

The Total Cross Section and its Diffractive Components

ABSTRACT

The total and diffraction dissociation *pp* cross sections at LHC and higher energies are predicted in a model based on a novel, fully unitarized parton-model approach in which diffraction is derived from inclusive proton PDFs and QCD color factors.

Classification of diffractive processes

- Basic diffractive processes: **SD** Single Dissociation (Single Diffraction)
- SD_1 or SD_2 , depending on which incoming *p* dissociates **DD D**ouble **D**issociation (**D**ouble **D**iffraction)

CD Central Dissociation (\rightarrow DPE: Double Pomeron Exchange) **Combined diffractive processes:**

- *multigap* diffraction with two or more gaps other than CD e.g. SDD: SD plus CD (predicted & studied at CDF)
- **SDDS:** 4-gap diffraction: fully predicted (can be studied at LHC)
- This poster is organized in four columns:
- **Abstract / Introduction** (this column)
- □ The **RENORM** model **RENOR**malization **Model**
- Diffraction at CDF: validation of RENORM by data
- Diffraction at LHC: predictions and comparisons with data

INTRODUCTION

DIFFRACTION IN QCD



BASIC AND EXAMPLES OF COMBINED DIFFRACTIVE PROCESSES

acronym	basic diffractive processes
$\overline{\mathbf{SD}_{ar{p}}}$	$\bar{p}p \to \bar{p} + \operatorname{gap} + [p \to X_p],$ gap
\mathbf{SD}_p	$\bar{p}p \to [\bar{p} \to X_{\bar{p}}] + \operatorname{gap} + p,$
DD	$\bar{p}p \to [\bar{p} \to X_{\bar{p}}] + \operatorname{gap} + [p \to X_p],$
DPE	$\bar{p}p \to \bar{p} + \operatorname{gap} + X_c + \operatorname{gap} + p,$
	2-gap combinations of SD and DD
$\mathbf{SDD}_{\bar{p}}$	$\bar{p}p \to \bar{p} + \operatorname{gap} + X_c + \operatorname{gap} + [p \to X_p],$
\mathbf{SDD}_p	$\bar{p}p \to [\bar{p} \to X_{\bar{p}}] \operatorname{gap} + X_c + \operatorname{gap} + p.$

The SDD cross sections are predicted by the *RENORM* model ->

THE RENORM MODEL



Diffractive, elastic, and the total pp cross sections based on the **RENORM** model are now implemented in PYTHIA8.

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In all the above cases there is excellent agreement between *RENORM*-based MBR and data.

RENORM has been implemented in PYTHIA8 in collaboration with Robert Ciesielski from The Rockefeller University.

