

Status Report for Run 1-7c joint fit Update

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Overview

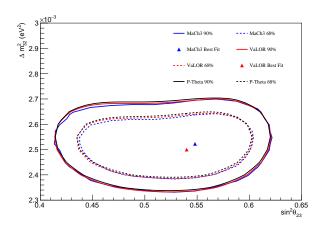
- ► All three analyses are updating to the full Run 1-7c dataset
- FHC POT: 7.48 $\times\,10^{20}$ RHC POT: 7.47 $\times\,10^{20}$
- Working to get the joint fit ready by ICHEP
- Asimov rates have been compared and found to be in good agreement
- Will show comparisons between the three groups asimov contours and first data results today
- All results are with reactor constraint



Asimov comparisons

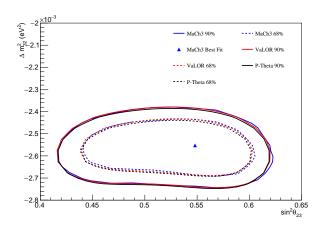


Comparison of Asimov Set 1 2D contours - NH



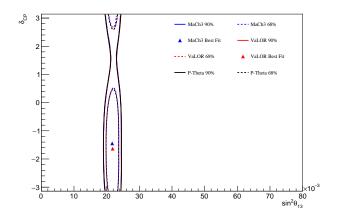


Comparison of Asimov Set 1 2D contours - IH



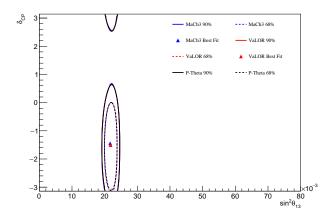


Comparison of Asimov Set 1 2D contours - NH



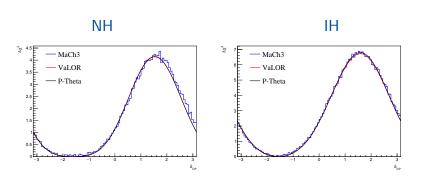


Comparison of Asimov Set 1 2D contours - IH





Comparison of Asimov Set 1 1D contours







Observed rates

Table: Number of data events in each of the SK samples for run 1–7c, compared to the prefit MC prediction (using oscillation parameters set 1).

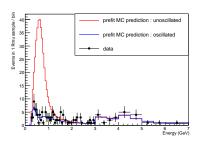
	FHC $1\mathrm{R}_{\mu}$	FHC $1R_{ m e}$	RHC $1\mathrm{R}_{\mu}$	RHC $1R_{ m e}$
Data	135	32	66	4
Prefit MC	136.21	28.75	64.40	6.01
Data/MC Ratio	1.00	1.11	1.02	0.67



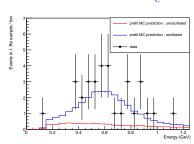


Observed spectra

u mode $1R_u$



ν mode $1R_{\rm e}$

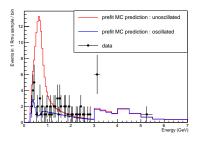




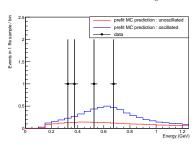


Observed spectra

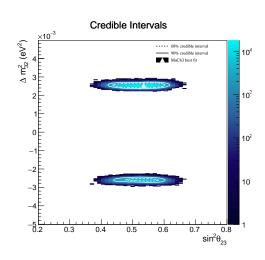
$\bar{\nu}$ mode $1R_{\mu}$



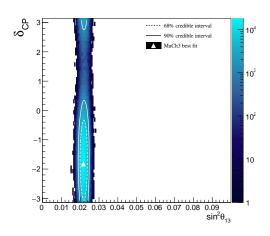
$\bar{\nu}$ mode $1R_{\rm e}$



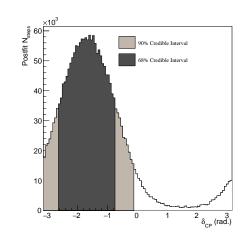










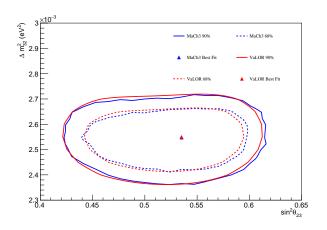




VaLOR-MaCh3 data fit comparison

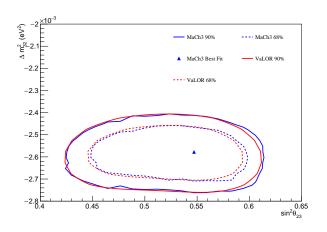


Comparison of Data 2D contours - NH



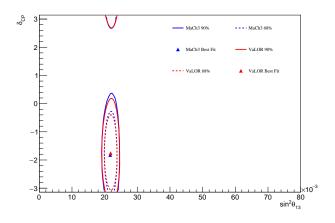


Comparison of Data 2D contours - IH



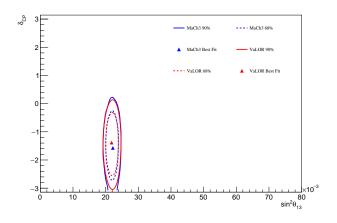


Comparison of Data 2D contours - NH



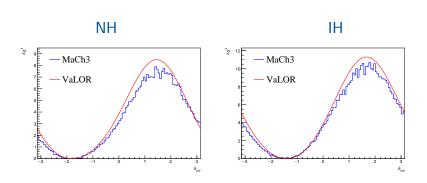


Comparison of Data 2D contours - IH





Comparison of Data 1D contours





- ▶ 2D and 1D Asimov contours compared between all three analyses
- Agreement between analyses is good
- Data results have been shown
- Similar differences between MaCh3 and VaLOR as in 1-7b seen
- Expected to be due to E_{rec} vs $E_{rec} \theta$ binning
- MaCh3 are working on implementing $E_{rec} \theta$ binning
- Still working on results without reactor constraint and updating TN