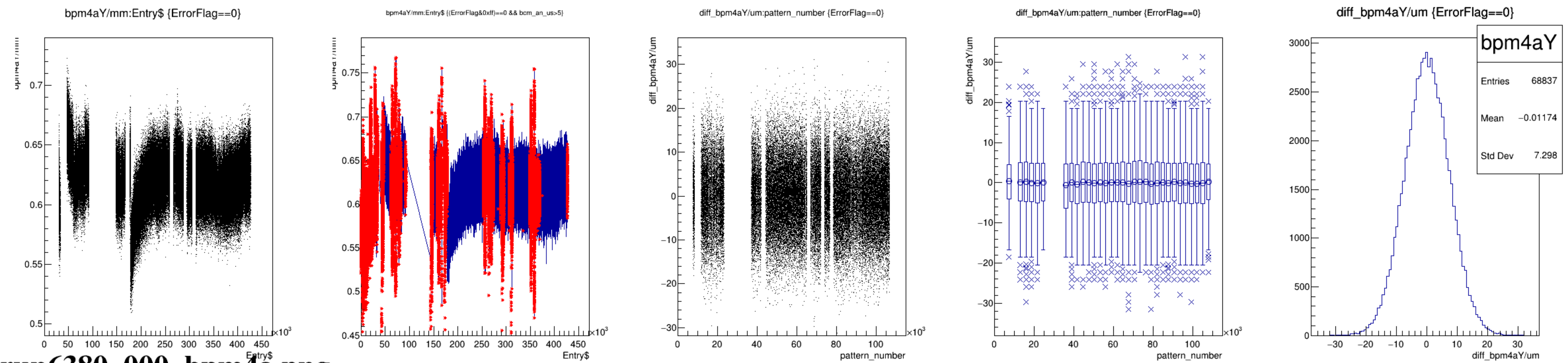
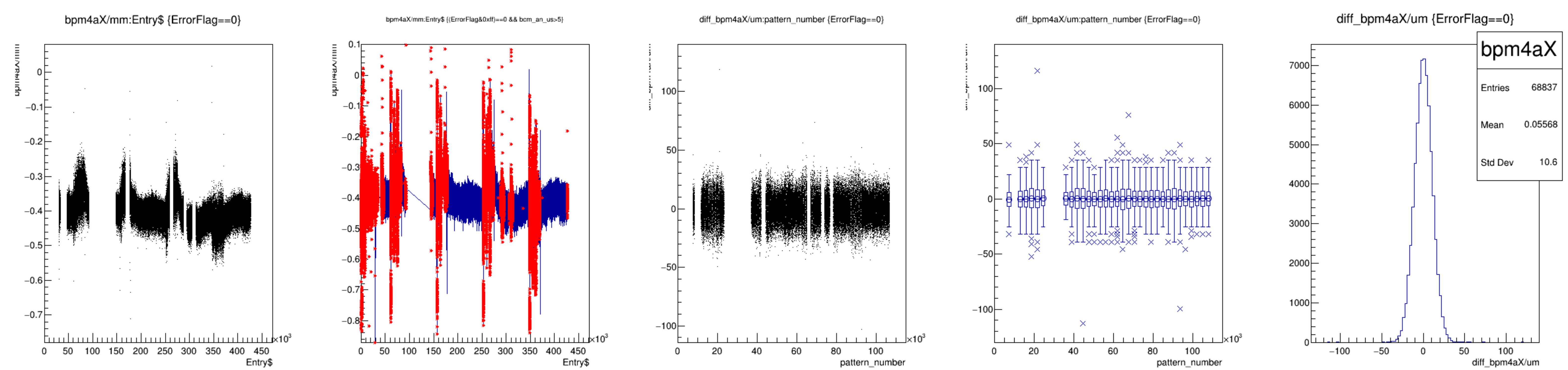
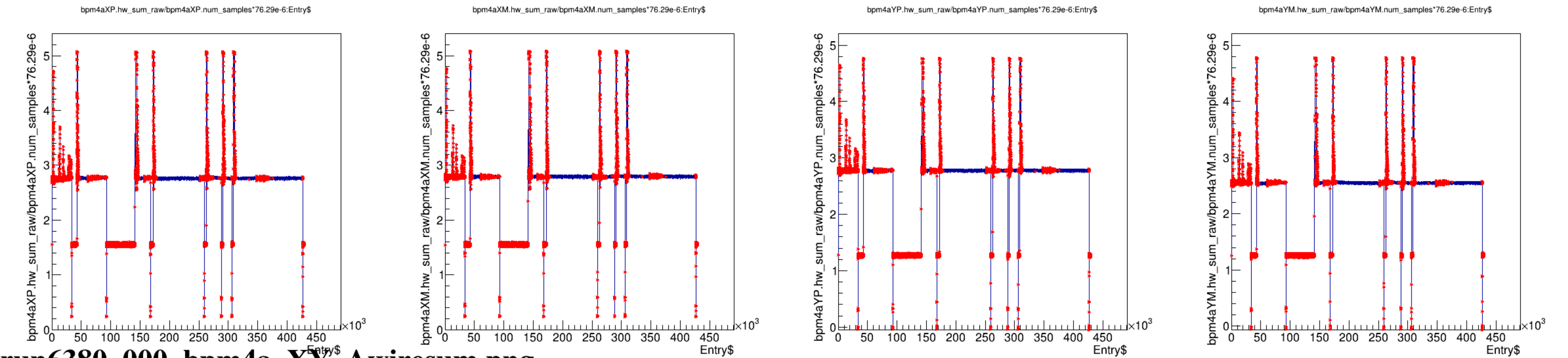
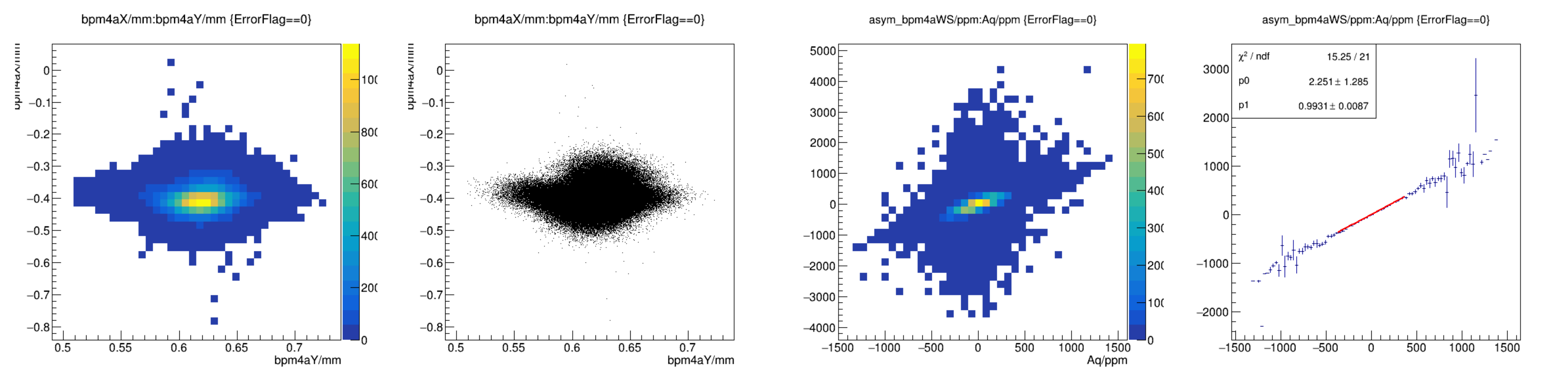
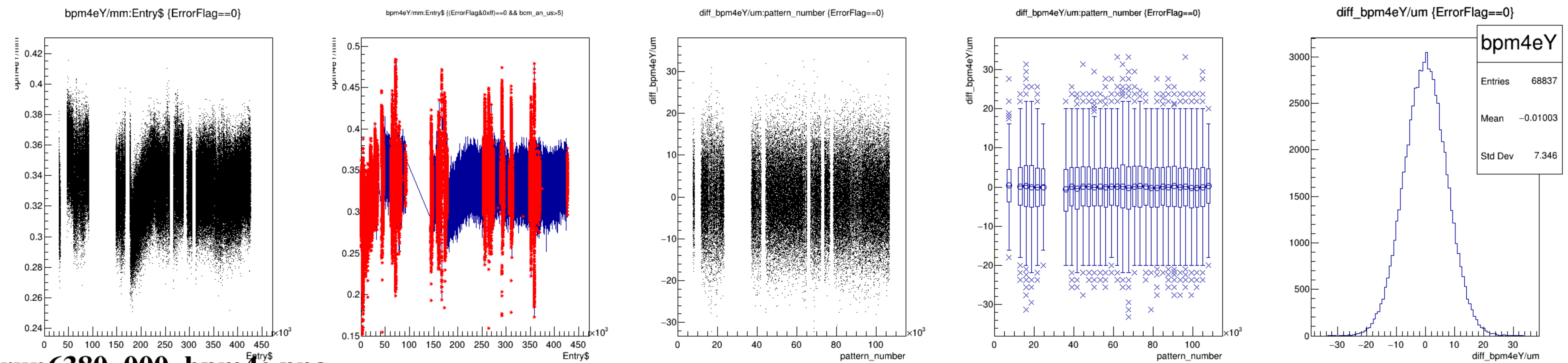
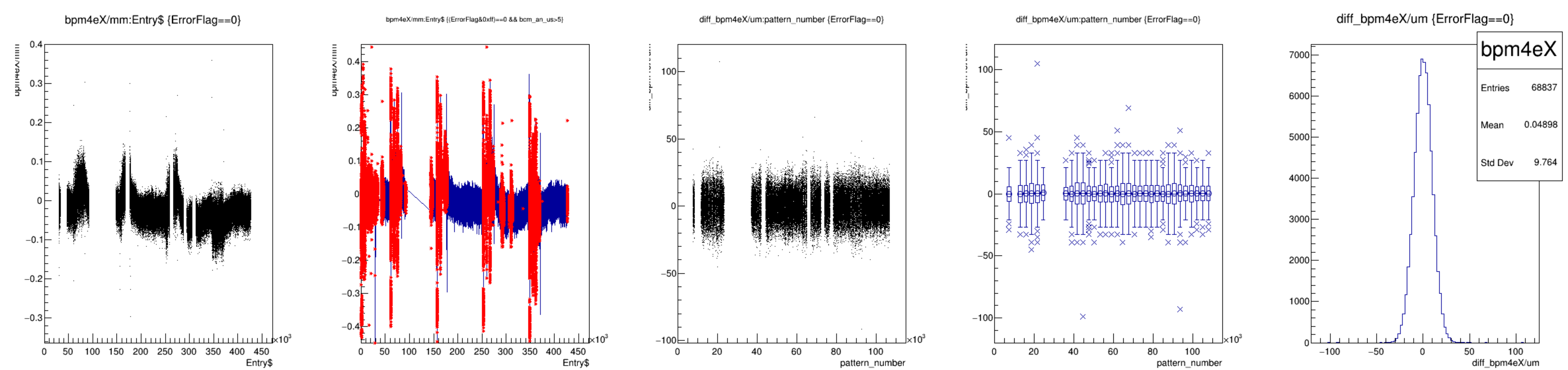
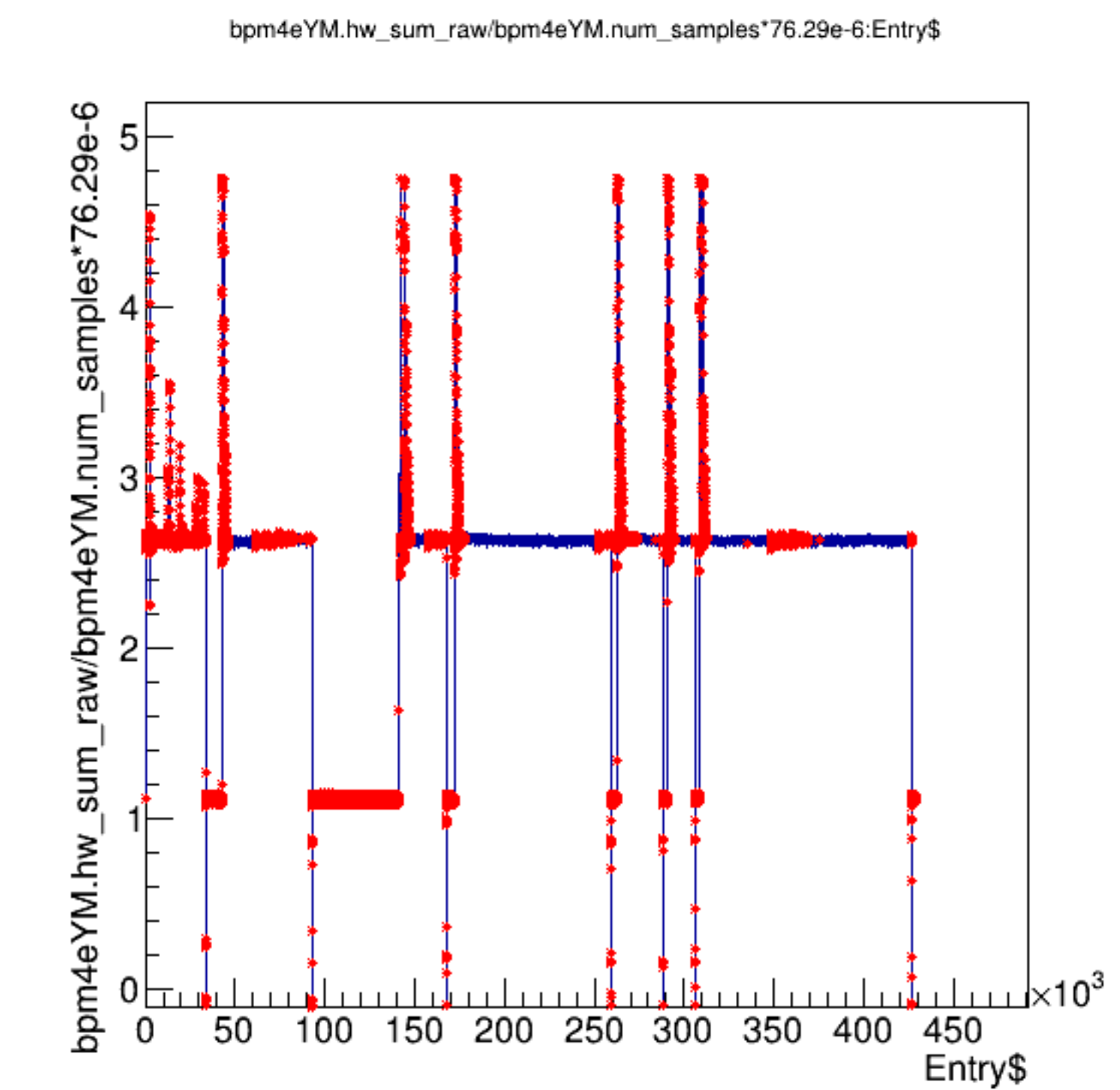
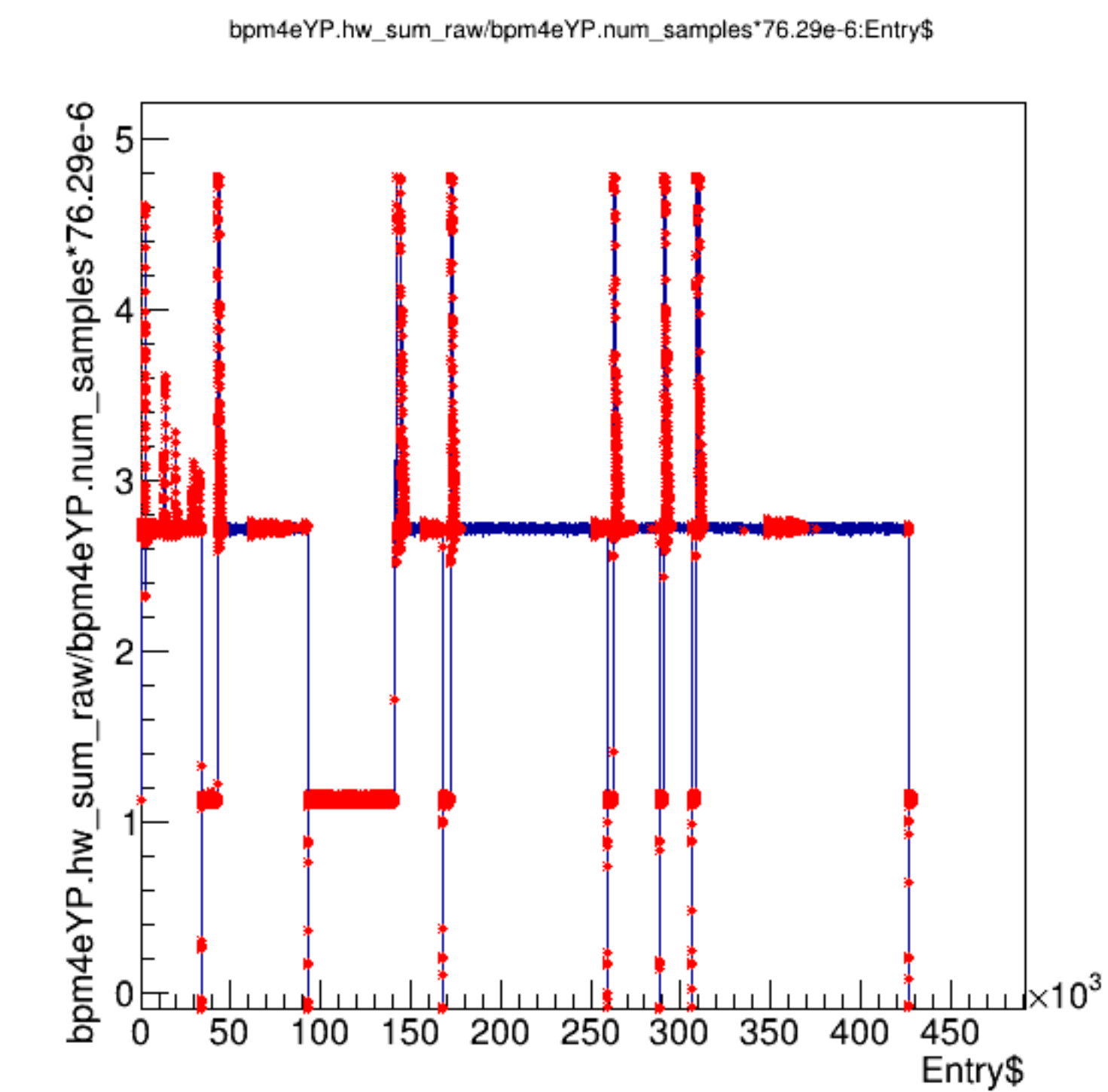
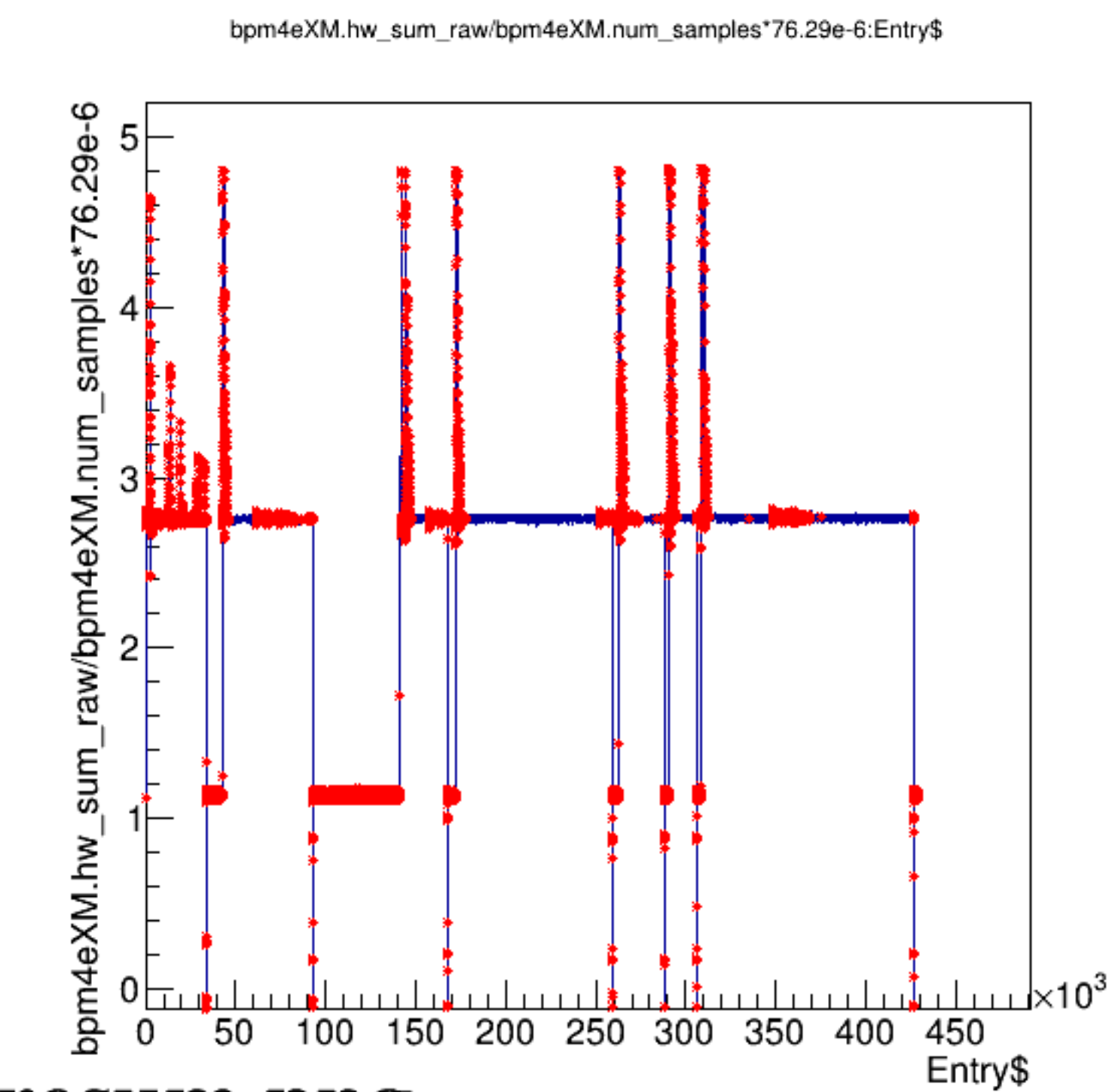
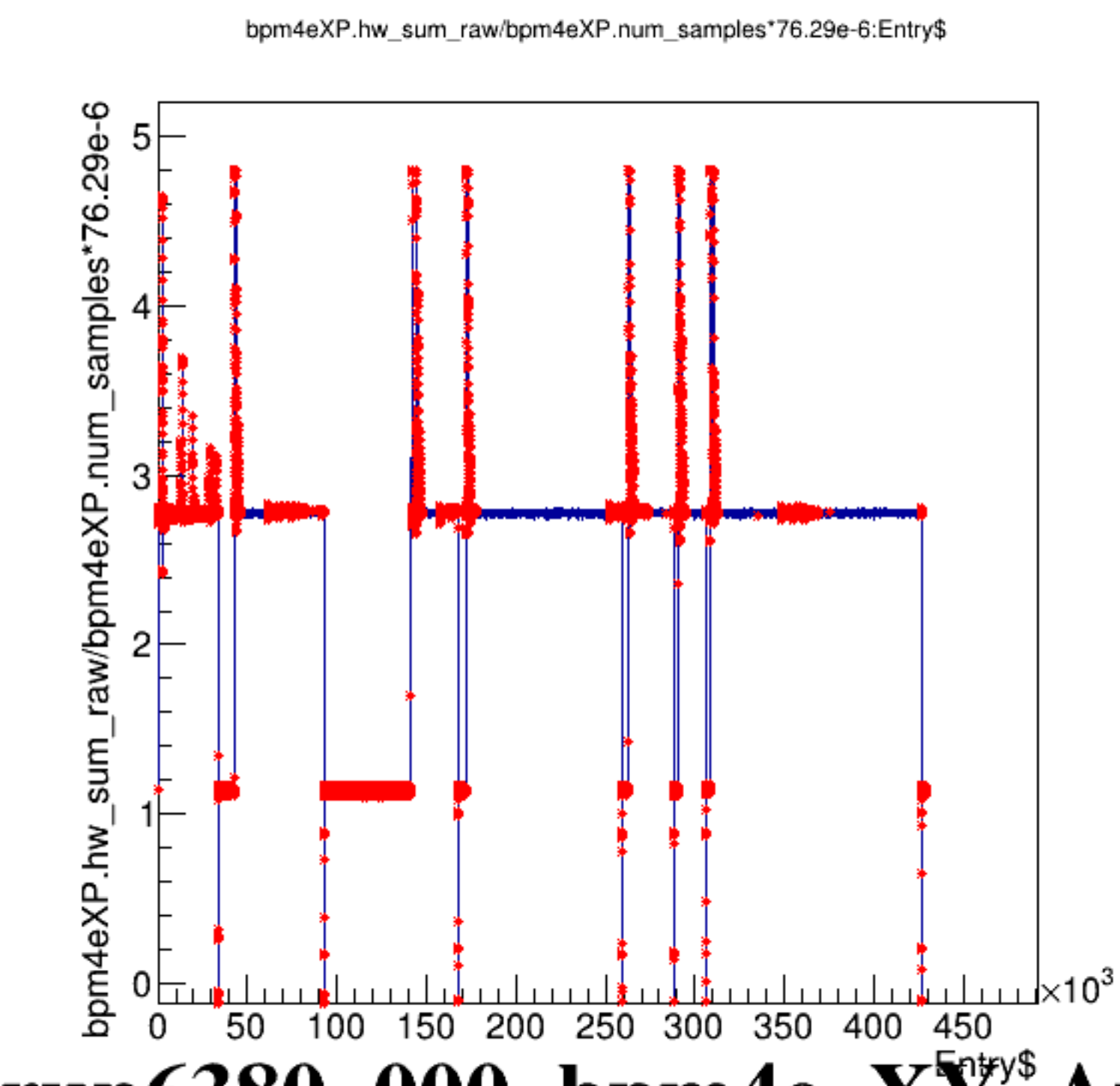
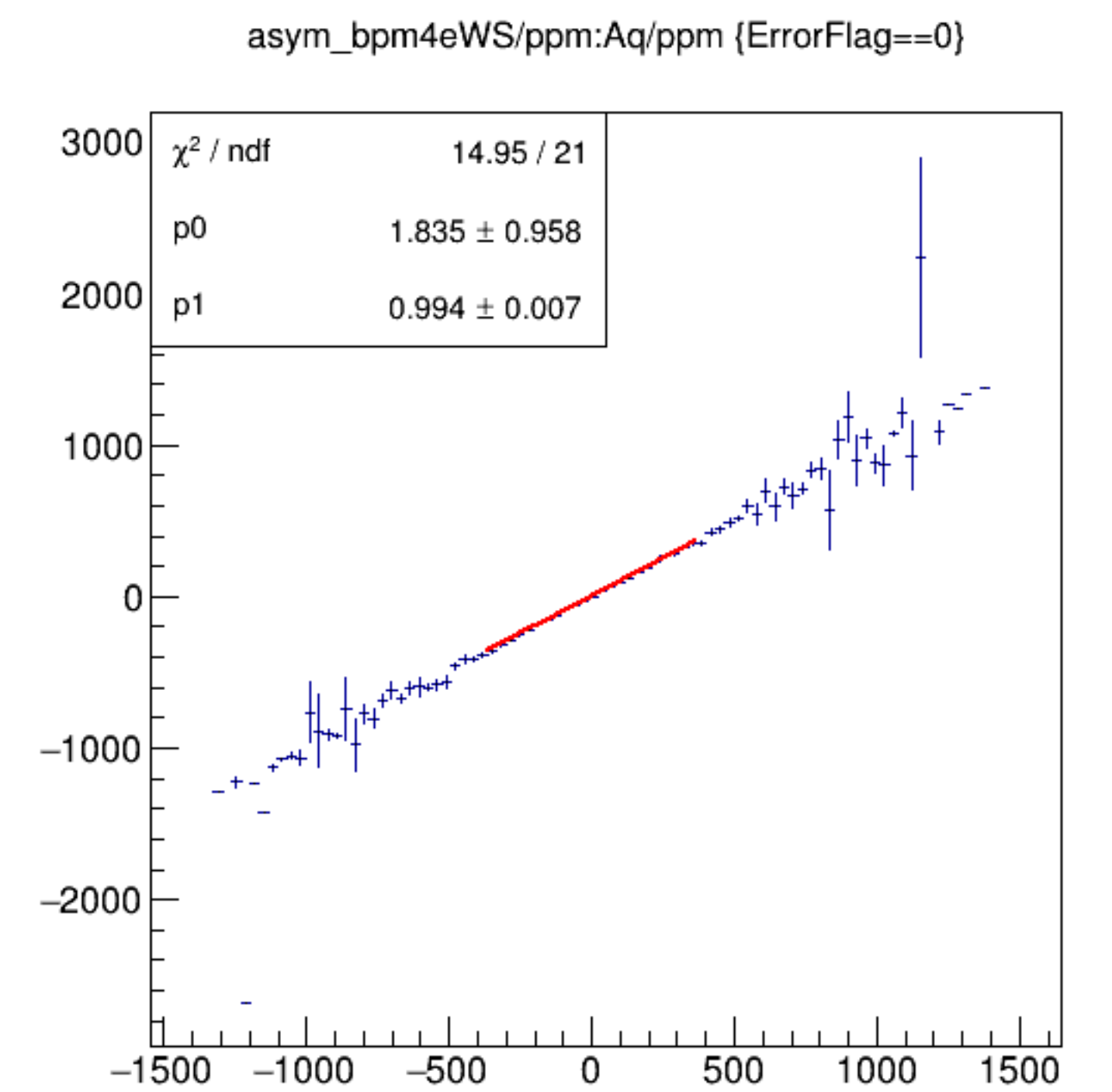
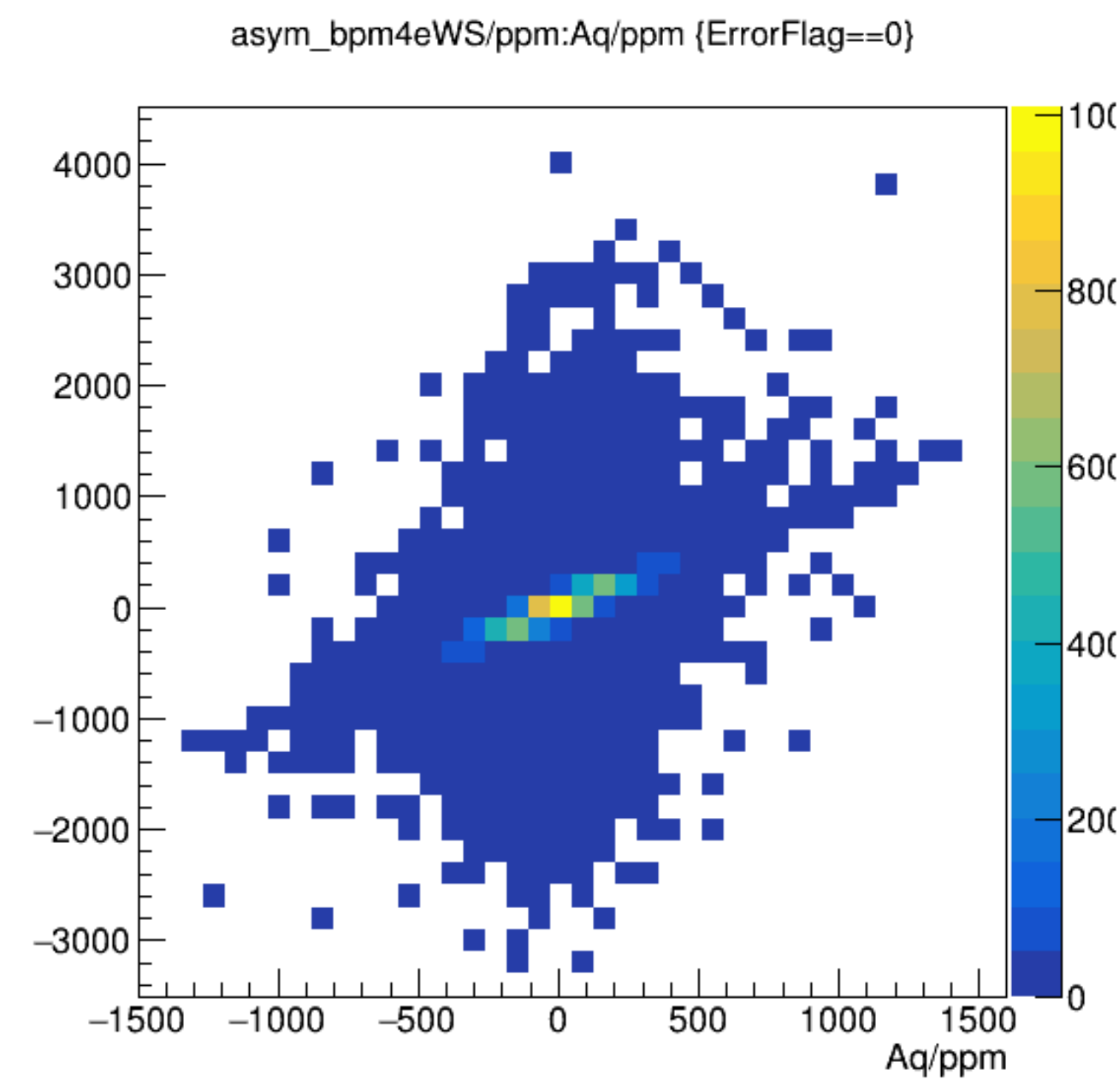
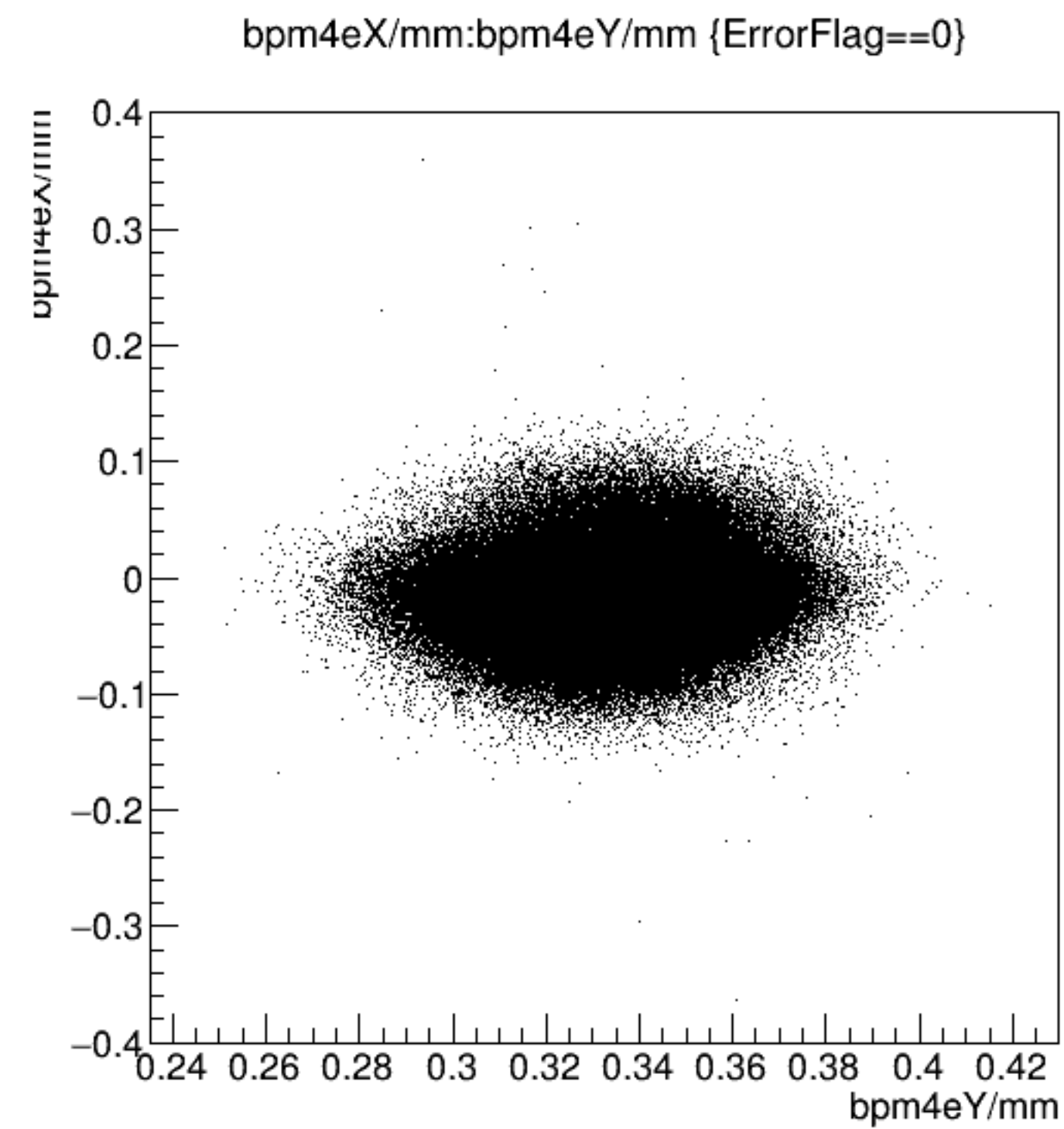
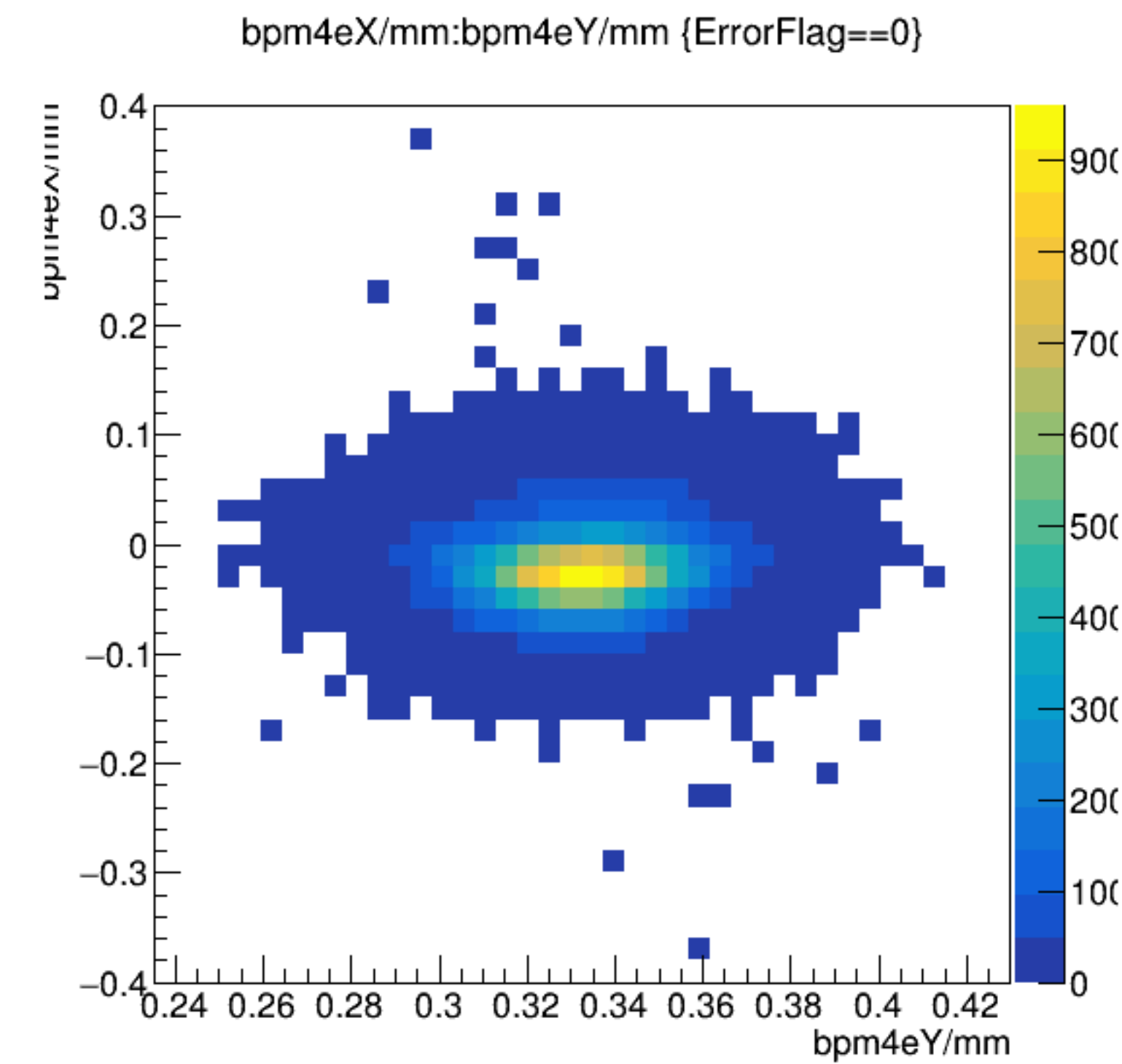




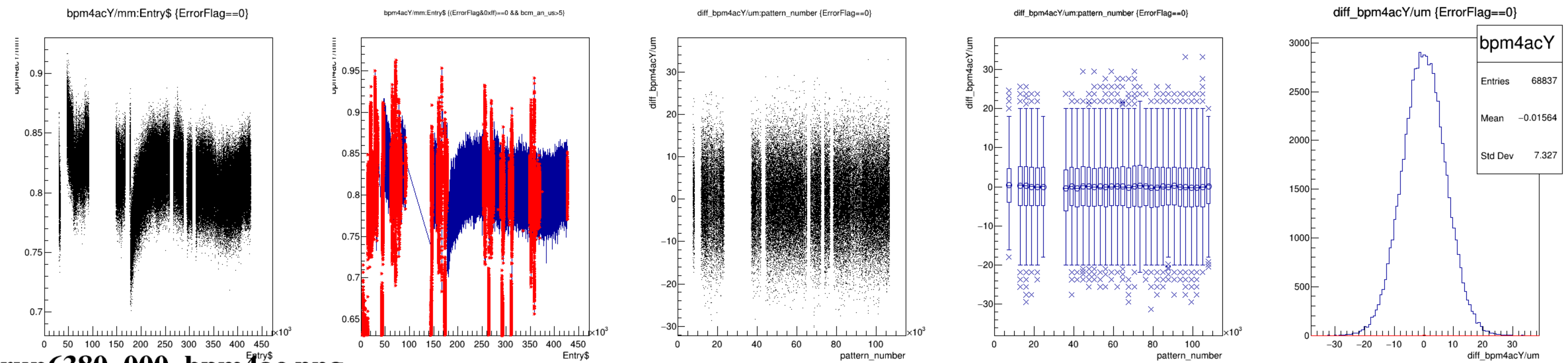
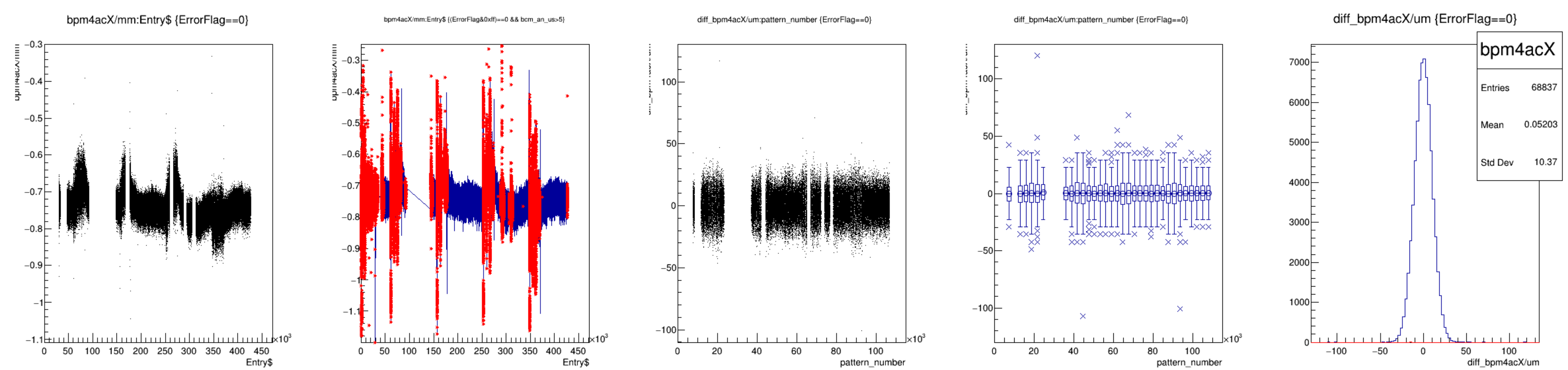


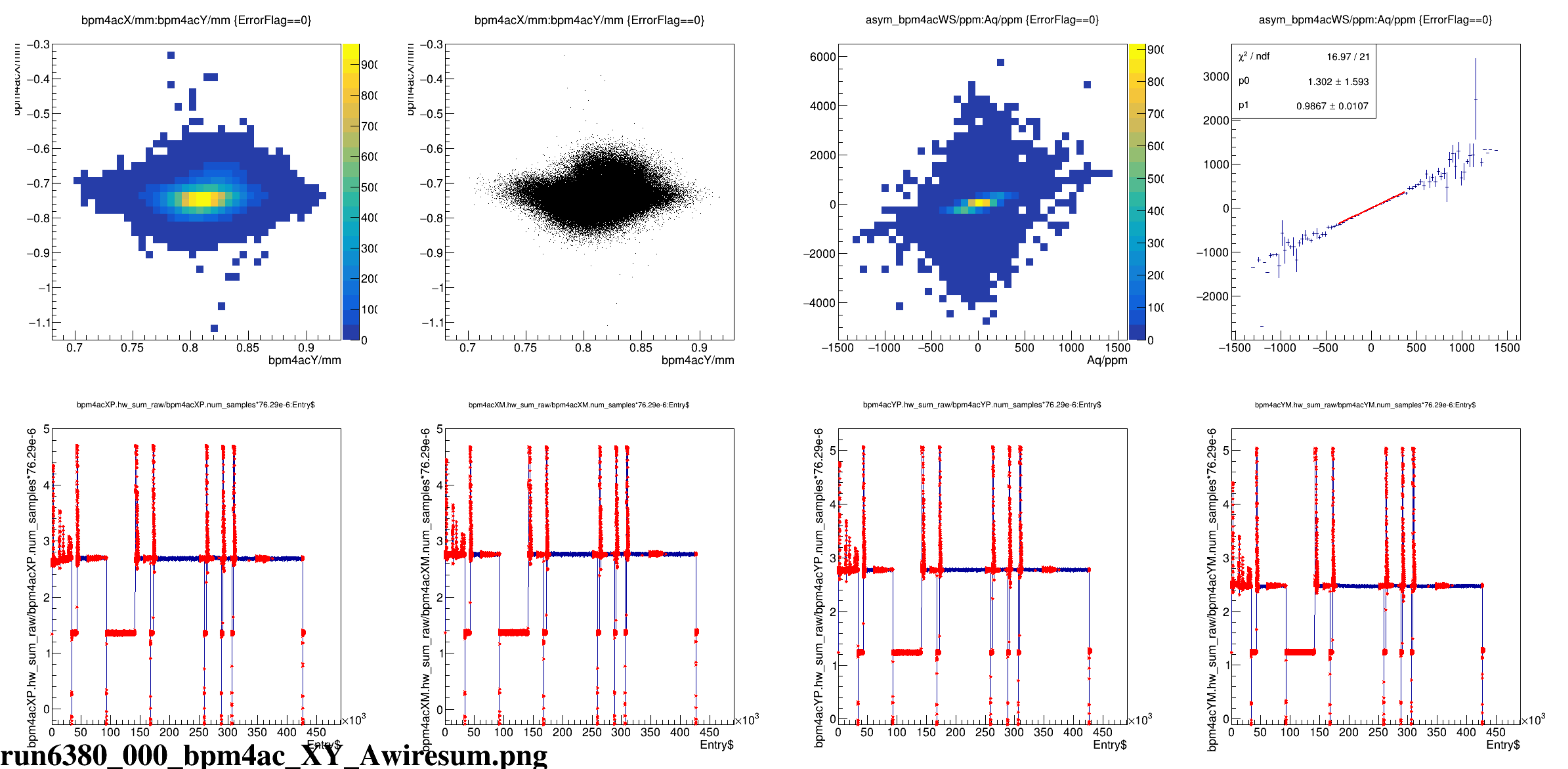
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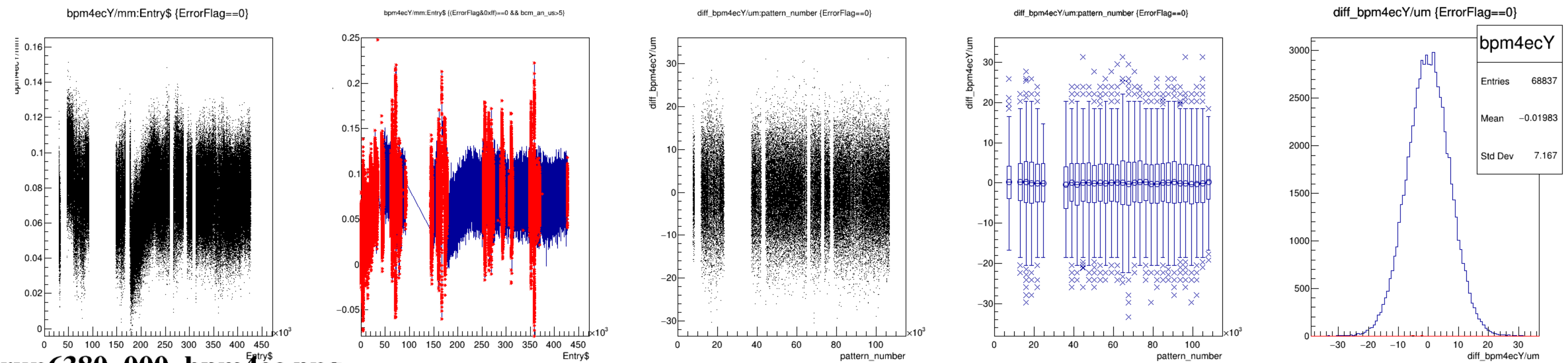
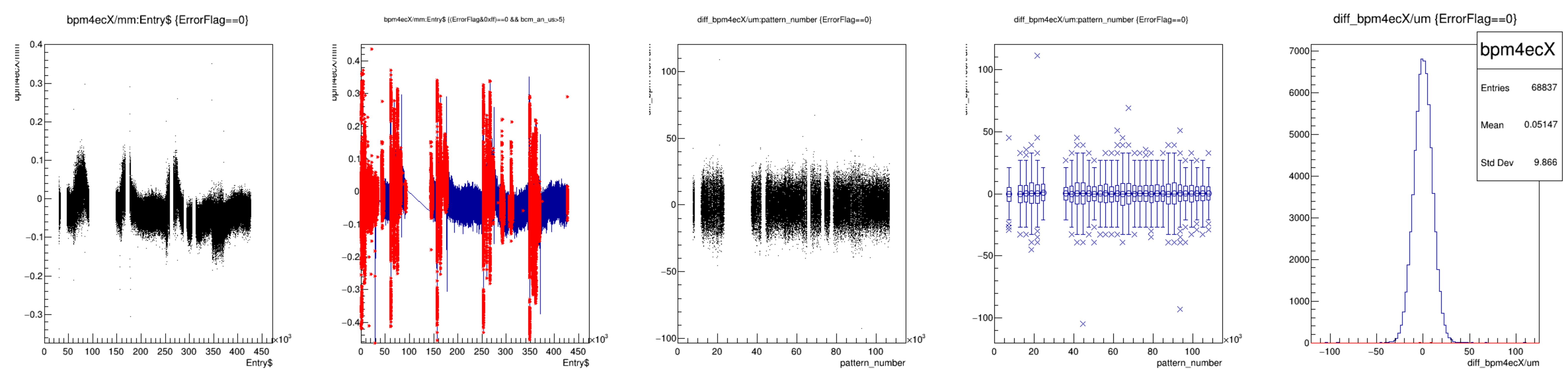


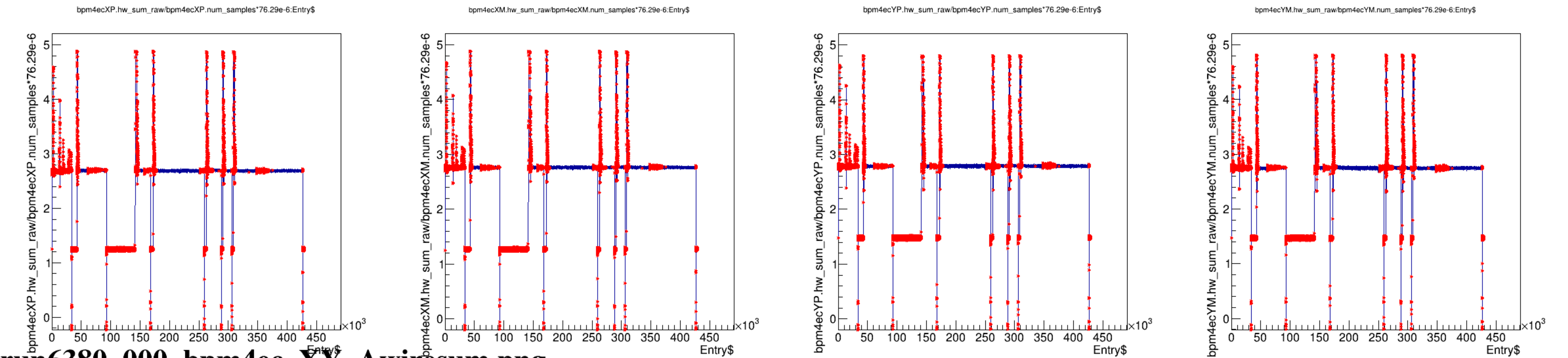
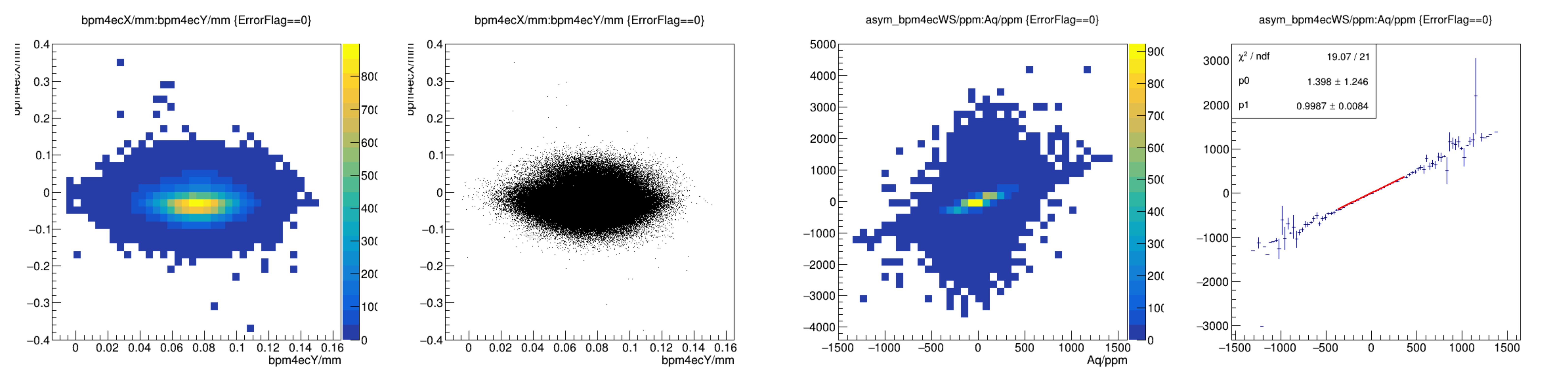


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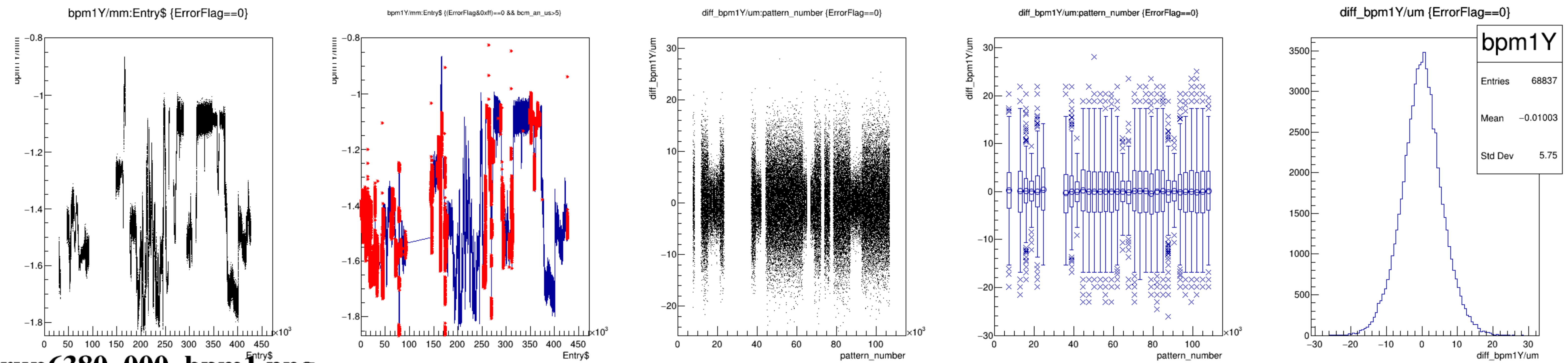
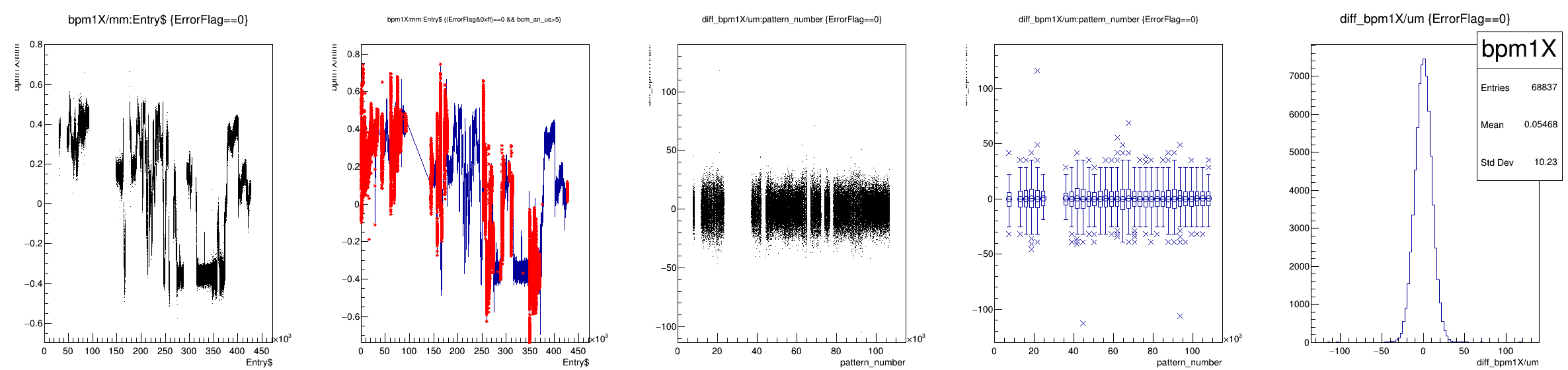


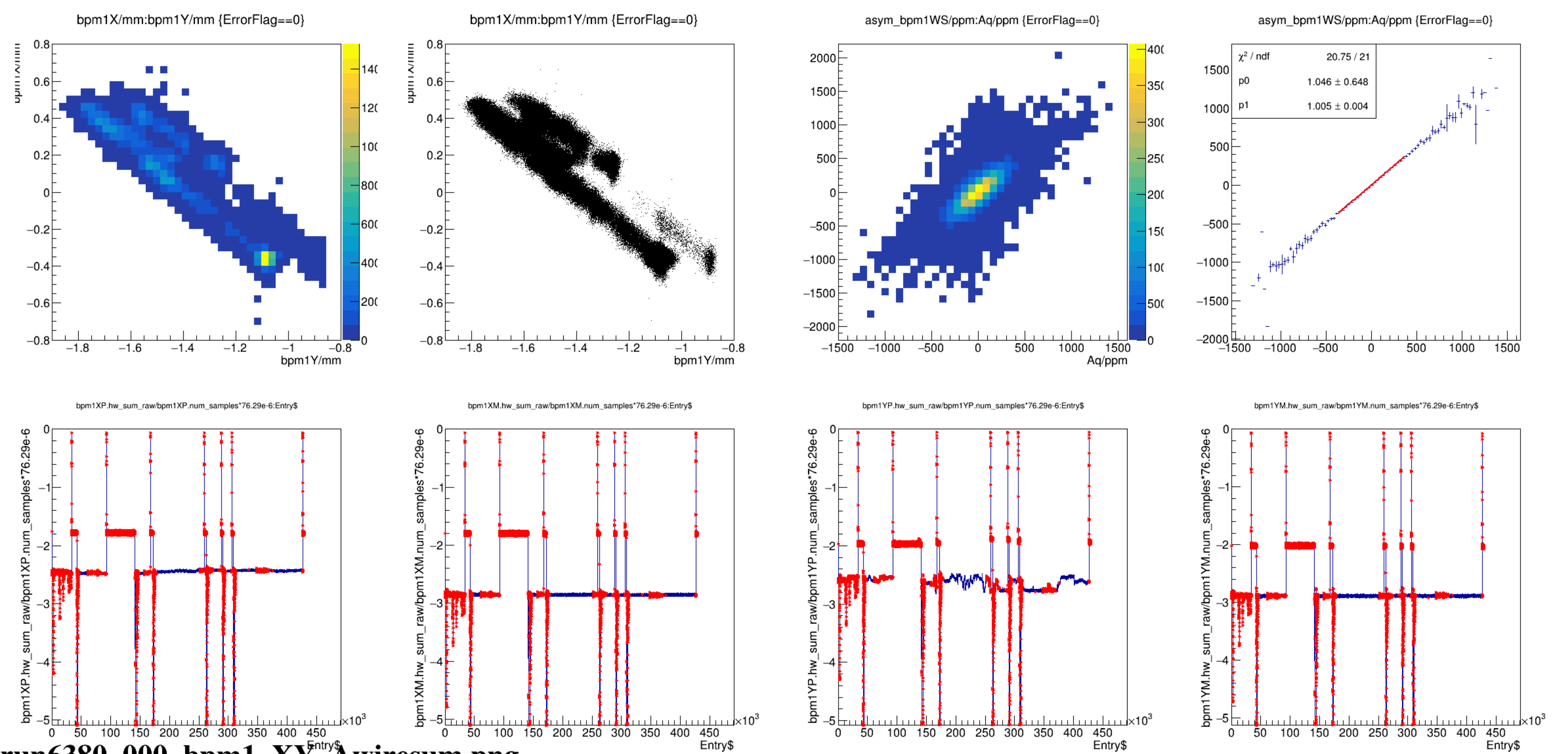


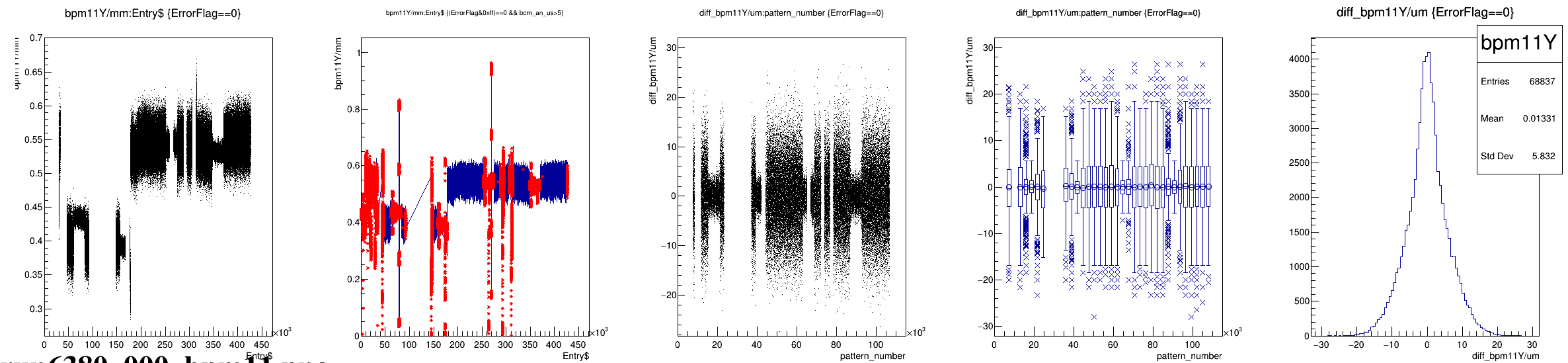
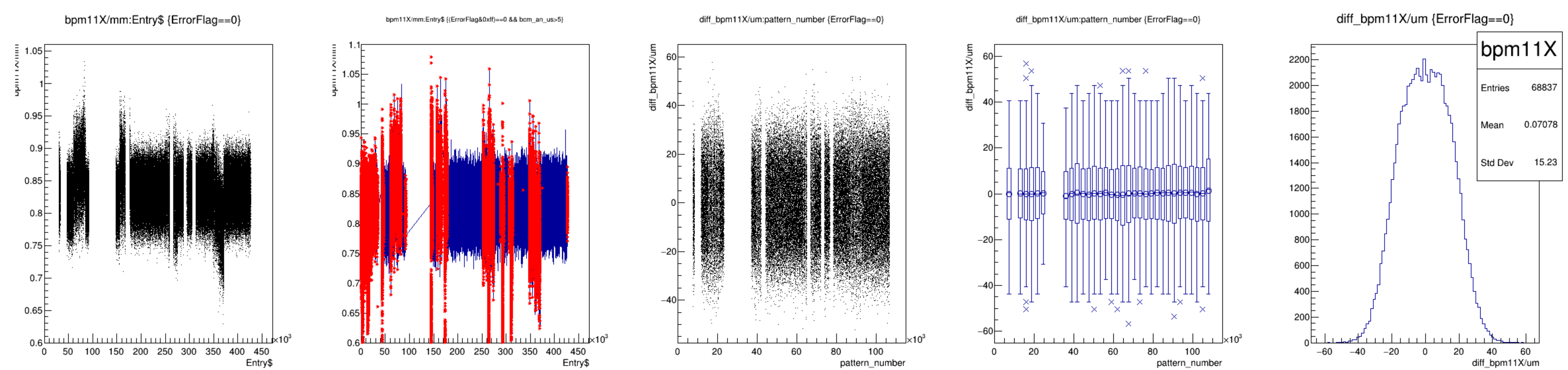


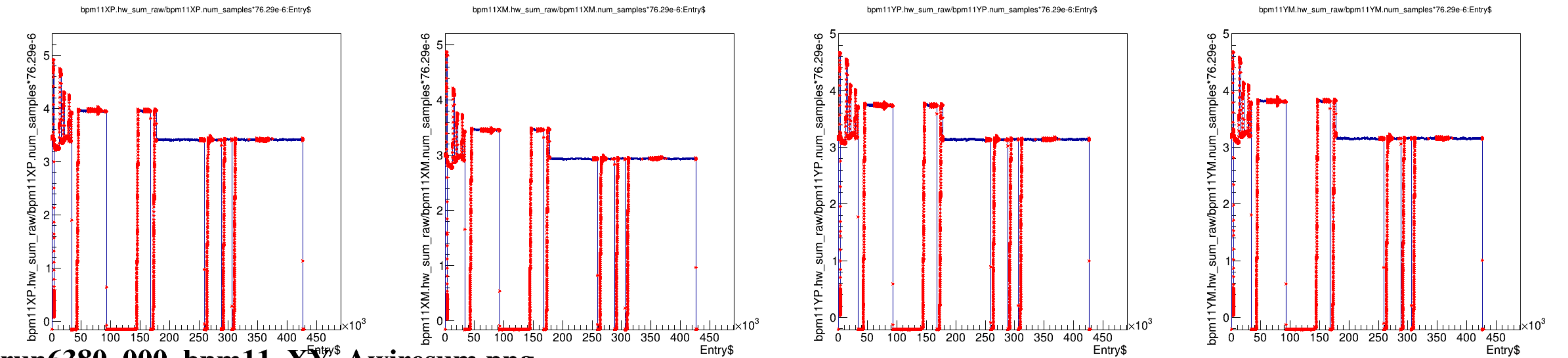
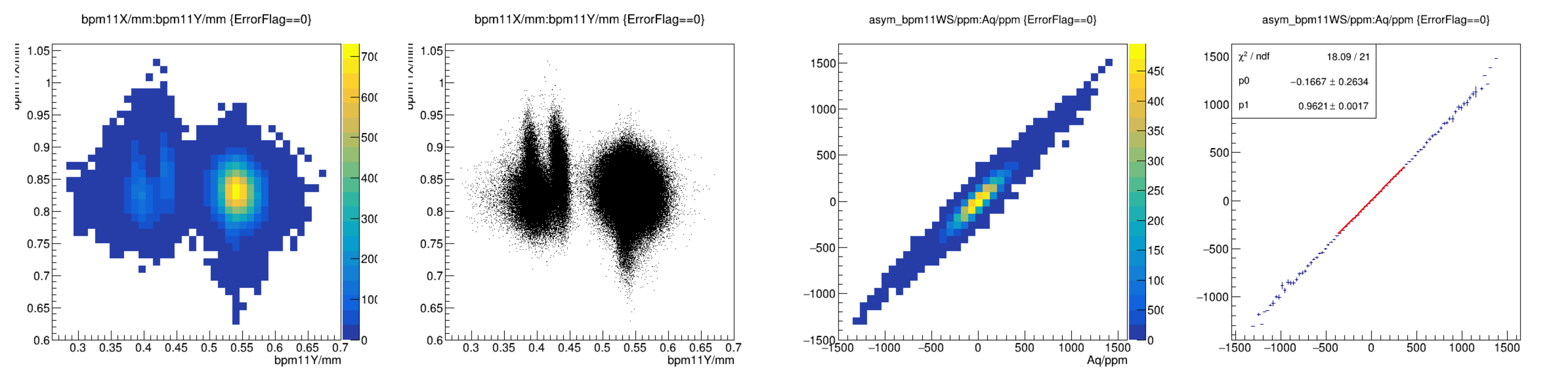


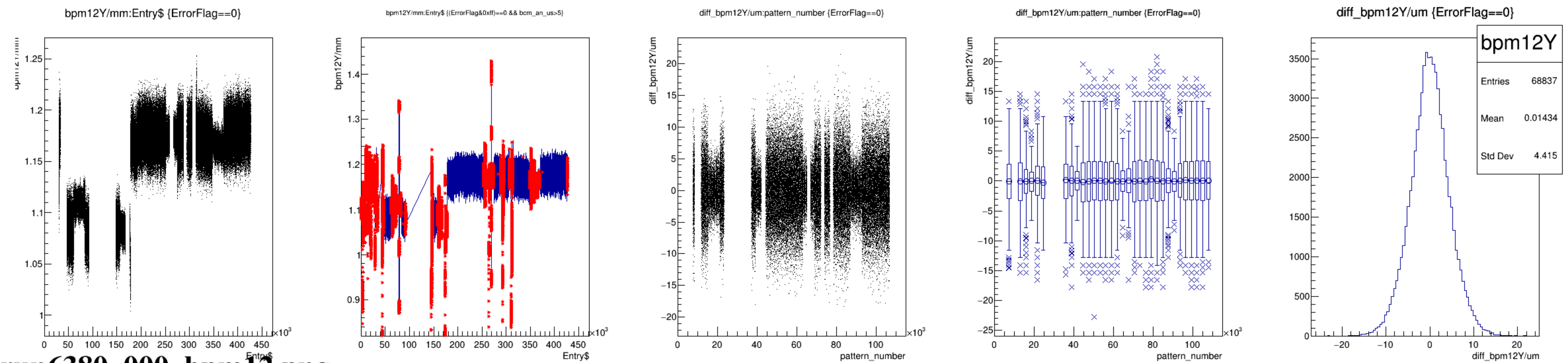
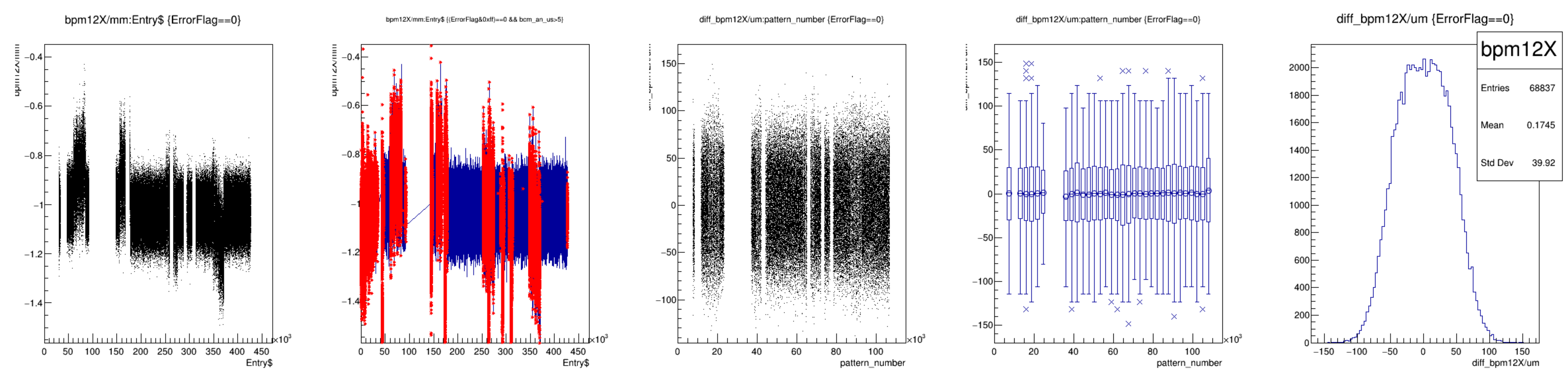
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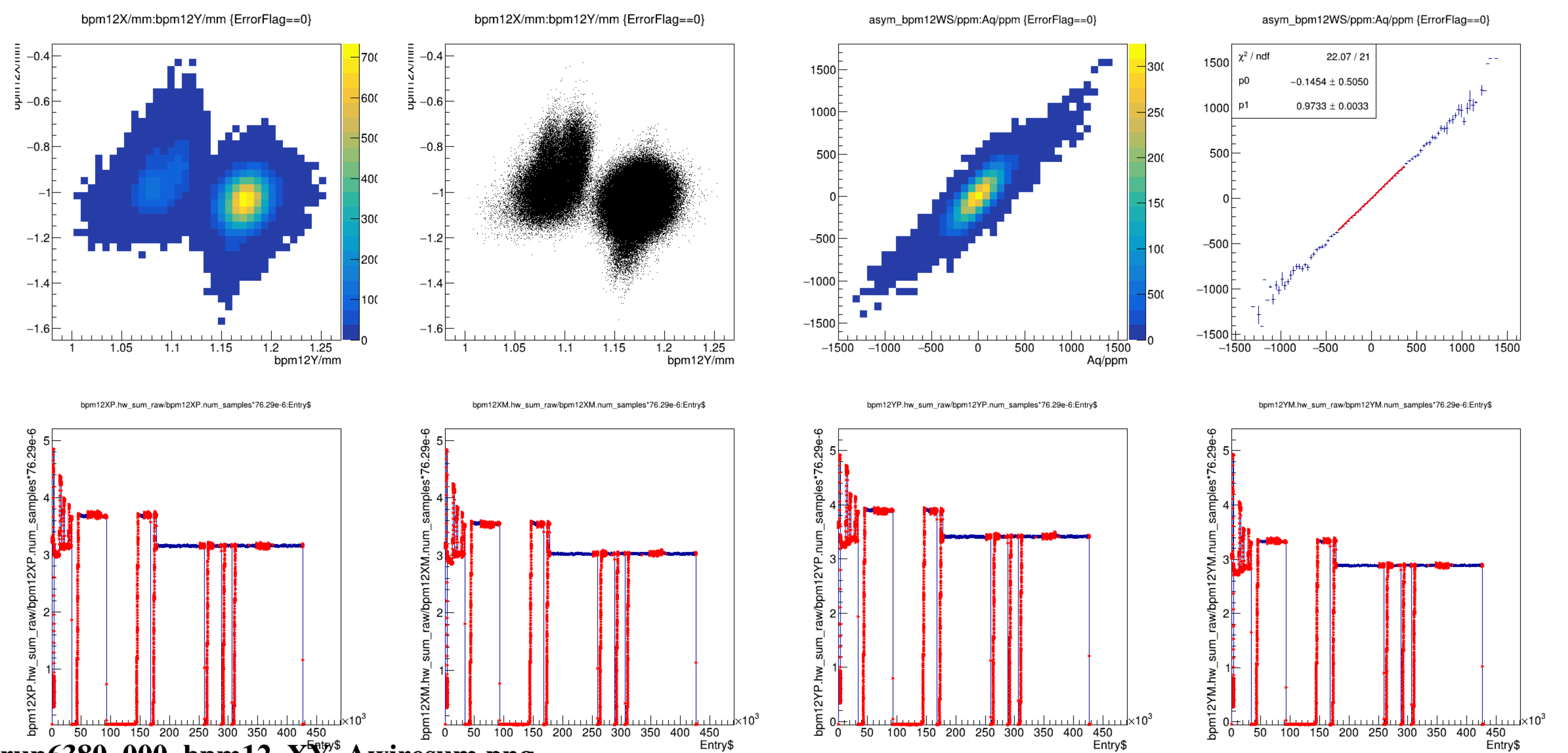


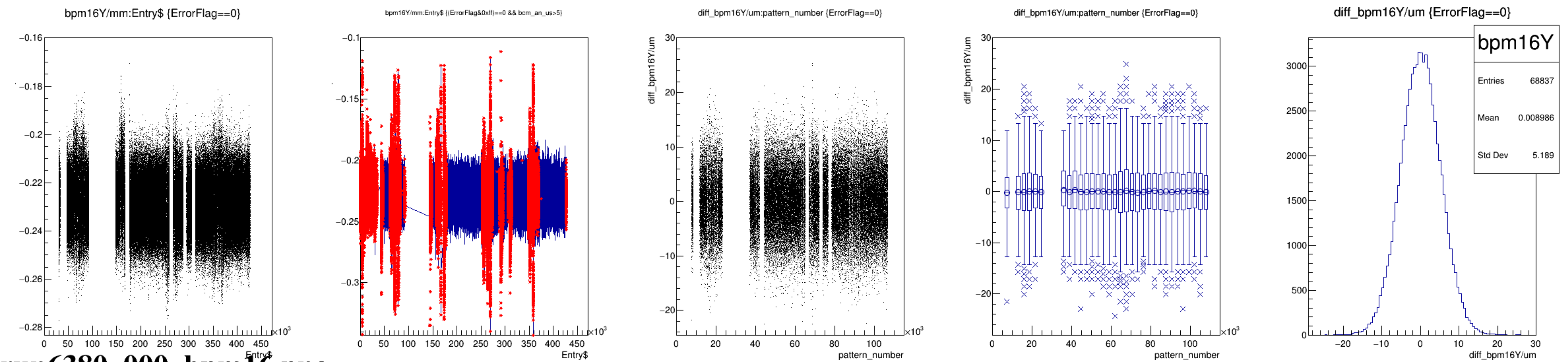
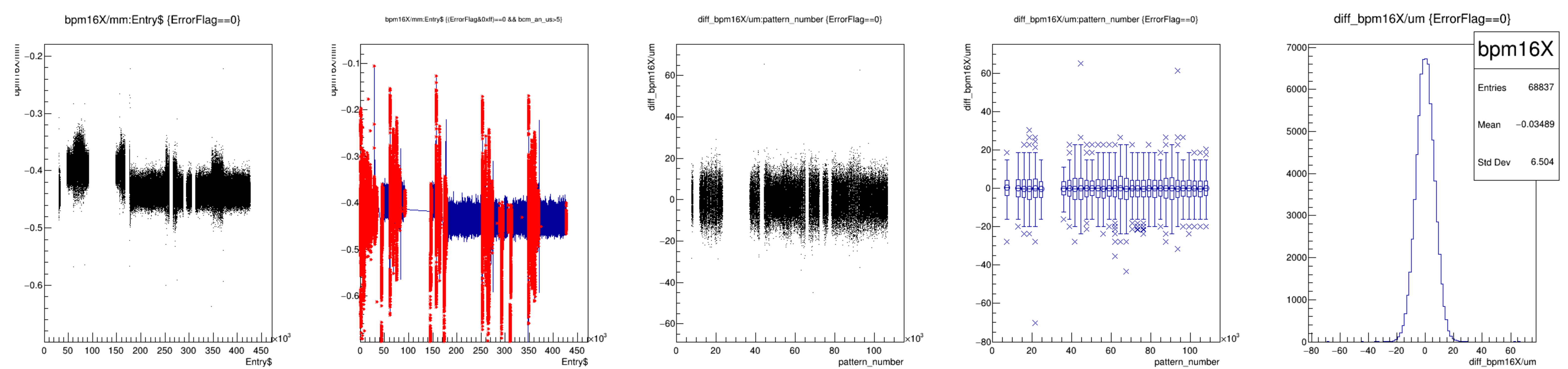


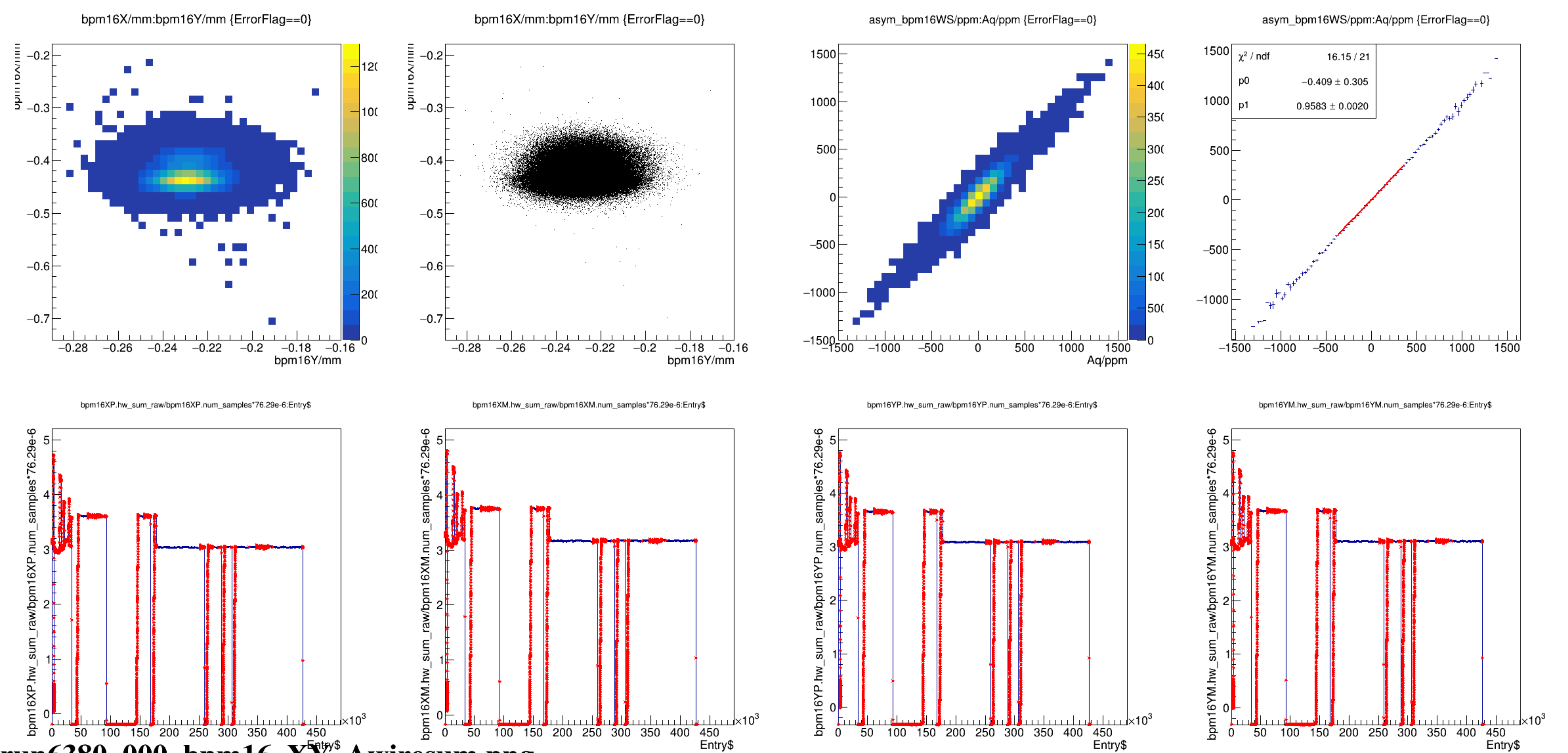


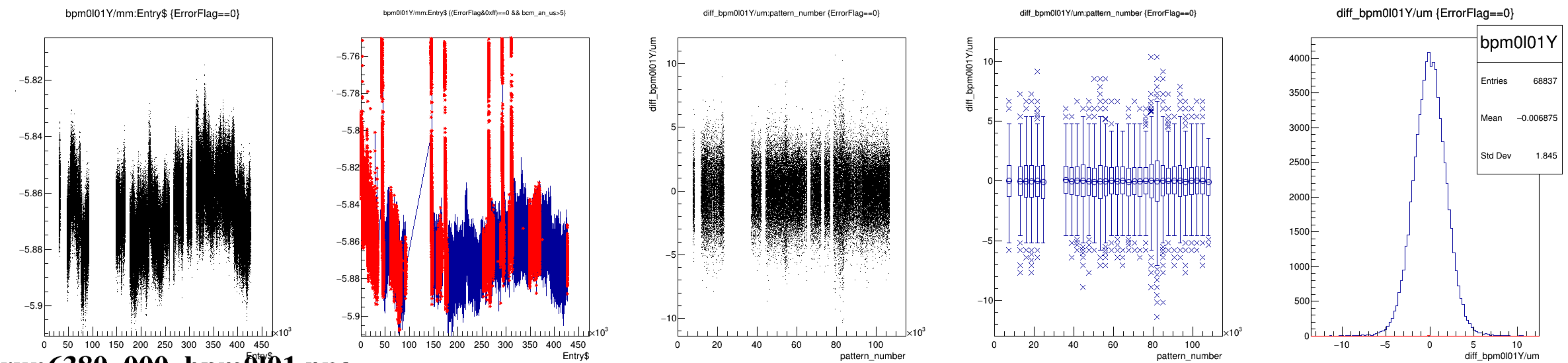
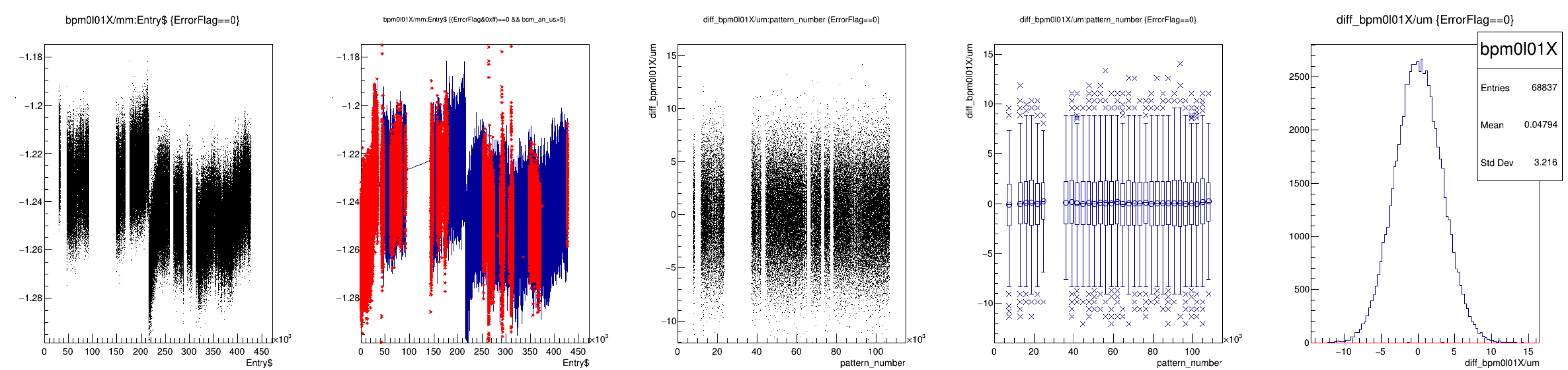


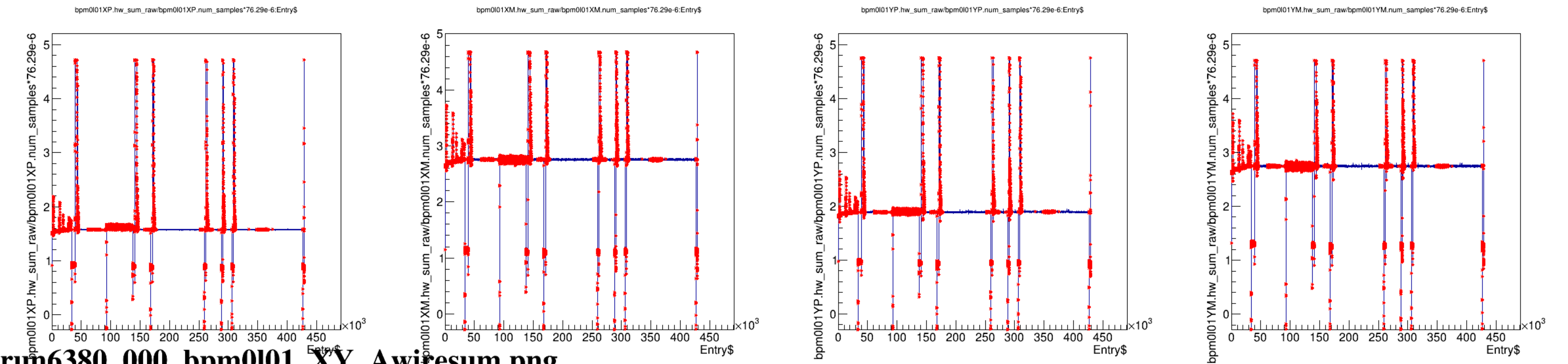
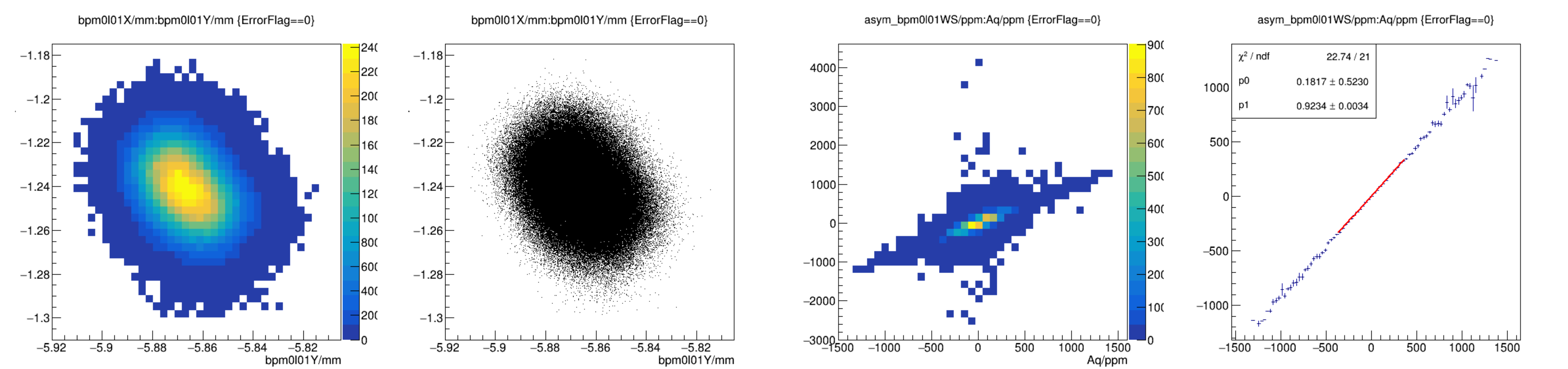


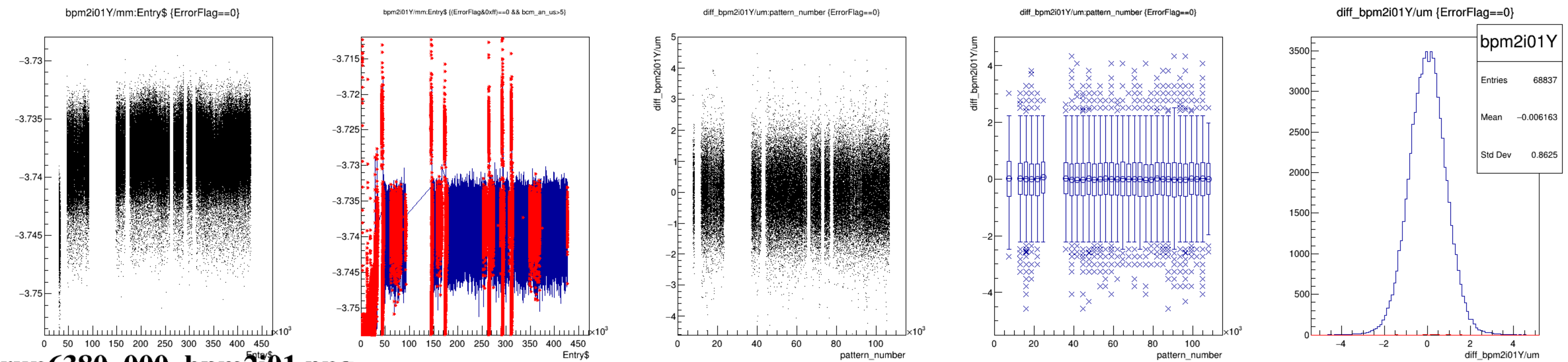
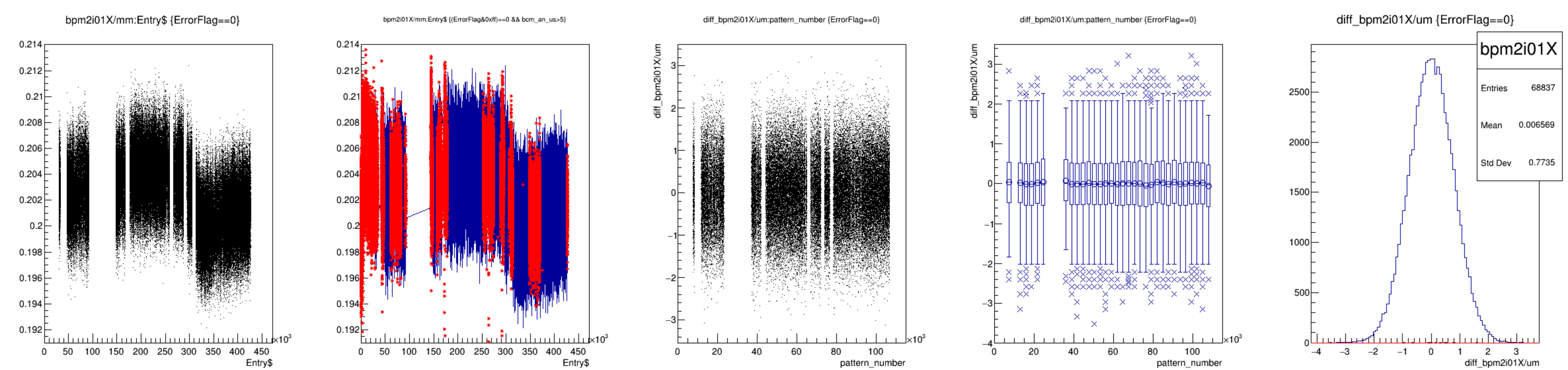


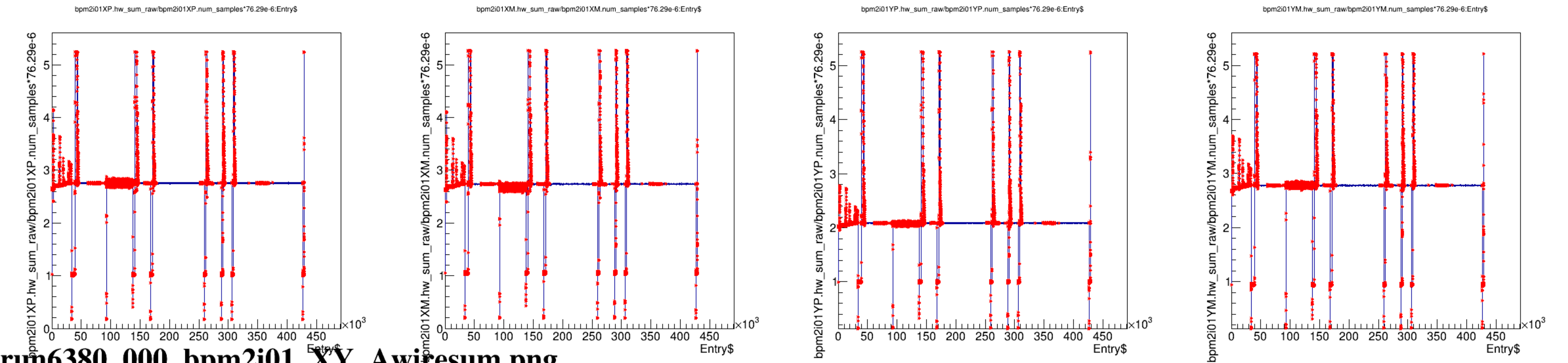
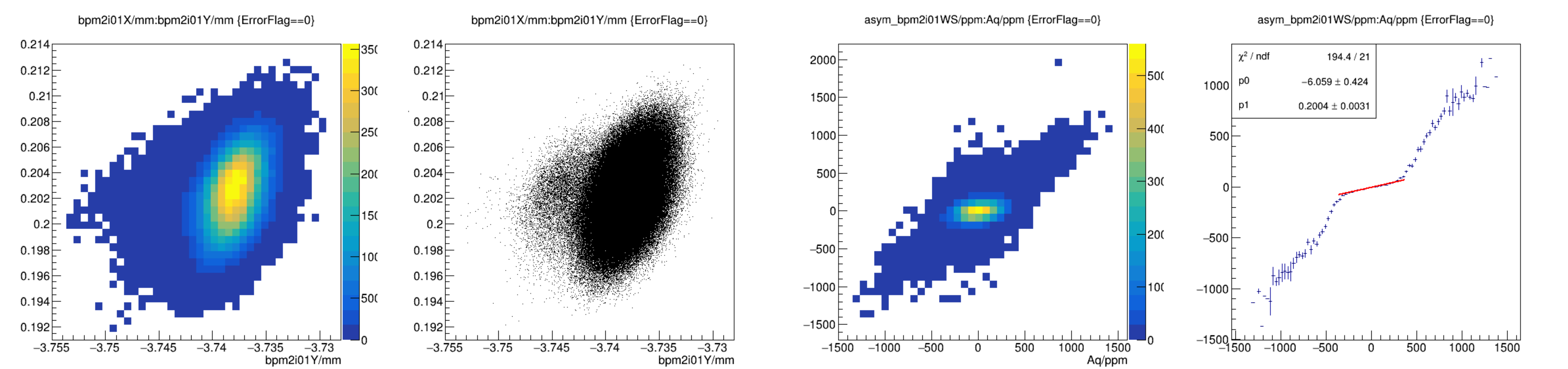




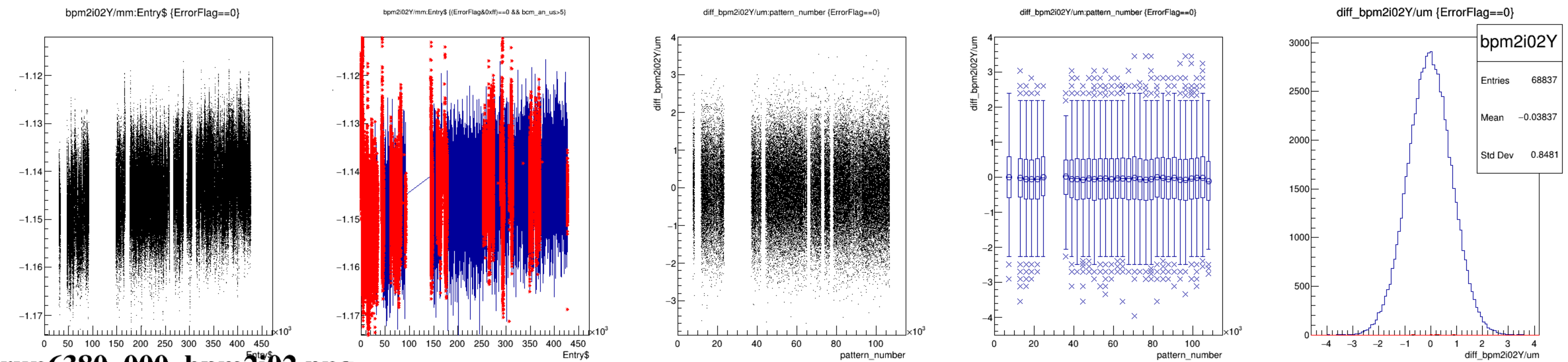
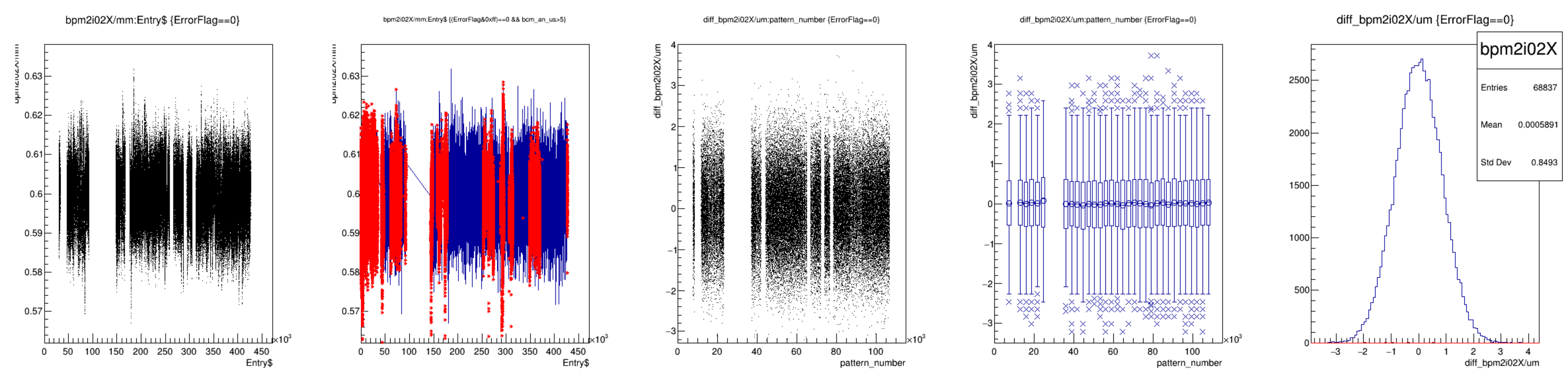


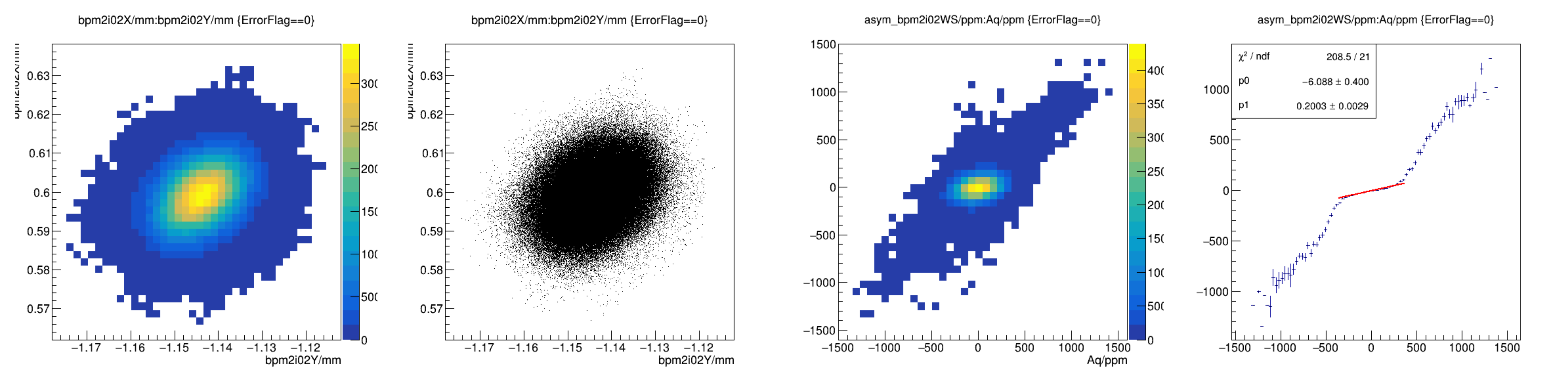


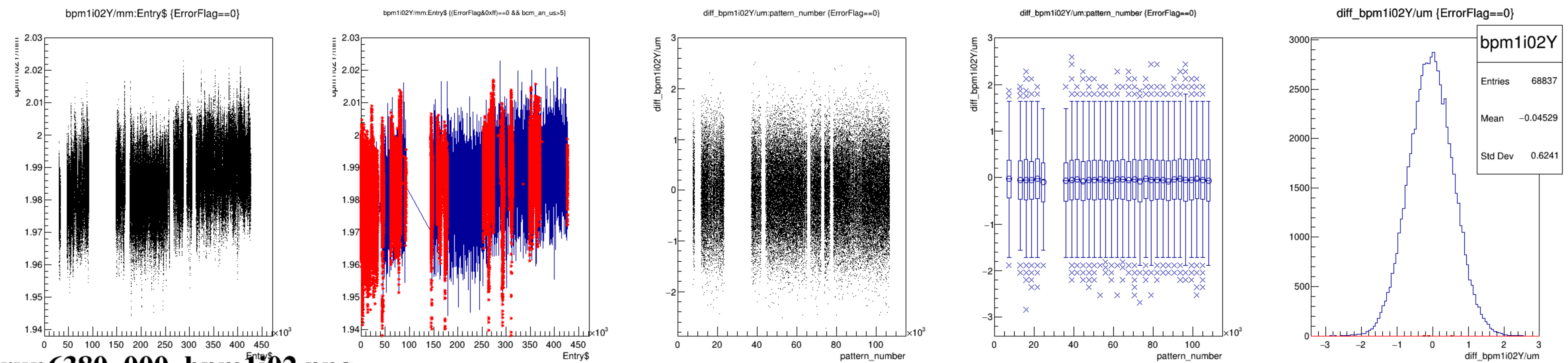
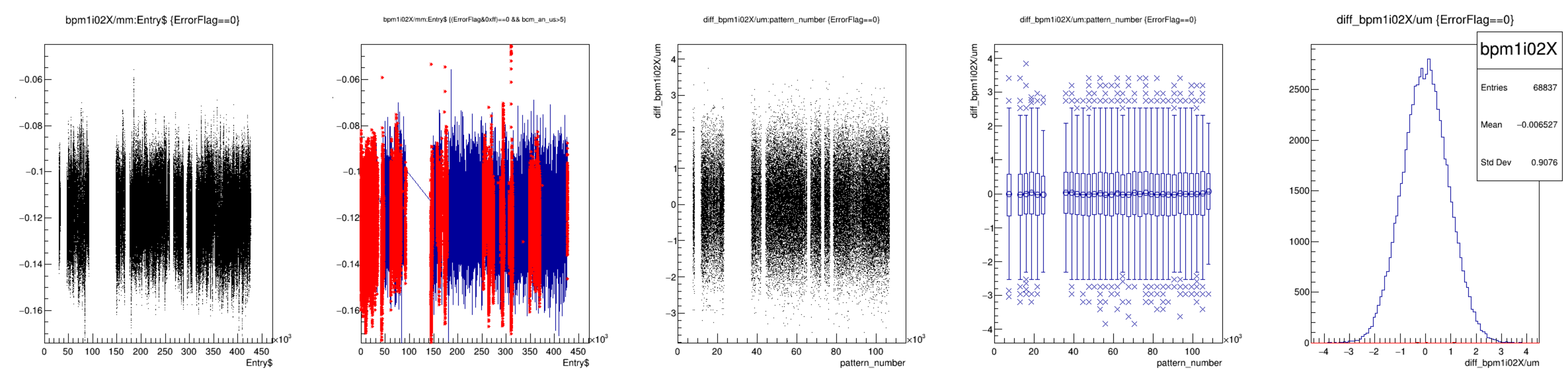




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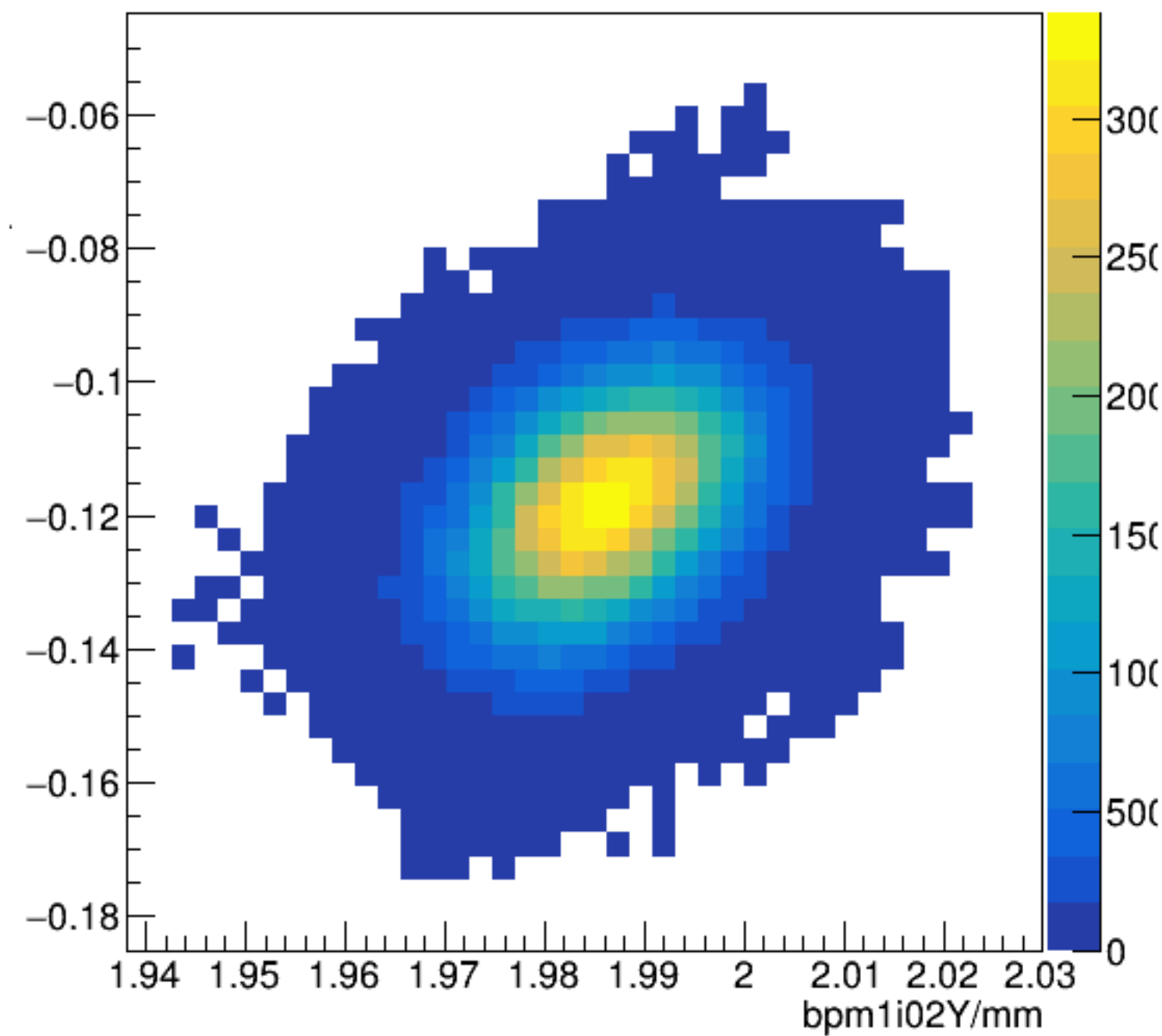




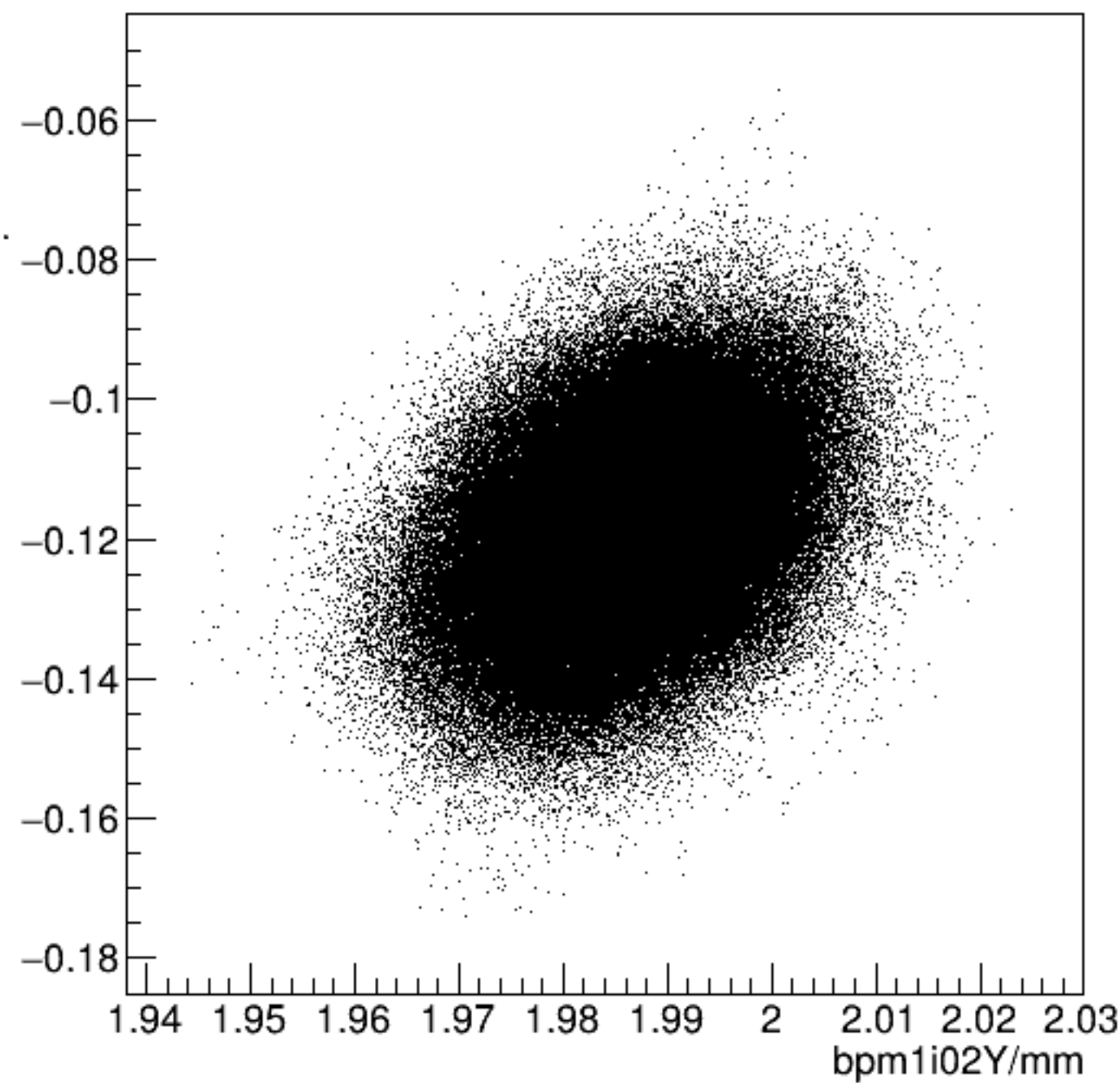


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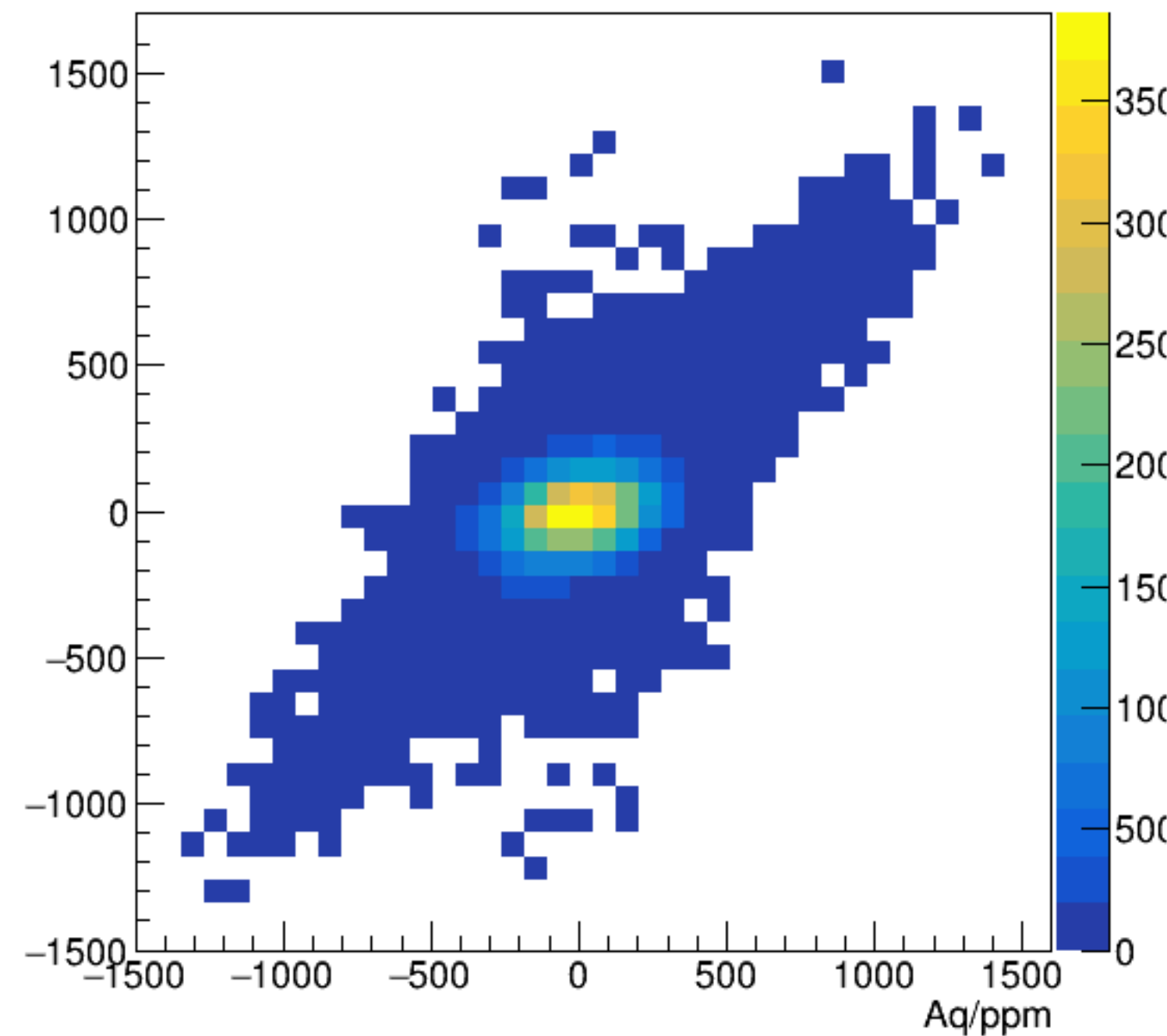
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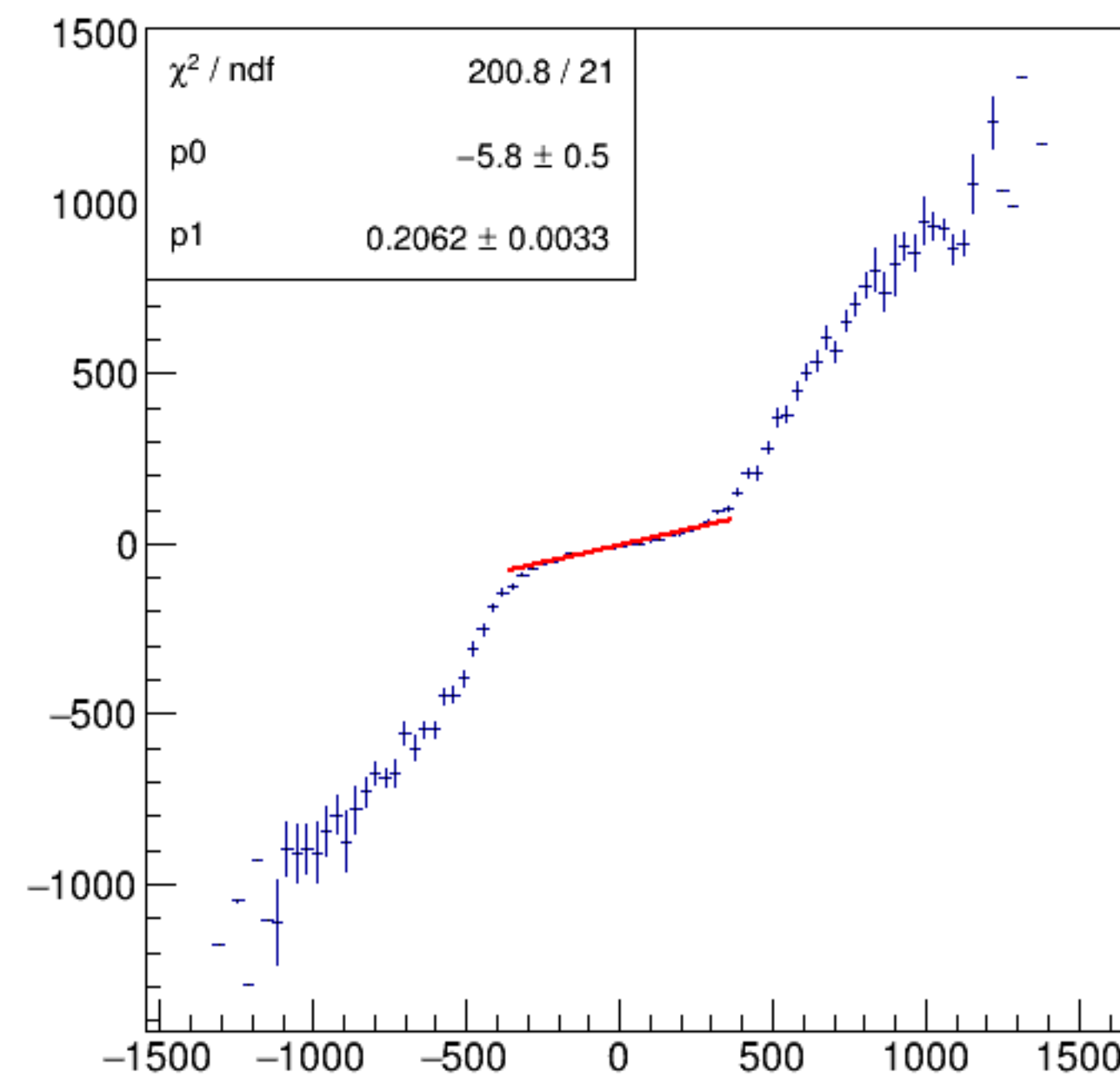
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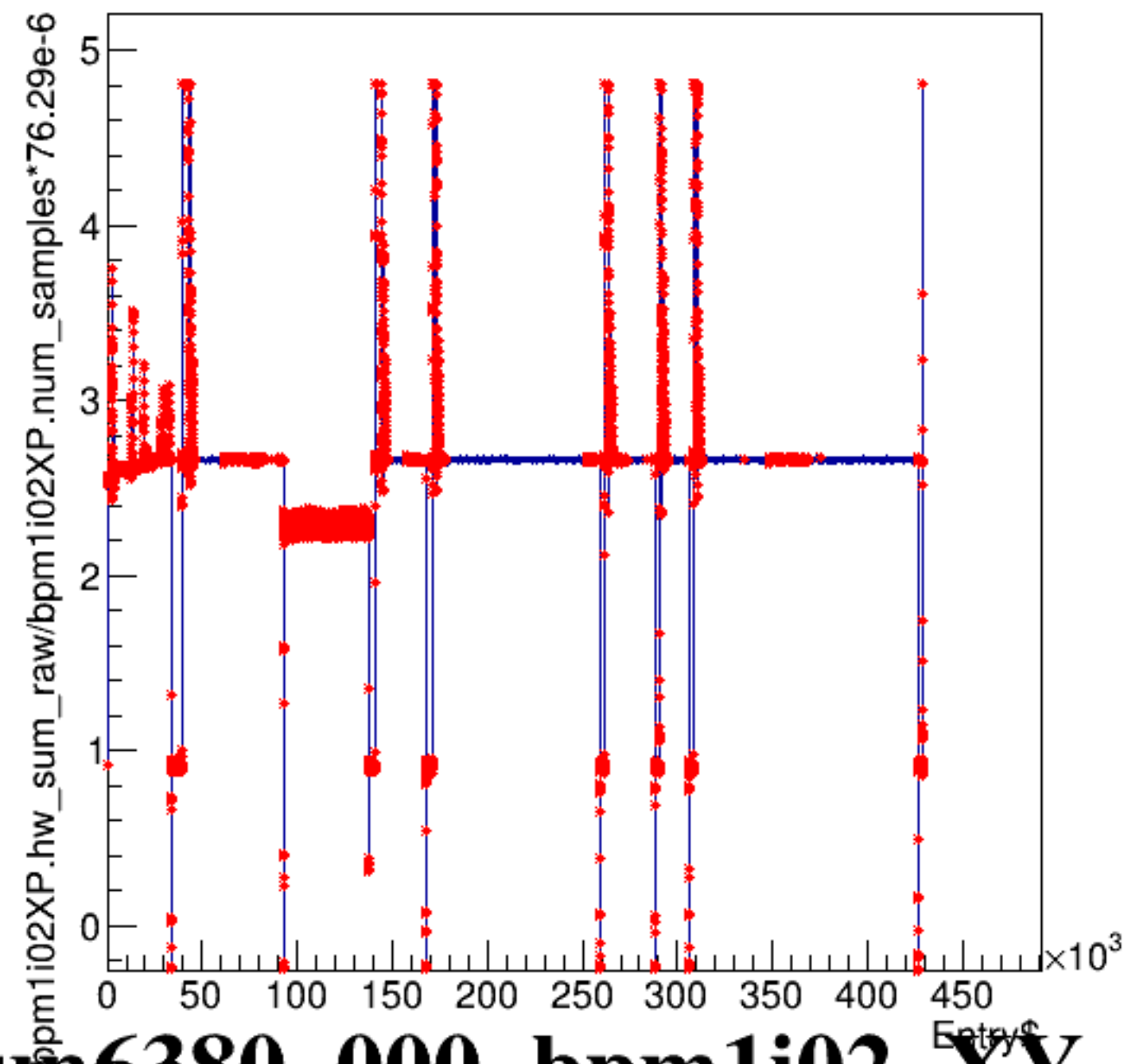
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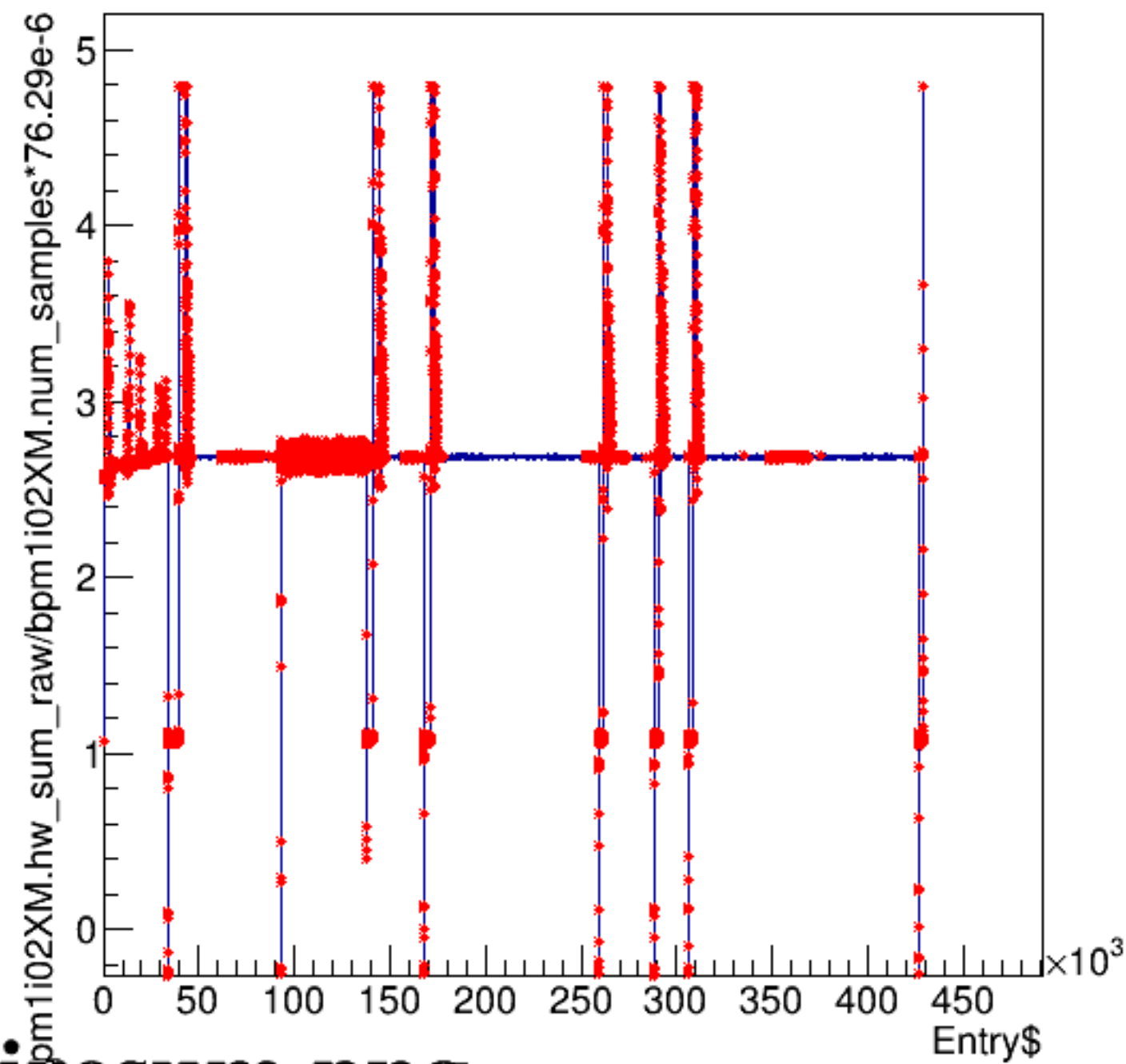
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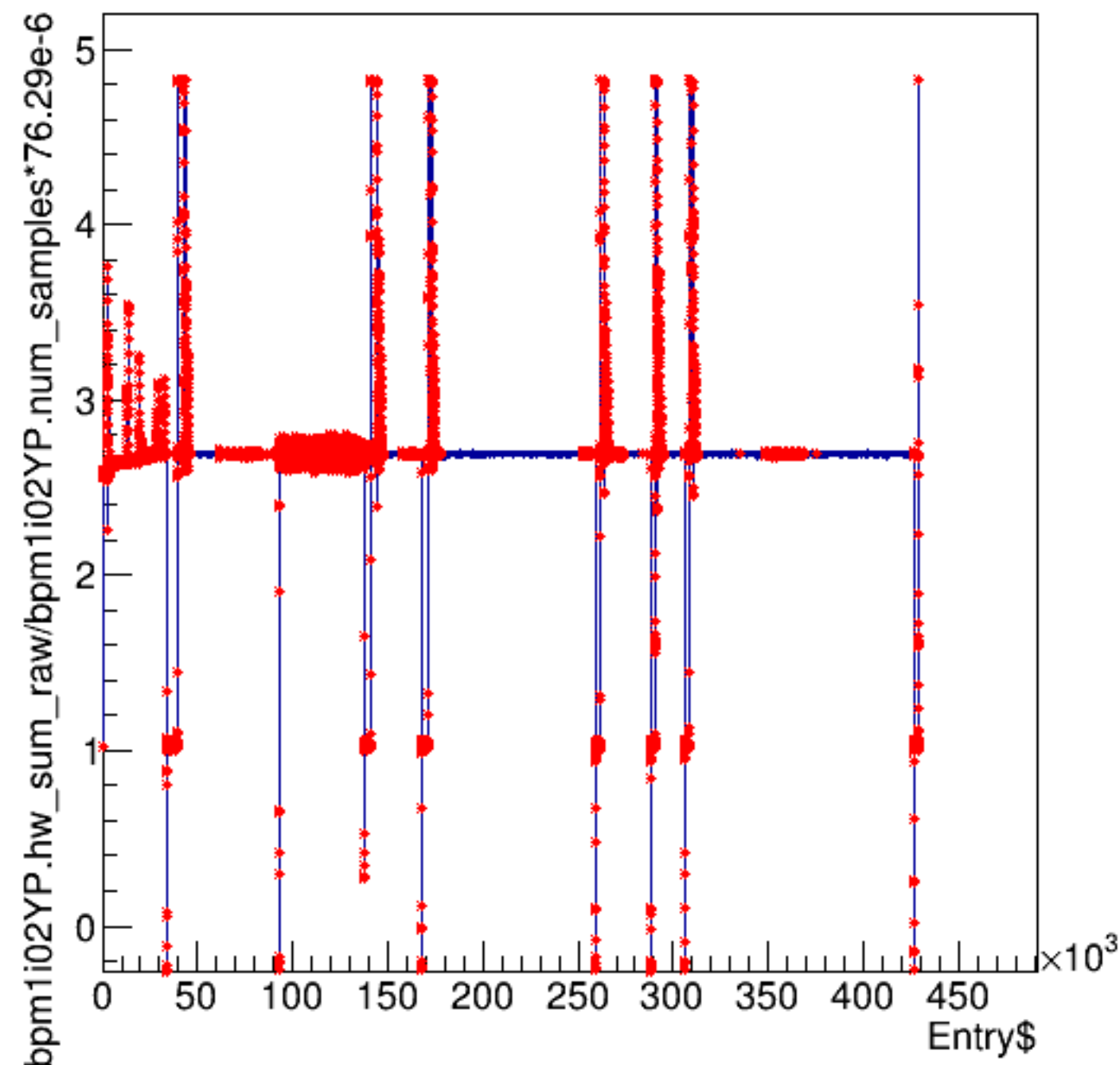
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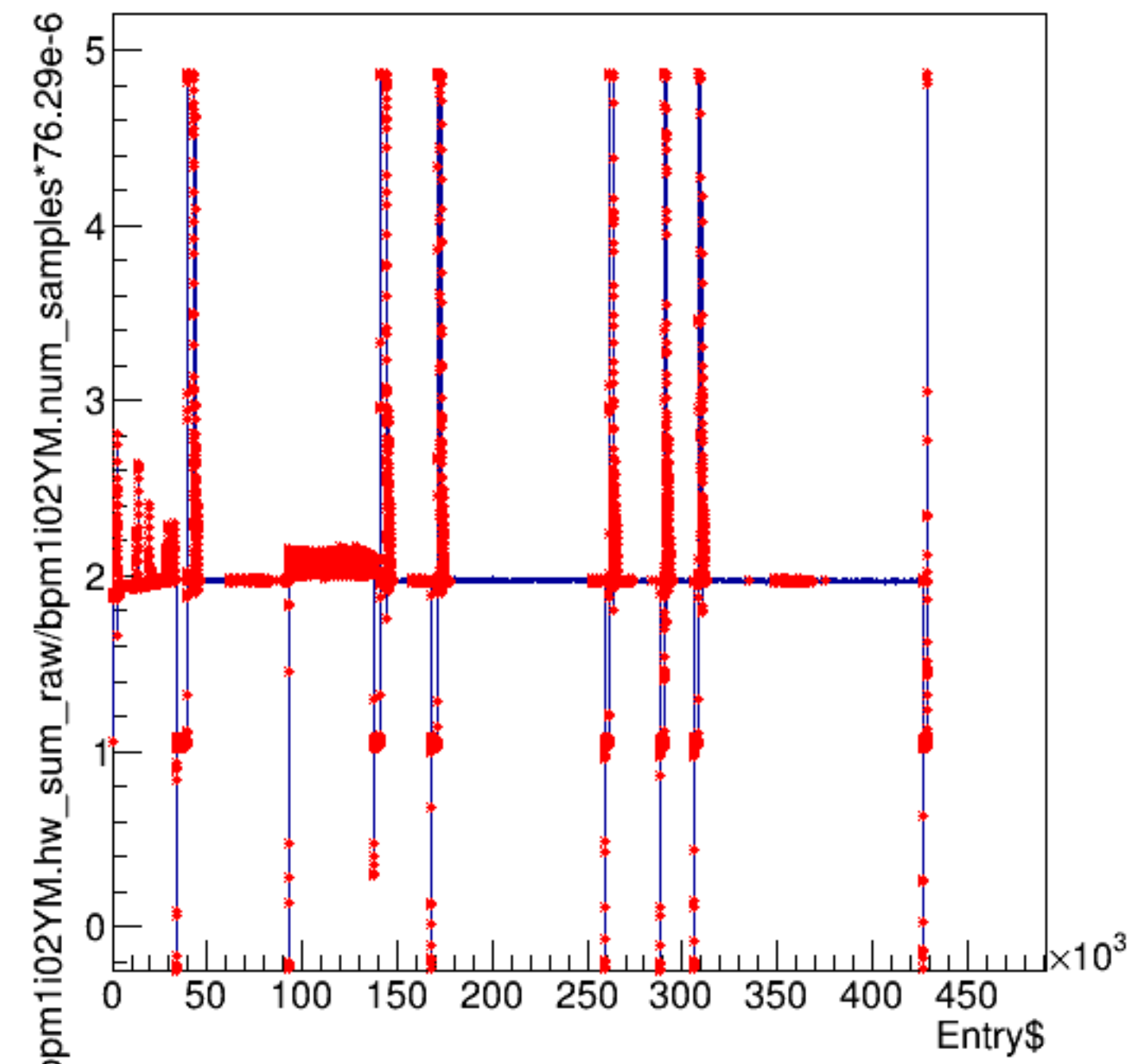
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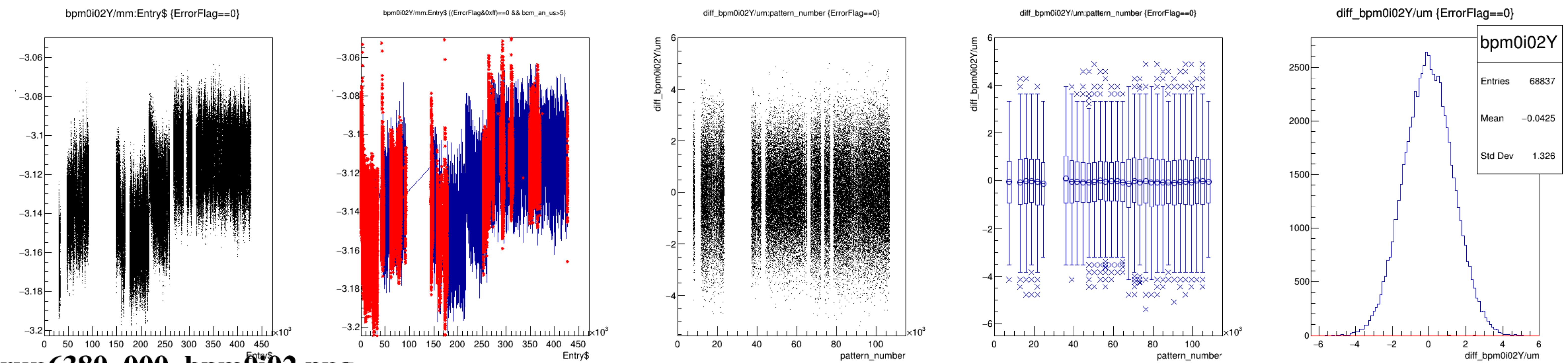
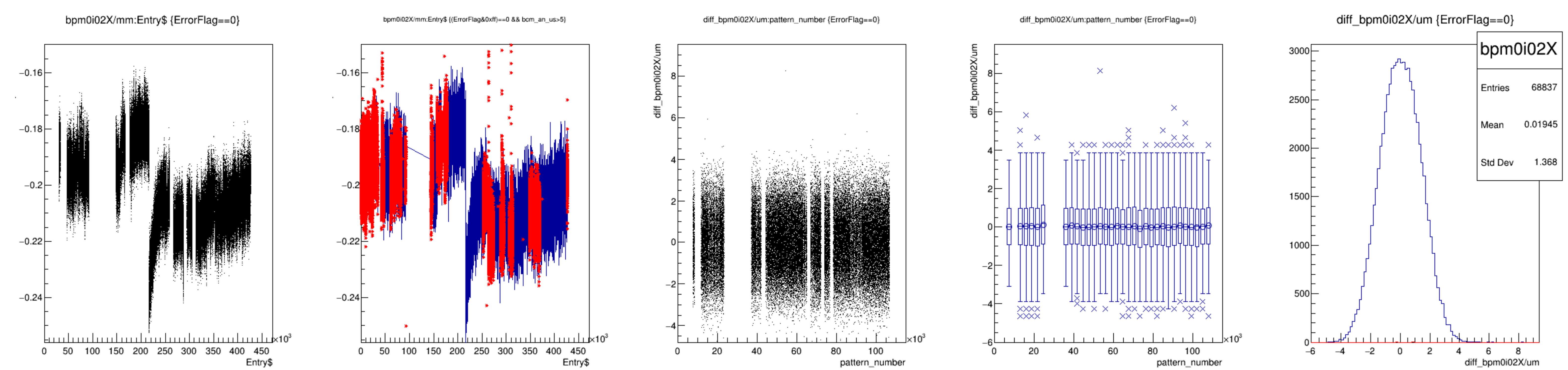


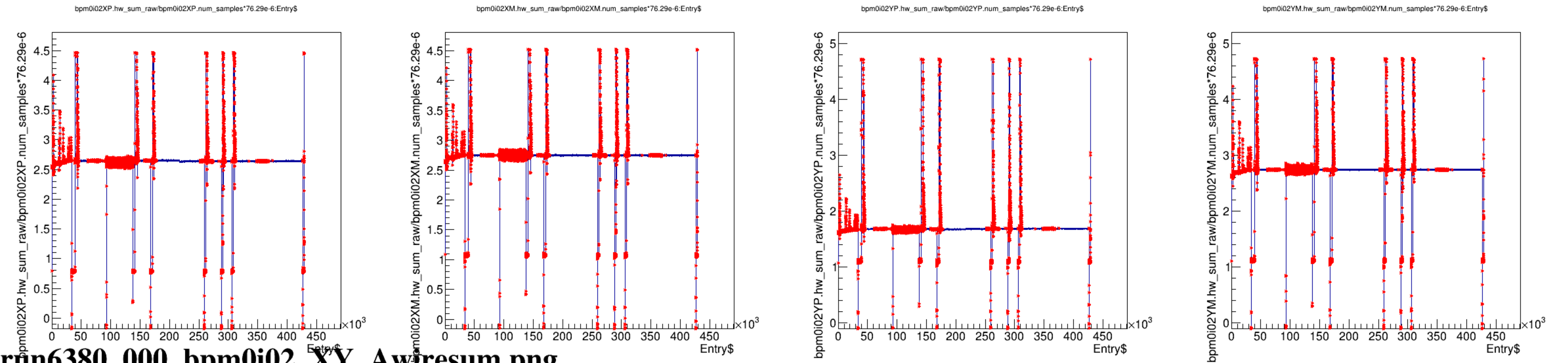
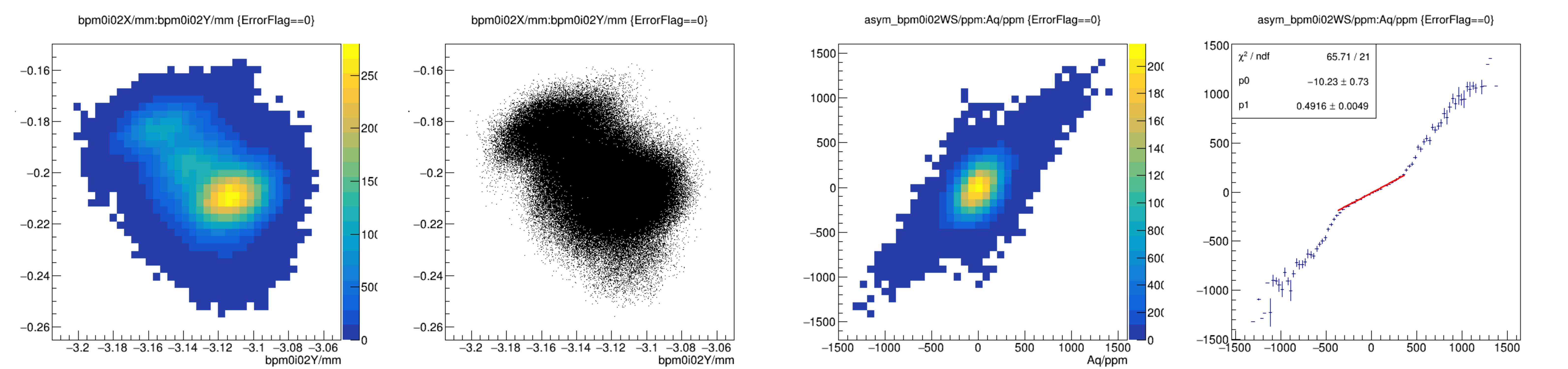
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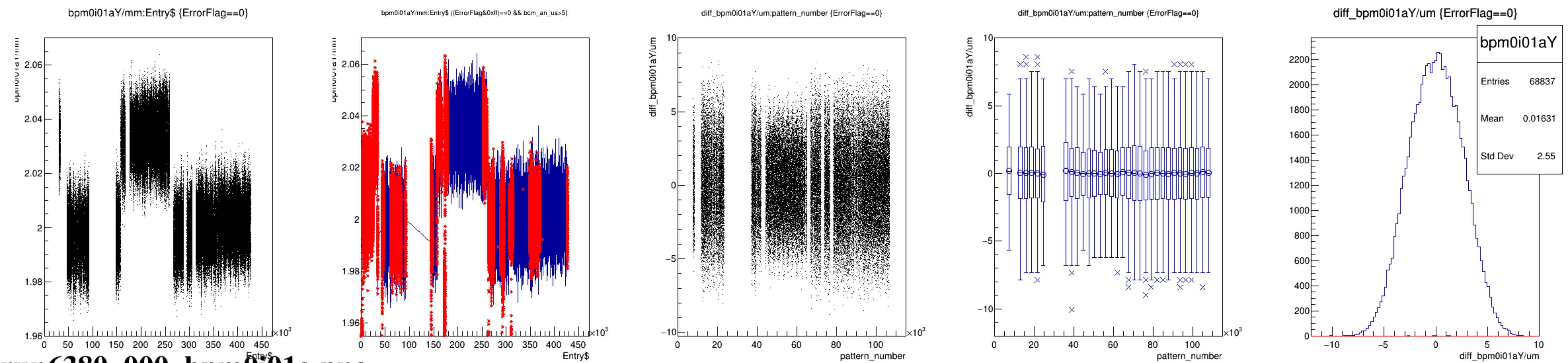
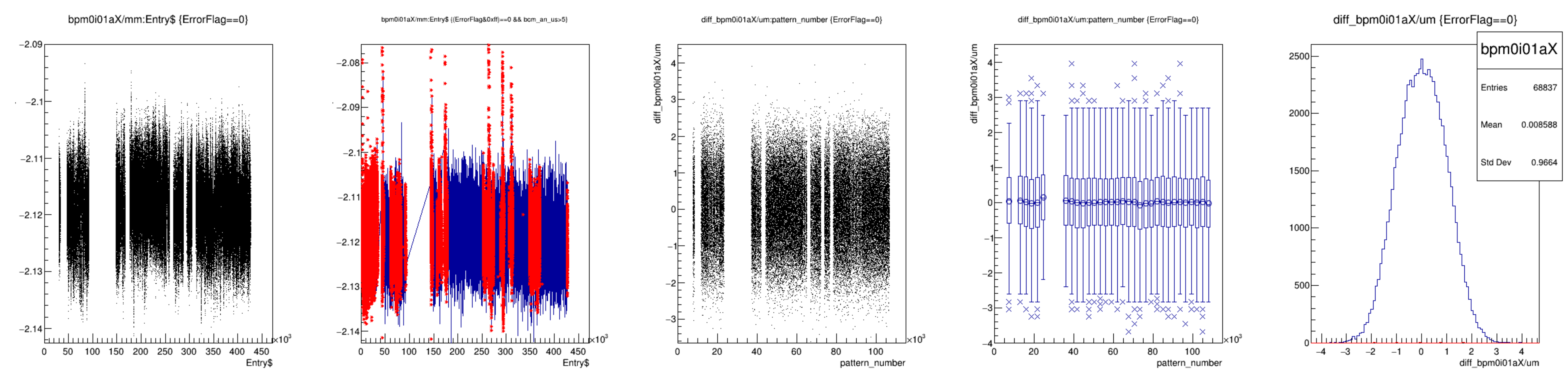


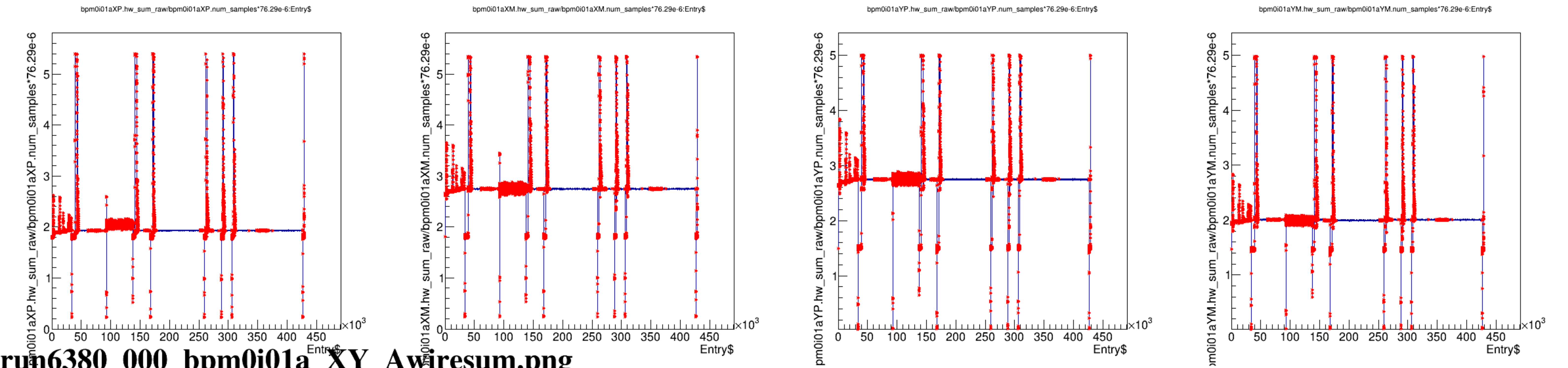
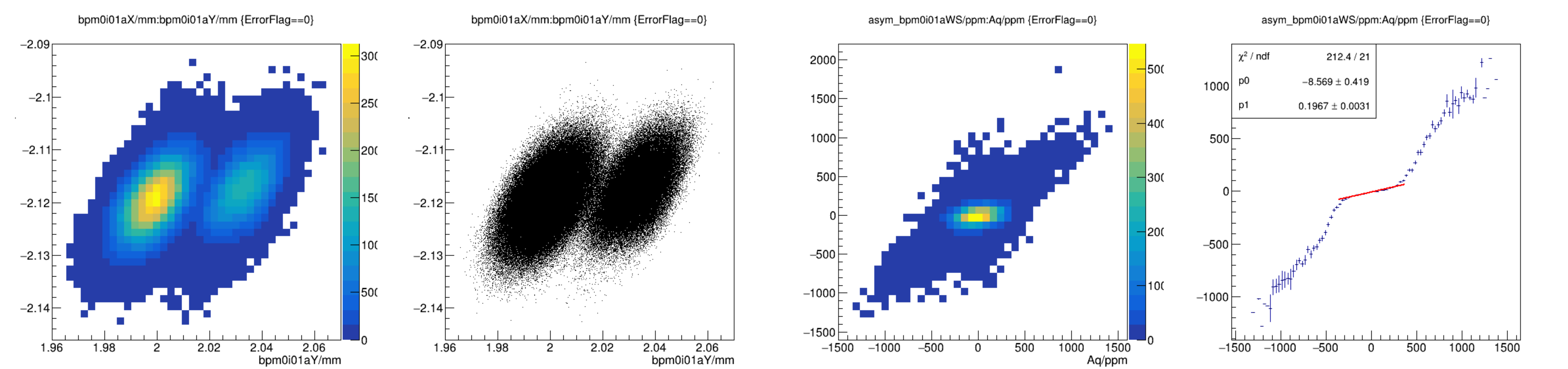
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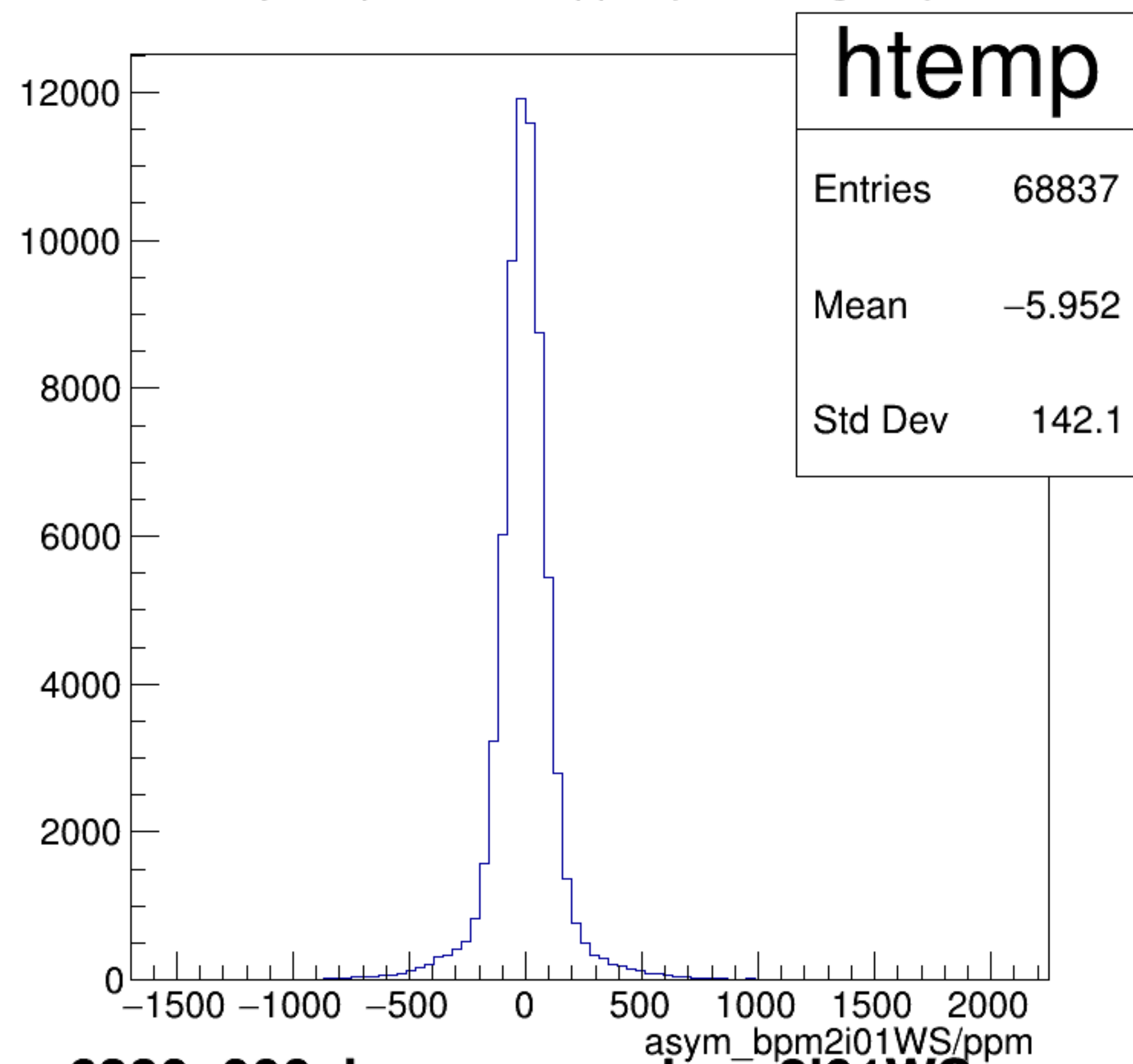




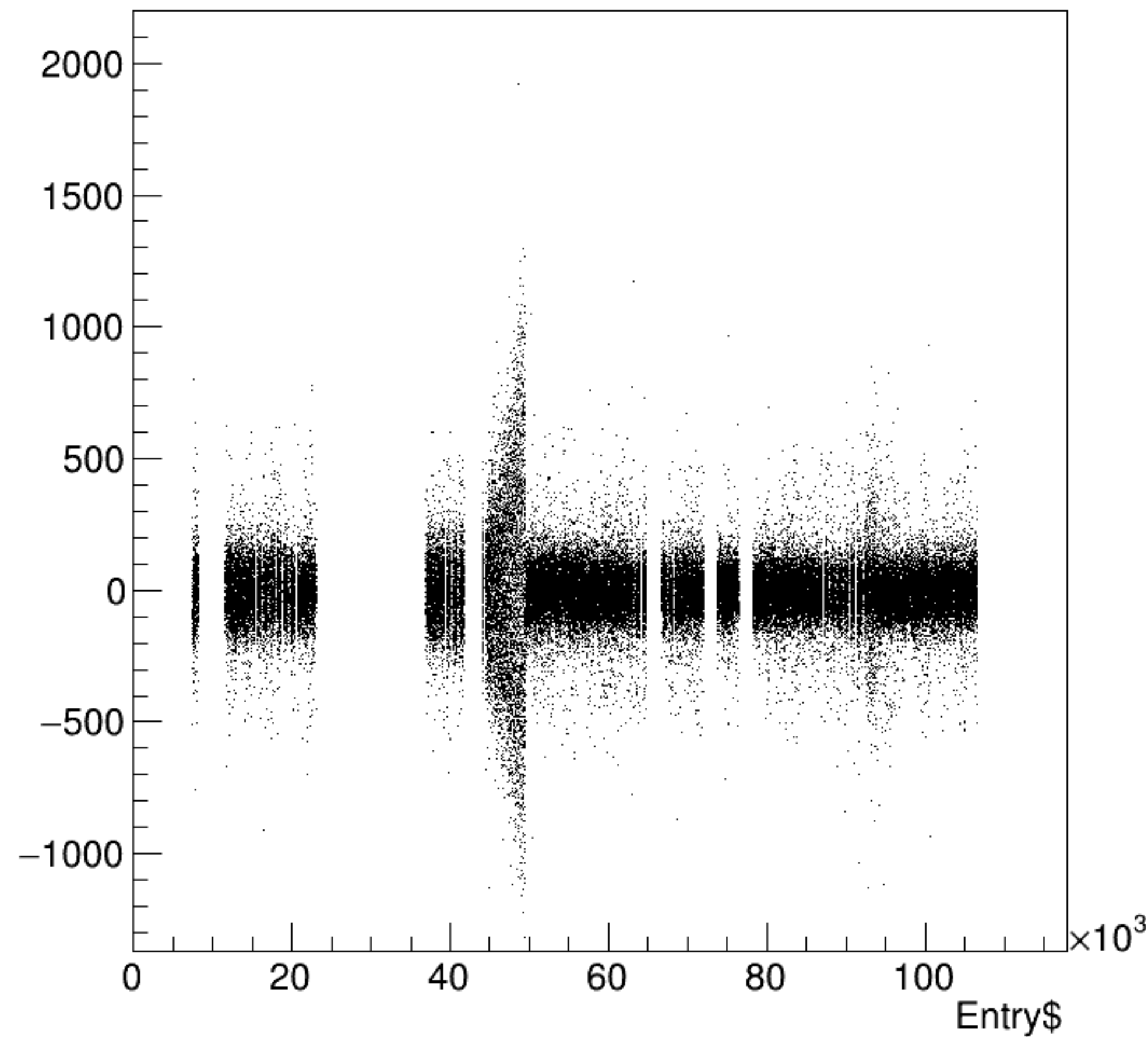




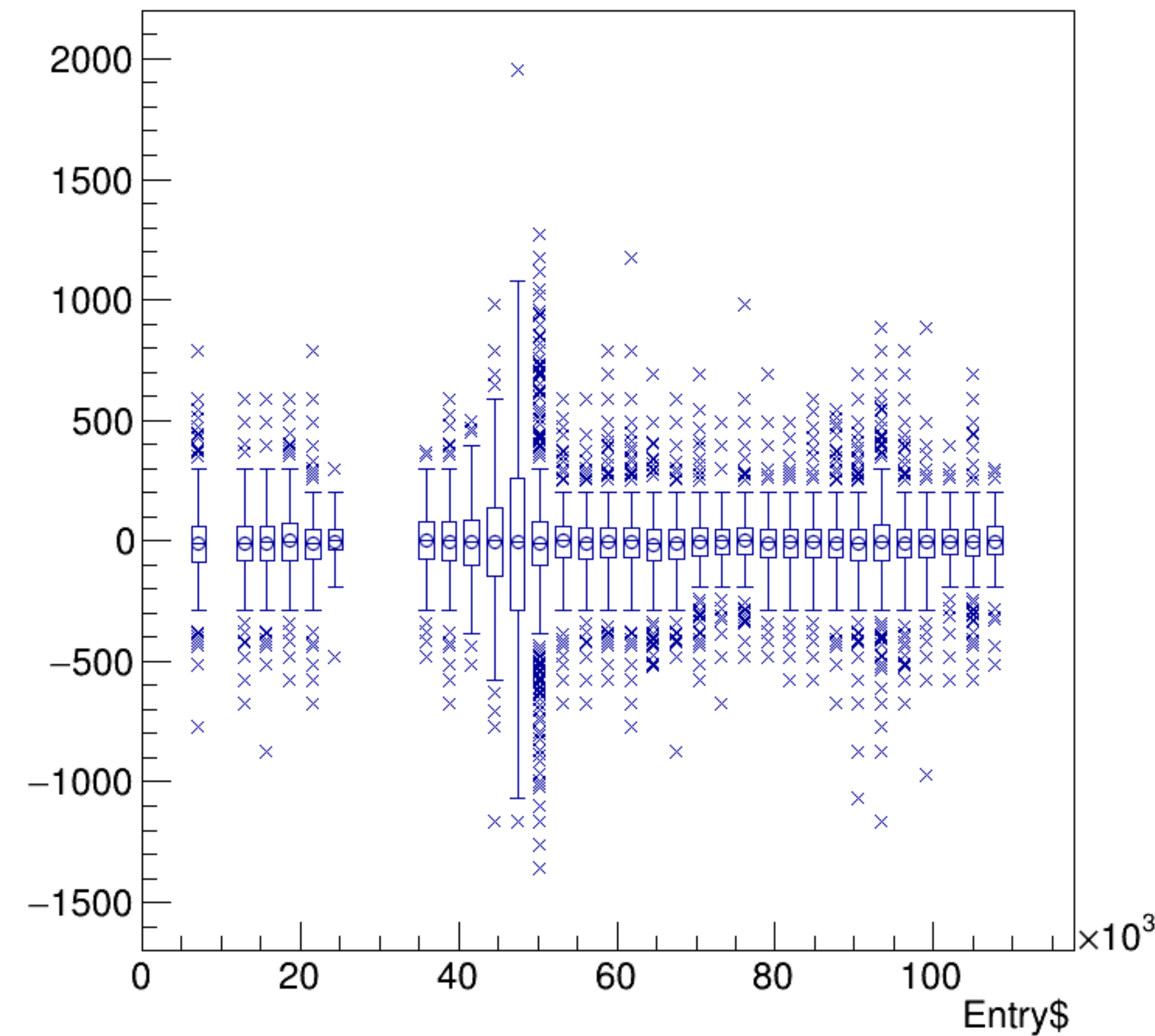
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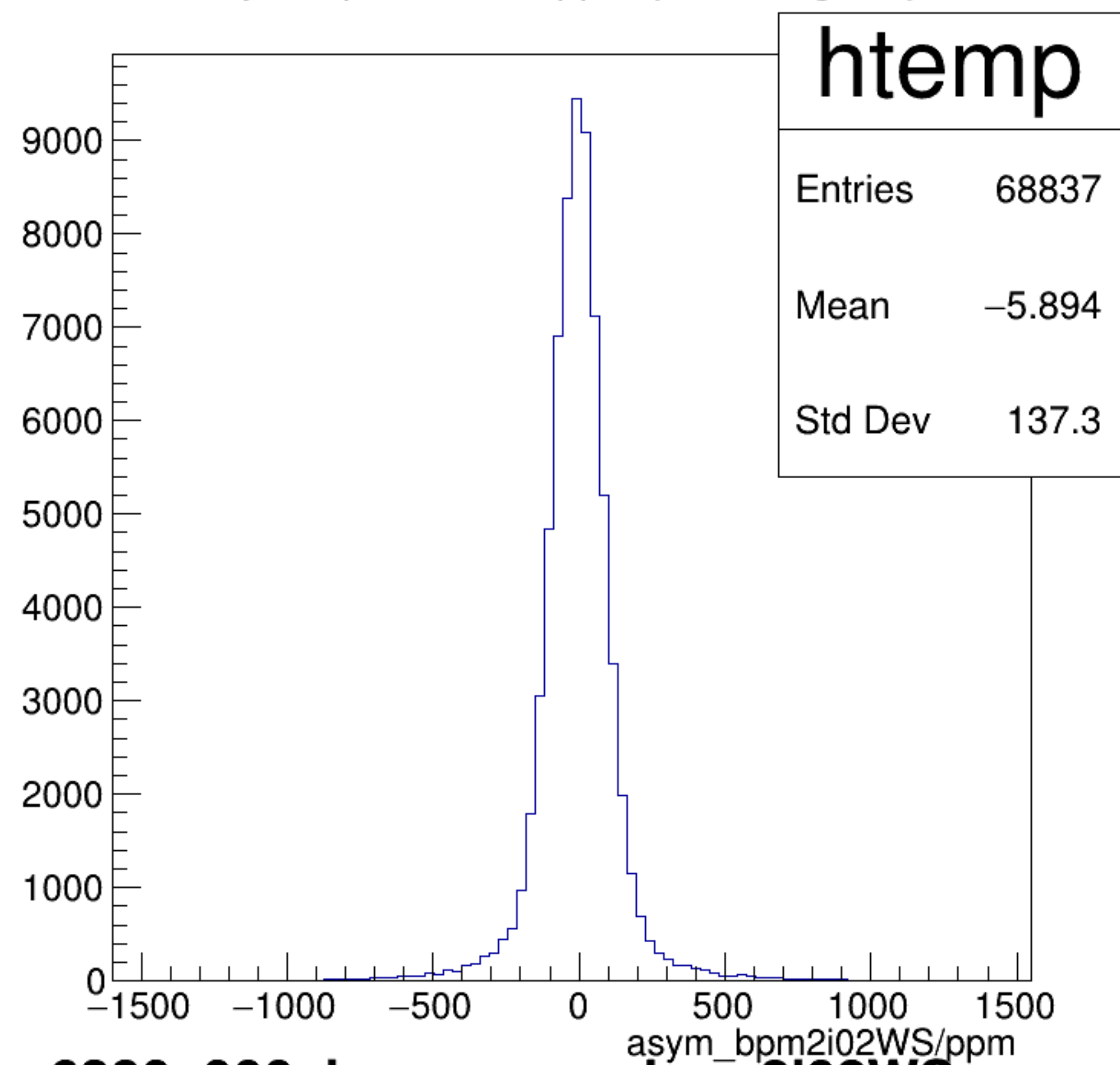
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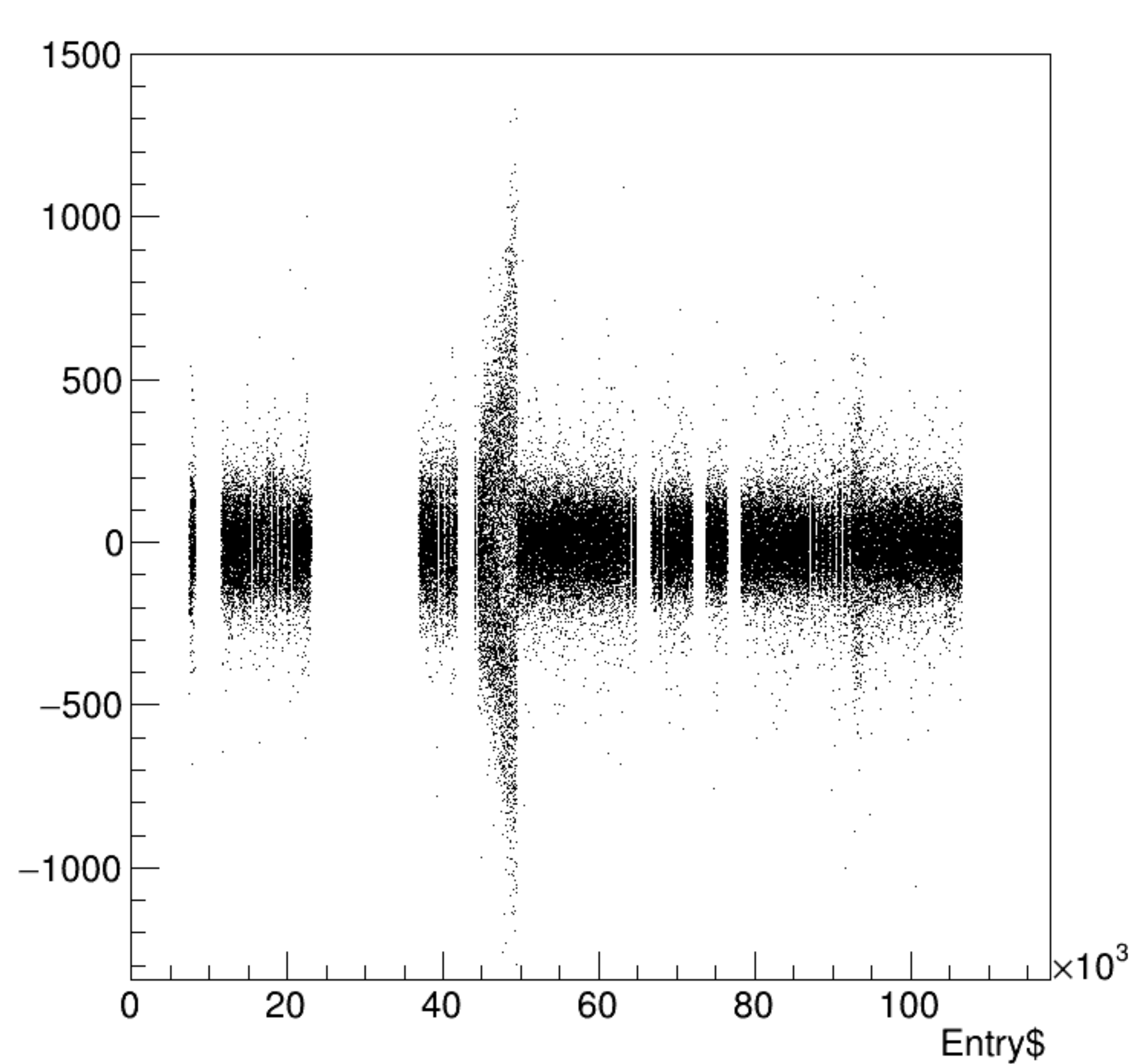


asym_bpm2i02WS/ppm {ErrorFlag==0}

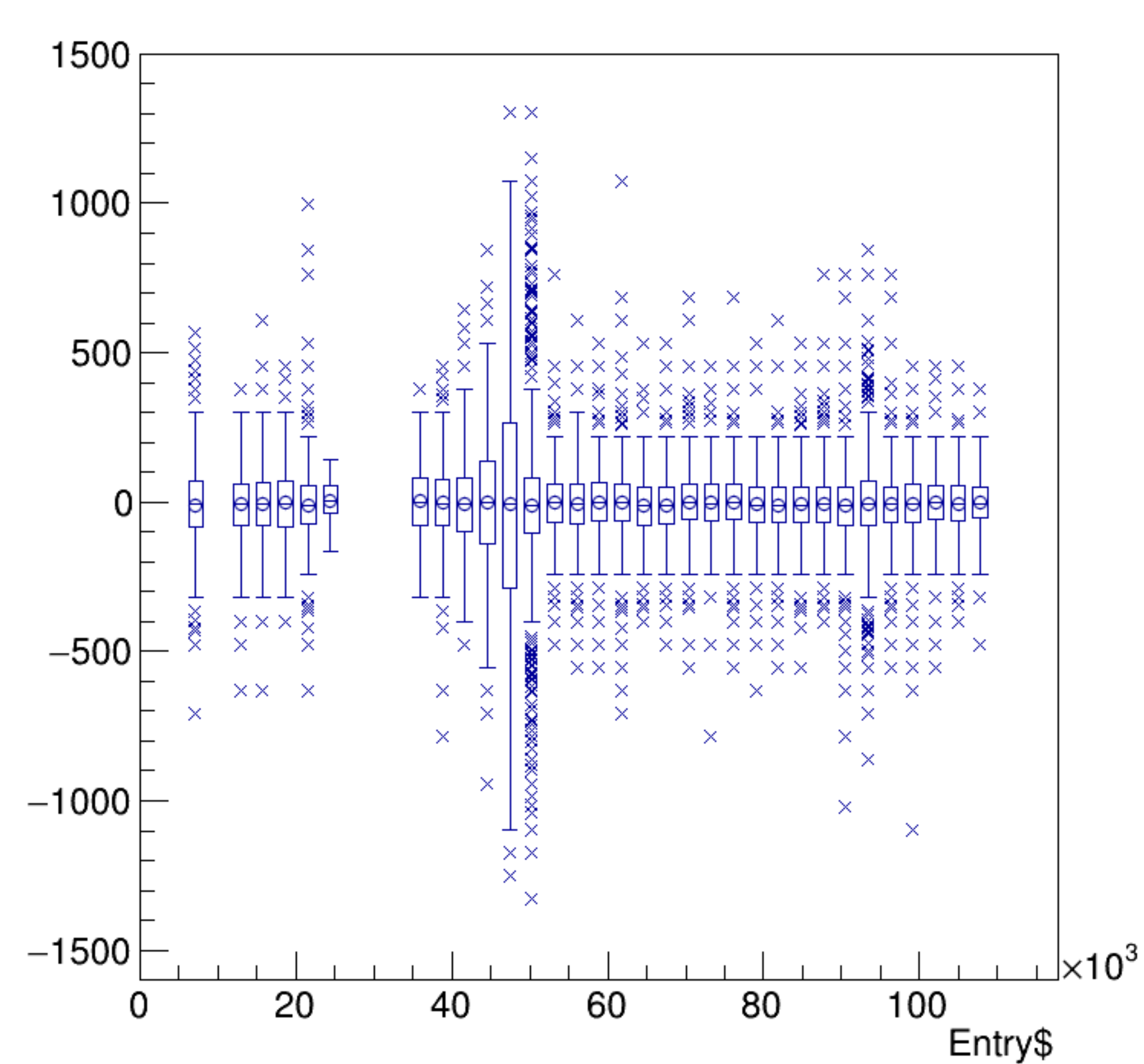


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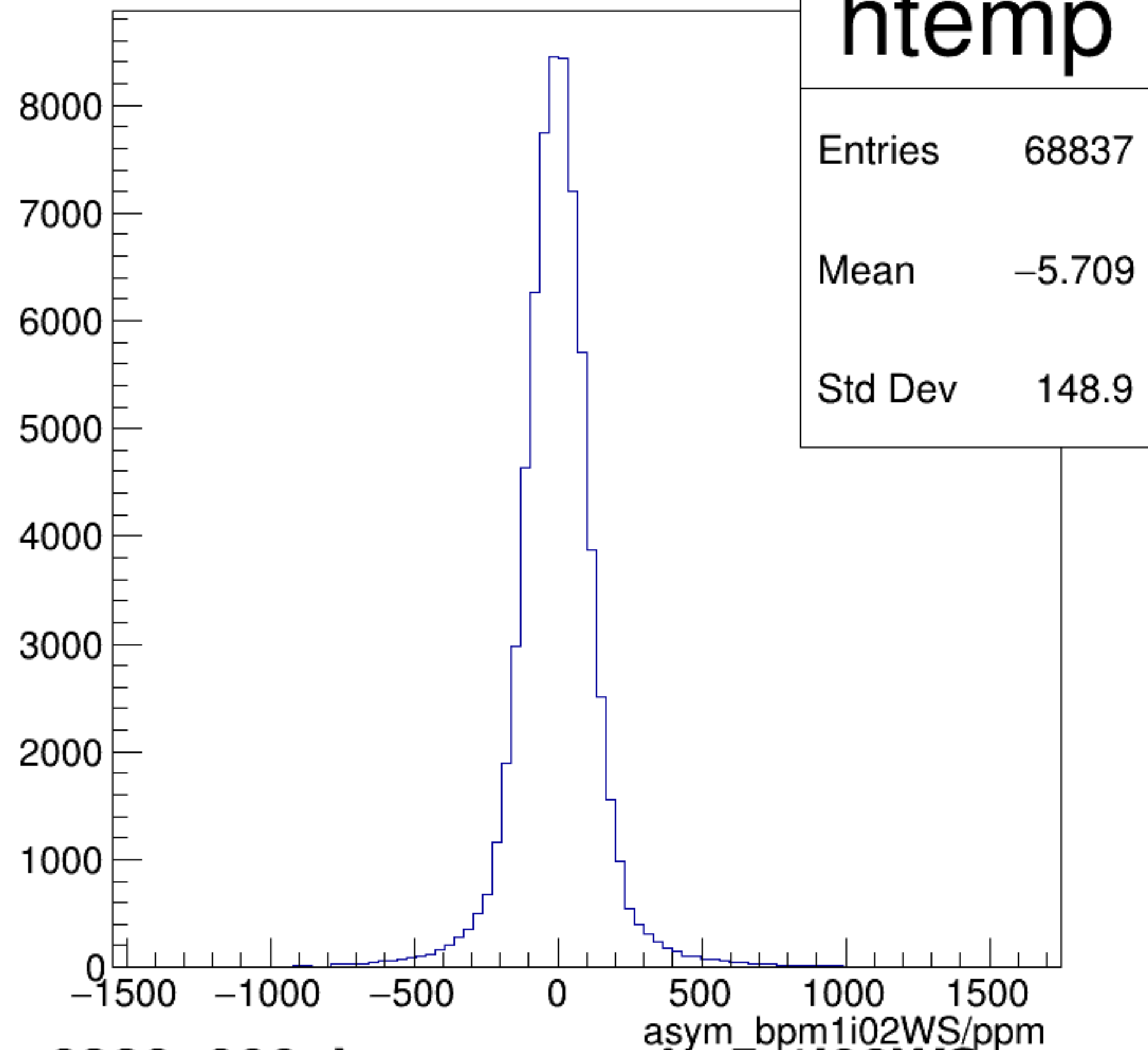
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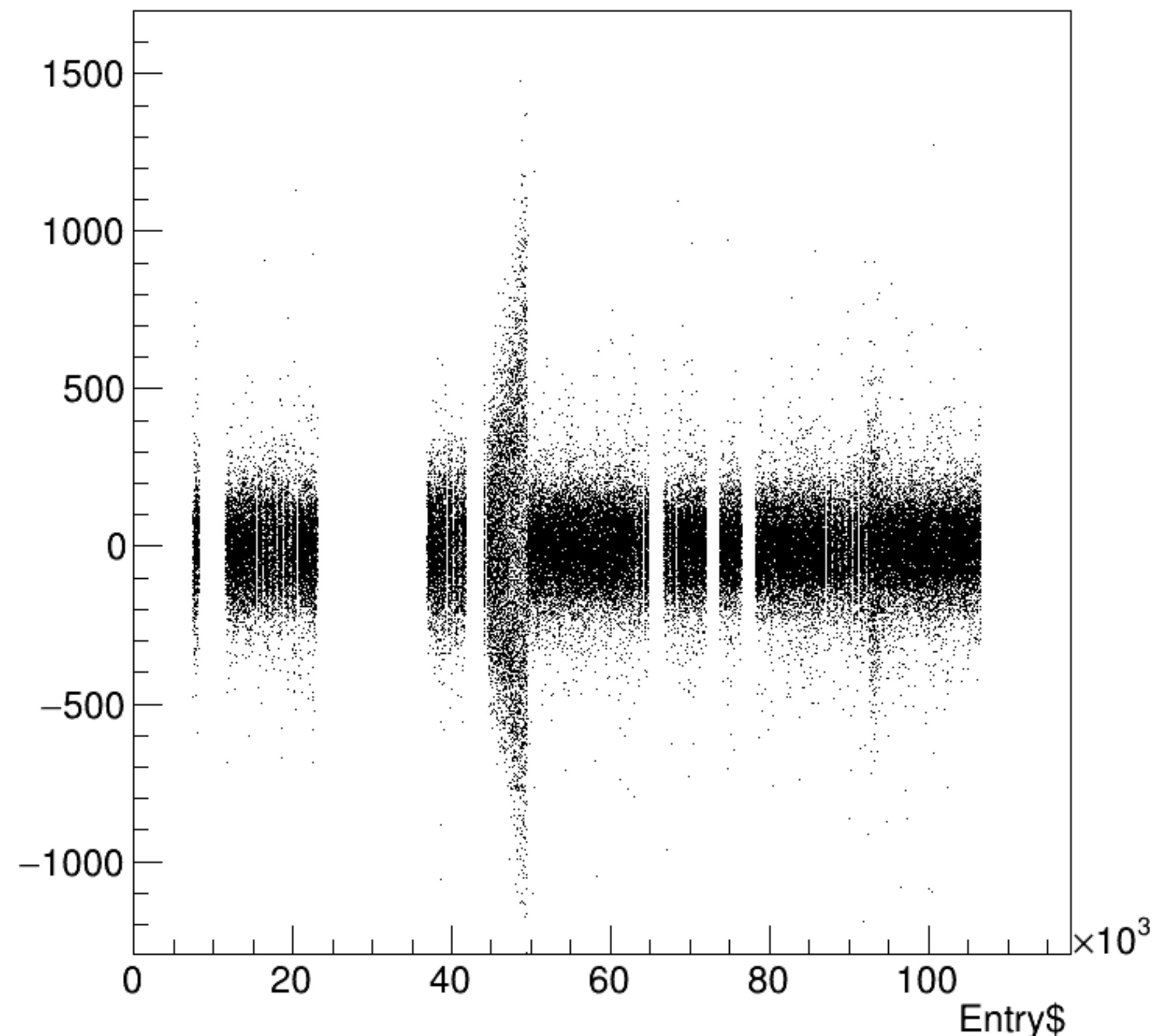


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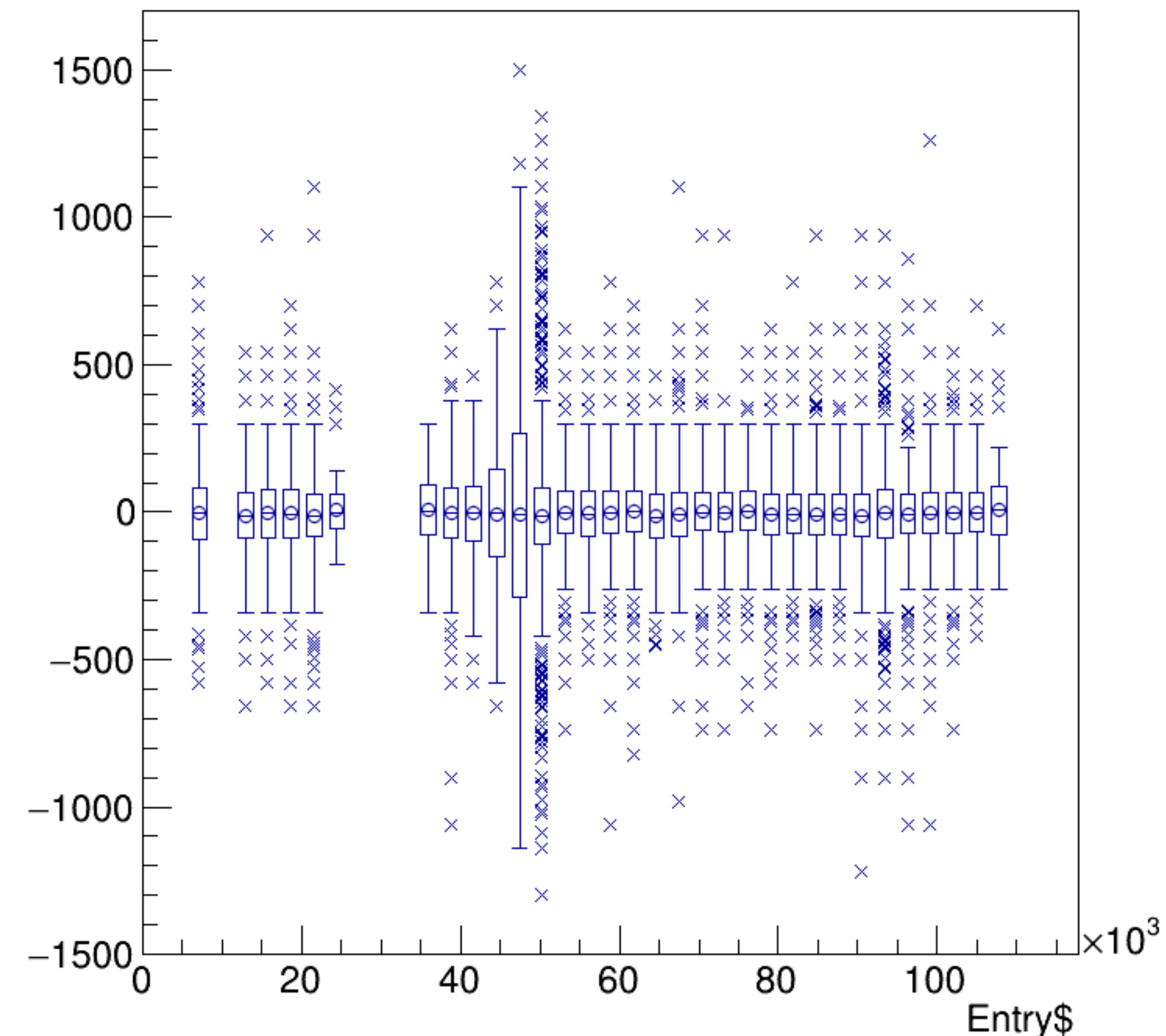


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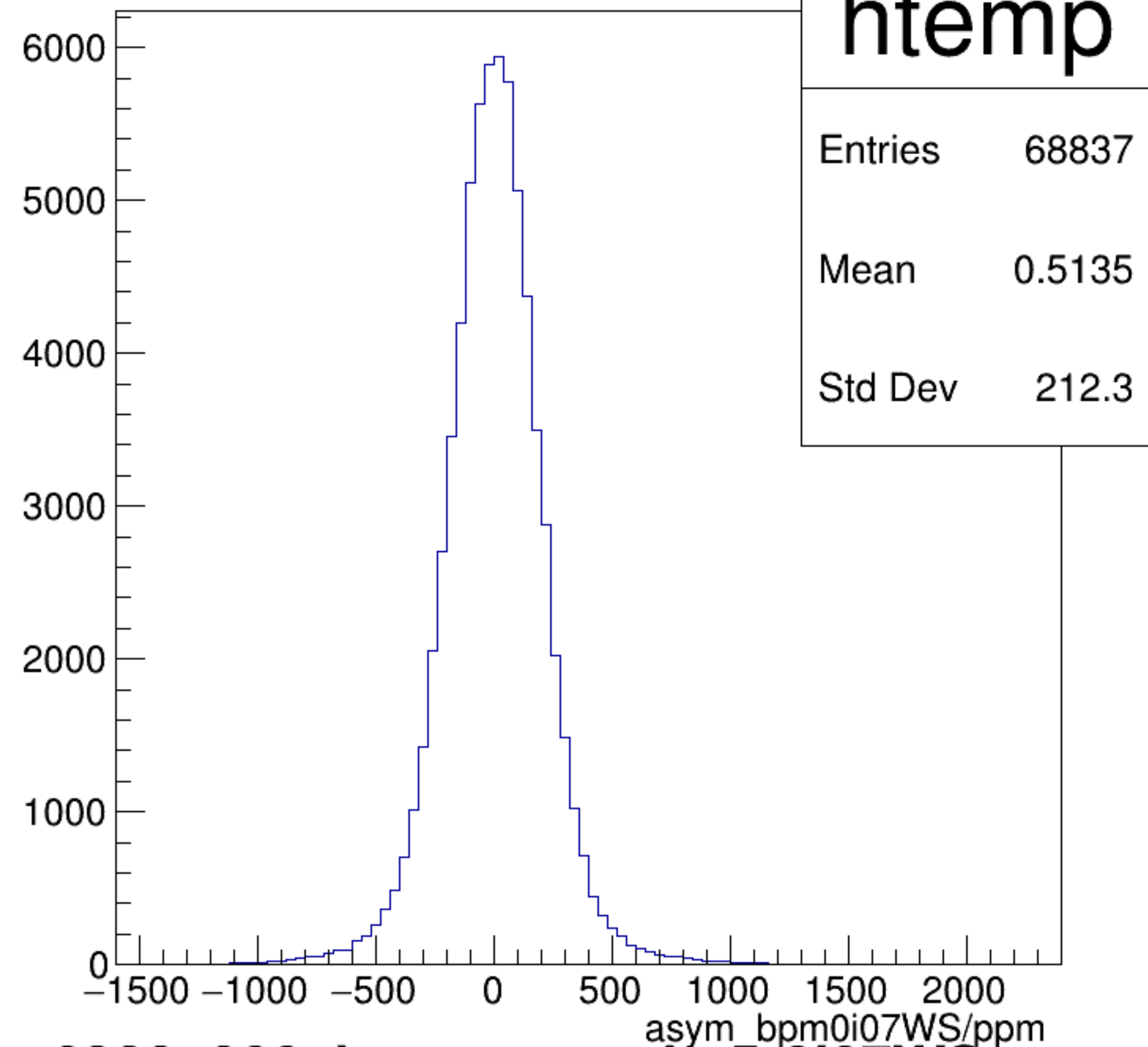
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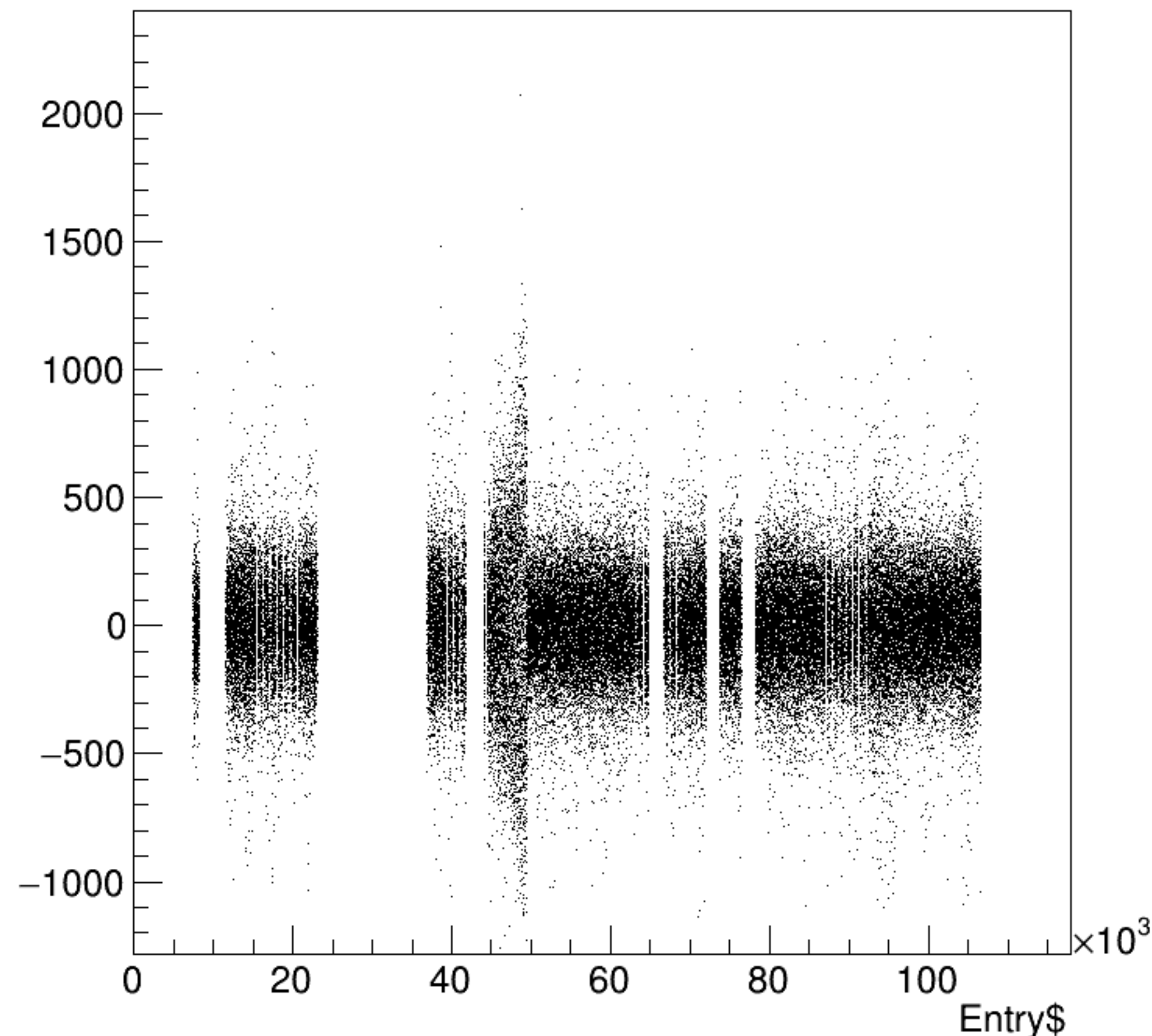


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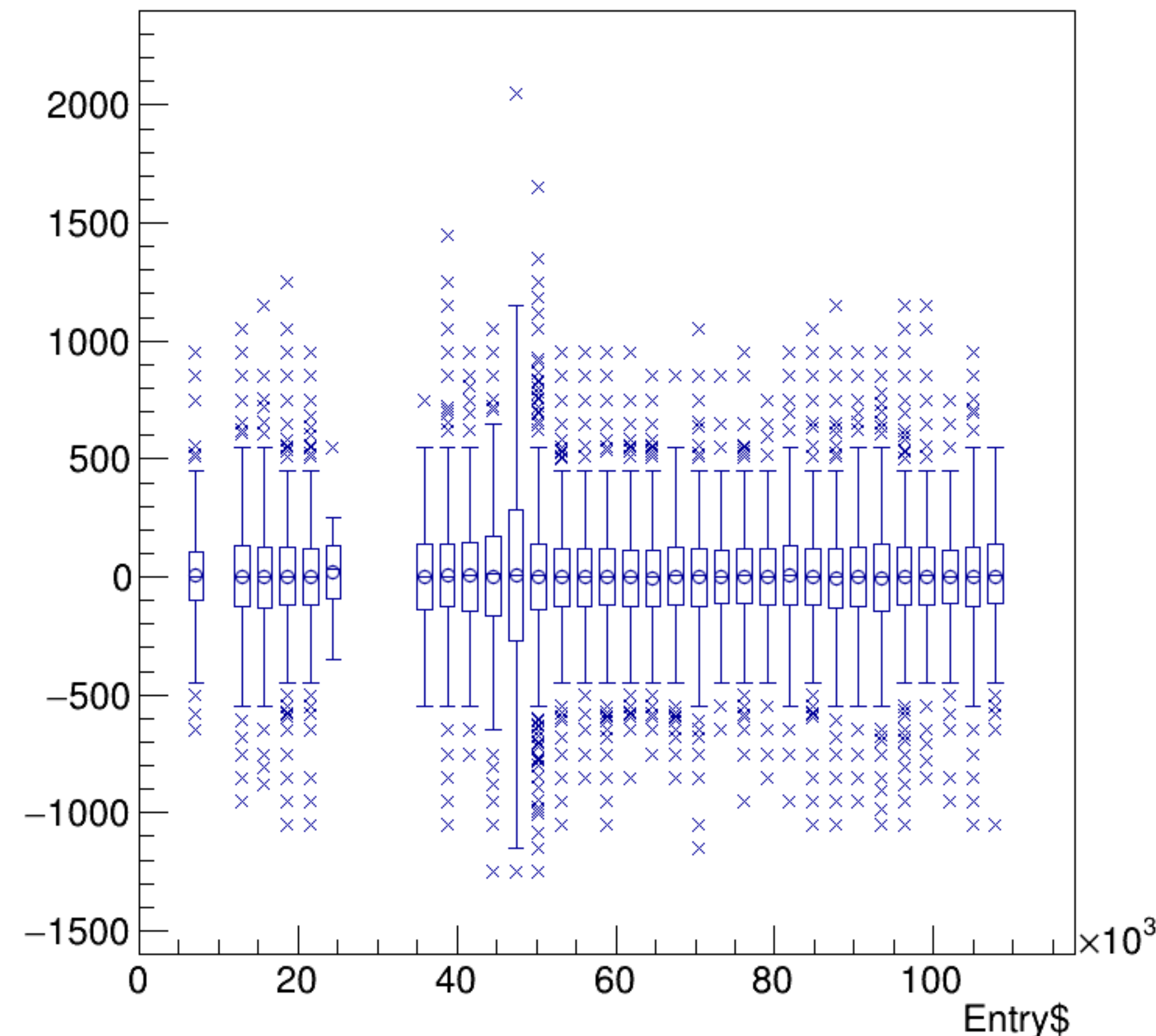


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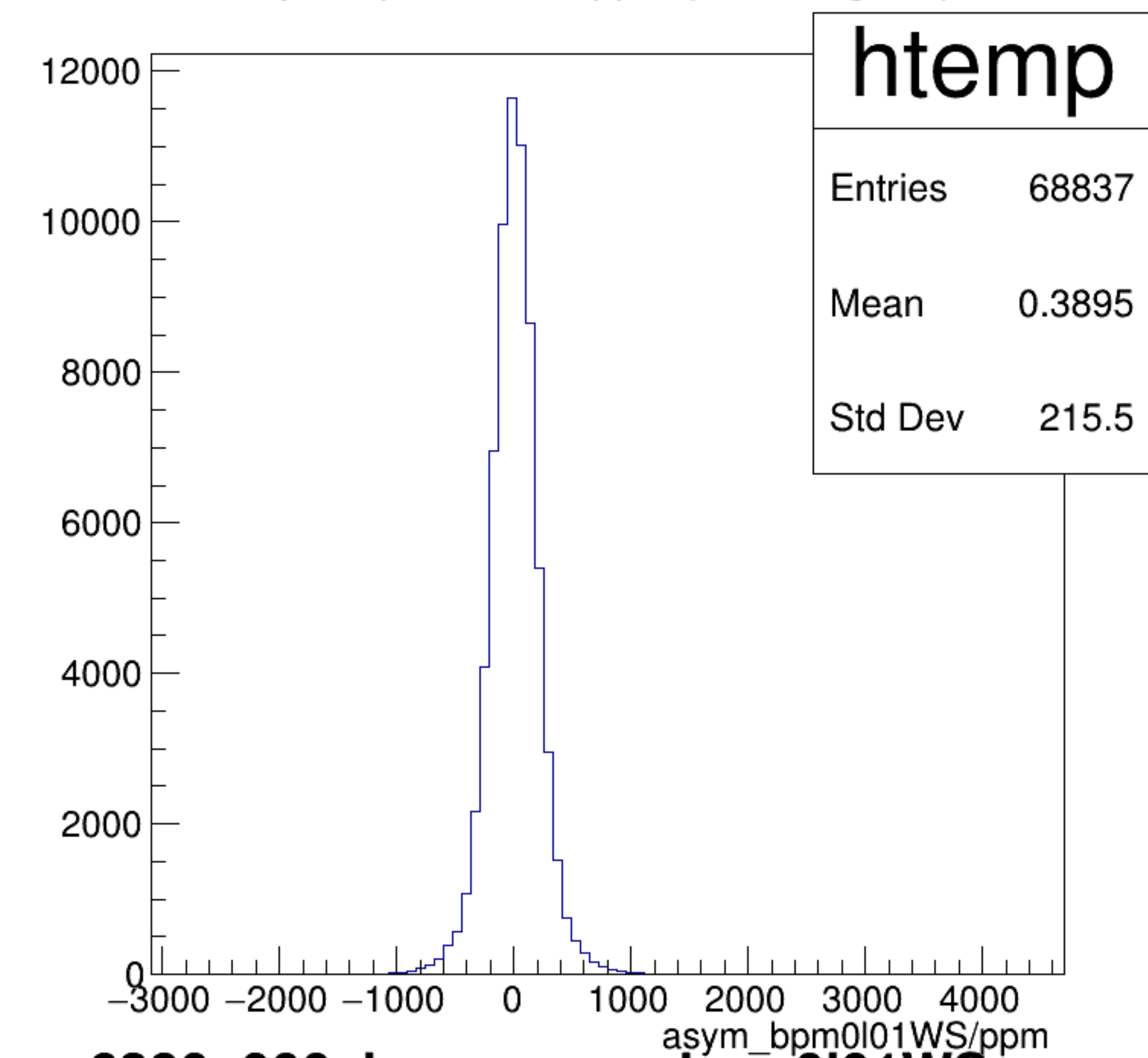
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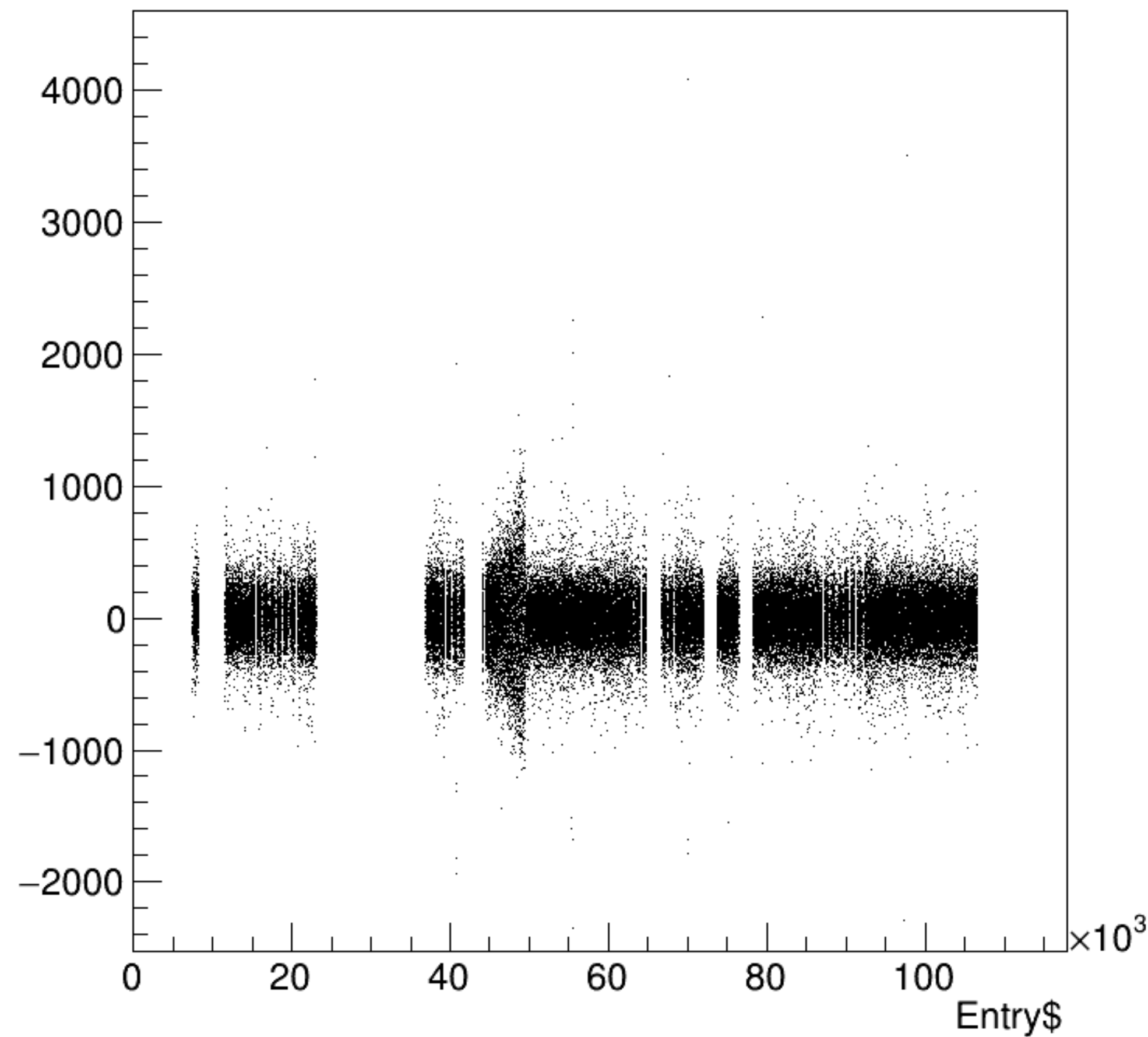


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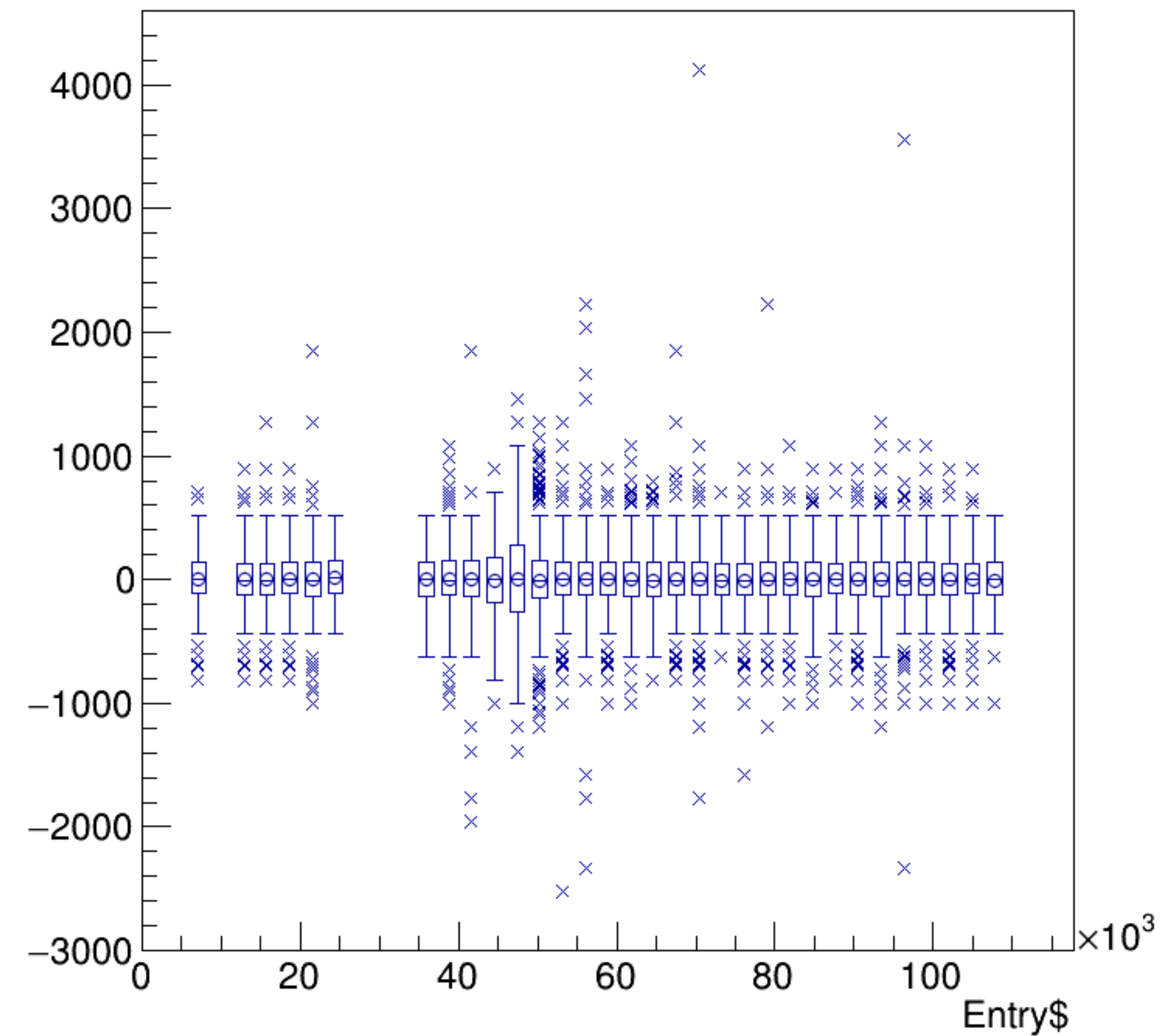


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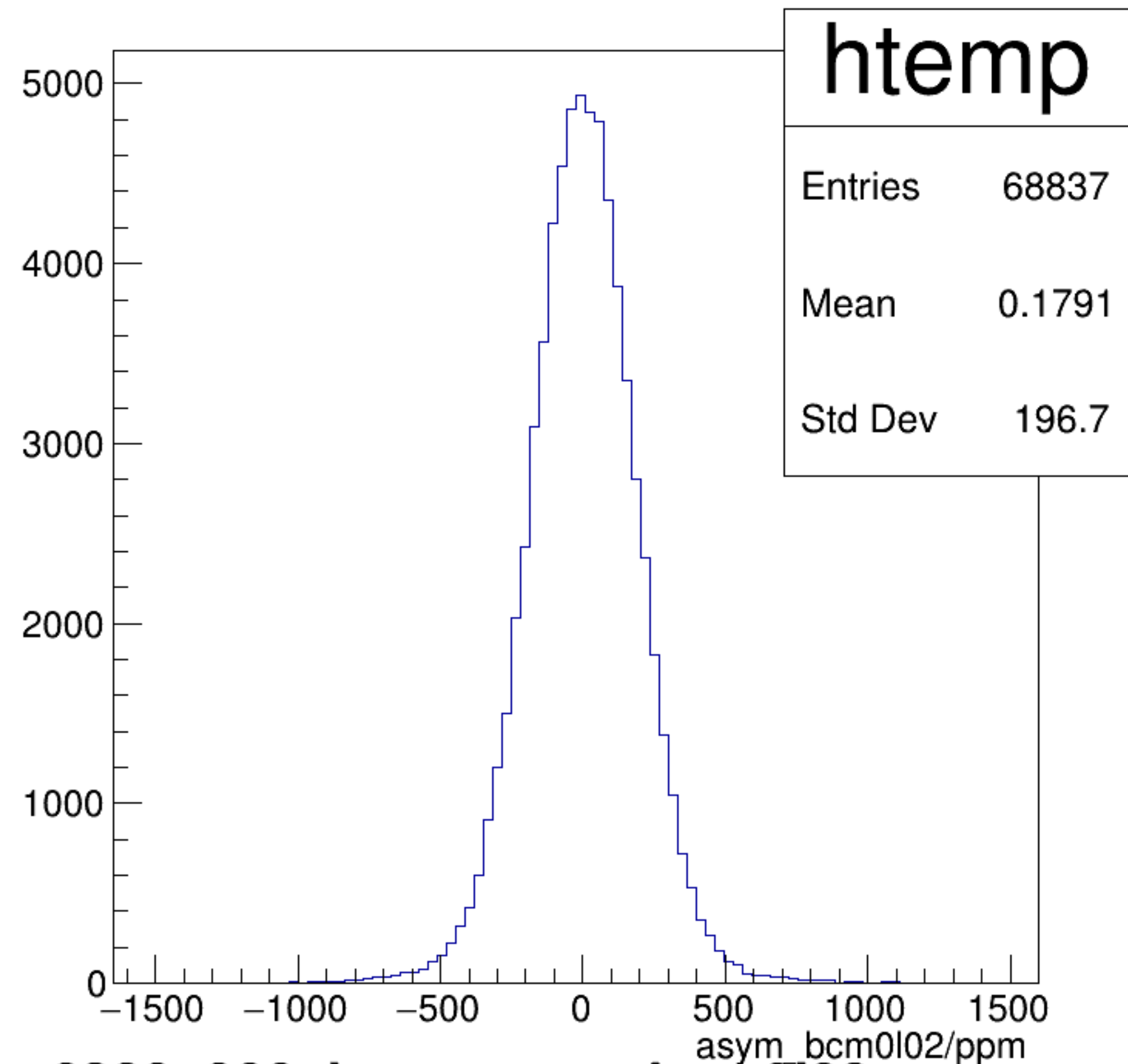
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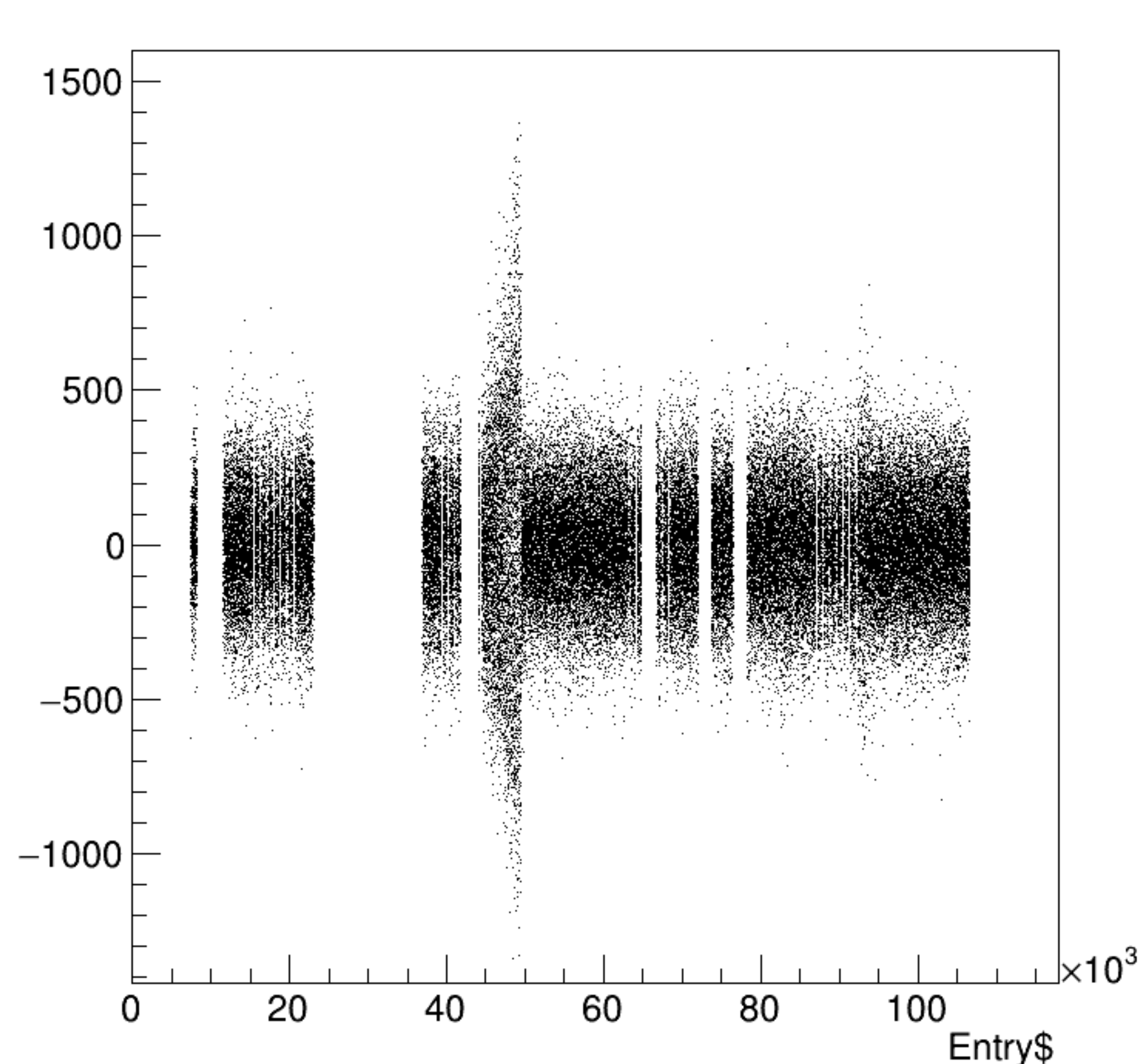
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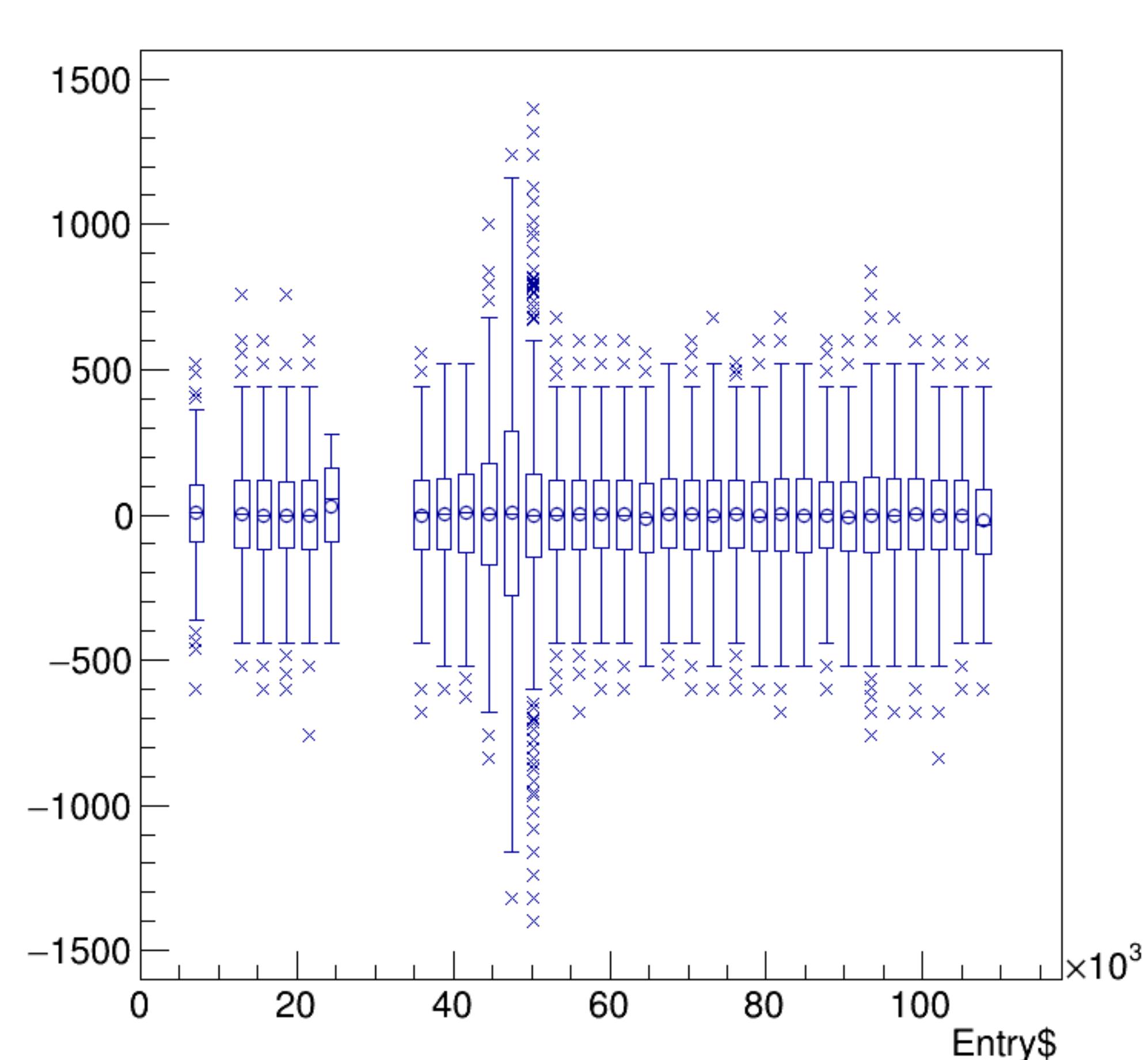
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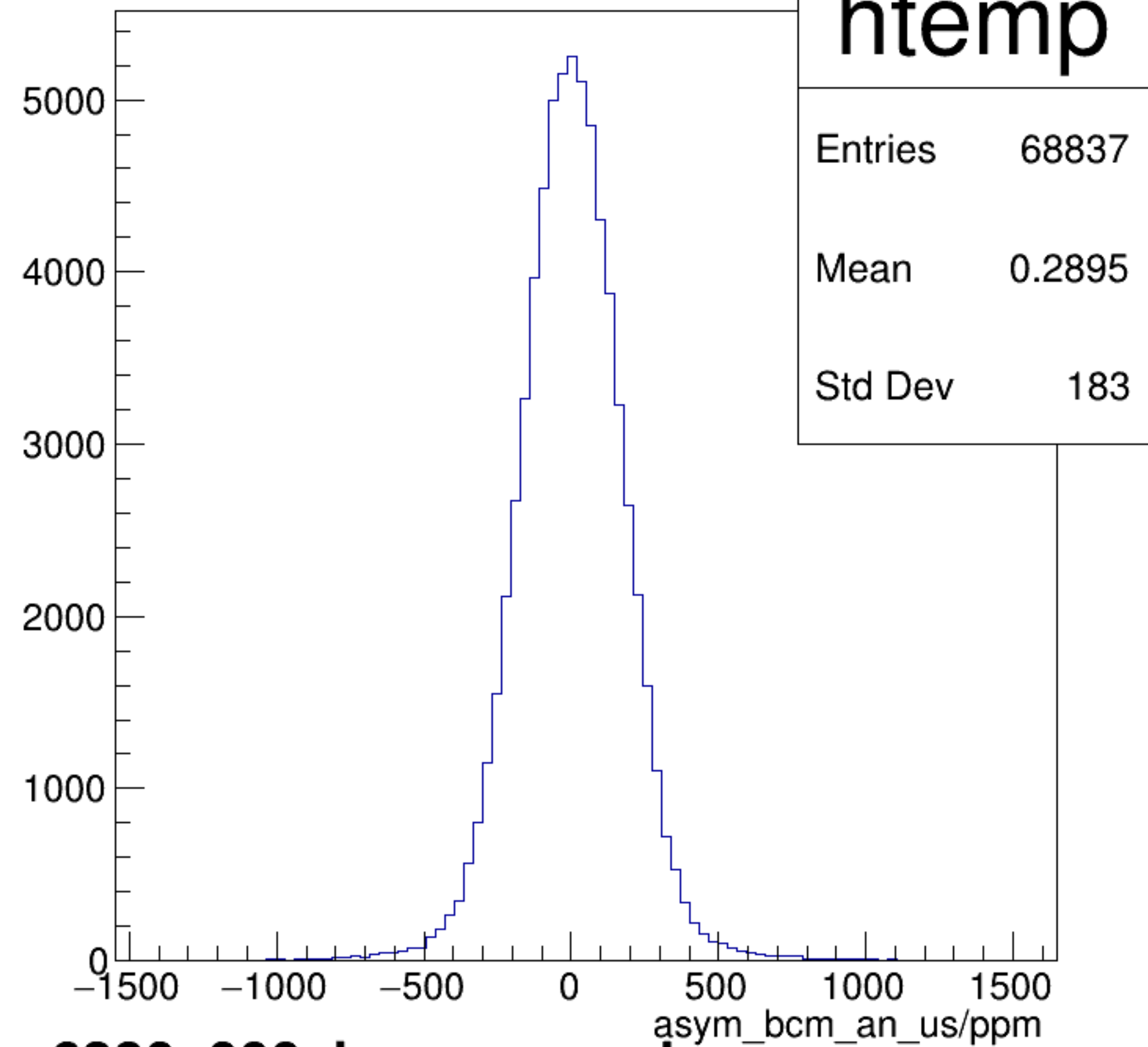
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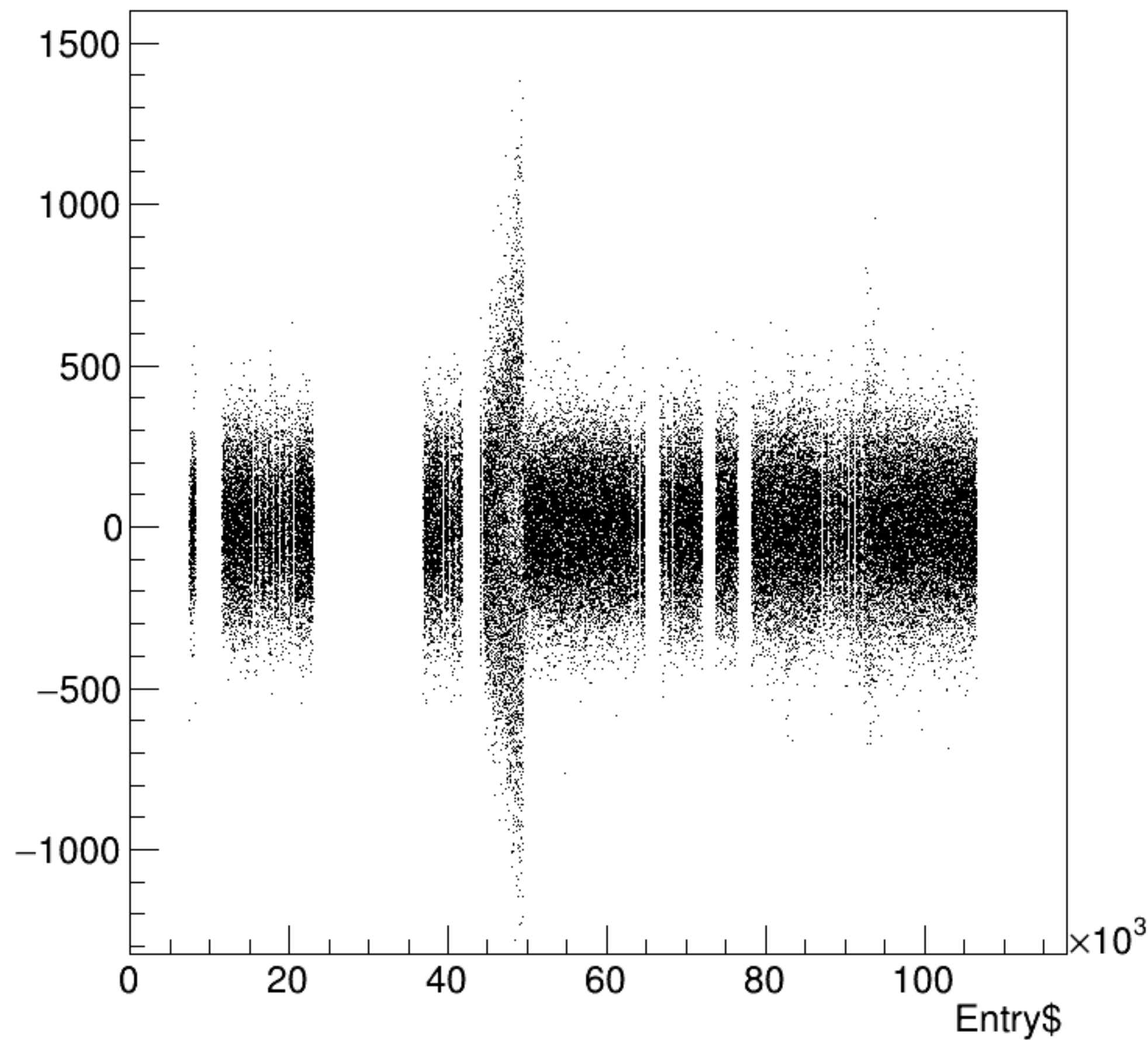


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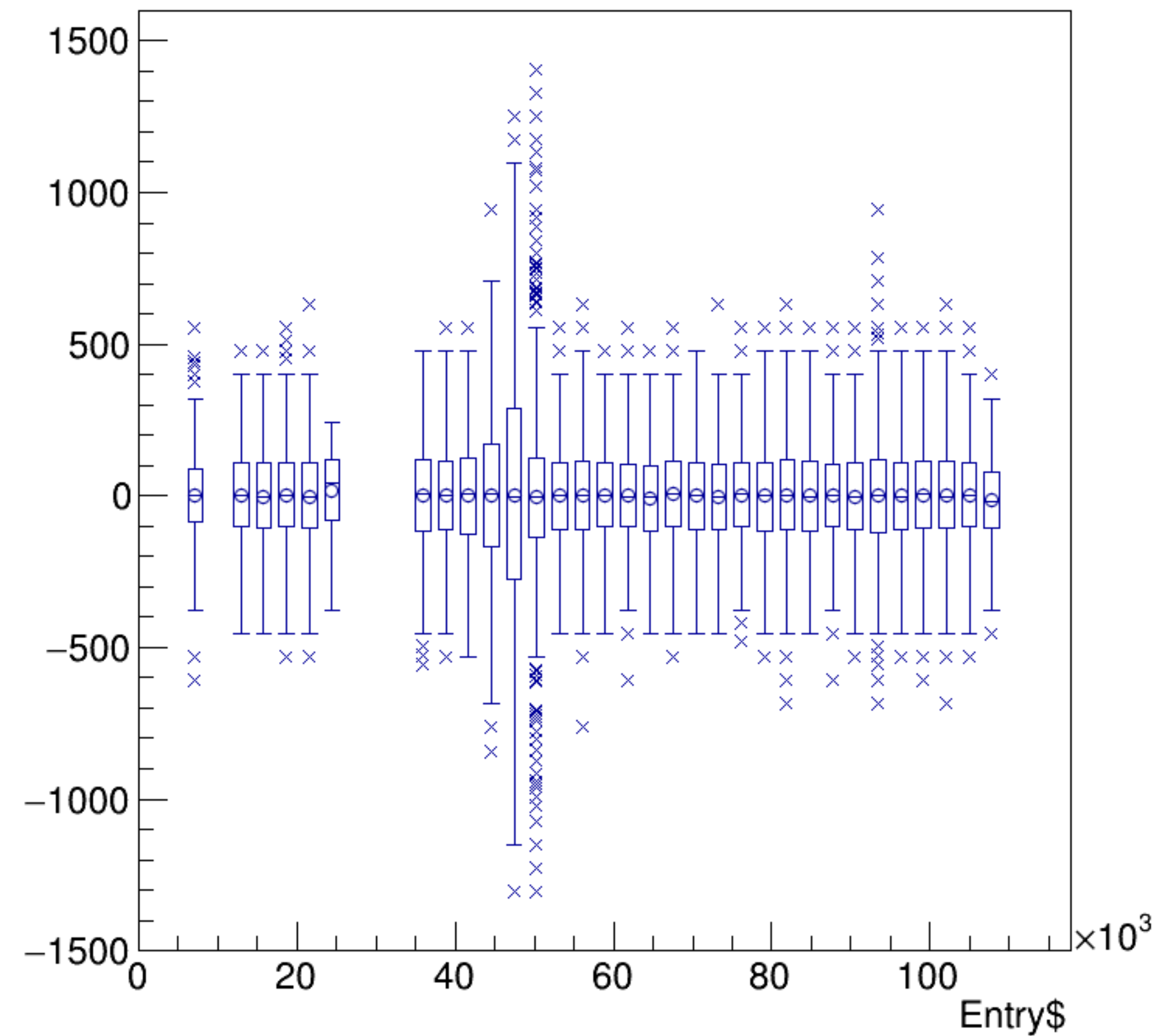


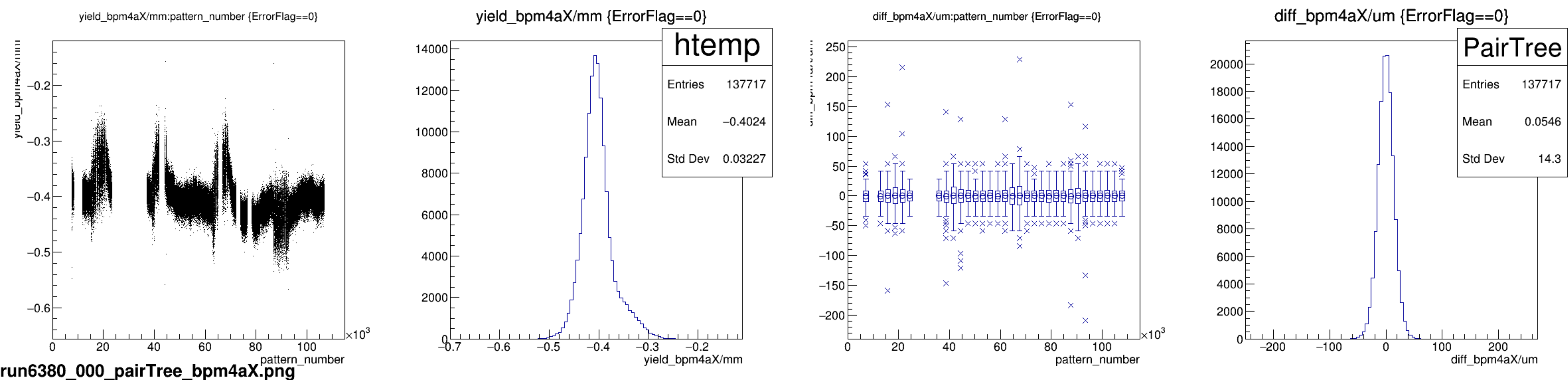
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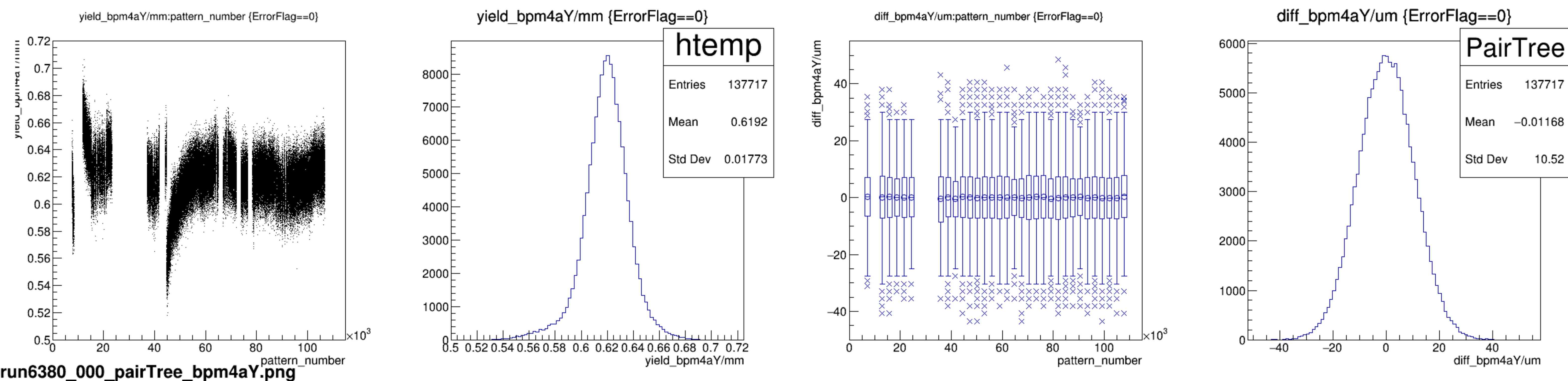
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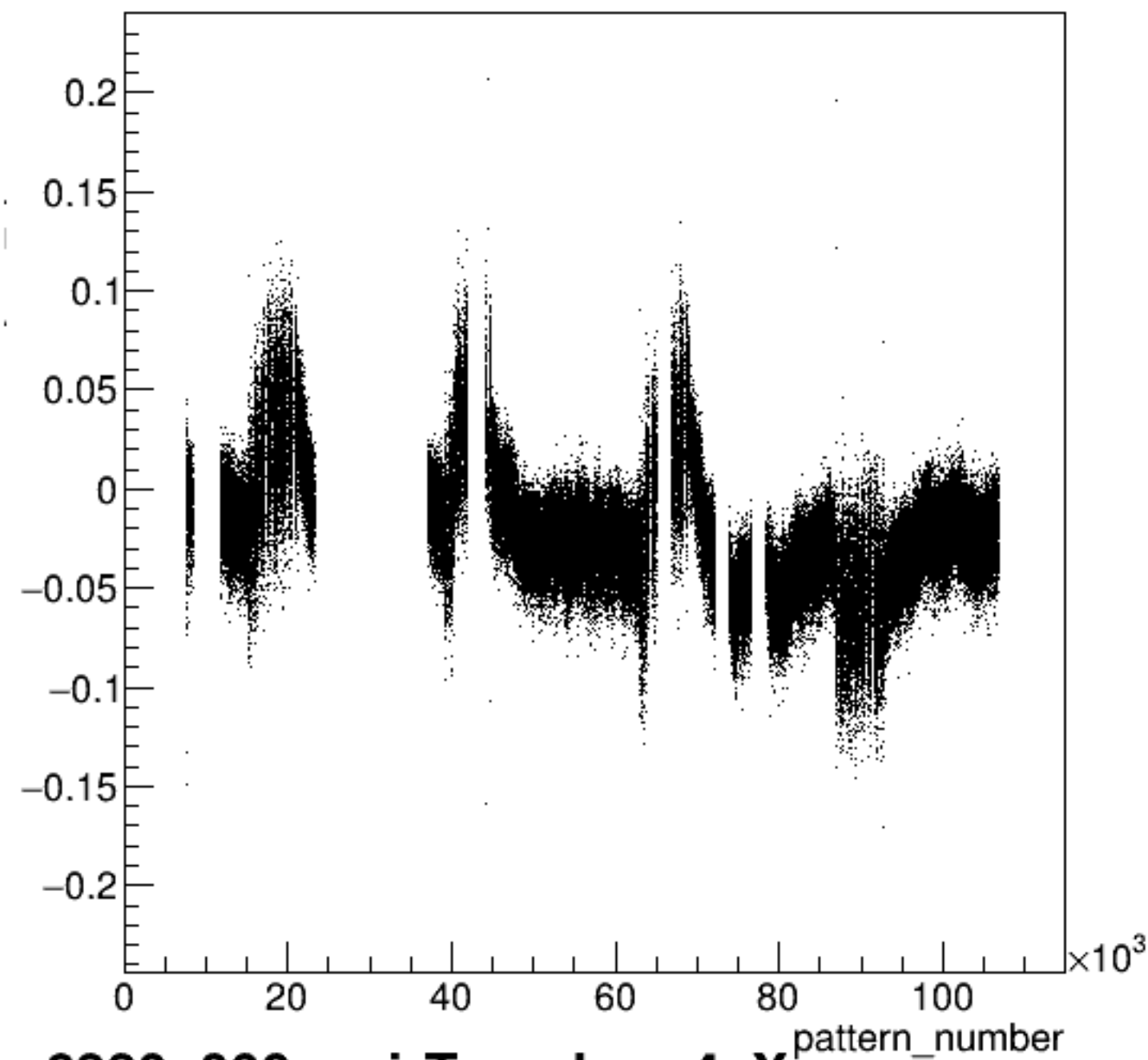
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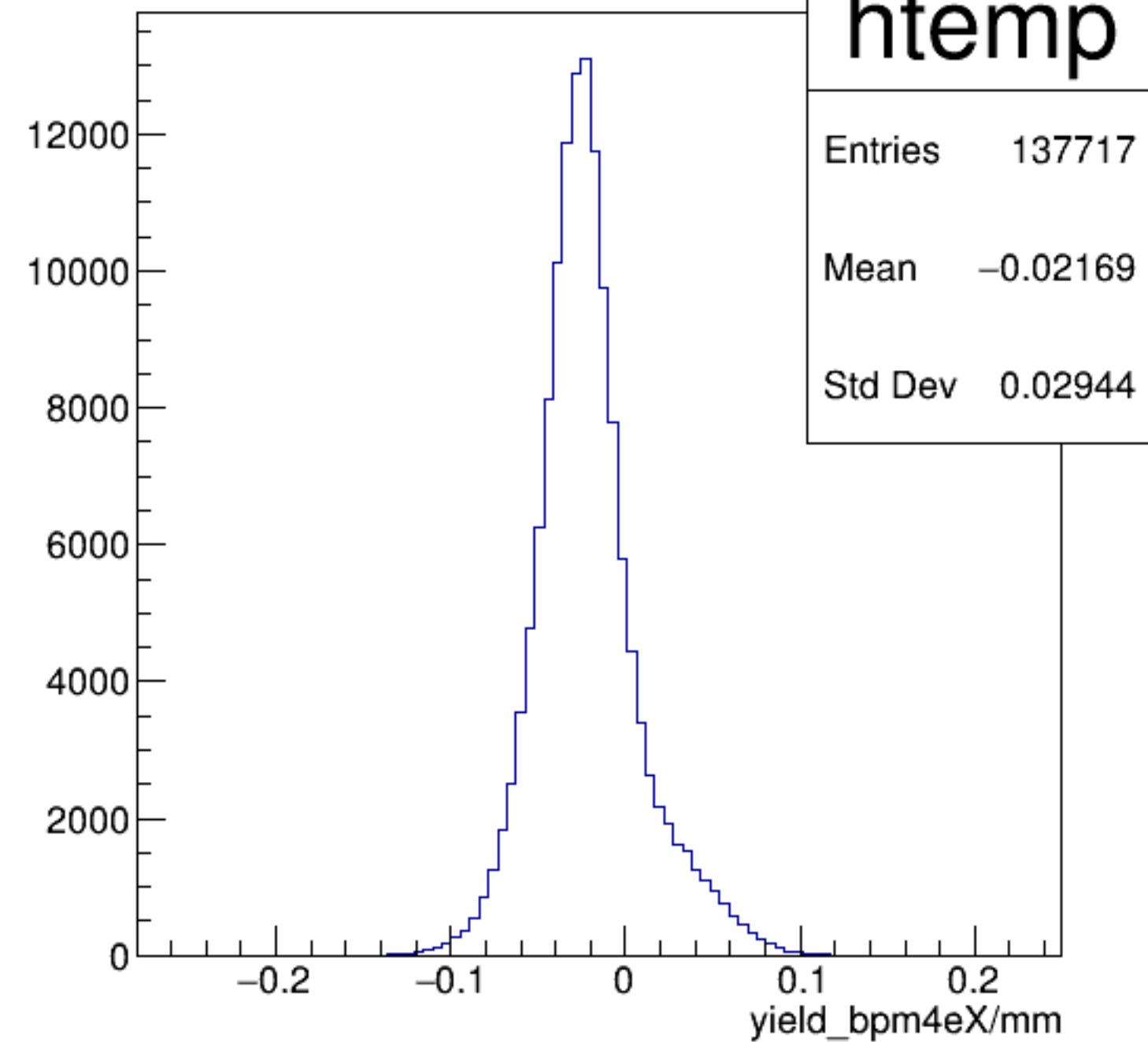


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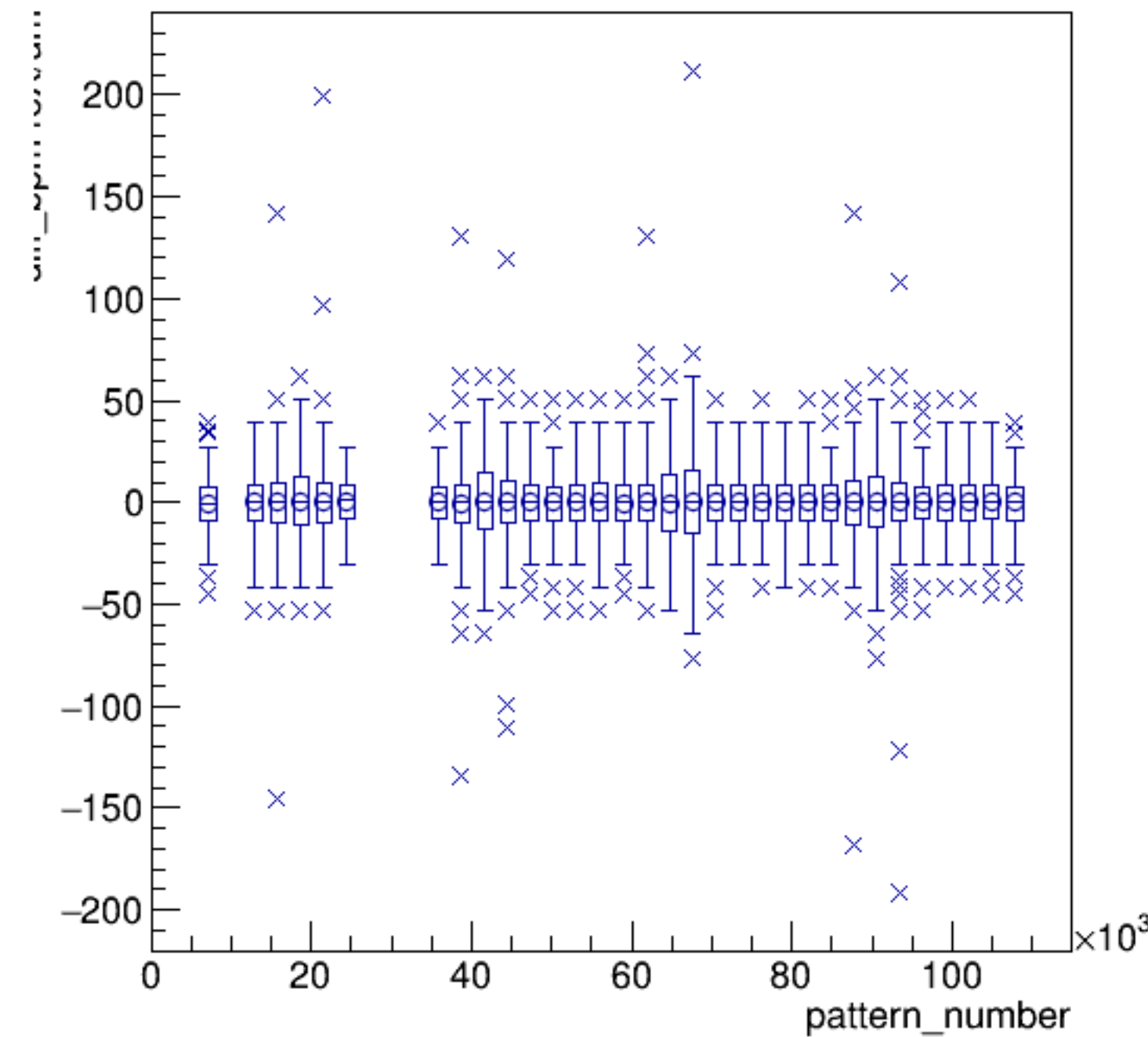


run6380_000_pairTree_bpm4eX.png

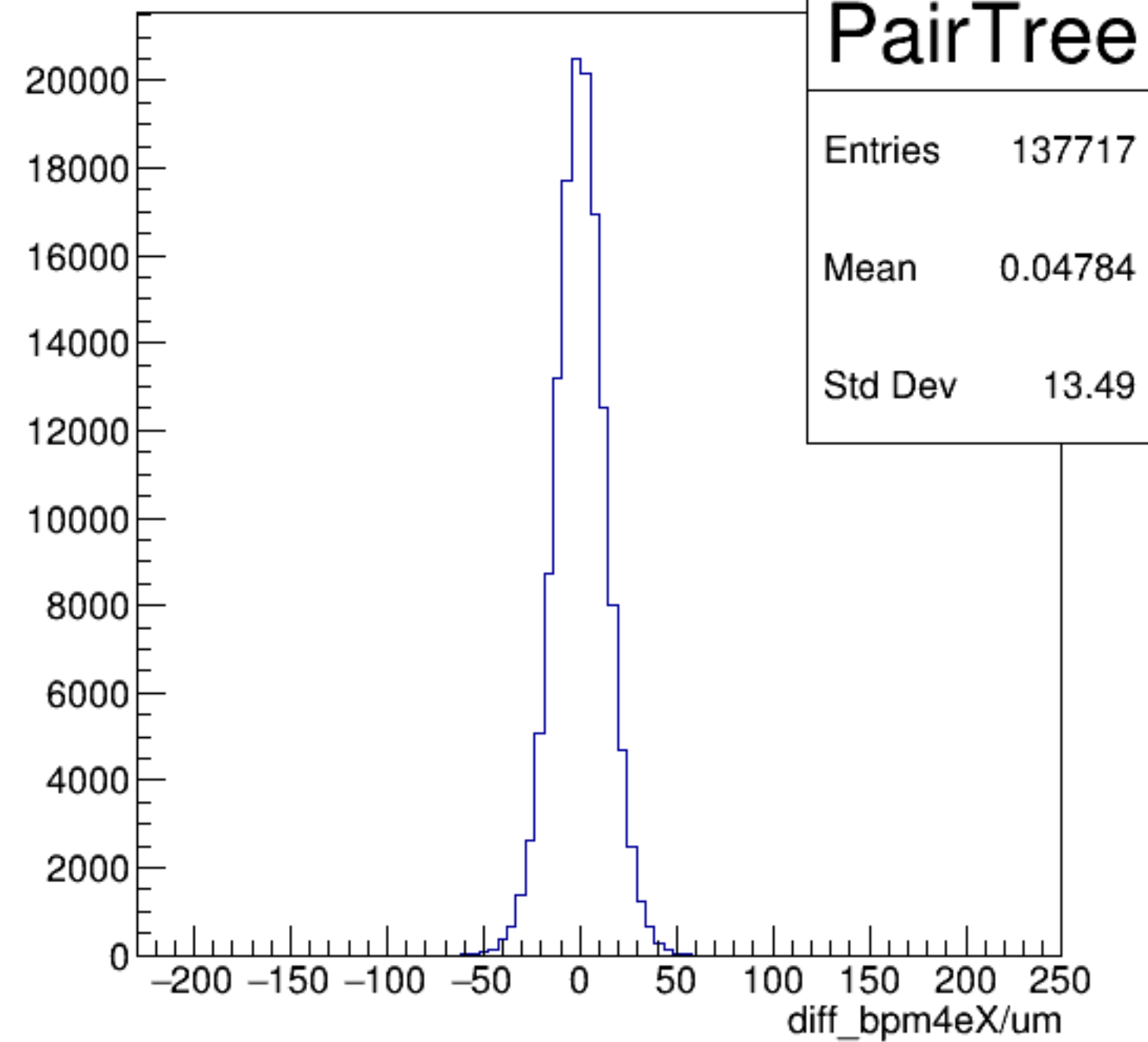
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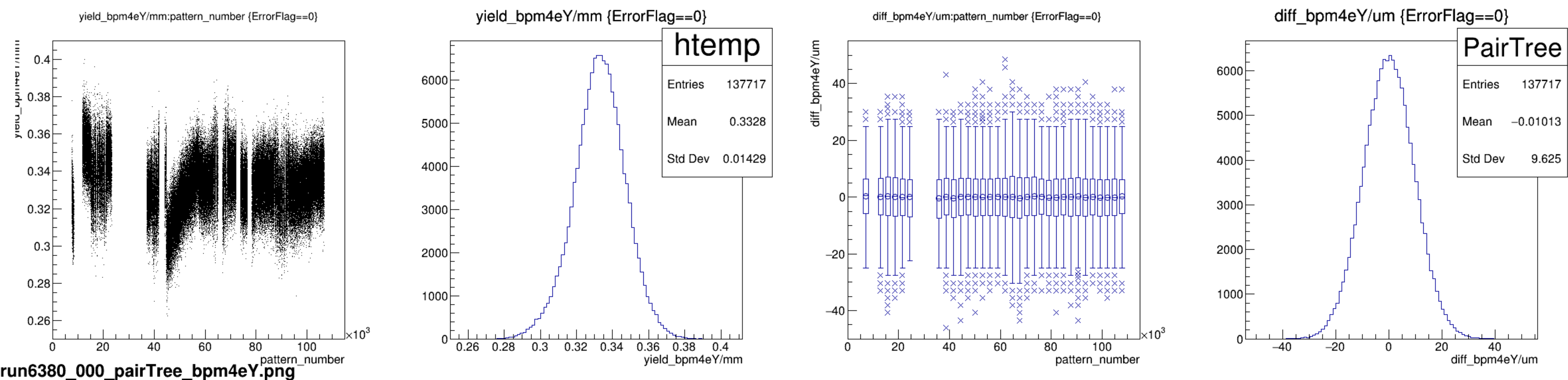


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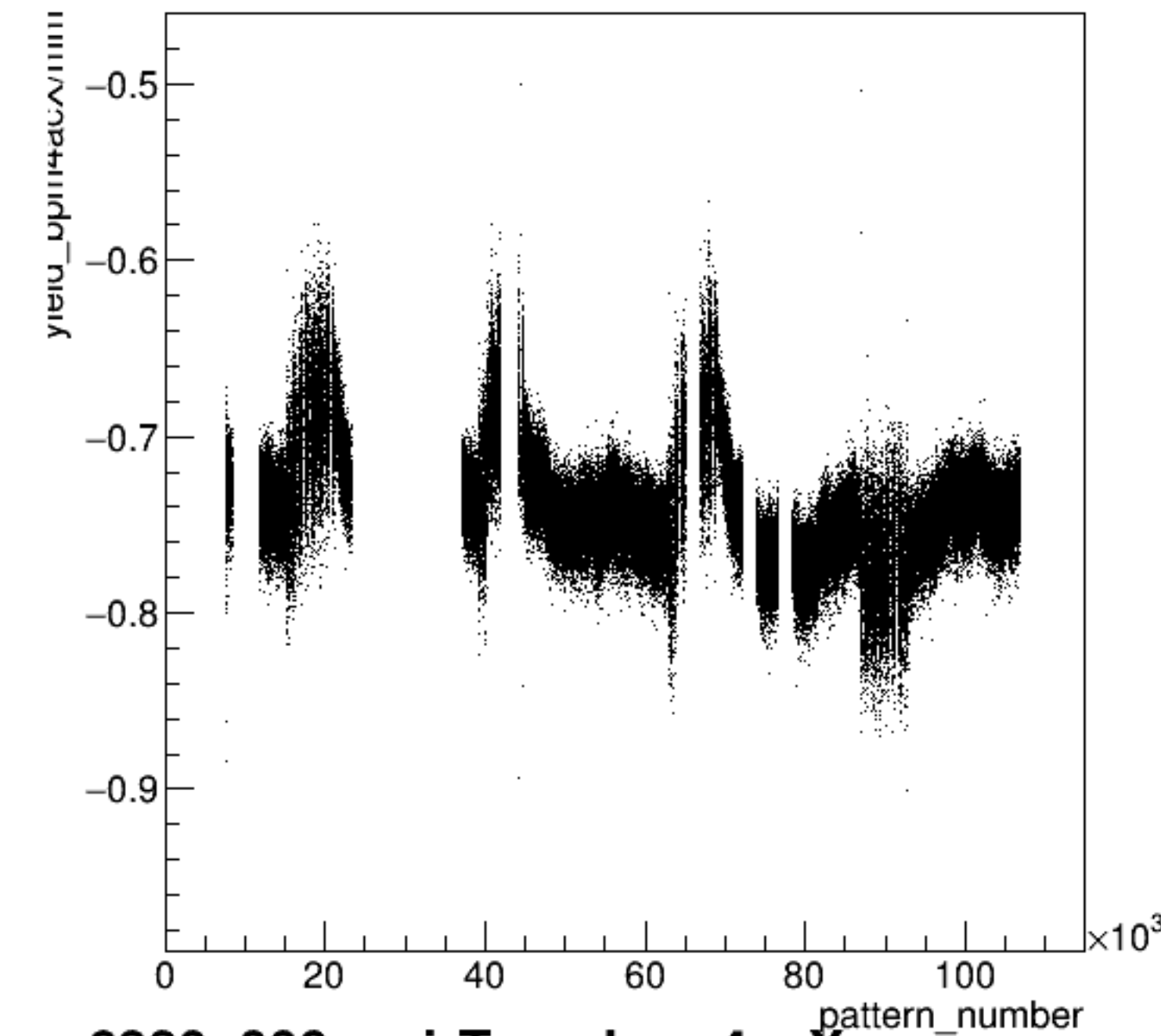


diff_bpm4eX/um {ErrorFlag==0}



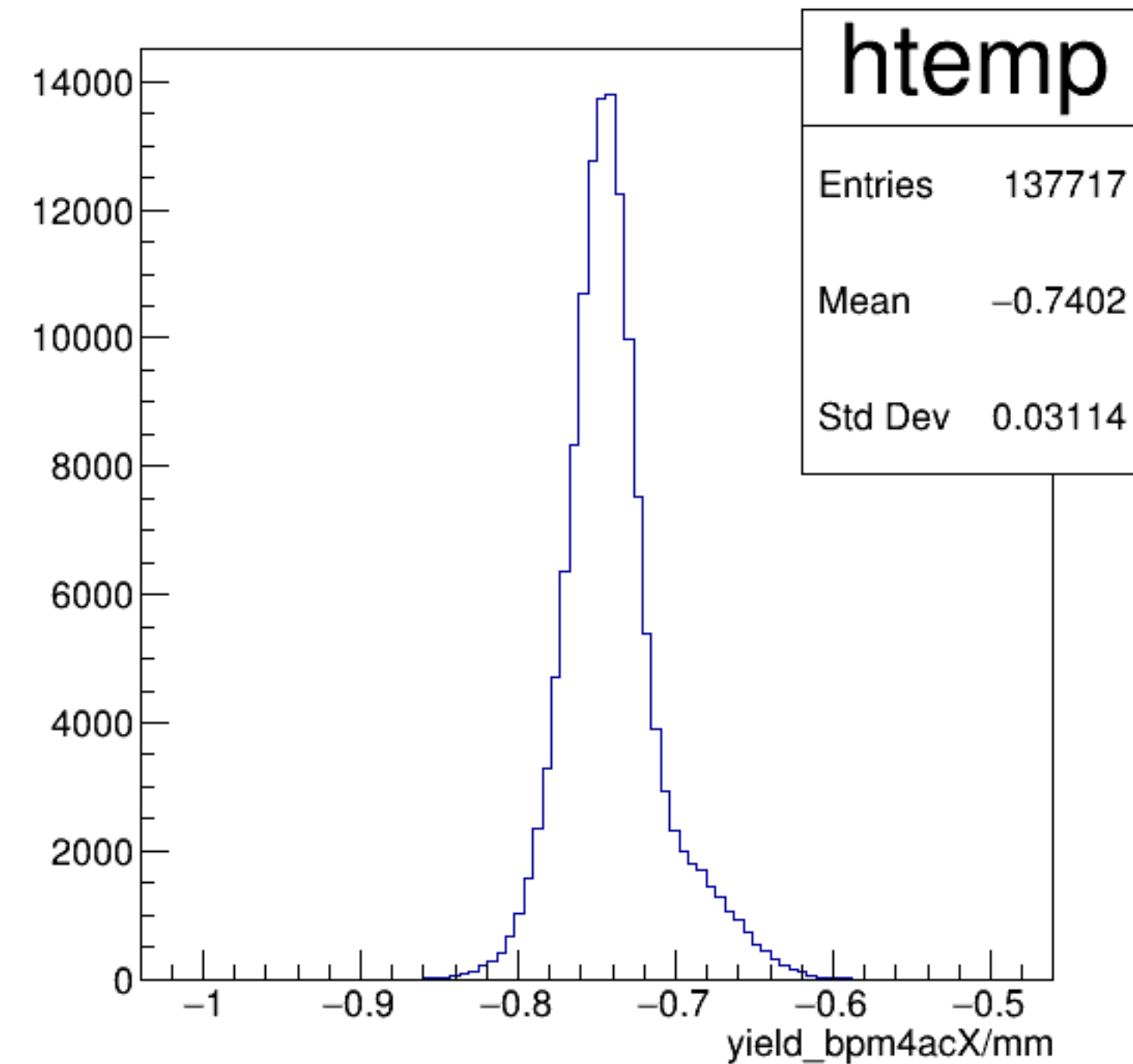


yield_bpm4acX/mm:pattern_number {ErrorFlag==0}

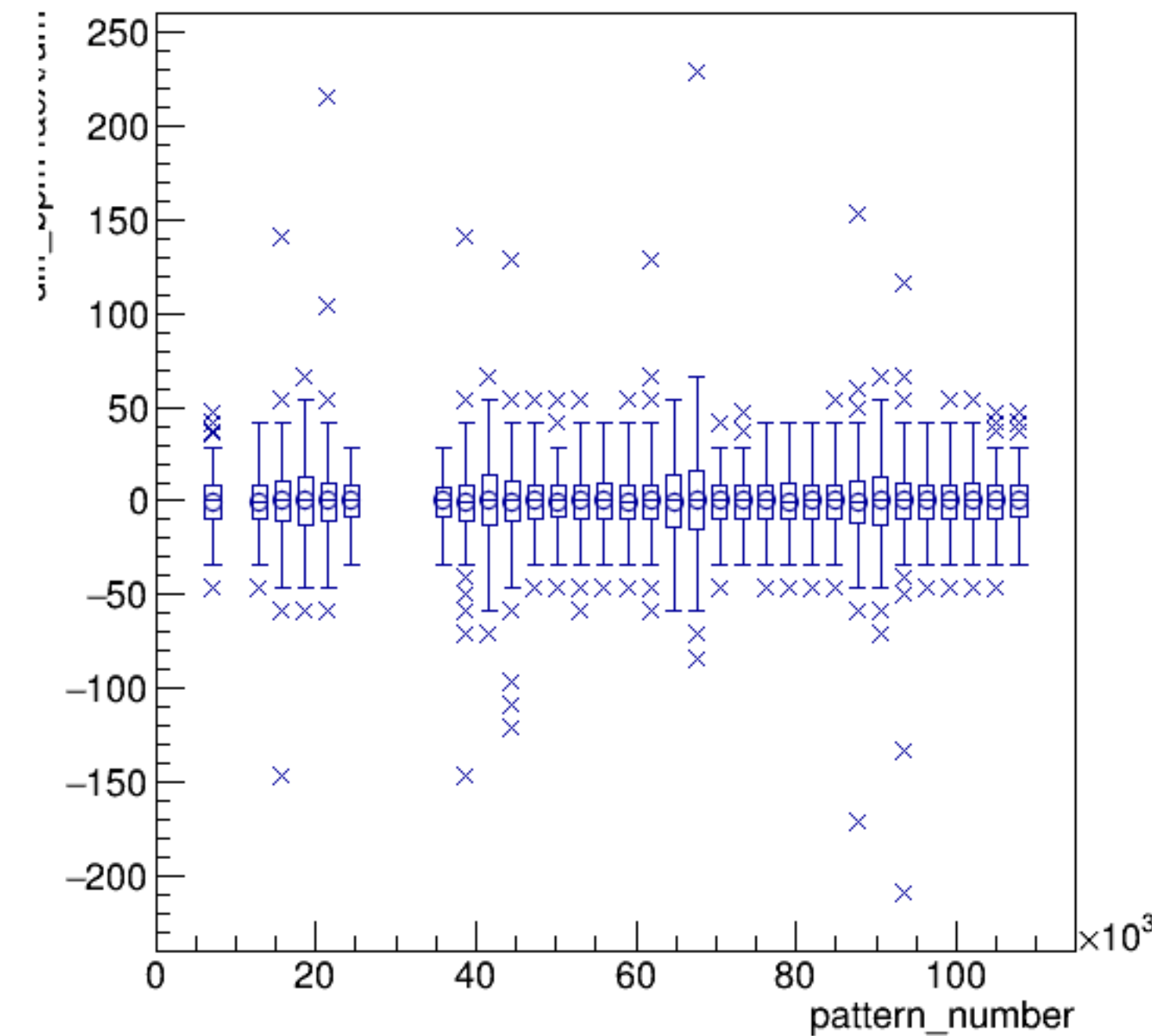


run6380_000_pairTree_bpm4acX.png

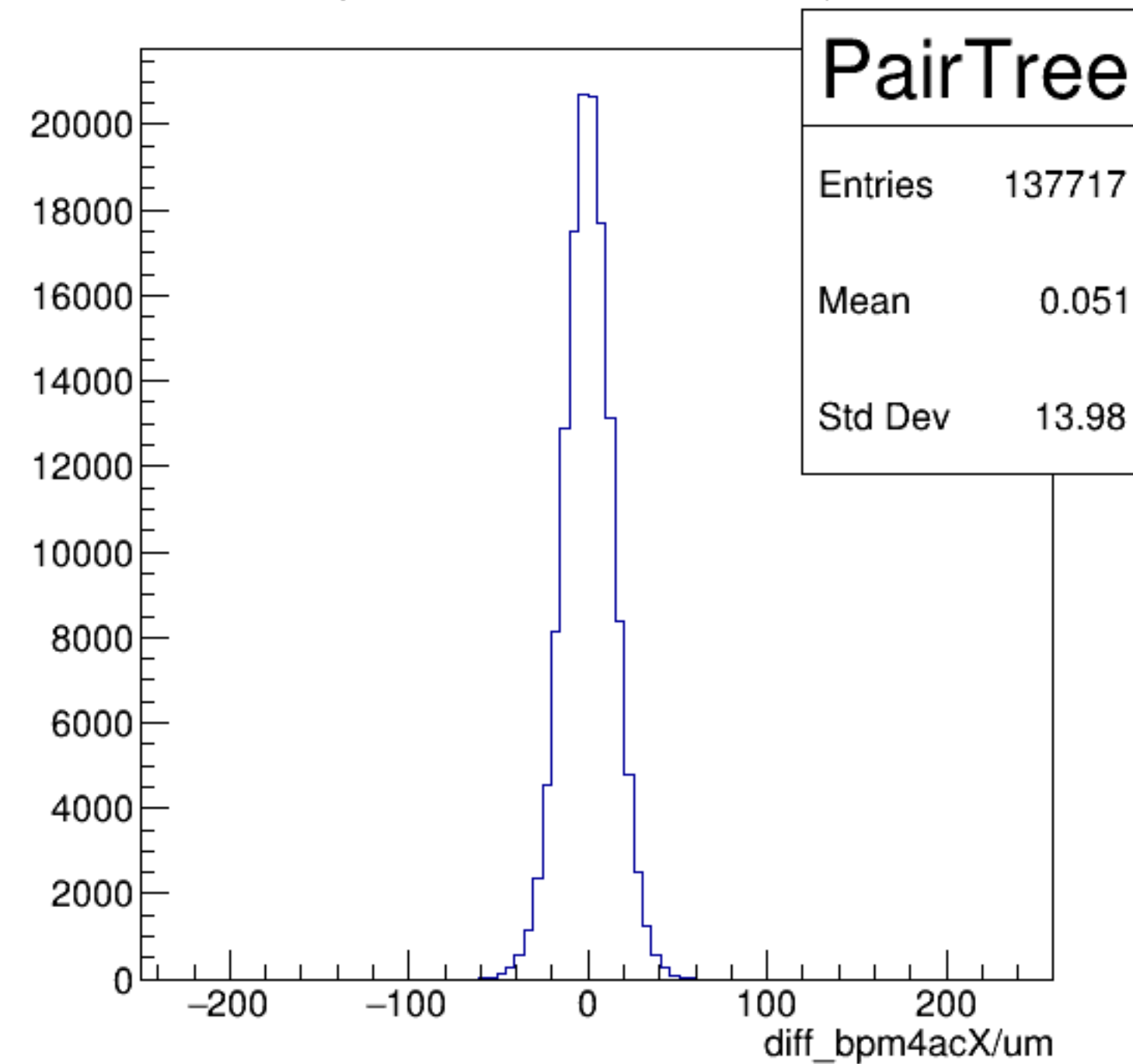
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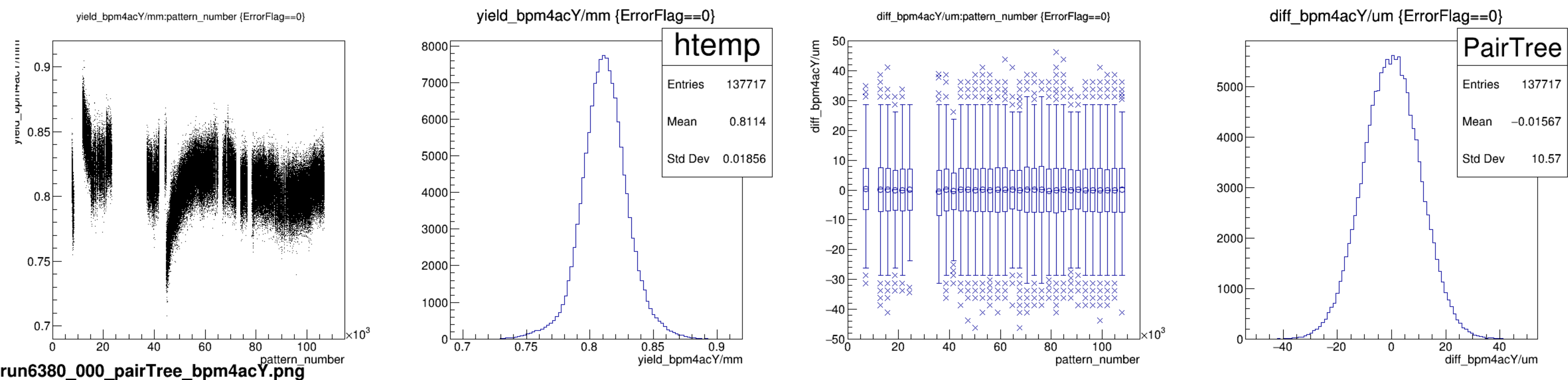


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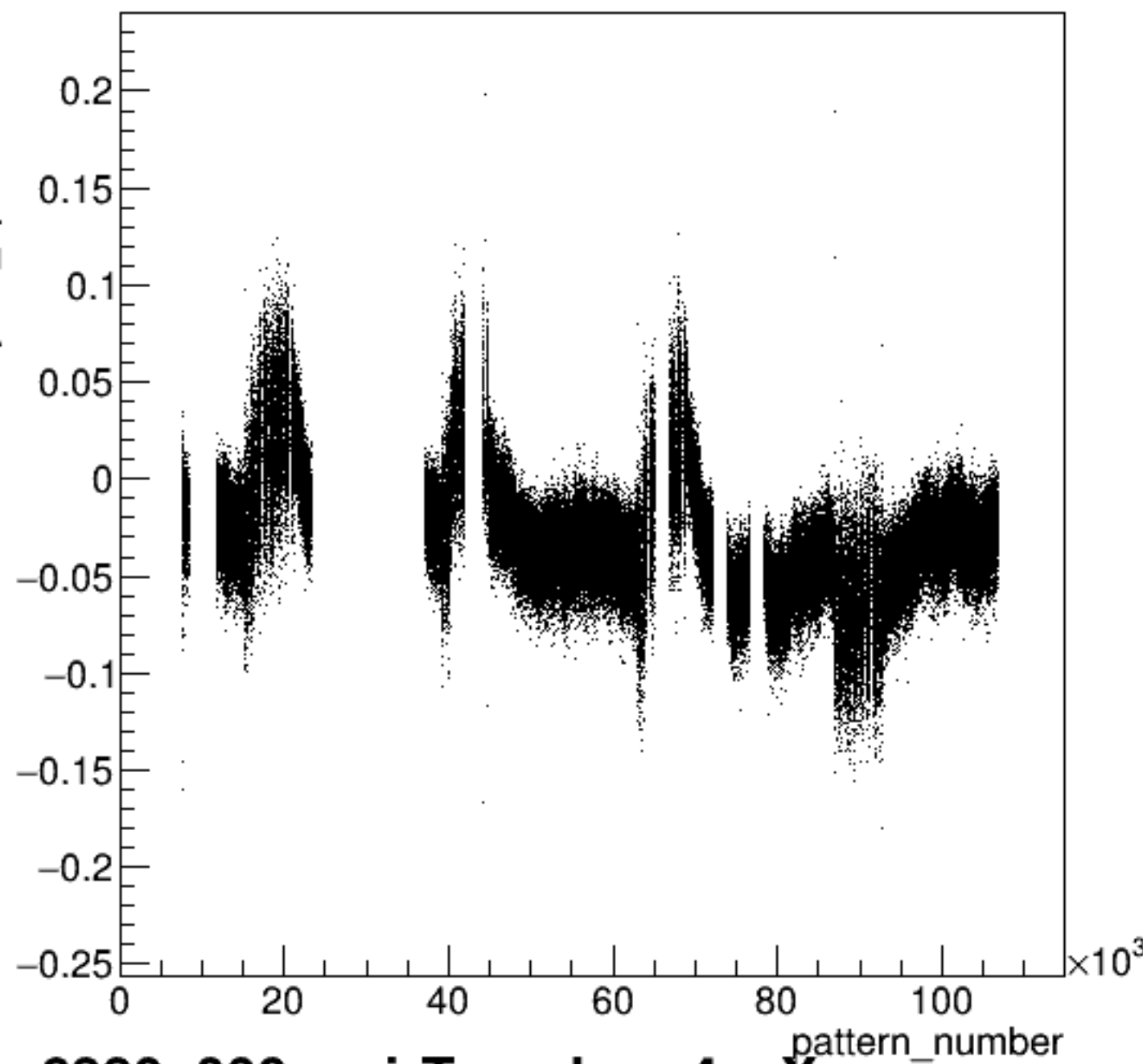


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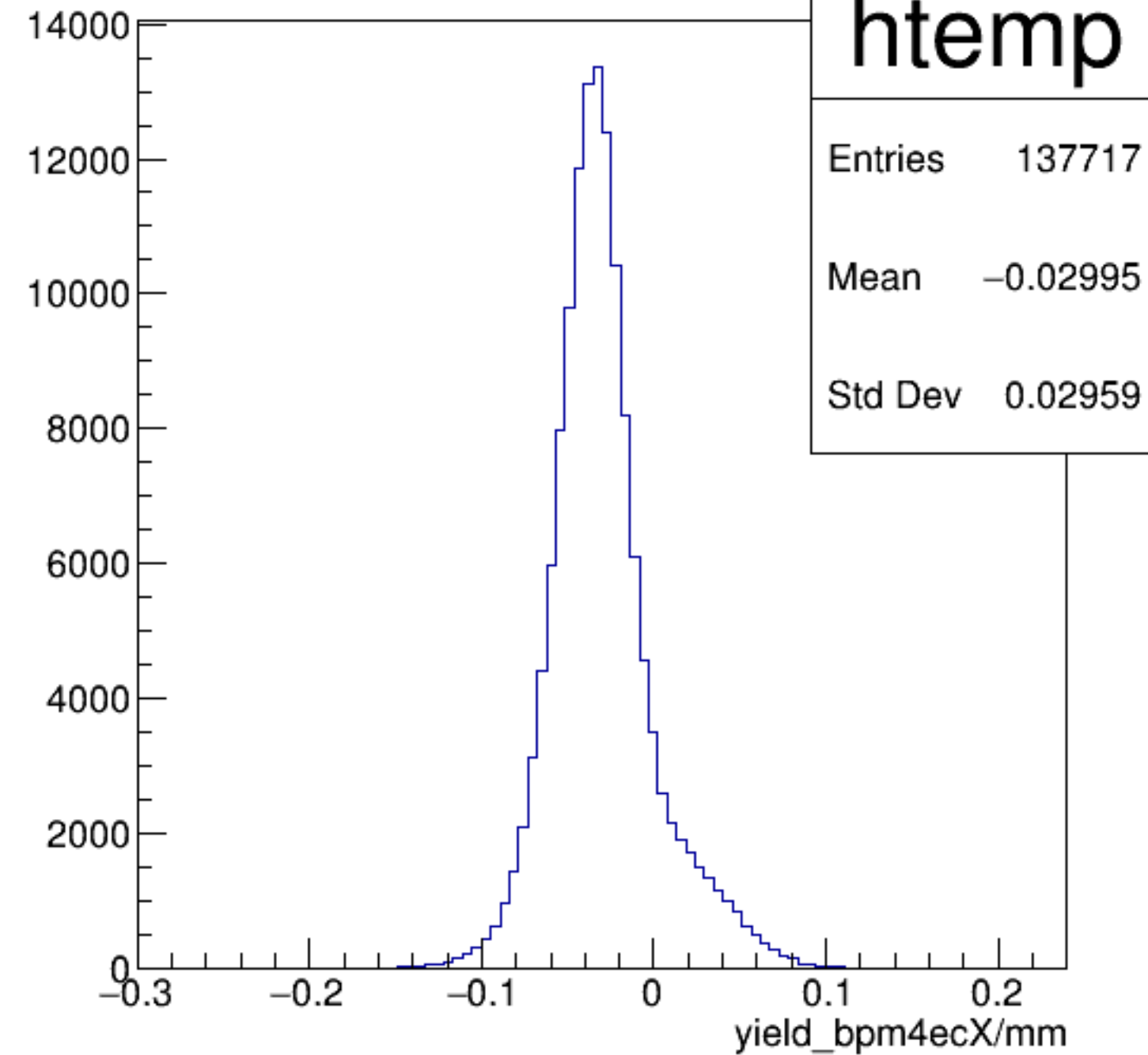


yield_bpm4ecX/mm:pattern_number {ErrorFlag==0}

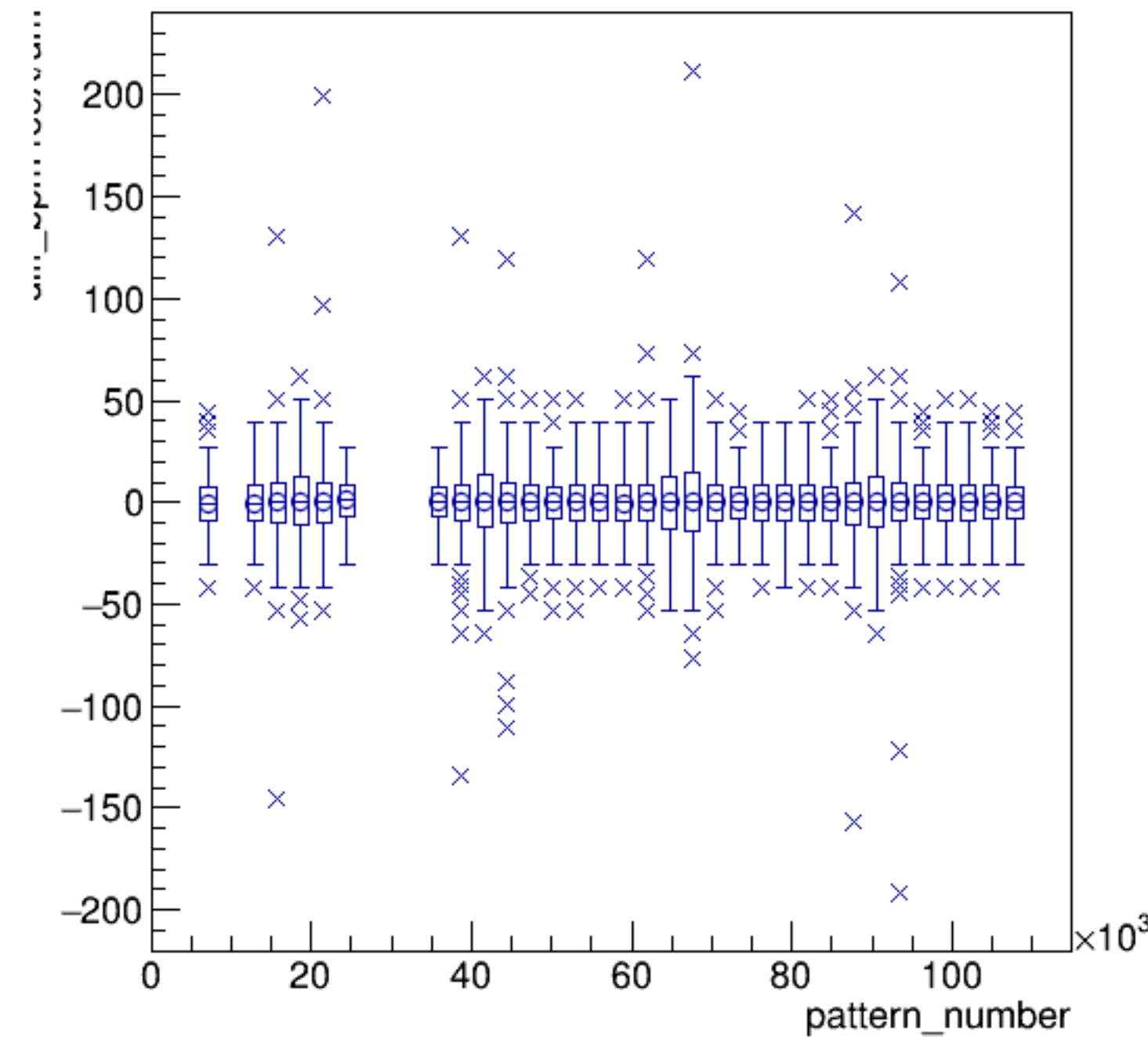


run6380_000_pairTree_bpm4ecX.png

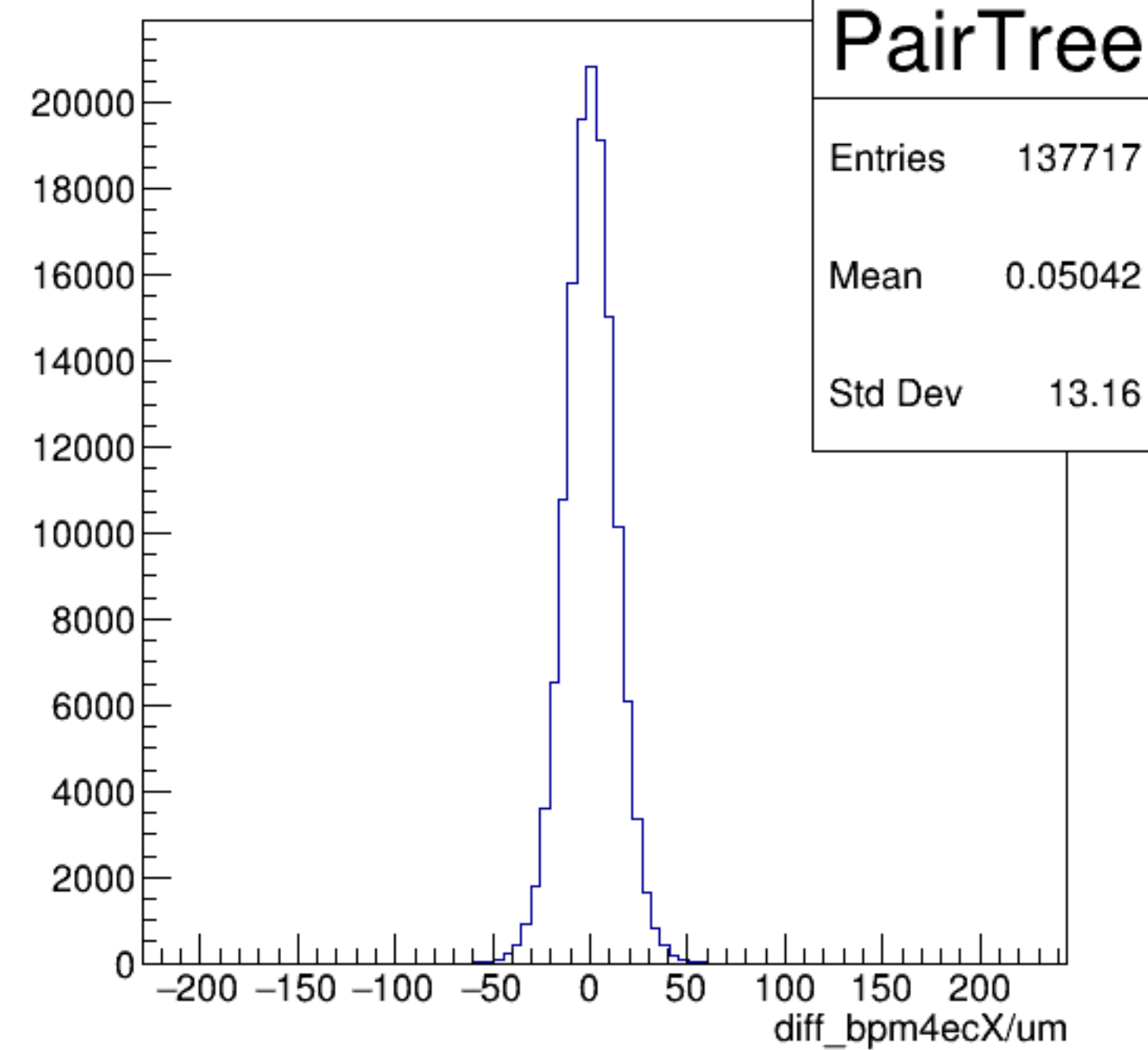
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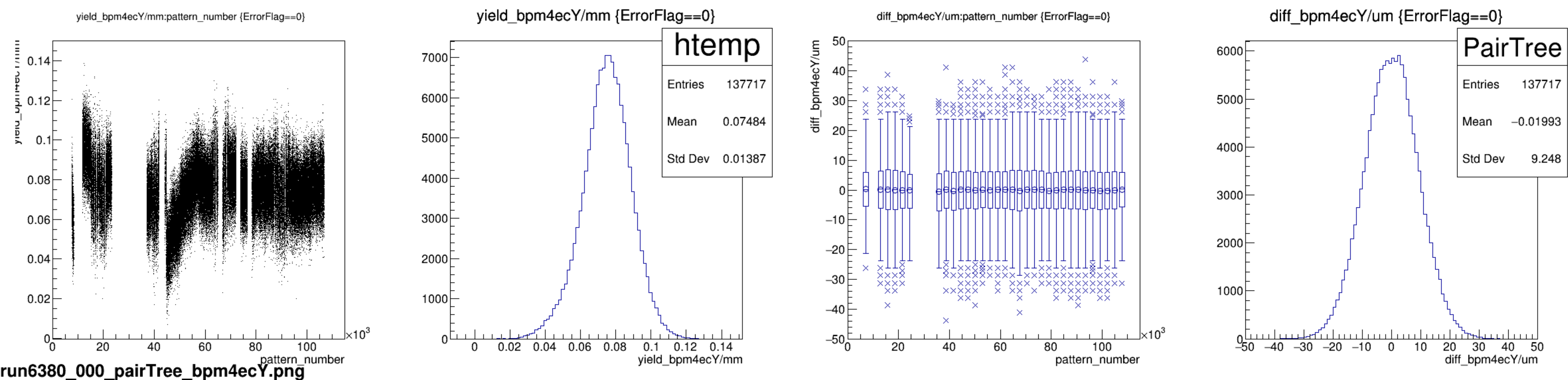


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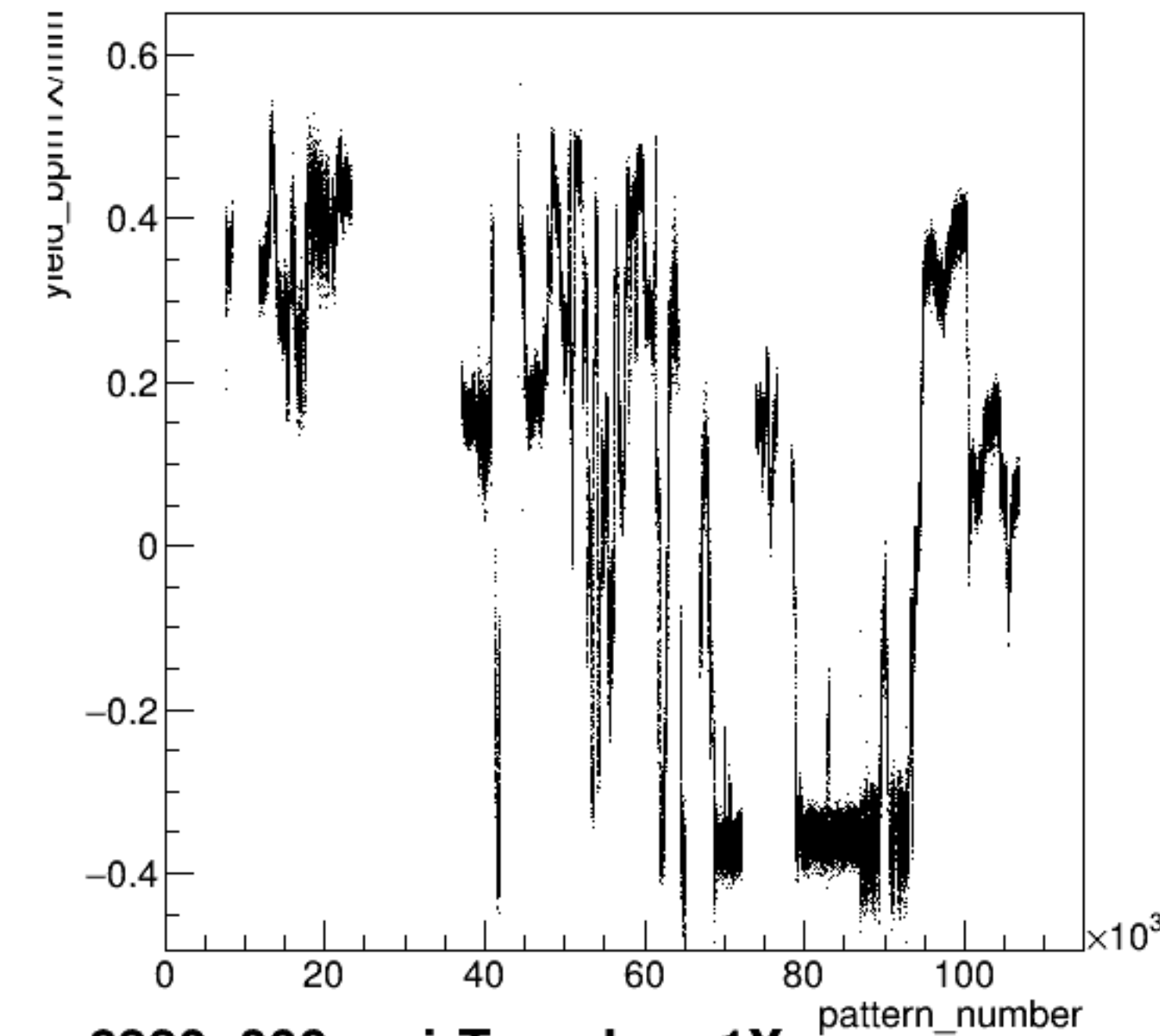


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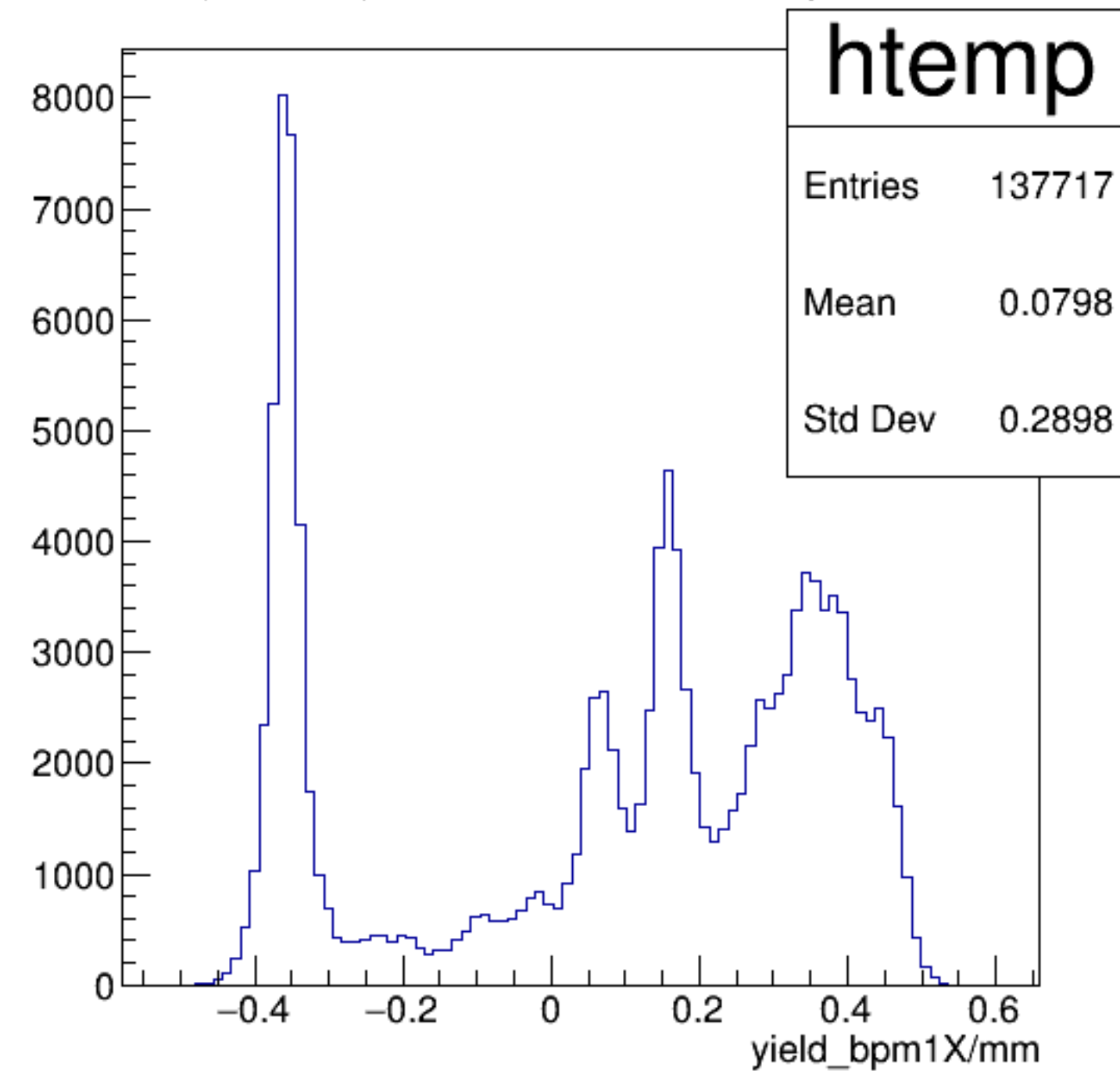


yield_bpm1X/mm:pattern_number {ErrorFlag==0}

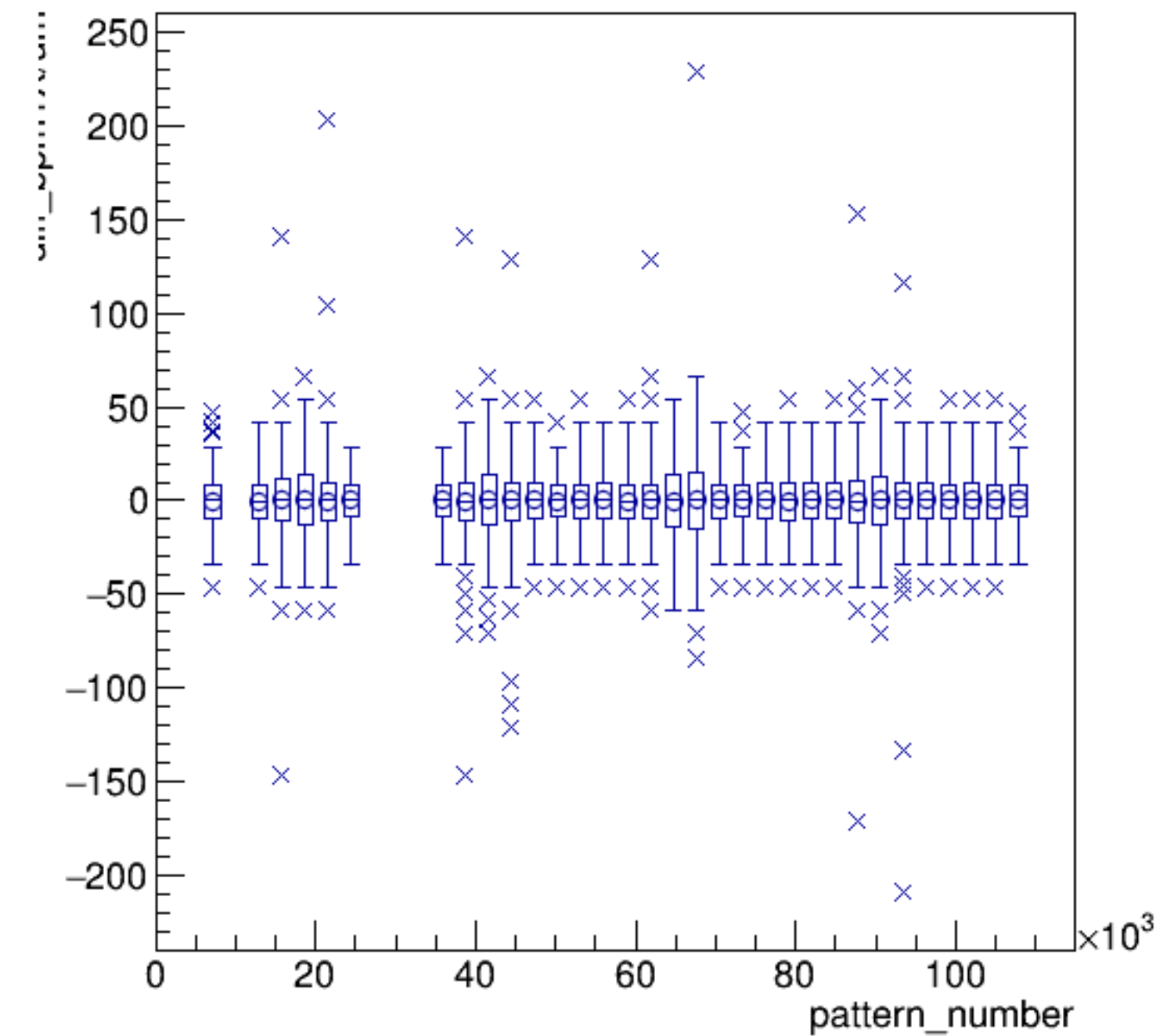


run6380_000_pairTree_bpm1X.png

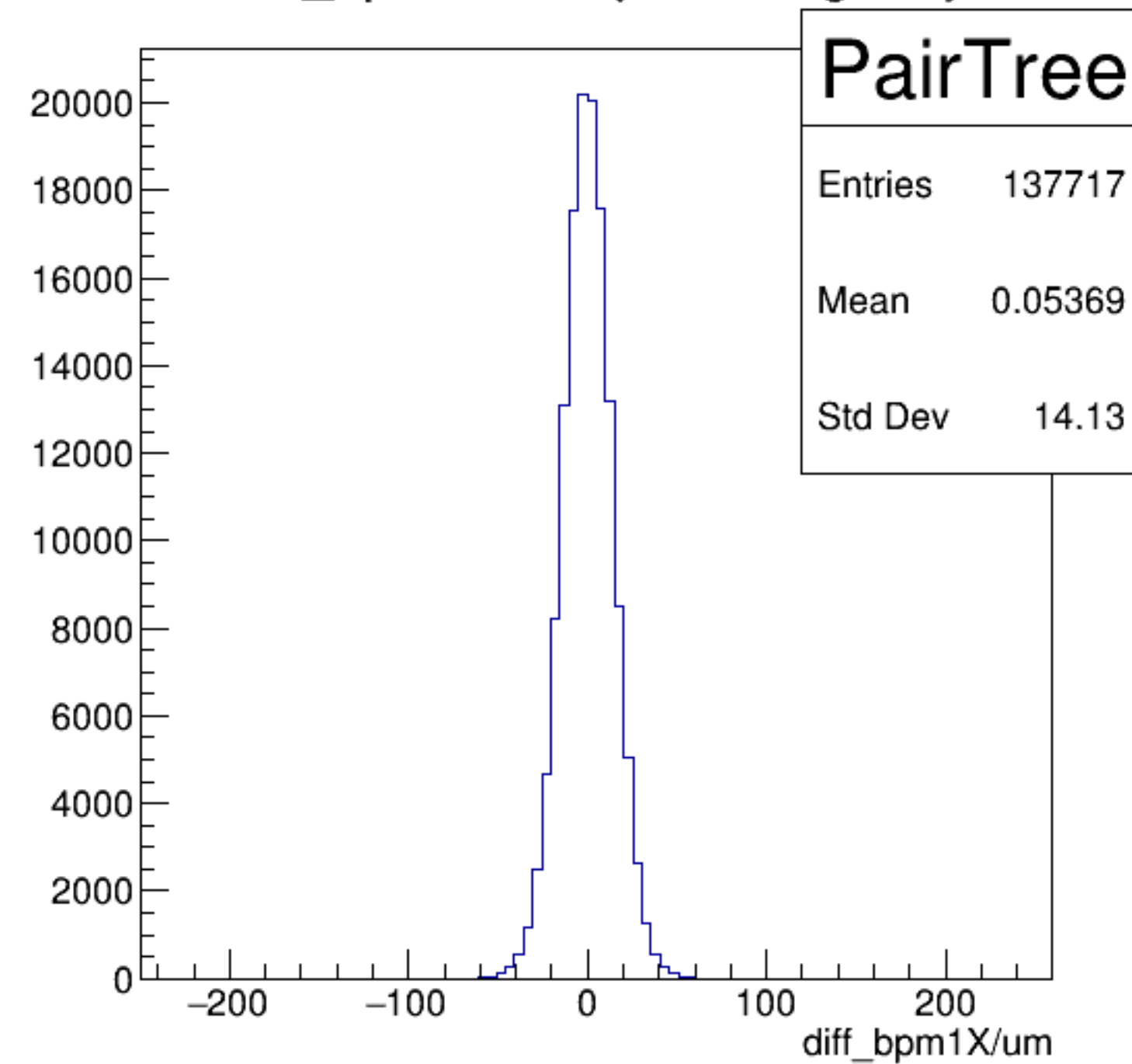
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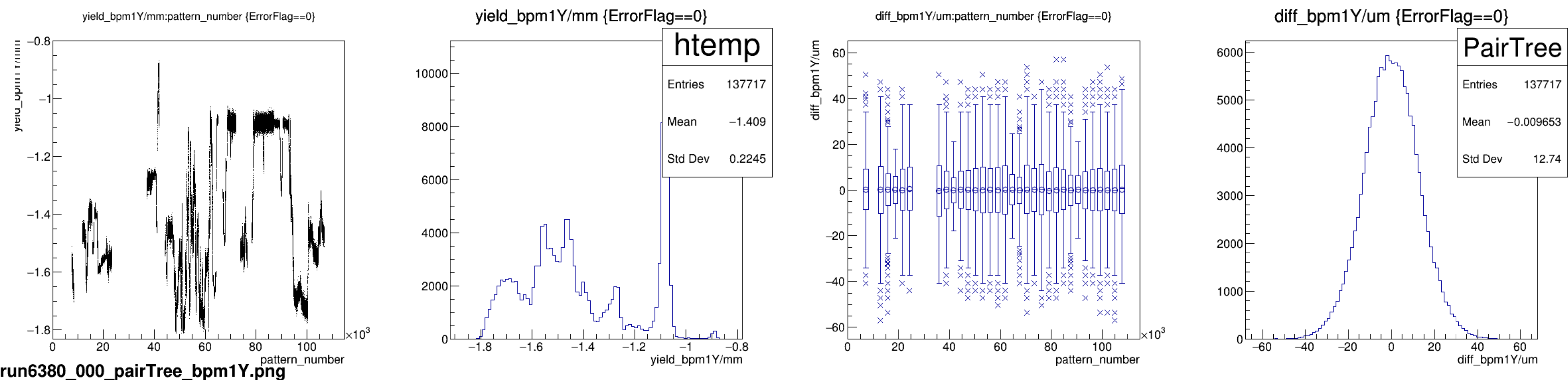


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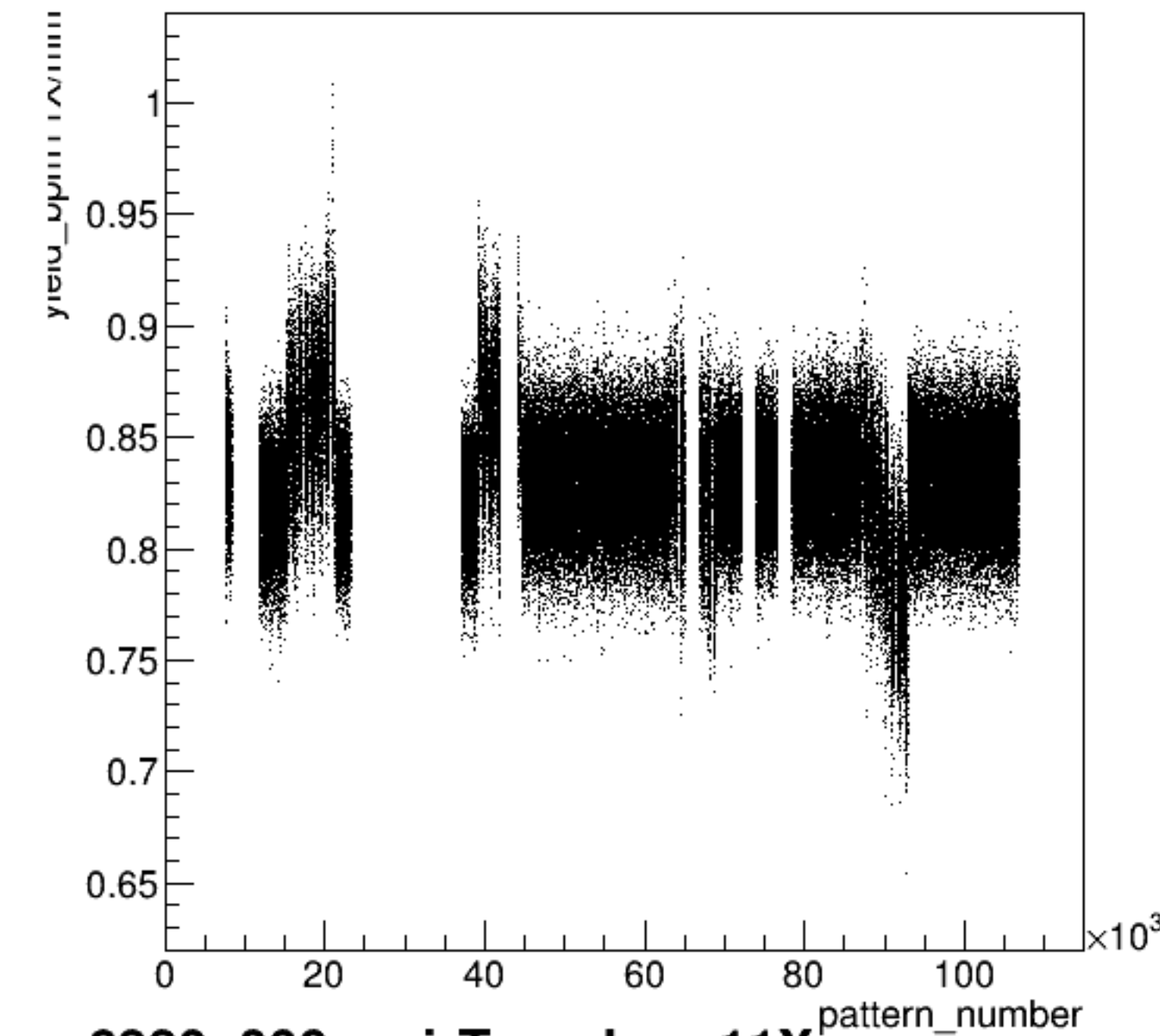


diff_bpm1X/um {ErrorFlag==0}



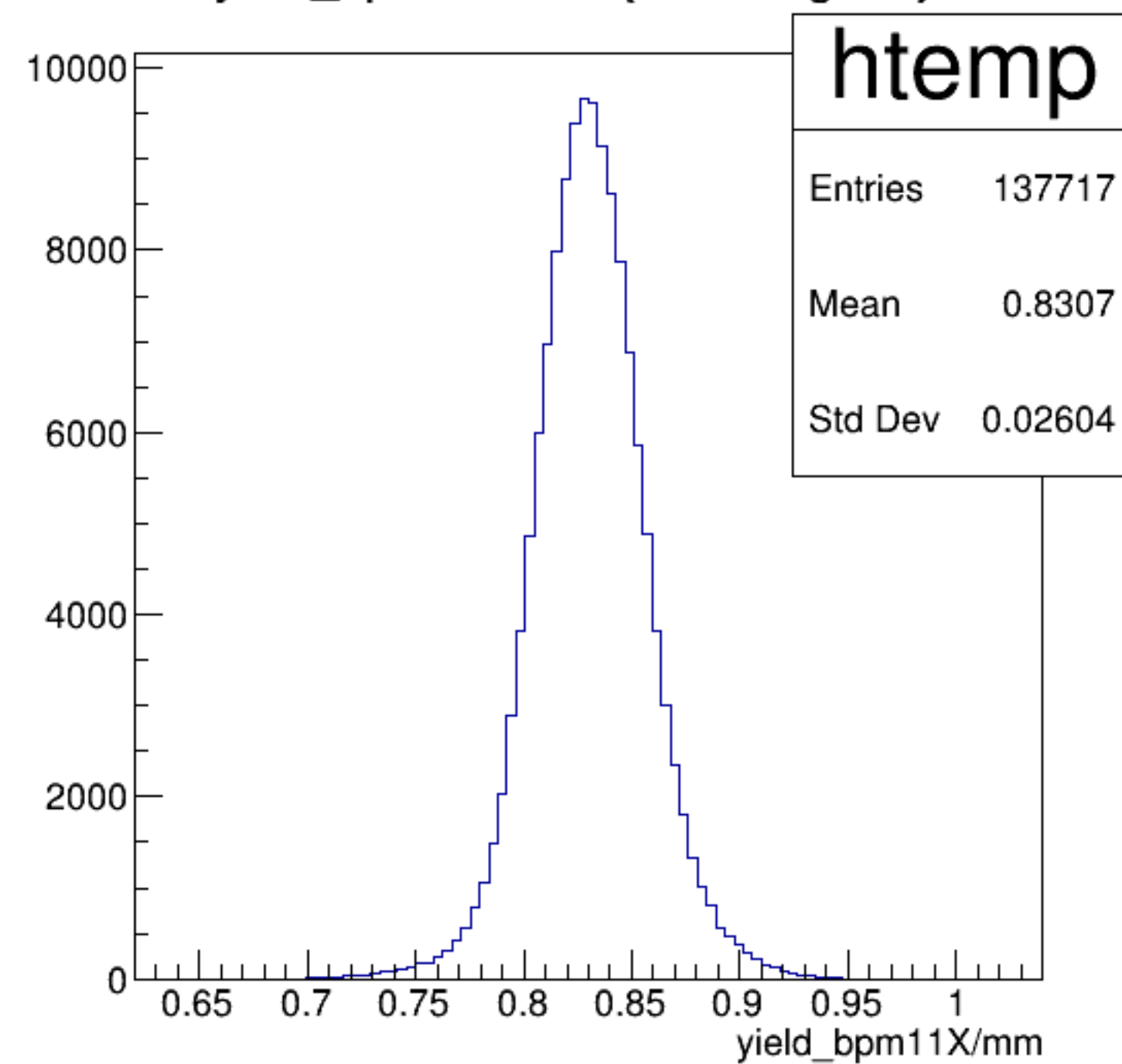


yield_bpm11X/mm:pattern_number {ErrorFlag==0}

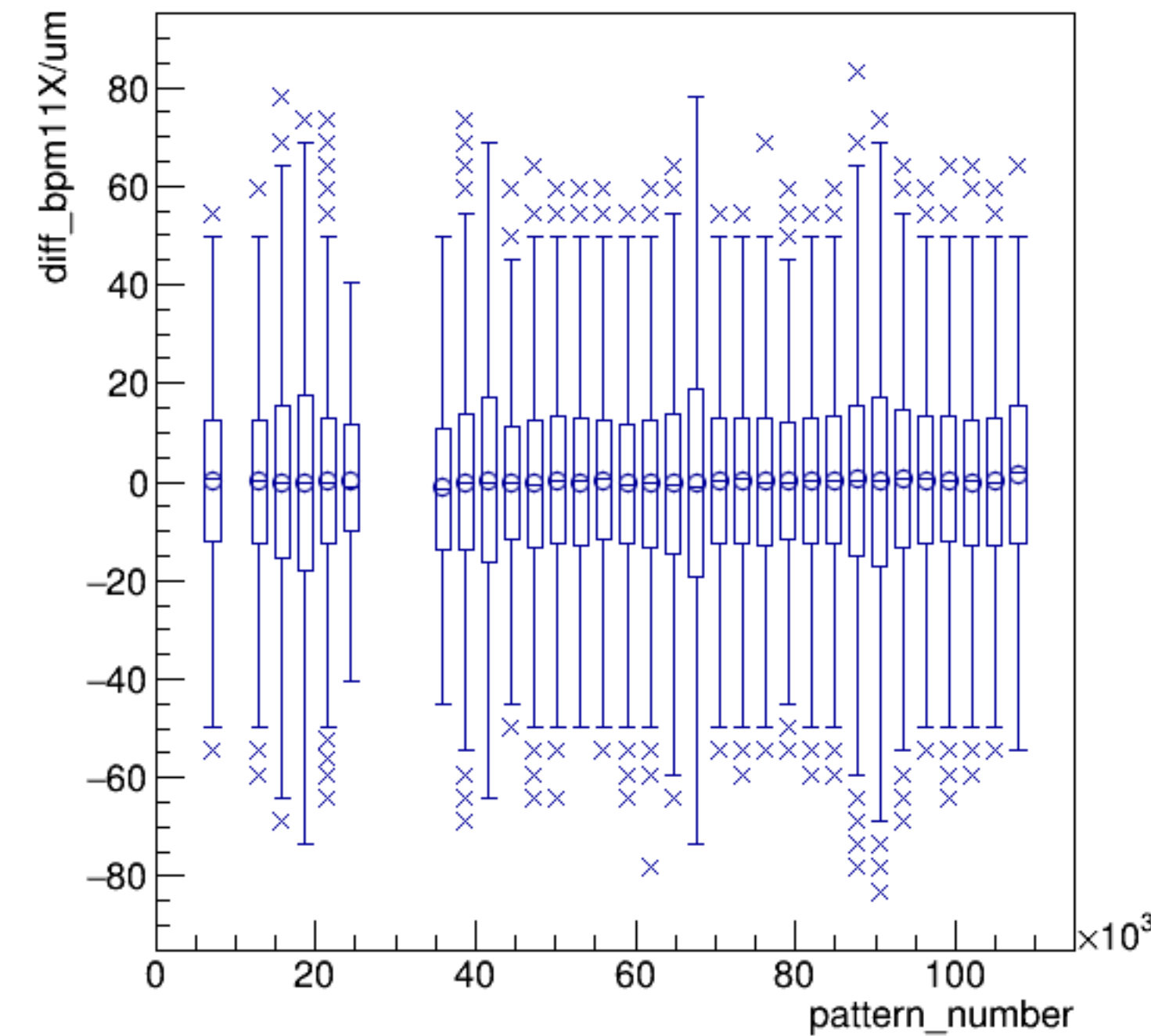


run6380_000_pairTree_bpm11X.png

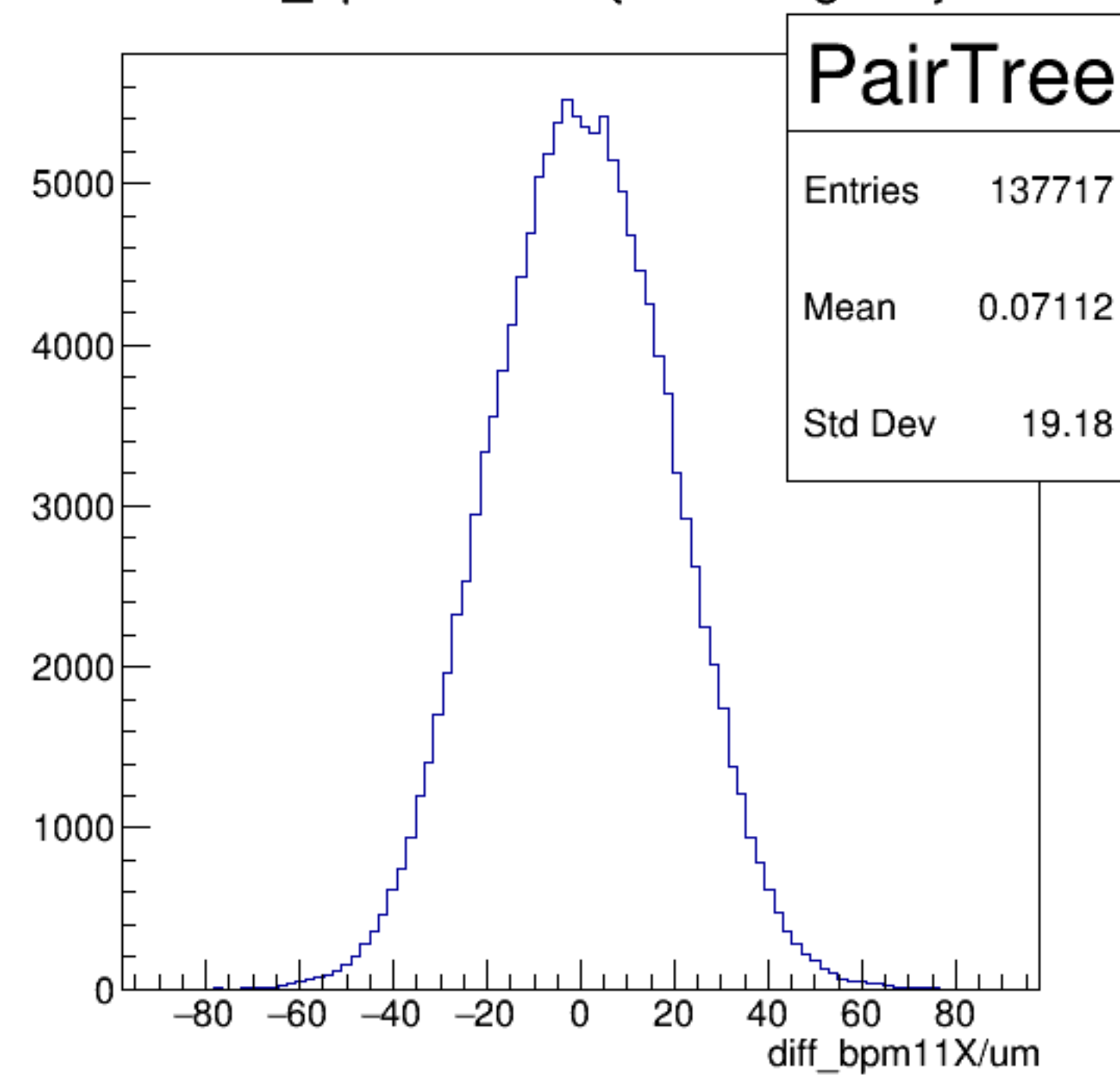
yield_bpm11X/mm {ErrorFlag==0}

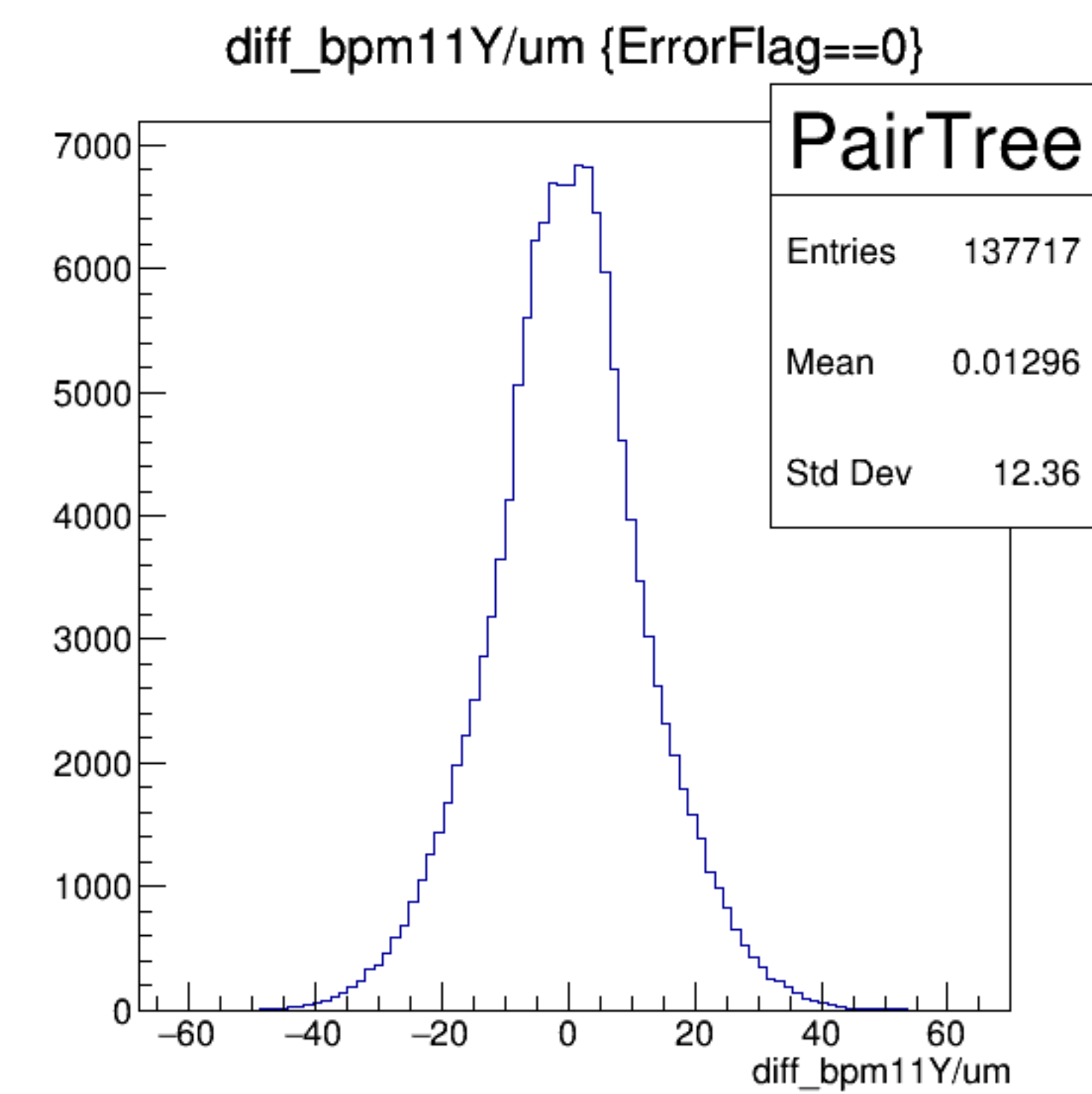
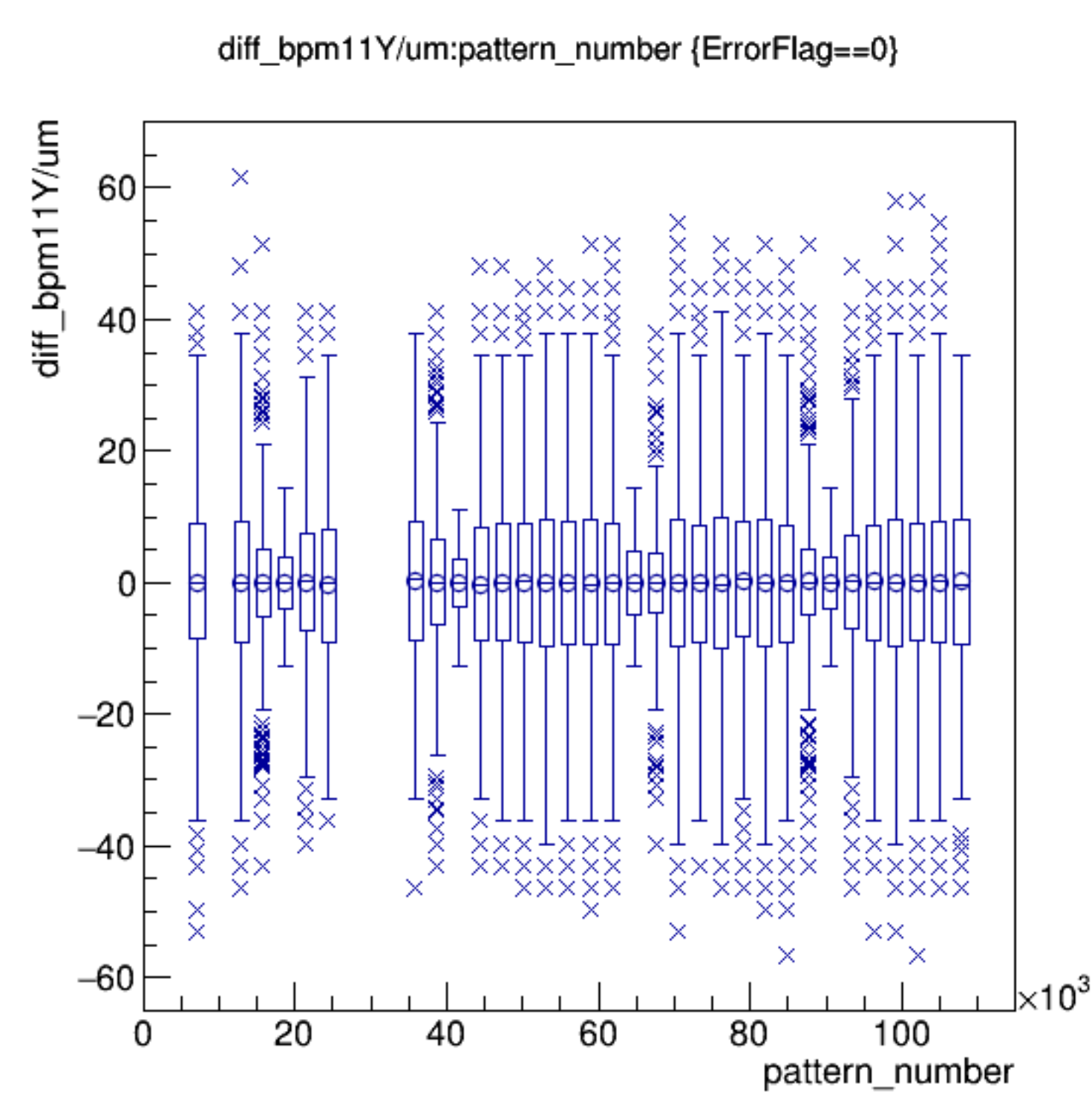
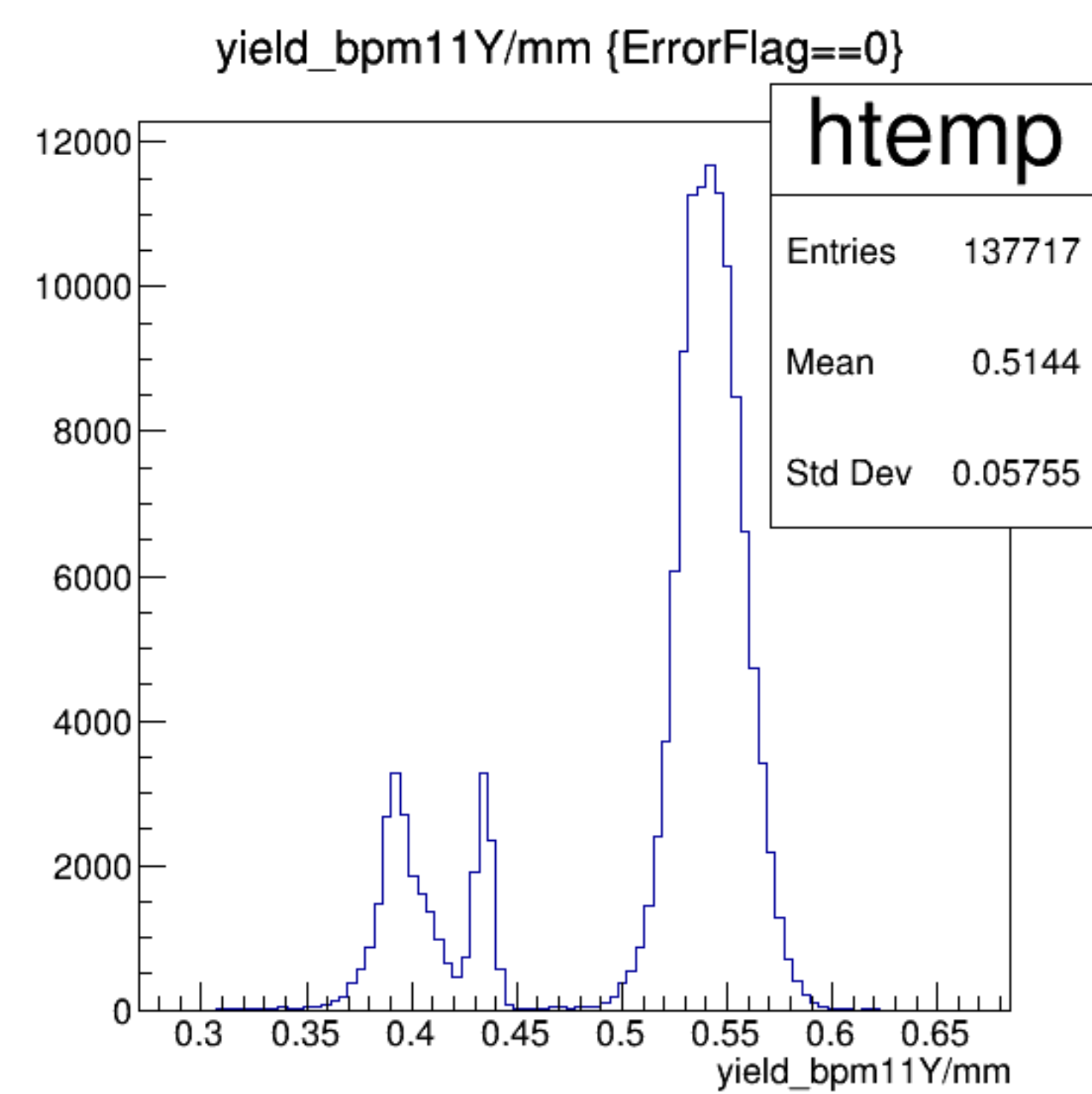
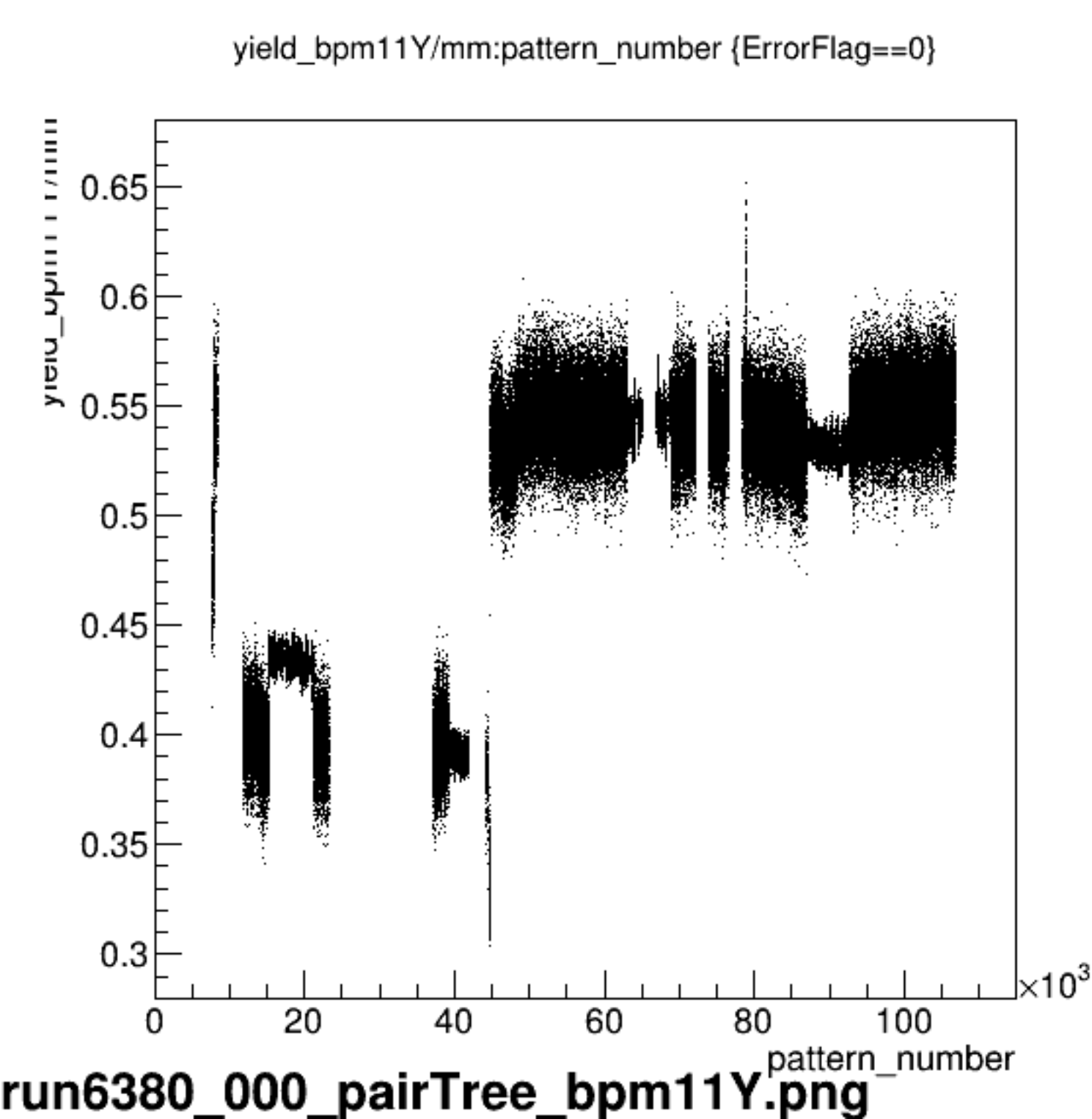


diff_bpm11X/um:pattern_number {ErrorFlag==0}

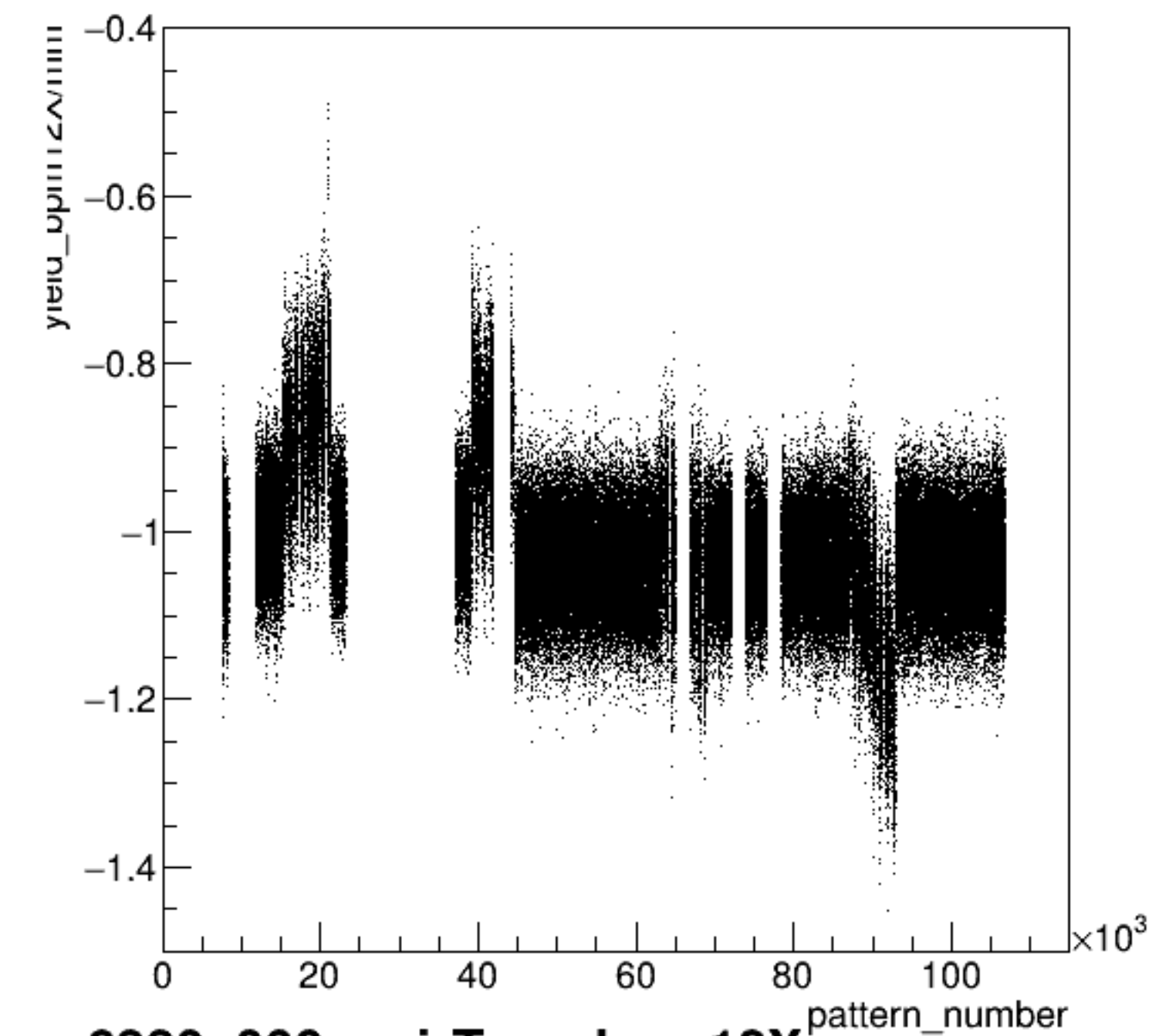


diff_bpm11X/um {ErrorFlag==0}



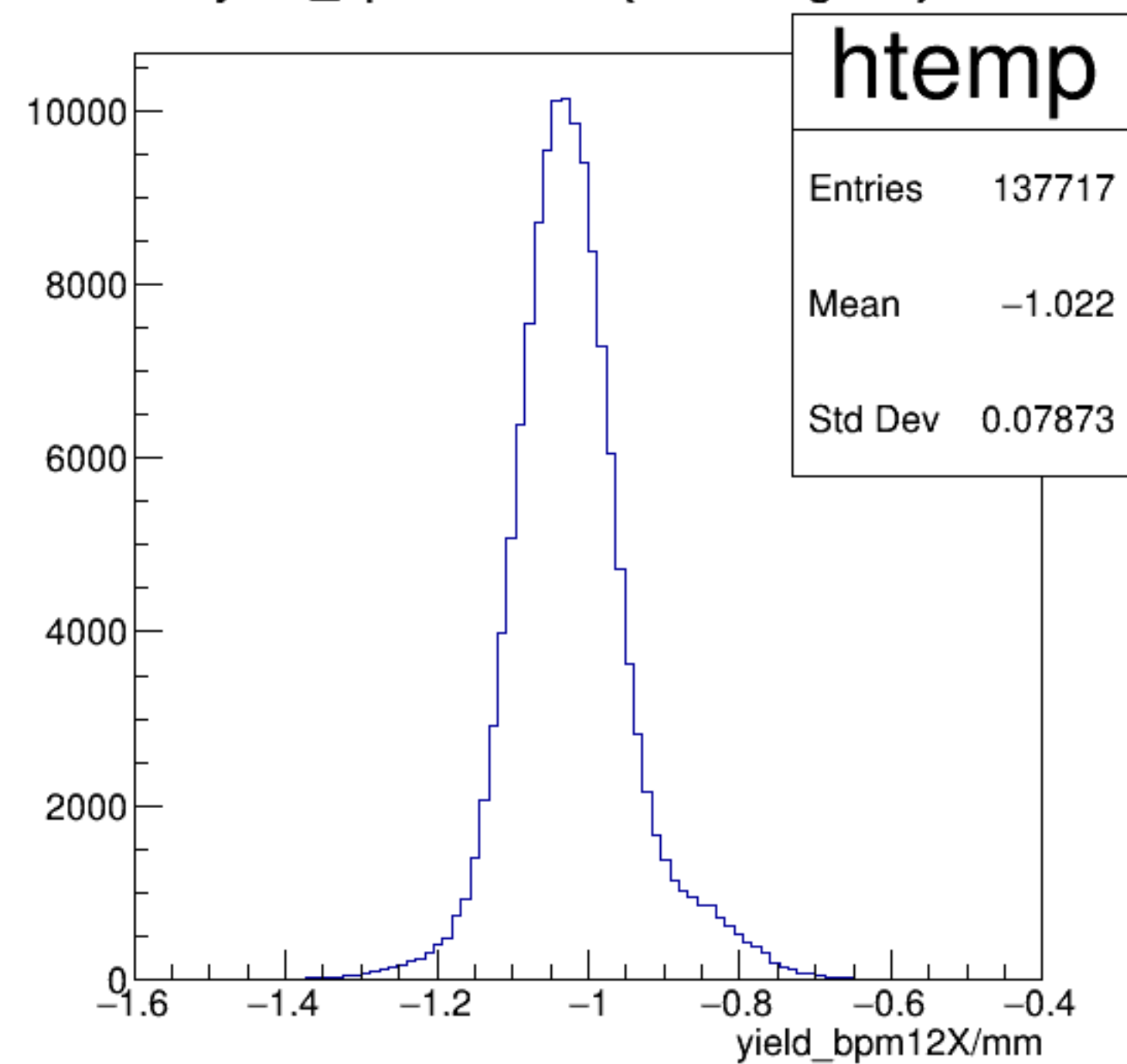


yield_bpm12X/mm:pattern_number {ErrorFlag==0}

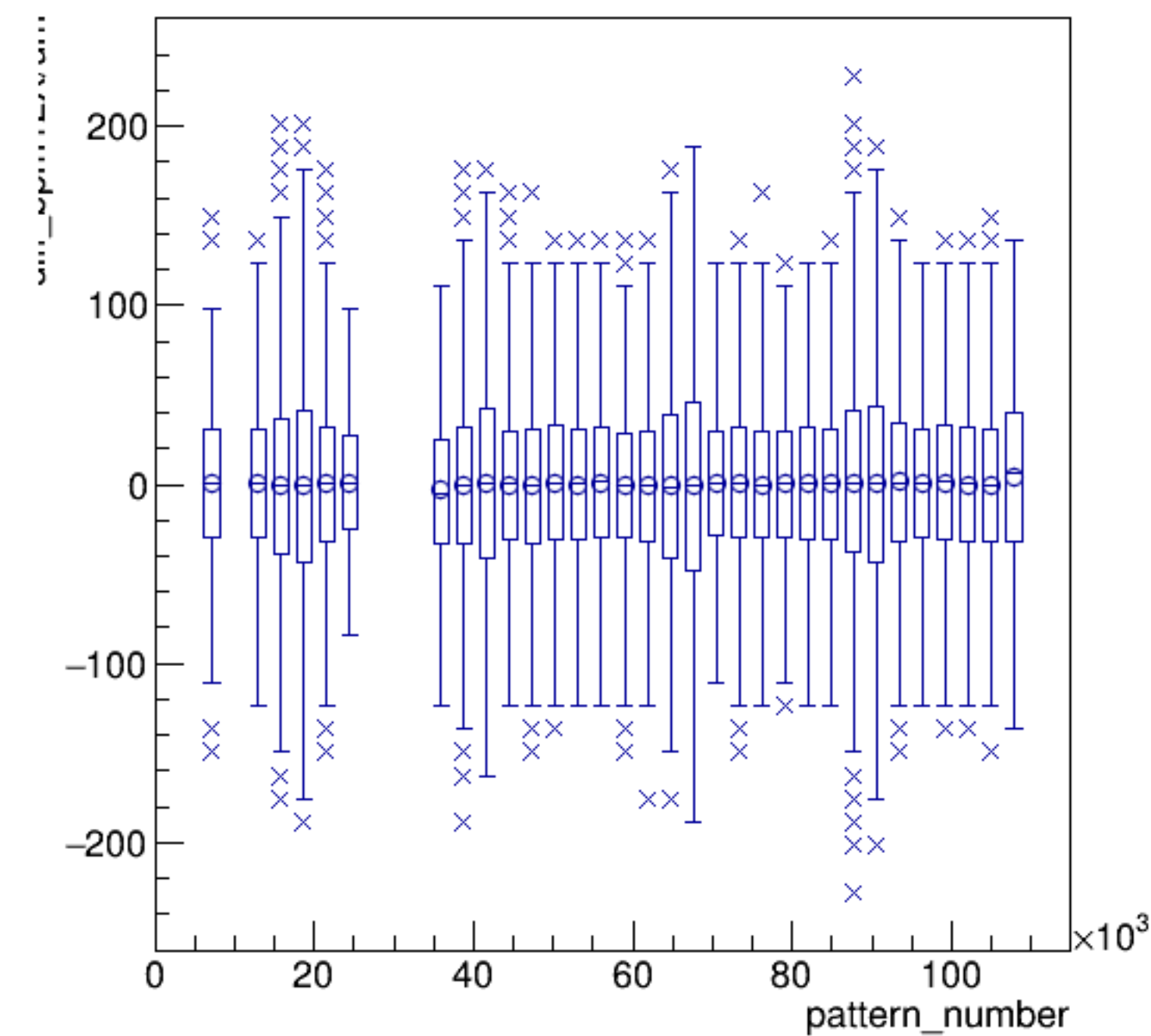


run6380_000_pairTree_bpm12X.png

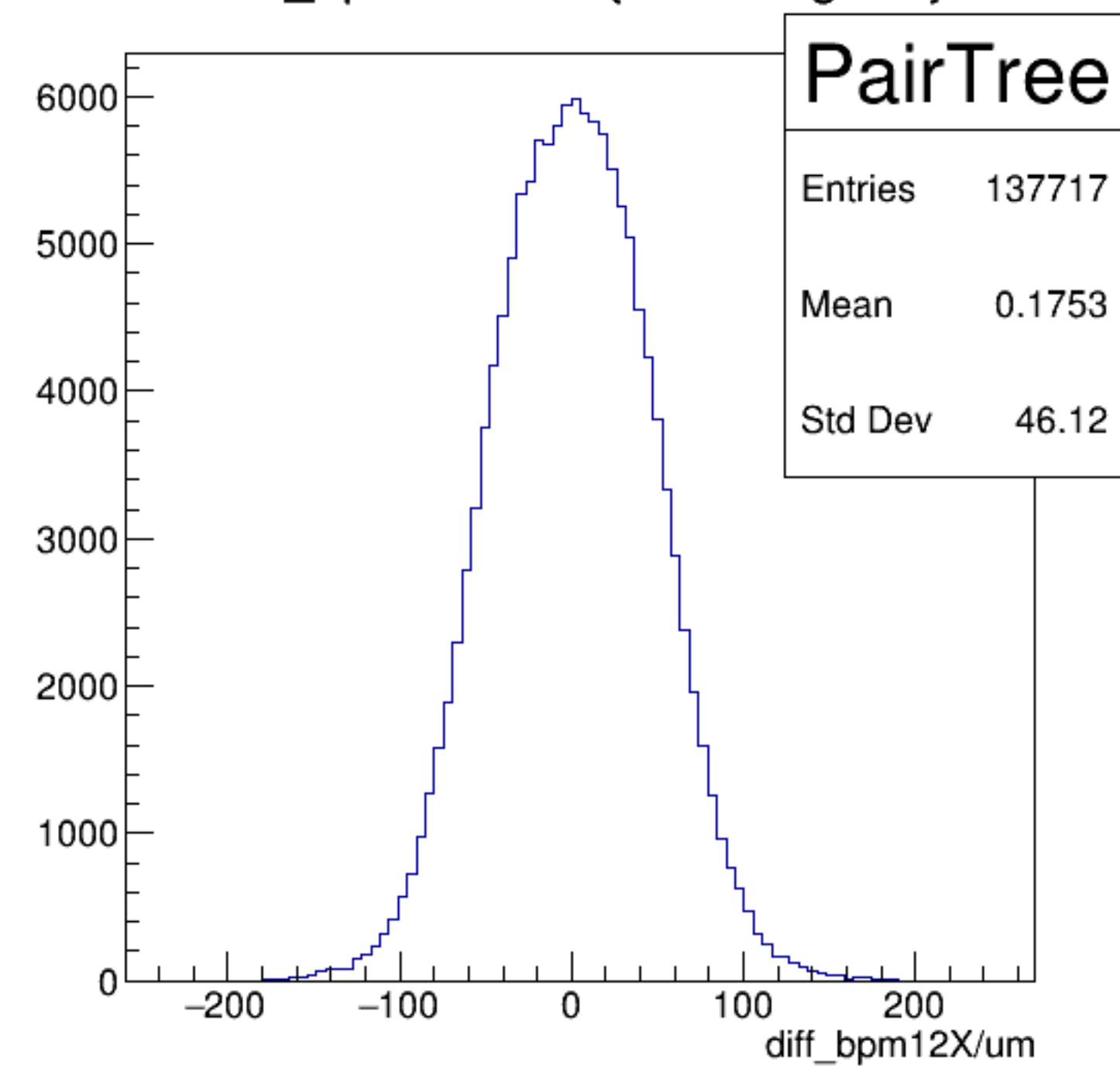
yield_bpm12X/mm {ErrorFlag==0}

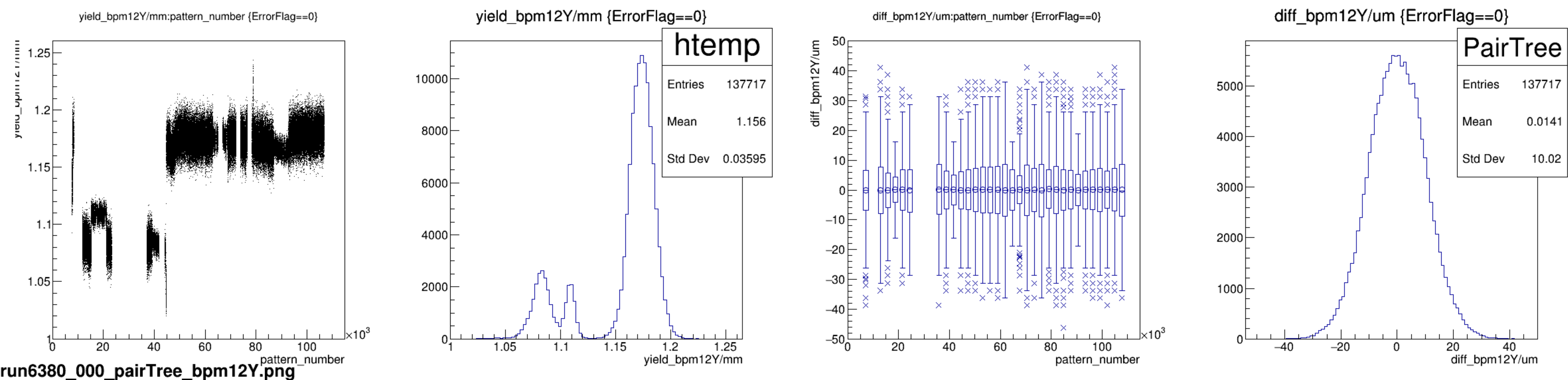


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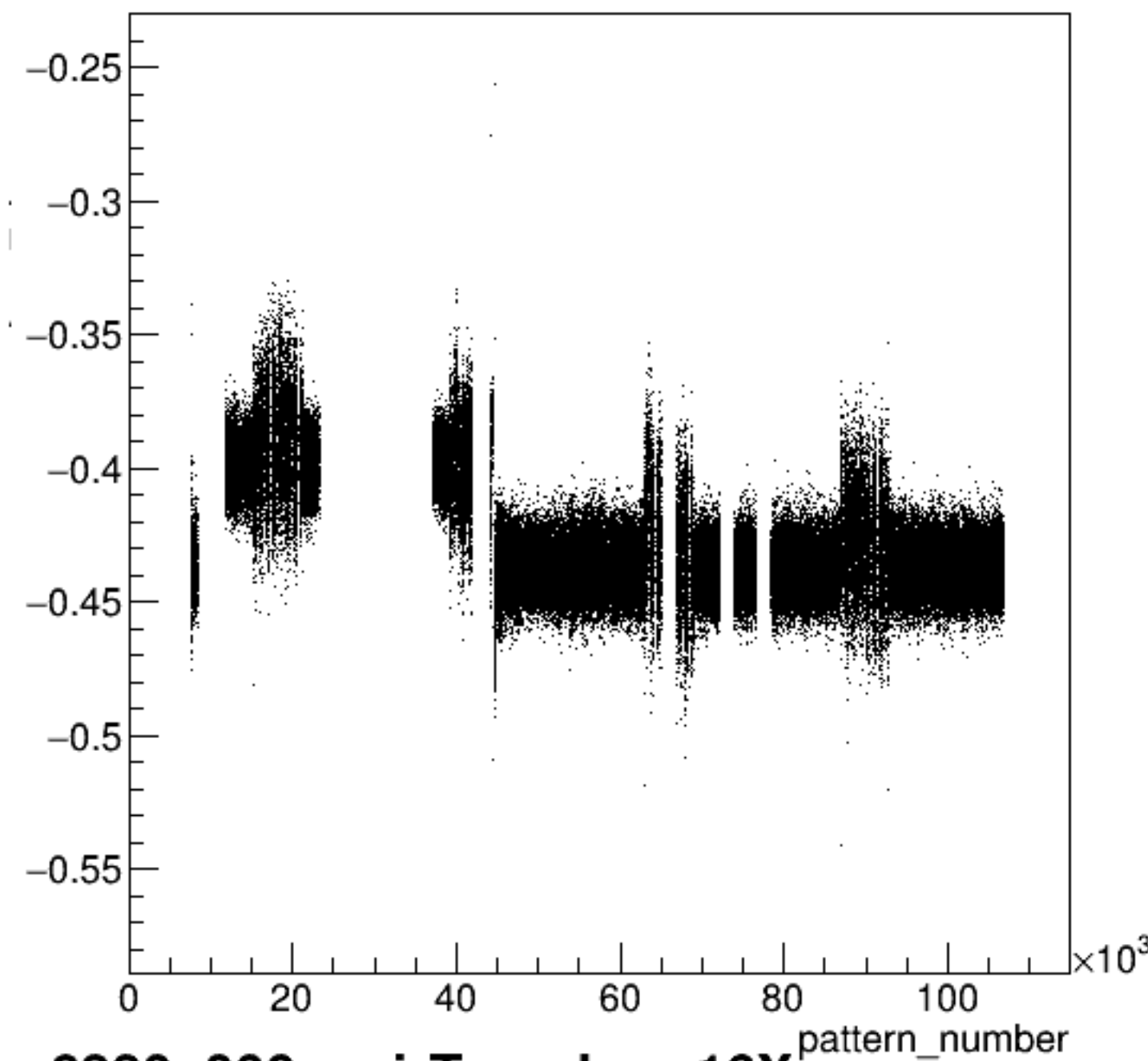


diff_bpm12X/um {ErrorFlag==0}



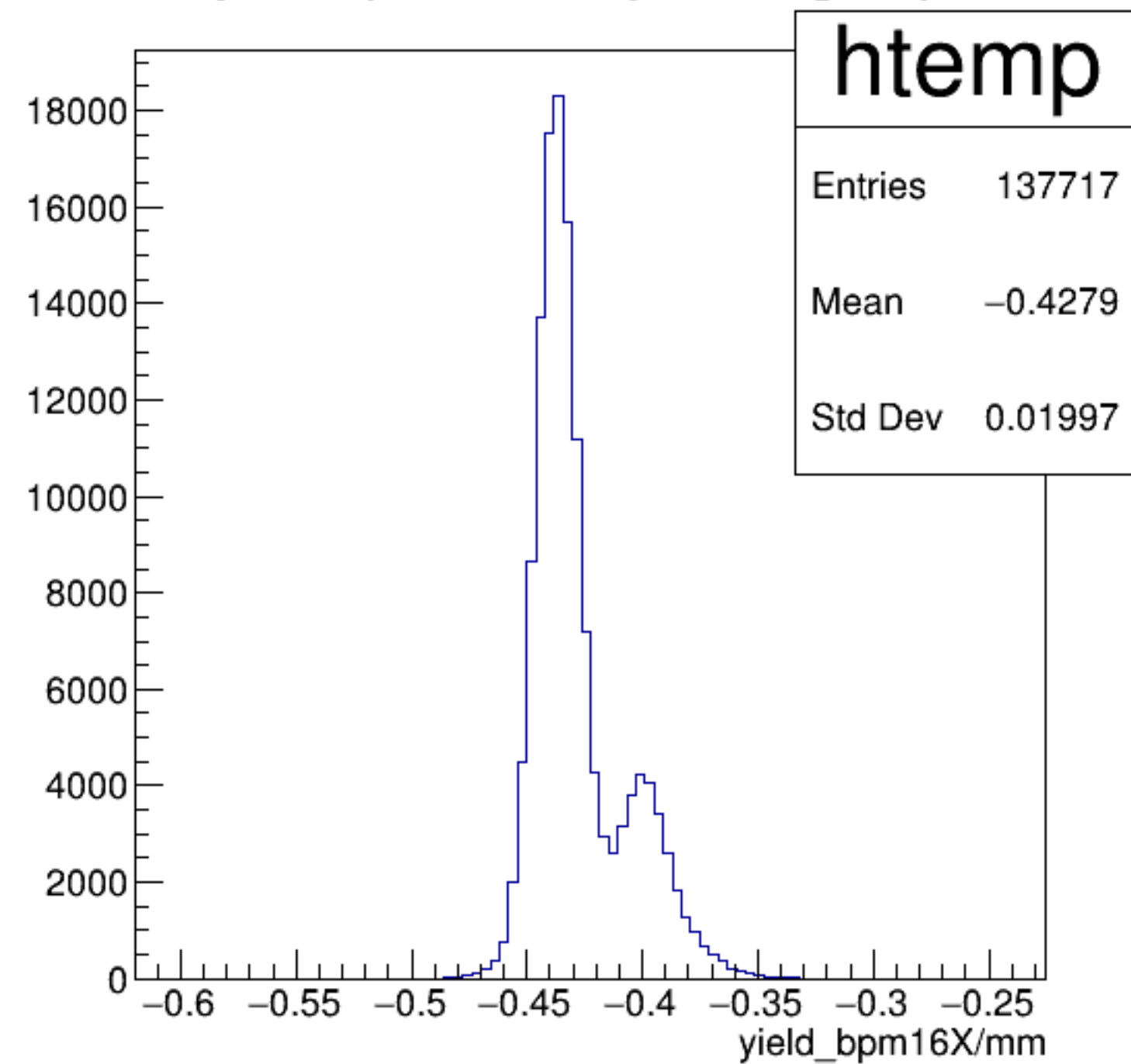


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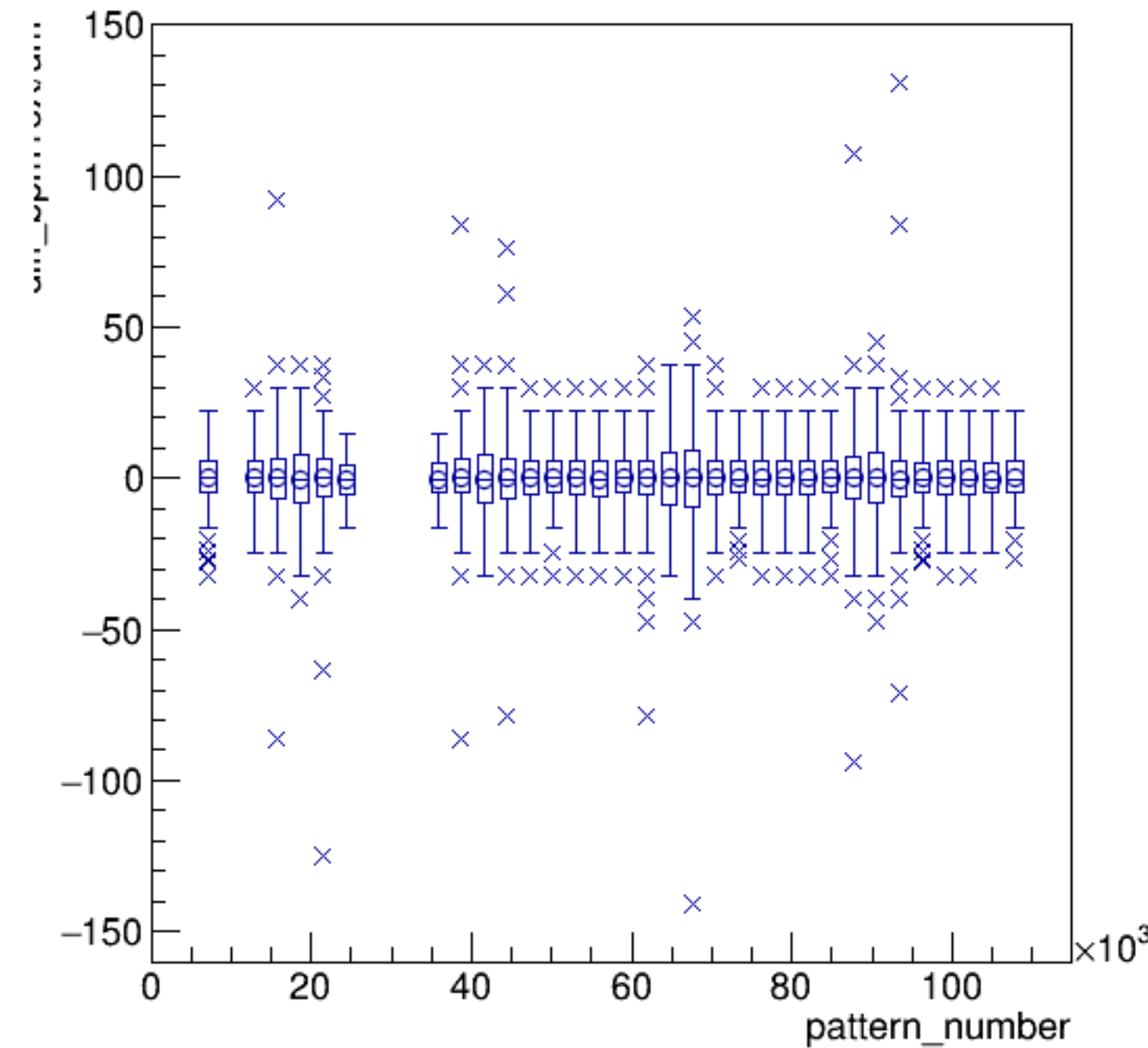


run6380_000_pairTree_bpm16X.png

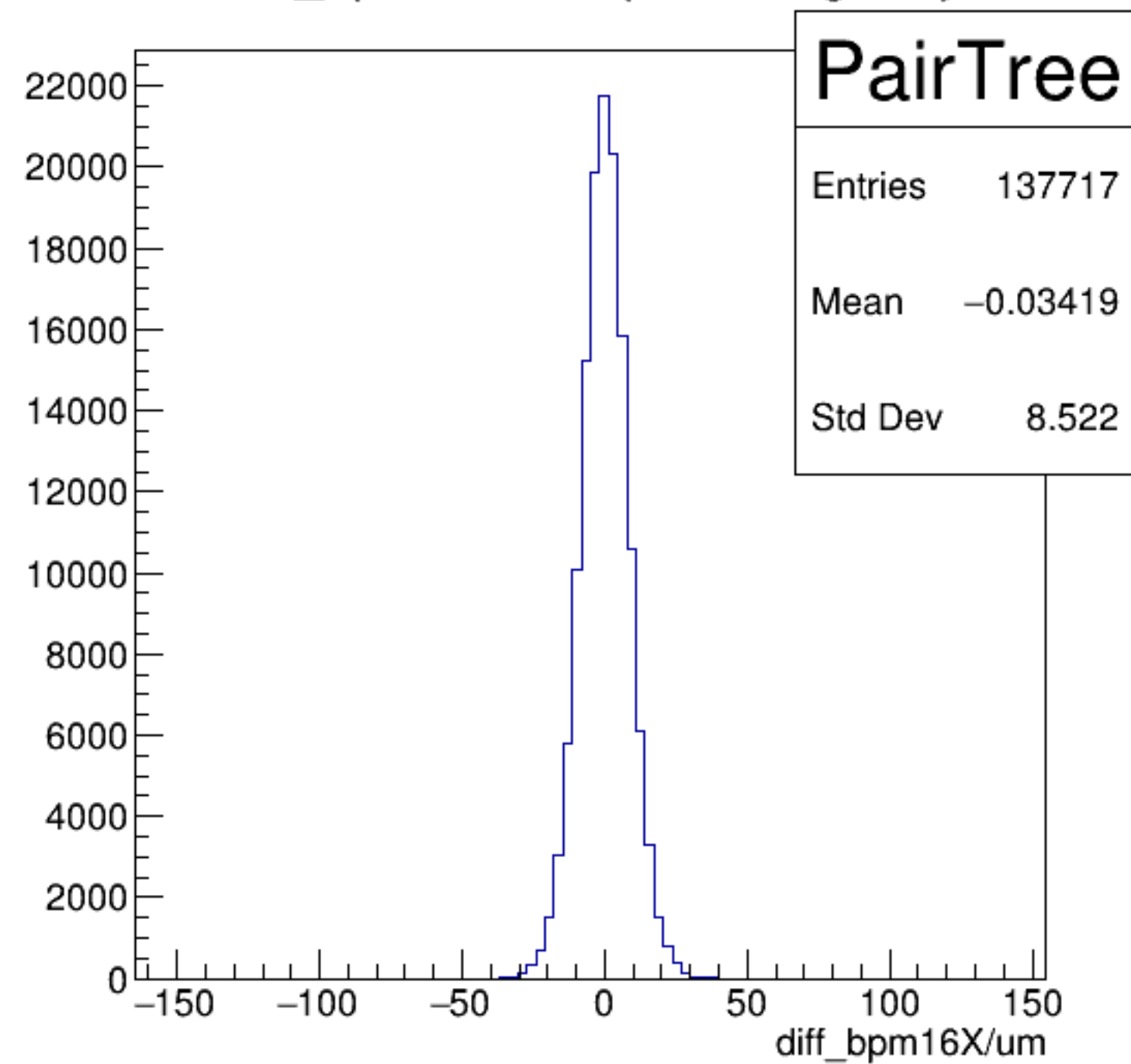
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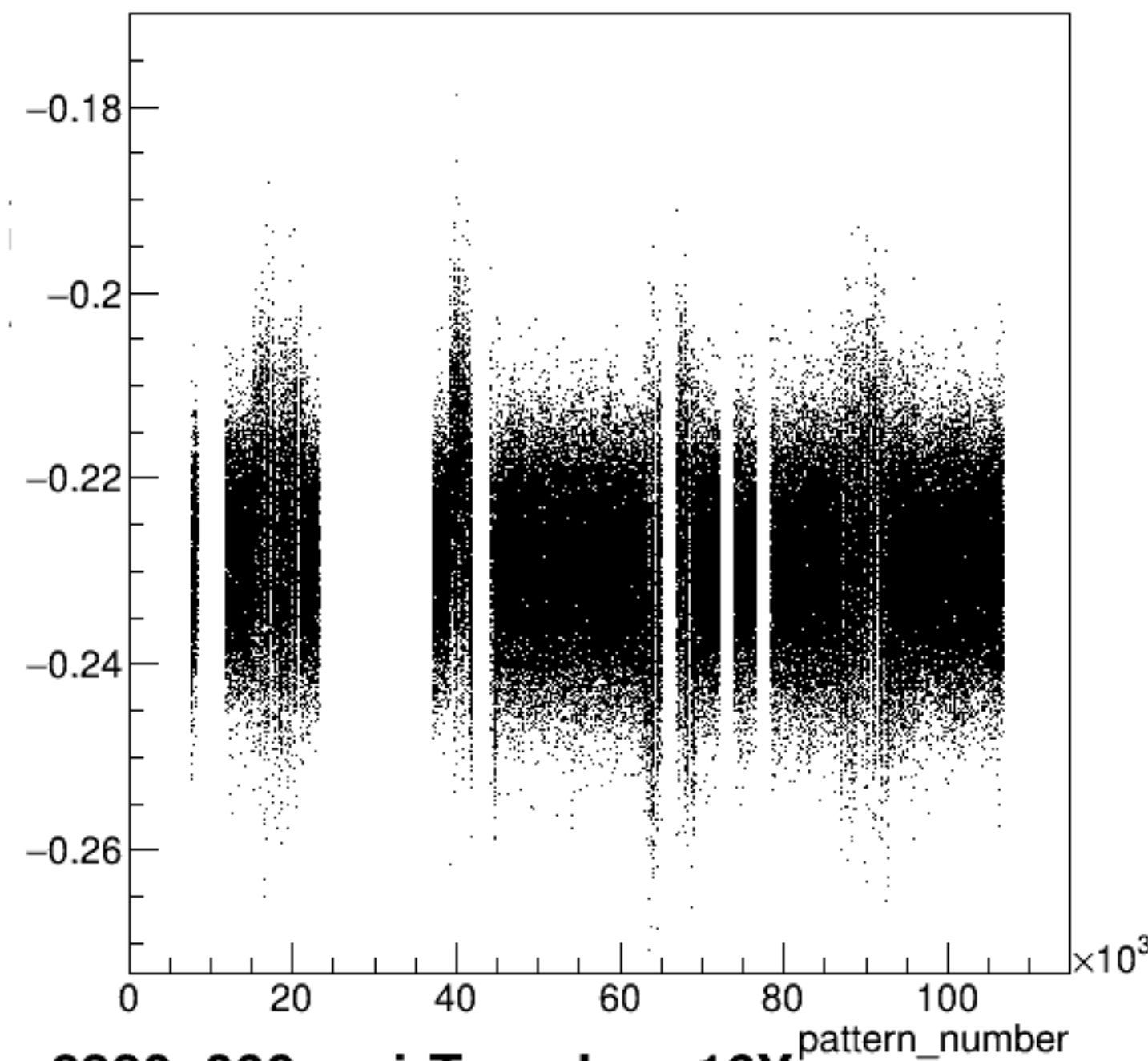
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diff_bpm16X/um {ErrorFlag==0}

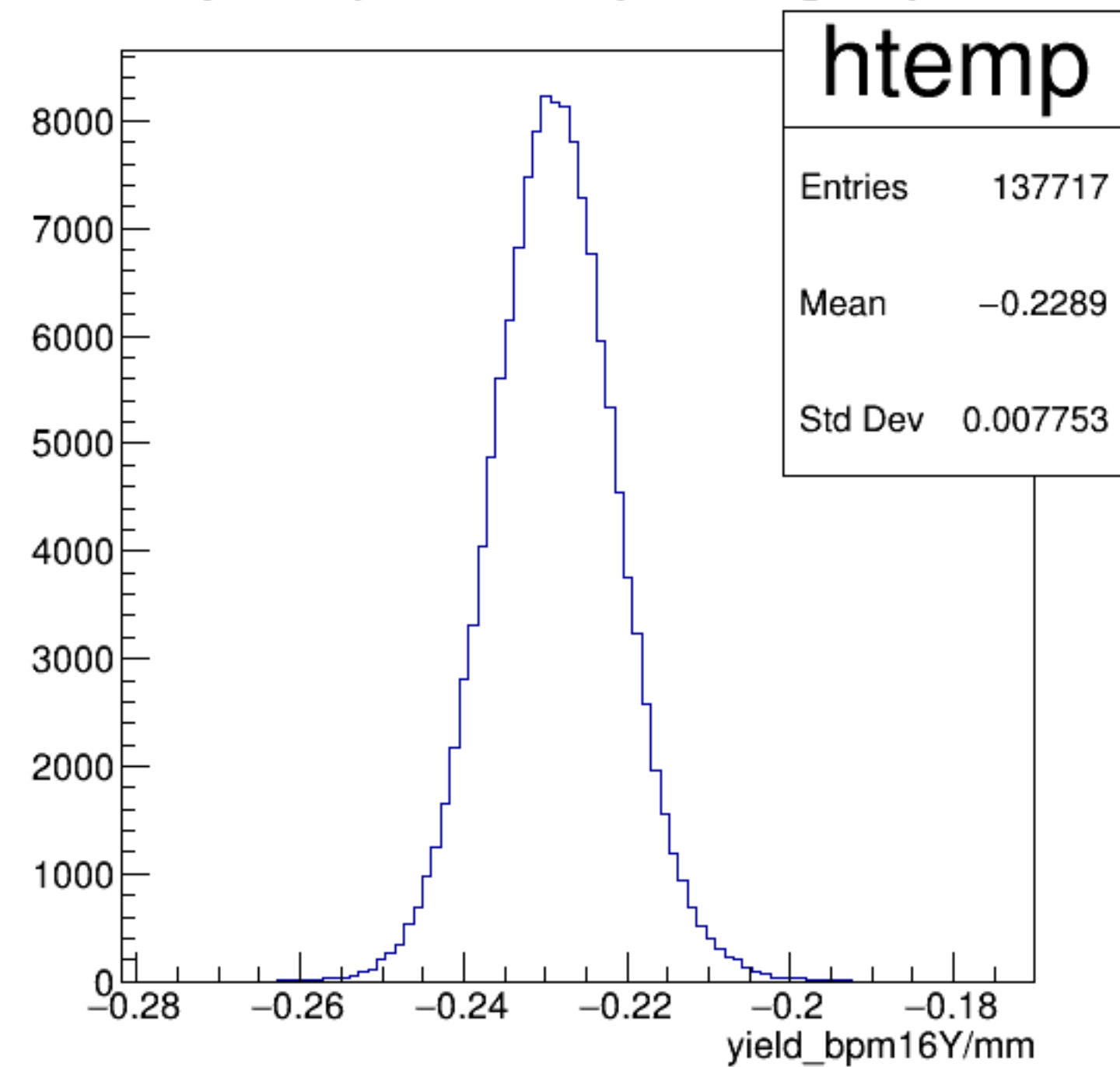


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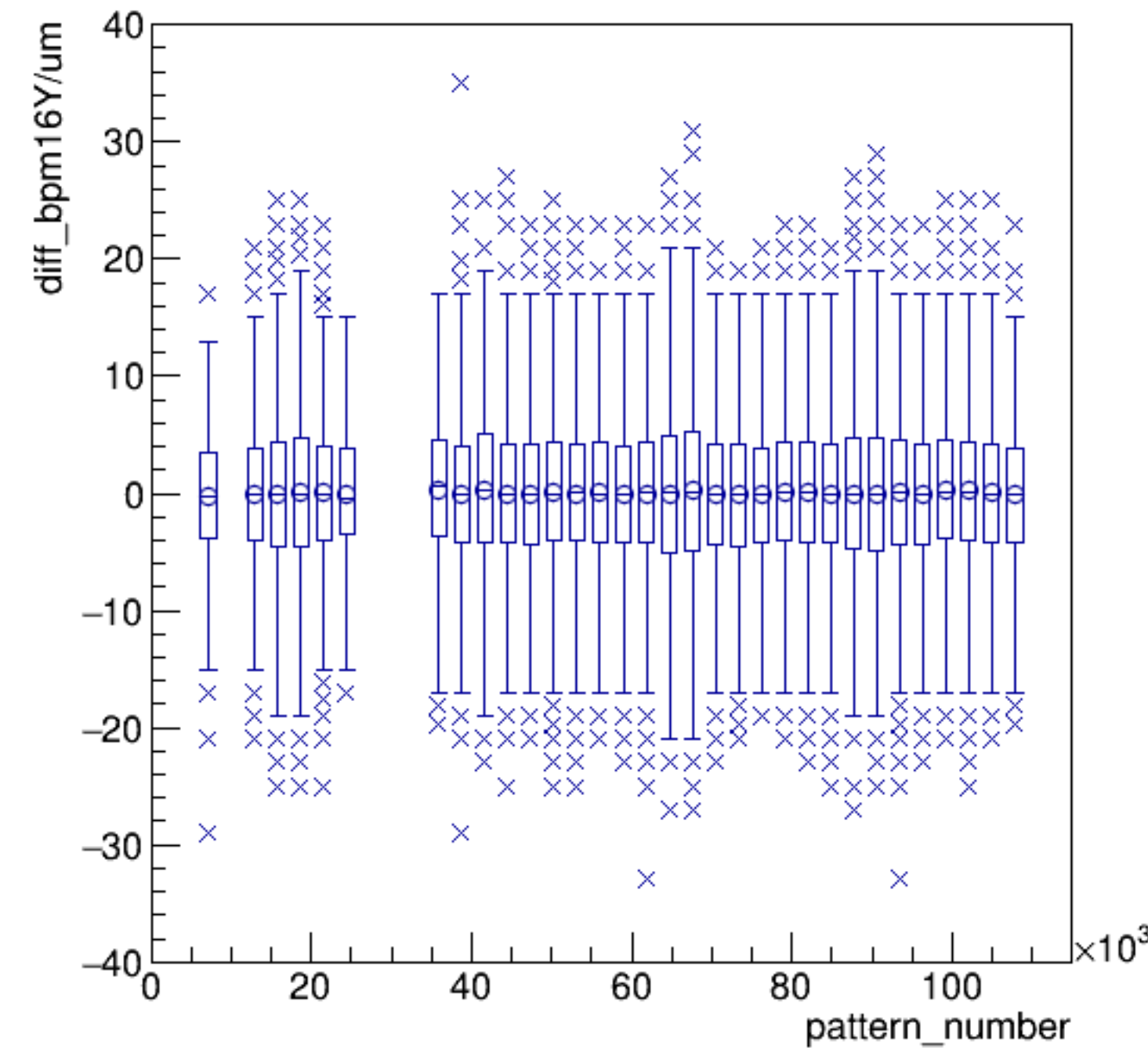


run6380_000_pairTree_bpm16Y.png

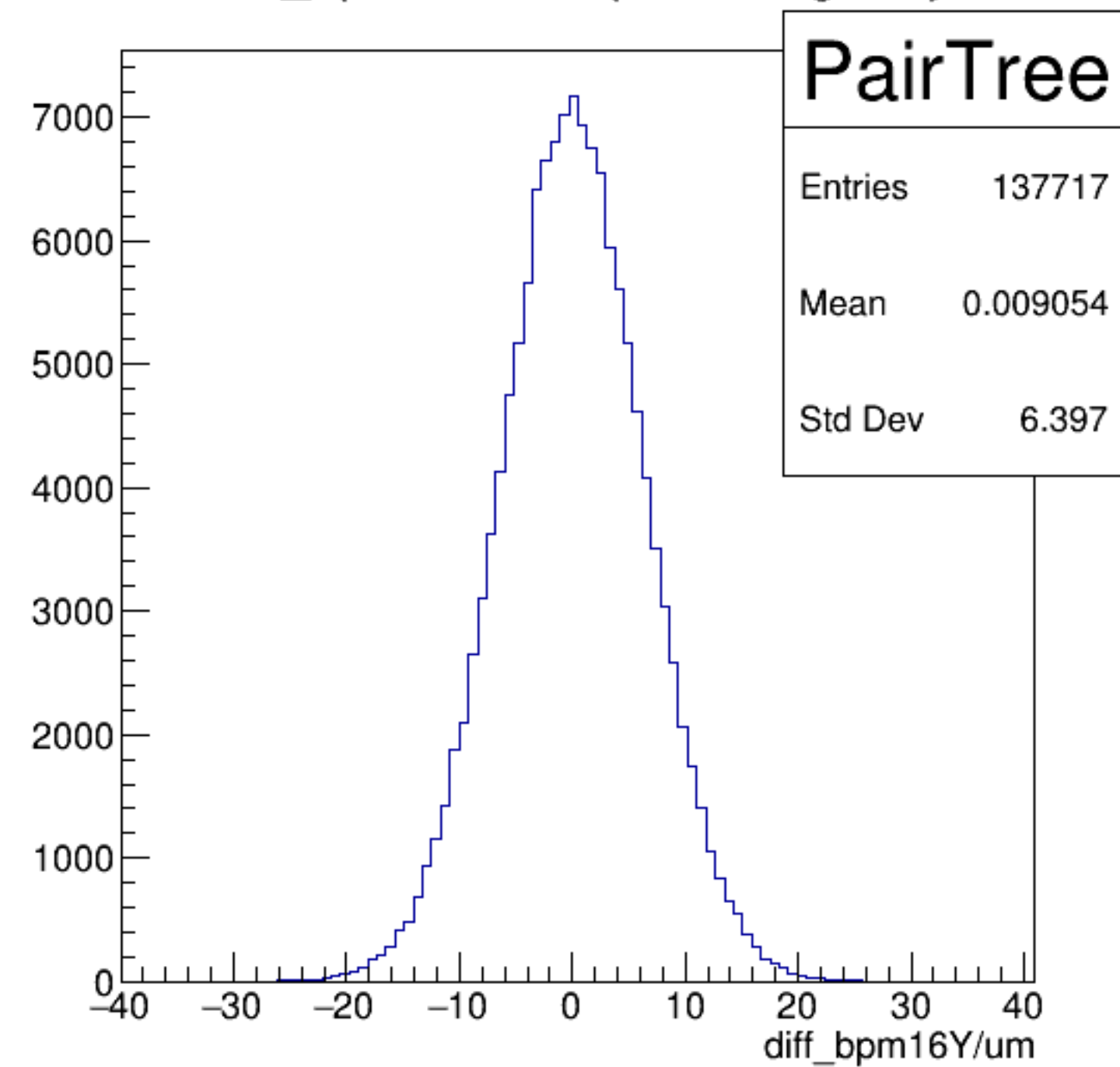
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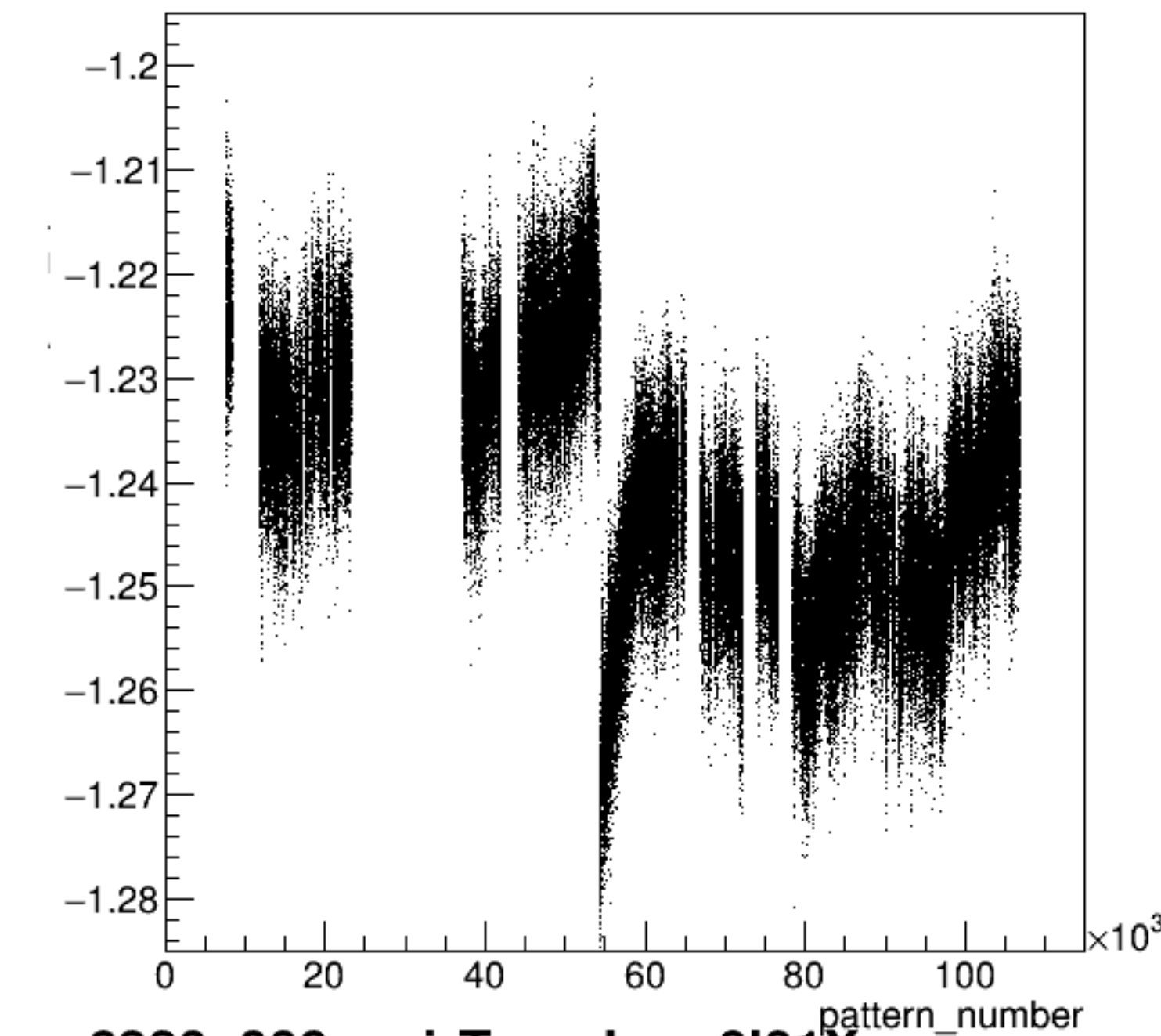
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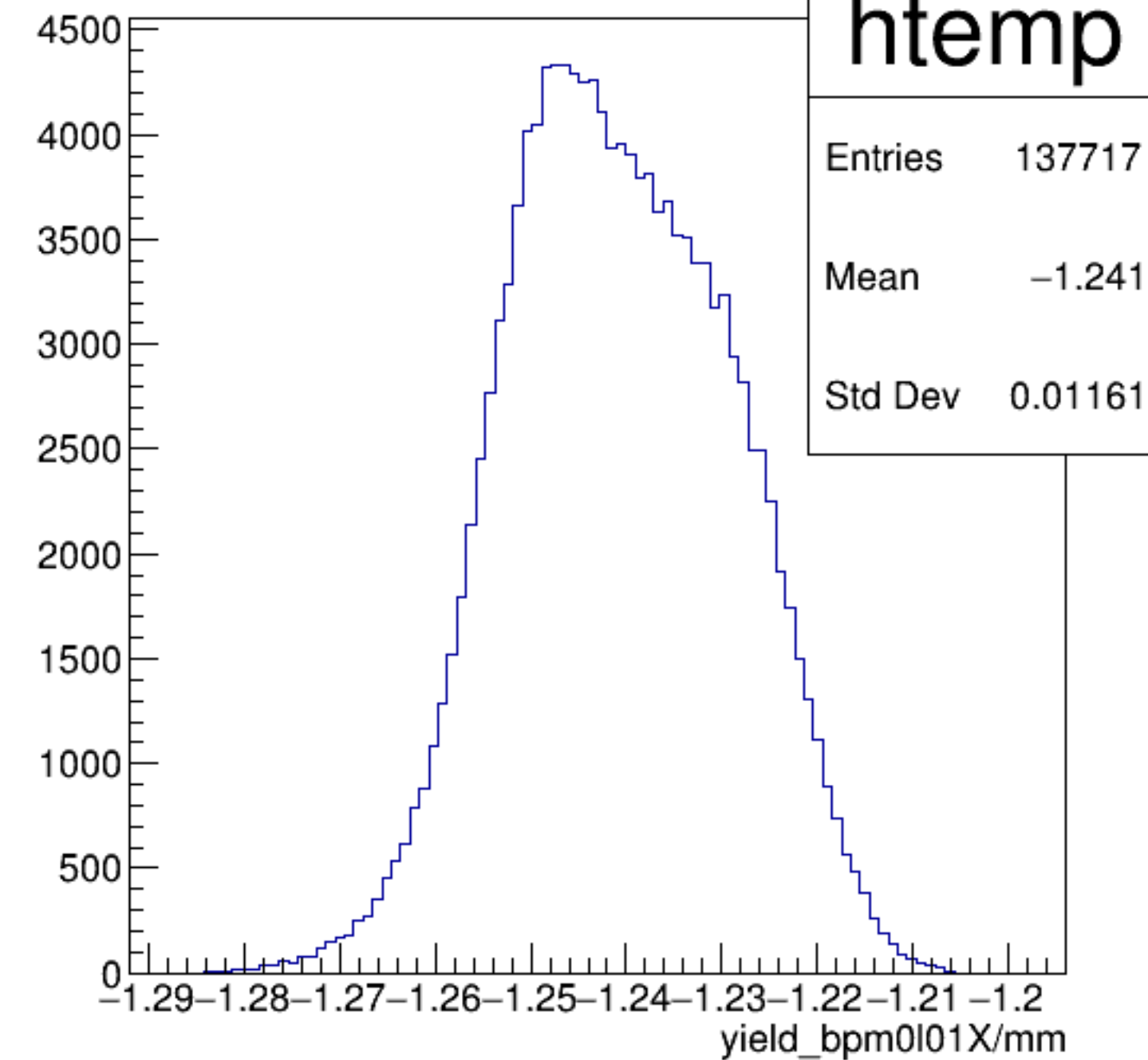


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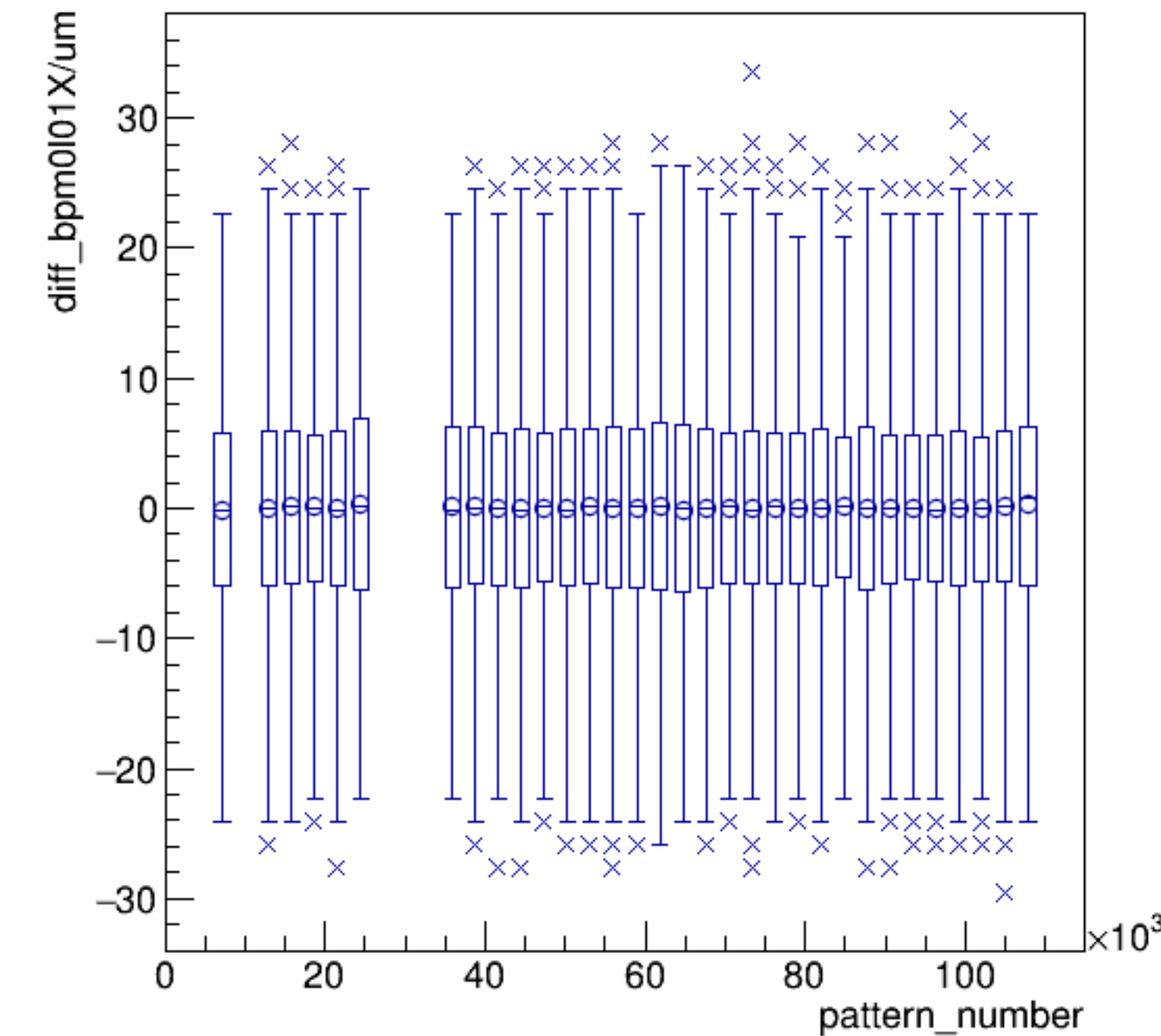


run6380_000_pairTree_bpm0l01X.png

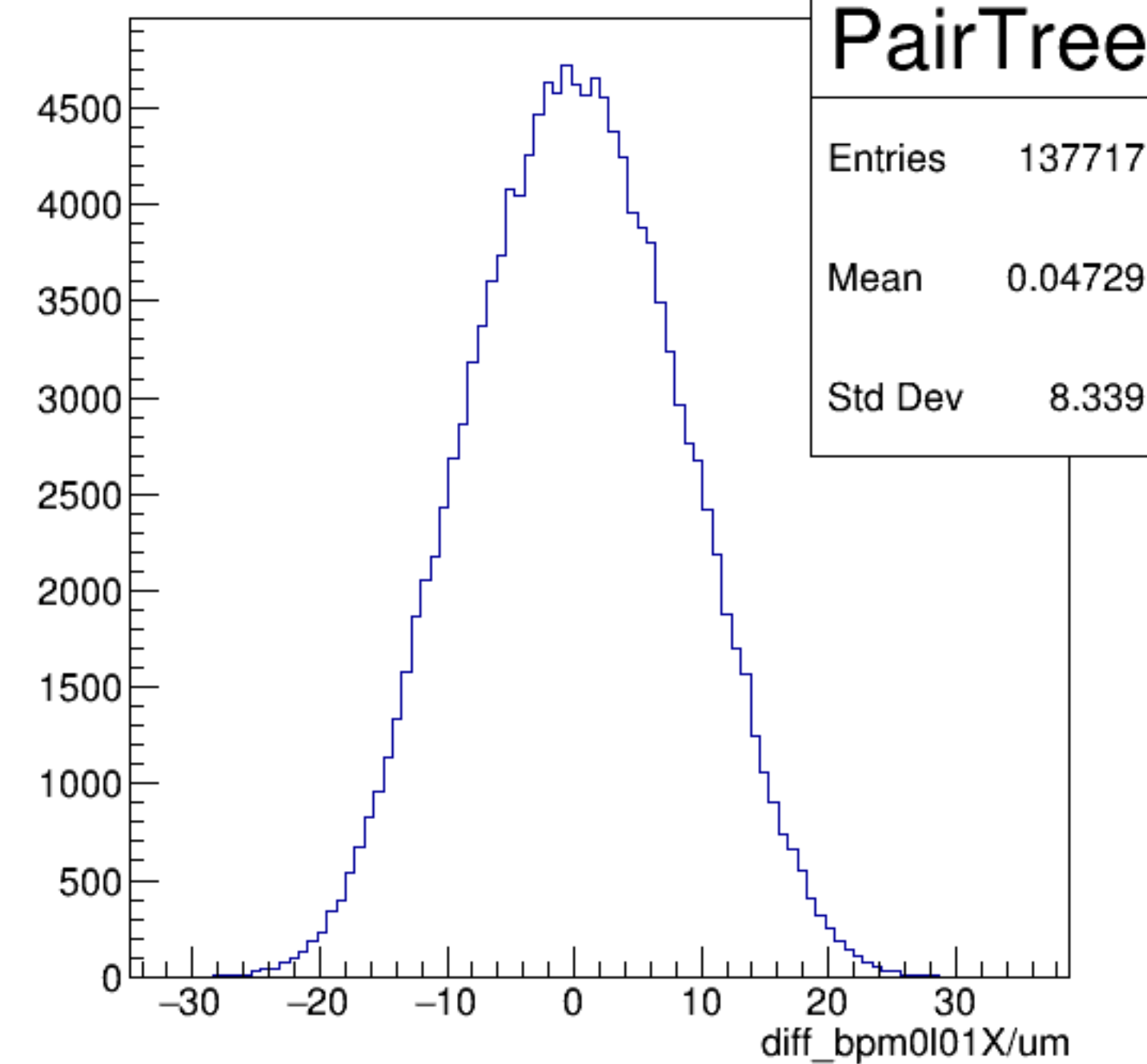
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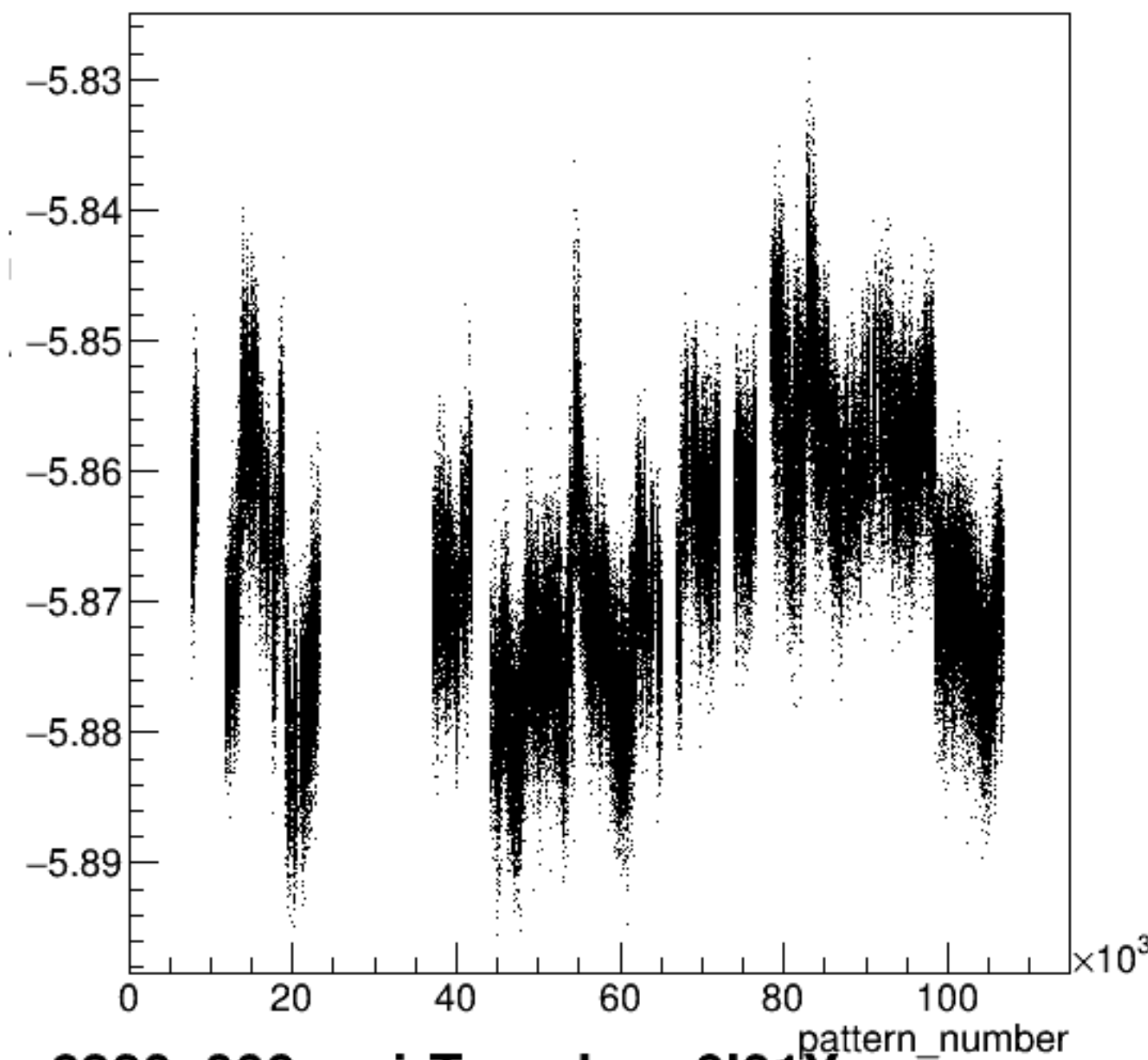
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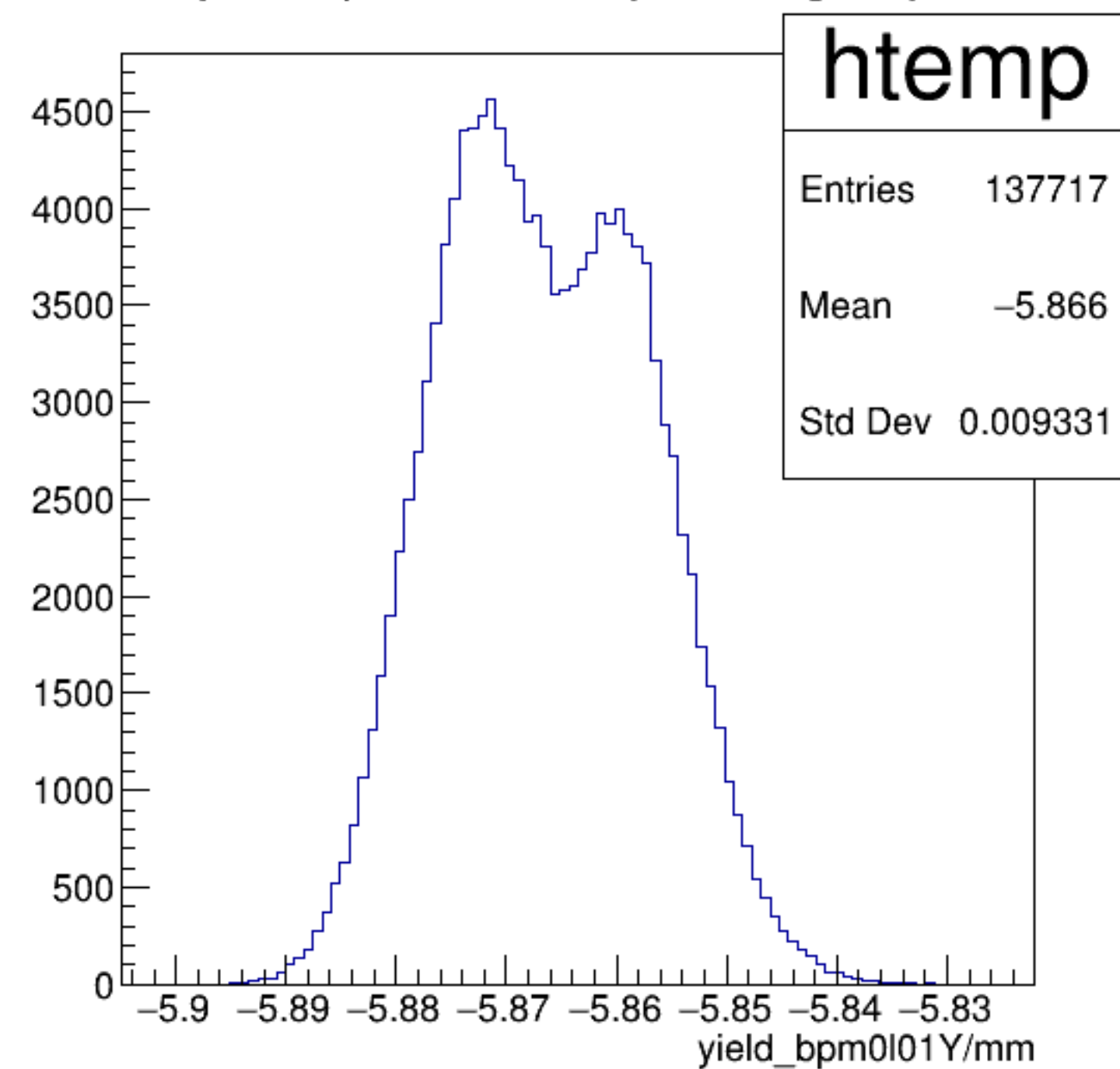
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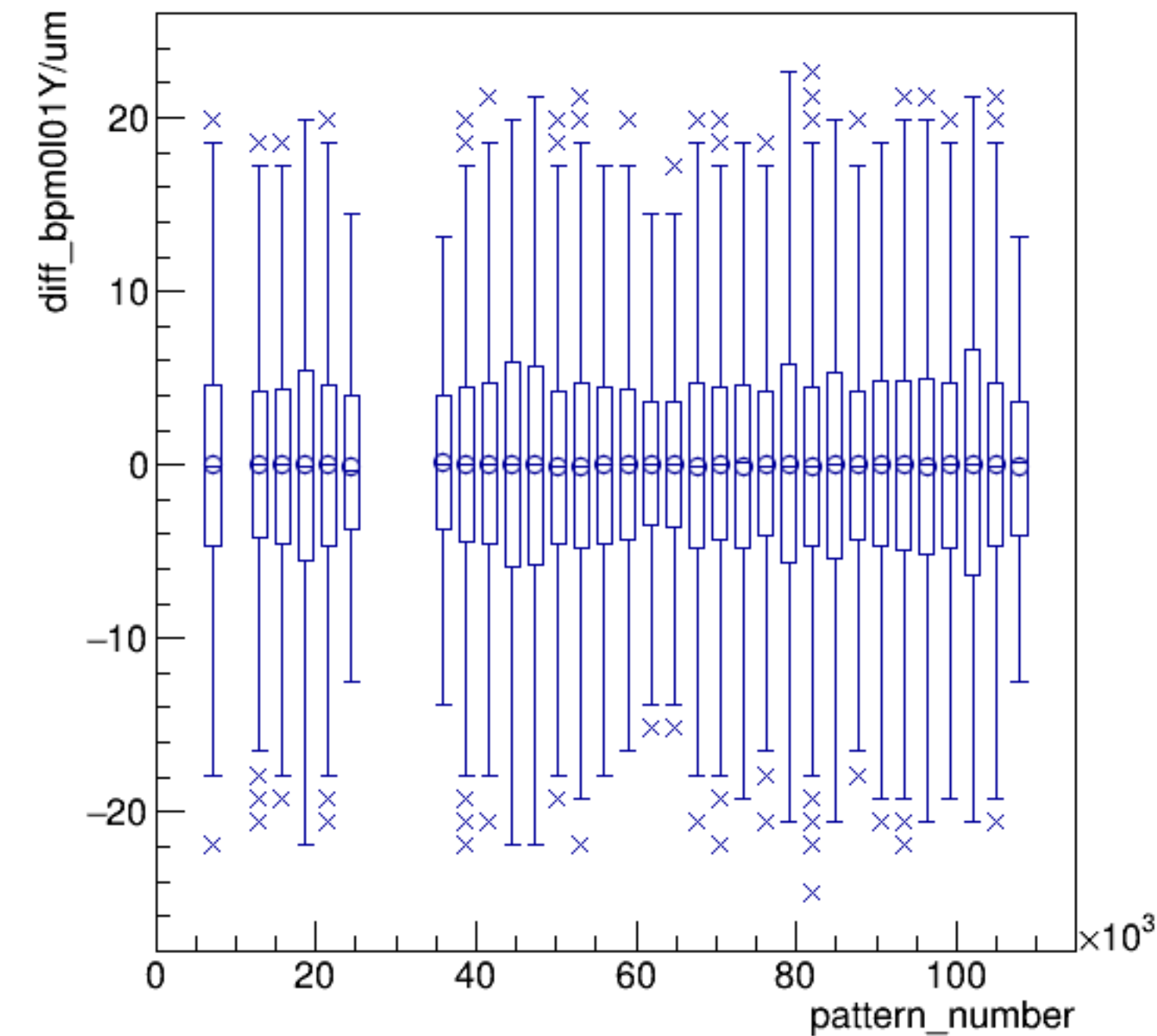
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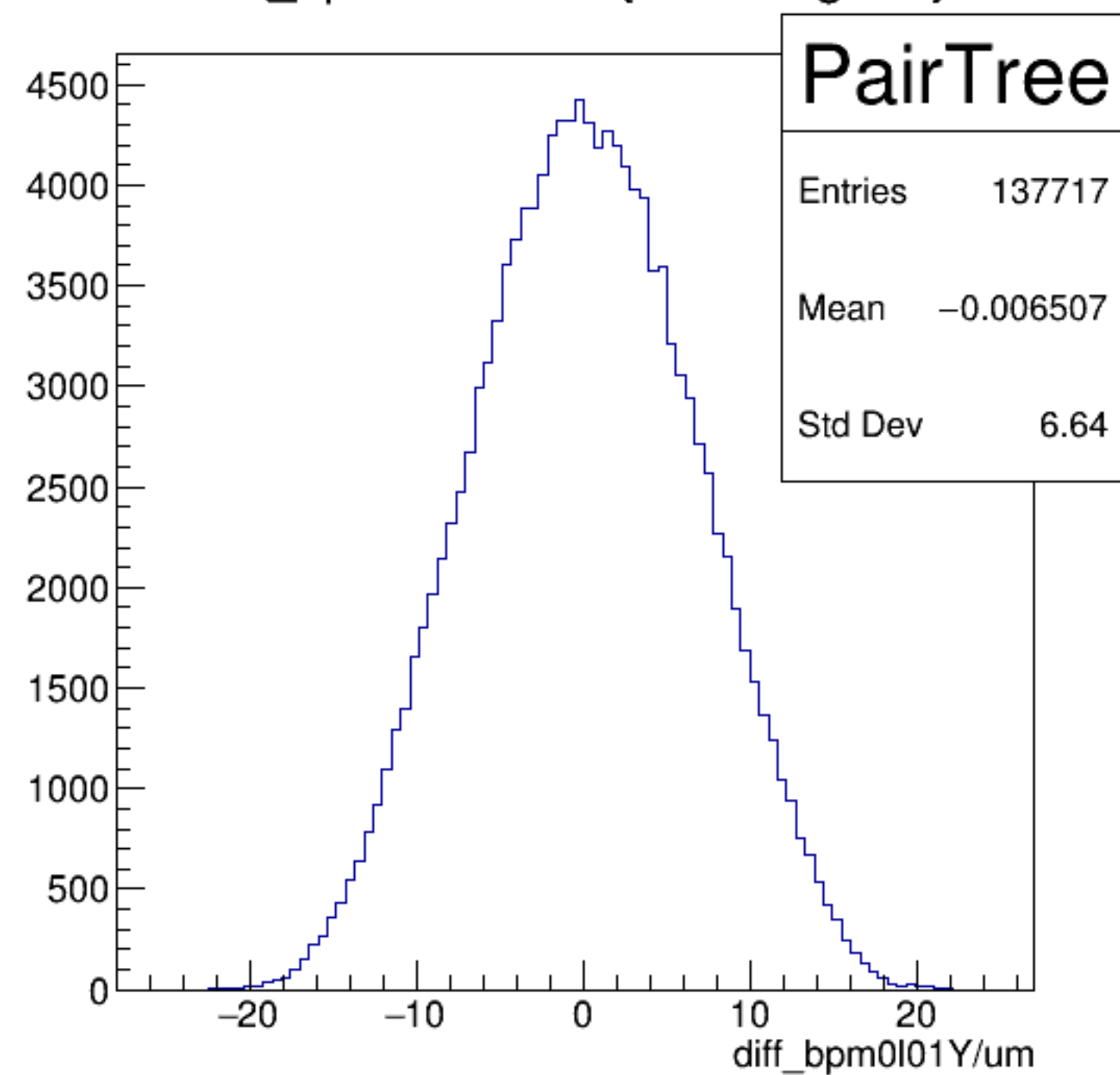
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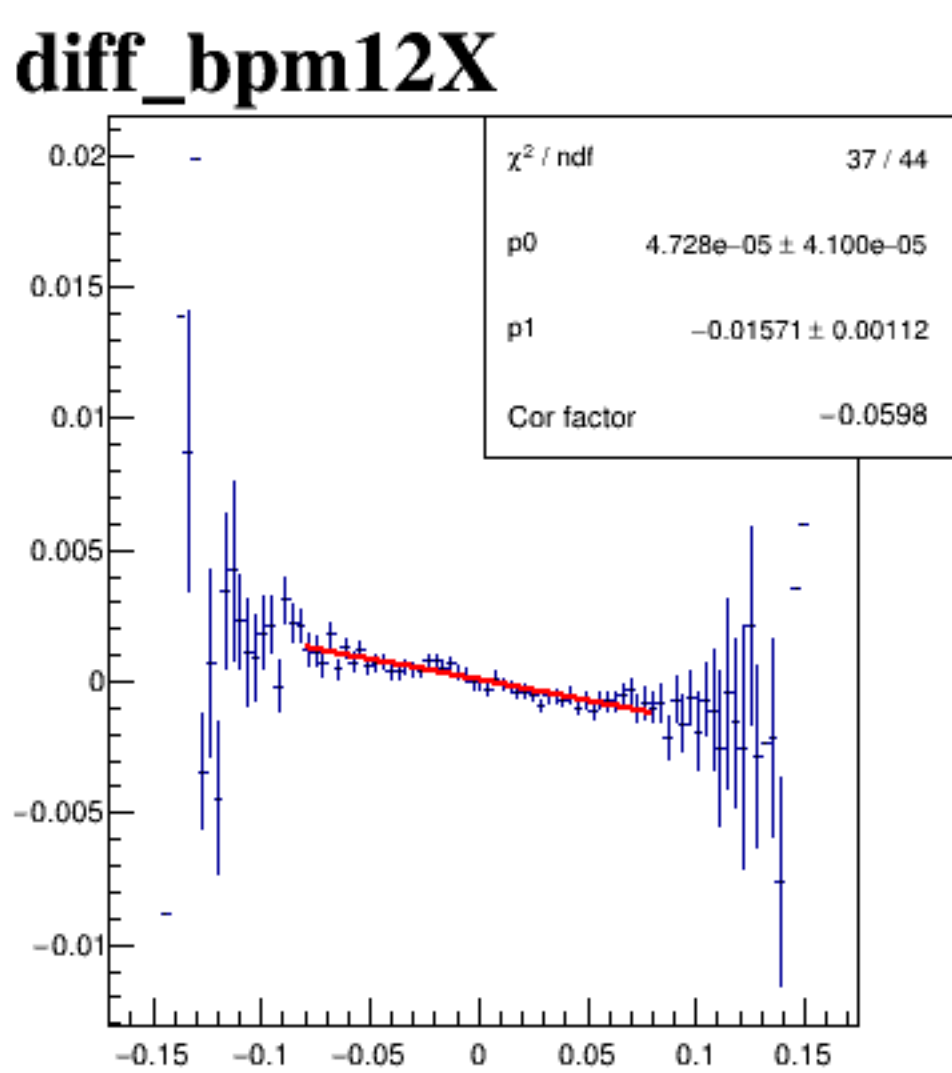
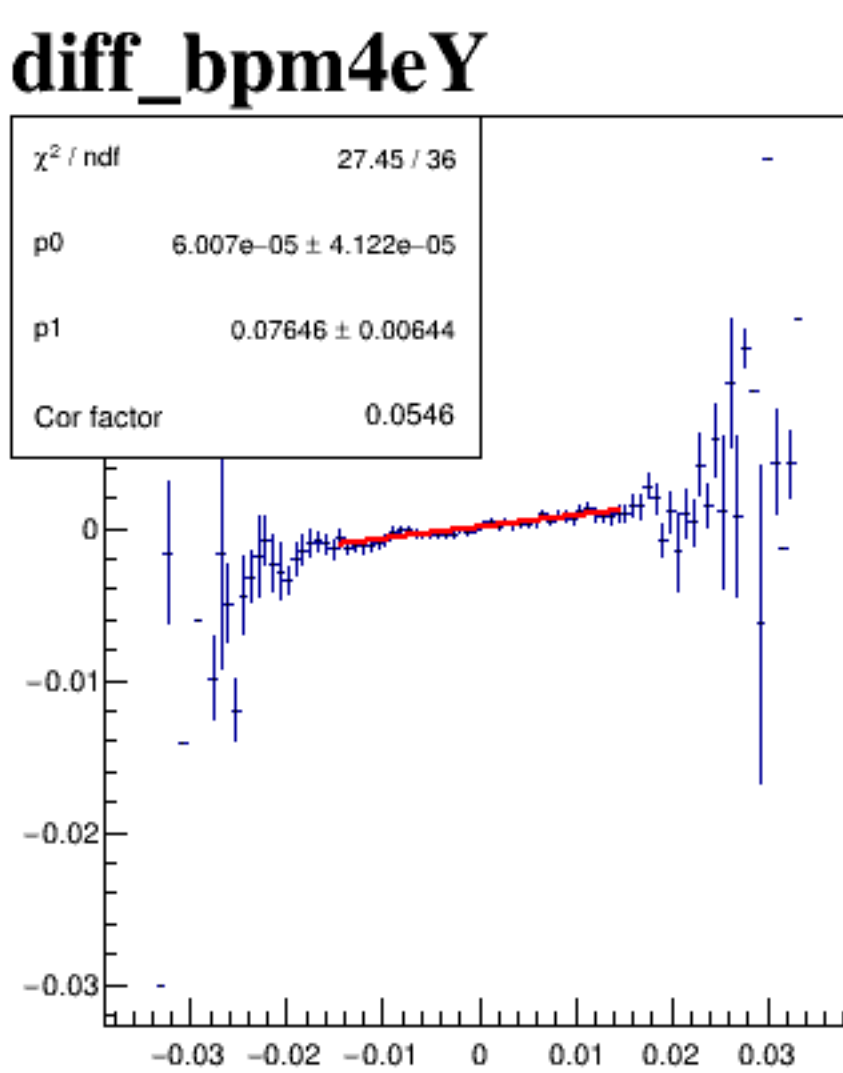
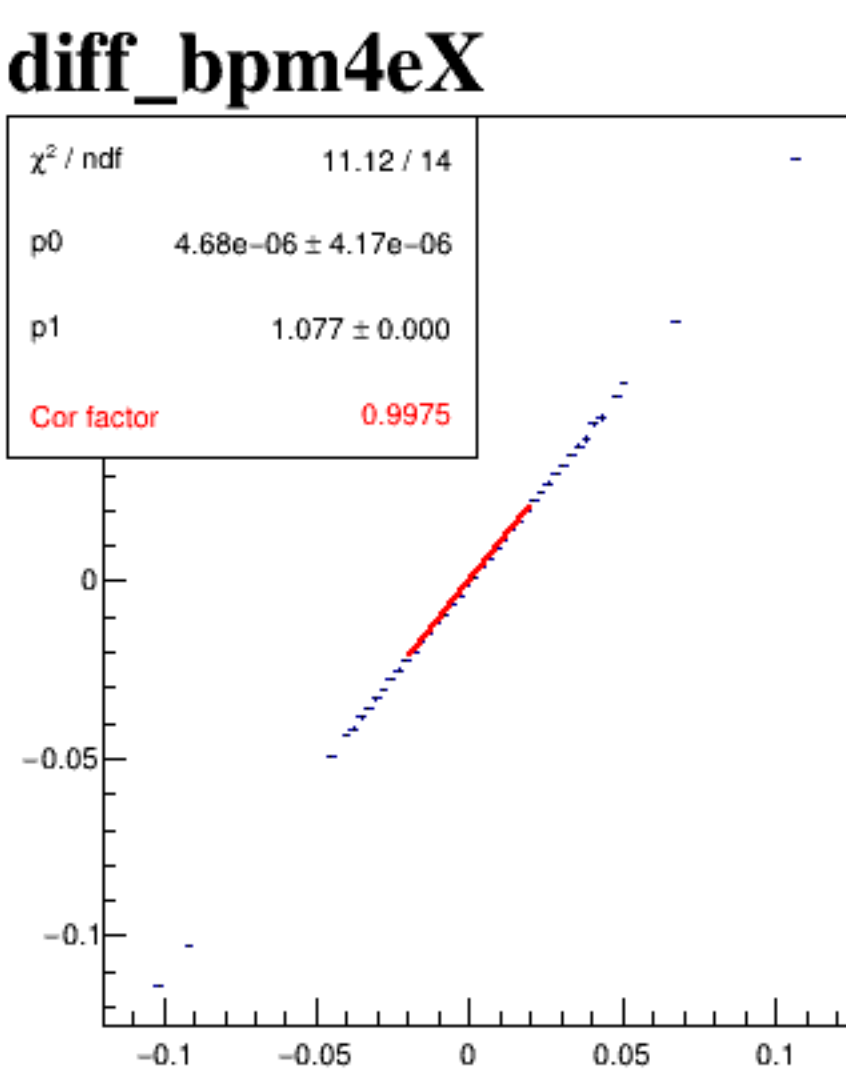
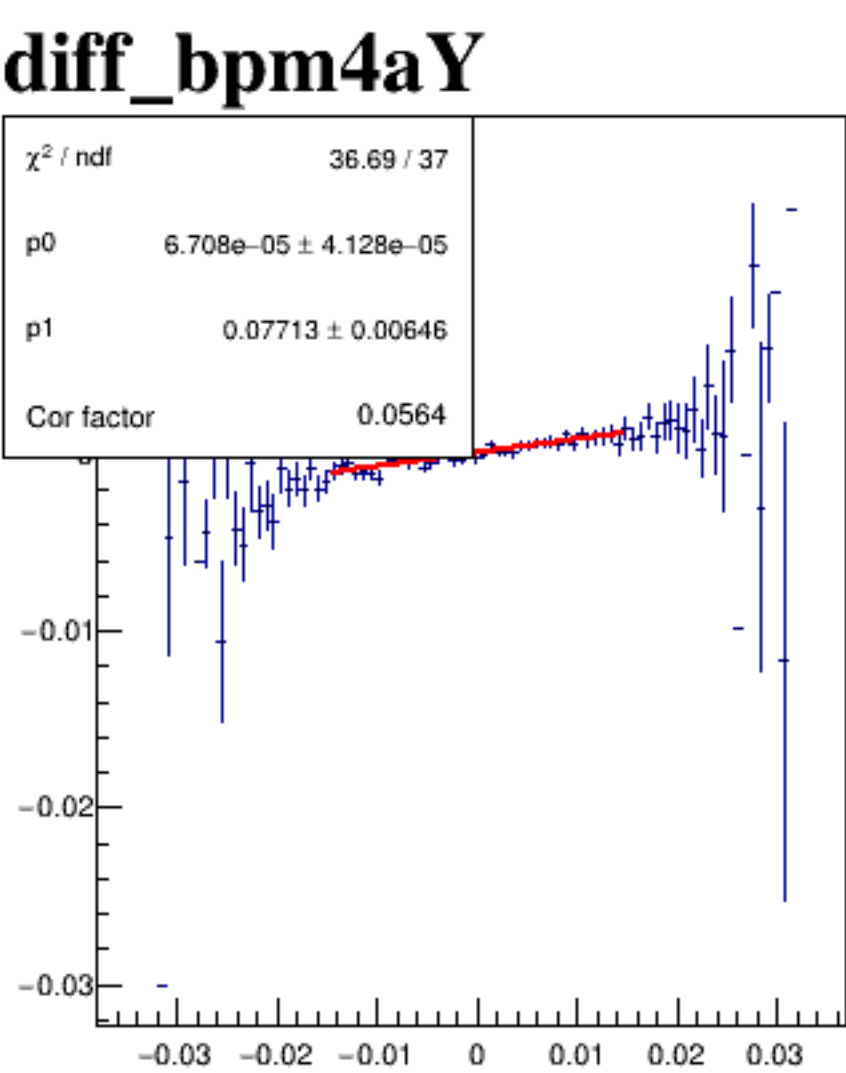
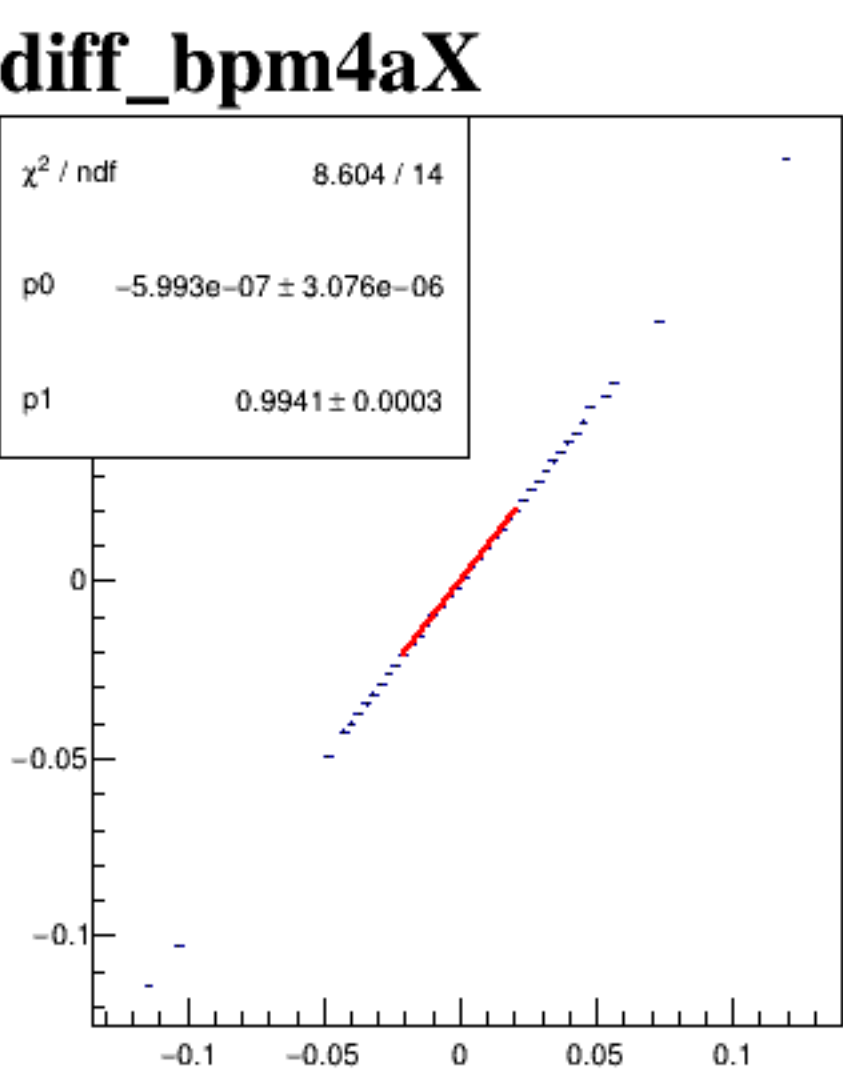


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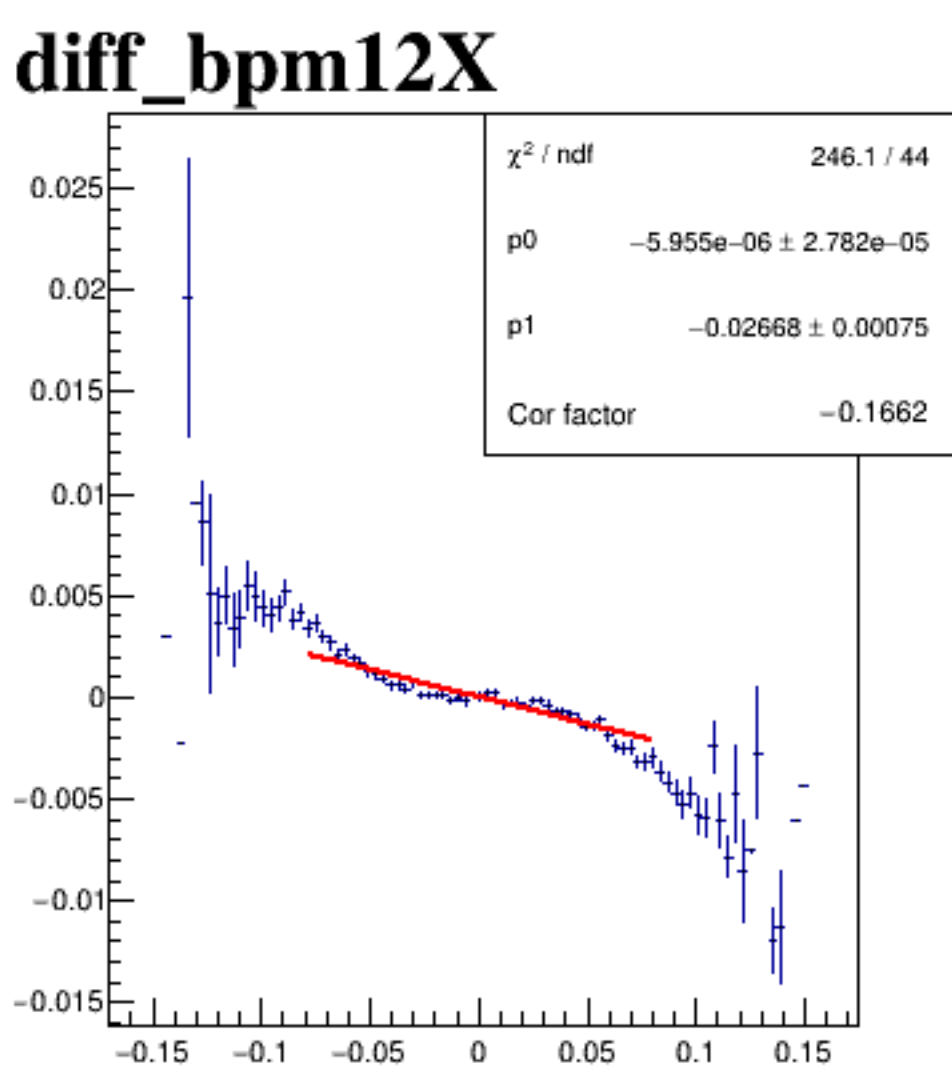
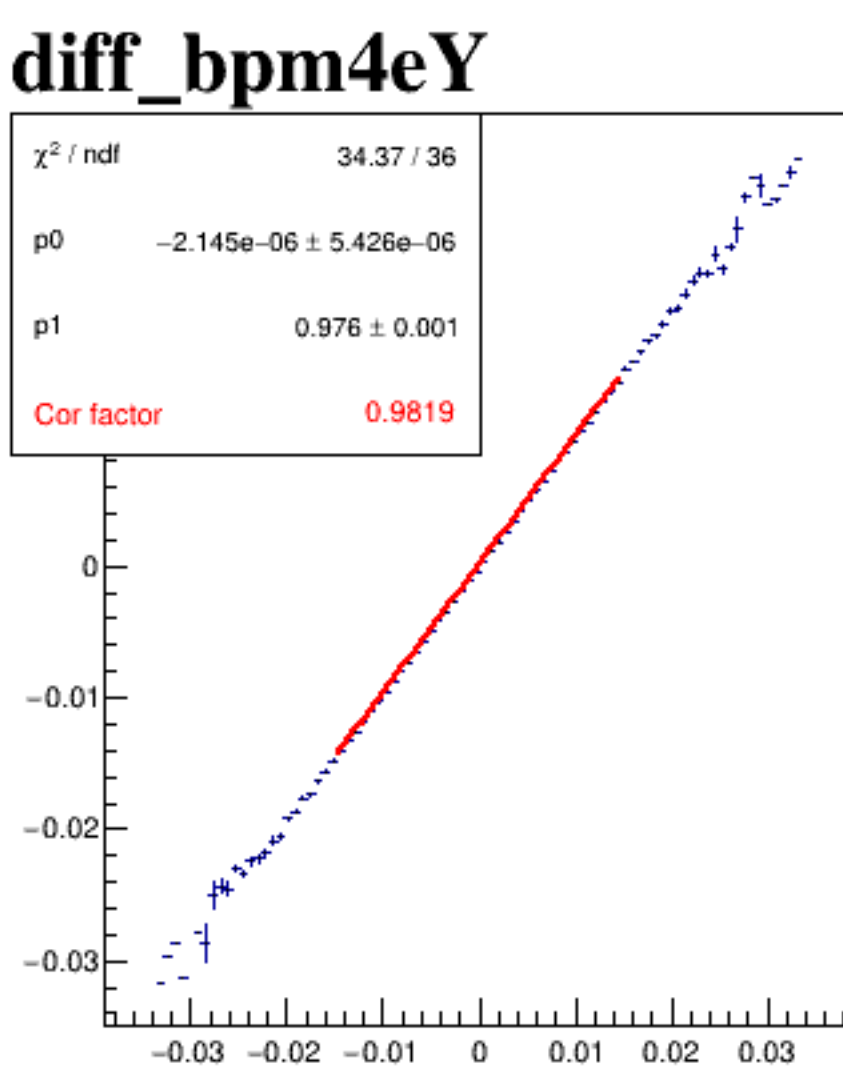
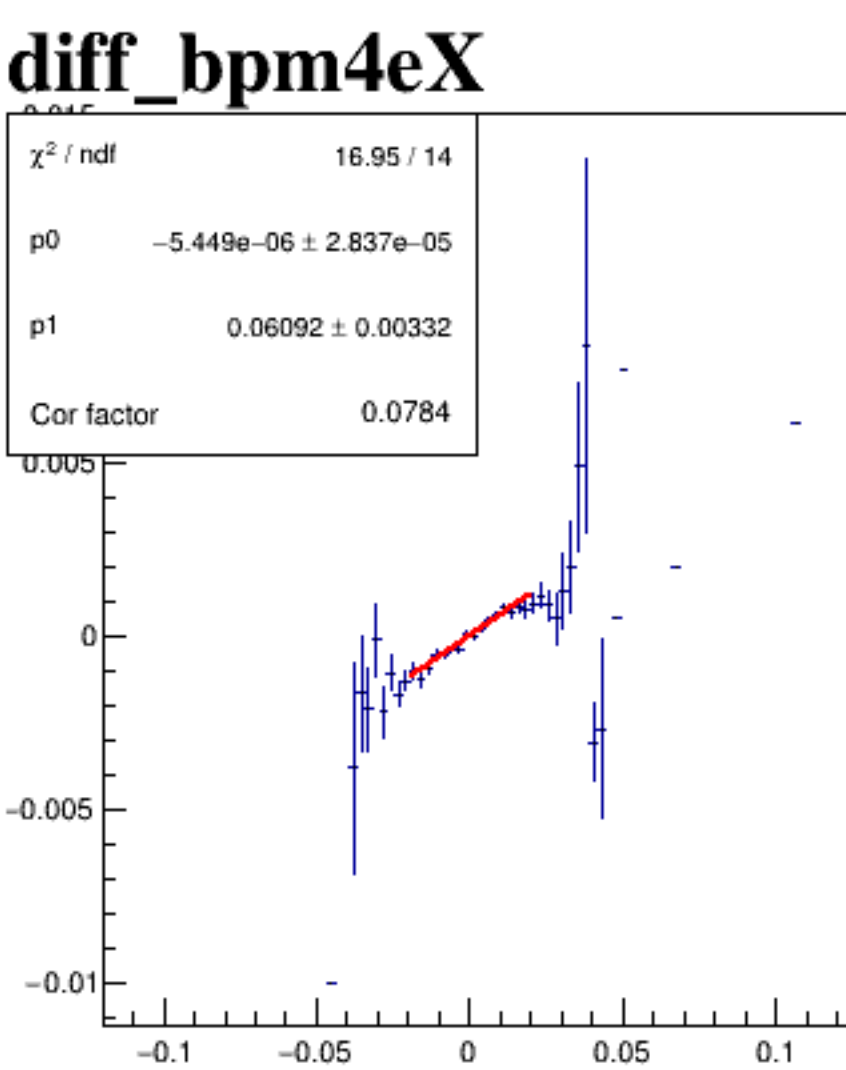
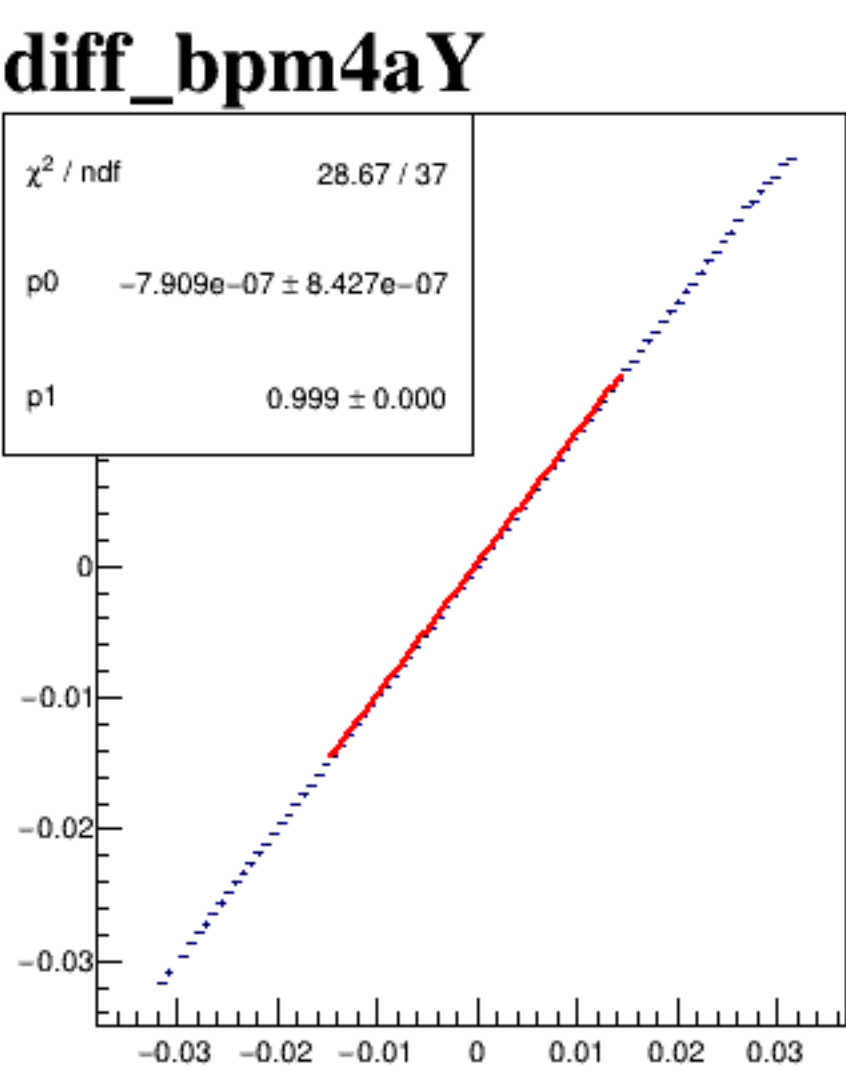
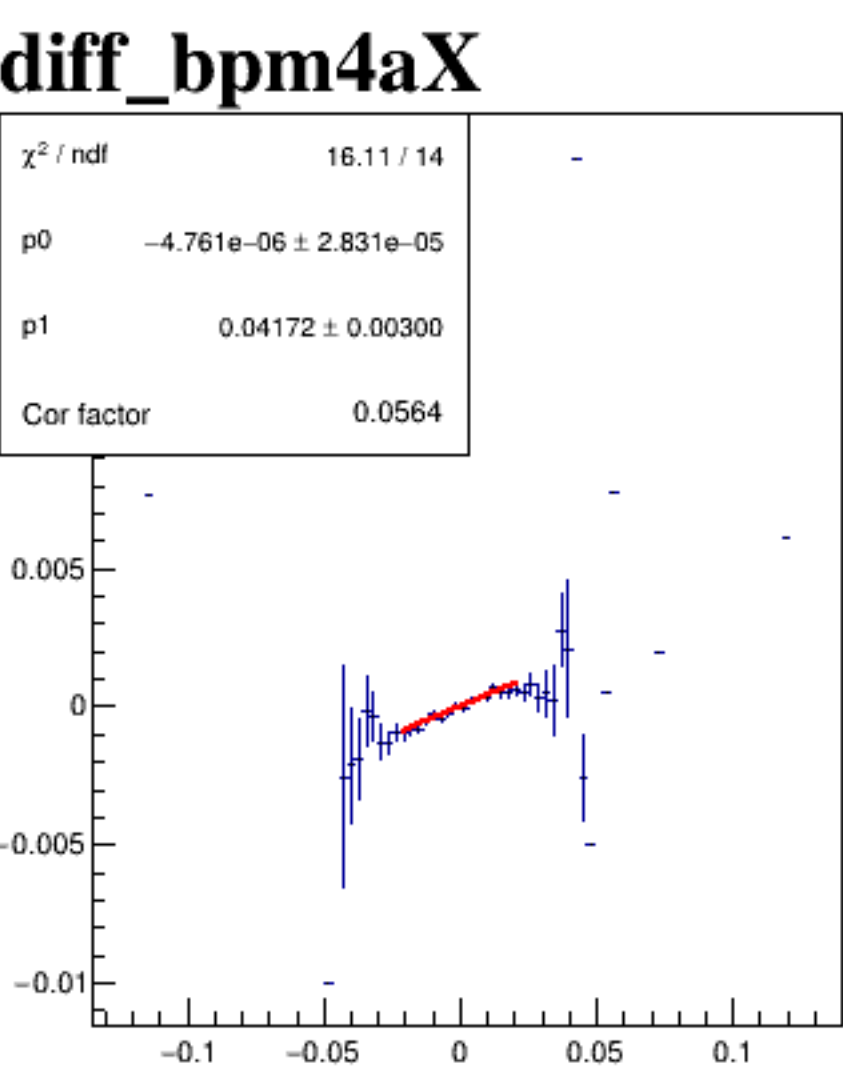


run6380_000_diff_bpm_mutual_correlation_COLZ_default.png

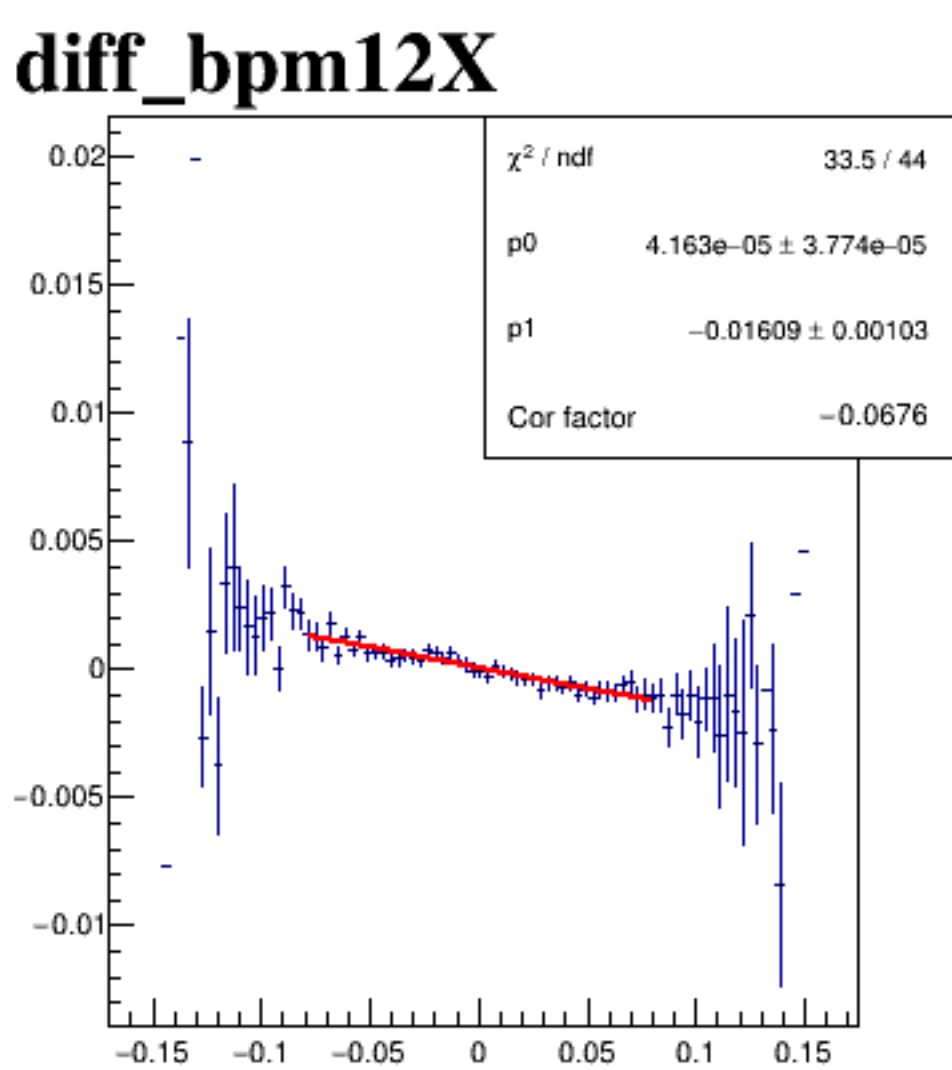
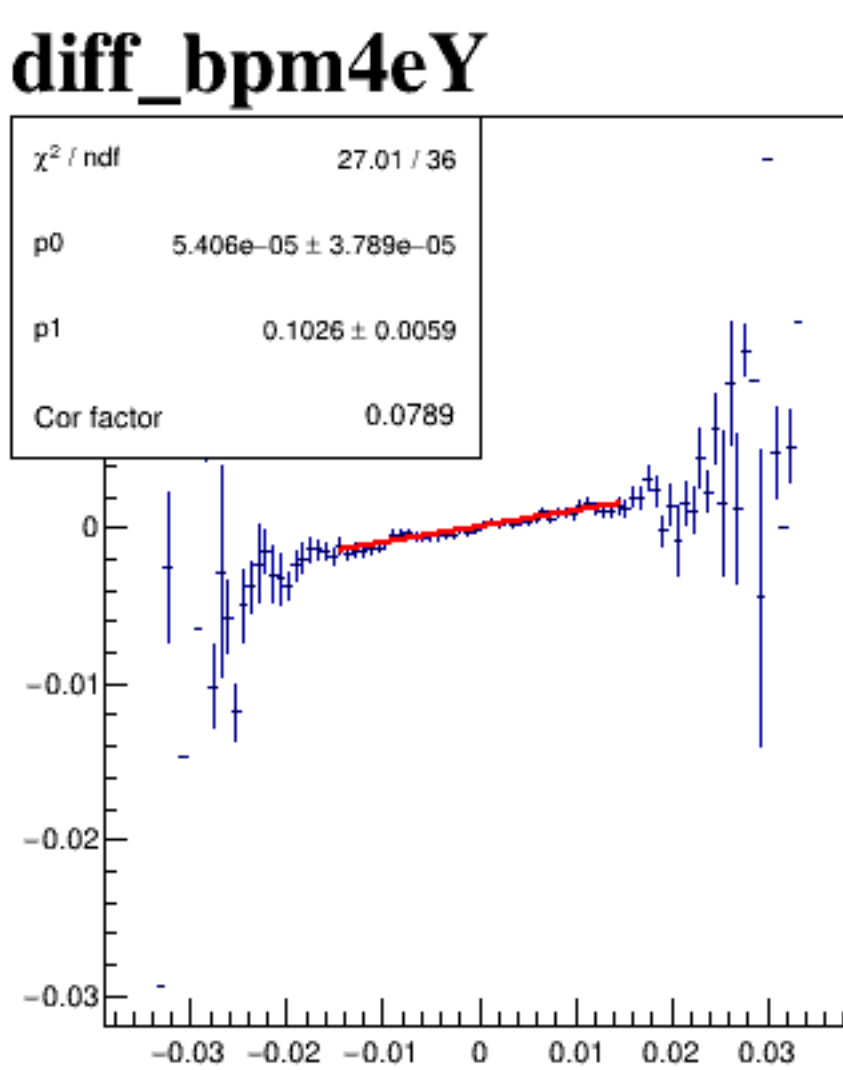
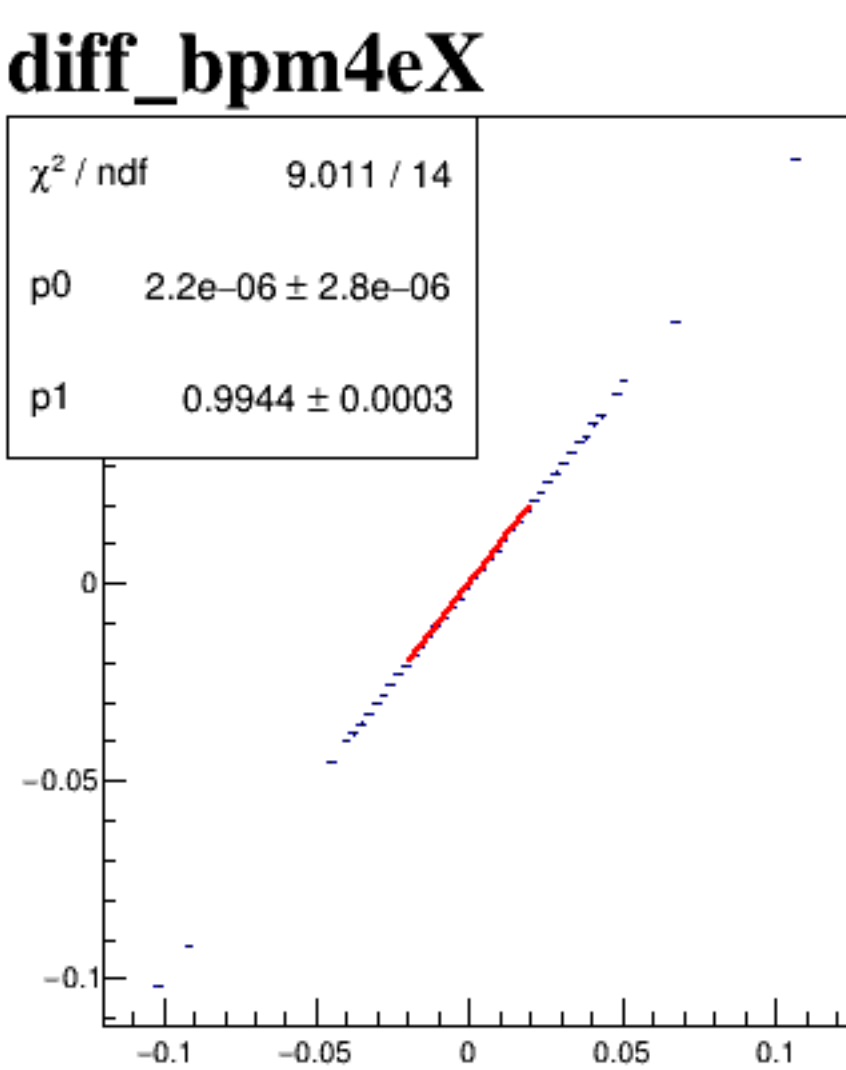
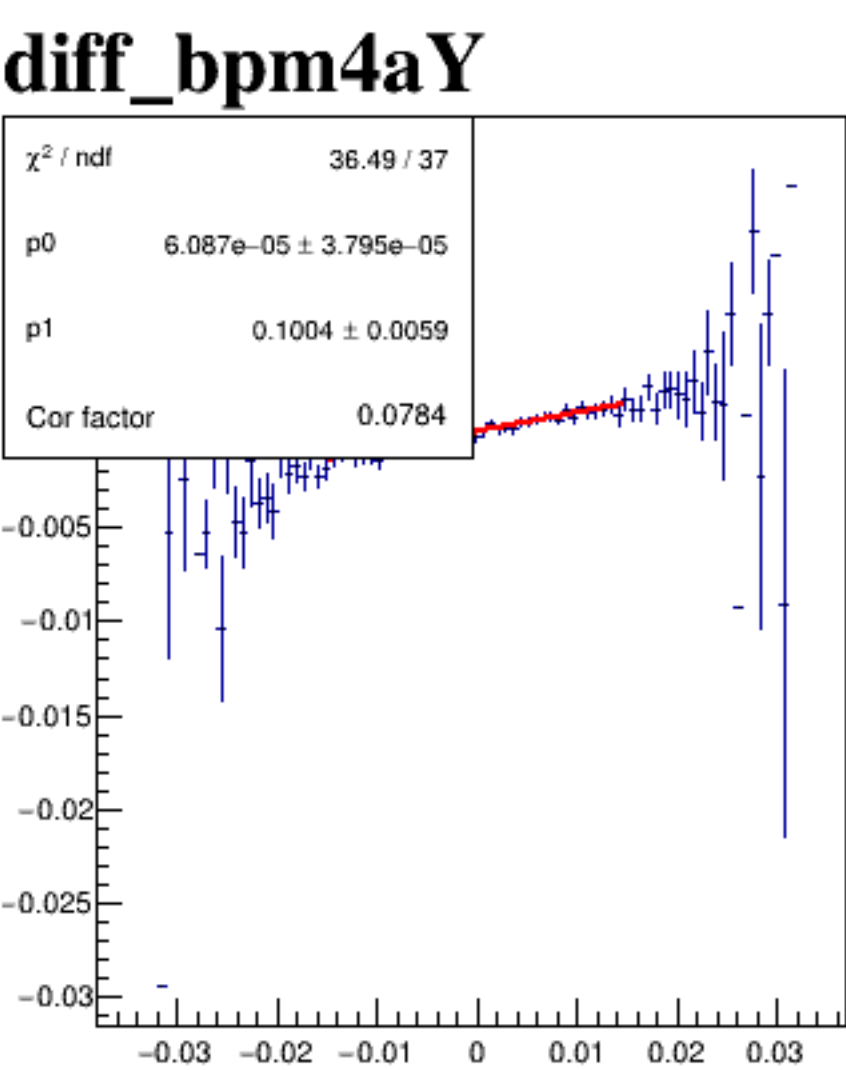
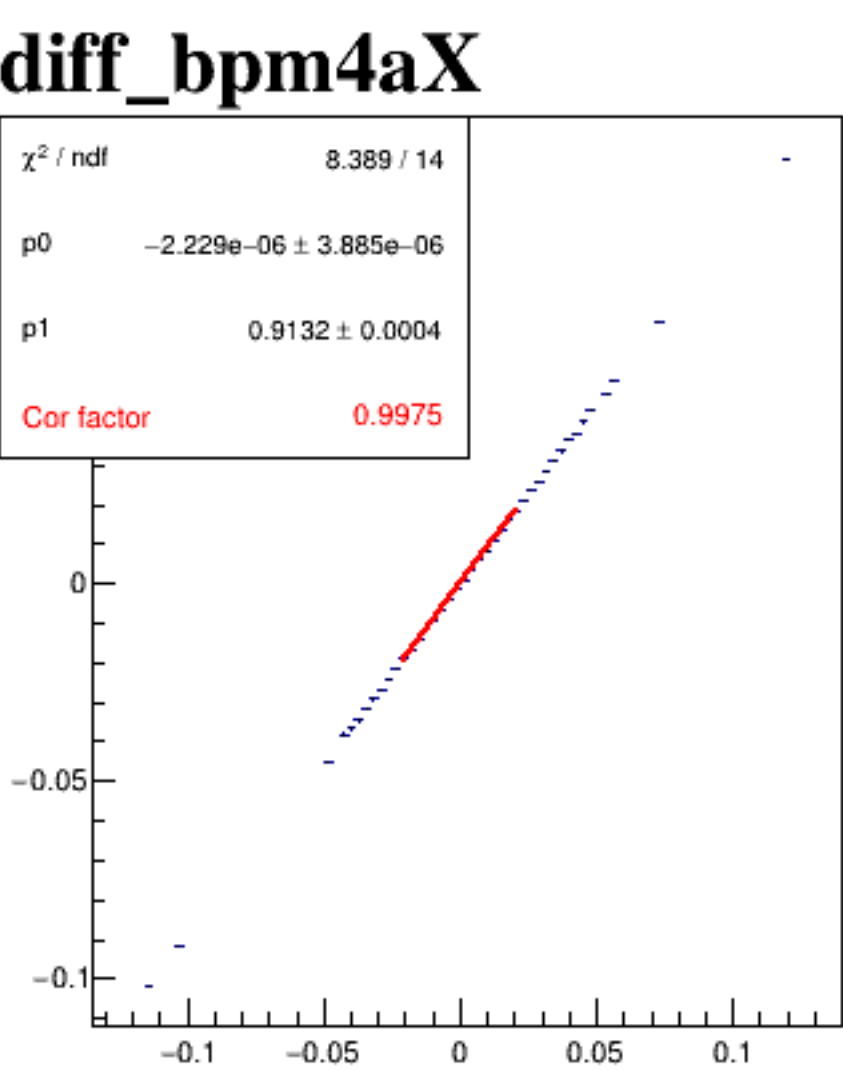
diff_bpm4aX



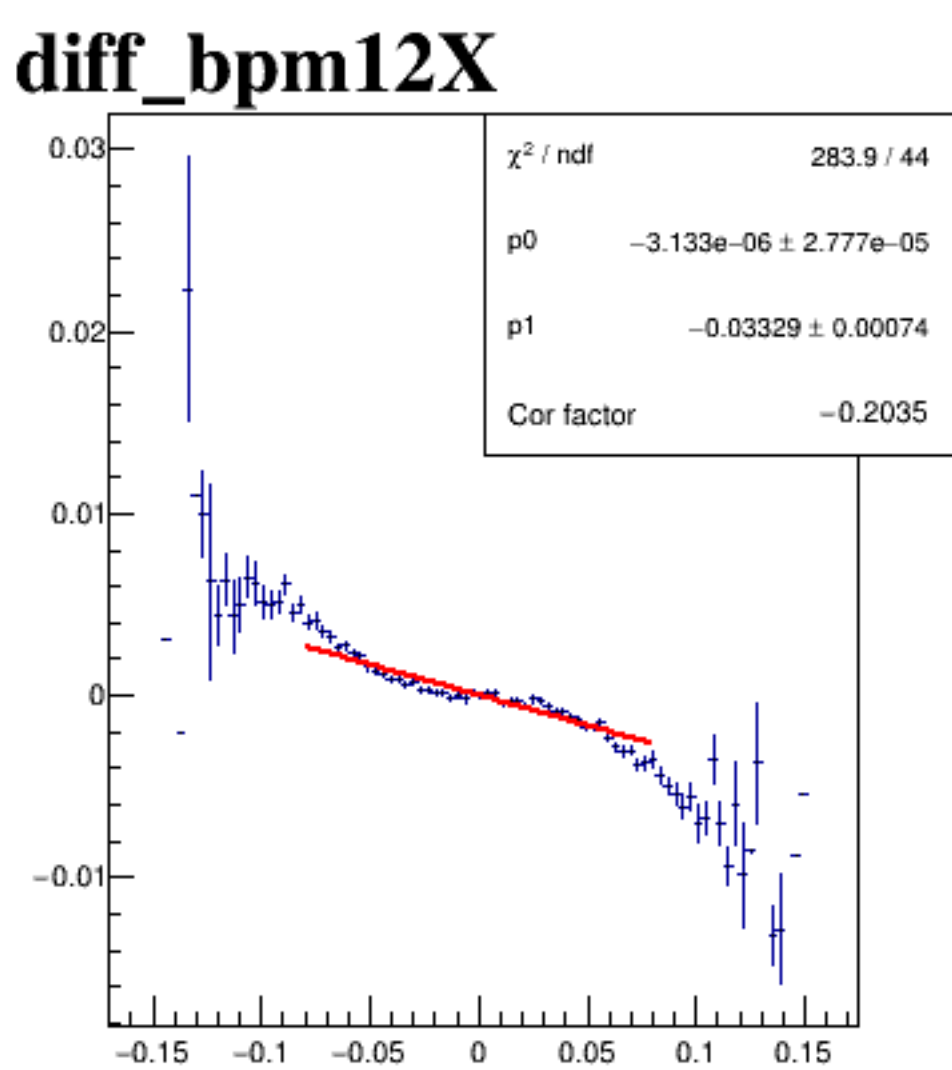
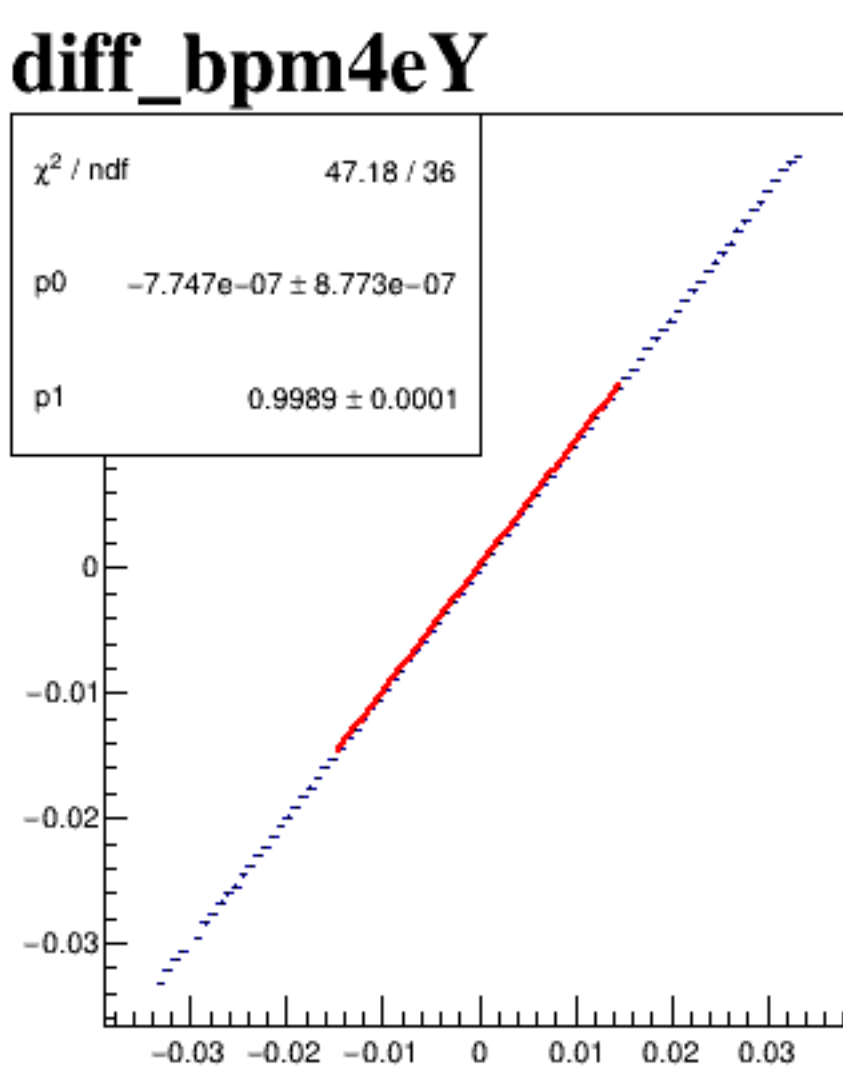
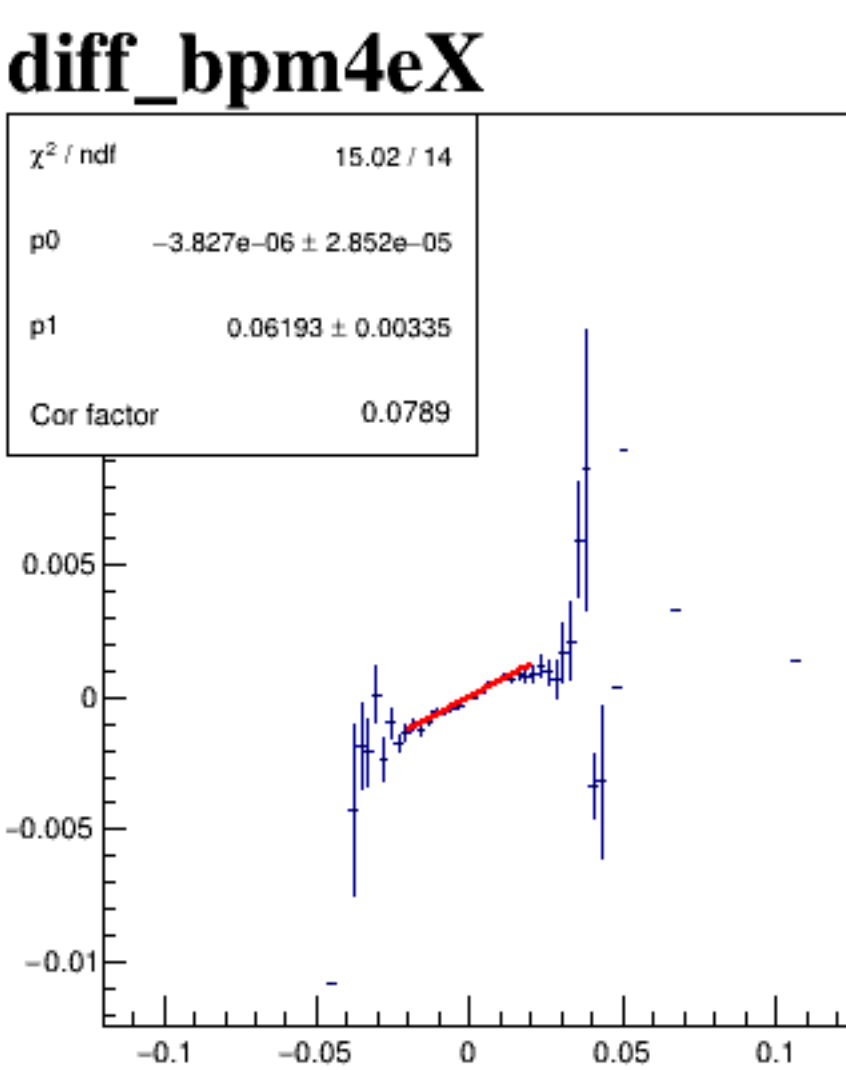
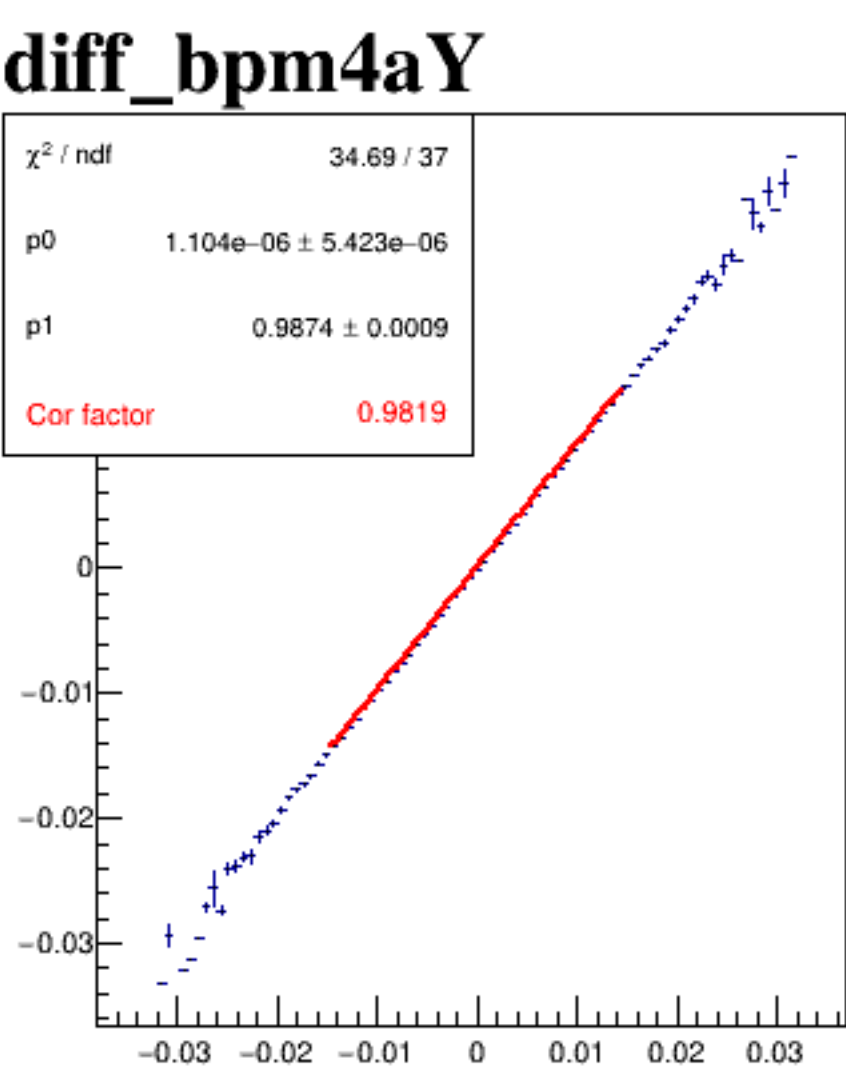
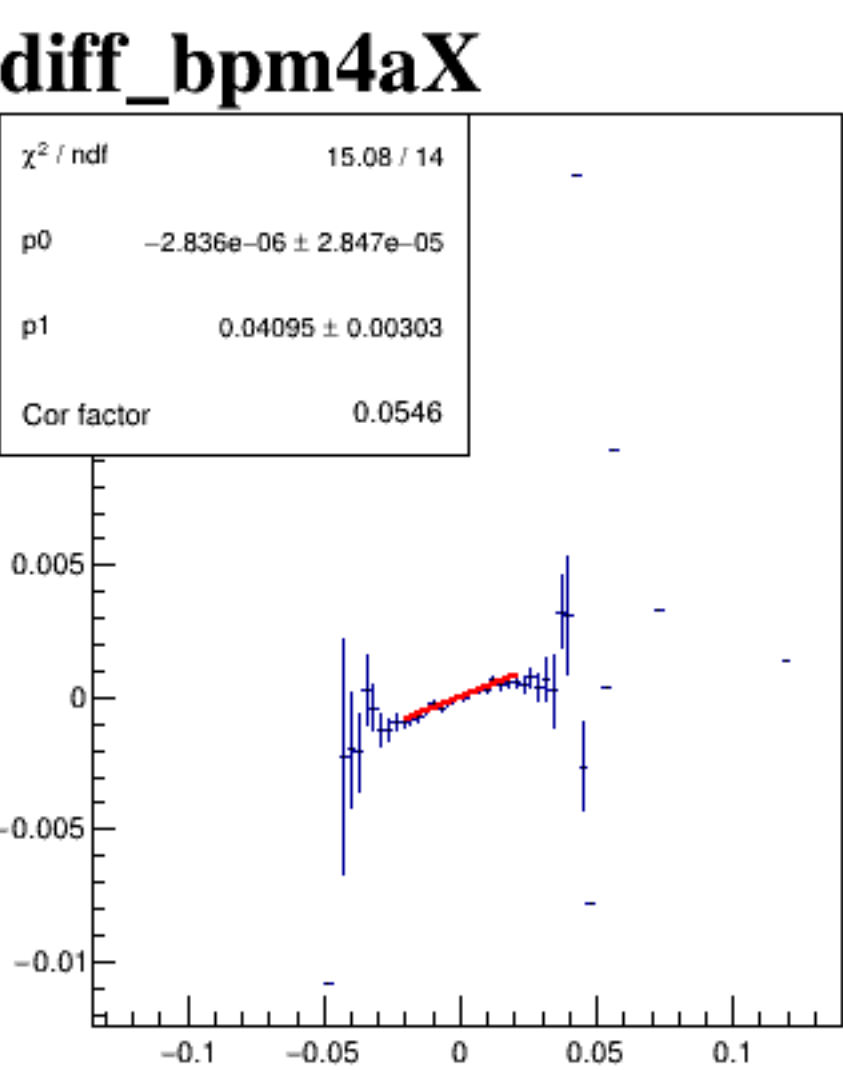
diff_bpm4aY



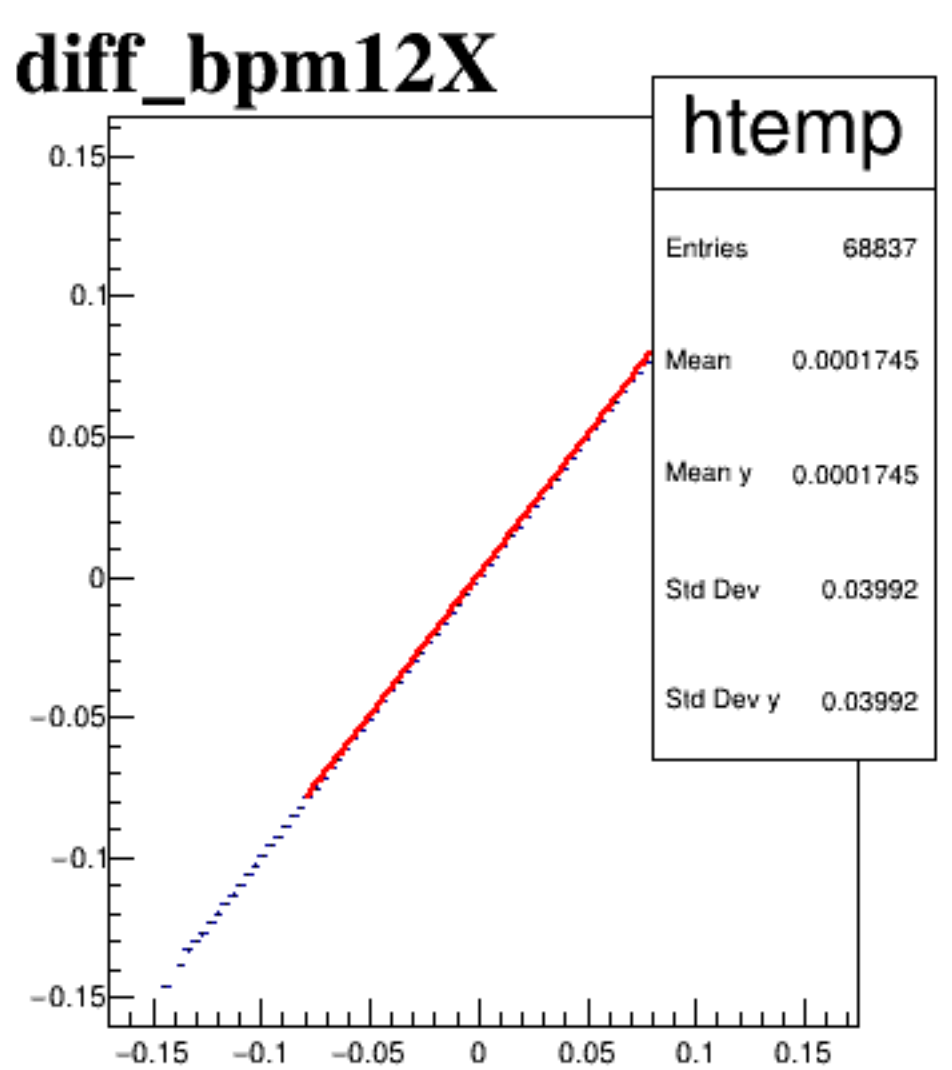
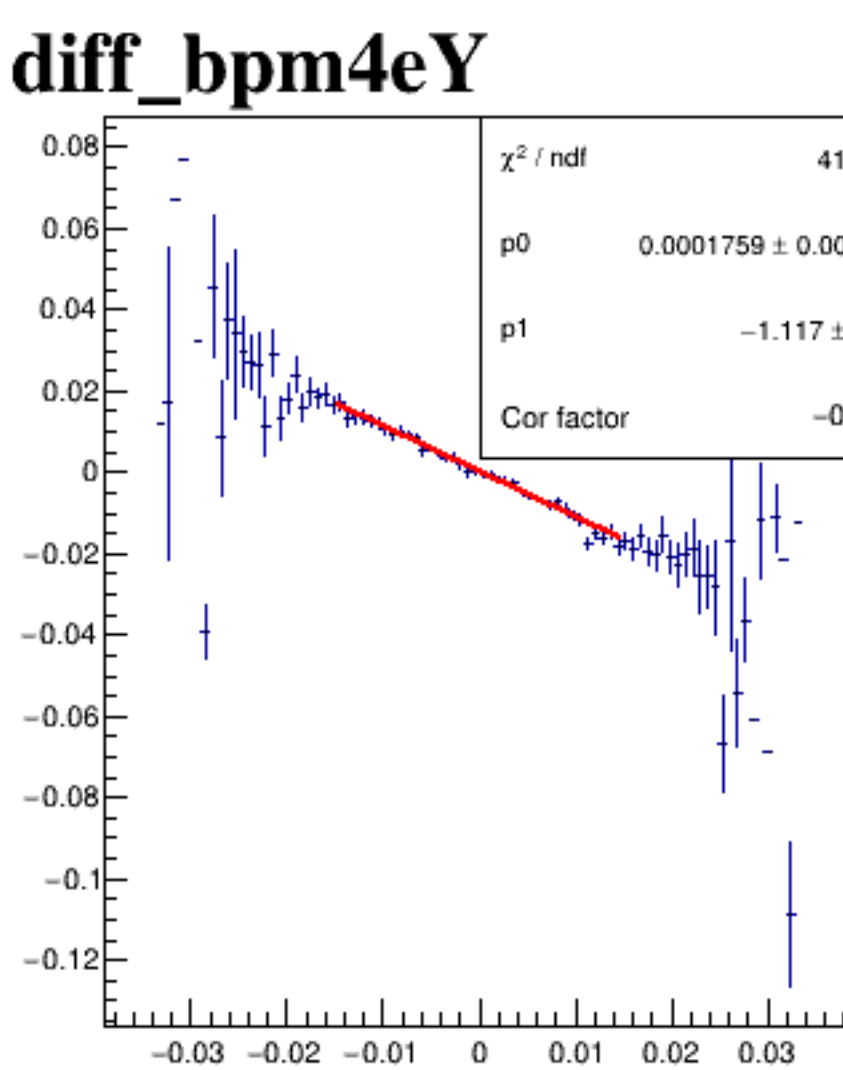
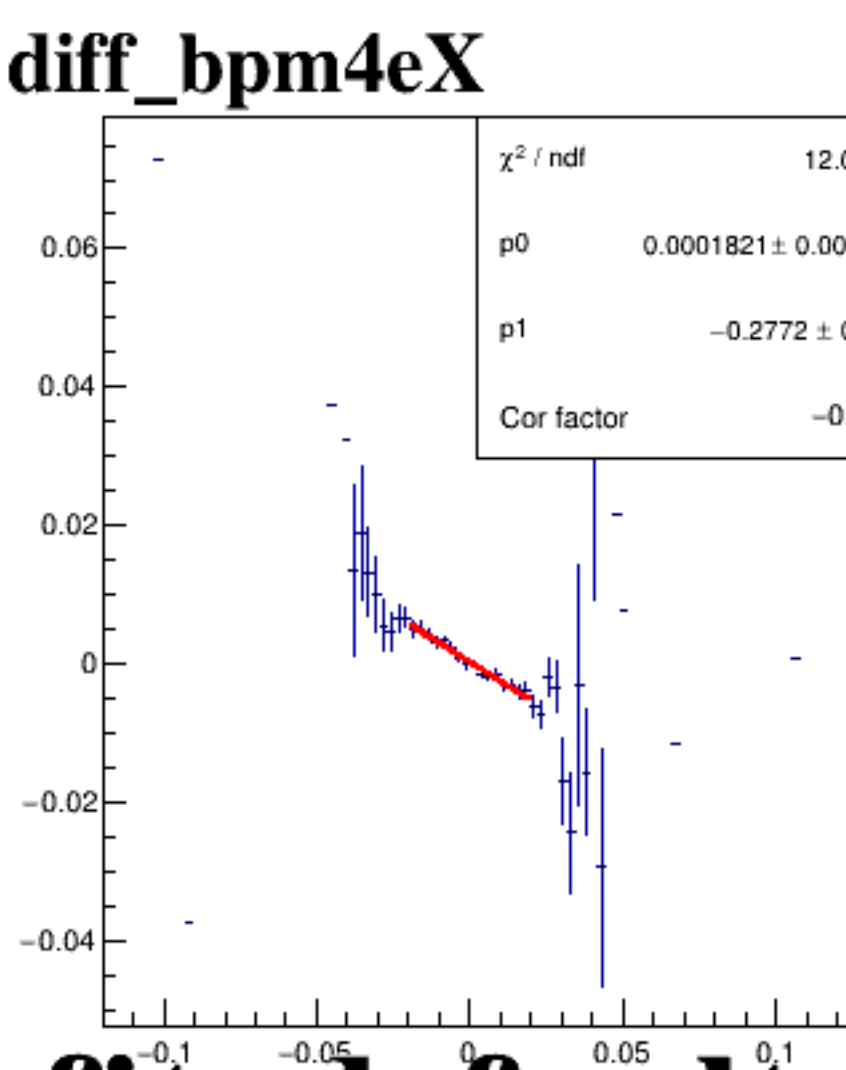
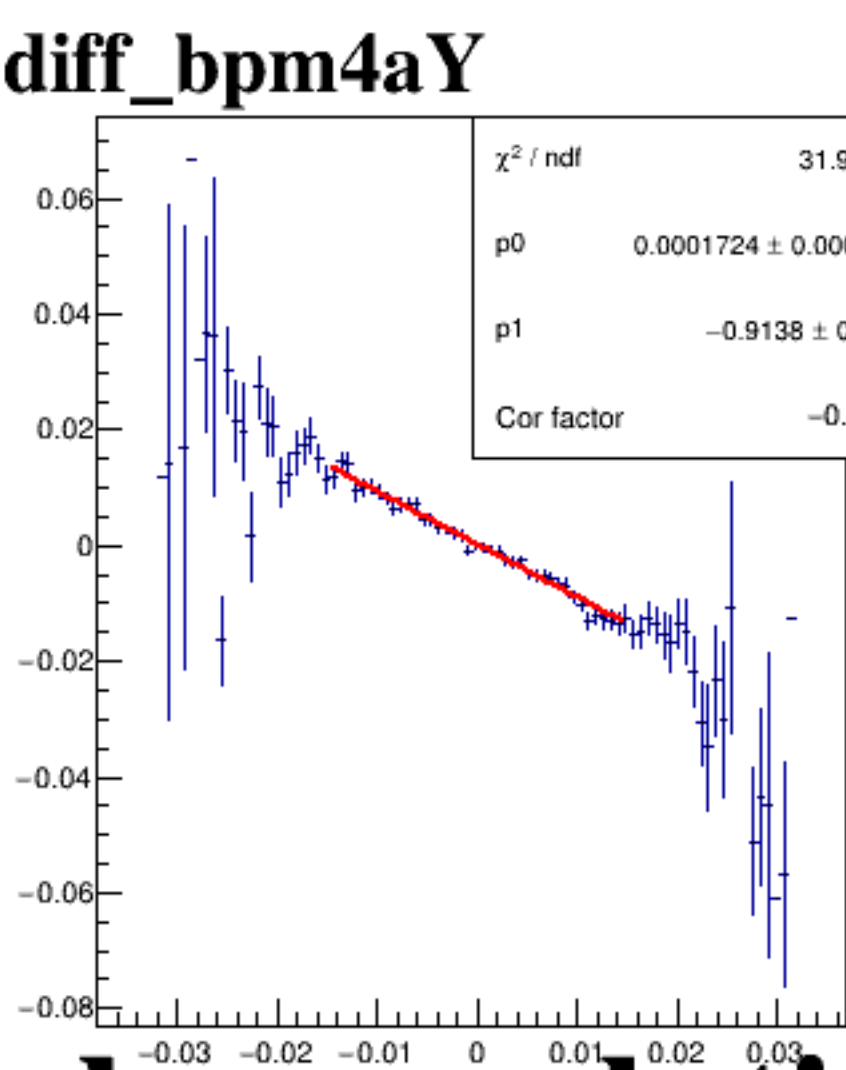
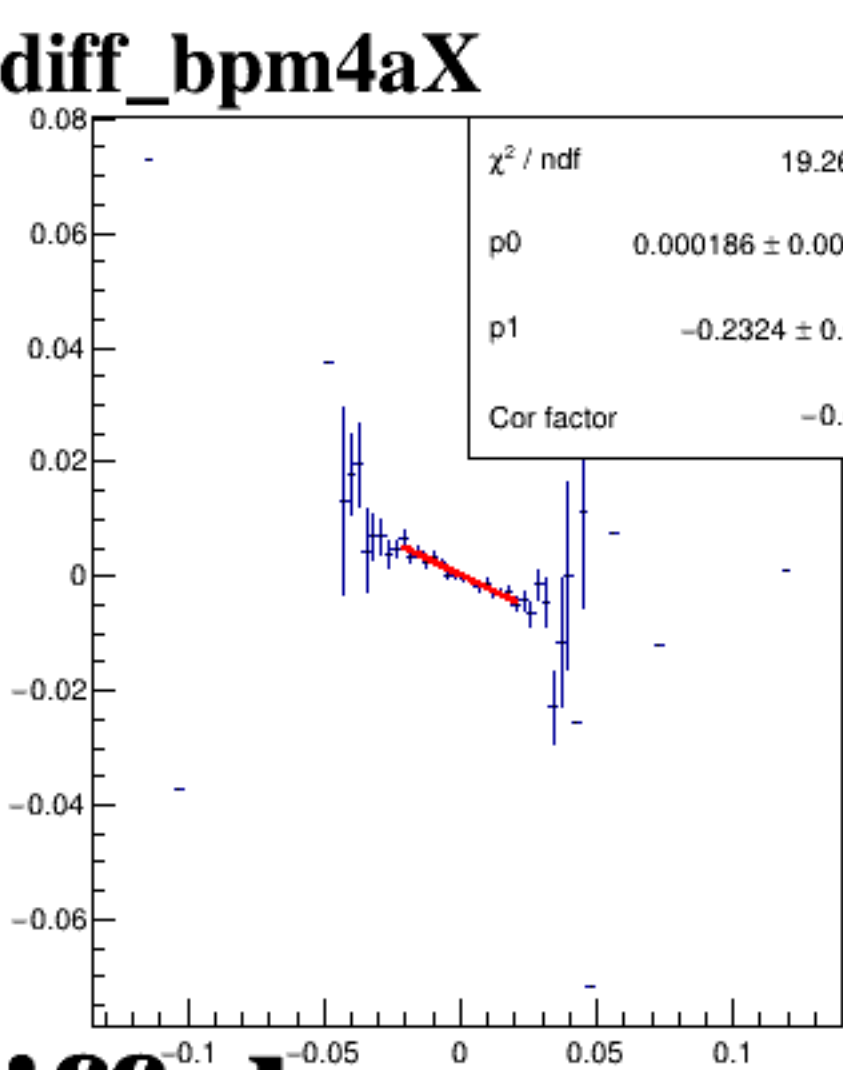
diff_bpm4eX



diff_bpm4eY



diff_bpm12X



run6380_000_diff_bpm_mutual_correlation_fit_default.png

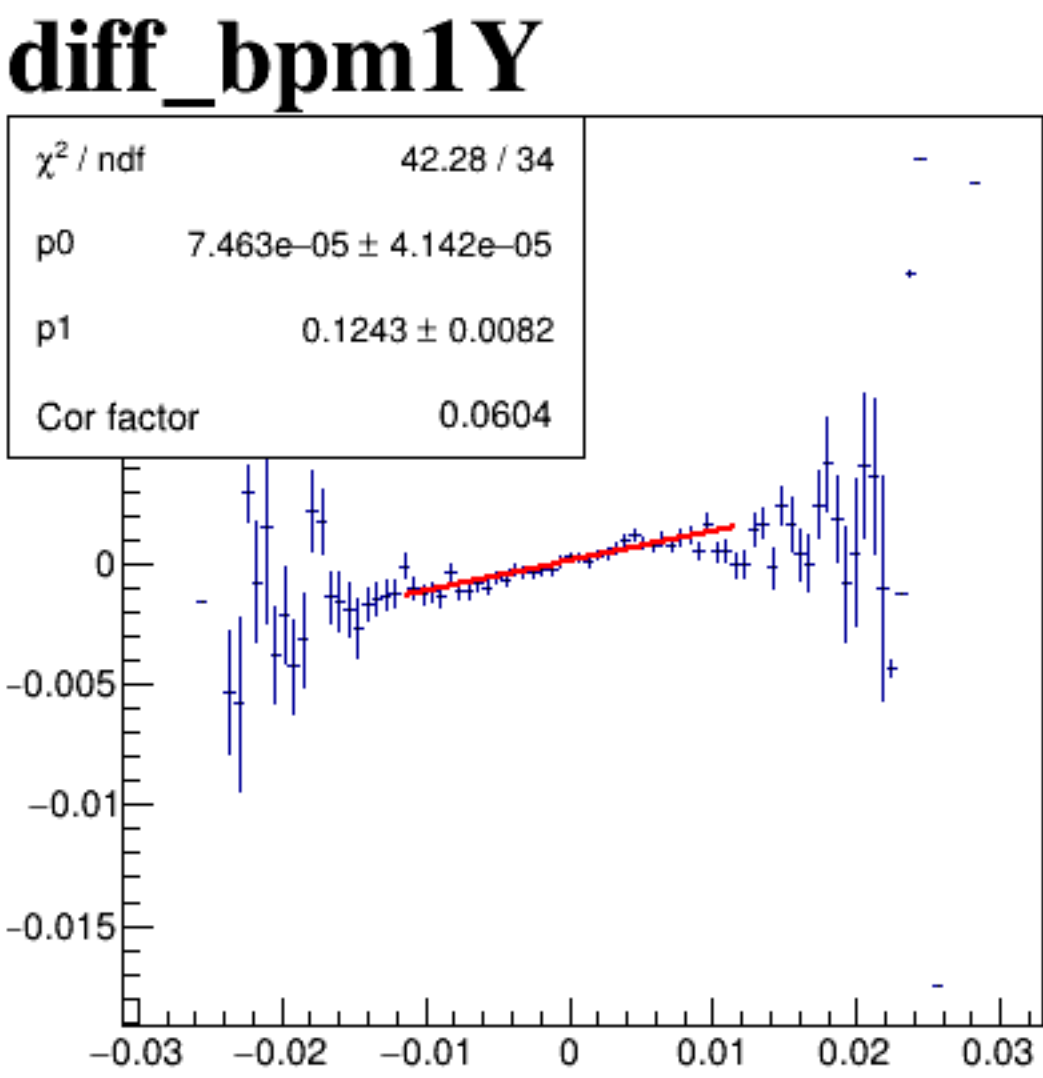
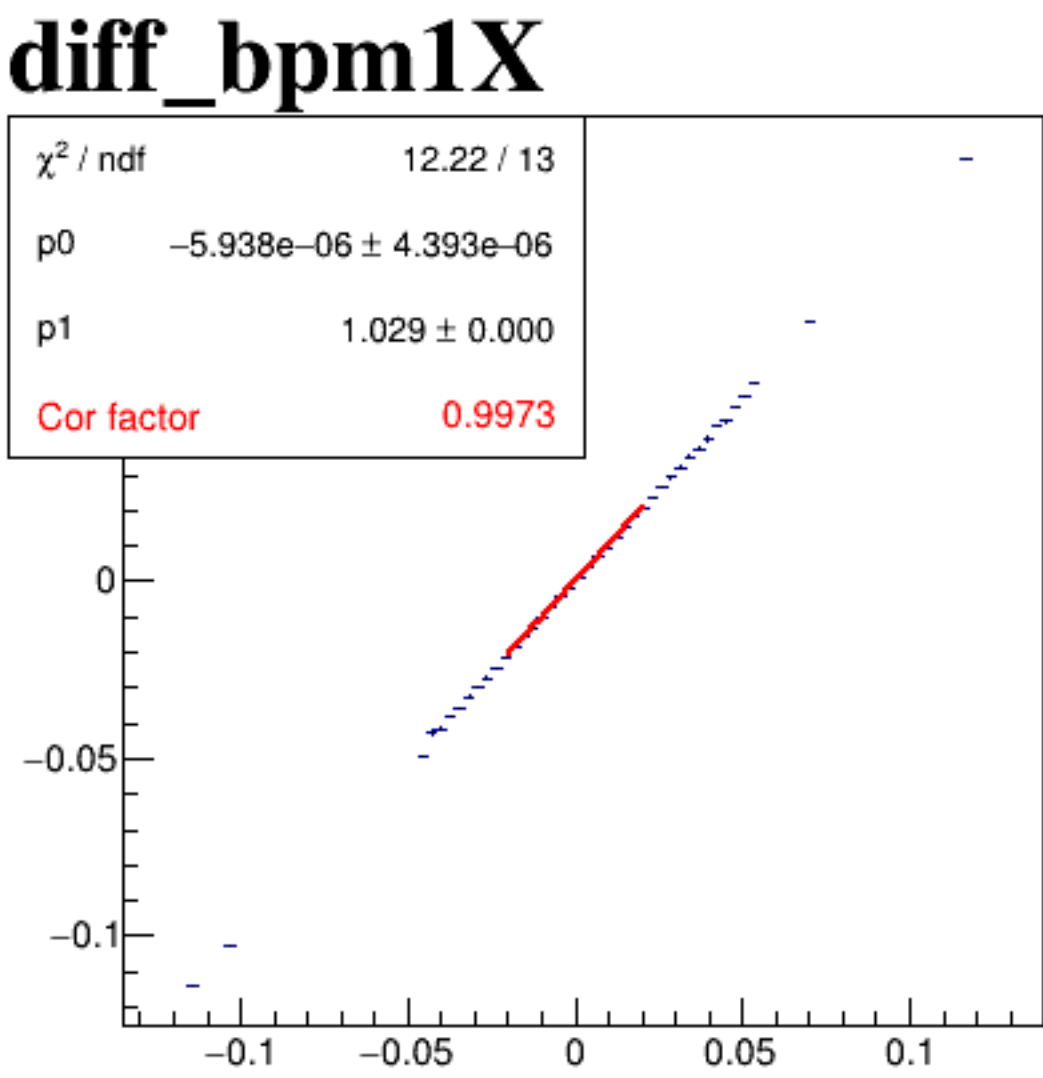
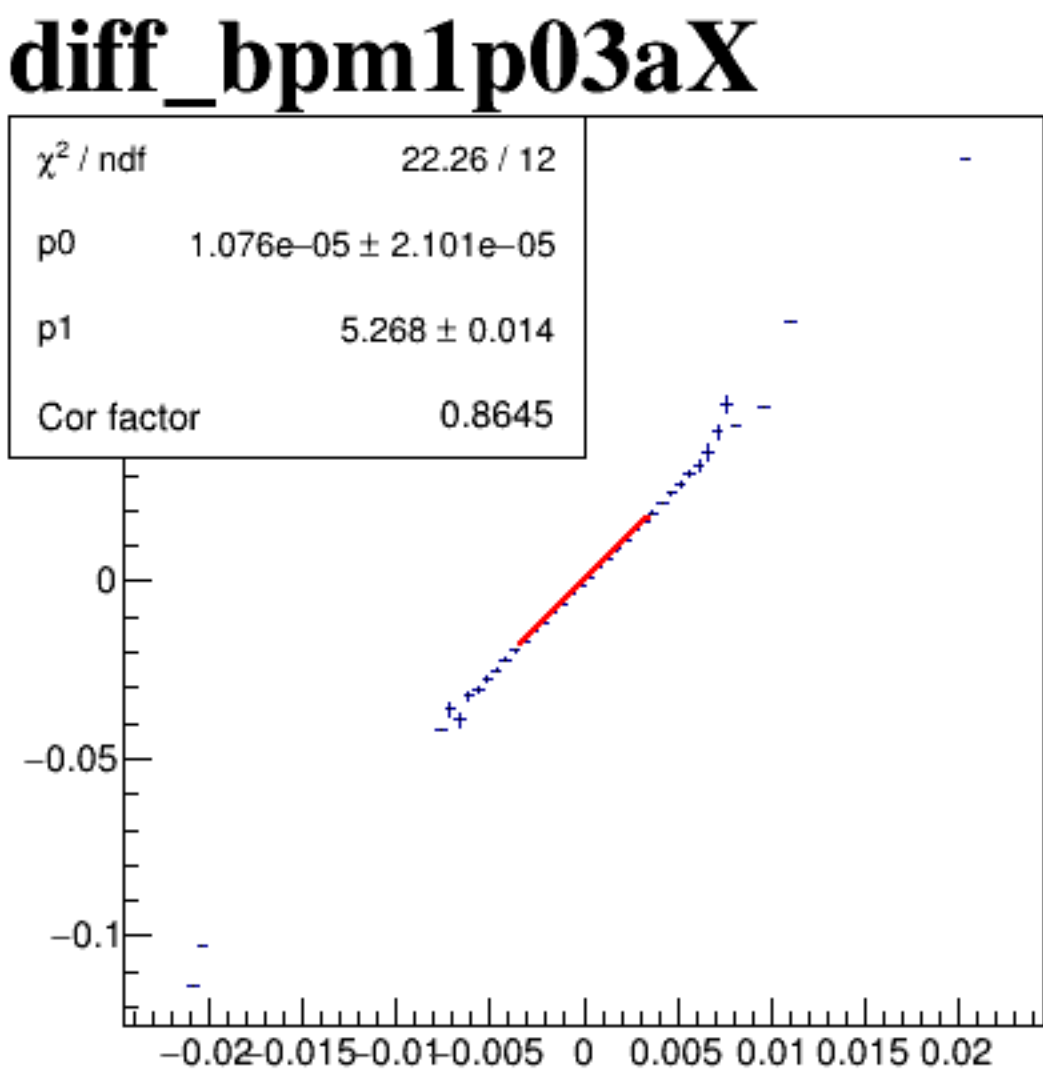
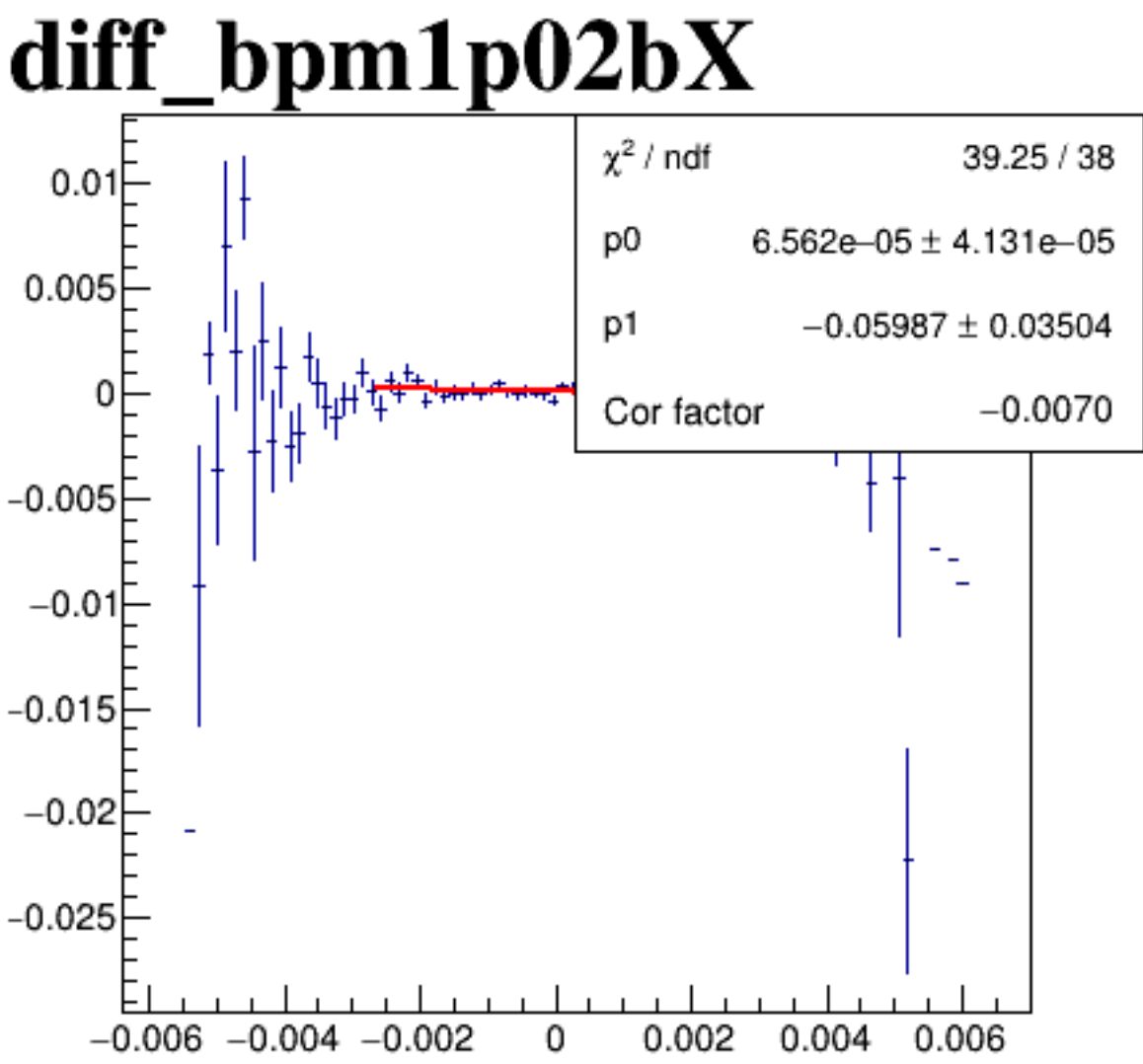
A scatter plot showing the relationship between the log2 fold change (y-axis) and the difference in binding per million (diff_bpm4aY, x-axis) for the 1000 most differentially expressed genes. The y-axis ranges from -0.1 to 0.1, and the x-axis ranges from -0.03 to 0.03. The data points are densely clustered around the origin (0,0), forming an elliptical shape elongated along the x-axis. There are a few outliers with higher log2 fold change values, particularly near the top of the plot.

A scatter plot showing the relationship between diff_bpm4ex on the x-axis and diff_bpm4ex on the y-axis. The data points are tightly clustered along a diagonal line, indicating a strong positive correlation. The axes range from -0.1 to 0.1.

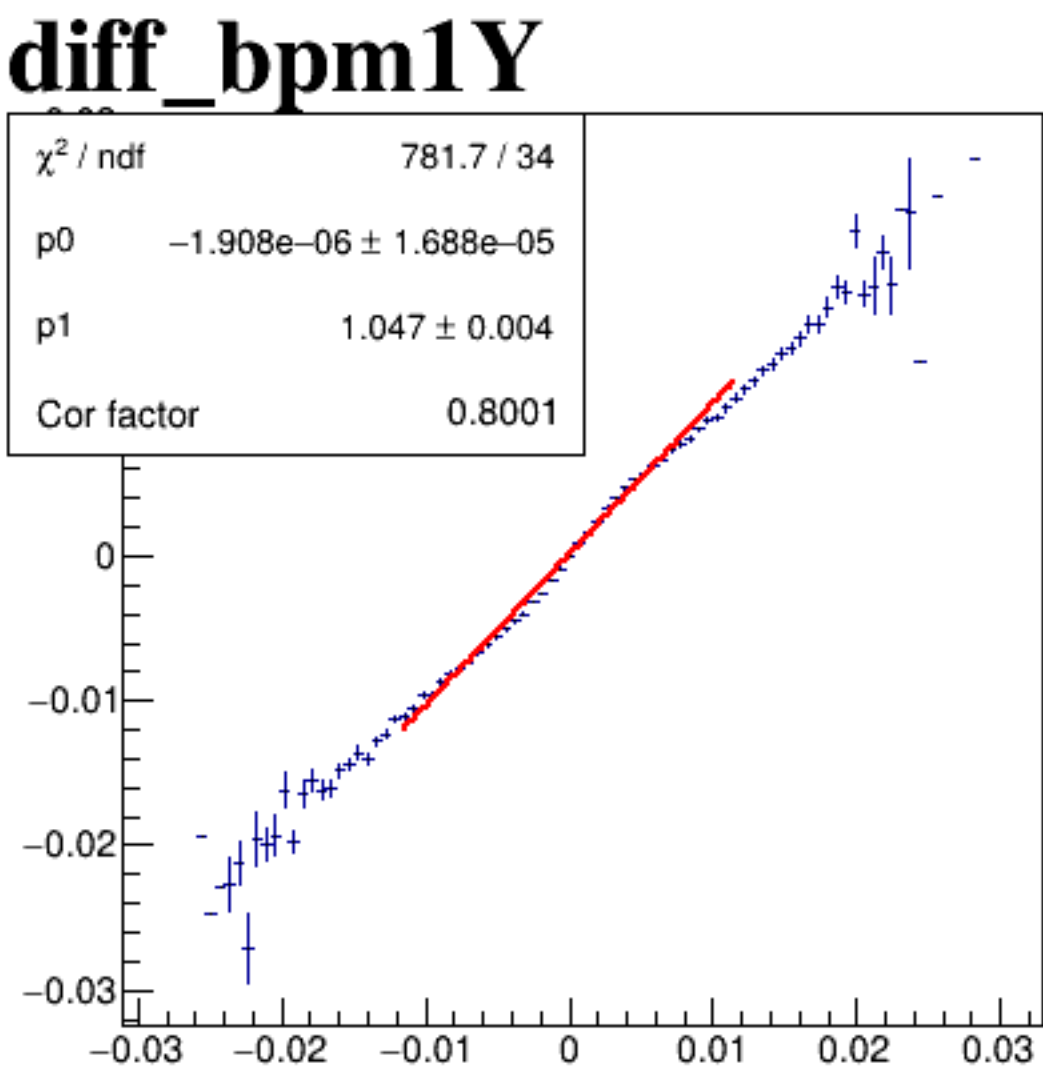
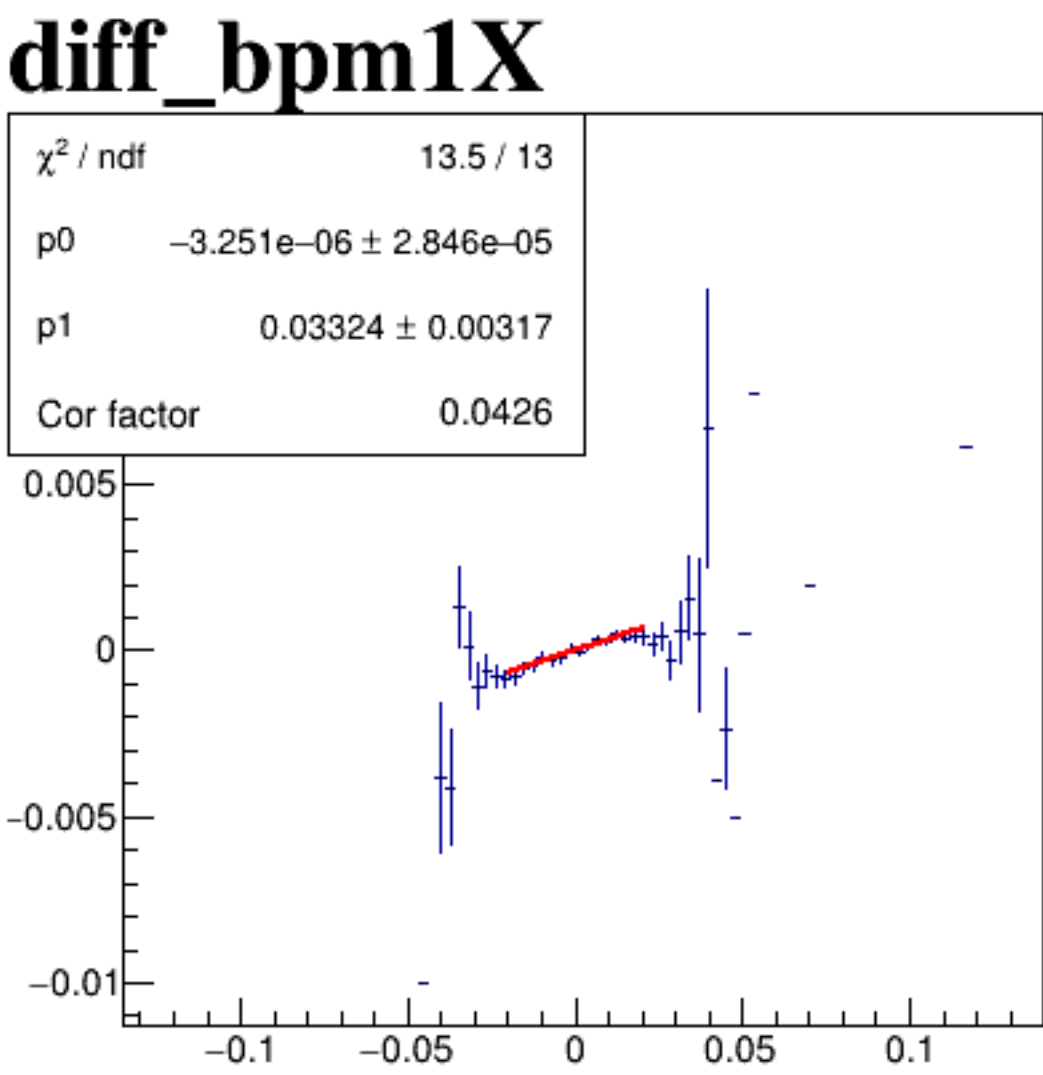
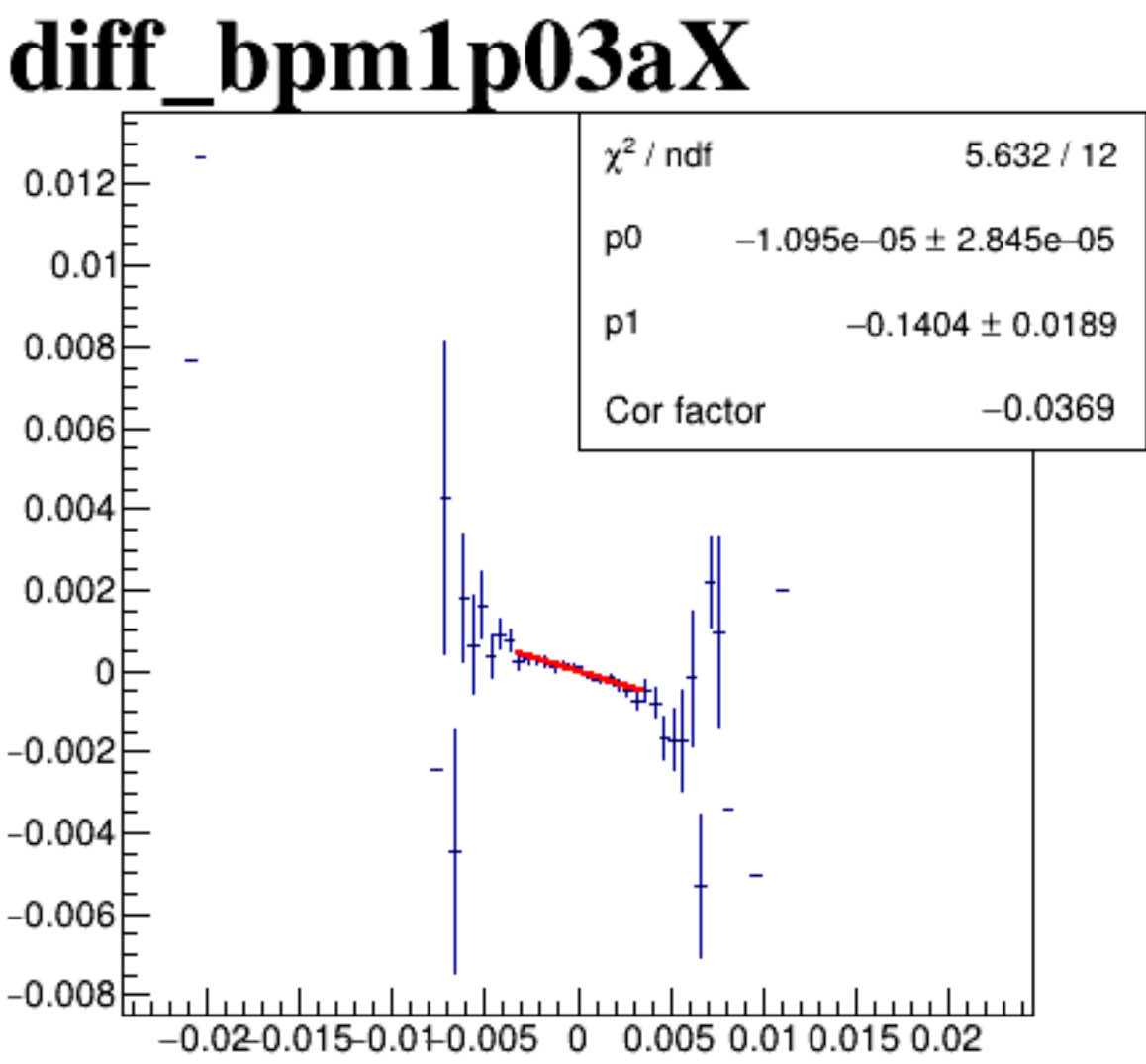
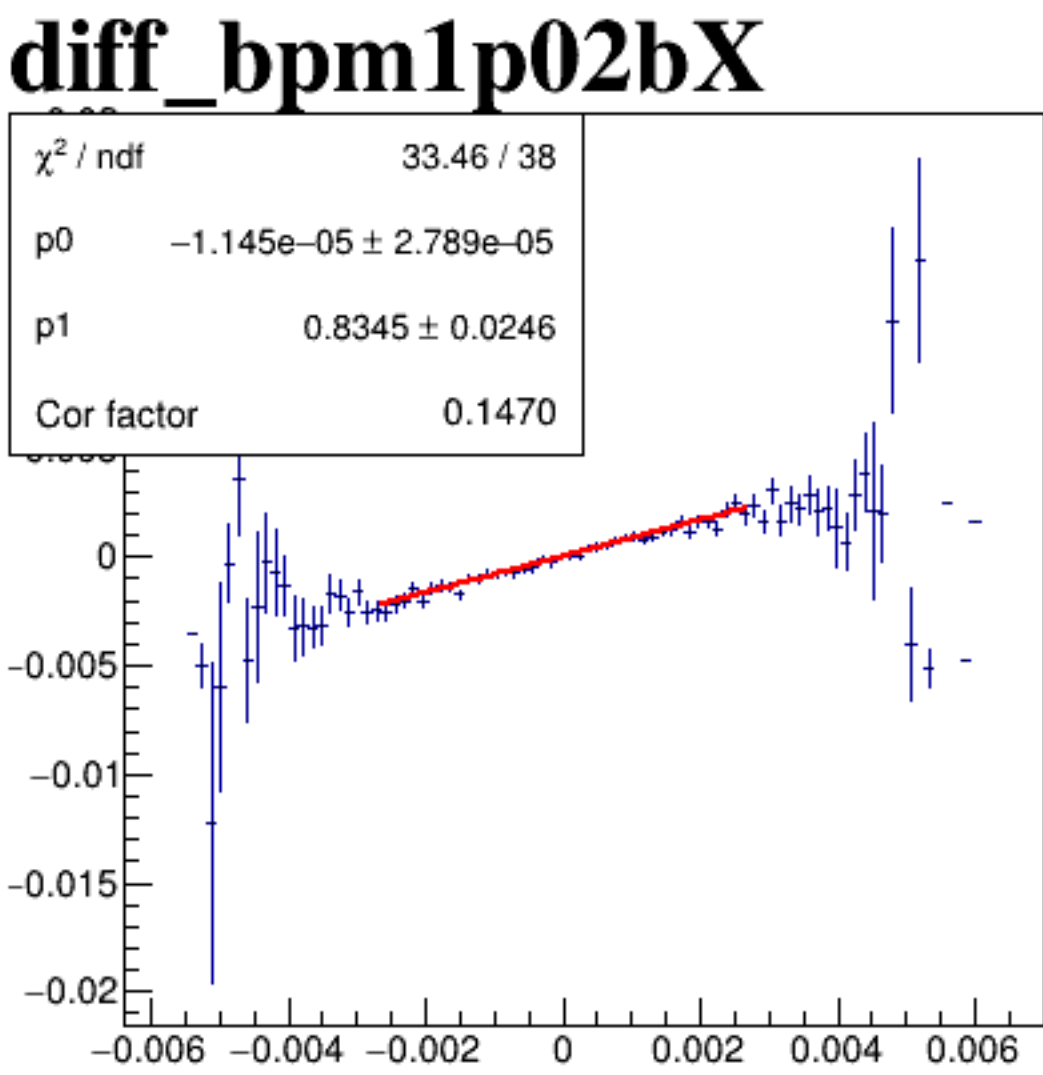
A scatter plot showing the correlation coefficient between the two models. The x-axis is labeled 'Correlation' and ranges from -0.03 to 0.03. The y-axis is labeled 'Correlation' and ranges from -0.15 to 0.15. The data points are densely clustered around the origin (0,0), forming a circular cloud of points. The density of points is highest in the center and decreases as they move away from the origin.

run6380_000_diff_bpm_mutual_correlation_scat_default.png

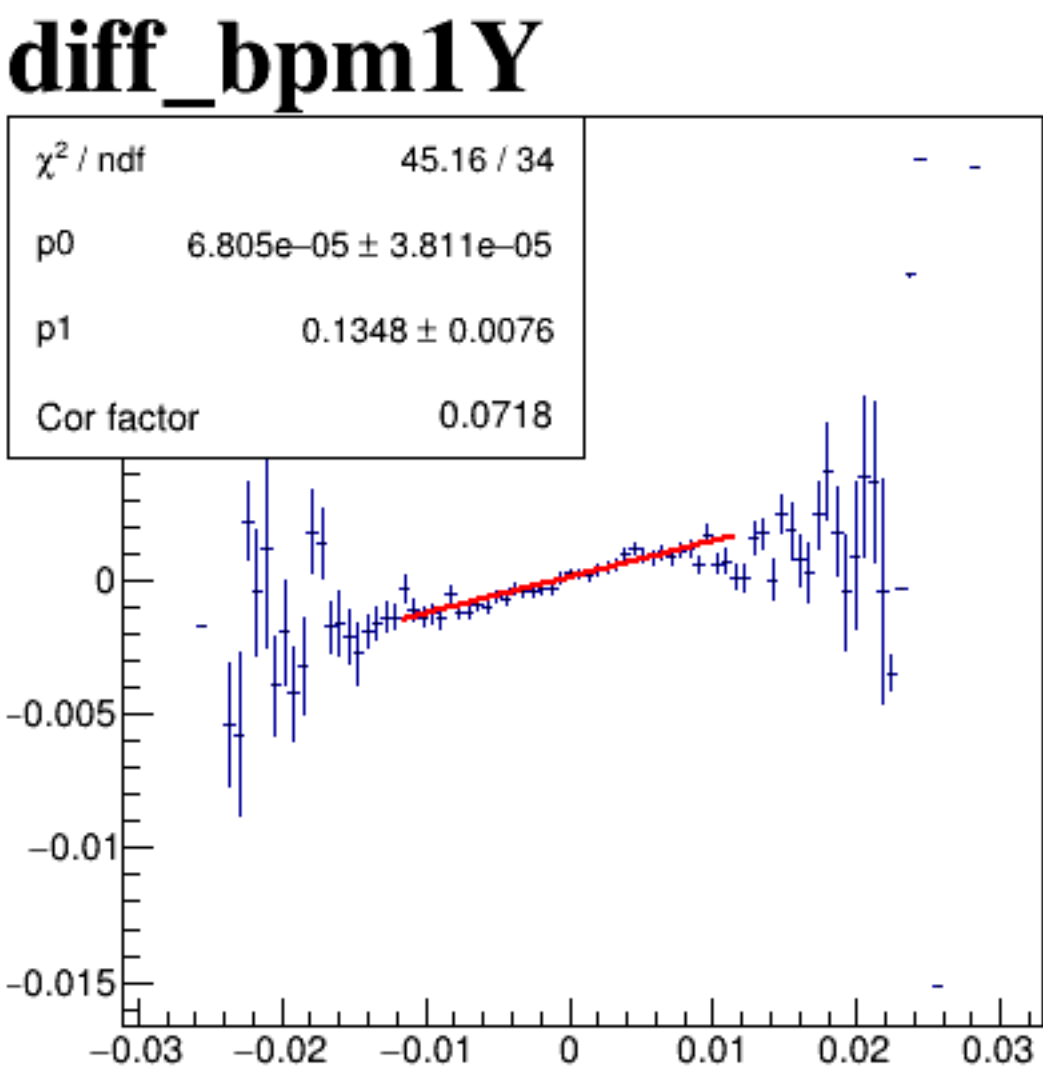
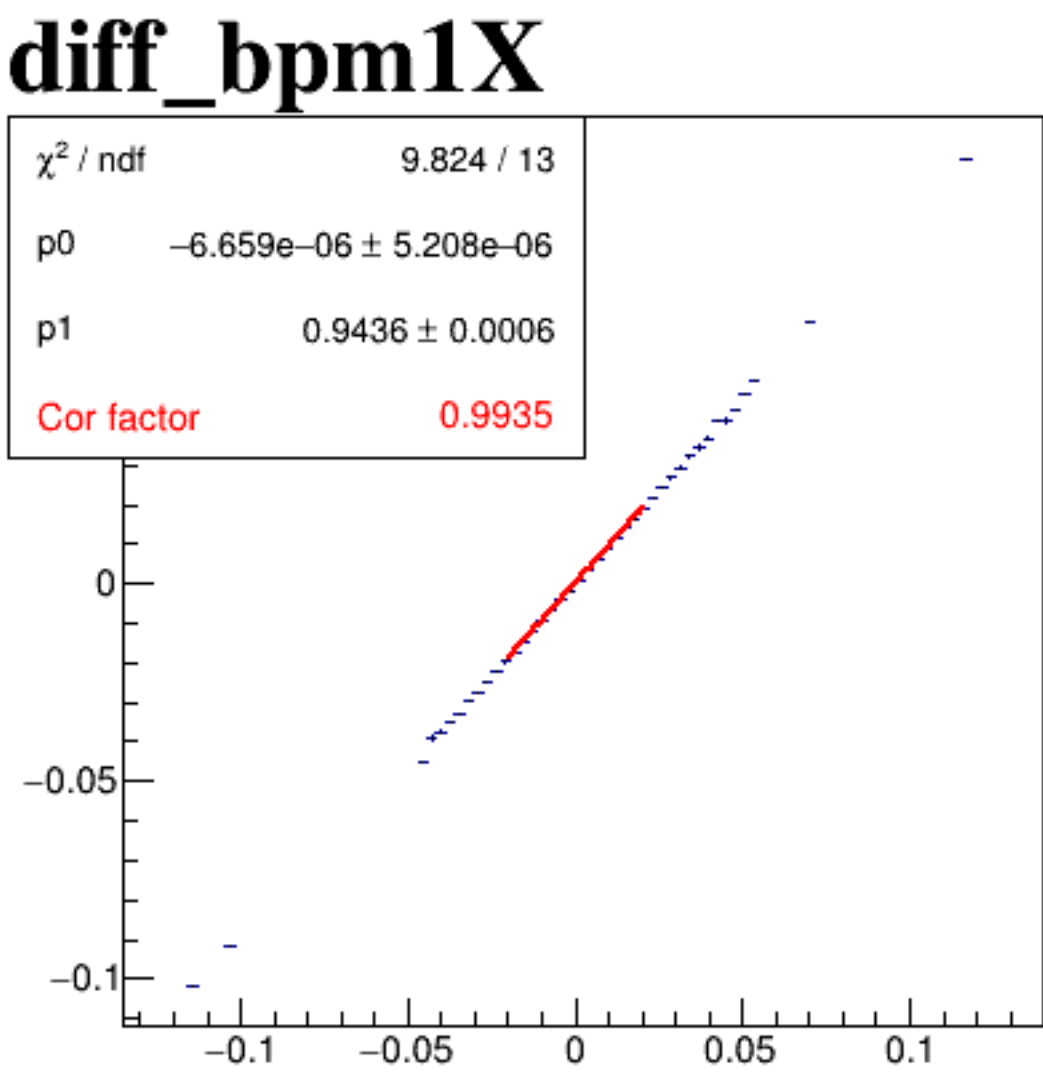
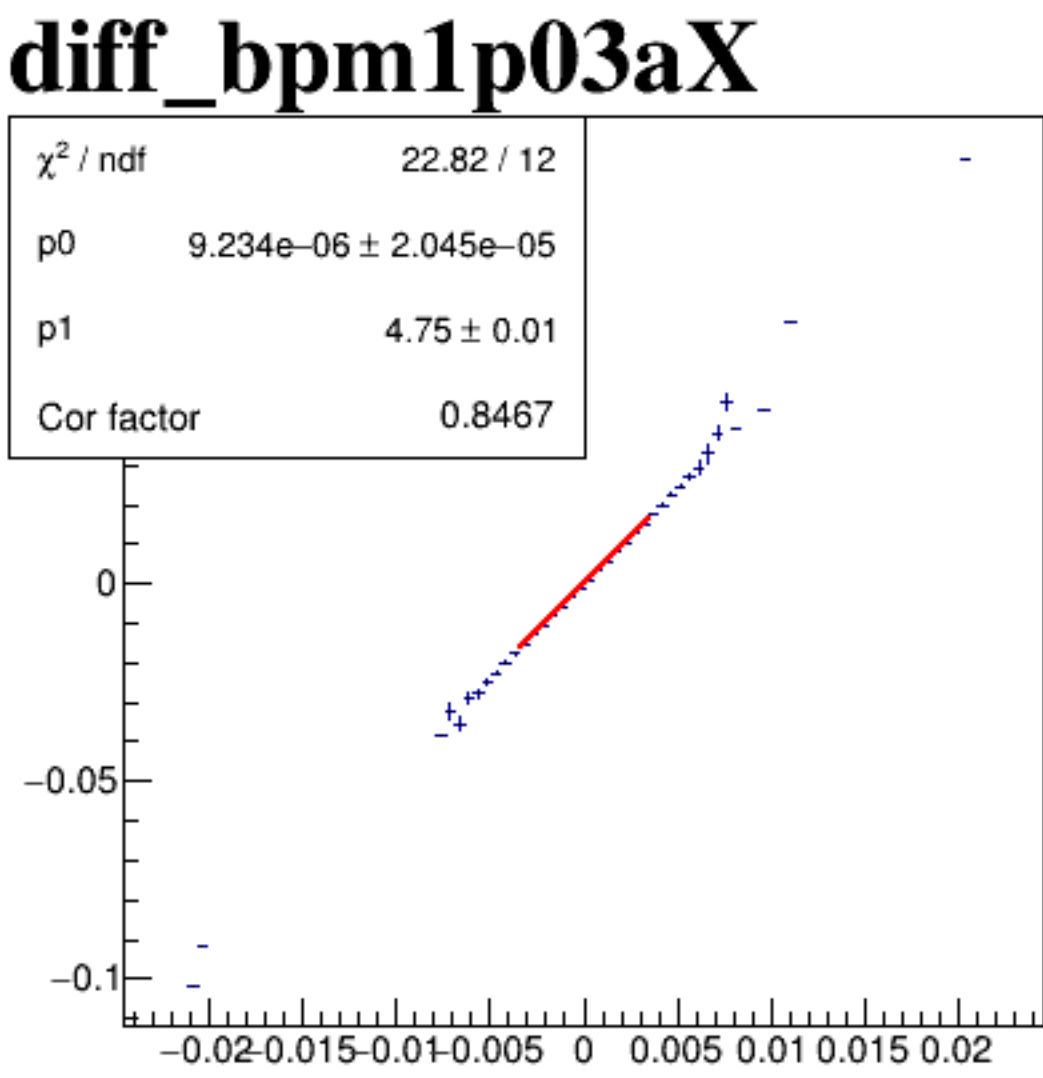
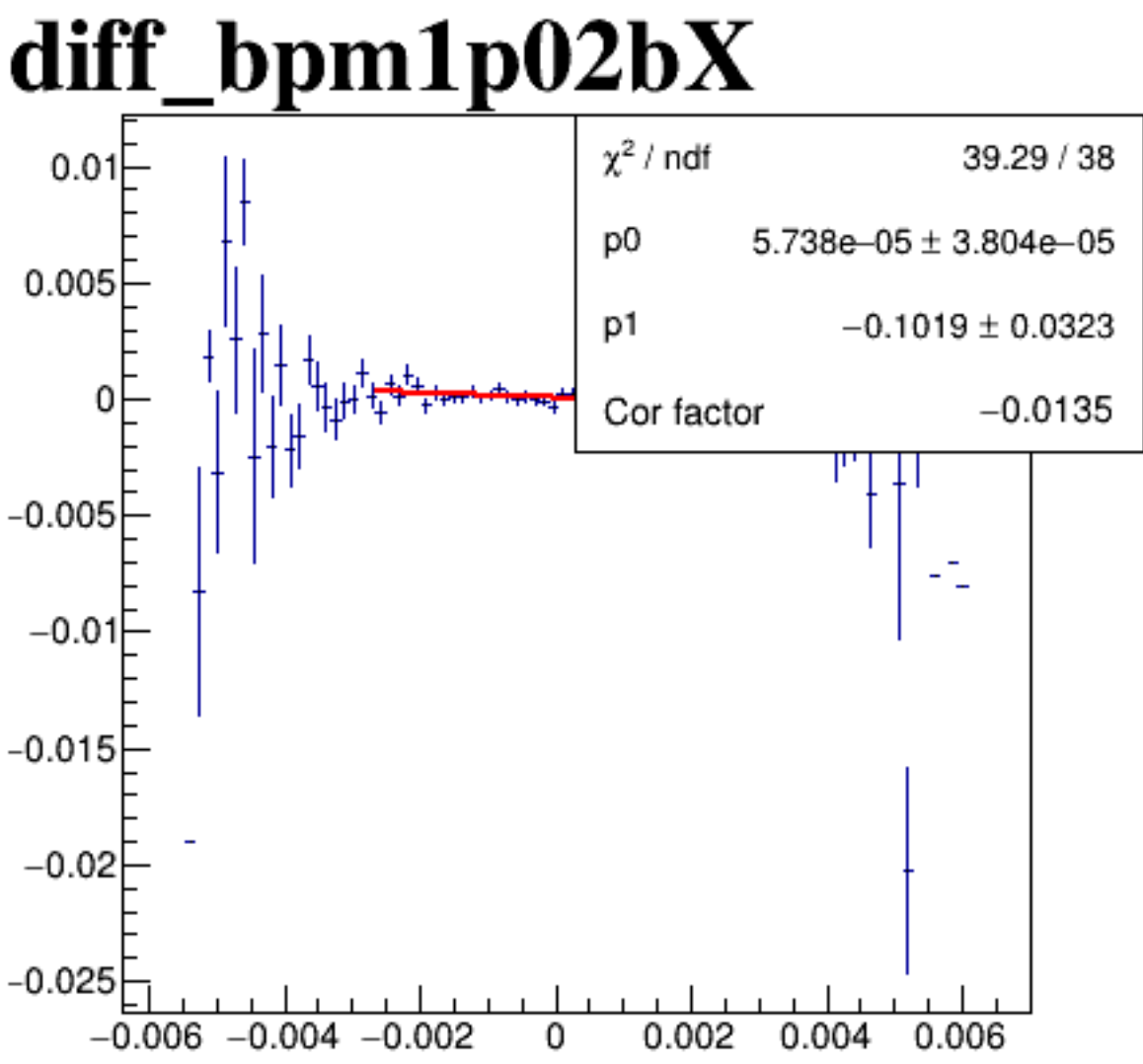
diff_bpm4aX



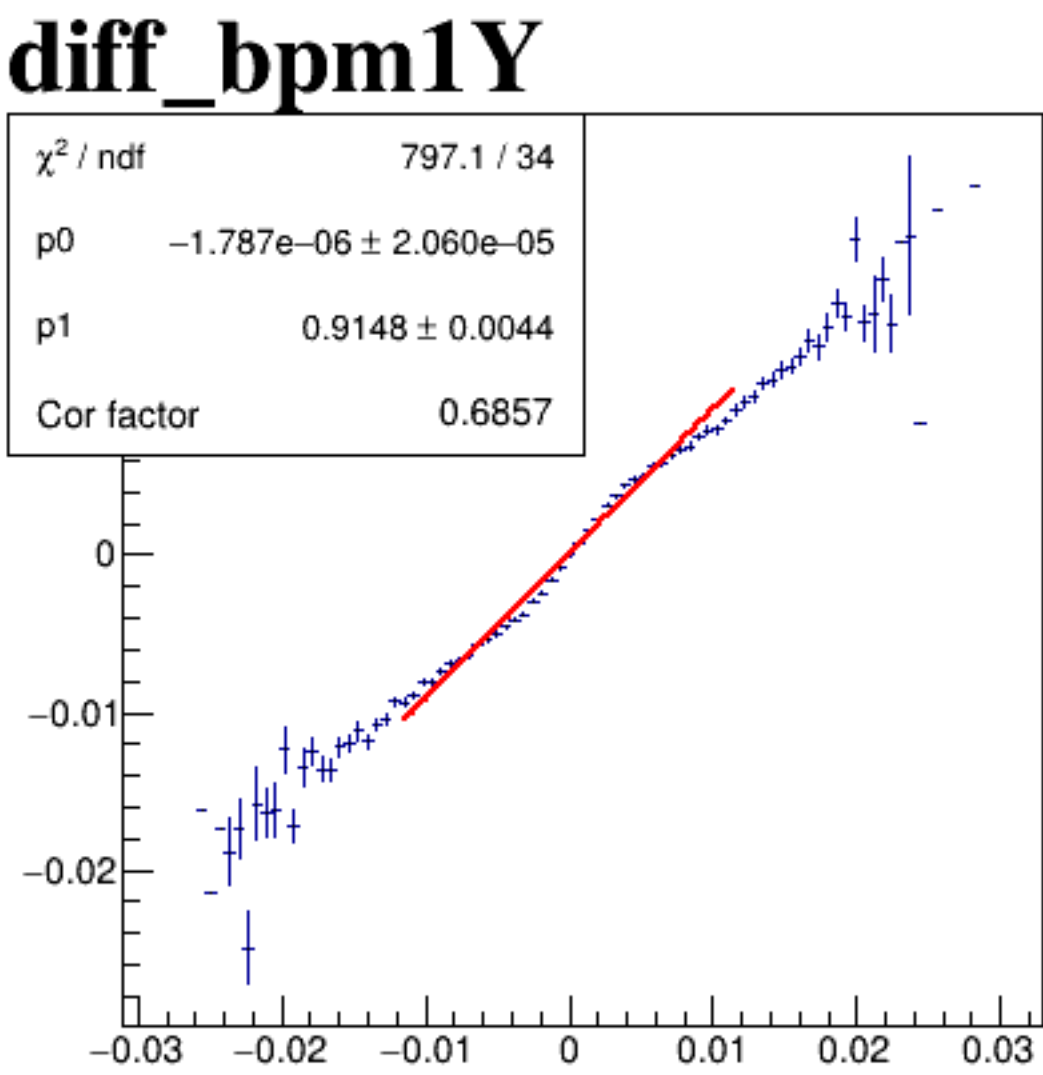
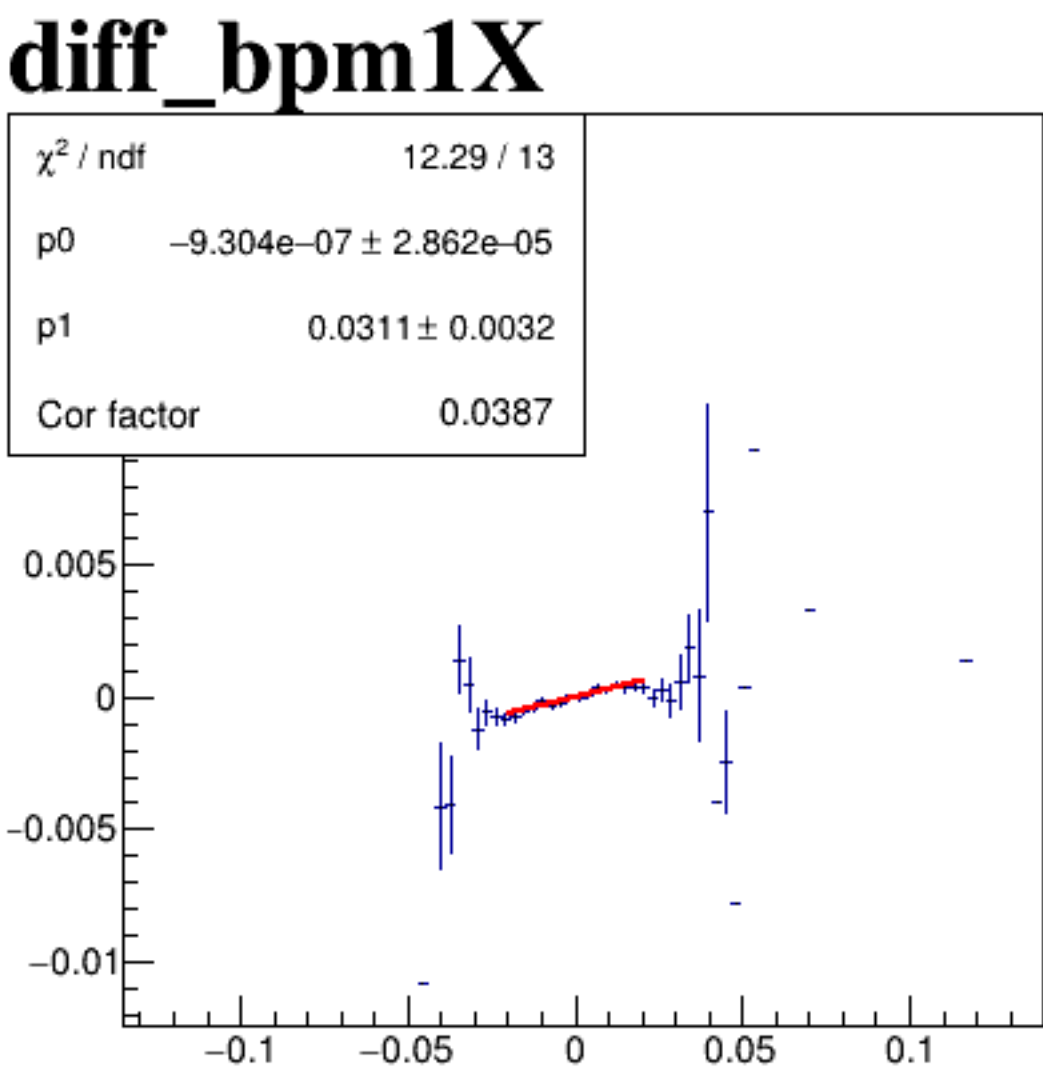
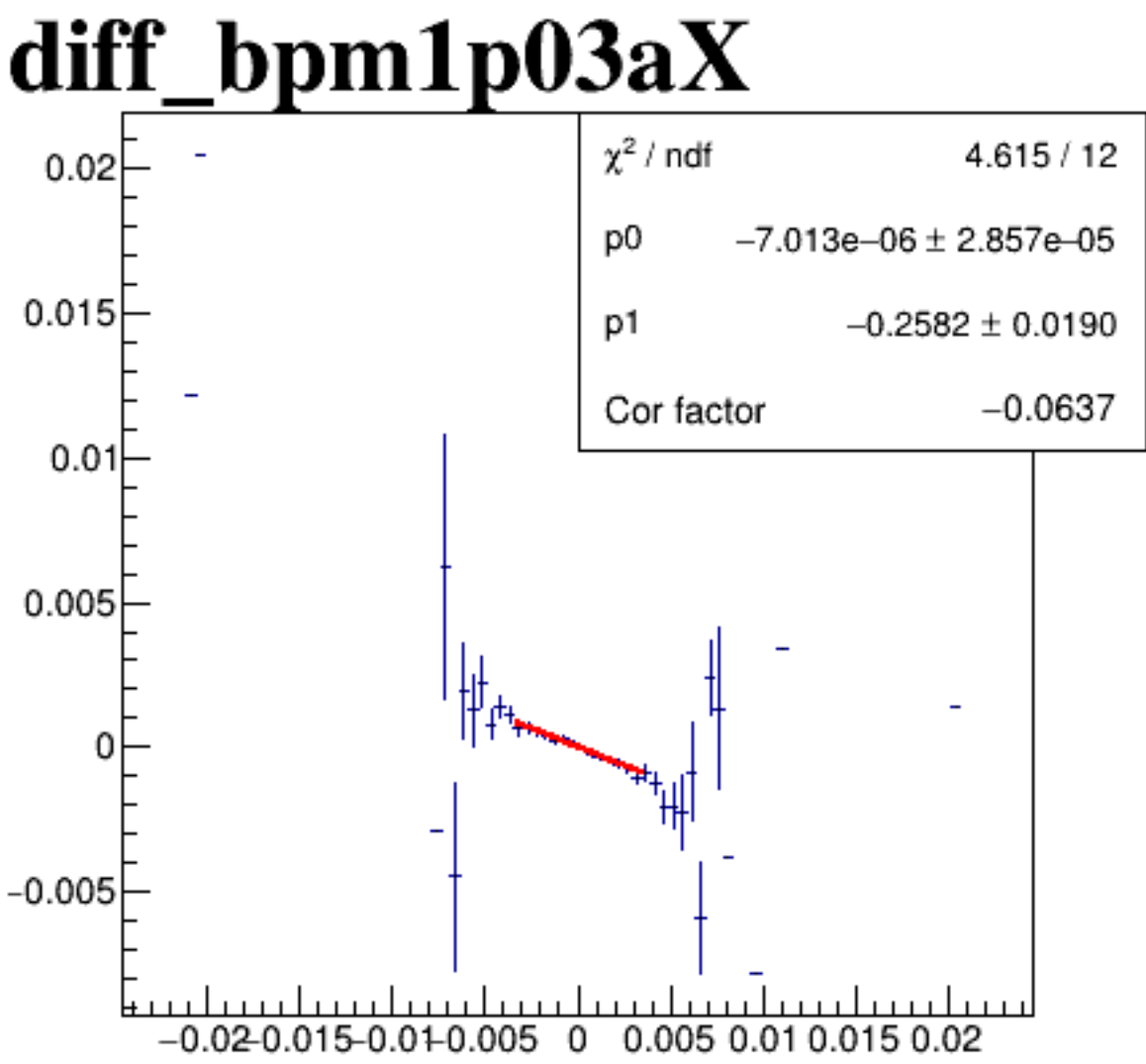
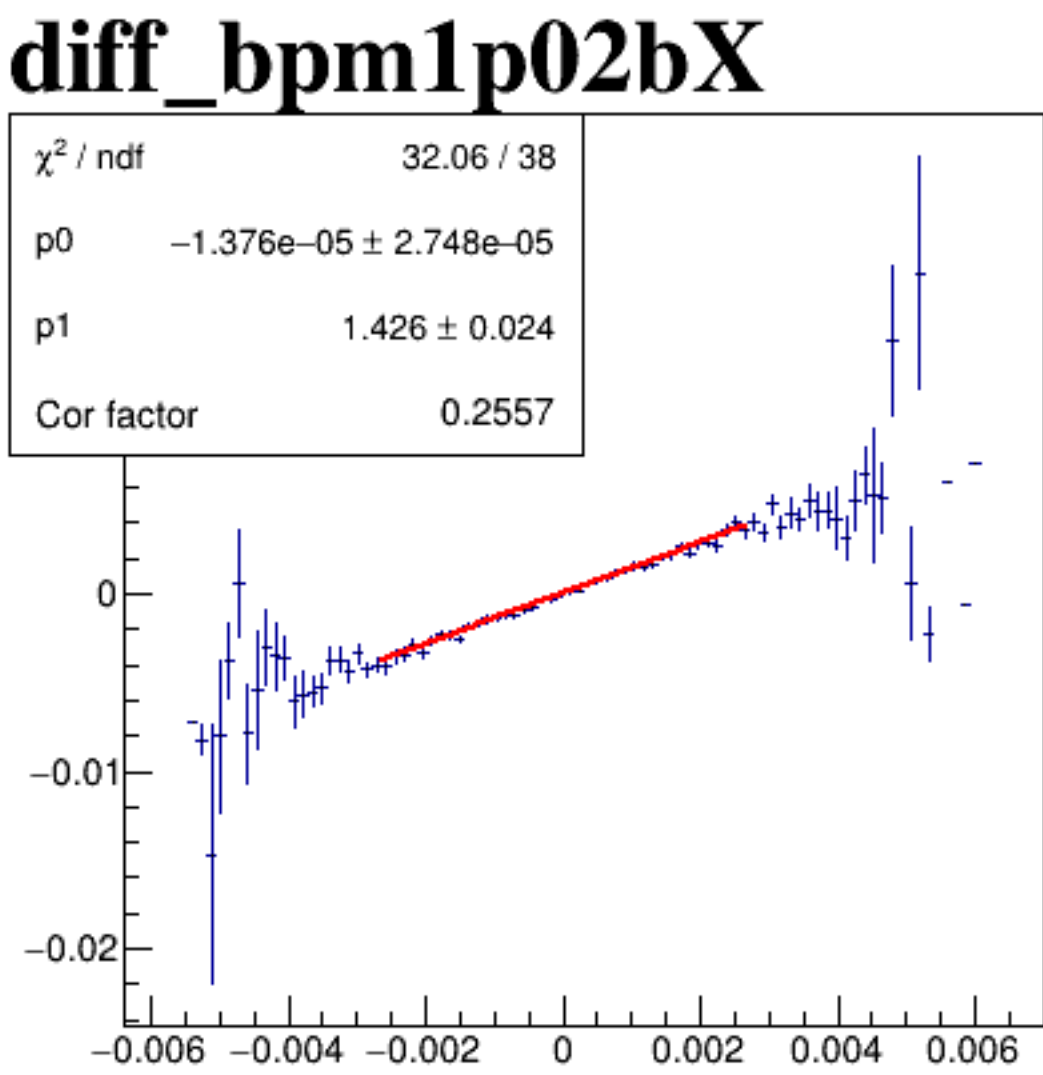
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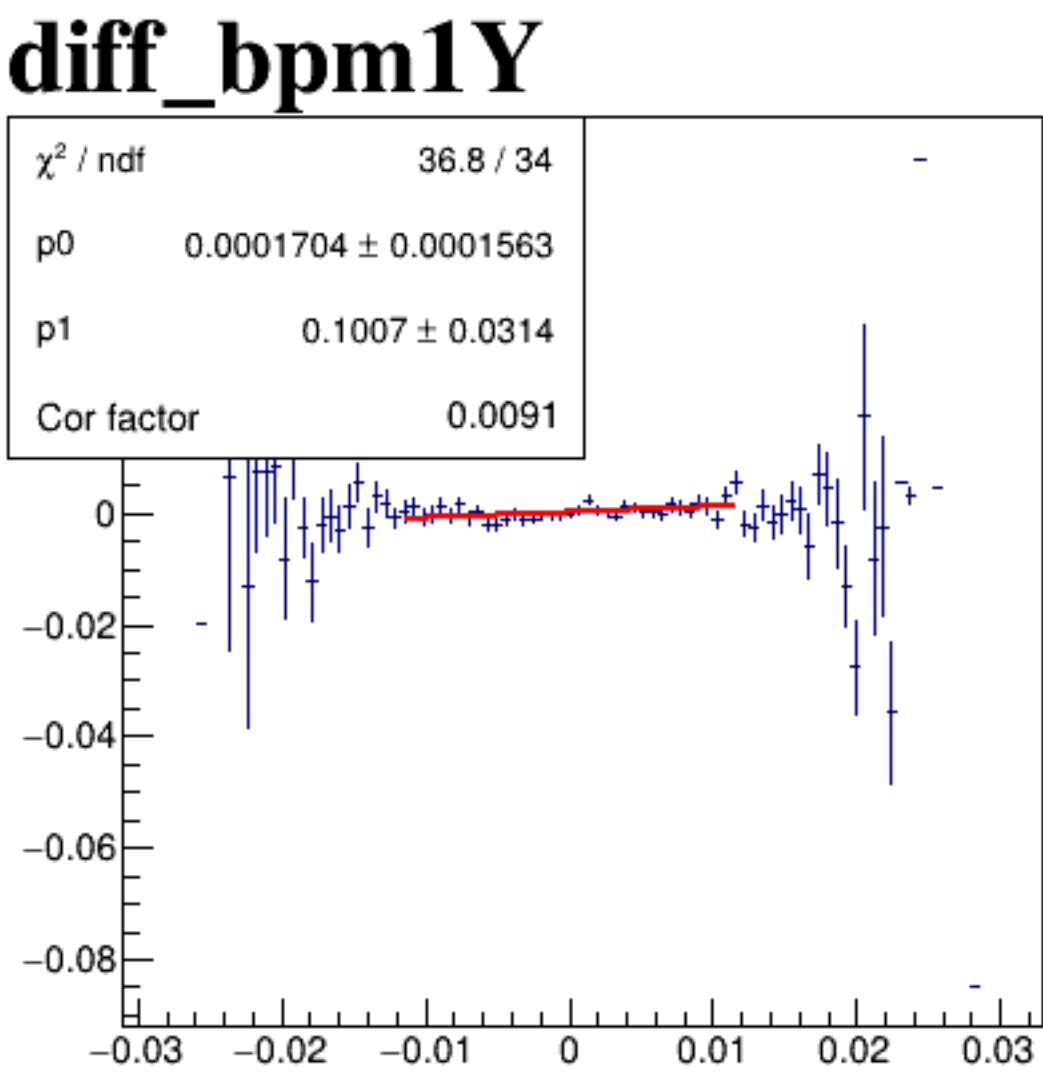
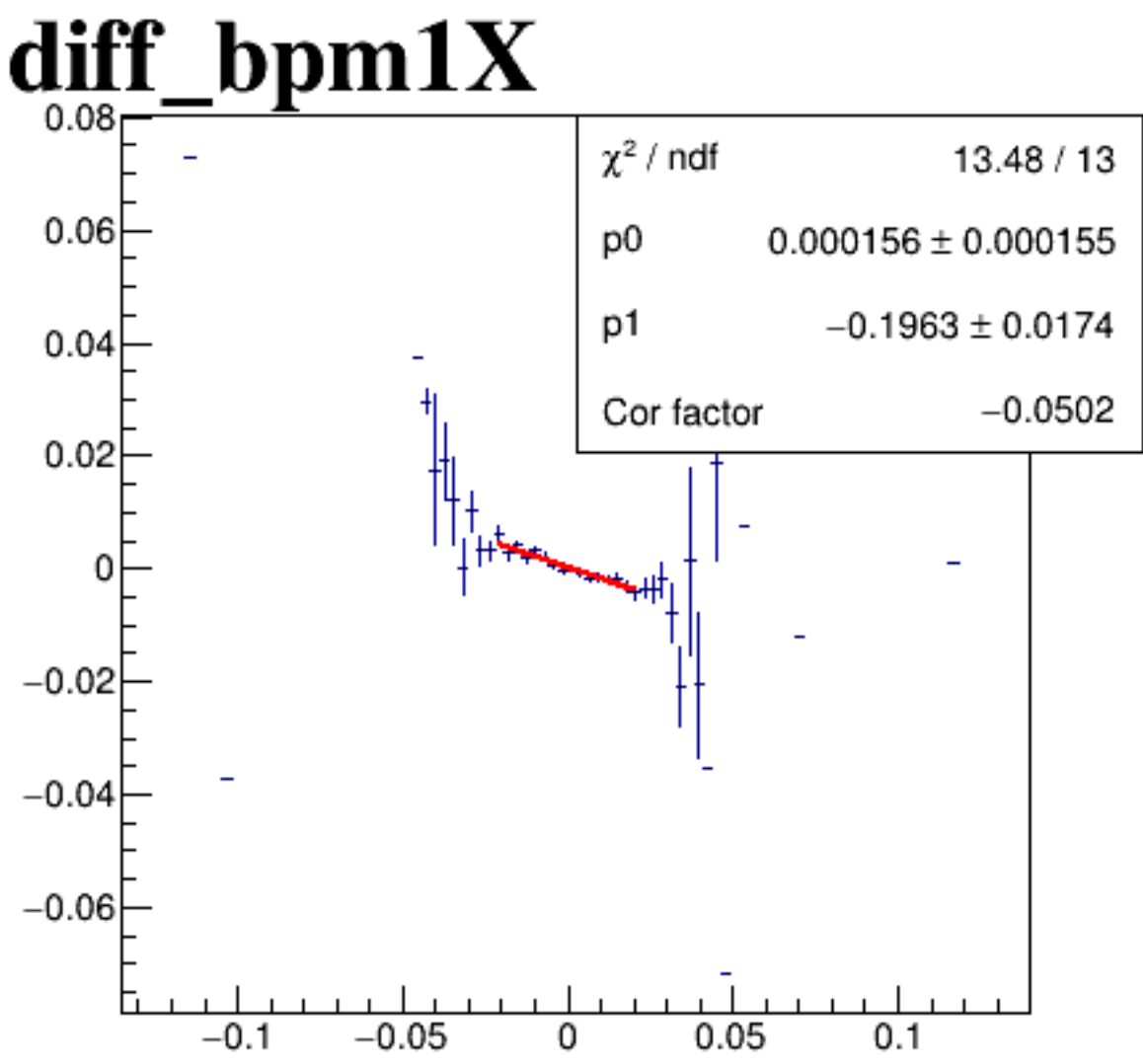
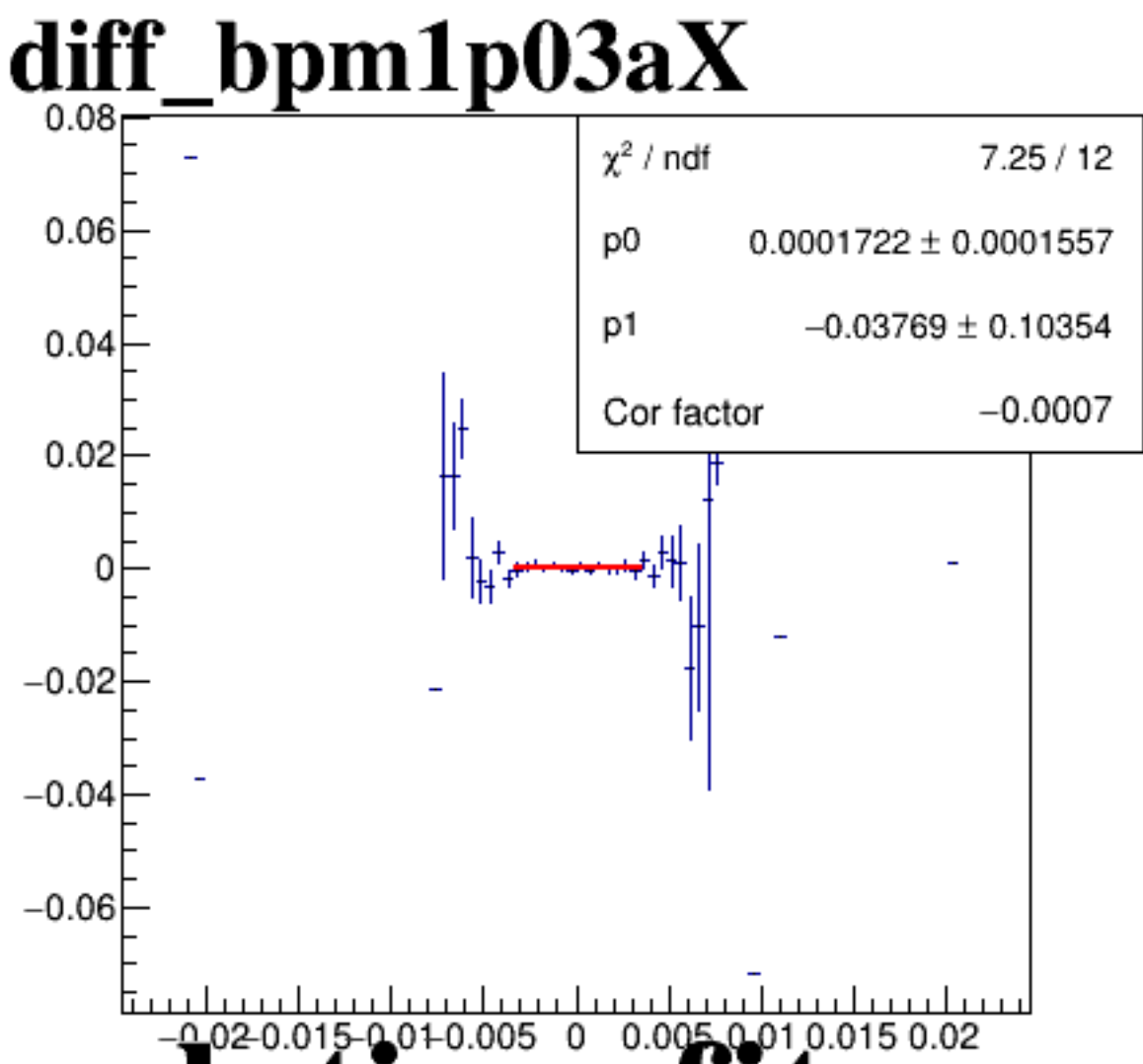
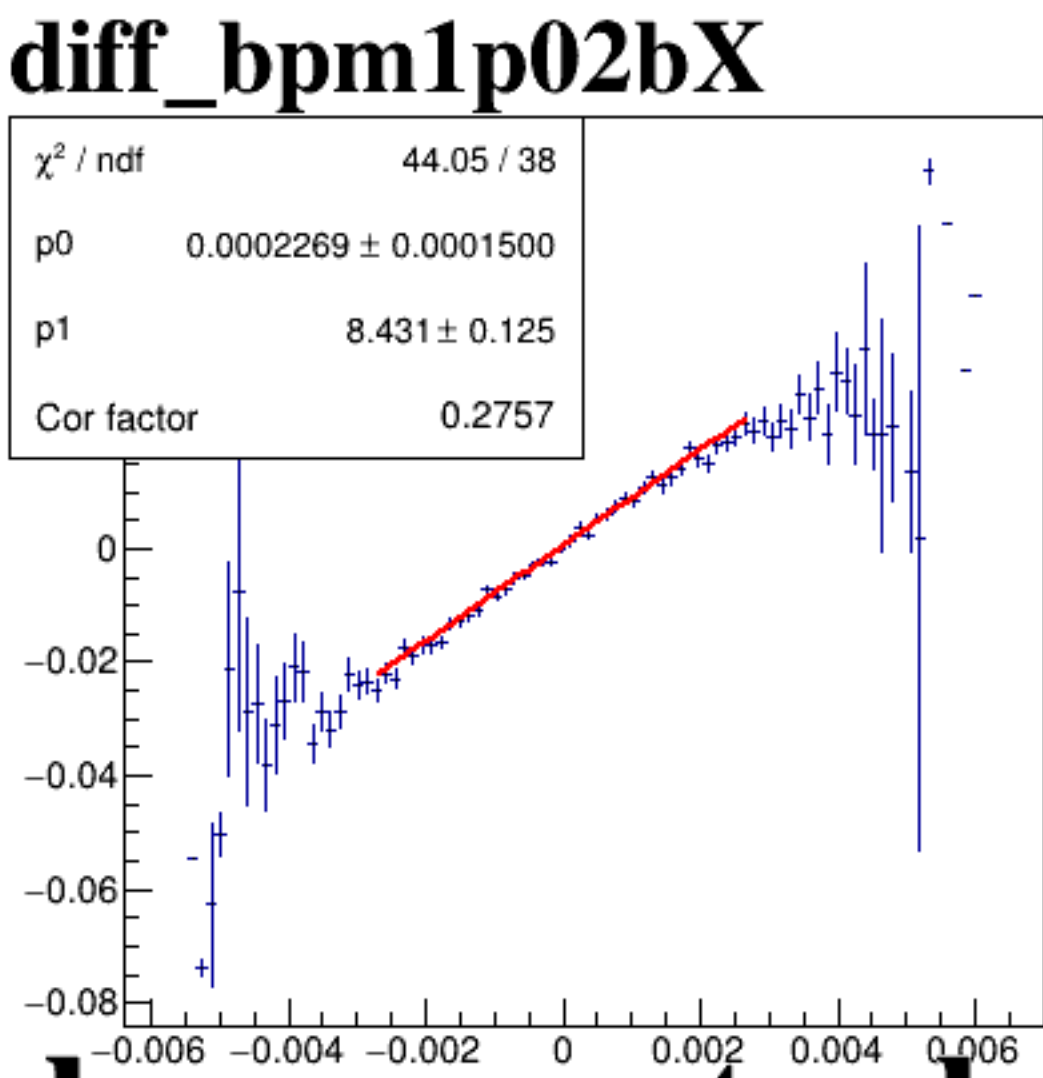
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diff_bpm4eY



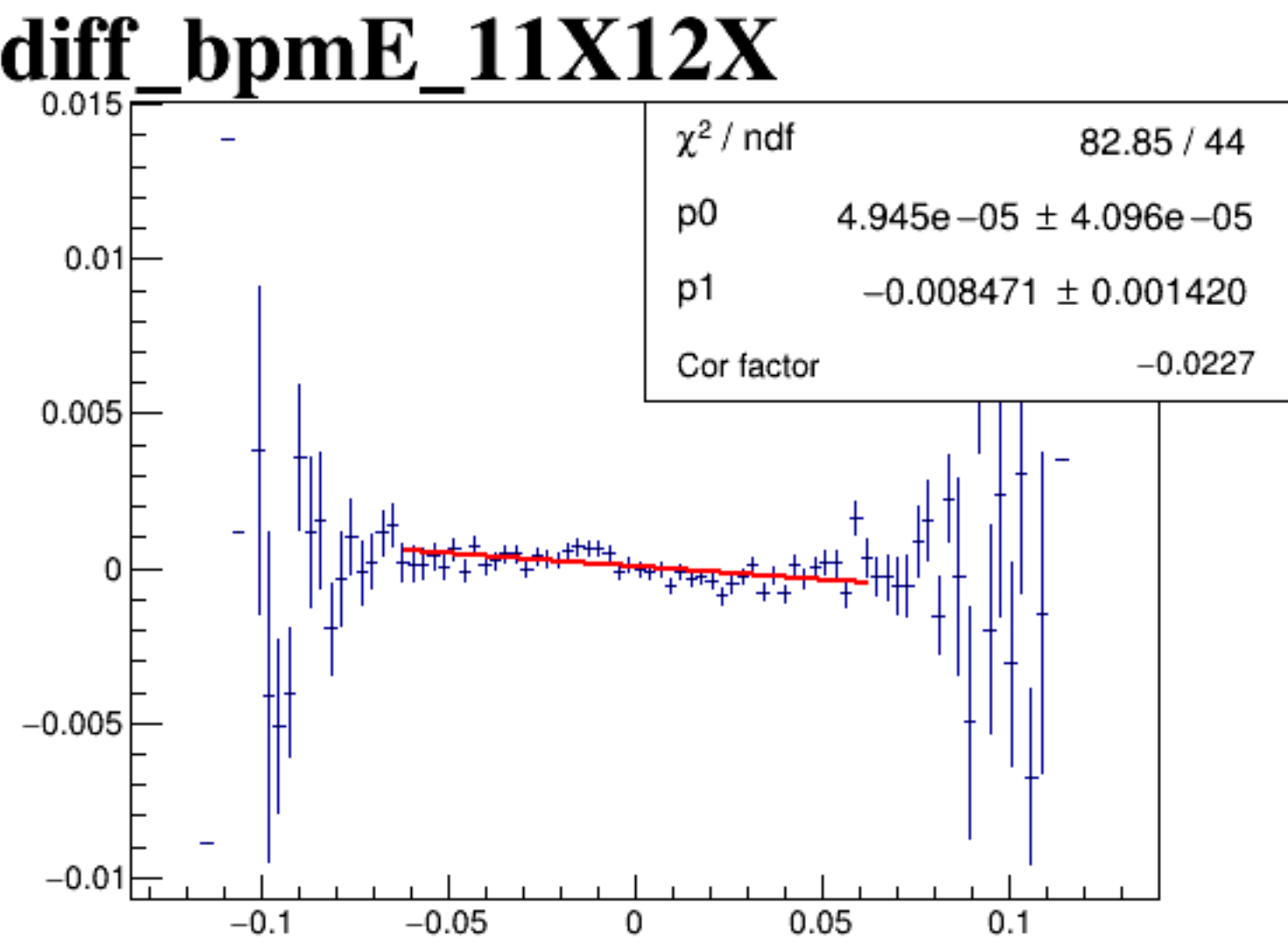
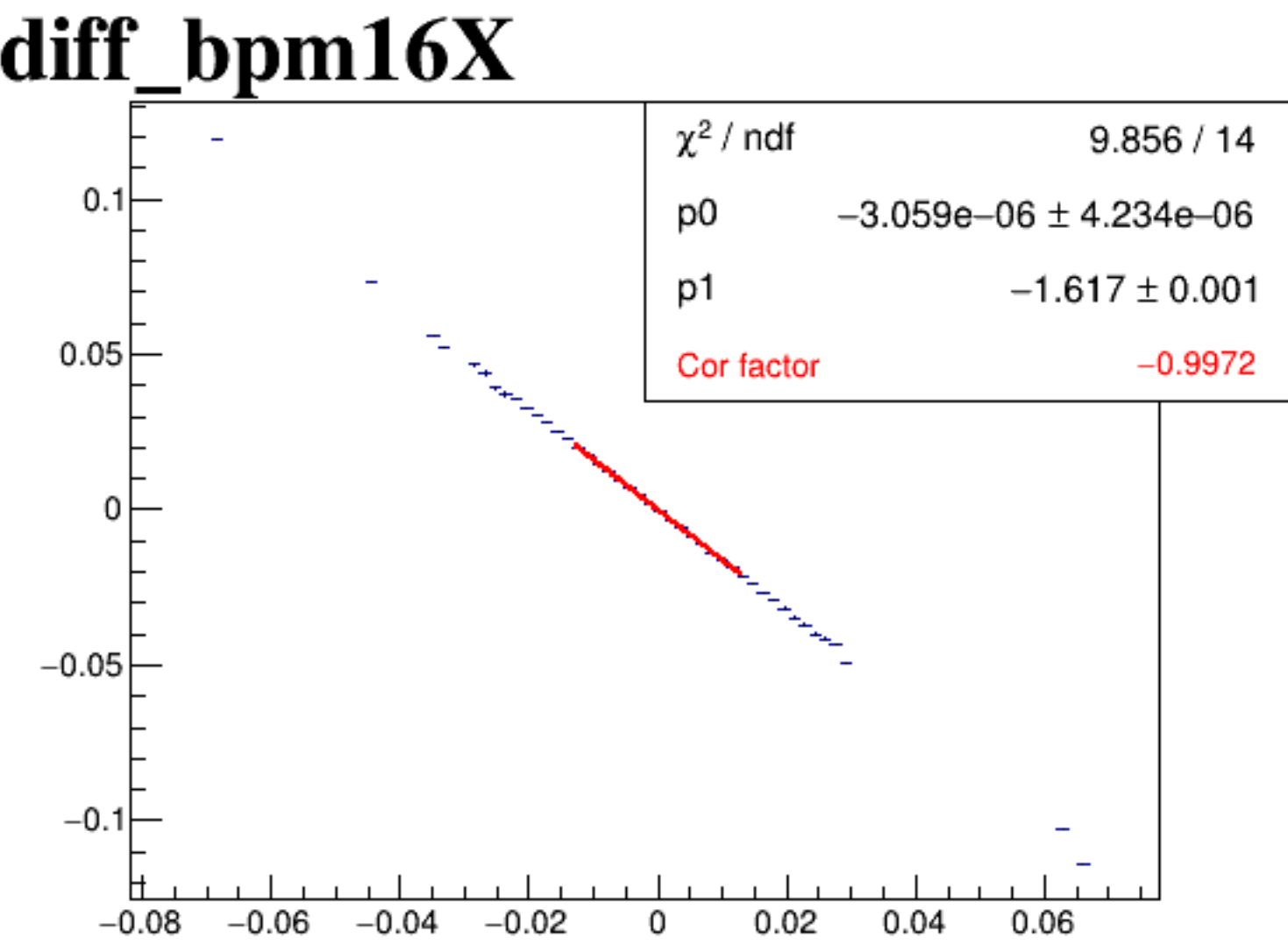
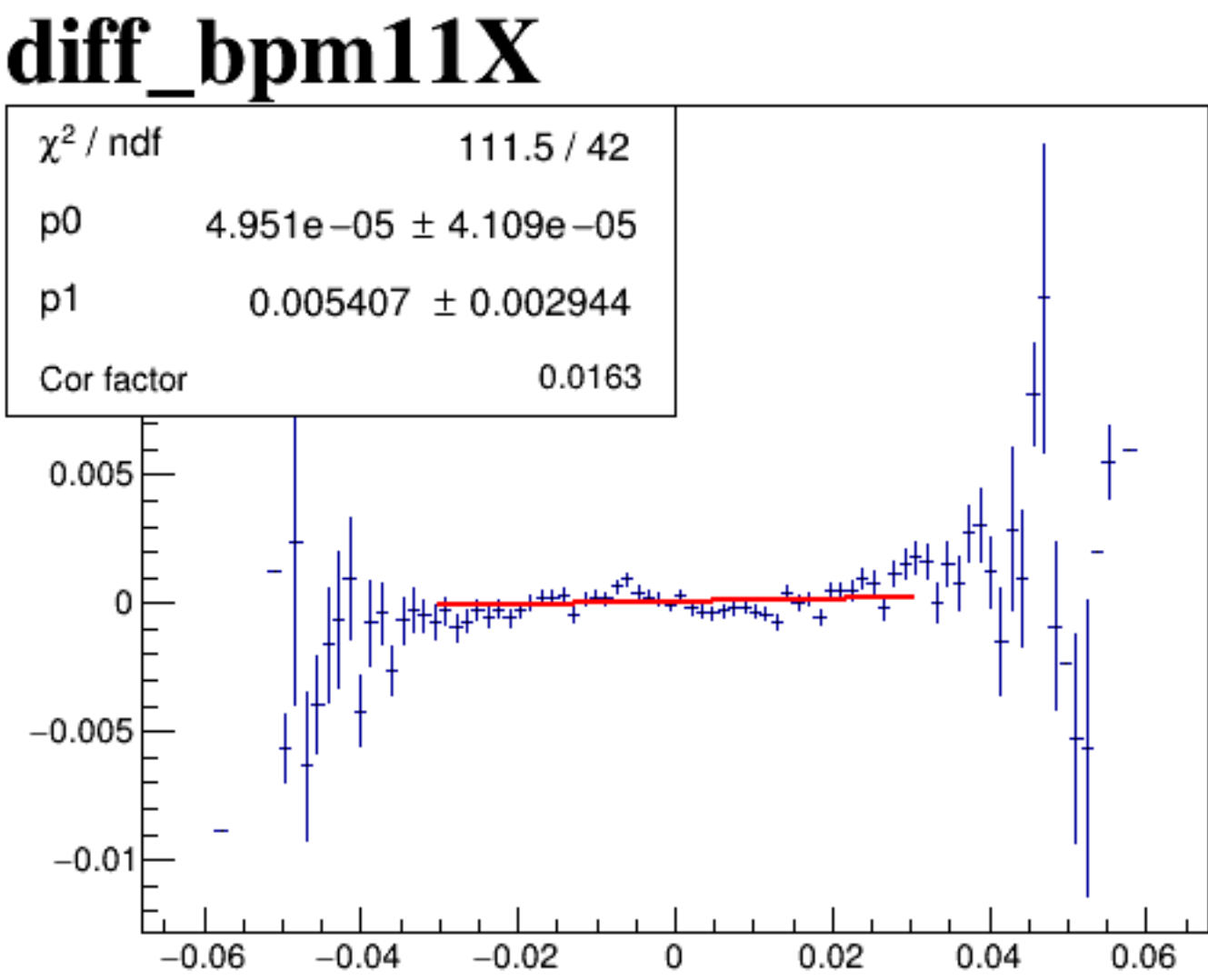
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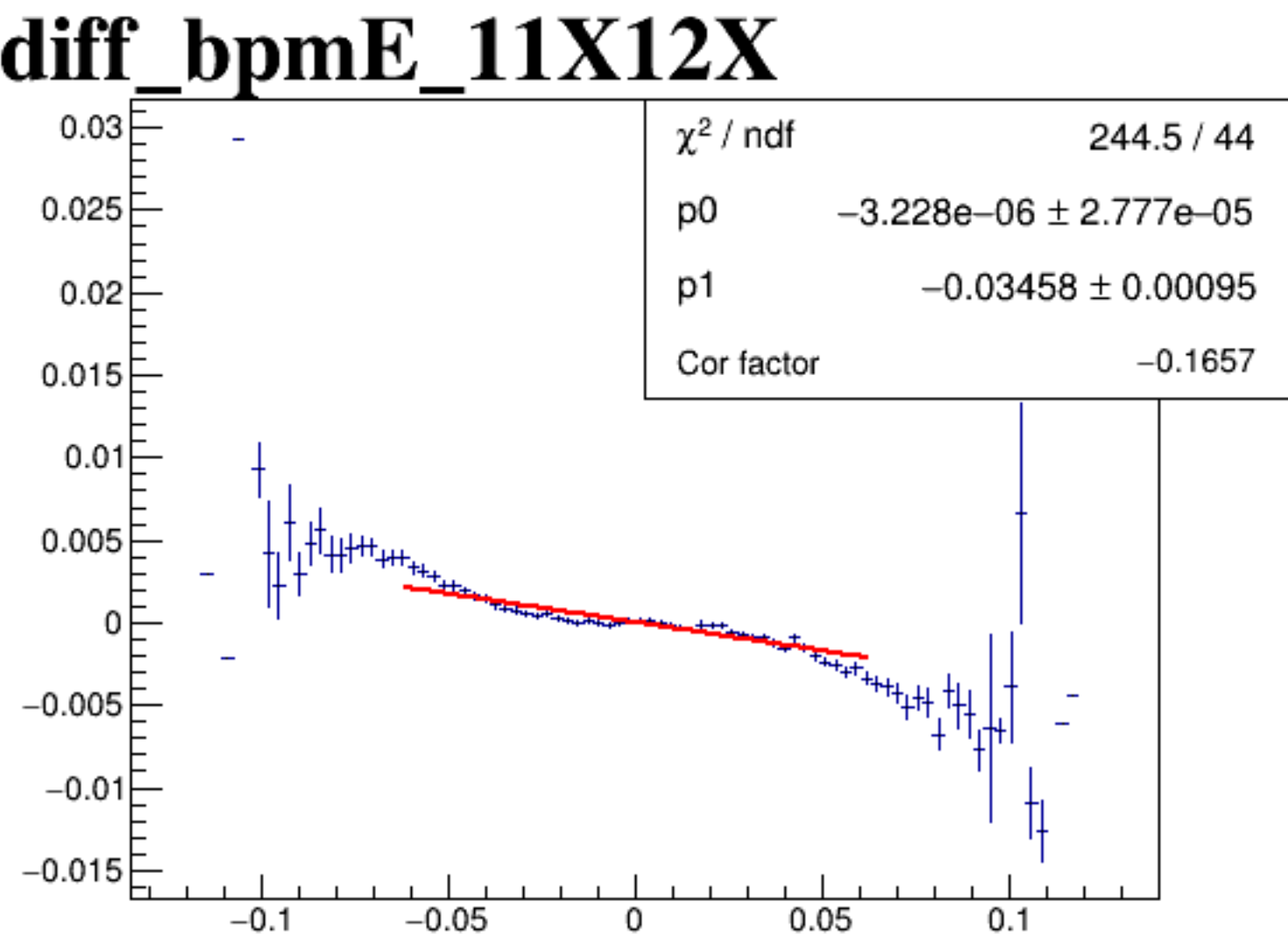
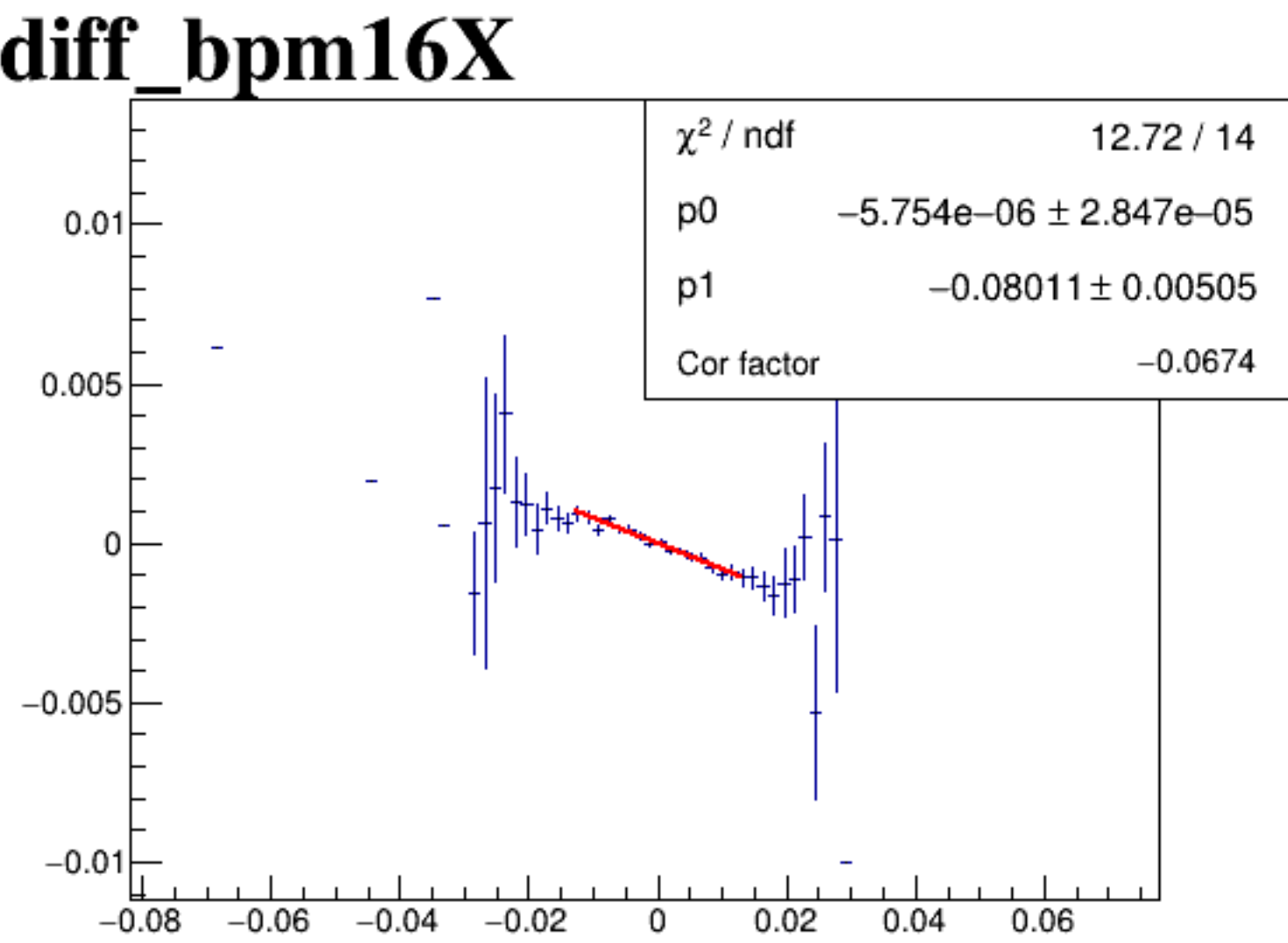
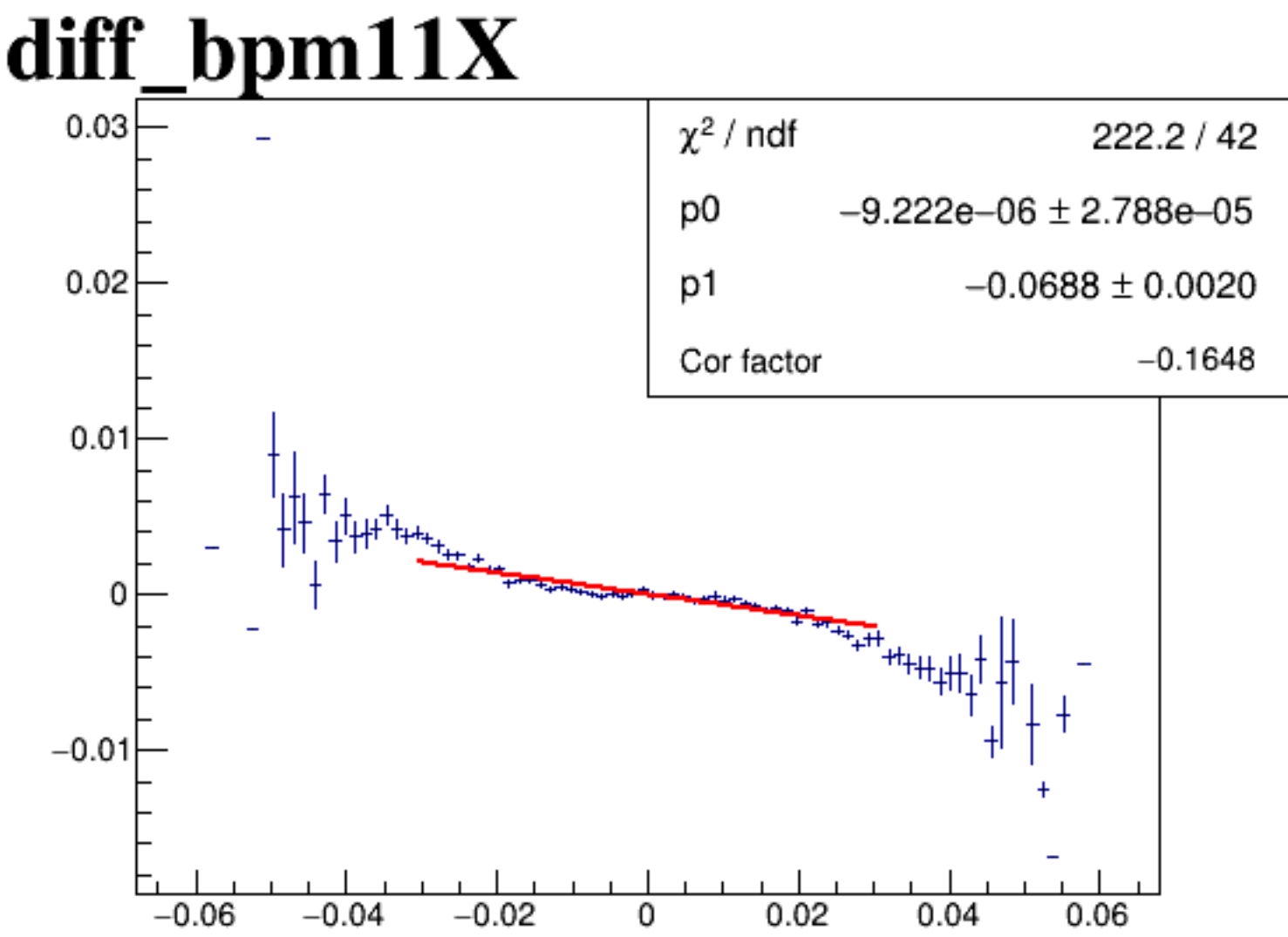
A scatter plot titled "diff_bpm1Y". The x-axis is labeled "diff_bpm1Y" and ranges from -0.03 to 0.03 with major ticks every 0.01. The y-axis ranges from -0.1 to 0.1 with major ticks every 0.05. The plot shows a dense, roughly elliptical cloud of points centered at (0,0), with the majority of points falling within the range [-0.01, 0.01] on the x-axis and [-0.05, 0.05] on the y-axis.

A scatter plot showing the relationship between two variables, both labeled `diff_bpm1Y` on the axes. The plot displays a dense, elongated cloud of points, indicating a strong positive correlation. The axes range from -0.03 to 0.03, with major ticks every 0.01 units. The data points are most concentrated along the diagonal line where the two variables are equal.

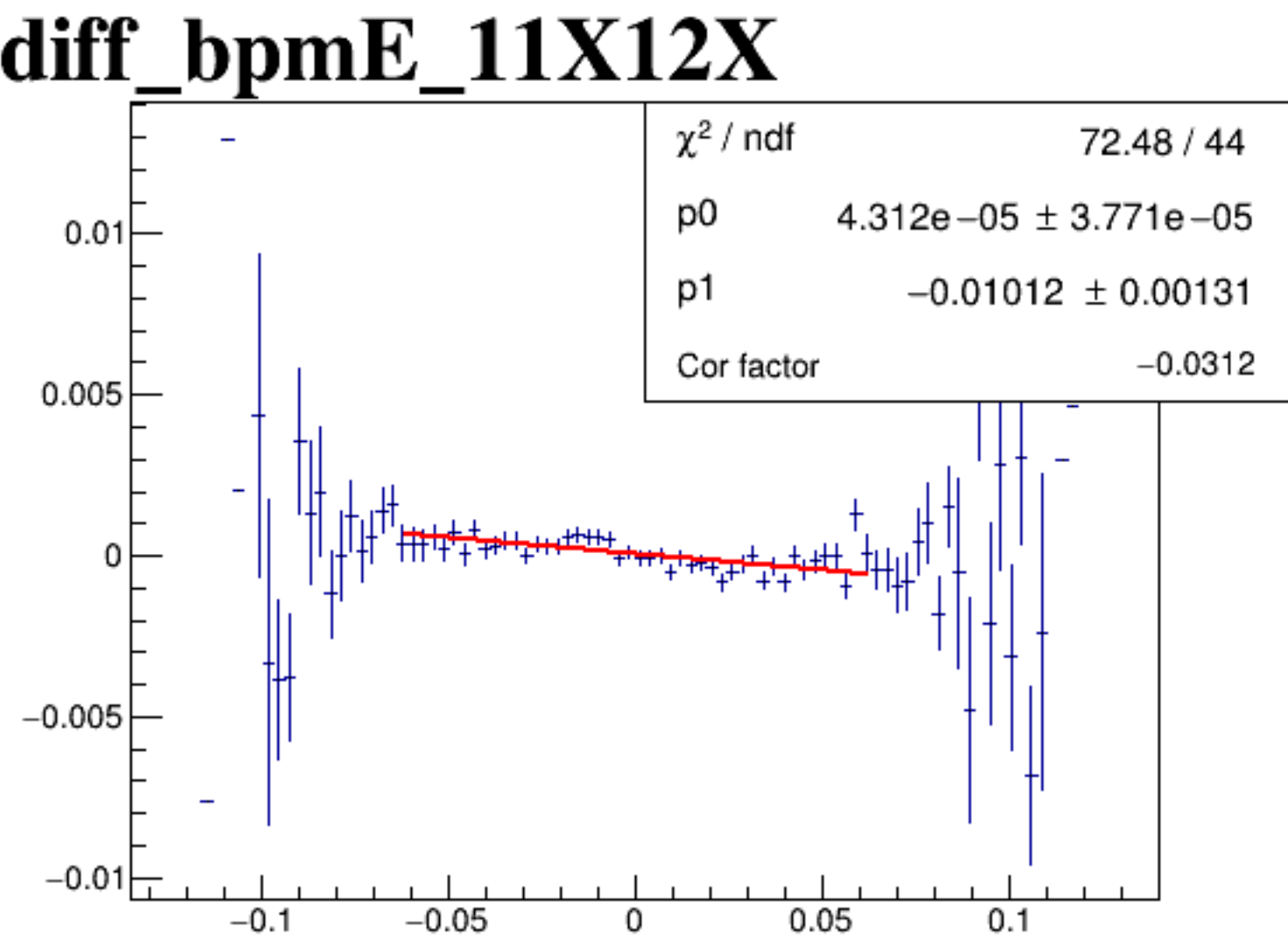
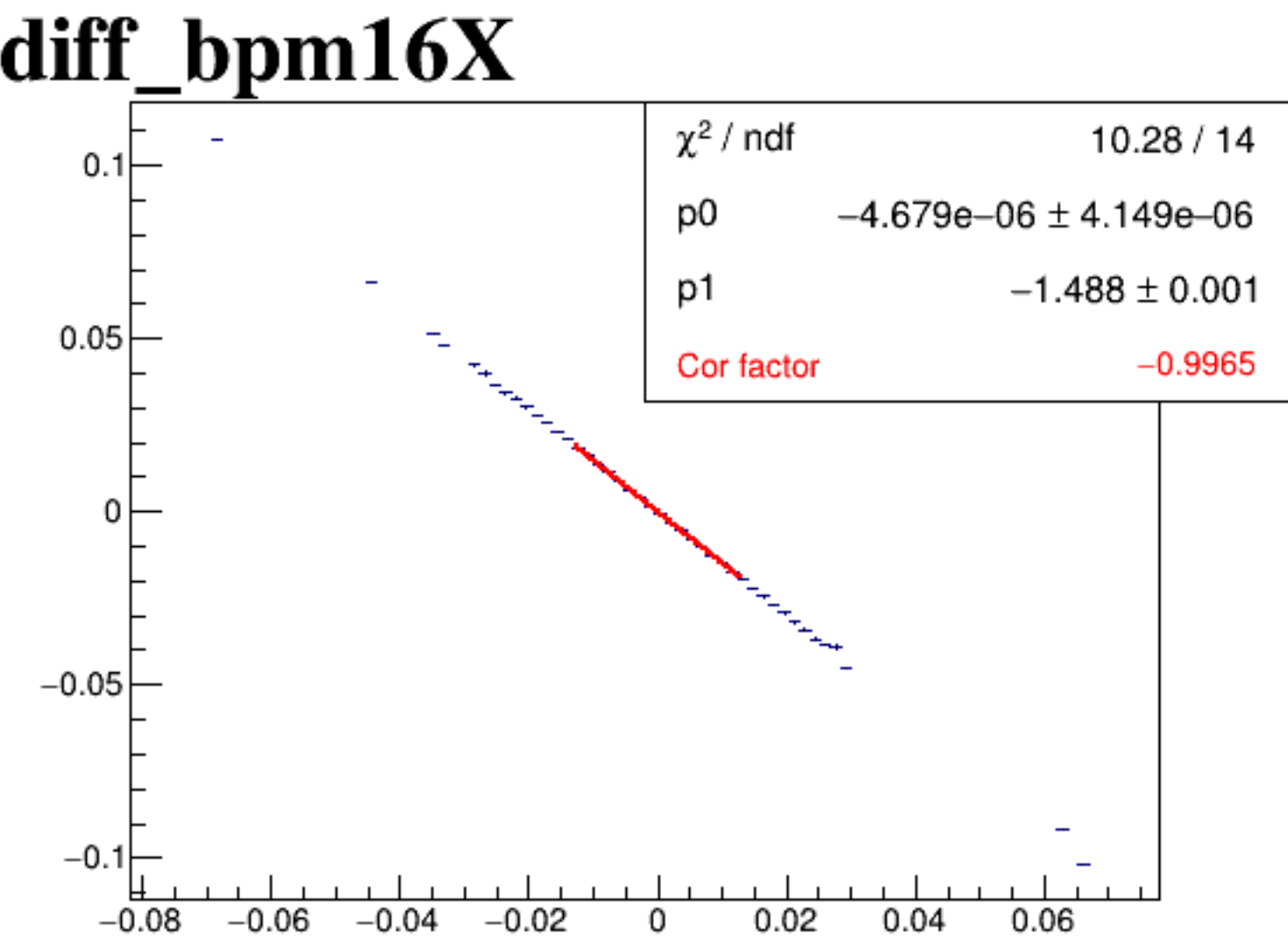
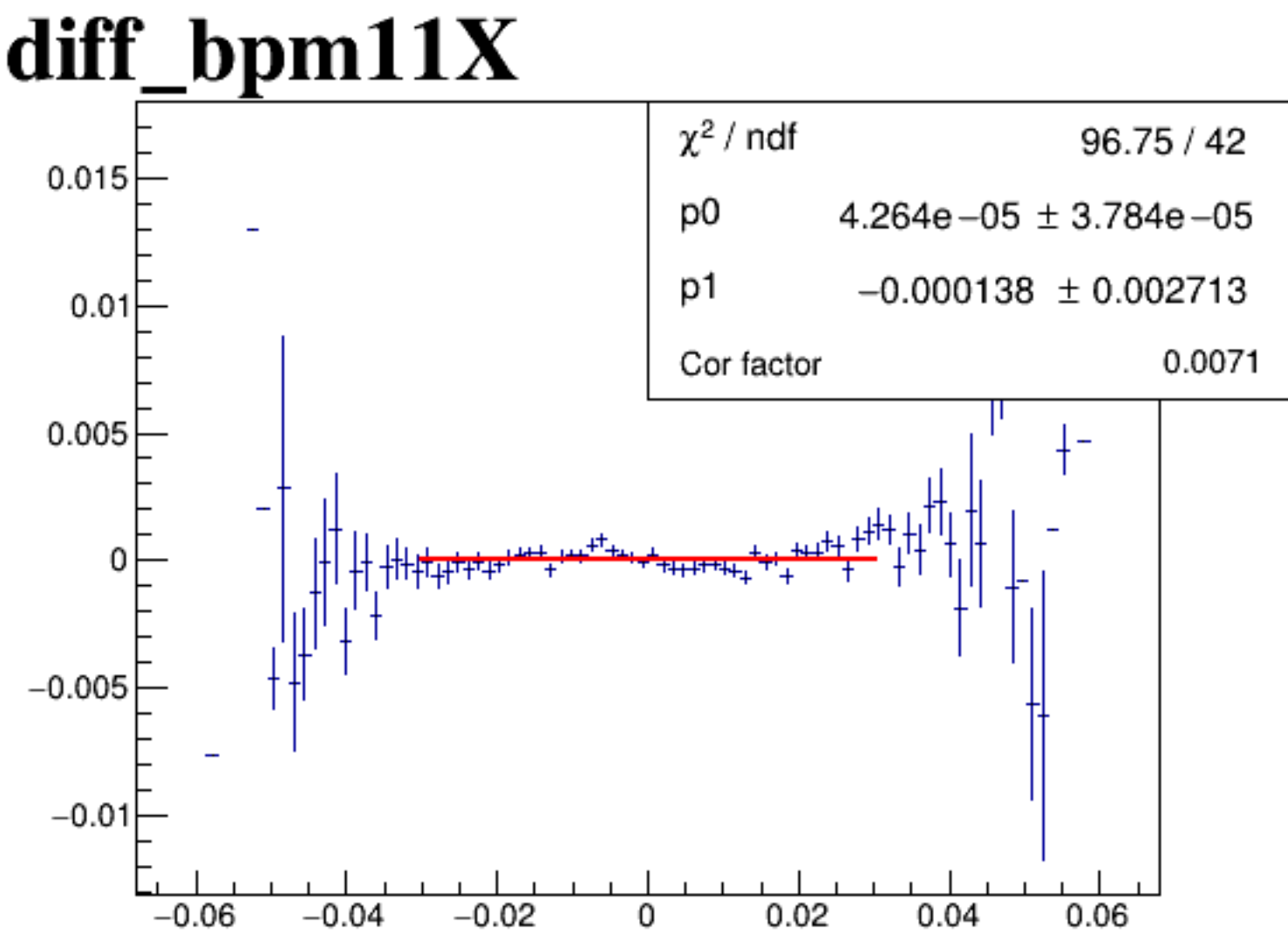
diff_bpm4aX



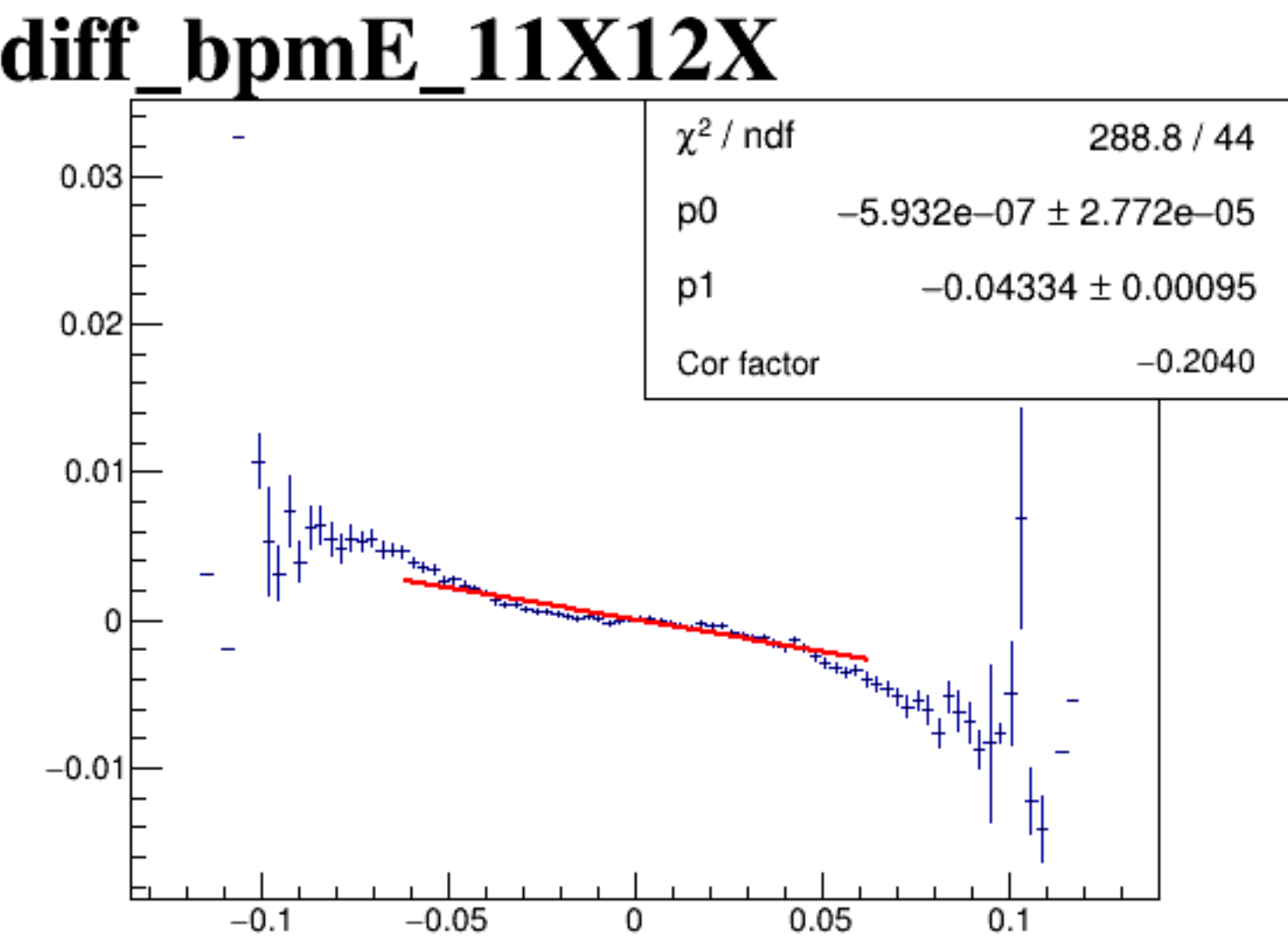
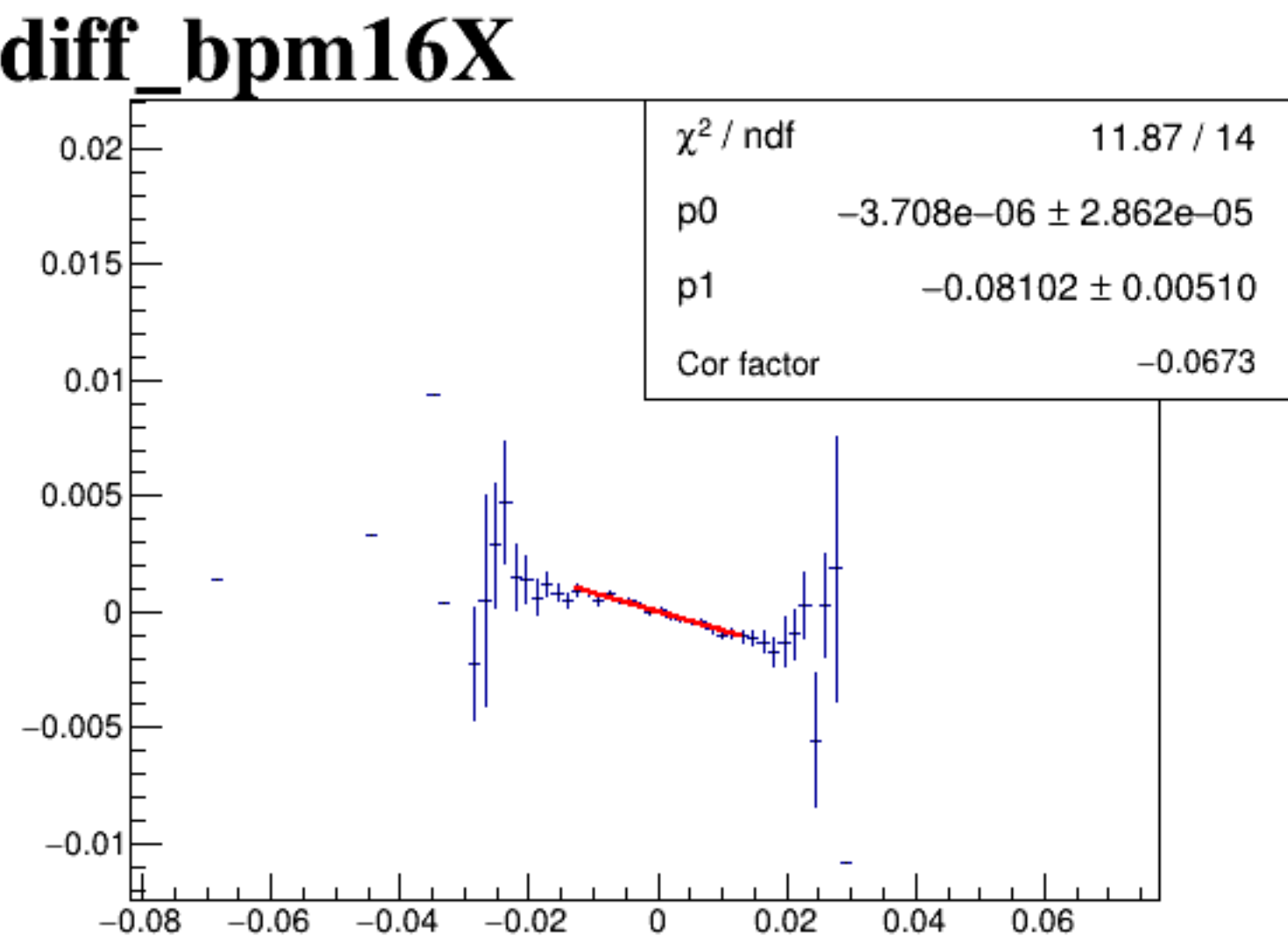
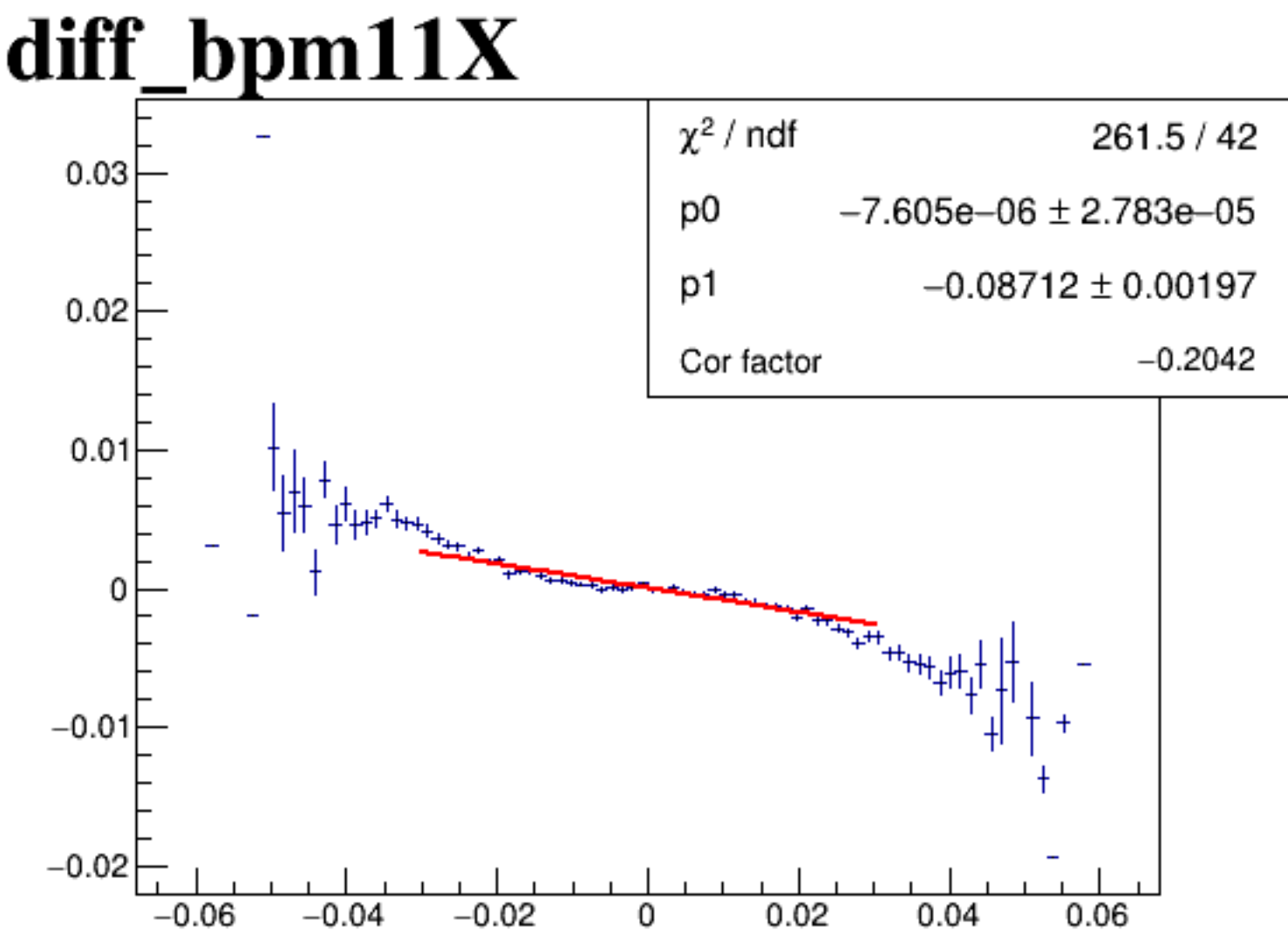
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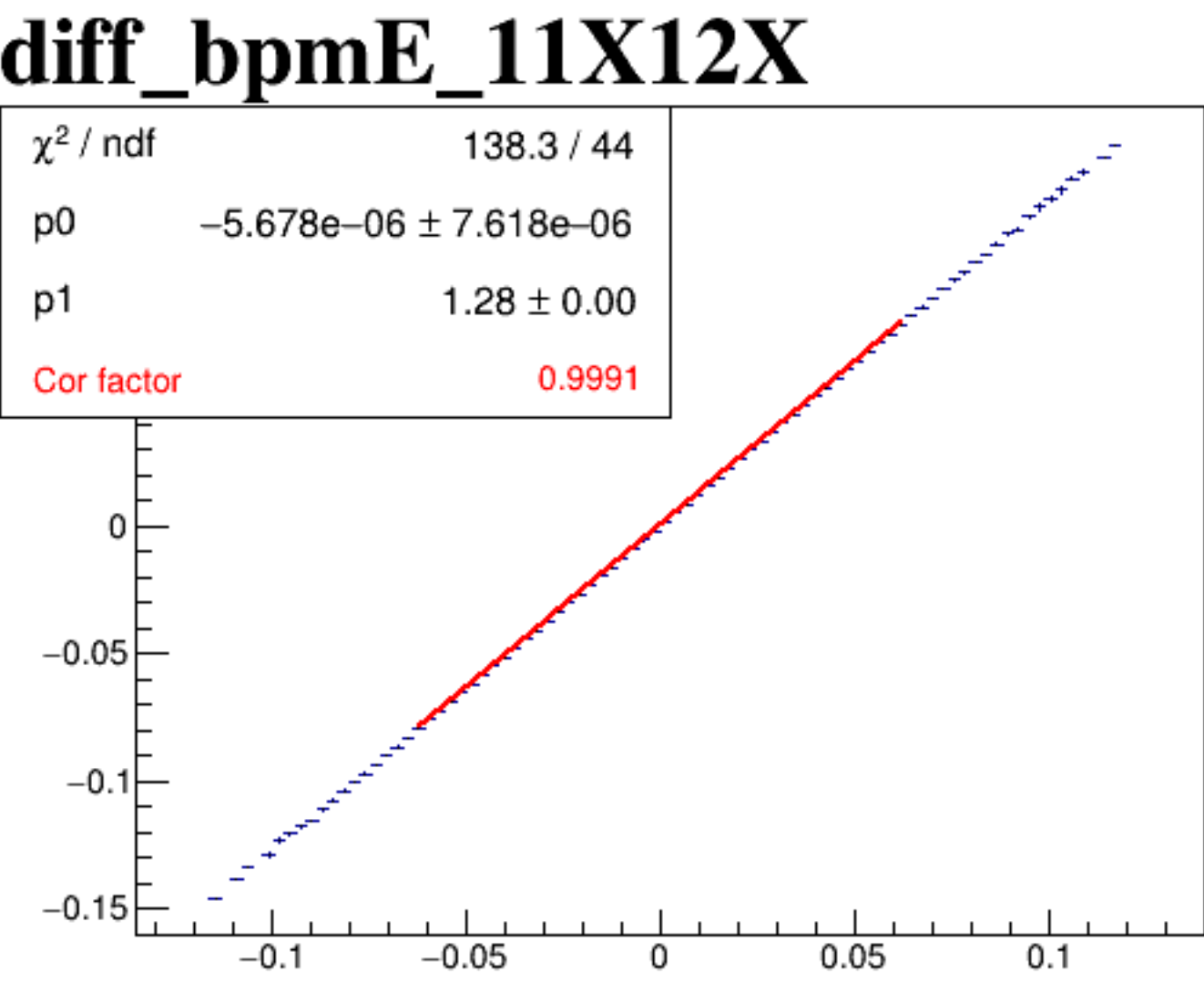
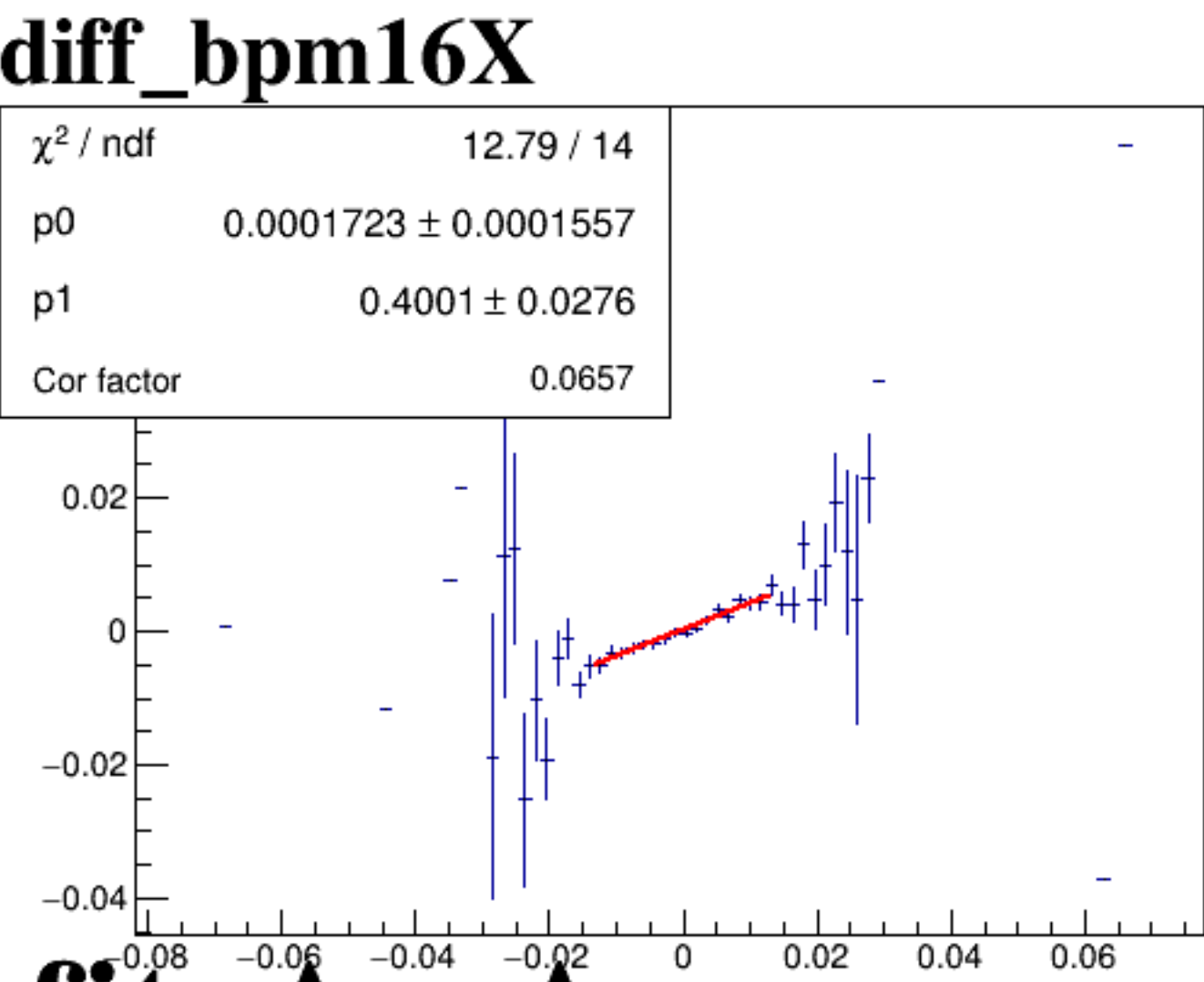
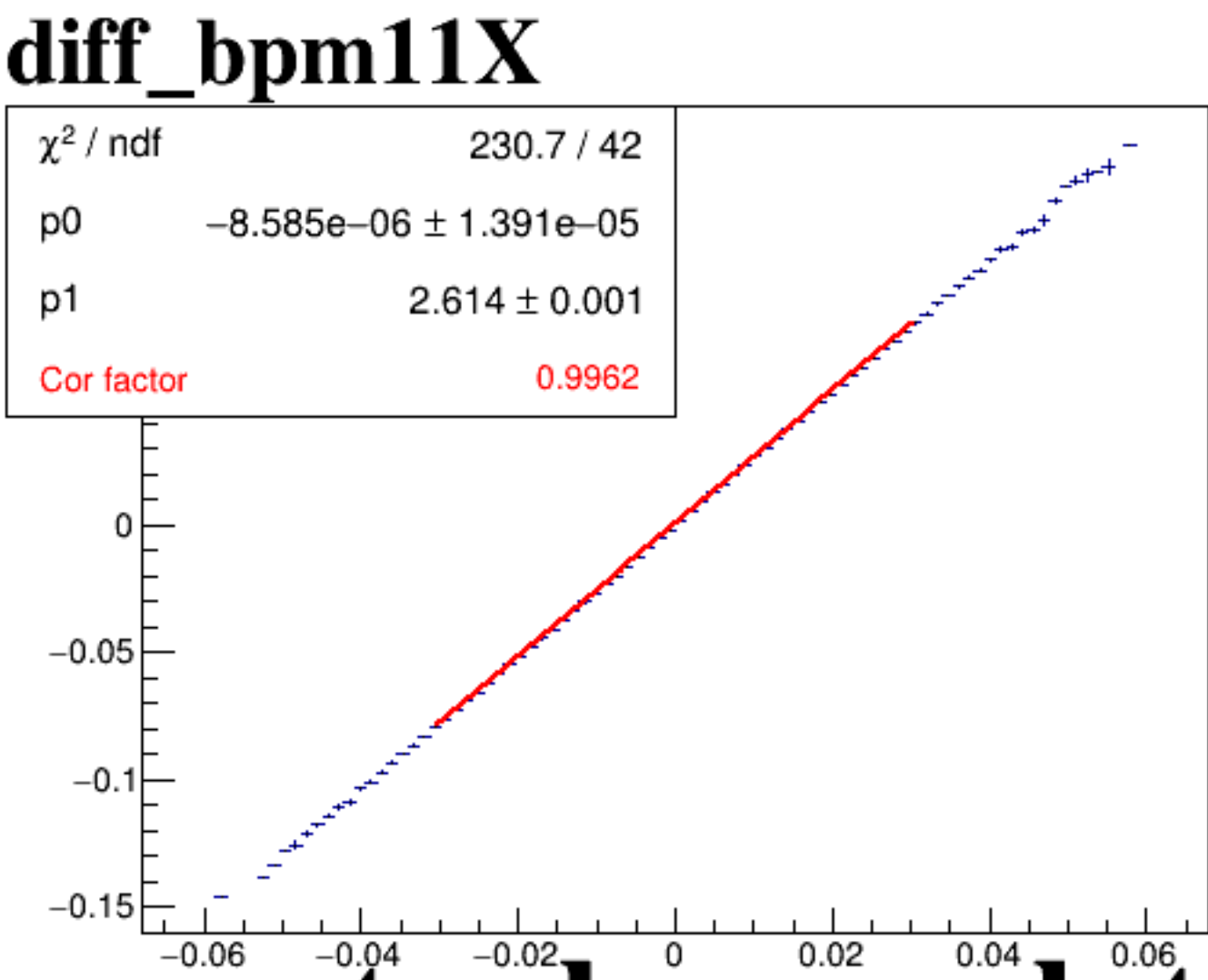
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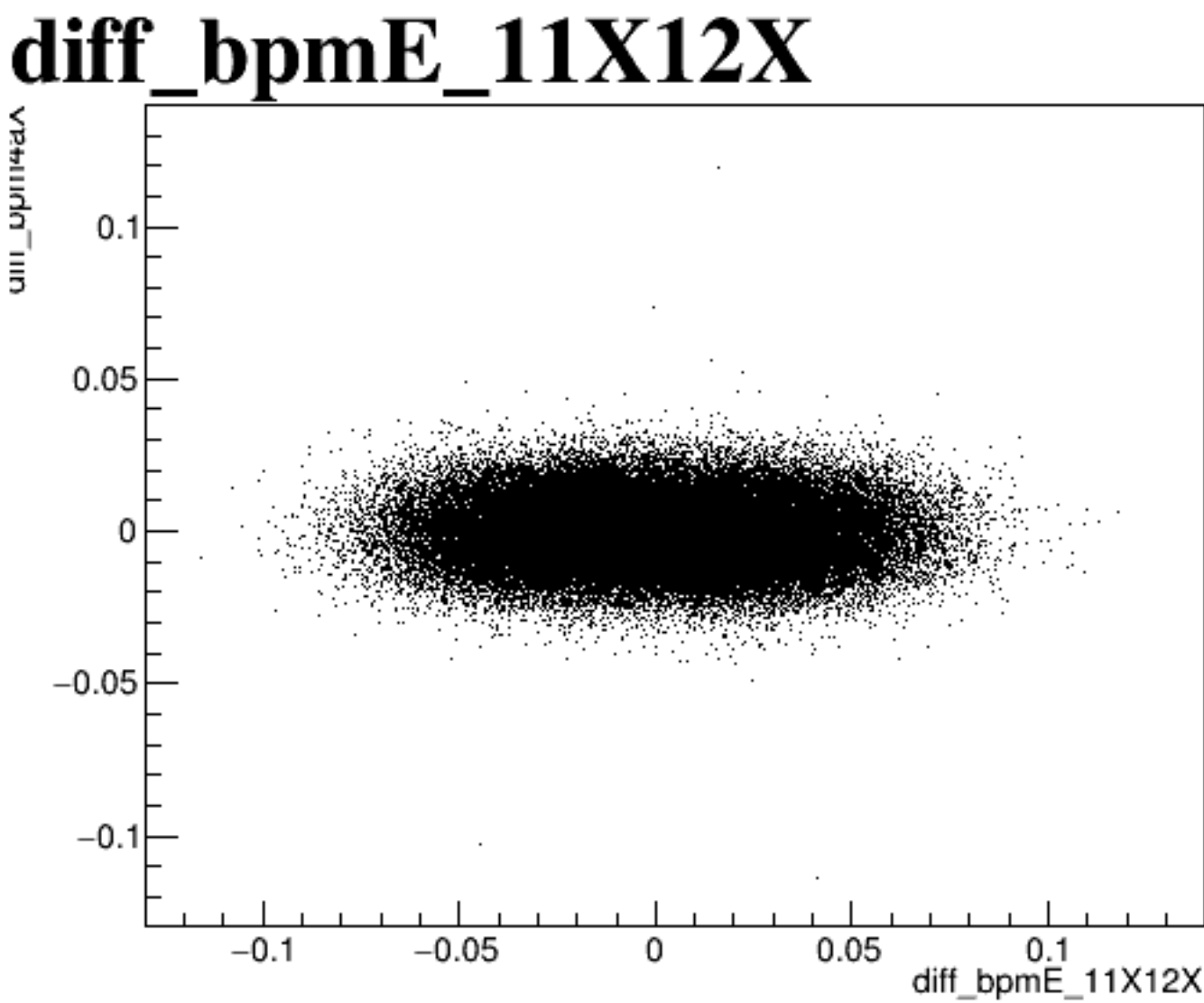
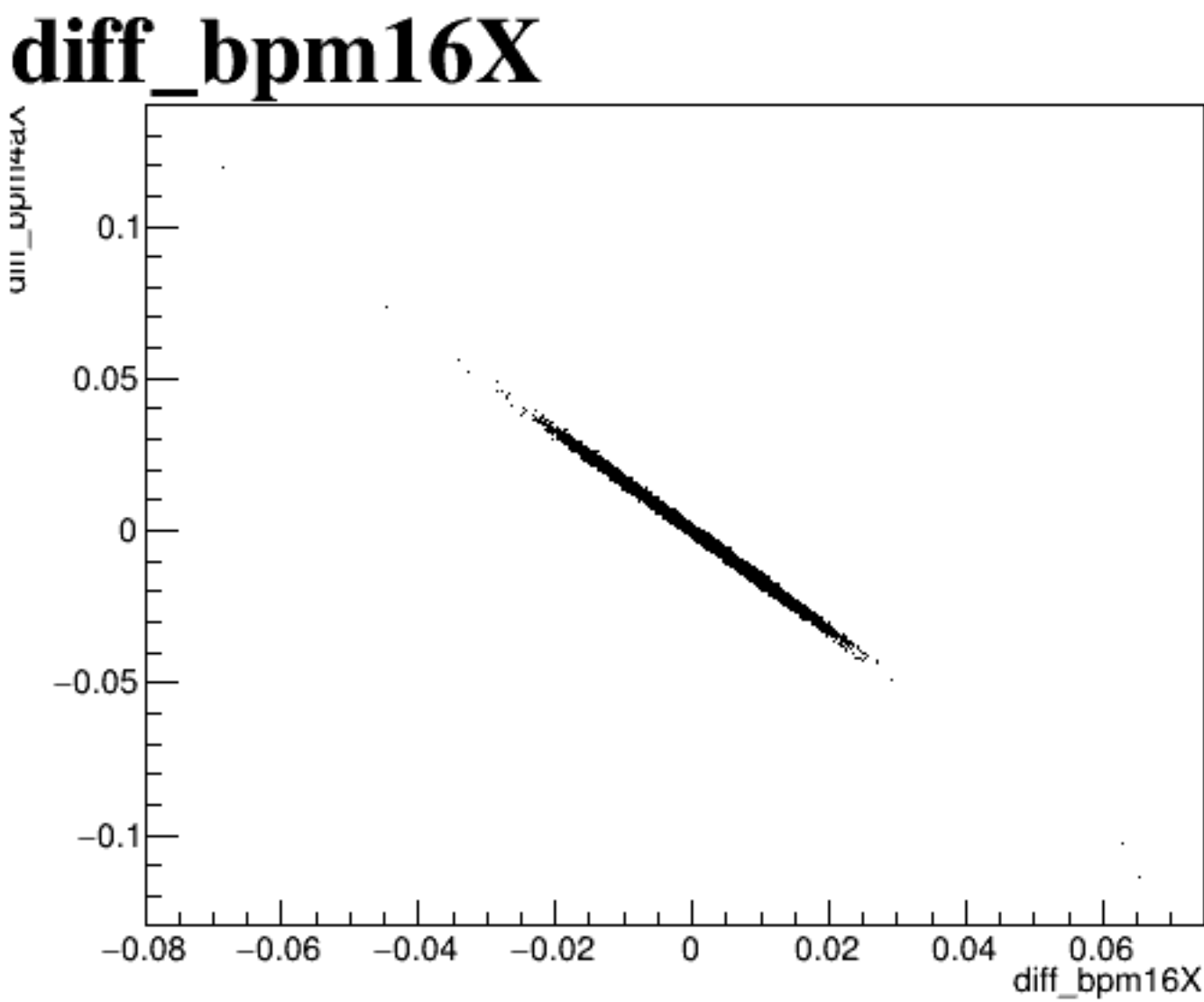
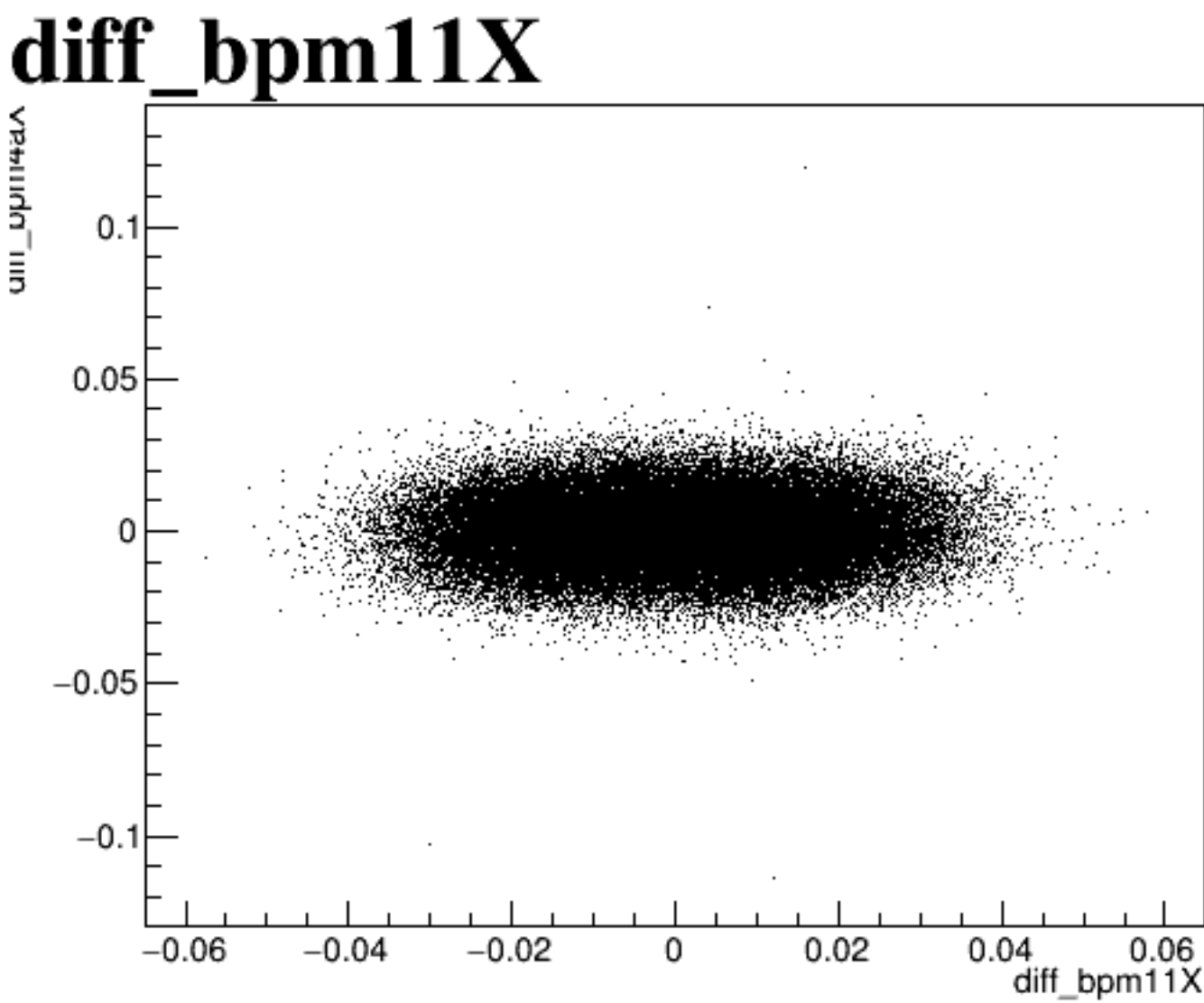
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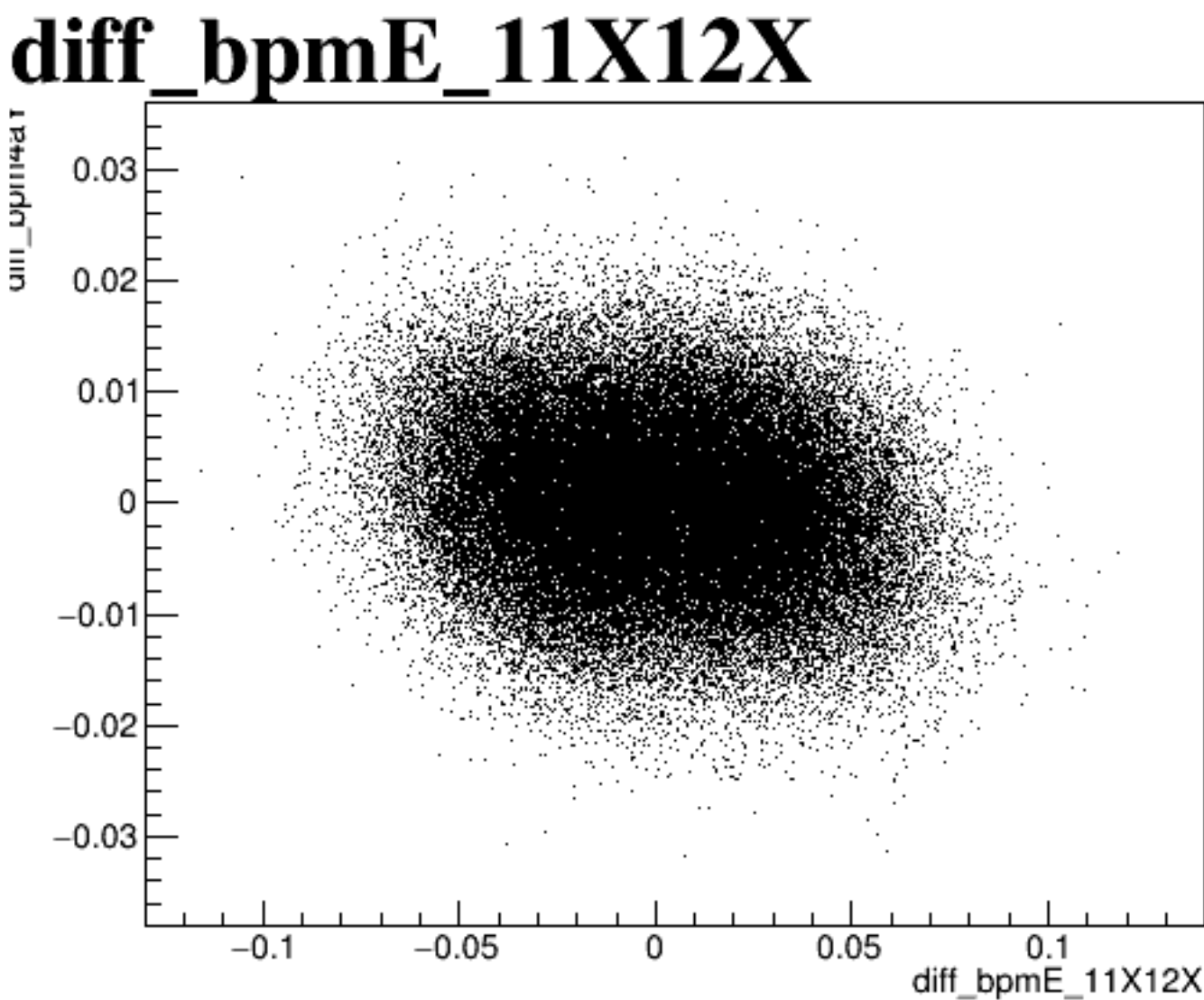
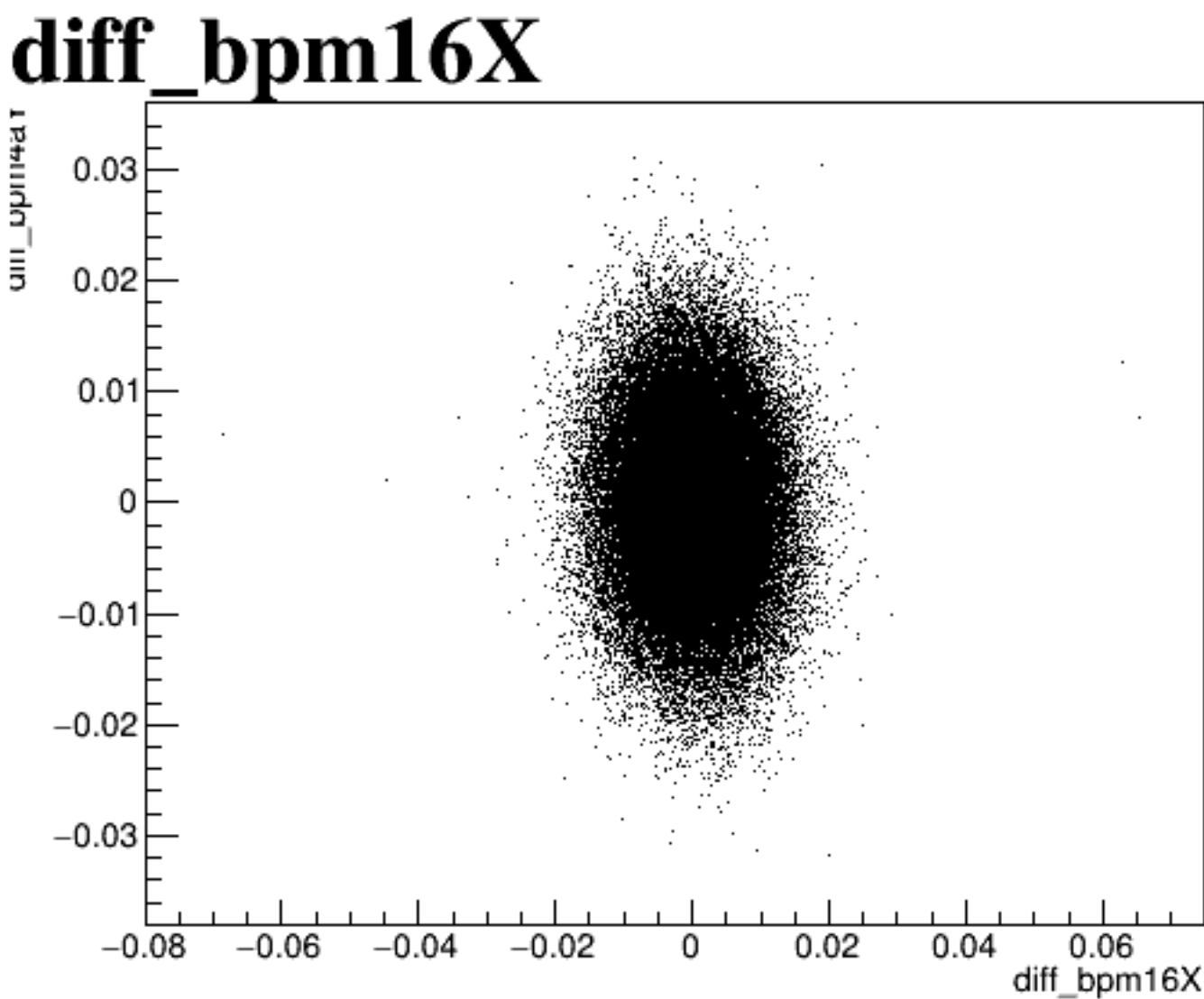
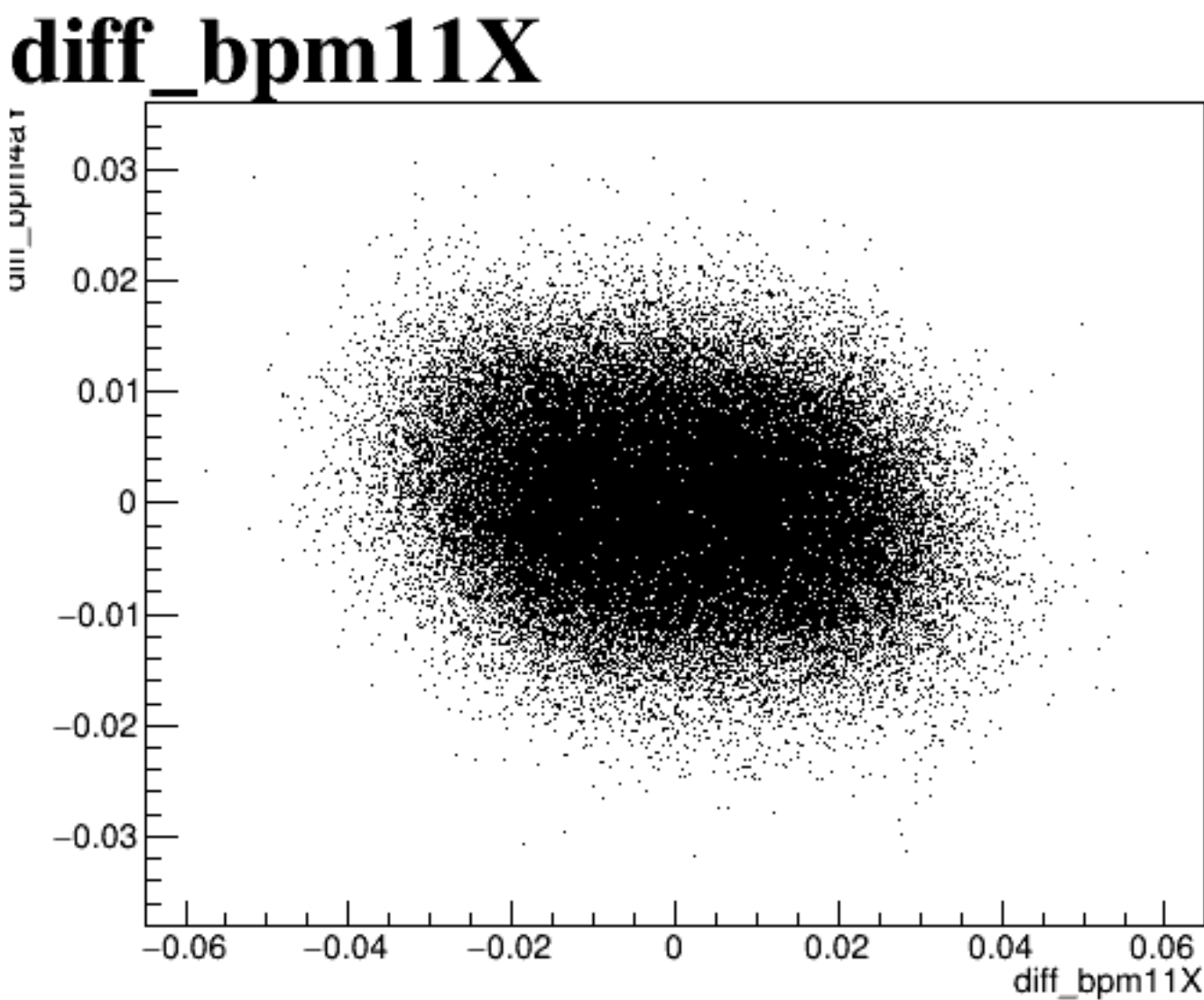
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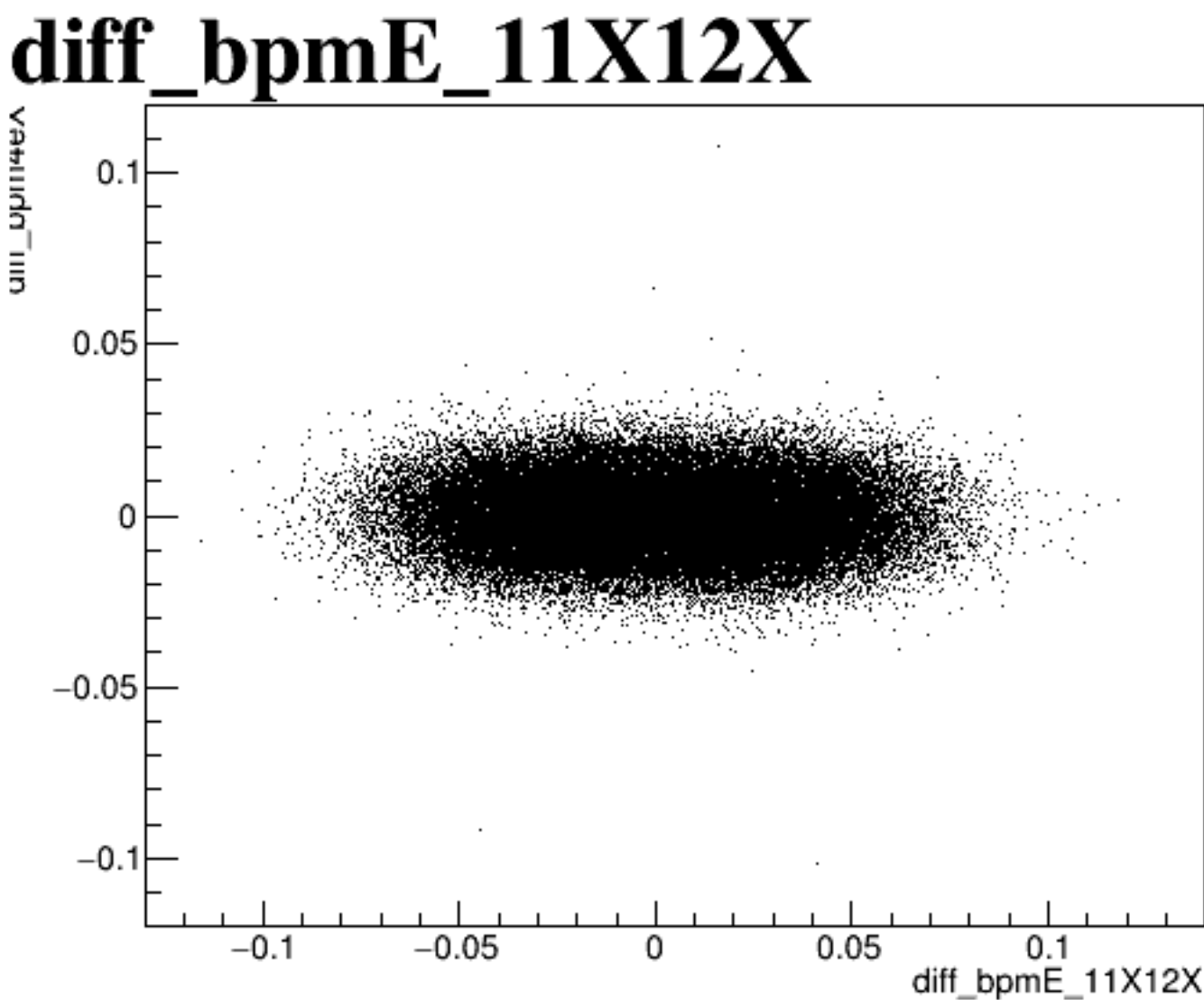
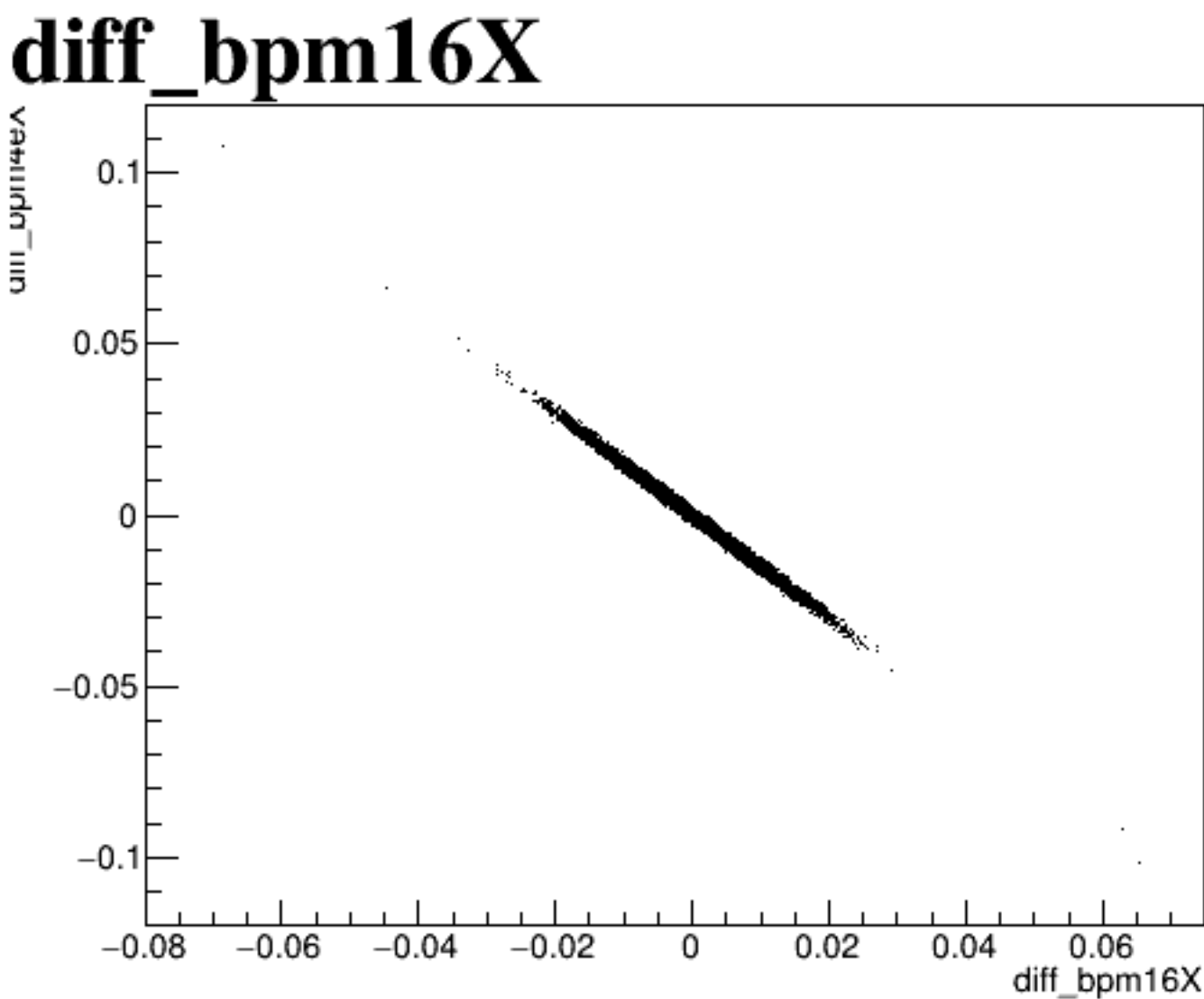
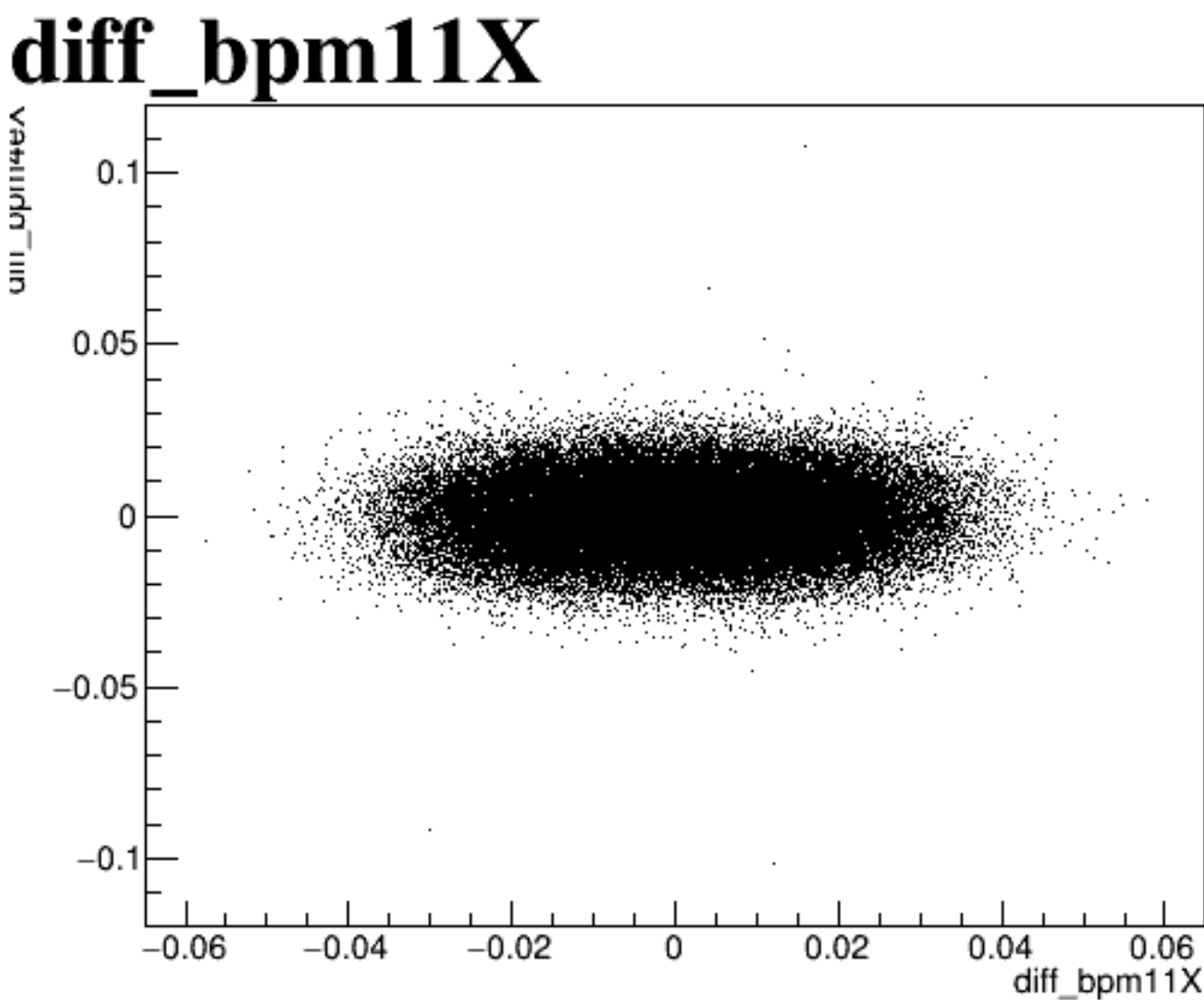
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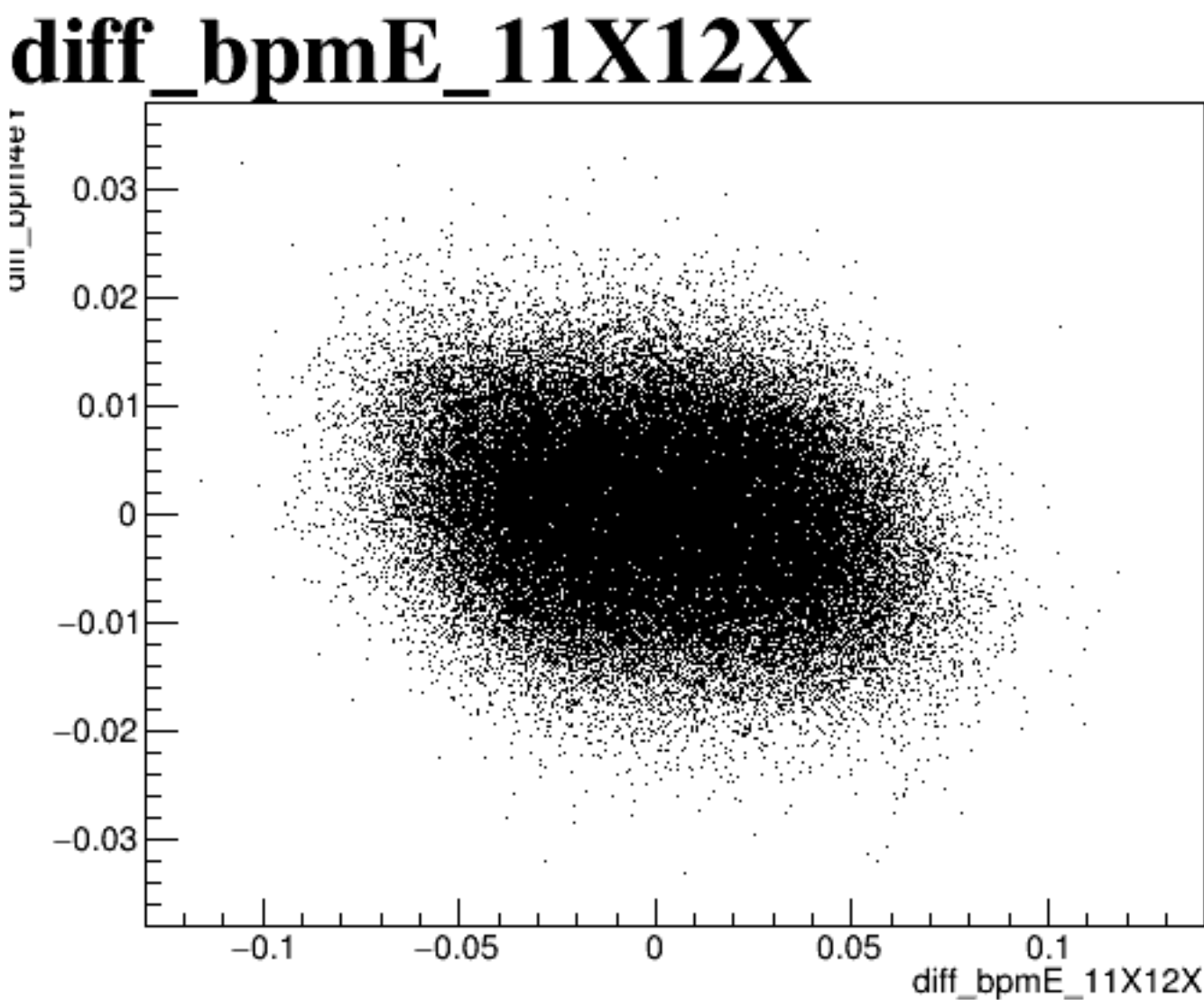
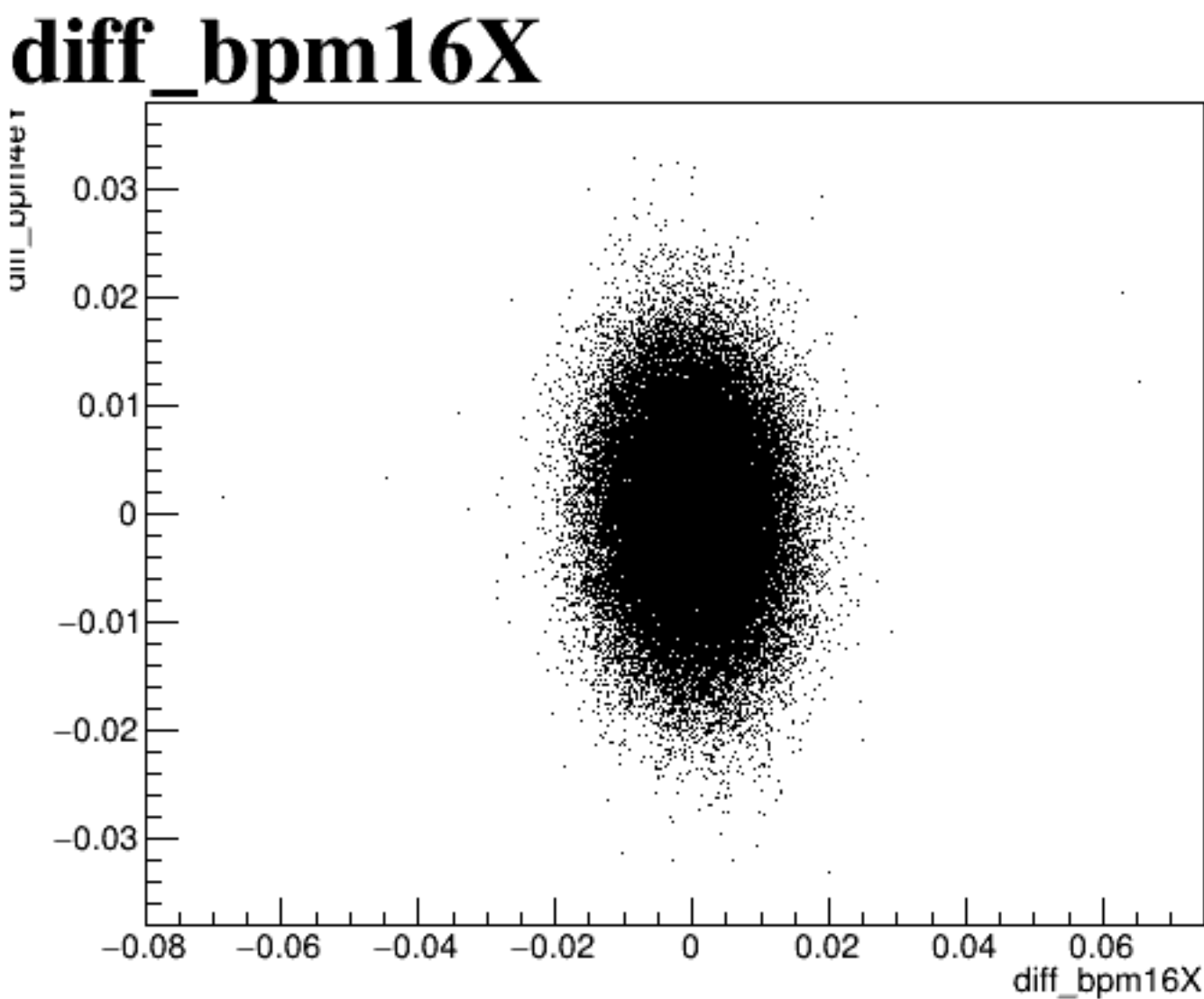
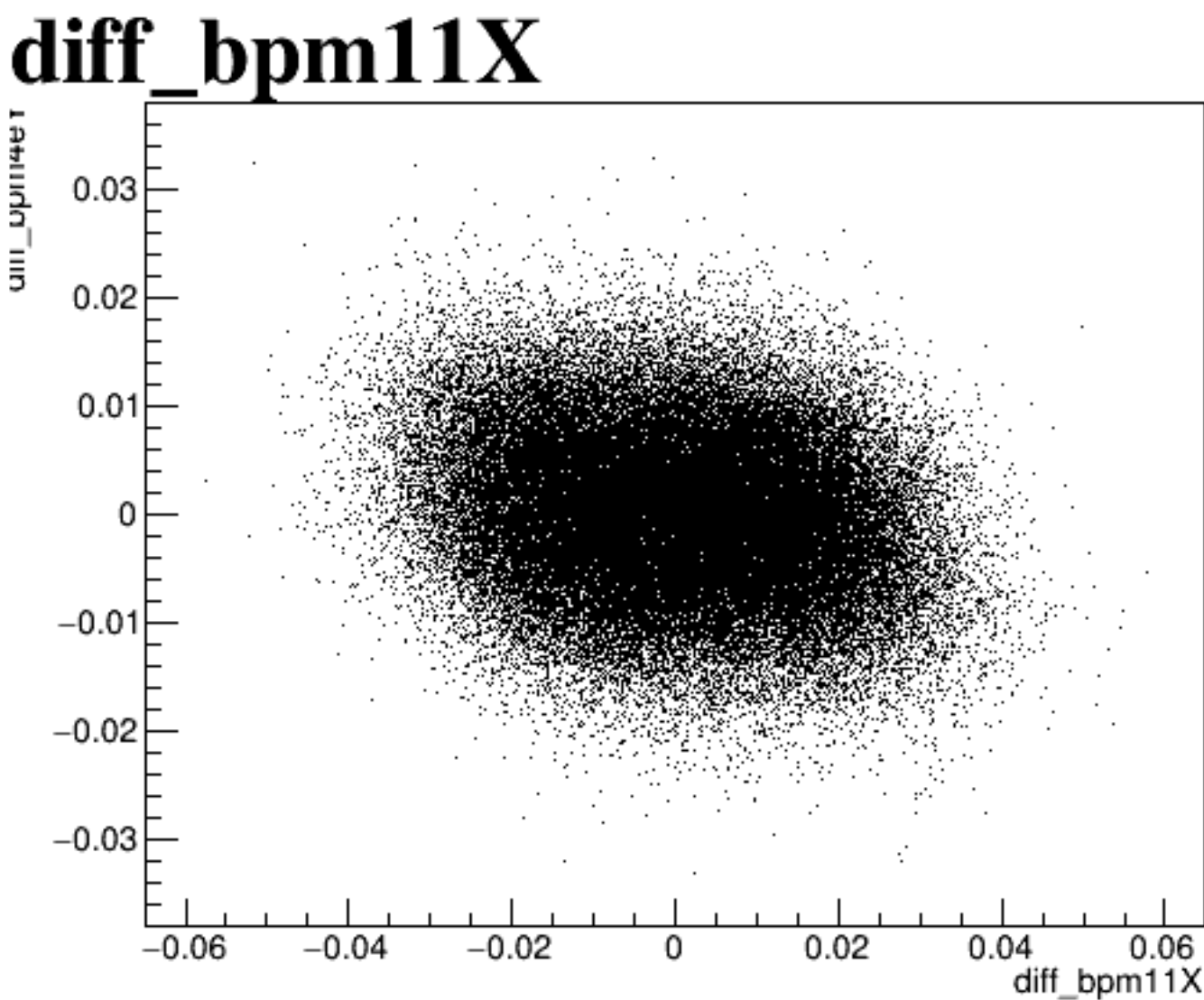
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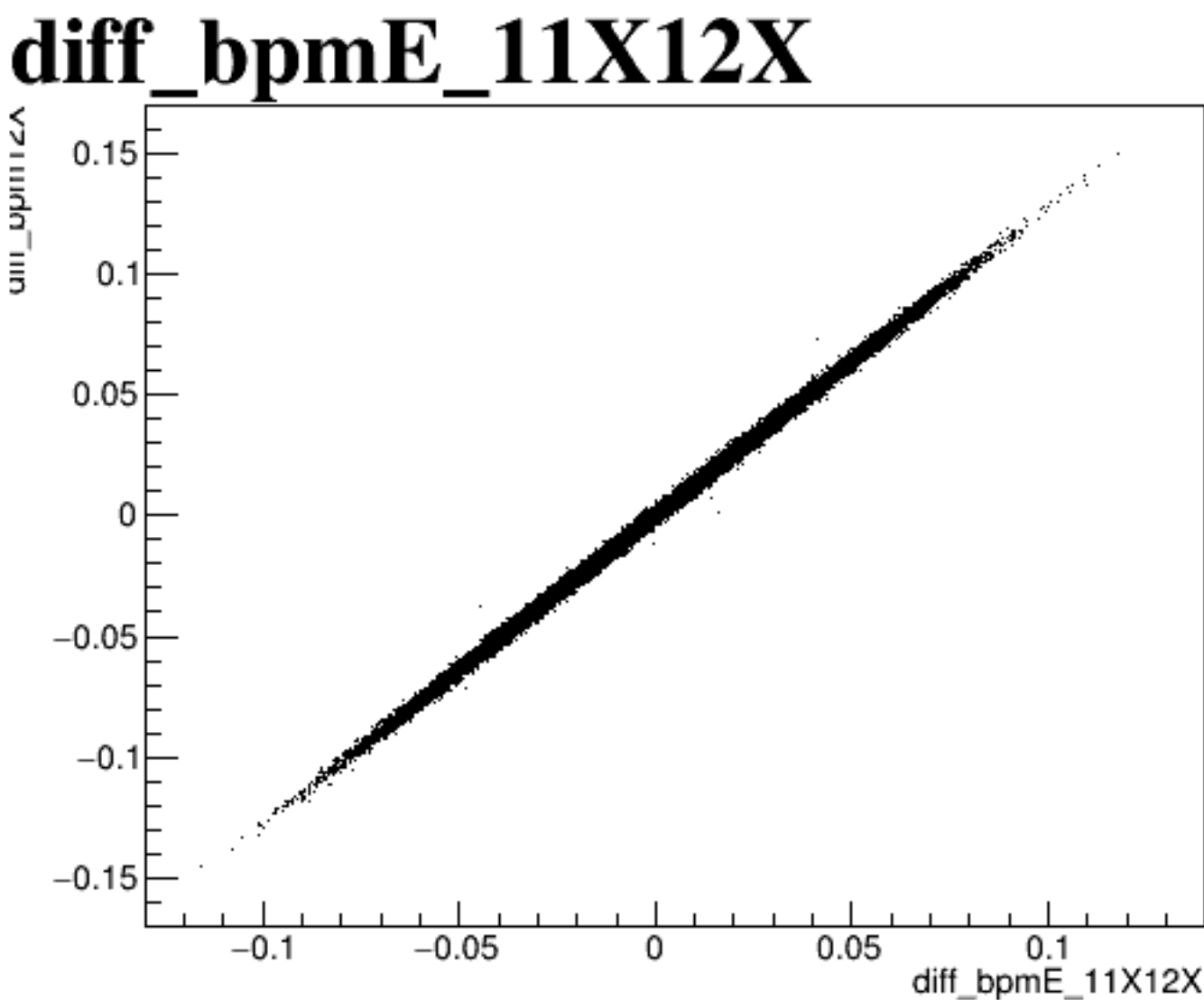
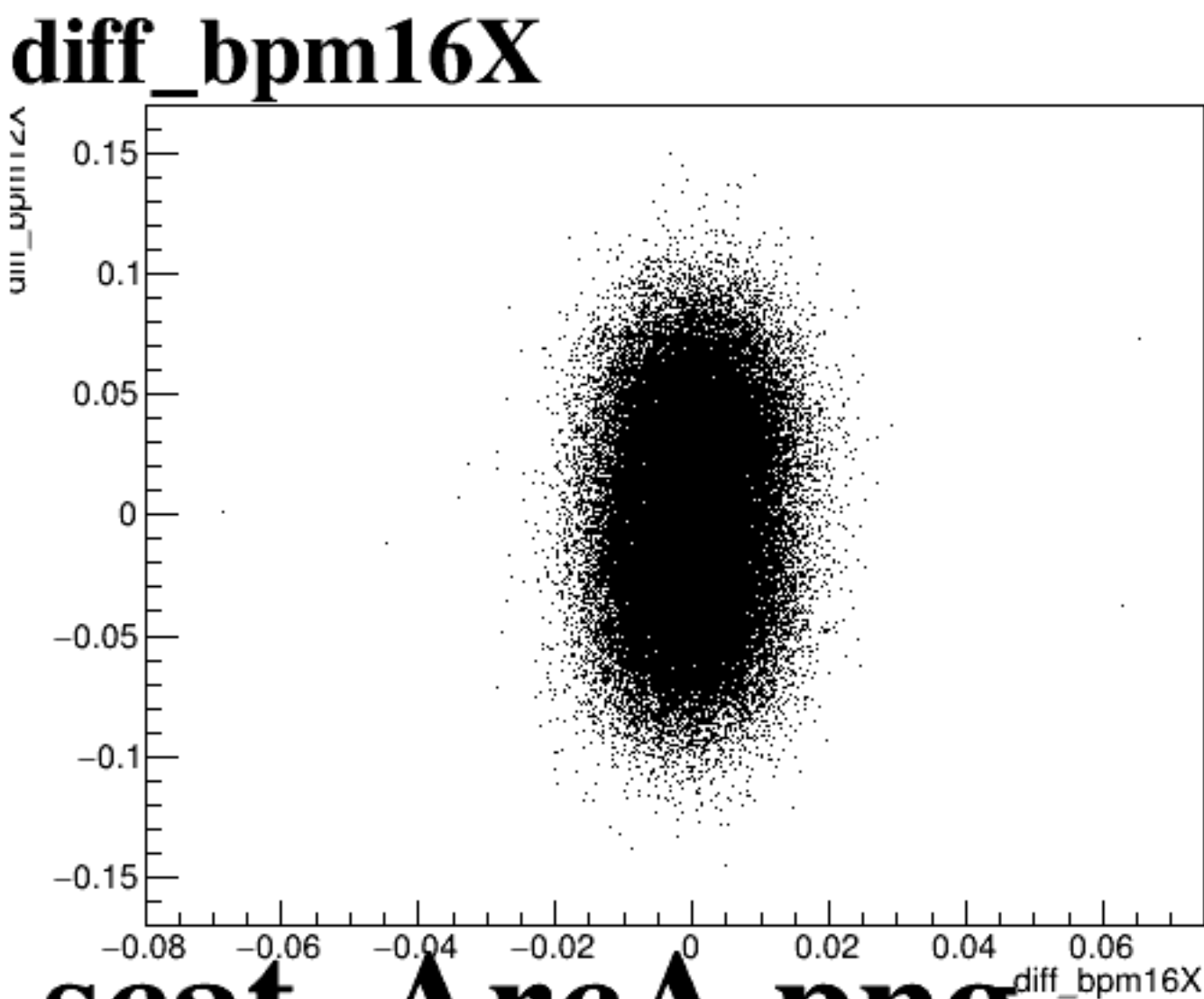
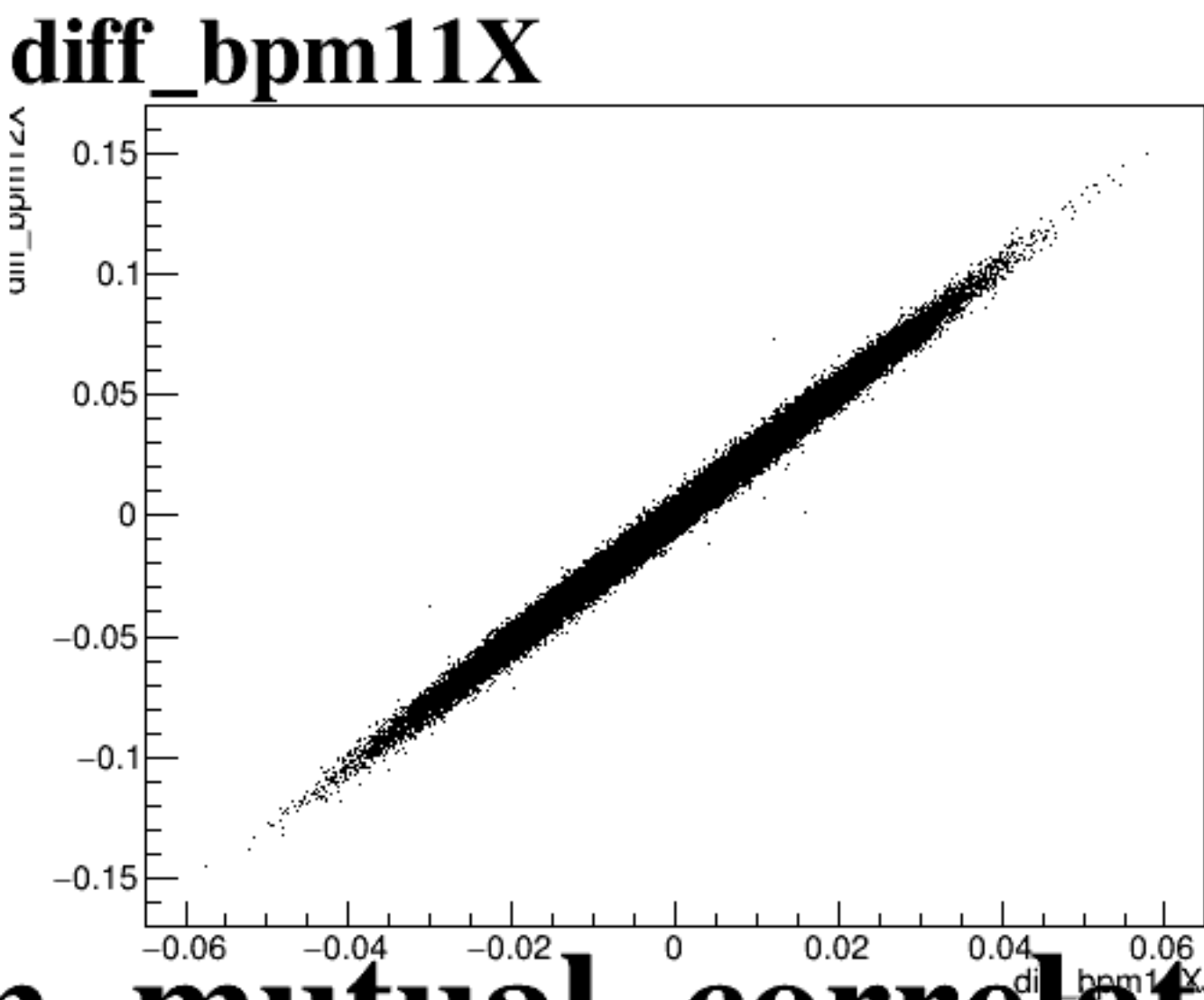
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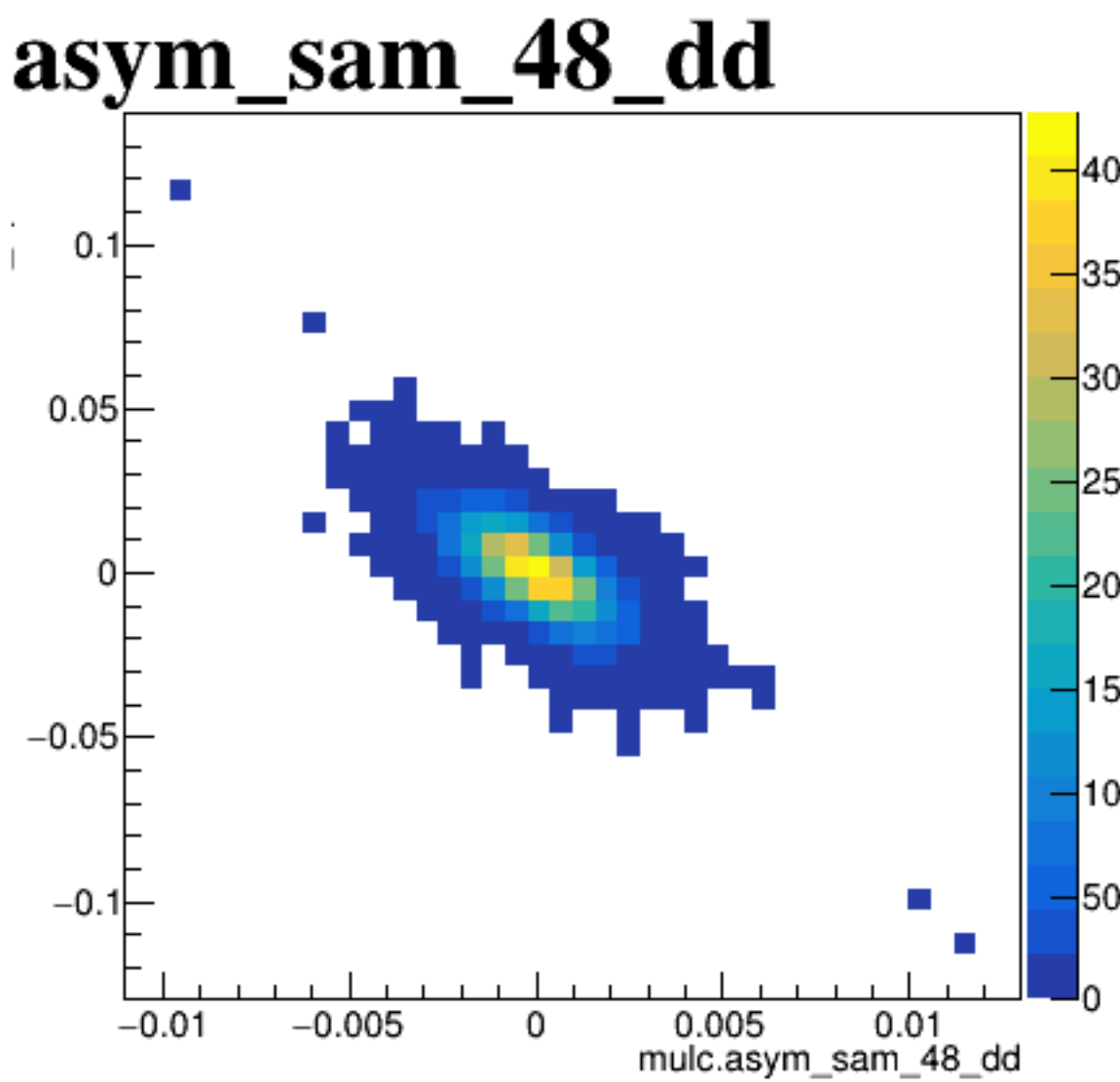
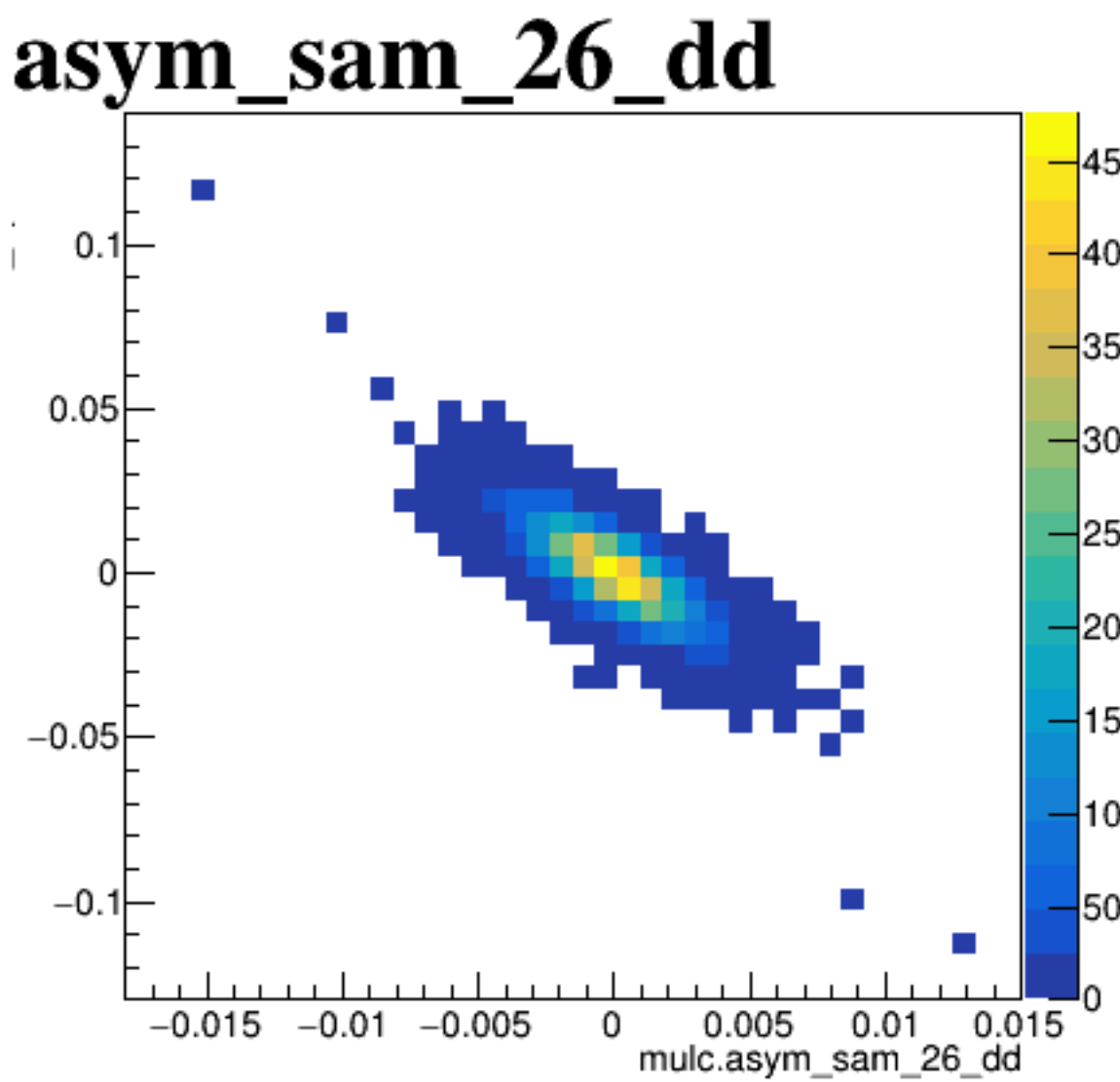
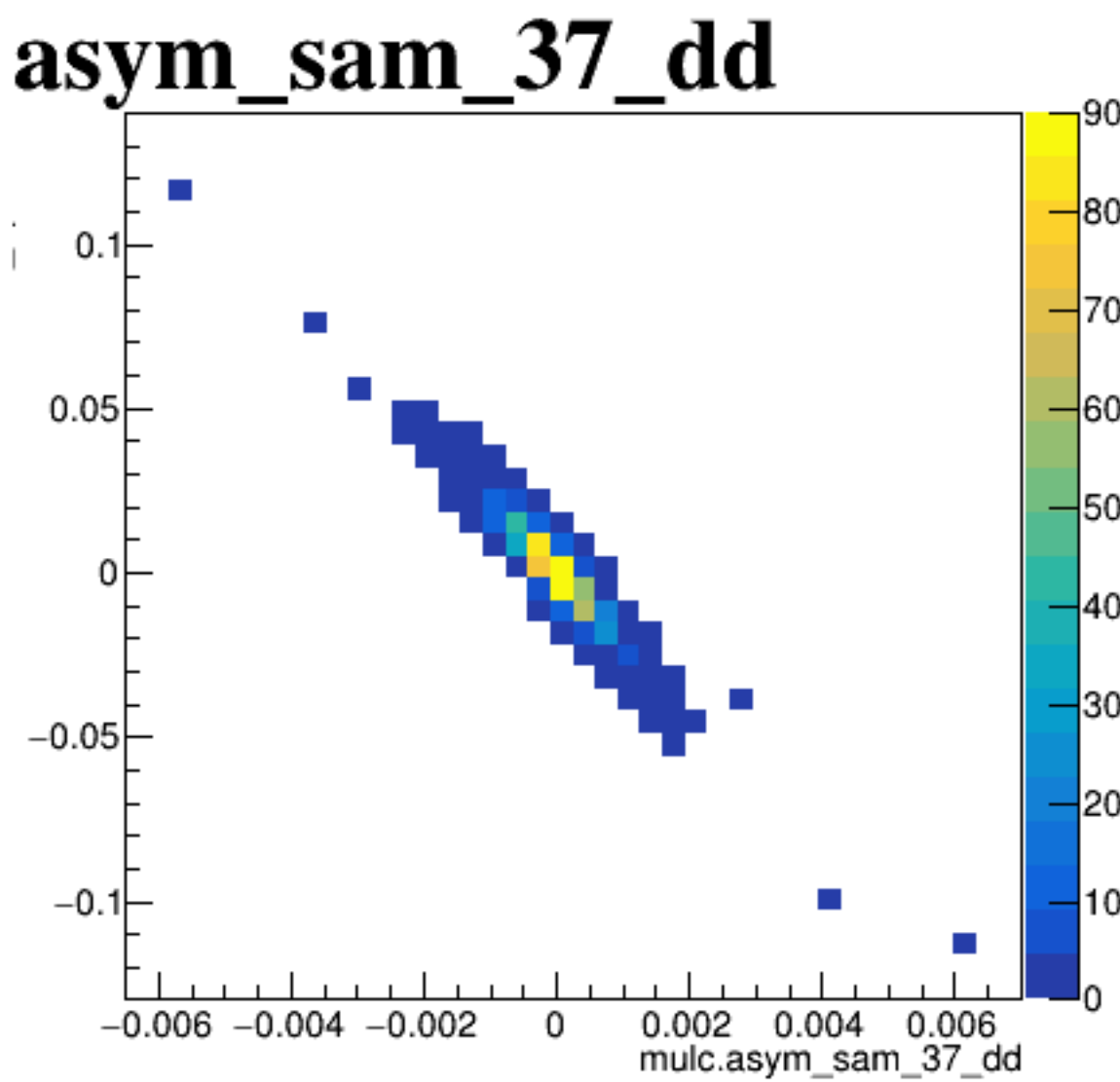
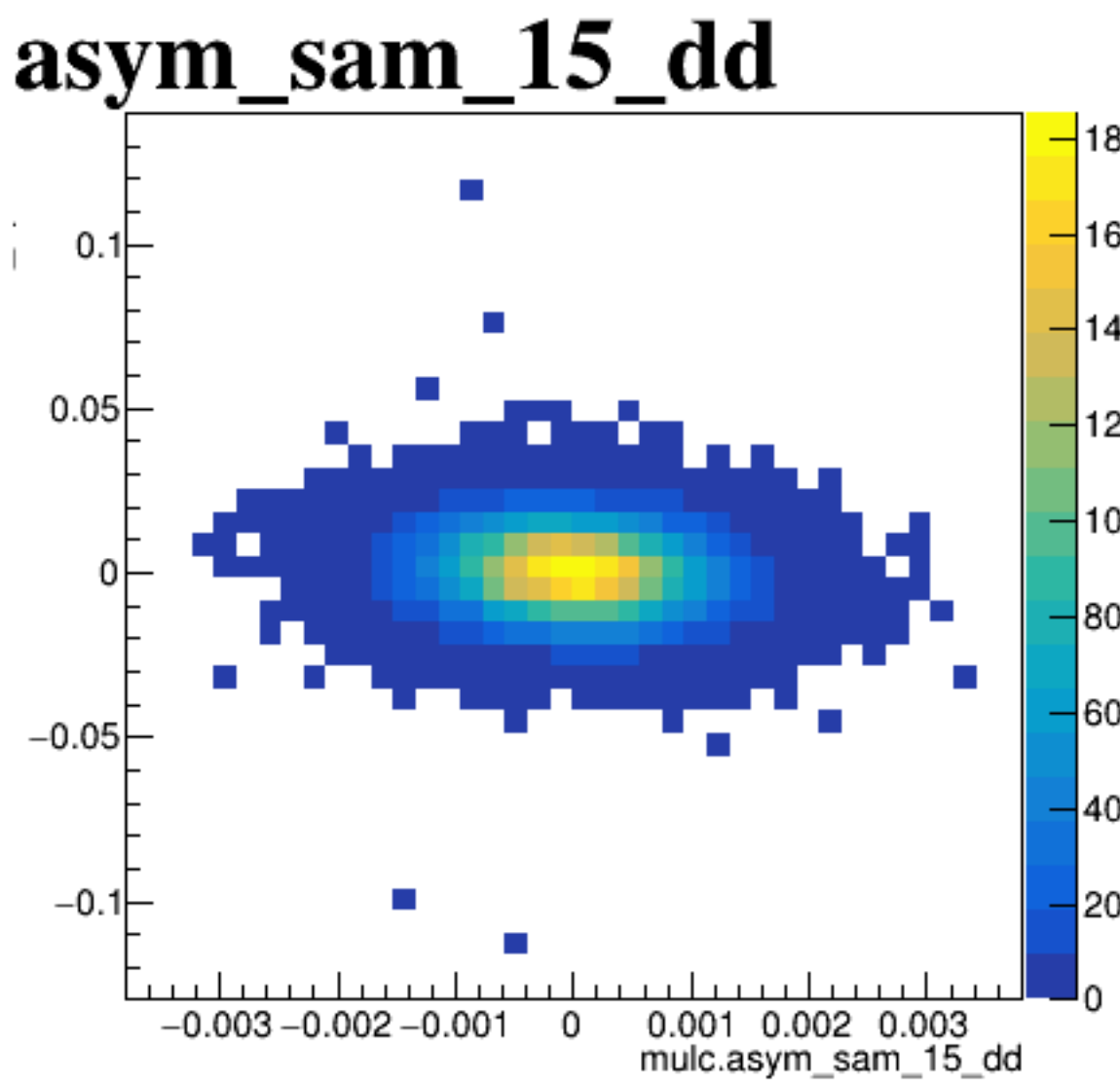
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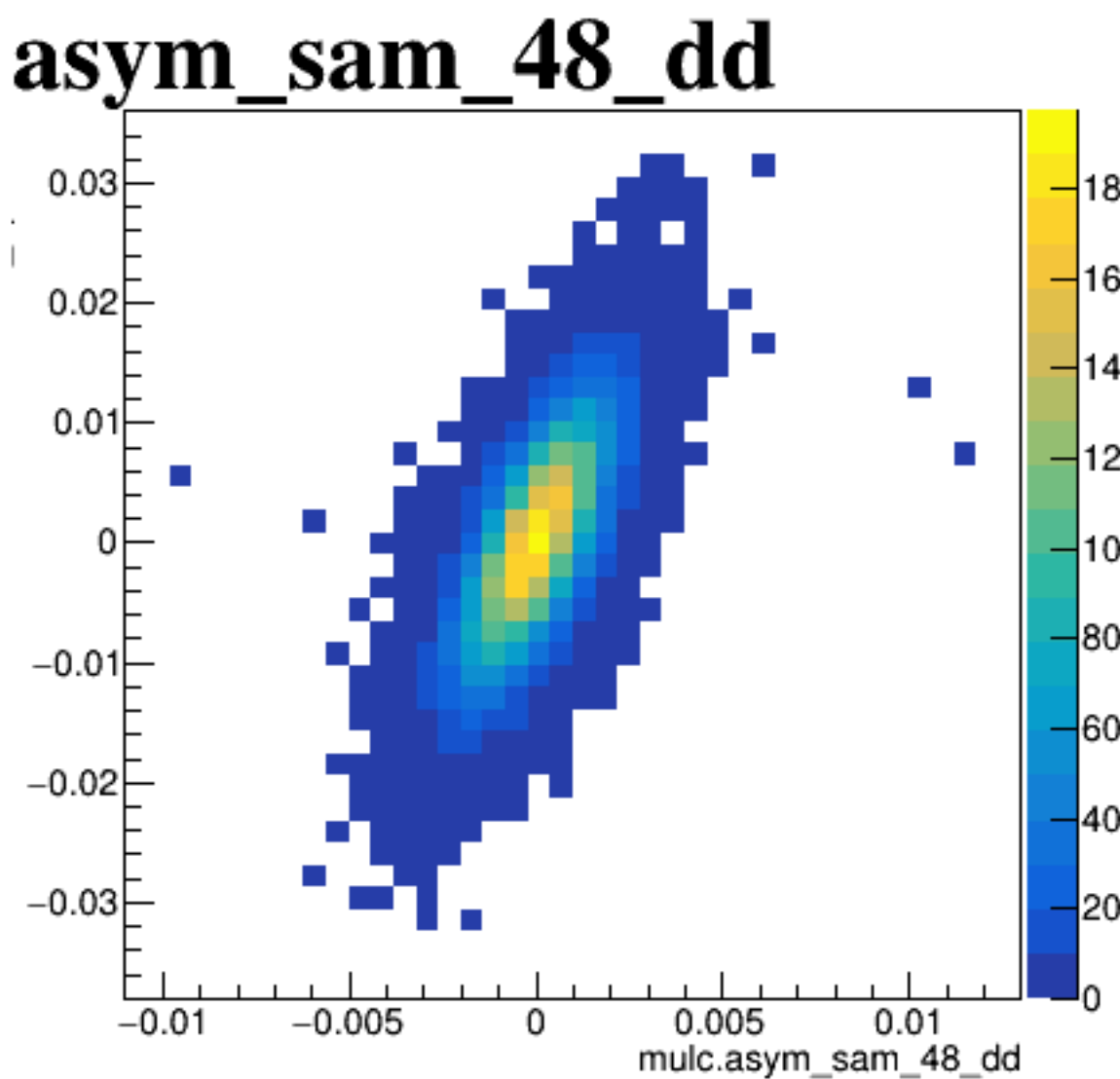
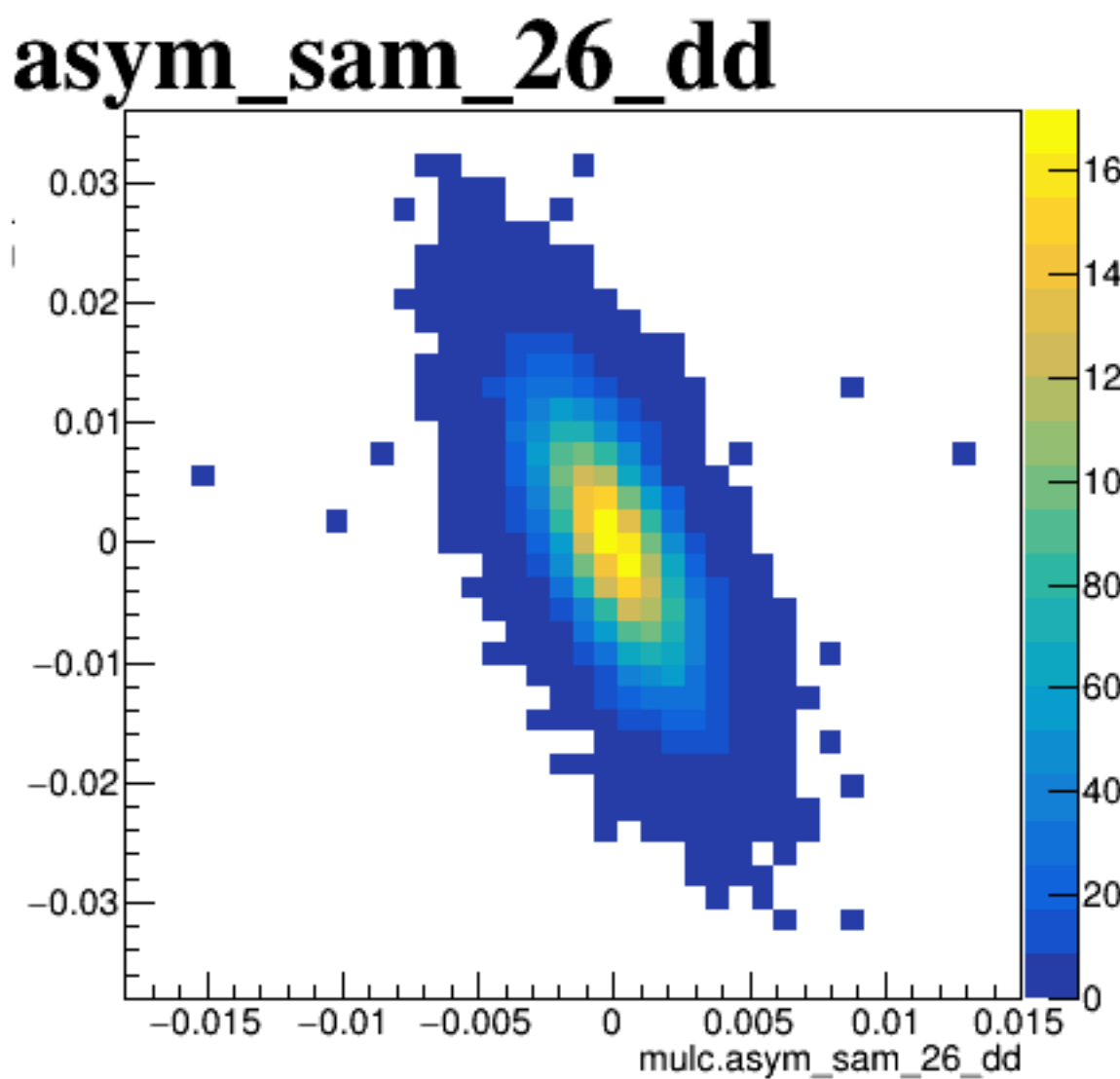
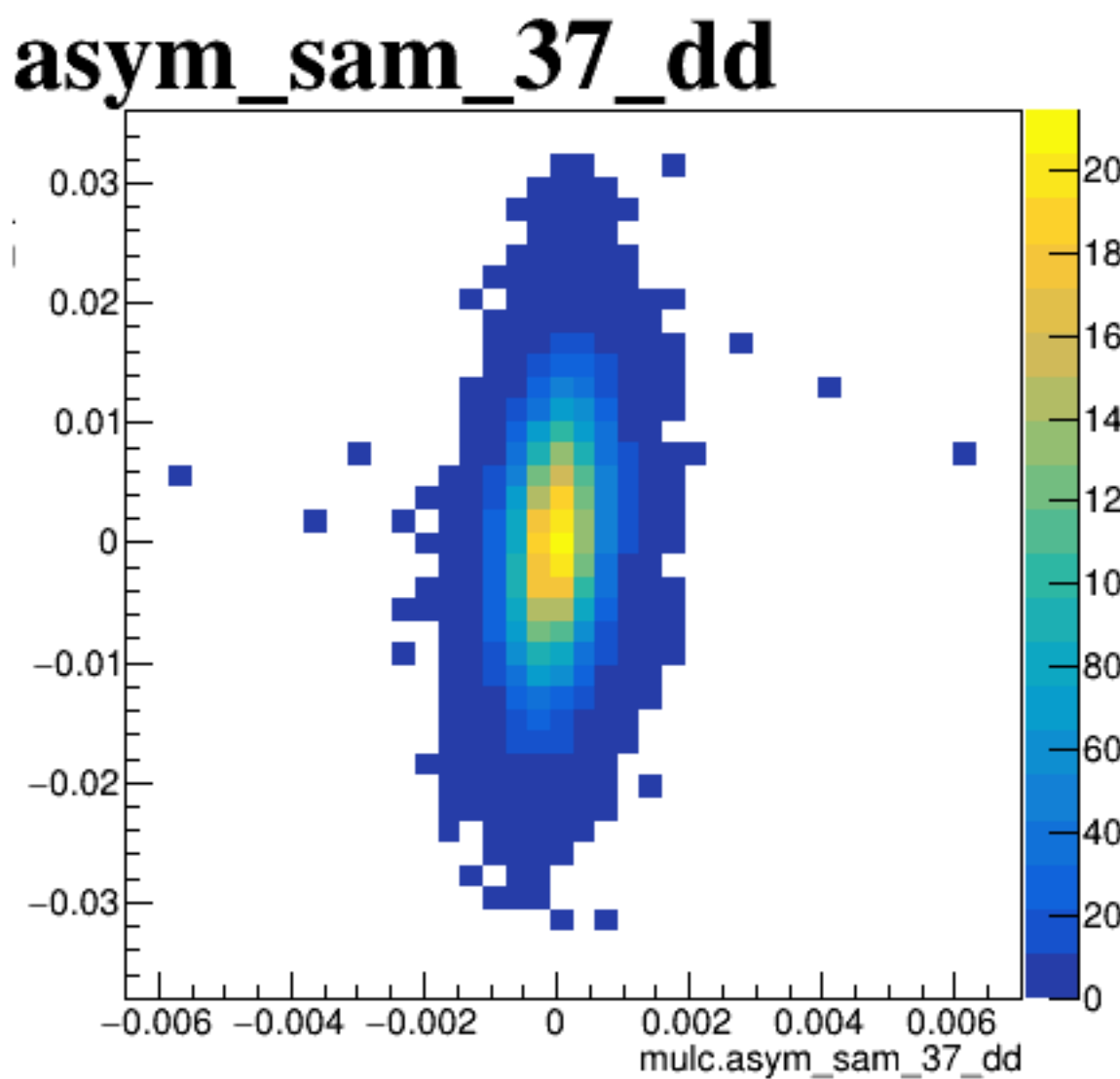
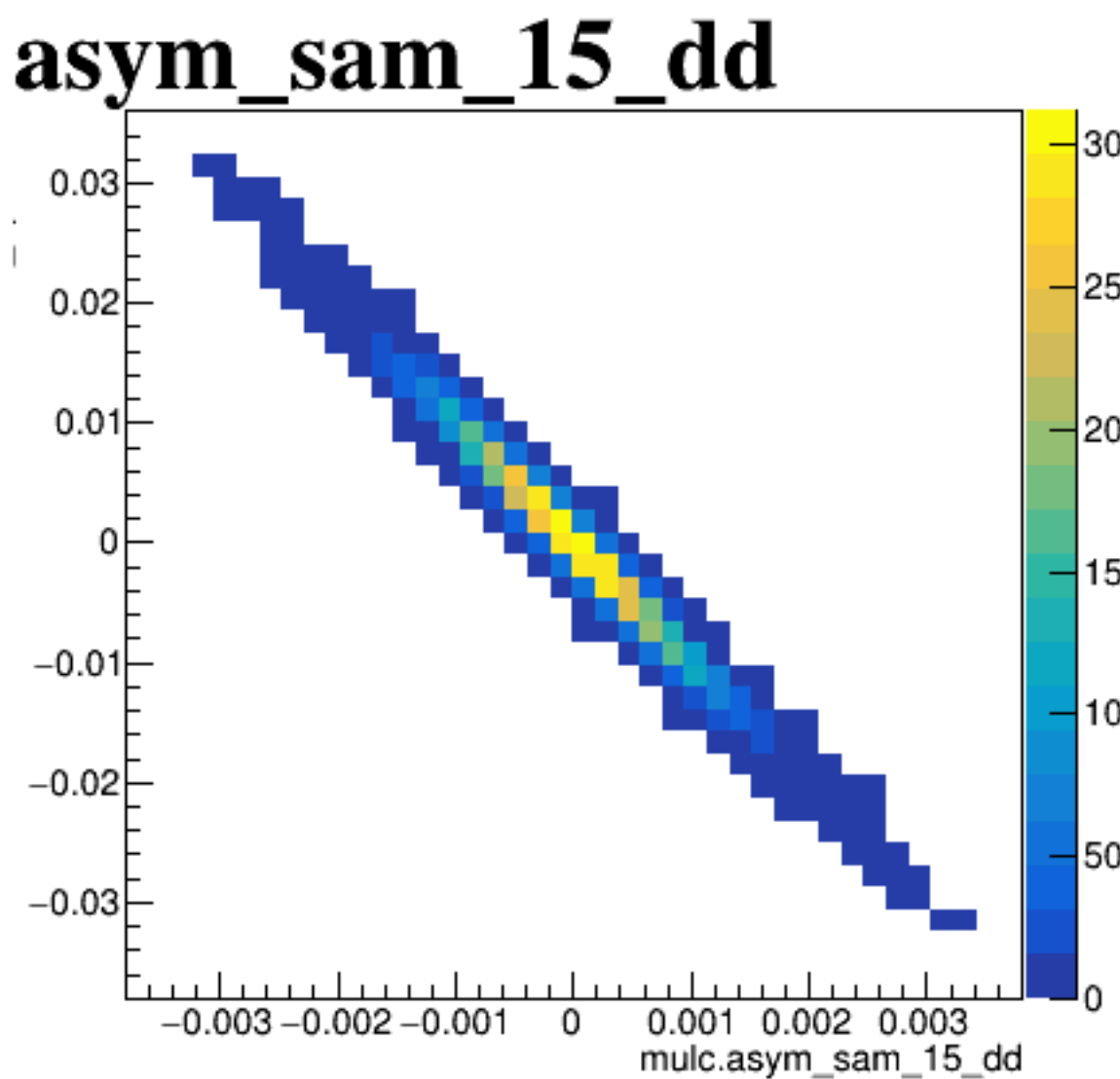
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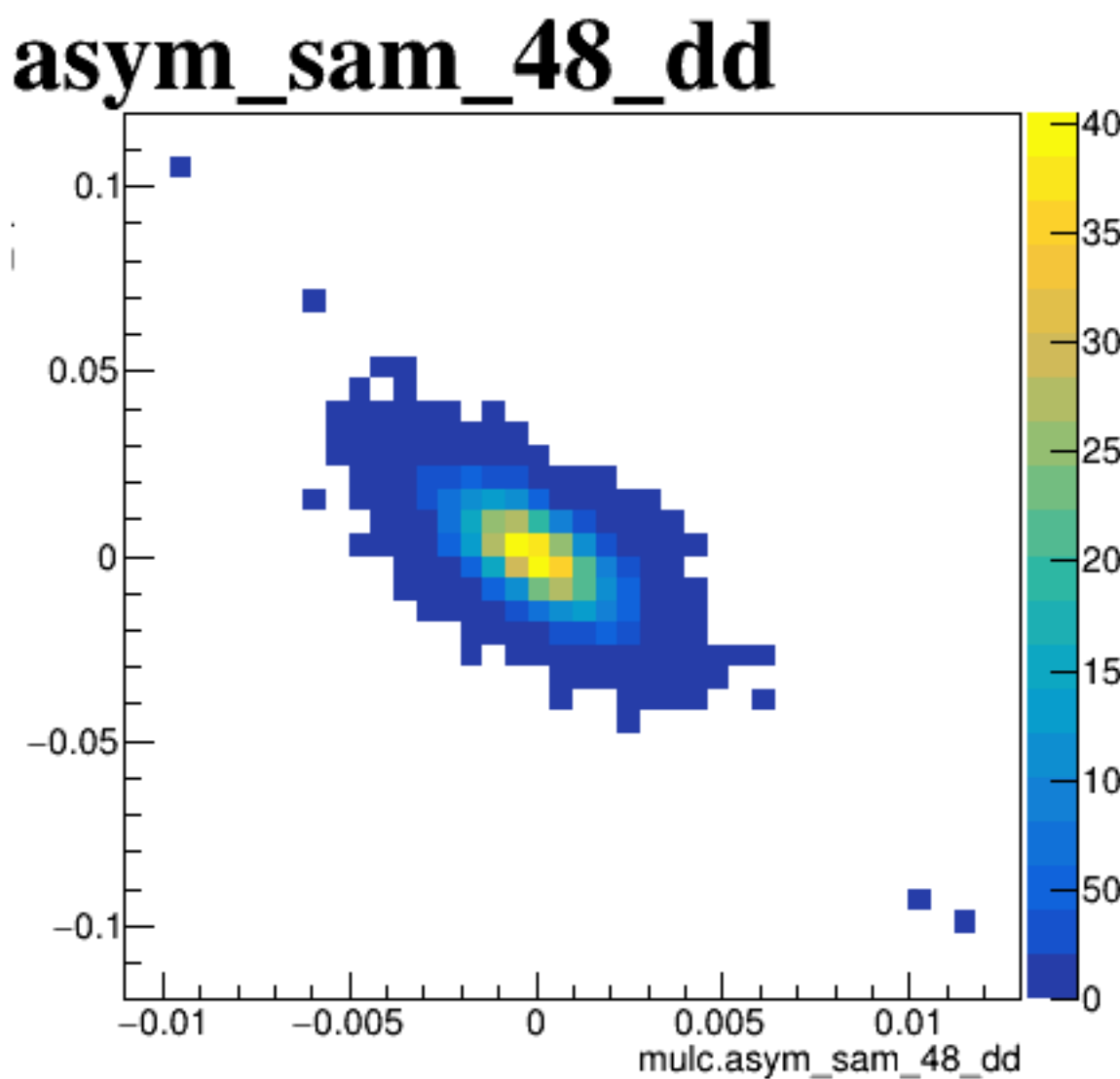
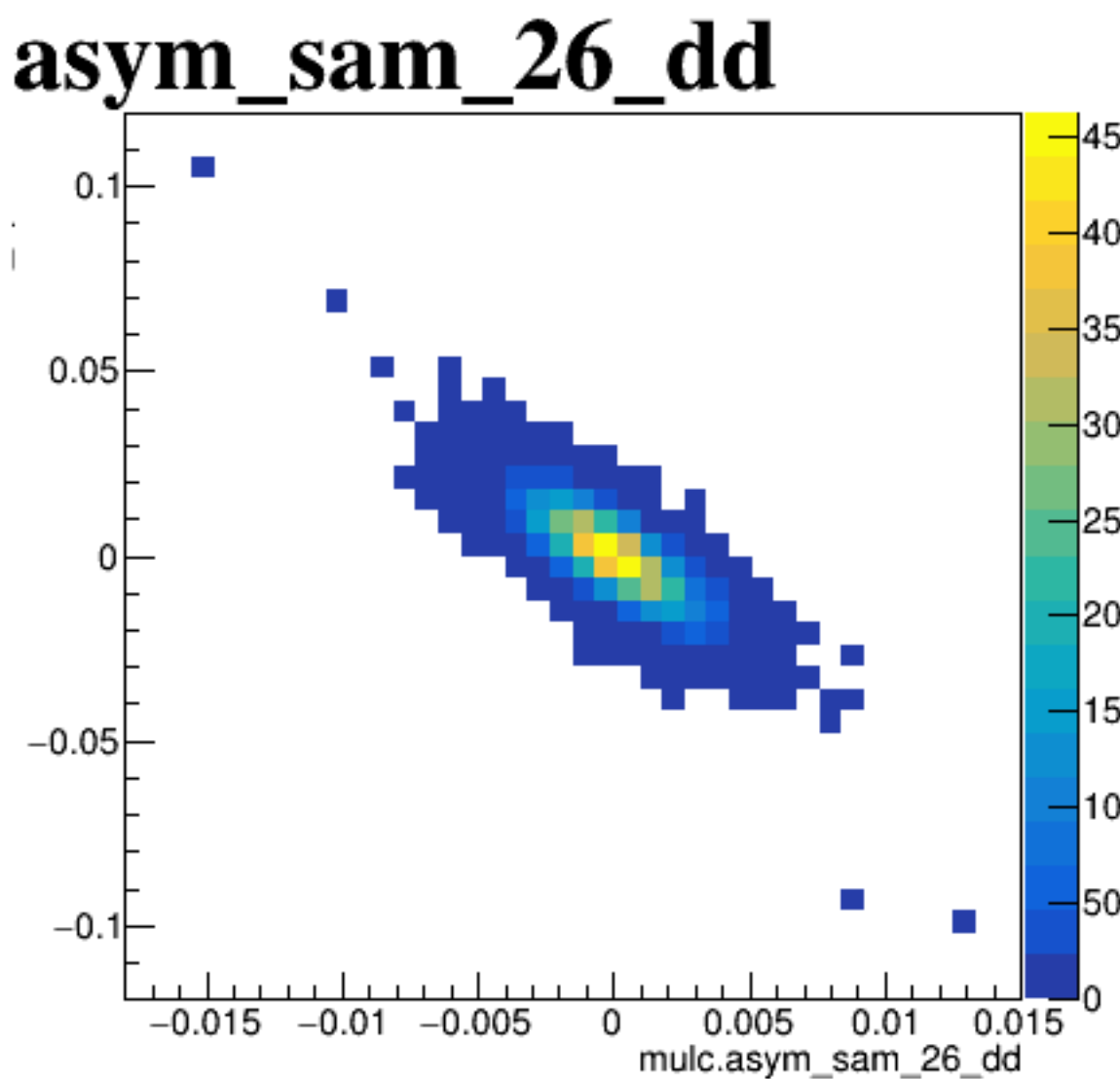
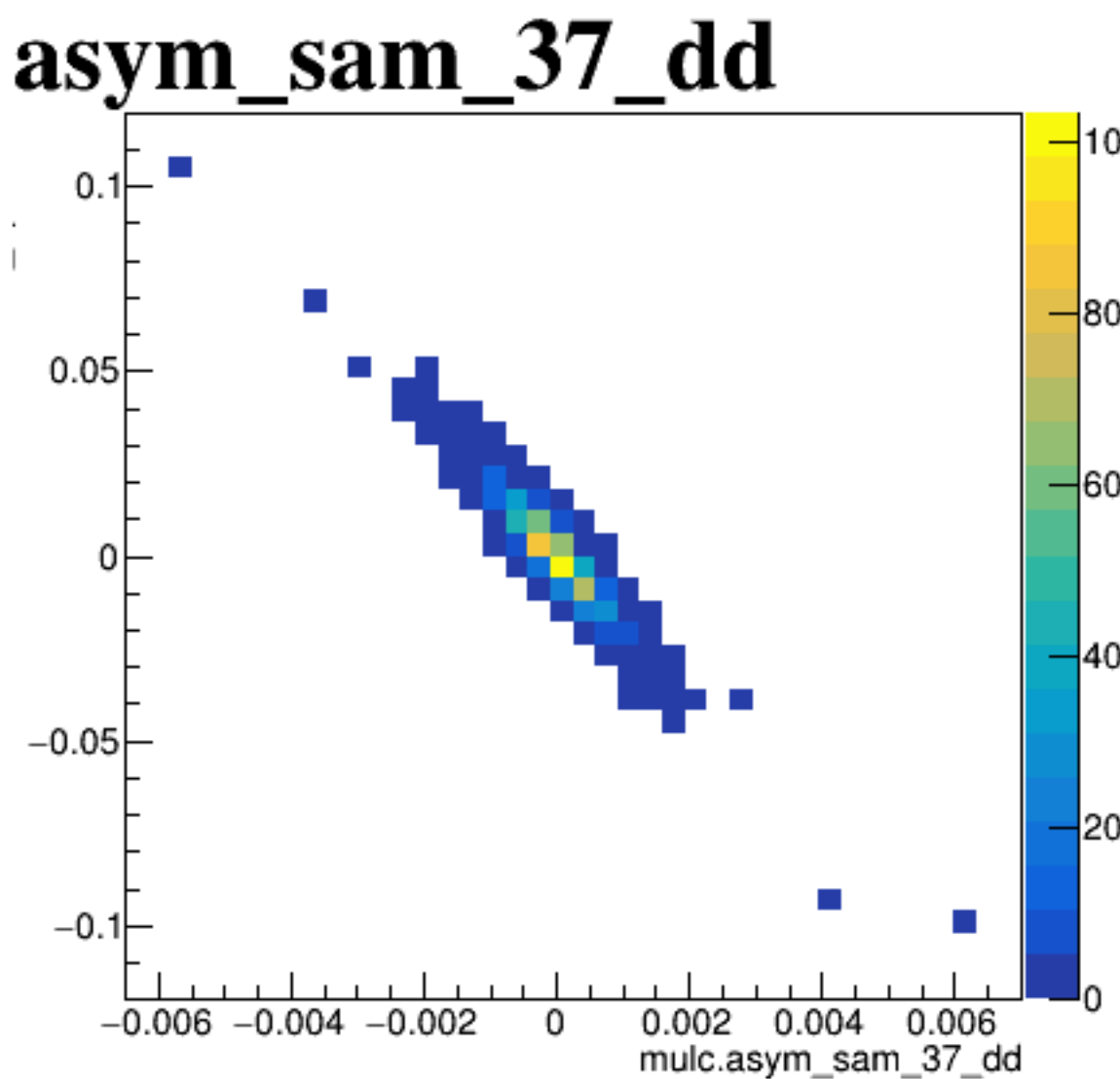
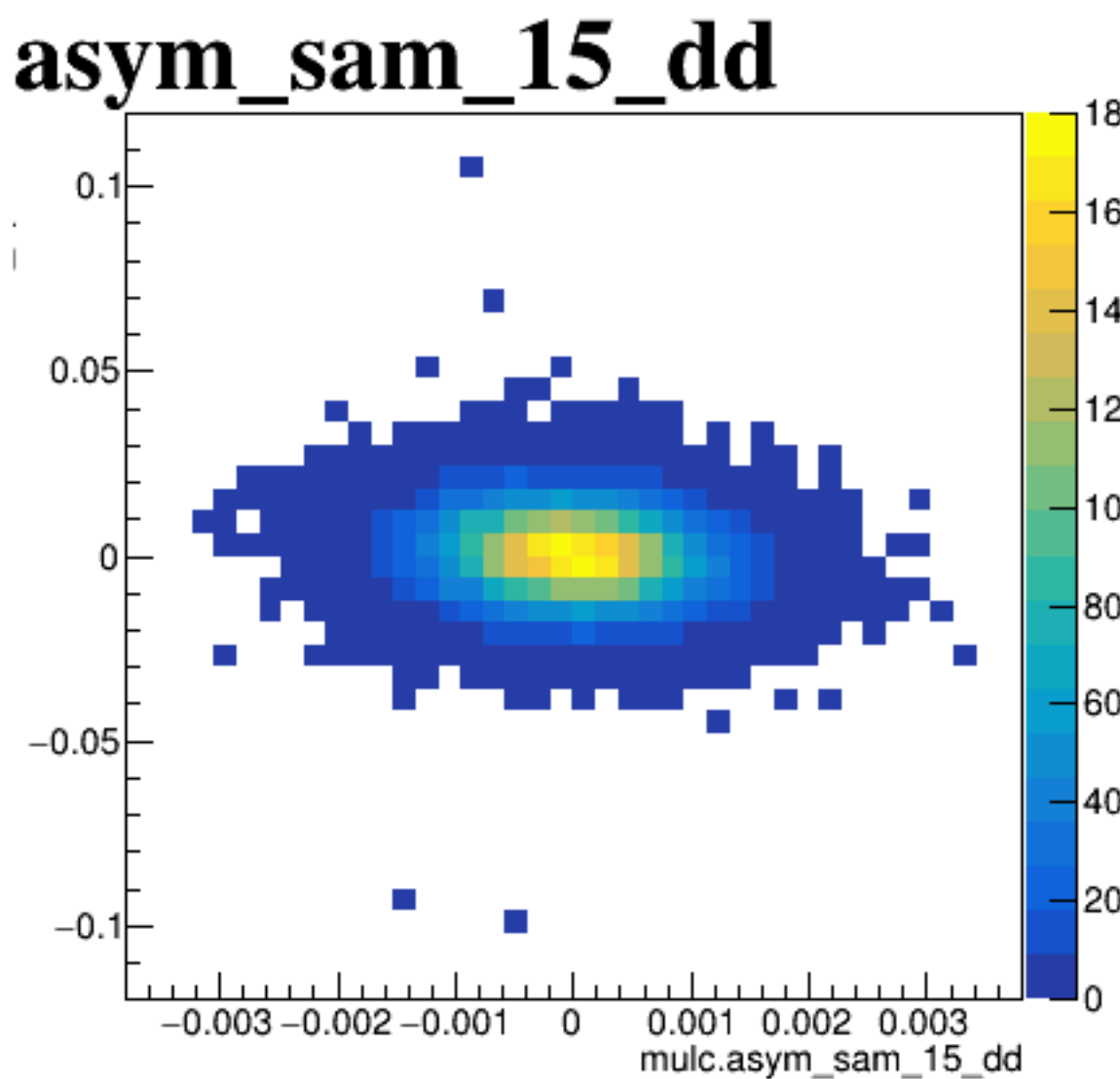
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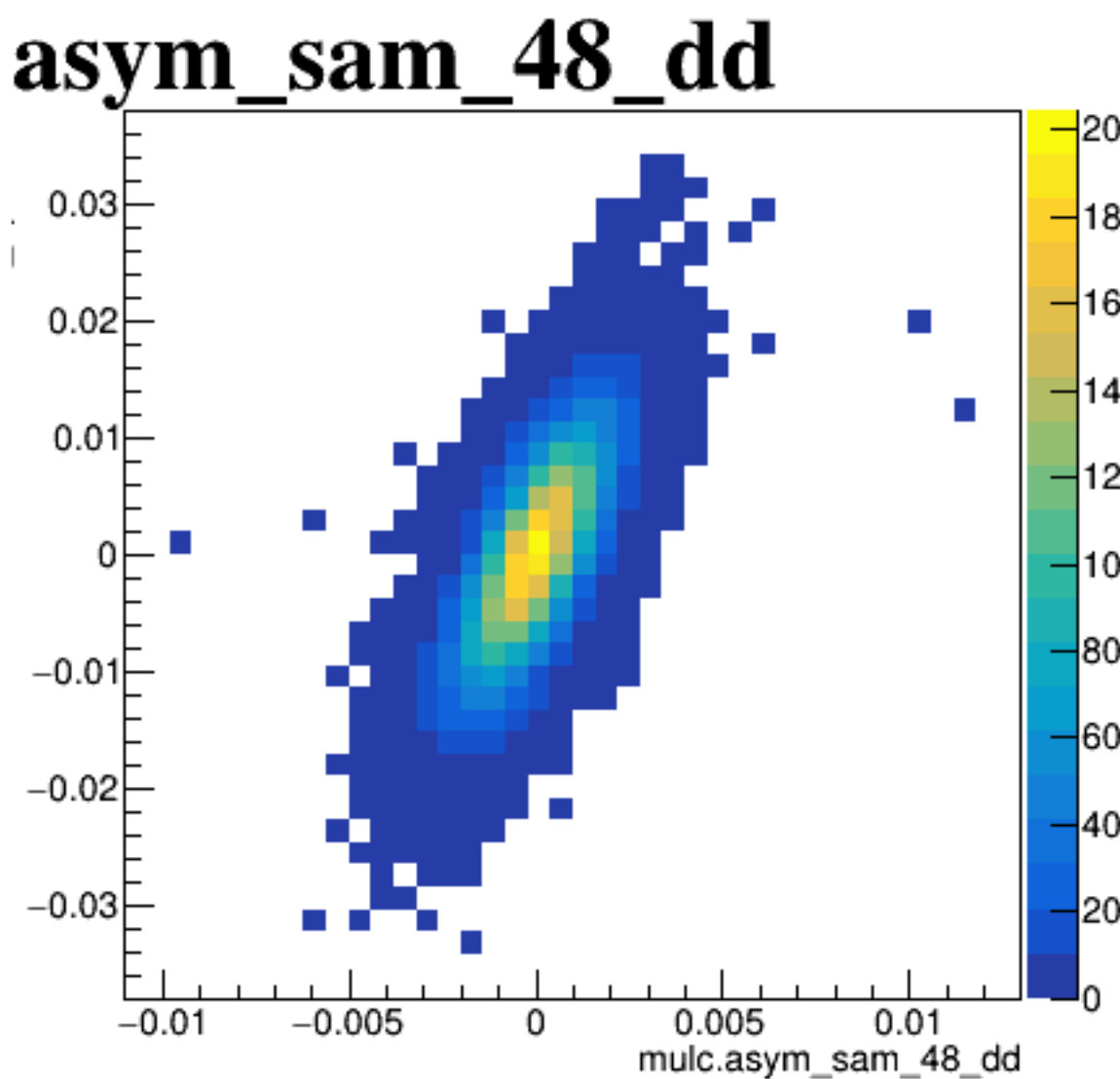
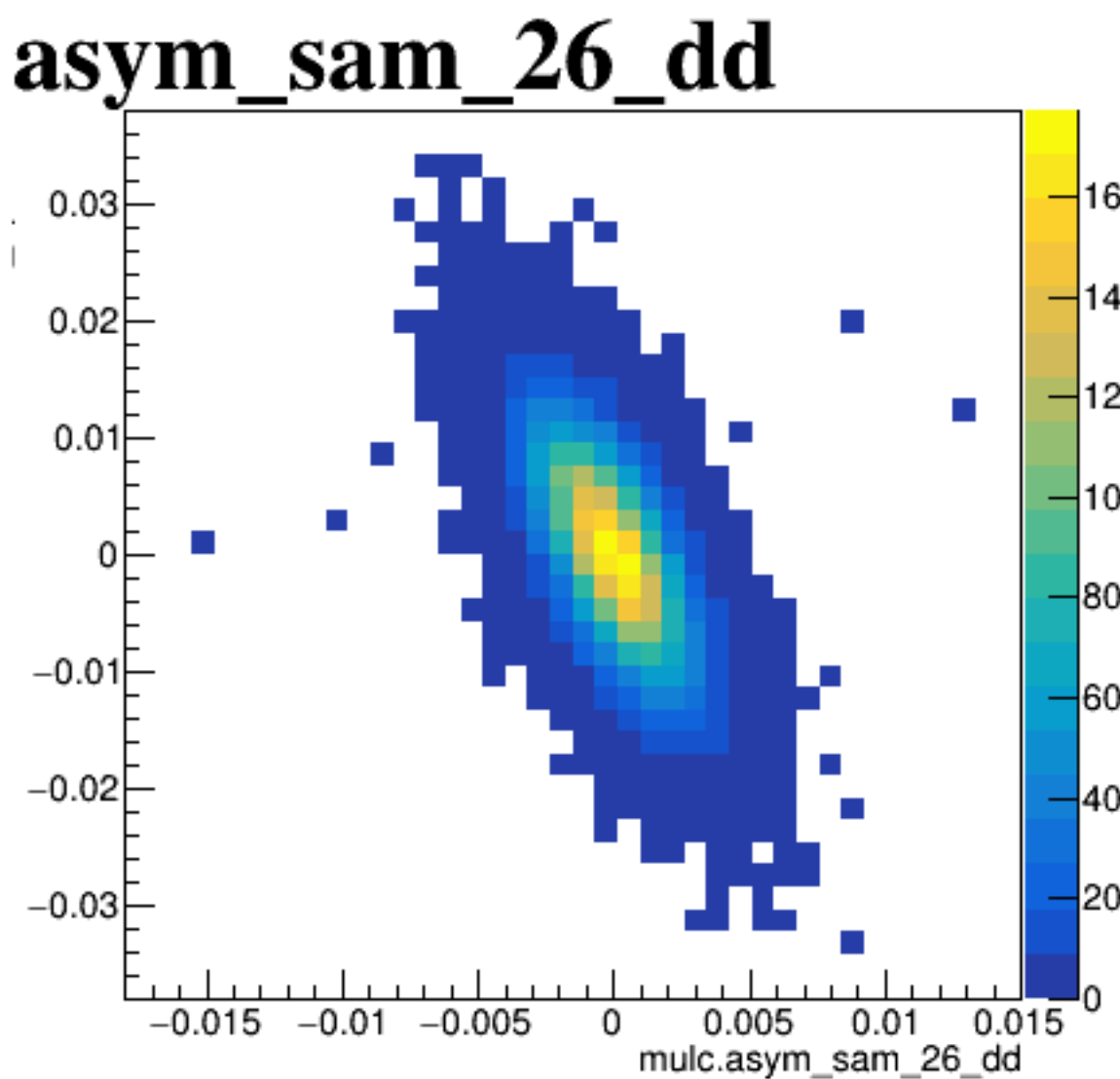
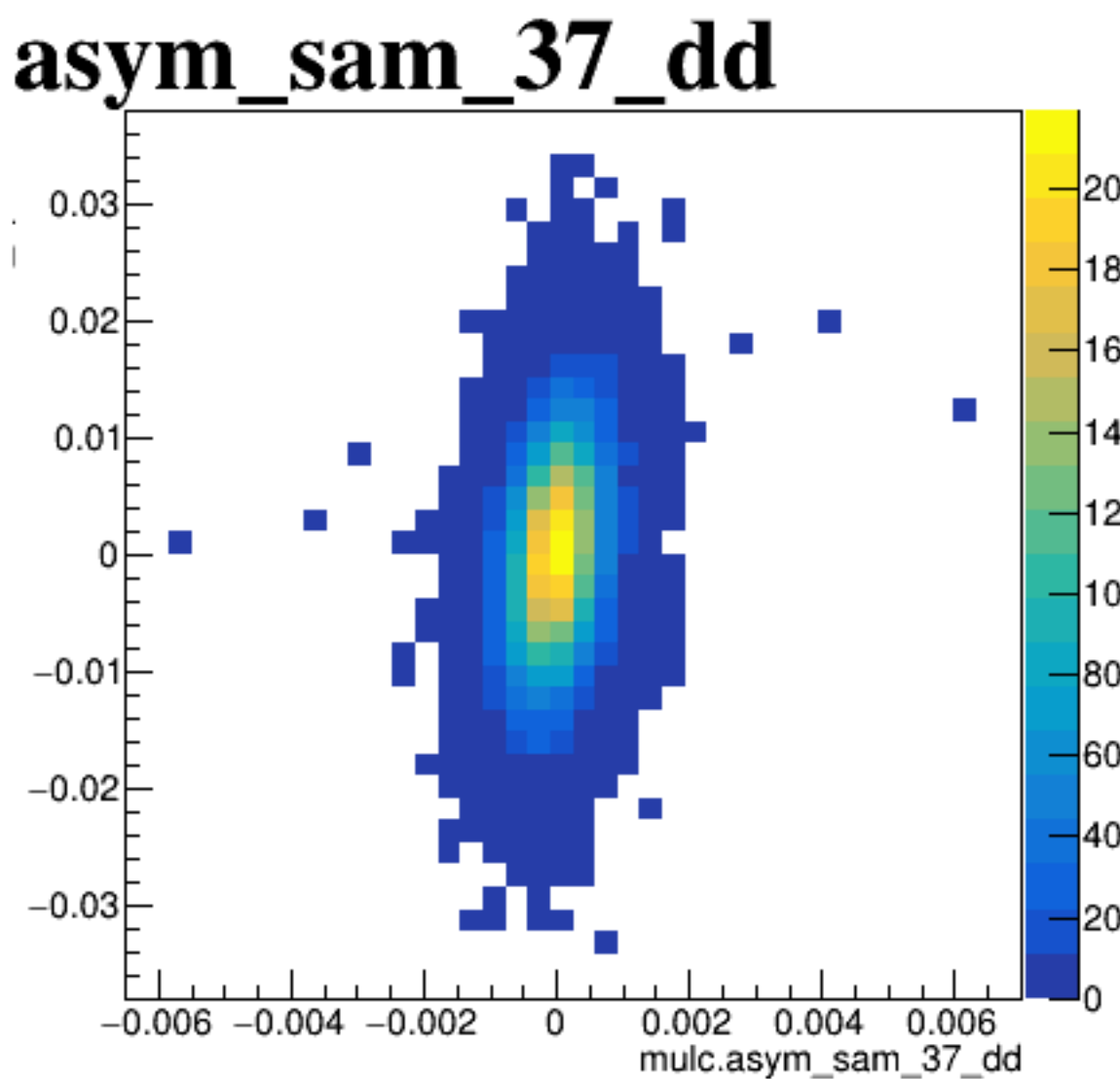
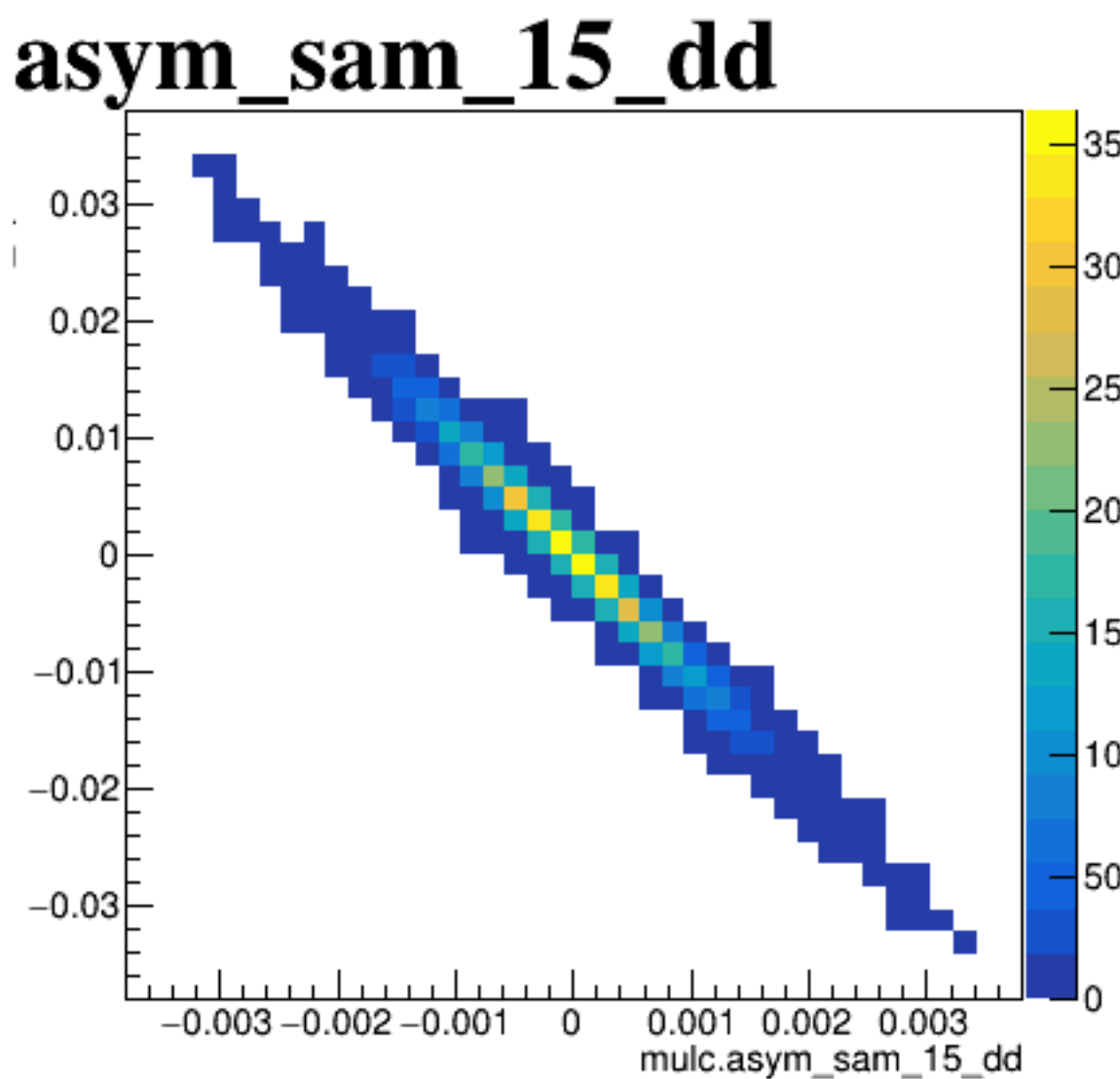
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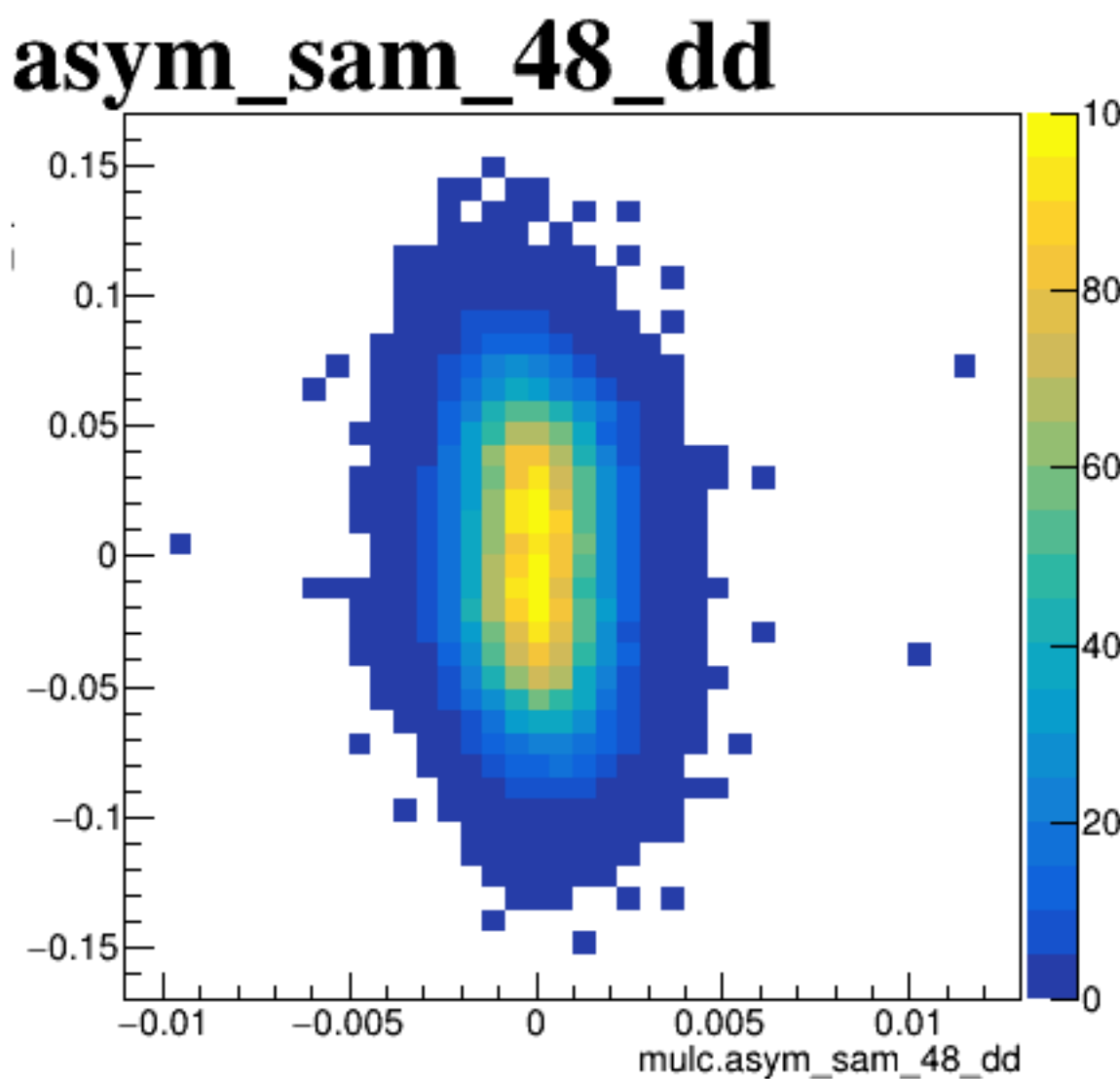
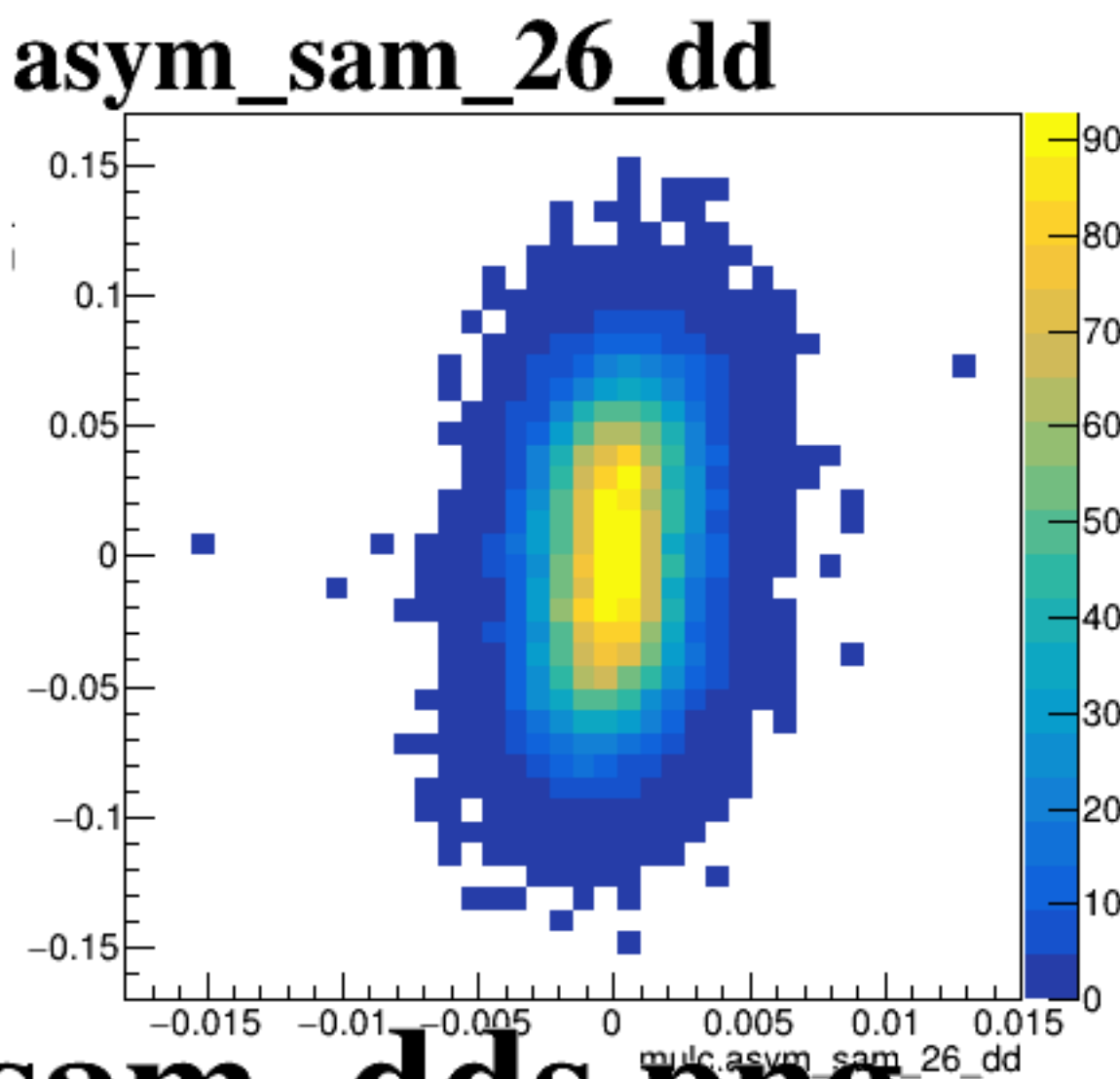
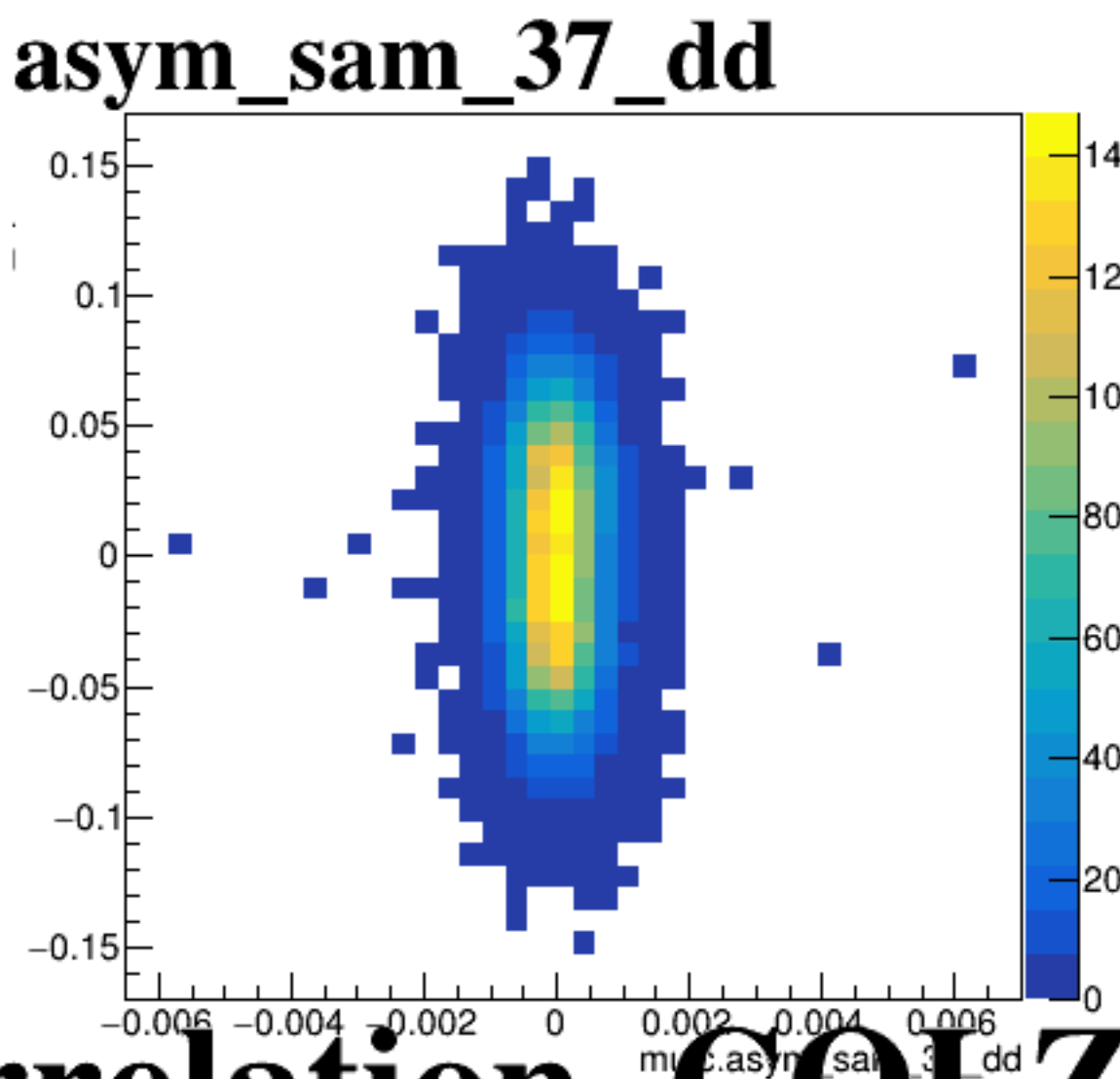
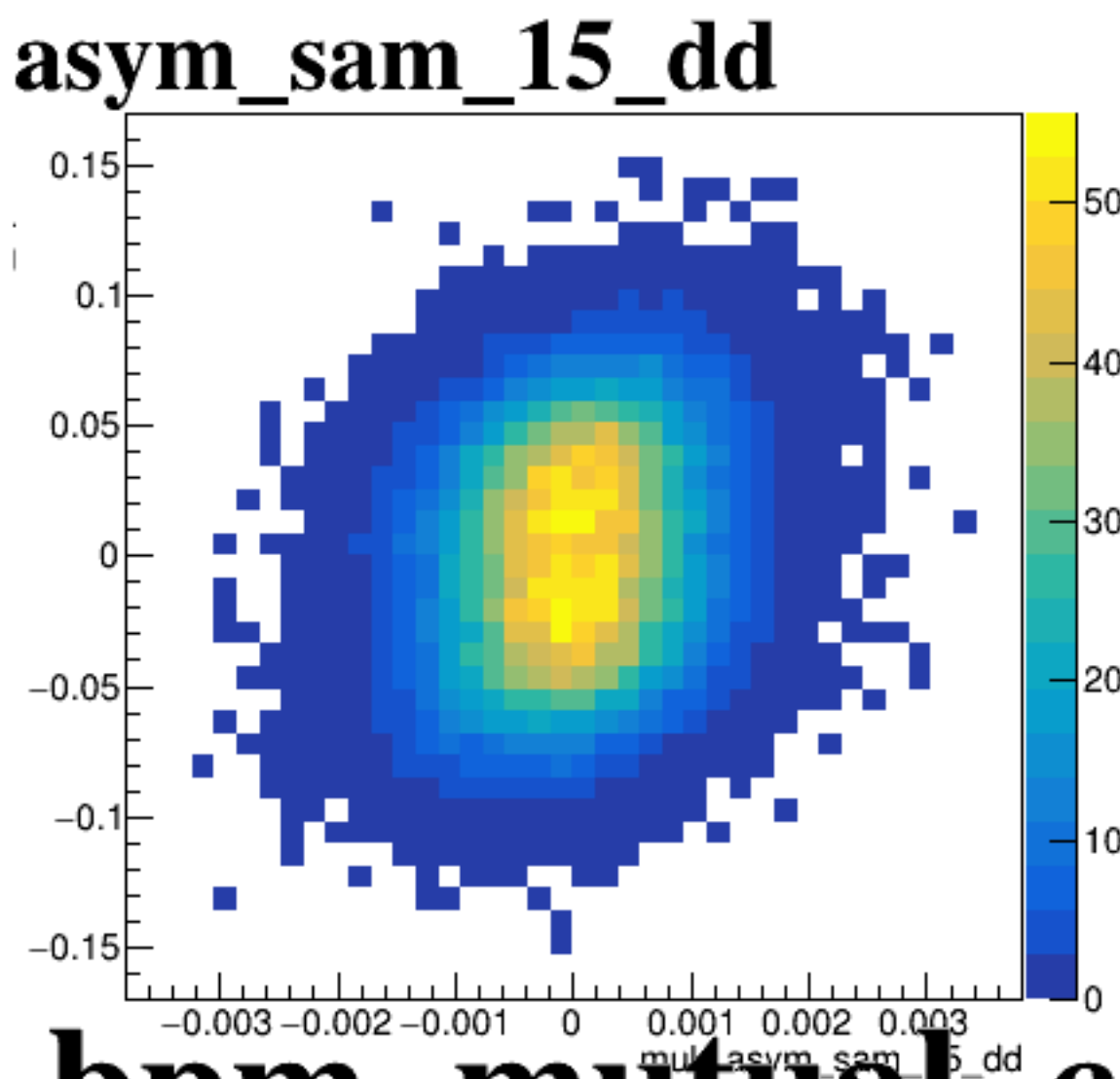
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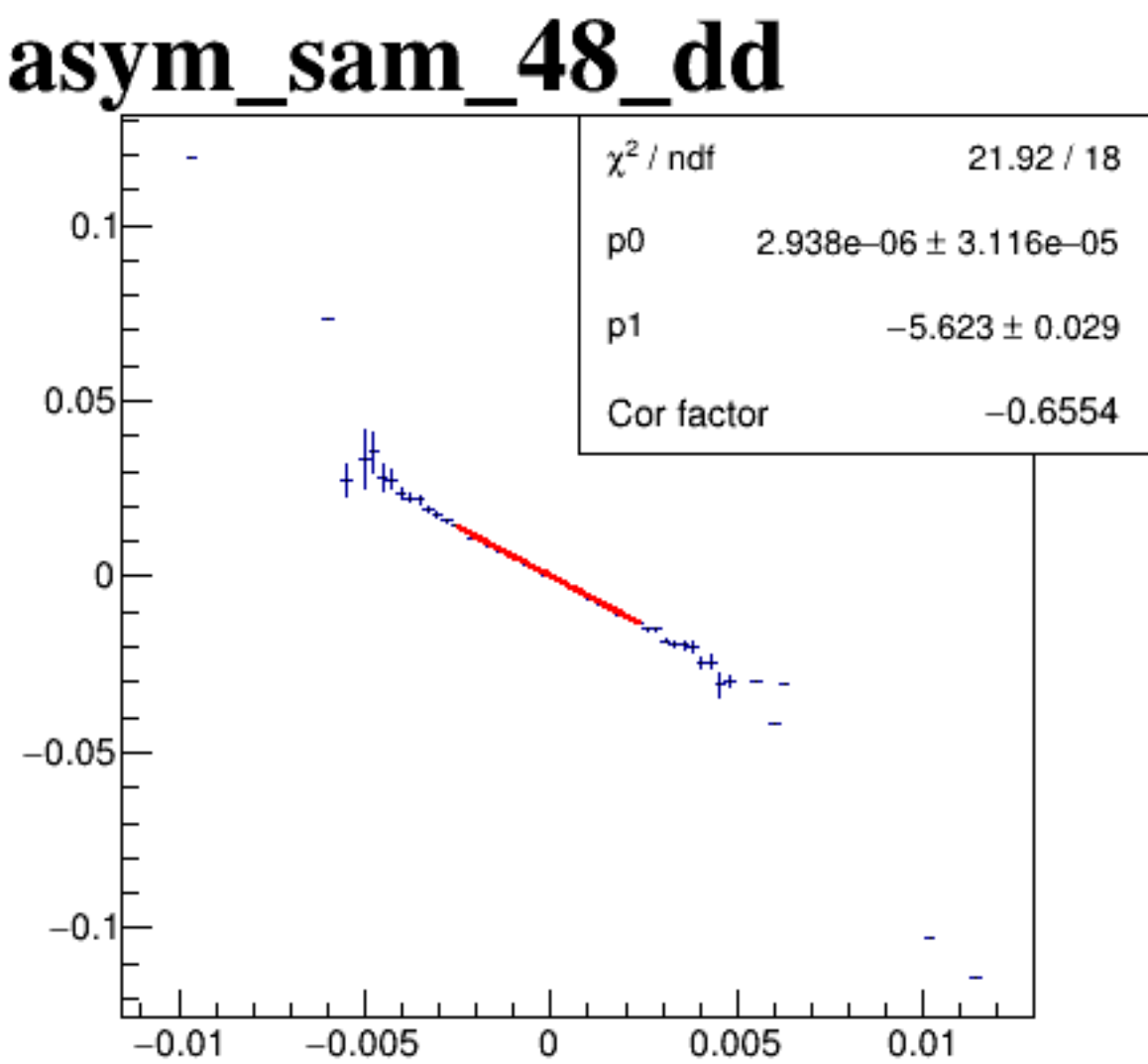
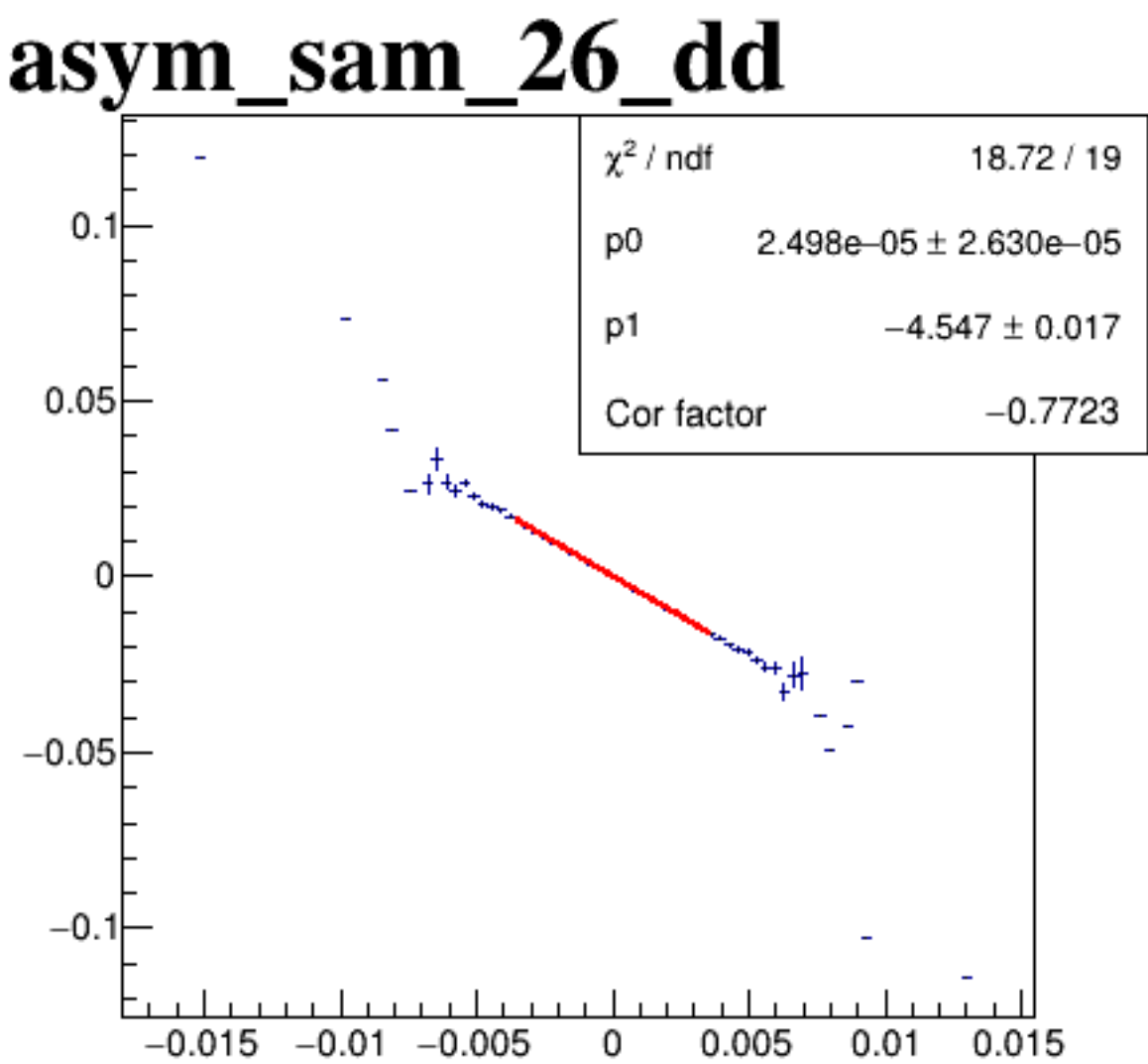
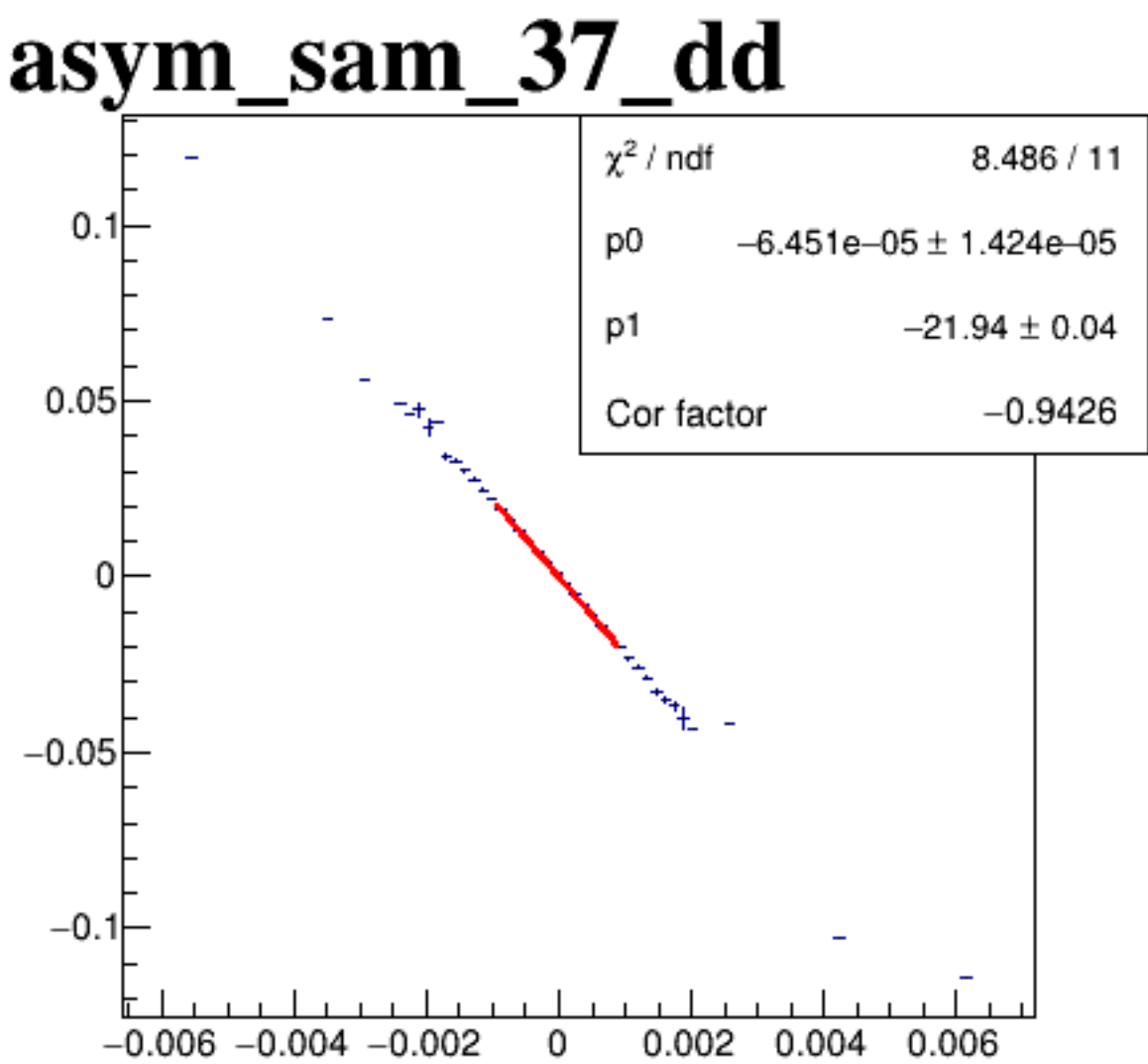
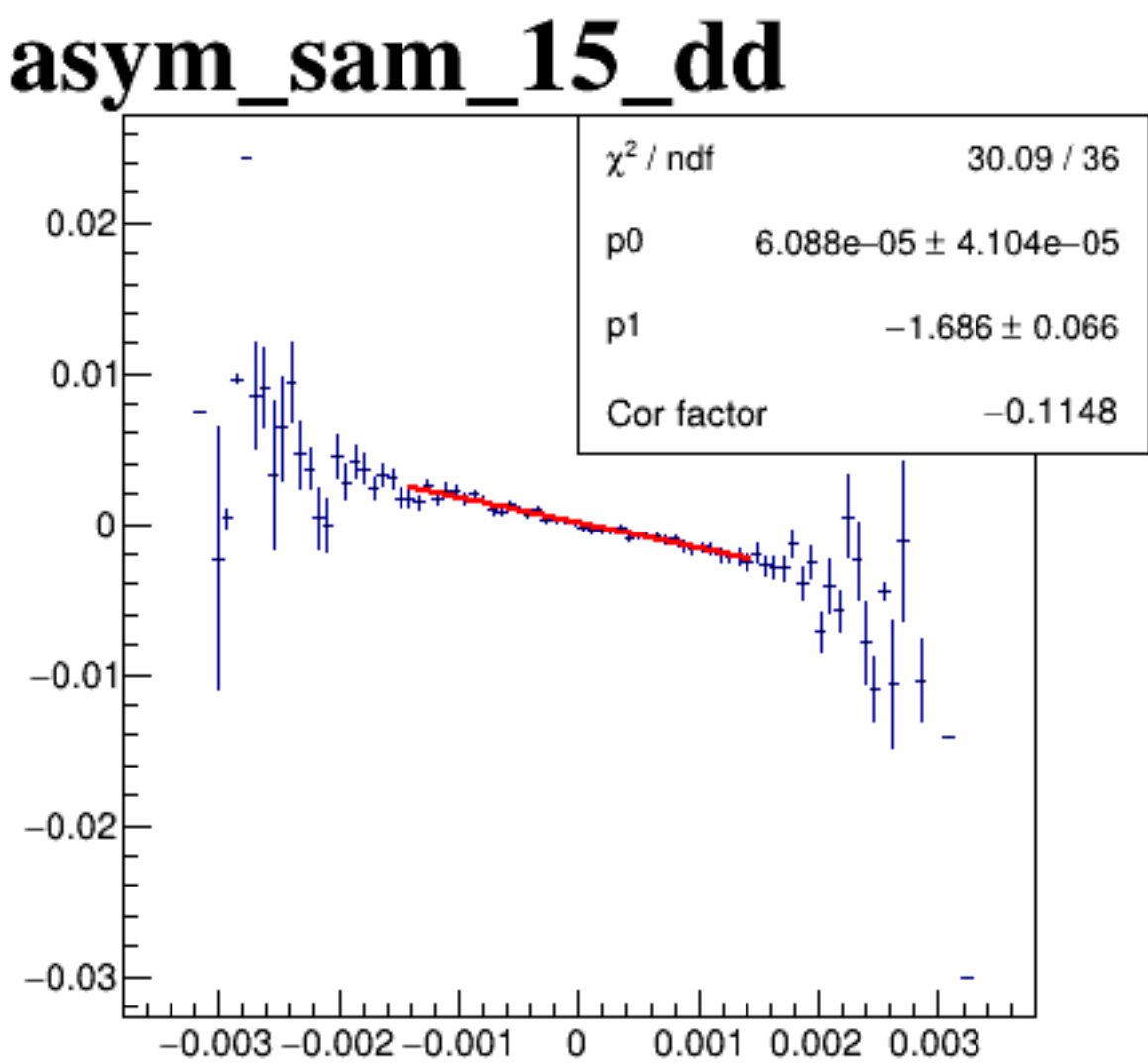
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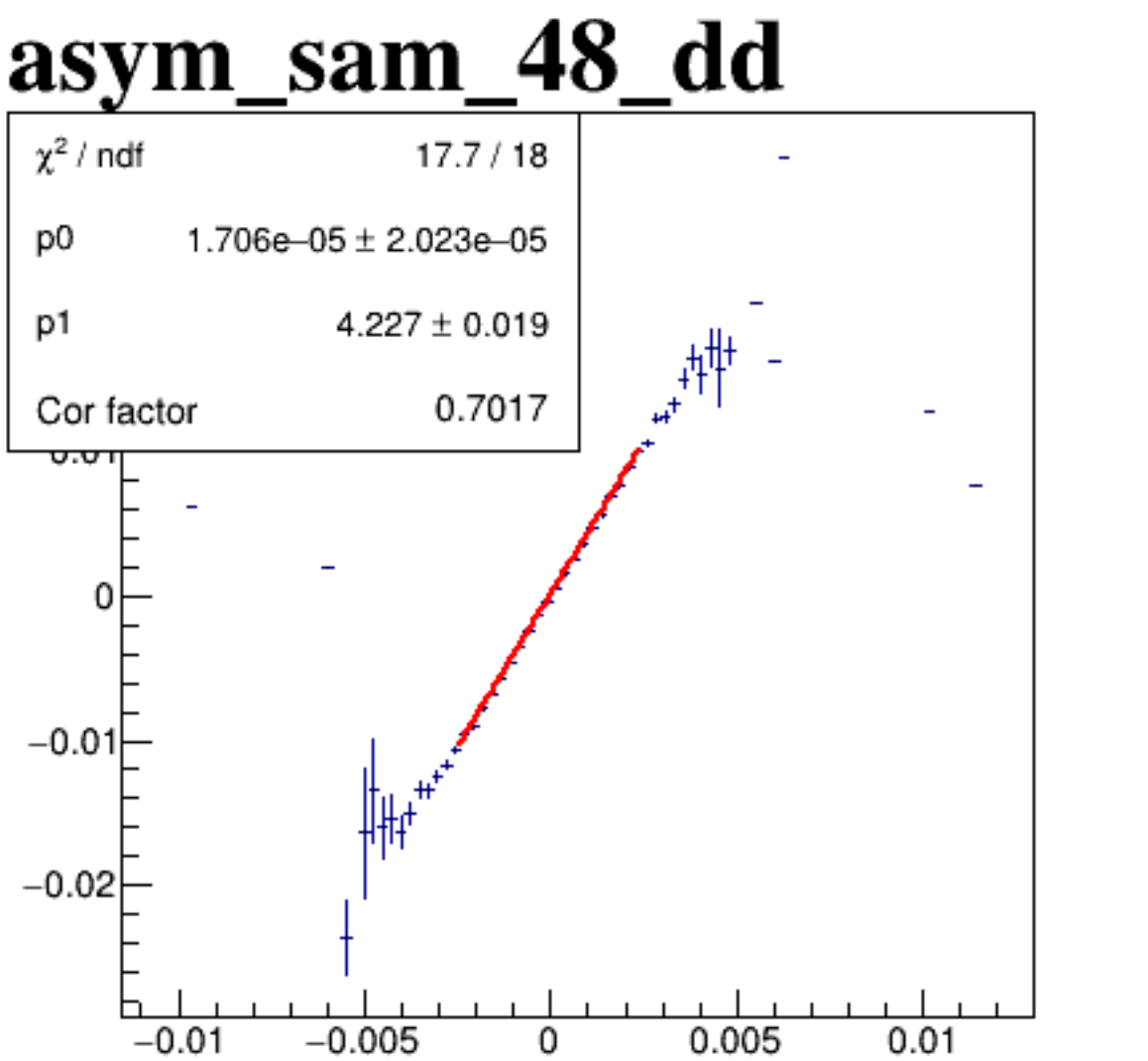
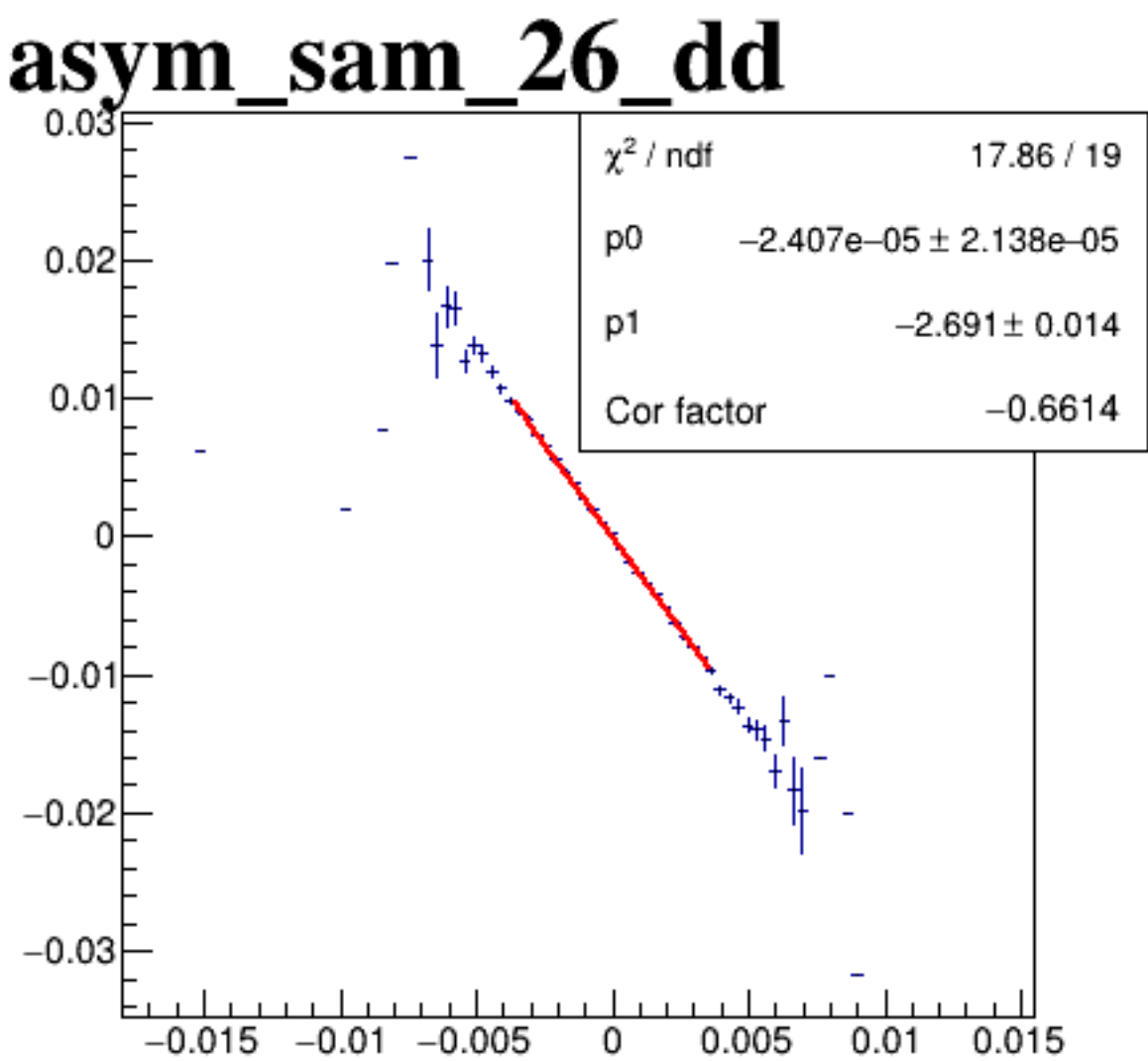
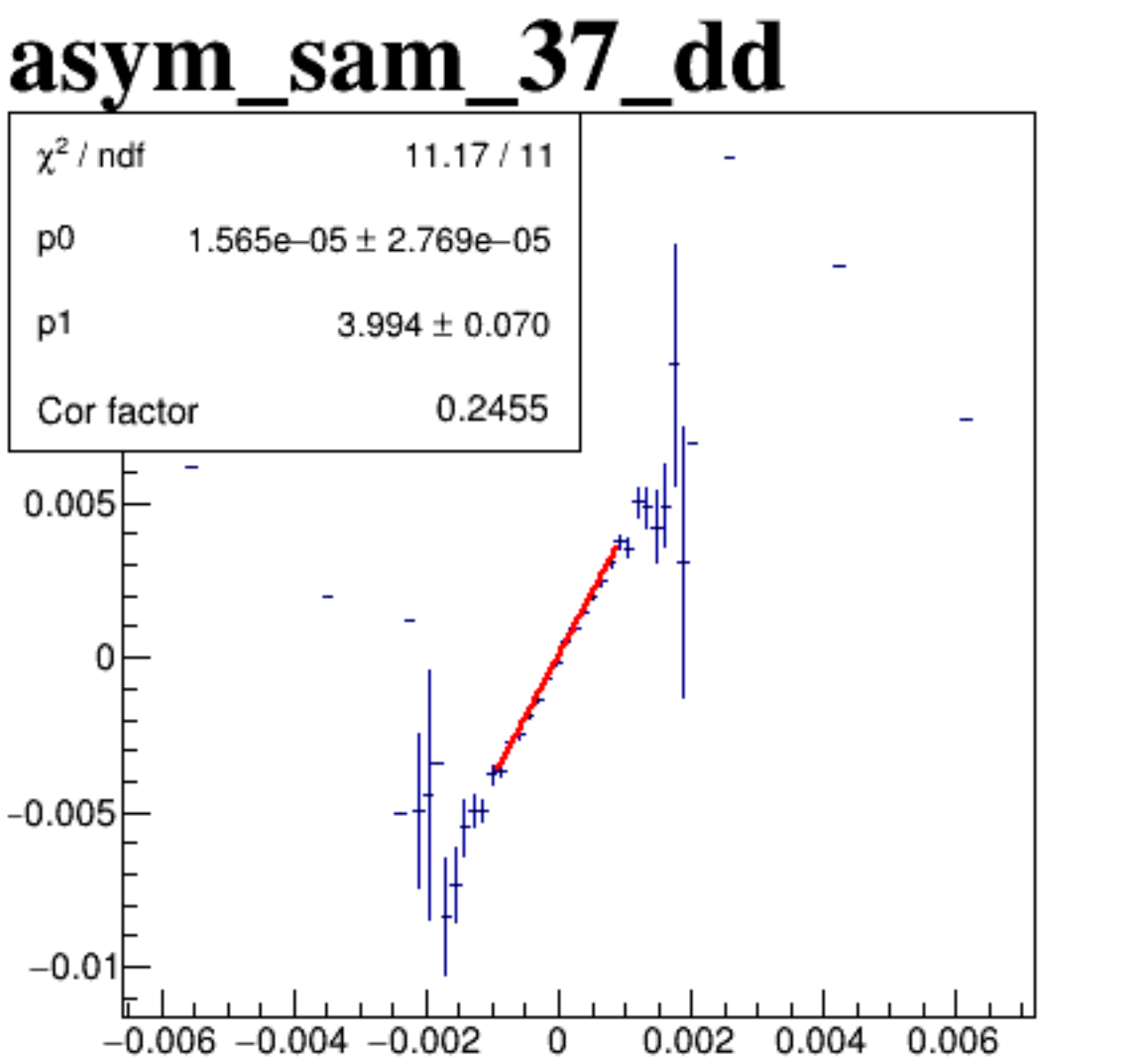
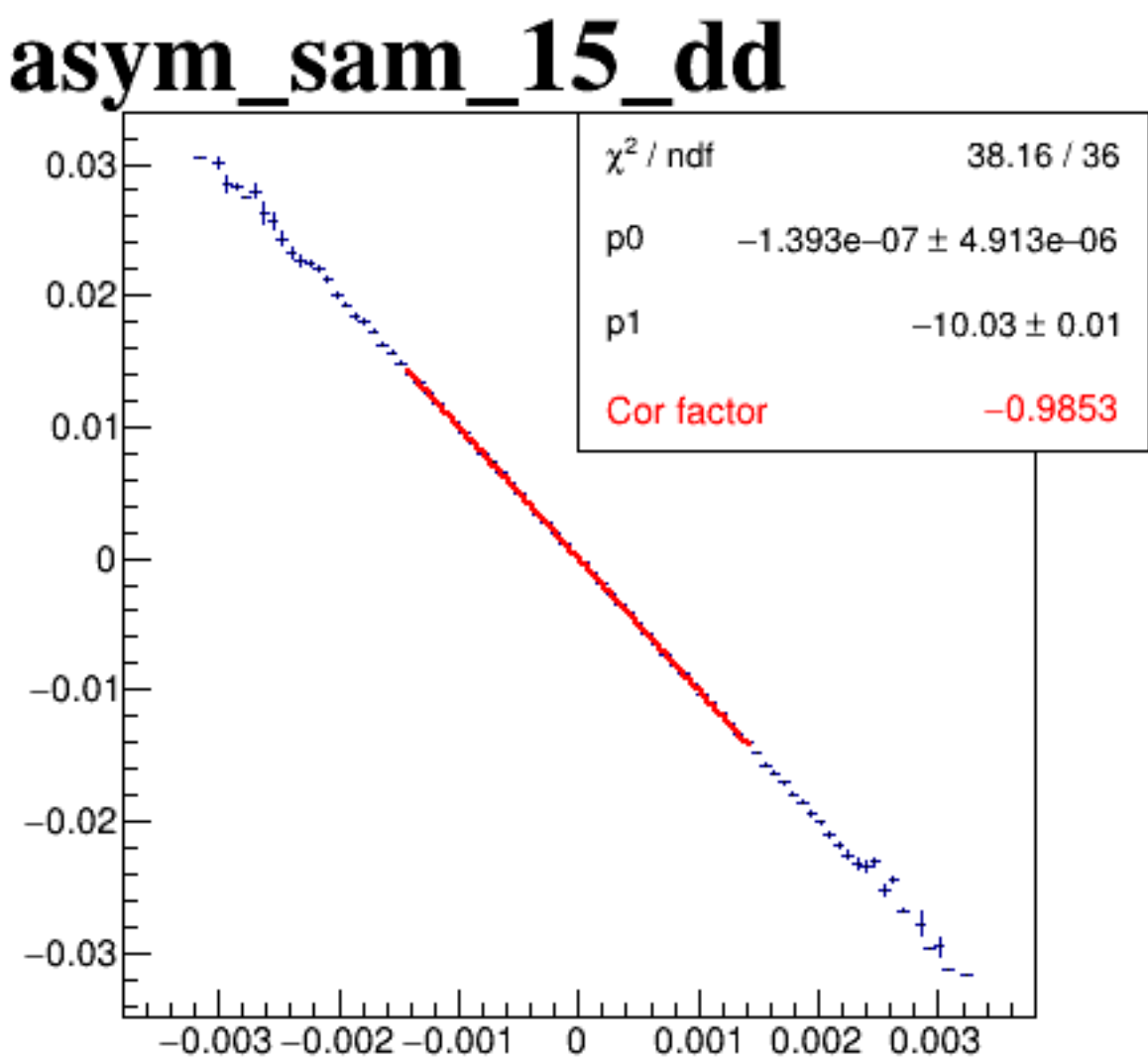
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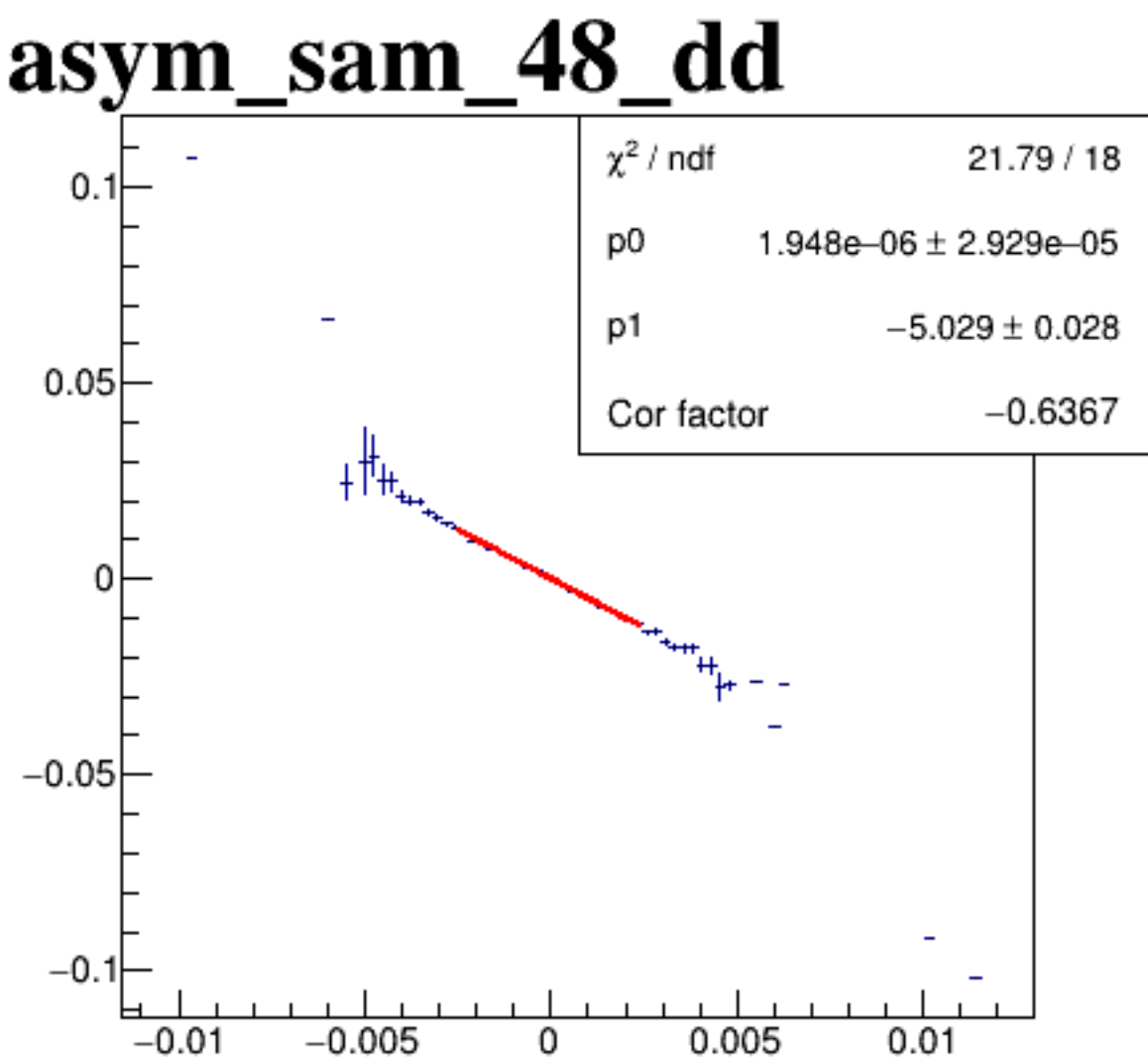
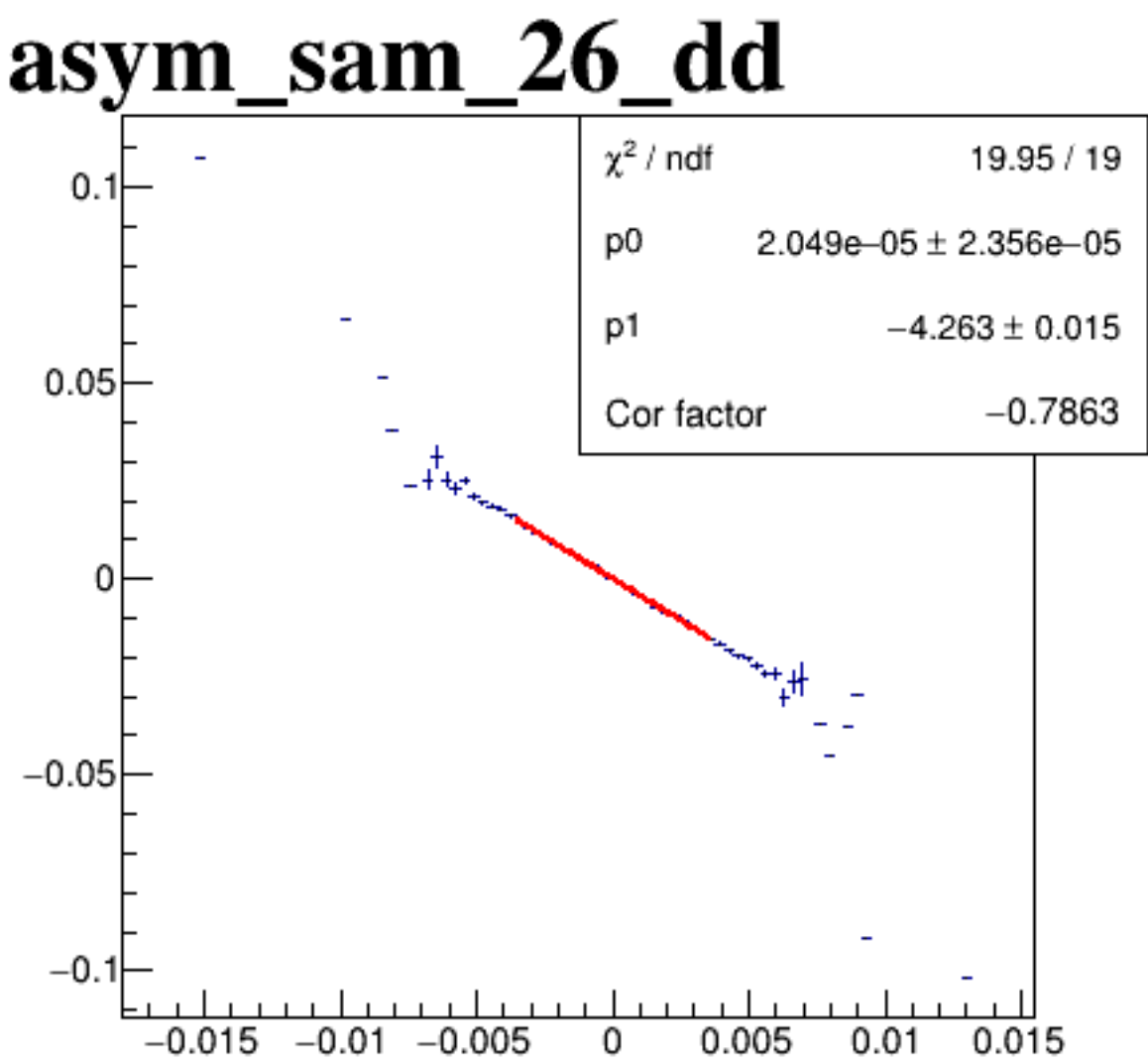
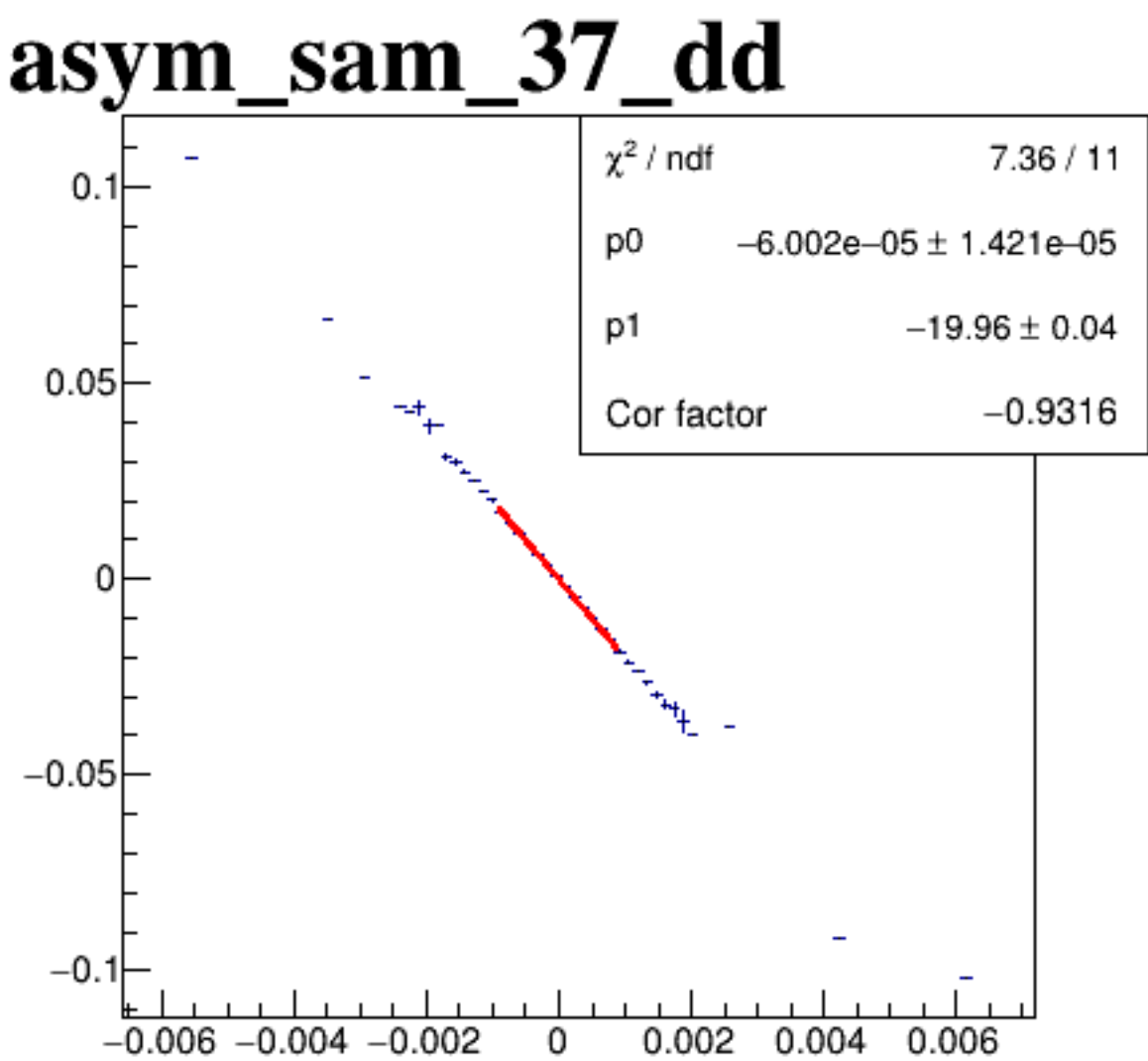
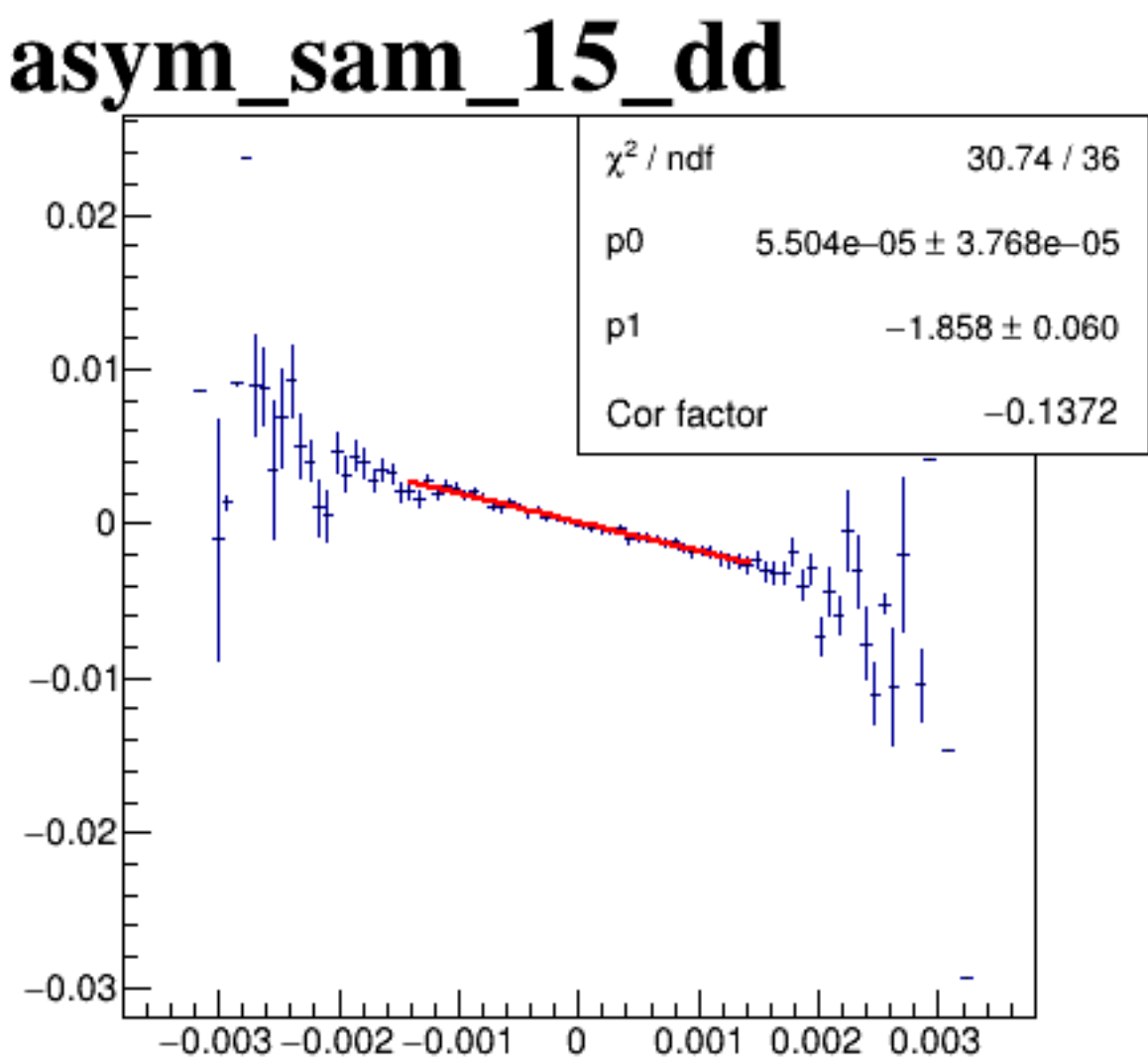
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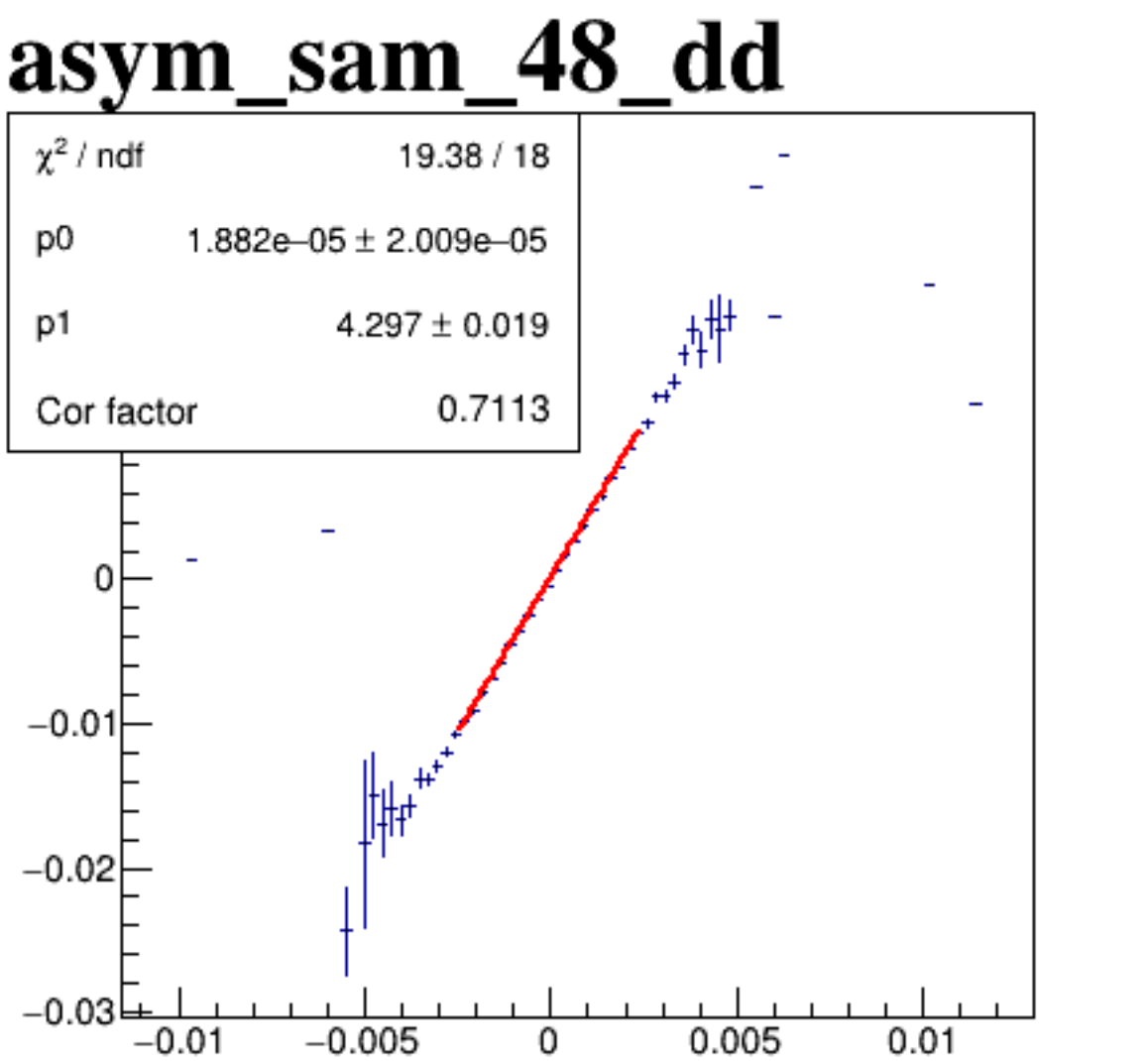
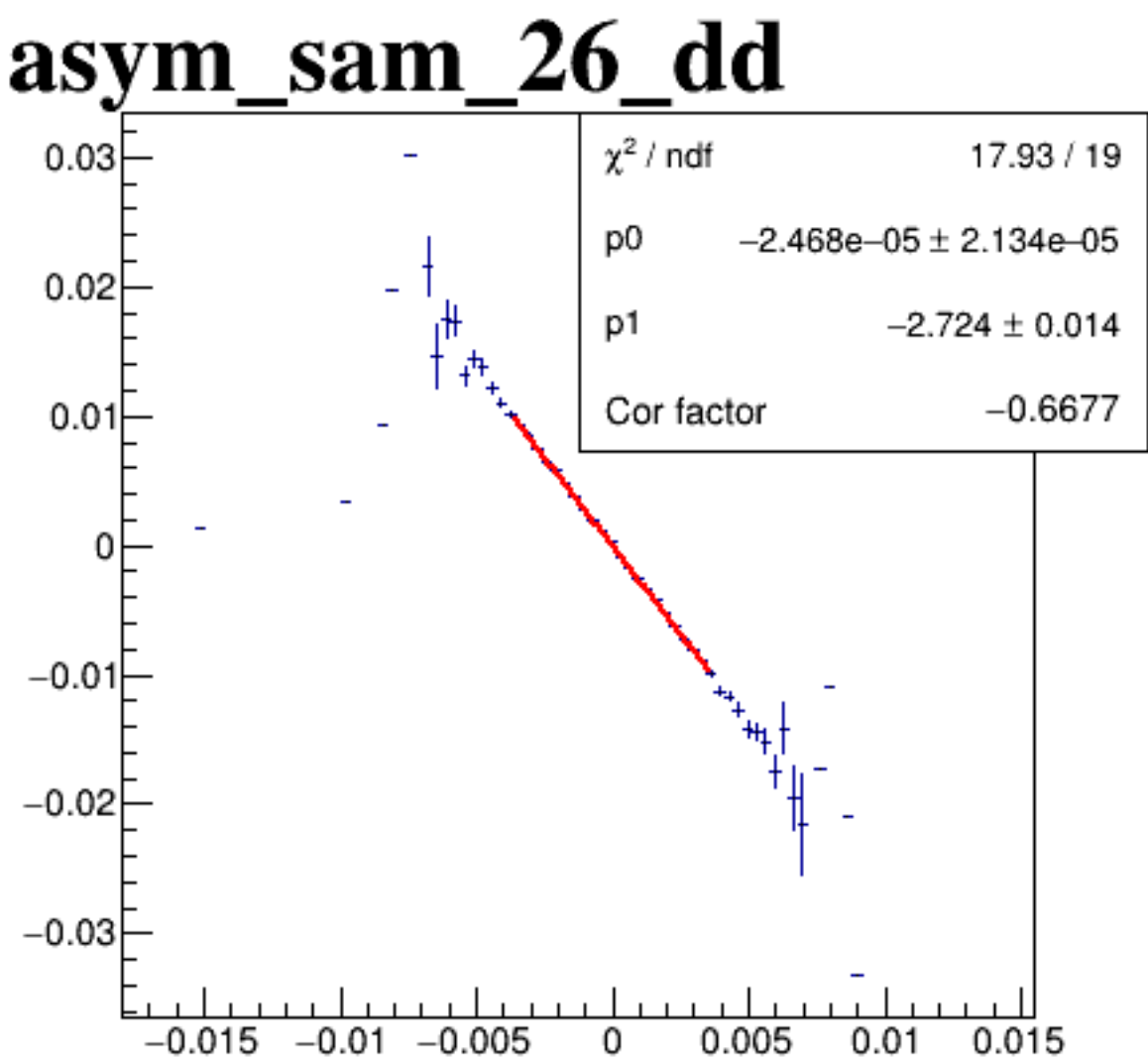
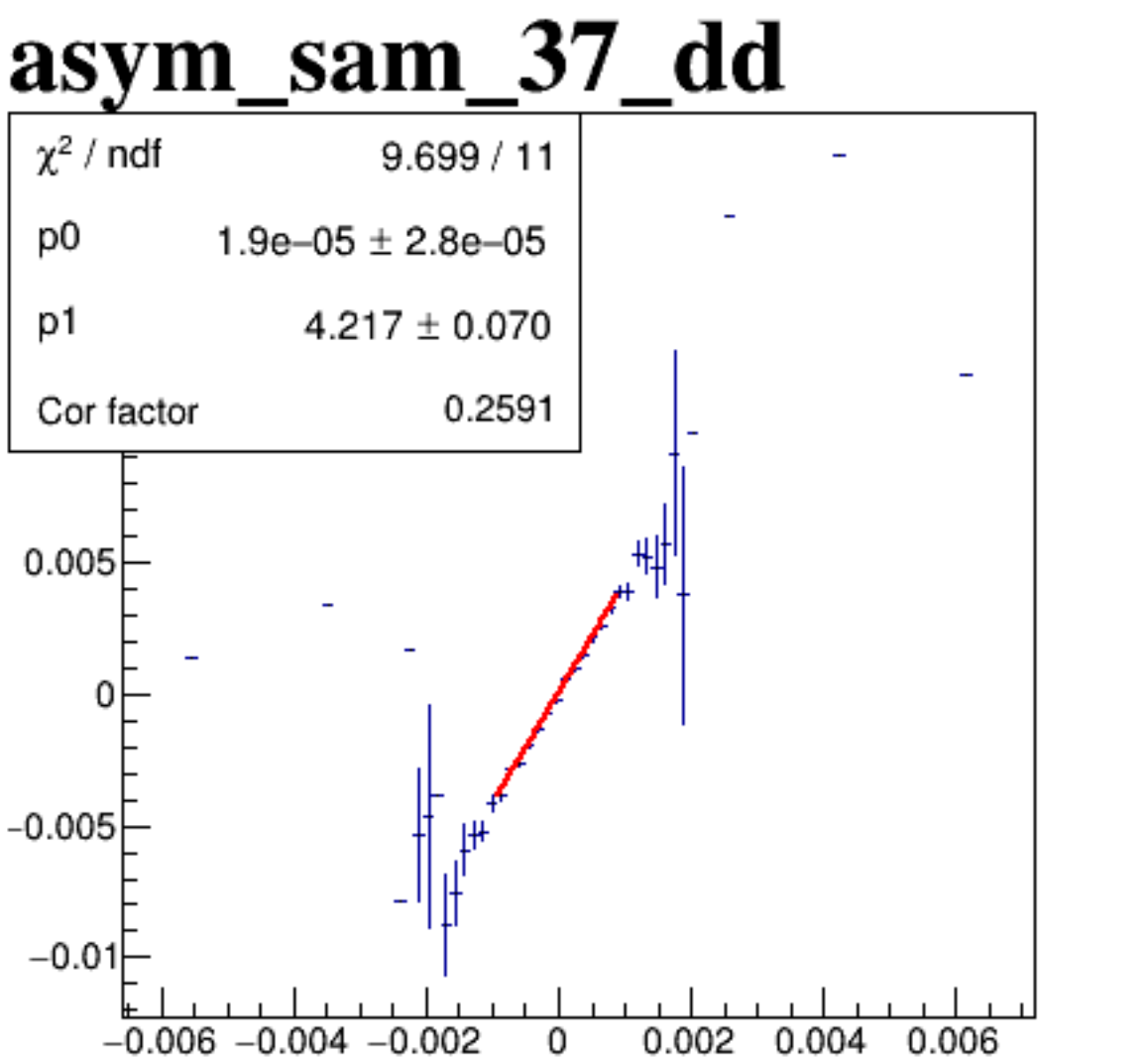
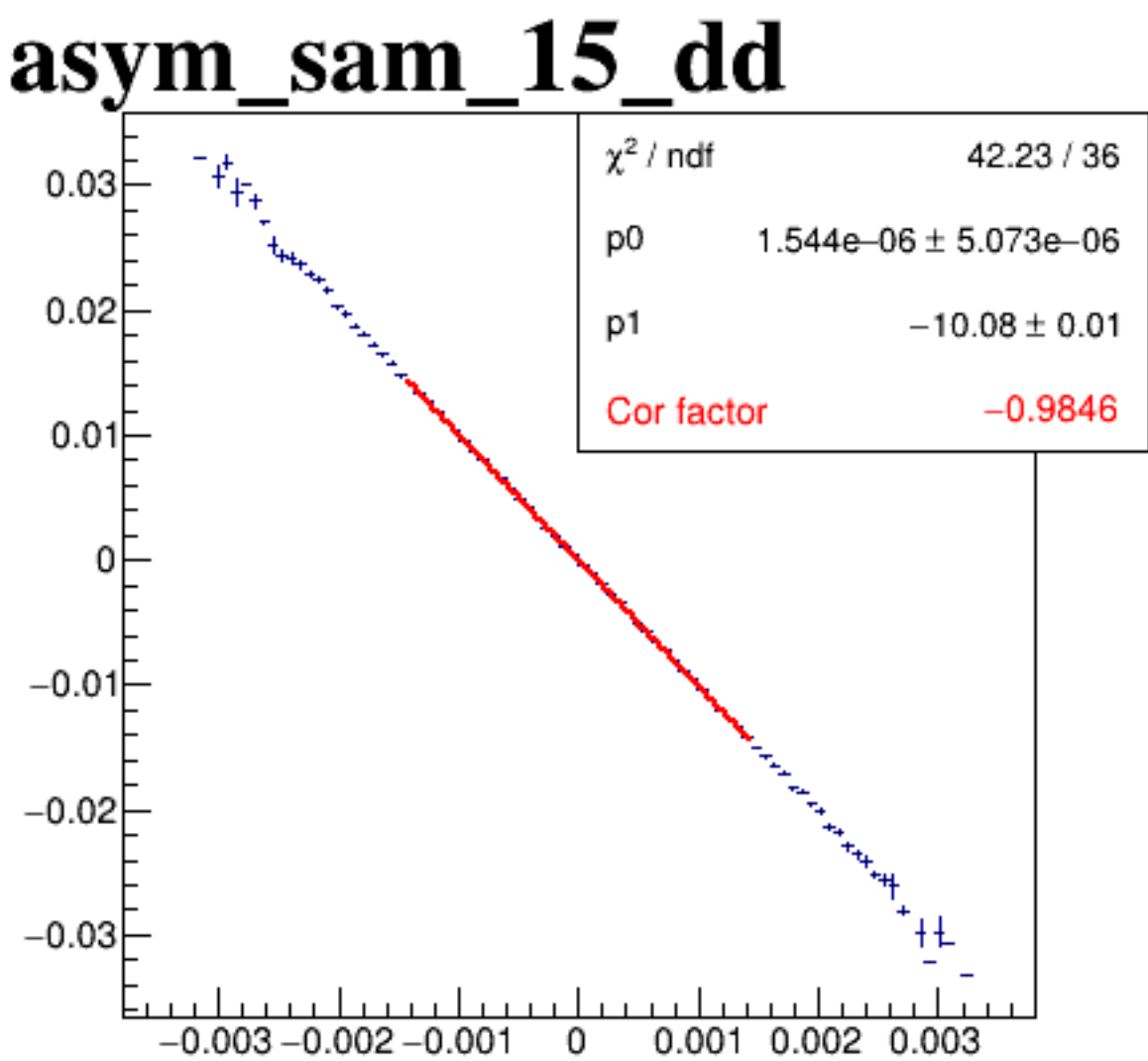
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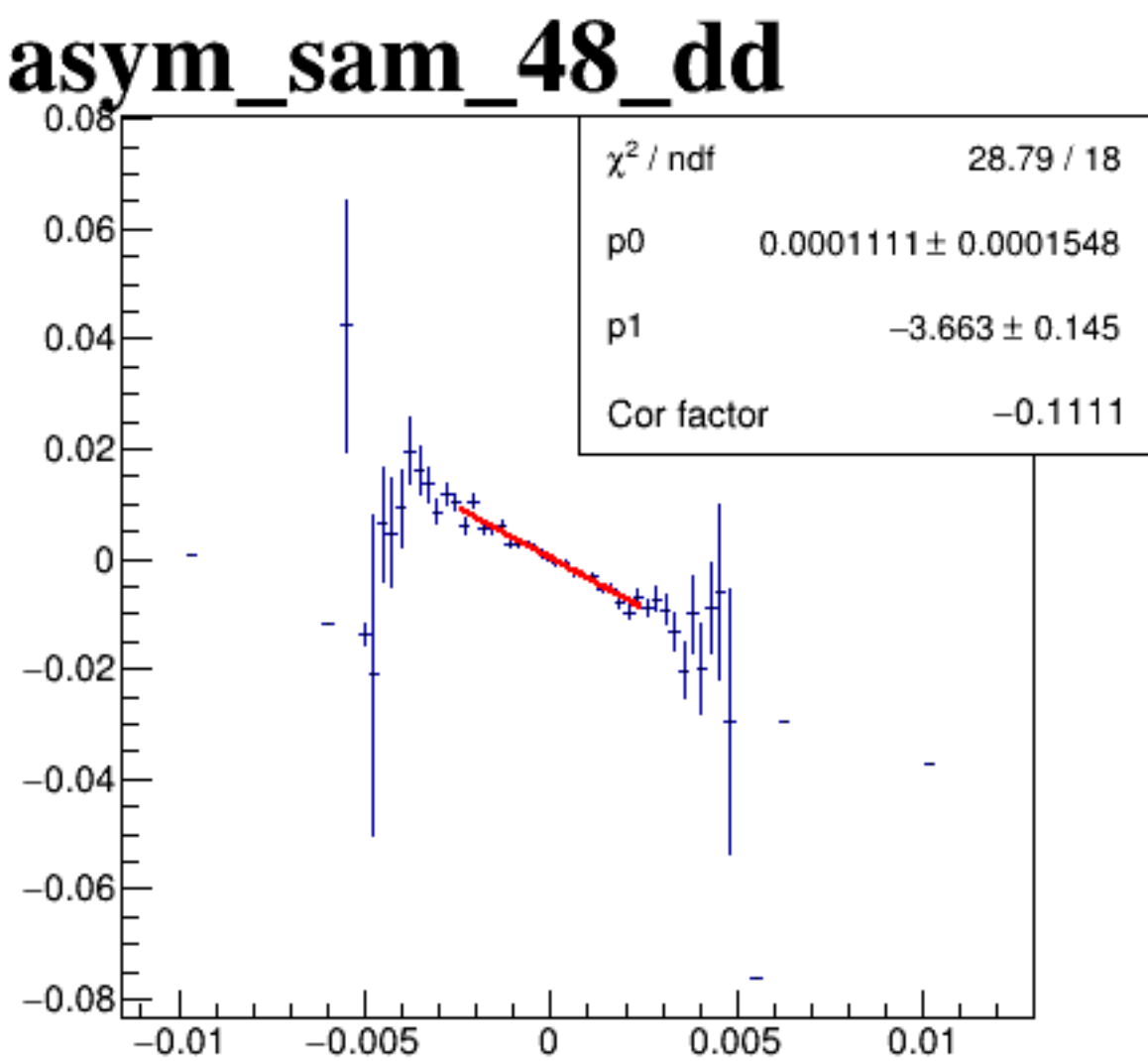
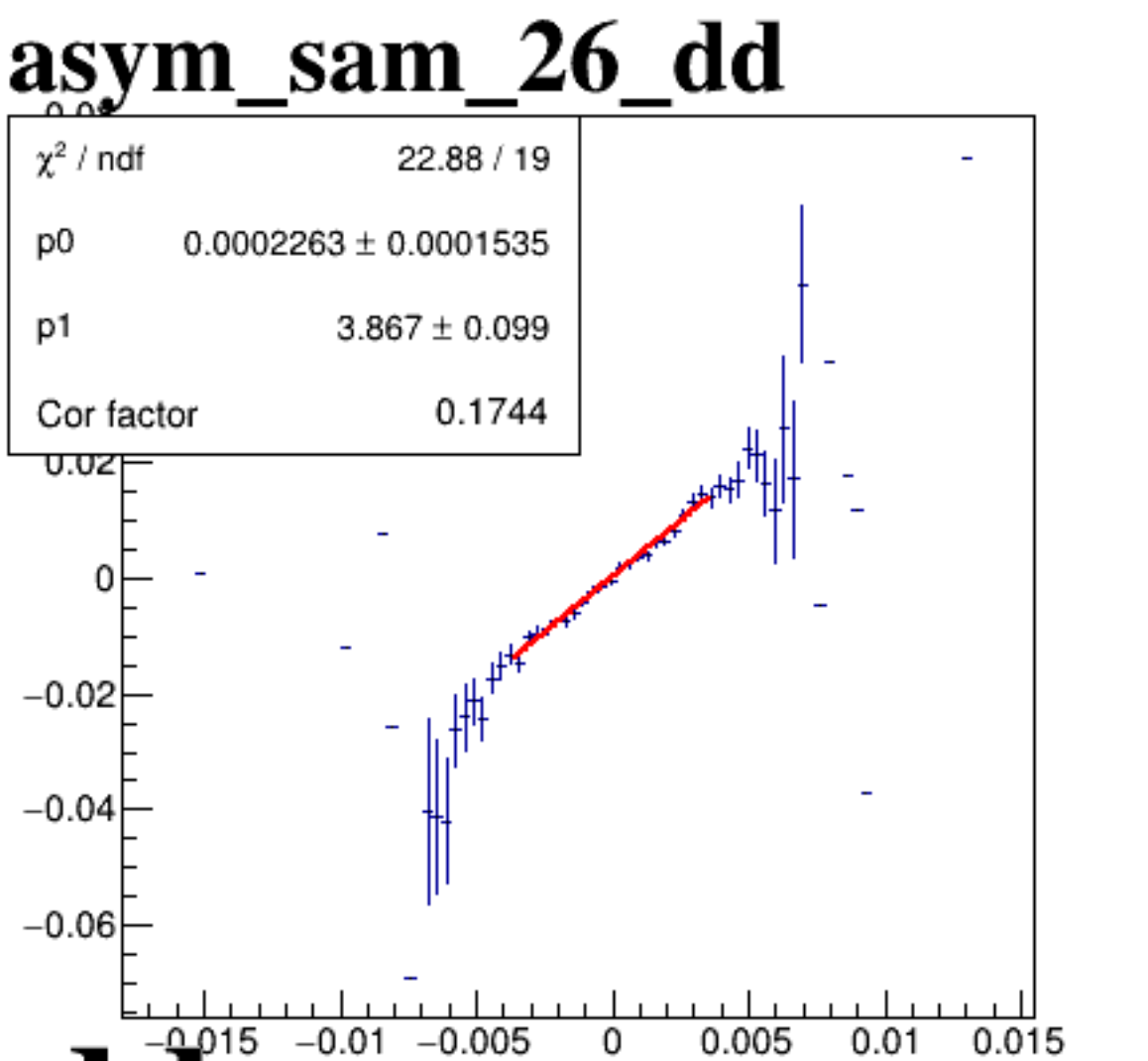
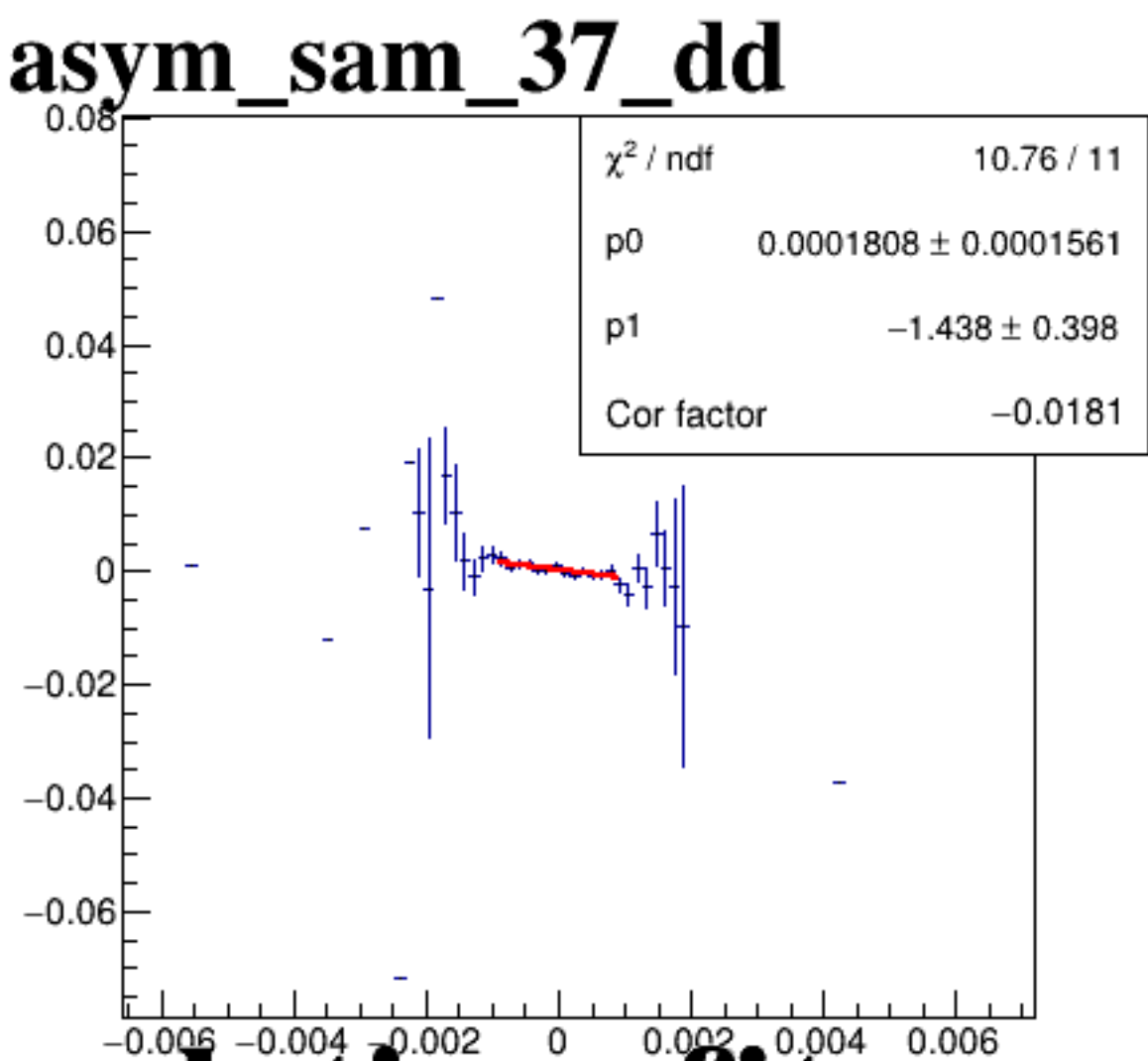
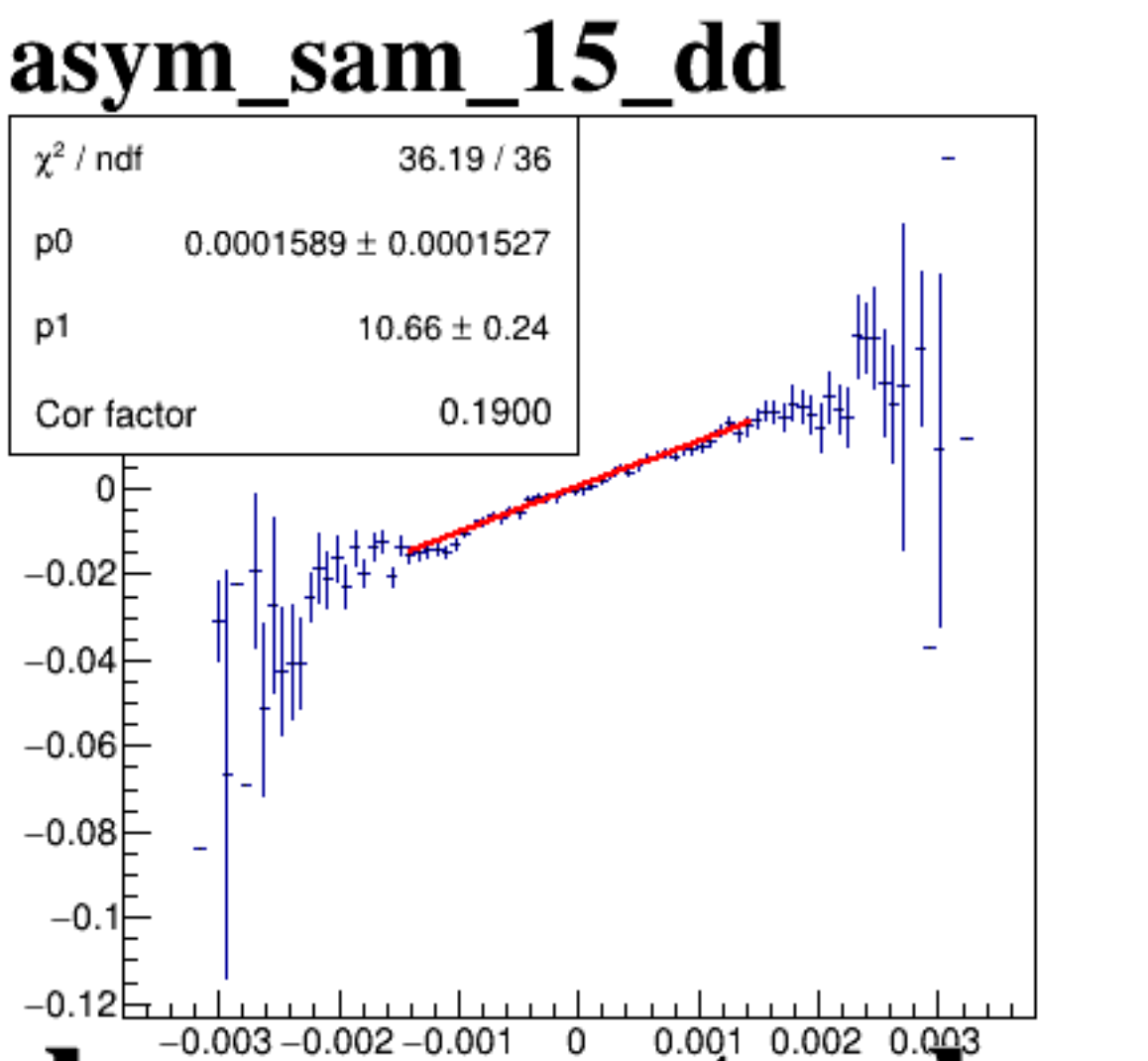
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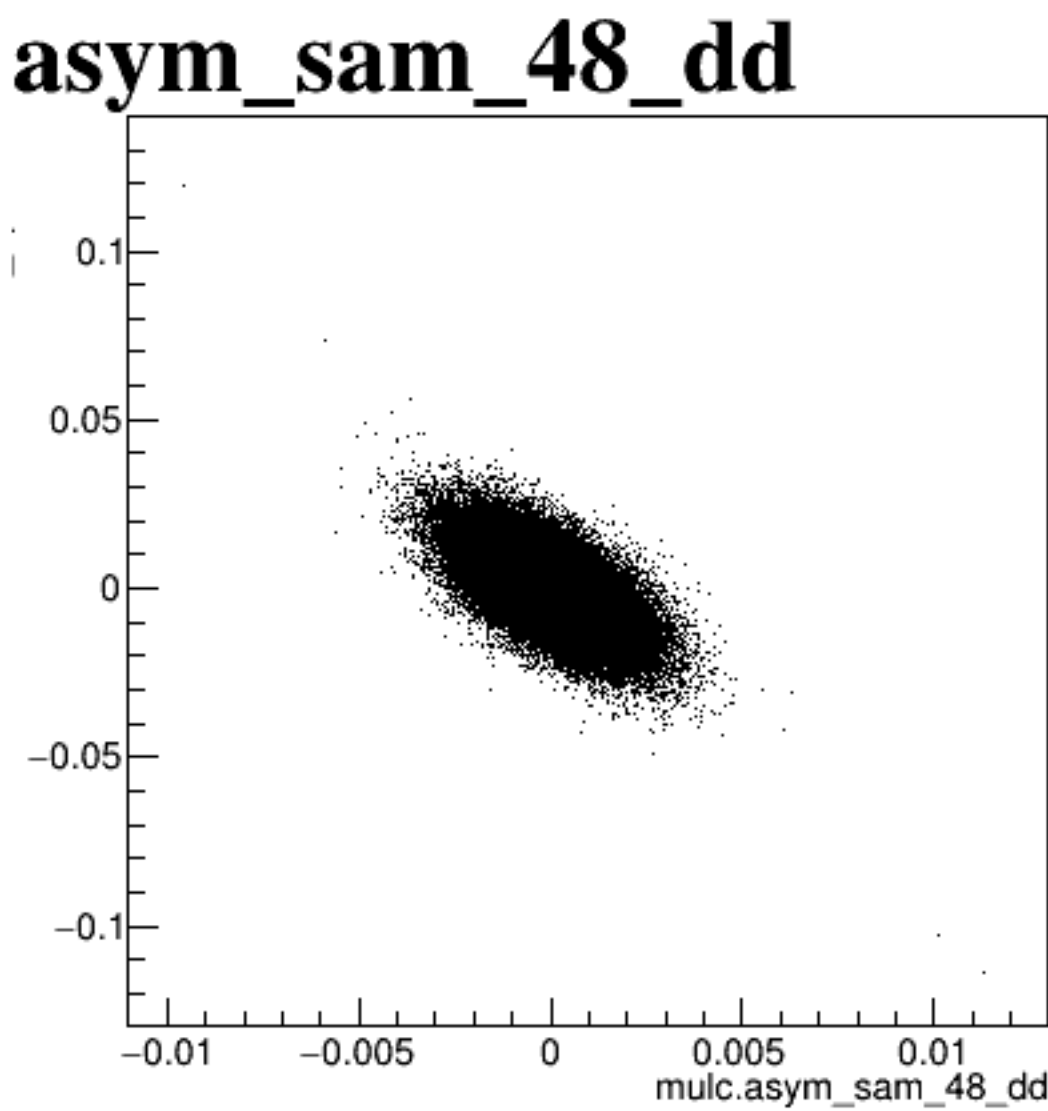
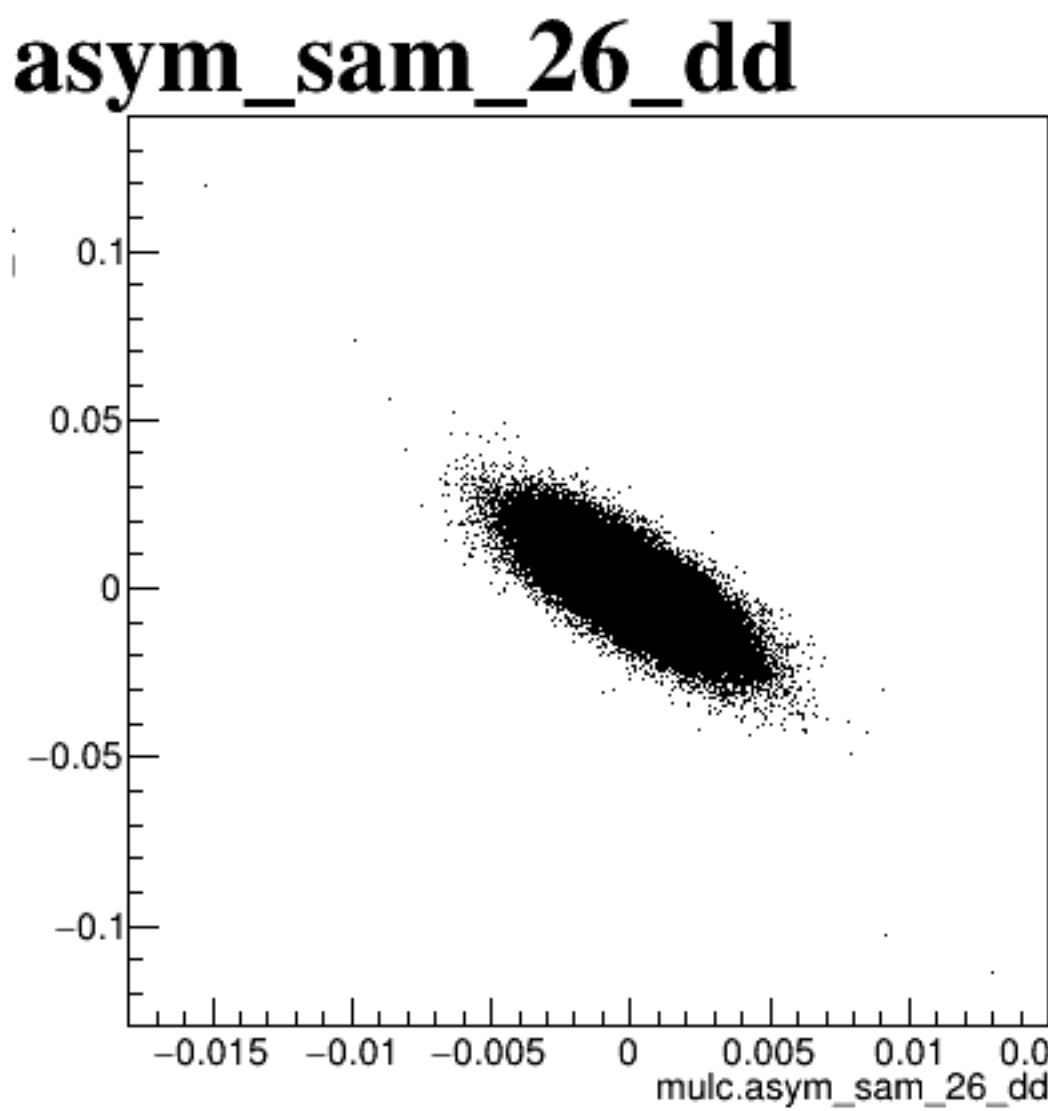
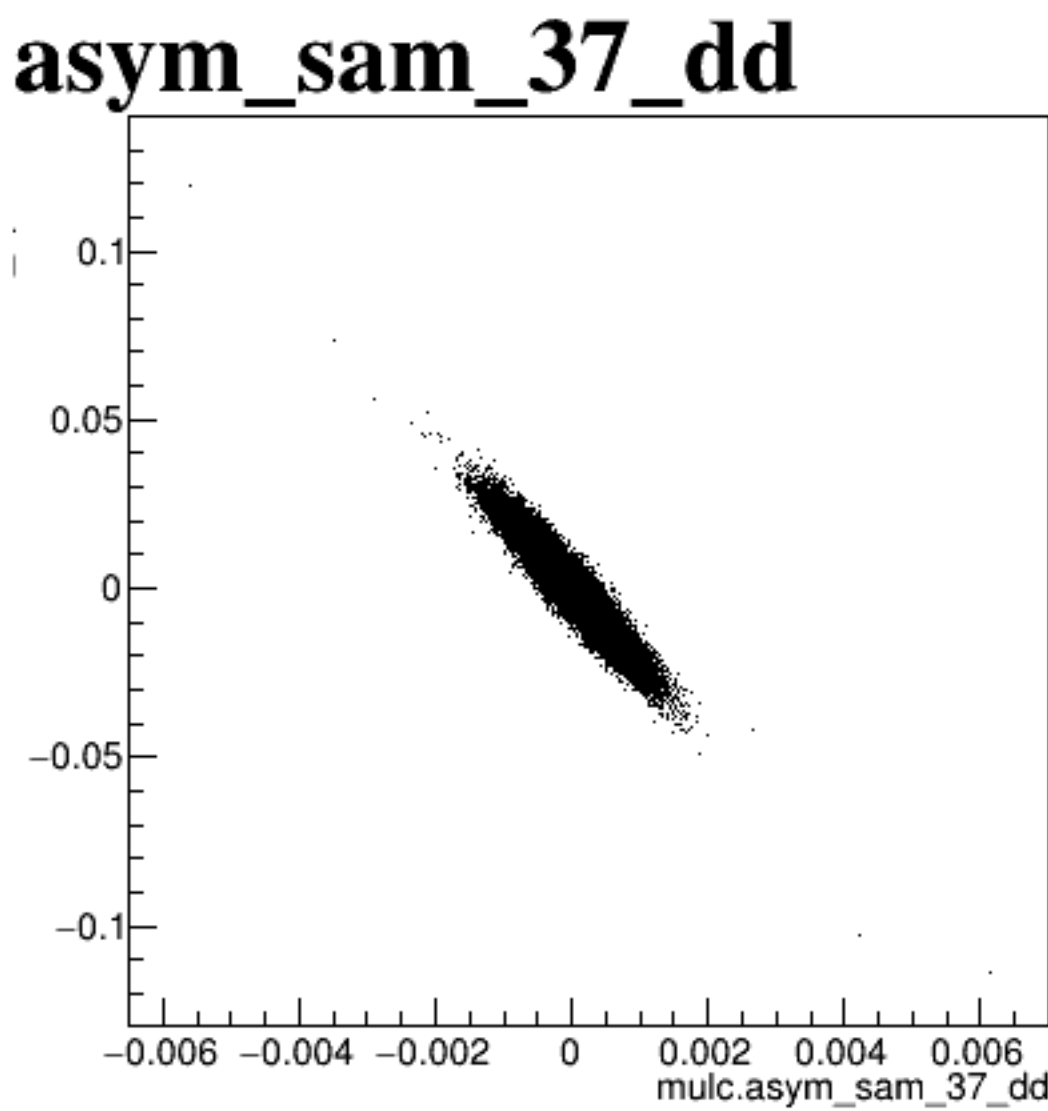
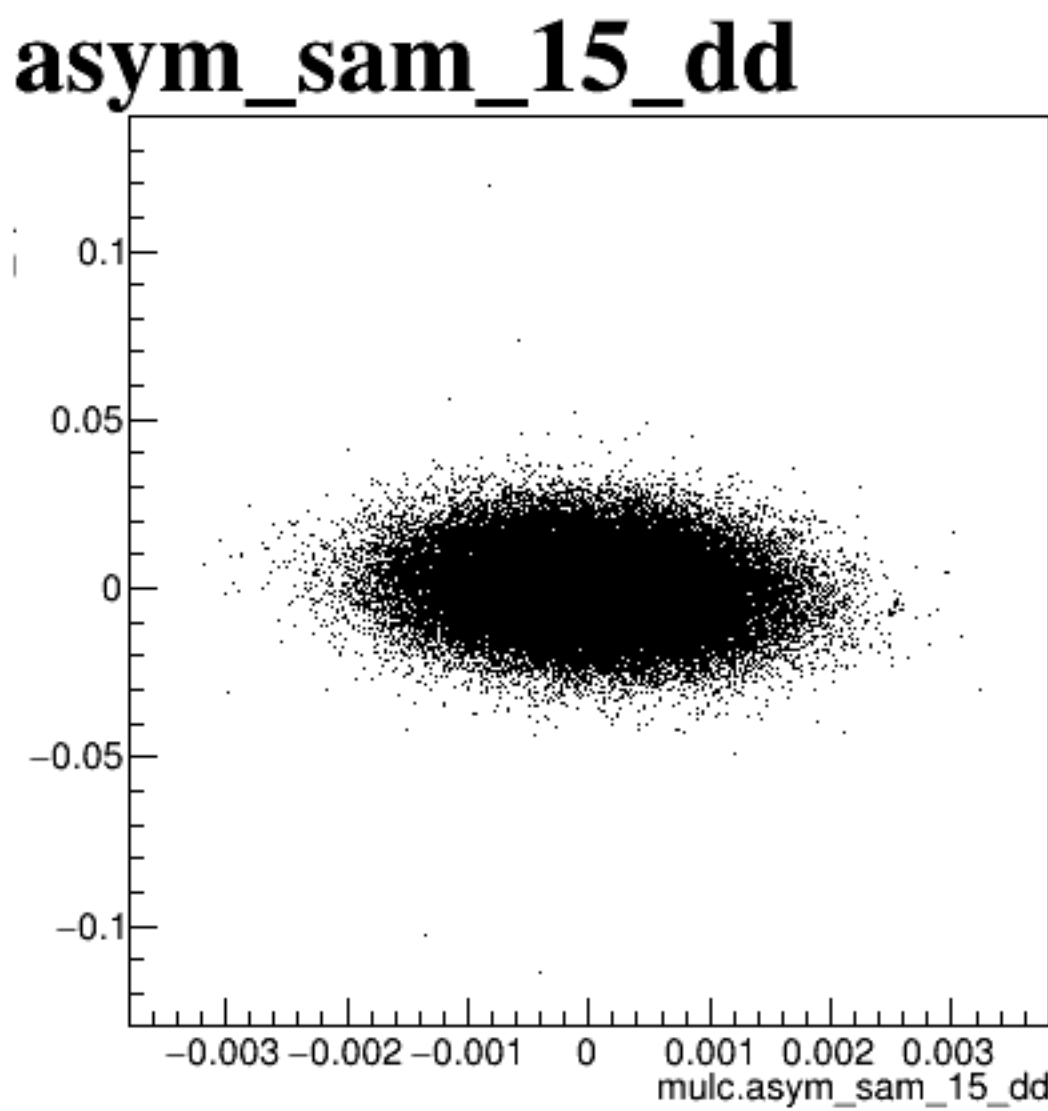
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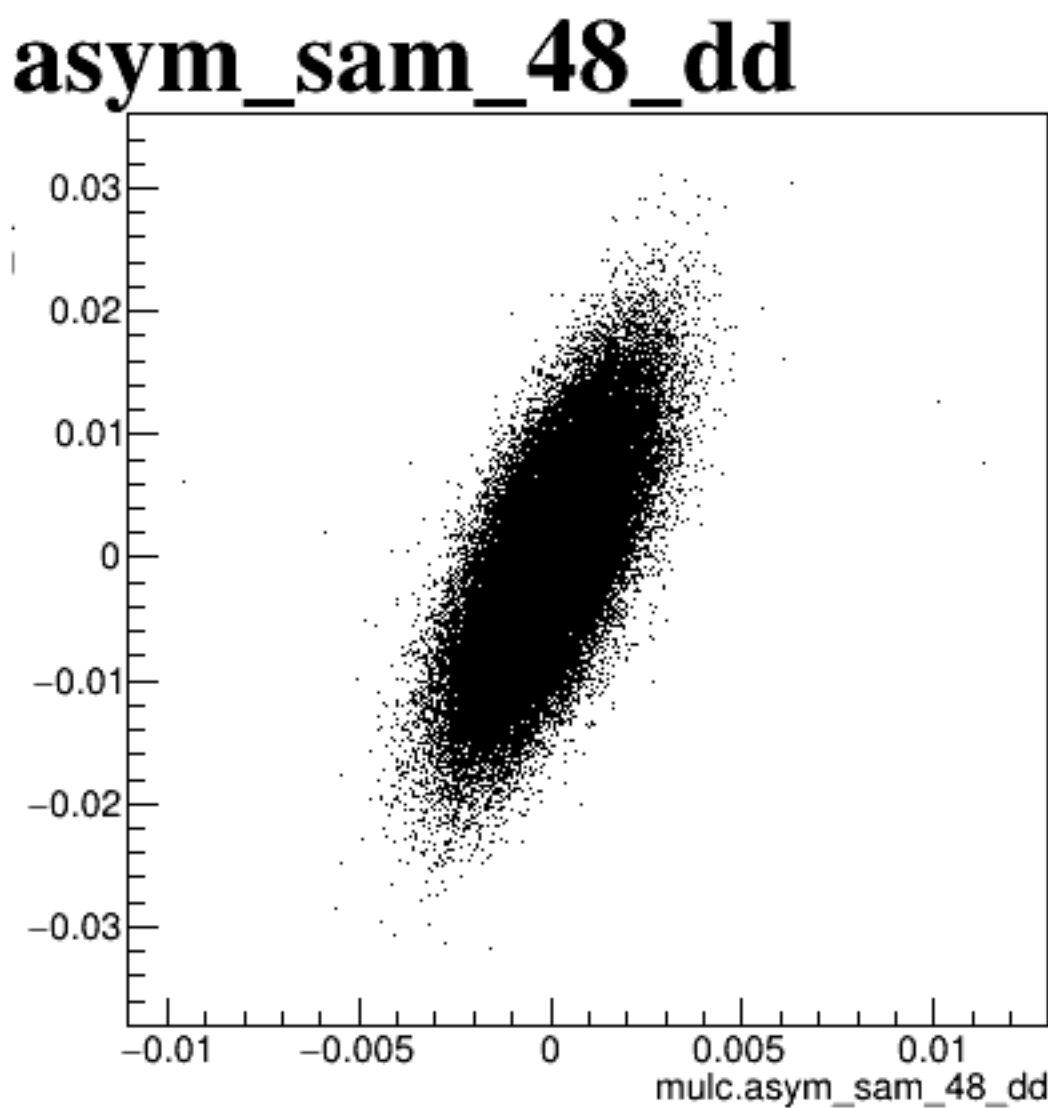
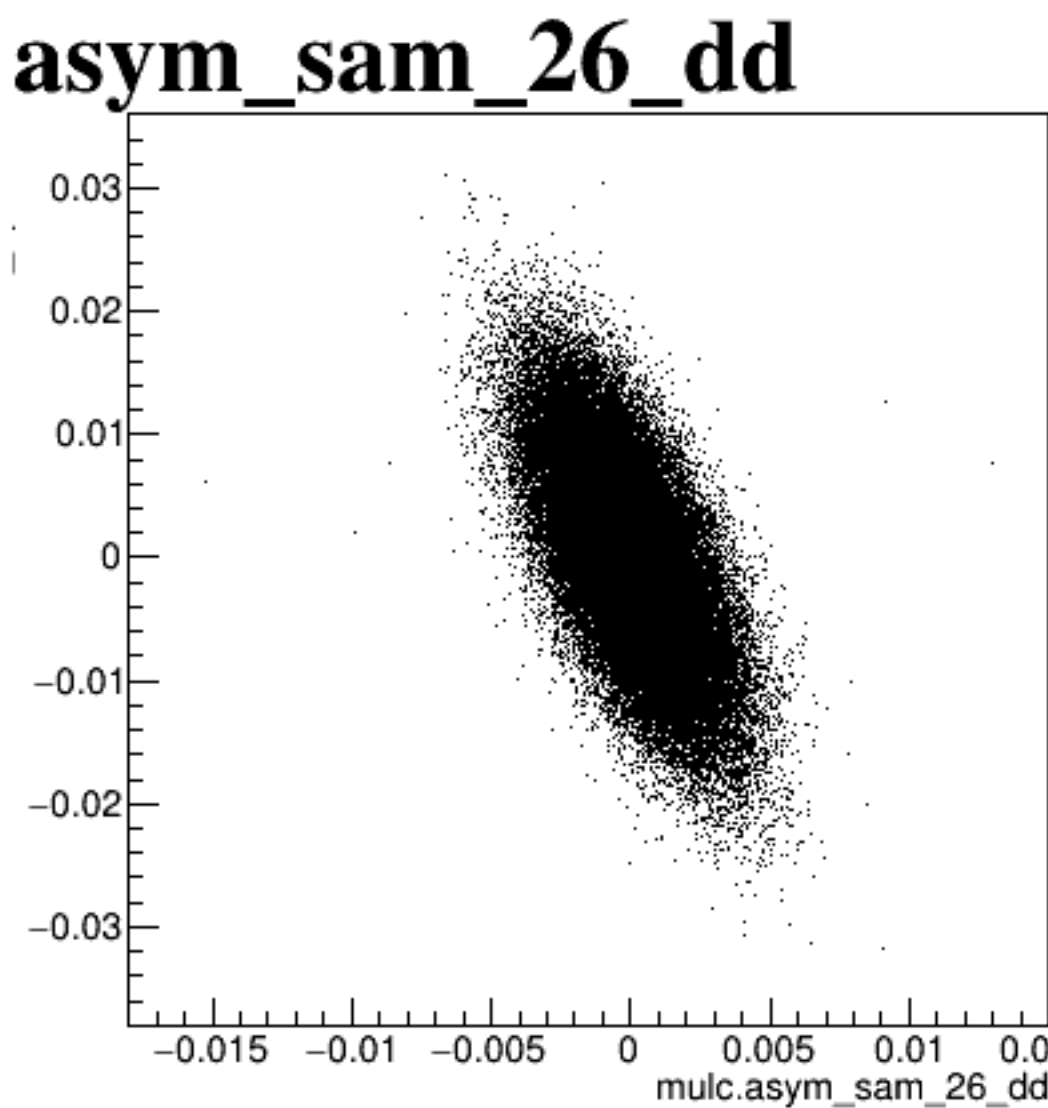
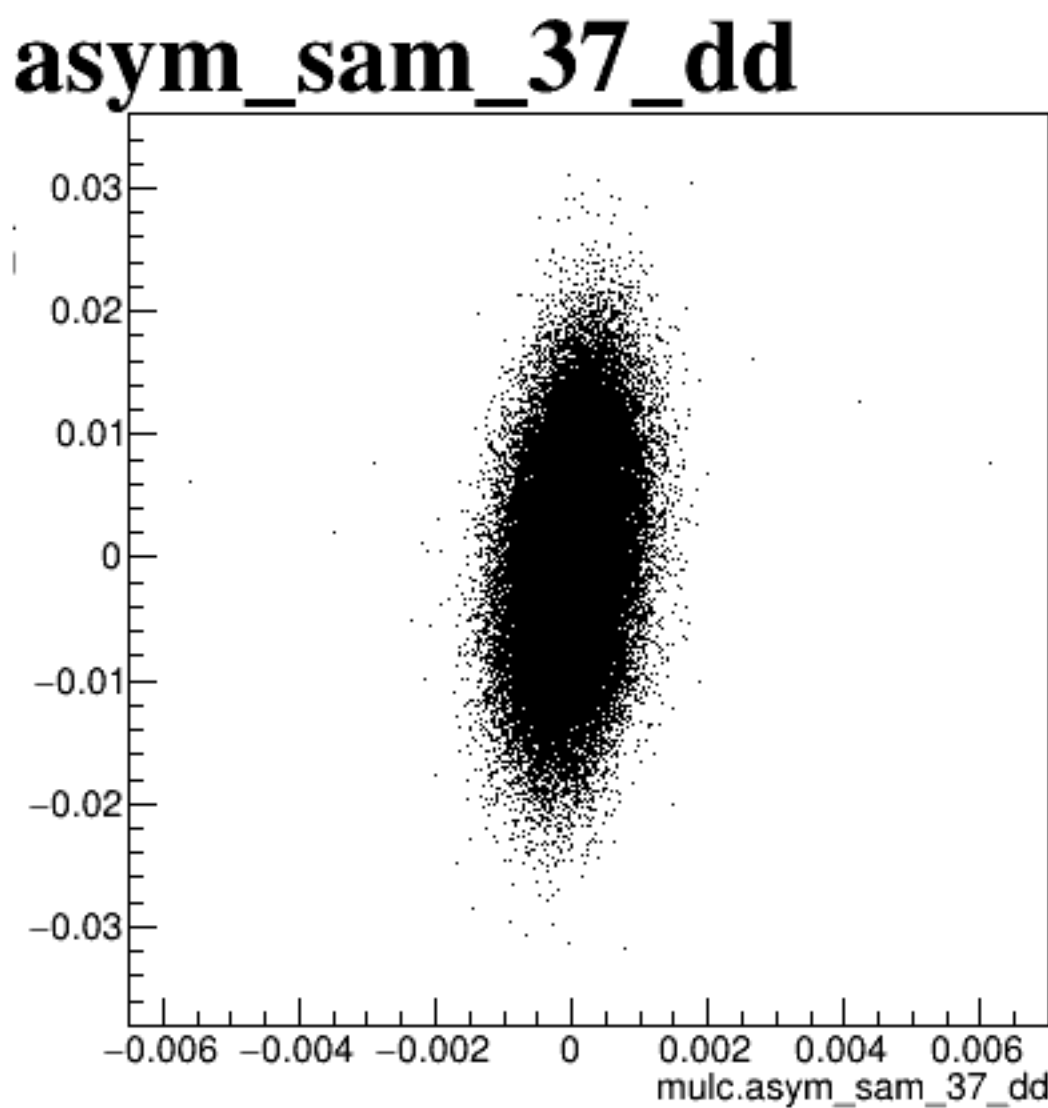
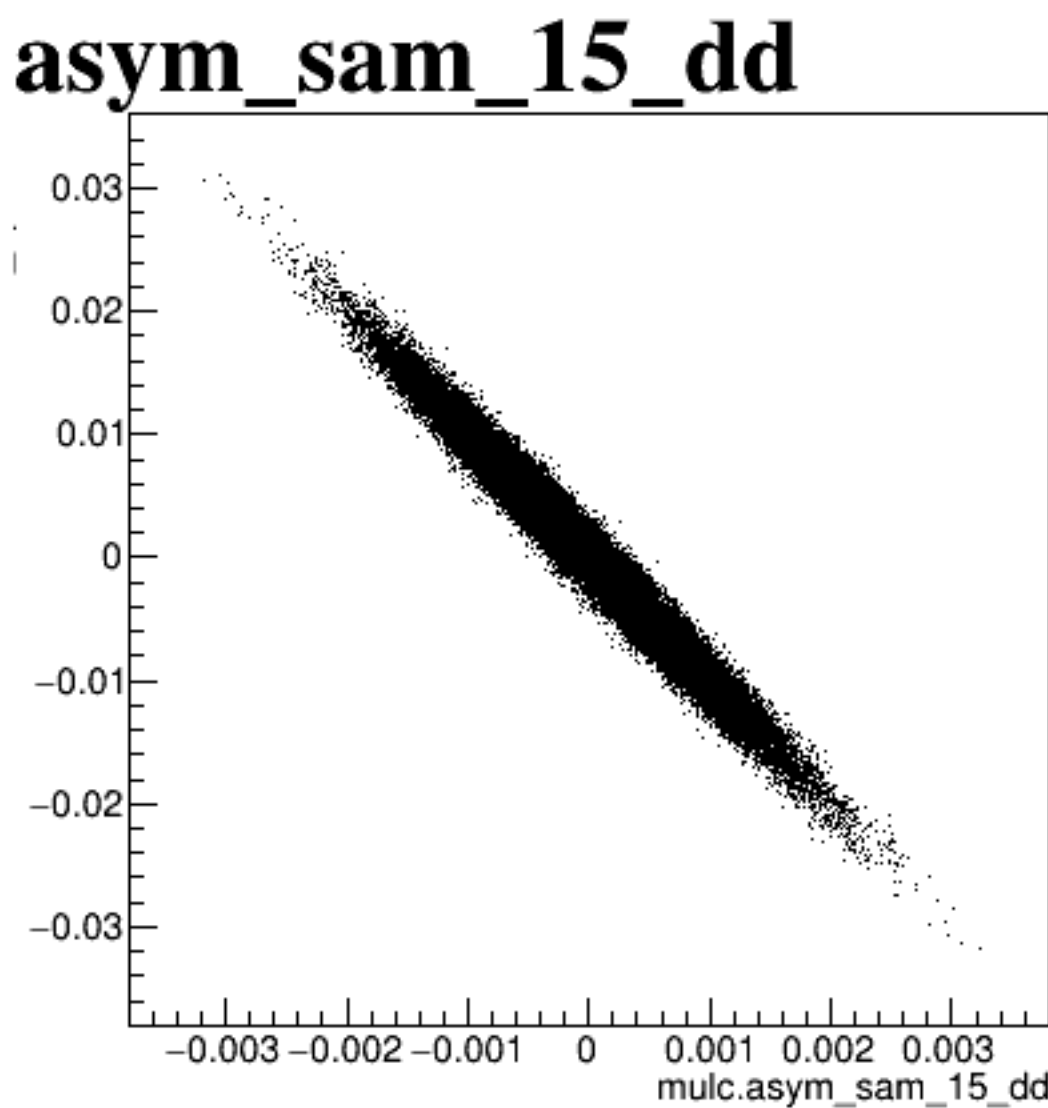
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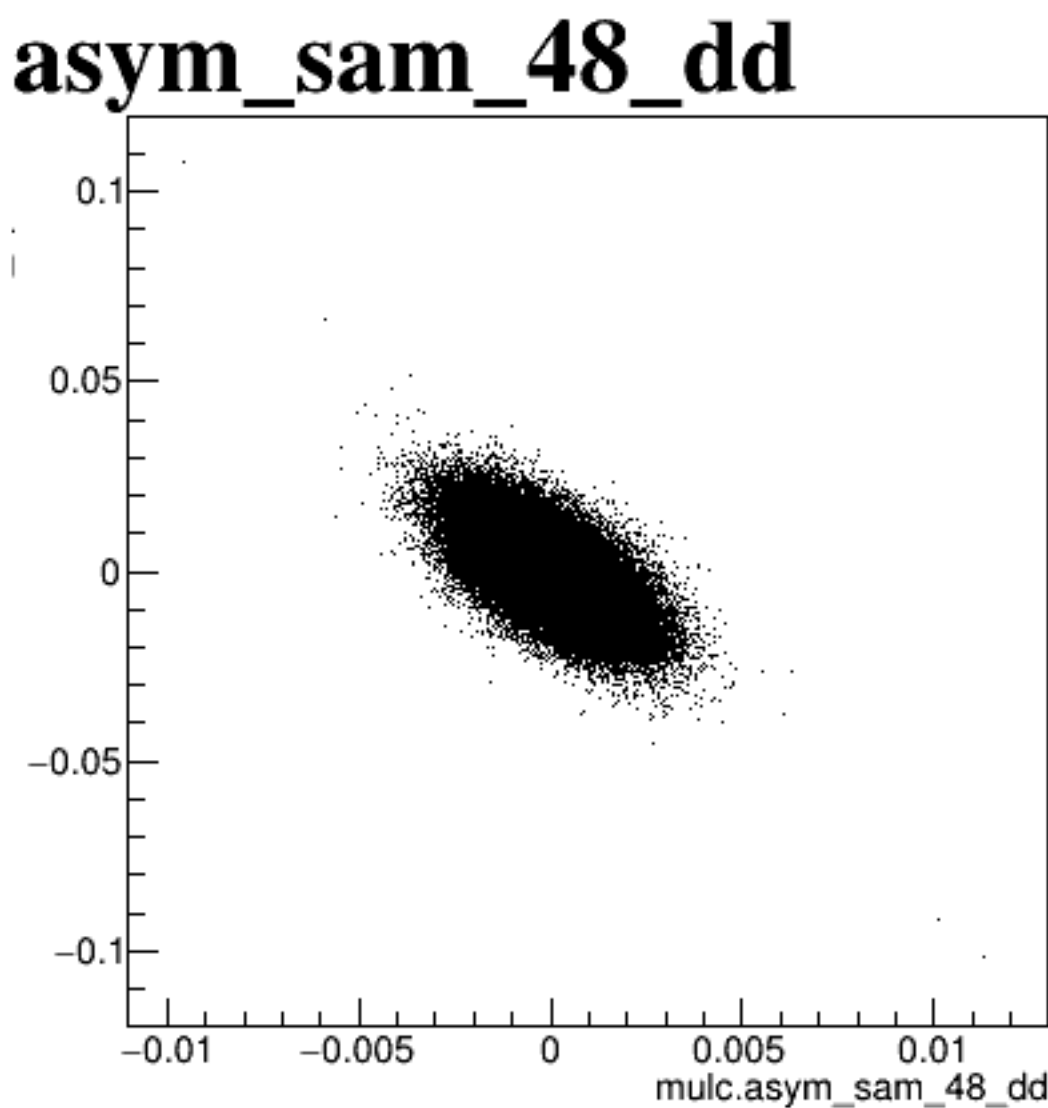
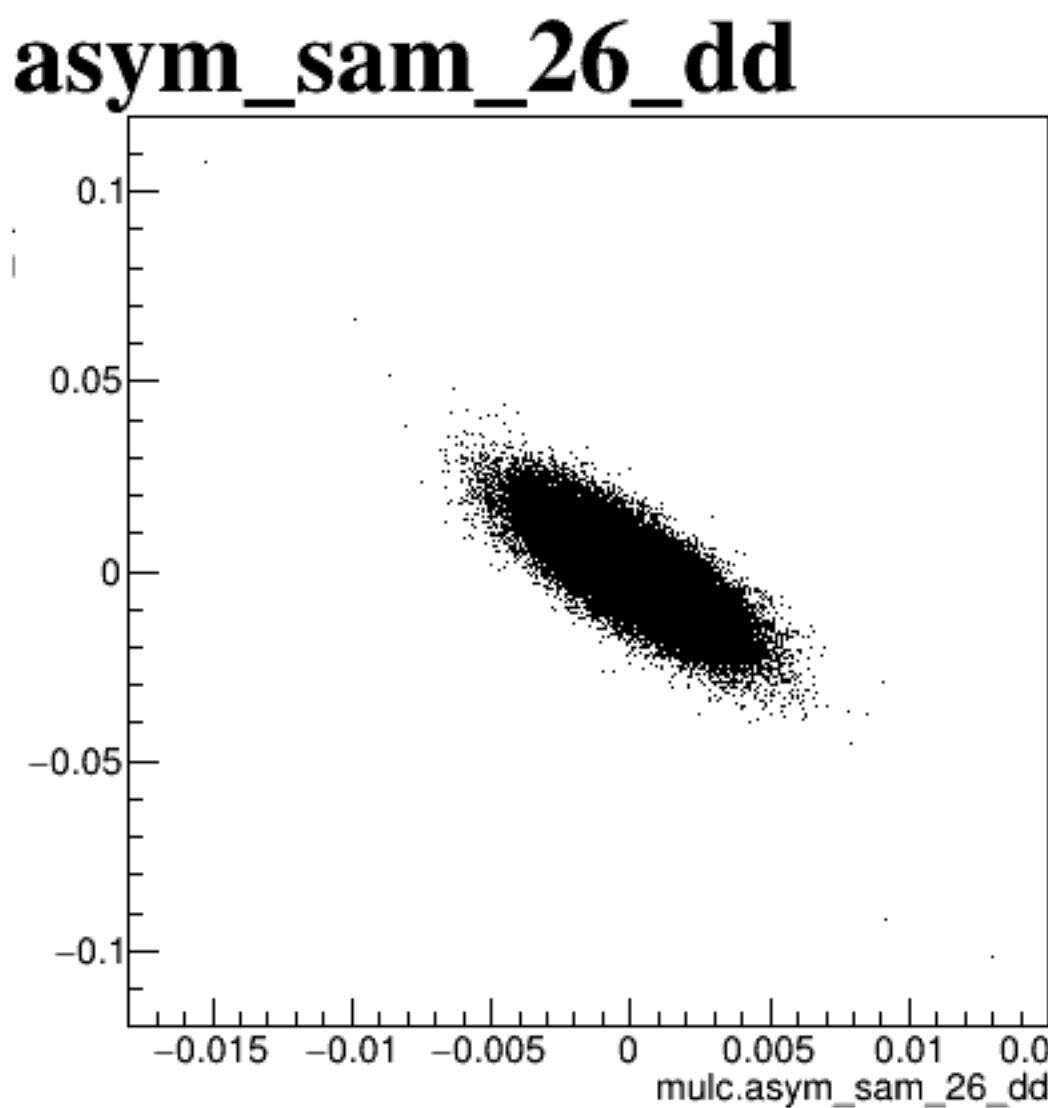
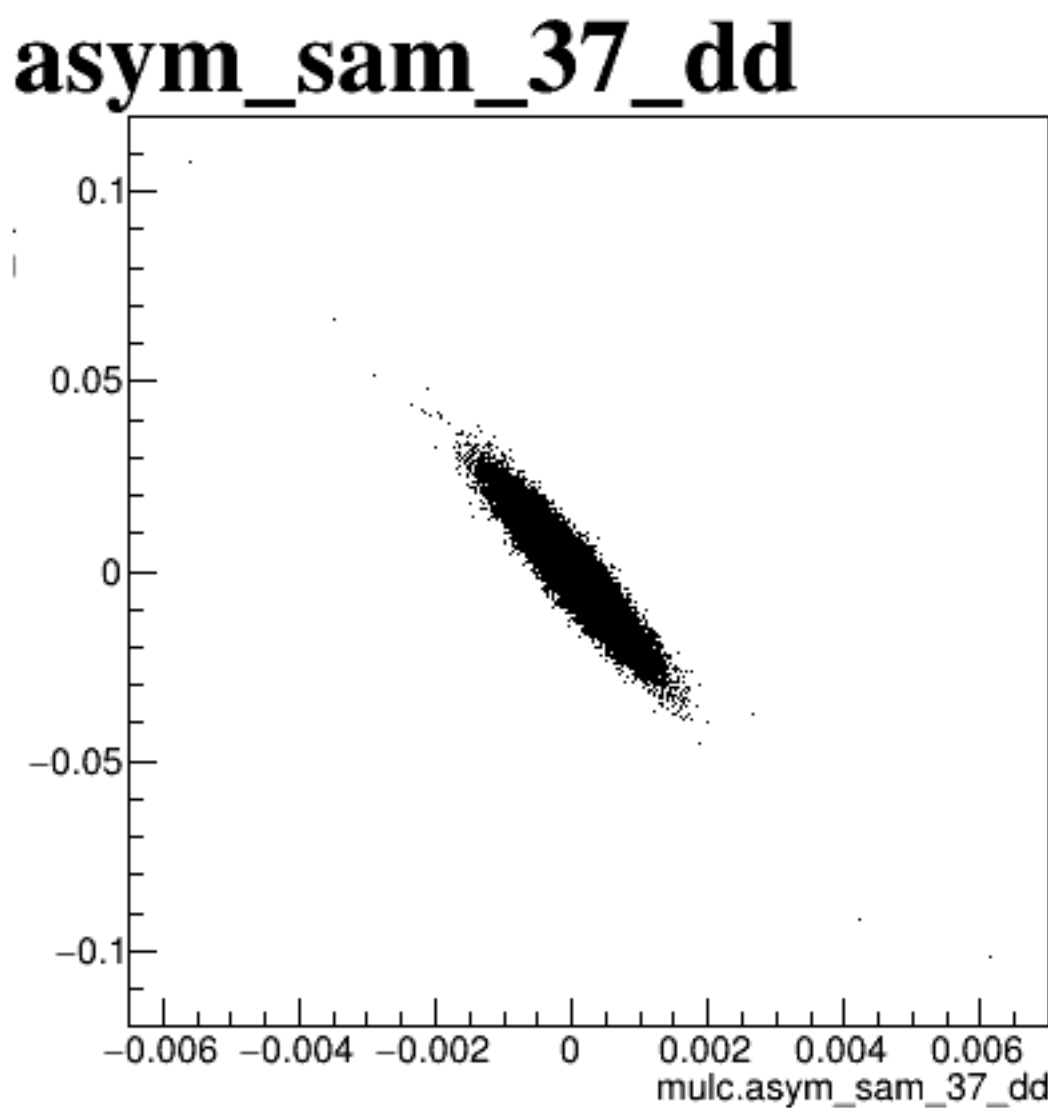
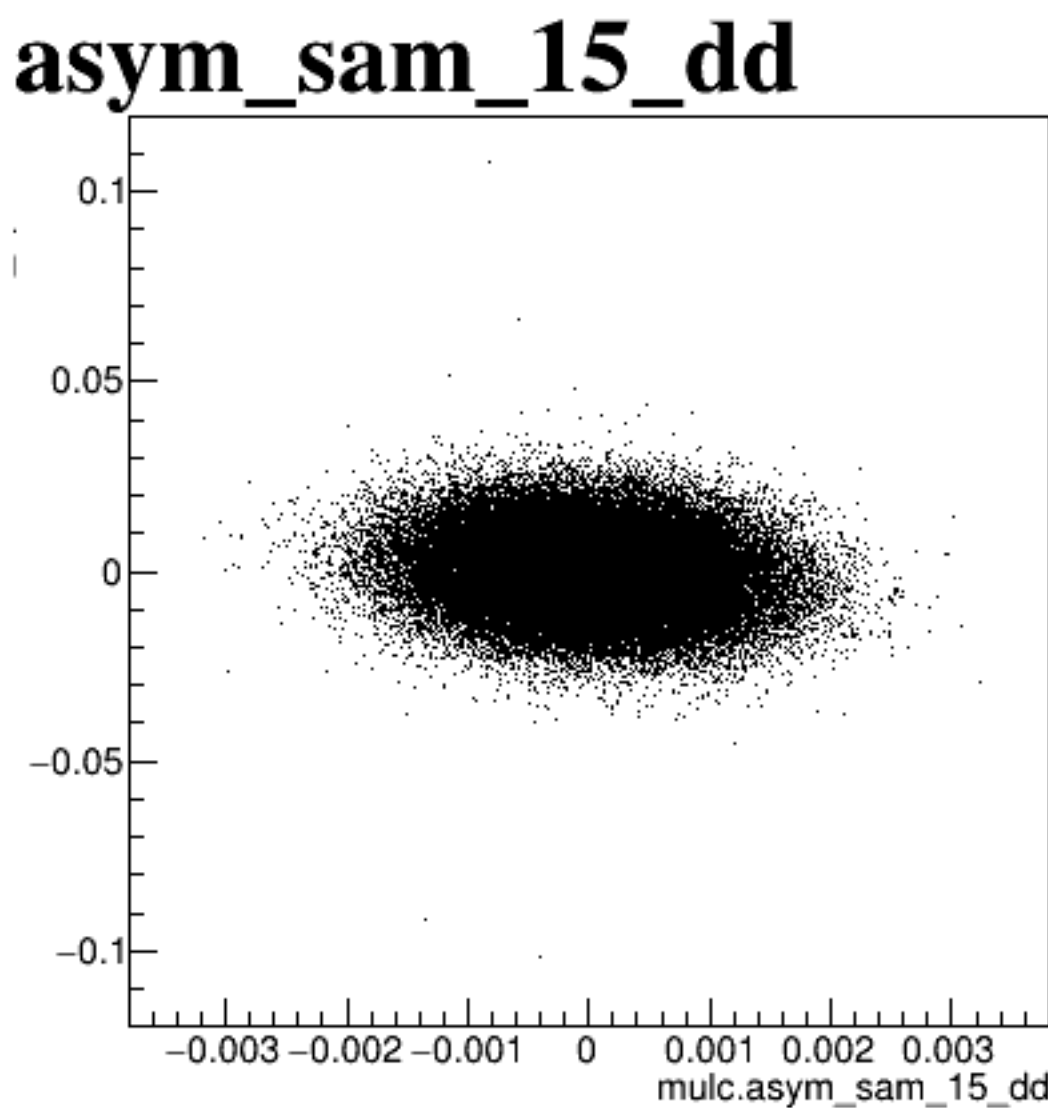
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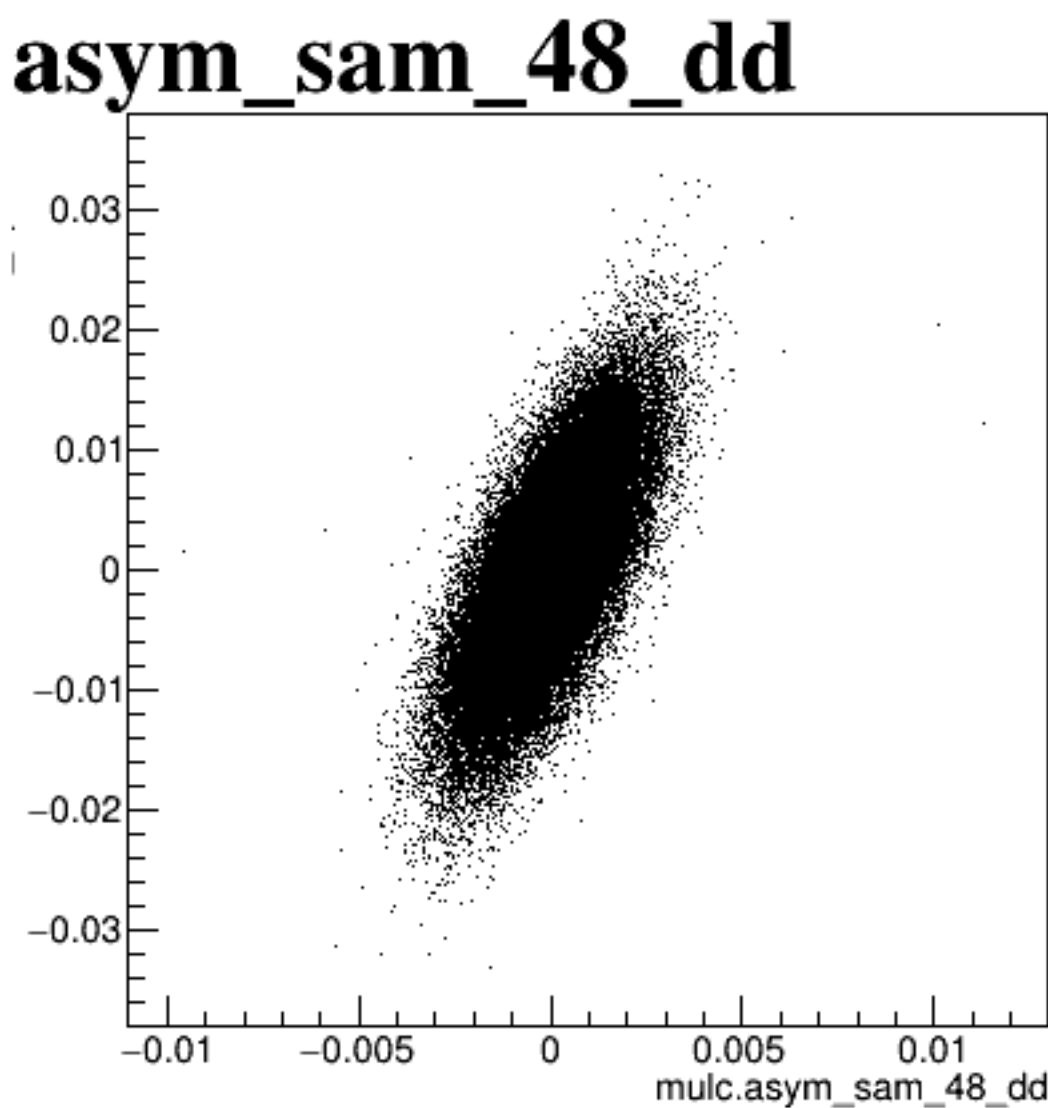
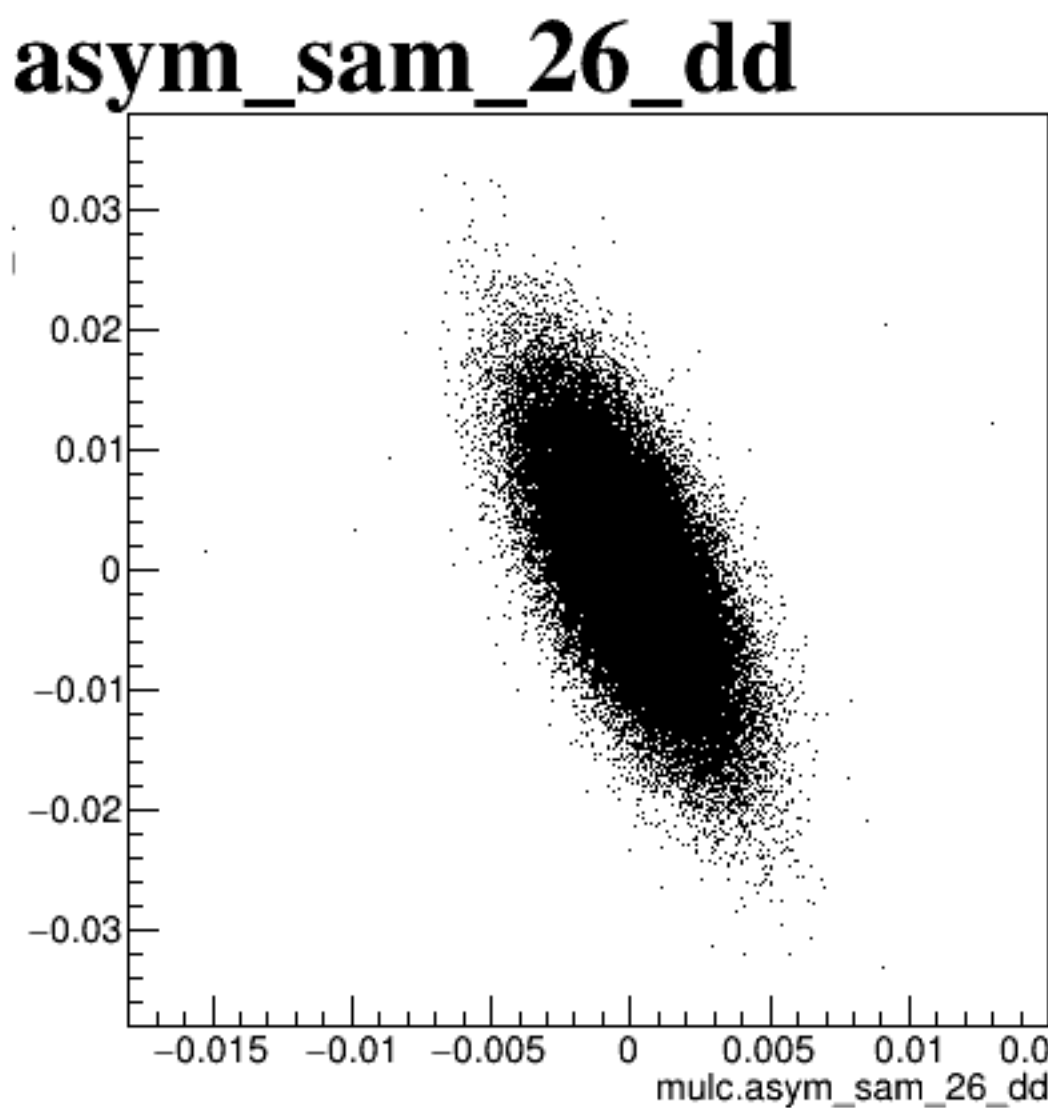
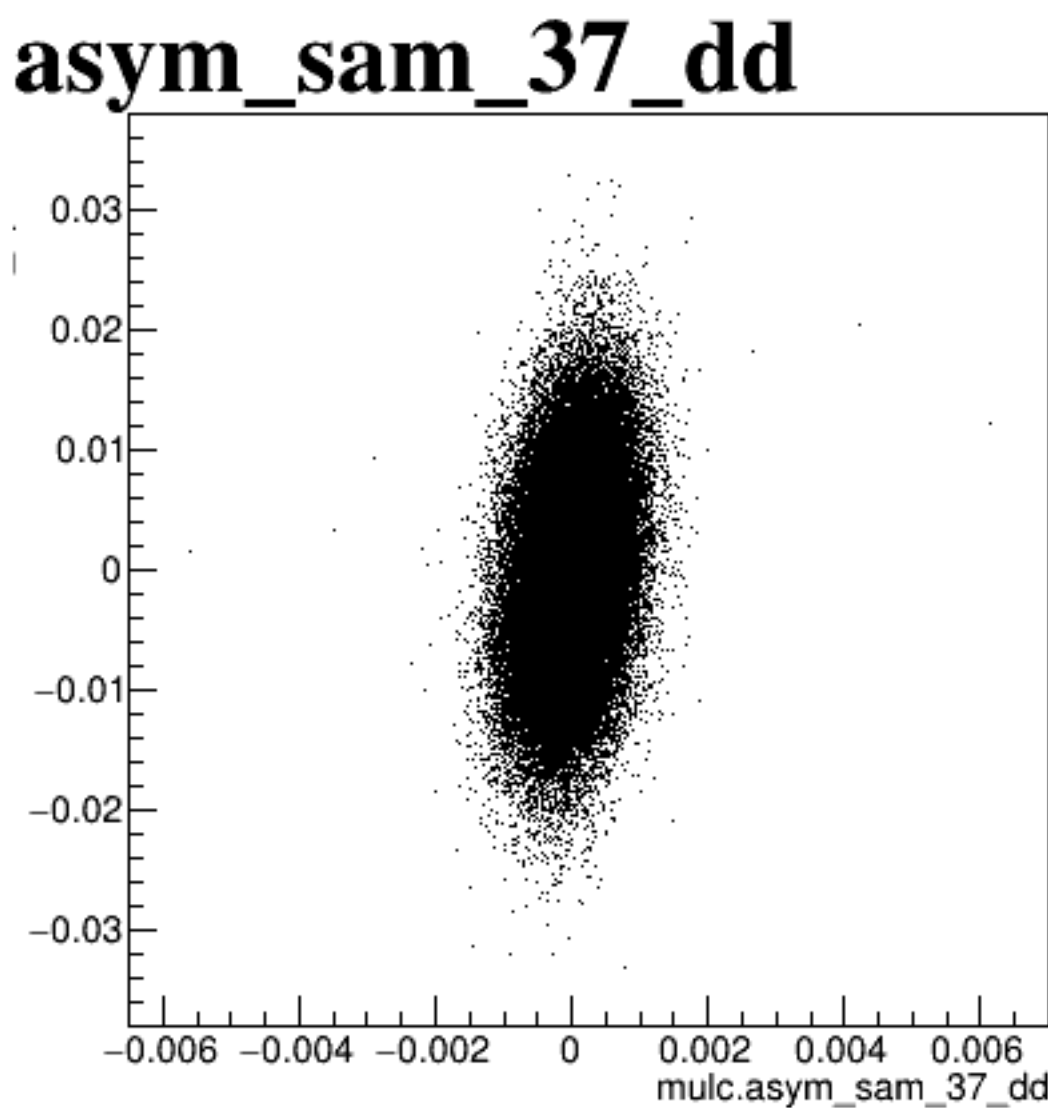
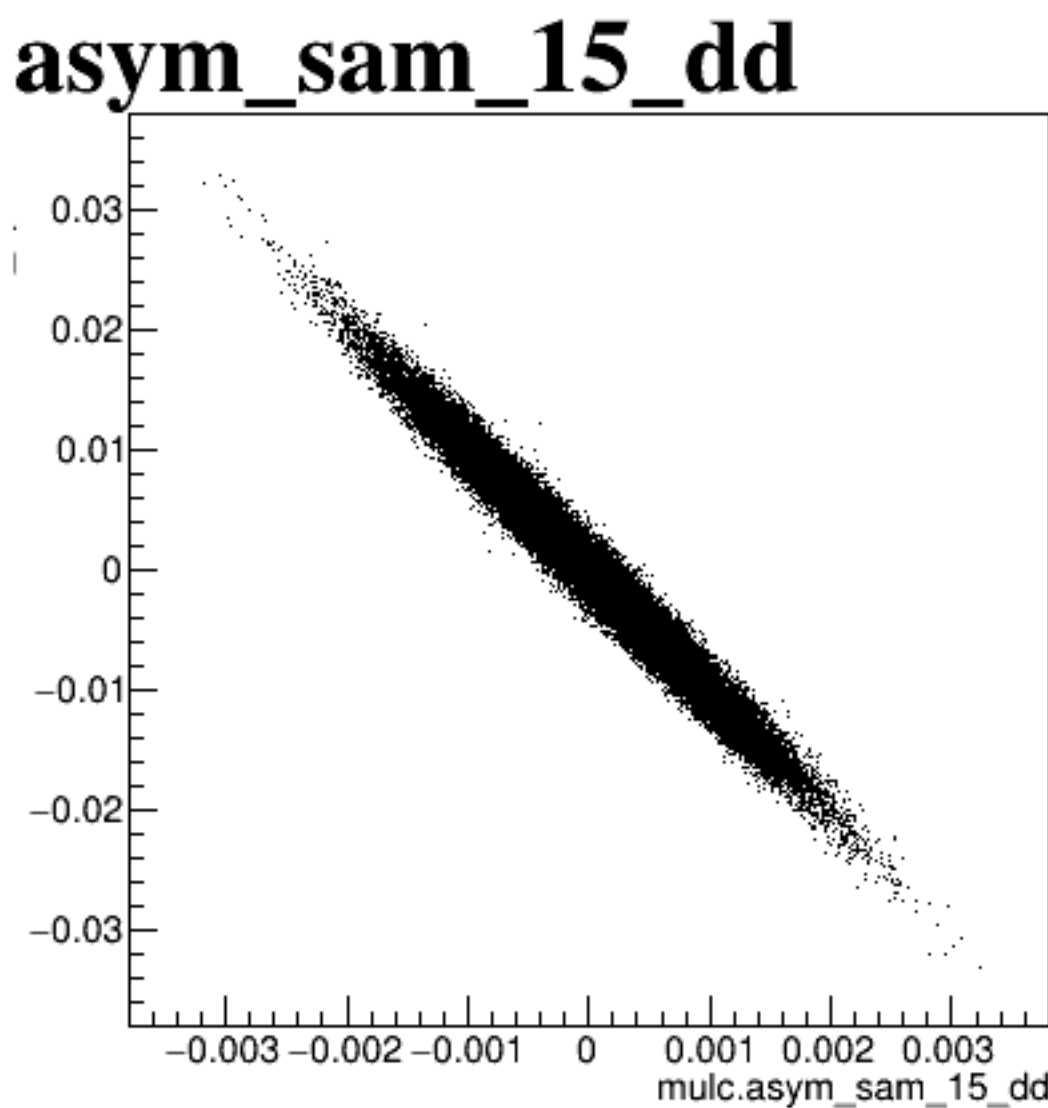
diff_bpm4aY



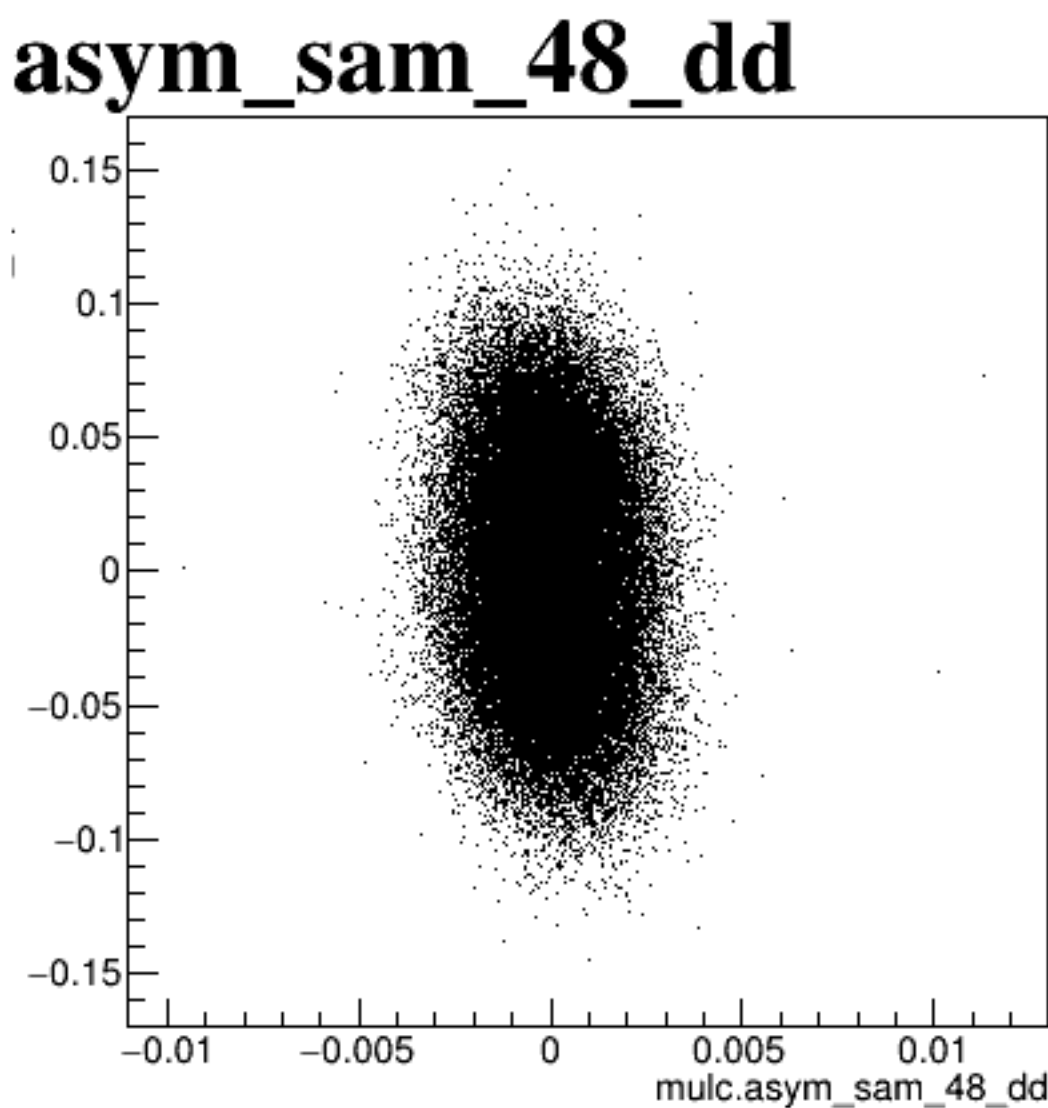
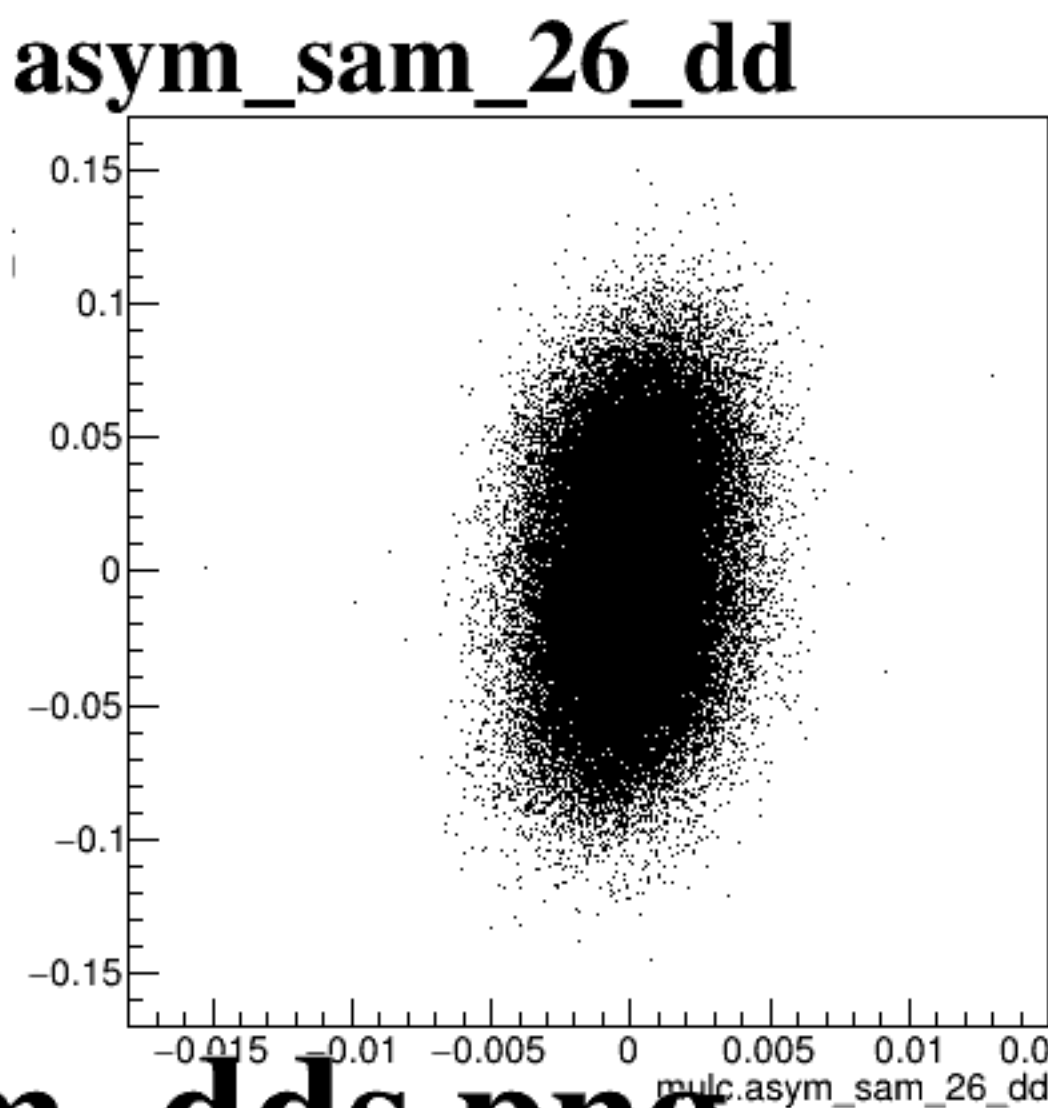
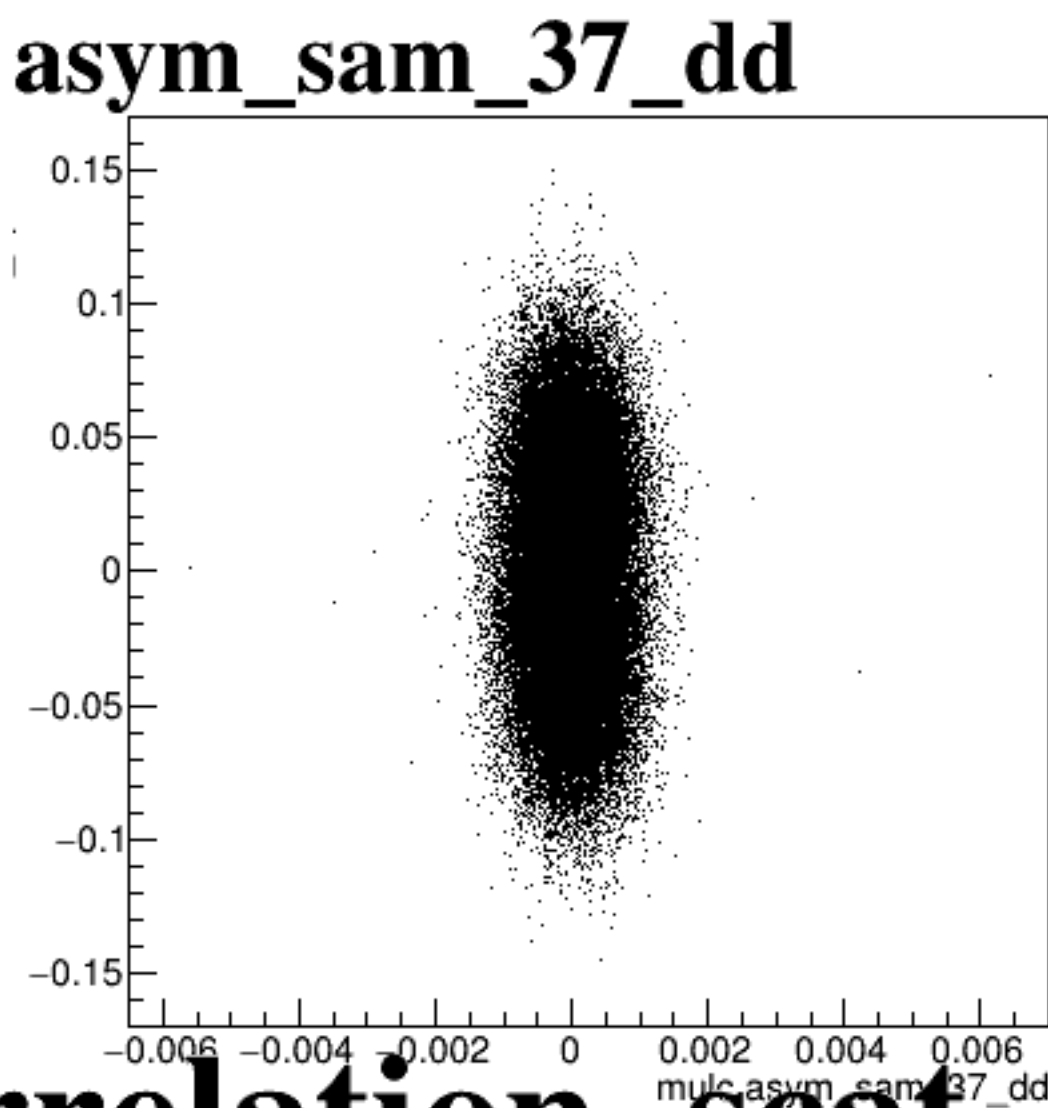
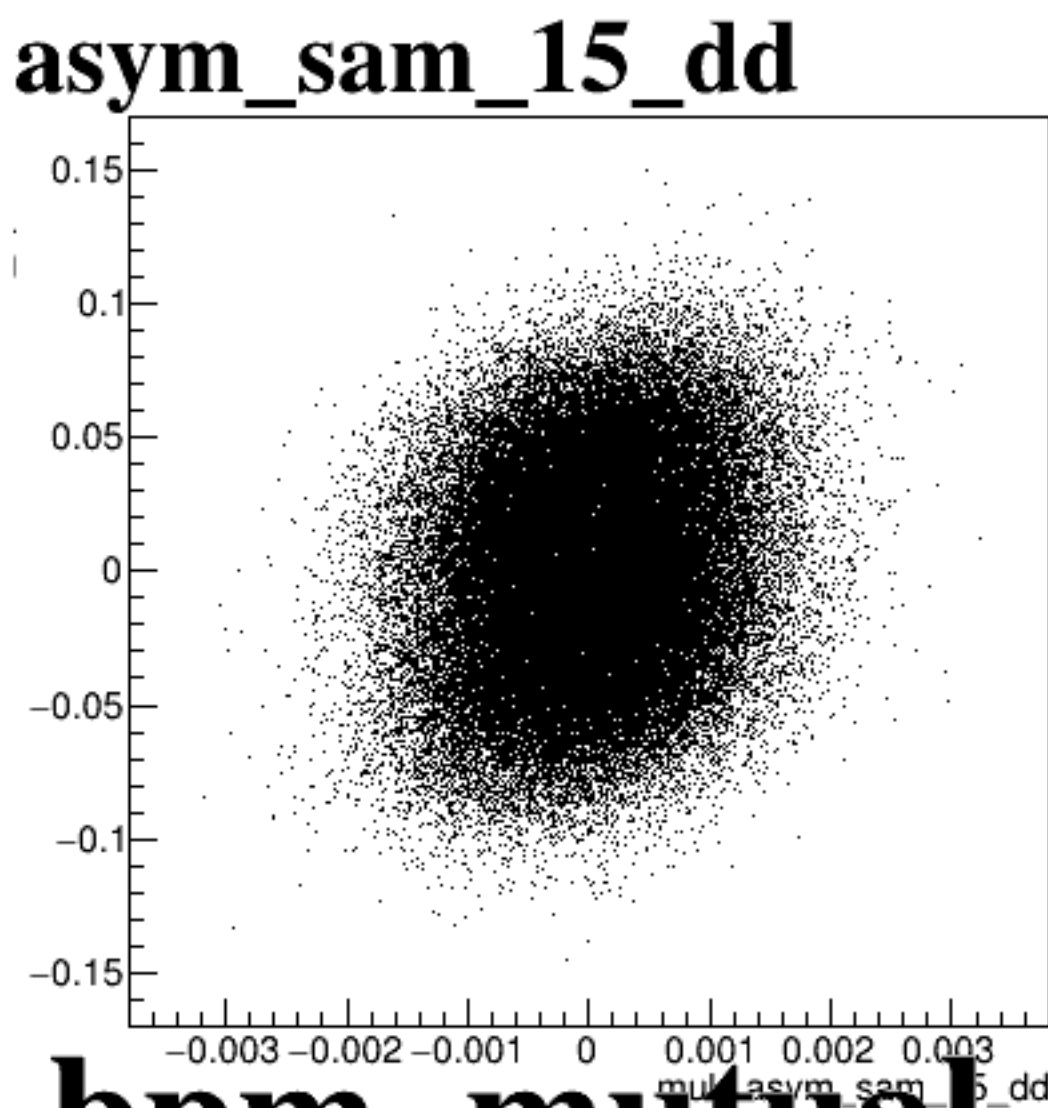
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diff_bpm4eY

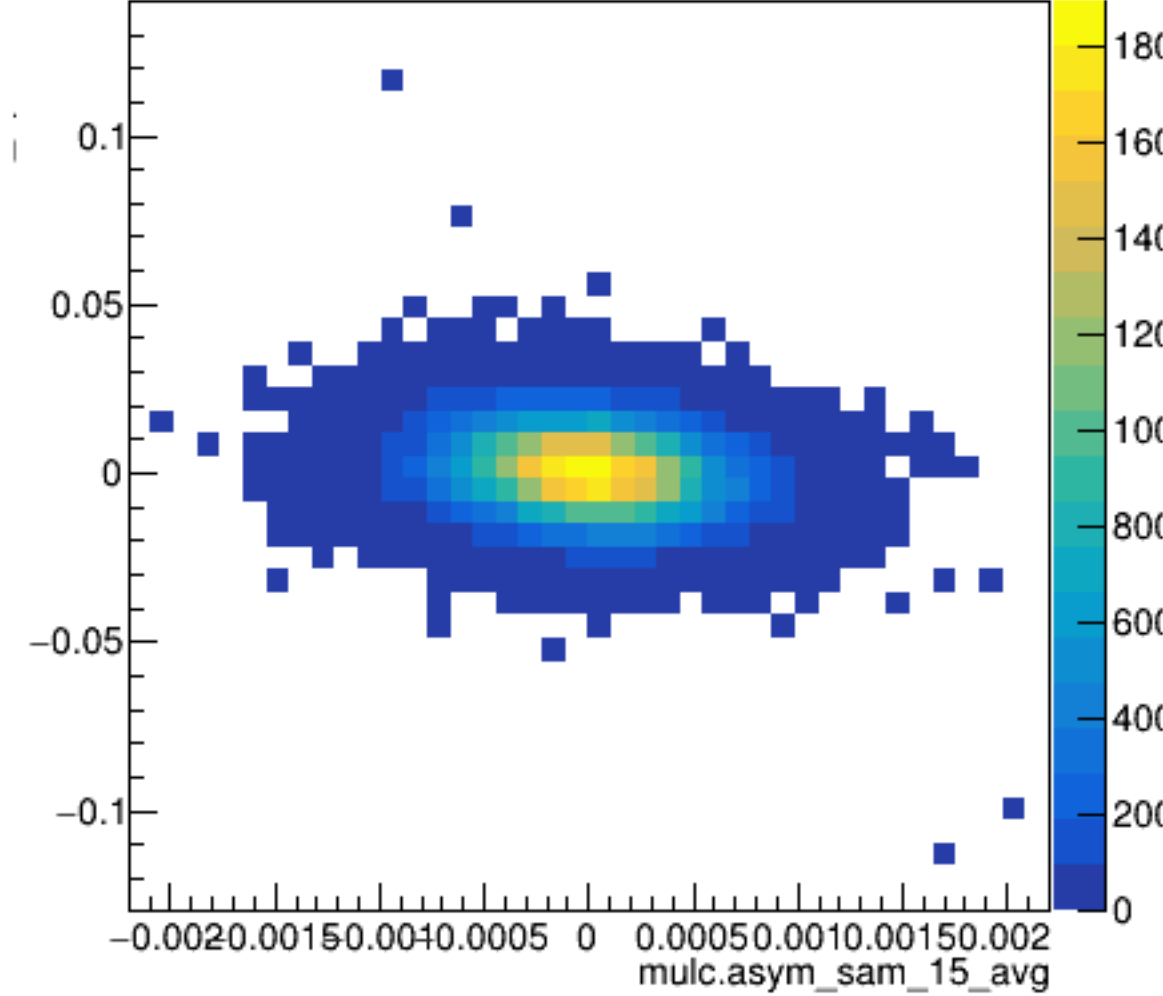


diff_bpm12X

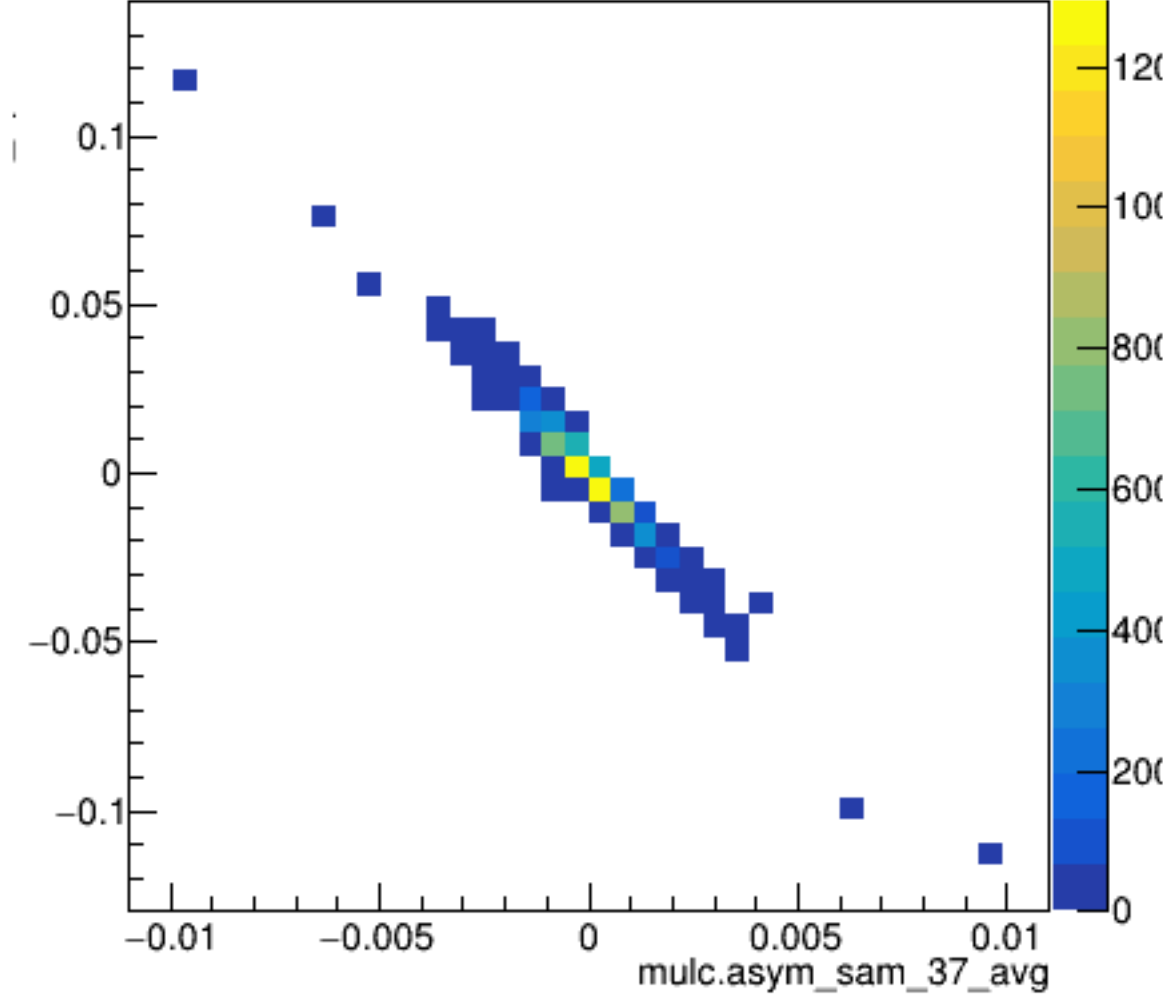


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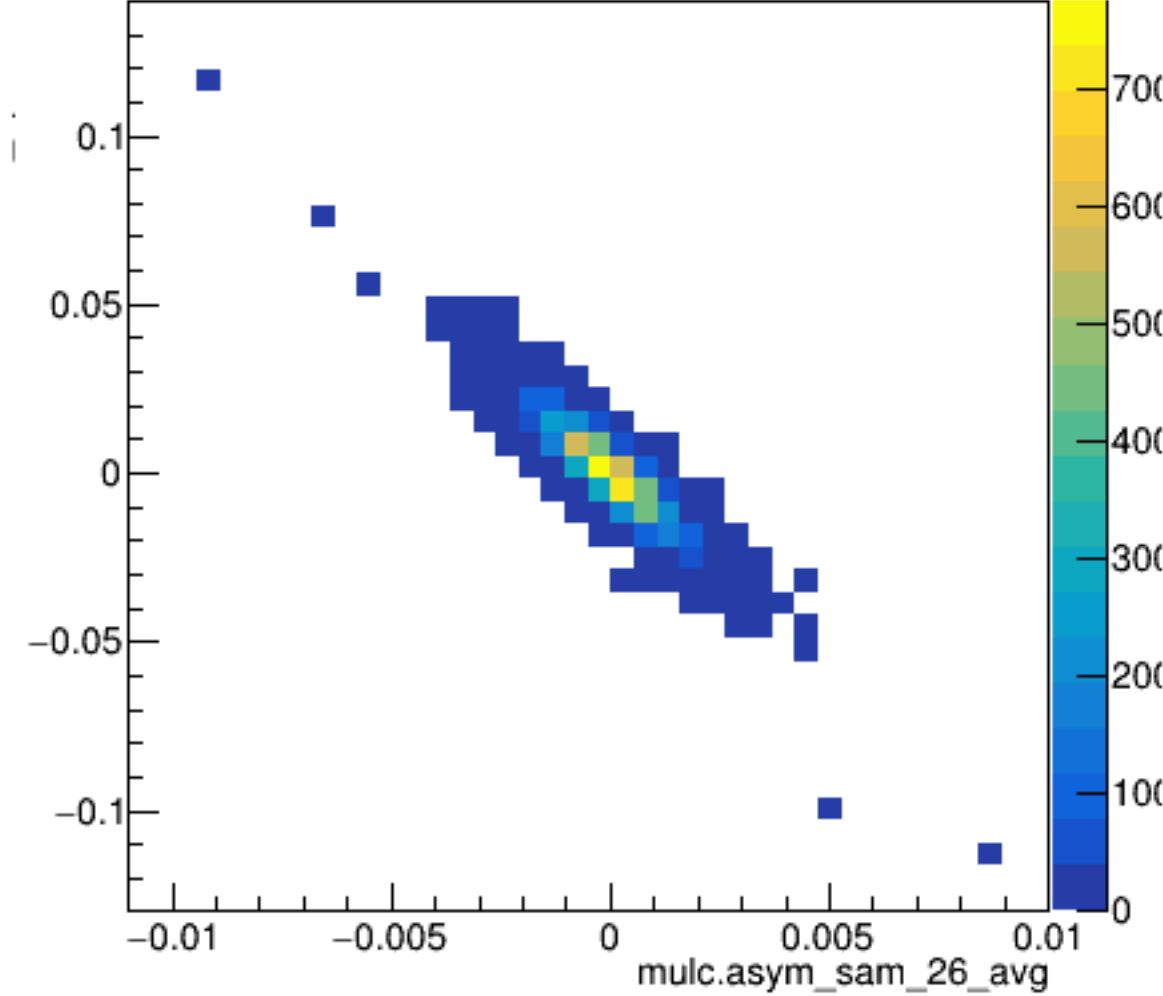
asym_sam_15_avg



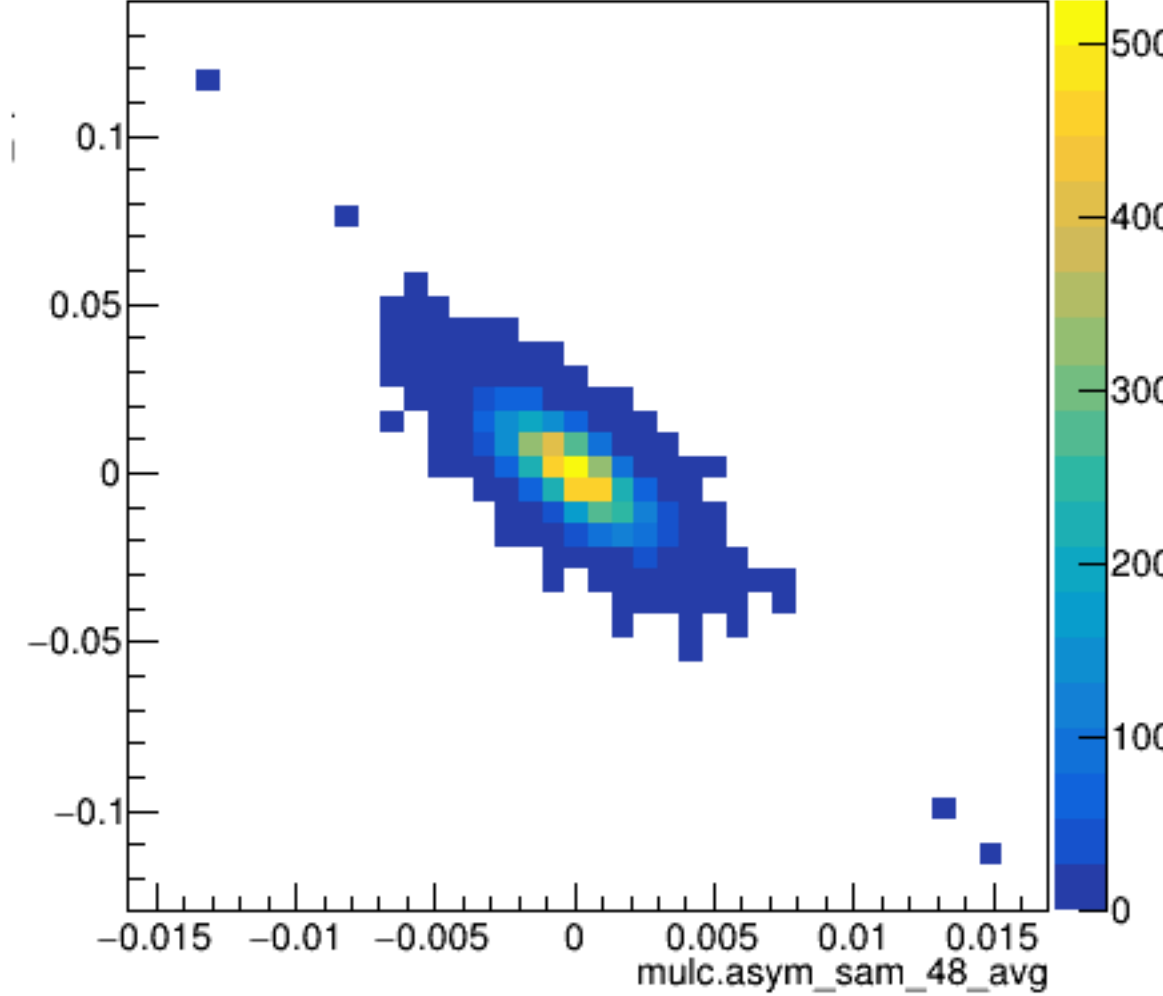
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asym_sam_26_avg

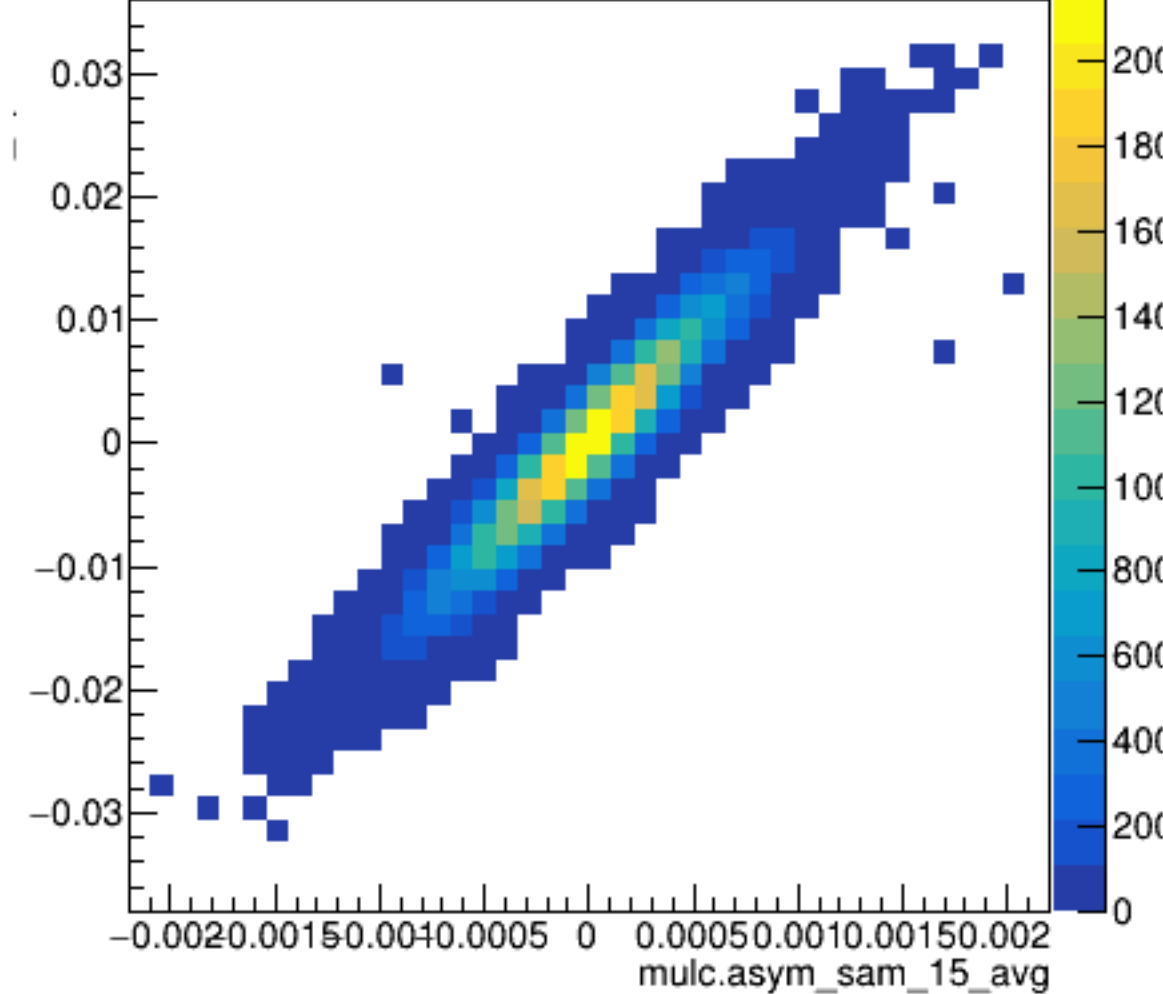


asym_sam_48_avg

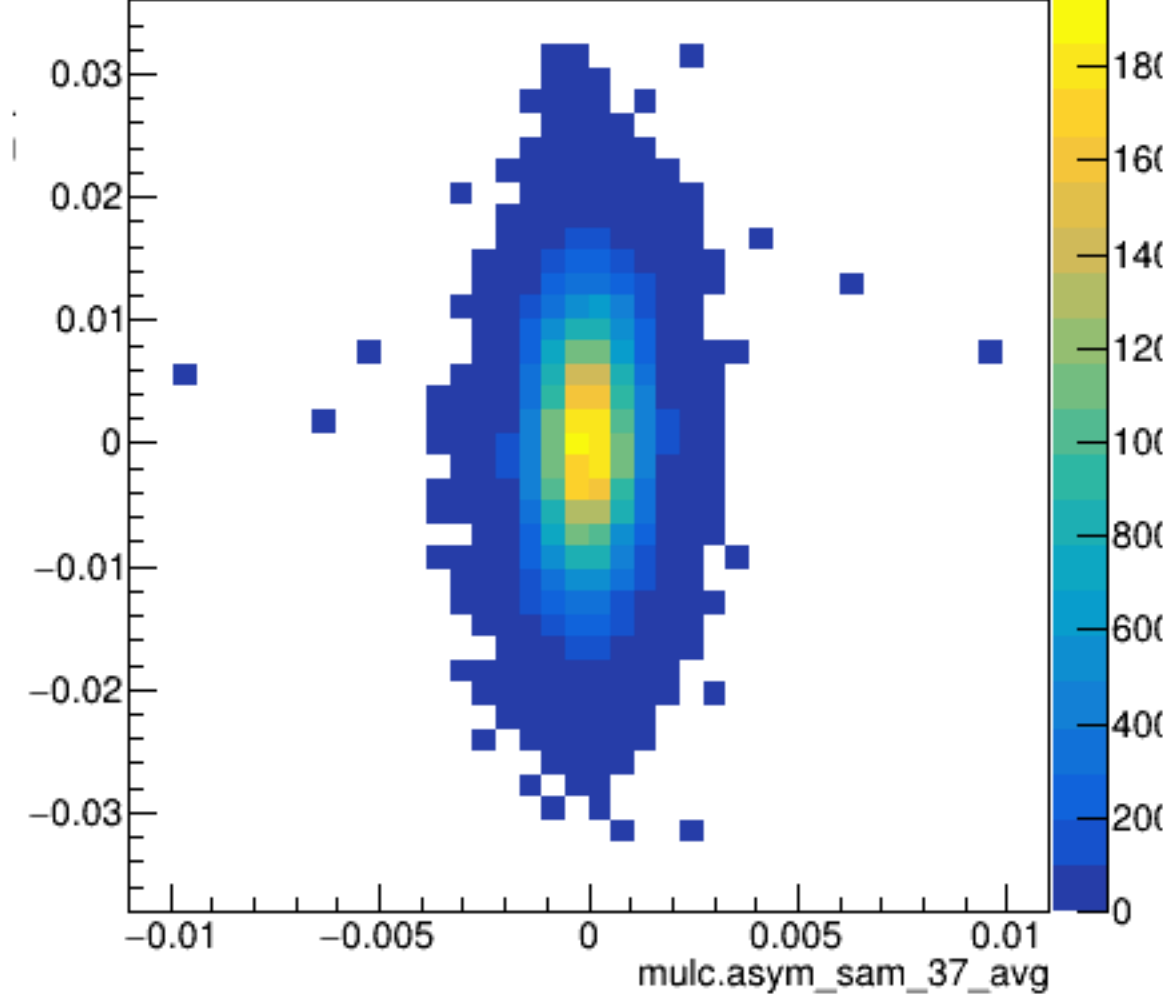


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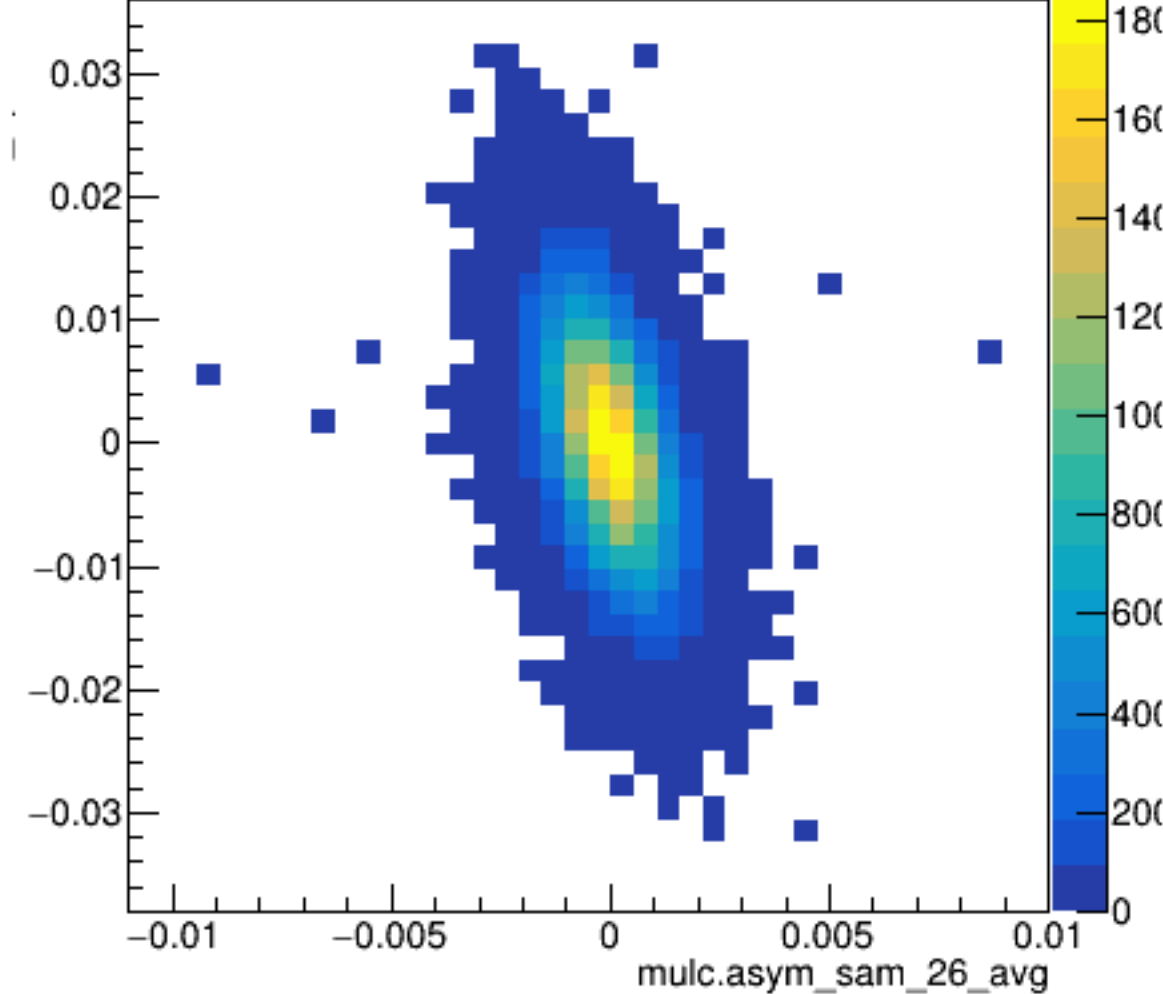
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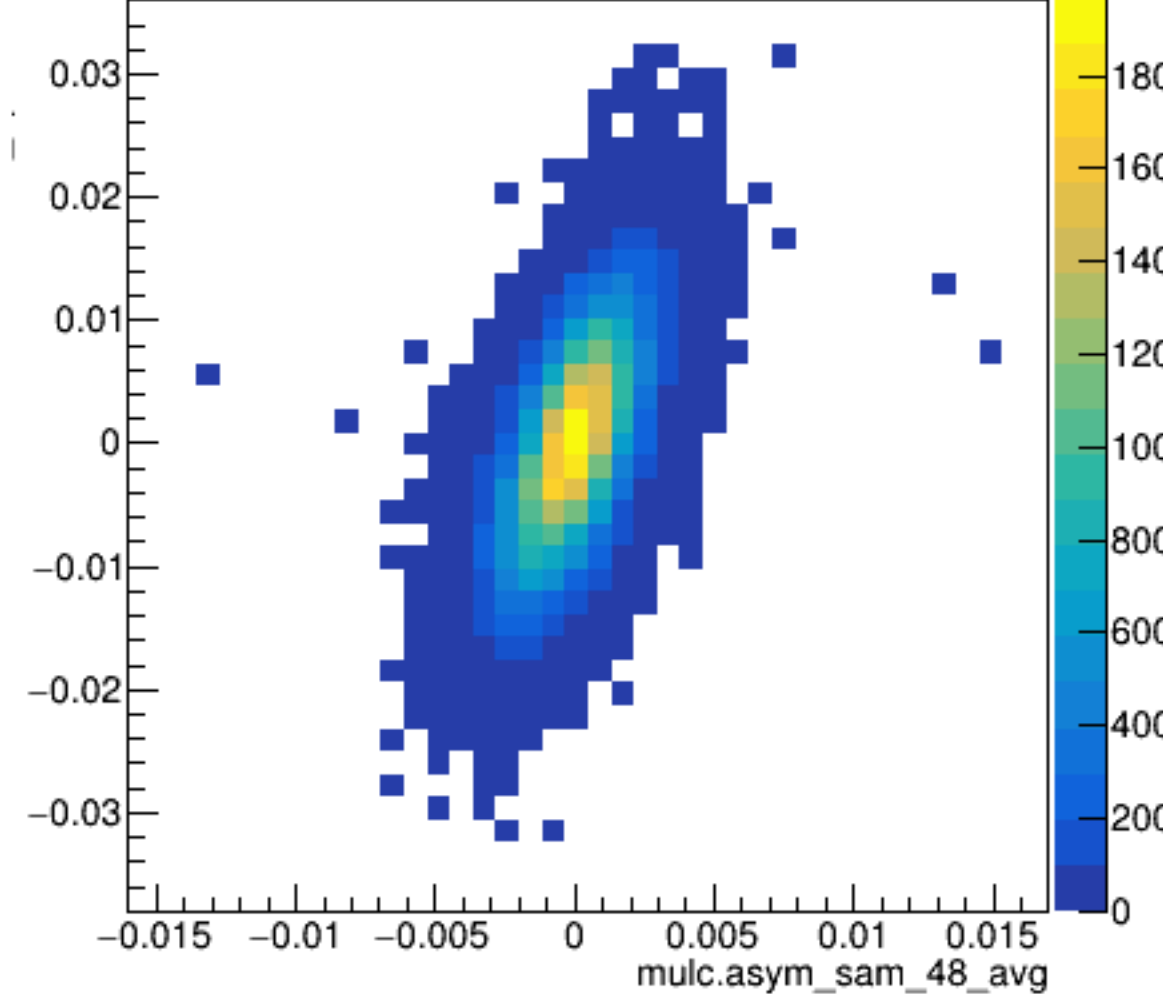
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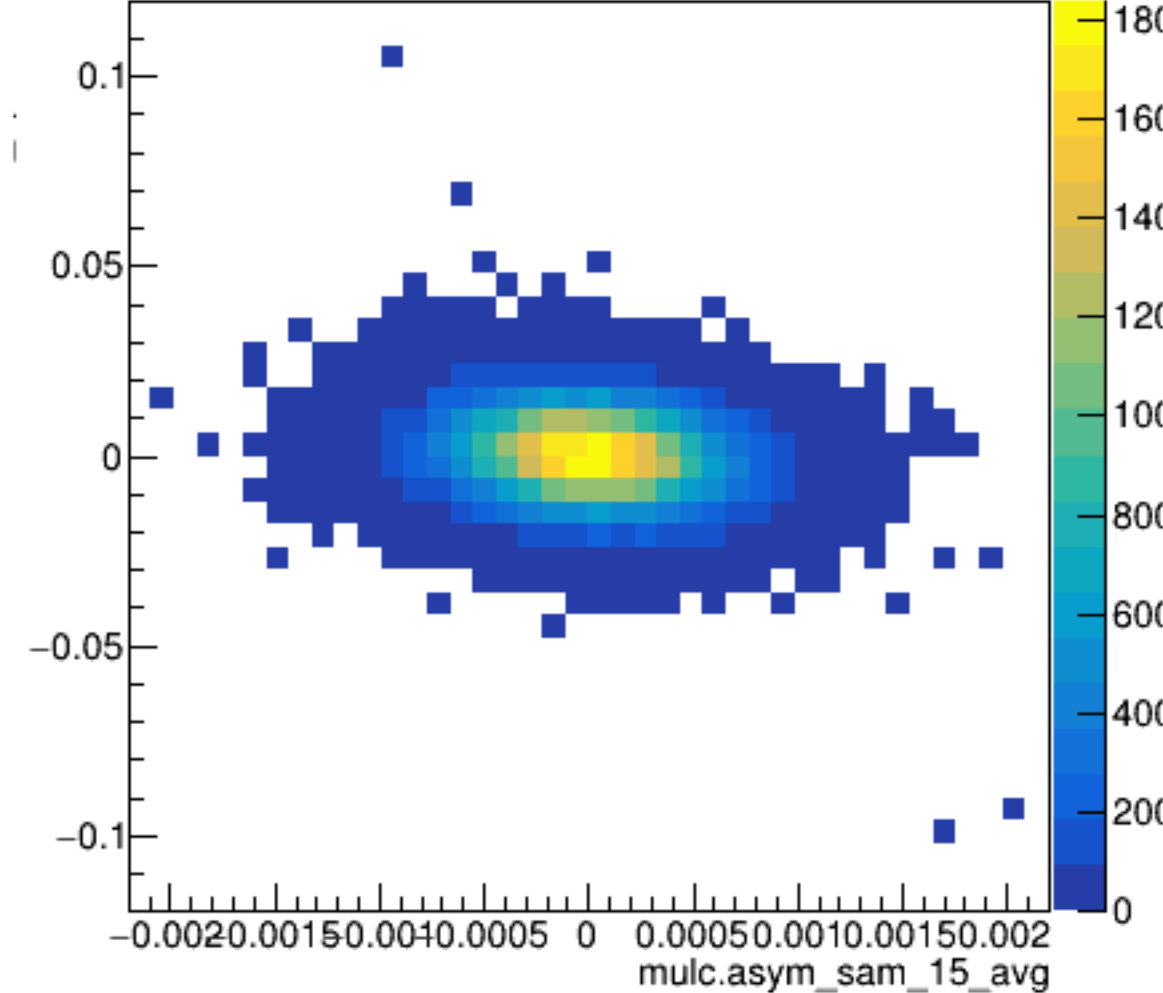


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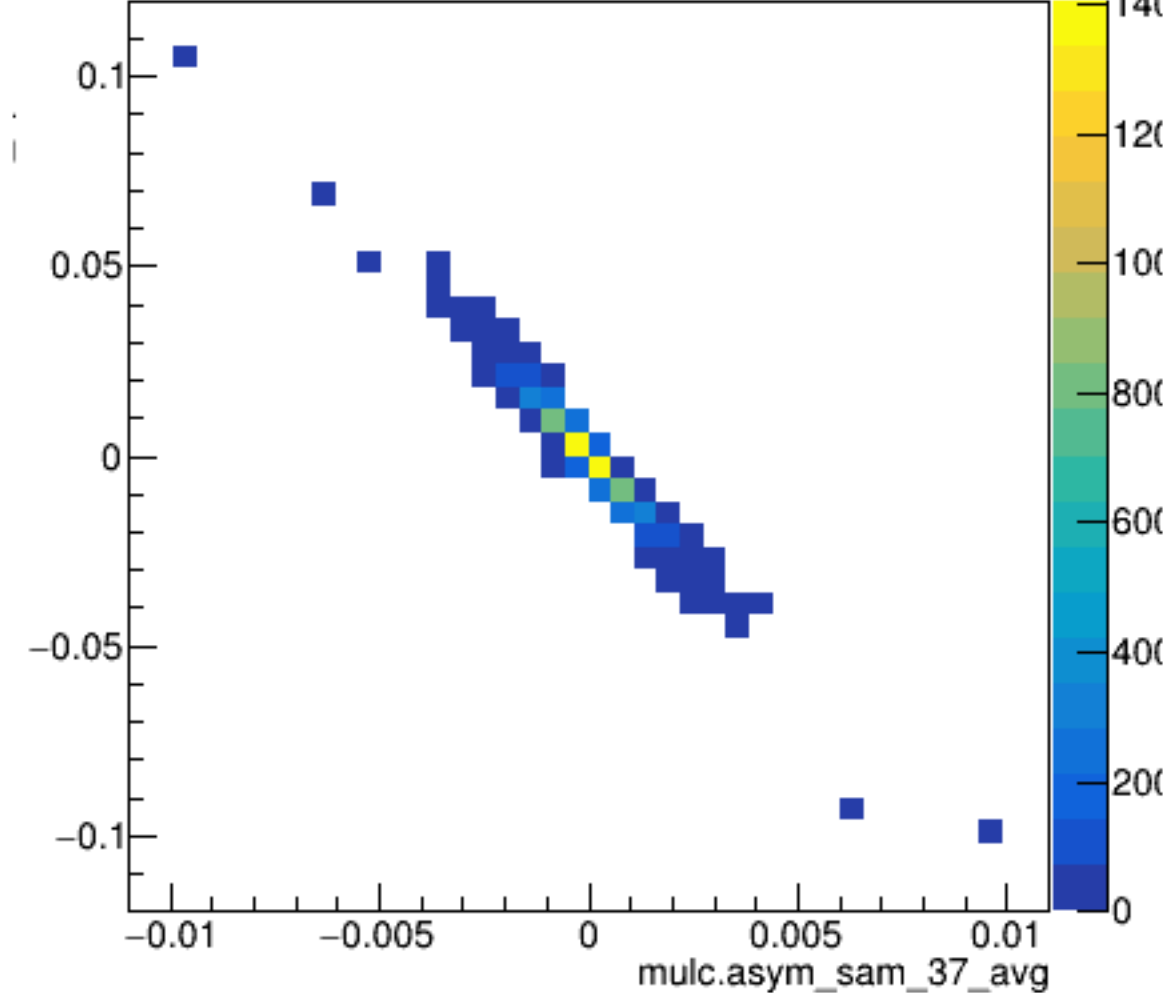


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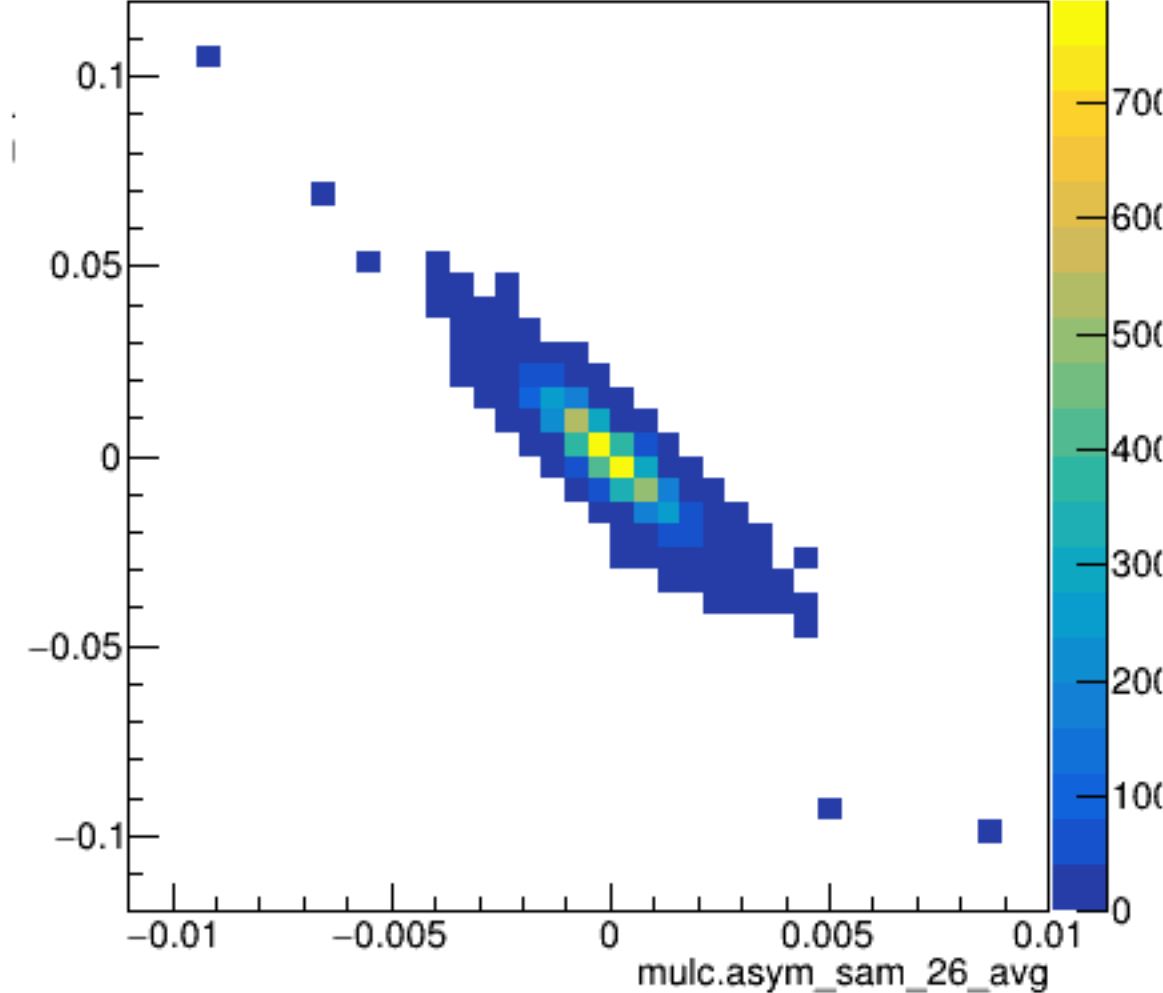
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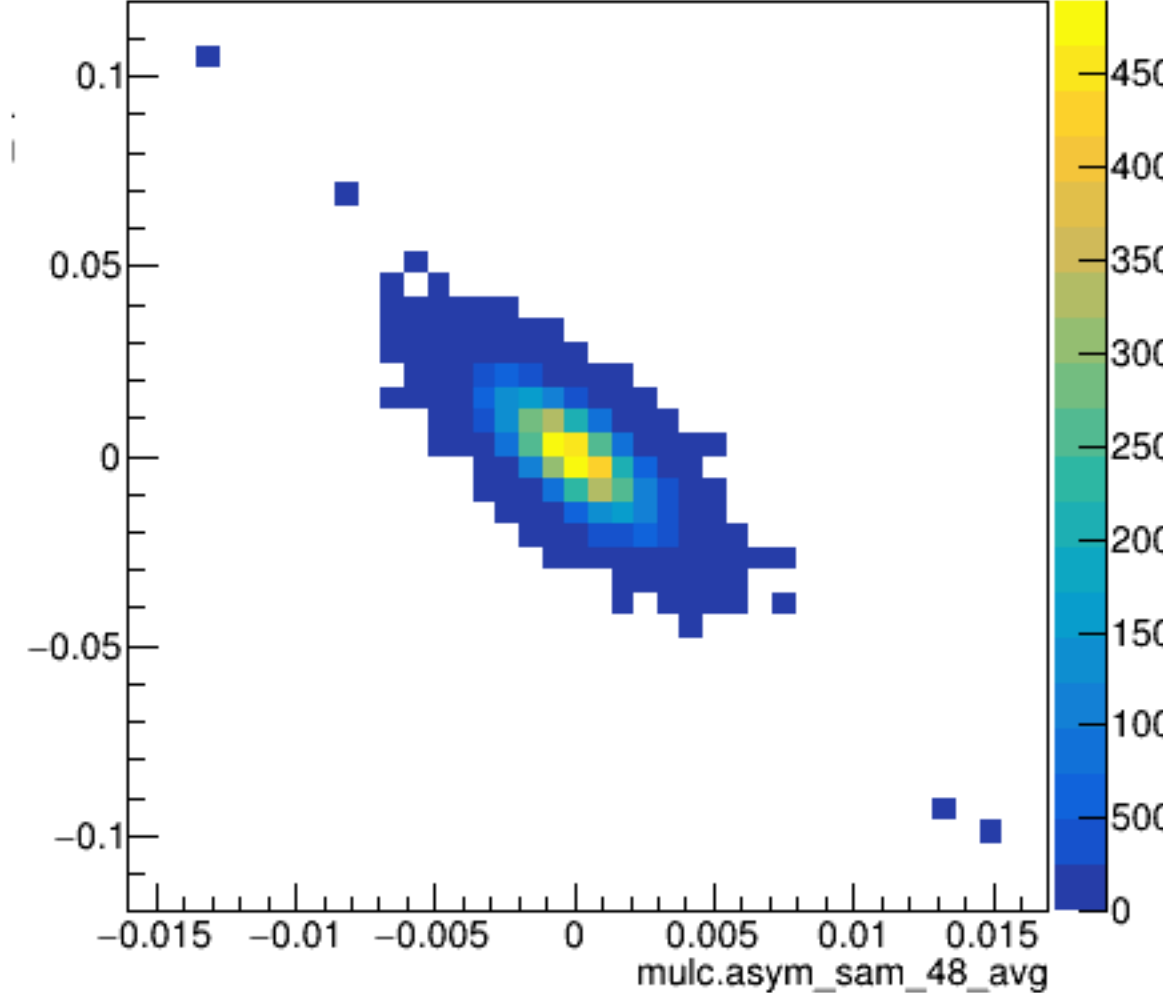
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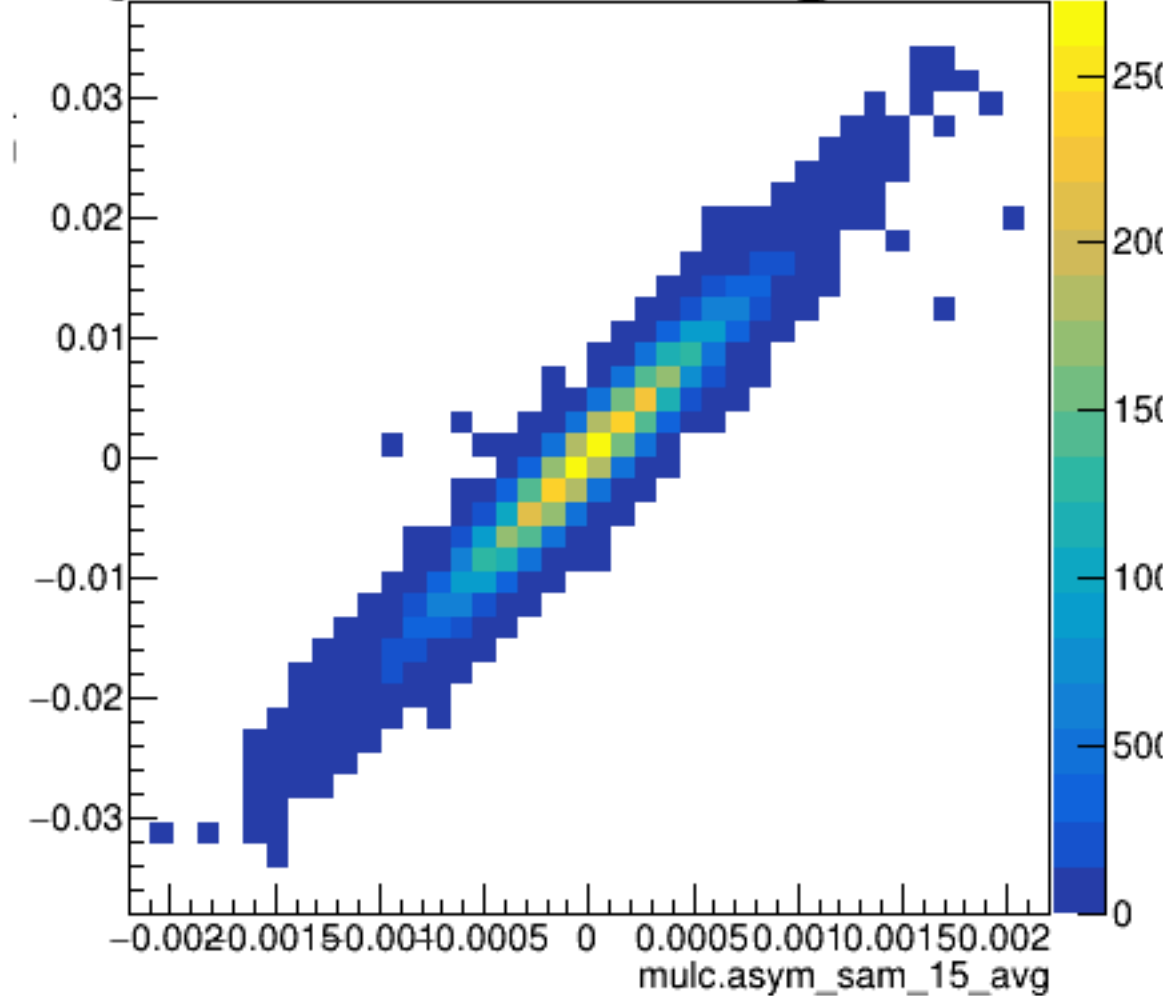


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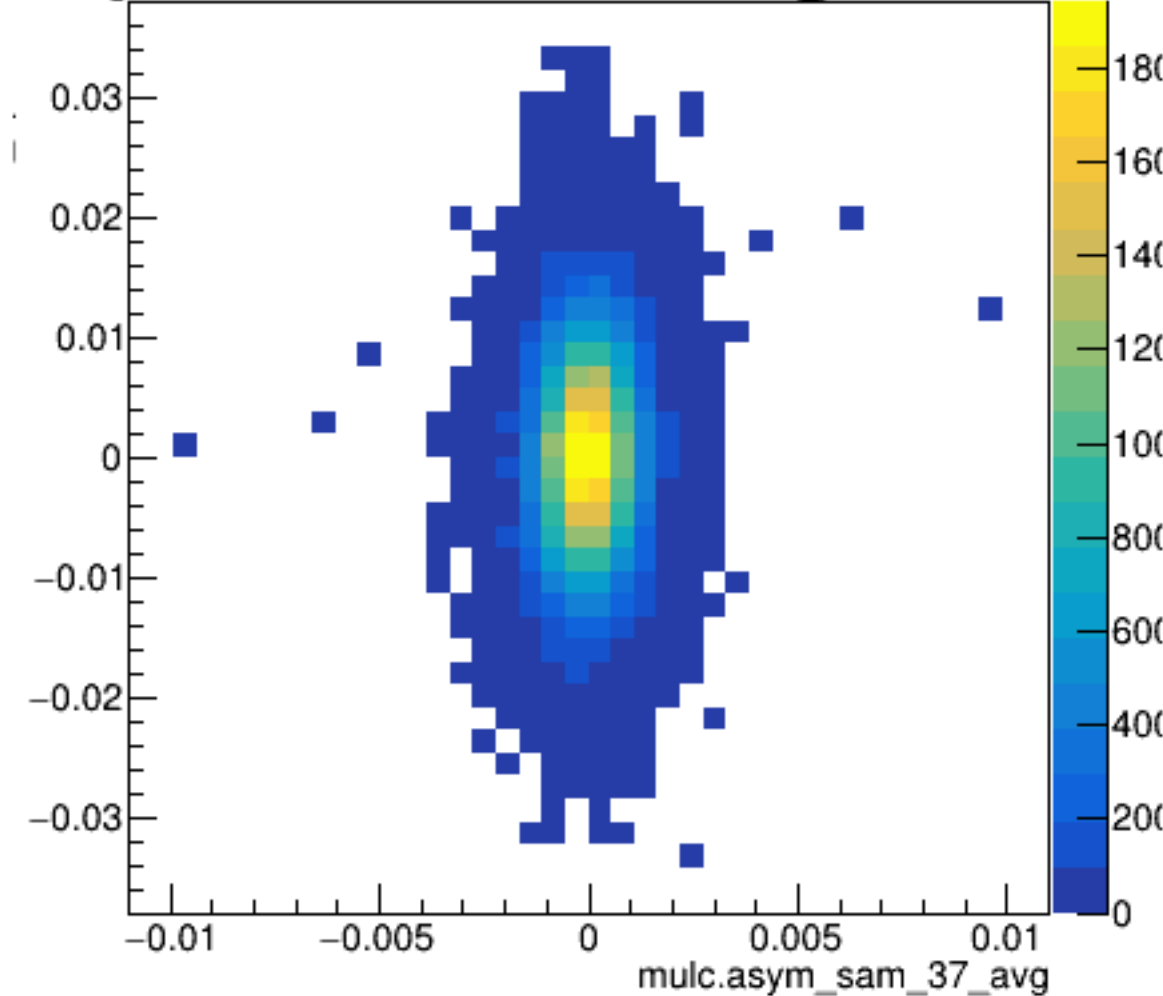


diff_bpm4eY

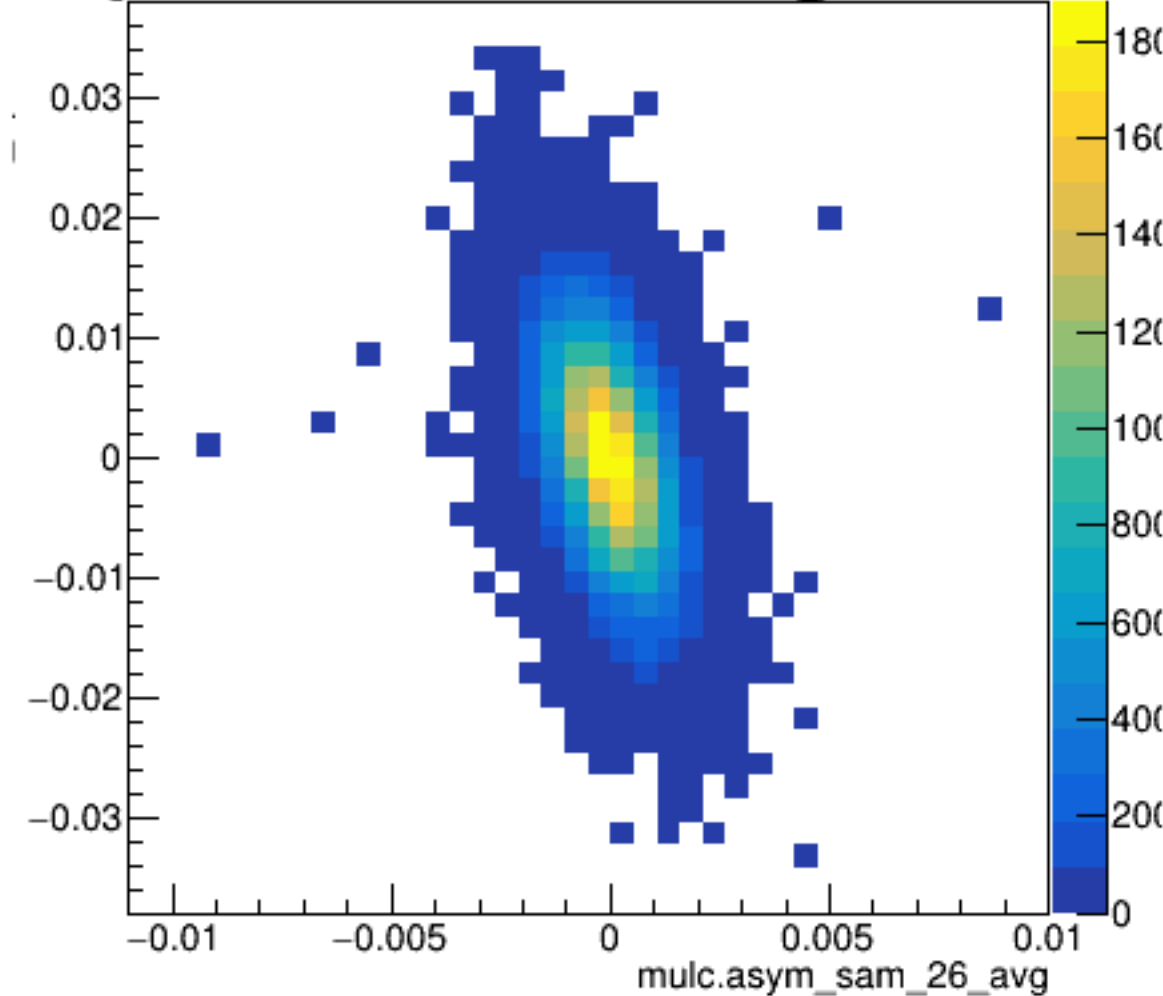
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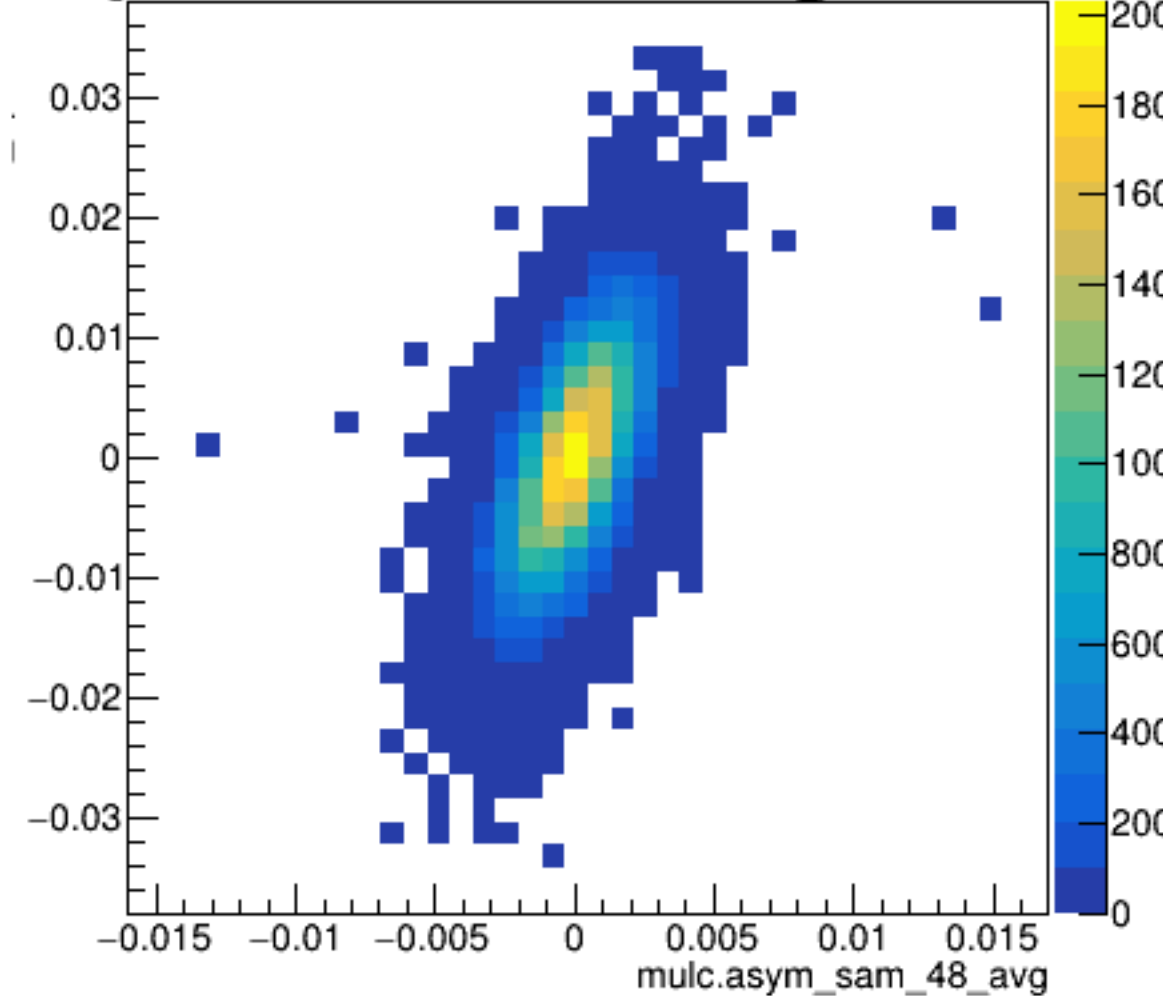
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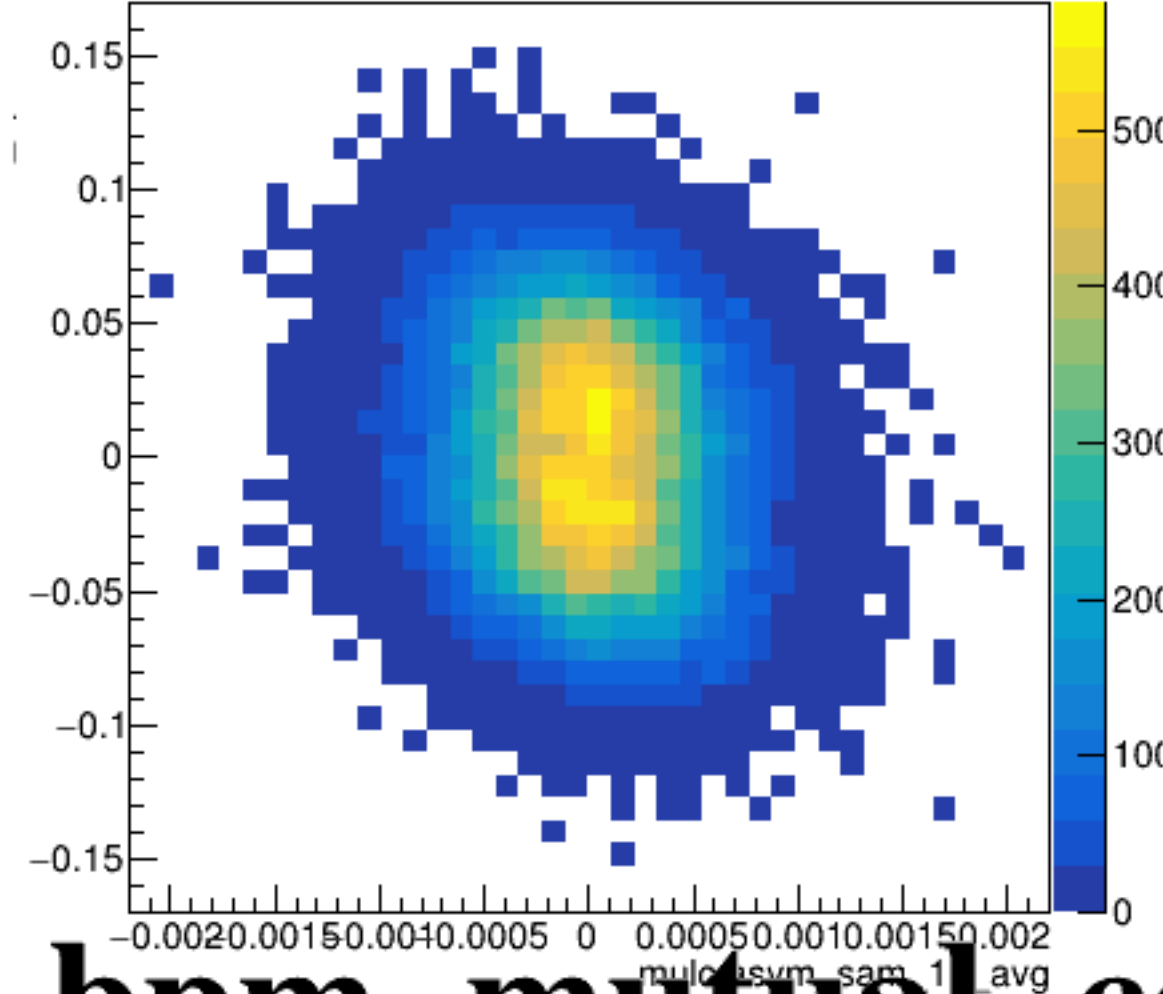


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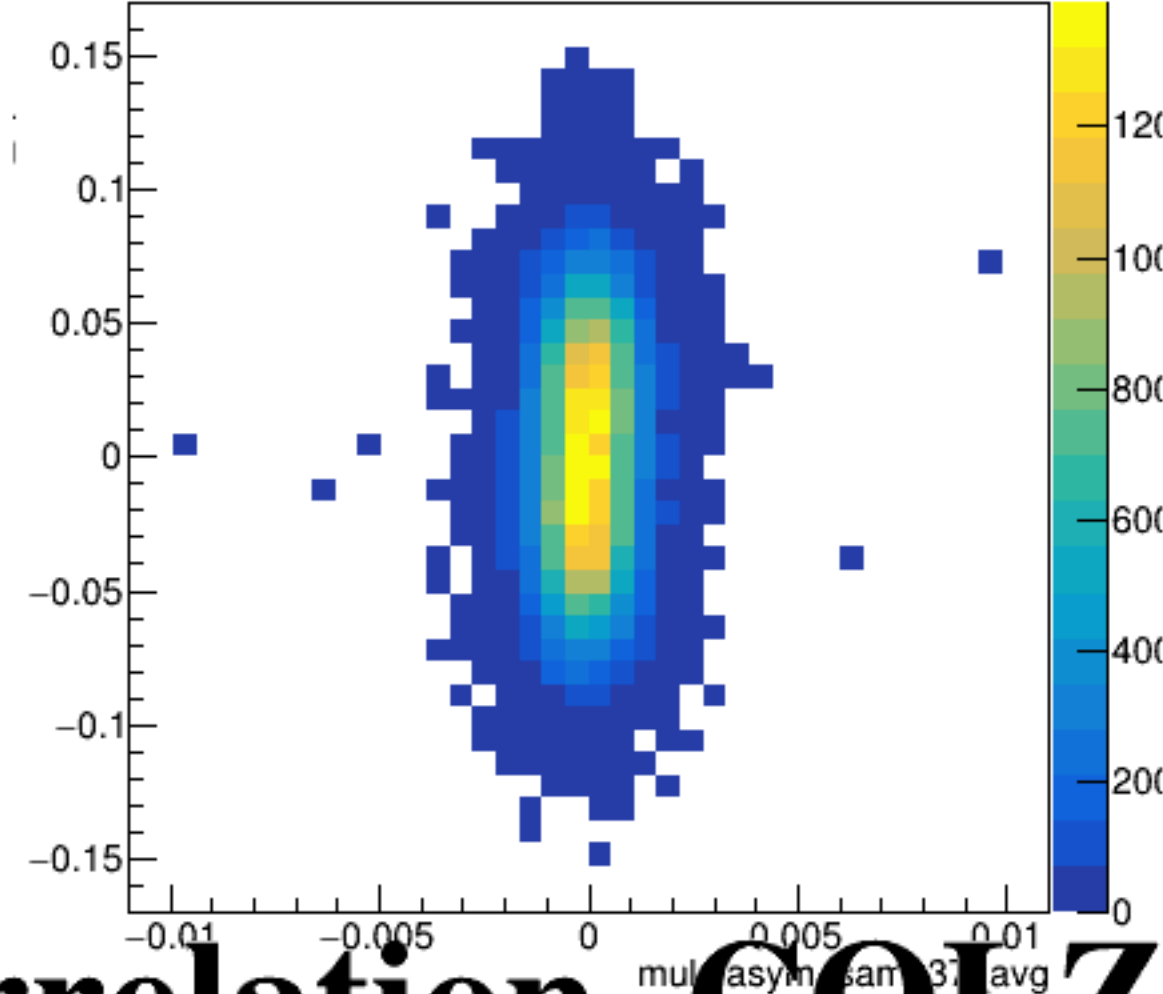


diff_bpm12X

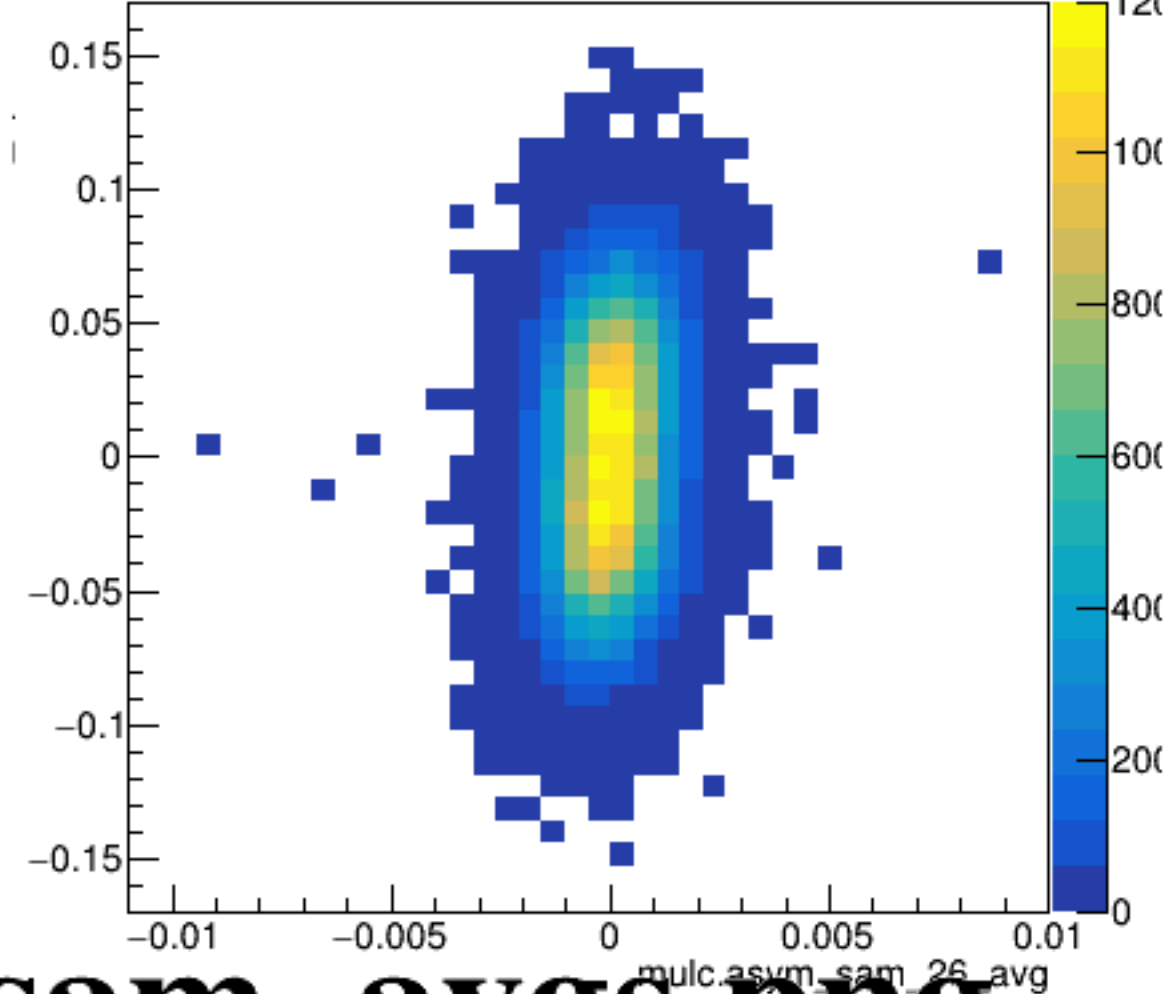
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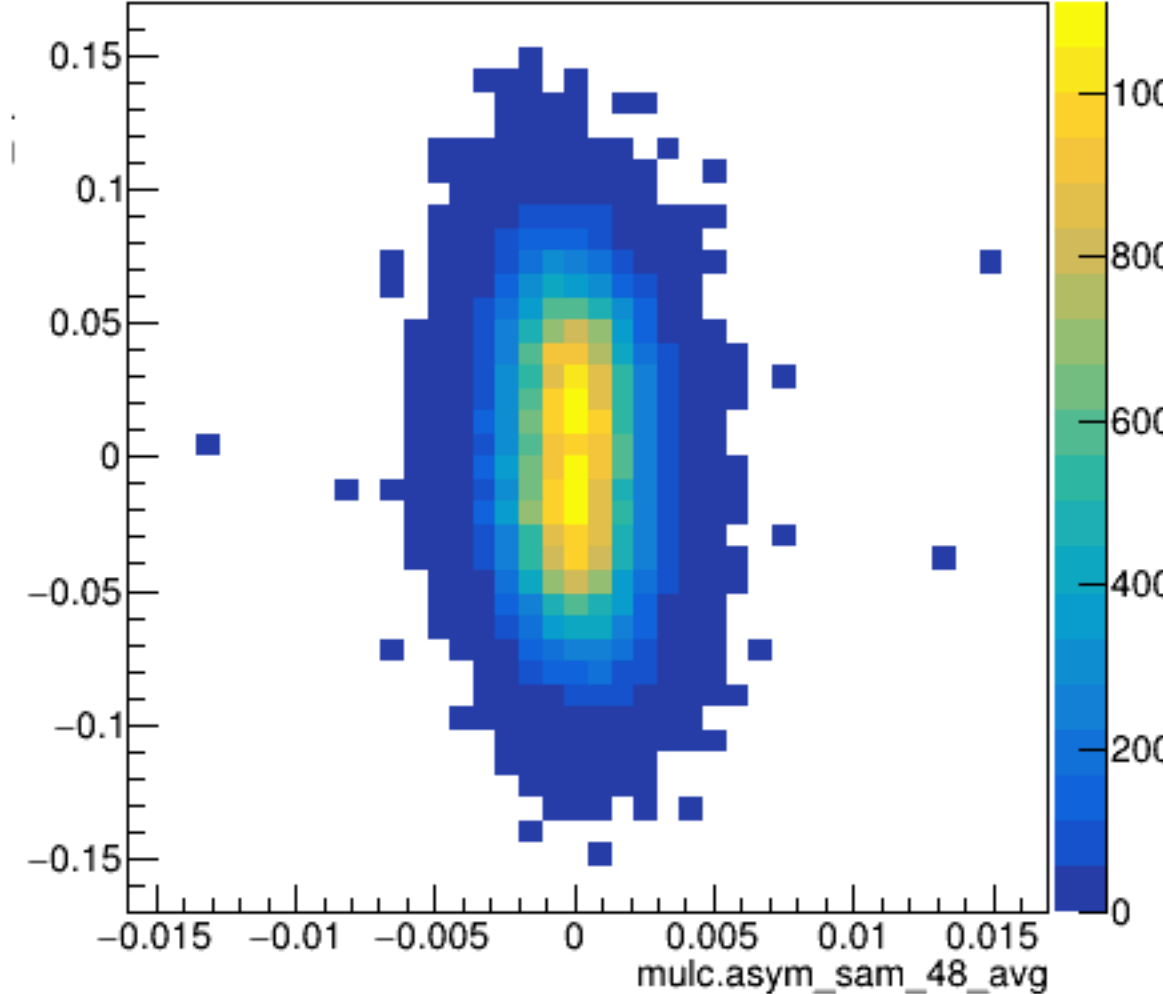
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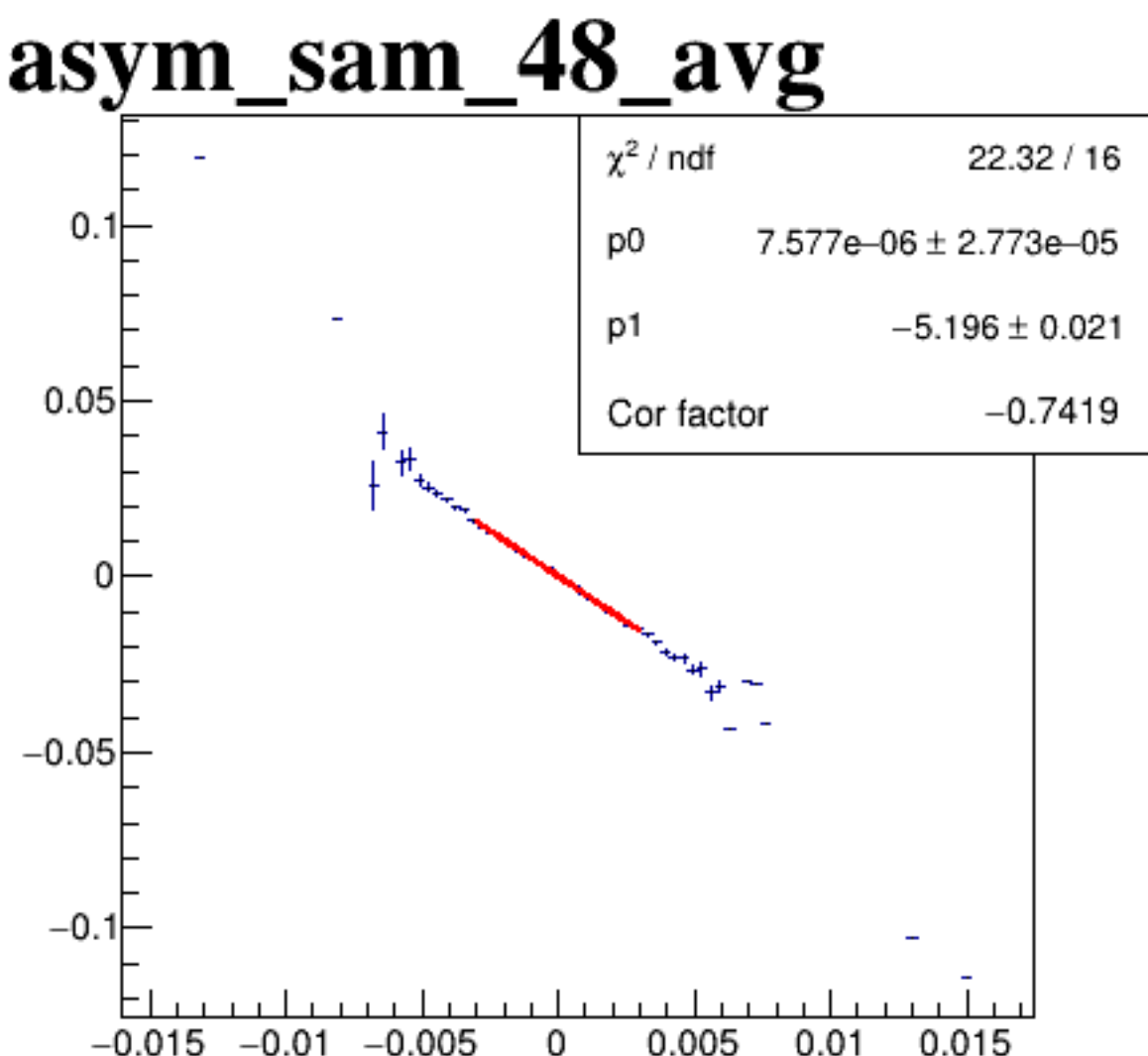
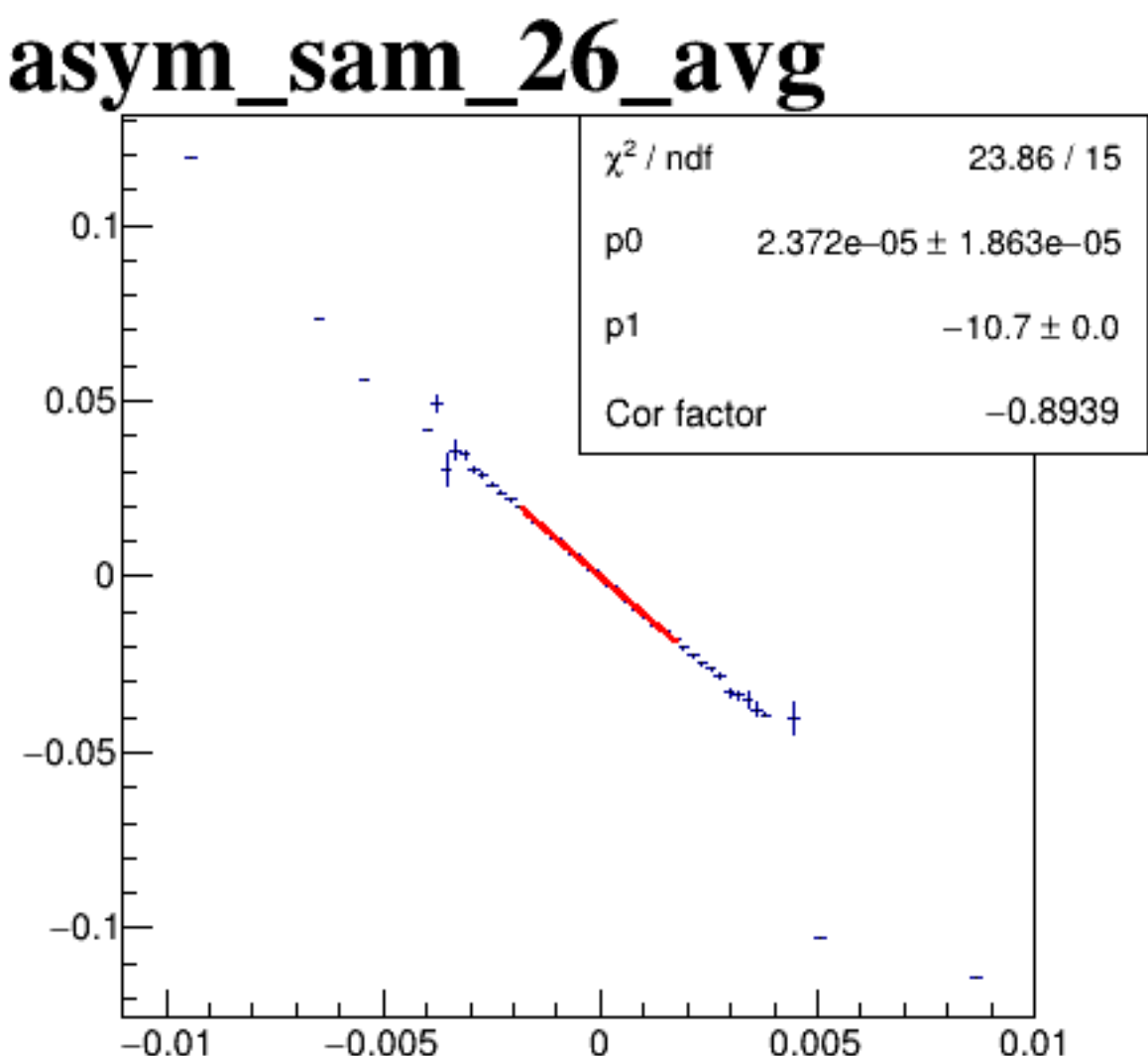
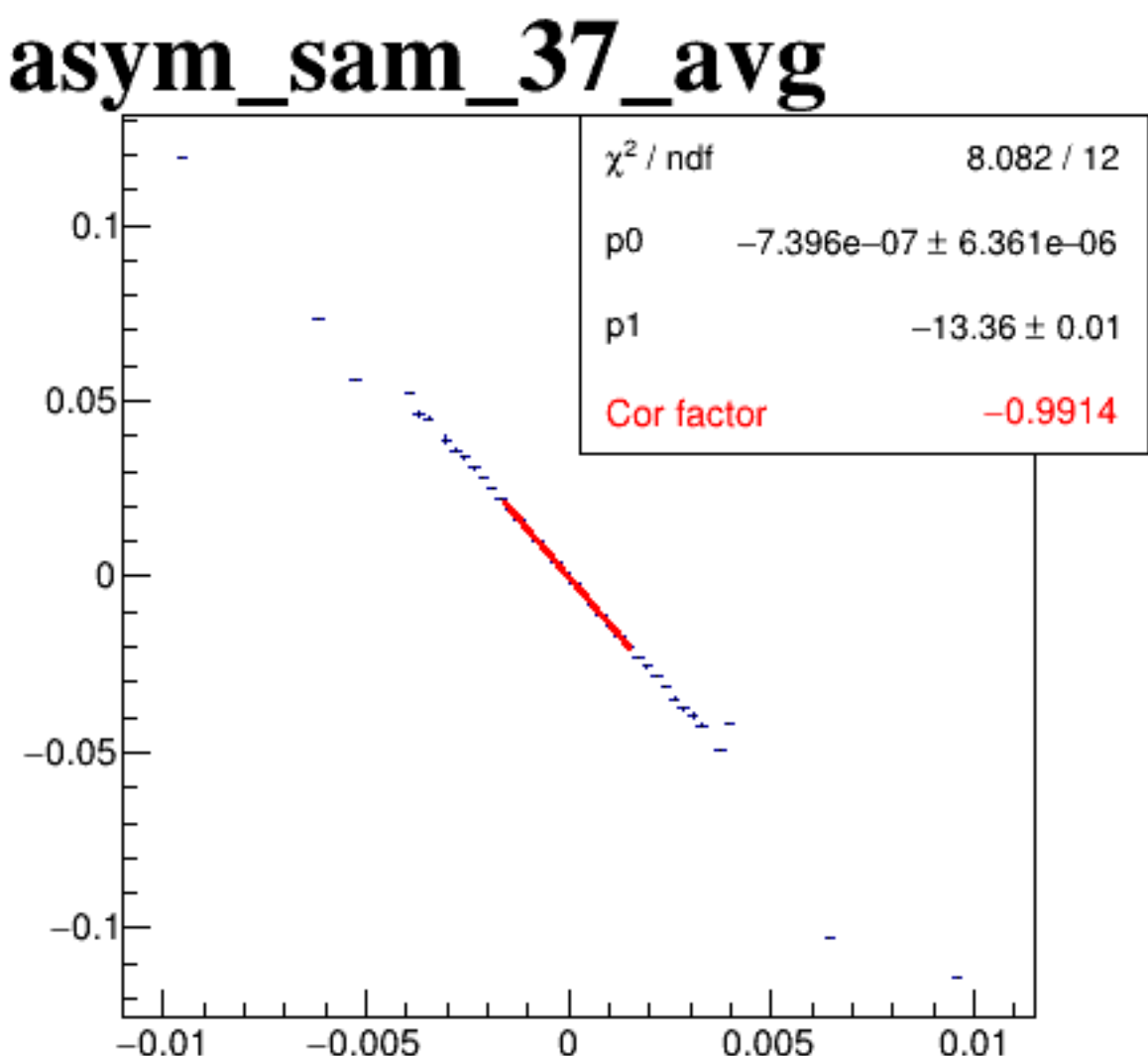
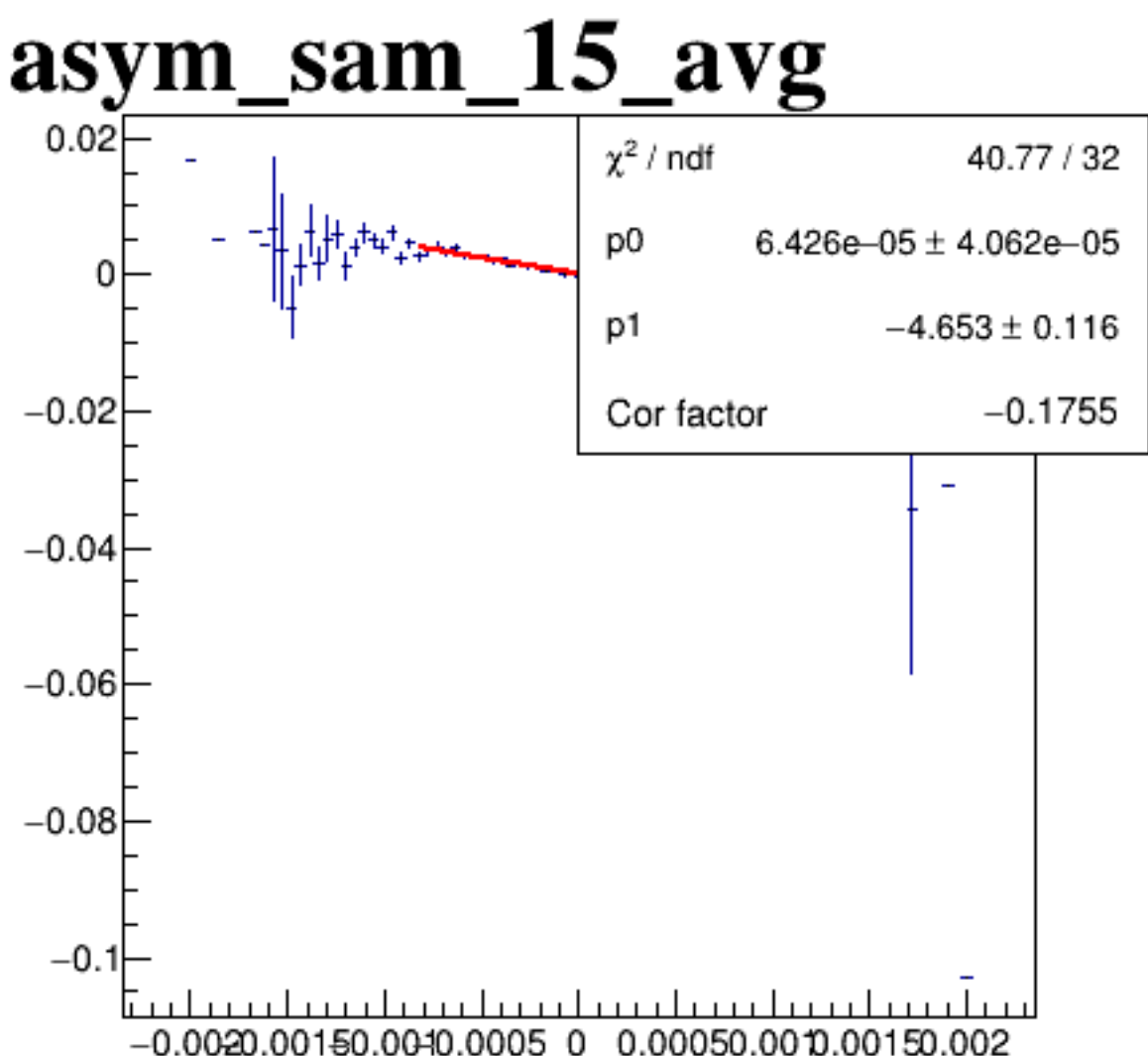
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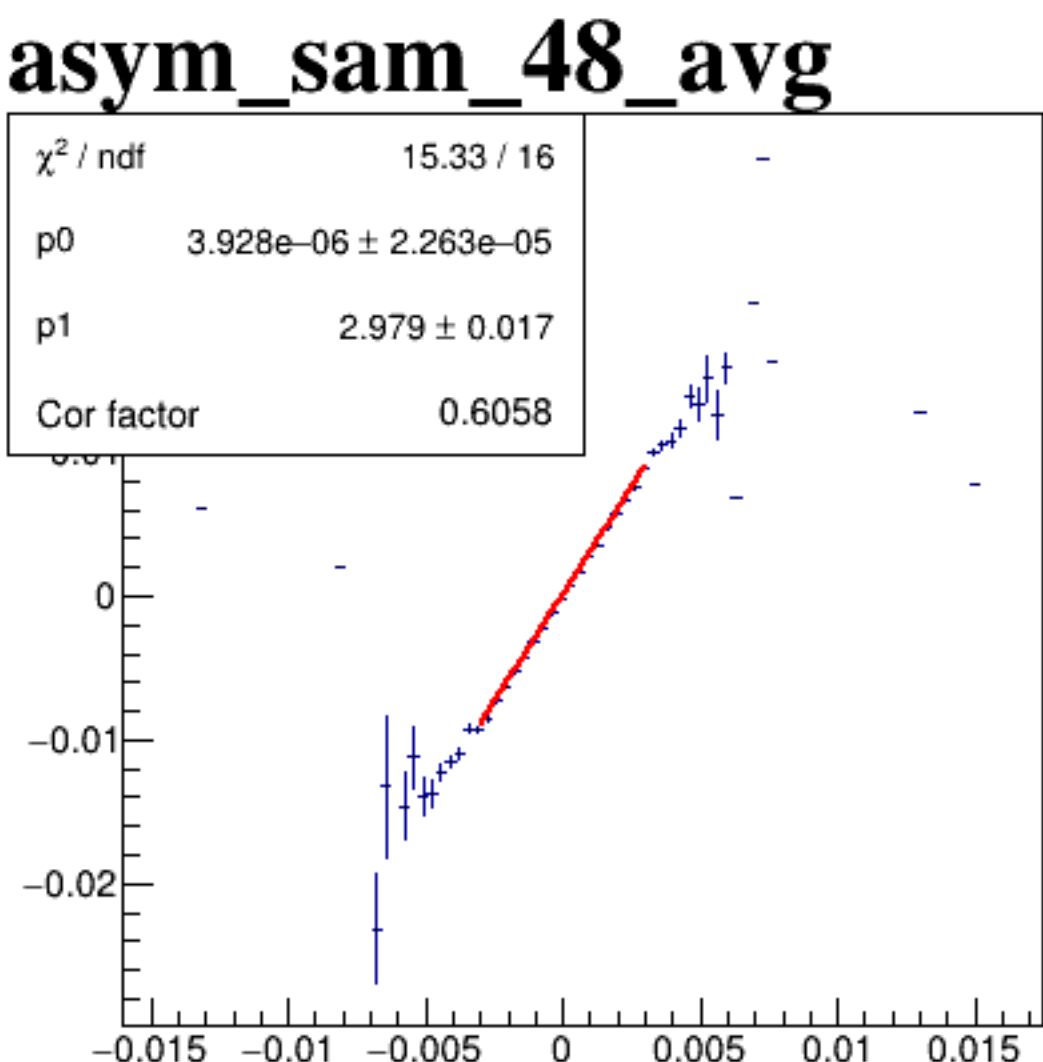
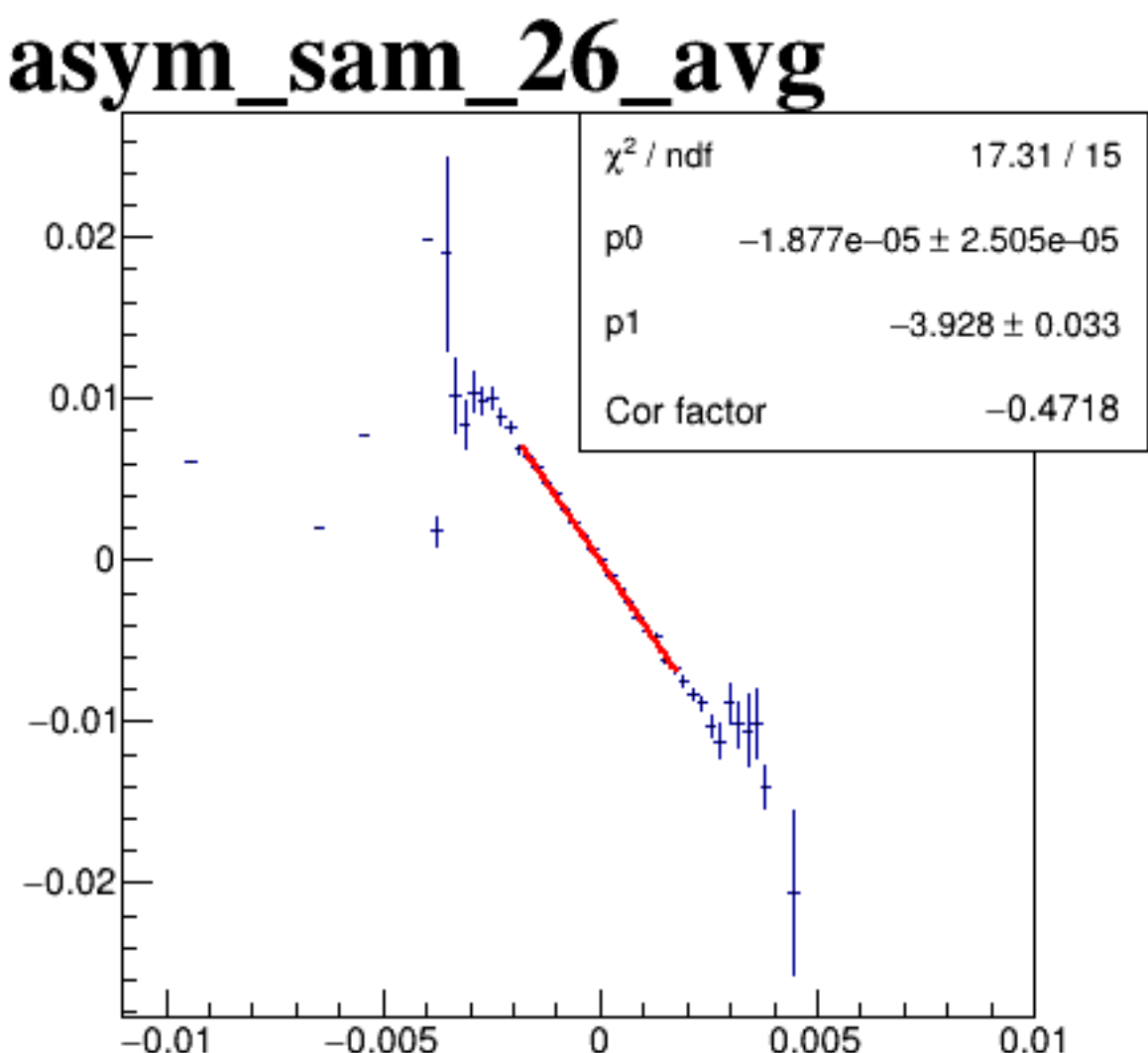
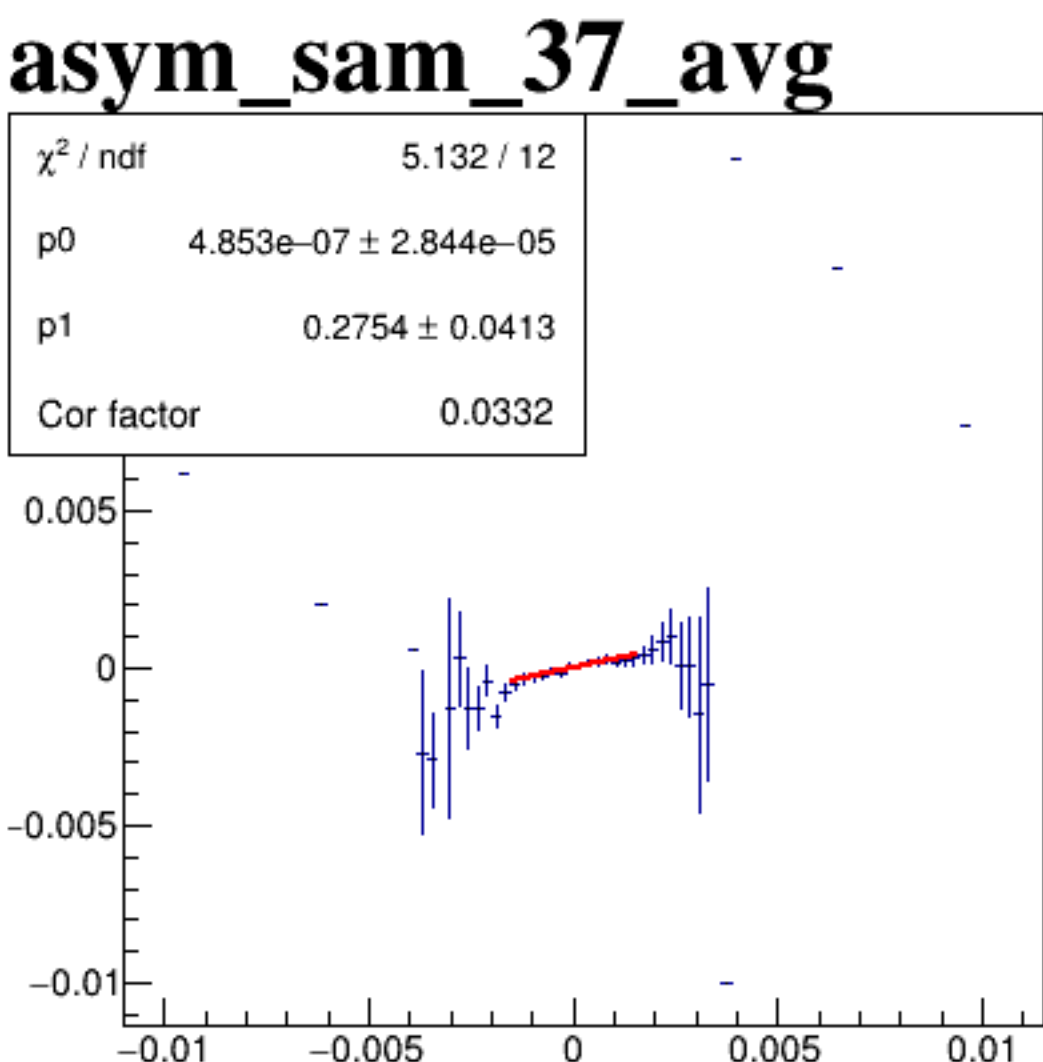
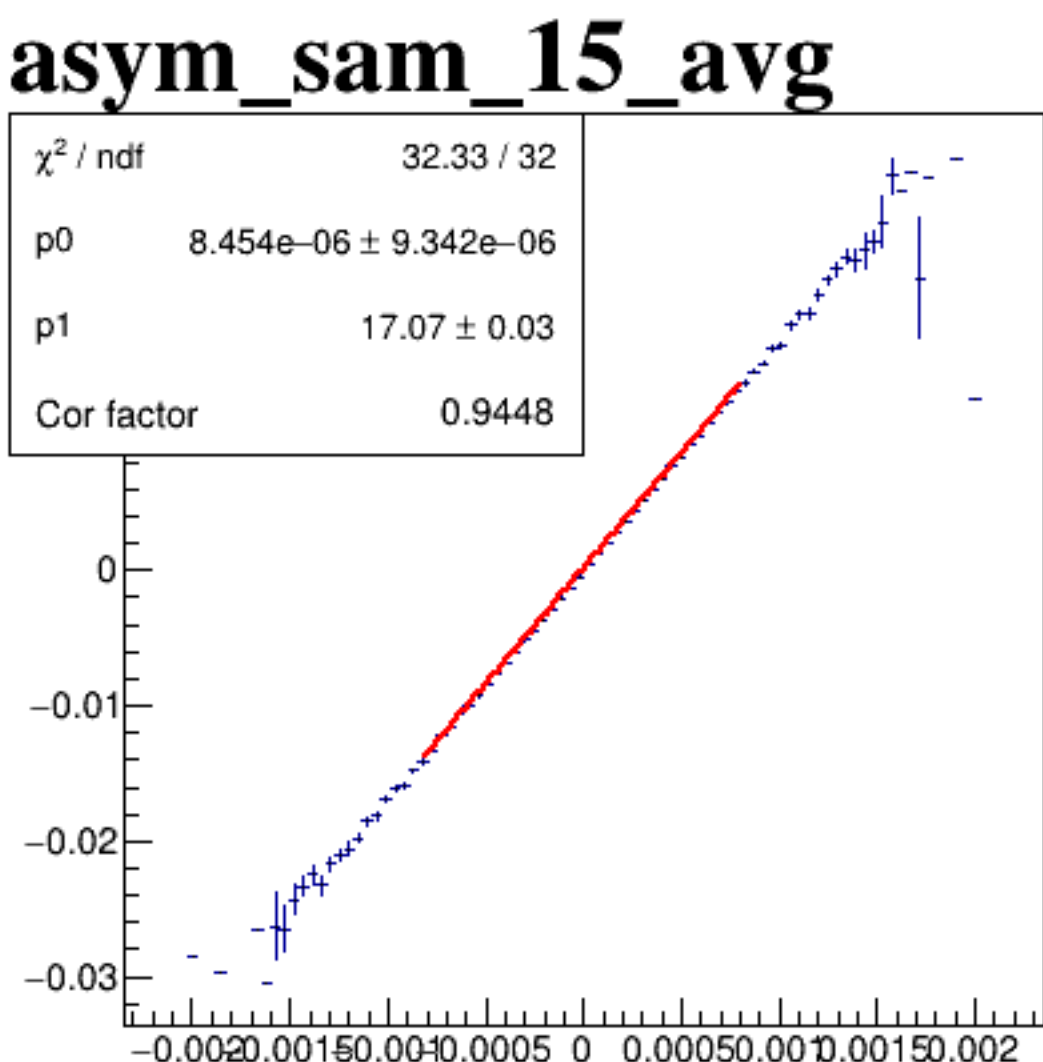
asym_sam_48_avg



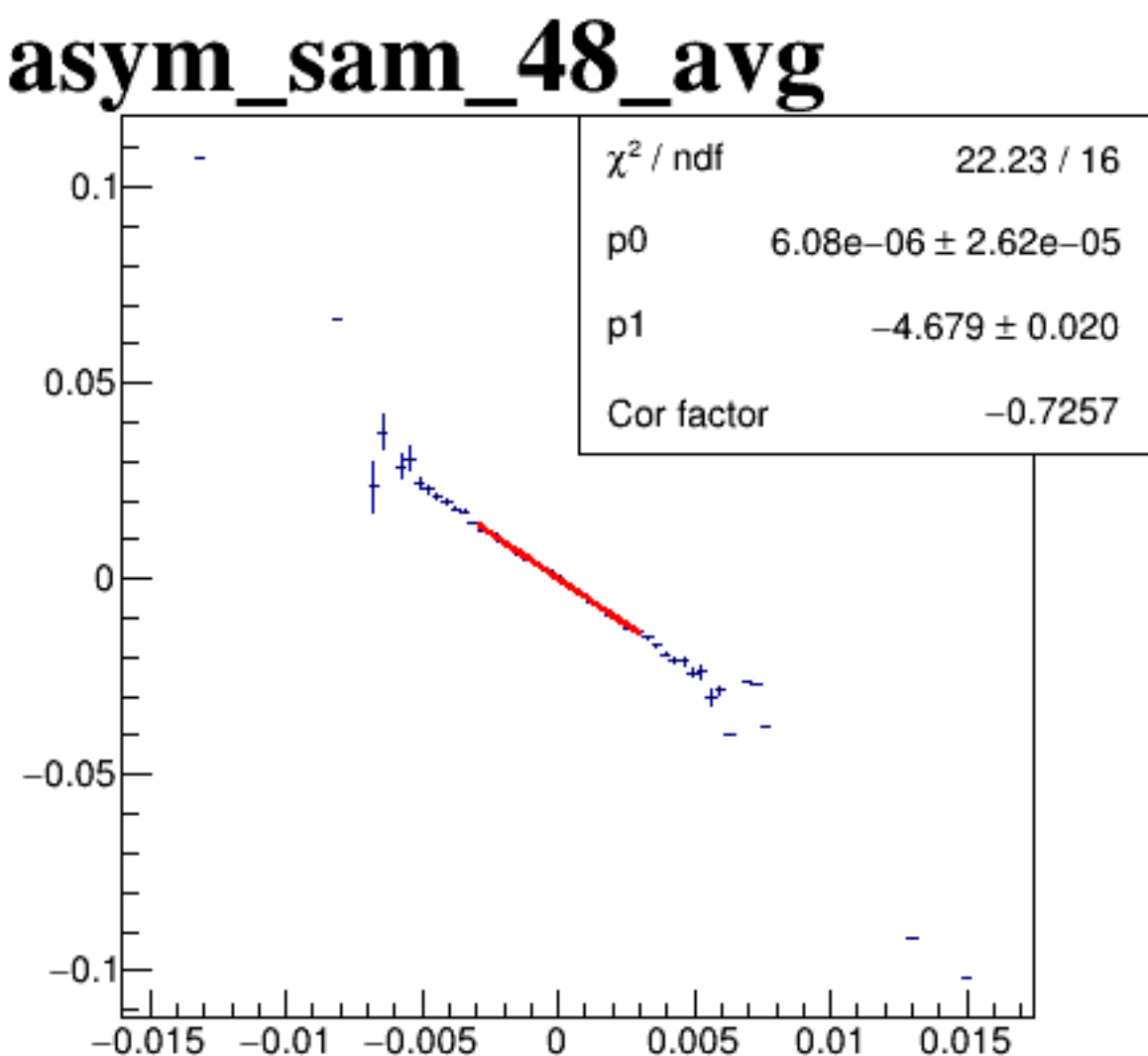
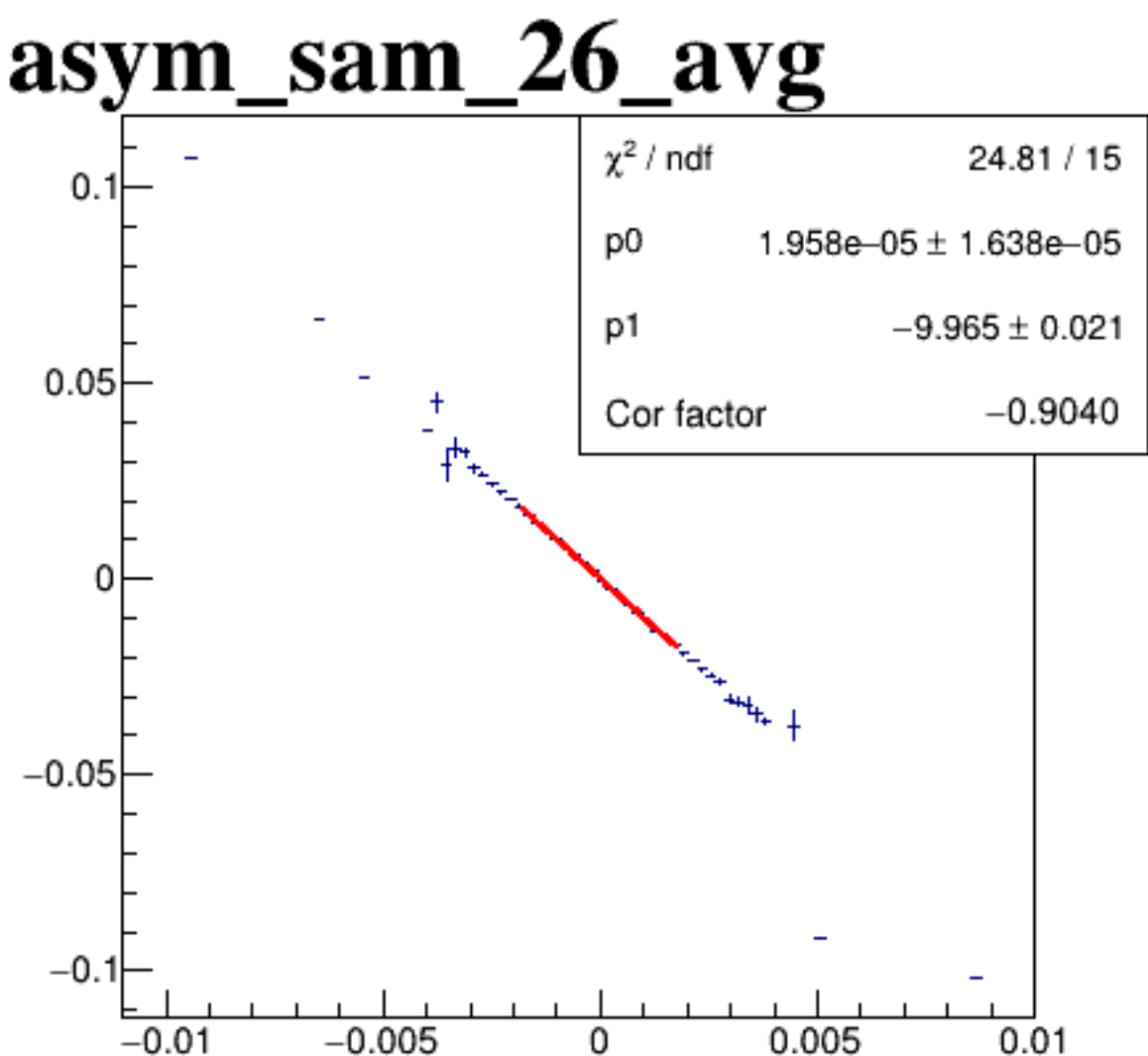
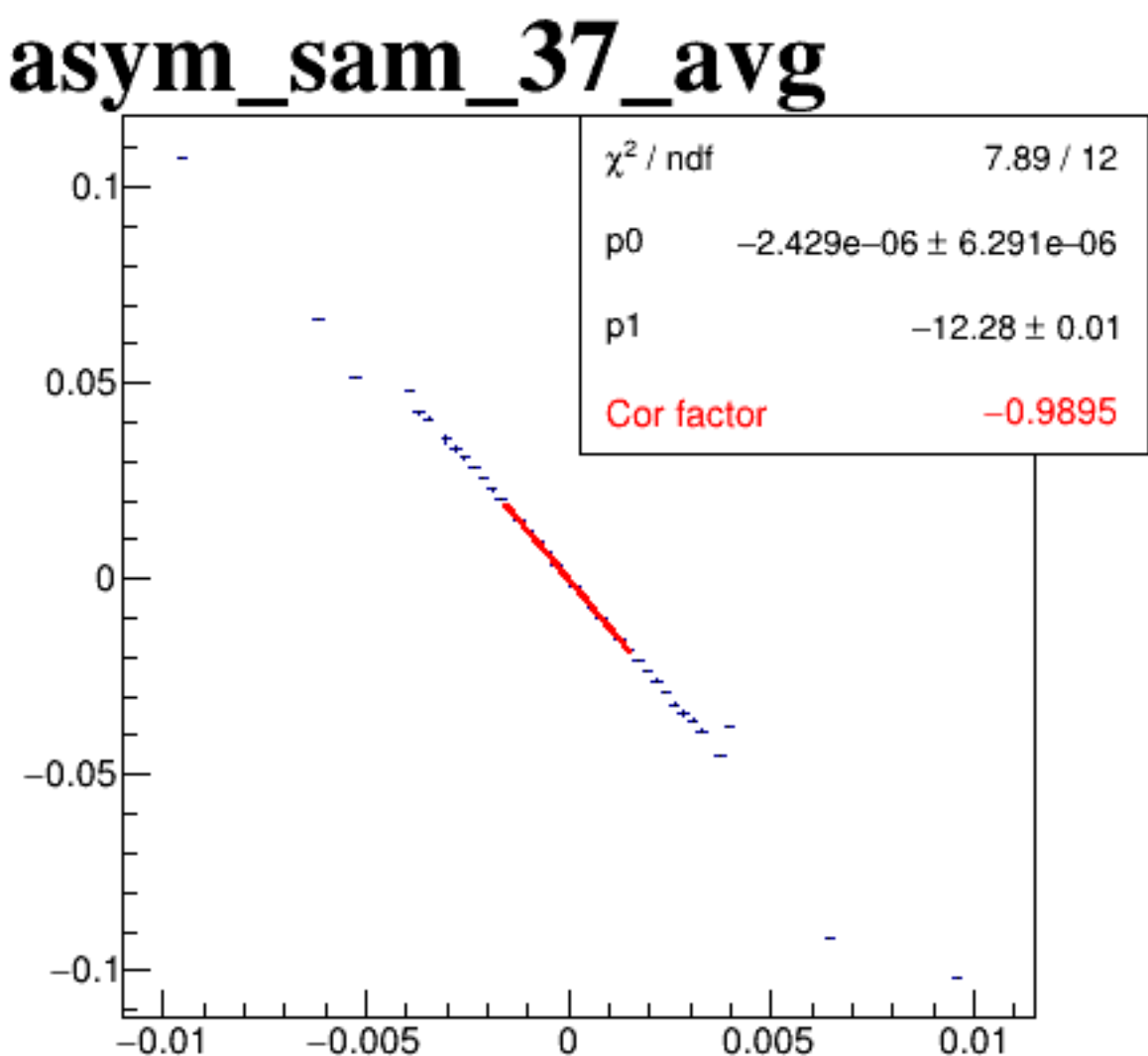
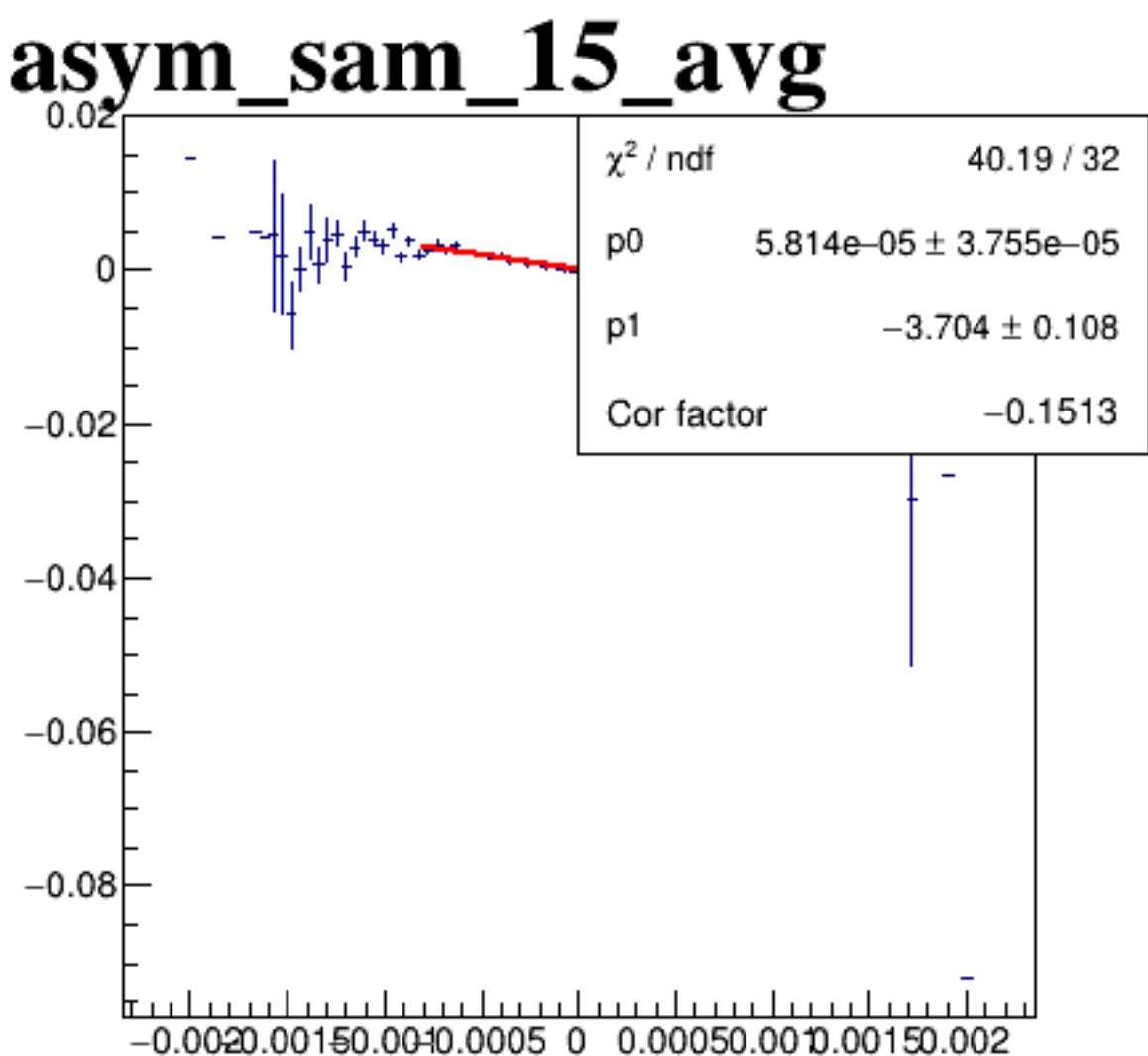
diff_bpm4aX



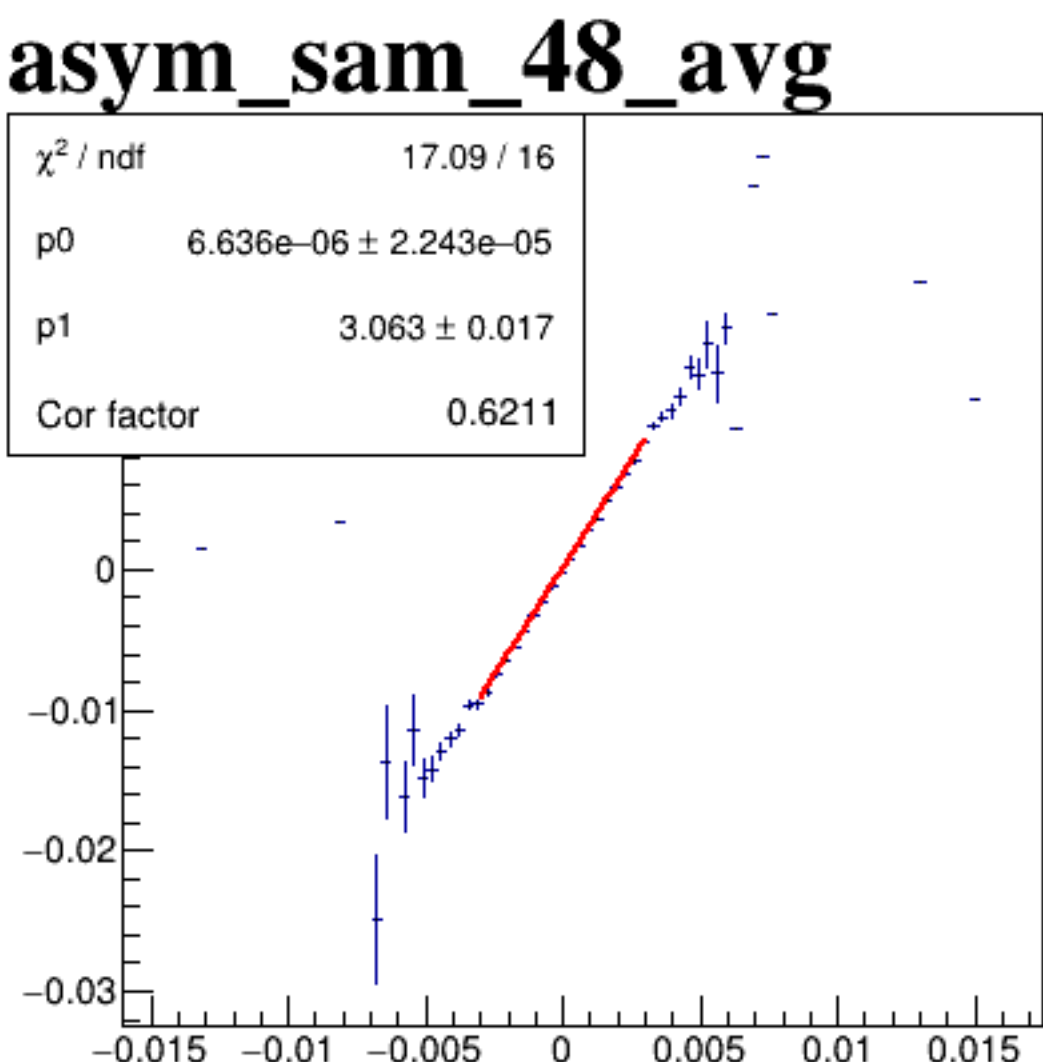
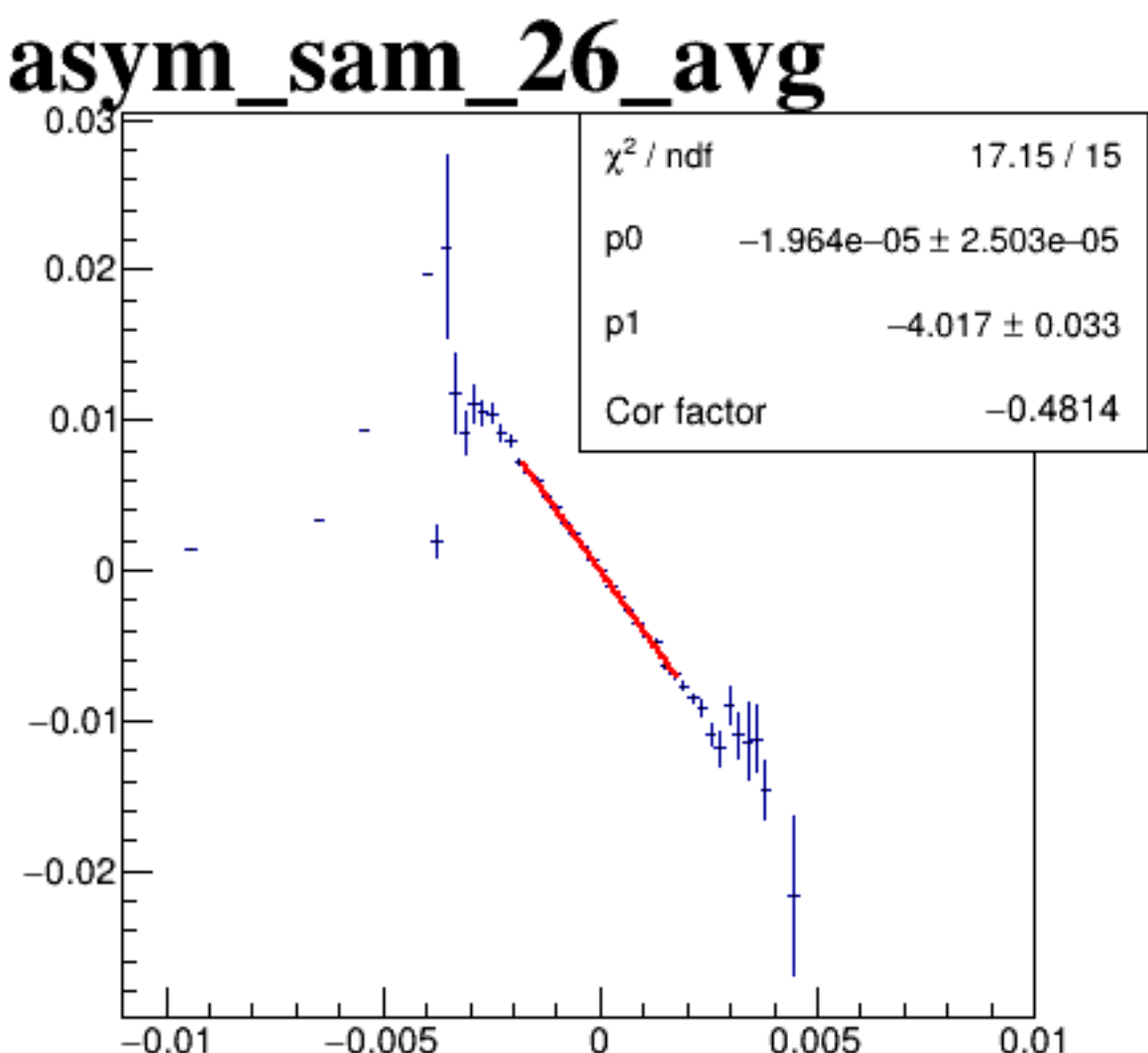
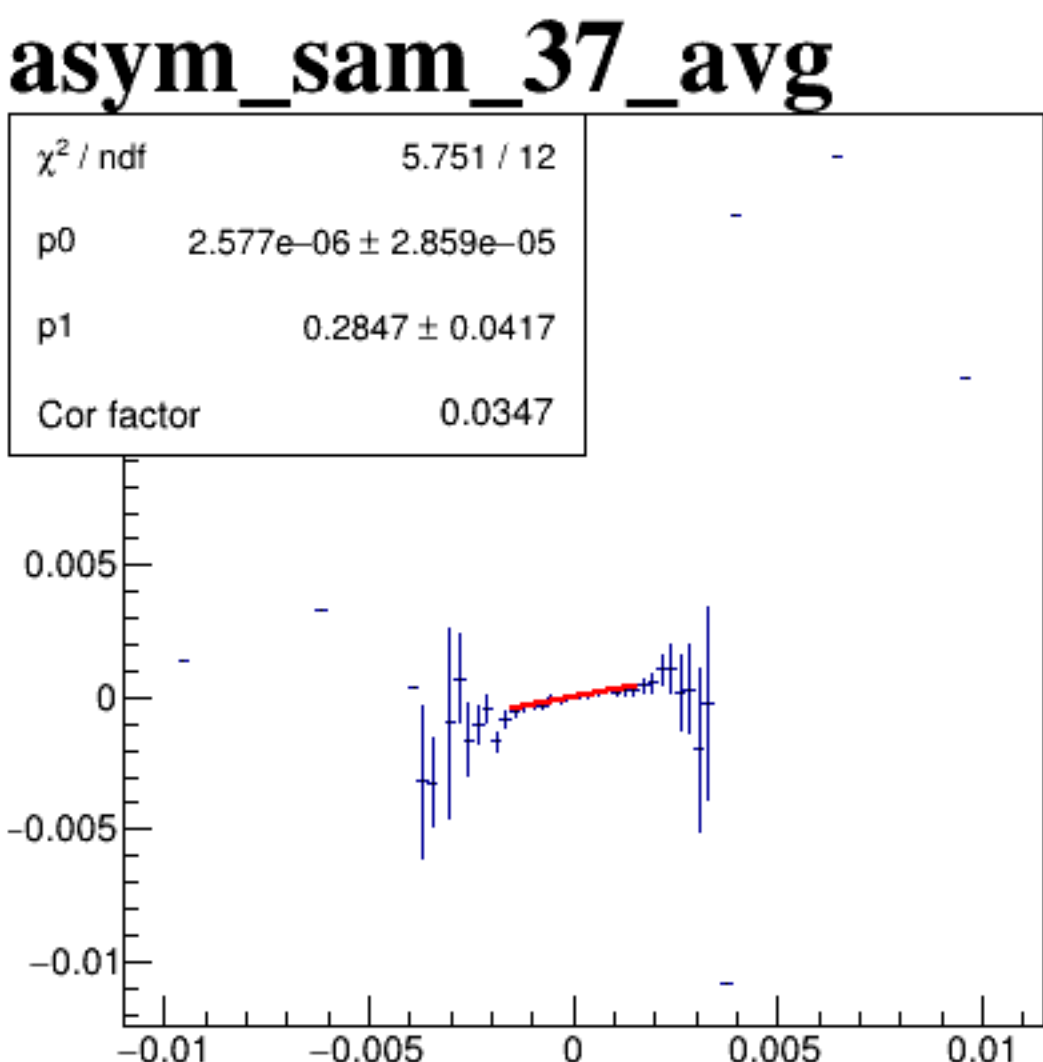
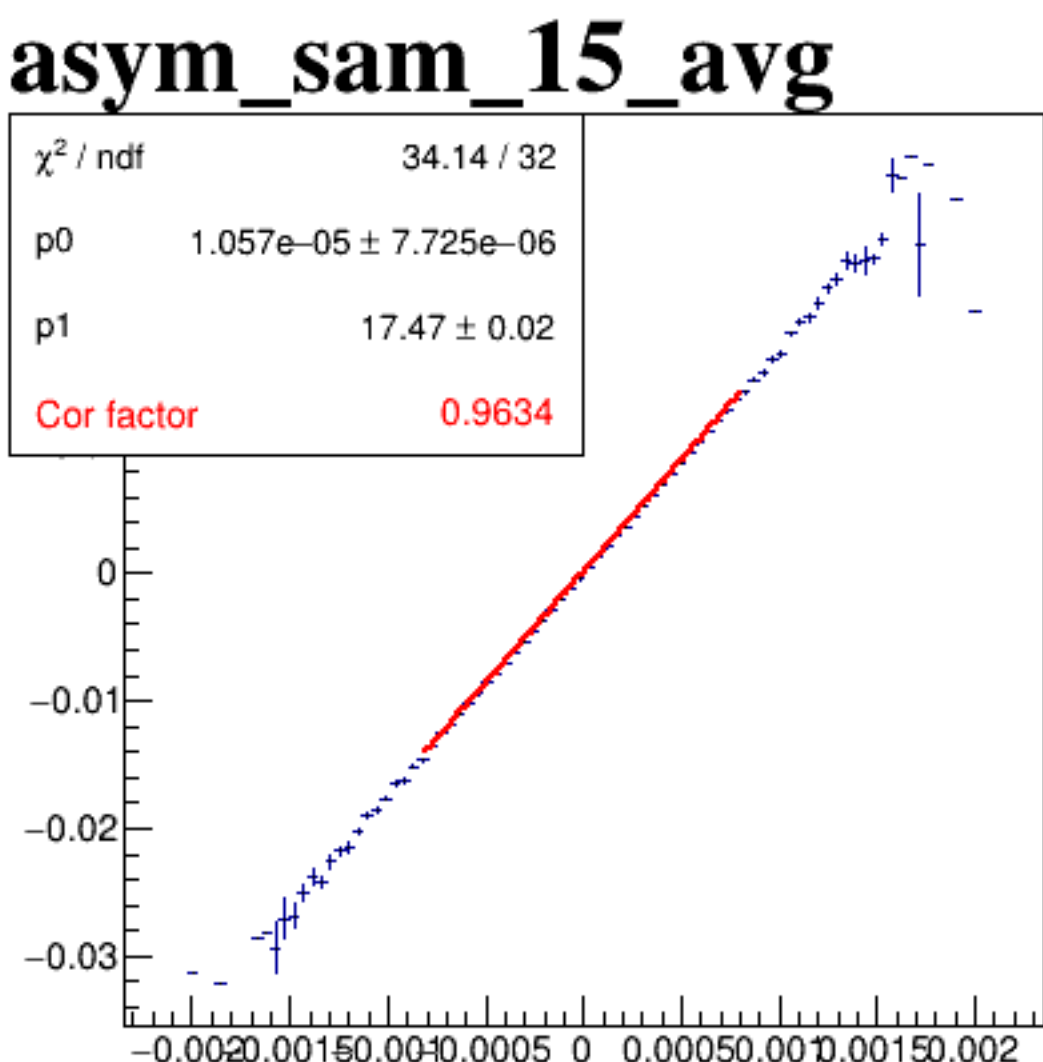
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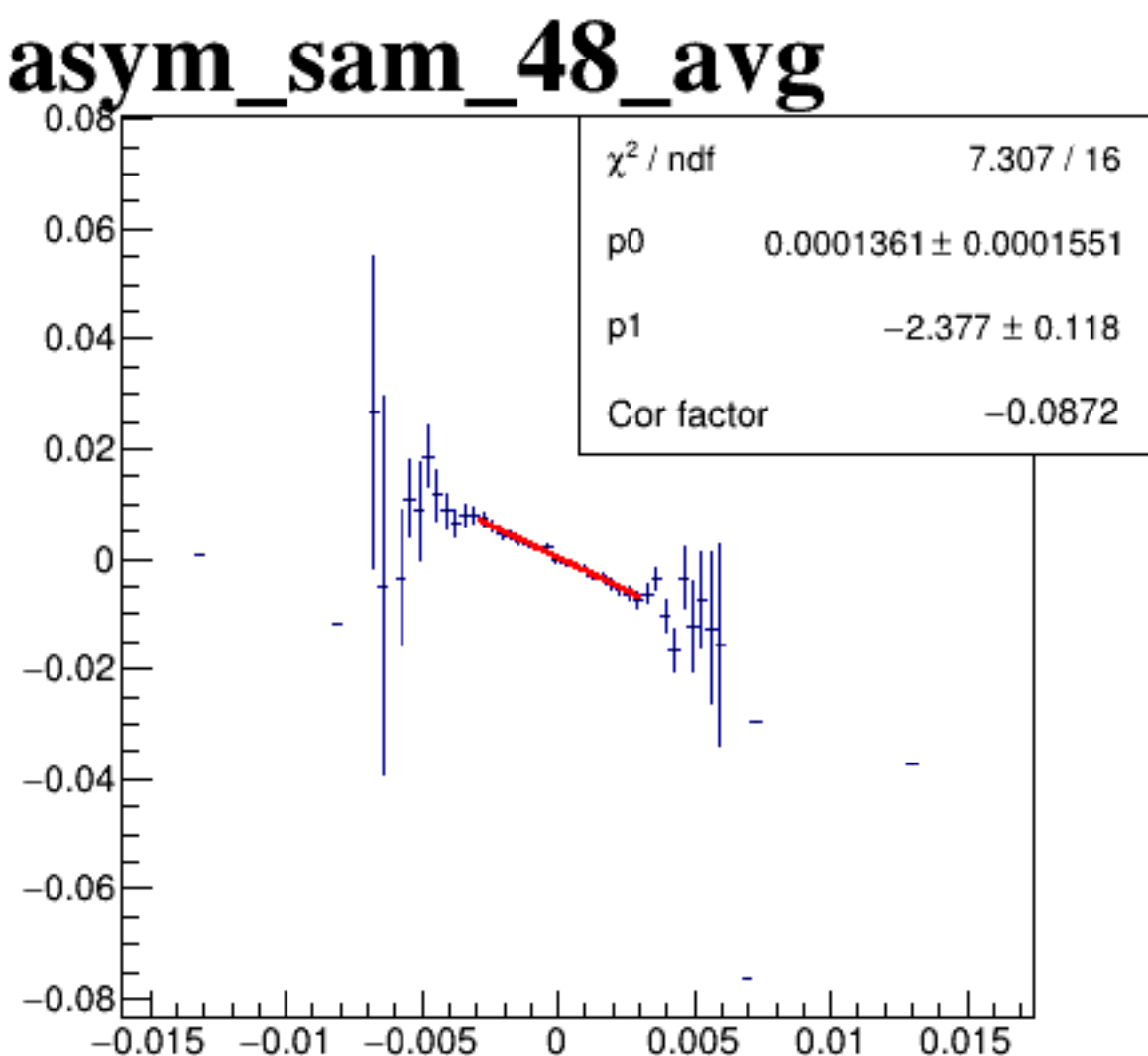
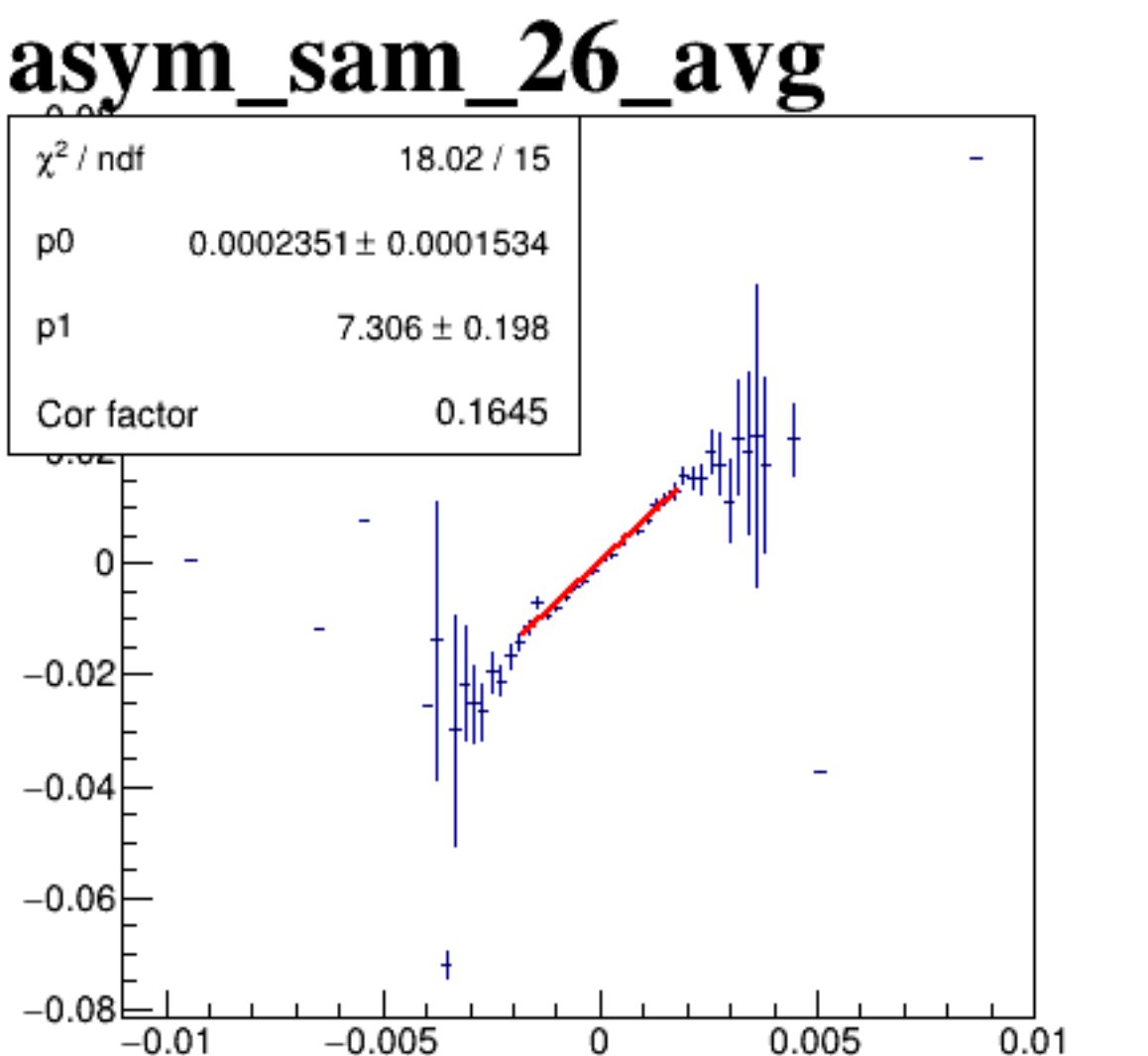
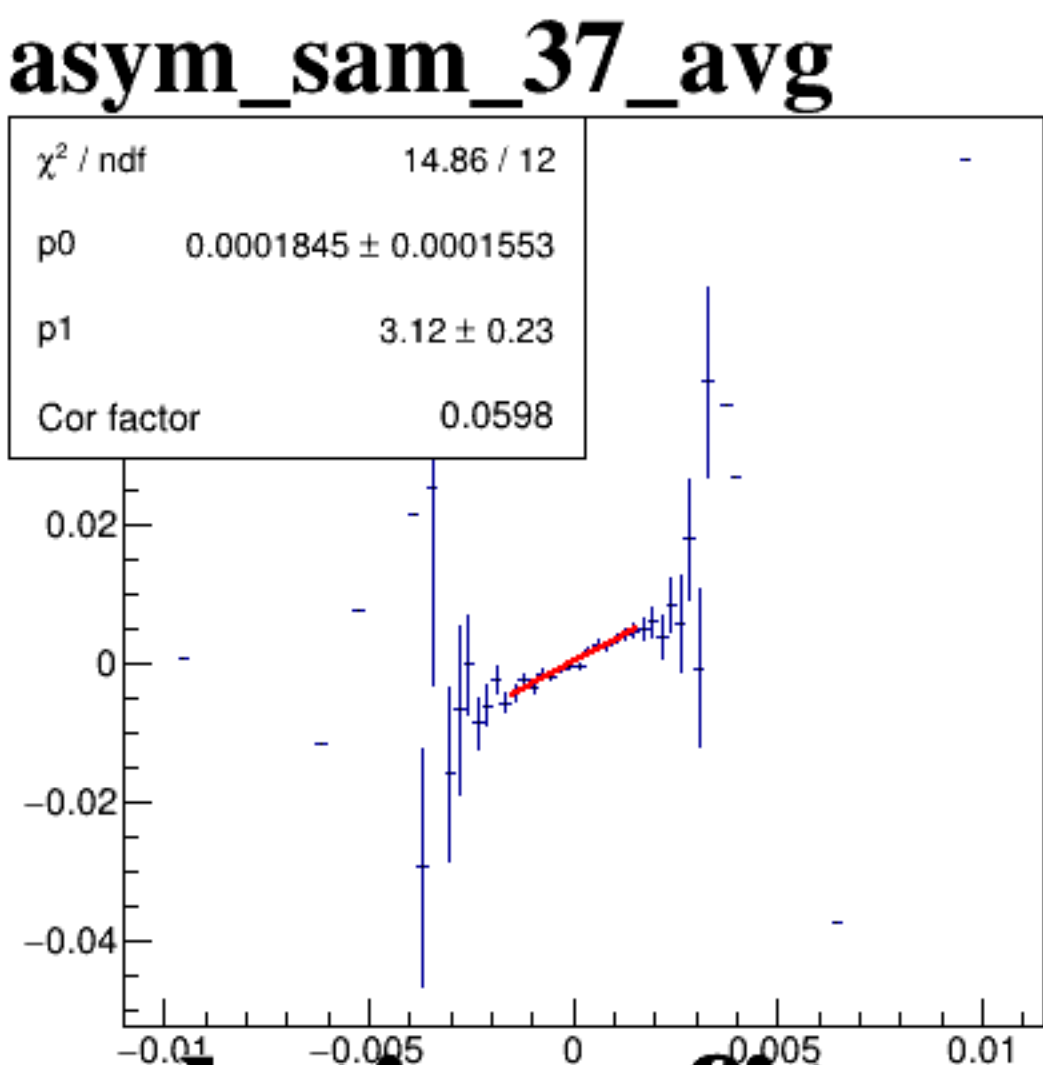
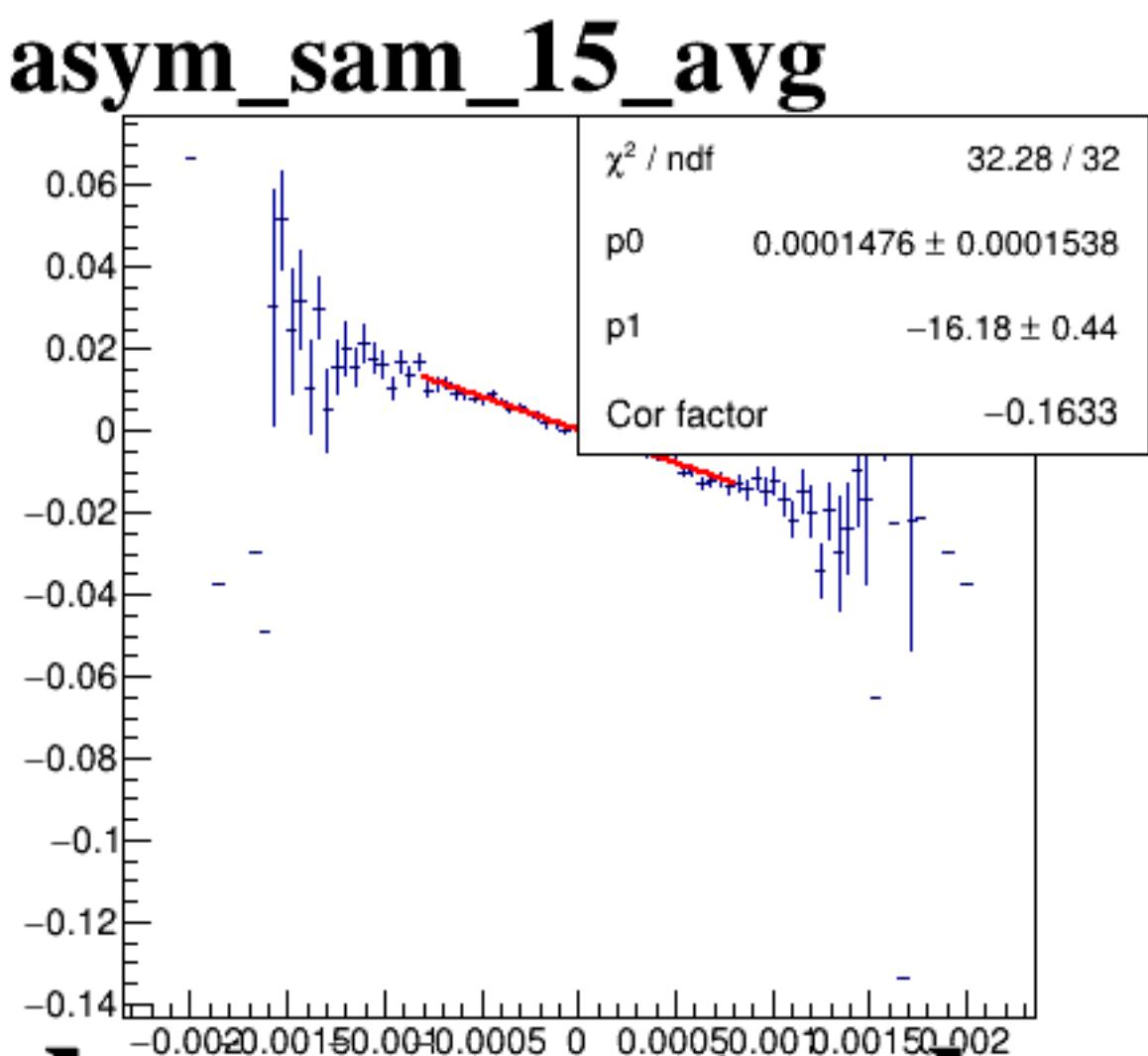
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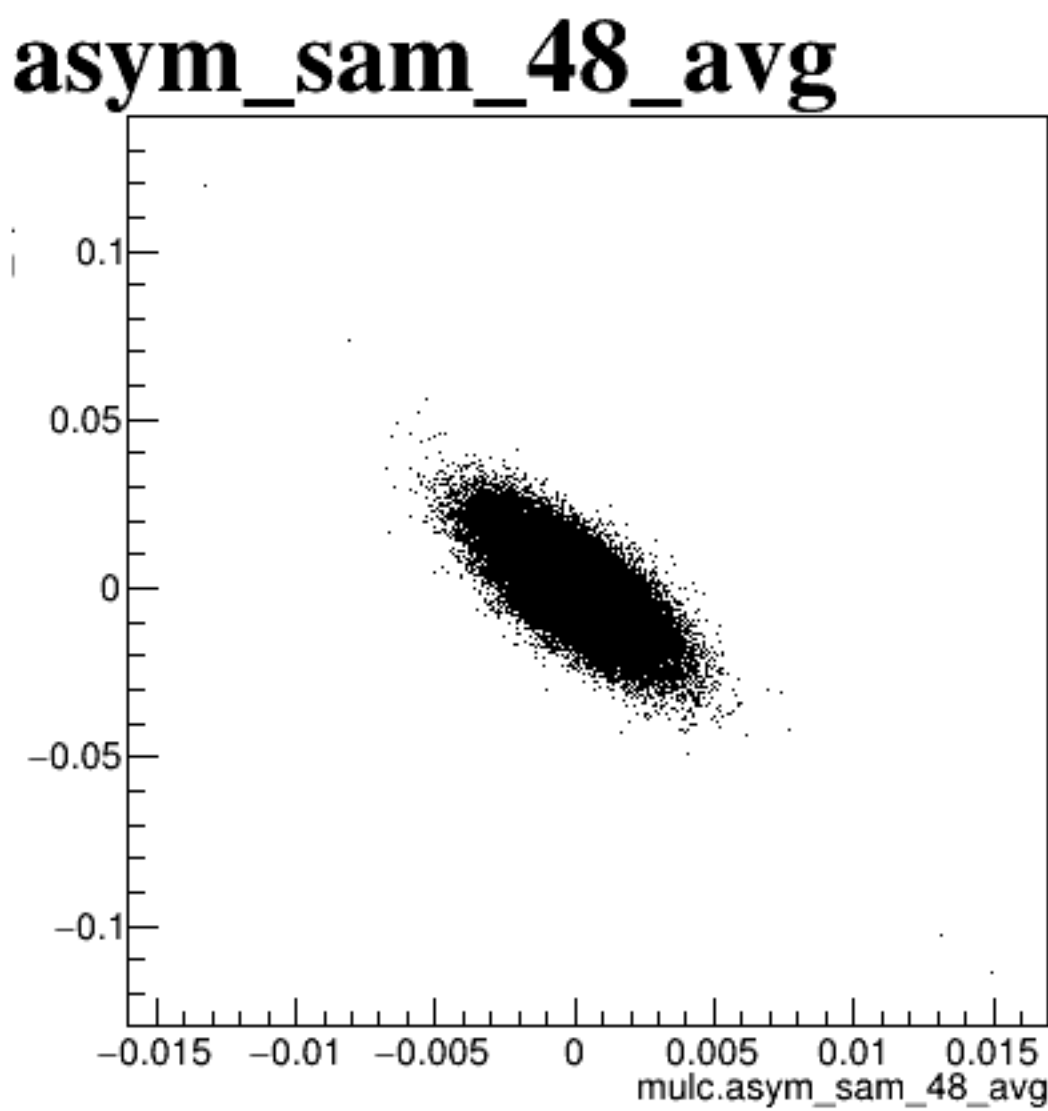
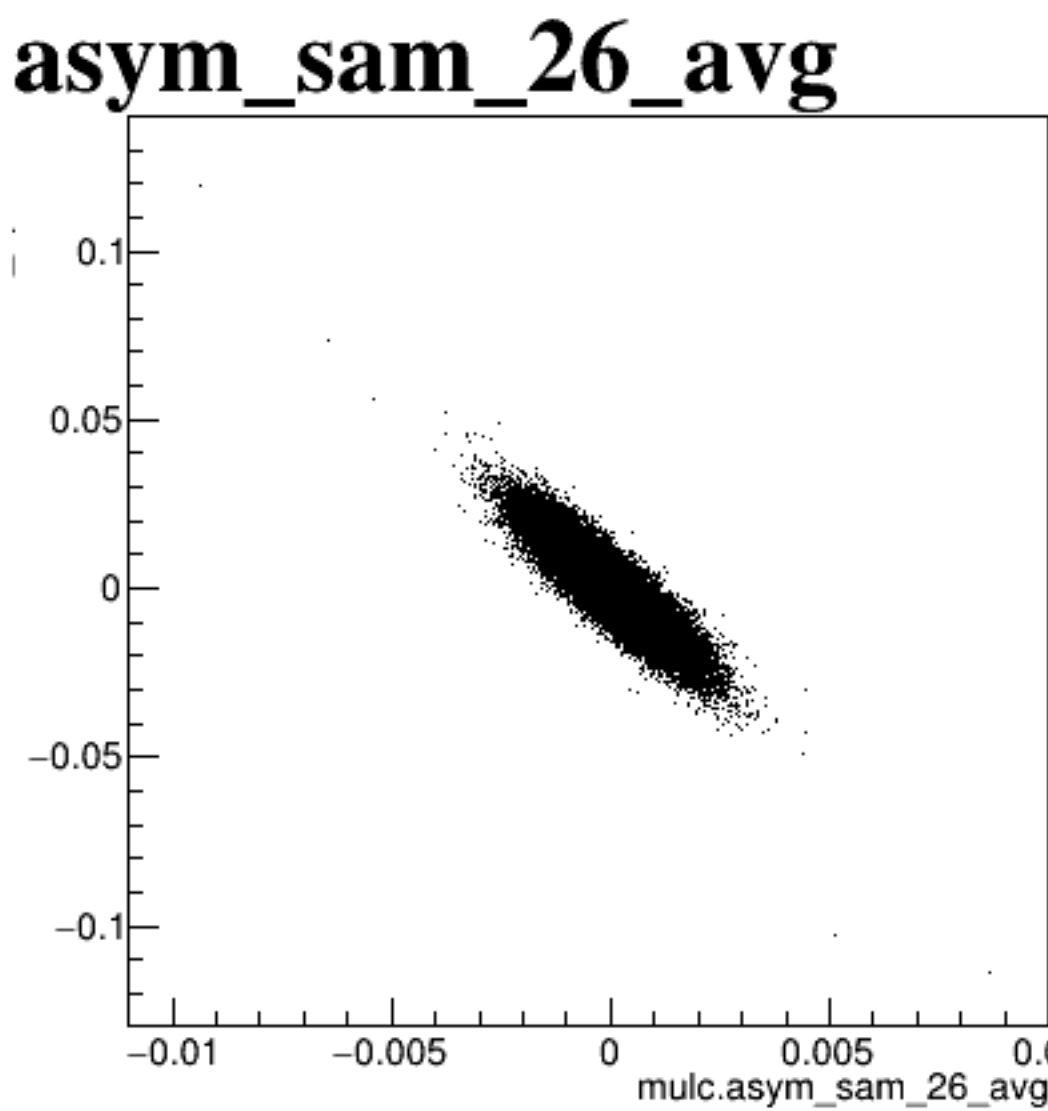
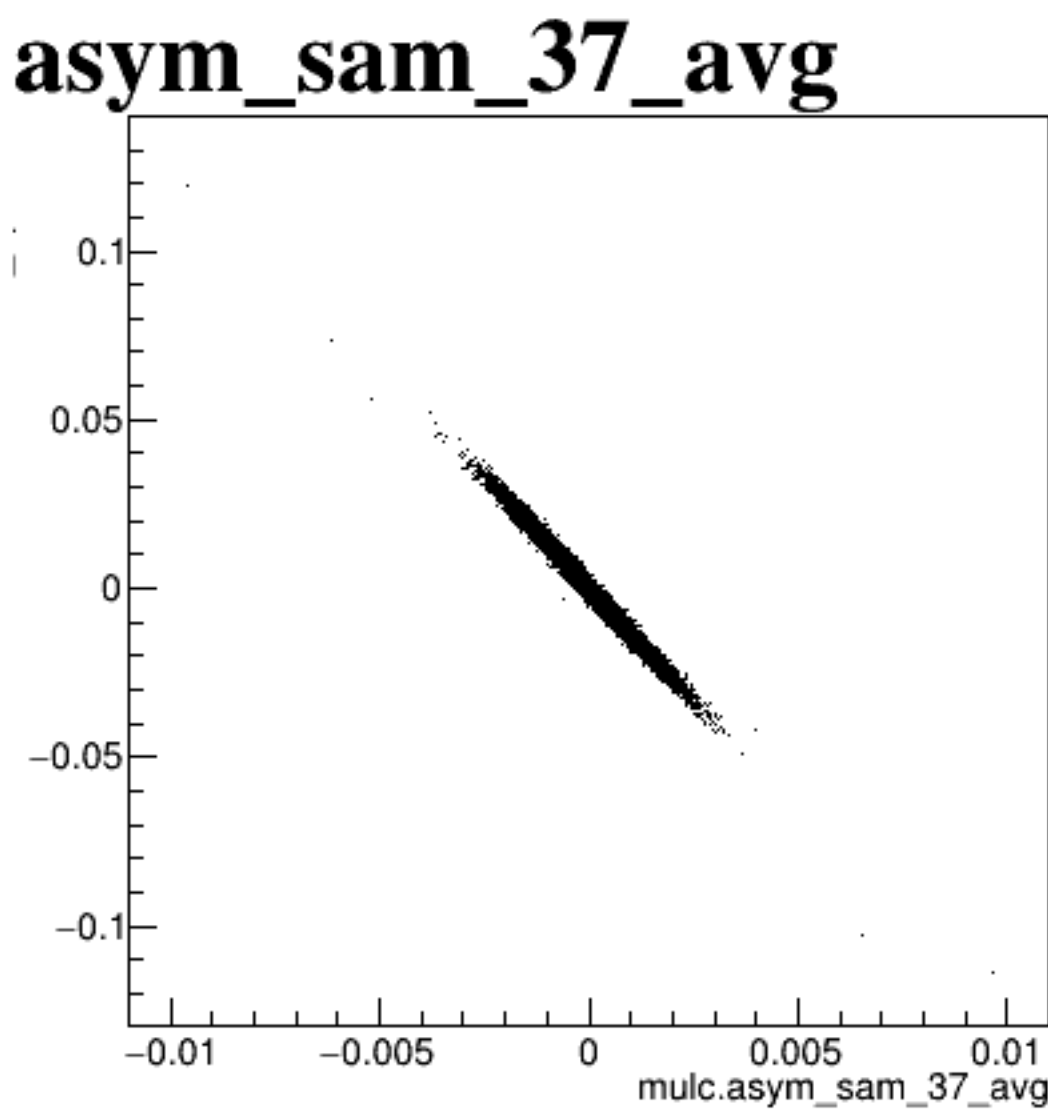
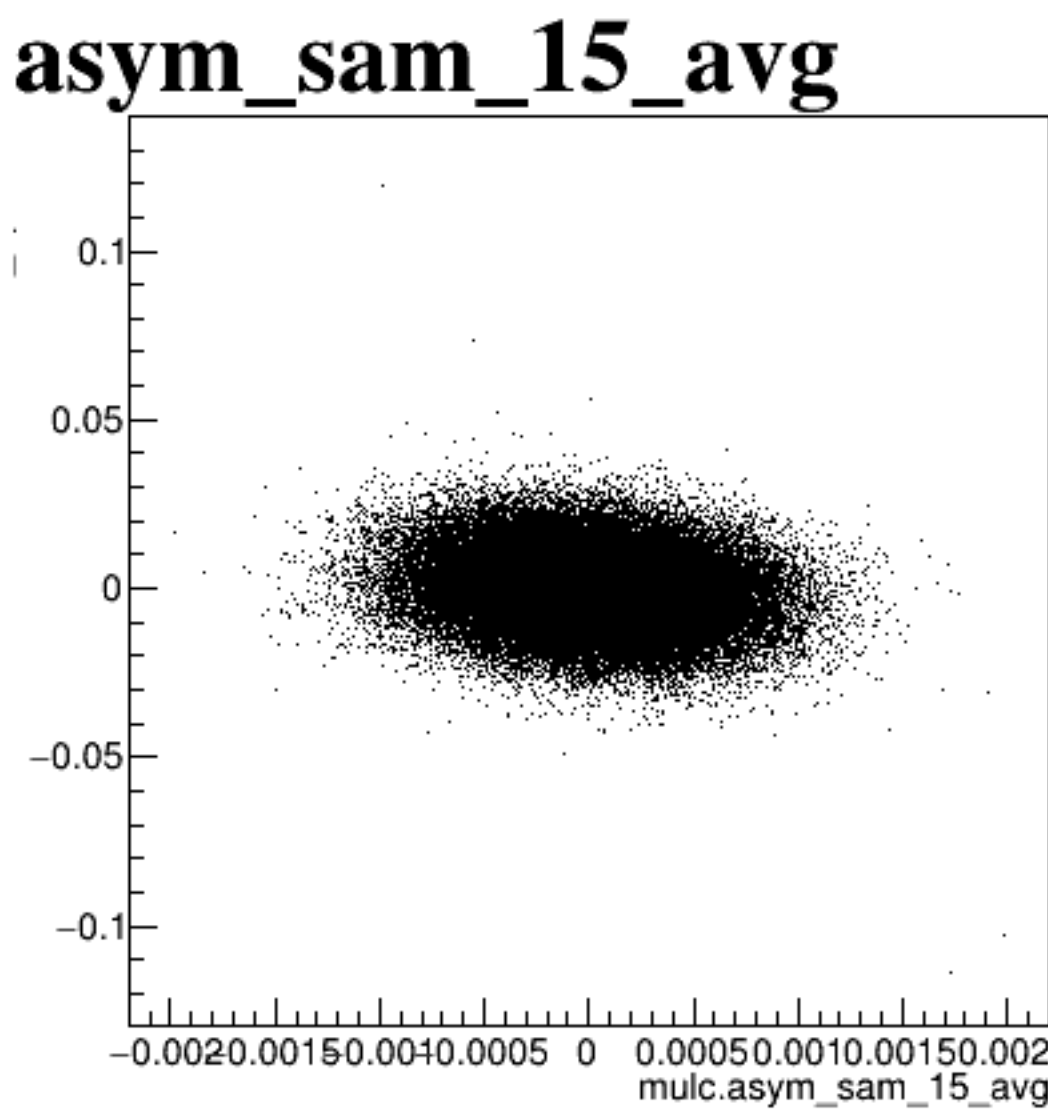
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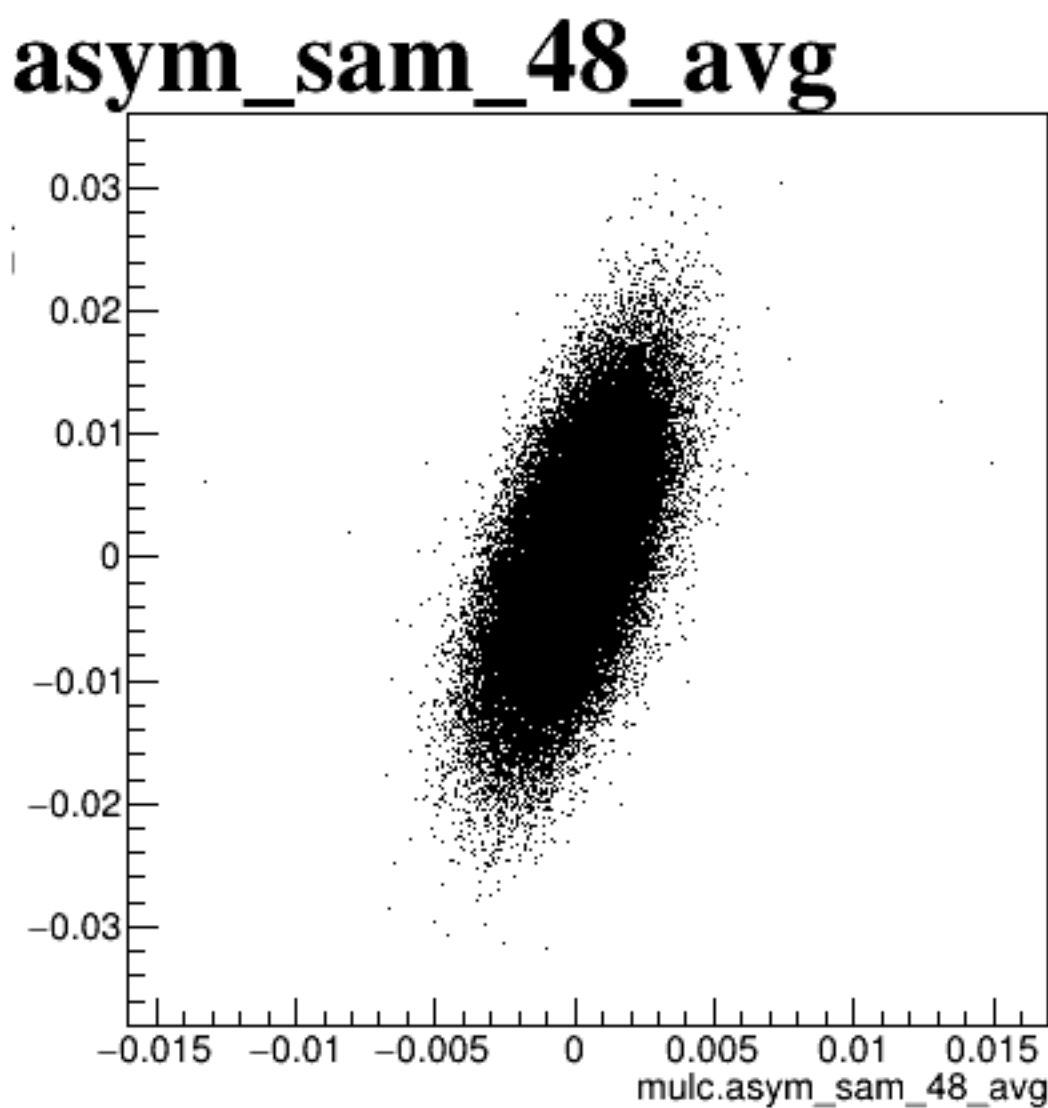
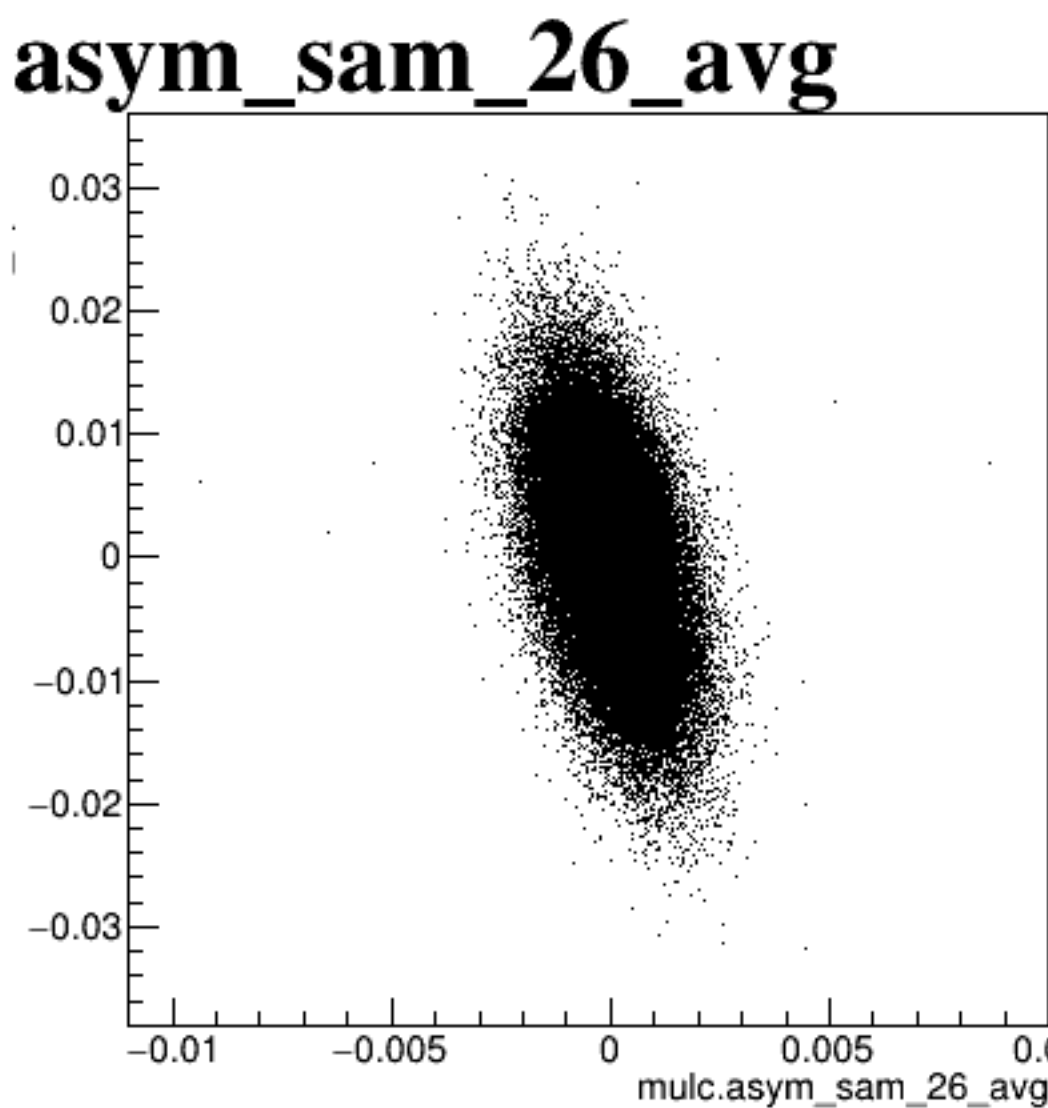
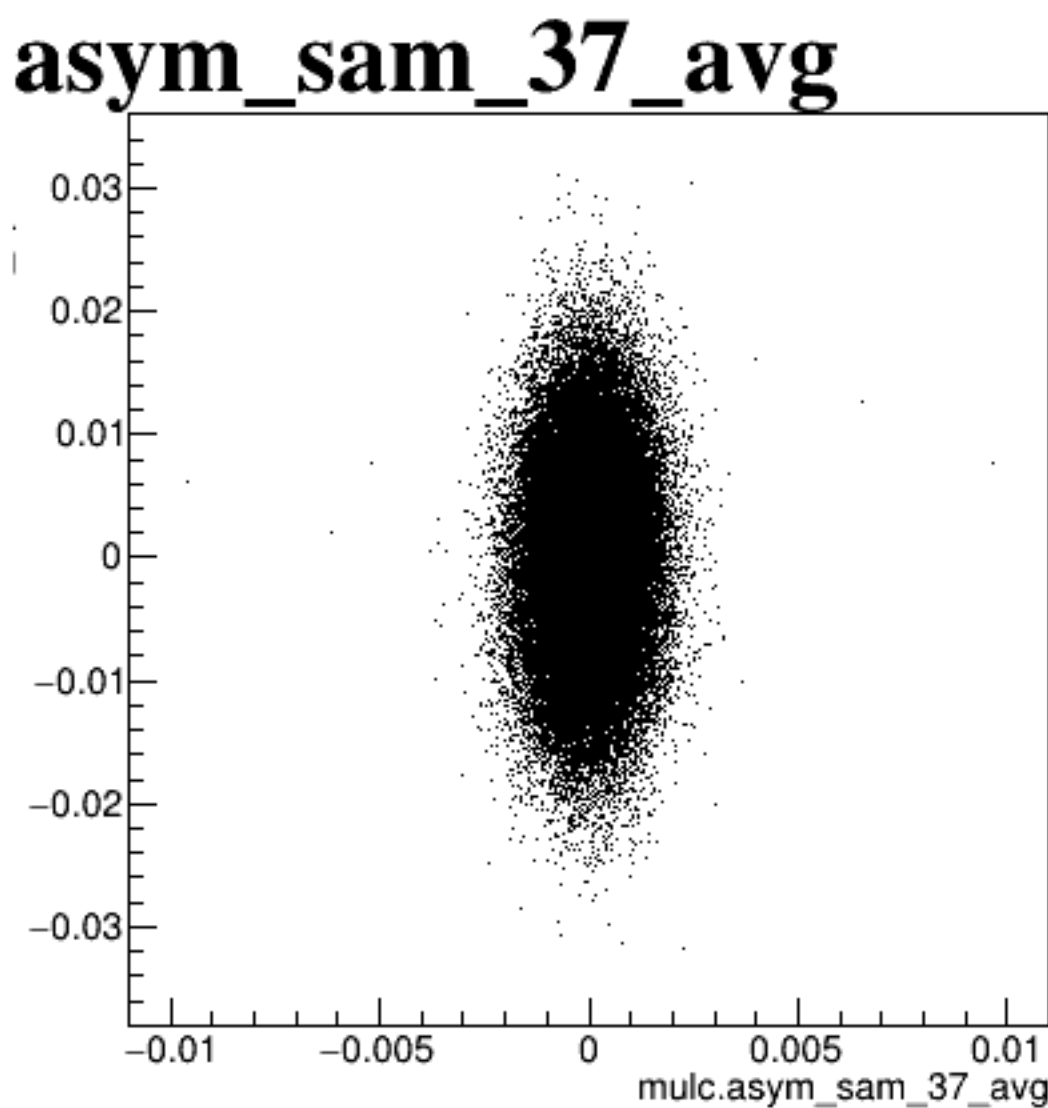
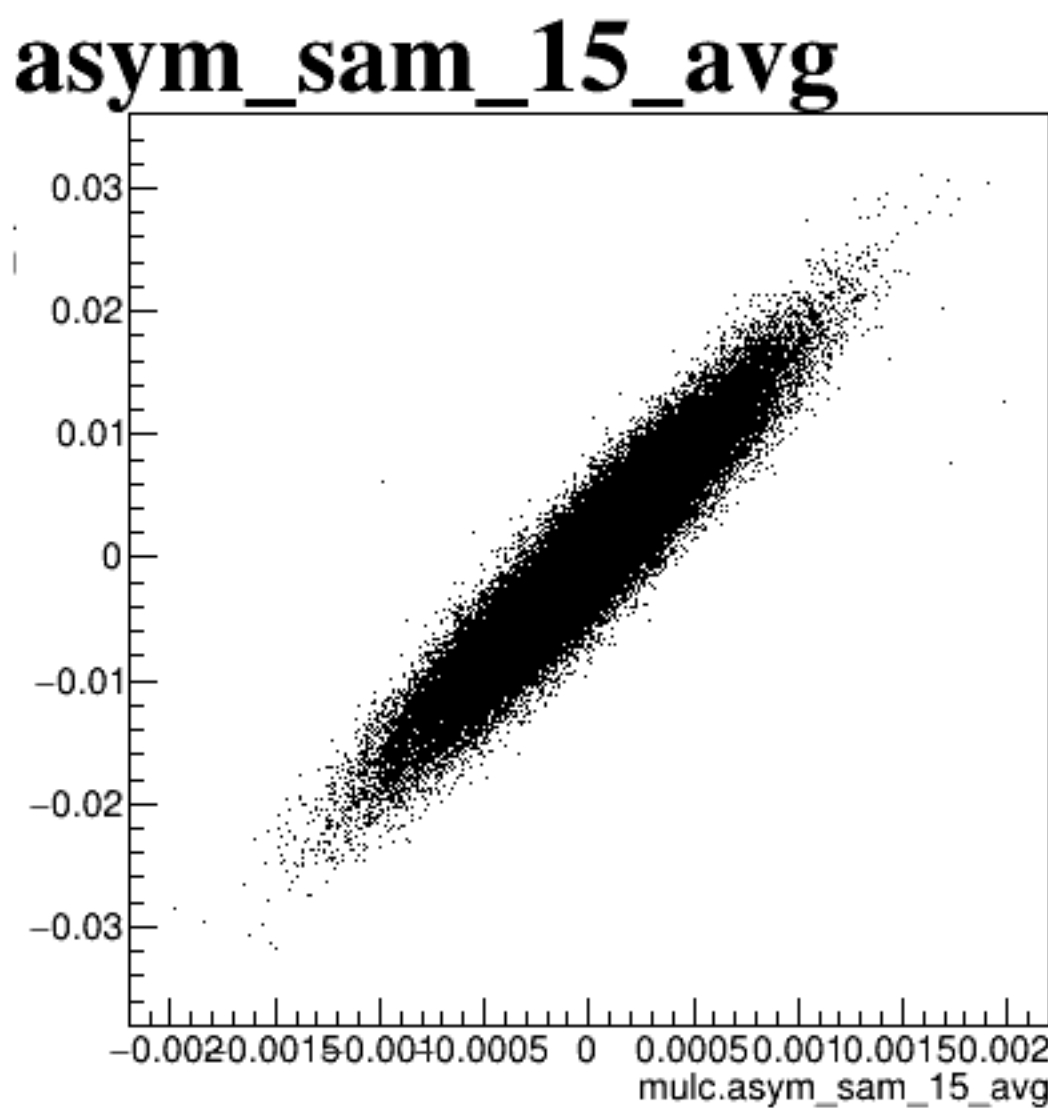
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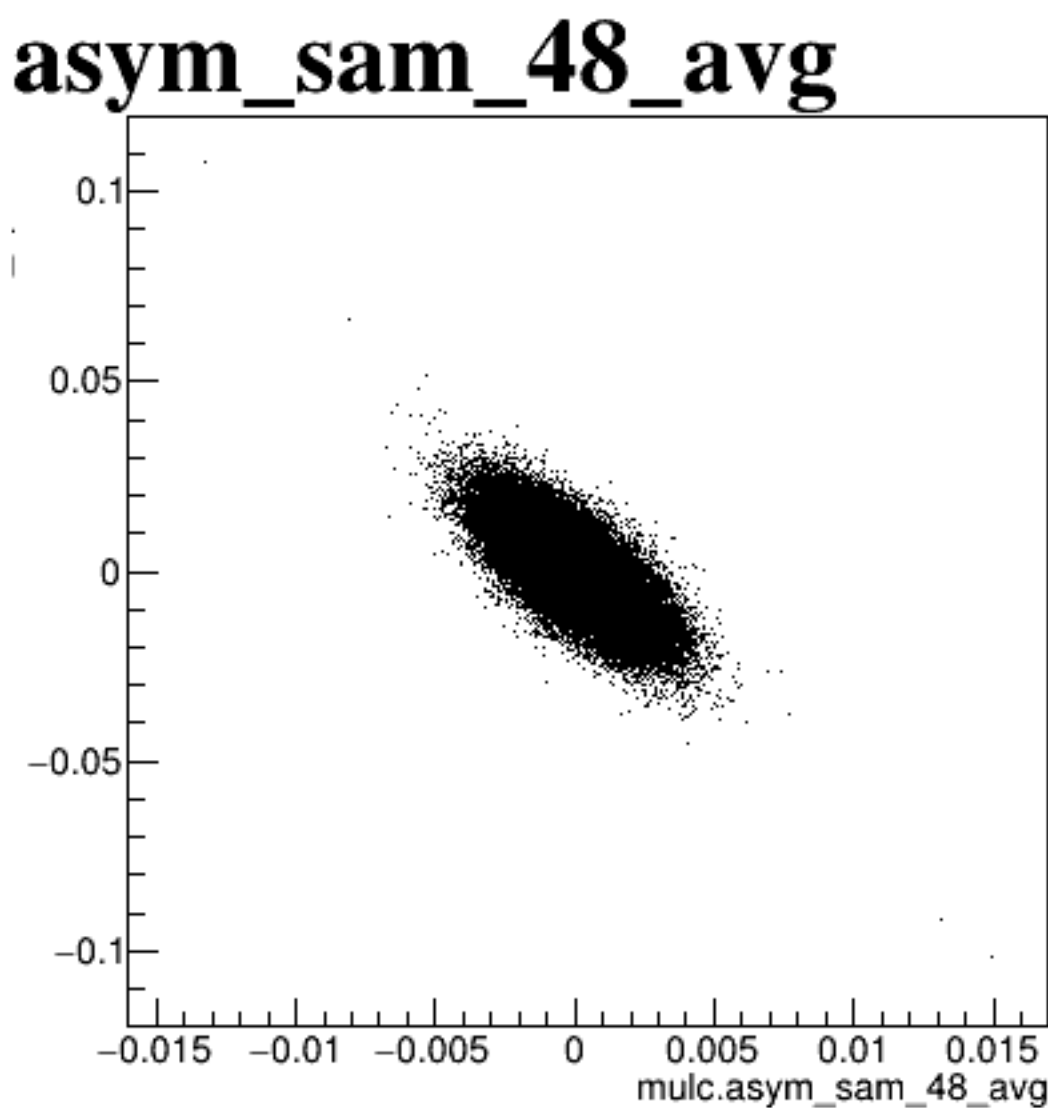
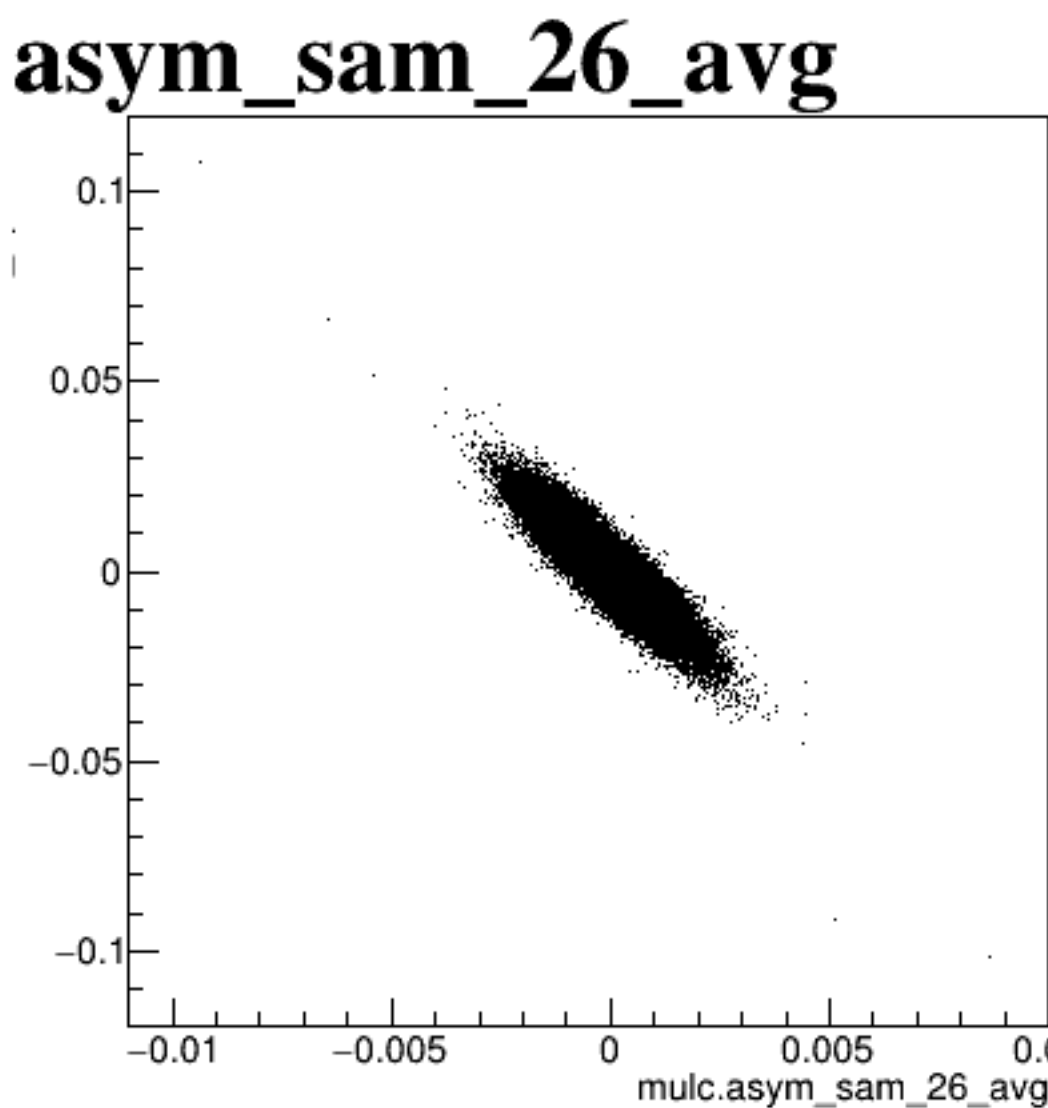
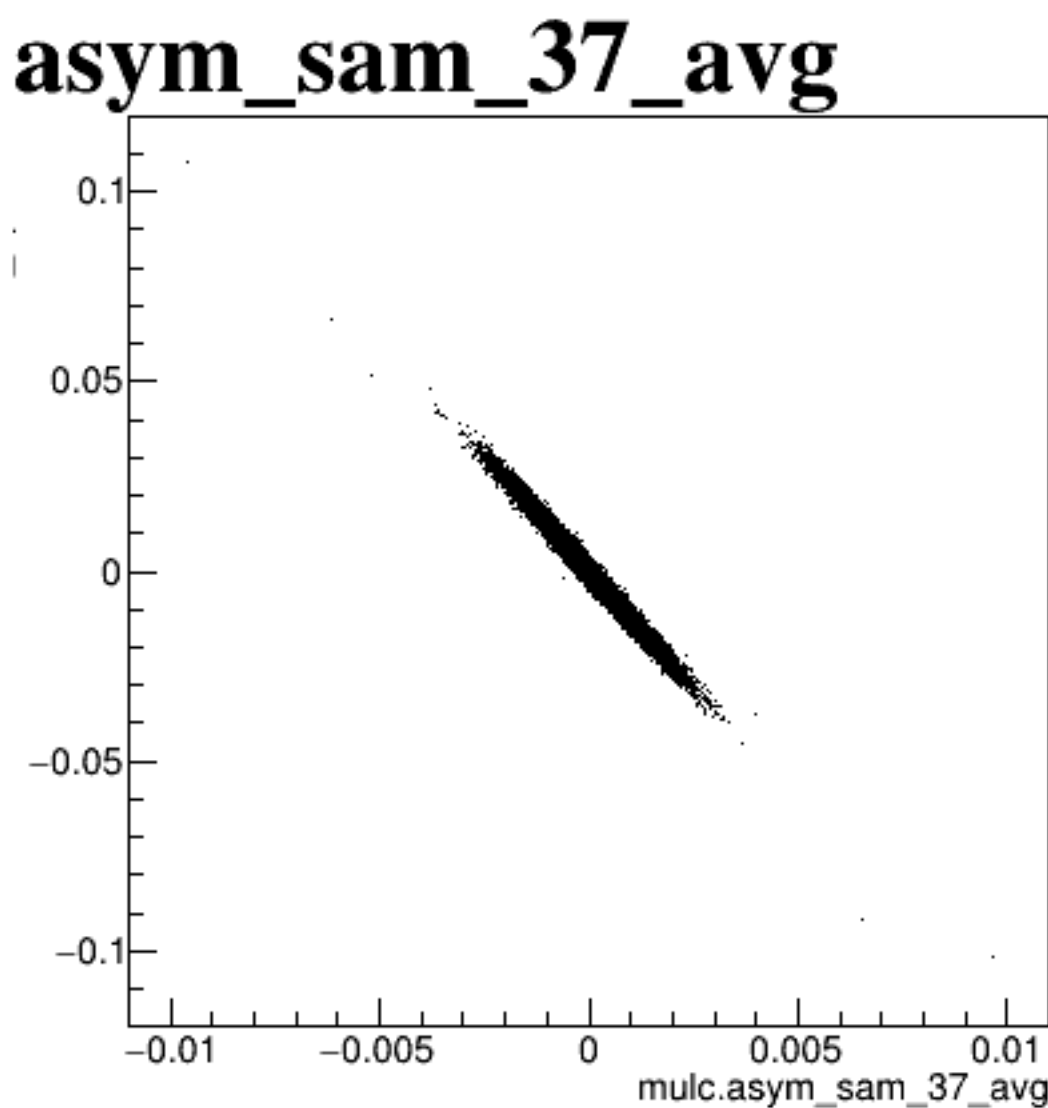
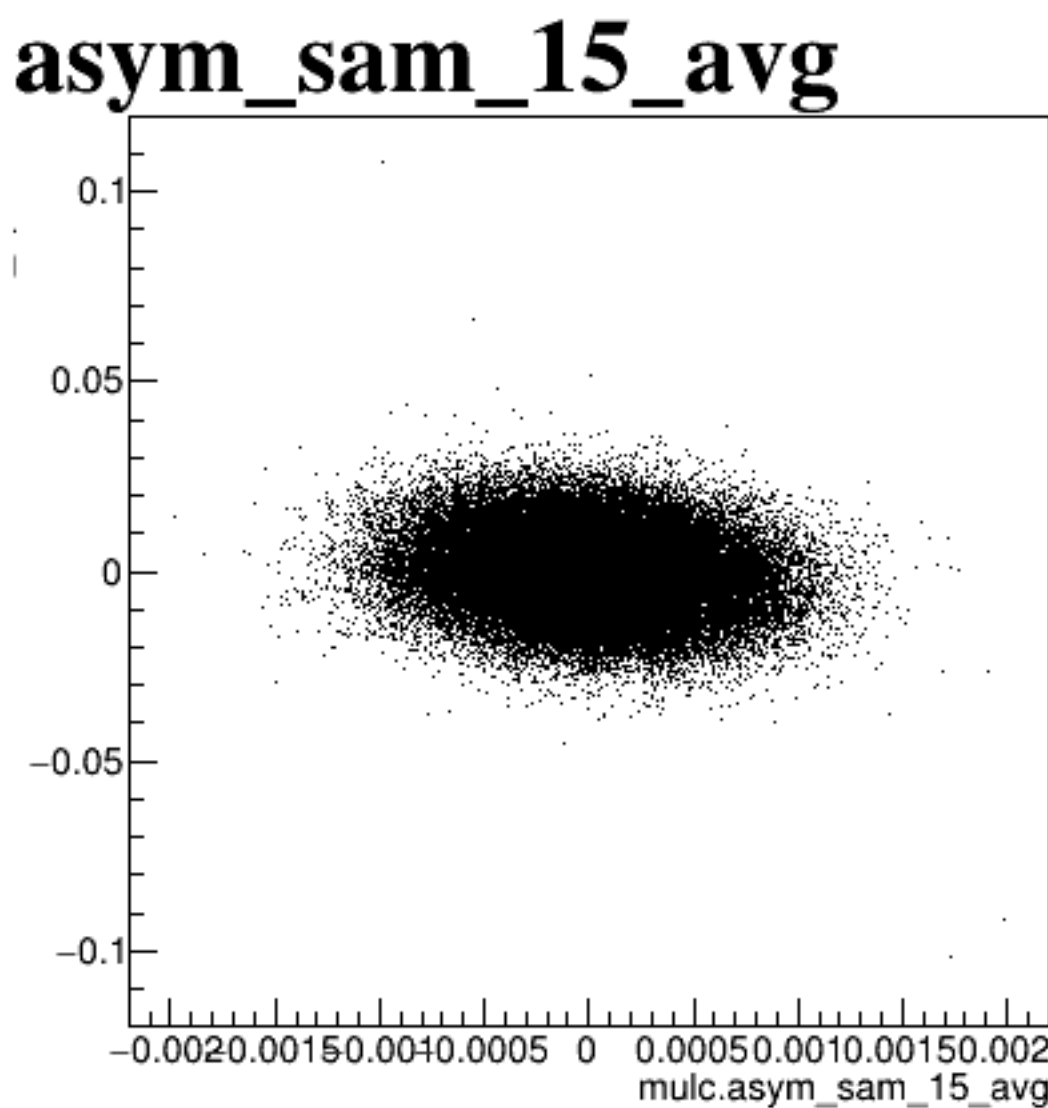
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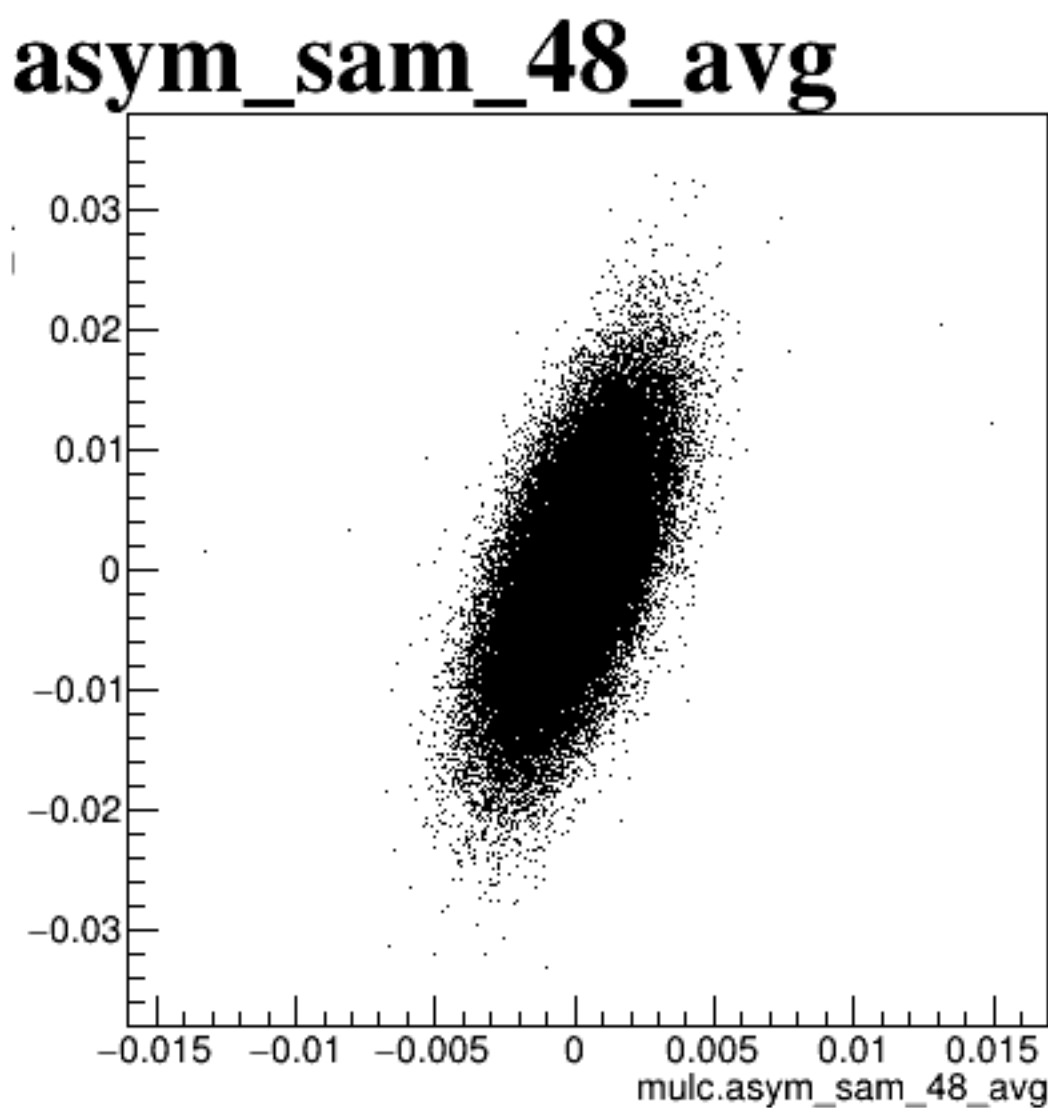
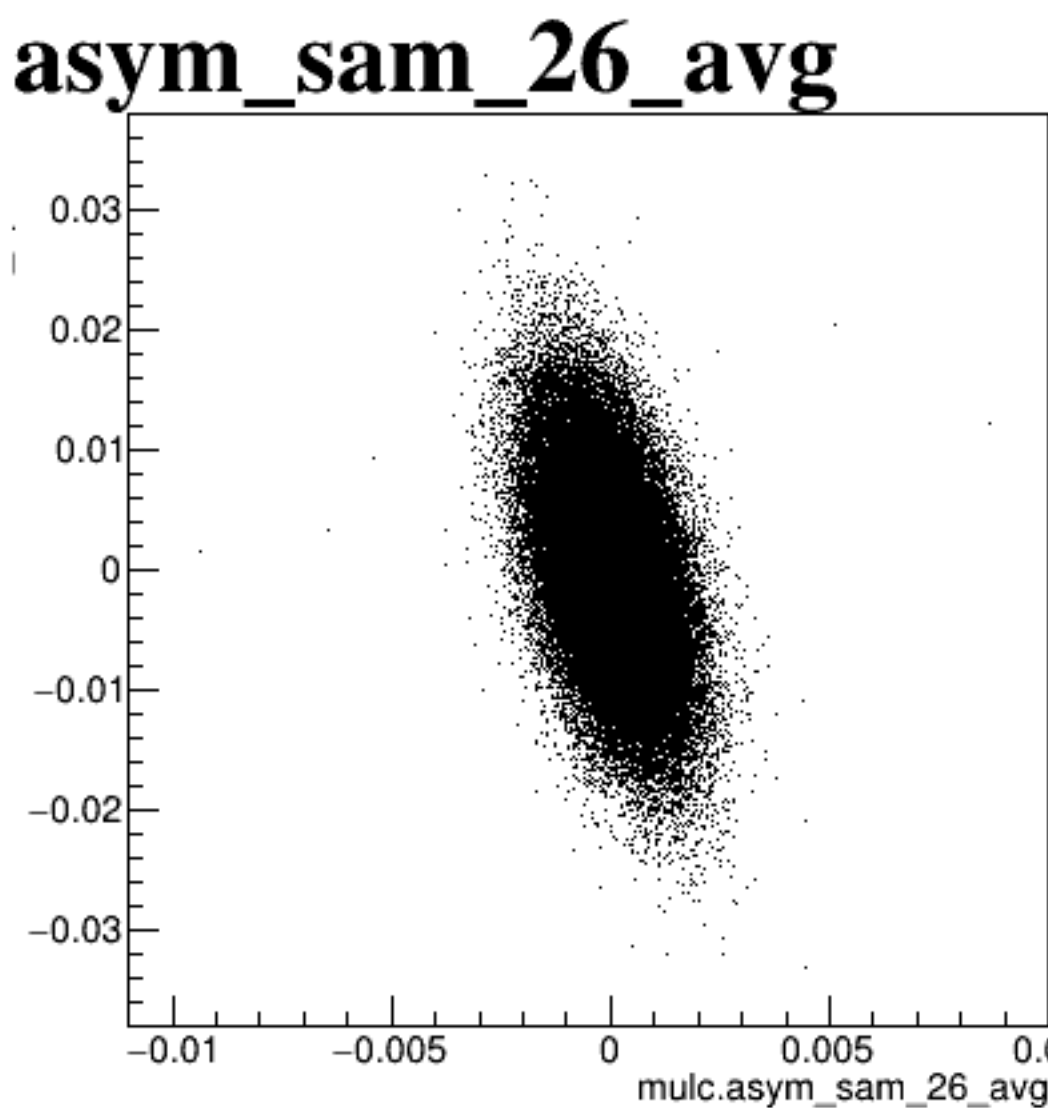
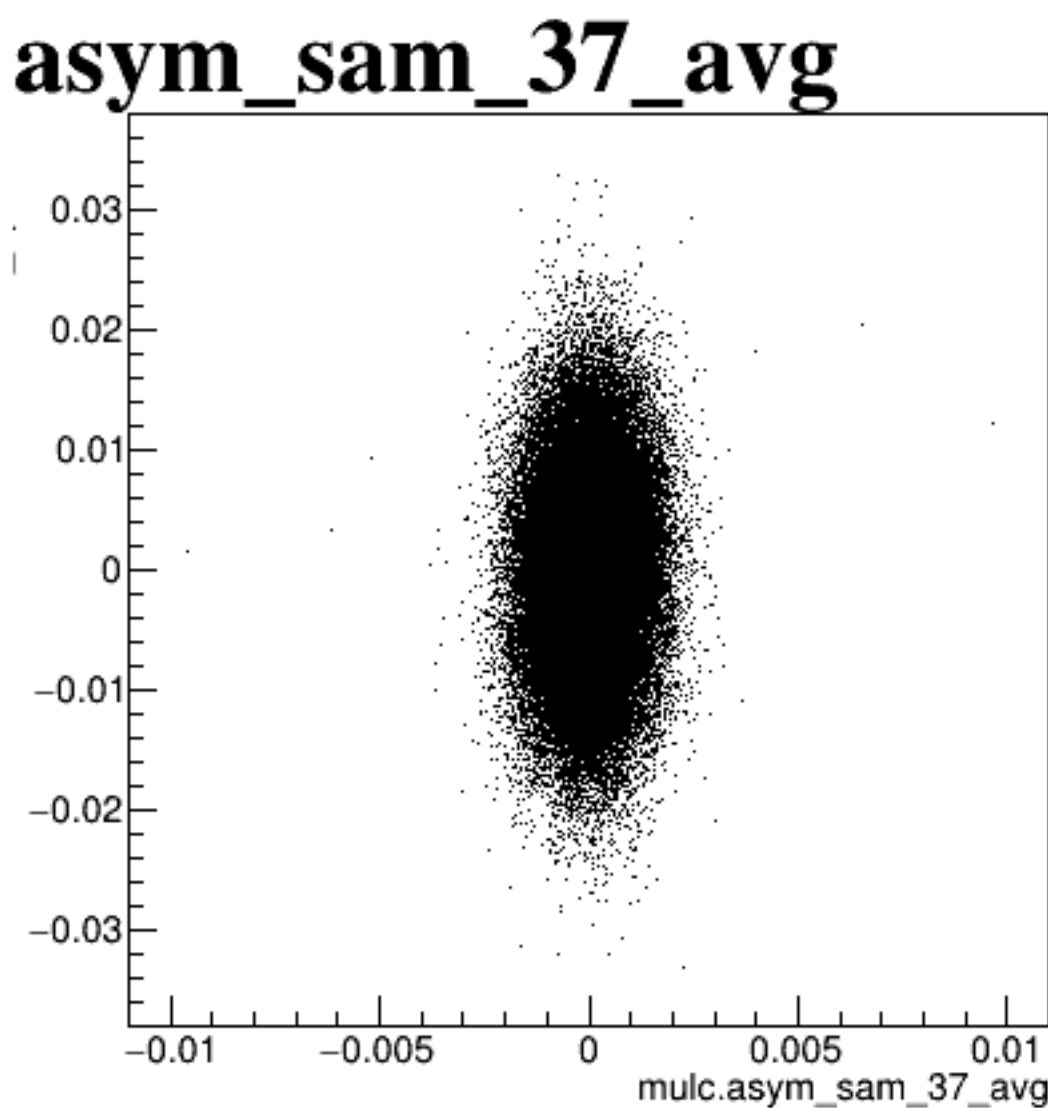
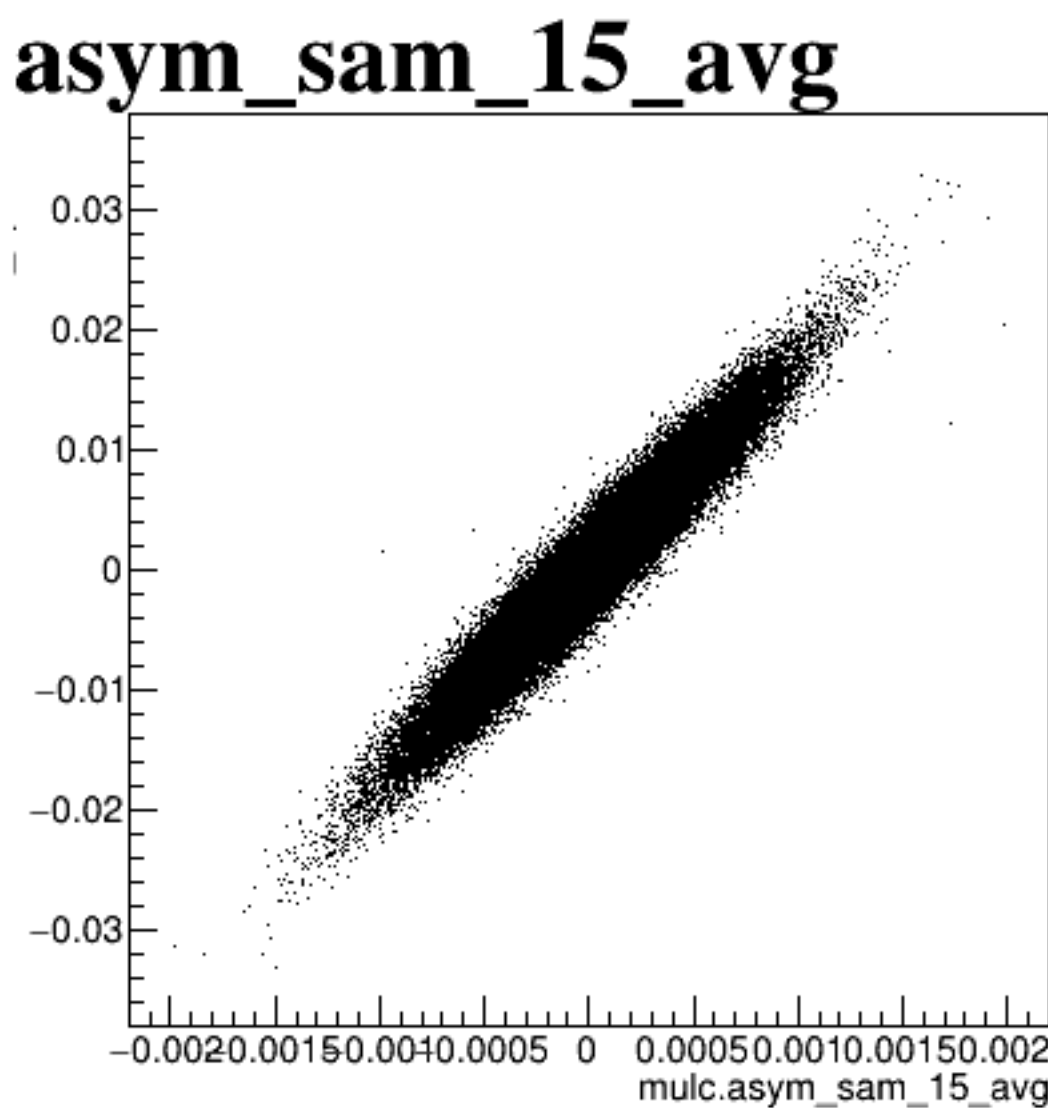
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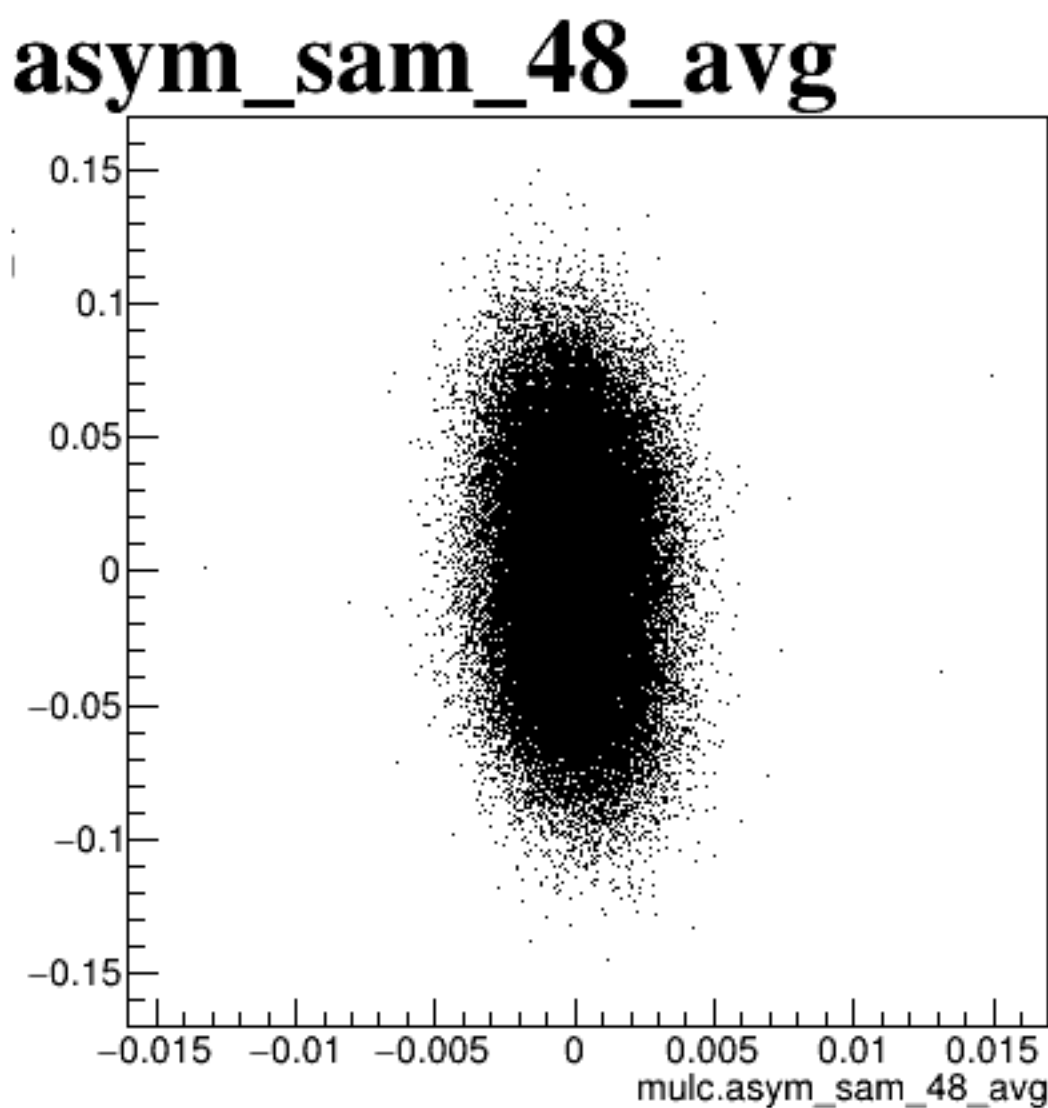
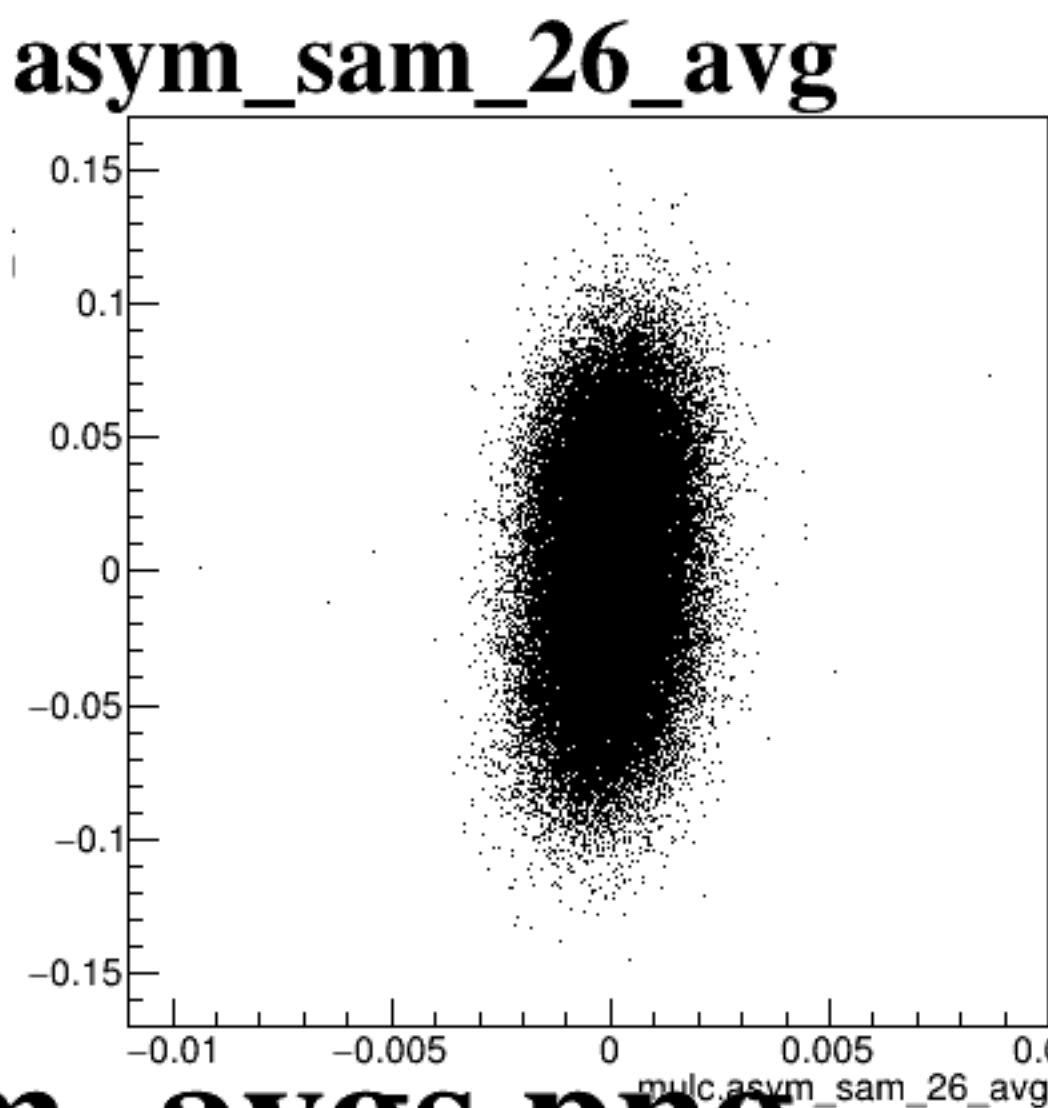
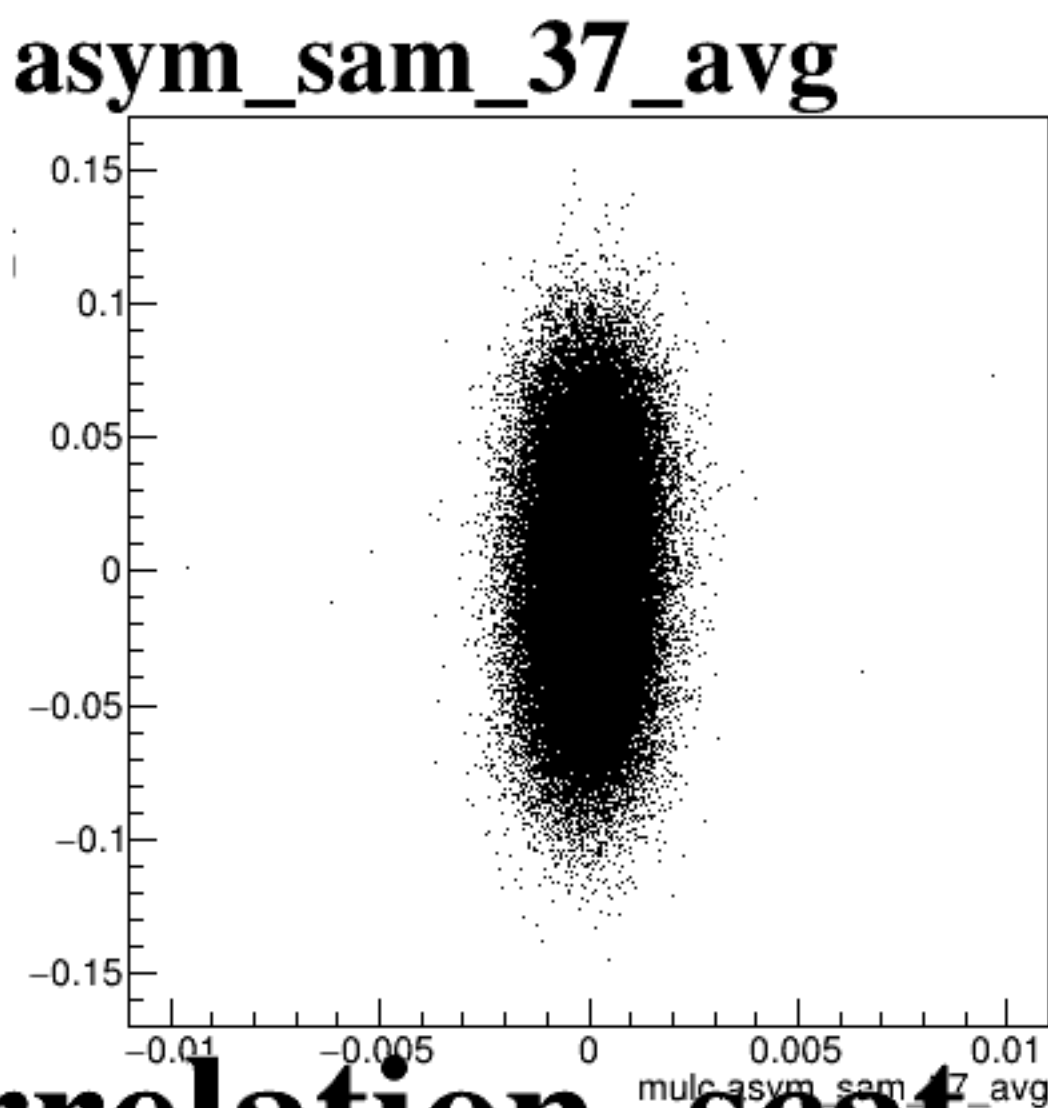
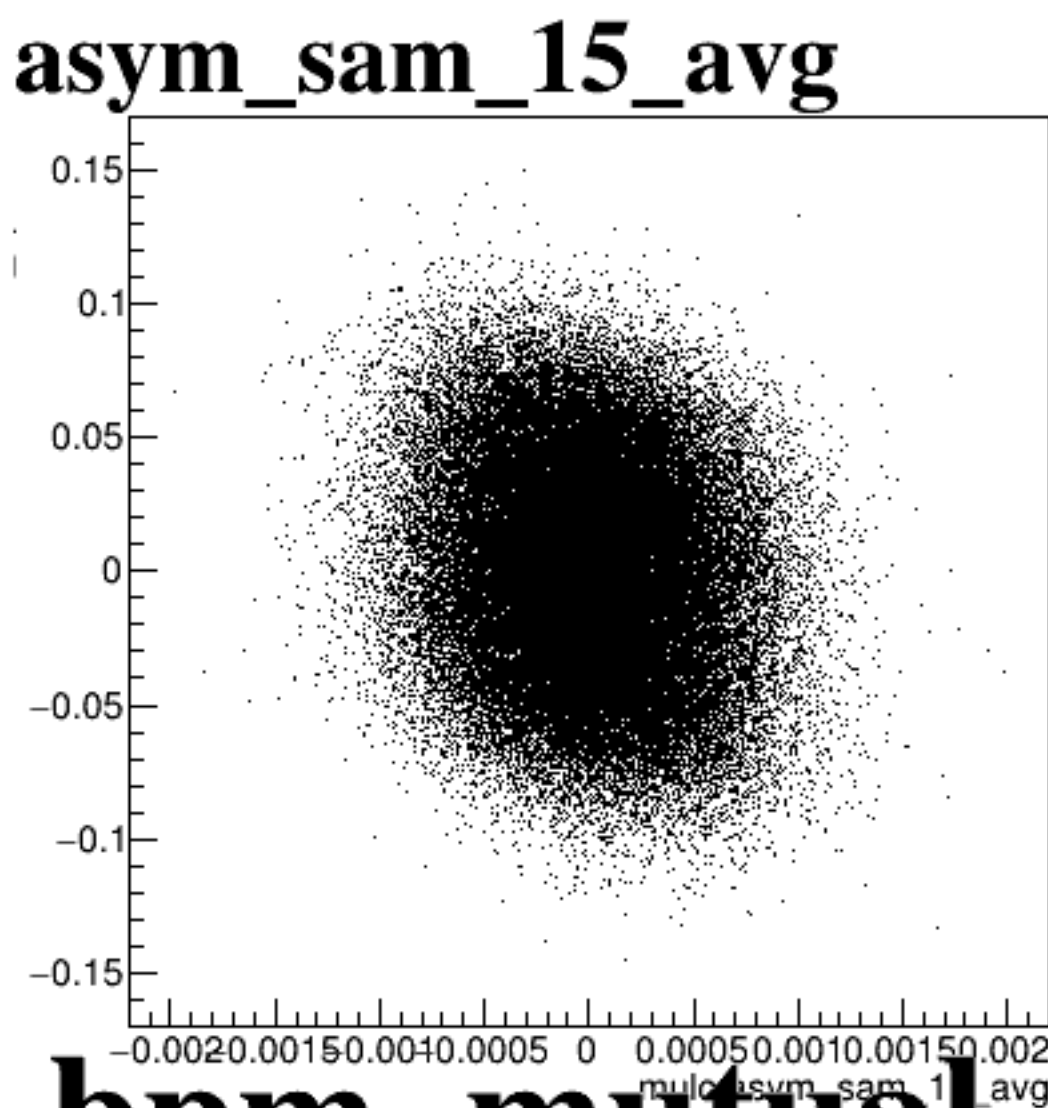
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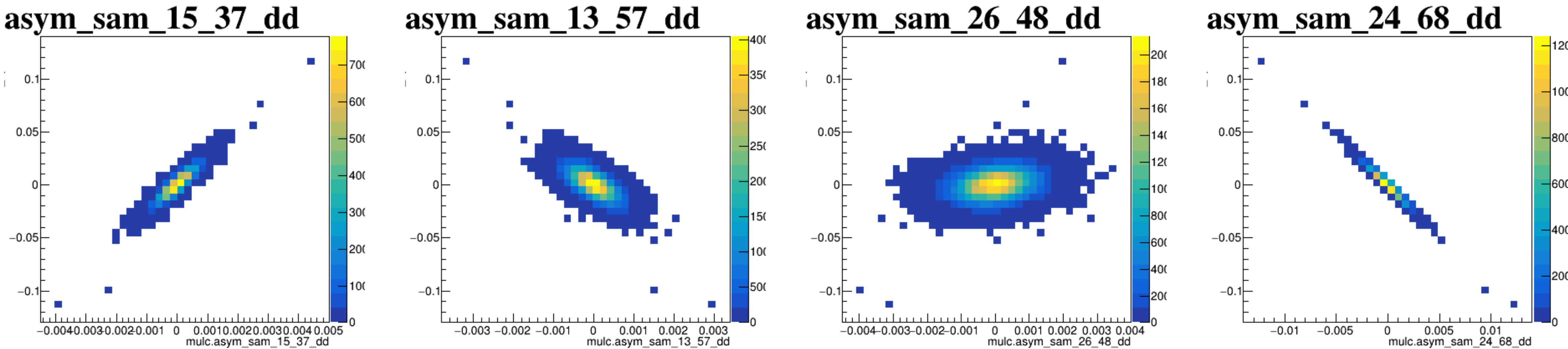
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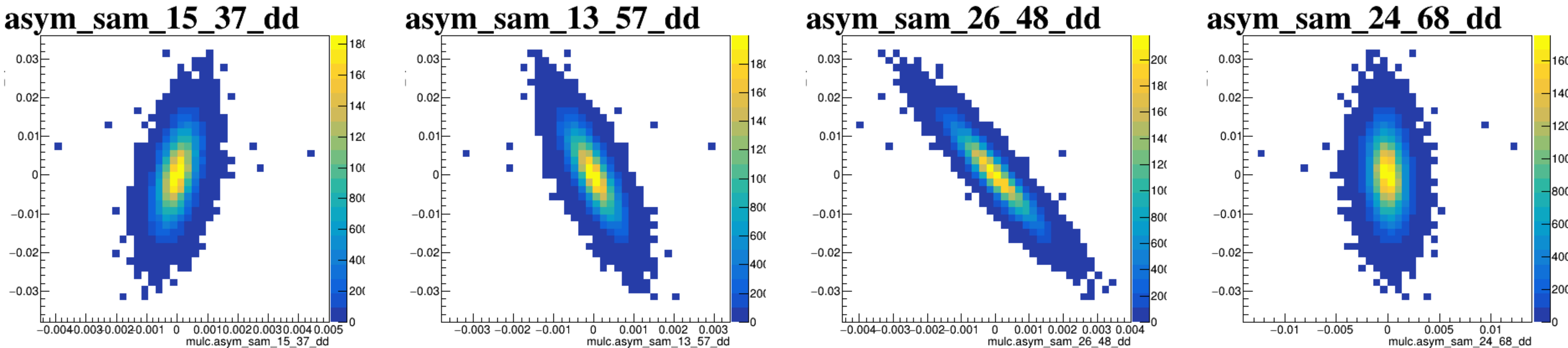
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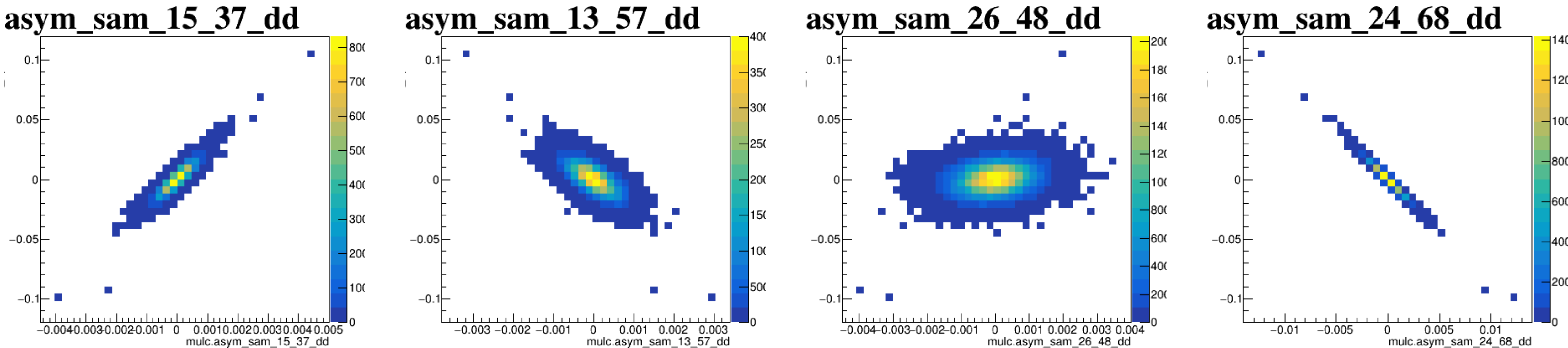
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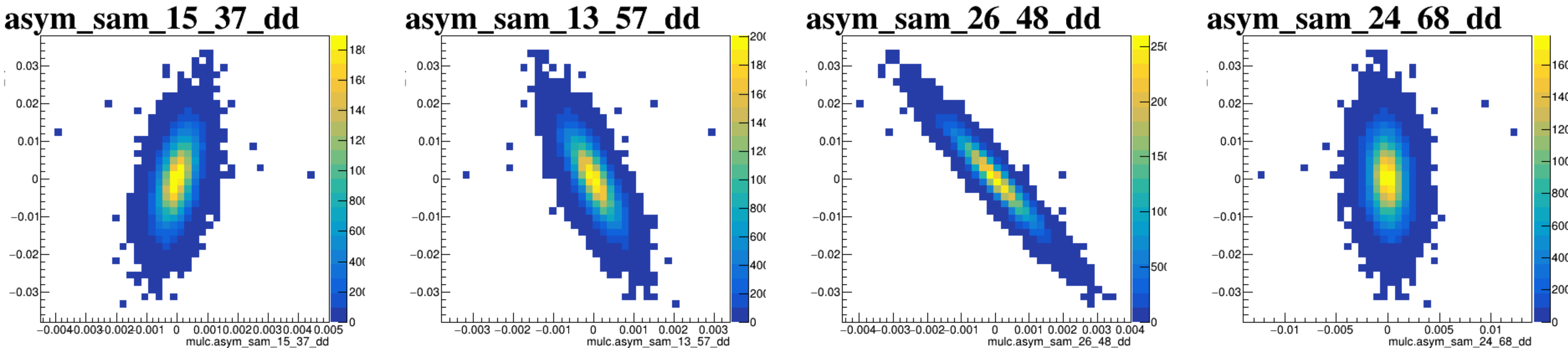
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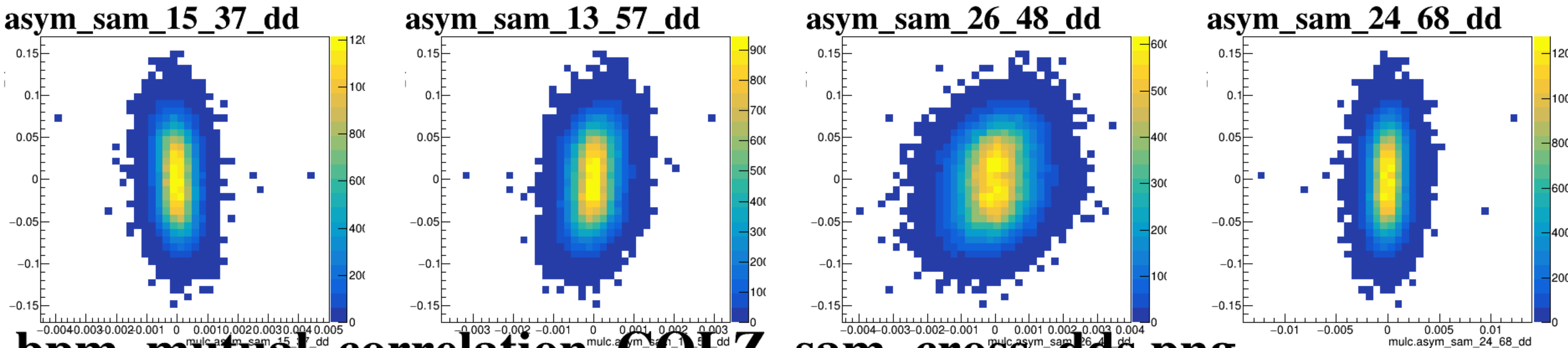
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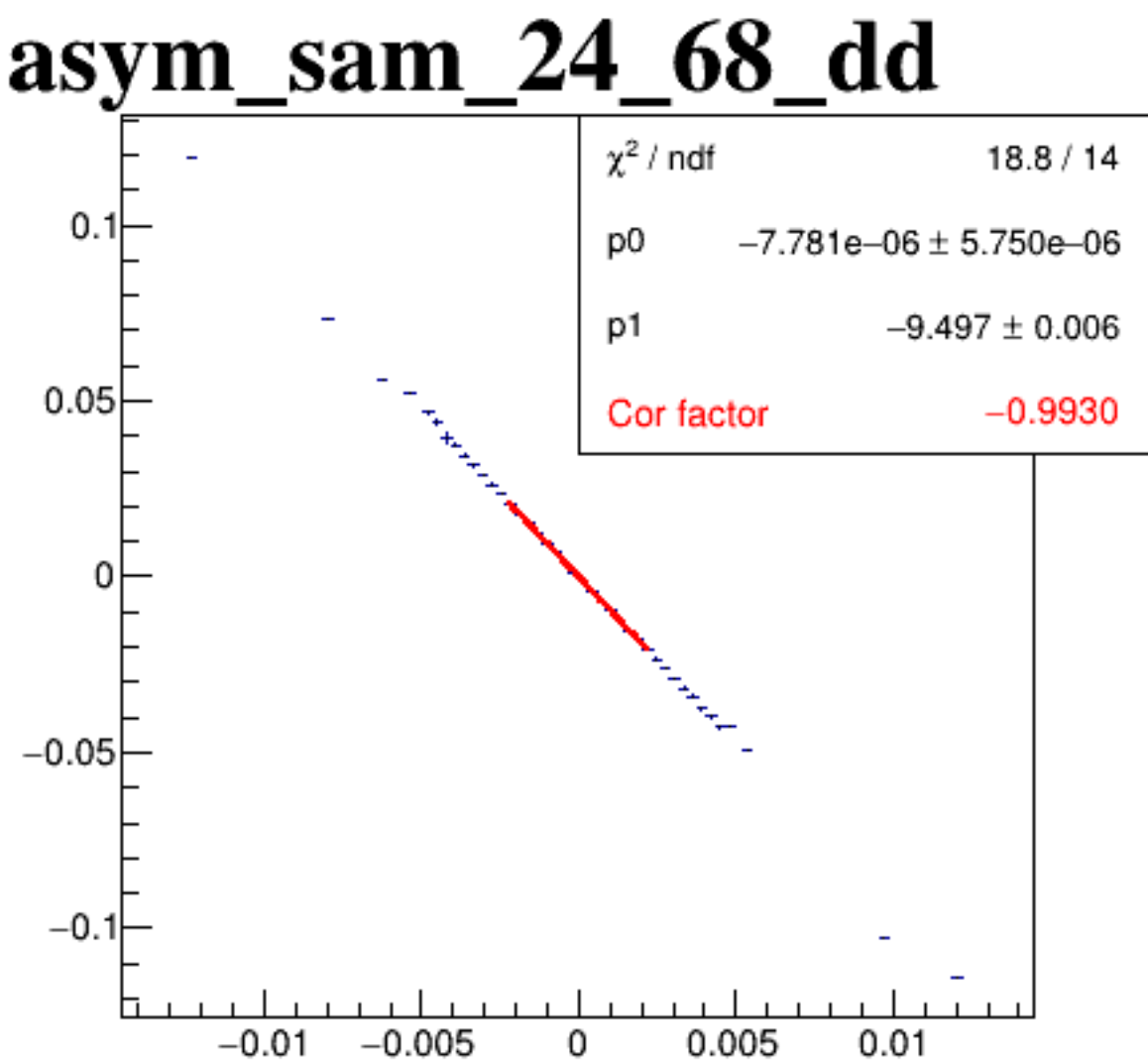
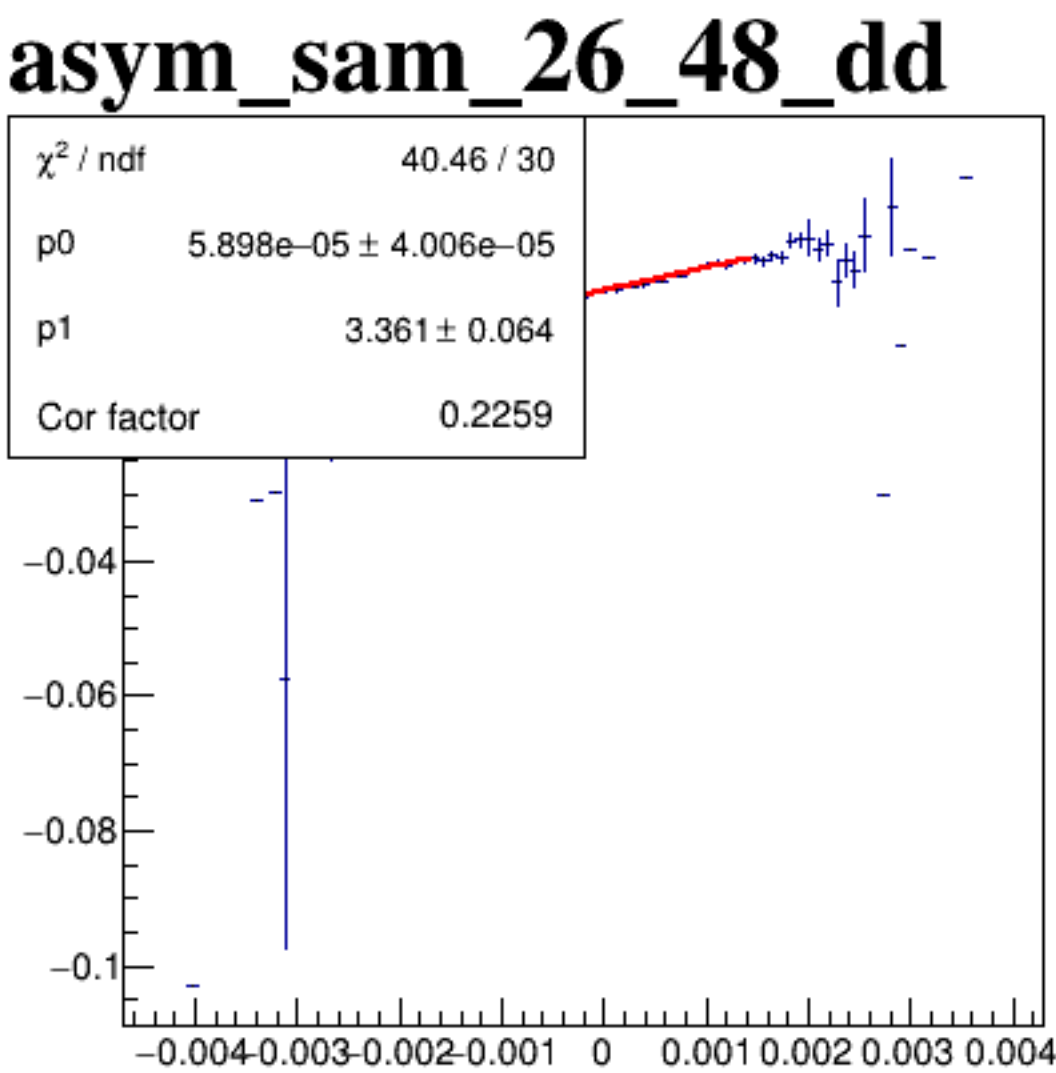
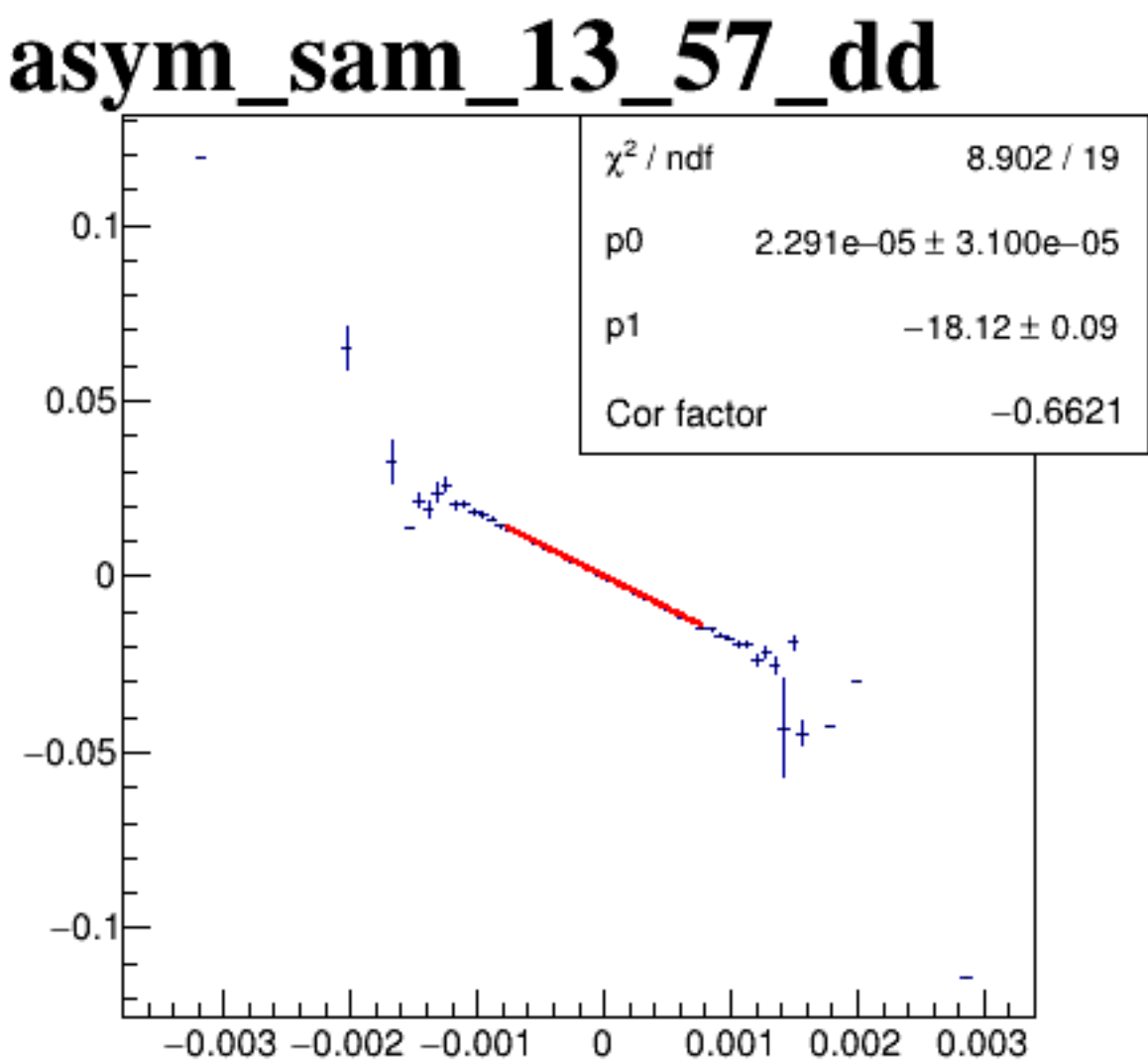
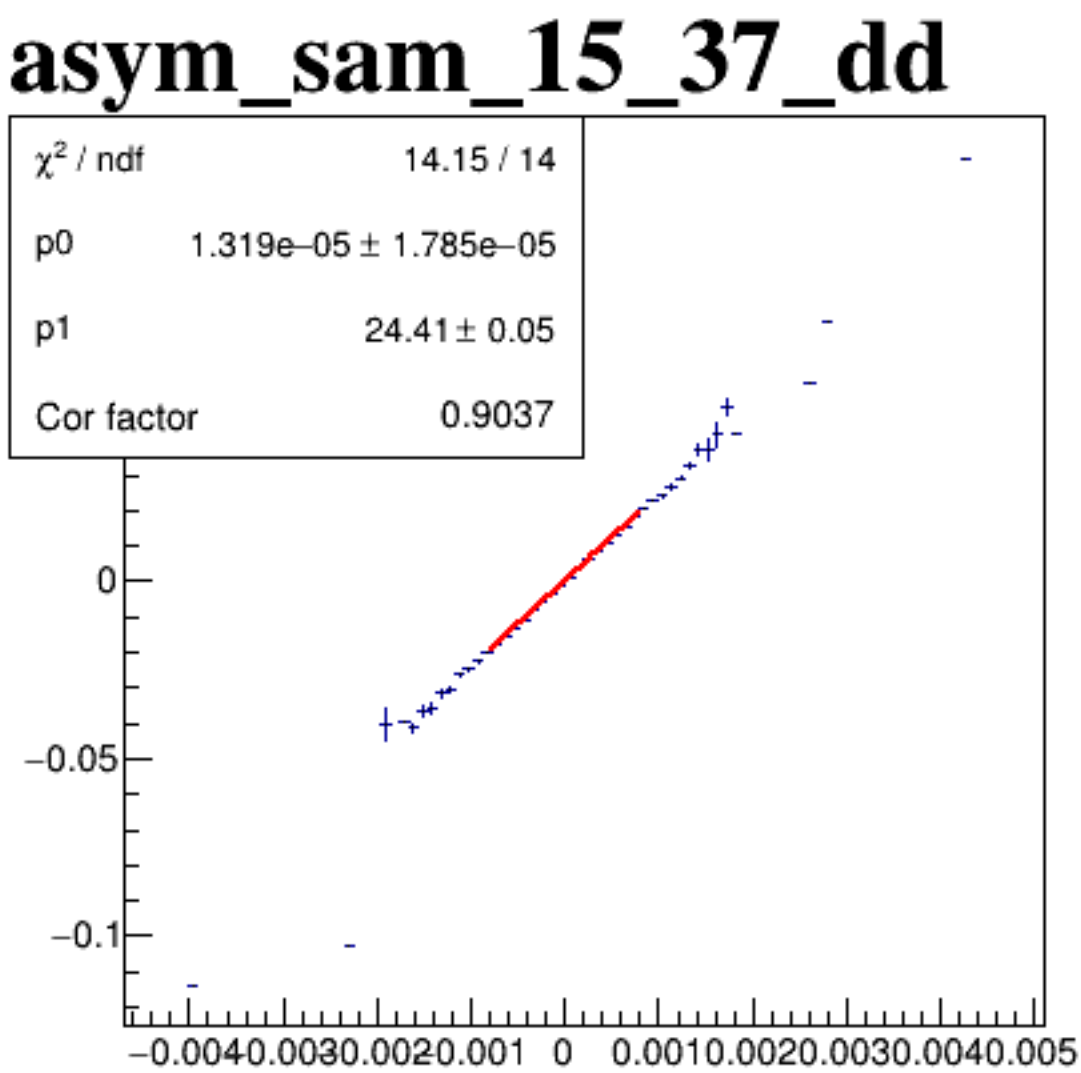
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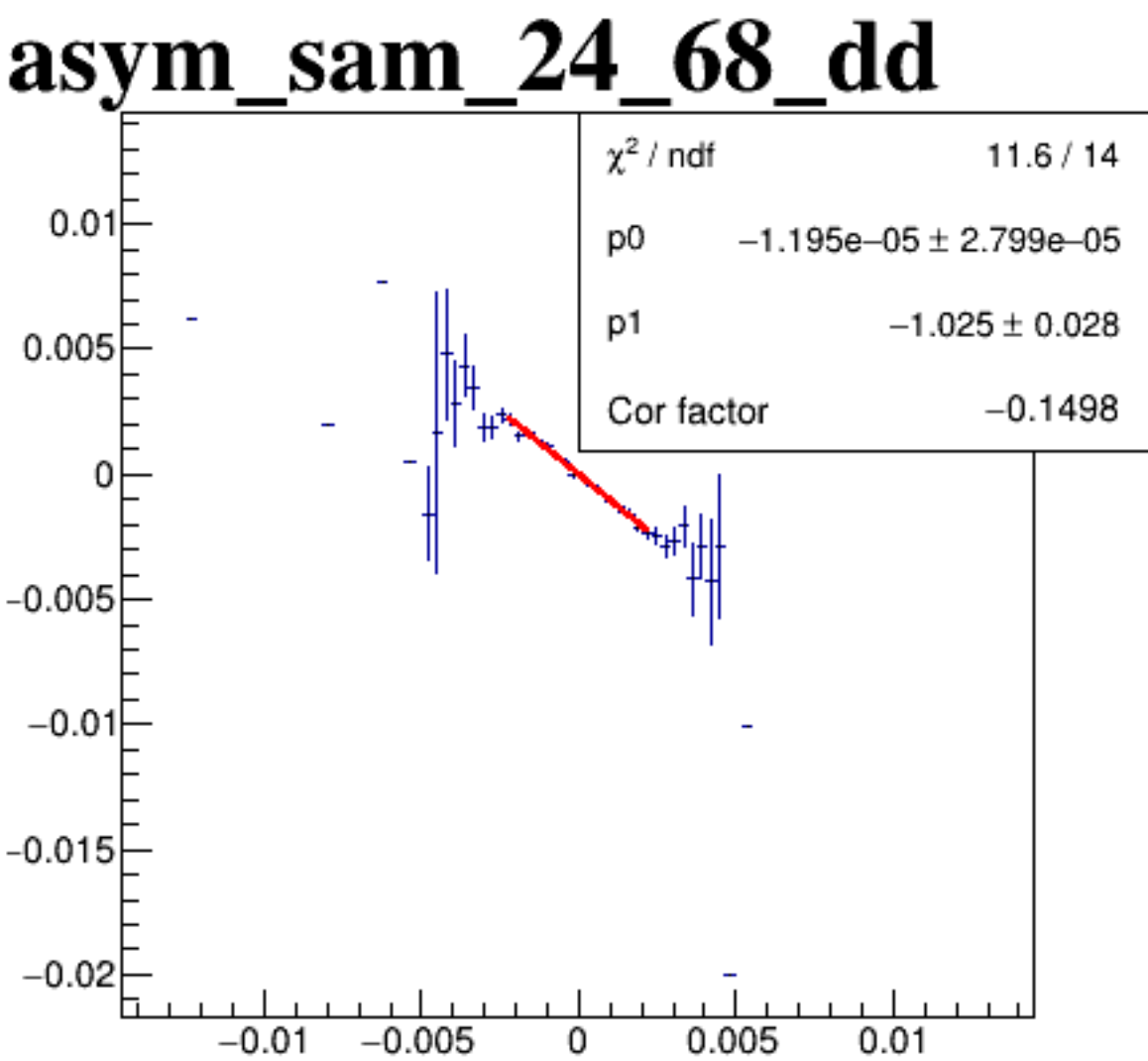
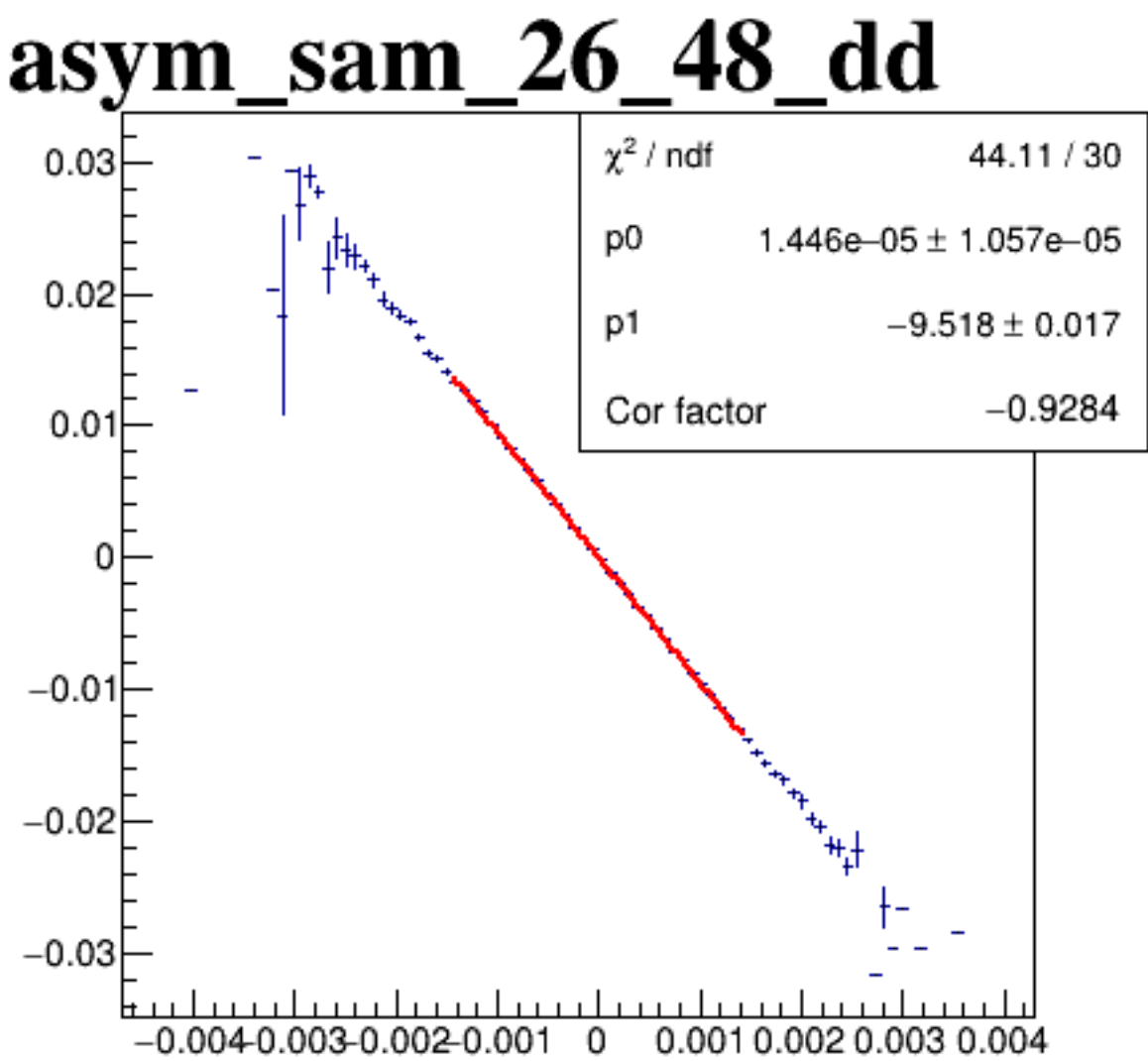
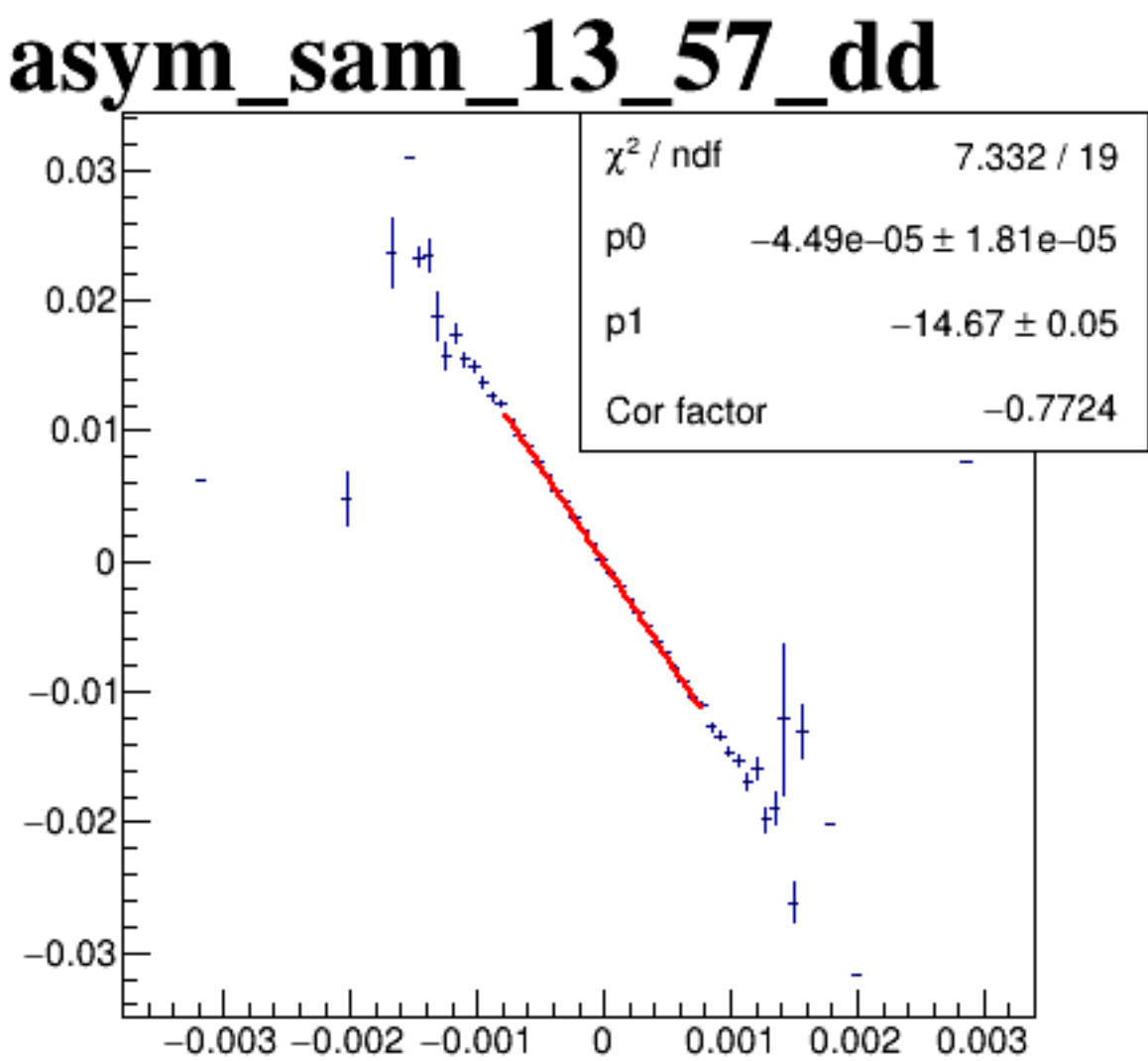
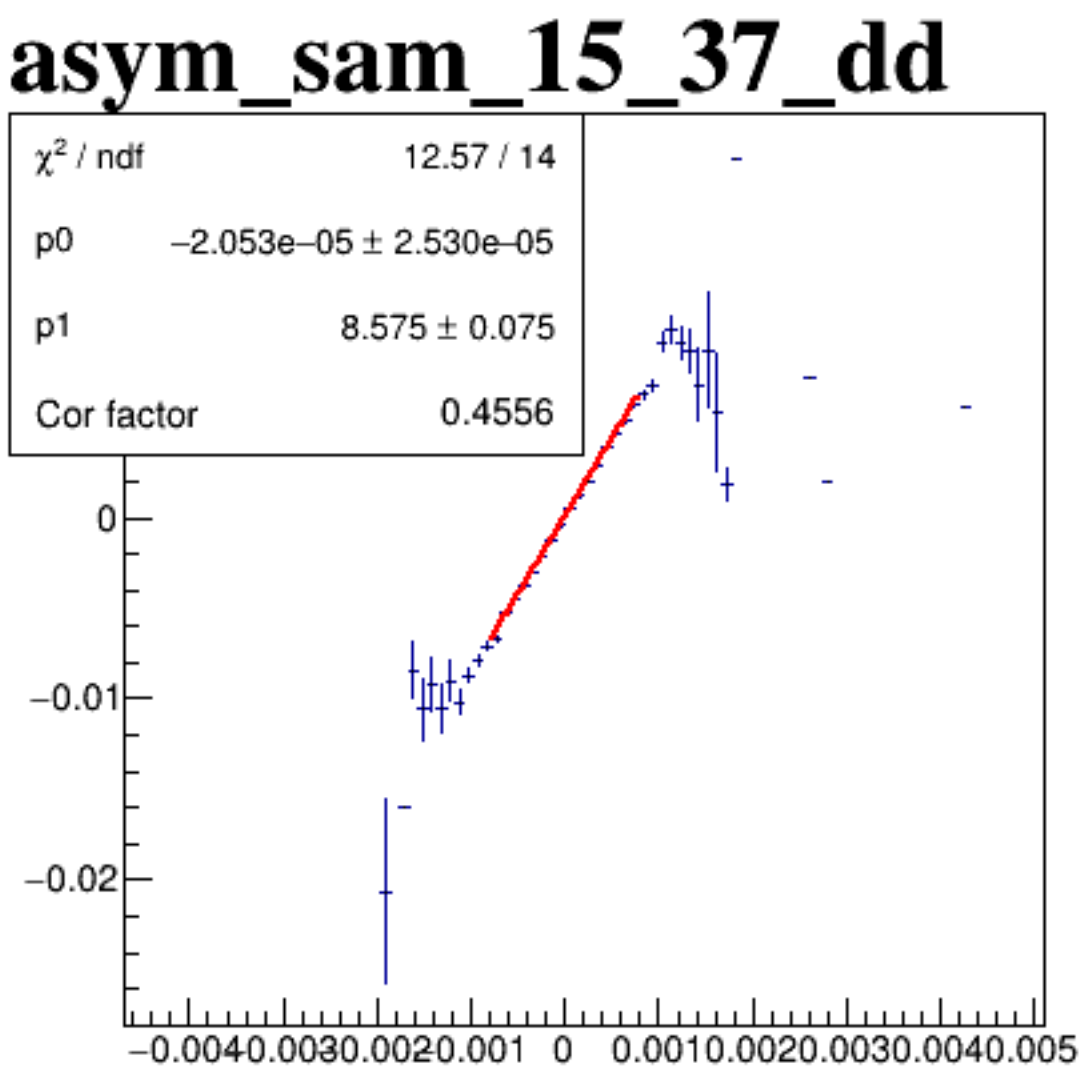
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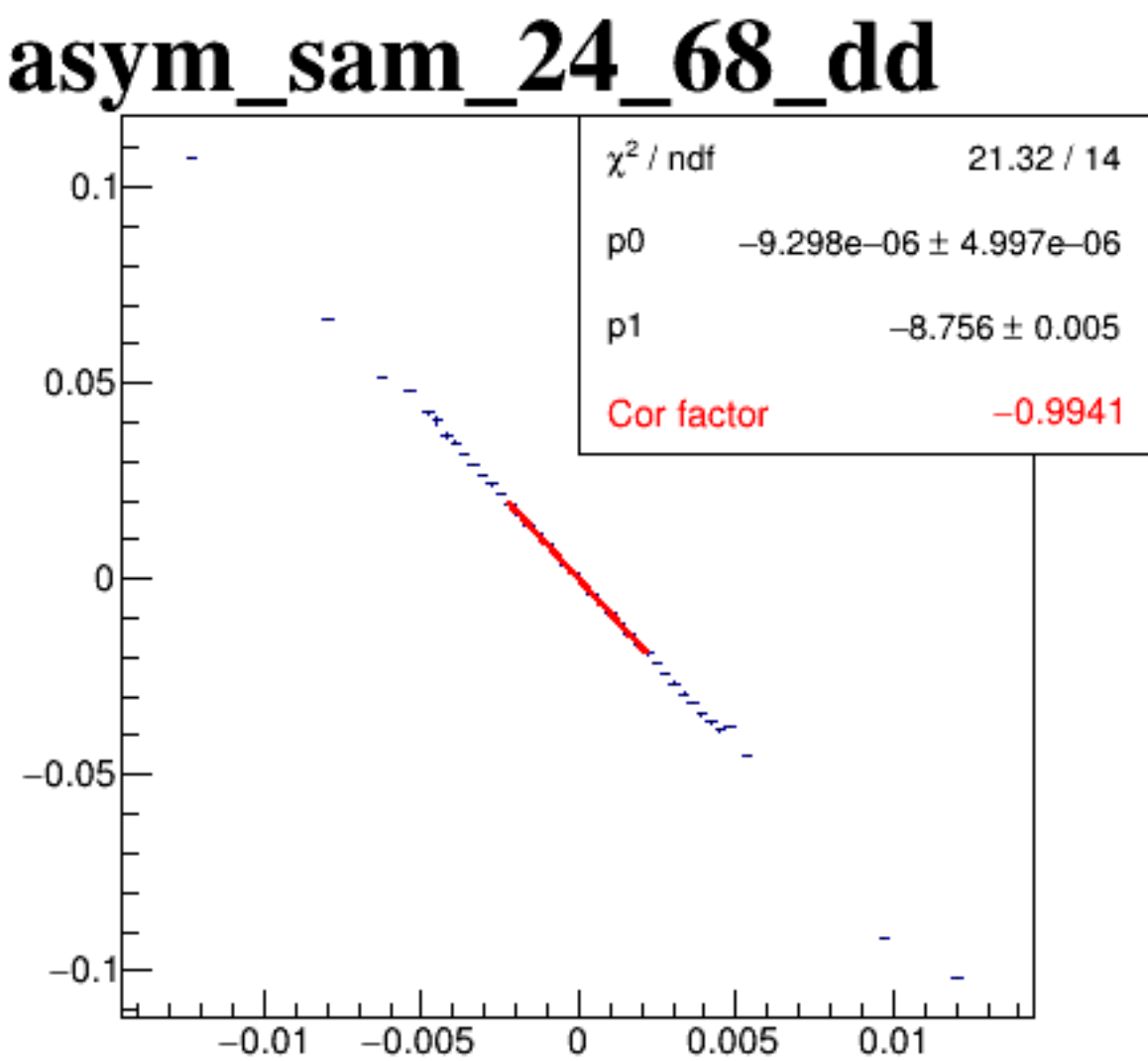
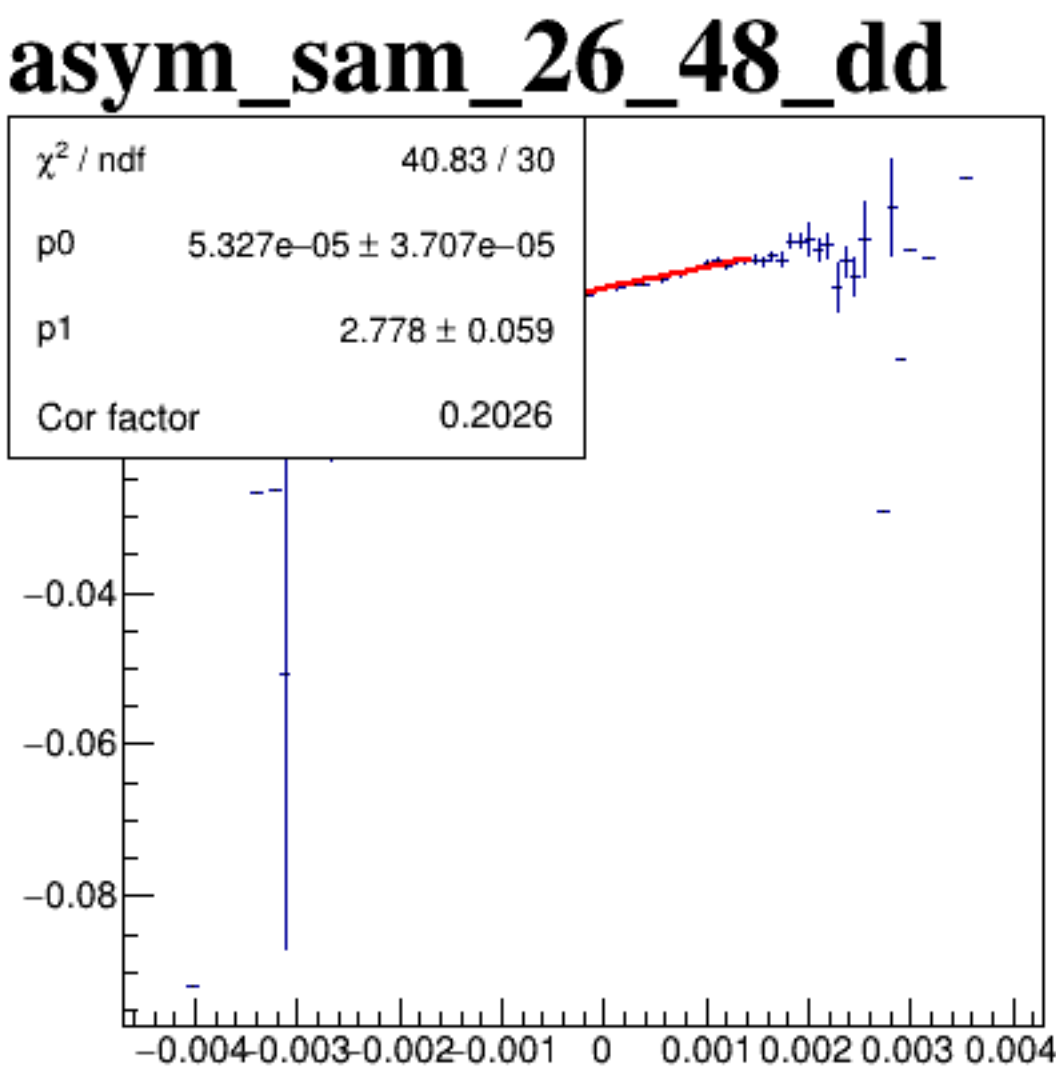
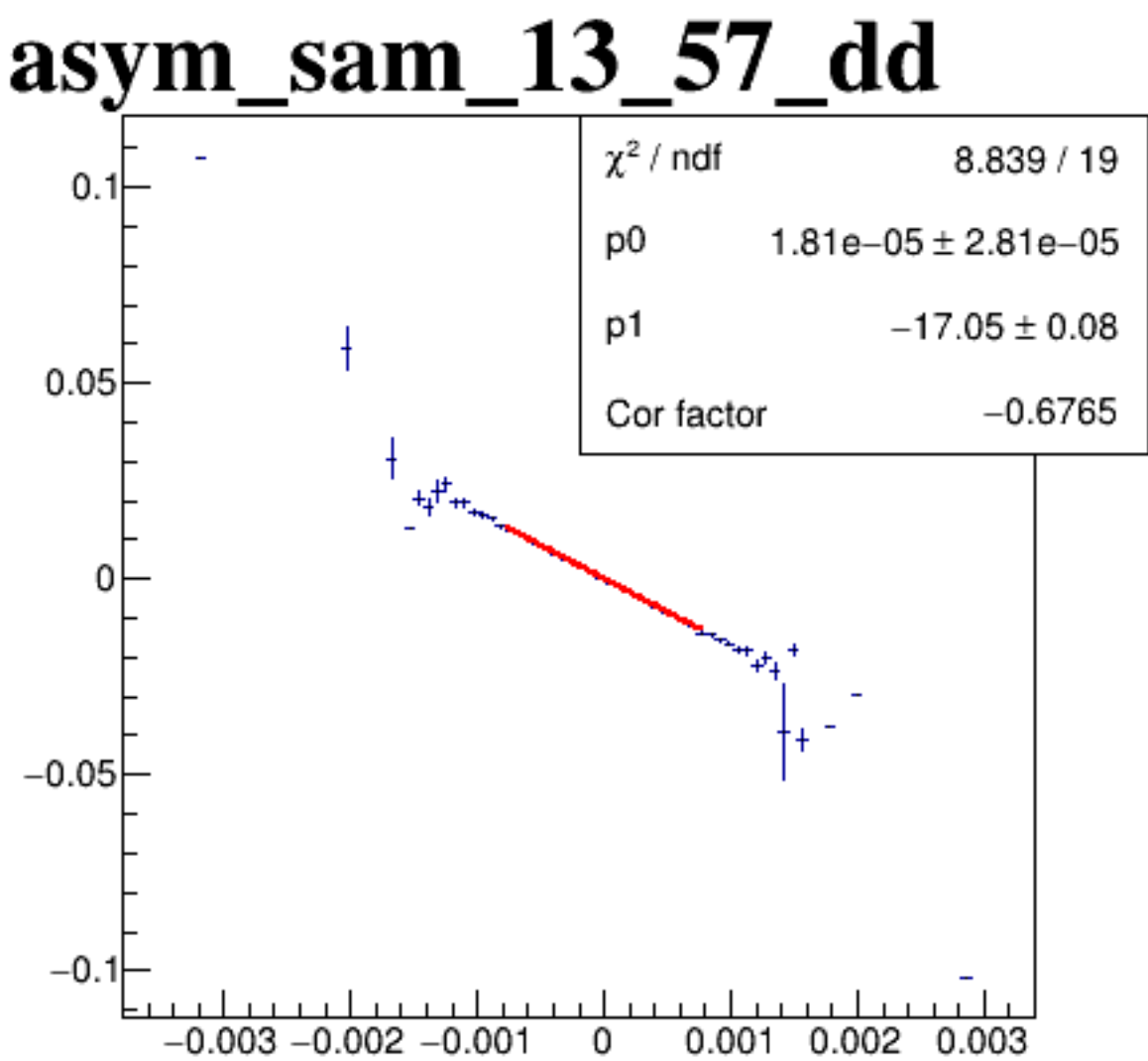
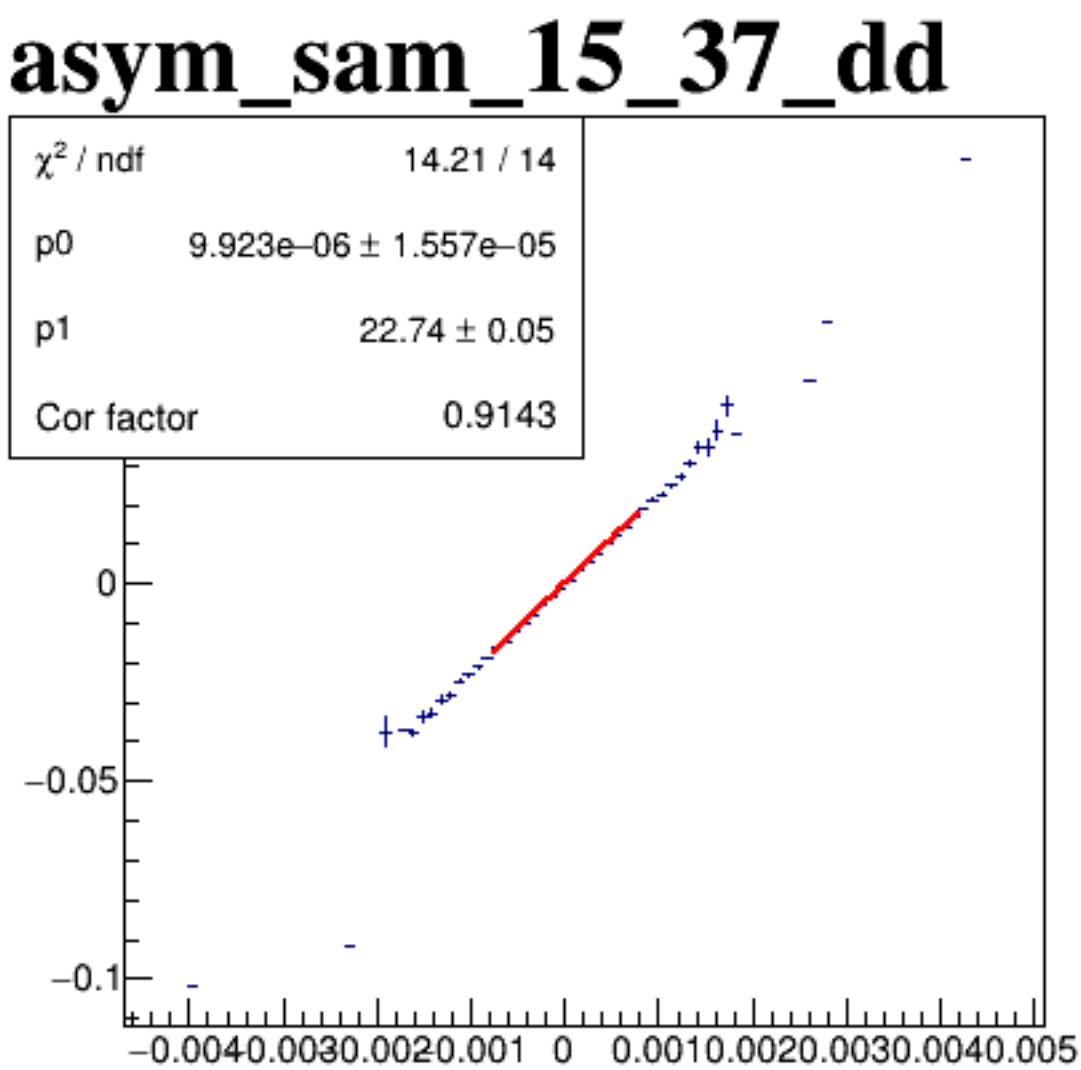
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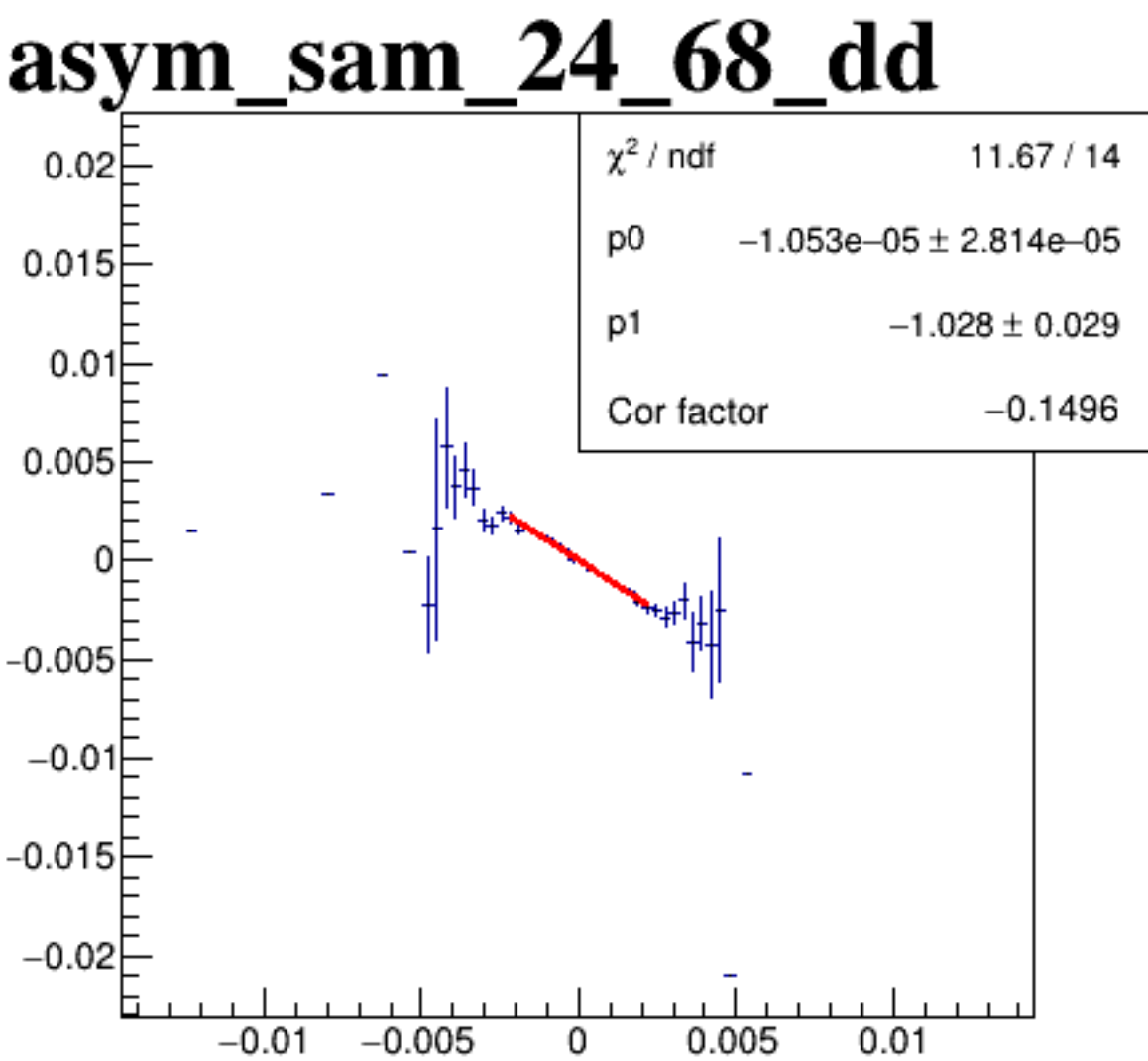
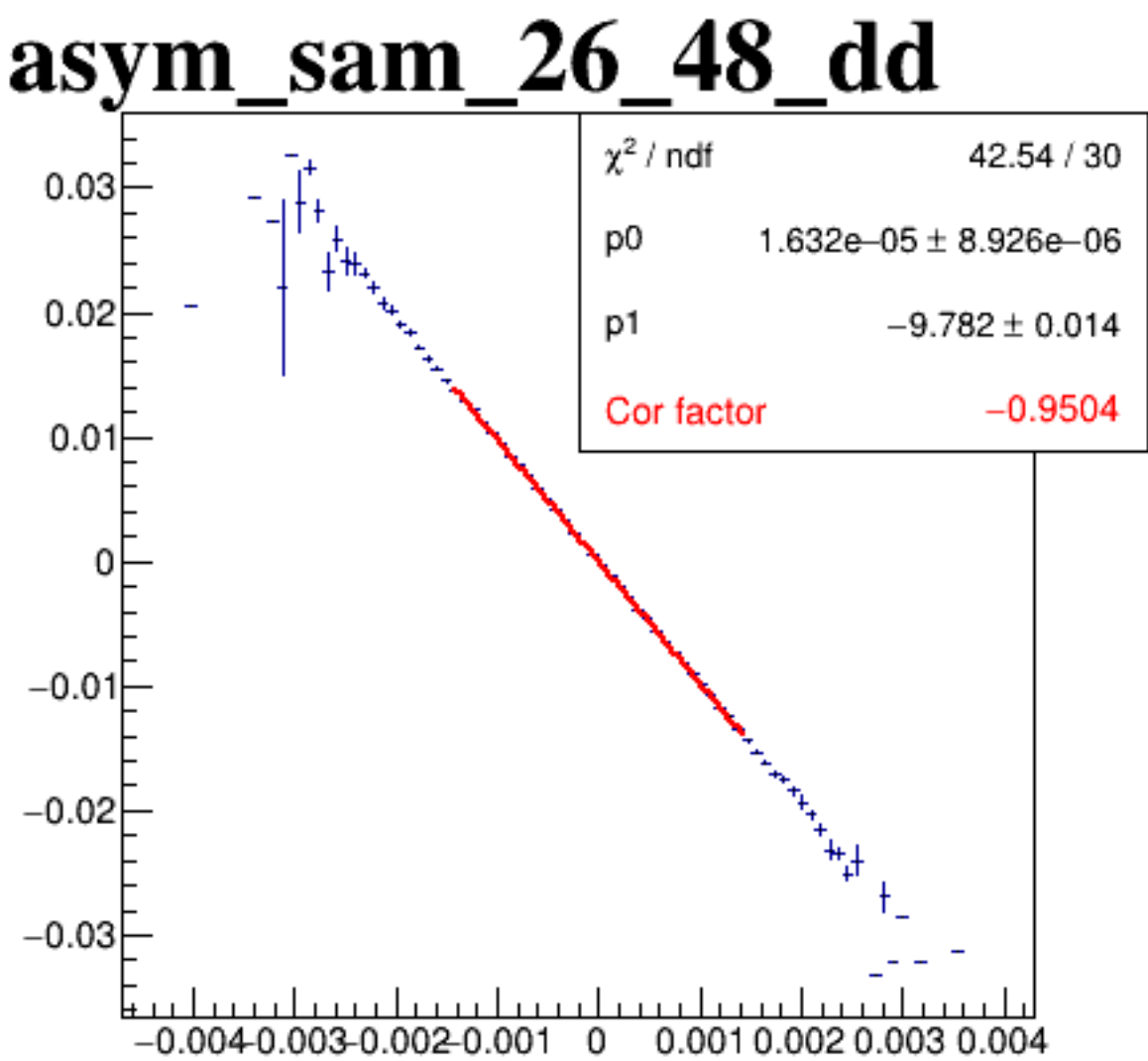
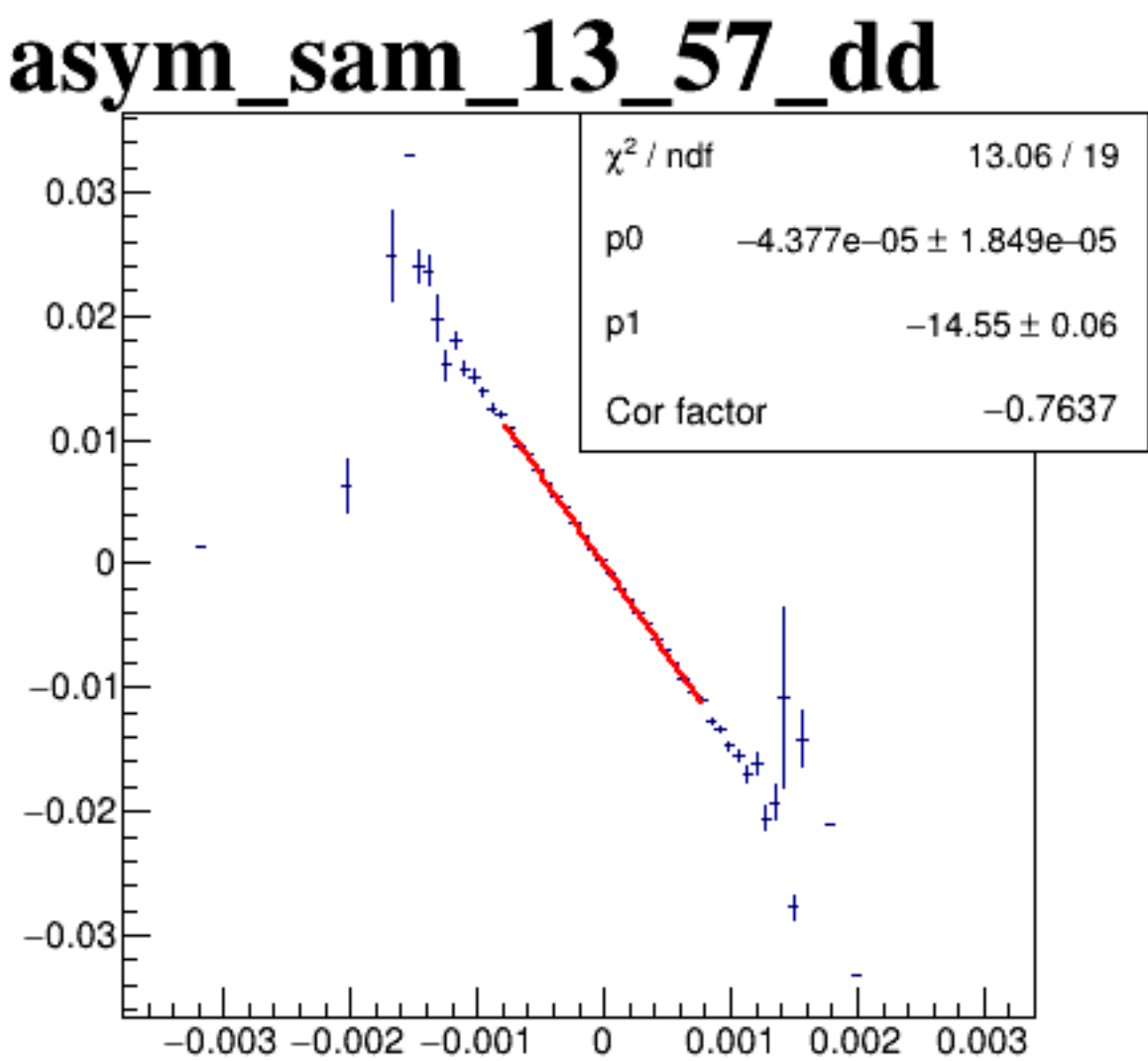
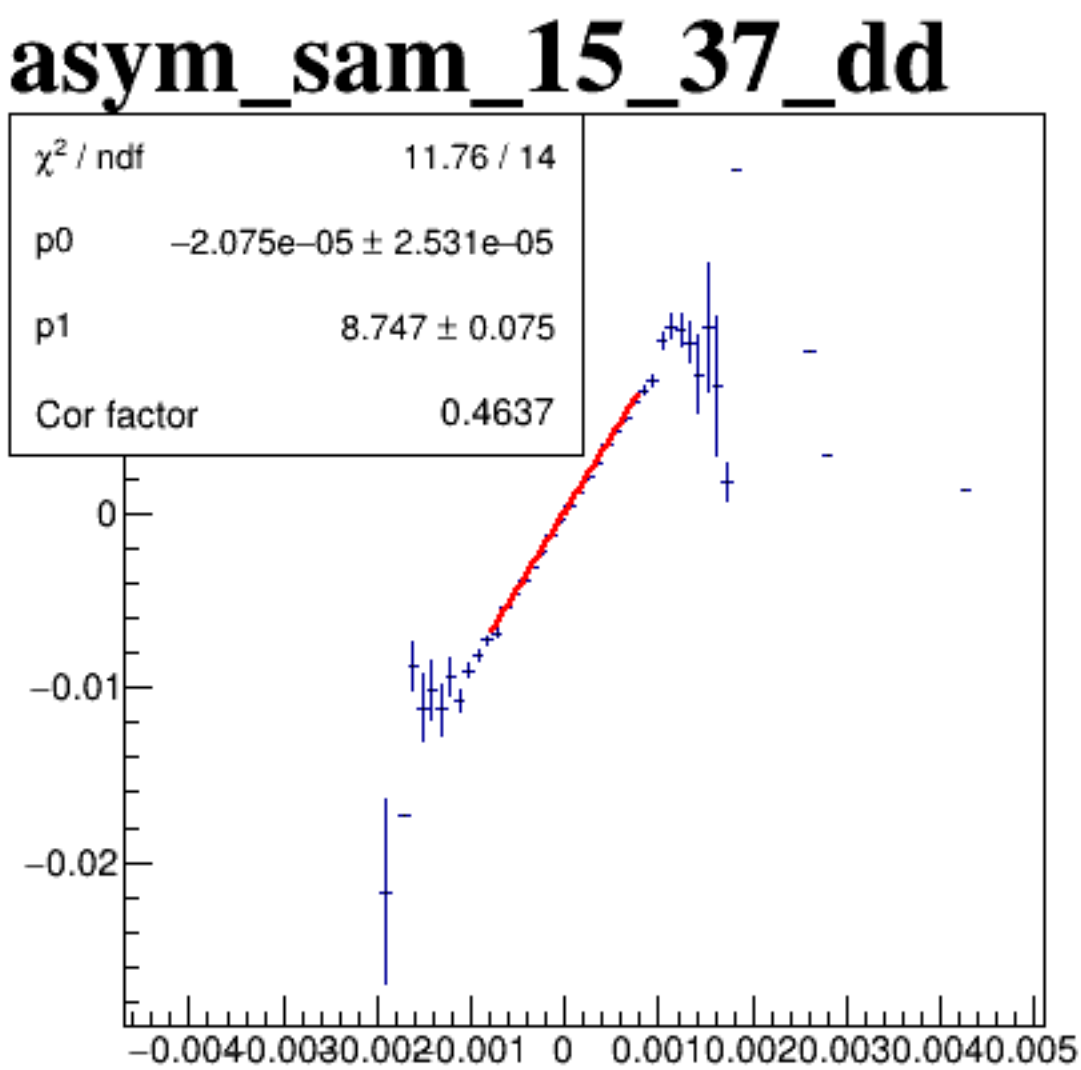
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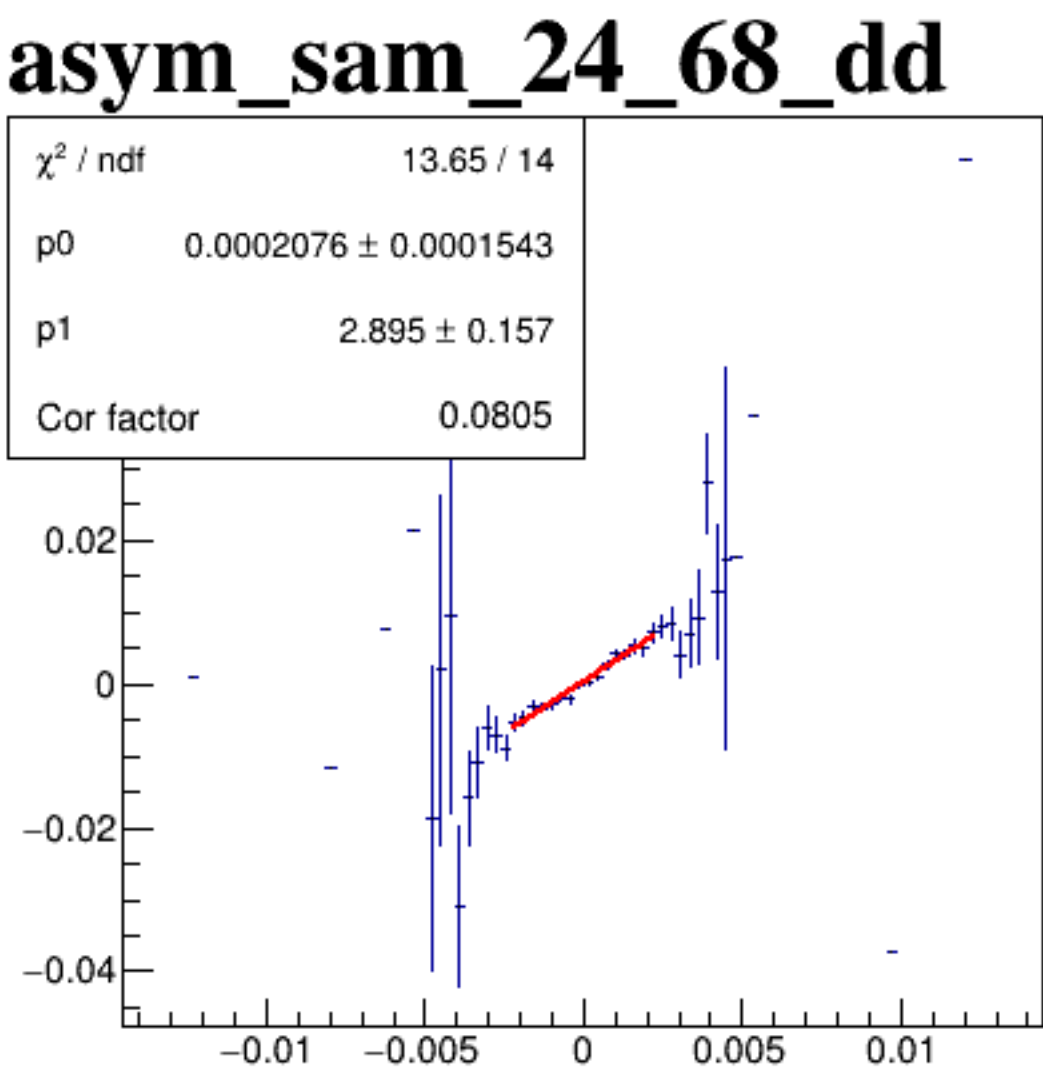
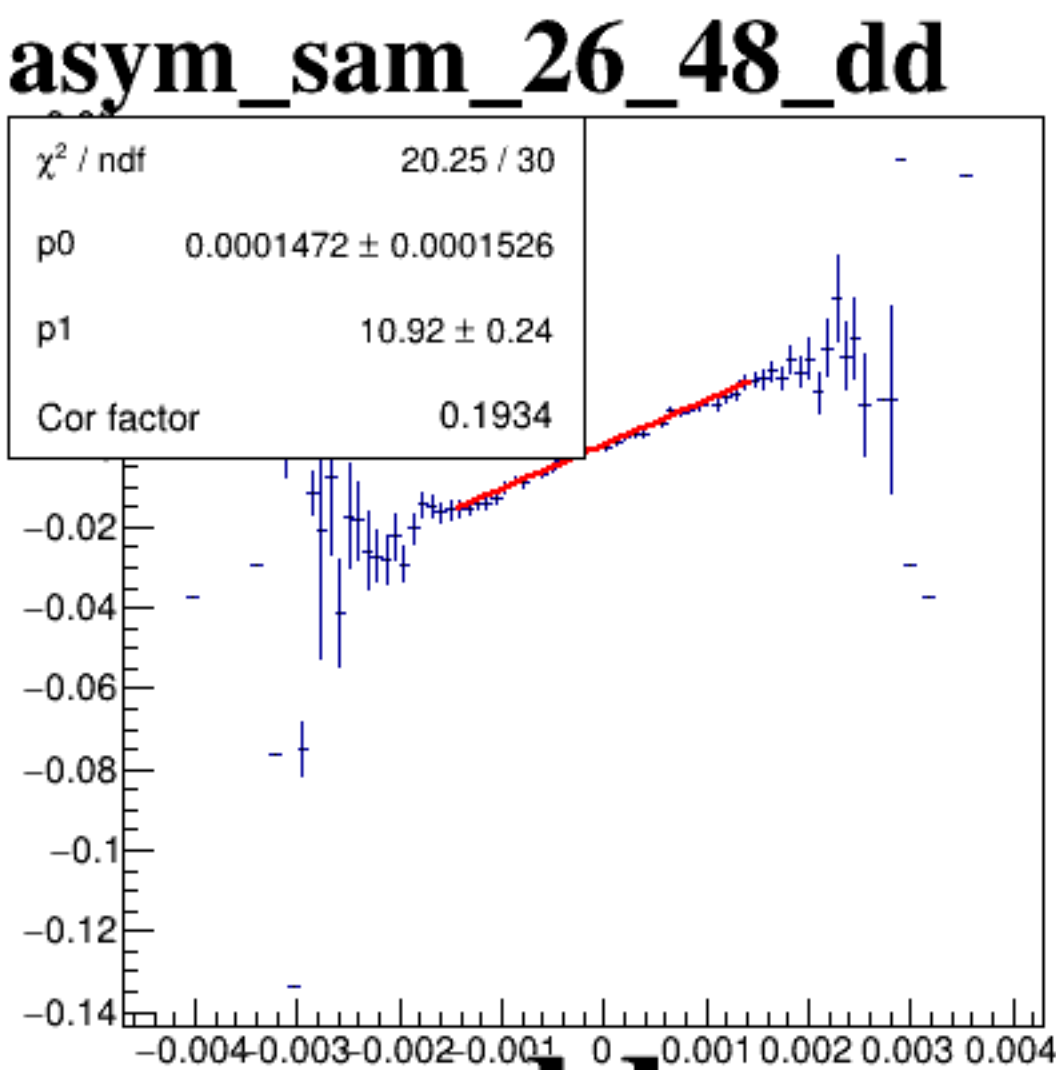
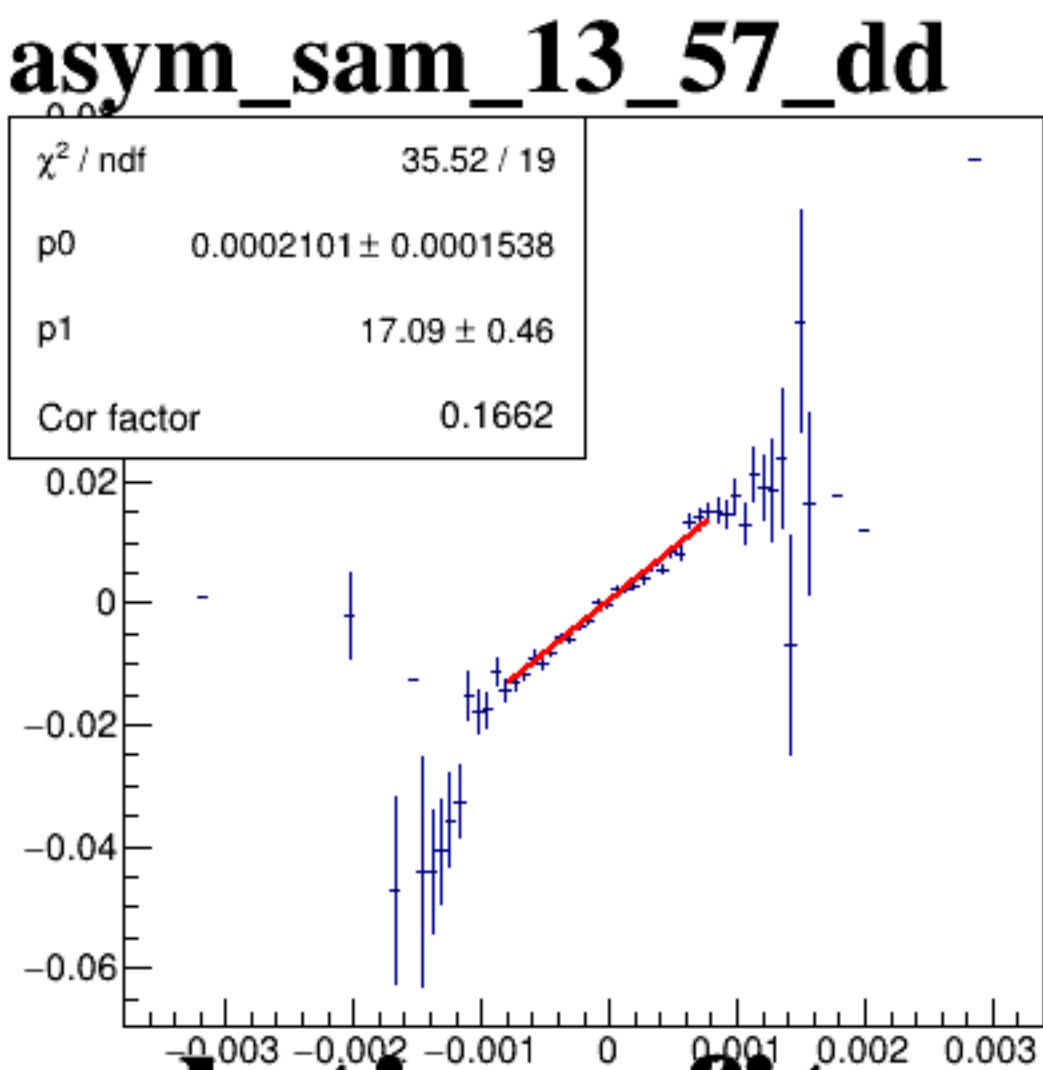
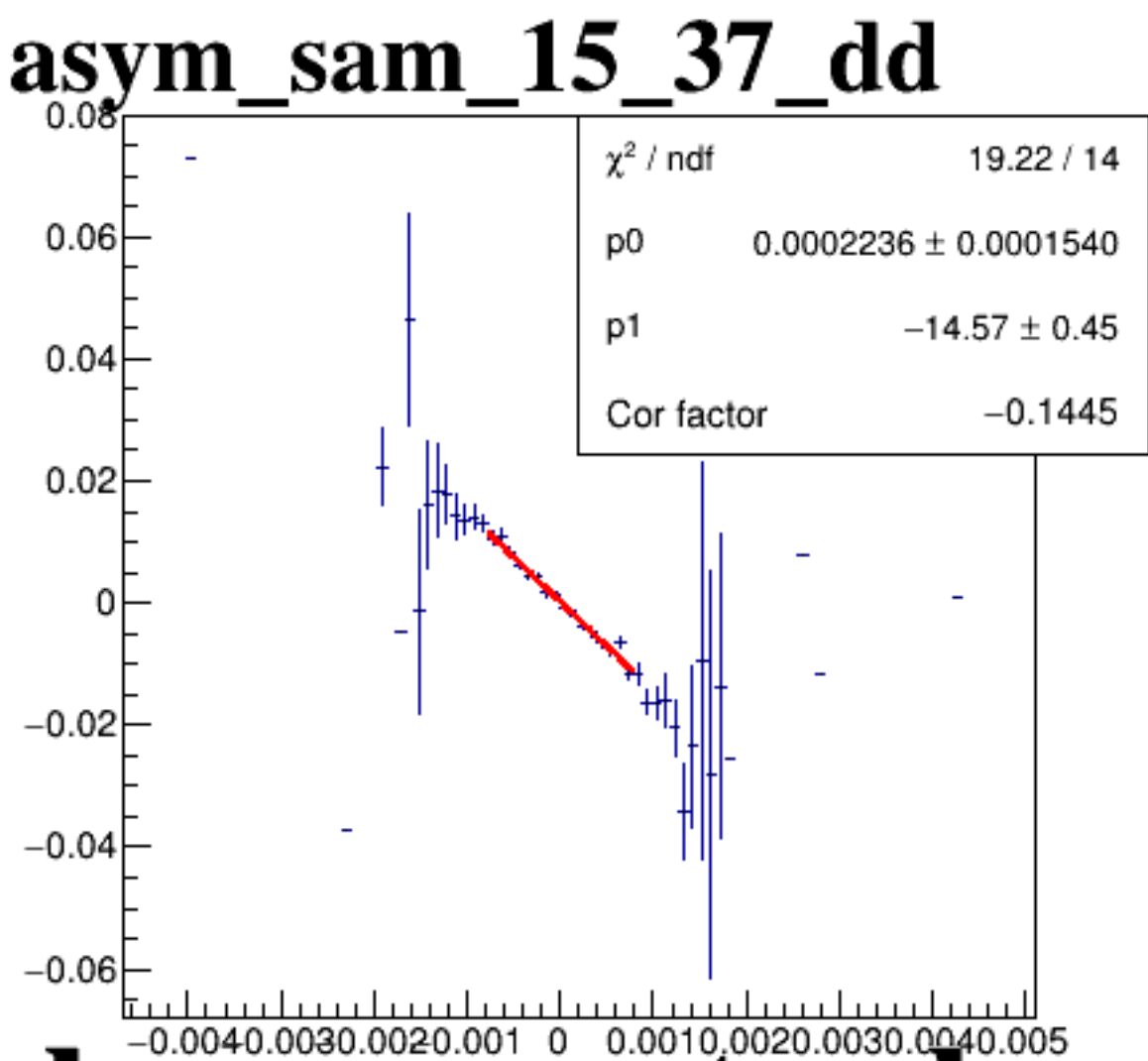
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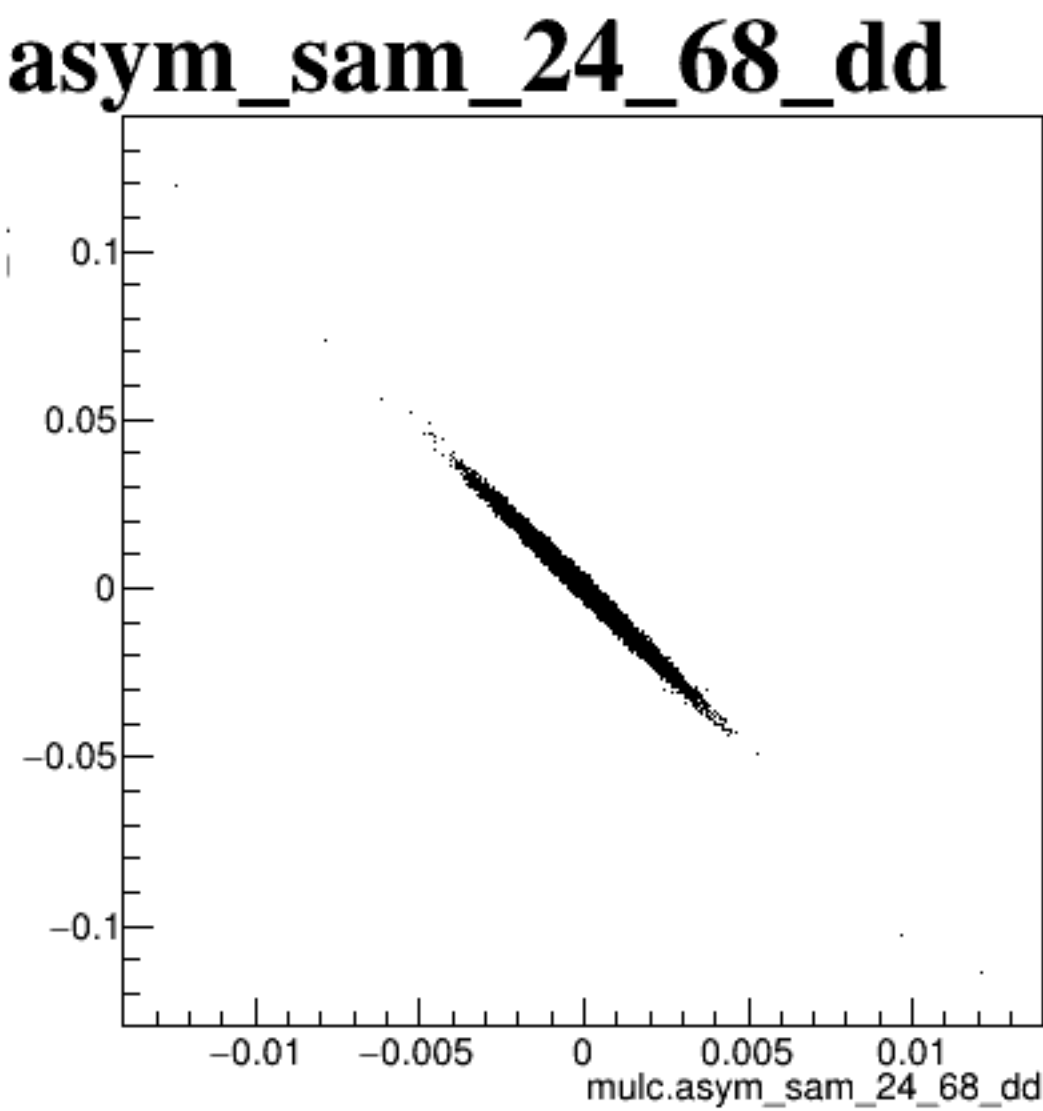
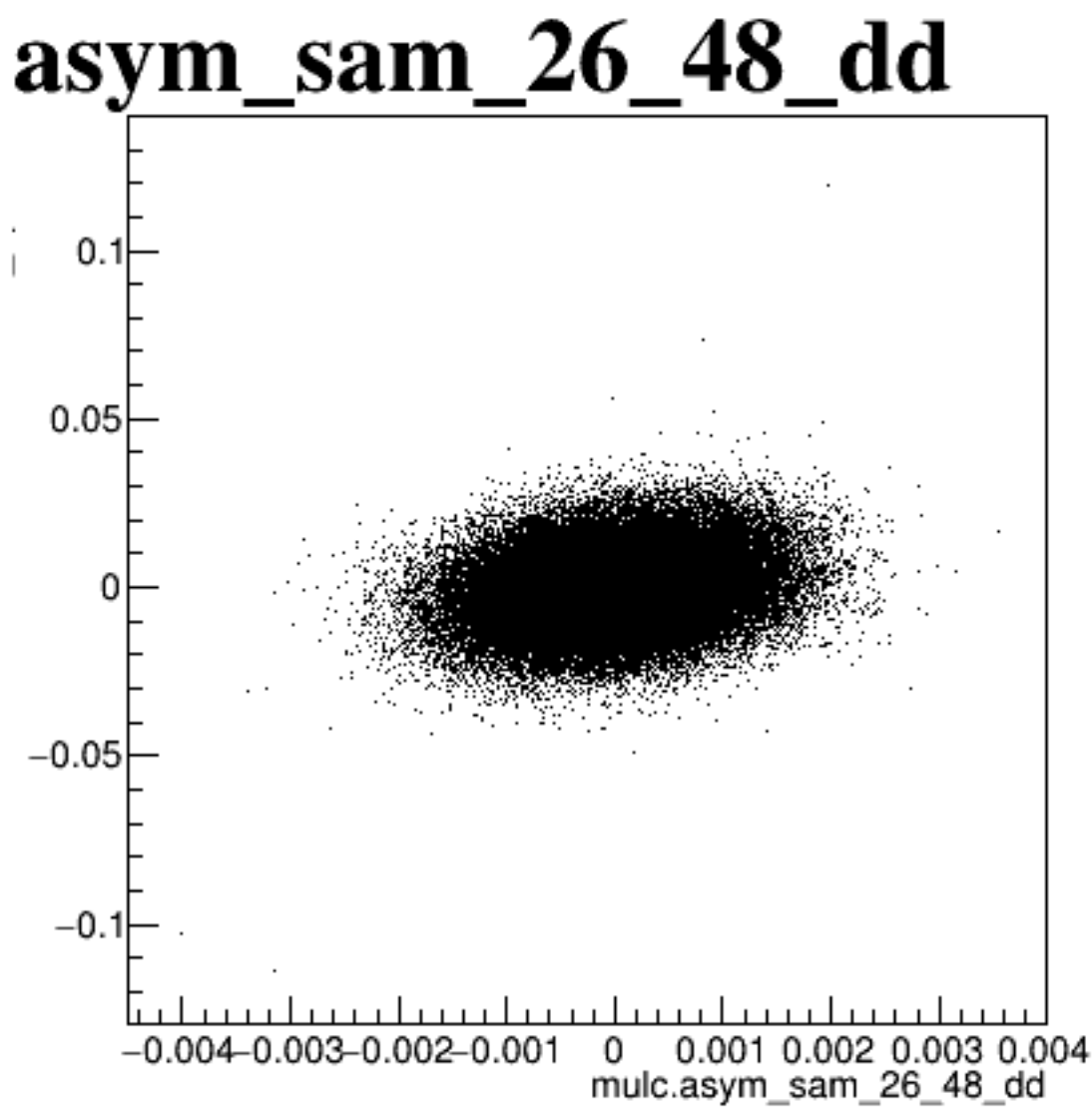
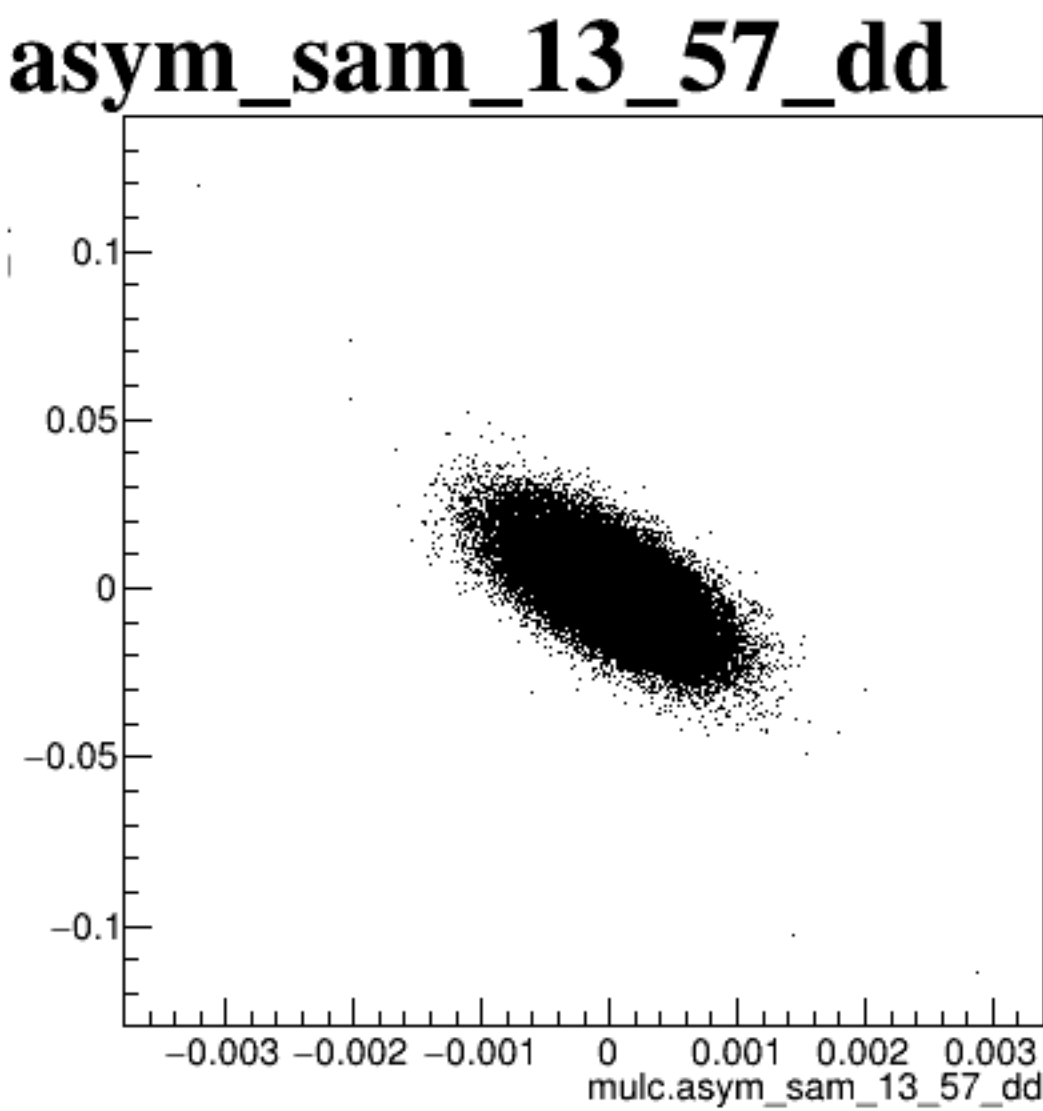
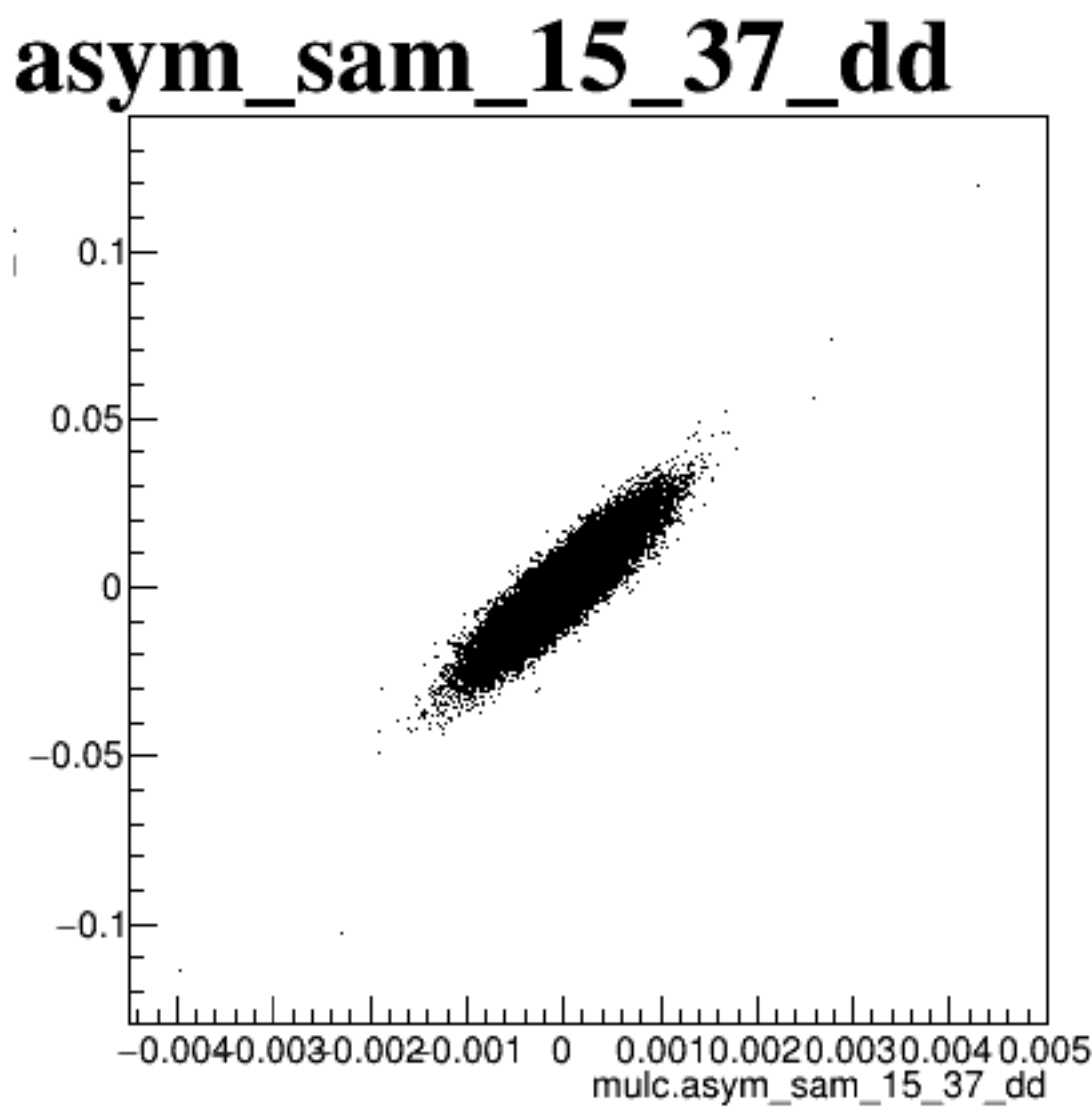
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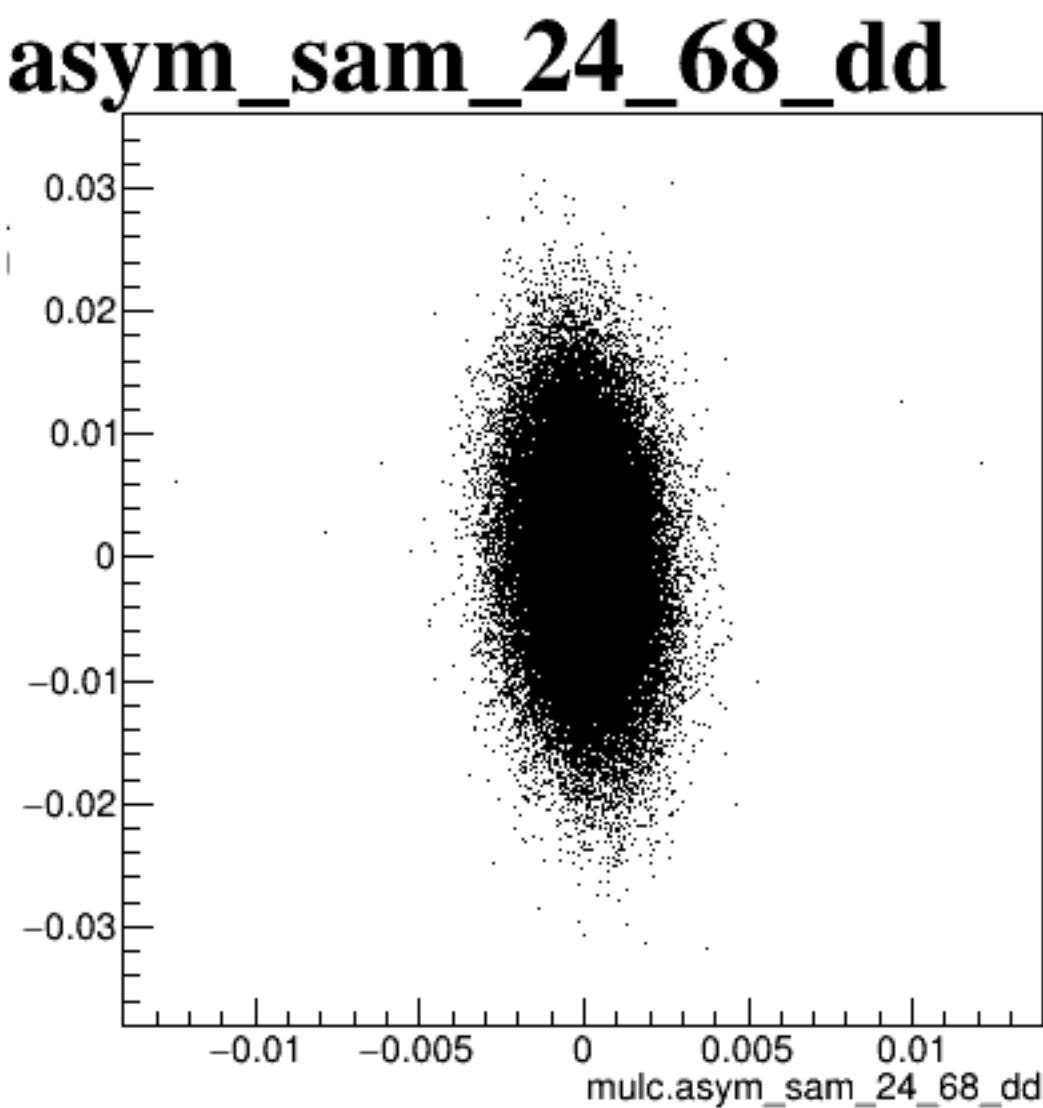
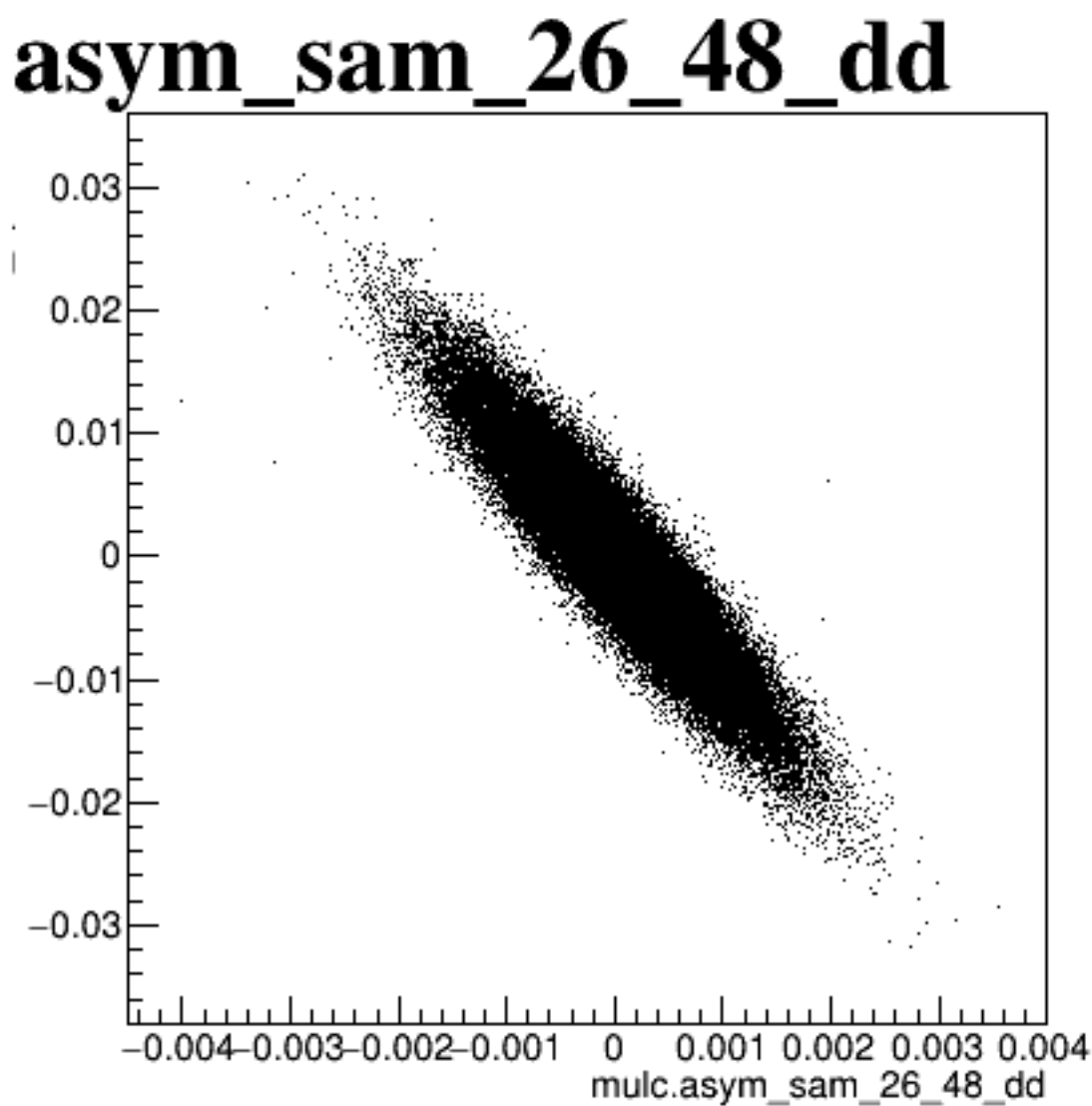
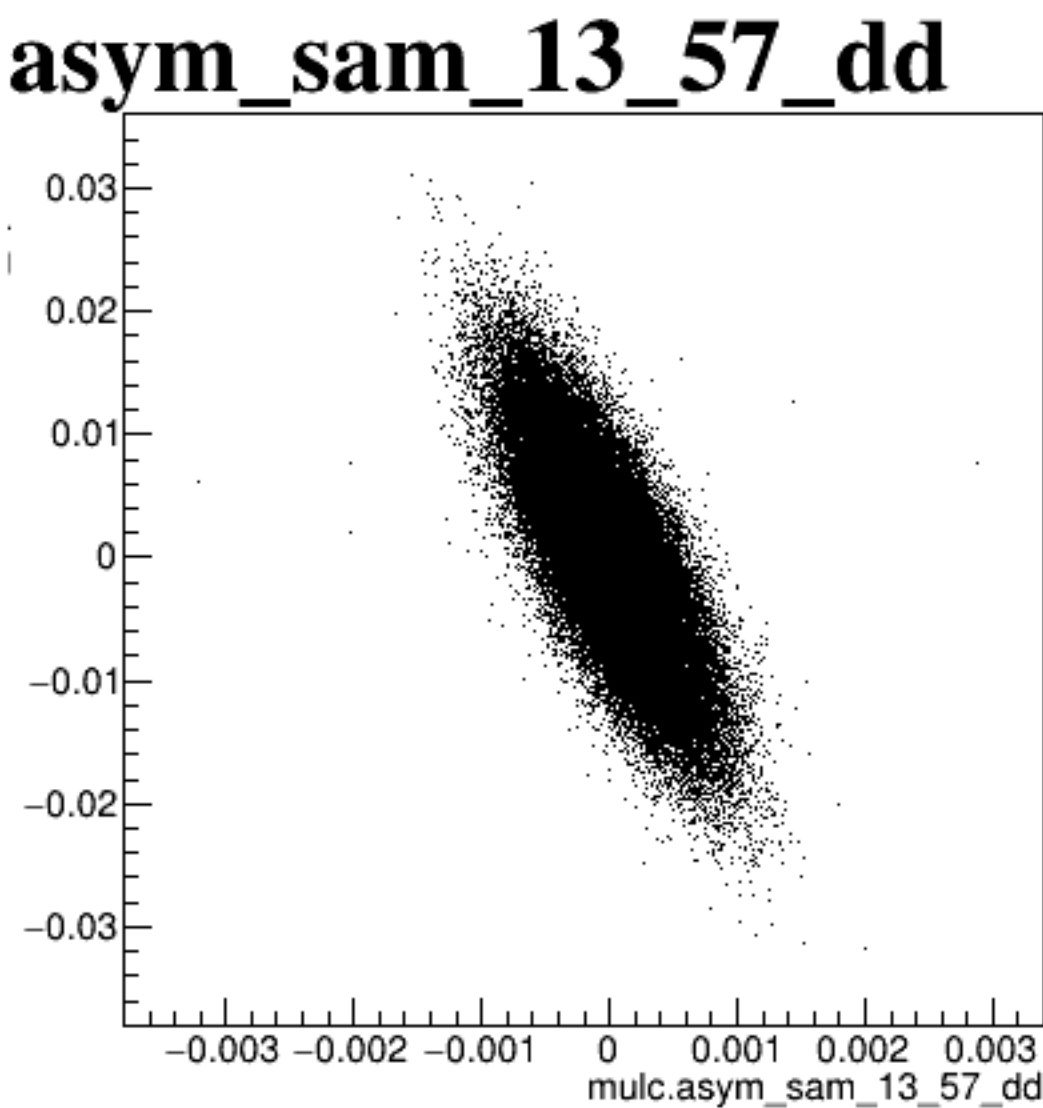
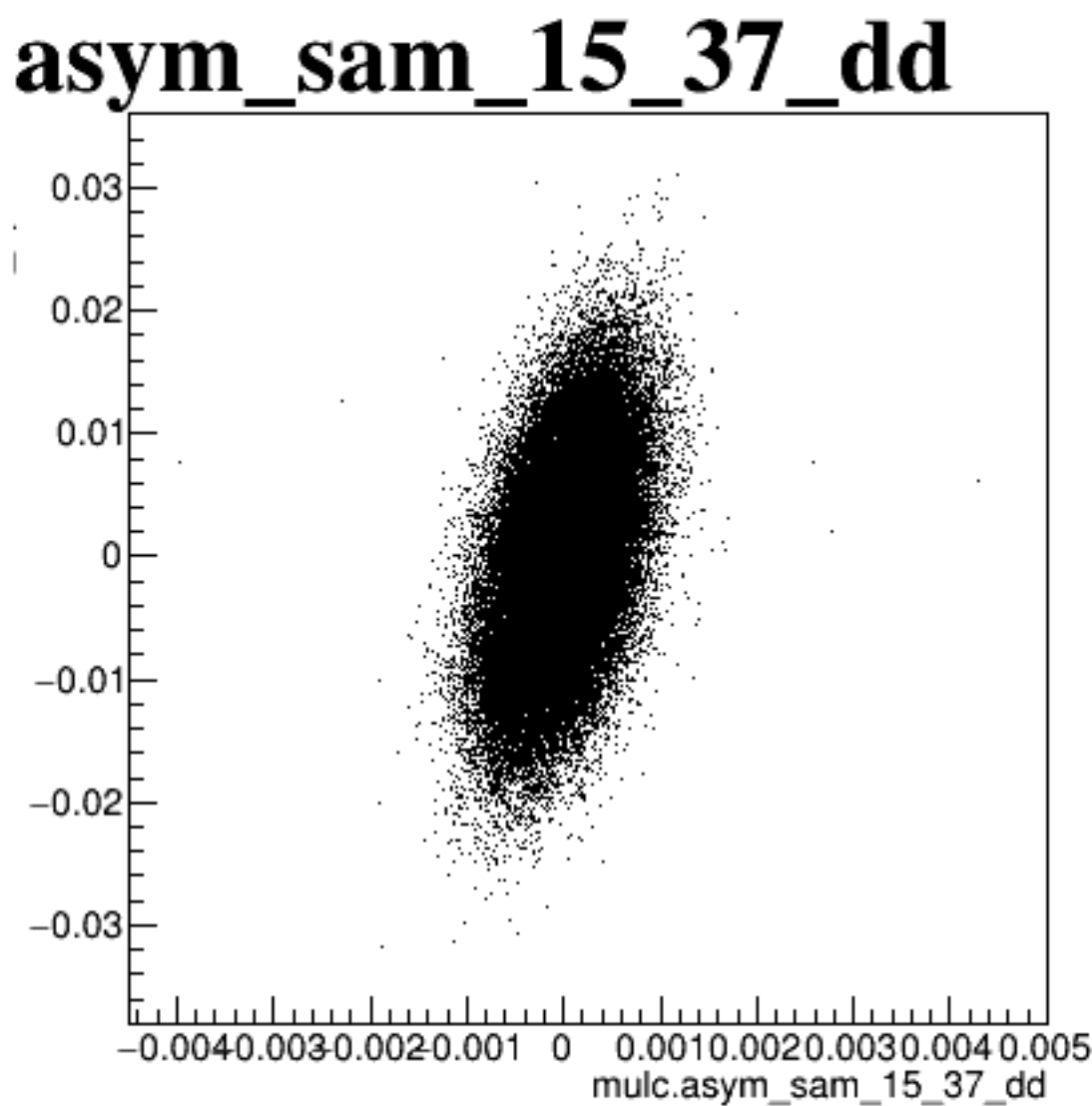
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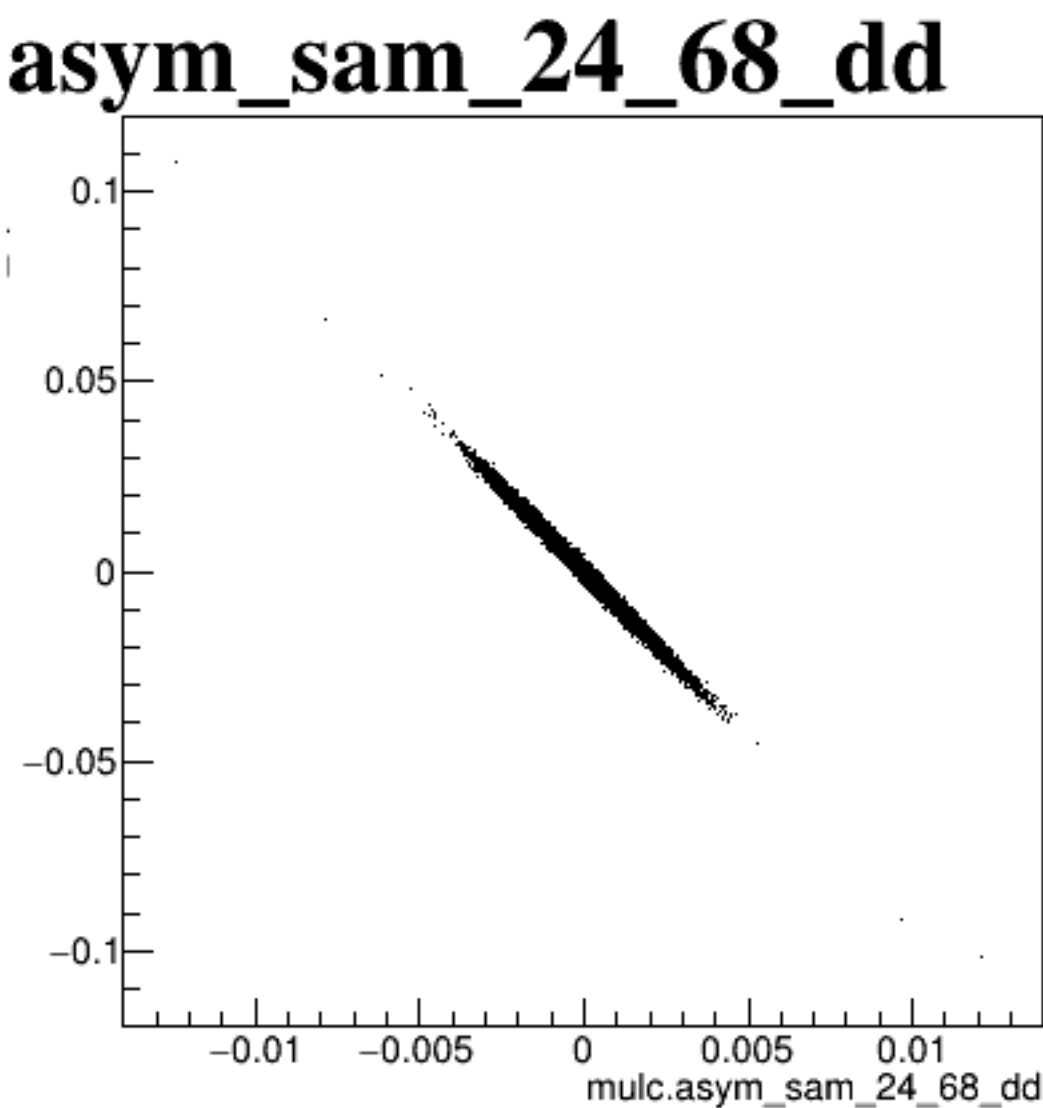
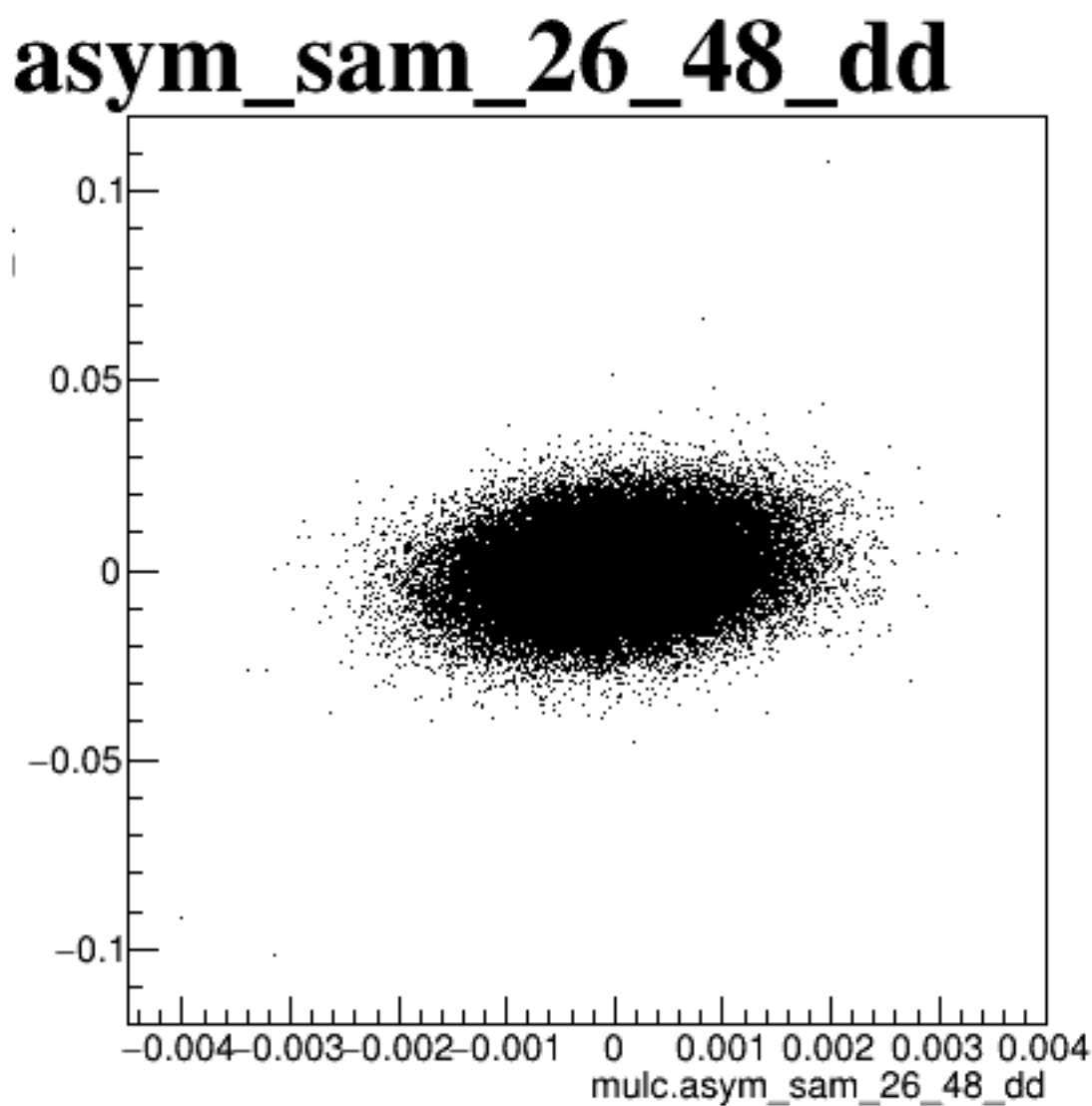
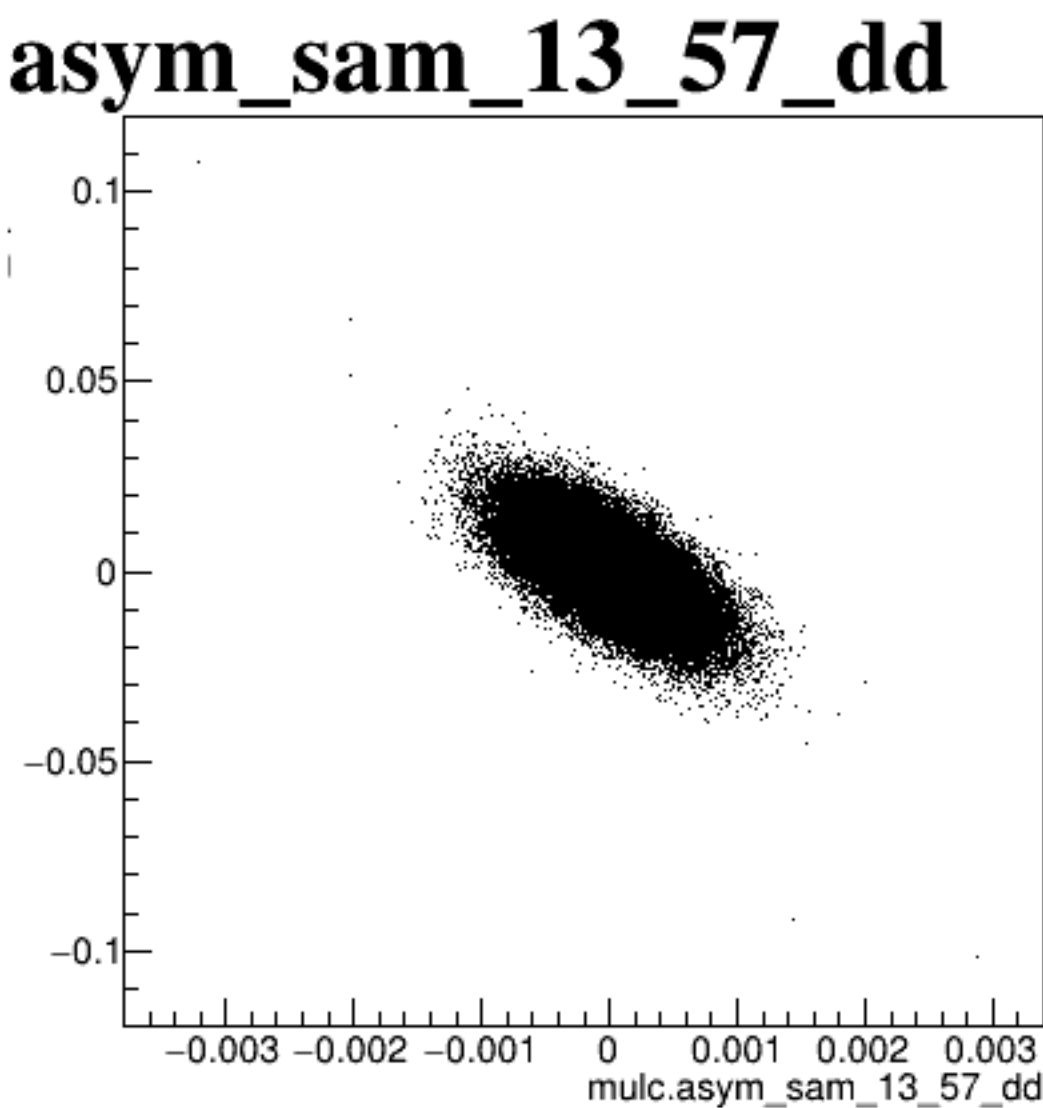
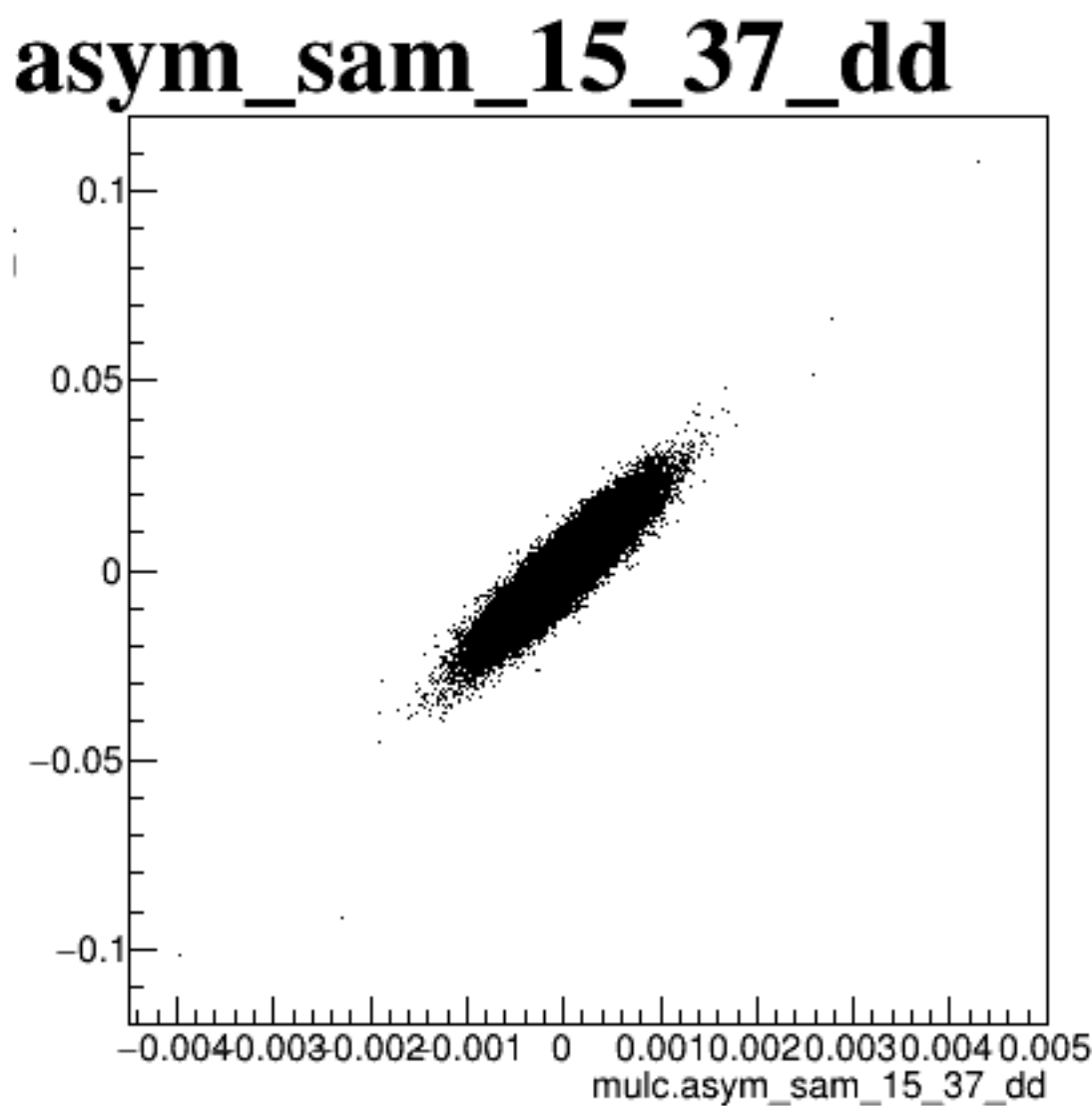
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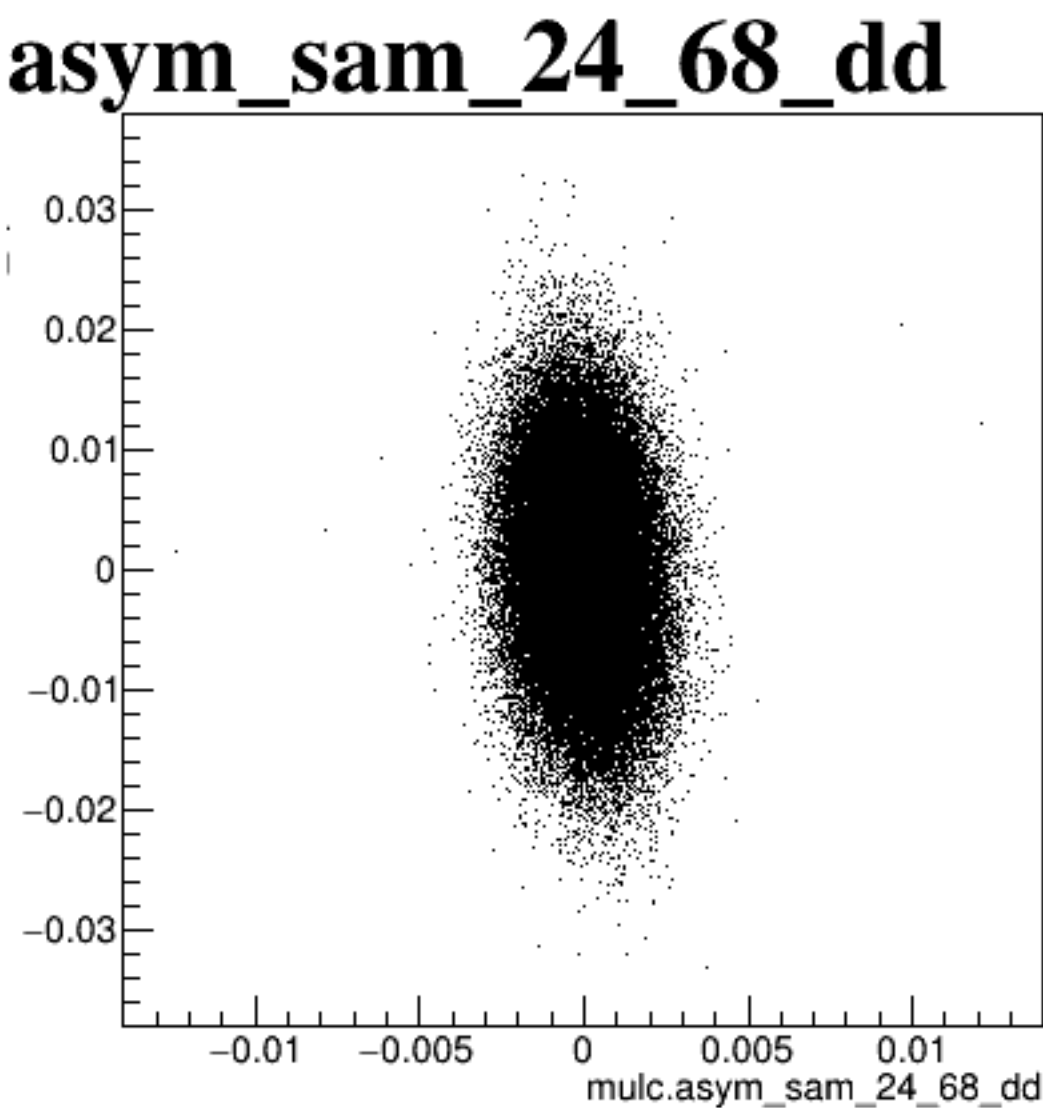
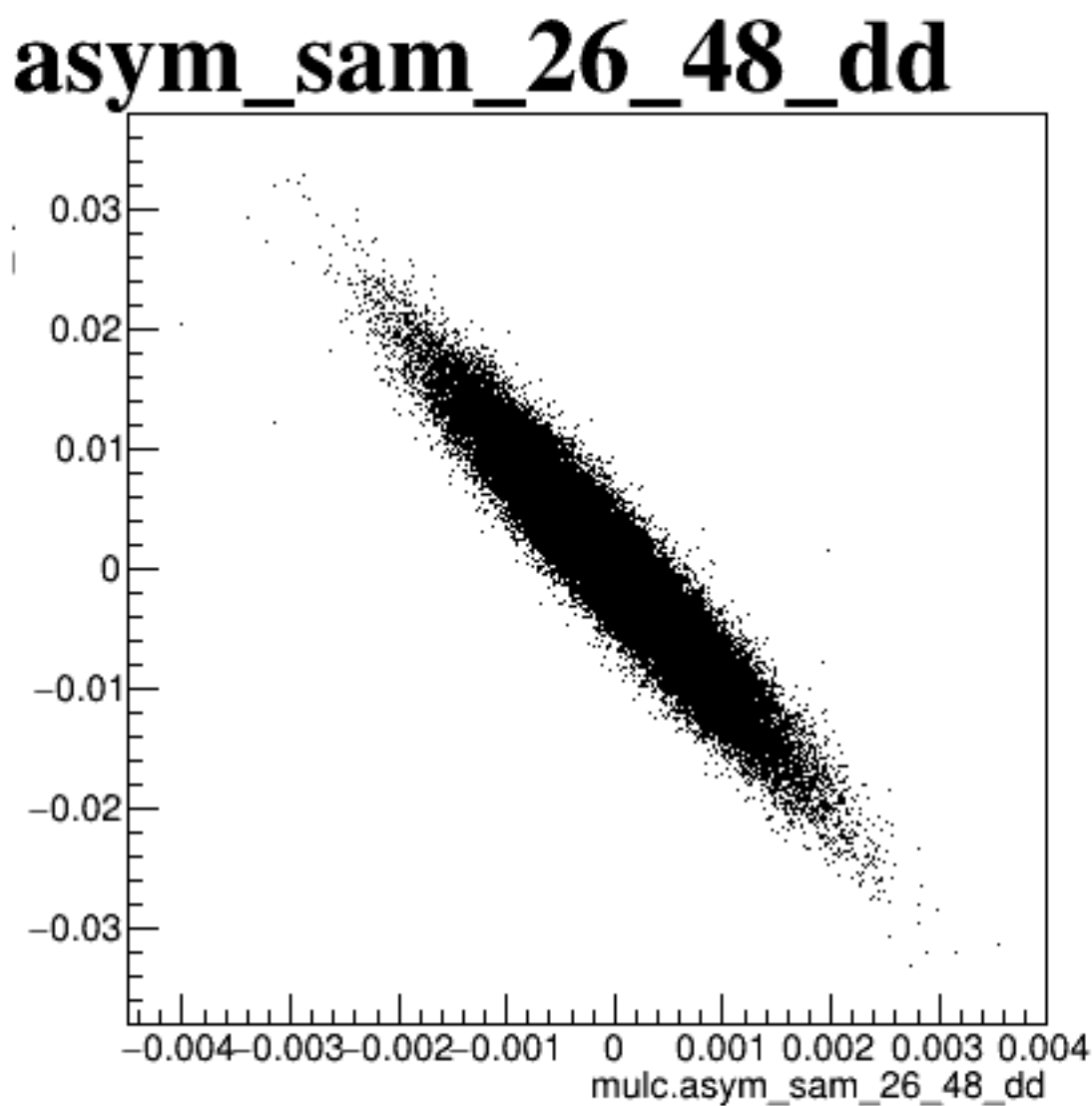
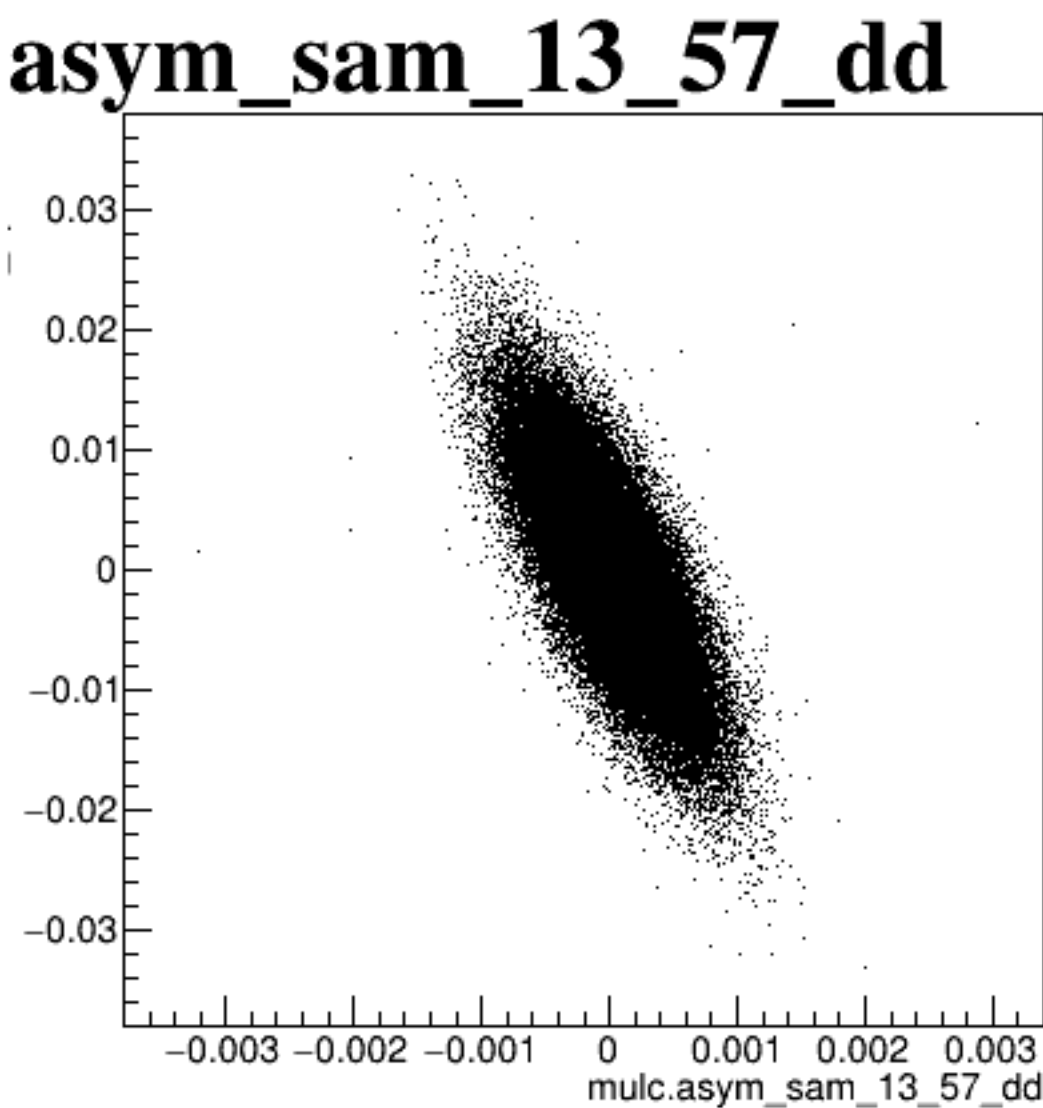
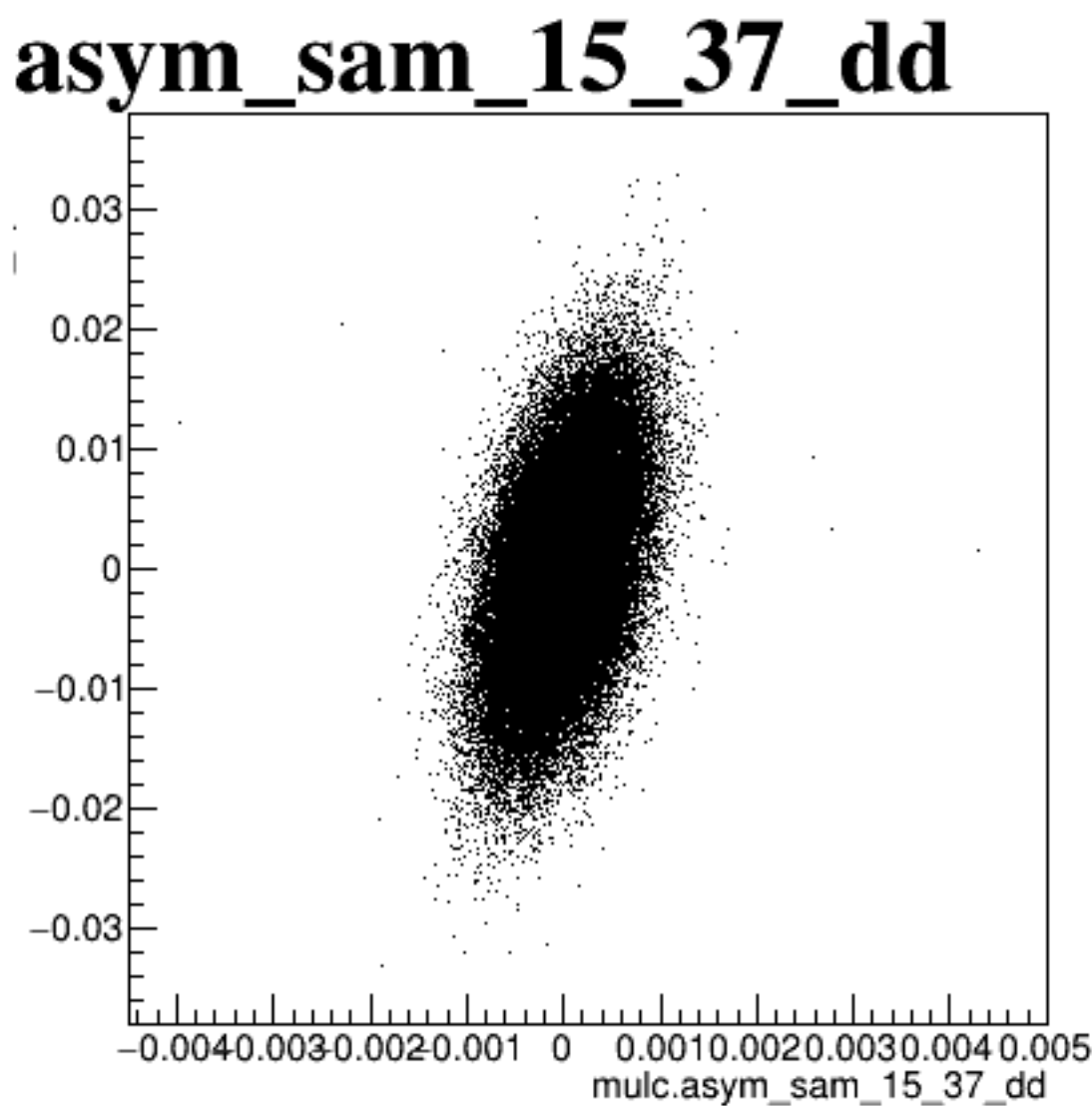
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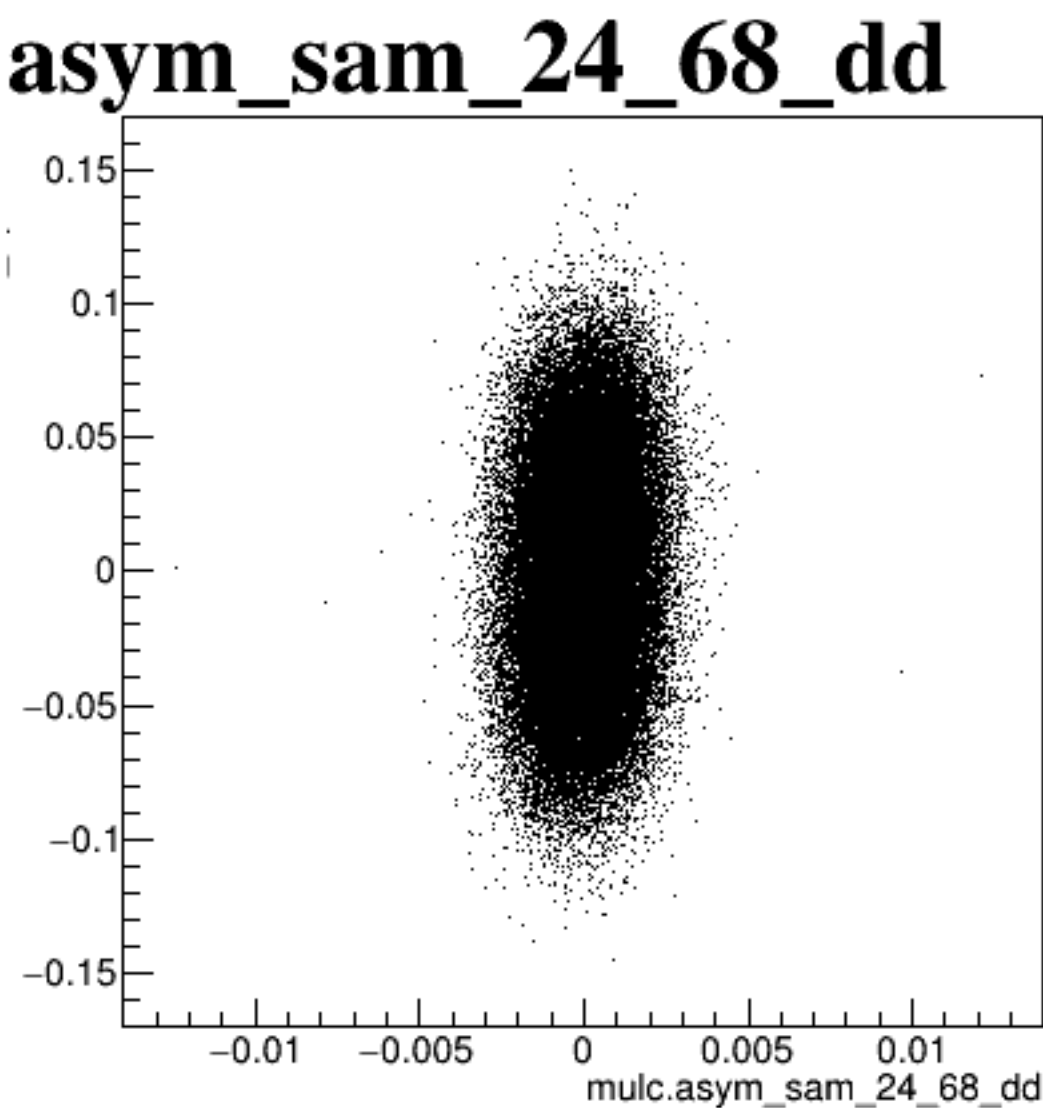
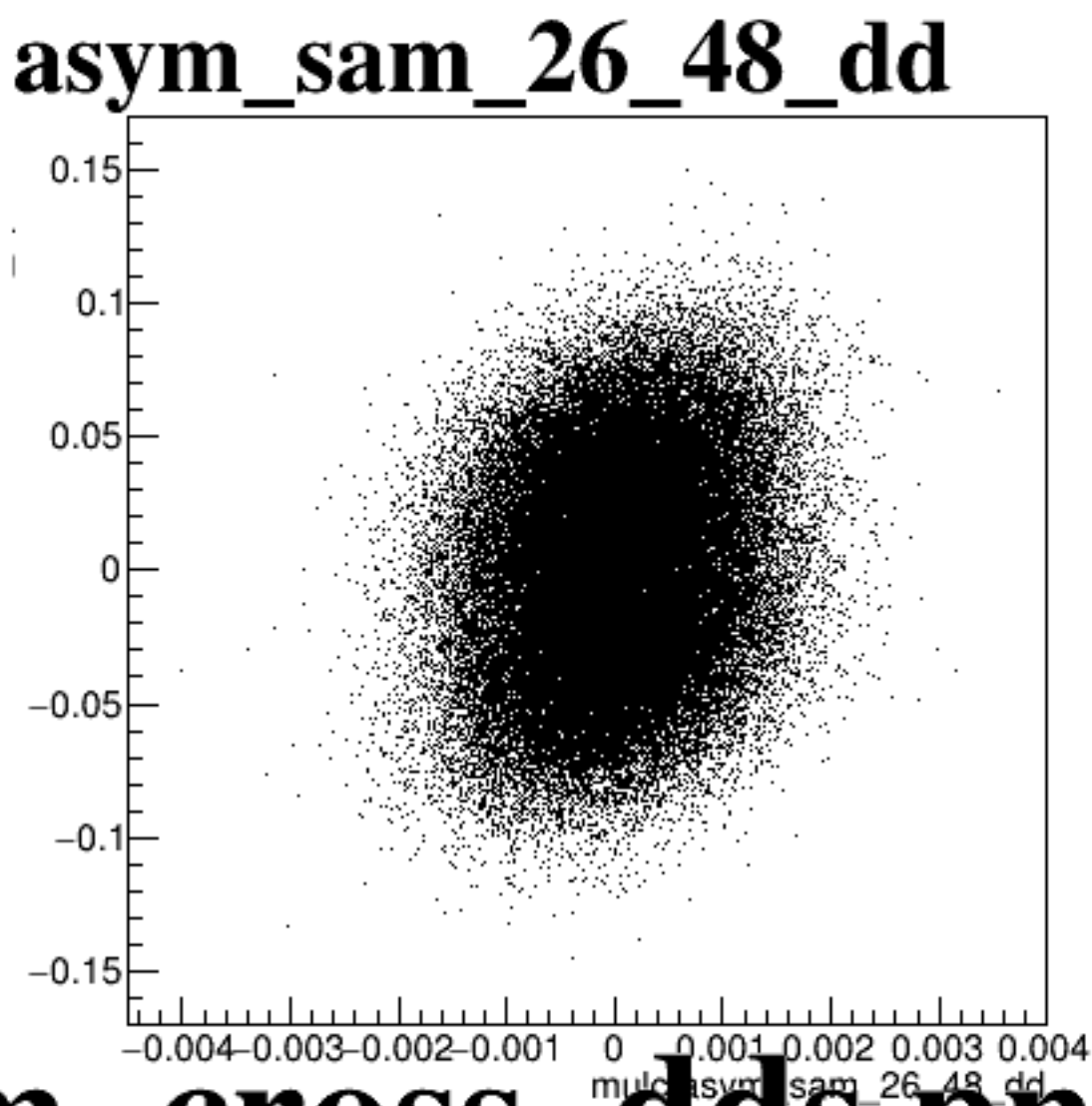
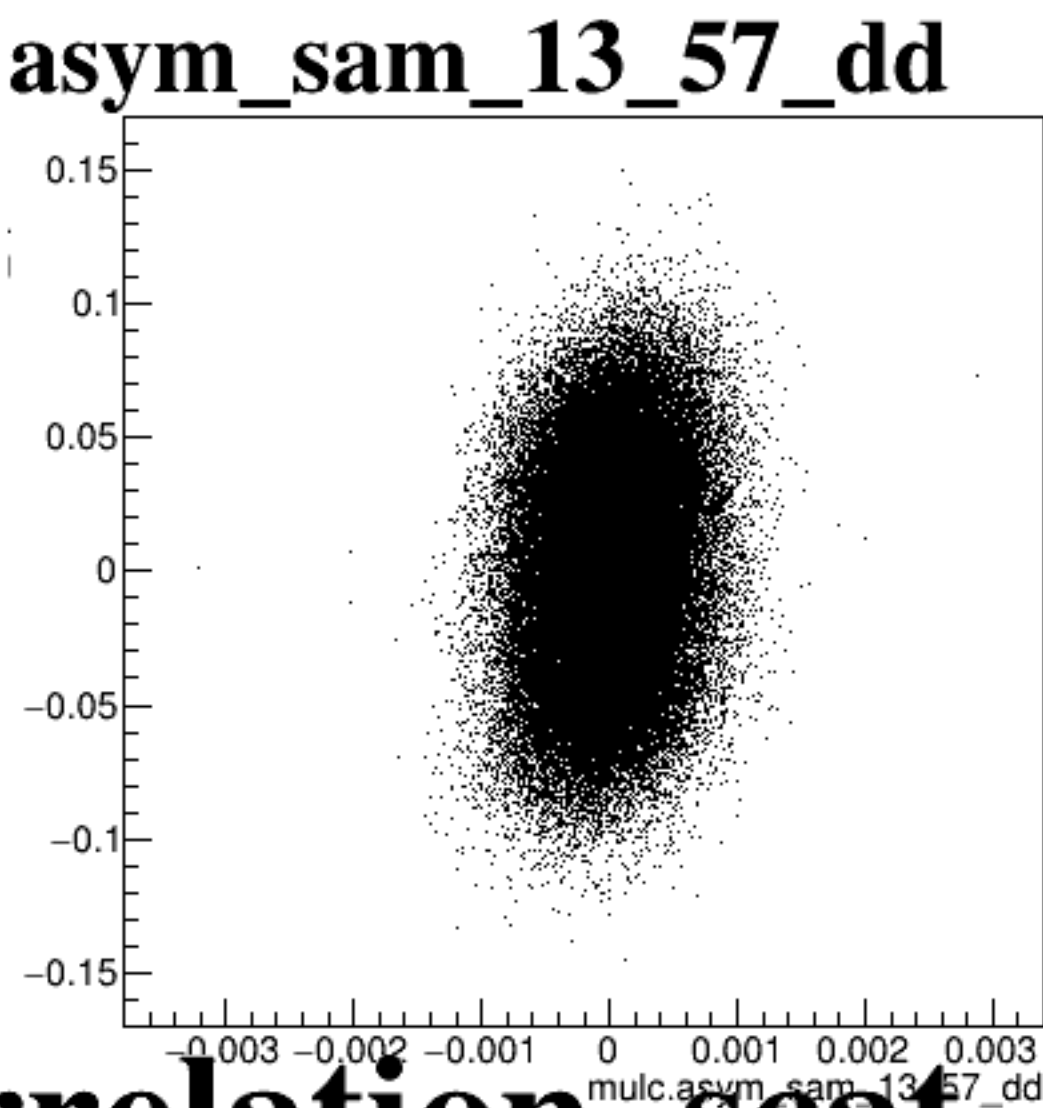
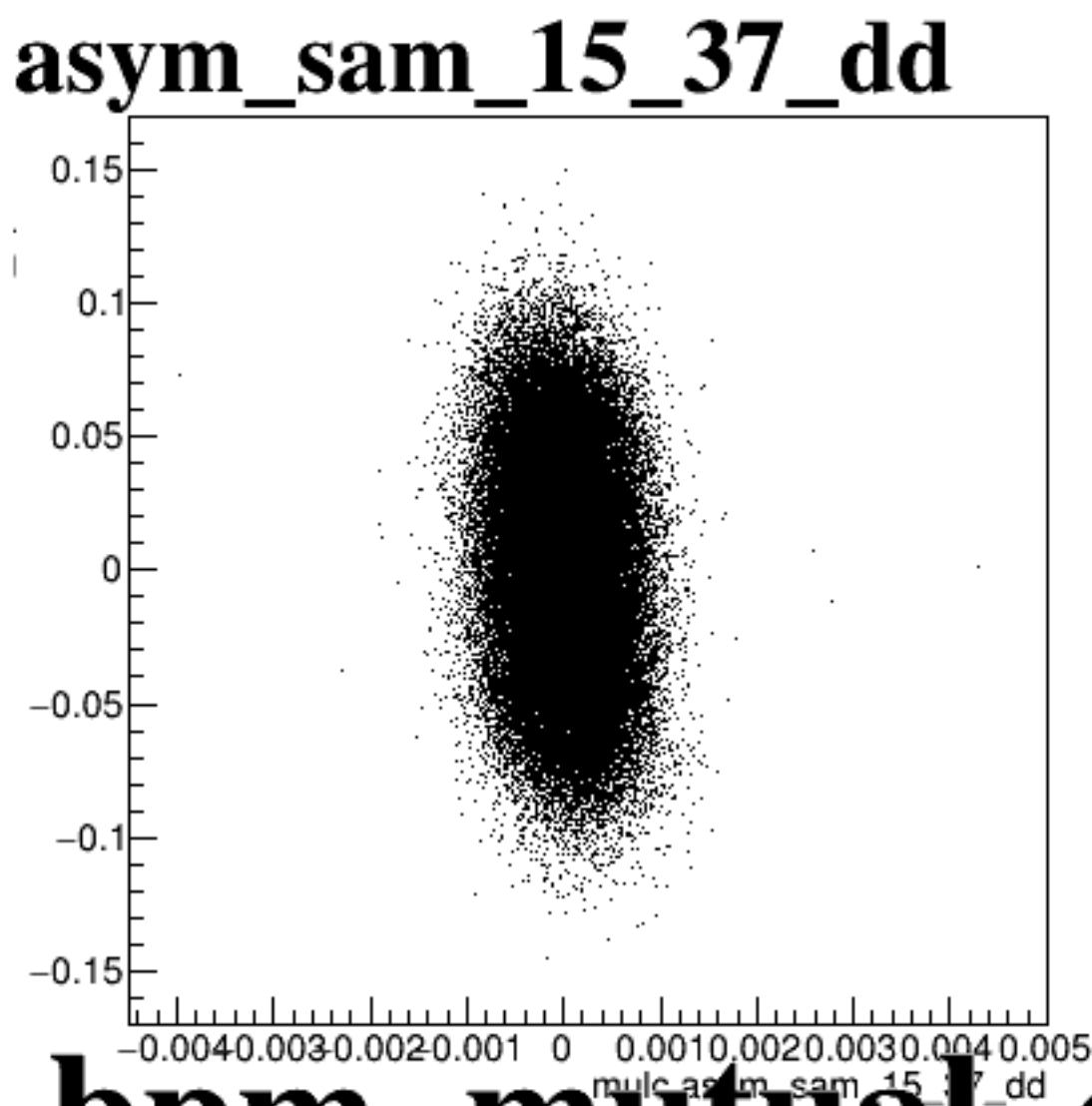
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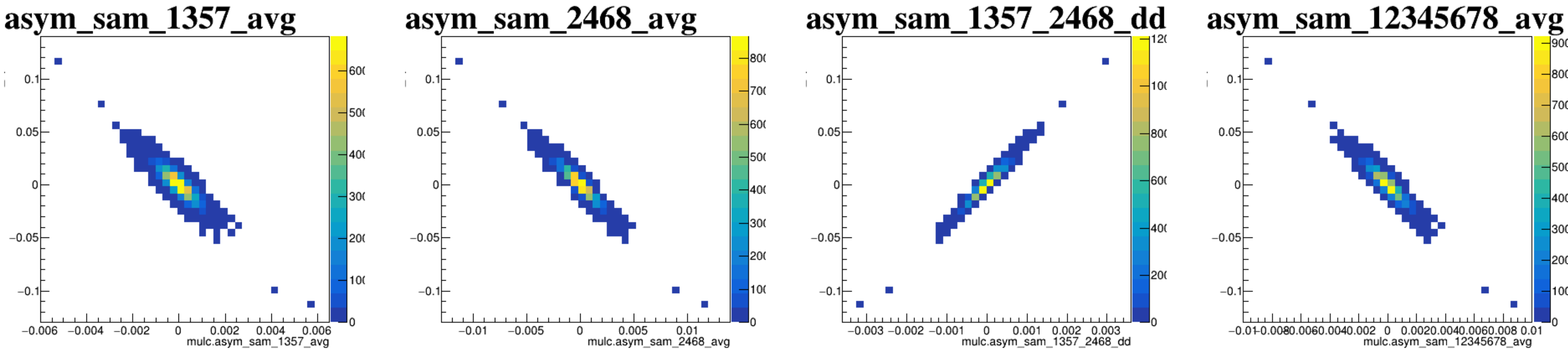
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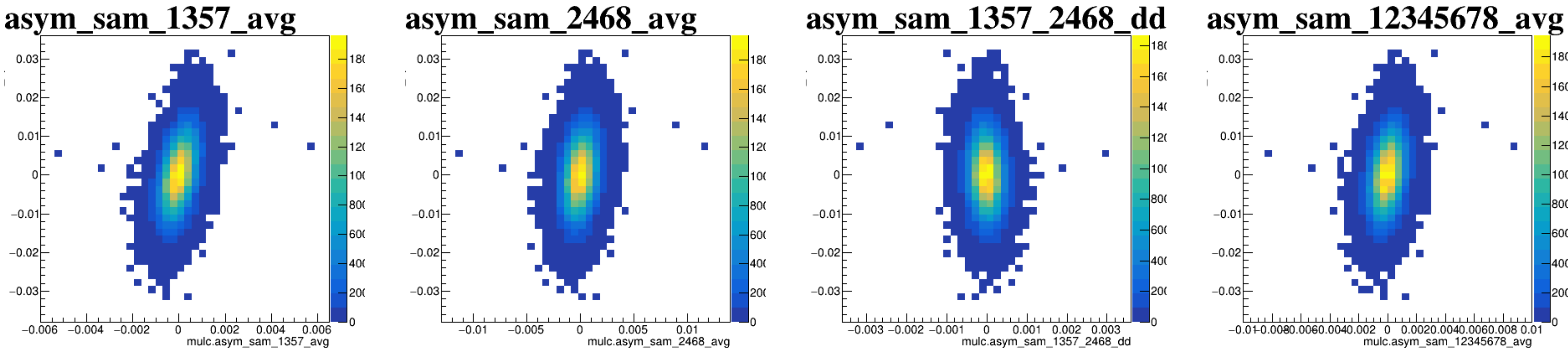
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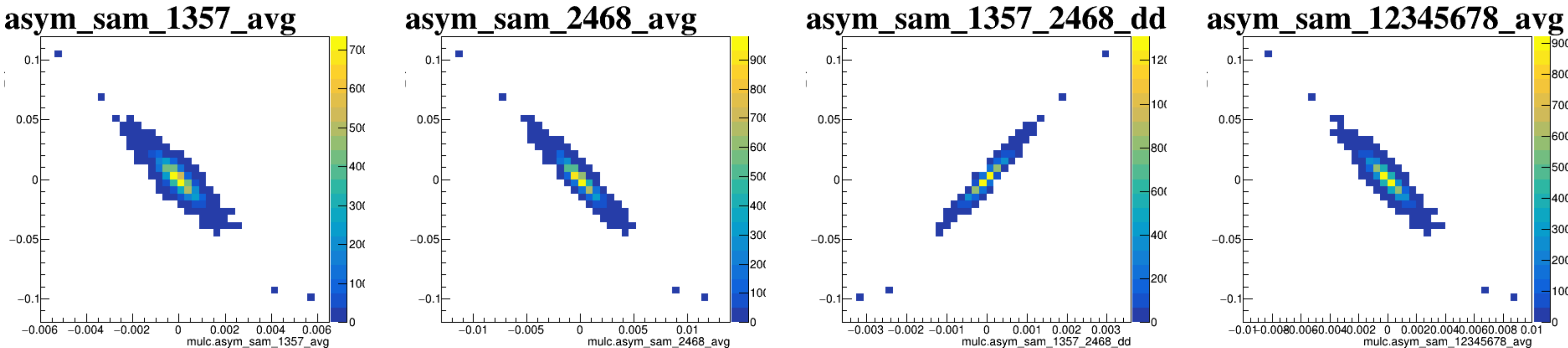
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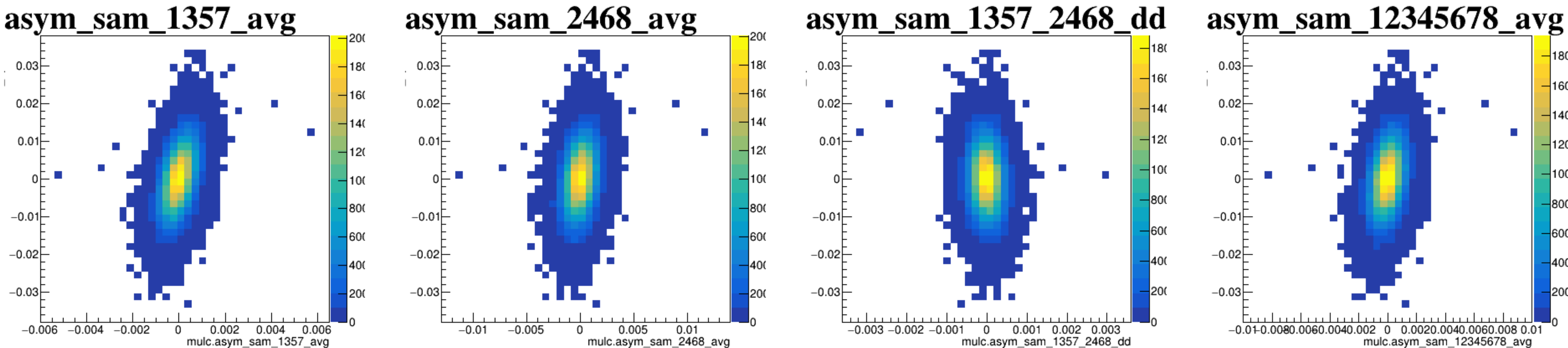
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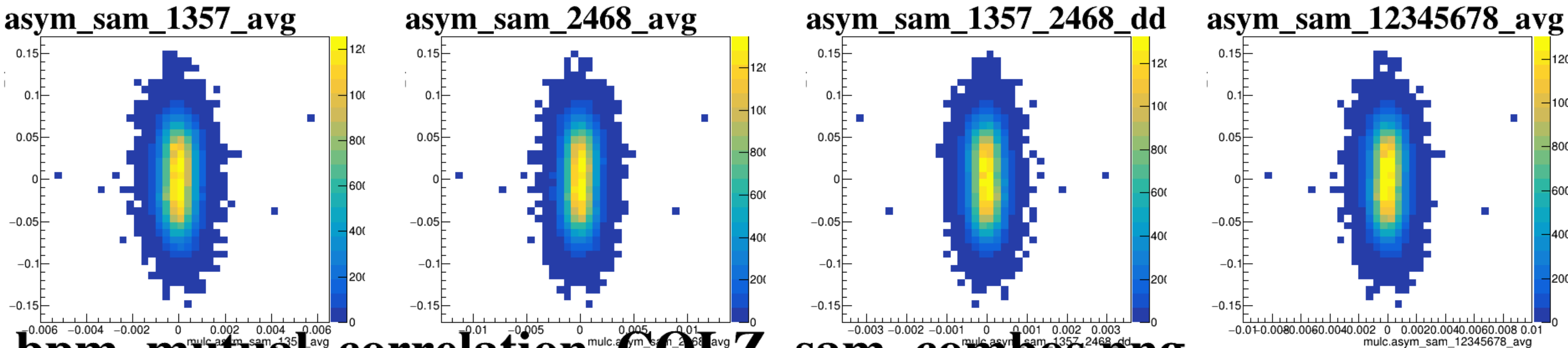
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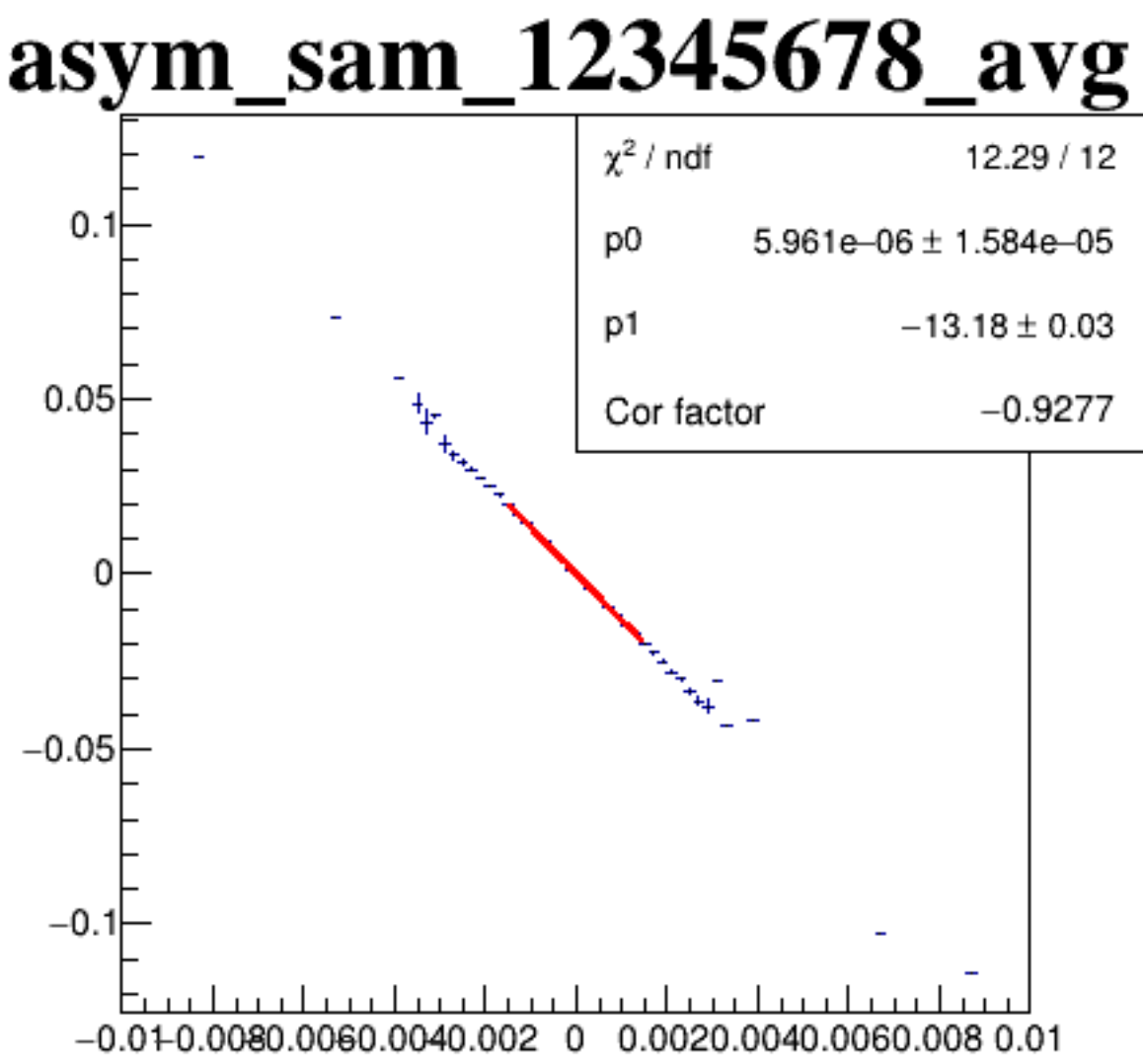
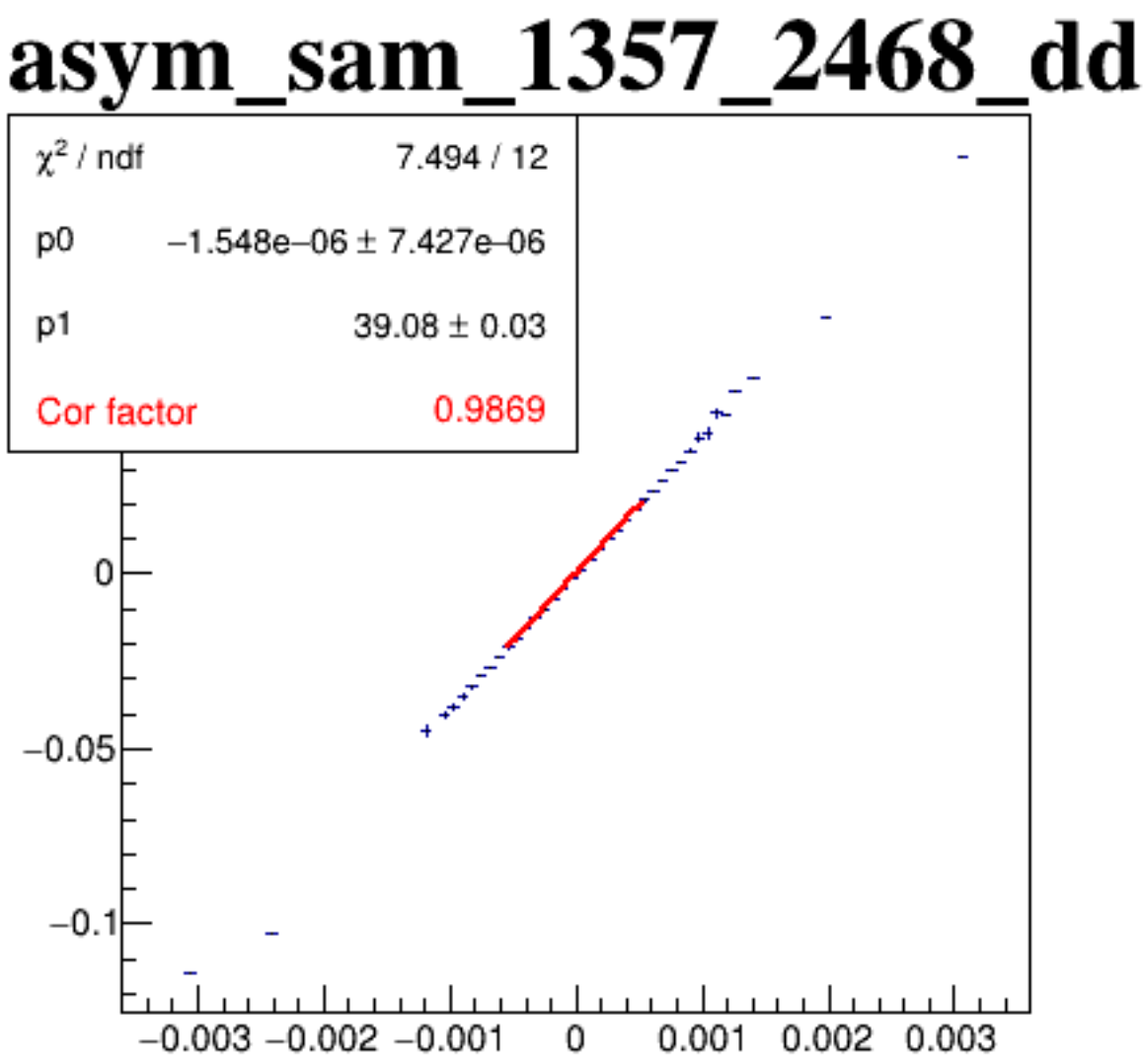
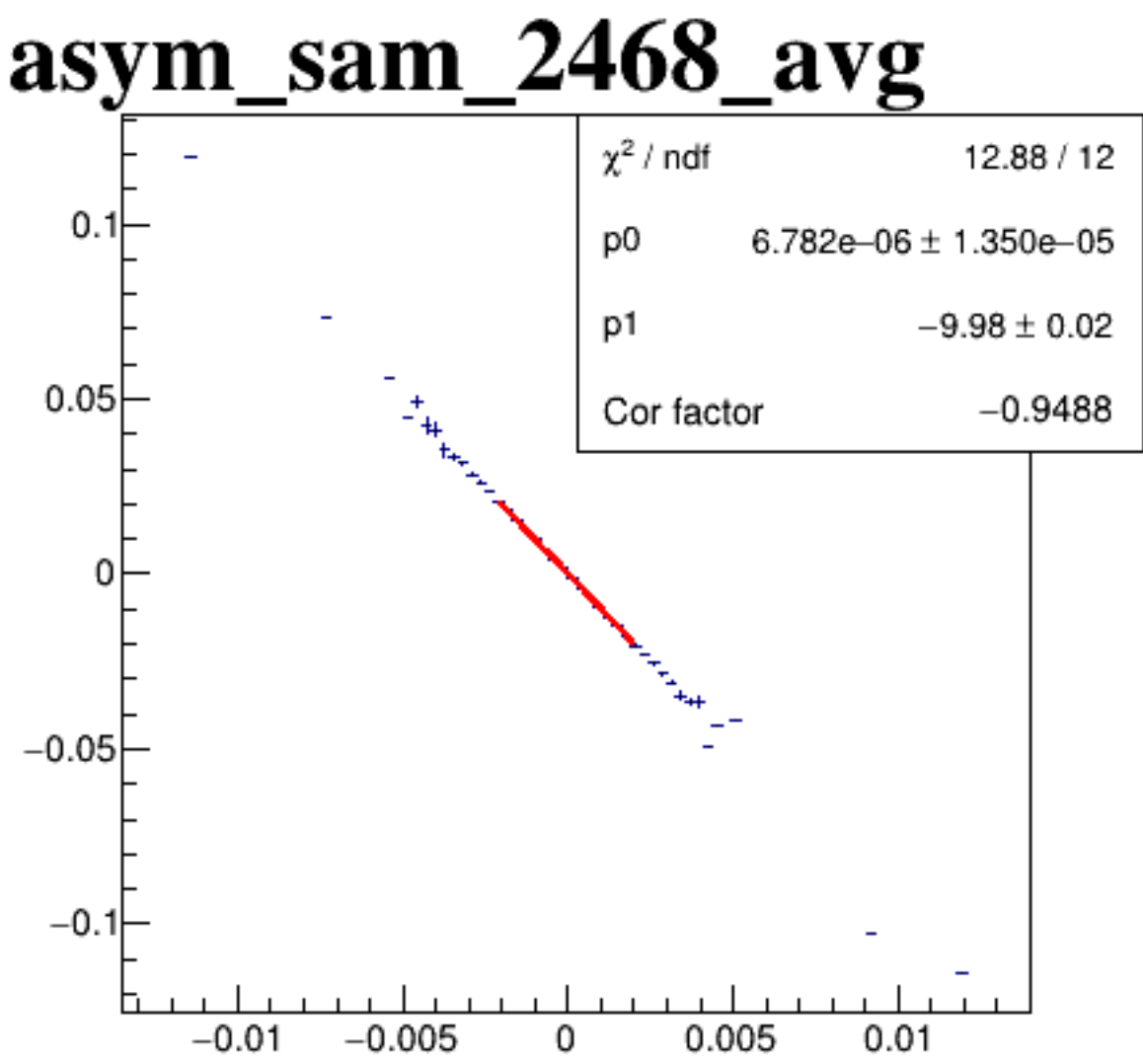
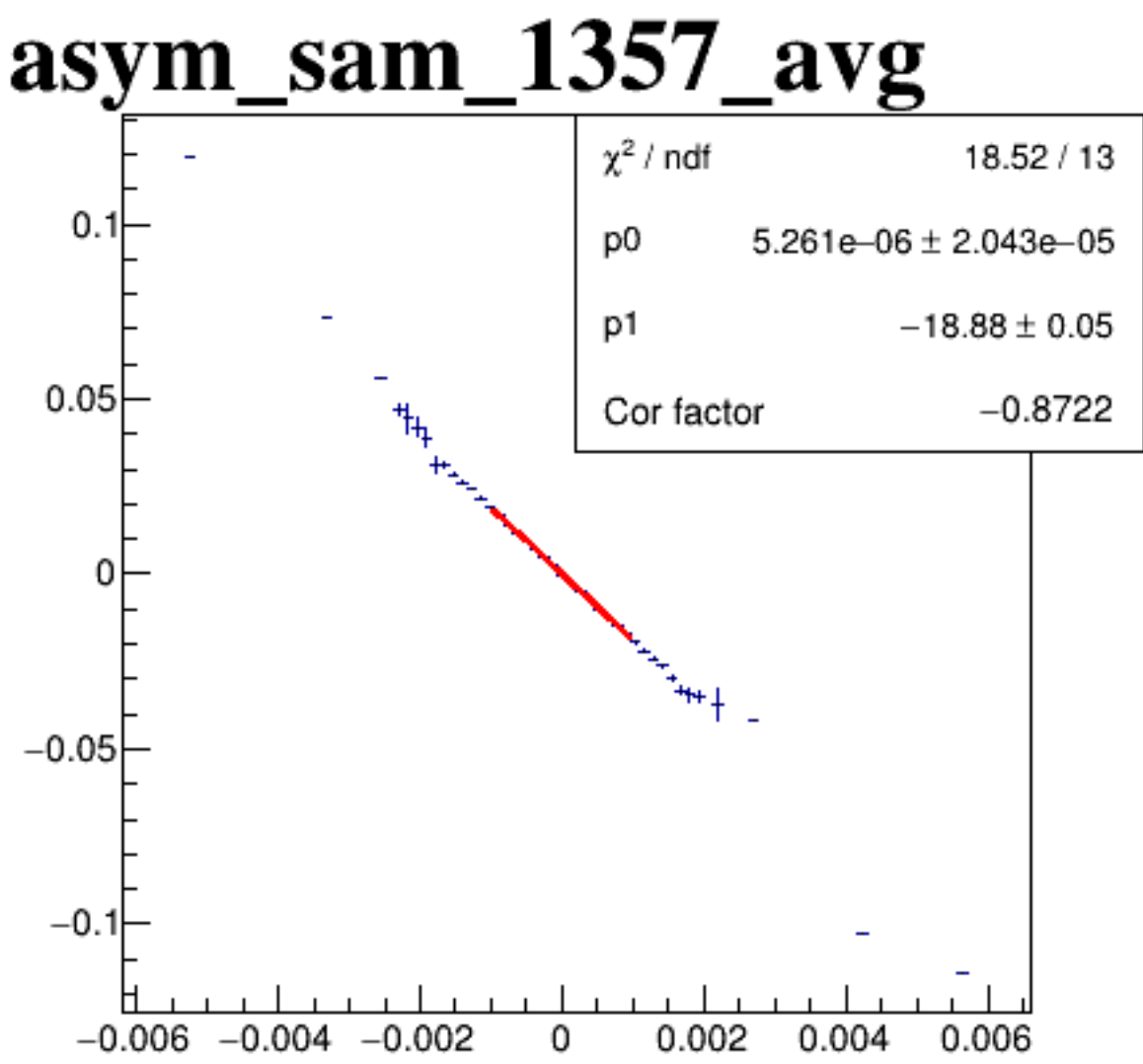
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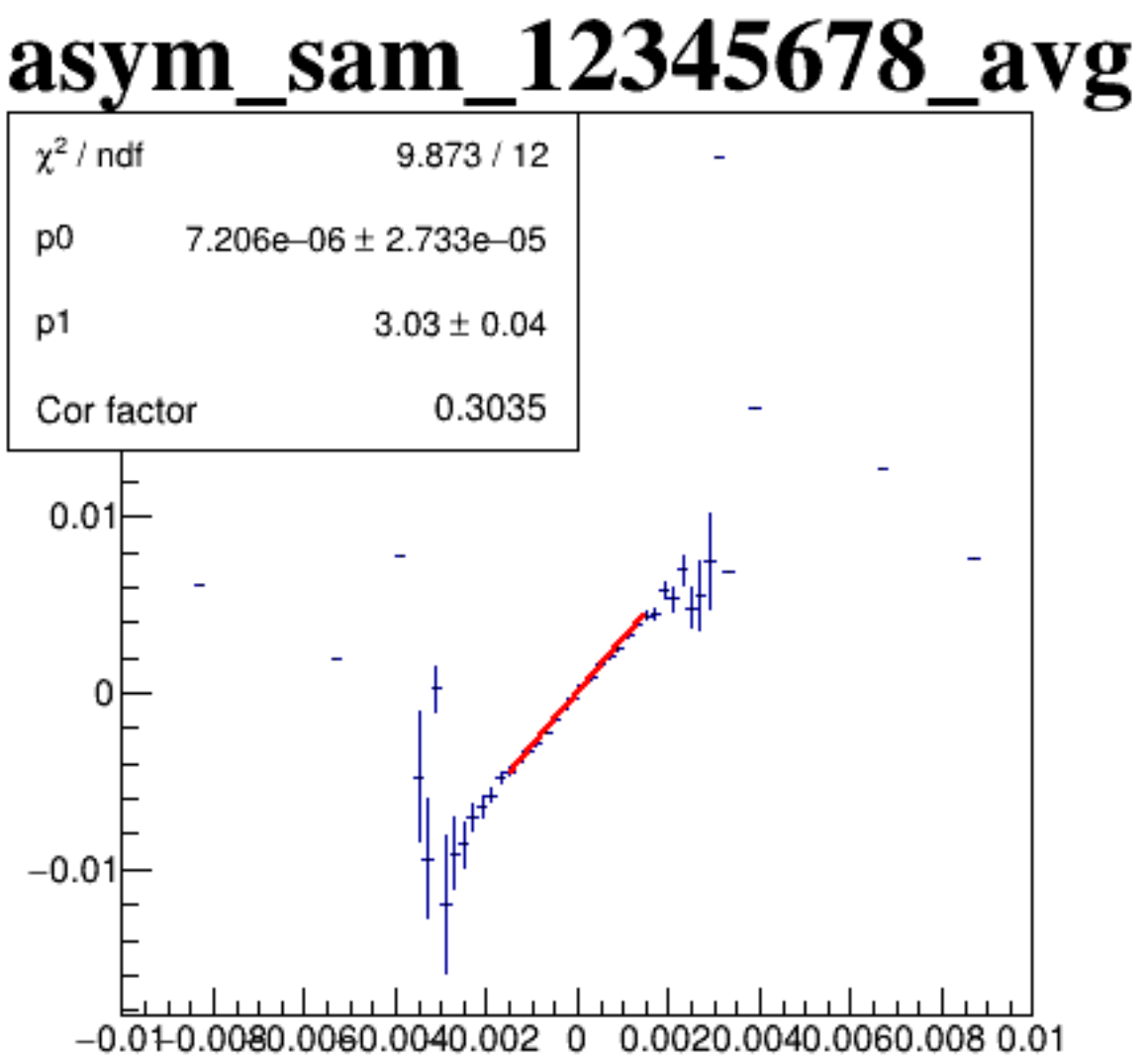
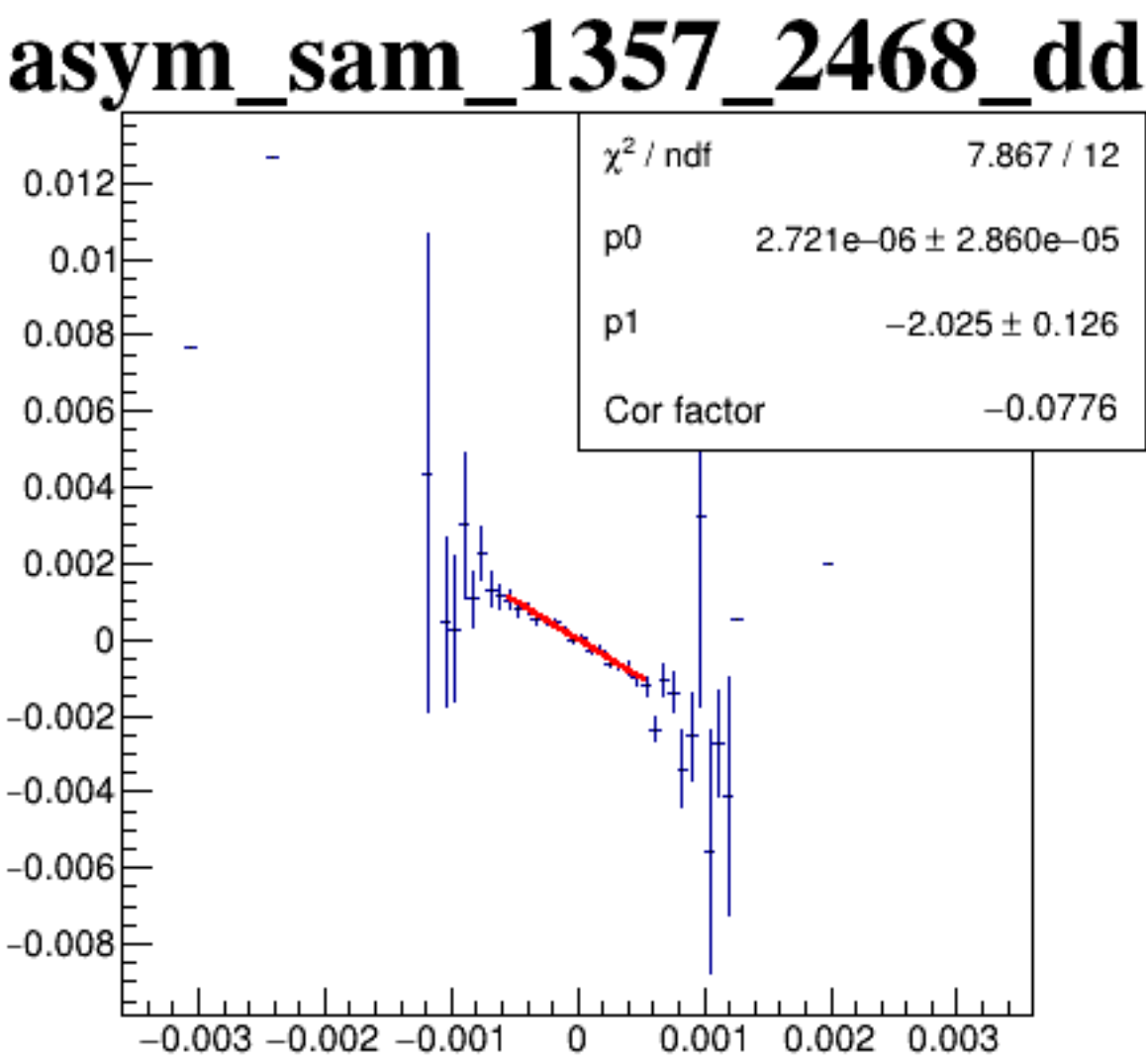
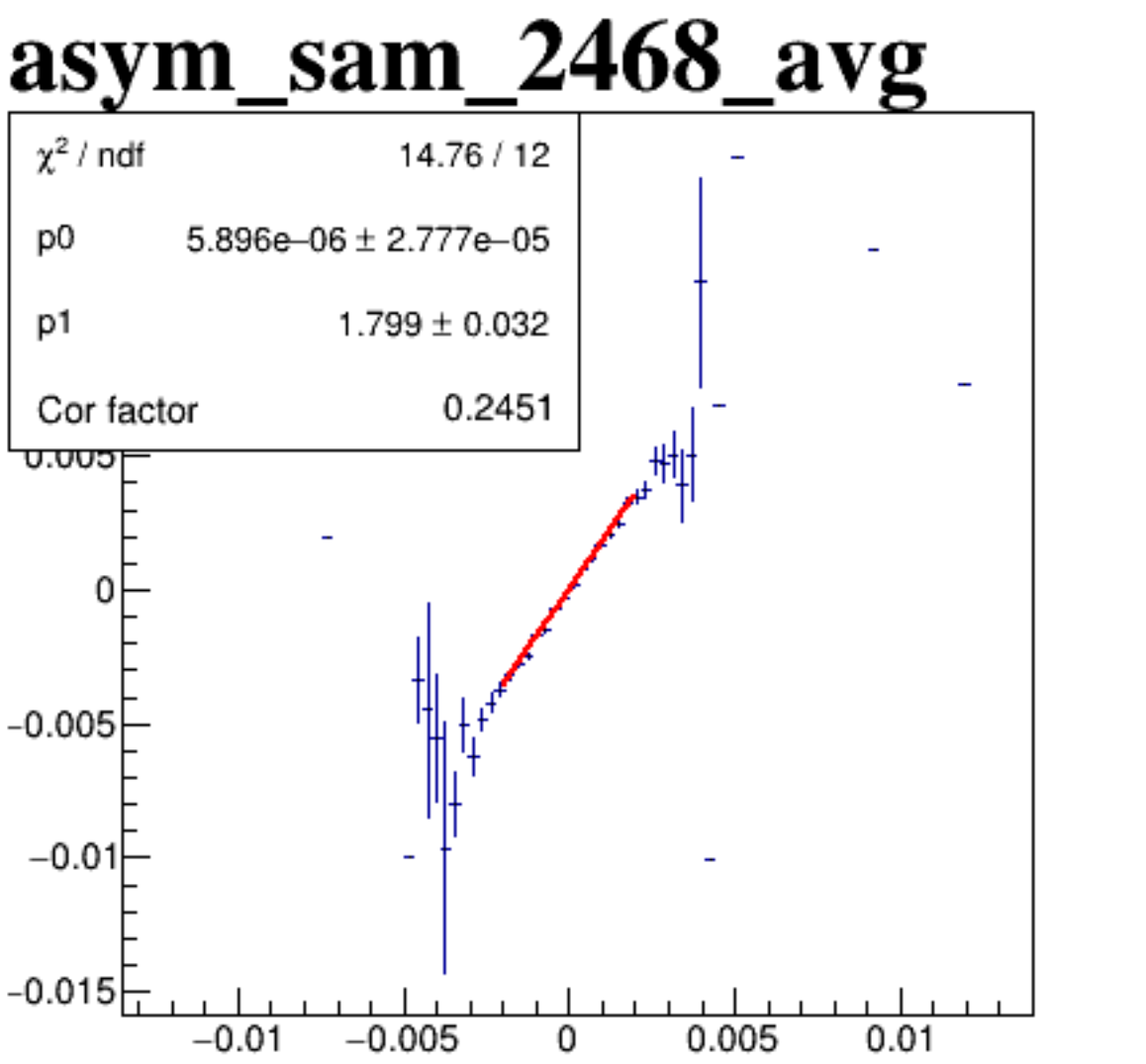
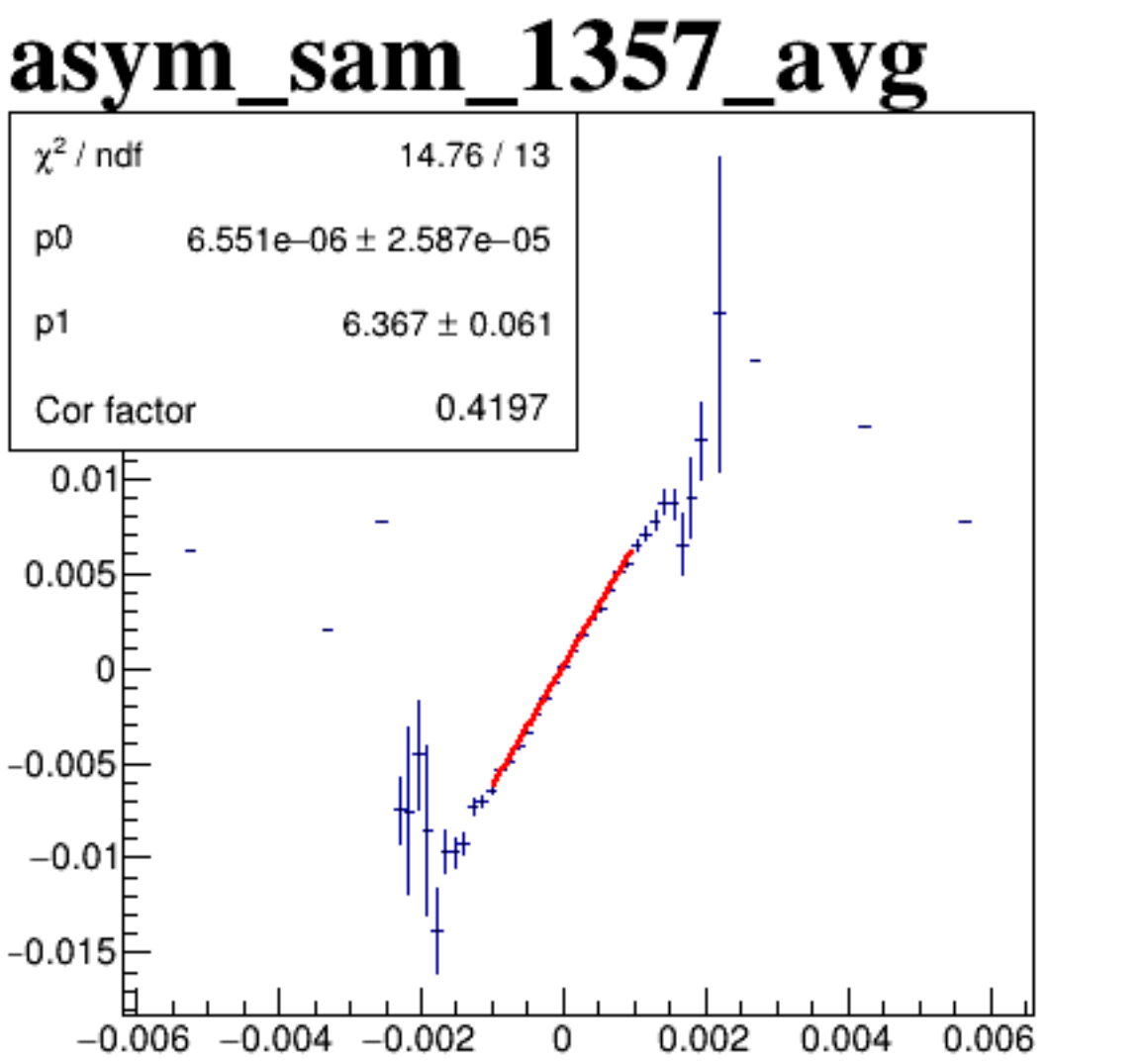
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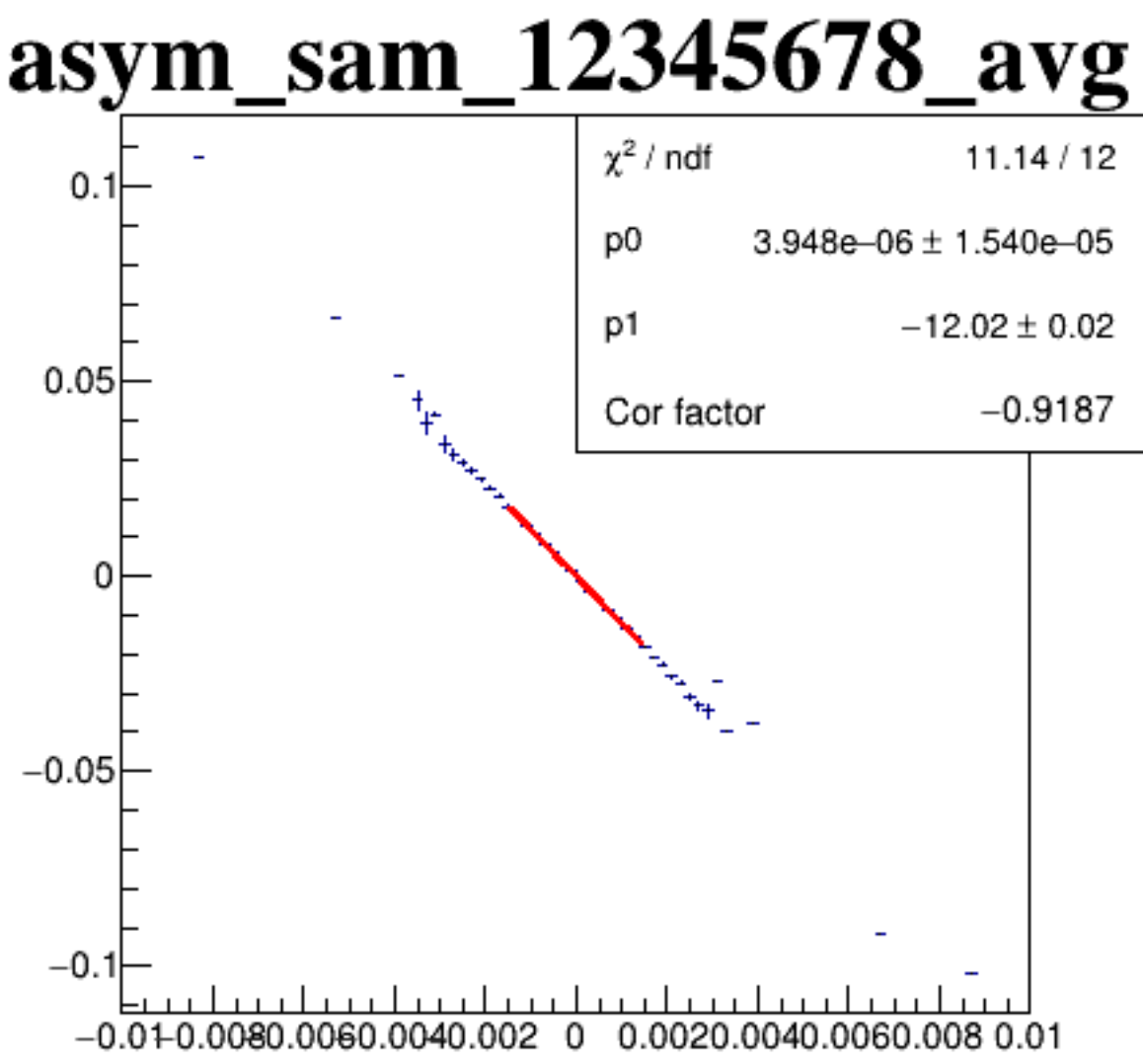
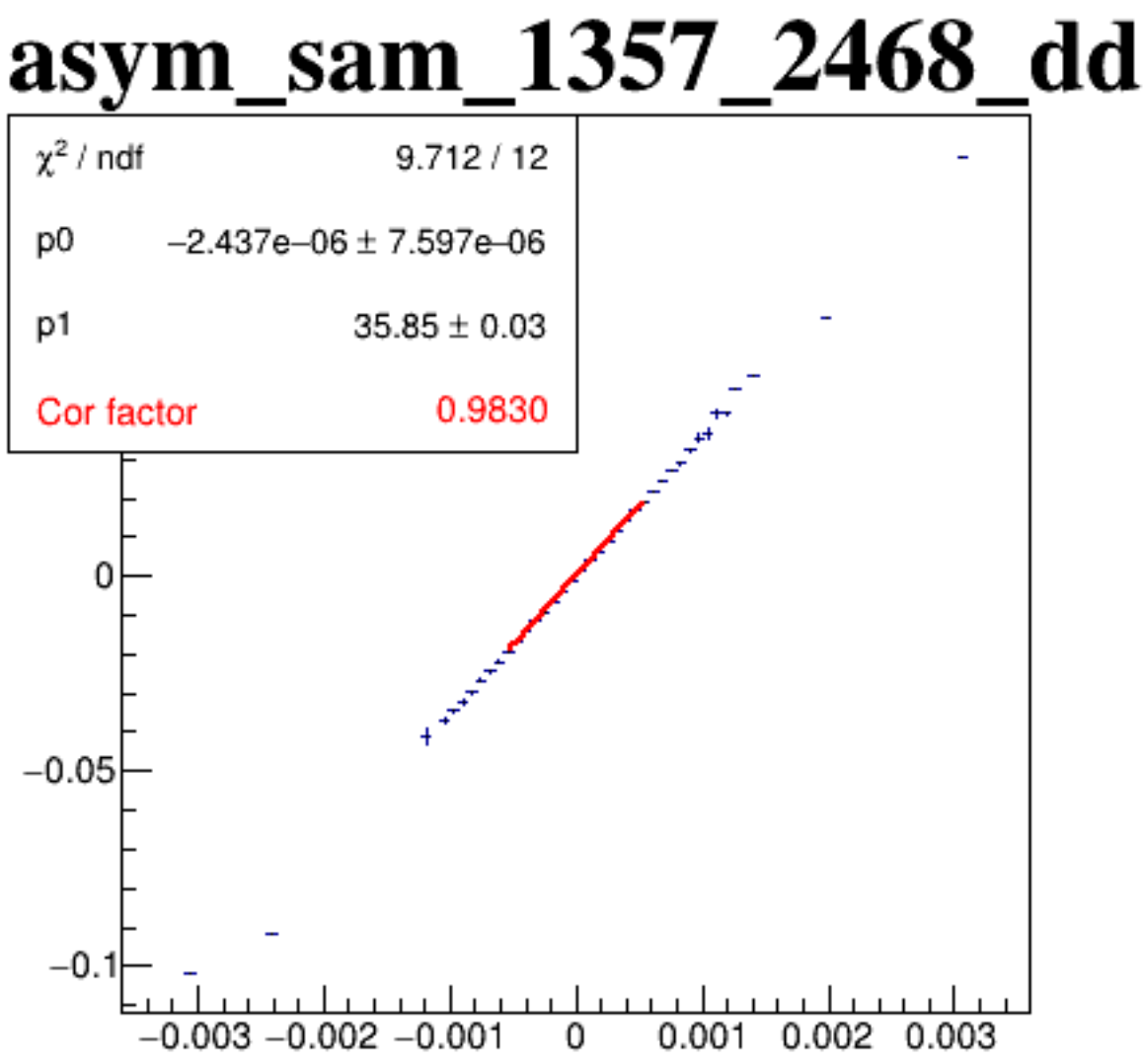
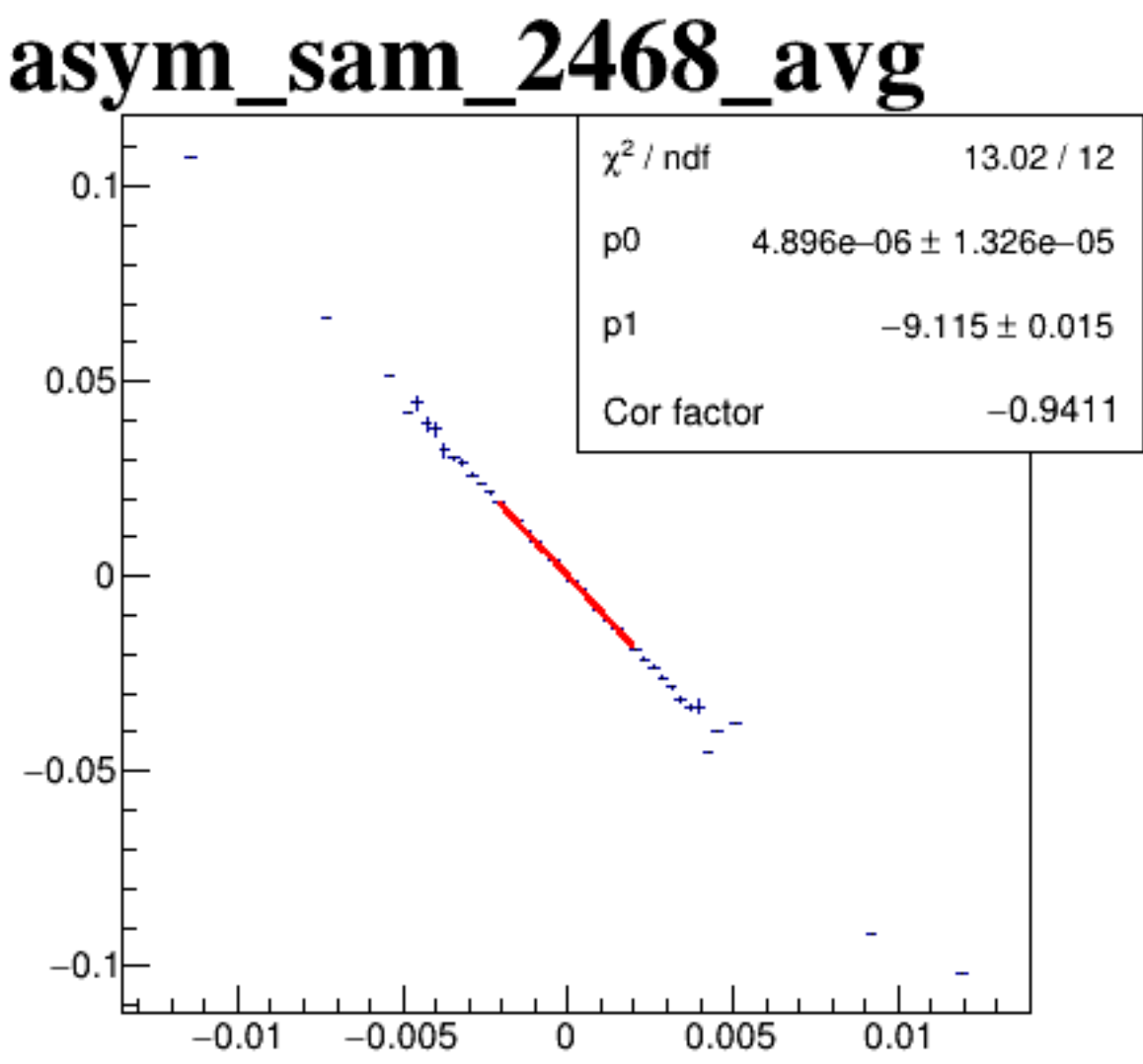
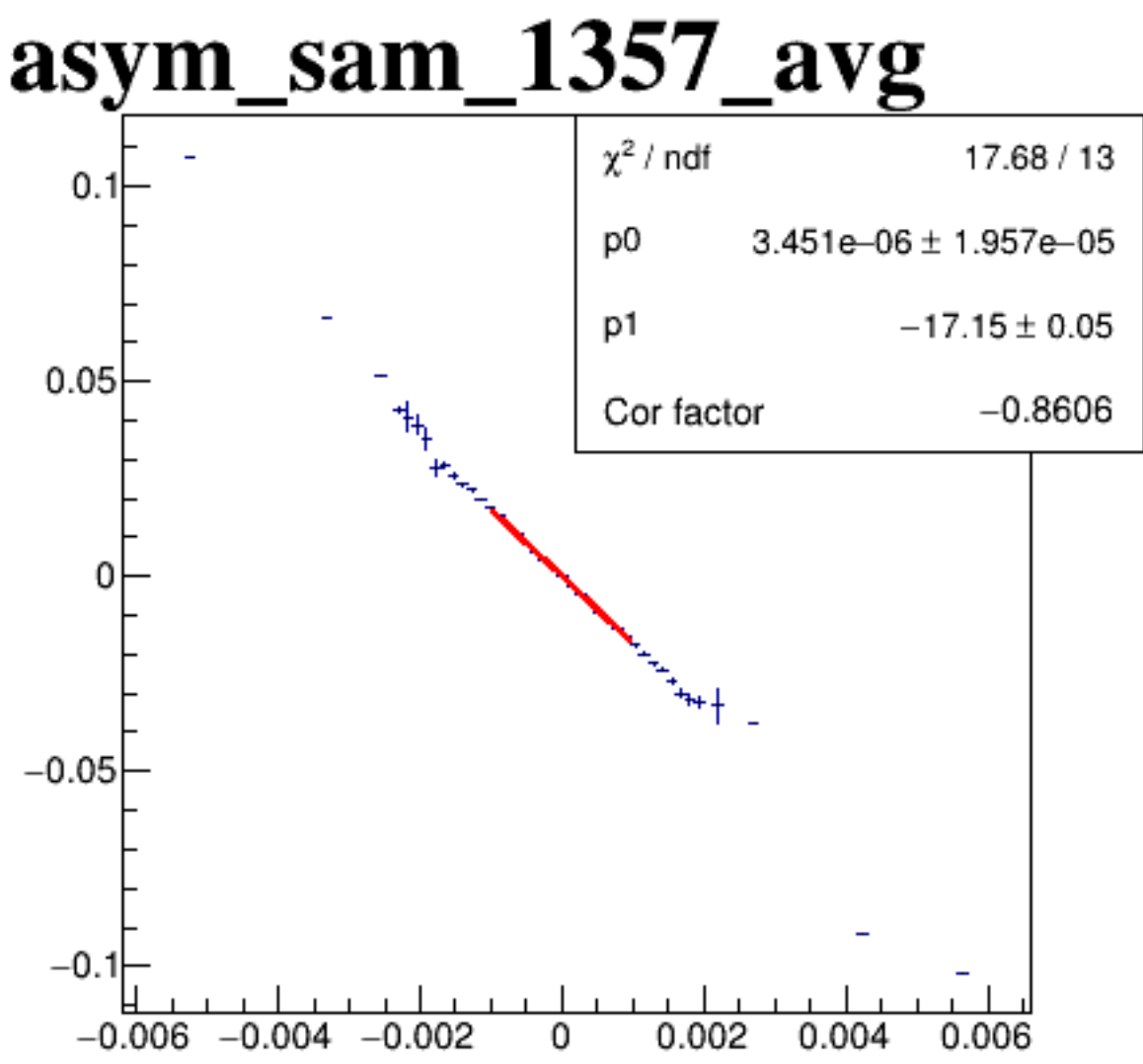
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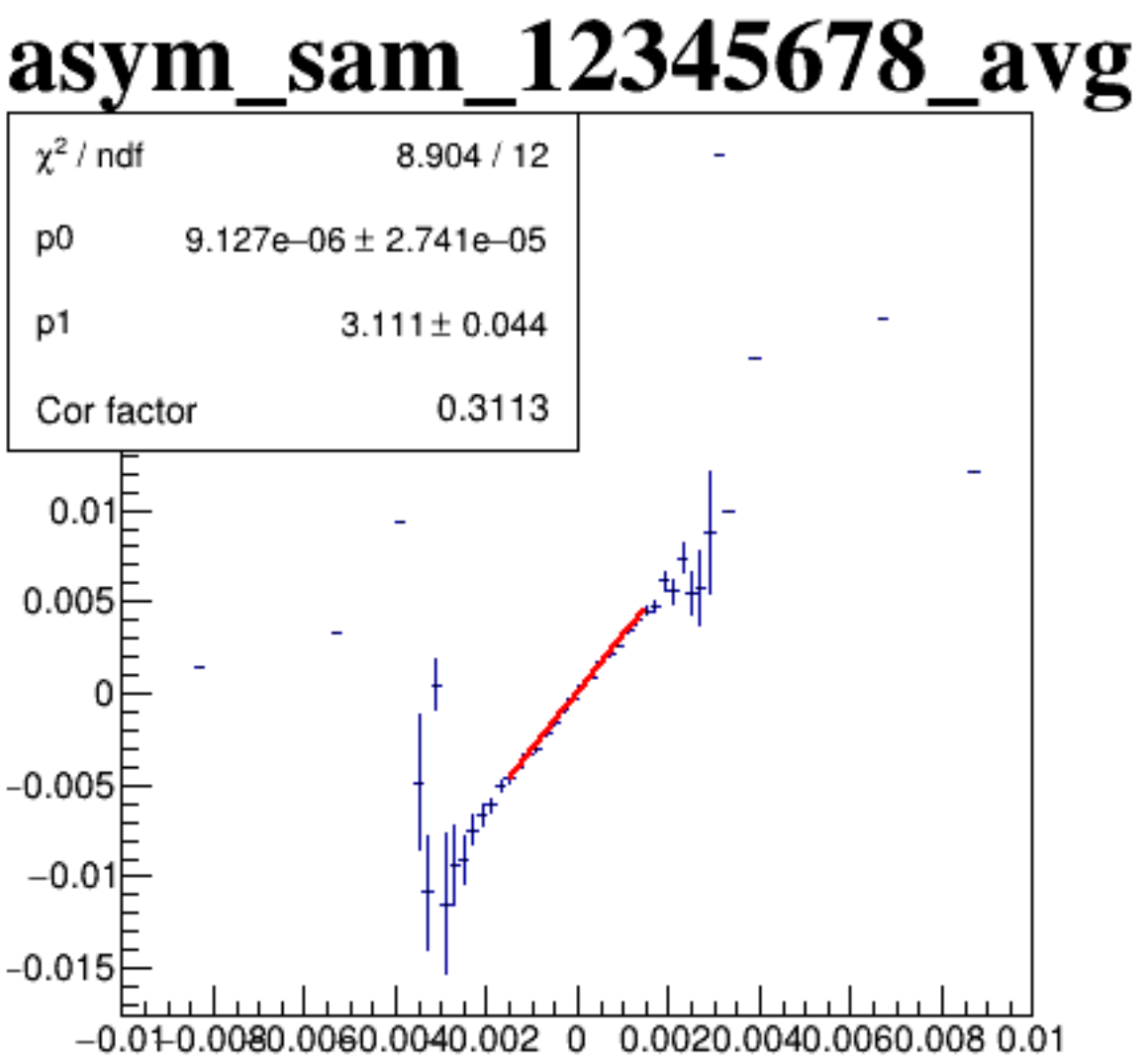
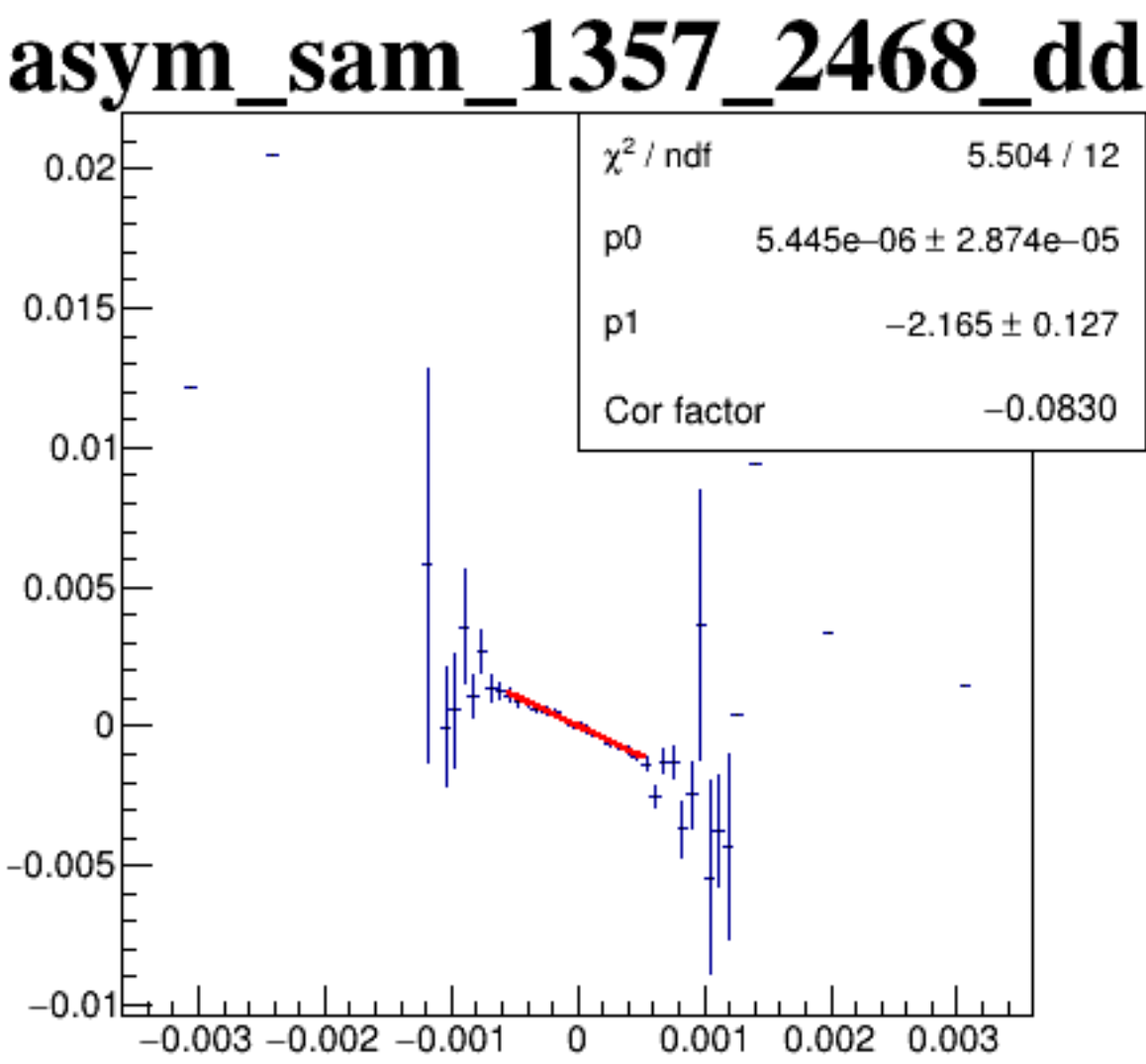
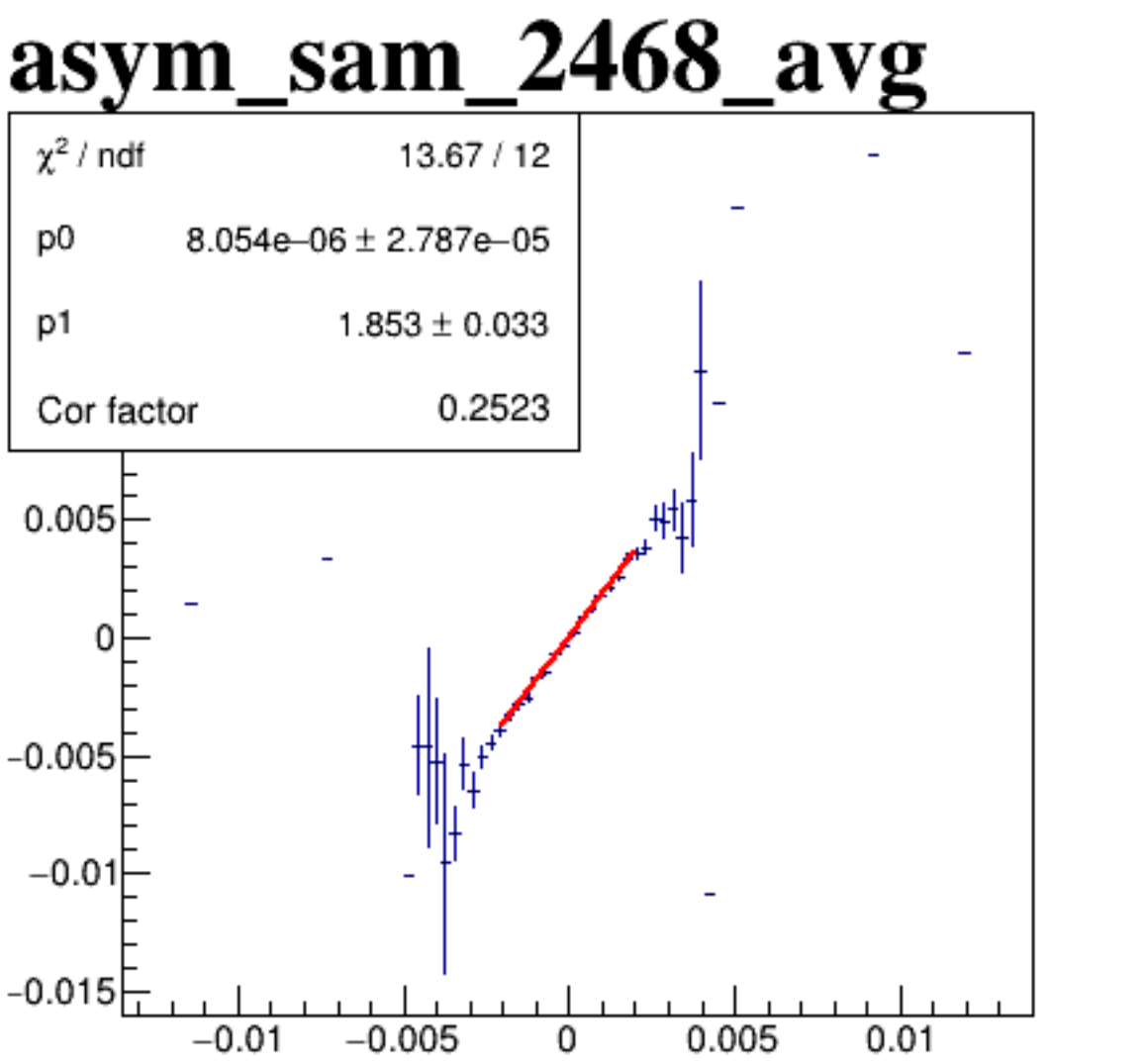
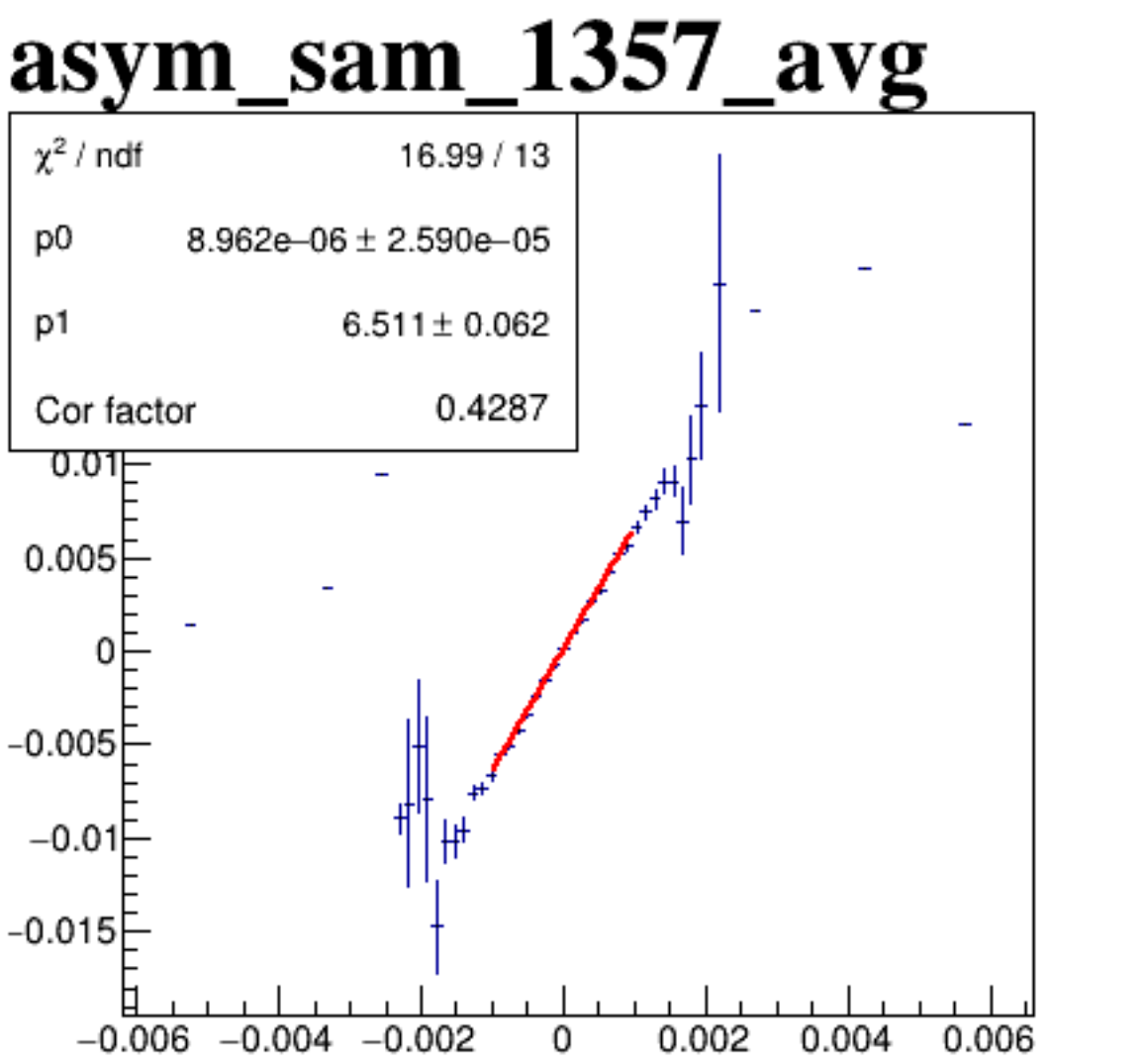
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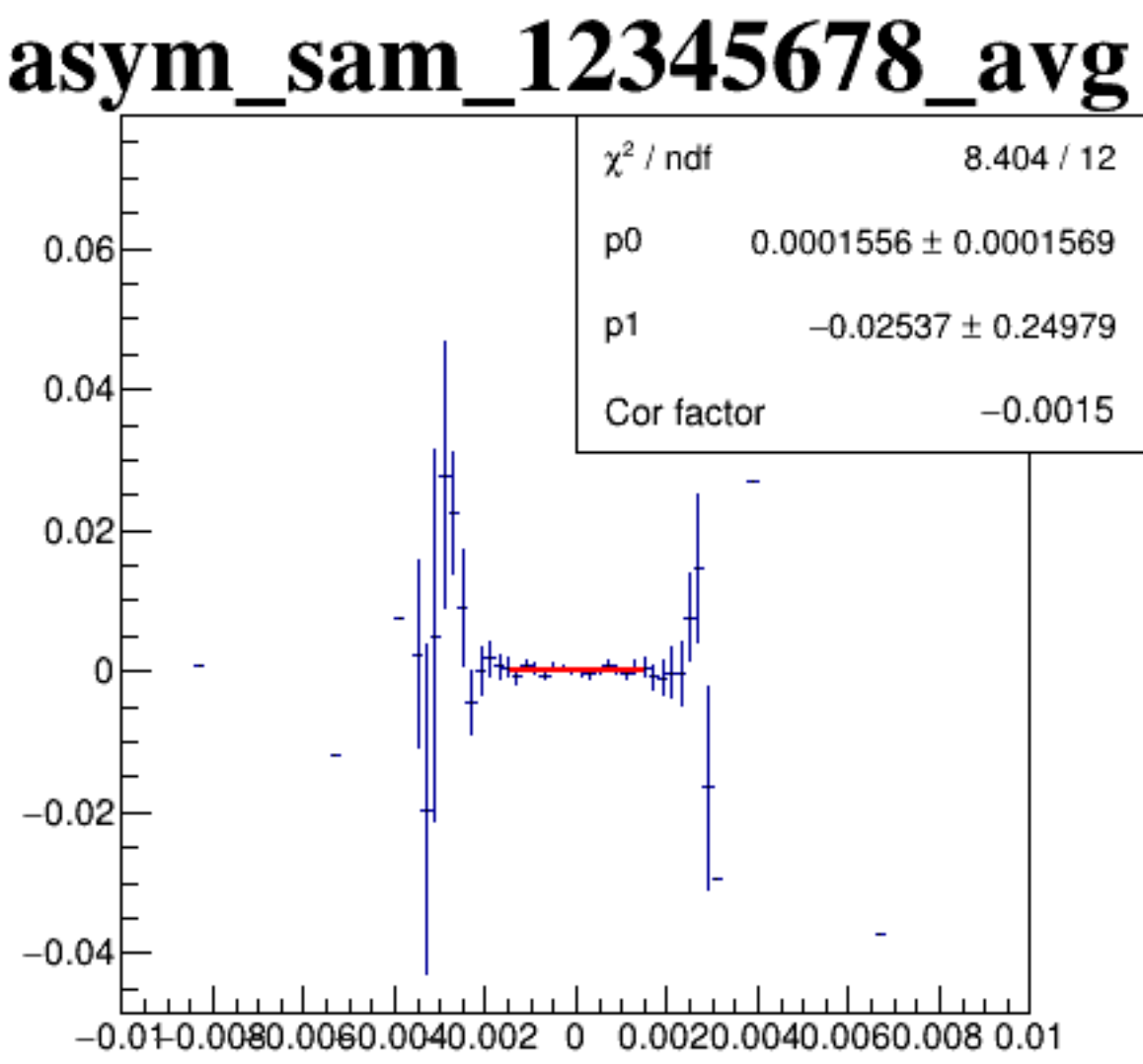
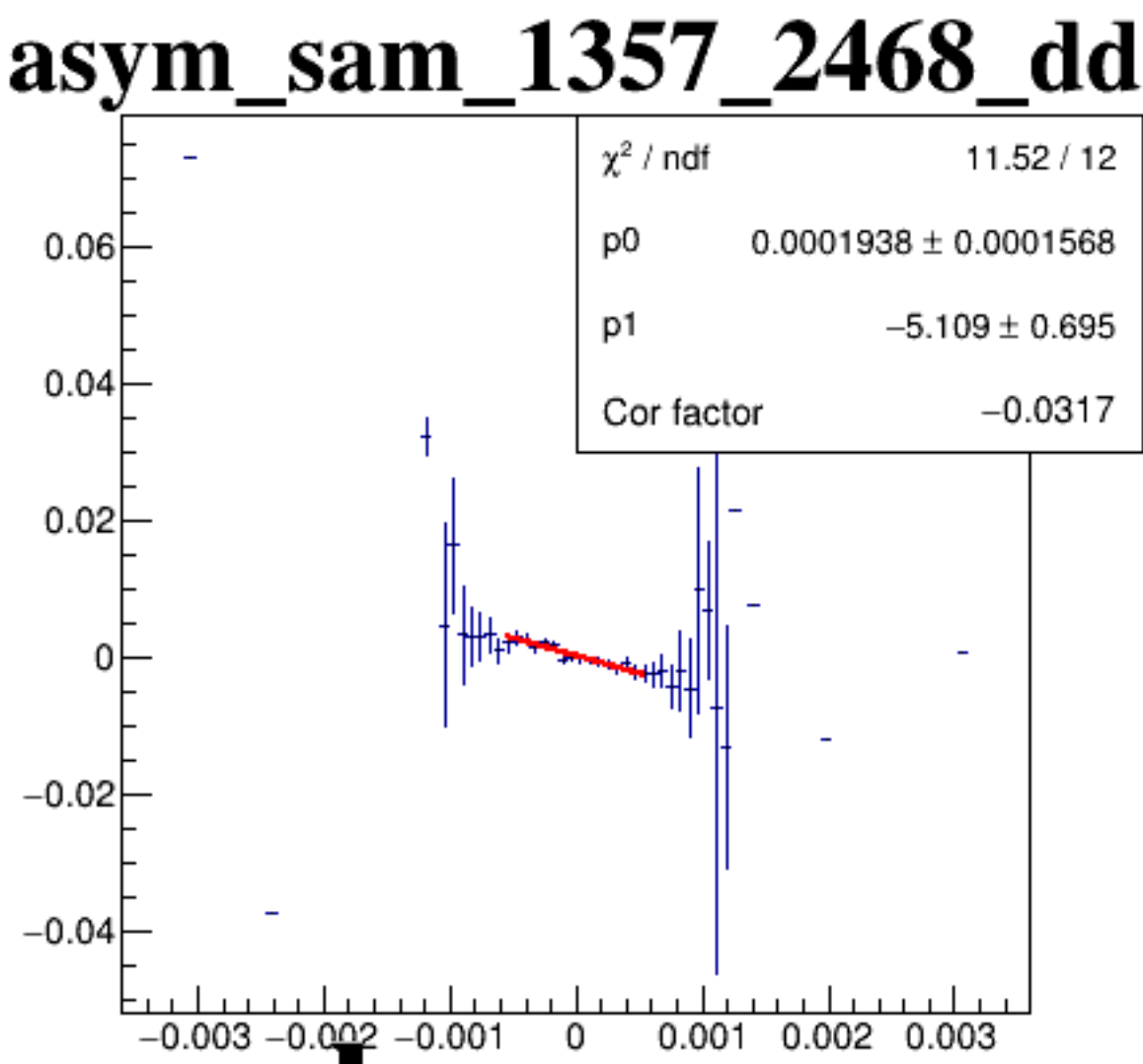
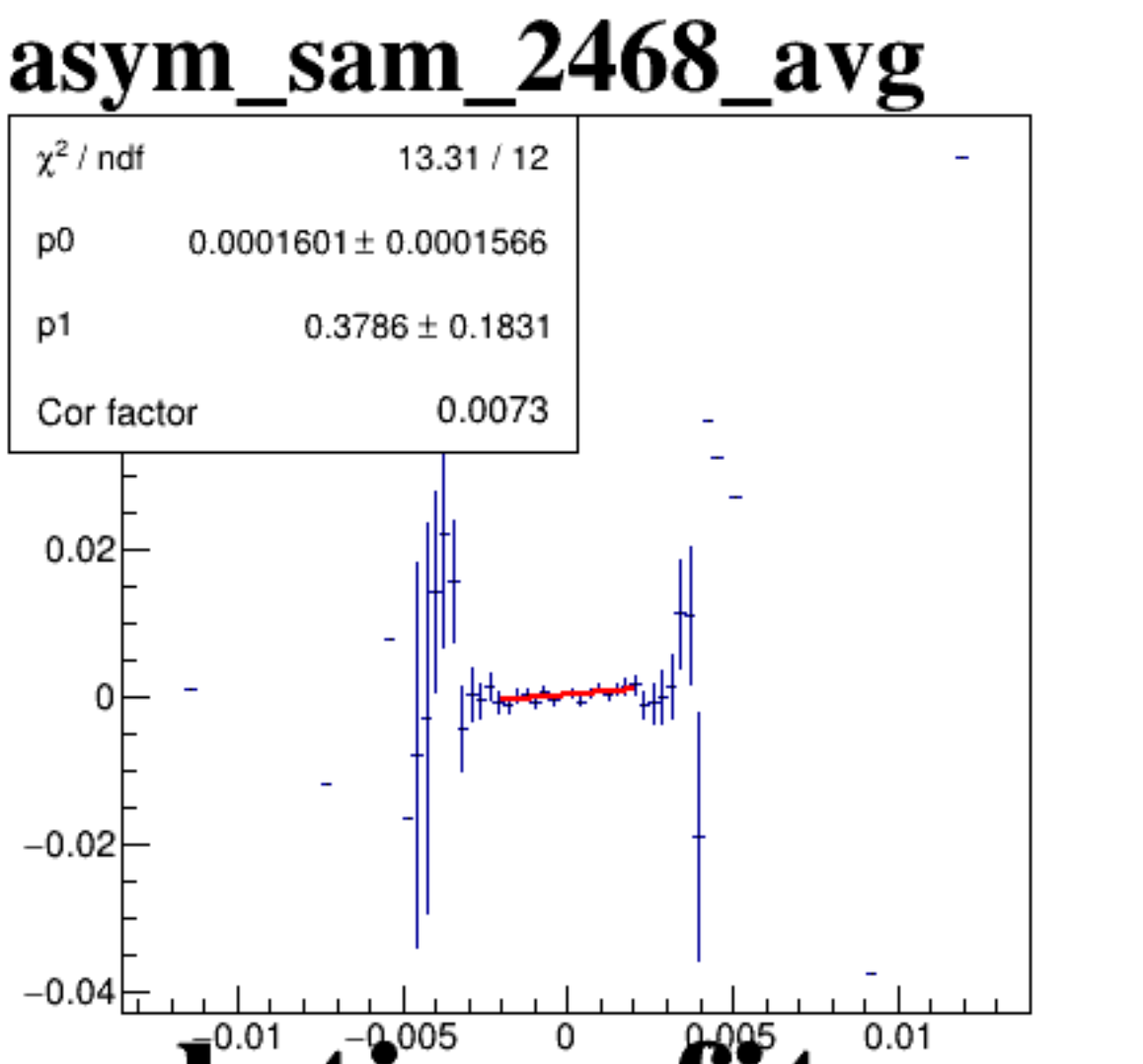
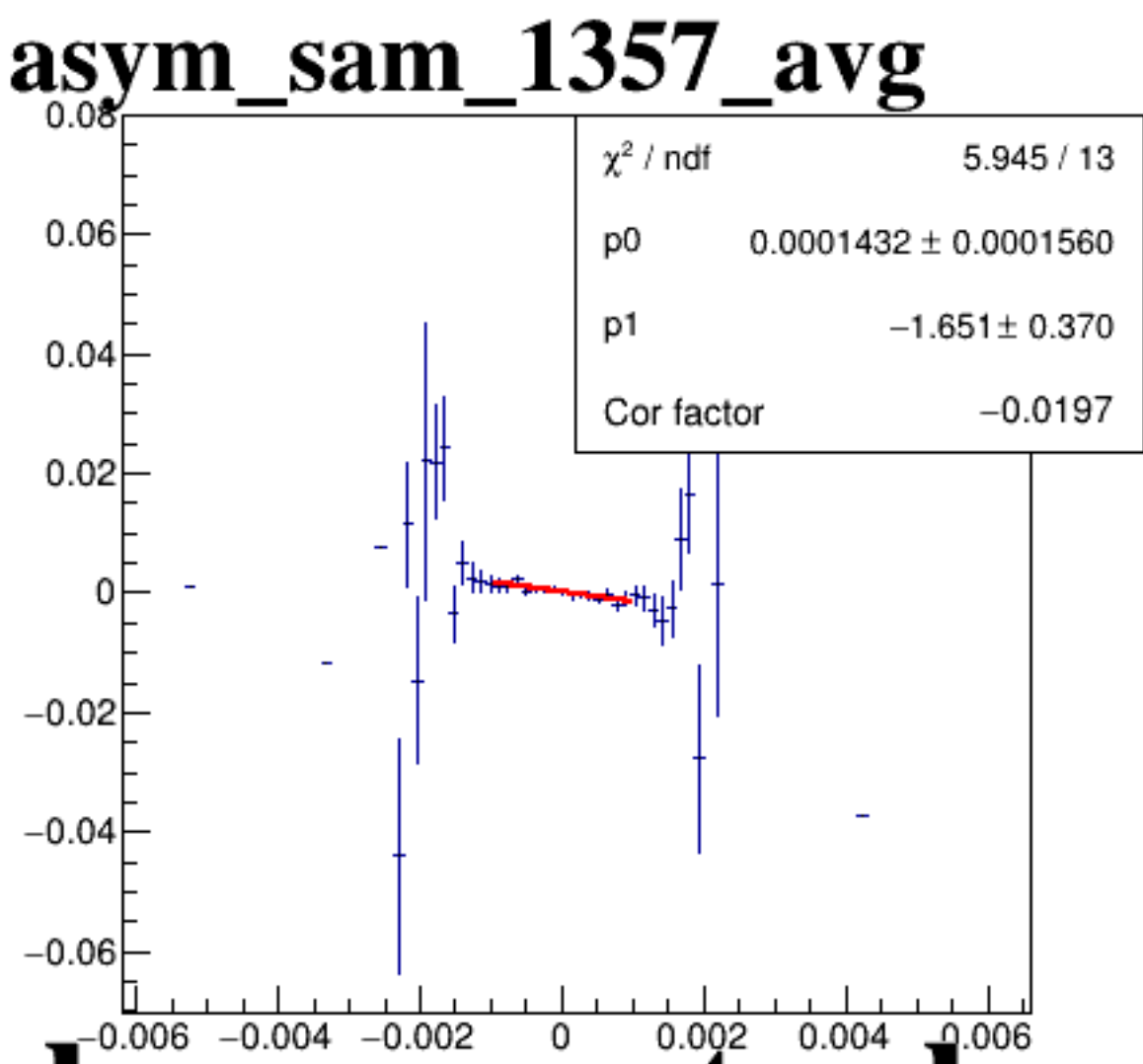
diff_bpm4eX



diff_bpm4eY



diff_bpm12X



A scatter plot showing the relationship between `mulc.asym_sam_12345678_avg` (x-axis) and `asym_sam_12345678_avg` (y-axis). The x-axis ranges from -0.01 to 0.01, and the y-axis ranges from -0.03 to 0.03. The data points form a dense, vertically elongated elliptical cloud centered at (0,0), indicating a positive correlation.

A scatter plot showing the relationship between `asym_sam_12345678_avg` (y-axis) and `mulc.asym_sam_12345678_avg` (x-axis). The y-axis ranges from -0.1 to 0.1, and the x-axis ranges from -0.01 to 0.01. The data points form a dense, elongated cluster along the diagonal line $y = x$, indicating a strong positive correlation between the two variables.

asym_sam_12345678_avg

A scatter plot showing a dense, vertically elongated cloud of points centered around (0,0). The x-axis is labeled 'mulc.asym_sam_12345678_avg' and ranges from -0.01 to 0.01. The y-axis ranges from -0.03 to 0.03.

A scatter plot showing the relationship between `asym_sam_12345678_avg` (y-axis) and `mulc.asym_sam_12345678_avg` (x-axis). The y-axis ranges from -0.15 to 0.15 with major ticks every 0.05. The x-axis ranges from -0.01 to 0.01 with major ticks every 0.002. The data points form a dense, vertically elongated elliptical cloud centered at (0,0), indicating a strong positive correlation.