



Meson Spectroscopy at COMPASS

Methods for Amplitude Analysis

Sebastian Neubert

Technische Universität München

Hirschegg, January 2011





The COMPASS Experiment

Diffractive Pion Dissociation

Amplitude Analysis Formalism

PWA Model Selection

- Mass Independent Fit

- Bayesian Model Evaluation

- Waveset Exploration

Mass Dependent Fit

ROOTPWA Analysis Toolkit



The COMPASS Experiment

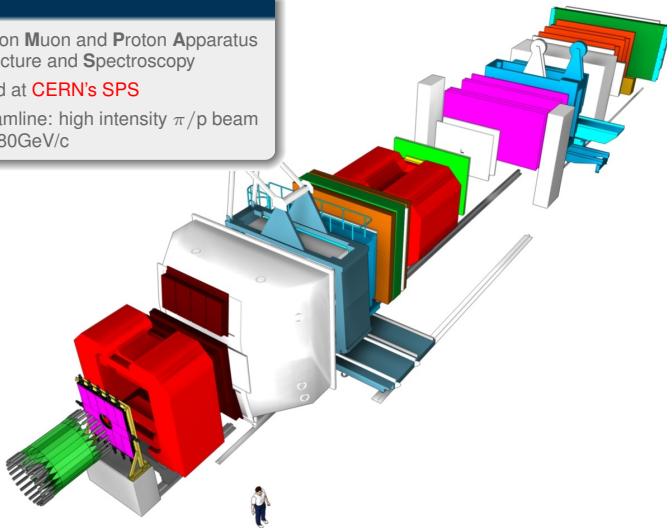
Searching for Gluonic Contributions to the Meson Spectrum



Technische Universität München

Overview

- **CO**mmun **MU**on and **P**roton **A**pparatus for **S**tructure and **S**pectroscopy
- Located at **CERN's SPS**
- M2-beamline: high intensity π/p beam up to 280GeV/c





The COMPASS Experiment

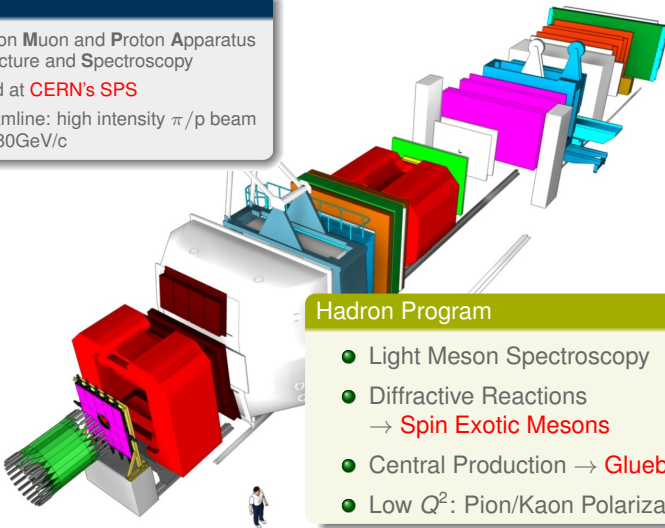
Searching for Gluonic Contributions to the Meson Spectrum



Technische Universität München

Overview

- **CO**mmun **MU**on and **P**roton **A**pparatus for **S**tructure and **S**pectroscopy
- Located at **CERN's SPS**
- M2-beamline: high intensity π/p beam up to 280GeV/c



Hadron Program

- Light Meson Spectroscopy
- Diffractive Reactions
→ **Spin Exotic Mesons**
- Central Production → **Glueballs**
- Low Q^2 : Pion/Kaon Polarizabilities

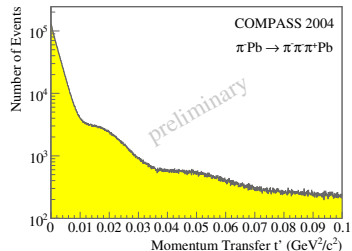
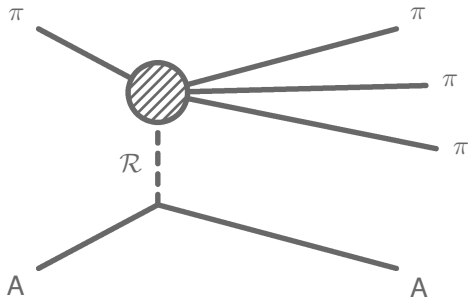


Diffraction Pion Dissociation

Example: 3 Pion Final State



Technische Universität München



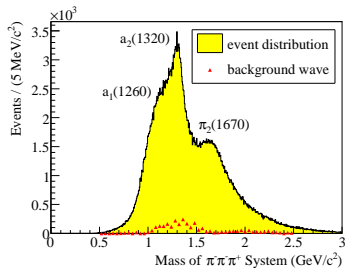
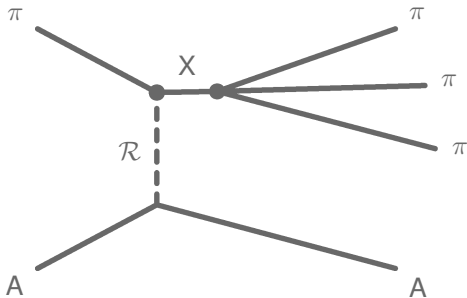


Diffraction Pion Dissociation

Example: 3 Pion Final State



Technische Universität München



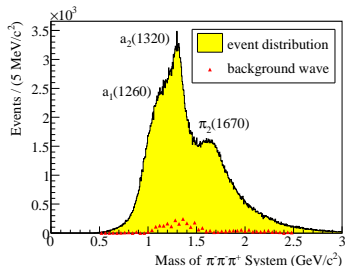
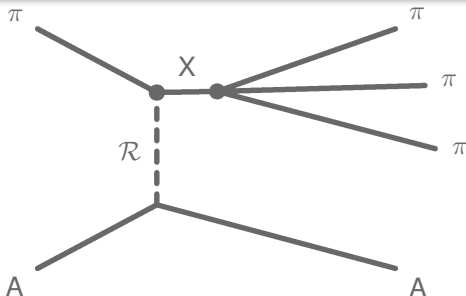


Diffractive Pion Dissociation

Example: 3 Pion Final State



Technische Universität München



X: $J^{PC} M^{\epsilon}$ decay amplitude in *reflectivity* base

Implement parity conservation:

$$\psi_{JM}^{\epsilon} = c(M) \left[\psi_{JM}(\tau) - \epsilon P(-1)^{J-M} \psi_{J(-M)}(\tau) \right]$$

$$\epsilon = \pm 1 \quad M \in [0..J] \quad c(M > 0) = \frac{1}{\sqrt{2}} \quad c(M = 0) = \frac{1}{2}$$



PWA Formalism Overview

2Stage Isobar-Model Fit



Technische Universität München

STEP 1: Mass-Independent PWA

- Fit angular distributions + isobar systems in independent mass bins

$$\mathcal{I}(\tau, m) = \sum_{\epsilon=\pm 1} \sum_{r=1}^{N_r} \left| \sum_{\alpha} T_{\alpha r}^{\epsilon} \psi_{\alpha}^{\epsilon}(\tau, m) \right|^2$$

- Production amplitude
- Decay amplitude



PWA Formalism Overview

2Stage Isobar-Model Fit



STEP 1: Mass-Independent PWA

- Fit angular distributions + isobar systems in independent mass bins

$$\mathcal{I}(\tau, m) = \sum_{\epsilon=\pm 1} \sum_{r=1}^{N_r} \left| \sum_{\alpha} T_{\alpha r}^{\epsilon} \psi_{\alpha}^{\epsilon}(\tau, m) \right|^2$$

- Production amplitude
- Decay amplitude

STEP 2: Mass-Dependent χ^2 fit \rightarrow Extract Resonance Parameters

- Parameterization of spin-density matrix elements $\sum_r T_{ir}^{\epsilon} T_{jr}^{\epsilon*}(m_x)$
- Takes into account interference terms
- Coherent background for some waves



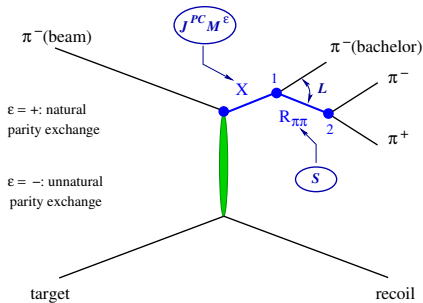
Decay Parameterization: The Isobar Model



Technische Universität München

Chain of successive 2-body decays

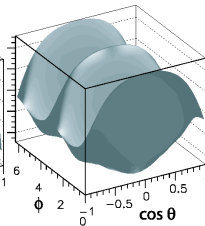
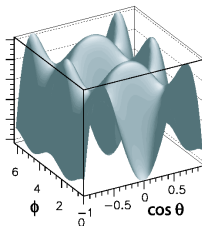
Model n-body decay by a chain of successive 2-body decays:



Example angular distributions:

$X(2^-) \rightarrow f_2(1275)\pi$

$f_2(1275) \rightarrow \pi\pi$



- For fixed n-body mass m there are $3n - 4$ parameters (angles, intermediate state masses)
- Parameterization of isobar subsystems

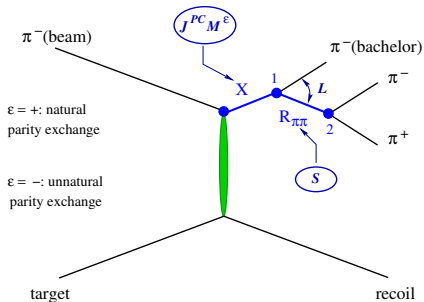


Decay Parameterization: The Isobar Model



Chain of successive 2-body decays

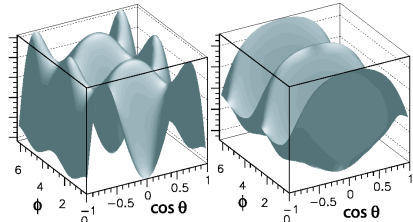
Model n-body decay by a chain of successive 2-body decays:



Example angular distributions:

$X(2^+) \rightarrow f_2(1275)\pi$

$f_2(1275) \rightarrow \pi\pi$



Known shortcomings

- For fixed n-body mass m there are many possible partial waves (angles, intermediate state masses)
- Parameterization of isobar subsystems
- Unitarity violation
- Rescattering effects
- \rightarrow Potential for improvement
- Input from theory needed (see e. g. talk by B. Kubis)



Mass Independent Amplitude Fit

Intensity distribution parameterization



Technische Universität München

Intensity distribution \mathcal{I} as a function of decay-kinematic variables τ :

$$\mathcal{I}(\tau) = \sum_{\epsilon=\pm 1} \sum_r \left| \sum_{\alpha \in M} T_{\alpha r}^{\epsilon} \bar{\psi}_{\alpha}^{\epsilon}(\tau) \right|^2$$

- Finite waveset M
- Production amplitude
- Decay amplitude



Mass Independent Amplitude Fit

Intensity distribution parameterization



Technische Universität München

Intensity distribution \mathcal{I} as a function of decay-kinematic variables τ :

$$\mathcal{I}(\tau) = \sum_{\epsilon=\pm 1} \sum_r \left| \sum_{\alpha \in M} T_{\alpha r}^{\epsilon} \bar{\psi}_{\alpha}^{\epsilon}(\tau) \right|^2$$

- Finite waveset M
- Production amplitude
- Decay amplitude

The likelihood \mathcal{L} to observe (a specific set of) N events in a bin with finite acceptance $\eta(\tau)$ (assuming a model M , parameters T_{ir}^{ϵ}) is:

$$P(\text{Data} | T_{ir}, M) = \mathcal{L} = \left[\frac{\bar{N}^N}{N!} e^{-\bar{N}} \right] \prod_i \frac{\mathcal{I}(\tau_i) \eta(\tau_i) f(\tau_i)}{\underbrace{\int \mathcal{I}(\tau) \eta(\tau) d\rho(\tau)}_{=\bar{N}}} \quad \text{with} \quad d\rho(\tau) = f(\tau) d\tau$$



Mass Independent Amplitude Fit

Definition of LogLikelihood Function



Technische Universität München

$$\mathcal{L} = \left[\frac{\bar{N}^N}{N!} e^{-\bar{N}} \right] \prod_i^N \frac{\mathcal{I}(\tau_i)}{\bar{N}} \eta(\tau_i) f(\tau_i) = \frac{1}{N!} \prod_i^N \mathcal{I}(\tau_i) \cdot \prod_i^N \eta(\tau_i) f(\tau_i) \cdot e^{-\bar{N}}$$



Mass Independent Amplitude Fit

Definition of LogLikelihood Function



Technische Universität München

$$\mathcal{L} = \left[\frac{\bar{N}^N}{N!} e^{-\bar{N}} \right] \prod_i^N \frac{\mathcal{I}(\tau_i)}{\bar{N}} \eta(\tau_i) f(\tau_i) = \frac{1}{N!} \prod_i^N \mathcal{I}(\tau_i) \cdot \prod_i^N \eta(\tau_i) f(\tau_i) \cdot e^{-\bar{N}}$$

Taking the logarithm leads to and inserting for \bar{N}

$$\ln \mathcal{L} = -N \ln N + \sum_i^N \eta(\tau_i) f(\tau_i) + \sum_i^N \ln \mathcal{I}(\tau_i) - \int \mathcal{I}(\tau) \eta(\tau) d\rho(\tau)$$



Mass Independent Amplitude Fit

Definition of LogLikelihood Function



$$\mathcal{L} = \left[\frac{\bar{N}^N}{N!} e^{-\bar{N}} \right] \prod_i^N \frac{\mathcal{I}(\tau_i)}{\bar{N}} \eta(\tau_i) f(\tau_i) = \frac{1}{N!} \prod_i^N \mathcal{I}(\tau_i) \cdot \prod_i^N \eta(\tau_i) f(\tau_i) \cdot e^{-\bar{N}}$$

Taking the logarithm leads to and inserting for \bar{N}

$$\ln \mathcal{L} = -N \ln N + \sum_i^N \eta(\tau_i) f(\tau_i) + \sum_i^N \ln \mathcal{I}(\tau_i) - \int \mathcal{I}(\tau) \eta(\tau) d\rho(\tau)$$

drop $(-N \ln N + \sum_i^N \eta(\tau_i) f(\tau_i))$ and insert intensity parameterization

$$\ln \mathcal{L} = \sum_{n=1}^{N_{\text{events}}} \ln \left[\sum_{\epsilon, r} \sum_{\alpha, \beta \in M} T_{\alpha r}^{\epsilon} T_{\beta r}^{\epsilon*} \bar{\psi}_{\alpha}^{\epsilon}(\tau_n) \bar{\psi}_{\beta}^{\epsilon}(\tau_n)^* \right] - \sum_{\epsilon, r} \sum_{\alpha, \beta \in M} T_{\alpha r}^{\epsilon} T_{\beta r}^{\epsilon*} I A_{\alpha\beta}^{\epsilon}$$



Mass Independent Amplitude Fit

Definition of LogLikelihood Function



$$\mathcal{L} = \left[\frac{\bar{N}^N}{N!} e^{-\bar{N}} \right] \prod_i^N \frac{\mathcal{I}(\tau_i)}{\bar{N}} \eta(\tau_i) f(\tau_i) = \frac{1}{N!} \prod_i^N \mathcal{I}(\tau_i) \cdot \prod_i^N \eta(\tau_i) f(\tau_i) \cdot e^{-\bar{N}}$$

Taking the logarithm leads to and inserting for \bar{N}

$$\ln \mathcal{L} = -N \ln N + \sum_i^N \eta(\tau_i) f(\tau_i) + \sum_i^N \ln \mathcal{I}(\tau_i) - \int \mathcal{I}(\tau) \eta(\tau) d\rho(\tau)$$

drop $(-N \ln N + \sum_i^N \eta(\tau_i) f(\tau_i))$ and insert intensity parameterization

$$\ln \mathcal{L} = \sum_{n=1}^{N_{\text{events}}} \ln \left[\sum_{\epsilon, r} \sum_{\alpha, \beta \in M} T_{\alpha r}^{\epsilon} T_{\beta r}^{\epsilon*} \bar{\psi}_{\alpha}^{\epsilon}(\tau_n) \bar{\psi}_{\beta}^{\epsilon}(\tau_n)^* \right] - \sum_{\epsilon, r} \sum_{\alpha, \beta \in M} T_{\alpha r}^{\epsilon} T_{\beta r}^{\epsilon*} IA_{\alpha\beta}^{\epsilon}$$

With acceptance-corrected phase space integral

$$IA_{\alpha\beta}^{\epsilon} = \int \bar{\psi}_{\alpha}^{\epsilon}(\tau) \bar{\psi}_{\beta}^{\epsilon}(\tau)^* \eta(\tau) d\tau$$



Which waves to include into the waveset?



Which waves to include into the waveset?

Avoid overfitting



Which waves to include into the waveset?

Avoid overfitting

→ Data driven method



How to Measure the Goodness of a Model



Marginal Likelihood Definition

Bayes' Theorem (for the Model Probability after Observation)

$$P(M_k|\text{Data}) = \frac{P(\text{Data}|M_k)P(M_k)}{\sum_{k'} P(\text{Data}|M_{k'})P(M_{k'})}$$

with model-priors $P(M_k)$ $\sum_{k'} P(M_{k'}) = 1$



Bayes' Theorem (for the Model Probability after Observation)

$$P(M_k | \text{Data}) = \frac{P(\text{Data} | M_k) P(M_k)}{\sum_{k'} P(\text{Data} | M_{k'}) P(M_{k'})}$$

with model-priors $P(M_k)$ $\sum_{k'} P(M_{k'}) = 1$

Marginal Likelihood or Evidence

$$P(\text{Data} | M_k) = \int \underbrace{P(\text{Data} | T^k, M_k)}_{\mathcal{L}} \underbrace{P(T^k | M_k)}_{\text{Prior}} dT^k$$

$P(T^k | M_k)$ contains any pre-knowledge on the model-parameters T

- Marginalization ($= \int dT$) is not trivial in high-dimensional spaces
- Numerically stable is only the **Log**Likelihood



The Occam Factor Approximation

David J. C. MacKay, 2003 "Information Theory, Inference and Learning Algorithms"



Technische Universität München

$$P(\text{Data}|M_k) = \int \underbrace{P(\text{Data}|T^k, M_k)}_{\mathcal{L}} \underbrace{P(T^k|M_k)}_{\text{Prior}} dT^k$$



The Occam Factor Approximation

David J. C. MacKay, 2003 "Information Theory, Inference and Learning Algorithms"



Technische Universität München

$$P(\text{Data}|M_k) = \int \underbrace{P(\text{Data}|T^k, M_k)}_{\mathcal{L}} \underbrace{P(T^k|M_k)}_{\text{Prior}} dT^k$$

Approximate with Laplace's method:

$$P(\text{Data}|M_k) \approx P(\text{Data}|T_{\text{ML}}^k, M_k) \cdot \underbrace{P(T_{\text{ML}}^k|M_k) \cdot \sqrt{(2\pi)^d |\mathbf{C}_{T|\text{Data}}|}}_{\text{Occam factor}}$$



The Occam Factor Approximation

David J. C. MacKay, 2003 "Information Theory, Inference and Learning Algorithms"



Technische Universität München

$$P(\text{Data}|M_k) = \int \underbrace{P(\text{Data}|T^k, M_k)}_{\mathcal{L}} \underbrace{P(T^k|M_k)}_{\text{Prior}} dT^k$$

Approximate with Laplace's method:

$$P(\text{Data}|M_k) \approx P(\text{Data}|T_{\text{ML}}^k, M_k) \cdot \underbrace{P(T_{\text{ML}}^k|M_k) \cdot \sqrt{(2\pi)^d |\mathbf{C}_{T|\text{Data}}|}}_{\text{Occam factor}}$$

- $P(\text{Data}|T_{\text{ML}}^k, M_k)$ LogLikelihood at maximum likelihood solution T_{ML}
- $|\mathbf{C}_{T|\text{Data}}|$ determinant of covariance matrix
- Dimension of parameter space: d



The Occam Factor Approximation

David J. C. MacKay, 2003 "Information Theory, Inference and Learning Algorithms"



Technische Universität München

$$P(\text{Data}|M_k) = \int \underbrace{P(\text{Data}|T^k, M_k)}_{\mathcal{L}} \underbrace{P(T^k|M_k)}_{\text{Prior}} dT^k$$

Approximate with Laplace's method:

$$P(\text{Data}|M_k) \approx P(\text{Data}|T_{\text{ML}}^k, M_k) \cdot \underbrace{P(T_{\text{ML}}^k|M_k) \cdot \sqrt{(2\pi)^d |\mathbf{C}_{T|\text{Data}}|}}_{\text{Occam factor}}$$

- $P(\text{Data}|T_{\text{ML}}^k, M_k)$ LogLikelihood at maximum likelihood solution T_{ML}
- $|\mathbf{C}_{T|\text{Data}}|$ determinant of covariance matrix
- Dimension of parameter space: d

Logarithmic evidence:

$$\ln P(\text{Data}|M_k) \approx \ln P(\text{Data}|T_{\text{ML}}^k, M_k) + \ln P(T^k|M_k) + \ln \sqrt{(2\pi)^d |\mathbf{C}_{T|D}|}$$



Log-Evidence

$$\ln P(\text{Data}|M_k) \approx \ln \mathcal{L}_{ML} + \ln \sqrt{(2\pi)^d |\mathbf{C}_{T|\text{Data}}|} - \ln V_T^k + \sum_{i \in M} \ln S_i$$

where V_T^k is the (prior) volume of parameter space

- Models (=wavesets) compared through the Bayes-Factor

$$B_{12} = \frac{P(\text{Data}|M_1)}{P(\text{Data}|M_2)}$$

- Interpretation according to Kass&Raftery:

$2 \ln B_{12}$	B_{12}	Evidence
0 to 2	1 to 3	Not worth mentioning
2 to 6	3 to 20	Positive
6 to 10	20 to 150	Strong
> 10	> 150	Very strong

Kass, Raftery, *Bayes Factors*, J. Am. Stat. Assoc. 90 (1995) 773

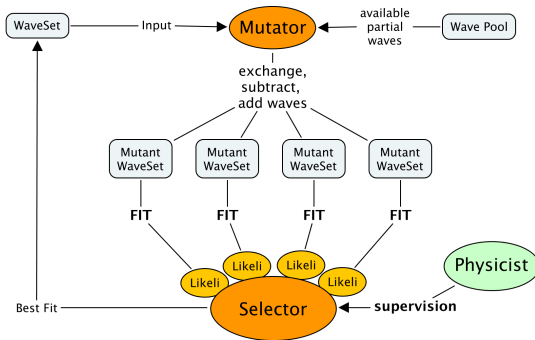


Automatic Waveset Exploration

Genetic Algorithm



Technische Universität München



Strategies:

- Start with population of small (2-15 waves) wavesets (adding waves)
- Start with diverse population (10 - 80 waves)
- Start with population of large wavesets (not done yet)

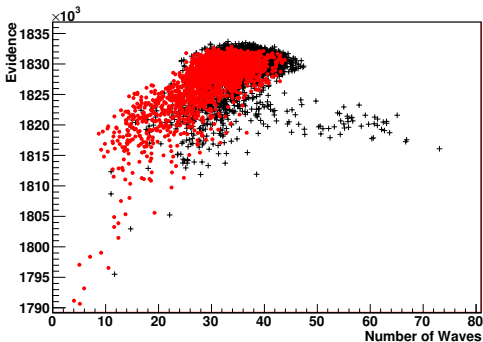


Automatic Waveset Exploration

Genetic Algorithm – 50 generations, population size 50



Technische Universität München



Number of waves optimizes at around 35

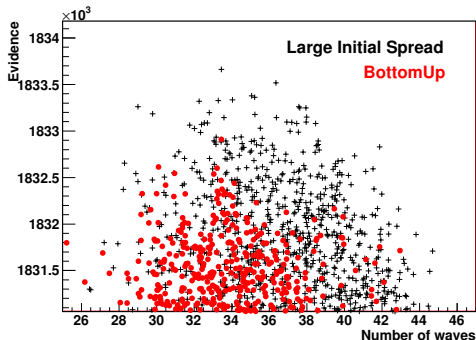


Automatic Waveset Exploration

Genetic Algorithm – 50 generations, population size 50



Technische Universität München



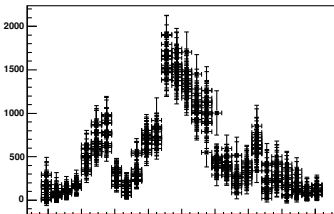
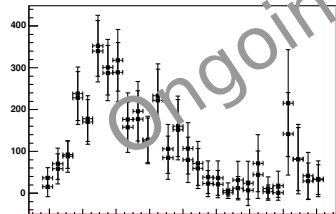
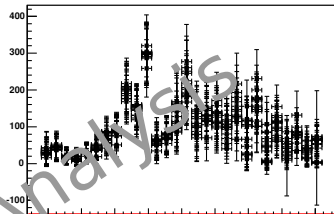
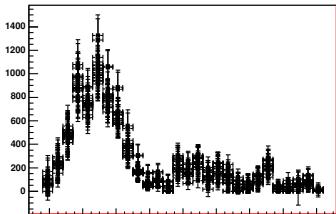
- Diverse initial population run \rightarrow better results
- Typical log-Evidence differences: 30-100



Example: Top 20 Fits from Genetic Search



Technische Universität München



Ongoing Analysis



Example Mass Dependent Fit

$$T_i^\epsilon T_j^{\epsilon*} = \rho_{ij}^\epsilon(m) = \left(\sum_k C_{ik}^\epsilon BW_k(m) \sqrt{\int |\psi_i^\epsilon|^2 d\tau} \right) \left(\sum_l C_{jl}^\epsilon BW_l(m) \sqrt{\int |\psi_j^\epsilon|^2 d\tau} \right) \quad (1)$$

with Breit-Wigner amplitude:

$$BW_{ik}(m, M_0, \Gamma_0) = \frac{M_0 \Gamma_0}{m^2 - M_0^2 + i\Gamma_{tot}(m)M_0} \quad (2)$$

and dynamic width:

$$\Gamma_{tot}(m) = \sum_n \gamma_n \frac{\rho_n(m)}{\rho_n(M_0)} \quad \rho_n(m) \sim \int |\psi_i^\epsilon|^2 dq \quad \sum \gamma_n = \Gamma_0 \quad (3)$$

and background terms:

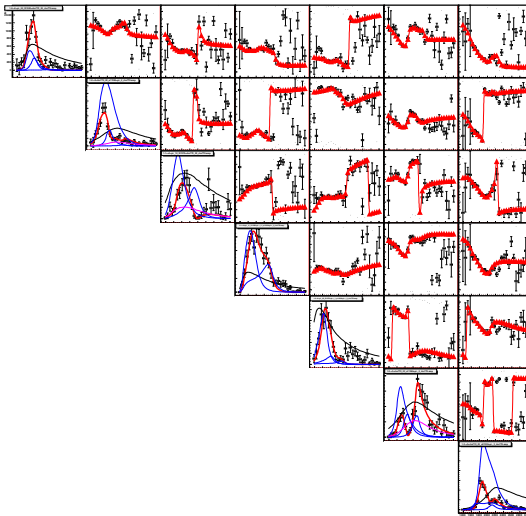
$$bkg(m) = e^{-\alpha q} \quad q - \text{Breakup momentum} \quad (4)$$



Fit Results Overview - Spin Density Matrix



7 waves, 8 resonances





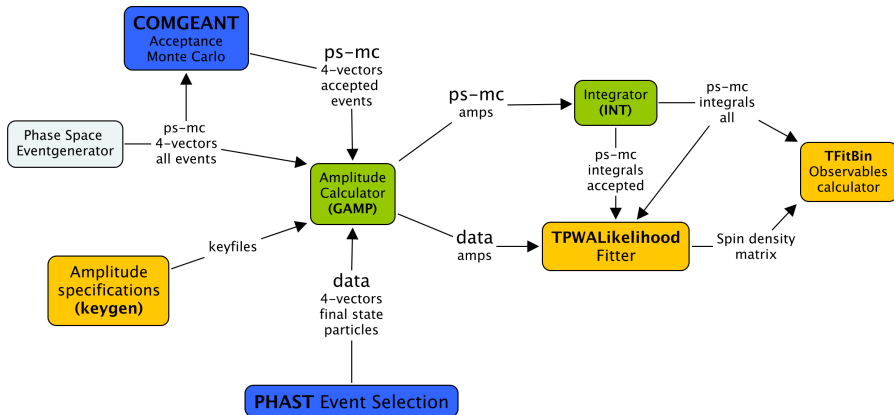
ROOTPWA: Open Source Analysis Toolkit



Technische Universität München

<http://sourceforge.net/projects/rootpwa>

- Based on BNL code “pwa2000”
- Largely rewritten
- Workflow for mass-dependent fit:





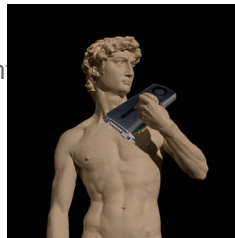
Main Features:

- **Amplitude calculator** for diffractive production (Helicity Form.)
- General **Amplitude Framework** upcoming (B. Grube)
- MC generators (diffraction)
- Numerical tools
 - MC integrator
 - **Fitters**
 - **Genetic Optimization**
- Resonance parameterizations (under development)
- Visualization & Plotting tools (ROOT-based)
- **CUDA support**



Main Features:

- **Amplitude calculator** for diffractive production (Helicity Form.)
- General **Amplitude Framework** upcoming (B. Grube)
- MC generators (diffraction)
- Numerical tools
 - MC integrator
 - **Fitters**
 - **Genetic Optimization**
- Resonance parameterizations (under development)
- Visualization & Plotting tools (ROOT-based)
- **CUDA support**





ROOTPWA Graphical User Interface

Developed by P. Jasinski



Technische Universität München

The screenshot displays the ROOTPWA GUI within an Eclipse IDE. The main window, titled 'rootpwa main window', contains a 'Kpipi PWA analysis' section with buttons for 'Load', 'Modify', 'New', 'Save', and 'Save as'. Below this is a progress bar for the analysis steps:

- set up workspace: 100%
- fit fit phase space events: 100%
- run MC acceptance analysis: 100%
- filter data into bins: 100%
- generate PWA keyfiles: 100%
- calculate PWA amplitudes: 100%
- integrate PWA amplitudes: 100%
- specify amplitudes for fit: 100%
- fit partial waves: 43%
- show results: 0%
- predict: 0%

The central plot area shows four graphs:

- Top-left: Intensity plot (Y-axis 0 to 18000) vs. \sqrt{s} (X-axis 1.7 to 1.9).
- Top-right: Phase Angle (deg) plot (Y-axis -200 to 200) vs. \sqrt{s} (X-axis 1.7 to 1.9).
- Bottom-left: Intensity plot (Y-axis 0 to 18000) vs. \sqrt{s} (X-axis 1.7 to 1.9).
- Bottom-right: Phase Angle (deg) plot (Y-axis -200 to 200) vs. \sqrt{s} (X-axis 1.7 to 1.9).

The terminal at the bottom shows the following commands and output:

```
File Edit View Terminal Tabs Help
>>> plotCoherence(): info: plotting
using selection criterion '(max
running TTree::Draw() expressi
'Mon_Dec_20_09_53_43_2010_', ''
scanning fit result Wed_Dec_15_17_
>>> plotIntensity(): info: plotting
&& (massBinCenter() <= 2990)'
running TTree::Draw() expressi
K_ .amp_0' on tree 'Wed_Dec_15_17_1
maximum intensity for graph 1-
>>> plotPhase(): info: plotting phi
using selection criterion '(max
running TTree::Draw() expressi
ec_15_17_15_33_2010_', ''
>>> plotIntensity(): info: plotting
10) && (massBinCenter() <= 2990)' &&
running TTree::Draw() expressi
01_pi_ .amp_0' on tree 'Wed_Dec_15_
maximum intensity for graph 1-
```



Summary

- ROOTPWA is one of 2 PWA programs used at COMPASS
- **2 step analysis:**
 - 1 Fit angular correlations with Isobar Model decay
 - 2 Parameterize dynamics → resonance extraction
- **Genetic search** for waveset exploration
- Open source toolkit <http://sourceforge.net/projects/rootpwa>



Summary

- ROOTPWA is one of 2 PWA programs used at COMPASS
- **2 step analysis:**
 - 1 Fit angular correlations with Isobar Model decay
 - 2 Parameterize dynamics → resonance extraction
- **Genetic search** for waveset exploration
- Open source toolkit <http://sourceforge.net/projects/rootpwa>

Outlook

- Improvements in amplitude parameterizations
- Study of non-resonant contributions (Deck effect)
- Theory input needed (Rescattering etc.)
- Status of Analyses and Results → **Talk by B. Ketzer tomorrow**