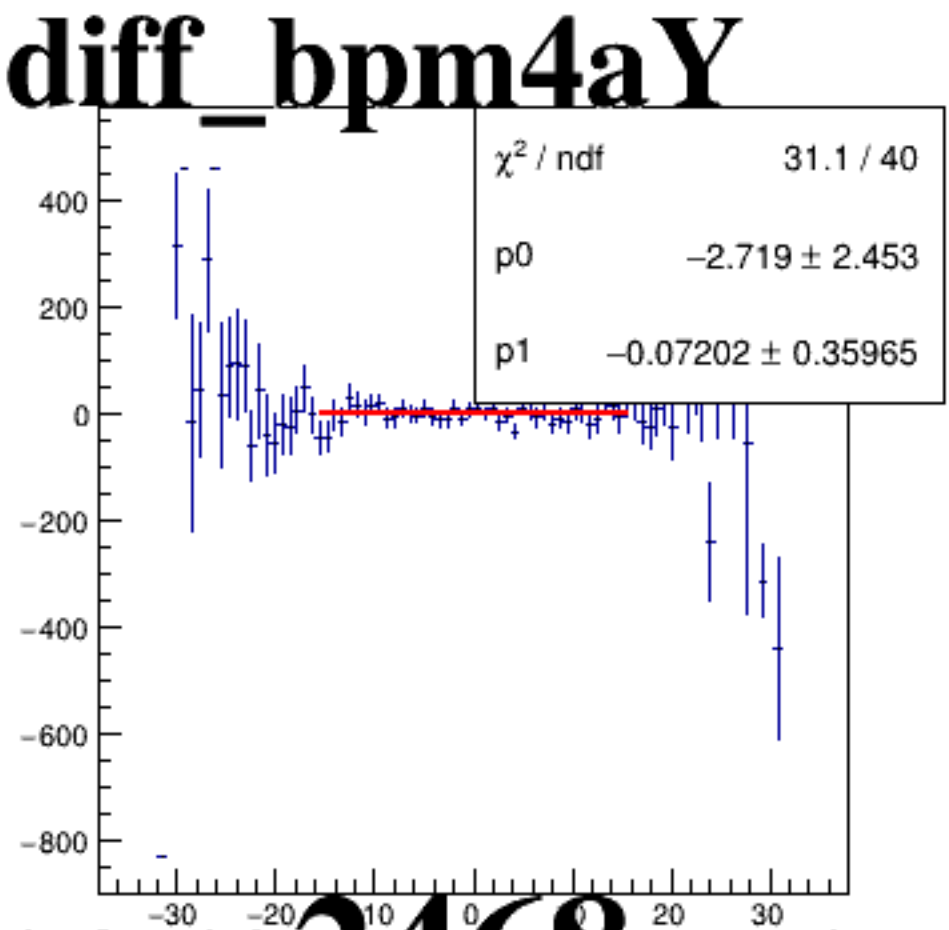
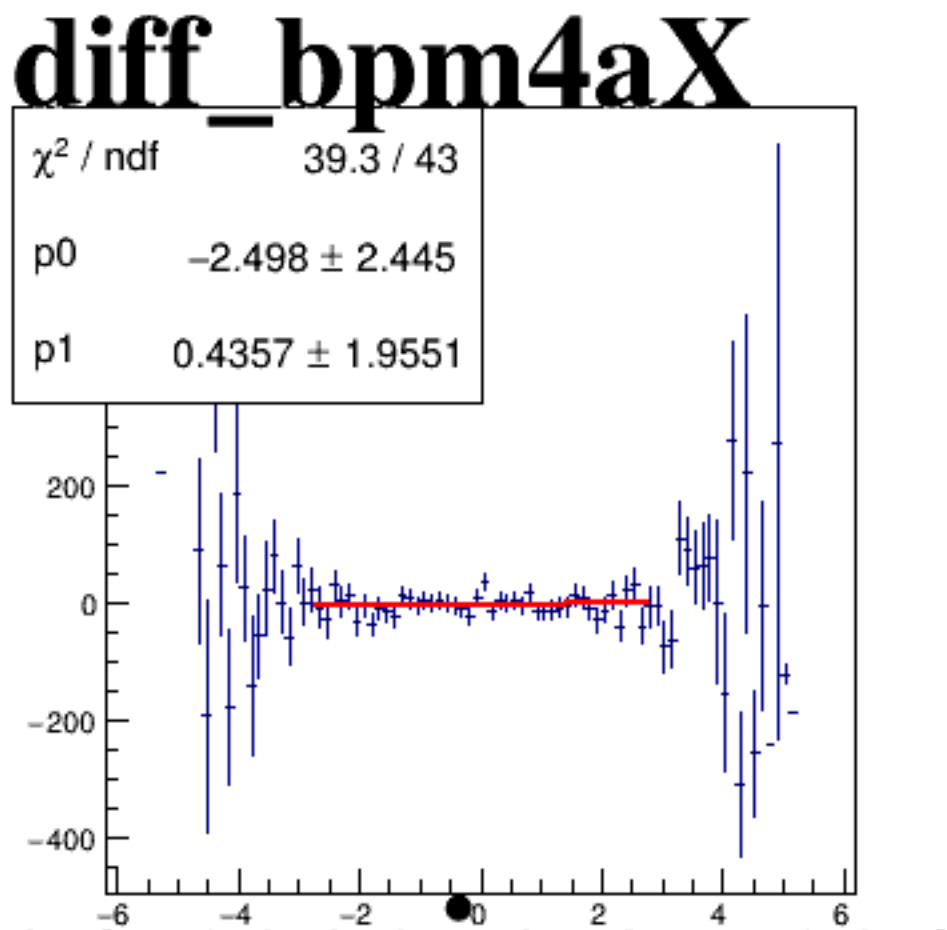
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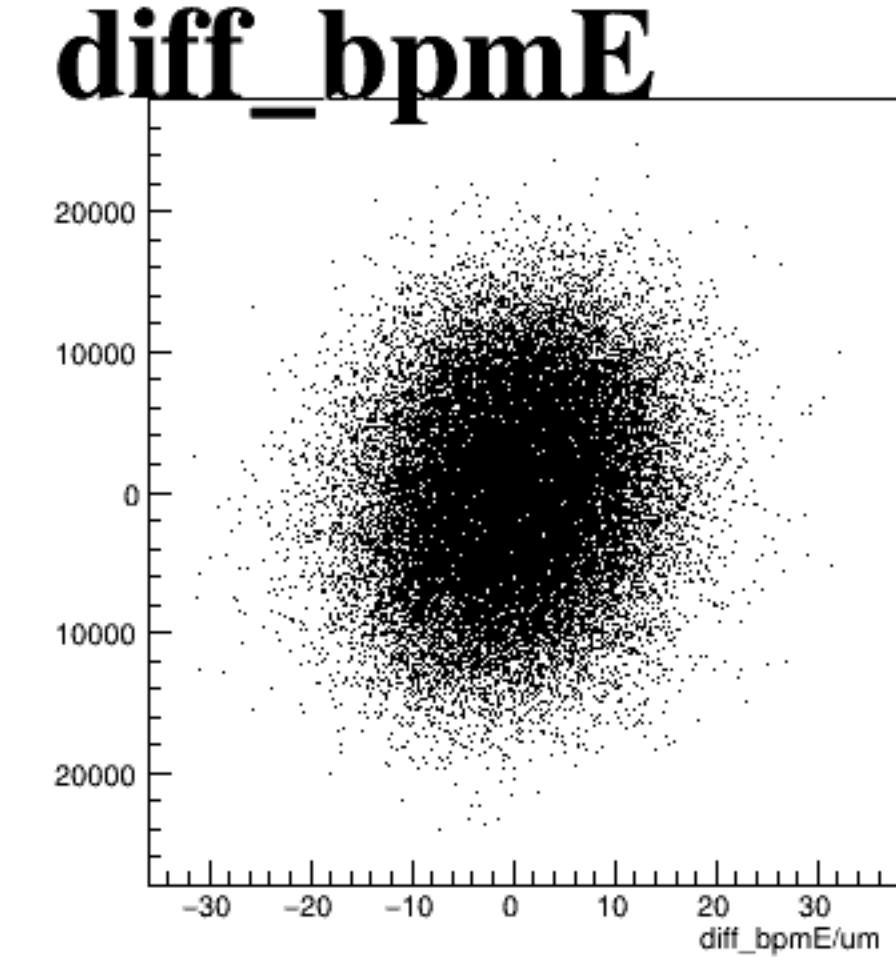
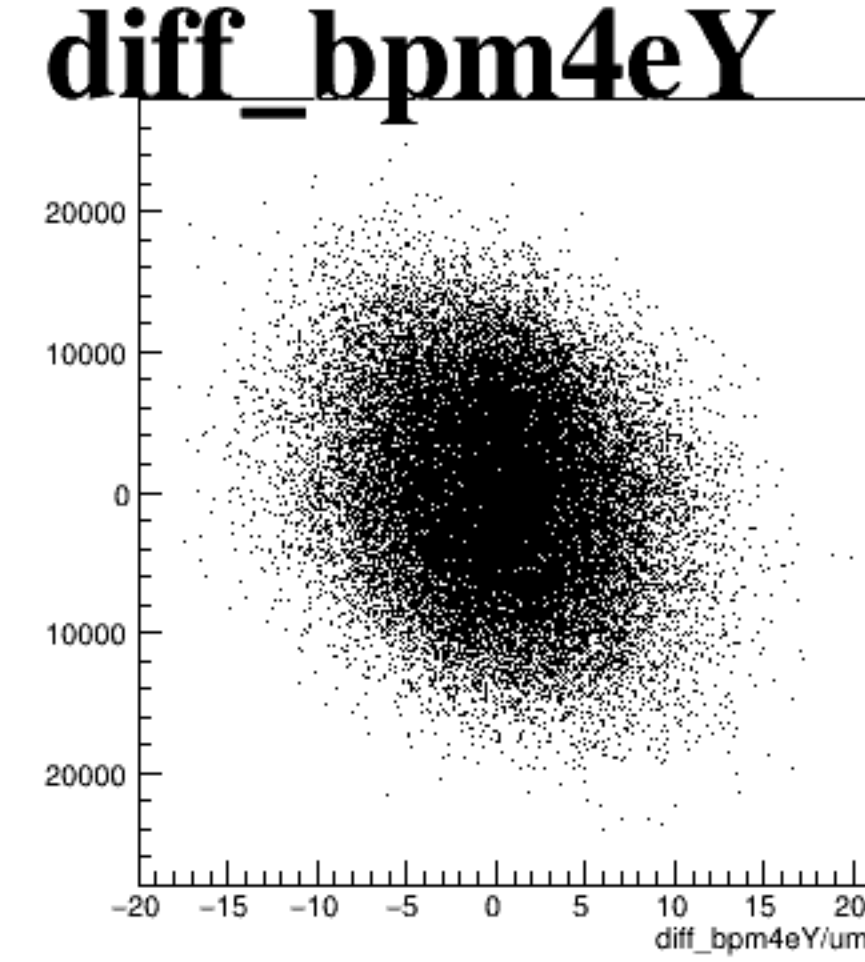
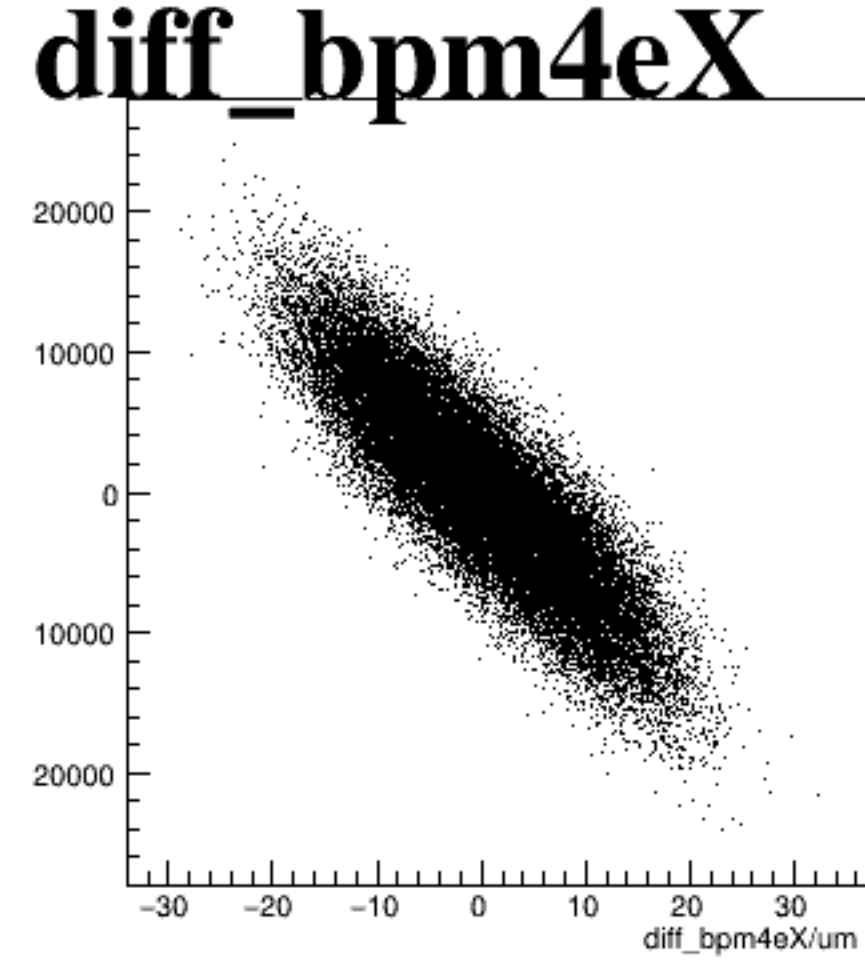
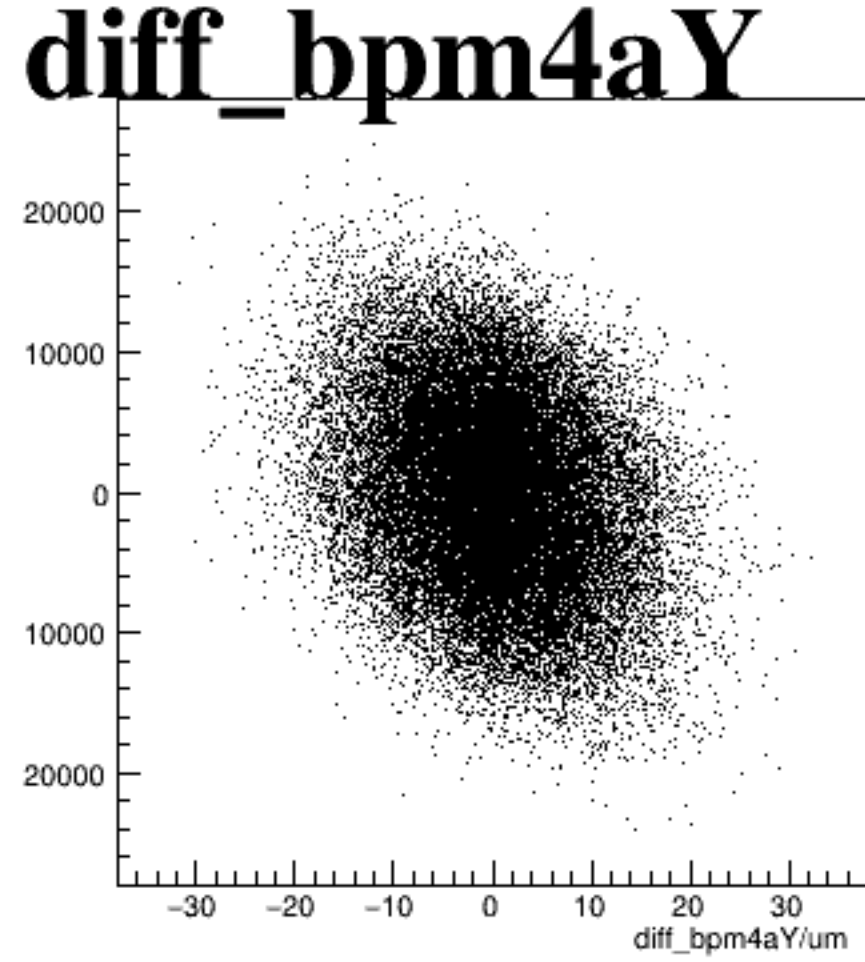
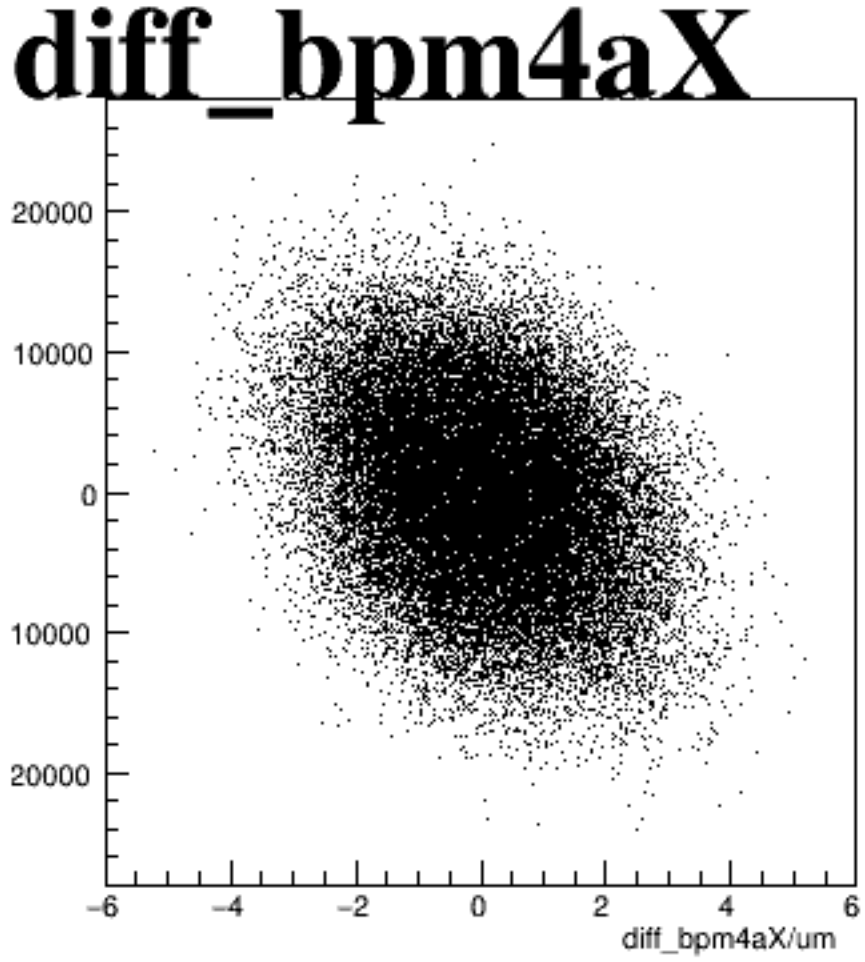
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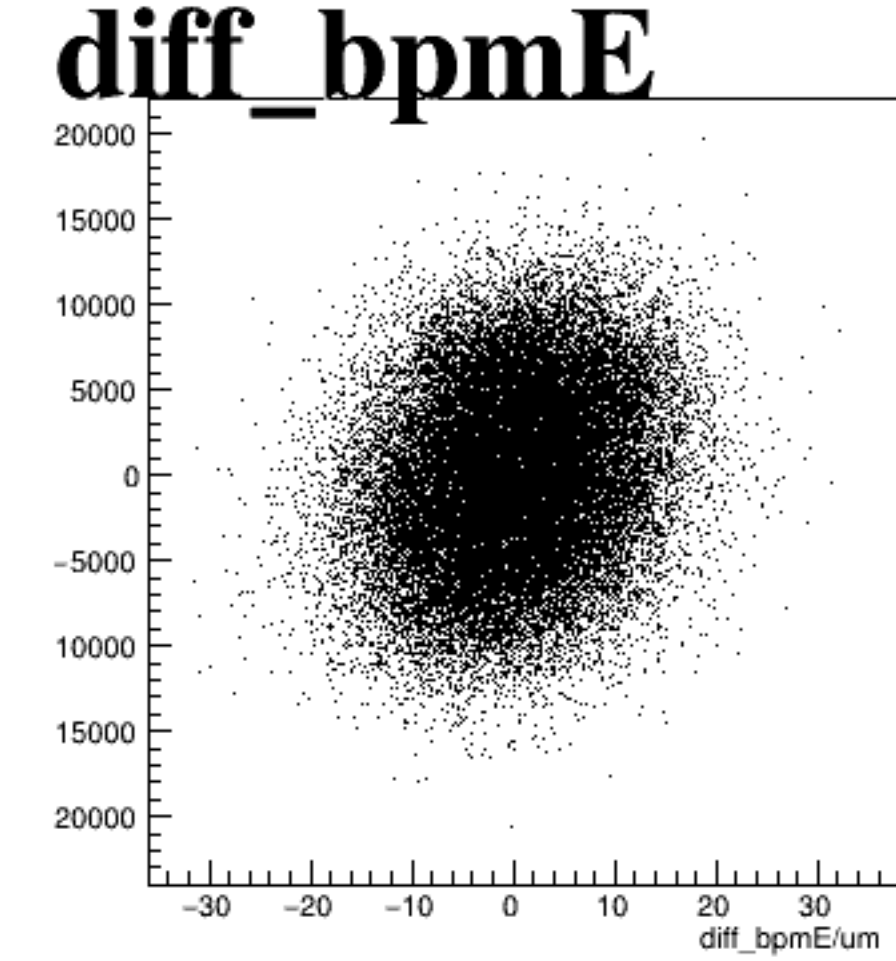
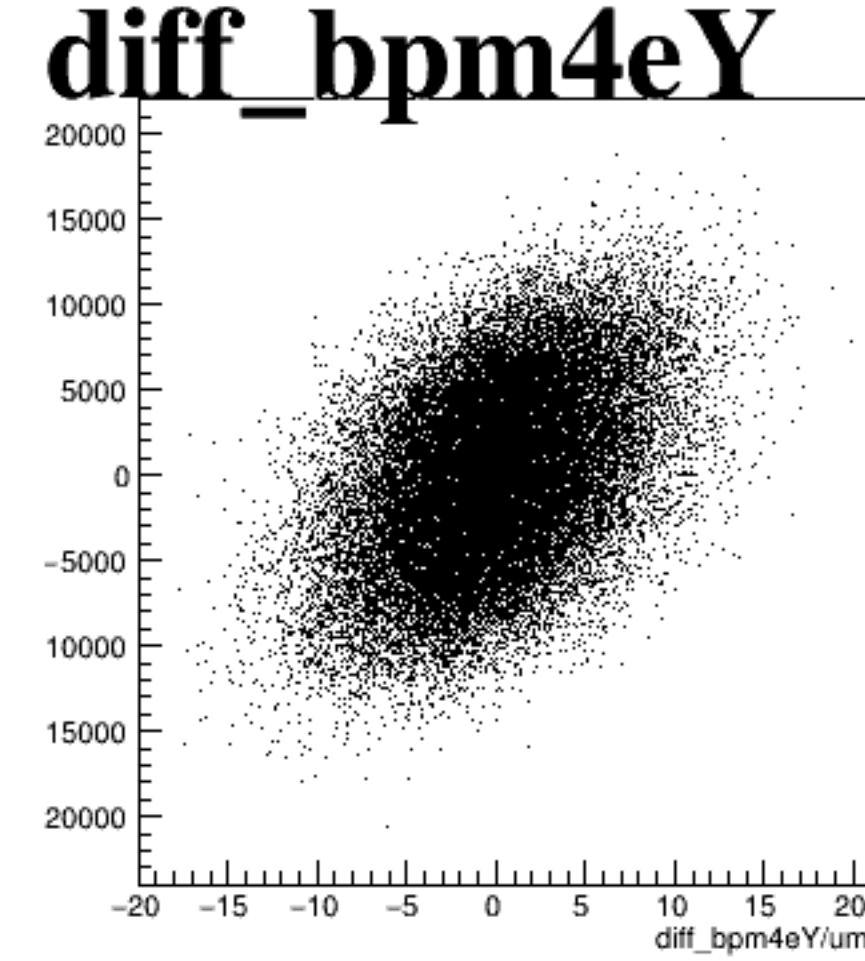
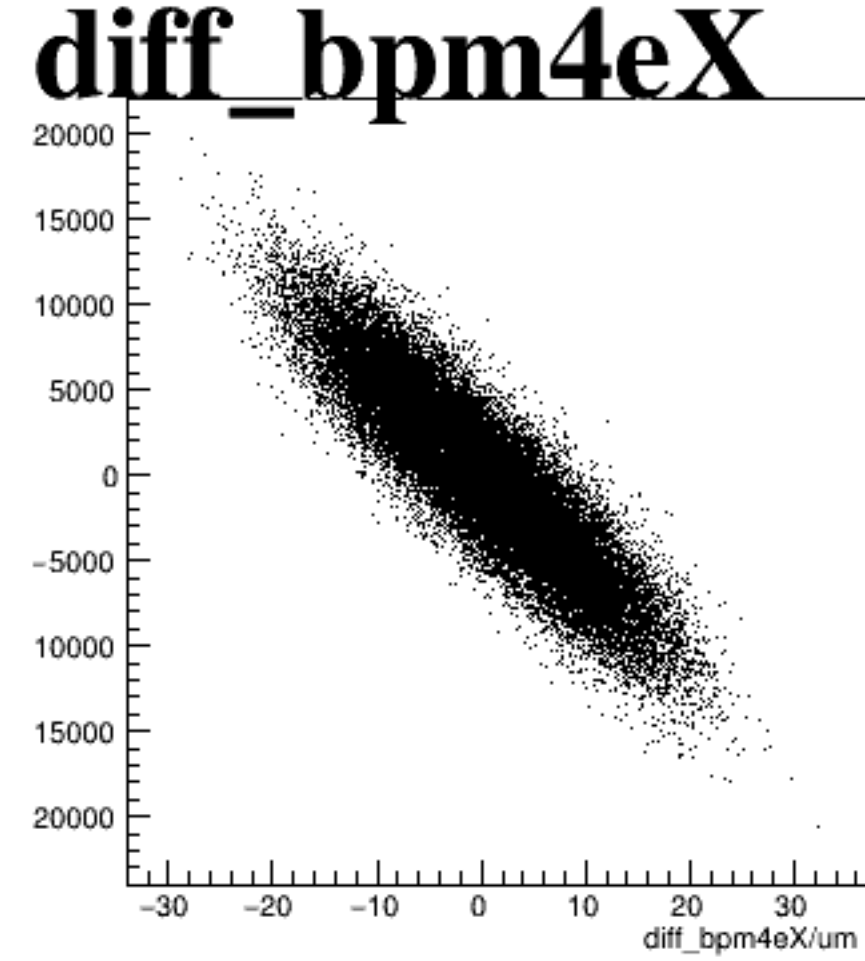
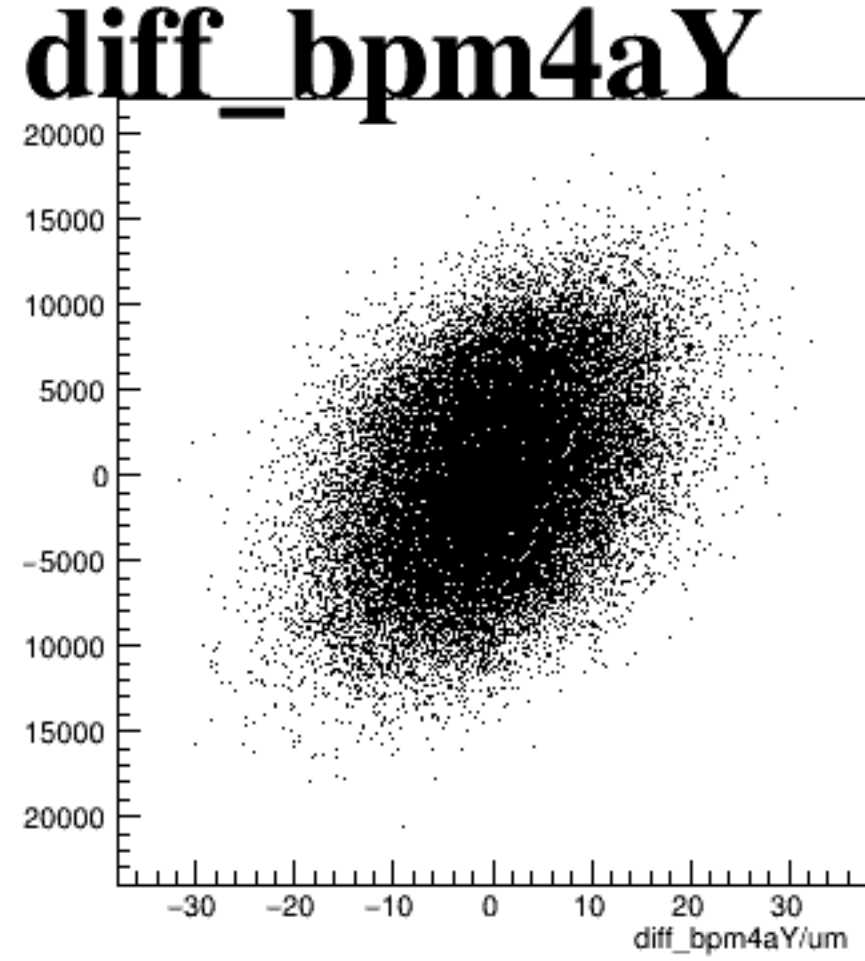
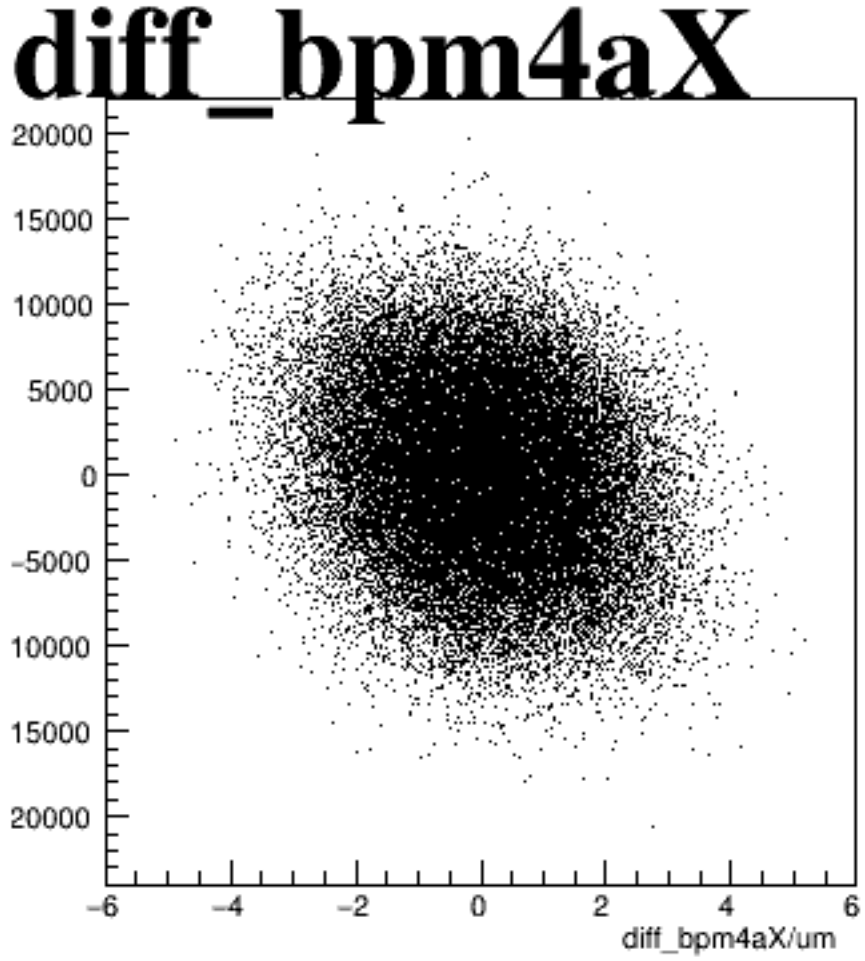
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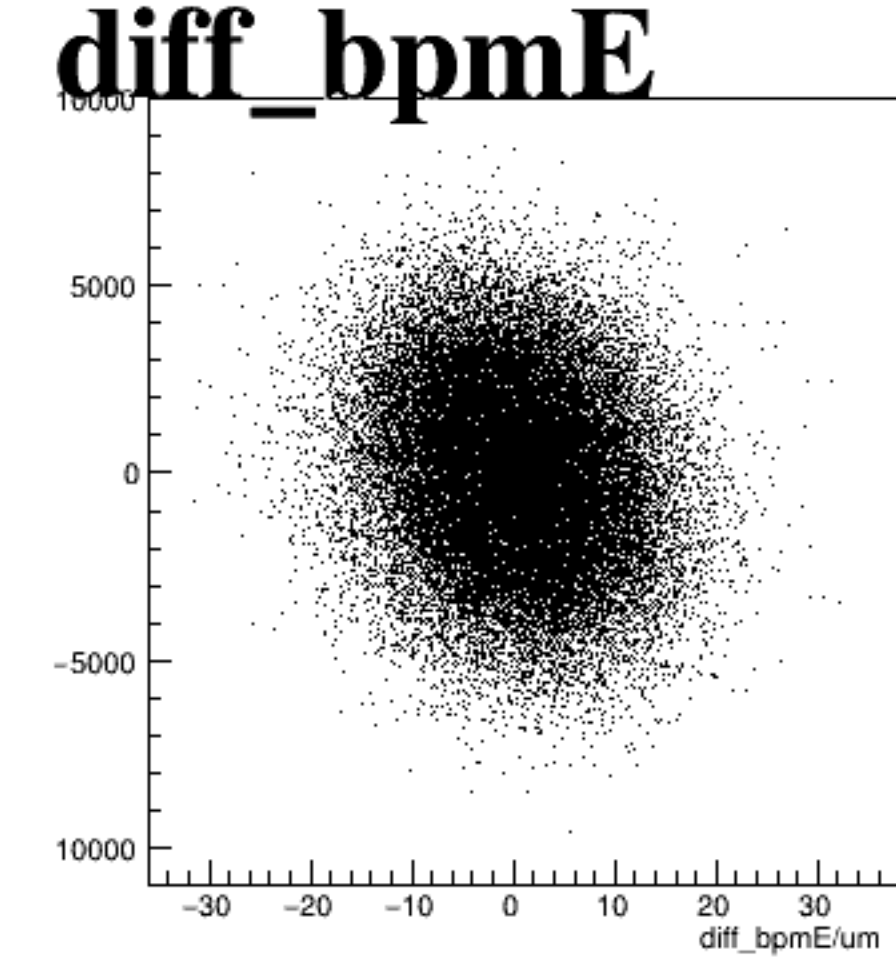
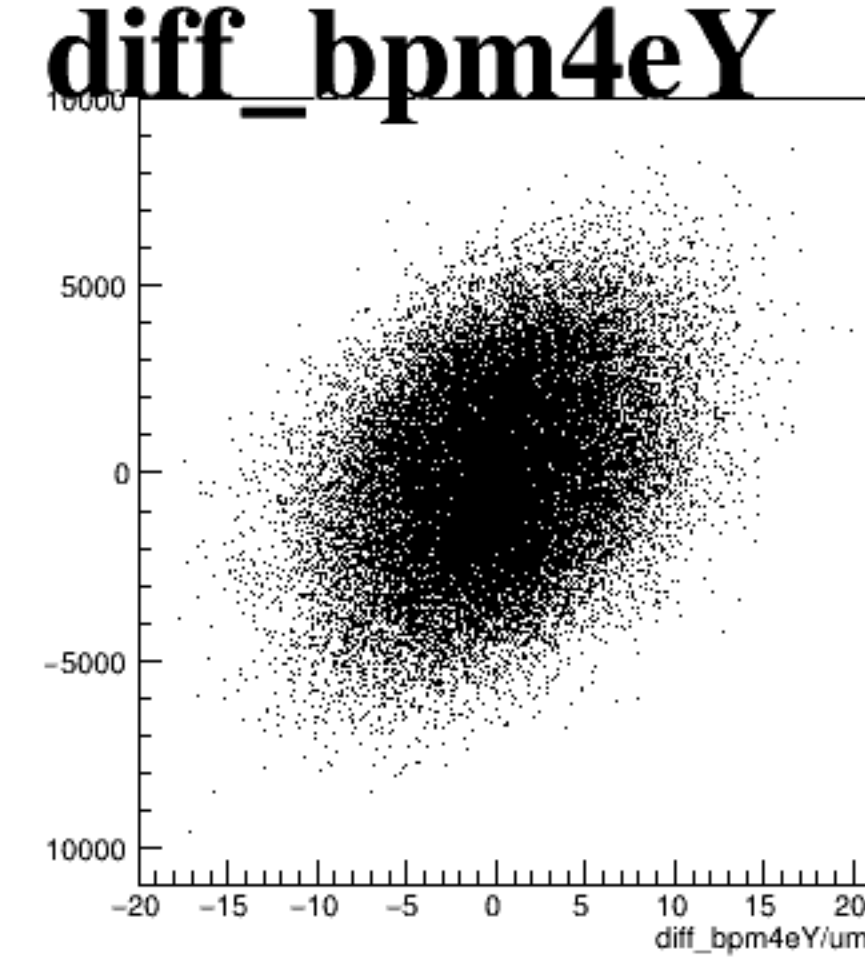
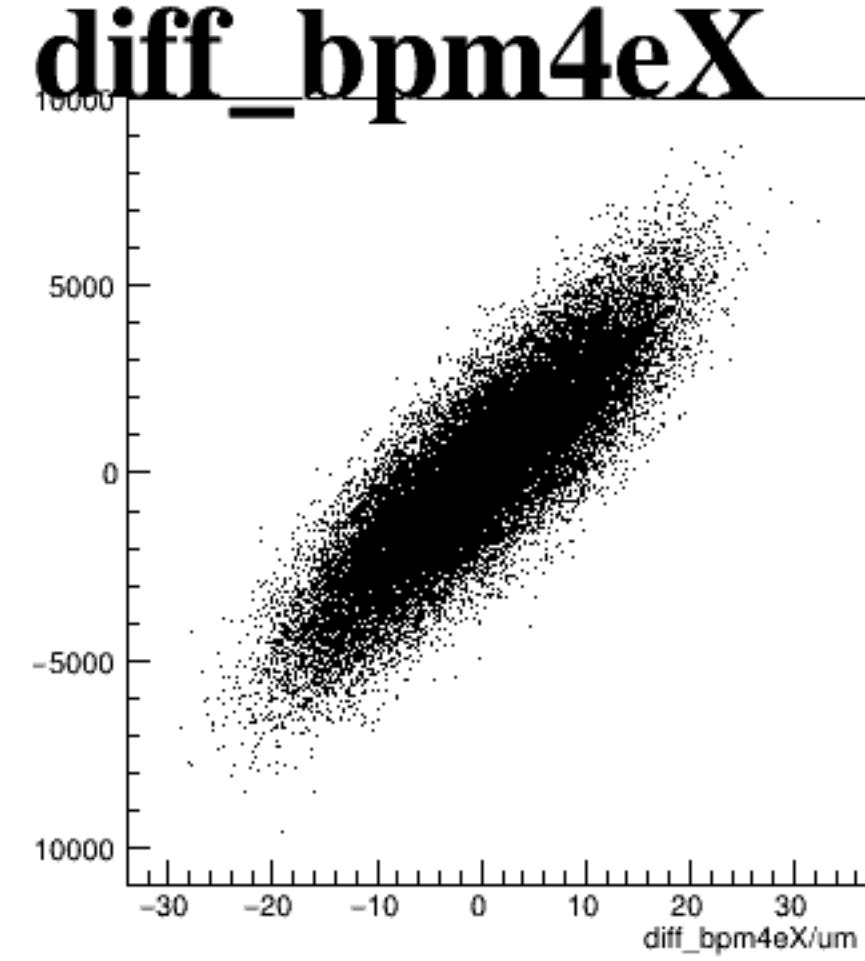
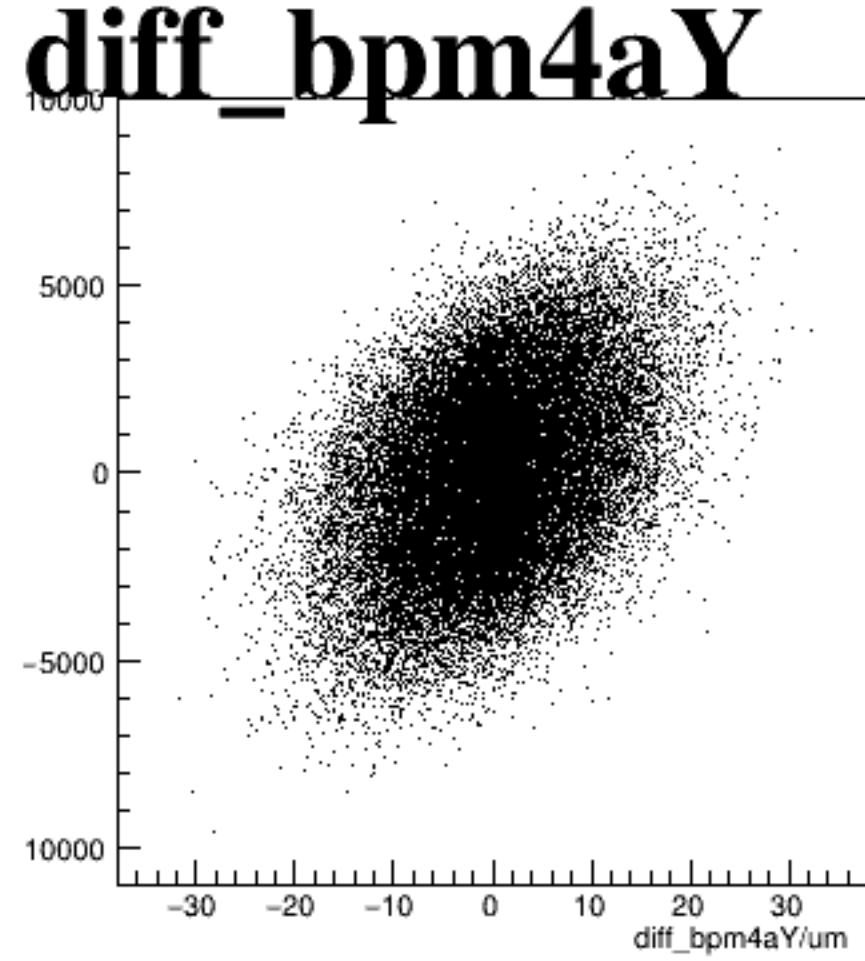
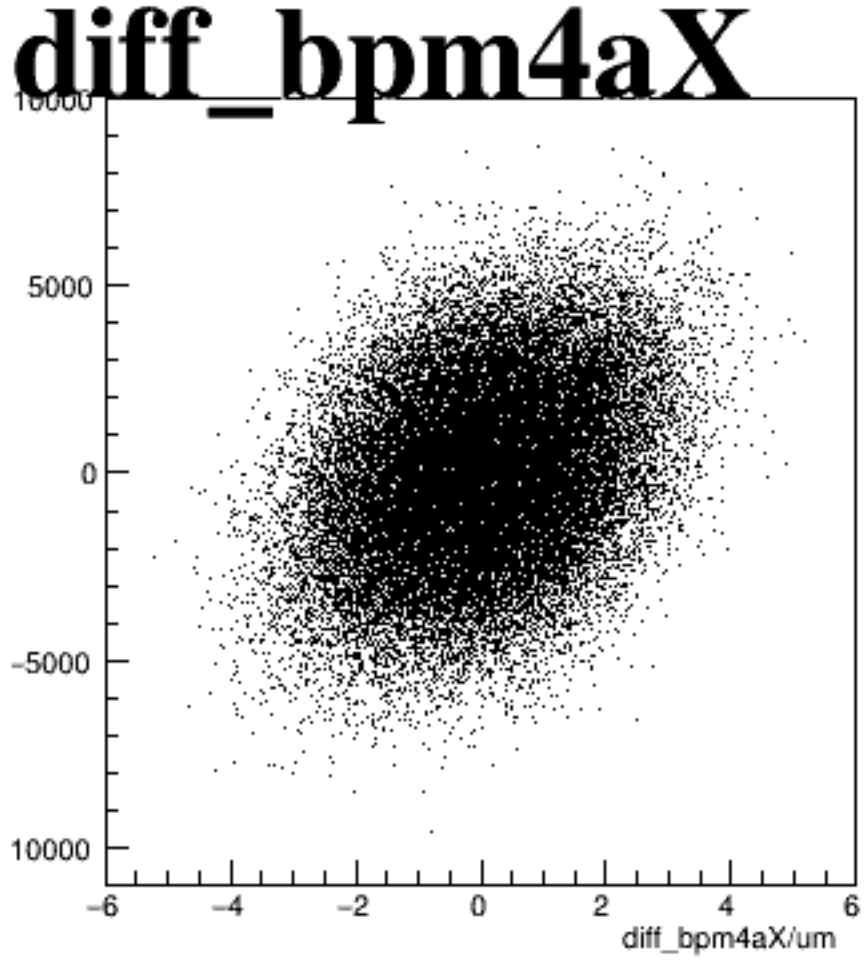
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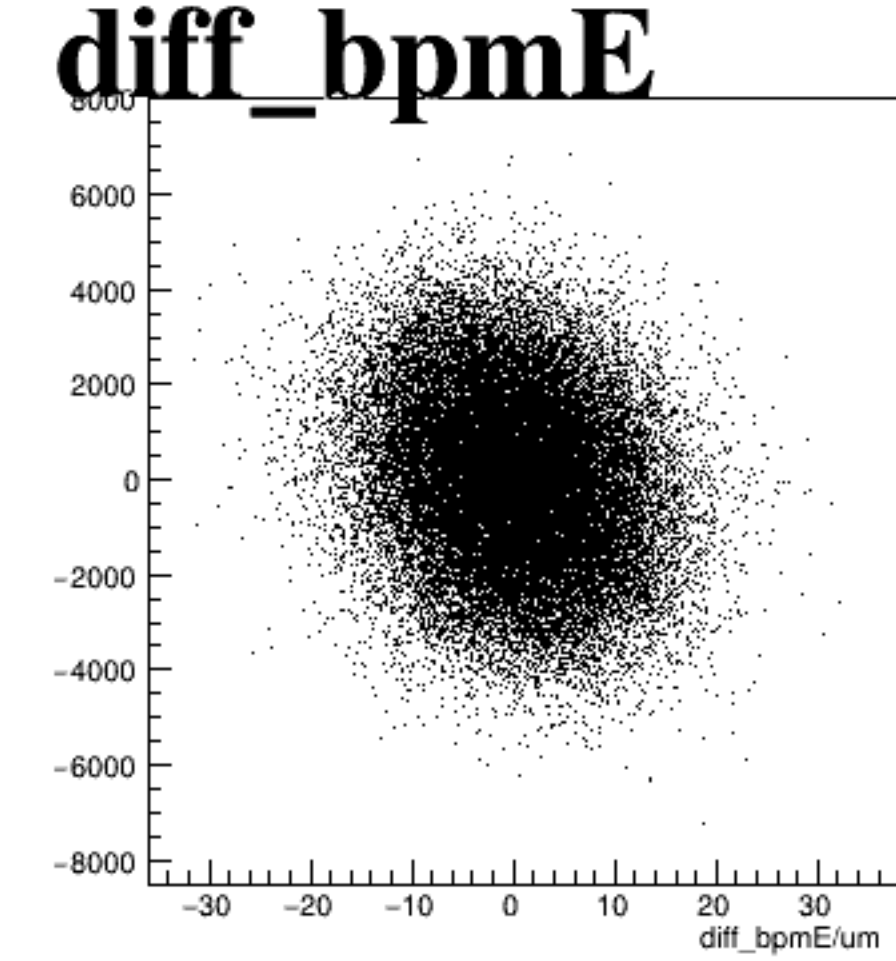
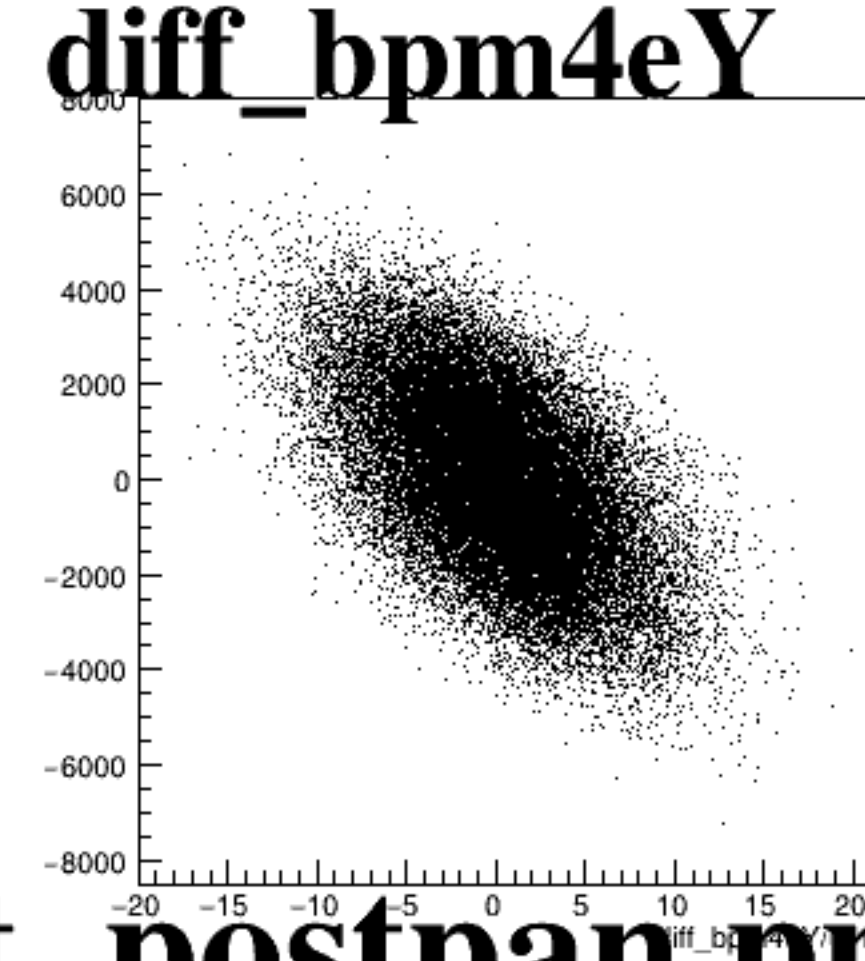
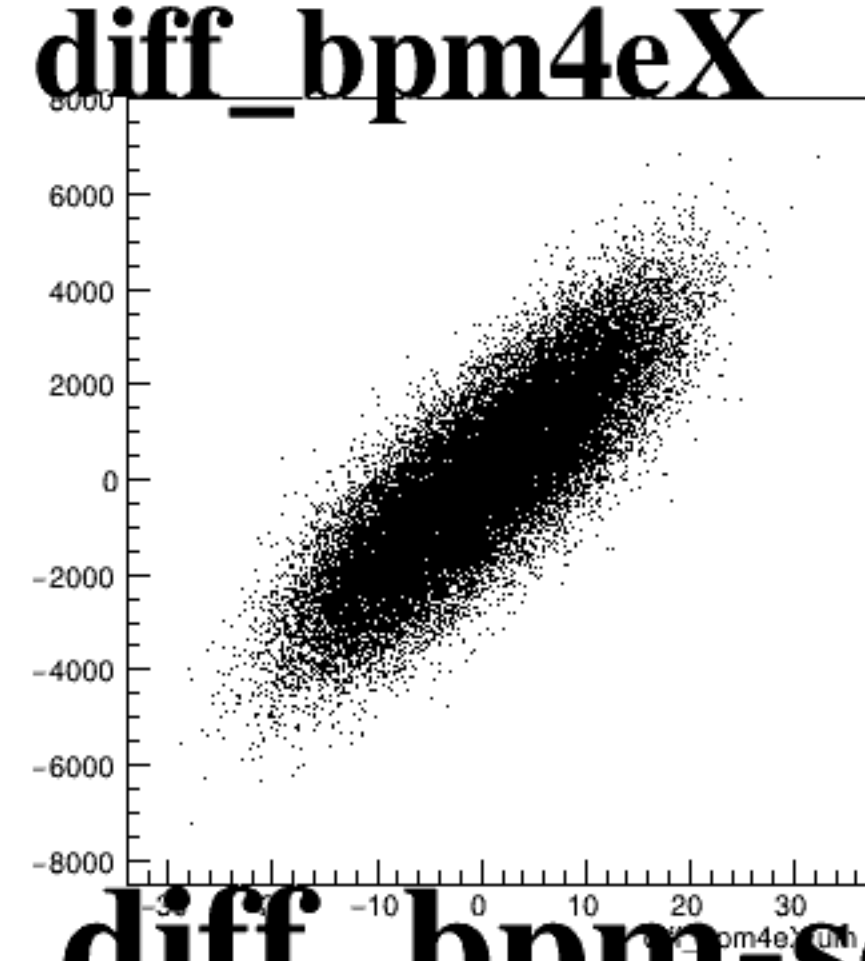
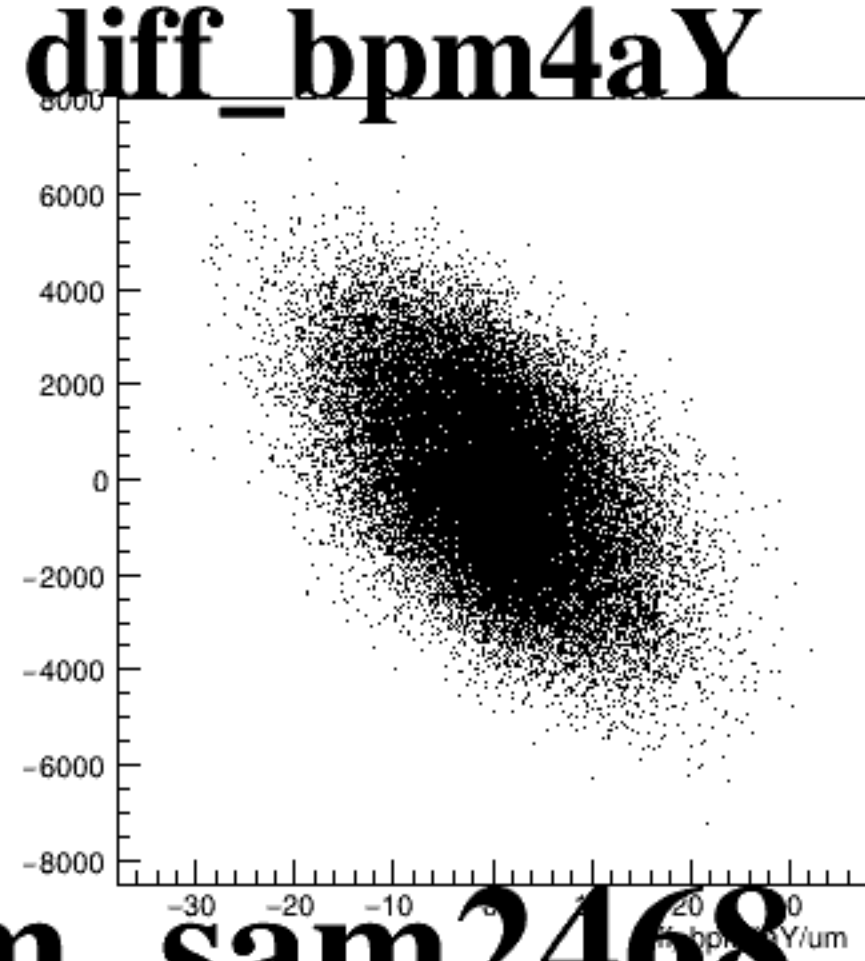
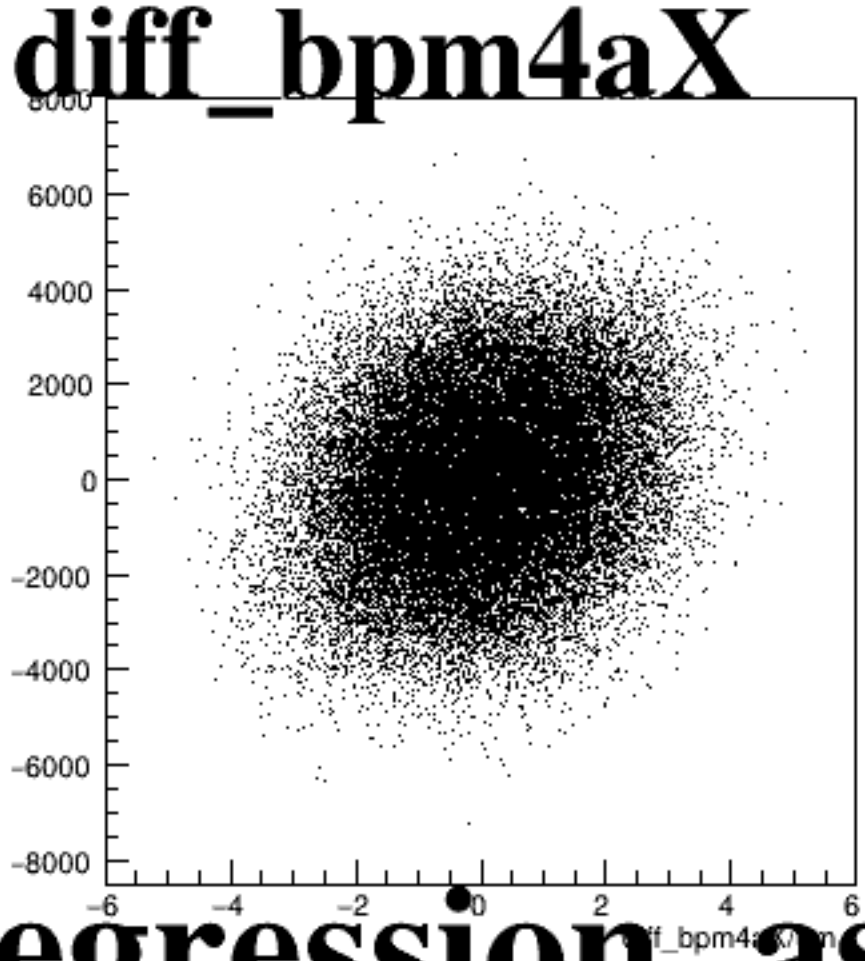
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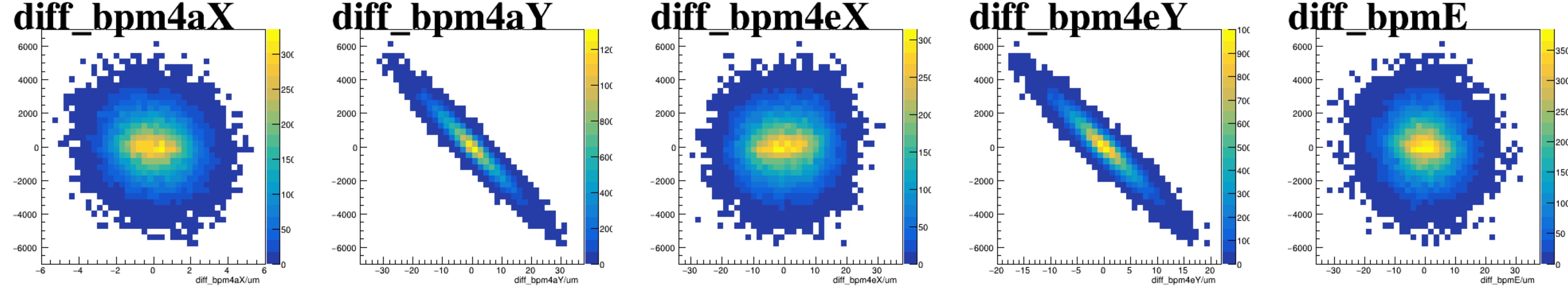
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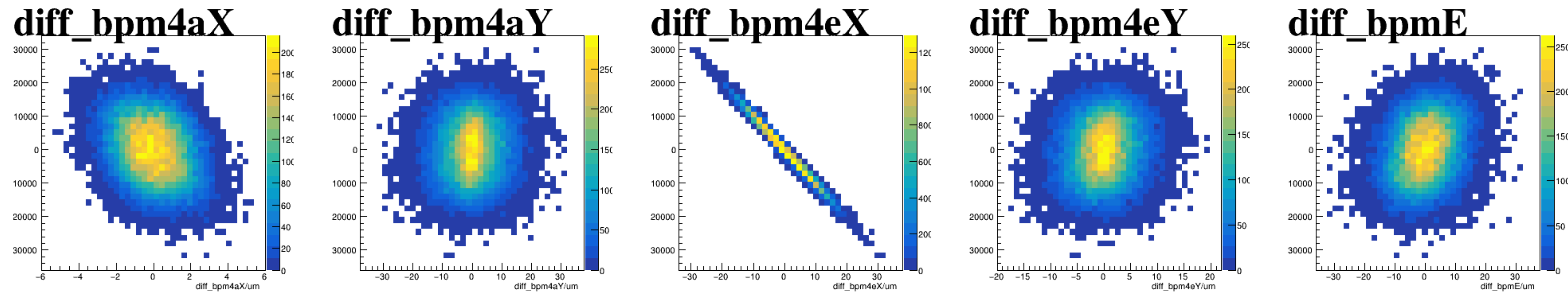
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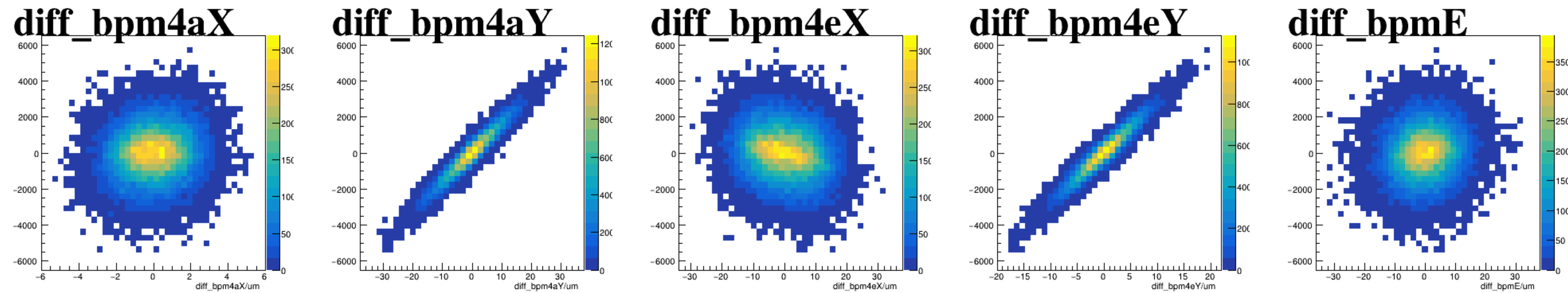
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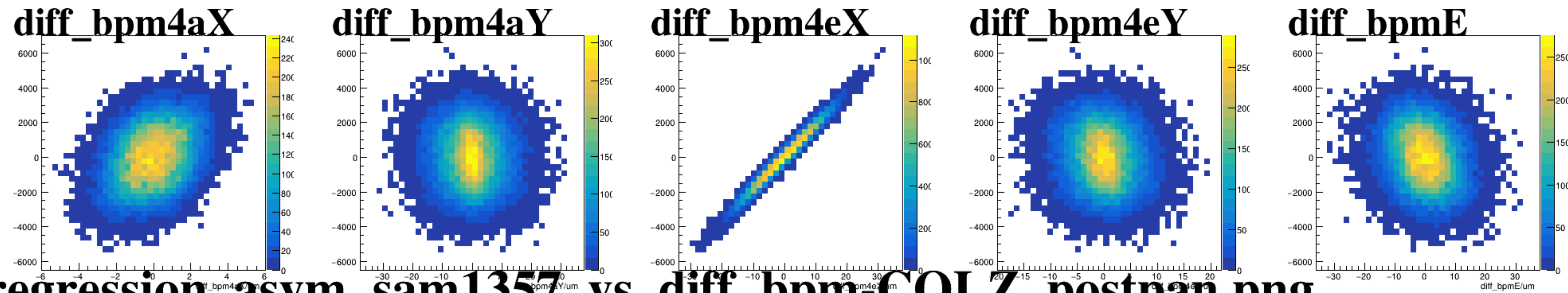
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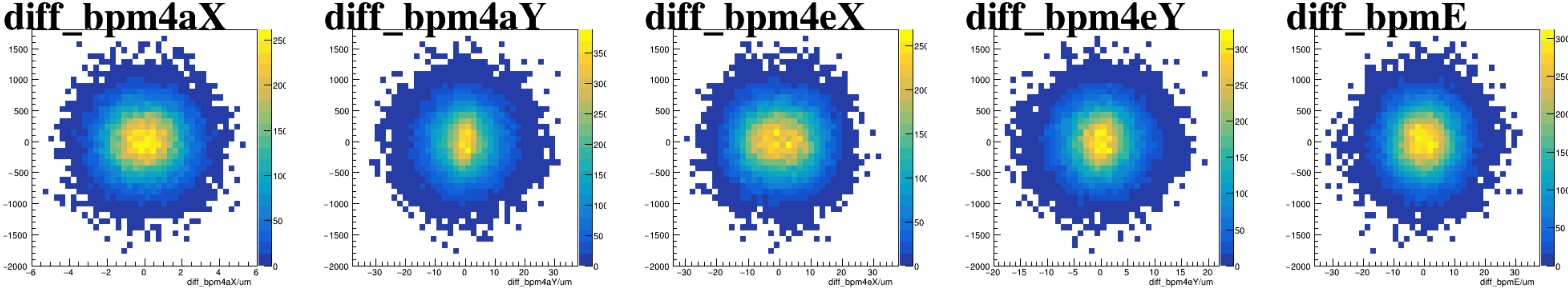
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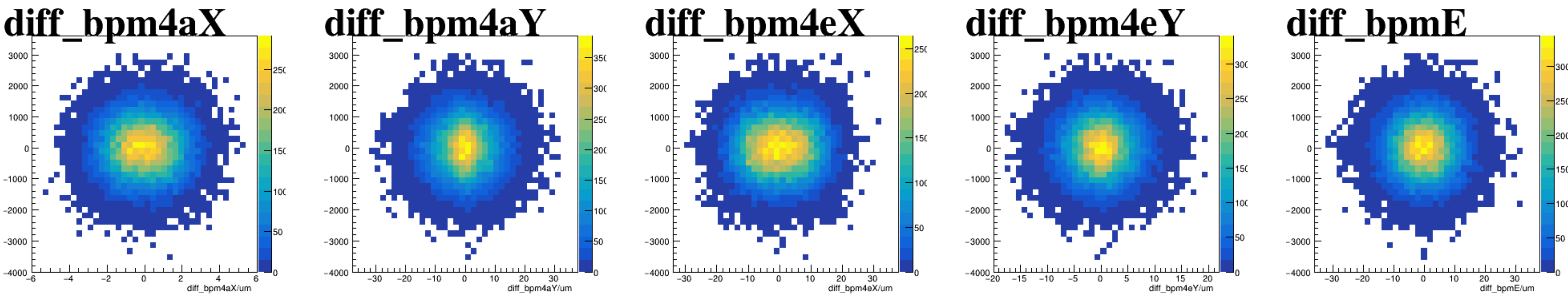
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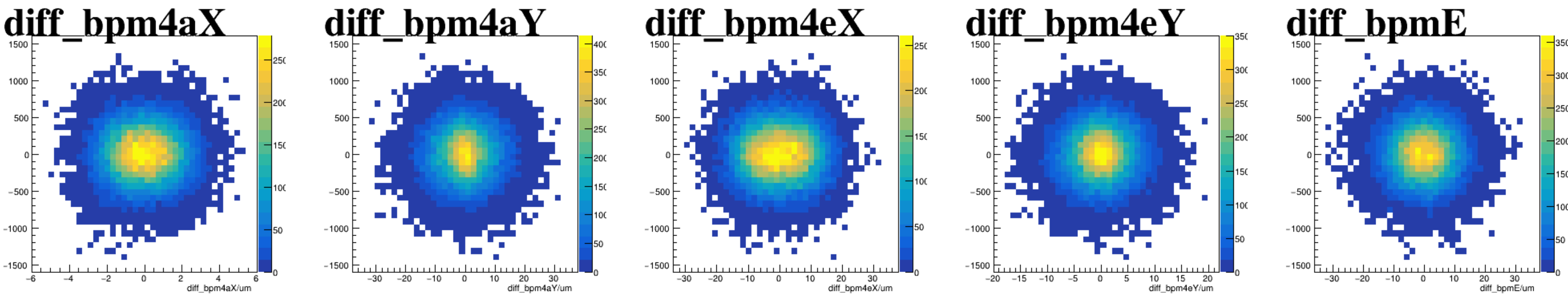
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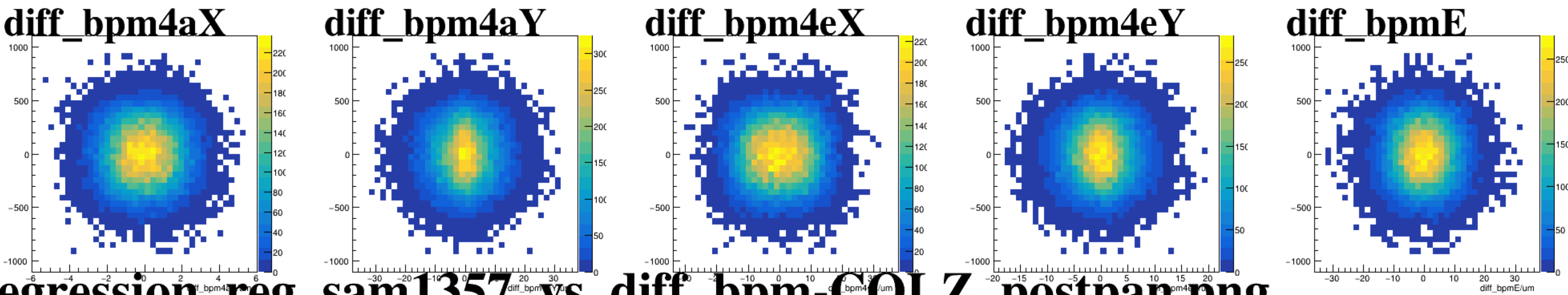
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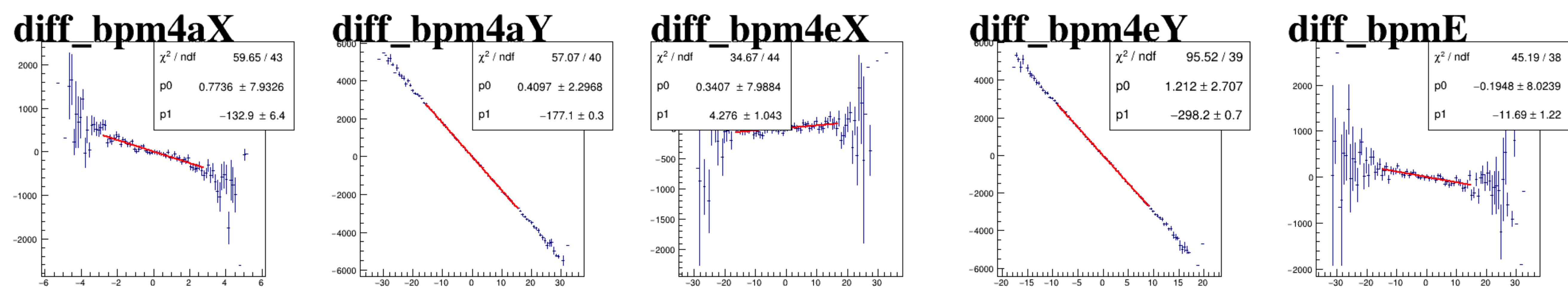
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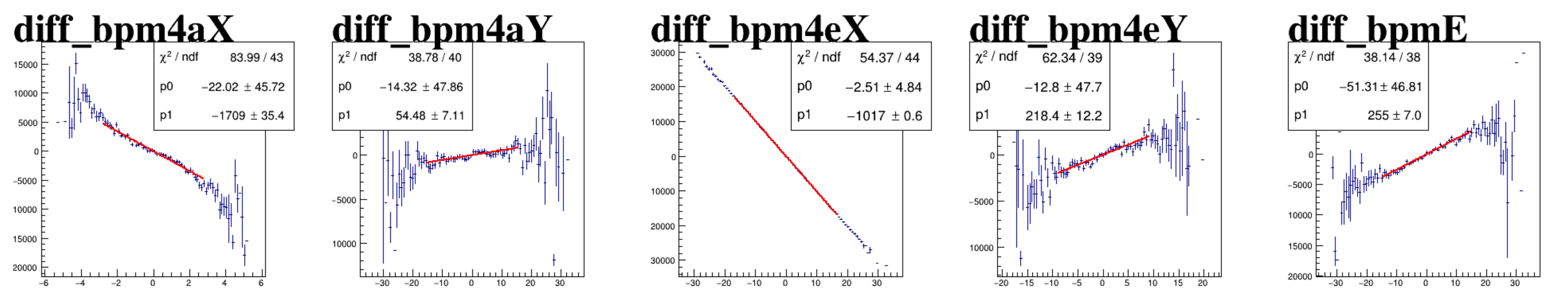
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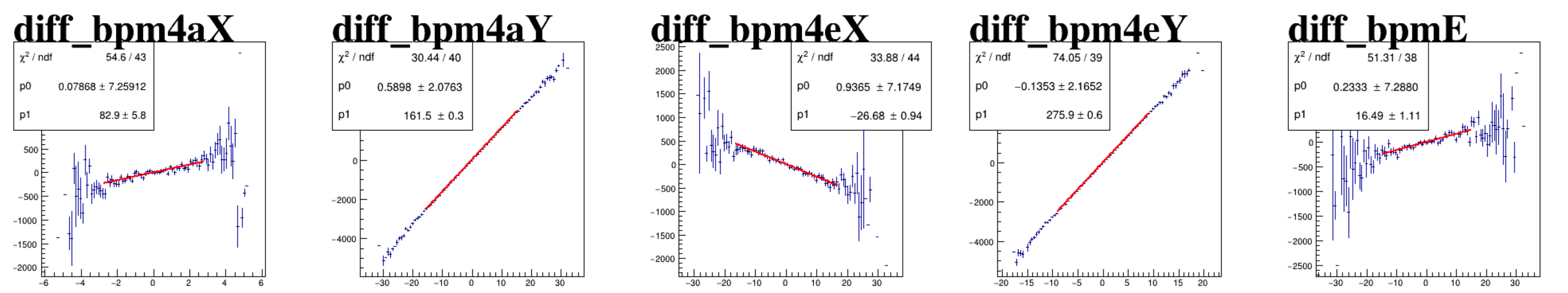
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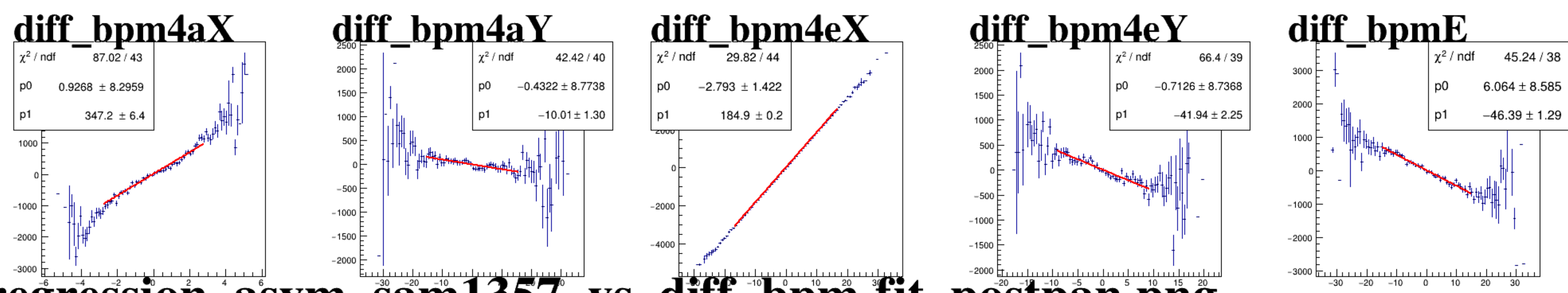
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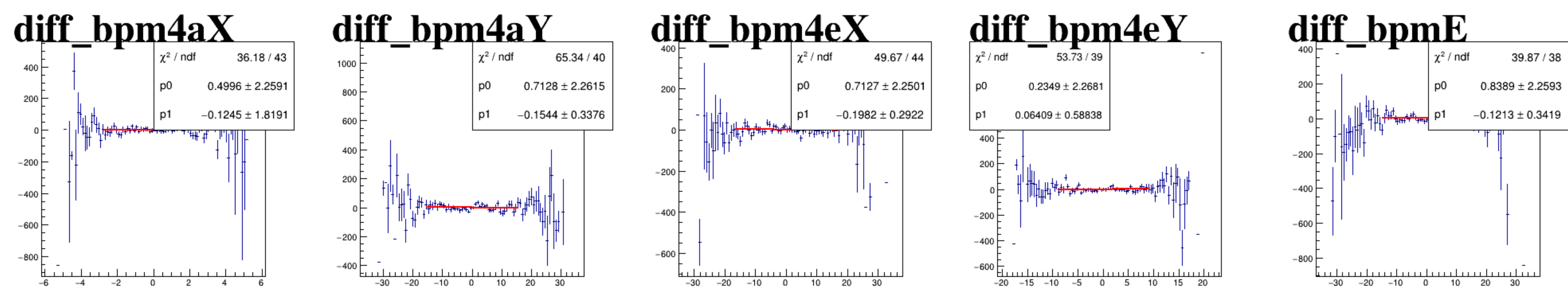
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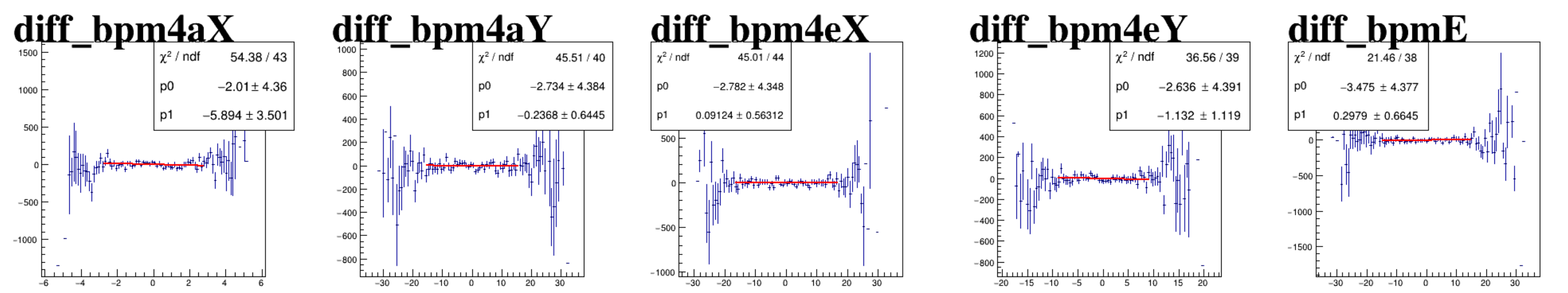
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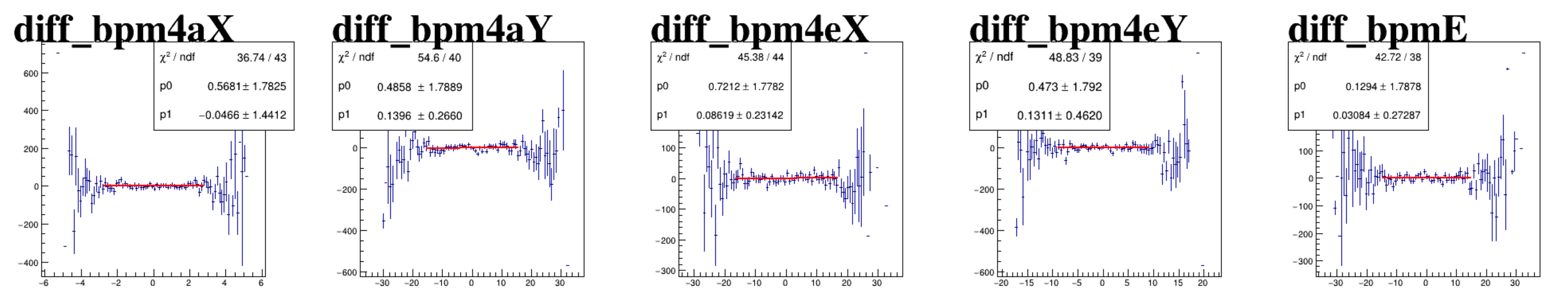
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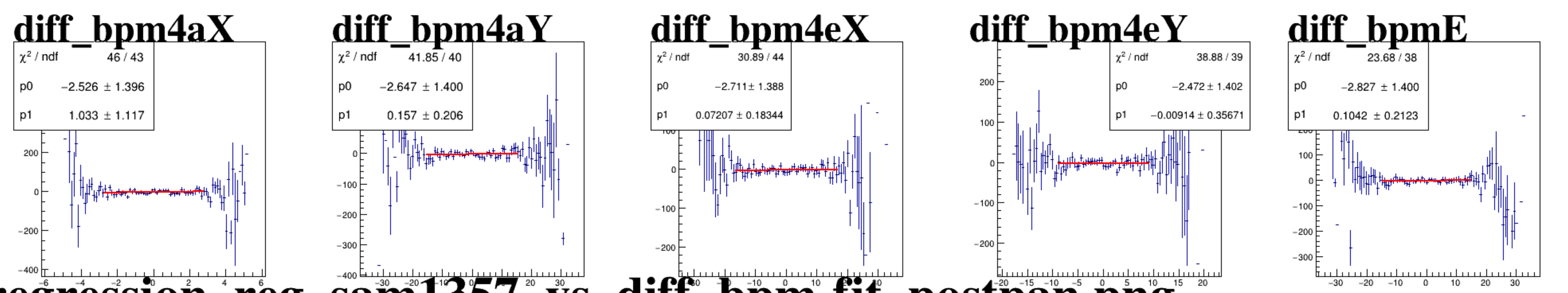
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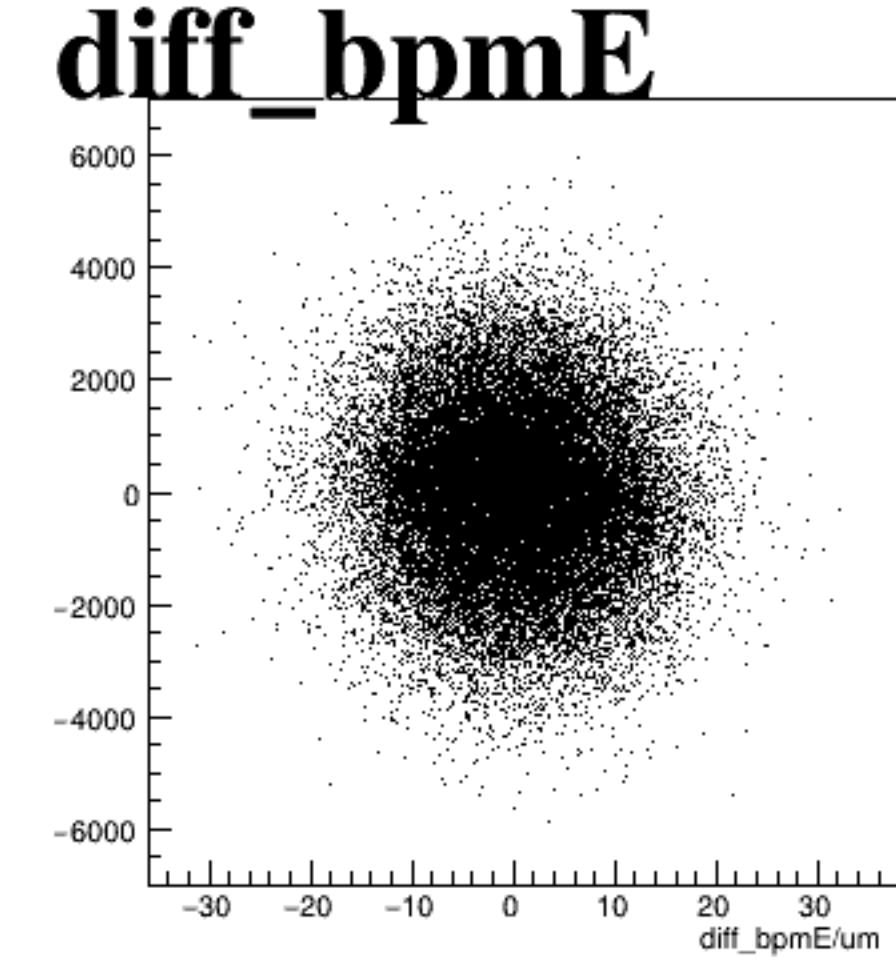
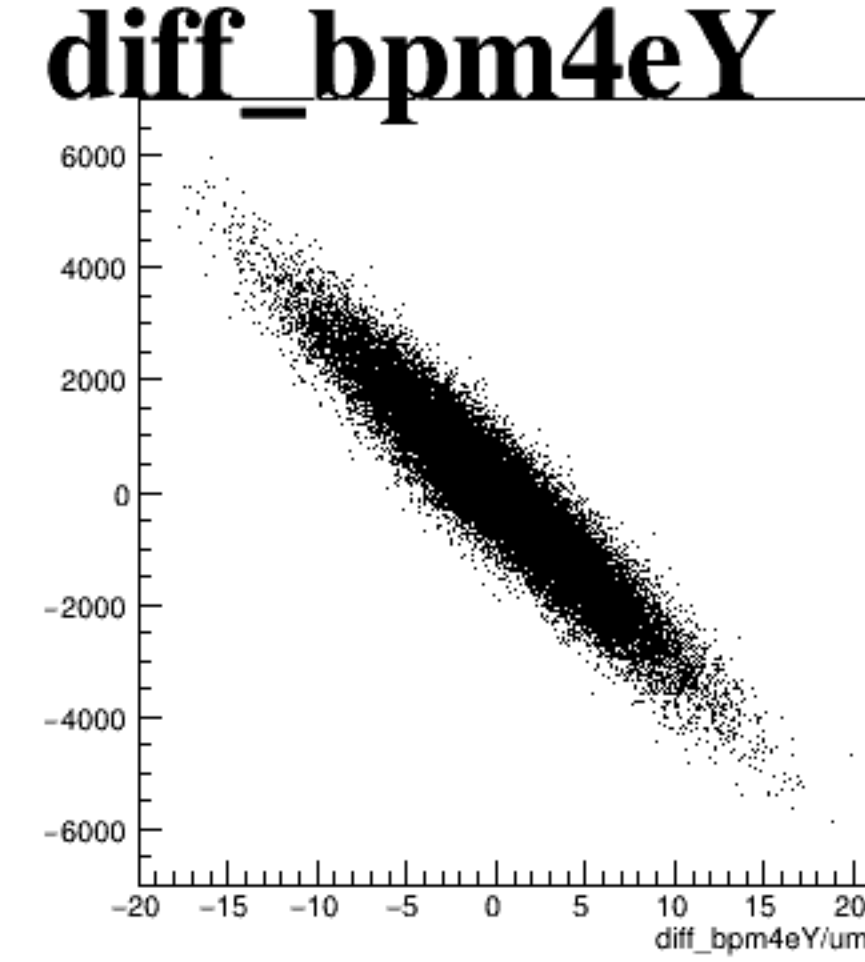
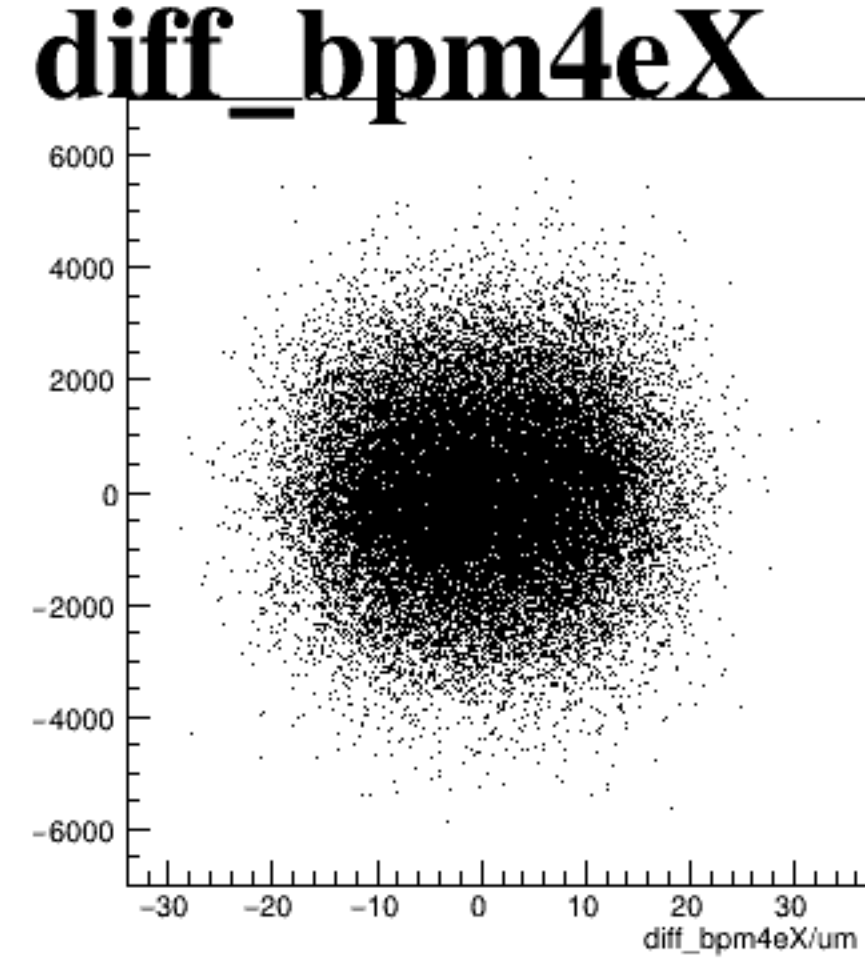
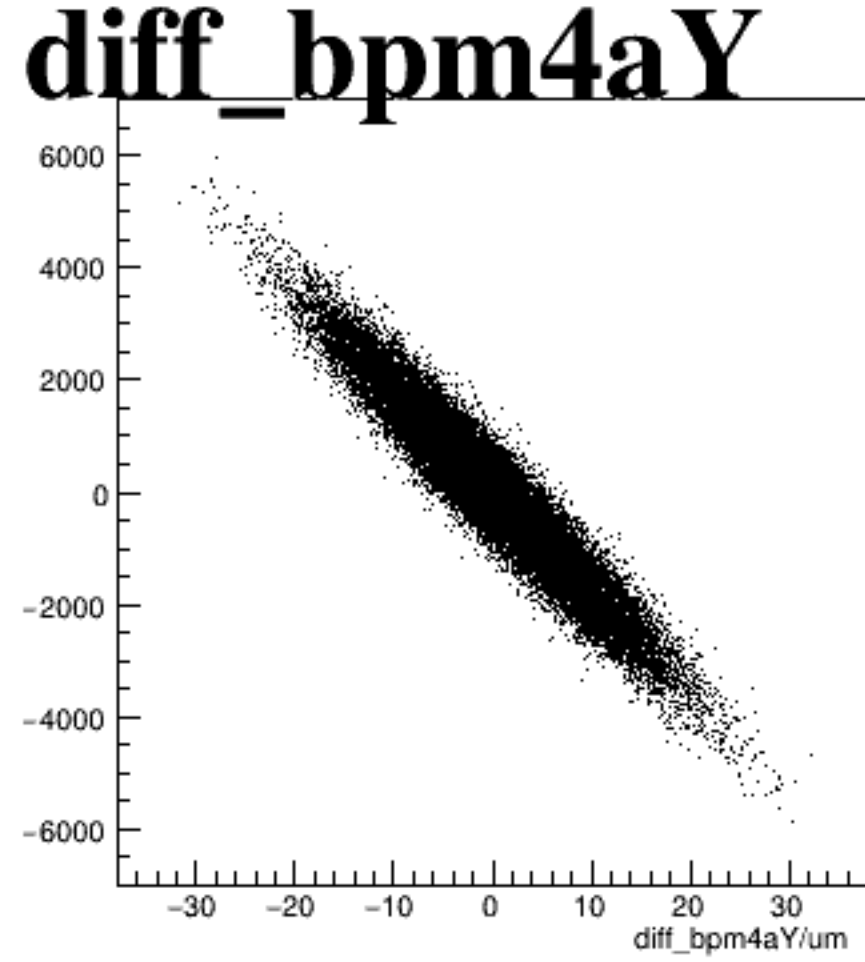
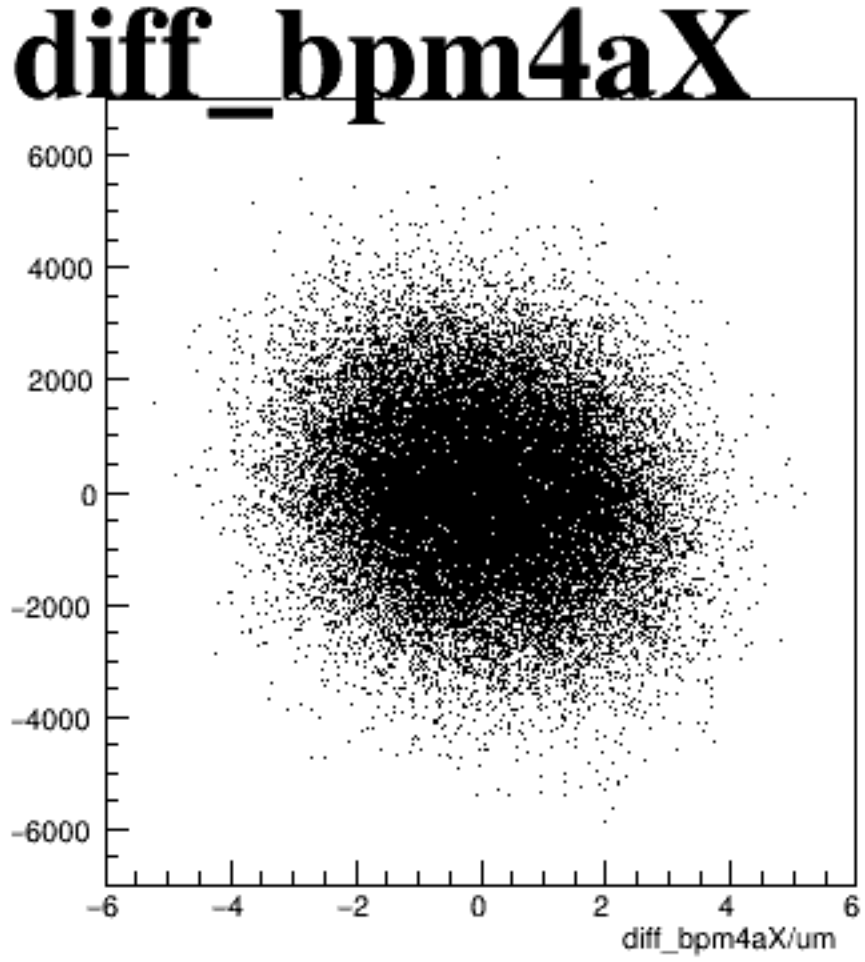
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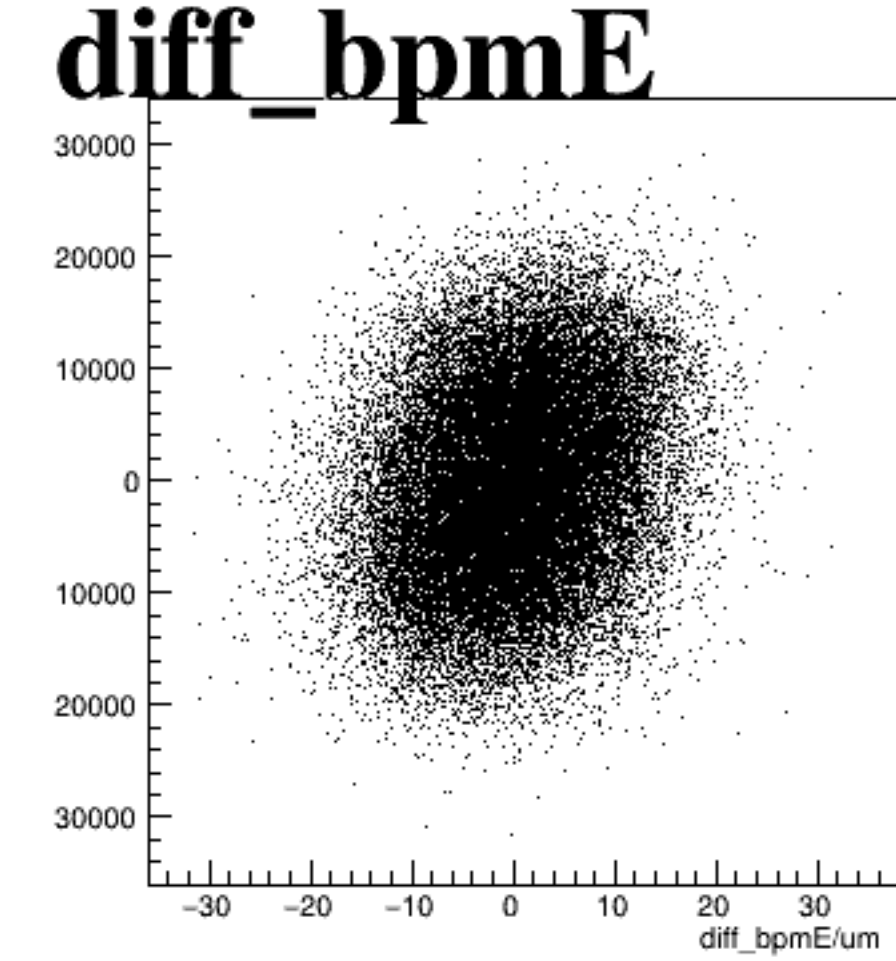
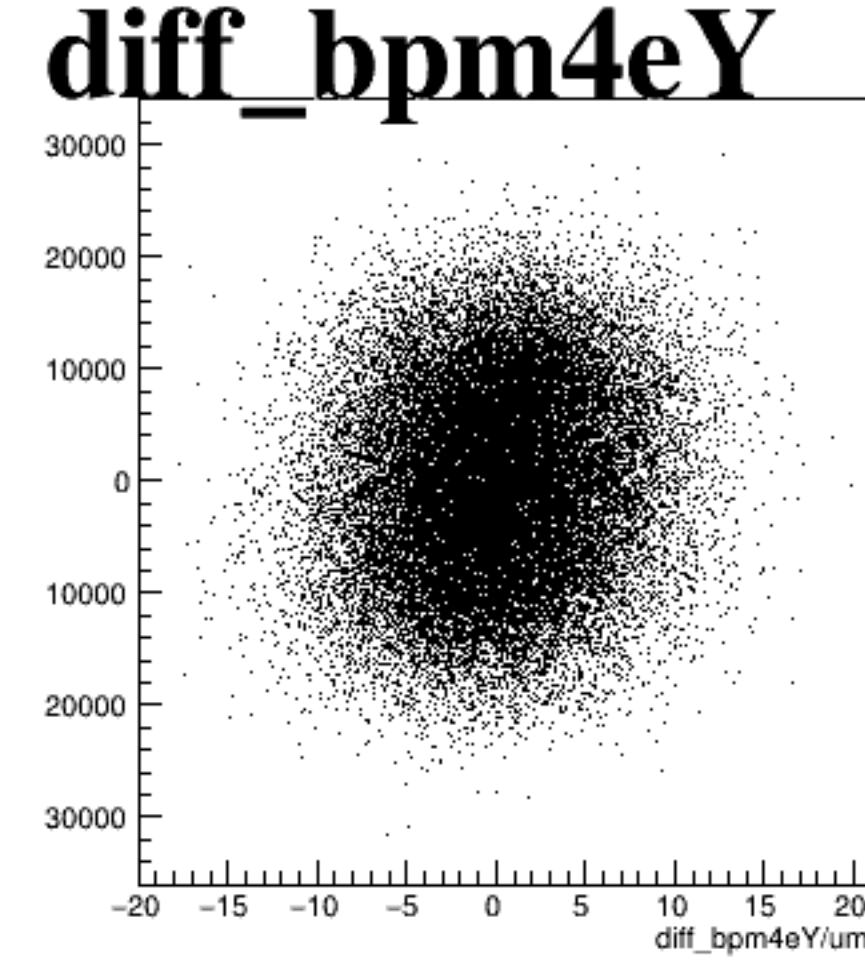
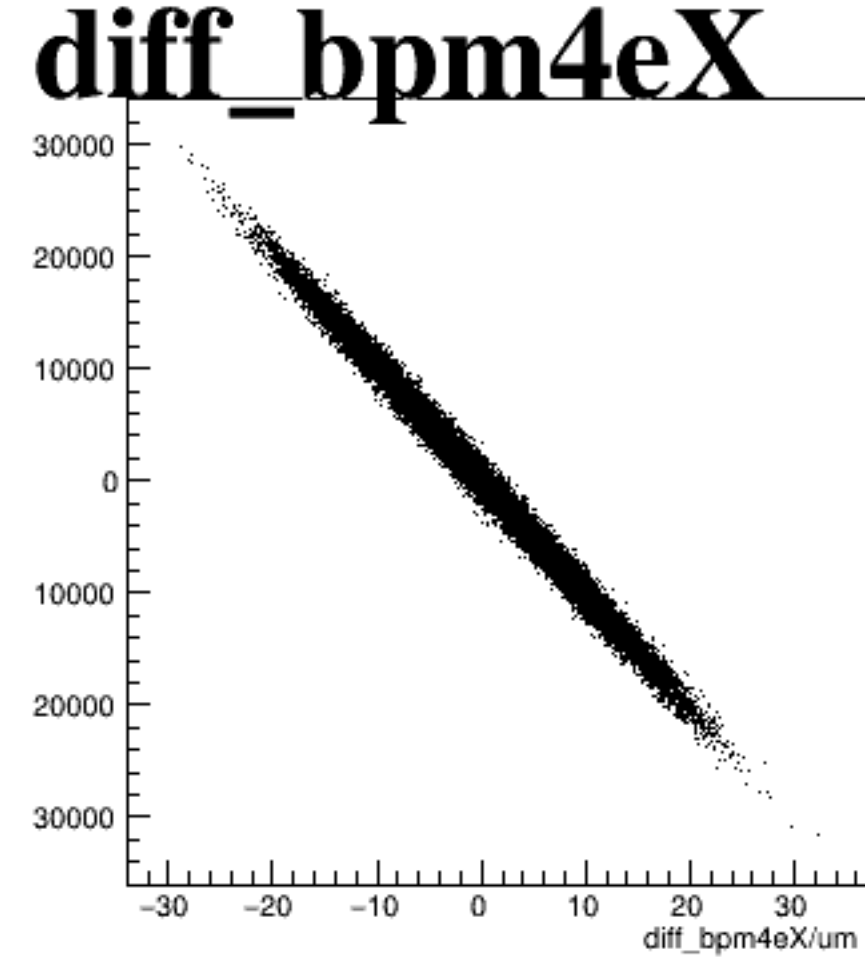
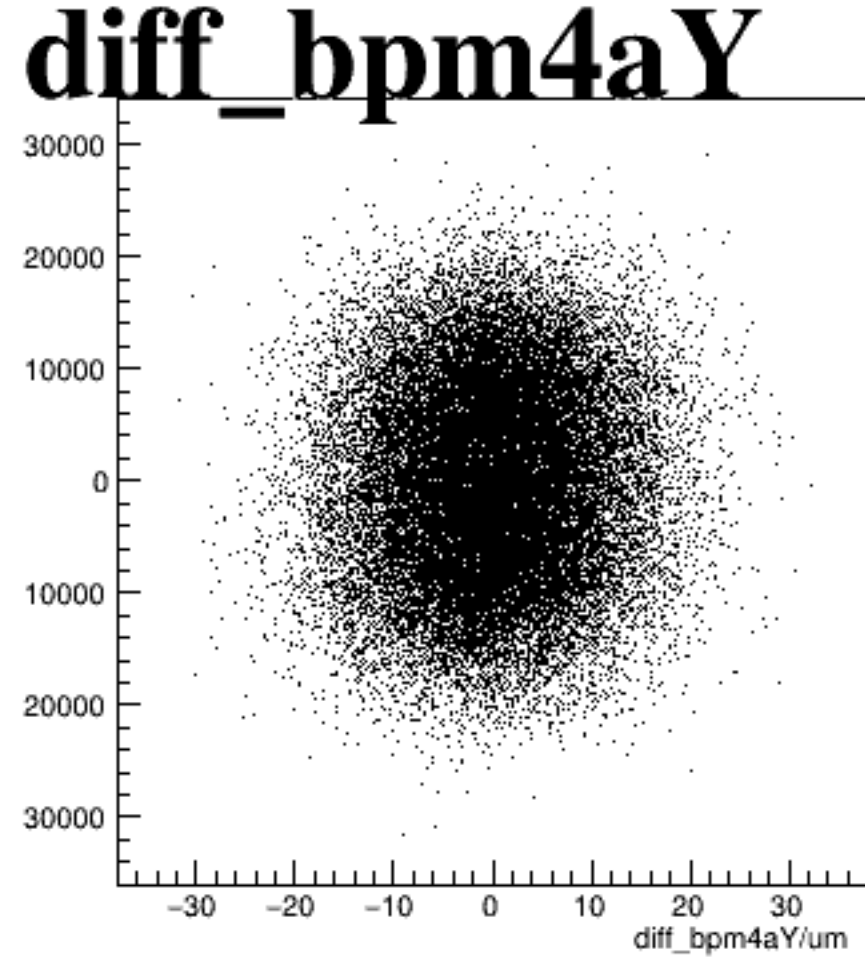
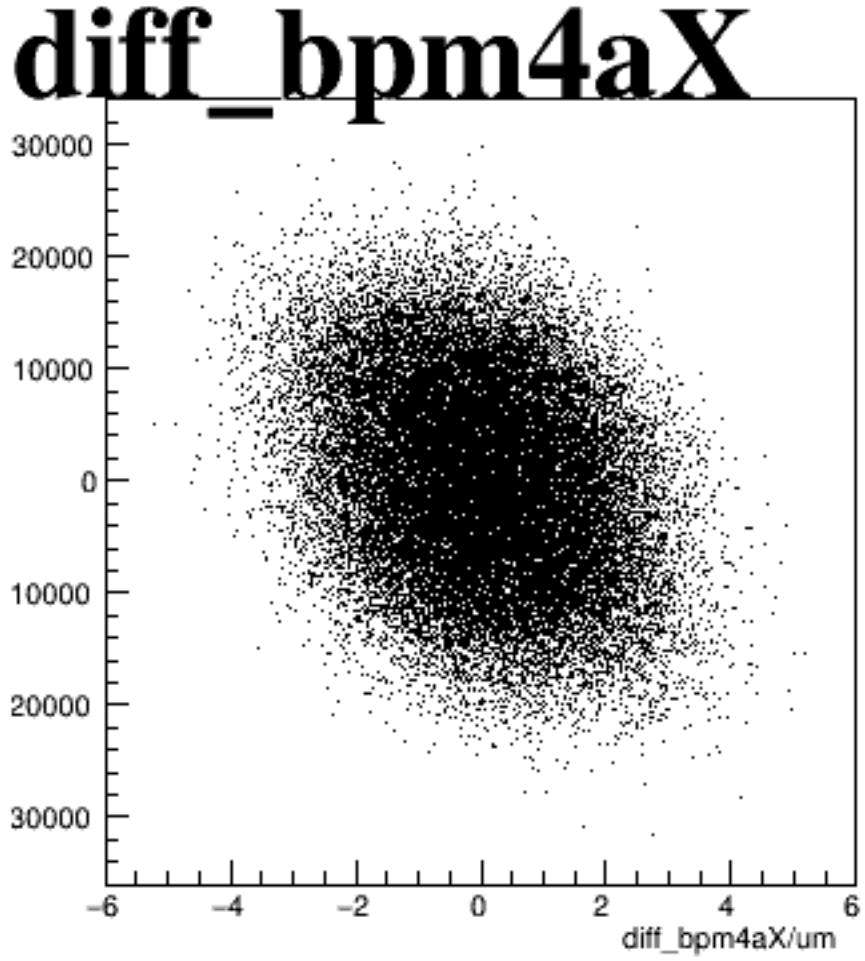
reg_asym_sam7



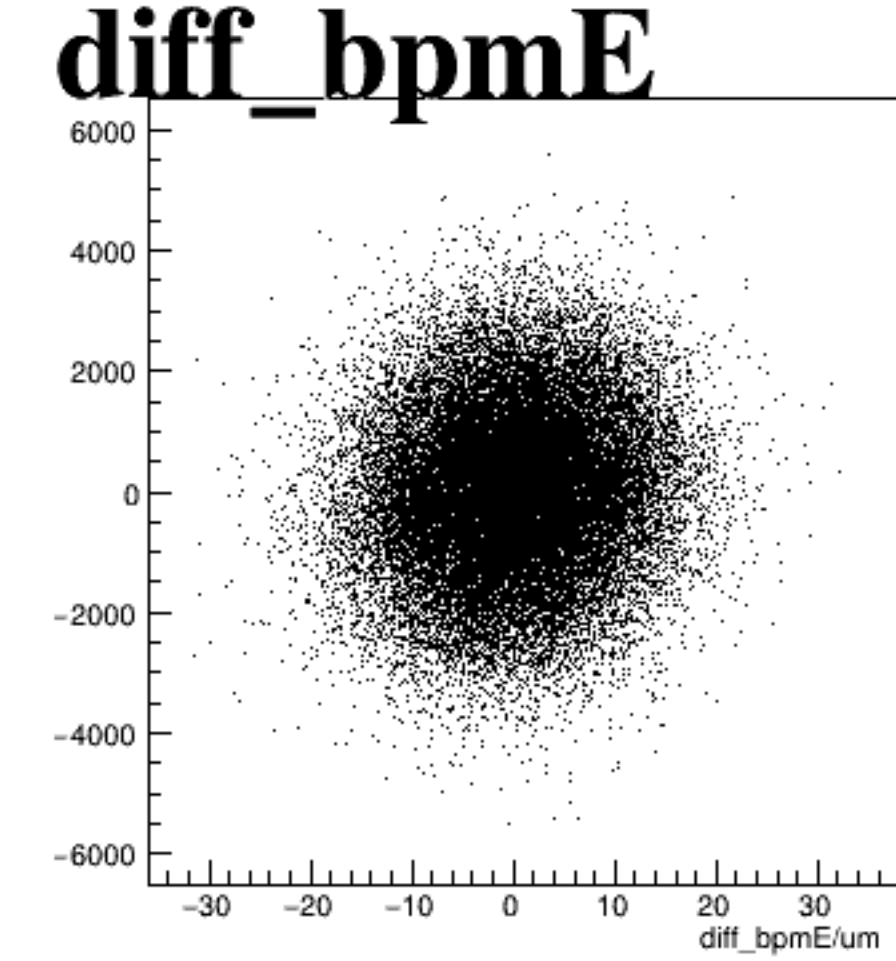
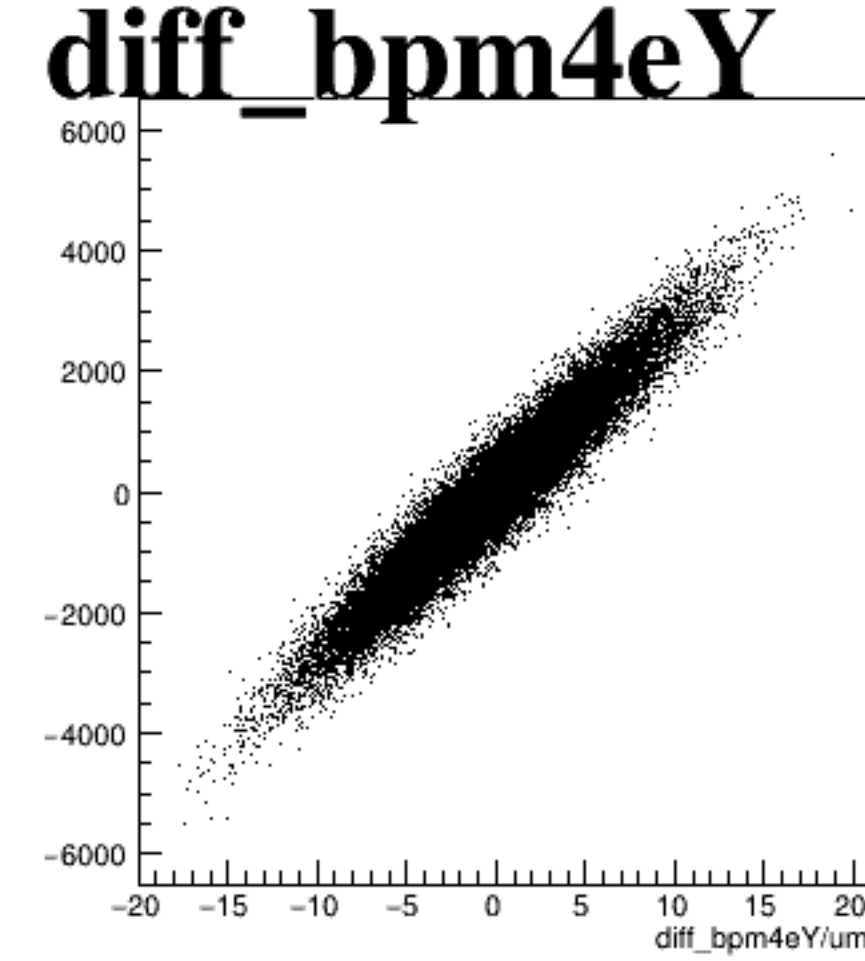
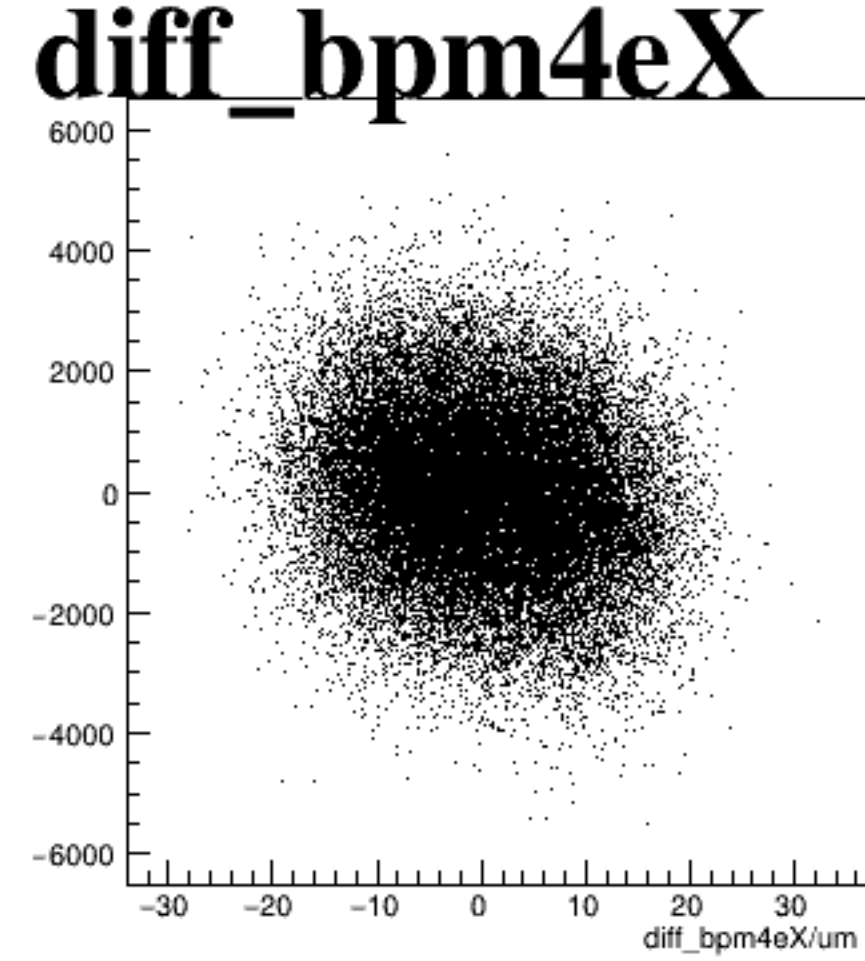
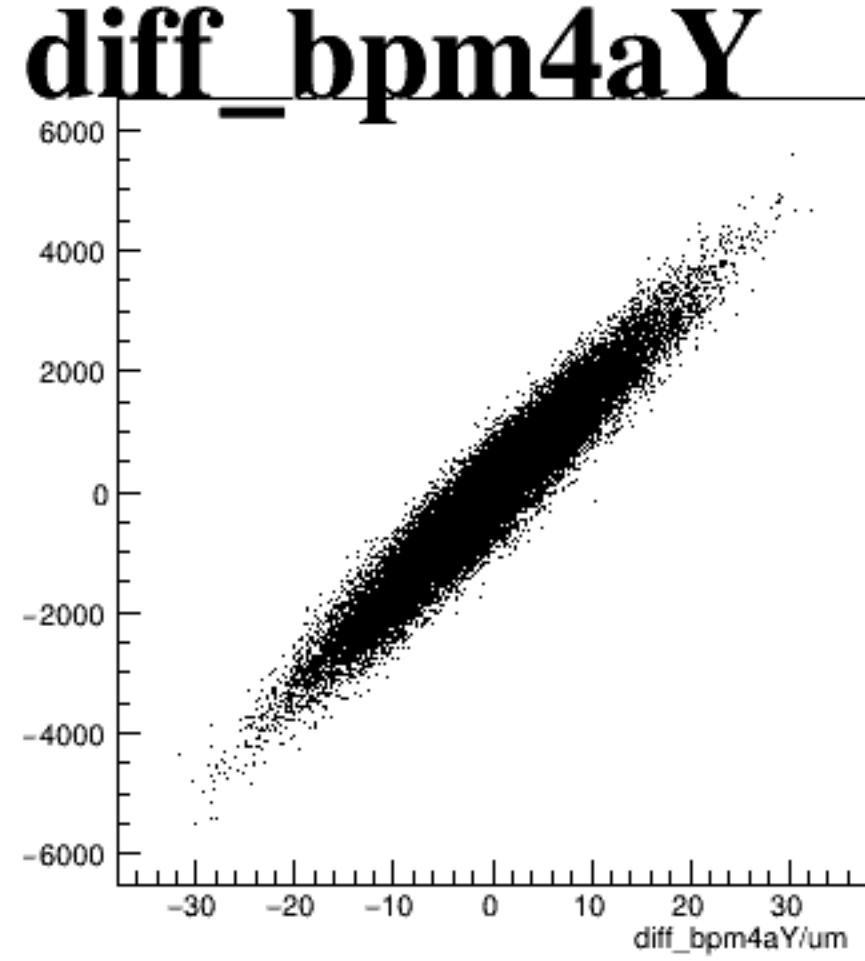
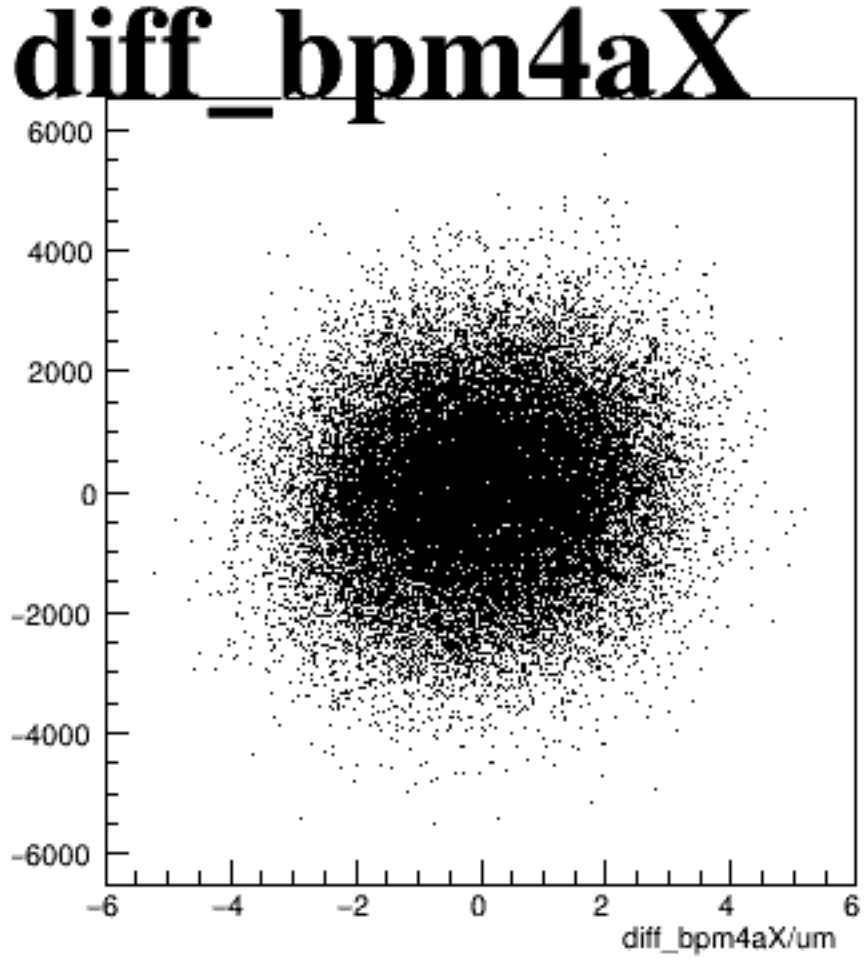
asym_sam1



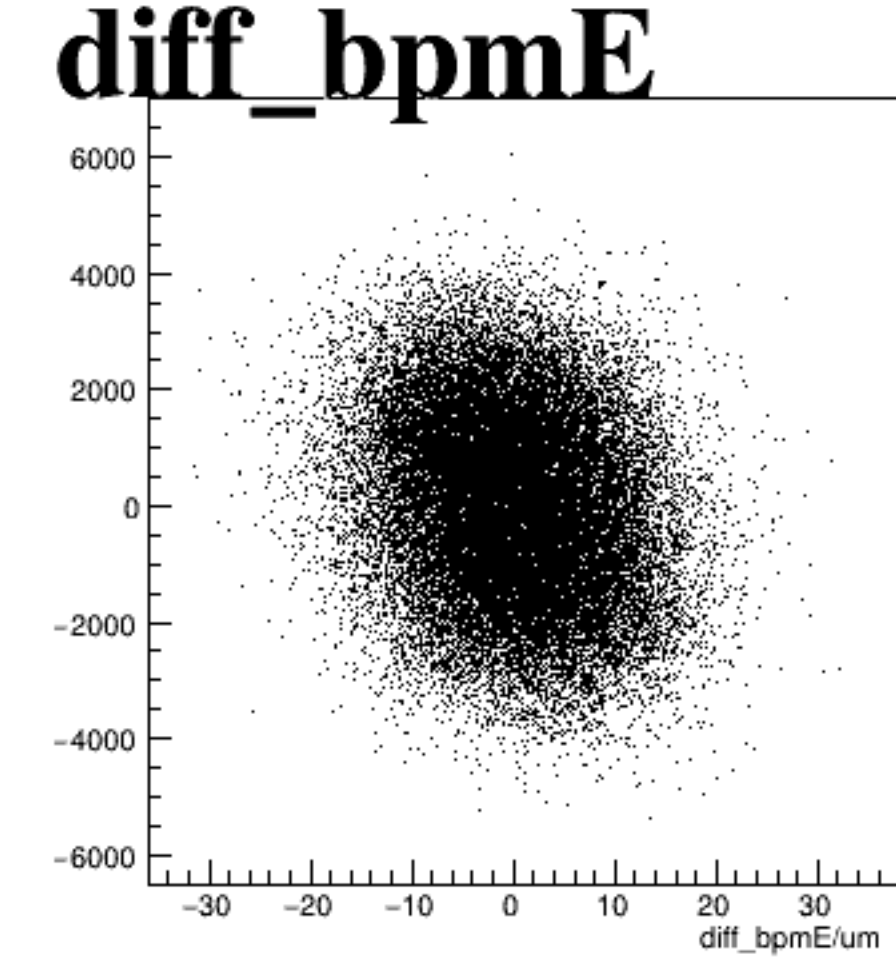
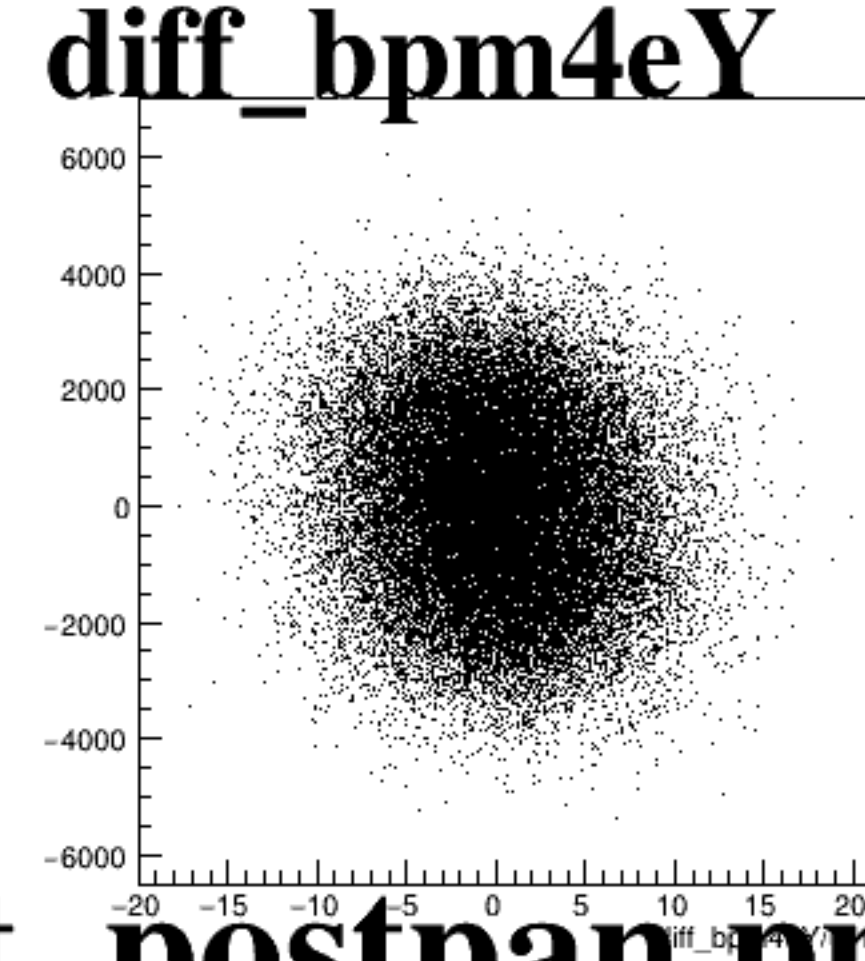
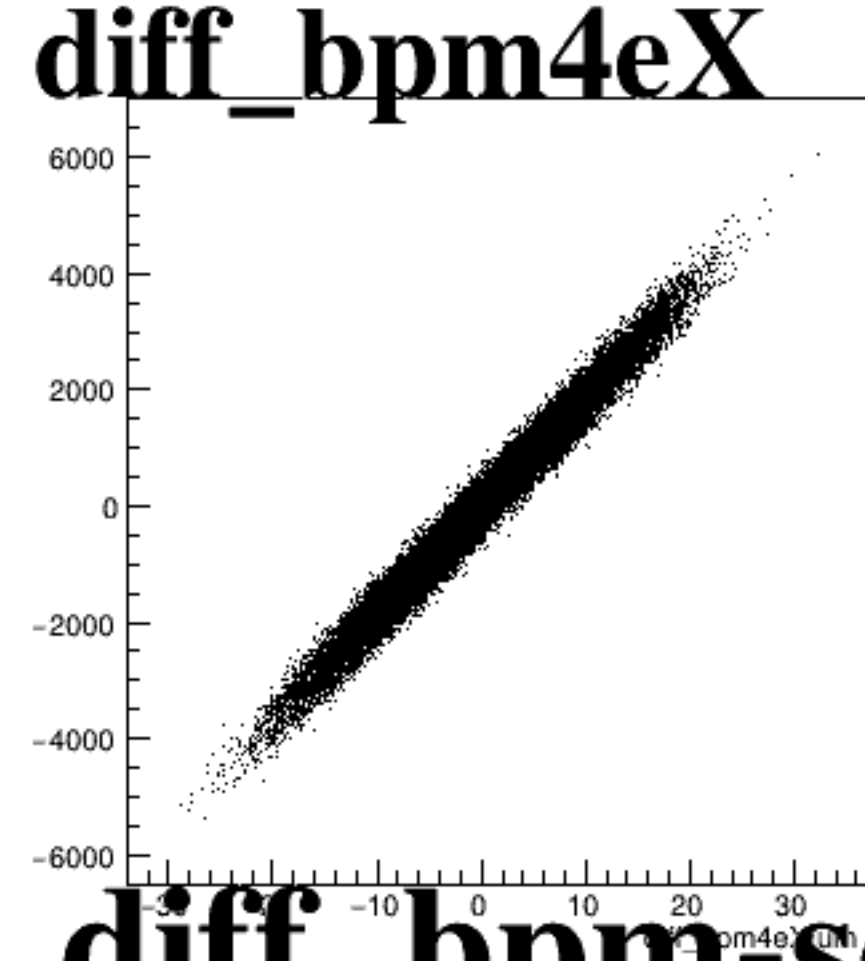
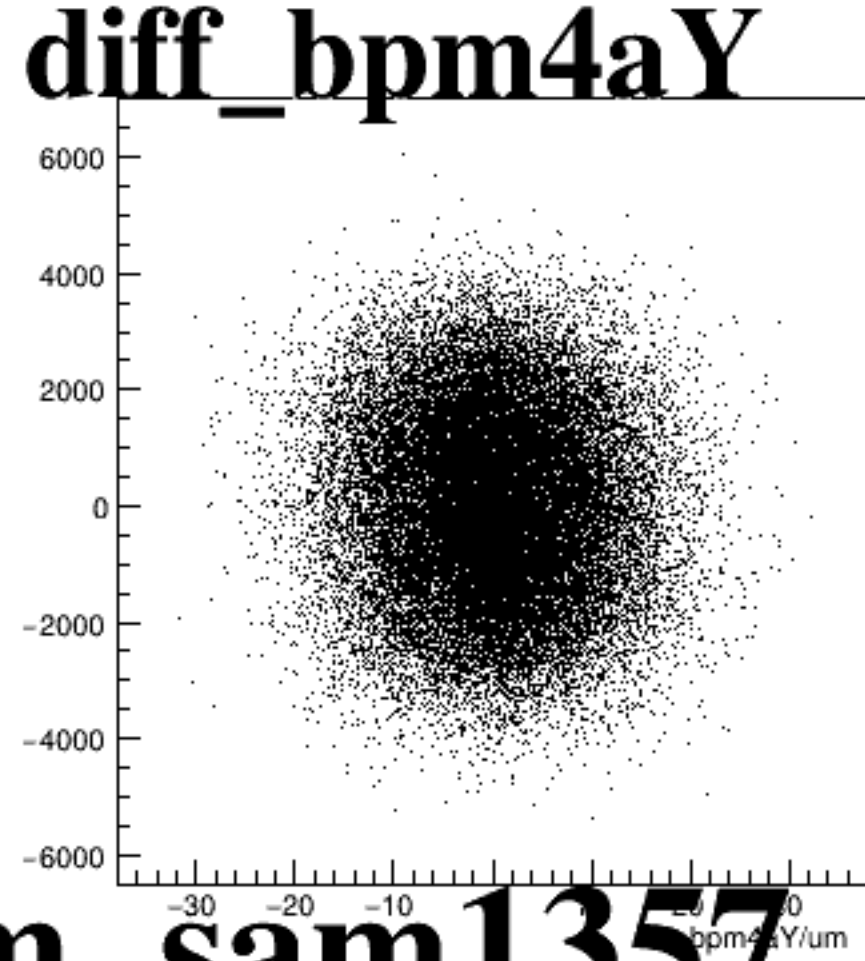
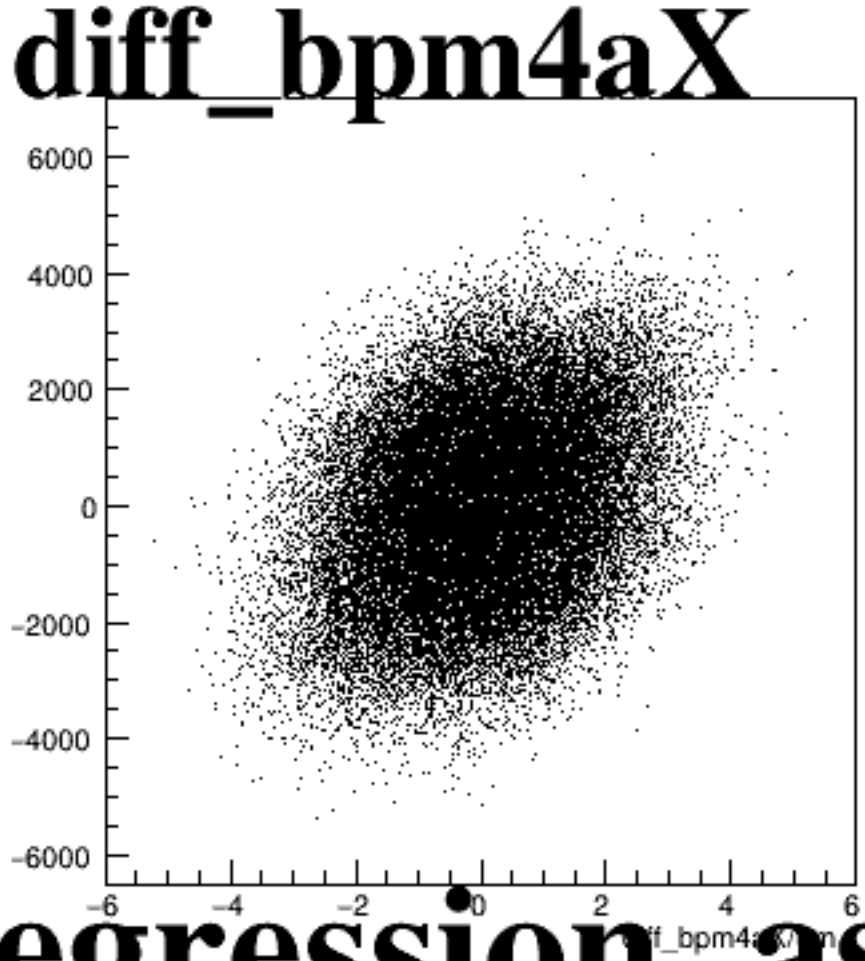
asym_sam3



asym_sam5



asym_sam7



The figure consists of five scatter plots arranged horizontally, each showing the distribution of differences between BPM4 and other models. The plots are titled: **diff_bpm4aX**, **diff_bpm4aY**, **diff_bpm4eX**, **diff_bpm4eY**, and **diff_bpmE**. Each plot has a y-axis ranging from -1500 to 1500 and an x-axis ranging from -6 to 6 for **diff_bpm4aX**, and -30 to 30 for the others. The data points are densely clustered around the origin, indicating a high degree of agreement between the models.

The figure consists of five scatter plots arranged horizontally. Each plot shows a dense cloud of points centered around zero, indicating a good fit or agreement between the compared variables. The y-axis for all plots ranges from -1000 to 1000. The x-axis labels are partially visible at the bottom of the plots.

- diff_bpm4aX**: The x-axis is labeled `diff_bpm4aX/um`.
- diff_bpm4aY**: The x-axis is labeled `diff_bpm4aY/um`.
- diff_bpm4eX**: The x-axis is labeled `diff_bpm4eX/um`.
- diff_bpm4eY**: The x-axis is labeled `diff_bpm4eY/um`.
- diff_bpmE**: The x-axis is labeled `diff_bpmE/um`.