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A GPU-based framework for multivariate analysis in particle physics

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Outline

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- The GooStats framework
- Motivation: Borexino spectrum fit
- Package design
- Systematic uncertainty estimation
- Conclusion

GooStats hosted on GitHub



https://github.com/GooStats/GooStats.git

📮 GooStats	O Unwatch → 2					
<> Code	Issues 0	n Pull requests 0	Projects 0	🗉 Wiki	Insights	Settings
Releases	Tags					
Pre \$ v5.0.0-a •• 1	-release Ilpha-p2 - 82054b7 Verified	Simplify CM DingXuefeng released Assets 2 Source code (zip) Source code (tar.gz) Simplify CMakeLis Add simpleFit an	akeLists.t d this on Nov 25, 20 ts.txt d statAnalysis	t Xt 18 · 3 commi	ts to master si	nce this release

What and Why?



- A convenient GPU multivariate analysis framework
- Easy spectrum fitting

./fit toyMC.cfg out exposure=500



Figure 7. Left: screen shot of fit result summary. Right: produced figure in the file format of pdf.

GooStats: performance



• Speed up: around 200 times



Table 1. Comparison of fitting time between GooStats and original software used by the Borexino collaboration. T_{tot} : total execution time. N: the number of iterations taken to converge in MINUIT. T_{it} : average execution time per iteration. Speed up: $T_{\text{it}}(\text{CPU})/T_{\text{it}}(\text{GPU})$.

	CPU			GPU			
Туре	AMD Opteron(TM) Processor 6238			nVidia Tesla K20m			
Size	$T_{\rm tot}$ (s)	N	$T_{\rm it}~({\rm ms})$	$T_{\rm tot}$ (s)	N	$T_{\rm it}~({\rm ms})$	speed up
400	27.6	1128	24.4	0.181	1346	0.135	181
350	29.4	1331	22.1	0.156	1294	0.121	183
300	22.5	1239	18.2	0.243	1995	0.122	149

Motivation: Borexino spectrum fit

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• Motivation: spectral-fitting tool for Borexino analysis



Analytical Response Function



$$M: f(E) \mapsto g(\text{charge}) = \int_0^{E_{\text{end}}} dE \cdot f(E) \cdot \text{RPF} \left[\text{charge}; \mu(E), \text{var}(\mu)\right]$$

- Analytical shape of spectrum of mono-energetic events
 - Momentum based approximation
 - Match the average (energy scale + non-linearity model)
 - Match the variance (energy resolution model)
 - ... (-> simplified)
 - More: "Mask", "pile-up" etc...
- We can simplify because
 - Borexino response is simple: small FV in center, low energies => no irregular tail
 - We are not sensitive.. => small systematics
 - Fit full MC to get the bias introduced in simplification

Multivariate Likelihood (Davini method)



Scaling factor introduced to remove bias.

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Package Design

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Middle Layer between GooFit and User module



Parameter Synchronization



- Synchronize fit parameters in joint analysis
- Supported by tree type internal data structure



TTree Output



- All fit parameters saved in TTree output automatically
- More quantities can be added as lambda functions in the output builder

```
root [1] .ls
TFile**
                test_tree.root
TFile*
                test_tree.root
 KEY: TTree
                fit_result;1
                                Fit result of GooStats
root [2] fit_result->Show(0)
====> EVENT:0
 default.NReactor = 3.02987
 default.NReactor err = 0.0159146
 default.Nbkg
                 = 1.02405
 default.Nbkg_err = 0.0162459
 default.U235
                 = 0.5
 default.U235_err = 0
 default.U238
                 = 0.2
 default.U238_err = 0
 default.Pu239
                 = 0.1
 default.Pu239_err = 0
 default.U241
                 = 0.2
 default.U241_err = 0
                 = 1300
 default.LY
 default.LY_err = 0
                 = 2.78788
 default.qc1
 default.qc1_err = 0
                 = -0.528003
 default.qc2
 default.qc2_err = 0
 default.v1
                 = 0.3
 default.v1 err = 0
 default.vT
                 = 5
 default.vT_err = 0
 default.Reactor_dEvis = 1078.28
 default.Reactor_dEvis_err = 13.2291
 chi2
                 = 390.448
 NDF
                 = 397
 likelihood
                 = 1883.96
```

Statistical analysis modules



- Adding tasks of statistical analysis by registering new modules inheriting abstract classes
- A few pre-installedt modules: ScanPar, DiscoveryTest..

```
PrepareData *data = new PrepareData();
SimpleFit *fit = new SimpleFit();
DiscoveryTest *discovery = new DiscoveryTest();
ana->registerModule(inputManager);
ana->registerModule(data);
ana->registerModule(fit);
ana->registerModule(fit);
ana->registerModule(discovery);
ana->registerModule(outManager);
```

Physics modules

Solar Neutrinos

JUNO style





- create an ensemble of models according to experiment precision
 - Fit the data with varying models through their coordinates
 - Generate pseudo-experiments with varying models and take the width of distribution of best-fit

Method 1: fit with varying models



- Full MC during each iteration of the fit, vary kb / absorption length spectrum etc. re-simulate and produce new pdf on the fly —> when one day computer is fast enough
 - ~200, 000 CPU x years per fit
- Semi-analytical: analytical non-linearity model + response Matrix
 - ~30 minutes per fit
- Full analytical
 - ~2 hours per fit

Problem with scaling response matrix

- Systematic uncertainties of LY can be included by scaling the response matrix
- This could be dangerous if the interested parameters is correlated with the resolution parameters
- When LY is scaled, the resolution should also be changed

 $v (^7 Be)$



correlation plot from full analytical fit

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48.5 (-2.1 +2.3)

 $inj 48.00 \pm 0.07$

Correlation with NL/resolution



• Using full analytical response function, we can see the correlation with detector responses

[1] M. Agostini et al., "First Simultaneous Precision Spectroscopy of \$pp\$, \$^7\$Be, and \$pep\$ Solar Neutrinos with Borexino Phase-II," pp. 1–8, Jul. 2017.

correlation with NL





correlation with res



Method 2: Monte Carlo method



pseudo-experiment spectra without distortion —> statistical sensitivity

pseudo-experiment spectra with distortion —> statistical systematic uncertainty

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Conclusion



- GooStats, a GPU based multivariate analysis framework is introduced.
- Statistical analysis module is easy to be implemented as needed. TTree/figure output provided.
- Full analytical response function has advantage that it treated the NL and resolution in a coherent way when evaluating the systematic uncertainties
- The systematic uncertainties can also be evaluated using Monte Carlo method by looking at change of width of best fit with/without distortion