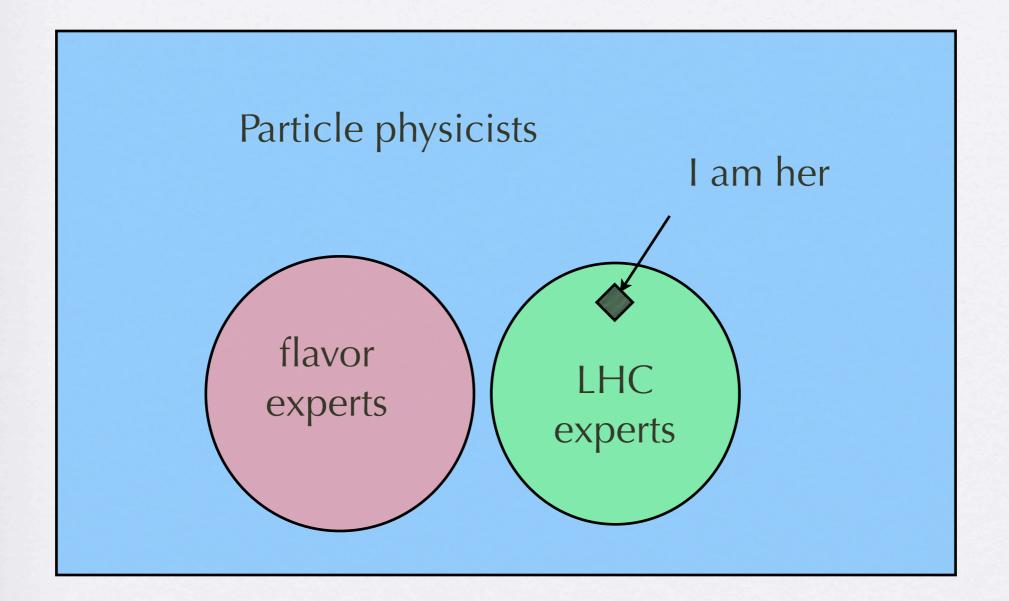
SUSY at LHC connection to flavor physics for "flavor physics in the era of LHC" Mihoko M. Nojiri

> YITP, Kyoto University Moving to KEK from Jan, 2006



### Starting from excuses....



Whatever I talk about "flavor physics" can be wrong. (It might be more useful if you ask numbers you want from LHC directly to me) Also forgive me about improper references.

#### Why LHC is related to "flavor physics"

- It sets scale of the new physics
- It measures important parameters. For the case of supersymmetry, it is
  - B physics : charged higgs, sbottom and stop at LHC
  - LFV: determination and slepton masses, direct searches
  - μ, tanβ

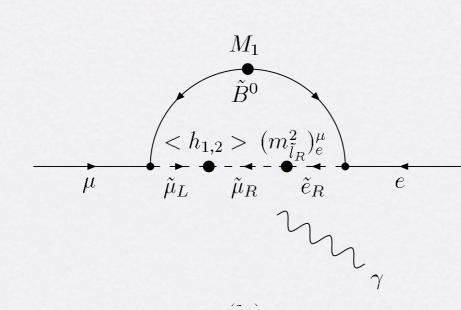
#### Note : We know nothing about that now

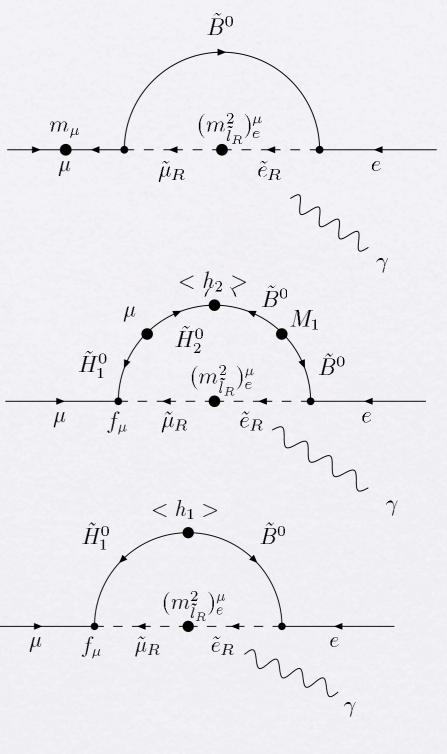
#### What do we want to know for LFV?

- What we need (for SU(5) GUT)
  - tanβ: Yukawa coupling
  - m<sub>R</sub> (12)
  - $M_{1}$ ,  $\mu$ : neutralino mass matrix

 $M_2$  and  $m_L$  for  $SU(5)+v_R(see-saw)$ 

Distinguish Left and Right sleptons is important

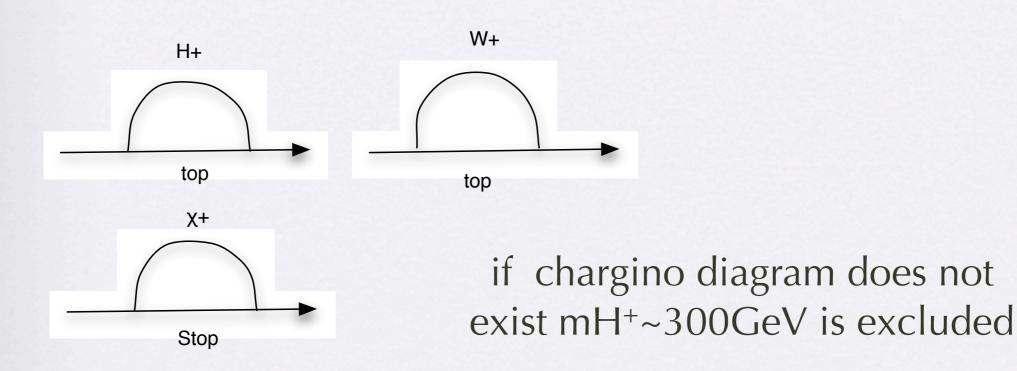




### what do we want to know

## for B physics

- many competing diagrams
- Cancellations.
- Ex:b →sγ: What is charged Higgs mass, stop mass and mixing angle?



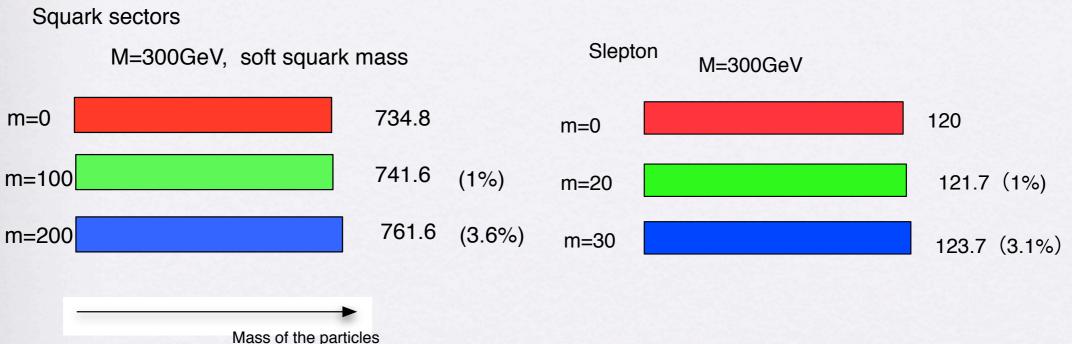
# How to put LHC constraint in the flavor analysis

- Most Flavor studies is performed in the form of scatter plot to show the deviation from SM.
- LHC will come. You may want to put minimum and model independent constraint to your analysis. What are they?
- Most of LHC analysis have been done in a few points in MSUGRA. How generic are they? You confused..

#### Access to the Flavor structure

We have seen in Franks talk that we might have high sensitivity in squark and slepton masses.

(1% for squarks, and O(1GeV) for slepton mass differences)



#### What does this means?

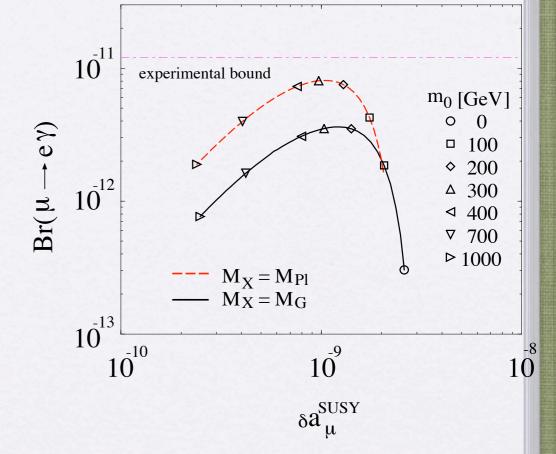
gaugino mass dominates the squark masses and it is universal. All fancy flavor effects are reduced.

$$\left[\frac{10 \text{ TeV}}{m_{\tilde{q},\tilde{g}}}\right]^2 \left[\frac{\Delta m_{\tilde{q}_{12}}^2/m_{\tilde{q}}^2}{0.1}\right]^2 \lesssim 1 .$$

K bound is not too difficult to satisfy if m<M at GUT scale

#### Flavor and LHC

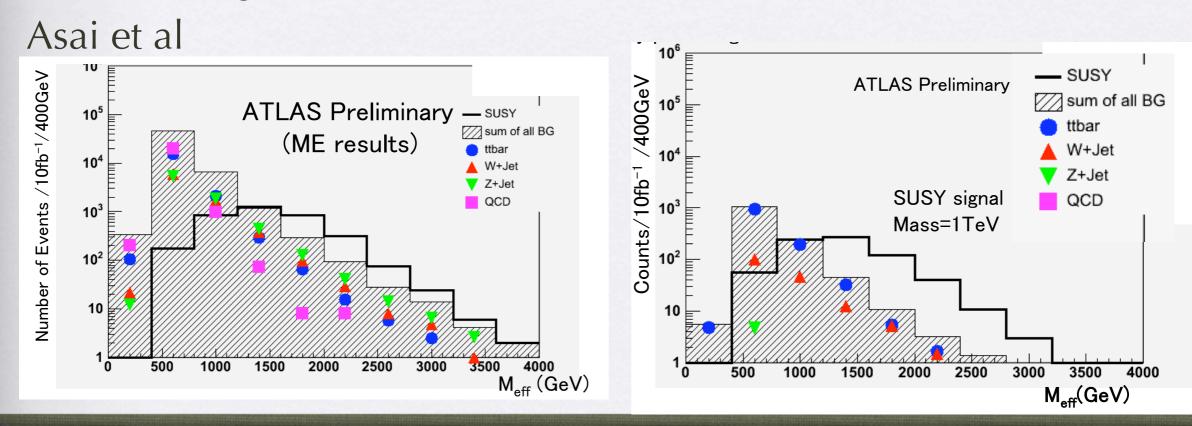
- The effect of non-diagonal scalar mass m(ij) at GUT scale (but of the order of diagonal scalar mass) will be suppressed at lower scale.
   δm<sup>2</sup>/m<sup>2</sup> < 0.01 is possible for m«M</li>
- LFV is large for m~M or m>M regions.
- LHC measurements are good when m«M. No clean result for the other MSUGRA points.



Hisano and Tobe(2001)  $M_2=250$  GeV, tan $\beta=10$ SUSY sea-saw model  $V_{13}=0.05$ ,  $Y_t=Y_{vT}$ 

#### SUSY scale determination for m»M

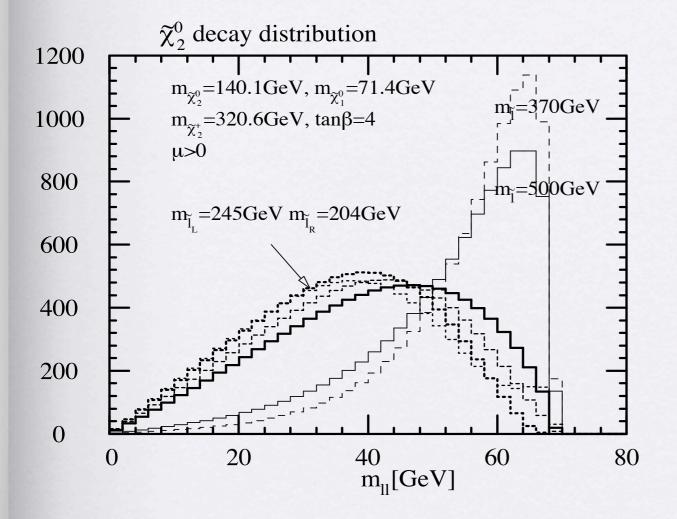
- Peak position of M<sub>eff</sub> reflects 2xM<sub>SUSY</sub>, but we have more SM background than originally thought.
- M<sub>eff</sub> (One lepton), same strategy but model dependent?
- 3 body decay of gluino. less efficient?
- We only measure gluino masses. squark decays immediately to gluino and hard to reconstruct.



#### Slepton masses and $\tilde{\chi}_2^0$ decay distribution

• m» M: all 2 body decays into slepton is closed

• virtual process  $\tilde{\chi}_2^0 \rightarrow ll \tilde{\chi}_1^0$  ? The decay distribution depends on left hand slepton masses in MSUGRA



when  $\tilde{l}_R$  is open, virtual  $\tilde{l}_L$  contribution appear beyond the two body end points.

 $\Gamma(\tilde{\chi}_2^0 \to l l \tilde{\chi}_1^0) / \Gamma(\tilde{\chi}_2^0 \to l \tilde{l}_R \to l l \tilde{\chi}_1^0)$ 

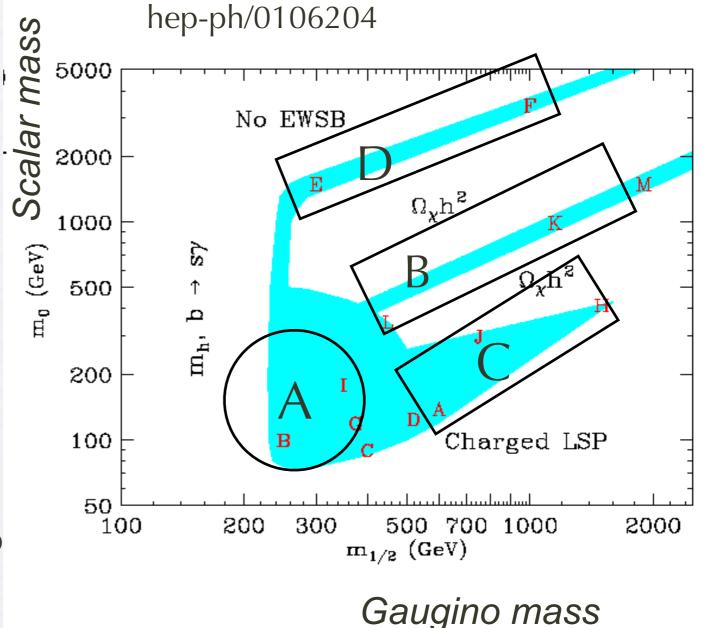
 $\Gamma(3 \text{ body})/\Gamma(\text{two bodies}) = 0.05$  for SPS1a.

#### Life with cosmological constraints

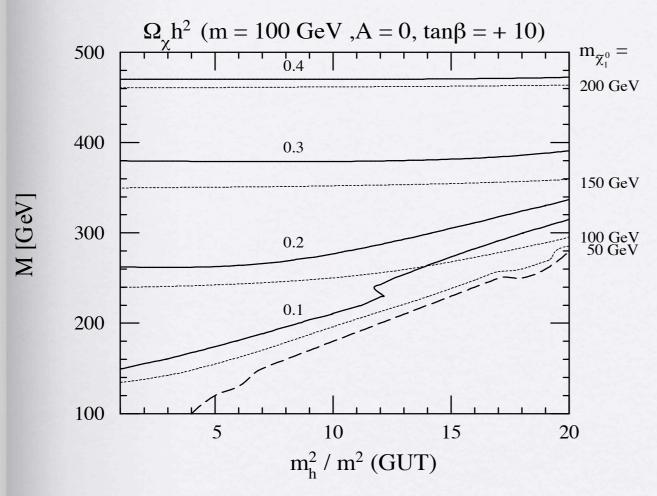
A) Bulk region: Bino like. Slepton exchange sets  $\Omega$ 

 $\Omega h^2 \propto m_{\tilde{l}}^4/m_{\tilde{\chi}}^2$ too large mass density B) Higgs pole effect near mH=2m<sub>\chi</sub>

C)  $\tilde{\tau}\tilde{\chi}$  co-annihilation D) focus point region significant higgsino-gaugino mixing



## MSUGRA might be too much



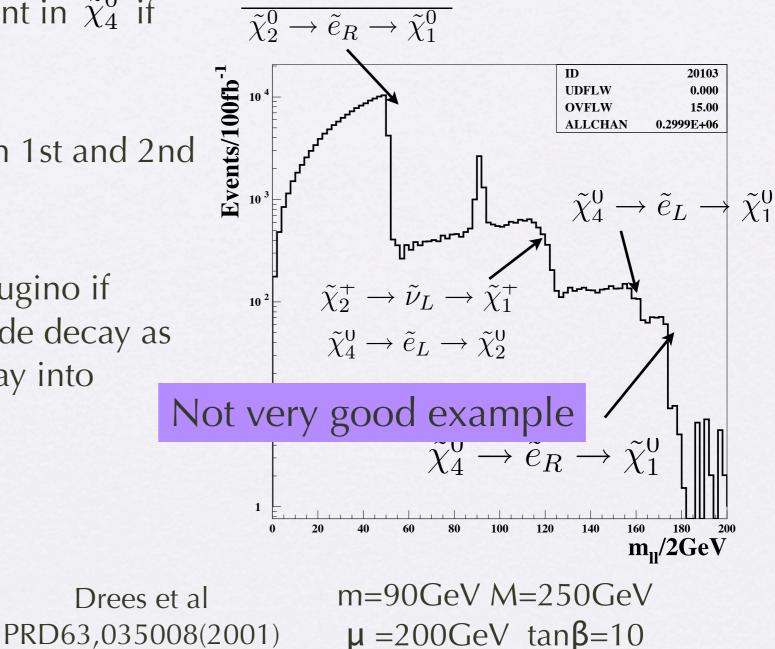
simultaneously affects LFV through 1)LR mixing of sfermions 2) Higgsino mass insertions

- µ parameter is determined so that
   v<sub>H</sub> is correct.
- So if m<sub>H</sub> at GUT scale is much higher than m<sub>16</sub>, MSUGRA prediction for µ can be evaded. More LSP pair annihilation into W, h
- m<sub>H</sub> (1,2) can be tuned independently so that higgs pole effect is enhanced.

Some recent benchmarks hep-ph 058198 for CMS (light mH

## life with cosmological constraint What happens if $\mu$ is smaller

- More gaugino component in *χ*<sup>0</sup><sub>4</sub> if M<sub>2</sub>~μ
- mass difference between 1st and 2nd ino smaller
- heavier ino becomes gaugino if M<sub>2</sub>(M<sub>1</sub>)>µ, longer cascade decay as squark dominantly decay into gaugino (heavier ino)
- several ino signature



## Example of tanβ determination scalar-muon LR mixing

if  $\tilde{\chi}_2^0 \sim \widetilde{W}$  then decay width  $\Gamma(\tilde{\chi}_2^0 \rightarrow \tilde{l}_R l)$  is so suppressed.

Small left component of scalar muon affects decay branching ratios strongly.

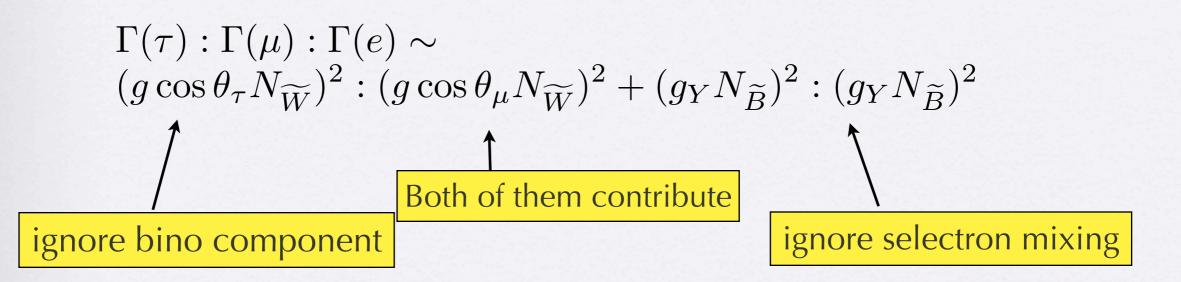
 $\tilde{\mu}_1 \sim \tilde{\mu}_R + \epsilon_L \tilde{\mu}_L \qquad \epsilon_L \sim m_\mu (\mu \tan \beta + A_\mu) / (m_L^2 - m_R^2)$ 

$$\begin{aligned} \Gamma(l) &\propto L^2 + R^2 \\ L &= g \cos \theta_l N_{\widetilde{W}^2} \\ R &= g_Y \sin \theta_l N_{\widetilde{B}^2} \end{aligned}$$

tanβ	Br(e)	Br(μ) /Br(e)	S (300fb-1)
10	6.3%	1.04	5.6
20	1.2%	1.17	7.8

Goto et al (2004) PRD70:075016

#### Importance of relative branching ratio



relative branching ratio is important. Note systematics are very different

 $\mu$ : clean, outer muon system

e: inner tracking +E<sub>cal</sub>

 $\tau$ : decay into hadron, jet isolation ,  $v_{\tau}$  missing, efficiency~50%

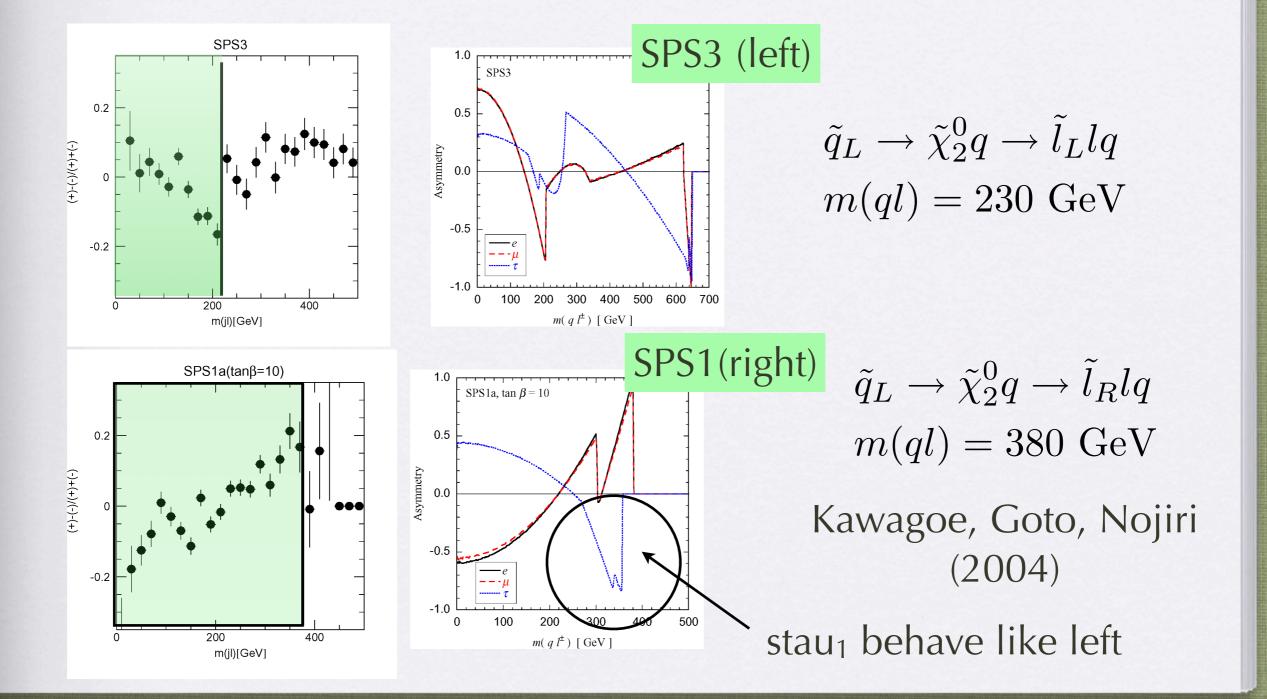
For 3 body decay and  $\mu > M_2 > M_1$ , left hand slepton dominates....

## Left or right??

- at LHC( pp collider)  $\sigma$  (squark)»  $\sigma$ (anti-squark)
- MSUGRA case  $\tilde{q} \rightarrow \tilde{\chi}_2^0 q \rightarrow \tilde{l} l q \rightarrow \tilde{\chi}_1^0 q l l$
- wino-like ino produced from left-hand squarks
- wino is polarized(in average). it decays into lepton /antilepton equally (Majorana). lepton/anti-lepton correlation to wino(jet) direction is opposite. charge asymmtry.
- NOTE, slepton further decay into lepton. look into the distribution near the jl edge( it may not be end point)

## Left or right, simulations

• m(jl) distribution tell us combination of the chirality of squark and slepton in the cascade decays.



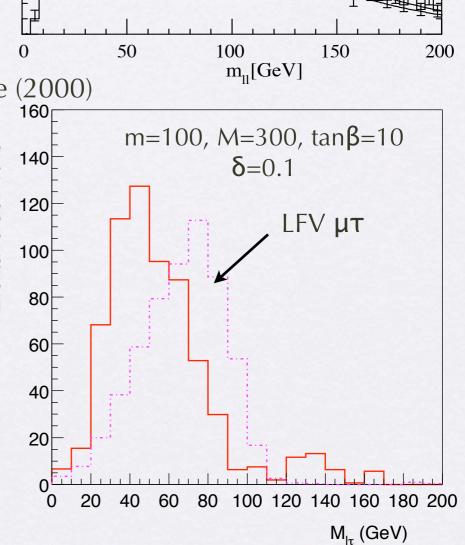
## Flavor violation in slepton decays

PRD65:116002 200 Hisano et al • LFV in 2 body decay 150 number of events /bin 100 • signal: edge in  $e\mu$  distribution 50 shoulder in  $e\tau$ ,  $\tau\mu$  distribution 0 50 Hinchiliffe (2000) Loop process: 160 I Events/10 GeV/10 fb<sup>-1</sup> GIM like suppression and 140

cancellation among diagrams.

$$BR(\tau \to \mu \gamma) \approx 1.1 \times 10^{-6} \left(\frac{\delta}{1.4}\right)^2 \left(\frac{100 \, {\rm GeV}}{M_{\tilde{\ell}}}\right)^4$$

$$\delta \equiv M_{\mu\tau}^2/M_L^2$$



hist: eu w/o LFV T: eu with LFV

-: bg fit

#### L= 100fb<sup>-1</sup>, $\Delta$ m=1.2GeV, sin2 $\theta$ =0.5

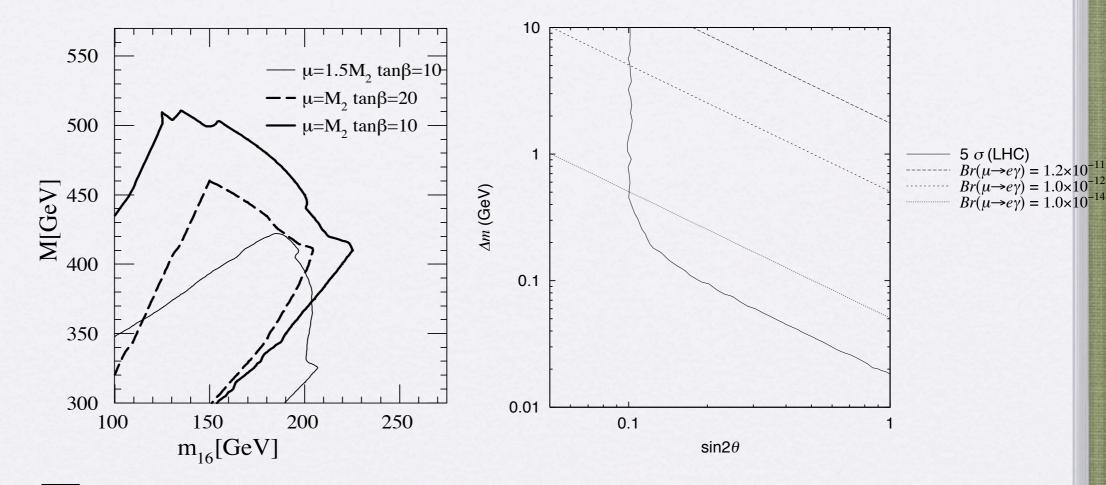
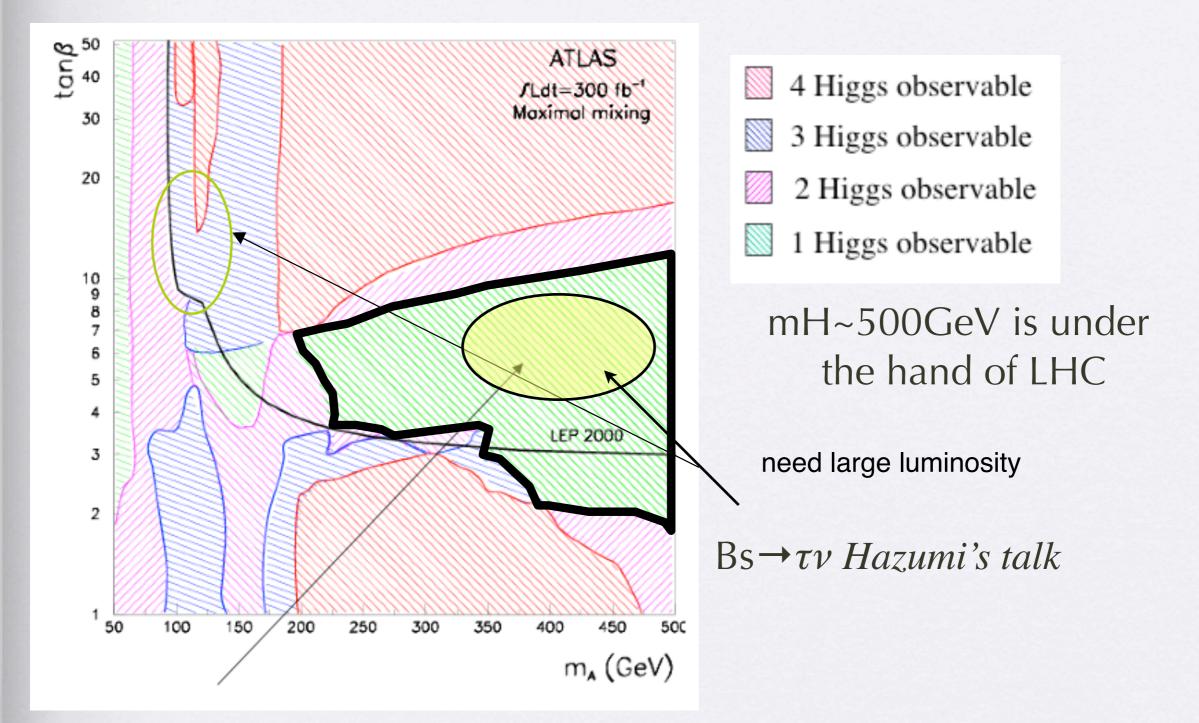


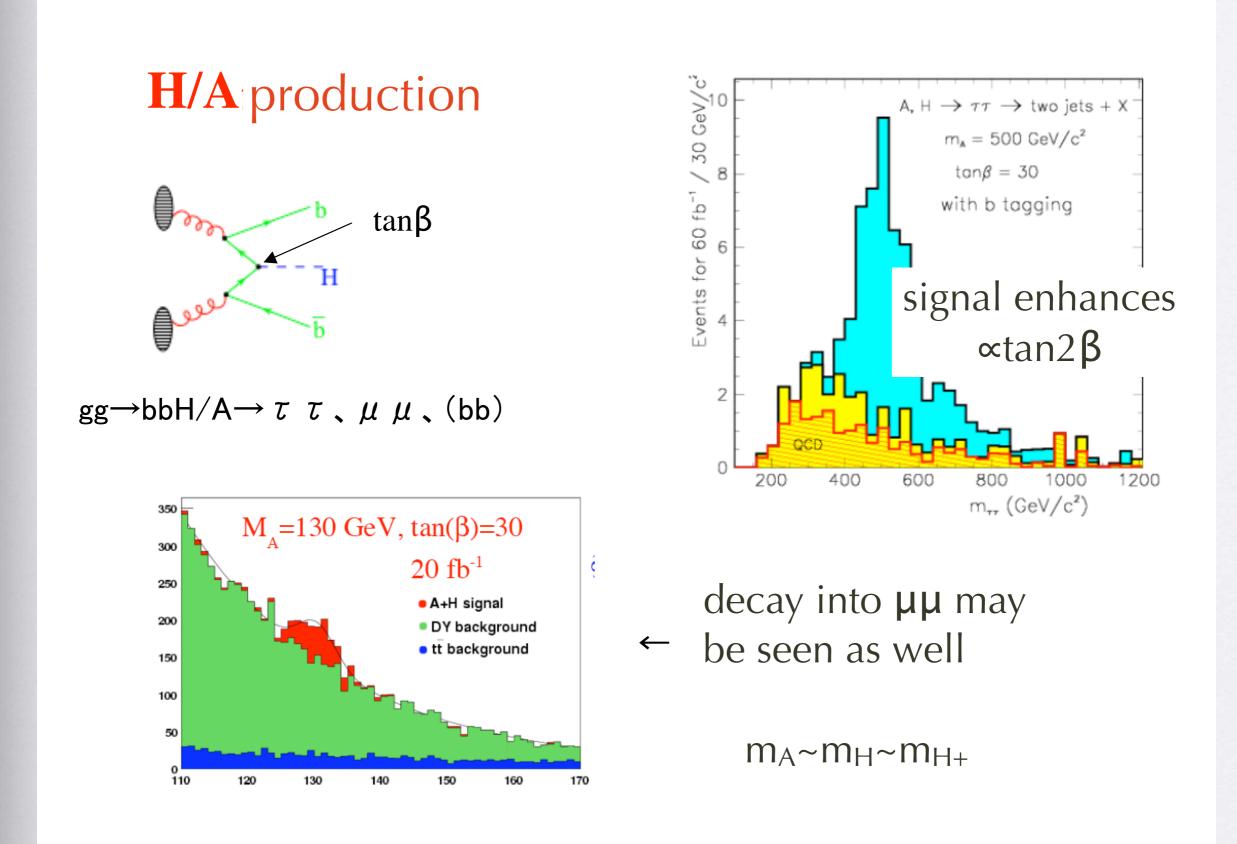
Figure 7:  $\sqrt{\Delta\chi^2} = 5$  contours for the LFV discovery. The thick solid line is for  $\mu = 1.5M_2$  and  $\tan\beta = 10$  in the cMSSM, the thick dashed line for  $\mu = M_2$  and  $\tan\beta = 20$ , and the solid line for  $\mu = M_2$  and  $\tan\beta = 10$ . We fix the  $\tilde{e}_R - \tilde{\mu}_R$  mixing angle  $\theta$  as  $\sin 2\theta = 0.5$  and the slepton mass difference  $\Delta m = 1.2$  GeV at the GUT scale.

 $\mu = M_2$ 

### LHC and MSSM Higgs in connection with B physics

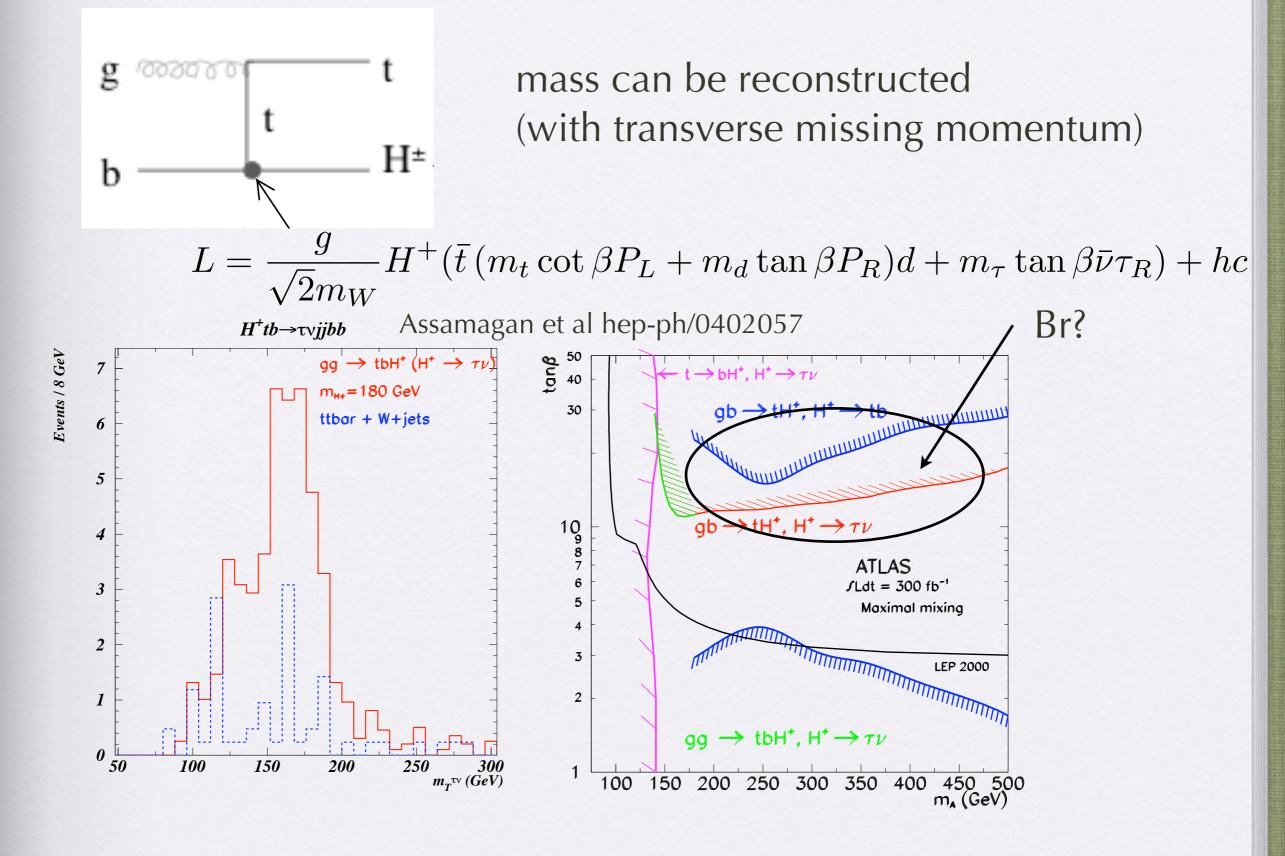


Only one higgs doublet



Here we are assuming all SUSY particles are heavy. Any loopholes or more information?

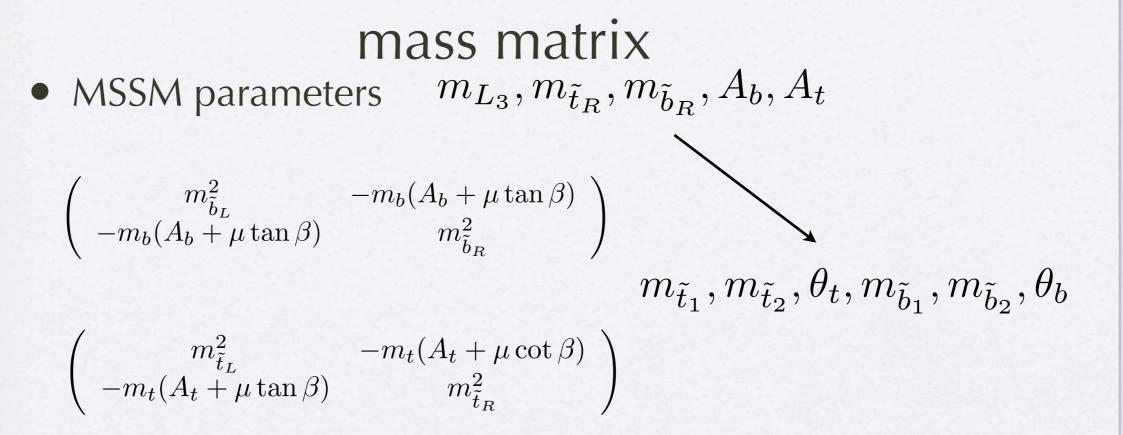
## charged higgs searches at LHC



### connection to B physics

- Example  $b \rightarrow s \gamma$ , large cancellation among diagrams
- You will have an access of the charged higgs mass, branching ratio.
- What can you say about chargino loop contribution?
   Stop mass and mixings (2-3)
- ✓ What is the info on stop from LHC??

#### 3rd generation squark

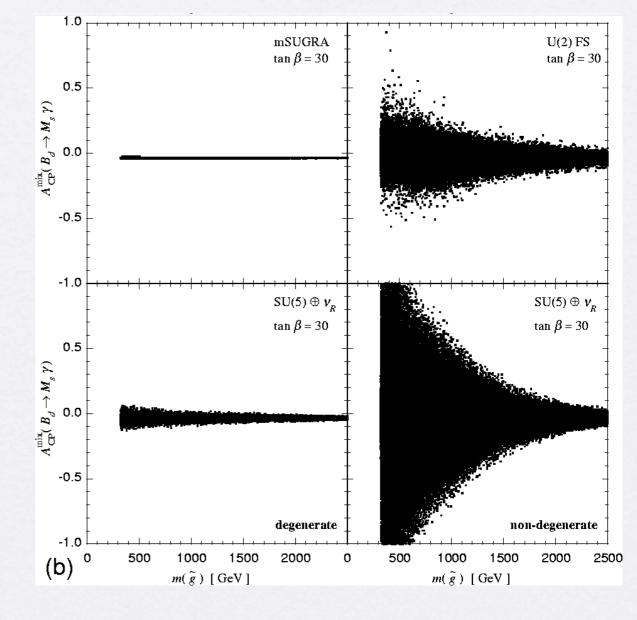


In MSUGRA: The 1st and 2nd squark mass are heavier than 3rd due to RGE running and mixings. SUSY events contain many b jets. b tagging efficiency is 60%

Production : direct production from gluon(small) decay from gluino (dominant if open)

## non-MSUGRA boundary condition in 3rd generation in B physics

- You may find surprise in B flavor violation process—this may comes from....
  - GUT scale neutrino mass assuming Y<sub>t</sub>-Y<sub>ντ</sub>
  - Non universal boundary condition
- stop and sbottom mass may also depends on such thing

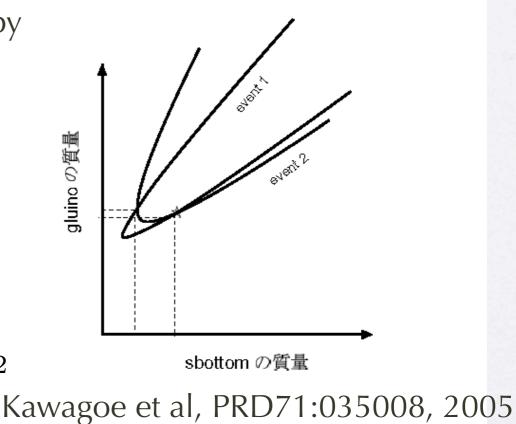


Goto et al PRD70:035012(2004)

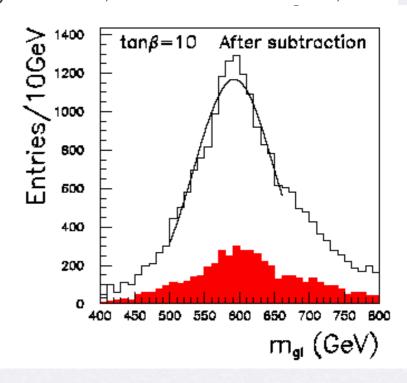
## Exact treatment for gluino and sbottom reconstruction

Longest cascade decays that can be "solved" event by event even though LSP is missing

$$\begin{split} m_{\tilde{\chi}}^2 &= p_{\chi}^2 \\ m_{\tilde{l}}^2 &= (p_{\chi} + p_{l_1})^2 \\ m_{\tilde{\chi}_2}^2 &= (p_{\chi} + p_{l_1} + p_{l_2})^2 \\ m_{\tilde{b}_1}^2 &= (p_{\chi} + p_{l_1} + p_{l_2} + p_{j_1})^2 \\ m_{\tilde{g}}^2 &= (p_{\chi} + p_{l_1} + p_{l_2} + p_{j_1} + p_{j_2})^2 \\ \end{split}$$

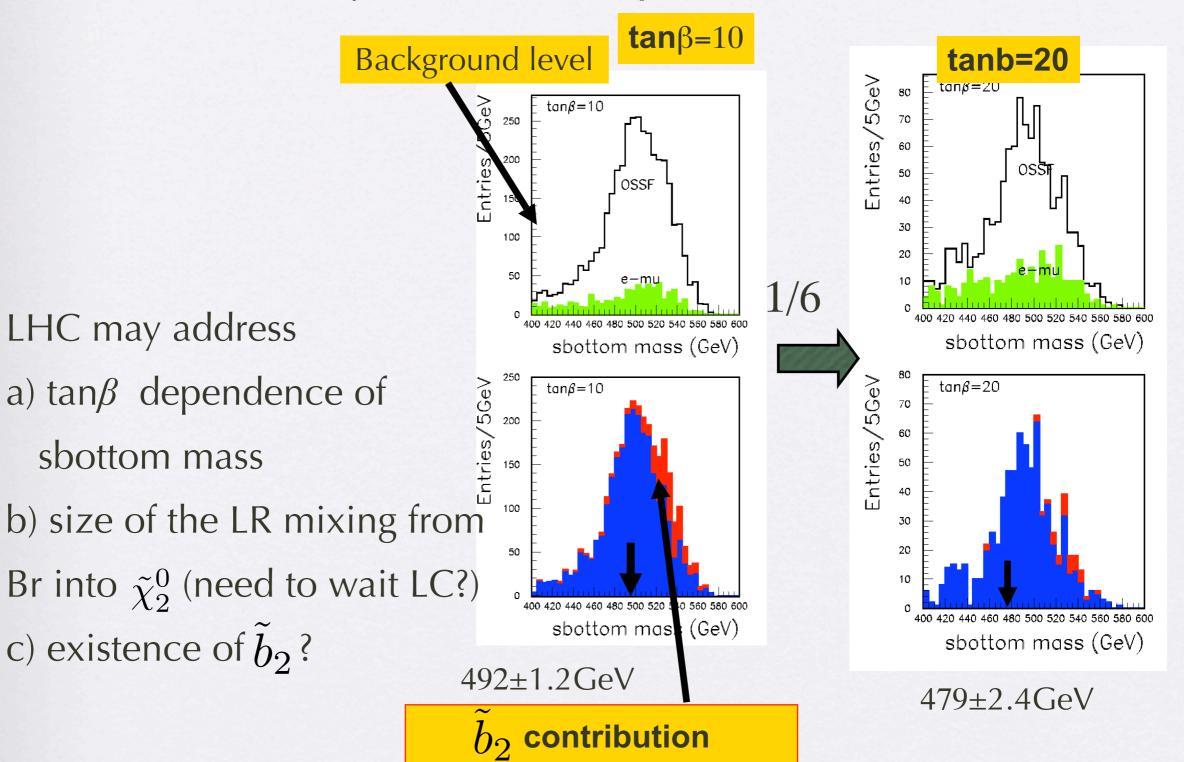


- Assume we know lighter masses
- 3 mass constraints for p(LSP) -> one degree of freedom in 2dim gluino and squark mass space
- 2 events-> mass fixed
- After mass fix, the missing momentum is solved



## sbottom mass reconstruction

(two b jets and two lepton channel )



Kawagoe, MMN, Polesello 04

## Flavor violation in sbottom decays at LHC?

- $\epsilon_b = 60\%$ , not impressive
- take high  $p_T$  jet which comes from LFV sbottom decay+ b jet from gluino  $\rightarrow$  b sb<sub>1</sub> might work(using "mass relation" as cut).
- for O(1000) bbll decay, 360 must be tagged. for full flavor violation it is only 90 events.

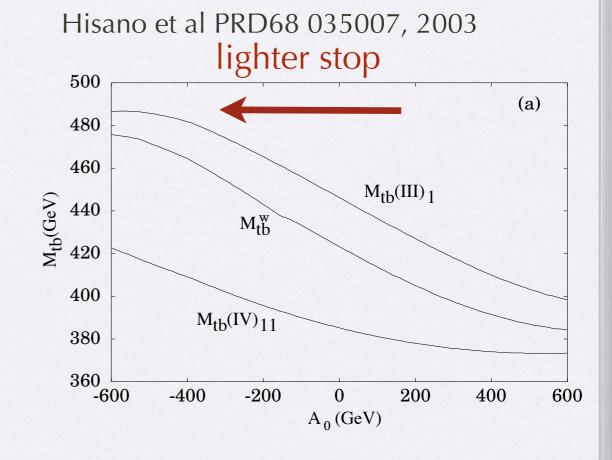
	No flavor violation $\epsilon_{b}=0.6$	full violation $\epsilon_{b}=1$	full violation $\epsilon_b=0.6$
bbll	0.6x0.6=0.36	0.25	0.09
jbll	0.6x0.4=0.24	0.25	0.21
bjll	0.6x0.4=0.24	0.25	0.21
jjll	0.4x0.4=0.16	0.25	0.49

#### scalar top at LHC

$$\tilde{g} \to \tilde{t}t \to bt\tilde{\chi}^{\pm} \Rightarrow Br(\tilde{t}) \quad M_{tb}(\tilde{t}) \\
\tilde{g} \to \tilde{b}b \to bt\tilde{\chi}^{\pm} \Rightarrow Br(\tilde{b}) \quad M_{tb}(\tilde{b})$$

- two subsequent cascade decays give tb end point. It is not dominated by single process.
- hadronic top decay can be reconstructed .
- The tb end point give you information of stop mass.

$$M_{tb}^{w} = \frac{Br(\tilde{t})M_{tb}(\tilde{t}) + Br(\tilde{b})M_{tb}(\tilde{b})}{Br(\tilde{t}) + Br(\tilde{b})}$$

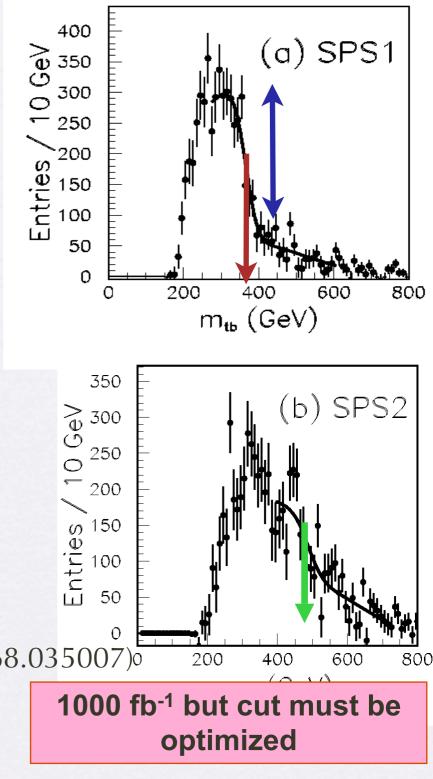


#### Difference between two body and three body

**Biggest branching ratio** 

 $\tilde{g} \rightarrow (t\tilde{t} \text{ or } b\tilde{b}) \rightarrow tb\tilde{\chi}_1^{\pm}$ 

**SPS1a:** edge with  $\Delta M_{tb} \sim 4 \text{ GeV}$  for  $100 \text{ fb}^{-1}$ height h and edge M<sub>tb</sub> may be used to understand stop sector **SPS2** :(focus points M=300GeV) No edge as expected Lower bound of stop and sbottom mass? Limited statistics but distribution may reflect  $m_{\widetilde{g}} - m_{\widetilde{\chi}_2^+} \sim 480 \text{GeV}$  $m_{\tilde{g}} - m_{\tilde{\chi}_1^+} \sim 560 \text{GeV}^{(\text{Hisano et al } \text{PRD68.035007})^\circ}$ 



## Endpoint reconstruction

What will we see if we put this constrain to B rare decays?

- weighted end point is reconstructed correctly by the fit over wide region of parameter space.
  - A1 A2: a msugra point but A changed maximally (m=100GeV, M=300GeV tanβ=10
  - T1 T2 stop mass moved by changed stop\_R mass
  - B, C, I, G from paper hep-ph/0106203
  - E1 E2, gluino decays only to stop and top.

