

Precise reconstruction of sparticle masses without ambiguities

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Precise reconstruction of sparticle masses without ambiguities

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ABSTRACT: We critically reexamine the standard applications of the method of kinematical endpoints for sparticle mass determination. We consider the typical decay chain in supersymmetry (SUSY) $\tilde{q} \rightarrow \tilde{\chi}_2^0 \rightarrow \tilde{\ell} \rightarrow \tilde{\chi}_1^0$, which yields a jet j , and two leptons ℓ_n^\pm and ℓ_f^\mp . The conventional approaches use the upper kinematical endpoints of the individual distributions $m_{j\ell\ell}$, $m_{j\ell(lo)} = \min\{m_{j\ell_n}, m_{j\ell_f}\}$ and $m_{j\ell(hi)} = \max\{m_{j\ell_n}, m_{j\ell_f}\}$, all three of which suffer from parameter space region ambiguities and may lead to multiple solutions for the SUSY mass spectrum. In contrast, we do not use $m_{j\ell\ell}$, $m_{j\ell(lo)}$ and $m_{j\ell(hi)}$, and instead propose a new set of (infinitely many) variables whose upper kinematic endpoints exhibit reduced sensitivity to the parameter space region. We then outline an alternative, much simplified procedure for obtaining the SUSY mass spectrum. In particular, we show that the four endpoints observed in the three distributions $m_{\ell\ell}^2$, $m_{j\ell_n}^2 \cup m_{j\ell_f}^2$ and $m_{j\ell_n}^2 + m_{j\ell_f}^2$ are sufficient to completely pin down the squark mass $m_{\tilde{q}}$ and the two neutralino masses $m_{\tilde{\chi}_2^0}$ and $m_{\tilde{\chi}_1^0}$, leaving only a discrete 2-fold ambiguity for the slepton mass $m_{\tilde{\ell}}$. This remaining ambiguity can be easily resolved in a number of different ways: for example, by a single additional measurement of the kinematic endpoint of any one out of the many remaining 1-dimensional distributions at our disposal, or by exploring the correlations in the 2-dimensional distribution of $m_{j\ell_n}^2 \cup m_{j\ell_f}^2$ versus $m_{\ell\ell}^2$. We illustrate our method with two examples: the LM1 and LM6 CMS study points. An additional advantage of our method is the expected improvement in the accuracy of the SUSY mass determination, due to the multitude and variety of available measurements.

KEYWORDS: Supersymmetry Phenomenology

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1 Introduction

SUSY is a primary target of the LHC searches for new physics beyond the Standard Model (BSM). In SUSY models with conserved R -parity the superpartners are produced in pairs

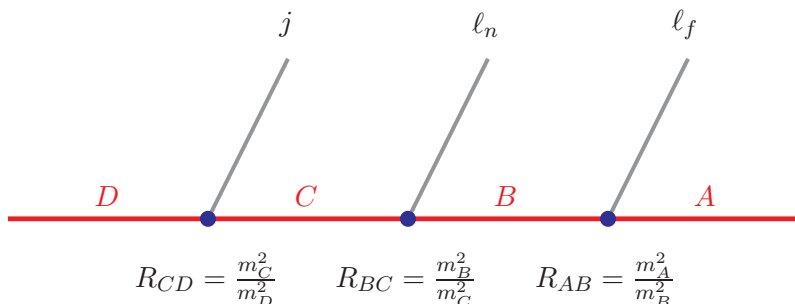


Figure 1. The typical cascade decay chain under consideration in this paper. Here D , C , B and A are new BSM particles, while the corresponding SM decay products are: a QCD jet j , a “near” lepton ℓ_n^\pm and a “far” lepton ℓ_f^\mp . This chain is quite common in SUSY, with the identification $D = \tilde{q}$, $C = \tilde{\chi}_2^0$, $B = \tilde{\ell}$ and $A = \tilde{\chi}_1^0$ ($\tilde{\chi}_2^0$) is the first (second) lightest neutralino. In what follows we shall quote our results in terms of the D mass m_D and the three dimensionless squared mass ratios R_{CD} , R_{BC} and R_{AB} defined in eq. (1.6).

and each one decays through a cascade decay chain down to the lightest superpartner (LSP). If the LSP is the lightest neutralino $\tilde{\chi}_1^0$, it escapes detection, making it rather difficult to reconstruct directly the preceding superpartners and thus measure their masses and spins. In recognition of this fact, in recent years there has been an increased interest in developing new techniques for mass [1–49] and spin [50–76] measurements in such SUSY-like missing energy events.

Roughly speaking, there are three basic types of mass determination methods in SUSY.¹ In this paper we concentrate on the classic method of kinematical endpoints [1]. Following the previous SUSY studies, for illustration of our results we shall use the generic decay chain $D \rightarrow jC \rightarrow j\ell_n^\pm B \rightarrow j\ell_n^\pm \ell_f^\mp A$ shown in figure 1. Here D , C , B and A are new BSM particles with masses m_D , m_C , m_B and m_A . Their corresponding SM decay products are: a QCD jet j , a “near” lepton ℓ_n^\pm and a “far” lepton ℓ_f^\mp . This decay chain is quite common in SUSY, with the identification $D = \tilde{q}$, $C = \tilde{\chi}_2^0$, $B = \tilde{\ell}$ and $A = \tilde{\chi}_1^0$, where \tilde{q} is a squark, $\tilde{\ell}$ is a slepton, and $\tilde{\chi}_1^0$ ($\tilde{\chi}_2^0$) is the first (second) lightest neutralino. However, our analysis is not limited to SUSY only, since the chain in figure 1 also appears in other BSM scenarios, e.g. Universal Extra Dimensions [77]. For concreteness, we shall assume that all three decays exhibited in figure 1 are two-body, i.e. we shall consider the mass hierarchy

$$m_D > m_C > m_B > m_A > 0. \quad (1.1)$$

This presents the most challenging case, in which one has to determine all four masses m_D , m_C , m_B and m_A .

The idea of the kinematic endpoint method is very simple. Given the SM decay

¹For a recent study representative of each method, see refs. [43, 47, 49].

products j , ℓ_n and ℓ_f exhibited in figure 1, form the invariant mass² of every possible combination, $m_{\ell\ell}$, $m_{j\ell_n}$, $m_{j\ell_f}$, and $m_{j\ell\ell}$, plot the resulting distributions and measure the corresponding upper kinematic endpoints [1, 7, 12]

$$(m_{\ell\ell}^{\max})^2 = m_D^2 R_{CD} (1 - R_{BC}) (1 - R_{AB}); \tag{1.2}$$

$$(m_{j\ell_n}^{\max})^2 = m_D^2 (1 - R_{CD}) (1 - R_{BC}); \tag{1.3}$$

$$(m_{j\ell_f}^{\max})^2 = m_D^2 (1 - R_{CD}) (1 - R_{AB}); \tag{1.4}$$

$$(m_{j\ell\ell}^{\max})^2 = \begin{cases} m_D^2(1 - R_{CD})(1 - R_{AC}), & \text{for } R_{CD} < R_{AC}, & \text{case (1, -),} \\ m_D^2(1 - R_{BC})(1 - R_{AB}R_{CD}), & \text{for } R_{BC} < R_{AB}R_{CD}, & \text{case (2, -),} \\ m_D^2(1 - R_{AB})(1 - R_{BD}), & \text{for } R_{AB} < R_{BD}, & \text{case (3, -),} \\ m_D^2(1 - \sqrt{R_{AD}})^2, & \text{otherwise,} & \text{case (4, -).} \end{cases} \tag{1.5}$$

Here and below we follow the notation and conventions of ref. [47], i.e. we write all results in terms of an overall mass scale (given by the mass m_D of the heaviest BSM particle D) and three dimensionless squared mass ratios

$$R_{ij} \equiv \frac{m_i^2}{m_j^2}, \quad i, j \in \{A, B, C, D\}. \tag{1.6}$$

Note that there are only three independent ratios in (1.6). We shall take those to be R_{AB} , R_{BC} , and R_{CD} (see figure 1), and their definition domain will be the interval $(0, 1)$.³

In spite of their transparent theoretical meaning, the set of four endpoints (1.2)–(1.5) by themselves have (justifiably) never been used as the sole basis for a SUSY mass determination analysis. This is due to three generic problems, which are all very well known, and are separately reviewed in the next three subsections 1.1, 1.2 and 1.3. Our new approach to resolving these three problems, and the outline of the rest of the paper are presented in section 1.4.

1.1 Near-far lepton ambiguity

The first problem is that one cannot differentiate between the “near” and “far” leptons ℓ_n and ℓ_f on an event-by-event basis. Since all decays in figure 1 are prompt, both leptons point back to the primary interaction vertex and there is no way to tell which came first and which came second. Consequently, one cannot *separately* construct the individual $m_{j\ell_n}$ and $m_{j\ell_f}$ invariant mass distributions, whose upper endpoints would be given by (1.3) and (1.4). This problem has motivated most of the previous invariant mass studies in the literature,

²We shall see below that the formulas simplify considerably if we use invariant masses *squared* instead. This distinction is not central to our analysis.

³As seen in eq. (1.5), at times we shall also utilize one or more of the other three ratios, R_{AC} , R_{AD} and R_{BD} , whenever this will lead to a simplification of the formulas. Of course, the latter three ratios are related to our preferred set $\{R_{AB}, R_{BC}, R_{CD}\}$ due to the transitivity property $R_{ij}R_{jk} = R_{ik}$.

beginning with [7], to introduce an alternative definition of the two $j\ell$ distributions, simply by ordering the two $m_{j\ell}$ entries in each event by invariant mass as follows

$$m_{j\ell(lo)} \equiv \min \{m_{j\ell_n}, m_{j\ell_f}\}, \tag{1.7}$$

$$m_{j\ell(hi)} \equiv \max \{m_{j\ell_n}, m_{j\ell_f}\}. \tag{1.8}$$

Both of the newly defined quantities $m_{j\ell(lo)}$ and $m_{j\ell(hi)}$ also exhibit upper kinematic endpoints ($m_{j\ell(lo)}^{\max}$ and $m_{j\ell(hi)}^{\max}$, correspondingly). Since the *individual* $m_{j\ell(lo)}$ and $m_{j\ell(hi)}$ distributions are observable, their endpoints are experimentally measurable and can be related to the underlying SUSY mass spectrum as follows [7, 12]

$$\left(m_{j\ell(lo)}^{\max}\right)^2 = \begin{cases} \left(m_{j\ell_n}^{\max}\right)^2, & \text{for } (2 - R_{AB})^{-1} < R_{BC} < 1, & \text{case } (-, 1), \\ \left(m_{j\ell(eq)}^{\max}\right)^2, & \text{for } R_{AB} < R_{BC} < (2 - R_{AB})^{-1}, & \text{case } (-, 2), \\ \left(m_{j\ell(eq)}^{\max}\right)^2, & \text{for } 0 < R_{BC} < R_{AB}, & \text{case } (-, 3); \end{cases} \tag{1.9}$$

$$\left(m_{j\ell(hi)}^{\max}\right)^2 = \begin{cases} \left(m_{j\ell_f}^{\max}\right)^2, & \text{for } (2 - R_{AB})^{-1} < R_{BC} < 1, & \text{case } (-, 1), \\ \left(m_{j\ell_f}^{\max}\right)^2, & \text{for } R_{AB} < R_{BC} < (2 - R_{AB})^{-1}, & \text{case } (-, 2), \\ \left(m_{j\ell_n}^{\max}\right)^2, & \text{for } 0 < R_{BC} < R_{AB}, & \text{case } (-, 3); \end{cases} \tag{1.10}$$

where

$$\left(m_{j\ell(eq)}^{\max}\right)^2 = m_D^2 (1 - R_{CD}) (1 - R_{AB}) (2 - R_{AB})^{-1} \tag{1.11}$$

and $m_{j\ell_n}^{\max}$ and $m_{j\ell_f}^{\max}$ were already defined in (1.3) and (1.4), correspondingly. With this approach, the original set of 4 endpoints in eqs. (1.2)–(1.5) is replaced by

$$m_{\ell\ell}^{\max}, m_{j\ell\ell}^{\max}, m_{j\ell(lo)}^{\max}, m_{j\ell(hi)}^{\max}. \tag{1.12}$$

In contrast to this conventional approach in the literature, we shall adopt a very different attitude towards resolving the problem of the near-far lepton ambiguity. We will do the simplest possible thing, namely, we shall do nothing. We shall never ask the question “which lepton was ℓ_n and which one was ℓ_f ?”. We shall also *not* use the ordering (1.7), (1.8). Instead, we shall simply take the two $m_{j\ell}$ entries in each event, and always treat them in a symmetric fashion. For example, any observable invariant mass distribution that we will build out of the two measured quantities $m_{j\ell_n}$ and $m_{j\ell_f}$ should be invariant under the symmetry

$$m_{j\ell_n} \leftrightarrow m_{j\ell_f}. \tag{1.13}$$

The advantages of our approach may not be immediately obvious at this point, but will become clear in the process of our mass determination analysis in section 3 below.

1.2 Insufficient number of measurements

The second problem associated with the original set of four measurements (1.2)–(1.5), as well as the alternative set (1.12), is that the measured endpoints may not all be independent from each other. Indeed, there are certain regions of parameter space where one finds the following correlation [12]

$$(m_{j\ell\ell}^{\max})^2 = (m_{j\ell(hi)}^{\max})^2 + (m_{\ell\ell}^{\max})^2. \quad (1.14)$$

In this case, the four measurements (1.12) are clearly insufficient to pin down all four independent input parameters m_D , m_C , m_B and m_A . Therefore, one has to measure an additional independent endpoint. To this end, it has been suggested to consider the constrained distribution $m_{j\ell\ell(\theta>\frac{\pi}{2})}$, which exhibits a useful *lower* kinematic endpoint $m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min}$ [7]

$$\begin{aligned} \left(m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min}\right)^2 = \frac{1}{4}m_D^2 & \left\{ (1 - R_{AB})(1 - R_{BC})(1 + R_{CD}) \right. \\ & \left. + 2(1 - R_{AC})(1 - R_{CD}) - (1 - R_{CD})\sqrt{(1 + R_{AB})^2(1 + R_{BC})^2 - 16R_{AC}} \right\}. \end{aligned} \quad (1.15)$$

The distribution $m_{j\ell\ell(\theta>\frac{\pi}{2})}$ is nothing but the usual $m_{j\ell\ell}$ distribution over a subset of the original events, subject to the additional dilepton mass constraint

$$\frac{m_{\ell\ell}^{\max}}{\sqrt{2}} < m_{\ell\ell} < m_{\ell\ell}^{\max}. \quad (1.16)$$

In the rest frame of particle B , this cut implies the following restriction on the opening angle θ between the two leptons [6]

$$\theta > \frac{\pi}{2}, \quad (1.17)$$

thus justifying the notation for $m_{j\ell\ell(\theta>\frac{\pi}{2})}$.

The advantage of the “threshold” endpoint measurement (1.15) is that it is always independent of the other four measurements in (1.12). As a result, it would appear that the enlarged set of five kinematic endpoint measurements

$$m_{\ell\ell}^{\max}, m_{j\ell\ell}^{\max}, m_{j\ell(lo)}^{\max}, m_{j\ell(hi)}^{\max}, m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min} \quad (1.18)$$

should be in principle sufficient to determine all four unknown masses (see, however, ref. [47] and section 1.3 below).

Unfortunately, the “threshold” (1.15) also suffers from certain disadvantages, which are mostly of experimental nature. It is generally expected that the experimental precision on the determination of the *lower* kinematic endpoint (1.15) will be rather inferior compared to the precision on the other four *upper* kinematic endpoints (1.12) [12]. There are several generic reasons for such a pessimistic attitude. First, the region in the $m_{j\ell\ell(\theta>\frac{\pi}{2})}$ distribution near its lower endpoint (1.15) is rather sparsely populated, resulting in a shallow edge and sizable statistical errors. To make matters worse, the $m_{j\ell\ell(\theta>\frac{\pi}{2})}$ distribution near its lower edge is a convex function [19], which makes it even more difficult to tell where the

signal ends and the tails from various sources begin [12]. Finally, the low mass region of almost any invariant mass distribution in SUSY is generally associated with larger SM (as well as SUSY combinatorial) backgrounds compared to its high mass counterpart.

Overall we find all these disadvantages sufficiently convincing so that we will drop the measurement (1.15) altogether and will never use it in the course of our analysis in section 3 below. We will be justified in doing so, since the linear dependence problem (1.14), which has plagued previous studies and was the prime motivation for introducing the $m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min}$ measurement in the first place, will have no effect on our analysis. In fact, we will not be using the endpoint measurement $m_{j\ell(hi)}^{\max}$ (for the reasons given in the previous subsection 1.1) and we will not be using the endpoint measurement $m_{j\ell\ell}^{\max}$ (for the reasons given in the following subsection 1.3). Once these two problematic measurements are removed from consideration, the linear dependence problem (1.14) does not arise, and the “threshold” measurement (1.15) is not central to the analysis any more.

1.3 Parameter space region ambiguity

The third problem with the conventional set of measurements (1.18) is immediately obvious from the defining equations (1.5), (1.9) and (1.10) for the kinematic endpoints $m_{j\ell\ell}^{\max}$, $m_{j\ell(lo)}^{\max}$, and $m_{j\ell(hi)}^{\max}$, correspondingly. One can see that the relevant expressions are piecewise-defined functions, i.e. they depend on the values of the independent variables m_A , m_B , m_C and m_D . For example, there are four different cases for $m_{j\ell\ell}^{\max}$, and three different cases for the pair of $(m_{j\ell(lo)}^{\max}, m_{j\ell(hi)}^{\max})$. Altogether, these give rise to 9 different cases⁴ which must be separately considered [12, 47]. Of course, this represents a problem, since the masses are a priori unknown, and it is not clear which case is the relevant one. Barring any model-dependent assumptions, one is forced to consider all possibilities, obtain a solution for the spectrum, and only at the very end, test whether the solution falls within the parameter space applicable for the case at hand. This procedure may often result in several alternative solutions [12, 47, 78–82]. In fact, ref. [47] recently proved that there exists a sizable parameter space region in which even the full set of measurements (1.18) would always yield two alternative solutions, even under ideal experimental conditions. The problem is further exacerbated by the inevitable experimental errors on the measurements (1.18), which would allow for an even larger number of “fake” or “duplicate” solutions [47, 79, 80].

Having identified the root of the duplication problem as the piecewise definition of the mathematical formulas in (1.5), (1.9), (1.10), our solution to the problem will be again very simple and conservative. We will simply avoid using any kinematic endpoints which are given in terms of piecewise-defined expressions. This requirement automatically eliminates from consideration the three conventional endpoints $m_{j\ell\ell}^{\max}$, $m_{j\ell(lo)}^{\max}$, and $m_{j\ell(hi)}^{\max}$. Since we already gave up on $m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min}$ in the previous subsection, this leaves $m_{\ell\ell}^{\max}$ as the only measurement out of the conventional set (1.18) that we shall use in our analysis. This is perhaps the most drastic difference between our approach and all previous studies in the literature.

⁴The remaining 3 cases are always unphysical [12].

1.4 Posing the problem

In the previous three subsections we discussed each of the three generic theoretical⁵ problems with the previous applications of the kinematic endpoint method for mass determination. We are now ready to explicitly formulate our main goal in this paper. We aim to design a method for measuring the masses of the particles in the decay chain of figure 1, which is based on kinematic endpoint information, and satisfies the following requirements:

- It does not make use of any kinematic endpoints whose interpretation is ambiguous, i.e. whose expressions in terms of the physical masses are piecewise-defined functions.
- It does not make use of any *lower* kinematic endpoints such as the “threshold” $m_{j\ell\ell(\theta>\frac{\pi}{2})}^{\min}$, due to the experimental challenges with such measurements.
- It relies solely on 1-dimensional distributions, unlike the methods recently advertised in [18, 46, 47, 81, 82], which utilize 2-dimensional correlation plots. While the latter do provide a wealth of valuable information, they also typically require more data in order to obtain good enough statistics for drawing any robust conclusions from them. In contrast, the one-dimensional distributions should be available rather early on, and with sufficient statistics for endpoint measurements.

As already alluded to in the previous subsections, the first two requirements already eliminate four out of the five conventional inputs (1.18). Obviously, we will need to find a way to replace those with an alternative set of kinematic endpoint measurements which nevertheless satisfy the above requirements. In section 2 we introduce and investigate a new set of invariant mass variables whose upper endpoints can be useful for our analysis. Then in section 3 we outline our basic method, which makes use of some of these new variables. We illustrate our discussion in section 4 with two numerical examples: the LM1 and LM6 CMS study points. Section 5 is reserved for our conclusions. In appendix A we supply the analytic expressions for the shapes of the 1-dimensional invariant mass distributions used in our main analysis in section 3.1. Those results can be useful in improving the precision on the extraction of the kinematical endpoints.

2 New variables

In this section we propose a new set of invariant mass (squared) variables. As already explained in the Introduction, our variables should be composed of $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$ in a symmetric way, in accordance with (1.13). Consequently, any plotting manipulations or

⁵In addition, there are problems which are of experimental nature, e.g. identifying the correct jet and the correct lepton pair resulting from the decay chain in figure 1. There exists a set of standard experimental techniques which are aimed at overcoming these problems, e.g. the opposite flavor subtraction for the two leptons and the mixed event subtraction for the jet [83]. Wrong $\ell\ell$ and $j\ell$ pairings can also be identified and a posteriori removed whenever an invariant mass entry for $m_{\ell\ell}$, $m_{j\ell}$ or $m_{j\ell\ell}$ exceeds the corresponding kinematic endpoint $m_{\ell\ell}^{\max}$, $m_{j\ell(\text{hi})}^{\max}$ or $m_{j\ell\ell}^{\max}$. In what follows we shall assume that those preliminary steps have already been done and the samples we are dealing with have already been appropriately subtracted to remove the combinatorial background.

mathematical operations involving $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$ should obey the symmetry implied by eq. (1.13).

2.1 The union $m_{j\ell_n}^2 \cup m_{j\ell_f}^2$

We begin with the simplest case, where we postpone applying any mathematical operations to $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$, and instead simply plot them. The requirement of eq. (1.13) implies that the only possibility is to place both of them together on the same plot, in essence forming the union

$$m_{jl(u)}^2 \equiv m_{j\ell_n}^2 \cup m_{j\ell_f}^2 \tag{2.1}$$

of the individual $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$ distributions. Since each individual distribution is smooth and has a kinematic endpoint, the same two kinematic endpoints should be visible on the combined distribution $m_{jl(u)}^2$ as well.⁶ We shall denote the larger of the two endpoints with

$$\left(M_{jl(u)}^{\max}\right)^2 \equiv \max \left\{ \left(m_{j\ell_n}^{\max}\right)^2, \left(m_{j\ell_f}^{\max}\right)^2 \right\} \tag{2.2}$$

and the smaller of the two endpoints with

$$\left(m_{jl(u)}^{\min}\right)^2 \equiv \min \left\{ \left(m_{j\ell_n}^{\min}\right)^2, \left(m_{j\ell_f}^{\min}\right)^2 \right\}. \tag{2.3}$$

The newly introduced quantities $M_{jl(u)}^{\max}$ and $m_{jl(u)}^{\min}$ are nothing but the usual kinematic endpoints $m_{j\ell_n}^{\max}$ and $m_{j\ell_f}^{\max}$, given by (1.3) and (1.4), correspondingly. Of course, at this point we do not know which is which, and we have an apparent two-fold ambiguity: we can have either

$$M_{jl(u)}^{\max} = m_{j\ell_n}^{\max}, \quad m_{jl(u)}^{\min} = m_{j\ell_f}^{\min}, \quad \text{if } R_{AB} \geq R_{BC}, \tag{2.4}$$

or

$$M_{jl(u)}^{\max} = m_{j\ell_f}^{\max}, \quad m_{jl(u)}^{\min} = m_{j\ell_n}^{\min}, \quad \text{if } R_{AB} \leq R_{BC}. \tag{2.5}$$

Notice that both (2.2) and (2.3) are officially *upper* kinematic endpoints, and thus satisfy our basic requirements.

The benefits of our alternative treatment (2.1) in response to the near-far lepton ambiguity problem of section 1.1, are now starting to emerge. With the conventional ordering (1.7), (1.8) one has to deal with a *three-fold* ambiguity in the interpretation of the endpoints $m_{j\ell(lo)}^{\max}$ and $m_{j\ell(hi)}^{\max}$, as seen in eqs. (1.9), (1.10). Instead, the simple union (2.1) leads only to the *two-fold* ambiguity of eqs. (2.4), (2.5). More importantly, the analysis of section 3.1 below will reveal that in spite of the remaining two-fold ambiguity in eqs. (2.4), (2.5), one can nevertheless *uniquely* determine all three of the masses m_D , m_C and m_A ! We consider this to be one of the important results of this paper.

⁶For specific numerical examples, refer to section 4.

2.2 The product $m_{j\ell_n} \times m_{j\ell_f}$

In the remainder of this section, we shall construct new invariant mass squared variables out of the two entries $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$, simply by applying various mathematical operations on them in a symmetric fashion. We begin with the product

$$m_{j\ell(p)}^2 \equiv m_{j\ell_n} m_{j\ell_f} \tag{2.6}$$

whose endpoint is given by

$$\left(m_{j\ell(p)}^{\max}\right)^2 \equiv \begin{cases} \frac{1}{2} m_D^2 (1 - R_{CD}) \sqrt{1 - R_{AB}}, & \text{for } R_{BC} \leq 0.5, \\ m_D^2 (1 - R_{CD}) \sqrt{R_{BC}(1 - R_{BC})(1 - R_{AB})}, & \text{for } R_{BC} \geq 0.5. \end{cases} \tag{2.7}$$

Unfortunately, this endpoint also turns out to be piecewise-defined, thus failing one of our basic requirements from the Introduction. Therefore we shall not use this endpoint in the course of our analysis.

2.3 The sums $m_{j\ell_n}^{2\alpha} + m_{j\ell_f}^{2\alpha}$

Another possibility is to consider various sums, for example $m_{j\ell_n}^2 + m_{j\ell_f}^2$ or $(m_{j\ell_n} + m_{j\ell_f})^2$, as originally proposed in [18]. Here we generalize the discussion in [18] and introduce a whole set of new variables, $m_{j\ell(s)}^2(\alpha)$, labelled by the continuous parameter α , which are defined as

$$m_{j\ell(s)}^2(\alpha) \equiv \left(m_{j\ell_n}^{2\alpha} + m_{j\ell_f}^{2\alpha}\right)^{\frac{1}{\alpha}}. \tag{2.8}$$

Since α is a continuous parameter, in principle there are infinitely many $m_{j\ell(s)}$ variables! Notice that the conventional variables $m_{j\ell(lo)}^2$ and $m_{j\ell(hi)}^2$ from (1.7) and (1.8) are also included in our set, and are simply given by

$$m_{j\ell(lo)}^2 \equiv m_{j\ell(s)}^2(-\infty), \tag{2.9}$$

$$m_{j\ell(hi)}^2 \equiv m_{j\ell(s)}^2(\infty). \tag{2.10}$$

We see that our new set (2.8) is a very broad generalization of the conventional definitions (1.7) and (1.8), which just correspond to the two extreme cases $\alpha = \pm\infty$. Of course, the user is free to choose α at will, and any finite value of α will lead to a new variable $m_{j\ell(s)}^2(\alpha)$.

In order to make the new variables $m_{j\ell(s)}^2(\alpha)$ useful for mass spectrum studies, we need to provide the formulas for their kinematic endpoints $(m_{j\ell(s)}^{\max}(\alpha))^2$. These formulas are easy to derive, using the results from [47], and we present them in the next two subsections, where it is convenient to consider separately the following two cases: $\alpha \geq 1$ (in section 2.3.1) and $\alpha < 1$, but $\alpha \neq 0$ (in section 2.3.2).

2.3.1 Kinematic endpoints of $m_{j\ell(s)}^2(\alpha)$ with $\alpha \geq 1$

When one chooses a value of $\alpha \geq 1$, the $m_{j\ell(s)}^2(\alpha)$ endpoint is given by the following expression

$$\left(m_{j\ell(s)}^{\max}(\alpha \geq 1)\right)^2 \equiv \begin{cases} \left(m_{j\ell_f}^{\max}\right)^2, & R_{AB} \leq 1 - (1 - R_{BC})(1 - R_{BC}^\alpha)^{-\frac{1}{\alpha}}, \\ \left(m_{j\ell}^{\max}(\alpha)\right)^2, & R_{AB} \geq 1 - (1 - R_{BC})(1 - R_{BC}^\alpha)^{-\frac{1}{\alpha}}, \end{cases} \tag{2.11}$$

where $m_{j\ell_f}^{\max}$ was already defined in (1.4), and $m_{j\ell}^{\max}(\alpha)$ is a newly defined, α -dependent quantity

$$\left(m_{j\ell}^{\max}(\alpha)\right)^2 \equiv m_D^2(1 - R_{CD}) \left[R_{BC}^\alpha (1 - R_{AB})^\alpha + (1 - R_{BC})^\alpha \right]^{\frac{1}{\alpha}}. \quad (2.12)$$

As a cross-check, one can verify that in the limit $\alpha \rightarrow \infty$ the expression (2.11) reduces to (1.10), in agreement with (2.10). In that case, the upper line in (2.11) corresponds to options $(-, 1)$ and $(-, 2)$ in (1.10), where $m_{j\ell(hi)}^{\max} = m_{j\ell_f}^{\max}$, while the lower line in (2.11) corresponds to option $(-, 3)$ in (1.10), where $m_{j\ell(hi)}^{\max} = m_{j\ell_n}^{\max}$. Unfortunately, just like the product endpoint (2.7), the endpoint (2.11) is in general piecewise-defined, and does not meet our criteria.

However, there is one important exception, namely the case of $\alpha = 1$, in which we do get a singly defined function. According to the general definition (2.8), $m_{j\ell(s)}^2(\alpha = 1)$ is simply the sum of the two $m_{j\ell}^2$ entries in each event:

$$m_{j\ell(s)}^2(\alpha = 1) \equiv m_{j\ell_n}^2 + m_{j\ell_f}^2. \quad (2.13)$$

Using the identity

$$m_{j\ell\ell}^2 = m_{j\ell_n}^2 + m_{j\ell_f}^2 + m_{\ell\ell}^2, \quad (2.14)$$

(2.13) can be equivalently rewritten as

$$m_{j\ell(s)}^2(\alpha = 1) \equiv m_{j\ell\ell}^2 - m_{\ell\ell}^2. \quad (2.15)$$

To find the expression for its endpoint, one can set $\alpha = 1$ in (2.11), and then realize that the logical condition for executing the upper line becomes $R_{AB} \leq 0$, which is impossible, since the mass ratios R_{ij} in (1.6) are always positive definite. Therefore, the endpoint $m_{j\ell(s)}^{\max}(\alpha = 1)$ is always calculated according to the lower line in (2.11), which results in [18]

$$\left(m_{j\ell(s)}^{\max}(1)\right)^2 \equiv m_D^2(1 - R_{CD})(1 - R_{AC}). \quad (2.16)$$

Note that this endpoint is perfect for our purposes since the formula (2.16) is always unique, i.e. it is independent of the parameter space region. The variable $m_{j\ell(s)}^2(\alpha = 1)$ will thus play a crucial role in our analysis below.

2.3.2 Kinematic endpoints of $m_{j\ell(s)}^2(\alpha)$ with $\alpha < 1$ and $\alpha \neq 0$

Finally, in the case when $\alpha < 1$, but $\alpha \neq 0$, the $m_{j\ell(s)}^2(\alpha)$ endpoint is given by the following expression

$$\left(m_{j\ell(s)}^{\max}(\alpha < 1)\right)^2 \equiv \begin{cases} \left(m_{j\ell}^{\max}(\alpha)\right)^2, & R_{BC} \geq \left[1 + (1 - R_{AB})^{\frac{\alpha}{\alpha-1}}\right]^{-1}, \\ m_D^2(1 - R_{CD}) \left[1 + (1 - R_{AB})^{\frac{\alpha}{1-\alpha}}\right]^{\frac{1-\alpha}{\alpha}}, & R_{BC} \leq \left[1 + (1 - R_{AB})^{\frac{\alpha}{\alpha-1}}\right]^{-1}, \end{cases} \quad (2.17)$$

where $m_{j\ell}^{\max}(\alpha)$ was already defined in (2.12). Again as a cross-check, one can verify that in the limit $\alpha \rightarrow -\infty$ the expression (2.17) reduces to (1.9), in agreement with (2.9). In the $\alpha \rightarrow -\infty$ case, the upper line in (2.17) corresponds to option $(-, 1)$ in (1.9), where $m_{j\ell(lo)}^{\max} = m_{j\ell_n}^{\max}$, while the lower line in (2.17) corresponds to options $(-, 2)$ and $(-, 3)$ in (1.9), where $m_{j\ell(lo)}^{\max} = m_{j\ell(eq)}^{\max}$. Unfortunately, the endpoint function (2.17) is again piecewise-defined, and does not meet one of our basic criteria spelled out in the introduction.

In passing, we note that the special case of $\alpha = \frac{1}{2}$, which involves the *linear* sum of the two masses

$$m_{j\ell(s)}^2\left(\alpha = \frac{1}{2}\right) \equiv (m_{j\ell_n} + m_{j\ell_f})^2, \quad (2.18)$$

was previously explored in [18, 84]. In that case, from (2.17) we find for its endpoint

$$\left(m_{j\ell(s)}^{\max}\left(\frac{1}{2}\right)\right)^2 \equiv \begin{cases} m_D^2(1 - R_{CD}) \left(\sqrt{R_{BC}(1 - R_{AB})} + \sqrt{1 - R_{BC}}\right)^2, & R_{BC} \geq \frac{1 - R_{AB}}{2 - R_{AB}}, \\ m_D^2(1 - R_{CD})(2 - R_{AB}), & R_{BC} \leq \frac{1 - R_{AB}}{2 - R_{AB}}. \end{cases} \quad (2.19)$$

2.4 The difference $|m_{j\ell_n}^2 - m_{j\ell_f}^2|$

Finally, one can also consider a set of variables which involve the absolute value of differences between $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$. In analogy with (2.8), we can define another infinite set of variables

$$m_{j\ell(d)}^2(\alpha) \equiv \left| m_{j\ell_n}^{2\alpha} - m_{j\ell_f}^{2\alpha} \right|^{\frac{1}{\alpha}}. \quad (2.20)$$

Once again, the user is free to consider arbitrary values of α . However, this freedom is redundant, when it comes to the issue of the kinematic endpoints of the variables in (2.20). It is not difficult to see that the endpoints of $m_{j\ell(d)}^2(\alpha)$ are always given by

$$\left(m_{j\ell(d)}^{\max}(\alpha)\right)^2 \equiv \left(M_{j\ell(u)}^{\max}\right)^2 \quad (2.21)$$

and are in fact independent of α ! Therefore, for the purposes of our discussion, it is sufficient to consider just one particular value of α . In the following we shall only use $\alpha = 1$:

$$m_{j\ell(d)}^2(\alpha = 1) \equiv \left| m_{j\ell_n}^2 - m_{j\ell_f}^2 \right|, \quad (2.22)$$

which is the analogue of $m_{j\ell(s)}^2(\alpha = 1)$ defined in (2.13).

The result (2.21) implies that the endpoint of (2.22) does not contain any new amount of information, which was not already present in the two kinematic endpoints $M_{j\ell(u)}^{\max}$ and $m_{j\ell(u)}^{\max}$ discussed in section 2.1. Nevertheless, the independent measurement of $(m_{j\ell(d)}^{\max}(1))^2$ can still be very useful, since it will mark the location of $(M_{j\ell(u)}^{\max})^2$ on the $m_{j\ell(u)}^2$ distribution. Then one will be looking for the second endpoint $(m_{j\ell(u)}^{\max})^2$ to the left, i.e. in the region of smaller $m_{j\ell(u)}^2$ values.

This completes our discussion of the new invariant mass variables and their kinematic endpoints. For our basic proof-of-principle measurement technique presented in the next

section 3.1, we shall use only three of them, namely $M_{j\ell(u)}^{\max}$, $m_{j\ell(u)}^{\max}$, and $m_{j\ell(s)}^{\max}(\alpha = 1)$. However, the remaining variables are in principle just as good, their only disadvantage being that they failed our arbitrarily imposed condition at the beginning that the endpoint functions should all be region independent. Of course, one could, and in fact should, use all of the available kinematic endpoint information, which in a global fit analysis can only increase the experimental precision of the sparticle mass determination.

3 Theoretical analysis

3.1 Our method and the solution for the mass spectrum

Our starting point is the set of four measurements

$$m_{\ell\ell}^{\max}, M_{j\ell(u)}^{\max}, m_{j\ell(u)}^{\max}, m_{j\ell(s)}^{\max}(\alpha = 1) \tag{3.1}$$

in place of the conventional set (1.18). It is easy to verify that the measurements (3.1) are always independent of each other, and thus never suffer from the linear dependence problem discussed in section 1.2.

Given the set of four measurements (3.1), it is easy to solve for the mass spectrum. To simplify the notation, we introduce the following shorthand notation for the endpoints of the mass *squared* distributions

$$L \equiv (m_{\ell\ell}^{\max})^2, \quad M \equiv (M_{j\ell(u)}^{\max})^2, \quad m \equiv (m_{j\ell(u)}^{\max})^2, \quad S \equiv (m_{j\ell(s)}^{\max}(\alpha = 1))^2 \tag{3.2}$$

The solution for the mass spectrum is then given by

$$m_D^2 = \frac{Mm(L + M + m - S)}{(M + m - S)^2}; \tag{3.3}$$

$$m_C^2 = \frac{MmL}{(M + m - S)^2}; \tag{3.4}$$

$$m_B^2 = \begin{cases} \frac{ML(S-M)}{(M+m-S)^2}, & \text{if } R_{AB} \geq R_{BC}, \\ \frac{mL(S-m)}{(M+m-S)^2}, & \text{if } R_{AB} \leq R_{BC}; \end{cases} \tag{3.5}$$

$$m_A^2 = \frac{L(S - m)(S - M)}{(M + m - S)^2}. \tag{3.6}$$

It is easy to verify that the right-hand side expressions in these equations are always positive definite, so that one can safely take the square root and compute the linear masses m_D , m_C , m_B and m_A . Notice that in spite of the two-fold ambiguity (2.4), (2.5), the solution for m_D , m_C and m_A is unique! Indeed, the expressions for m_D , m_C and m_A are symmetric under the interchange $M \leftrightarrow m$. The remaining two-fold ambiguity for m_B is precisely the result of the ambiguous interpretation (2.4), (2.5) of the two $m_{j\ell(u)}^2$ endpoints, and is related to the symmetry under (1.13), or equivalently, under the interchange

$$R_{AB} \leftrightarrow R_{BC} . \tag{3.7}$$

In the next subsection we discuss several ways in which one can lift the remaining two-fold degeneracy for m_B which is due to (3.7).

Notice the great simplicity of this method. The expressions for (3.3), (3.4) and (3.6) are region independent and therefore one does not have to go through the standard trial and error procedure involving the 9 parameter space regions $(N_{j\ell\ell}, N_{j\ell})$ [12, 47] associated with the various interpretations of the endpoints $m_{j\ell\ell}^{\max}$, $m_{j\ell(lo)}^{\max}$ and $m_{j\ell(hi)}^{\max}$.

3.2 Disambiguation of the two solutions for m_B

The method outlined in section 3.1 allowed us to find the true masses of particles A , C and D , but yields two separate possible solutions for the mass m_B of particle B . We shall now discuss several ways of lifting the remaining two-fold degeneracy for m_B .

3.2.1 Invariant mass endpoint method

One possibility is to use an additional measurement of an invariant mass endpoint. Indeed, as shown in sections 1 and 2, there are still quite a few one-dimensional invariant mass distributions at our disposal, which we have not used so far. Those include the conventional distributions of $m_{j\ell\ell}^2$, $m_{j\ell(lo)}^2$ and $m_{j\ell(hi)}^2$, as well as the new distributions $m_{j\ell(p)}^2$, $m_{j\ell(s)}^2(\alpha)$ and $m_{j\ell(d)}^2(1)$ which we introduced in section 2. Which of them can be used for our purposes? Note that the duplication in (3.5) arose due to the symmetry (3.7), so that any kinematic endpoint which violates this symmetry will be able to distinguish between the two solutions.

Let us begin with the conventional distributions $m_{j\ell\ell}^2$, $m_{j\ell(lo)}^2$, $m_{j\ell(hi)}^2$ and $m_{j\ell(\theta>\frac{\pi}{2})}^2$, whose endpoints we did not use in our analysis so far. It is easy to check that $m_{j\ell\ell}^{\max}$, $m_{j\ell(hi)}^{\max}$ and $m_{j\ell(\theta>\frac{\pi}{2})}^{\min}$ are invariant under the interchange (3.7) and cannot be used for discrimination. However, $m_{j\ell(lo)}^{\max}$ is *not* symmetric under (3.7) and can do the job. In fact, one can show that the two duplicate solutions for m_B always⁷ give different predictions for $m_{j\ell(lo)}^{\max}$.

More importantly, many of our new variables from section 2 can provide an independent cross-check on the correct choice for the solution. For example, the kinematic endpoint (2.7) of the product variable $m_{j\ell(p)}^2$, also violates the symmetry (3.7) and distinguishes among the two solutions. The infinite set of variables $m_{j\ell(s)}^2(\alpha)$ can also be used, and for almost the whole range of $\alpha < 1$. To see this, in figure 2 we compare the predictions for the kinematic endpoints $m_{j\ell(s)}^{\max}(\alpha)$ of the real and fake solutions, for the two examples discussed in detail in section 4: (a) the LM1 CMS study point and (b) the LM6 CMS study point. The corresponding mass spectra are listed in table 1 below. For convenience, we plot versus the parameter

$$\phi \equiv \arctan \alpha, \tag{3.8}$$

which allows us to map the whole definition domain $(-\infty, \infty)$ for α into the finite region $(-\frac{\pi}{2}, \frac{\pi}{2})$ for ϕ . Figure 2 shows that for most of the allowed ϕ range, the two solutions predict different values for the kinematic endpoints $m_{j\ell(s)}^{\max}(\alpha)$. In fact, for $\phi < \frac{\pi}{4}$, the two

⁷The only exception is the trivial case of $R_{AB} = R_{BC}$, but then the two solutions for m_B coincide, and m_B is again uniquely determined.

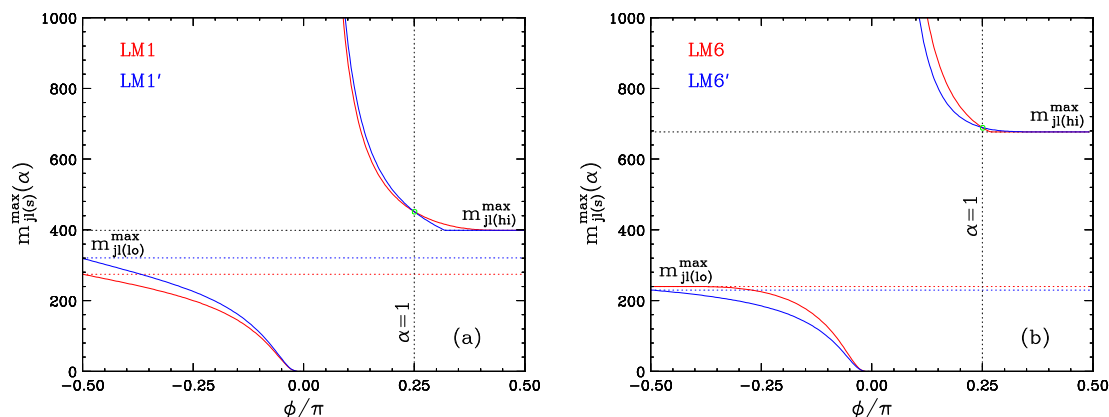


Figure 2. Comparison of the predictions for the kinematic endpoints $m_{j\ell(s)}^{\max}(\alpha)$ of the real and fake solutions, as a function of $\phi \equiv \arctan \alpha$ (in units of π), for the two examples discussed in detail in section 4: (a) the LM1 CMS study point and (b) the LM6 CMS study point. In each panel, the prediction of the real (fake) solution is plotted in red (blue). The vertical dotted line indicates the case of $\phi = \frac{\pi}{4}$ ($\alpha = 1$), for which the two solutions give an identical answer, marked with a green dot. The horizontal dotted lines show the corresponding asymptotic values $m_{j\ell(hi)}^{\max}$ and $m_{j\ell(lo)}^{\max}$, obtained at $\alpha \rightarrow \pm\infty$ ($\phi \rightarrow \pm\frac{\pi}{2}$).

predictions are always different, apart from the trivial case of $\phi = 0$ ($\alpha = 0$). Even for $\phi > \frac{\pi}{4}$, there still exists a range of ϕ , for which, at least theoretically, a discrimination can be made. The predictions are guaranteed to coincide only for $\phi = \frac{\pi}{4}$ ($\alpha = 1$) (as they should, see (3.1)), and for a certain range of the largest possible values of ϕ .

3.2.2 Invariant mass correlations

Another way to resolve the twofold ambiguity in our solution (3.5) is to simply go back to the original measurements of $M_{j\ell(u)}^{\max}$ and $m_{j\ell(u)}^{\max}$ and already at that point try to decide which of the two measured $m_{j\ell(u)}$ endpoints is $m_{j\ell_n}^{\max}$ and which one is $m_{j\ell_f}^{\max}$. As already discussed in [18, 46], this identification is in principle possible, if one considers the correlations which are present in the two-dimensional distribution $m_{j\ell(u)}^2$ versus $m_{\ell\ell}^2$. The basic idea is illustrated in figure 3, where we show scatter plots of $m_{j\ell(u)}$ versus $m_{\ell\ell}$, for the two examples used in figure 2 and discussed in detail later in section 4. Figure 3(a) (figure 3(b)) shows the result for the real (fake) solution corresponding to the LM1 study point, while figures 3(c) and 3(d) show the analogous results for the LM6 study point. In each plot we used 10,000 entries, which roughly corresponds to 20 fb⁻¹ (200 fb⁻¹) of data for the actual LM1 (LM6) SUSY study point. Here and below we show the ideal case where we neglect smearing effects due to the finite detector resolution, finite particle widths and combinatorial backgrounds. All of our plots are at the parton level (using our own Monte-Carlo phase space generator) and without any cuts. Notice that in order to avoid dealing with the large numerical values of the squared masses, we use a quadratic power scale on both axes, which allows us to preserve the simple shapes of the scatter plots when plotting versus the linear masses themselves.

Figure 3 shows that the combined distribution $m_{j\ell(u)}^2$ is simply composed of the two

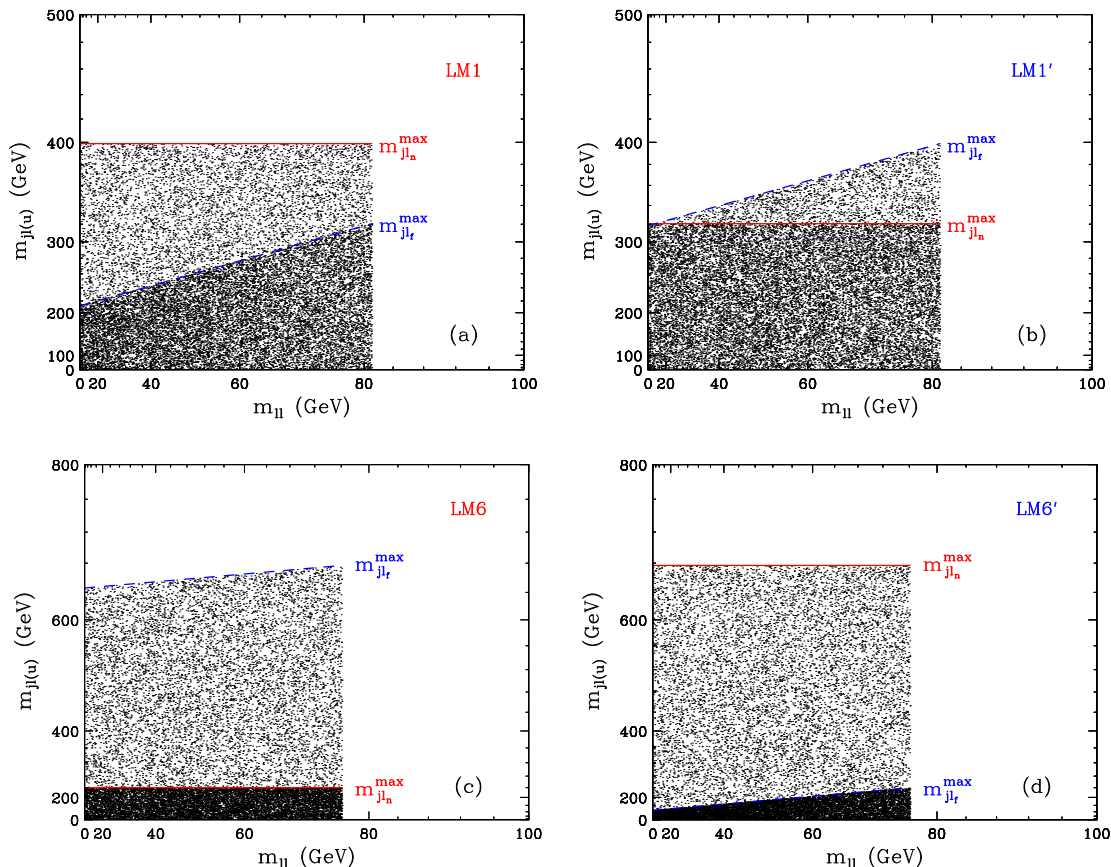


Figure 3. Predicted scatter plots of $m_{j\ell(u)}$ versus $m_{\ell\ell}$, for the case of the real and fake solutions for each of the two study points LM1 and LM6: (a) the real solution LM1; (b) the fake solution LM1'; (c) the real solution LM6; and (d) the fake solution LM6'. The red solid horizontal (blue dashed inclined) line indicates the conditional maximum $m_{j\ell_n}^{\max}(m_{\ell\ell})$ ($m_{j\ell_f}^{\max}(m_{\ell\ell})$) given by eq. (3.9) (eq. (3.10)). Each panel contains 10,000 entries. The results shown here are idealized in the sense that we neglect smearing effects due to the finite detector resolution, finite particle widths and combinatorial backgrounds. Notice the use of quadratic power scale on the two axes, which preserves the simple shapes of the scatter plots, even when plotted versus the linear masses $m_{j\ell(u)}$ and $m_{\ell\ell}$.

separate distributions $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$, but they are correlated *differently* with the dilepton distribution $m_{\ell\ell}^2$. In particular, let us concentrate on the conditional maxima $m_{j\ell_n}^{\max}(m_{\ell\ell})$ and $m_{j\ell_f}^{\max}(m_{\ell\ell})$, i.e. the maximum allowed values of $m_{j\ell_n}$ and $m_{j\ell_f}$, respectively, for a given fixed value of $m_{\ell\ell}$ [18, 46]. A close inspection of figure 3 shows that the values of $m_{j\ell_n}^2$ and $m_{\ell\ell}^2$ are uncorrelated, and as a result, the conditional maximum $m_{j\ell_n}^{\max}(m_{\ell\ell})$ does not depend on $m_{\ell\ell}$. In turn, this implies that the endpoint value $(m_{j\ell_n}^{\max})^2$ given in (1.3) can be obtained for any $m_{\ell\ell}^2$:

$$n \equiv (m_{j\ell_n}^{\max})^2 = [m_{j\ell_n}^{\max}(m_{\ell\ell})]^2 = m_D^2 (1 - R_{CD}) (1 - R_{BC}), \quad \forall m_{\ell\ell} \in [0, m_{\ell\ell}^{\max}] . \quad (3.9)$$

Because of (3.9), the shape of the $m_{j\ell_n}^2$ versus $m_{\ell\ell}^2$ scatter plot is a simple rectangle [18, 46]. This is confirmed by the plots in figure 3, where the (red) horizontal solid line indicates the constant value (3.9) for the conditional maximum $m_{j\ell_n}^{\max}(m_{\ell\ell})$.

In contrast, the values of $m_{j\ell_f}^2$ and $m_{\ell\ell}^2$ are correlated. The conditional maximum $m_{j\ell_f}^{\max}(m_{\ell\ell})$ does depend on the value of $m_{\ell\ell}$ as follows:

$$\left(m_{j\ell_f}^{\max}(m_{\ell\ell})\right)^2 = p + \frac{f-p}{L}m_{\ell\ell}^2, \tag{3.10}$$

where we introduce the shorthand notation used in [47]

$$f \equiv \left(m_{j\ell_f}^{\max}\right)^2 = m_D^2(1-R_{CD})(1-R_{AB}), \tag{3.11}$$

$$p \equiv R_{BC}f = m_D^2(1-R_{CD})R_{BC}(1-R_{AB}). \tag{3.12}$$

The absolute maximum of $m_{j\ell_f}^2$, which is given by (1.4) and denoted here by f , can only be obtained when $m_{\ell\ell}^2$ itself is at a maximum [18, 46]:

$$f \equiv \left[m_{j\ell_f}^{\max}(m_{\ell\ell}^{\max})\right]^2. \tag{3.13}$$

On the other hand, the conditional maximum $m_{j\ell_f}^{\max}(m_{\ell\ell})$ obtains its minimum value at $m_{\ell\ell}^2 = 0$ and corresponds to [18, 46]

$$p \equiv \left[m_{j\ell_f}^{\max}(0)\right]^2 \leq f. \tag{3.14}$$

Eqs. (3.13), (3.14) imply that the shape of the $m_{j\ell_f}^2$ versus $m_{\ell\ell}^2$ scatter plot is a right-angle trapezoid. This is confirmed by the plots in figure 3, where we mark with a (blue) dashed line the conditional maximum (3.10). With sufficient statistics, this difference in the kinematic boundaries may be observable, and would reveal the identity of $m_{j\ell_n}^{\max}$ and $m_{j\ell_f}^{\max}$ [18, 46]. Once the individual $m_{j\ell_n}^{\max}$ and $m_{j\ell_f}^{\max}$ are known, the solution for the mass spectrum is unique – see e.g. appendix A in [47]. Of course, in cases where $p \sim f$, namely $R_{BC} \sim 1$, it may be difficult in practice to tell which of the two boundaries in the scatter plot is inclined and which one is horizontal.⁸ One example of this sort is offered by point LM6, which has $R_{BC} = 0.91$ and leads to a rather flat $m_{j\ell_f}^{\max}(m_{\ell\ell})$ function, as seen in figure 3(c).

An alternative and somewhat related method will be to investigate the shapes of the one-dimensional distributions themselves [85]. In appendix A we provide the analytical expressions for the shapes of the four invariant mass distributions $m_{\ell\ell}^2$, $m_{j\ell(u)}^2$, $m_{j\ell(s)}^2(1)$ and $m_{j\ell(d)}^2(1)$ used in our basic analysis from section 3.1. Given what we have already seen in figure 3, it is not surprising that the true and the fake solutions predict different shapes for the one-dimensional distributions as well. In the LM1 and LM6 examples considered below in section 4, this difference is particularly noticeable for the $m_{j\ell(u)}^2$ and $m_{j\ell(d)}^2(1)$ distributions (see figures 4(b), 4(d), 5(b) and 5(d)), and can be tested experimentally.

3.2.3 M_{T2} endpoint method

Let us note that if we identify particle A with the LSP, we have a rather peculiar situation, in which we *know* the LSP mass m_A , and we are unsure about the NLSP mass M_B , for

⁸A separate problem, which arises in the case of $p \sim f$, will be discussed below in section 4.1.

which we have to choose among two alternatives. This goes against the common lore which considers the LSP mass (in this case m_A) to be the least constrained among the masses appearing in the decay chain in figure 1. For example, the method of the Cambridge M_{T2} variable [2, 8] treats the LSP mass as a continuous unknown parameter. At this point of our analysis we already know the LSP mass, and we can use this knowledge to our advantage. For example, if we can collect a sufficient number of events of B pair-production, we can apply the idea of M_{T2} for the $B \rightarrow A$ decay as in the original M_{T2} proposal [2]. When we use for the trial LSP mass the known true value of m_A given by (3.6), the kinematic endpoint of the M_{T2} distribution will reveal the correct value of the mass m_B of the parent particle B , thus selecting the true solution in (3.5).

As emphasized in ref. [43], the M_{T2} endpoint method does not necessarily rely on A being the LSP (i.e. the very last particle in the decay chain) or B being the “grandparent” (i.e. the very first particle in the decay chain). For example, suppose that A decays further. In that case, one simply needs to apply the more general “subsystem” variable $M_{T2}^{(n,p,c)}$ [43] with A being the “child” particle: $c = A$. Similarly, the two B particles do not have to be the two grandparents initiating the decay chains: it is sufficient to consider $M_{T2}^{(n,p,c)}$ with $p = B$ and arbitrary n [43]. Finally, for the purposes of selecting the correct solution in (3.5) it is also possible to apply the subsystem variable $M_{T2}^{(n,p,c)}$ in a different way, where B is the child, and the parent is either D or C . In this case, we know the parent mass, which is respectively given by (3.3) or (3.4), and we are asking the question, which of the two test masses in (3.5) gives the correct answer for the M_{T2} endpoint.

4 Numerical examples

We shall now illustrate the ideas of the previous section with two specific numerical examples: the LM1 and LM6 SUSY study points in CMS [83]. The mass spectra at LM1 and LM6 are listed in table 1. Point LM1 is similar to benchmark point A (A') in ref. [86] (ref. [87]) and to benchmark point SPS1a in ref. [88]. Point LM6 is similar to benchmark point C (C') in ref. [86] (ref. [87]). The table also lists the corresponding duplicate solutions LM1' and LM6', which are obtained by interchanging $R_{BC} \leftrightarrow R_{AB}$, or equivalently, by replacing the mass of B via

$$m_B \rightarrow m'_B = \frac{m_A m_C}{m_B}. \tag{4.1}$$

It is interesting to note that LM1 and LM6 represent both sides of the ambiguity (3.7): at LM1, we have $R_{AB} > R_{BC}$ and correspondingly, $m_{j\ell_n}^{\max} > m_{j\ell_f}^{\max}$ and (2.4) applies. On the other hand, at LM6 we have $R_{AB} < R_{BC}$ and $m_{j\ell_n}^{\max} < m_{j\ell_f}^{\max}$, so that (2.5) applies. Another interesting difference is that at LM1 particle B is the *right-handed* slepton $\tilde{\ell}_R$, while at LM6 the role of particle B is played⁹ by the *left-handed* slepton $\tilde{\ell}_L$. Of course, to the extent that we are interested in kinematical features, this difference is not relevant, and particle B of the LM6 spectrum may very well have been the right-handed slepton instead.

⁹Although the right-handed slepton $\tilde{\ell}_R$ is also kinematically accessible at point LM6, the wino-like neutralino $\tilde{\chi}_2^0$ decays much more often to $\tilde{\ell}_L$ as opposed to $\tilde{\ell}_R$.

Variable	LM1	LM1'	LM6	LM6'
m_A (GeV)	94.9		158.15	
m_B (GeV)	118.9	143.35	291.0	165.65
m_C (GeV)	179.6		304.8	
m_D (GeV)	561.6		861.9	
R_{AB}	0.6370	0.4383	0.2954	0.9115
R_{BC}	0.4383	0.6370	0.9115	0.2954
R_{CD}	0.1023		0.1251	
$m_{\ell\ell}^{\max}$ (GeV)	81.10		76.12	
$M_{j\ell(u)}^{\max}$ (GeV)	398.8		676.8	
$m_{j\ell(u)}^{\max}$ (GeV)	320.6		239.8	
$m_{j\ell(s)}^{\max}(\alpha = 1)$ (GeV)	451.8		689.2	
$m_{j\ell}^{\max}$ (GeV)	451.8		689.2	
$m_{j\ell(\theta > \frac{\pi}{2})}^{\min}$ (GeV)	215.2		176.4	
$m_{j\ell(hi)}^{\max}$ (GeV)	398.8		676.8	
$m_{j\ell(s)}^{\max}(\alpha = 2)$ (GeV)	406.6	398.8	676.8	677.0
$m_{j\ell(s)}^{\max}(\alpha = 1.5)$ (GeV)	417.9	402.5	676.8	678.4
$m_{j\ell(s)}^{\max}(\alpha = 0.5)$ (GeV)	611.0	638.9	886.0	807.1
$m_{j\ell(s)}^{\max}(\alpha = -0.5)$ (GeV)	142.9	159.7	174.9	138.0
$m_{j\ell(s)}^{\max}(\alpha = -1)$ (GeV)	200.1	225.9	224.8	184.8
$m_{j\ell(lo)}^{\max}$ (GeV)	274.6	319.1	239.8	229.9
$m_{j\ell(p)}^{\max}$ (GeV)	292.0	319.4	393.7	310.9
$m_{j\ell_n}^{\max}$ (GeV)	398.8	320.6	239.8	676.8
$m_{j\ell_f}^{\max}$ (GeV)	320.6	398.8	676.8	239.8

Table 1. The relevant part of the SUSY mass spectrum for the LM1 and LM6 study points. The corresponding duplicated solutions LM1' and LM6' are obtained by interchanging $R_{BC} \leftrightarrow R_{AB}$ as in (3.7). In the table we also list the corresponding values for various invariant mass endpoints. The first four of those represent our basic set of measurements (3.1) discussed in detail in section 4.1, while the last two ($m_{j\ell_n}^{\max}$ and $m_{j\ell_f}^{\max}$) are not directly observable. The remaining invariant mass endpoints are considered in section 4.2. In the case of $m_{j\ell(s)}^{\max}(\alpha)$, we show several representative values for α . For the complete α variation, refer to figure 2. Recall that $m_{j\ell(s)}^{\max}(+\infty) = m_{j\ell(hi)}^{\max}$ and $m_{j\ell(s)}^{\max}(-\infty) = m_{j\ell(lo)}^{\max}$.

4.1 Mass measurements at points LM1 and LM6

Given the mass spectra in table 1, it is straightforward to construct and investigate the relevant invariant mass distributions. For the purposes of illustration, we shall ignore spin correlations, referring the readers interested in those effects to refs. [52, 58, 71]. We are

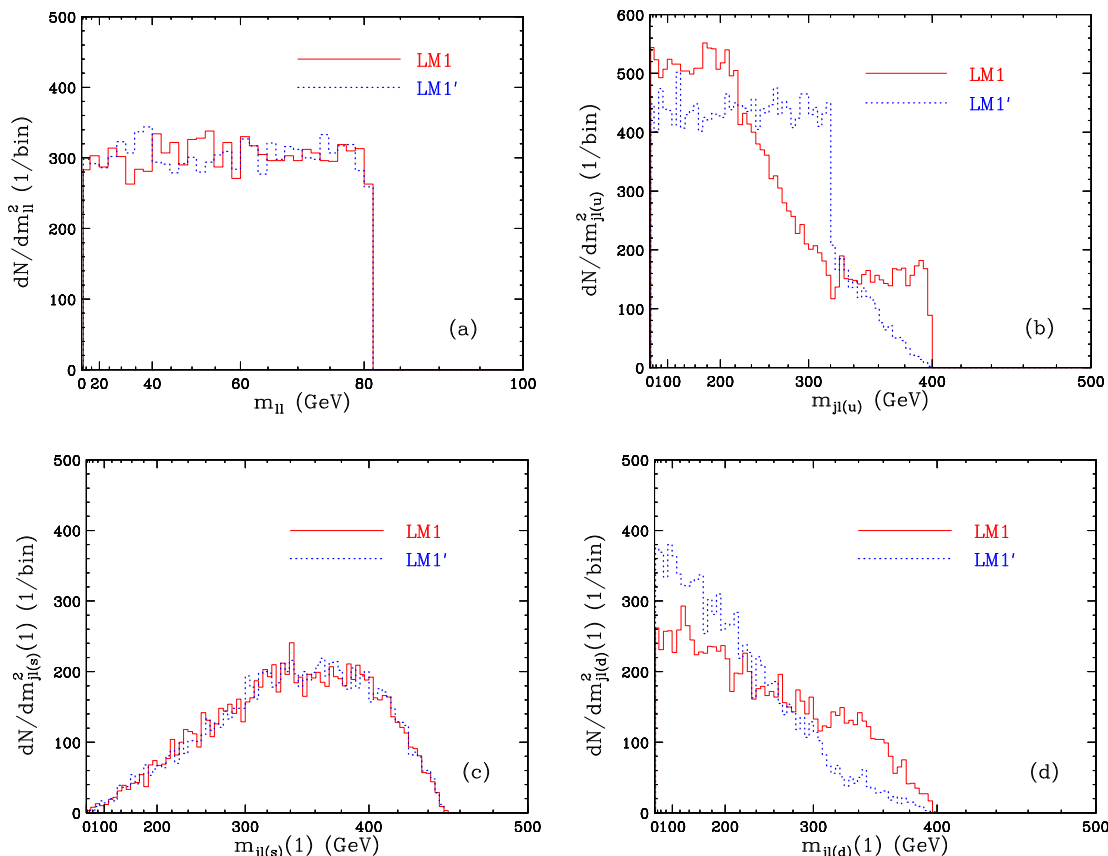


Figure 4. One-dimensional invariant mass distributions for the case of LM1 (red solid lines) and LM1' (blue dotted lines) spectra. The kinematic endpoints (3.1) used in our analysis in section 3.1 can be observed from these distributions as follows: $m_{\ell\ell}^{\max}$ is the upper kinematic endpoint of the $m_{\ell\ell}$ distribution in panel (a); $M_{j\ell(u)}^{\max}$ is the absolute upper kinematic endpoint seen in both the combined $m_{j\ell(u)}$ distribution in panel (b), or the difference distribution $m_{j\ell(d)}(1)$ in panel (d); $m_{j\ell(u)}^{\max}$ is the intermediate kinematic endpoint seen in panel (b); and $m_{j\ell(s)}^{\max}(\alpha = 1)$ is the upper kinematic endpoint of the $m_{j\ell(s)}(\alpha = 1)$ distribution in panel (c).

justified to do so for several reasons. First, our method relies on the measurement of kinematic endpoints, whose location is unaffected by the presence of spin correlations. Second, in the case of supersymmetry (which is really what we have in mind here), particle B is a scalar, which automatically washes out any spin effects in the $m_{\ell\ell}^2$ and $m_{j\ell_f}^2$ distributions. Furthermore, if particles D and their antiparticles \bar{D} are produced in equal numbers, as would be the case if the dominant production is from gg and/or $q\bar{q}$ initial state, any spin correlations in the $m_{j\ell_n}^2$ distribution are also washed out. Under those circumstances, therefore, the pure phase space distributions shown here are in fact the correct answer.

We begin our discussion with the four invariant mass distributions $m_{\ell\ell}^2$, $m_{j\ell(u)}^2$, $m_{j\ell(s)}^2(\alpha = 1)$ and $m_{j\ell(d)}^2(\alpha = 1)$, which form the basis of our method outlined in section 3.1. Figure 4 (figure 5) shows those four distributions for the case of study point LM1 (LM6). In each panel, the red (solid) histogram corresponds to the nominal spectrum (LM1 or LM6), while the blue (dotted) histogram corresponds to the “fake” solution

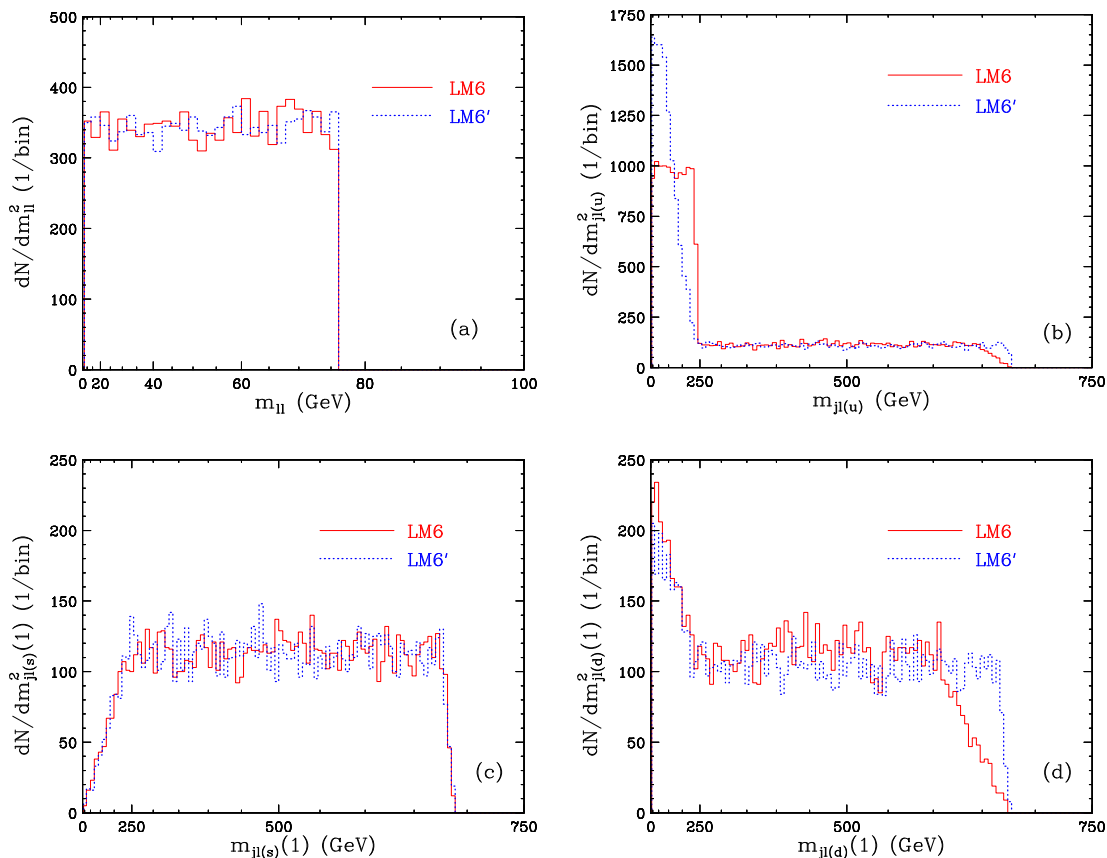


Figure 5. The same as figure 4, but for the LM6 mass spectrum (red solid lines) and the LM6' mass spectrum (blue dotted lines).

(LM1' or LM6'), which is obtained through the replacement (4.1). For all figures in this section, we use the same 4 samples of 10,000 events each, which were already used to make figure 3. Notice our somewhat unconventional way of filling and then plotting the histograms in this section. First, we show differential distributions in the corresponding mass squared, i.e. dN/dm^2 . This is done in order to preserve the connection to the analytical results in appendix A, which are written the same way. More importantly, the shapes of the one-dimensional histograms are much simpler in the case of dN/dm^2 as opposed to dN/dm [52, 58, 71]. In the next step, however, we choose to plot the thus obtained histogram versus the mass itself rather than the mass squared. This allows one to read off immediately the corresponding endpoint and compare directly to the values listed in table 1. It also keeps the x -axis range within a manageable range. However, since the histograms were binned on a mass squared scale, if we were to use a linear scale on the x -axis, we would get bins with varying size. This would be rather inconvenient and more importantly, would distort the nice simple shapes of the dN/dm^2 distributions. Therefore, we use a quadratic scale on the x -axis, which preserves the nice shapes and leads to a constant bin size on each plot.

Figures 4 and 5 illustrate how each one of the measurements (3.1) can be obtained.

For example, $m_{\ell\ell}^{\max}$ is the classic upper kinematic endpoint of the $m_{\ell\ell}$ distributions in figures 4(a) and 5(a). This endpoint is very sharp and should be easily observable. $M_{j\ell(u)}^{\max}$ is the absolute upper kinematic endpoint seen in the combined $m_{j\ell(u)}$ distribution in figures 4(b) and 5(b). Notice that the same endpoint can independently also be observed as the absolute upper kinematic limit of the difference distributions $m_{j\ell(d)}(1)$ shown in figures 4(d) and 5(d). The fact that there are two independent ways of getting to the endpoint $M_{j\ell(u)}^{\max}$ should allow for a reasonable accuracy of its measurement. Upon closer inspection of the combined $m_{j\ell(u)}$ distribution in figures 4(b) and 5(b), we also notice the intermediate kinematic endpoint $m_{j\ell(u)}^{\max}$ seen around 320 GeV in figure 4(b) and around 240 GeV in figure 5(b). Finally, $m_{j\ell(s)}^{\max}(\alpha = 1)$ is the upper kinematic endpoint of the $m_{j\ell(s)}(\alpha = 1)$ distribution shown in figures 4(c) and 5(c). It is also rather well defined, and should be well measured in the real data.

At this point we would like to comment on one potential problem which is not immediately obvious, but nevertheless has been encountered in practical applications of the invariant mass technique for SUSY mass determinations [85]. It has been noted that in the case of $p \sim f$ (see eqs. (3.11), (3.12)), the numerical fit for the mass spectrum becomes rather unstable. Given our analytical results in section 3.1, we are now able to trace the root of the problem. Notice that $p \sim f$ implies that $R_{BC} \sim 1$. In this limit, from eqs. (1.2), (1.3), (1.4) and (2.16) we find

$$\lim_{R_{BC} \rightarrow 1} (L) = 0, \quad \lim_{R_{BC} \rightarrow 1} (n) = 0, \quad \lim_{R_{BC} \rightarrow 1} (M + m - S) = 0. \quad (4.2)$$

This means that the functions (3.3)–(3.6) giving the solution for the mass spectrum will all behave as $\frac{0^2}{0^2}$, and, given the statistical fluctuations in an actual analysis, will have very poor convergence properties. We note that this problem is not limited to our preferred set of measurements (3.1) and is rather generic, but has been missed in most previous studies simply because the case of $R_{BC} \sim 1$ was rarely considered.

figures 4 and 5 reveal that, as expected, the real (red solid lines) and fake (blue dotted lines) solutions always give identical results for our basic set of four endpoint measurements (3.1). This is by design, and in order to discriminate among the real and the fake solution, we need additional experimental input, as discussed in section 3.2. Before we proceed with the disambiguation analysis in the next subsection, we should stress once again that the real and fake solutions agree on 75% of the relevant mass spectrum, i.e. they give the same values for the masses of particles D , C and A (see table 1). The only question mark at this point is, what is the mass of particle B . This issue is addressed in the following subsection.

4.2 Eliminating the fake solution for m_B

As already discussed in section 3.2, there are several handles which could discriminate among the two alternative values of m_B in the real and the fake solution. One possibility, reviewed in section 3.2.3, is to use additional independent measurements of M_{T2} kinematic endpoints. We shall not pursue this direction here, referring the interested readers to ref. [43] for details. Another possibility, discussed in section 3.2.2 and demonstrated

explicitly with figure 3, is to use the different correlations in the 2-dimensional invariant mass distributions $(m_{\ell\ell}^2, m_{j\ell_n}^2)$ and $(m_{\ell\ell}^2, m_{j\ell_f}^2)$. The near-far lepton ambiguity is avoided by studying the scatter plot of $(m_{\ell\ell}^2, m_{j\ell(u)}^2)$, shown in figure 3, which should be in principle sufficient to discriminate among the two alternatives.

In keeping with the main theme of this paper, in this subsection we shall concentrate on the third possibility, already suggested in section 3.2.1. We shall simply explore additional invariant mass endpoint measurements, which would hopefully discriminate among the two solutions for m_B . Figures 6 and 7 show several invariant mass distributions which have already been mentioned at one point or another in the course of our previous discussion. Figure 6 shows the following 6 distributions: (a) $m_{j\ell}^2$; (b) $m_{j\ell(hi)}^2$; (c) $m_{j\ell(p)}^2$; (d) $m_{j\ell(lo)}^2$; (e) $m_{j\ell(s)}^2(\alpha = -1)$ and (f) $m_{j\ell(s)}^2(\alpha = \frac{1}{2})$, for the LM1 mass spectrum (red solid lines) and its LM1' counterpart (blue dotted lines). Figure 7 shows the same 6 distributions, but for the LM6 and LM6' mass spectra. In both figures, we follow the same plotting conventions as in figures 4 and 5: we form the mass squared distribution dN/dm^2 , and then plot versus the corresponding linear mass m using a quadratic scale on the x -axis. Notice that the sum of the $m_{j\ell(hi)}^2$ distribution in figure 6(b) (figure 7(b)) and the $m_{j\ell(lo)}^2$ distribution in figure 6(d) (figure 7(d)) precisely equals the combined distribution $m_{j\ell(u)}^2$ in figure 4(b) (figure 5(b)). In order to be able to see this by the naked eye, we have kept the same x and y ranges on the corresponding plots.

As seen in figures 6 and 7, not all of the remaining invariant mass distributions are able to discriminate among the two m_B solutions. As explained in section 3.2.1, the suitable distributions are those whose endpoints violate the symmetry (3.7), which caused the m_B ambiguity in the first place. For example, figures 6(a) and 7(a) show that the endpoint of the $m_{j\ell\ell}^2$ distribution is the same for the real and the fake solution. This is to be expected, since the defining expression (1.5) for $m_{j\ell\ell}^{\max}$ is symmetric under (3.7). Figures 6(a) and 7(a) also show that even the *shapes* of the $m_{j\ell\ell}^2$ distributions for the real and fake solution are very similar. In spite of this, the observation of the $m_{j\ell\ell}^2$ endpoint can still be very useful, e.g. in reducing the experimental error on the mass determination.

Similar comments apply to the $m_{j\ell(hi)}^2$ distributions shown in figures 6(b) and 7(b). Here again the endpoint is a symmetric function of R_{AB} and R_{BC} , and the real and fake solutions predict identical endpoints. However, while the endpoints are the same, this time the shapes are not. The shape difference is more pronounced in the case of LM1 shown in figure 6(b), and less visible in the case of LM6 shown in figure 7(b).

The remaining four distributions shown in figures 6(c-f) and 7(c-f) already have different endpoints and can thus be used for discrimination among the real and fake solution for m_B . All of the endpoints in figures 6(c-f) and 7(c-f) are relatively sharp and should be measured rather well. One should not forget that in figures 6 and 7 we show $m_{j\ell(s)}^2(\alpha)$ distributions for only three representative values of α : $\alpha = -\infty$ in panels (d), $\alpha = -1$ in panels (e), and $\alpha = 0.5$ in panels (f). As seen in figure 2, there are infinitely many other choices for α , which would still exhibit different endpoints for the real and fake m_B solutions. Our conclusion is that through a suitable combination of additional endpoint measurements one would be able to tell apart the real solution for m_B from its fake cousin.

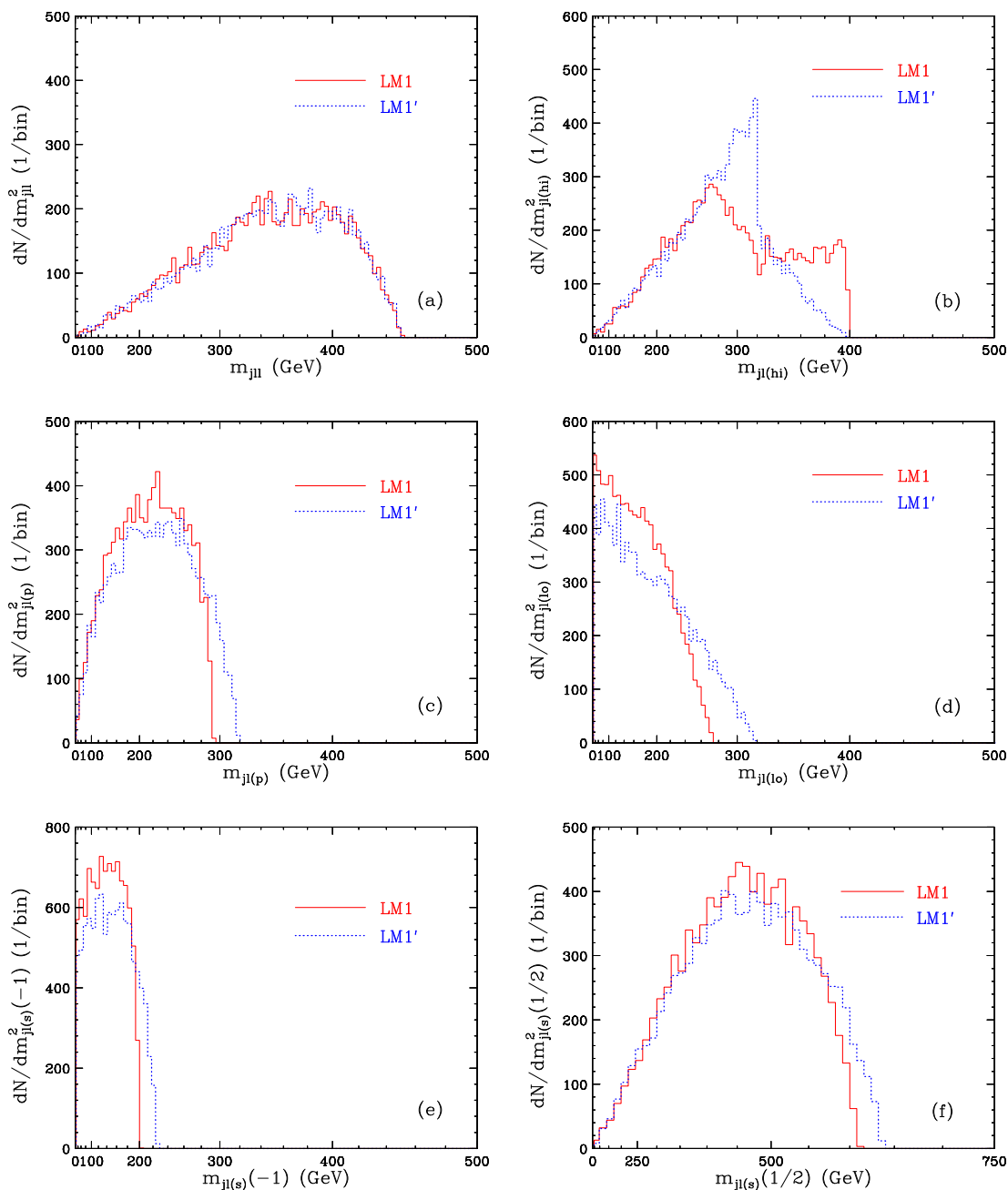


Figure 6. Some other one-dimensional invariant mass distributions of interest, for the case of the LM1 mass spectrum (red solid lines) and LM1' mass spectrum (blue dotted lines): (a) $m_{j\ell}^2$ distribution; (b) $m_{j\ell(hi)}^2$ distribution; (c) $m_{j\ell(p)}^2$ distribution; (d) $m_{j\ell(lo)}^2$ distribution; (e) $m_{j\ell(s)}^2(\alpha = -1)$ distribution; (f) $m_{j\ell(s)}^2(\alpha = \frac{1}{2})$ distribution. All distributions are then plotted versus the corresponding mass, on a quadratic scale for the x -axis.

5 Summary and conclusions

In this paper we revisited the classic technique for SUSY mass determinations via invariant mass endpoints. We set out to redesign the standard algorithm for performing these studies,

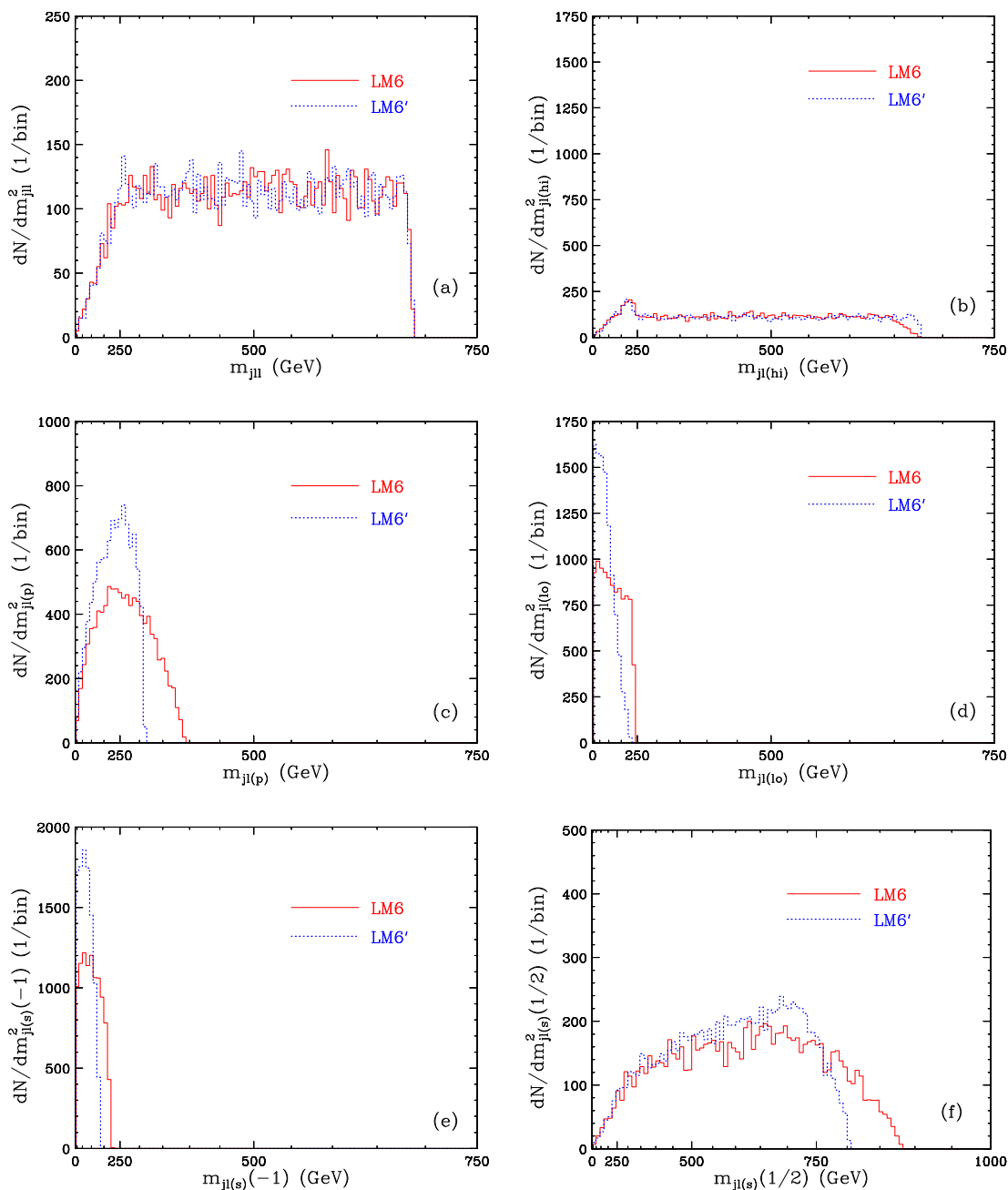


Figure 7. The same as figure 6, but for the LM6 mass spectrum (red solid lines) and the LM6' mass spectrum (blue dotted lines).

by pursuing two main objectives (see section 1.4):

- *Improving on the experimental precision of the SUSY mass determination.* For example, we required that our analysis be based exclusively on *upper* invariant mass endpoints, which are expected to be measured with a greater precision than the corresponding lower endpoints (a.k.a. thresholds). Consequently, we did not make use of the “threshold” measurement $m_{j\ell\ell}^{\min}(\theta > \frac{\pi}{2})$, which has been an integral part of most

SUSY studies since ref. [7]. In the same vein, we also demanded that we should not rely on any features observed in a two- or a three-dimensional invariant mass distribution — such measurements are expected to be less precise than the (upper) endpoints extracted from simple one-dimensional histograms.

- *Avoiding any parameter space region ambiguities.* It is well known that some of the invariant mass endpoints used in the conventional analyses are piecewise-defined functions. This feature may sometimes lead to multiple solutions for the SUSY mass spectrum in the “LHC inverse problem” [12, 47, 78–80]. In order to safeguard against this possibility, we conservatively demanded from the outset that none of our endpoint measurements be given by piecewise defined functions. This rather strict requirement rules out three of the standard endpoint measurements $m_{j\ell\ell}^{\max}$, $m_{j\ell(lo)}^{\max}$, and $m_{j\ell(hi)}^{\max}$.

In order to meet these objectives, in section 2 we proposed a set of new invariant mass variables whose upper kinematic endpoints can be alternatively used for SUSY mass reconstruction studies. Then in section 3 we outlined a simple analysis which was based on the particular set of four invariant mass variables (3.1), all of which satisfy our requirements. In section 3.1 we provided simple analytical formulas for the SUSY mass spectrum in terms of the four measured endpoints in eq. (3.1). Our solutions revealed a surprise: in spite of the two-fold ambiguity (2.4), (2.5) in the interpretation of two of our endpoints $M_{j\ell(u)}^{\max}$ and $m_{j\ell(u)}^{\max}$, the answer for three (m_D , m_C and m_A) out of the four SUSY masses is unique! The fourth mass (m_B) is also known, up to the two-fold ambiguity (4.1), which can be easily resolved by a variety of methods discussed and illustrated in sections 3.2 and 4.2. In section 4 we applied our technique to two specific examples — the LM1 and LM6 CMS study points.

Our method contains a number of elements which help in achieving our two main objectives. For example, the precision of the SUSY mass determination is expected to improve, due to the following factors:

1. *Precise knowledge of the whole shape of the invariant mass distribution.* In appendix A we list the analytical expressions for *all* differential invariant mass distributions used in our basic analysis from section 3.1: $m_{\ell\ell}^2$, $m_{j\ell(u)}^2$ and $m_{j\ell(s)}^2(1)$. We also provide the corresponding expression for the $m_{j\ell(d)}^2(1)$ distribution, whose upper endpoint offers an independent measurement of $M_{j\ell(u)}^{\max}$ (see eq. (2.21)). Finally, we also list the formula for the differential distribution of $m_{j\ell(p)}^2$, whose endpoint can be used for selecting the correct m_B solution, as shown in figures 6(c) and 7(c). The knowledge of the shape of the whole distribution is indispensable and greatly improves the accuracy of the endpoint extraction. In the absence of any analytical results like those in appendix A, one would be forced to use simple linear extrapolations, which would lead to a significant systematic error.
2. *The number of available measurements tremendously exceeds the number of unknown mass parameters.* In principle, in order to extract 4 mass parameters, one needs a set of 4 measurements, for which we chose (3.1). On the other hand, section 2 contains a number of additional variables, whose endpoints will also be measured, and possibly

even better than our basic set (3.1). The addition of these extra measurements cannot hurt, and can only improve the overall accuracy of the SUSY mass determinations.

3. *Improved precision on the endpoint measurements.* Clearly, not all invariant mass variables will have their endpoints measured with exactly the same precision — some endpoints will be measured better than others. This difference can be due to many factors, e.g. the slope of the distribution near the endpoint, the shape (convex versus concave) of the distribution near the endpoint, the actual location of the endpoint, the level of SM and SUSY combinatorial background near the endpoint, etc. Our analysis in section 3.1 was based on a specific set of 4 endpoint measurements (3.1), which were chosen due to the simplicity in their theoretical interpretation. However, these may not necessarily be *the best measured* endpoints. In fact one can already anticipate from figures 6 and 7 that the endpoints of some of the $m_{j\ell\ell}$, $m_{j\ell(l_0)}$, $m_{j\ell(p)}$ and $m_{j\ell(s)}(\alpha \neq 1)$ distributions might be measured even better. For example, the distributions in figures 6(c), 6(e) and 6(f) are all steeper near their endpoints, compared to the distribution in figure 4(c) that we used. By the same token, one might expect that the endpoints in figures 7(a), 7(d) and 7(e) will be measured more precisely than the upper endpoints of figures 5(b) and 5(d).
4. *Controlled selection of an optimum set of measurements.* Notice that the variable $m_{j\ell(s)}^2(\alpha)$ defined in eq. (2.8) depends on a *continuous* parameter α whose value can be dialed up by the experimenter at will. This has several advantages. For example, as we have seen in figure 2, the discriminating power of $m_{j\ell(s)}^2(\alpha)$ in rejecting the wrong solution in (3.5) depends on the value of α . Having obtained a preliminary information about the two competing solutions, one can then choose the optimum value (or a range of values) for α for a subsequent study. Similarly, after the initial solution for the mass spectrum has been obtained, one can analyze by Monte Carlo the shapes of the $m_{j\ell(s)}^2(\alpha)$ distributions as a function of α and select for further study specific values of α for which the corresponding endpoints $m_{j\ell(s)}^{\max}(\alpha)$ are expected to be measured with a much better experimental precision.

In meeting our second objective, our method shows a certain improvement on the theoretical side as well:

1. *Reduced sensitivity to the parameter space region.* All of the new variables introduced in section 2 exhibit milder sensitivity to the parameter space region, in comparison to the conventional endpoint $m_{j\ell\ell}^{\max}$. As can be seen from the formulas in section 2, the endpoint for each of our variables is given by at most two different expressions, as opposed to four in the case of $m_{j\ell\ell}^{\max}$. A notable exception is the variable $m_{j\ell(s)}(1)$, whose endpoint is actually uniquely predicted, and is independent of the parameter space region. We therefore strongly encourage the use of $m_{j\ell(s)}(1)$ in future analyses of SUSY mass determinations.
2. *Uniqueness of the solution.* It is worth emphasizing that with only the 4 measurements of eq. (3.1) we can already uniquely determine three out of the four masses

involved in the problem. Then, the addition of a fifth measurement, as discussed in sections 3.2.1 and 4.2, is sufficient to pin down all four of the SUSY masses. In contrast, with the conventional approach, one also starts with four measurements as in (1.12), but in the worst case scenario this results in infinitely many solutions, due to the linear dependence problem (1.14) discussed in section 1.2. Adding a fifth measurement as in (1.18) helps, but once again, the worst case scenario leads to two alternative solutions [47]. In order to resolve the remaining duplication, and thus guarantee uniqueness of the solution under any circumstances, one needs at least 6 measurements.

In conclusion, our main accomplishment in this paper was to expand the experimenter's arsenal with several new tools which can be used for SUSY mass determinations via kinematic endpoints. We believe that the variables suggested in section 2 and the shapes of their distributions listed in appendix A will eventually find their way into the actual experimental analyses after the discovery of SUSY (or any other new physics exhibiting the decay chain of figure 1).

Acknowledgments

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A Analytical expressions for the shapes of the invariant mass distributions

In this appendix we will provide the analytical expressions for the shapes of the invariant mass distributions $m_{\ell\ell}^2$, $m_{j\ell(u)}^2 \equiv m_{j\ell_n}^2 \cup m_{j\ell_f}^2$, $m_{j\ell(s)}^2(1) \equiv m_{j\ell_n}^2 + m_{j\ell_f}^2$, $m_{j\ell(d)}^2(1) \equiv |m_{j\ell_n}^2 - m_{j\ell_f}^2|$, and $m_{j\ell(p)}^2$. To simplify the expressions, we introduce the shorthand notation for the corresponding endpoints, which was already introduced in (3.2), (3.9), (3.11) and (3.12):

$$L \equiv (m_{\ell\ell}^{\max})^2 = m_D^2 R_{CD} (1 - R_{BC}) (1 - R_{AB}), \quad (\text{A.1})$$

$$n \equiv (m_{j\ell_n}^{\max})^2 = m_D^2 (1 - R_{CD}) (1 - R_{BC}), \quad (\text{A.2})$$

$$f \equiv (m_{j\ell_f}^{\max})^2 = m_D^2 (1 - R_{CD}) (1 - R_{AB}), \quad (\text{A.3})$$

$$p \equiv R_{BC} f = m_D^2 (1 - R_{CD}) R_{BC} (1 - R_{AB}). \quad (\text{A.4})$$

In this appendix, we shall ignore spin correlations and consider only pure phase space decays. General results including spin correlations for $m_{\ell\ell}^2$, $m_{j\ell_n}^2$ and $m_{j\ell_f}^2$ exist and can be found in [71]. We shall unit-normalize the $m_{\ell\ell}^2$, $m_{j\ell(s)}^2$, $m_{j\ell(d)}^2$ and $m_{j\ell(p)}^2$ distributions, to which each event contributes a single entry. In contrast, the union distribution $m_{j\ell(u)}^2$ has two entries per event, so it will be normalized to 2 instead. It is also convenient to write the distributions in terms of masses squared instead of linear masses. Of course, the two are trivially related by

$$\frac{dN}{dm} = 2m \frac{dN}{dm^2}. \quad (\text{A.5})$$

A.1 Dilepton mass distribution $m_{\ell\ell}^2$

The differential dilepton invariant mass distribution is given by

$$\frac{dN}{dm_{\ell\ell}^2} = \frac{1}{L}, \quad (\text{A.6})$$

which is unit-normalized:

$$\int_0^L dm_{\ell\ell}^2 \left(\frac{dN}{dm_{\ell\ell}^2} \right) = 1. \quad (\text{A.7})$$

A.2 Combined jet-lepton mass distribution $m_{j\ell(u)}^2$

The differential distribution for $u \equiv m_{j\ell(u)}^2$ is given by

$$\frac{dN}{du} = \theta(n-u) \theta(u) \frac{1}{n} + \theta(p-u) \theta(u) \frac{\ln(f/p)}{f-p} + \theta(f-u) \theta(u-p) \frac{\ln(f/u)}{f-p}, \quad (\text{A.8})$$

where $\theta(x)$ is the usual Heaviside step function

$$\theta(x) \equiv \begin{cases} 1, & x \geq 0, \\ 0, & x < 0. \end{cases} \quad (\text{A.9})$$

It is easy to verify the normalization condition

$$\int_0^M du \left(\frac{dN}{du} \right) = 2, \quad (\text{A.10})$$

where $M \equiv (M_{j\ell(u)}^{\max})^2$ was already defined in (3.2).

In figure 8(a) we cross-check the prediction of eq. (A.8) (blue dashed line) with the numerically obtained $m_{j\ell(u)}^2$ distribution in figure 4(b) (red solid line), for the case of study point LM1. We see that within the statistical errors, our formula is in perfect agreement with the numerical result.

A.3 Distribution of the sum $m_{j\ell(s)}^2(\alpha = 1)$

The differential distribution for $\sigma \equiv m_{j\ell(s)}^2(\alpha = 1)$ is given by

$$\begin{aligned} \frac{dN}{d\sigma} = \frac{1}{f-p} \left\{ \theta(m-\sigma) \theta(\sigma) \ln \left(\frac{fn}{fn-\sigma(f-p)} \right) \right. \\ \left. + \theta(M-\sigma) \theta(\sigma-m) \ln \left(\frac{M}{M-(f-p)} \right) \right. \\ \left. + \theta(n+p-\sigma) \theta(\sigma-M) \ln \left(\frac{fn-\sigma(f-p)}{p(n+p-f)} \right) \right\}, \quad (\text{A.11}) \end{aligned}$$

where $m \equiv (m_{j\ell(u)}^{\max})^2$ was defined in (3.2), and n , f and p were defined in (A.2)–(A.4). The normalization condition for (A.11) reads

$$\int_0^S d\sigma \left(\frac{dN}{d\sigma} \right) = 1, \quad (\text{A.12})$$

where S is defined in (3.2).

As a cross-check, figure 8(b) shows that our analytical formula in eq. (A.11) agrees with the numerical result from figure 4(c) for the LM1 study point.

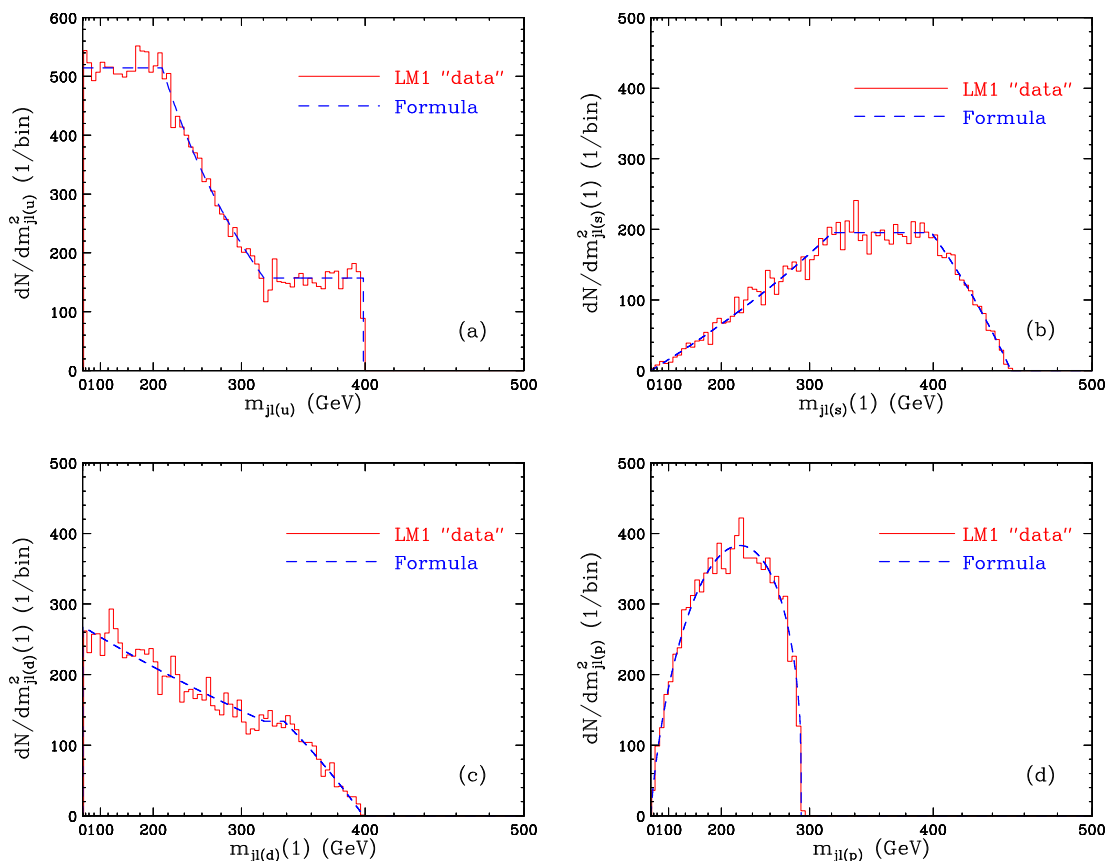


Figure 8. Comparison of the numerically obtained differential invariant mass distributions for study point LM1 (red solid lines) with the analytical results presented in this appendix (blue dashed lines): (a) the distribution of the combined jet-lepton mass $u \equiv m_{j\ell(u)}^2$ from figure 4(b) versus the analytical prediction of eq. (A.8); (b) the distribution of the sum $\sigma \equiv m_{j\ell(s)}^2(\alpha = 1)$ from figure 4(c) versus the analytical prediction of eq. (A.11); (c) the distribution of the difference $\Delta \equiv m_{j\ell(d)}^2(\alpha = 1)$ from figure 4(d) versus the analytical prediction of eqs. (A.15)–(A.19); (d) the distribution of the product $\rho \equiv m_{j\ell(p)}^2$ from figure 6(c) versus the analytical prediction of eqs. (A.22)–(A.23).

A.4 Distribution of the difference $m_{j\ell(d)}^2(\alpha = 1)$

The differential distribution for the difference $\Delta \equiv m_{j\ell(d)}^2(\alpha = 1)$ depends on the values of R_{BC} and R_{AB} . To simplify the notation, we define an antisymmetric function

$$L(x, y) = -L(y, x) \equiv \ln \left(\frac{nf + x(f - p)}{nf + y(f - p)} \right), \quad (\text{A.13})$$

which we heavily use in writing down the result for the differential Δ distribution. Notice that there are various equivalent ways to write down these formulas, due to the transitivity property

$$L(x, y) + L(y, z) = L(x, z). \quad (\text{A.14})$$

For $\Delta \equiv m_{j\ell(d)}^2(\alpha = 1)$ one needs to consider five separate cases:

- If $\frac{2}{3-R_{AB}} \leq R_{BC} < 1$, then

$$\frac{dN}{d\Delta} = \frac{1}{f-p} \left\{ \begin{aligned} &\theta(n-\Delta)\theta(\Delta) [L(0,-n) + L(-\Delta,-n)] \\ &+ \theta(p-n-\Delta)\theta(\Delta-n)L(0,-n) \\ &+ \theta(f-\Delta)\theta(\Delta-(p-n))L(f,\Delta) \end{aligned} \right\}. \quad (\text{A.15})$$

- If $\frac{1}{2-R_{AB}} \leq R_{BC} < \frac{2}{3-R_{AB}}$, then

$$\frac{dN}{d\Delta} = \frac{1}{f-p} \left\{ \begin{aligned} &\theta(p-n-\Delta)\theta(\Delta) [L(0,-n) + L(-\Delta,-n)] \\ &+ \theta(n-\Delta)\theta(\Delta-(p-n)) [L(f,\Delta) + L(-\Delta,-n)] \\ &+ \theta(f-\Delta)\theta(\Delta-n)L(f,\Delta) \end{aligned} \right\}. \quad (\text{A.16})$$

- If $R_{AB} \leq R_{BC} < \frac{1}{2-R_{AB}}$, then

$$\frac{dN}{d\Delta} = \frac{1}{f-p} \left\{ \begin{aligned} &\theta(n-p-\Delta)\theta(\Delta) [L(f,\Delta) + L(f,0)] \\ &+ \theta(n-\Delta)\theta(\Delta-(n-p)) [L(f,\Delta) + L(-\Delta,-n)] \\ &+ \theta(f-\Delta)\theta(\Delta-n)L(f,\Delta) \end{aligned} \right\}. \quad (\text{A.17})$$

- If $\frac{R_{AB}}{2-R_{AB}} \leq R_{BC} < R_{AB}$, then

$$\frac{dN}{d\Delta} = \frac{1}{f-p} \left\{ \begin{aligned} &\theta(n-p-\Delta)\theta(\Delta) [L(f,\Delta) + L(f,0)] \\ &+ \theta(f-\Delta)\theta(\Delta-(n-p)) [L(f,\Delta) + L(-\Delta,-n)] \\ &+ \theta(n-\Delta)\theta(\Delta-f)L(-\Delta,-n) \end{aligned} \right\}. \quad (\text{A.18})$$

- If $0 \leq R_{BC} < \frac{R_{AB}}{2-R_{AB}}$, then

$$\frac{dN}{d\Delta} = \frac{1}{f-p} \left\{ \begin{aligned} &\theta(f-\Delta)\theta(\Delta) [L(f,\Delta) + L(f,0)] \\ &+ \theta(n-p-\Delta)\theta(\Delta-f)L(f,0) \\ &+ \theta(n-\Delta)\theta(\Delta-(n-p))L(-\Delta,-n) \end{aligned} \right\}. \quad (\text{A.19})$$

The normalization condition now reads

$$\int_0^M d\Delta \left(\frac{dN}{d\Delta} \right) = 1. \quad (\text{A.20})$$

As before, in figure 8(c) we compare the prediction of our analytical formula in eqs. (A.15)–(A.19) to the numerical result obtained earlier in figure 4(d) for the LM1 study point, and we find very good agreement.

A.5 Distribution of the product $m_{j\ell(p)}^2$

Finally, for completeness we also list the differential distribution for the product variable (2.6), for which here we shall use the shorthand notation $\rho \equiv m_{j\ell(p)}^2$. To further simplify the notation, we define the function

$$X_{\pm}(\rho) \equiv \frac{\sqrt{n}}{2(f-p)} \left(\sqrt{n}f \pm \sqrt{f^2n + 4(p-f)\rho^2} \right), \quad (\text{A.21})$$

where n , f and p are defined as before in (A.2)–(A.4). There are two separate cases:

- If $R_{BC} \leq 0.5$, the ρ distribution is made up of two branches joining at $\rho = \sqrt{np}$ (see, for example the LM1 distribution in figure 6(c) and the LM6' distribution in figure 7(c))

$$\begin{aligned} \frac{dN}{d\rho} = \frac{2\rho}{nf} & \left\{ \theta(\sqrt{np} - \rho) \theta(\rho) \left[\ln\left(\frac{n}{p}\right) + 2 \ln\left(\frac{\rho}{X_-(\rho)}\right) \right] \right. \\ & \left. + \theta\left(\frac{f\sqrt{n}}{2\sqrt{f-p}} - \rho\right) \theta(\rho - \sqrt{np}) 2 \ln\left(\frac{X_+(\rho)}{X_-(\rho)}\right) \right\}. \end{aligned} \quad (\text{A.22})$$

- If $R_{BC} \geq 0.5$, there is a single branch, as illustrated by the LM1' distribution in figure 6(c) and the LM6 distribution in figure 7(c):

$$\frac{dN}{d\rho} = \frac{2\rho}{nf} \theta(\sqrt{np} - \rho) \theta(\rho) \left\{ \ln\left(\frac{n}{p}\right) + 2 \ln\left(\frac{\rho}{X_-(\rho)}\right) \right\}. \quad (\text{A.23})$$

In both of those cases, the normalization condition is

$$\int_0^{\rho^{\max}} d\rho \left(\frac{dN}{d\rho} \right) = 1, \quad (\text{A.24})$$

where ρ^{\max} is the corresponding $m_{j\ell(p)}^2$ endpoint defined in (2.7).

Figure 8(d) demonstrates that our analytical result (A.22) agrees well with the numerically derived $m_{j\ell(p)}^2$ distribution in figure 6(c) for the LM1 study point.

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