

# A new highly performing compression regime: the Laminar Bunching & GIOTTO, a code for extreme BD optimizations

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and PHD Marcello Rossetti Conti*

# The Laminar Bunching

## (1) The goal

prove **extreme high performances** in compact LINACs (~20 m)

$$E_n = 150 - 500 \text{ MeV}; \quad \varepsilon_{n,peak} \approx 0.3 - 1.0 \text{ mm} - \text{mrad};$$

$$I_{peak} \text{ up to: } 4 \text{ kA}; \quad \sigma_E < 50 \text{ keV}$$

Ultra bright & Ultra cold: **Dream beams**

## (2) The technique

Hybrid Laminar Velocity Bunching **or Laminar Bunching (LB)**

$\Leftrightarrow$

Velocity Bunching + **Drift** Laminar Bunching

## Important notes:

- ✓ charts show:  $\sigma_x$ ,  $\sigma_z$ ,  $\varepsilon_{nx}$ ,  $\sigma_E$
- ✓ ASTRA (\*) Simulations
- ✓ Optimizations made in GIOTTO (\*\*)

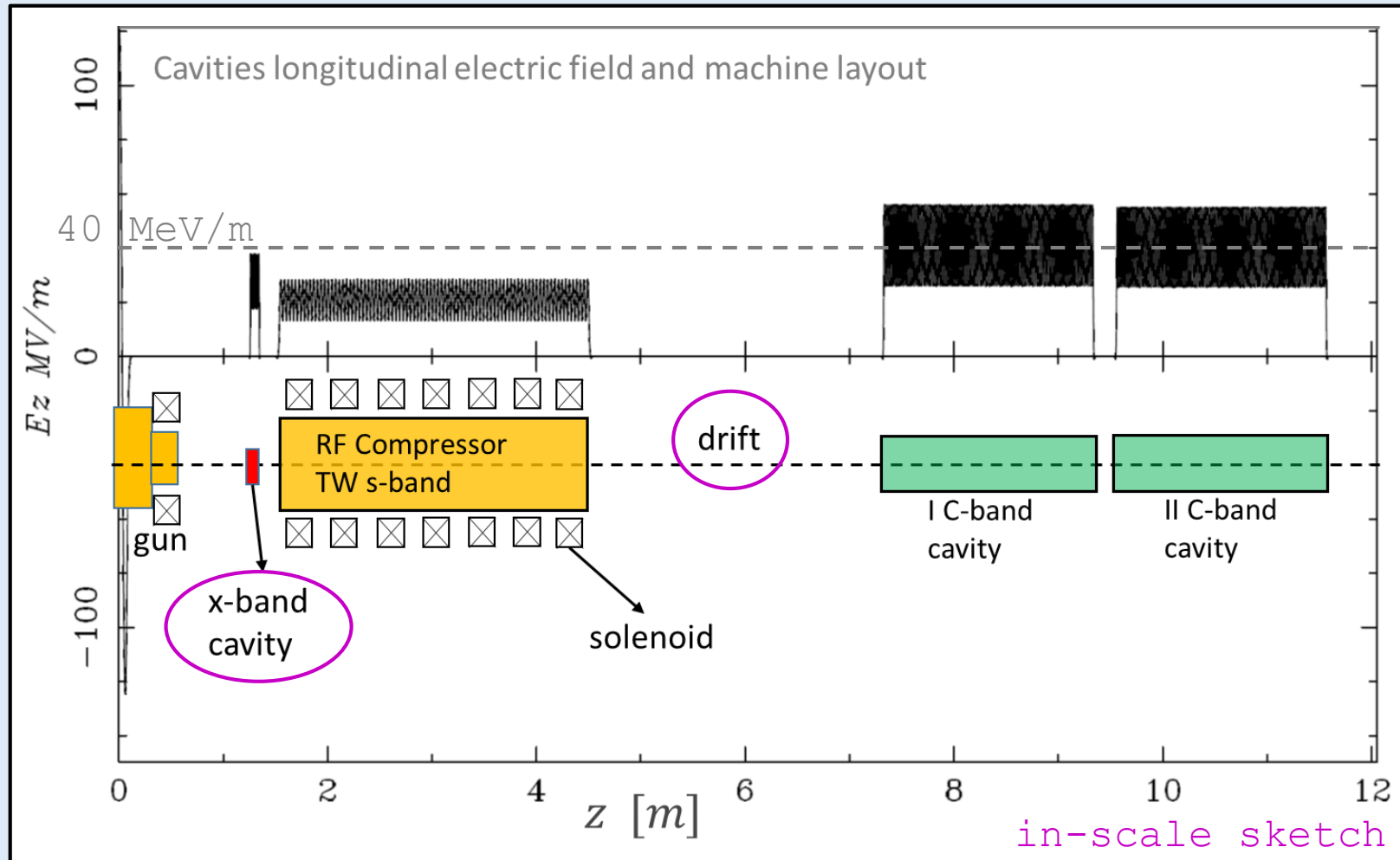
## Outline

- ❑ Ad-hoc Laminar Bunching LAYOUT
- ❑ Point out the Laminar Bunching effects
- ❑ Laminar Bunching / Velocity Bunching COMPARISON
- ❑ Some Beam Dynamics: Laminarity parameter

(\*) K. Floettmann, ASTRA—A space charge tracking algorithm, <http://www.desy.de/~mpyflo/>

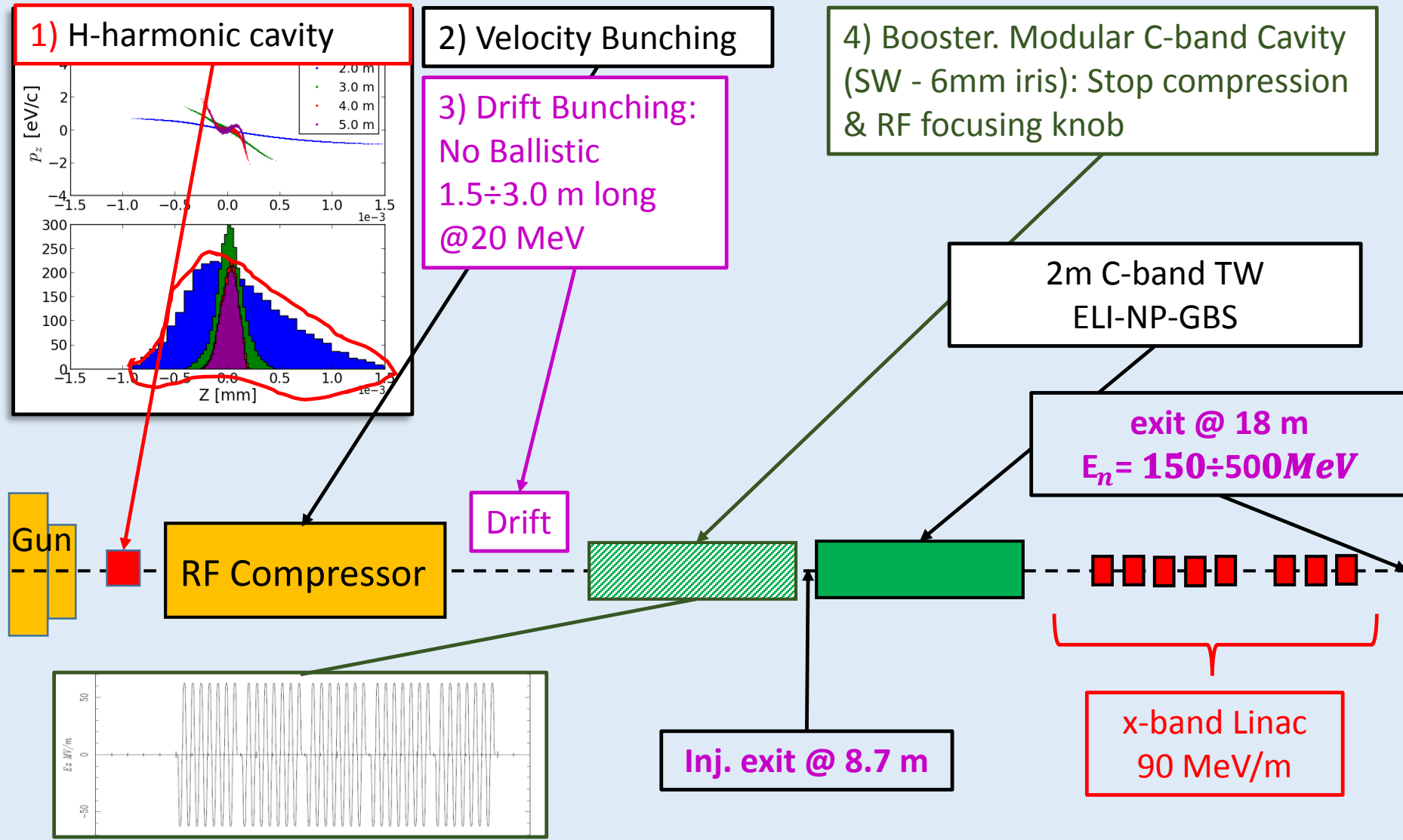
(\*\*) A. Bacci, et al. “GIOTTO: A Genetic Code for Demanding Beam-dynamics Optimizations”, doi: 10.18429/JACoW-IPAC2016-WEPOY03

# Ad-hoc layout for Laminar Bunching



# A compact machine layout working in Laminar Bunching

- 1) HighHarm-cavity current pre-correction;
- 2) Velocity Bunch.
- 3) Drift Laminar Bunching (balanced accordion effect);
- 4) RF-focusing tunable booster



# High harmonic cavity current pre-correction

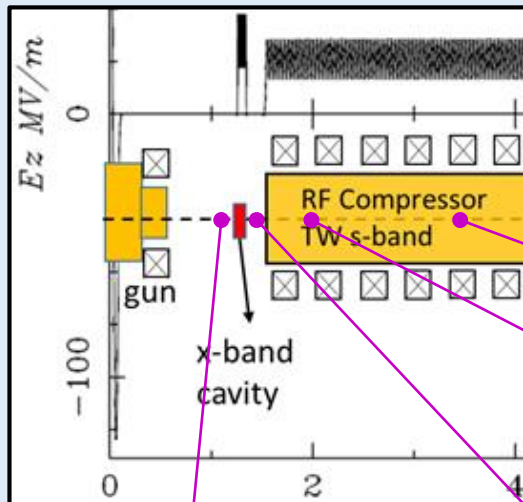


Table 1: C-band and X-band High Harmonic Correction Cavities Comparison and the Non-corrected VB Case

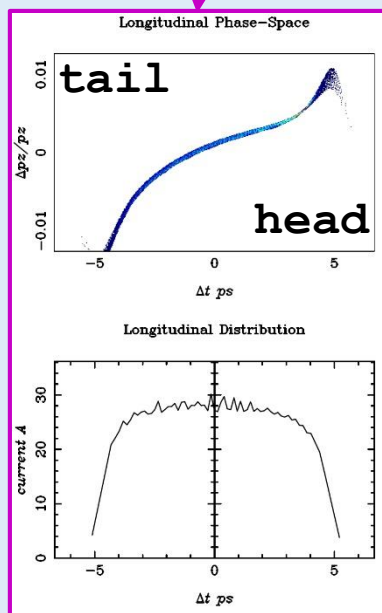
Type band	Grad. [MeV/m]	Decreased E [MeV]	$\sigma_z [\mu\text{m}]$	$\varepsilon_{n,x} [\mu\text{m}]$
Nothing	-	-	200	1.7
C-3 Cells	31	2.5	106	2.4
C-11 Cells	17.5	3.0	75	4.0
X-11 Cells	28	2.5	93	2.0
X-11 Cells	35	3.0	75	4.0

1.2 m

1.4 m

2.0 m

1.4 m



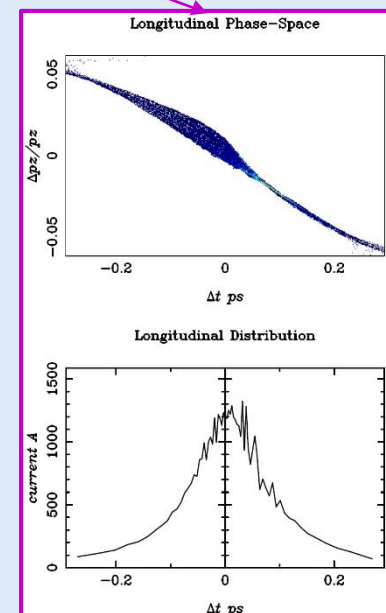
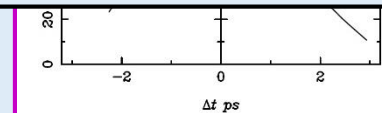
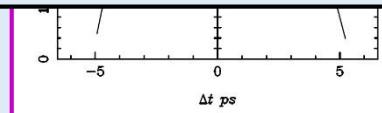
Many good effects:

- 1) RF curvature pre-correction
- 2) Current pre-correction
- 3) Starts the compression (\ chirp)

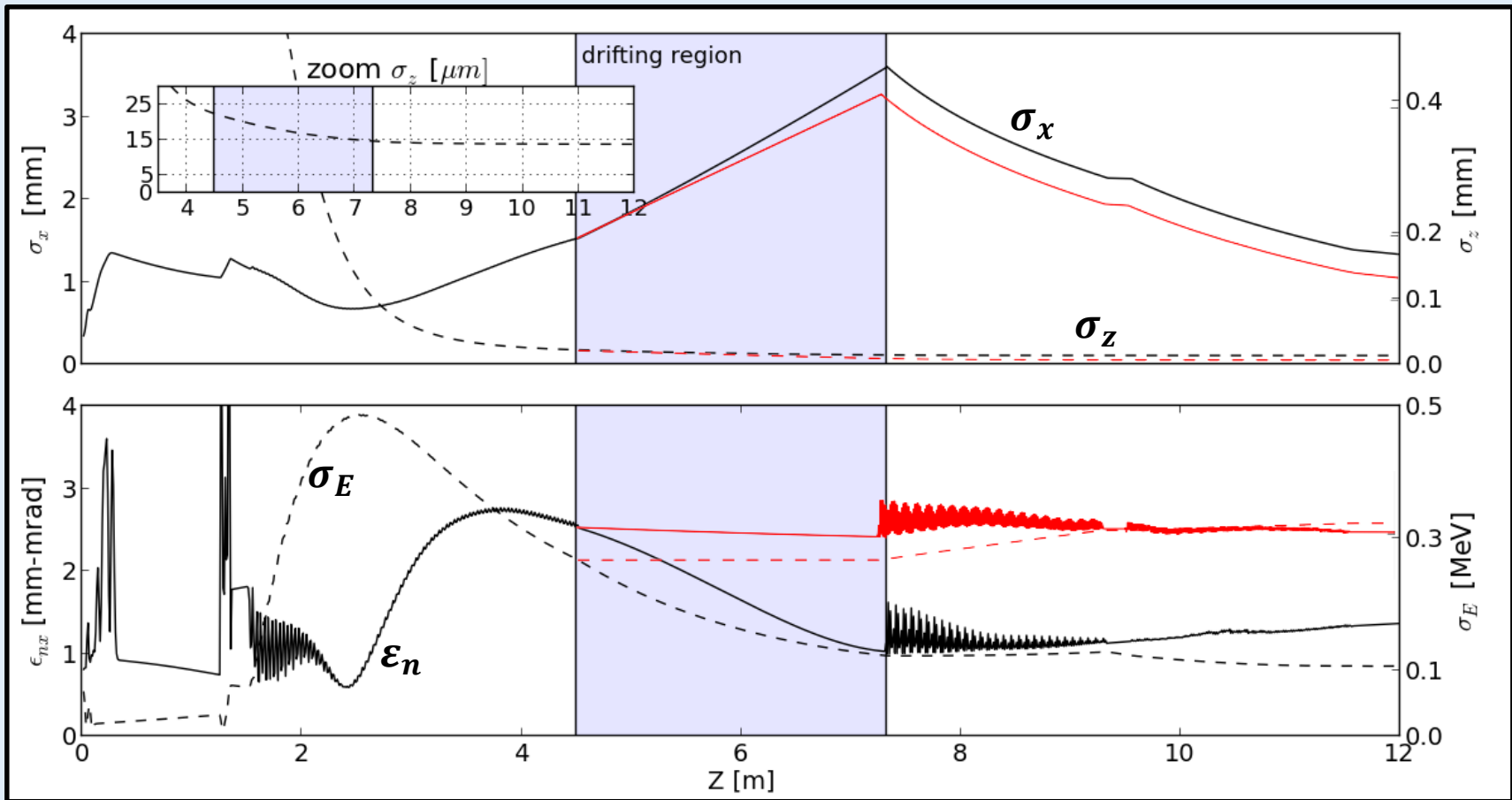
En decrease of 2.3 MeV on 6 MeV



- 4)  $\rho_z$  falls down favoring compression
- 5) A lower Energy favors VB



# $\sigma_x$ , $\sigma_z$ , $\epsilon_{nx}$ , $\sigma_E$ curves in Laminar Bunching



Turning off the SPACE charge  
from the drift onward

- $\sigma_x$  quasi linear rising
- $\sigma_z$  hyperbolic decreasing
- $\sigma_E$  a quasi full correction
- $\epsilon_n$  a quasi full correction

Both optimized by  
GIOTTO genetic  
algorithm

$$\sigma_Z, \varepsilon_{nx}, \sigma_E$$

Laminar Bunching (LB)  
Velocity Bunching (VB)  
only a comparison

LB & VB are  
relatives, but their  
final results are  
different.

The aim:  
outline LB peculiarities versus  
the VB known technique.

Both: Linear compression &  
same energies

LB works on the  
whole bunch dist.

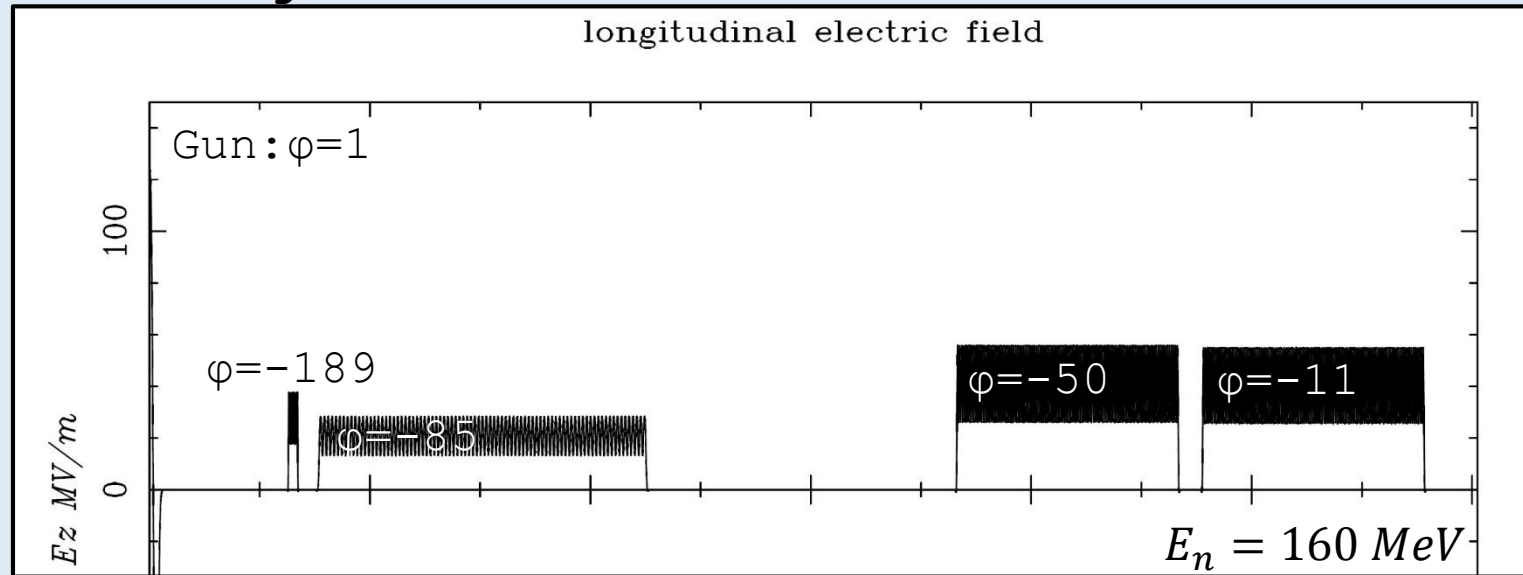
VB favors one spike  
@ bunch head



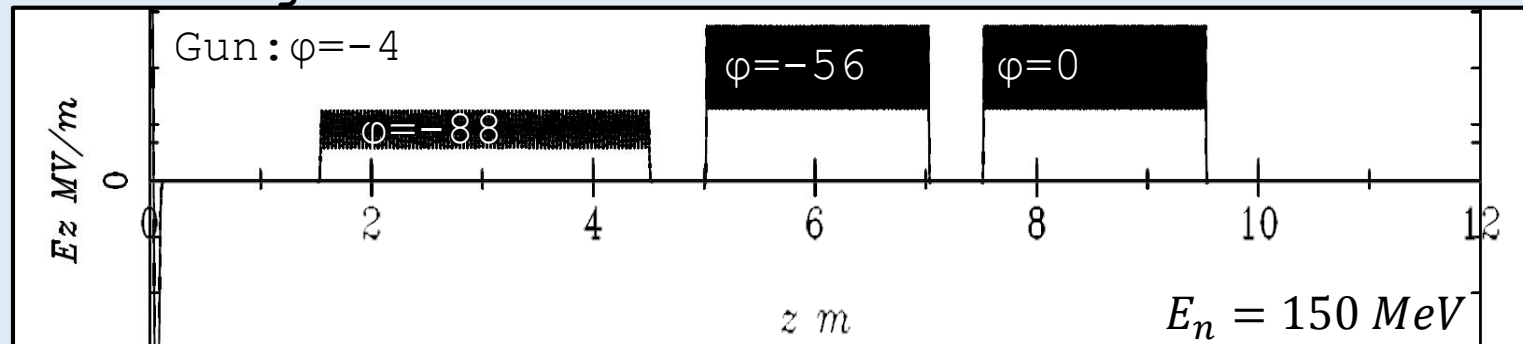
# Laminar & Velocity Bunching Layout

@ cathode for both LB & VB	$Q_b$ [pC]	$\tau_{Laser}$ [ps] flat-top	$\tau_{rising}$ [ps]	$\varepsilon_{th}$ [1 $\mu$ / mm]	$\sigma_x$ [ $\mu$ m]
	250	10	1	0.9	260

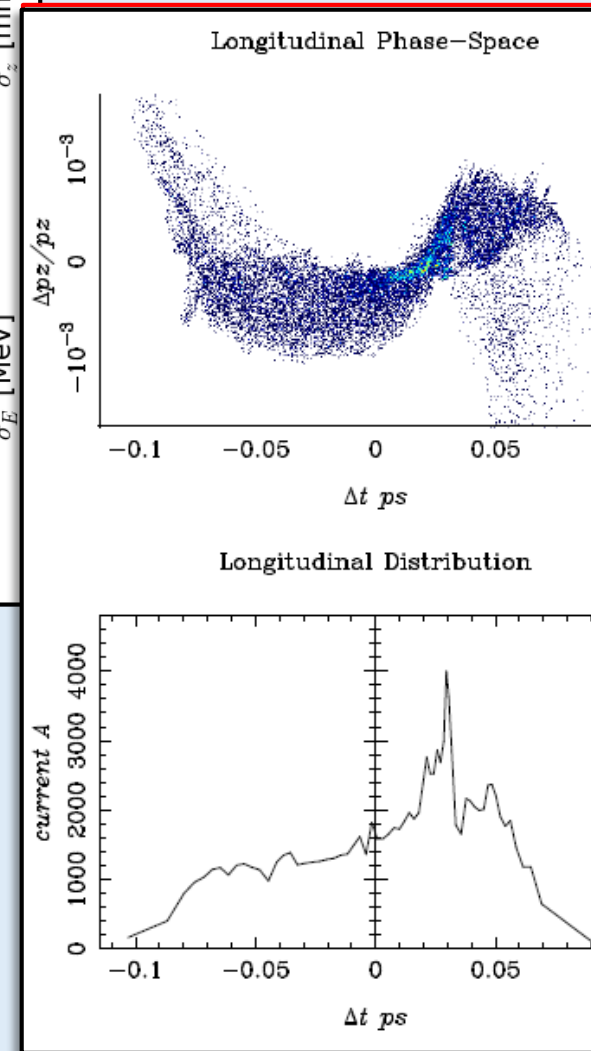
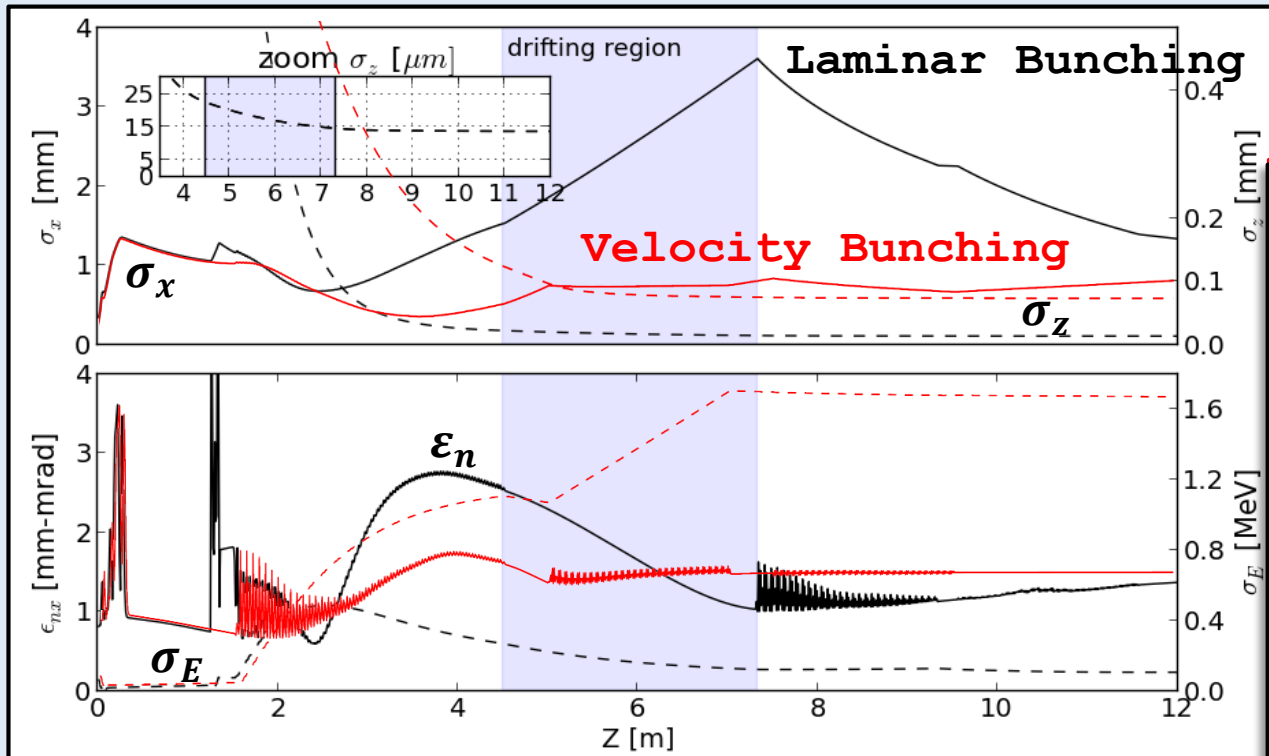
## L. Bunching



## V. Bunching

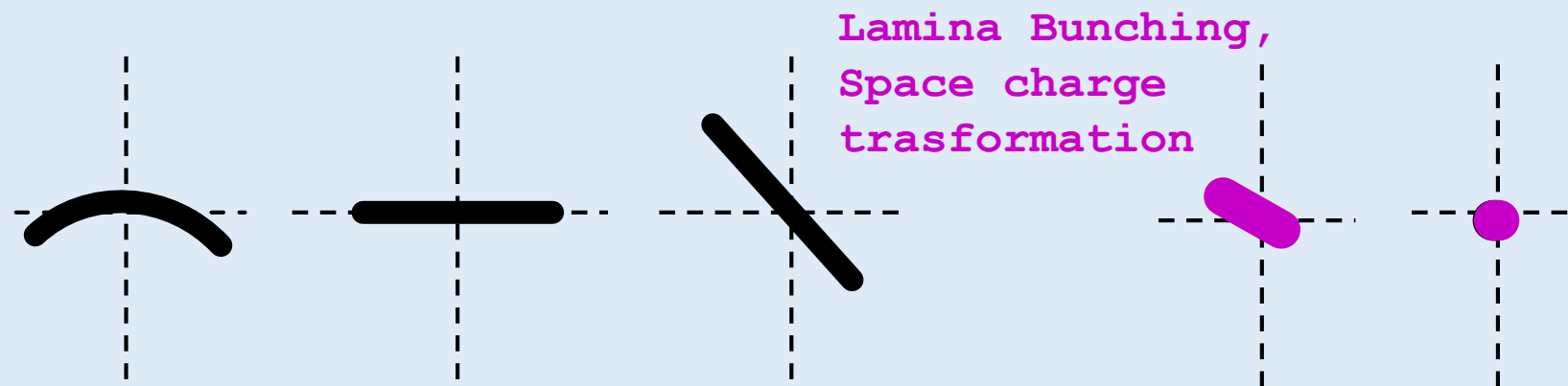
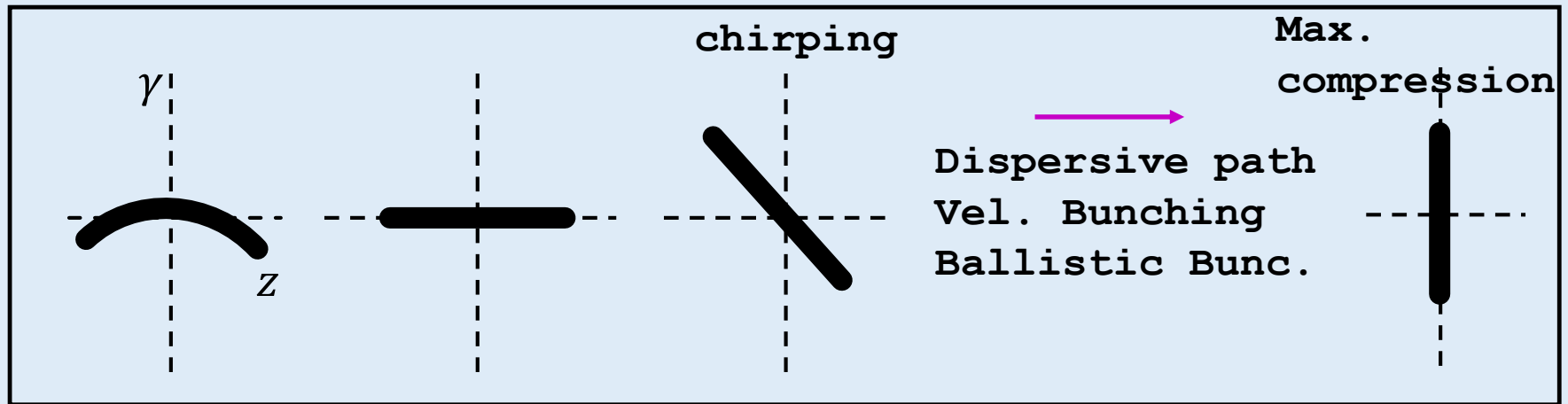


# Laminar " " & Velocity " " Bunching comparison



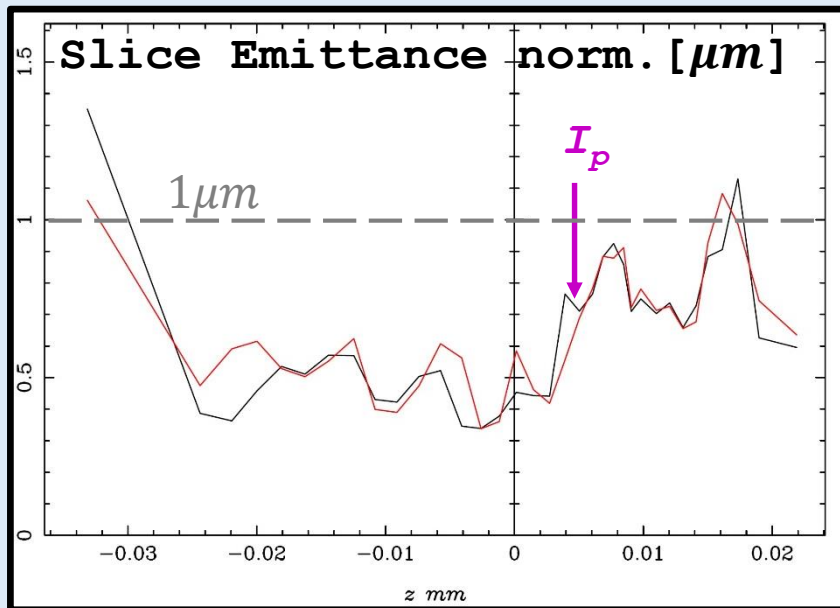
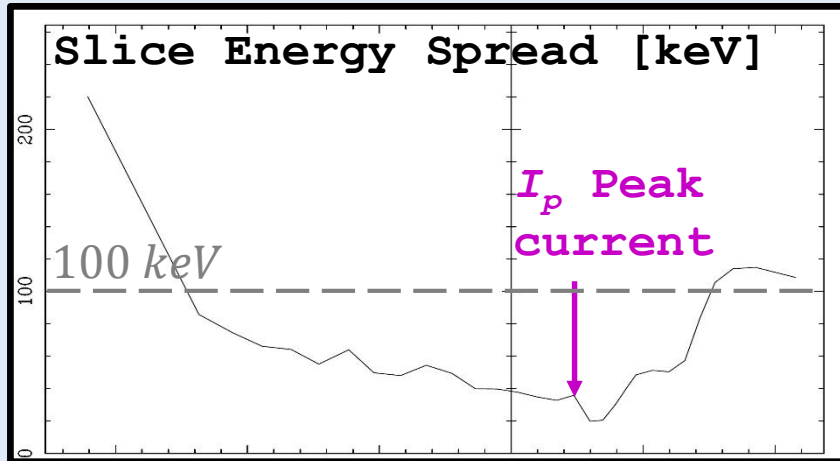
# A strong simplification of the longitudinal phase space modification

## Classic methods

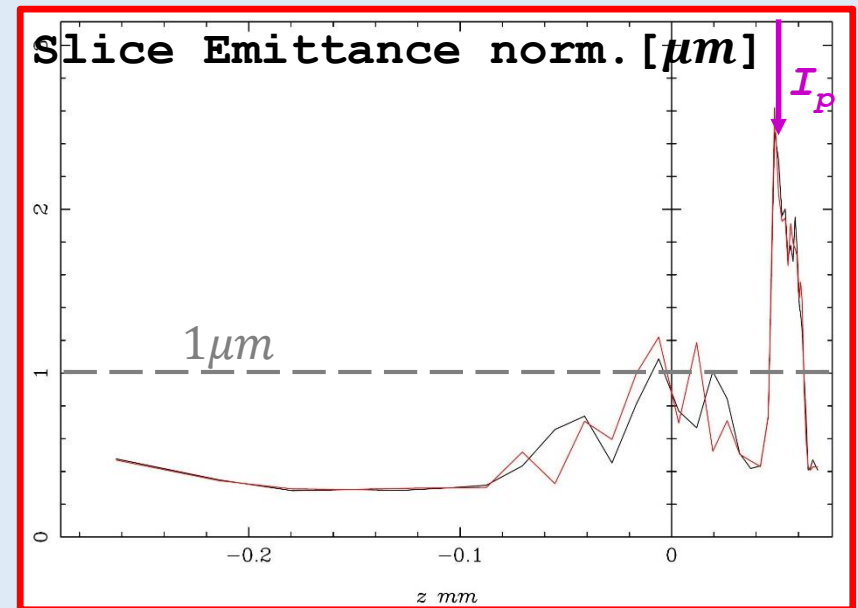
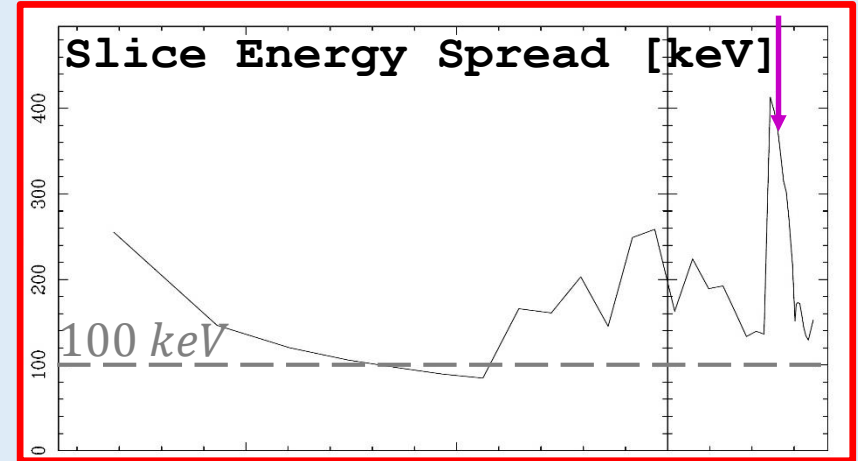


# Laminar "▒" & Velocity "▒" Bunching comparison

LB



LB



# Some Beam Dynamics

## Transverse envelope

$$\sigma'' + \frac{\gamma'}{\gamma} \sigma' + \left(\frac{k}{\gamma}\right)^2 \sigma = \frac{Q_c}{2I_A \gamma^3 \sigma_z \sigma} + \frac{\varepsilon_n^2}{\gamma^2 \sigma^3}$$

γ power of 3

## Long. envelope equation

$$\sigma_z'' + K_z \sigma_z + \frac{3\gamma' \sigma_z'}{\beta^2 \gamma} = \frac{Q_b c}{5\sqrt{5} I_A \beta^2 \gamma^4 \sigma_z \sigma} + \frac{\varepsilon_{nz}^2}{\beta^2 \gamma^6 \sigma_z^3}$$

Coupled by  $\sigma_x$

γ power of 4

## Laminar Parameters

$$\rho_{\perp} = \frac{Q_b c \sigma^2}{2I_0 \gamma \varepsilon_n^2}$$

Emittance  
compensations

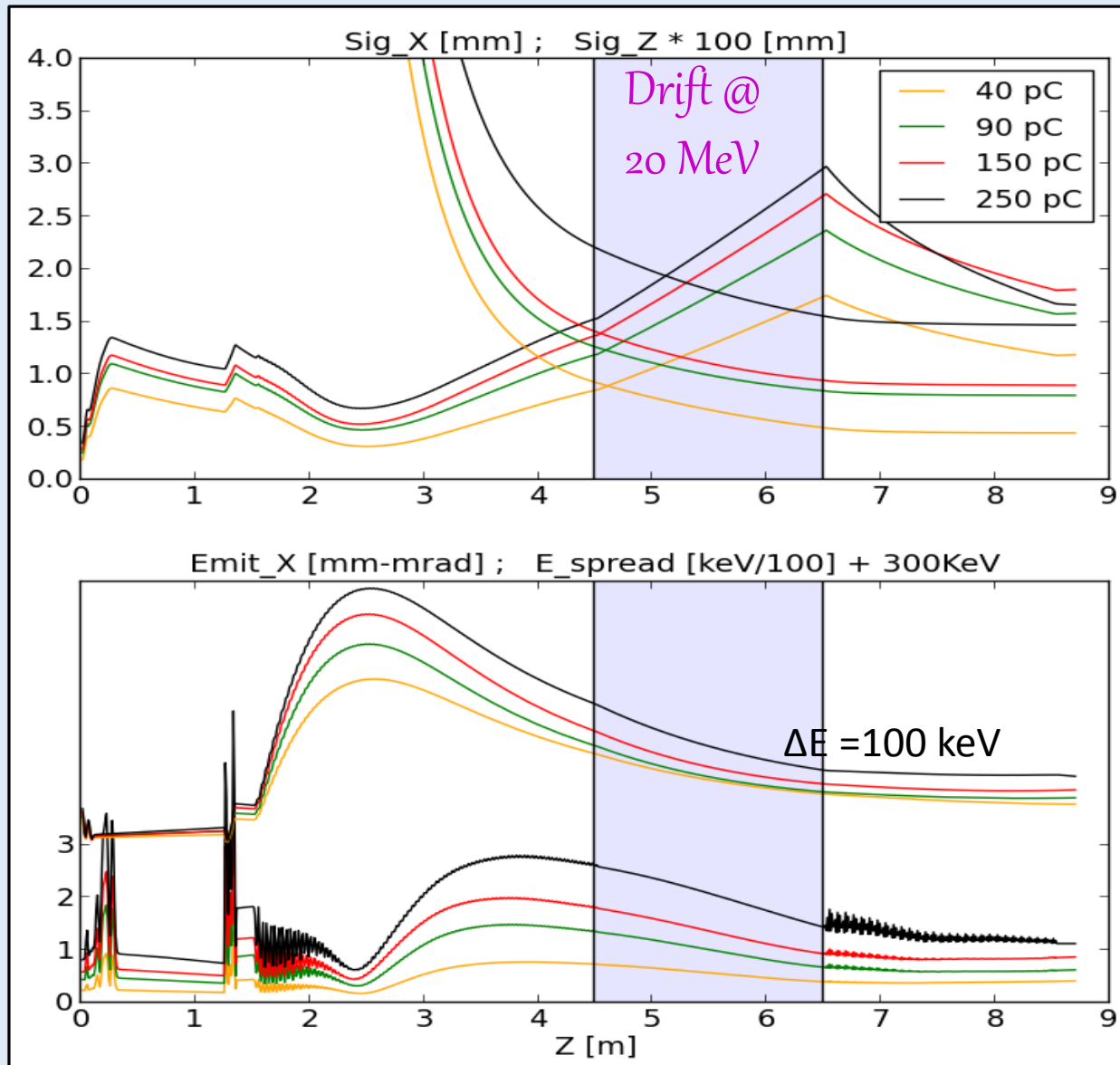
$$\rho_z = \frac{Q_b c (\gamma \sigma_z)^2}{I_0 \sigma \varepsilon_z^2}$$

1) Longitudinal  
compression

&

2) bunch stiffness  
respet to the  
compression

# LB performances VS. bunch-charge: 40;90;150;250 pC



@ booster end  
= 17 m (0.5 GeV)

$$Q_b = 40 \text{ pC}$$

$$\sigma_z = 4.2 \mu\text{m}$$

$$I_{\text{peak}} > 2.5 \text{ kA}$$

$$\epsilon_{\text{peak}} = 0.3 \mu\text{m}$$

$$B_{\text{peak}} = 3 \cdot 10^{16}$$

$$(I_p = 1.5 \text{ kA})$$

$$(\Delta\gamma/\gamma)_{@I_{\text{peak}}} \cong 8 \cdot 10^{-5}$$

# CONCLUSIONS

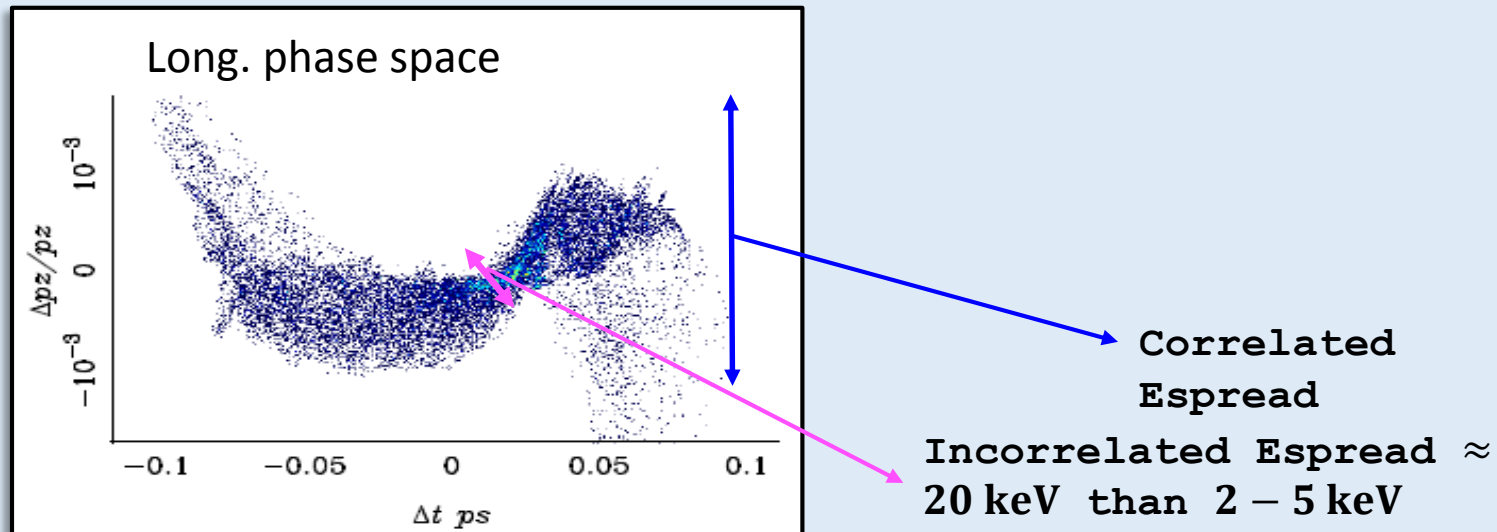
We saw a new compression technique: the Laminar Bunching

- ❑ A compression that works on the whole bunch distribution
- ❑ It is reproduced for a large range of charge: 40-250 pC
- ❑ ULTRA brightness and ULTRA low energy spread ( $10^{-4} \div 10^{-5}$ );  
A combination difficult to find!

- ❑ Drawback: Large envelopes for  $Q_b > 100$  pC.

An Ad-hoc large iris cavity can be used (rf-focusing knob)

- ❑ No Laser Heater (High un-correlated Energy Spread):



# Beam Dynamics study by using Genetic Algorithms

*Alberto Bacci and  
Marcello Rossetti Conti  
@ INFN-Milano*



# Outline

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## ❑ Genetic Algorithms (GAs) intro

## ❑ GAs applied to beamlines optimization

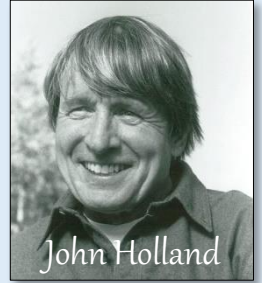
- From Beamlines to Chromosomes
- Following Genetic Laws: Fitness, Reproduction, Mutations, ...
- e.g.: SPARC beam line Optimization in Thomson case

## ❑ The GIOTTO code

- The GIOTTO Data-Based DB
- Inputs & Outputs
- Fitness function (or Idoneity) definition
- Optimizations & Statistics on some Specific Cases:
  - ✓ Ultra short bunches by Laminar Bunching
  - ✓ Comb bunches distributions

# □ Intro: Genetic Algorithms **Historical Notes**

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❖ 1970 John Holland - schemata theorem

❖ 1975 J. Holland publication: *"Adaptation in Natural and Artificial Systems: An Introductory Analysis with Application to Biology, Control, and Artificial Intelligence"*. **The Seminal work**

❖ 1975 K. De Jong (J. Holland's student), Thesis: *"An analysis of the behavior of a class of genetic adaptive systems"*.  
Broad applicability of GAs

❖ 1989 David Goldberg Book: *"Genetic Algorithms in Search, Optimization, and Machine Learning"*

It deals with **the topic at high level** and **is considered a milestone in GAs story**. It reports techniques like Multi Objective GA (MOGA), today very trendy.

# Intro: What are Genetic Algorithms (GAs)?

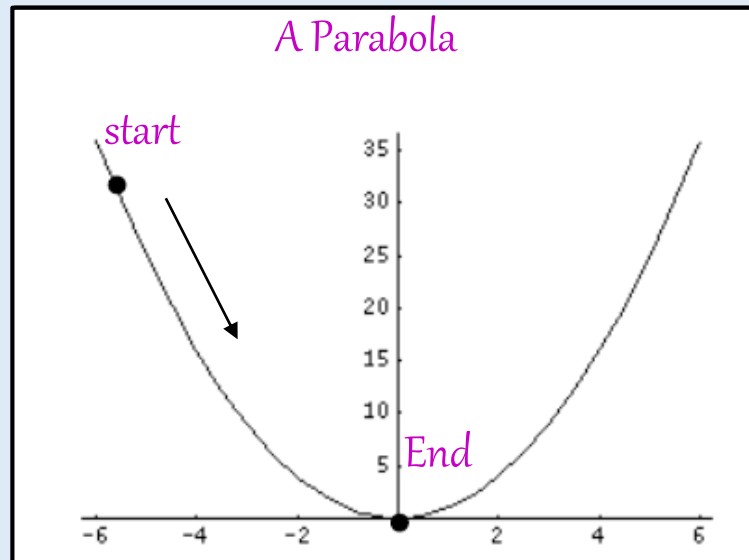
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**A Searching procedure** based on a **natural selection**  
(genetic laws)

Why choosing GAs versus other techniques?

A basic answer:

Newton-Raphson methods (or **variants**) are based on local information. The Scan moves in direction of local-maxima or local-minima



# Intro: Why Genetic Algorithms?

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Despite, Newton-Raphson methods can overcome the local solutions issue by some tricks ...

## **GAs are:**

- naturally able to manage the local solutions issue
  - naturally parallelizable
  - usable with a minimum mathematical effort
- by empirical results, show strong capability in problems where other methods fail

**pros and cons**  
**a suggested pub.**

## Genetic Algorithm Optimization Applied to Electromagnetics: A Review

Daniel S. Weile and Eric Michielssen, *Member, IEEE*

IEEE TRANSACTIONS ON ANTENNAS AND PROPAGATION, VOL. 45, NO. 3, MARCH 1997

# □ GAs in Beam Dynamics Optimizations

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**GAs give strong advantages in**

- **multi-dimensional problems** with variables strongly non-linearly correlated

**Two main examples :**

- **Space Charge & its non-linear nature:** correlates low energy beamline parameters
- **Also frozen beams** (space charge off), e.g.: **complex matching lines** (plasma accelerated beams)

**example:** **Thomson/Compton sources** (e.g. SPARC\_lab, STAR, ThomX, ELI-np, Munich Compact Light Source) **ask for :**

- **High spectral density** that means: **very low DE/E, low Emit**
- **High the photon flux** that means: **high  $Q_{\text{bunch}}$ , small spot size**

## □ ... or other examples

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considering **ultra short e-beams** (e.g. SPARC\_lab, LCLS, REAGE, XFEL, EUPRAXIA, ... ):

- **Femtosecond light pulses (FEL/X-FEL)**, Atoms in chemical reactions, phase-transition, *Photosynthesis Water Splitting* : timescale 1-100 fs [2014 “first snapshots of water splitting” by LCLS; ScienceDaily; Nature]
- **Plasma Wave Acceleration:**  $\lambda_{\text{plasma}}$  order of 30-600 fs . The **Witness** much shorter, the **Driver** (pwfa) comparable to  $\lambda_{\text{plasma}}$
- **Femtosecond Electron Diffraction (FED)**  
**Molecular or atomic motion movies:** phase transitions, ..., . Timescale: few 10s of fs. Relativistic case:  $E_b \approx 5\text{MeV}$ ,  $Q_b \approx 100\text{fC}$ ,  $\varepsilon_{\text{tr}} < 0.1$  mm-mrad,  $\sigma_z < 30\mu\text{m}$  (100fs)
- **THz radiation (by CoTrRad)**  
0.1 up to tens THz is of great interest for both **longitudinal electron beam diagnostics** (fs scale) and **spectroscopy** in pump-and-probe experiments ....

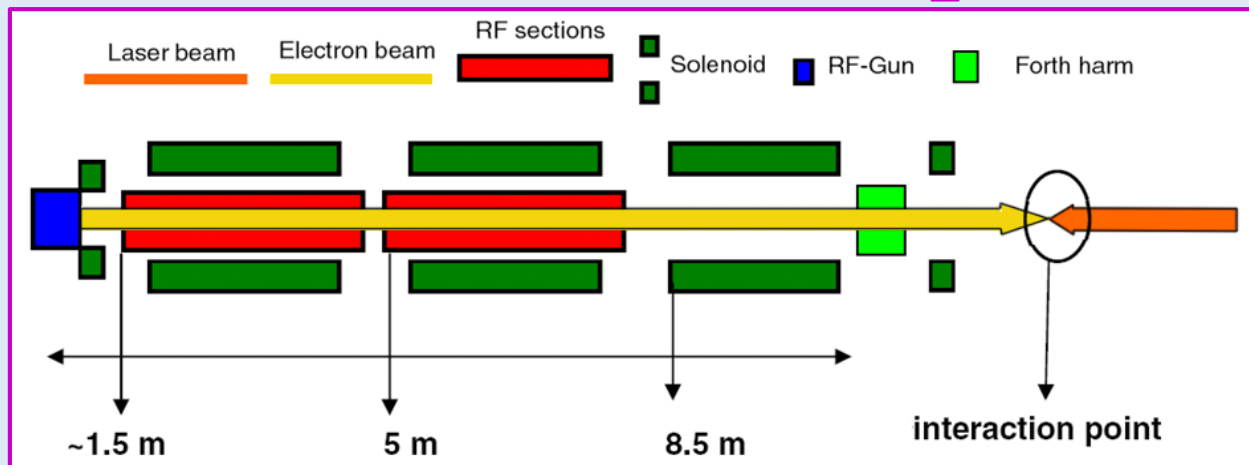
# Genetic Algorithms applied to Beam-Line Optimization

# ❑ From **beamlines** to **chromosomes**

Genetic laws work on **Chromosomes** ==> **Chromosomes** are made of **genes** (parameters)

Beamline setup  $\Leftrightarrow$  A genes array **.or.** One Chromosome

A BeamLine (exp. Thomson @ SPARC\_lab)



=

Chromosome

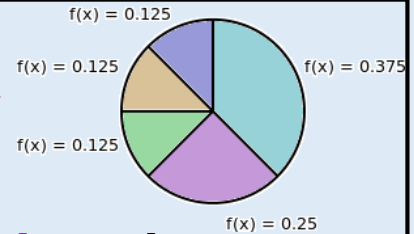
Gene	Reference case
$dE_g / dz$ (MV/m)	120
$\phi_g$ (°)	0
$B_g$ (T)	0.2707
$dE_1 / dz$ (MV/m)	13.4
$\phi_1$ (°)	-30
$B_1$ (T)	0.12
$z_1$ (m)	1.322
$dE_2 / dz$ (MV/m)	6.55
$\phi_2$ (°)	88
$B_2$ (T)	0.1145
$z_2$ (m)	5
$B_3$ (T)	0.1145
$z_3$ (m)	8.5
$\phi_{IVH}$ (°)	180
$z_{IVH}$ (m)	11.7



# Following genetic laws: the Fitness Function

Gene	Reference case
$dE_g / dz$ (MV/m)	120
$\phi_g$ (°)	0
$B_g$ (T)	0.2707
$dE_1 / dz$ (MV/m)	13.4
$\phi_1$ (°)	-30
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$\phi_2$ (°)	88
$B_2$ (T)	0.1145
$z_2$ (m)	5
$B_3$ (T)	0.1145
$z_3$ (m)	8.5
$\phi_{IH}$ (°)	180
$z_{IH}$ (m)	11.7

To pass generation → in generation:  
 -> Selection: bluffed Roulette wheel  
 -> Mutations

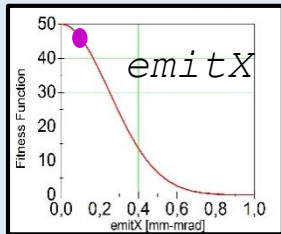


... & others methods & tricks. The rule: closest to Nature, best performances

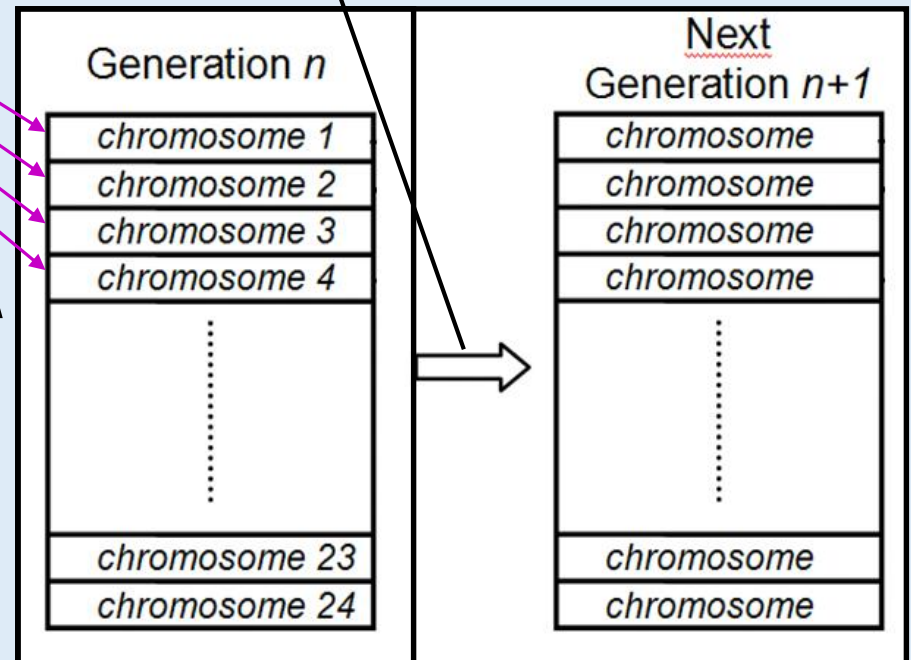
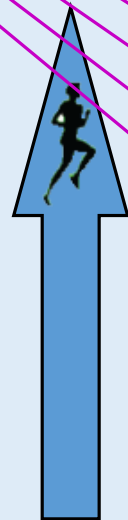
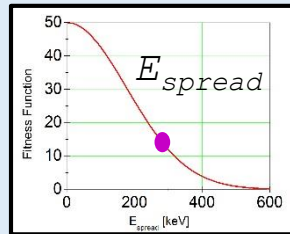
Starts from random generation

Chromosomes  sorting

$$\text{runner icon} == 50 \cdot e^{-\left(\frac{emitX}{0.35}\right)^2} + 50 \cdot e^{-\left(\frac{E_{spread}}{250}\right)^2}$$



+

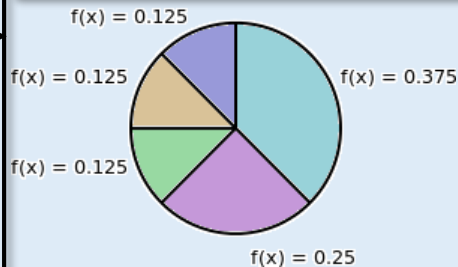


# ➤ Following genetic laws: Reproduction & Mutation

Chromosome Sorting

$$0 < \text{runner} < 100$$

By the Roulette Wheel:  
**two** Chromosomes



$$\sum f_p(x) = 1$$

Generation  $n$

chromosome 1  
chromosome 2  
chromosome 3  
chromosome 4

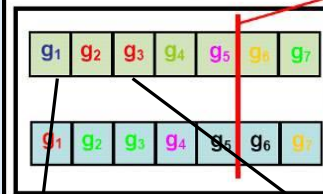
chromosome 23  
chromosome 24

Astra

99.  
97.  
90.  
85.  
...  
60.  
58.

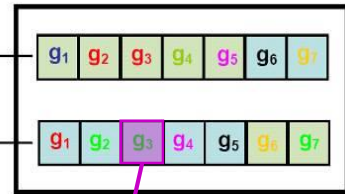
Reproduction by cross-over oper.

Elements of  $n^{\text{th}}$  generation



cross-over  
random  
function

Elements of the  $n+1^{\text{th}}$  generation



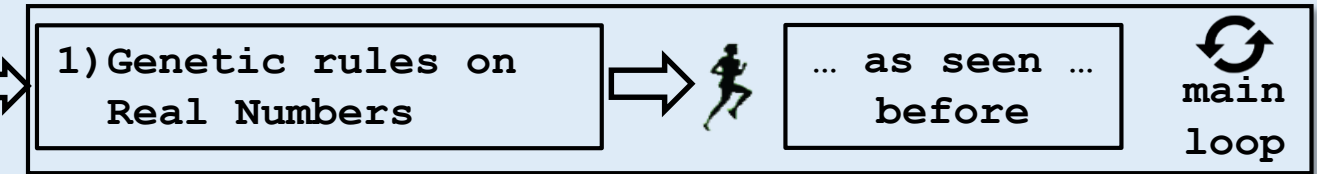
$\text{Gun}_{\Phi} \text{ inj.}$

$\text{Gun}_{\text{gradient}}$

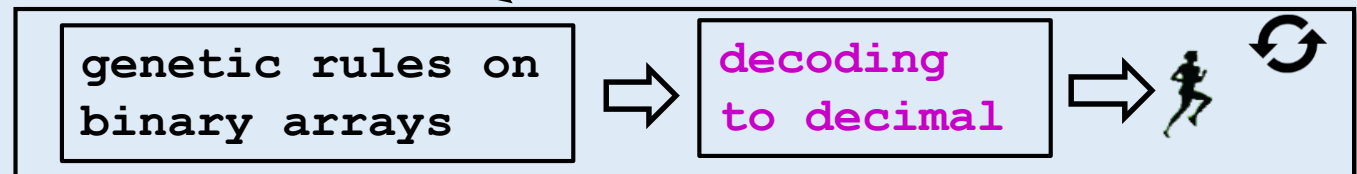
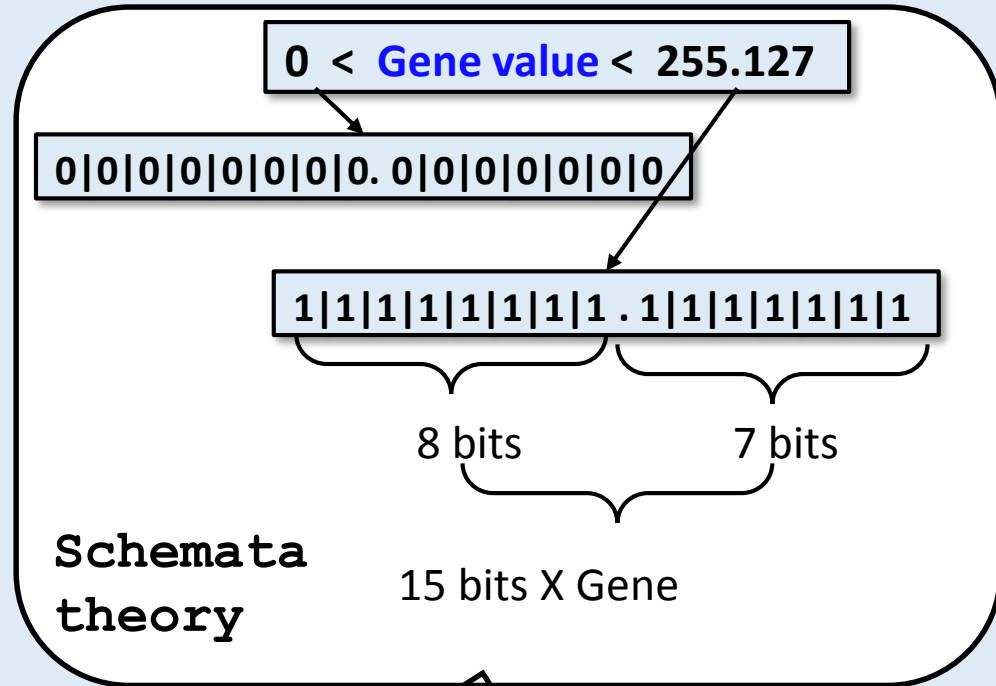
Mutation, with  
probability  $< 1\%$

# ➤ Following genetic laws: **real coding** & the **binary one**

Chromosome	
Gene	Reference case
$dE_g / dz \text{ (MV/m)}$	120
$\phi_g (^{\circ})$	0
$B_g \text{ (T)}$	0.2707
$dE_1 / dz \text{ (MV/m)}$	13.4
$\phi_1 (^{\circ})$	-30
$B_1 \text{ (T)}$	0.12
$z_1 \text{ (m)}$	1.322
$dE_2 / dz \text{ (MV/m)}$	6.55
$\phi_2 (^{\circ})$	88
$B_2 \text{ (T)}$	0.1145
$z_2 \text{ (m)}$	5
$B_3 \text{ (T)}$	0.1145
$z_3 \text{ (m)}$	8.5
$\phi_{IVH} (^{\circ})$	180
$z_{IVH} \text{ (m)}$	11.7

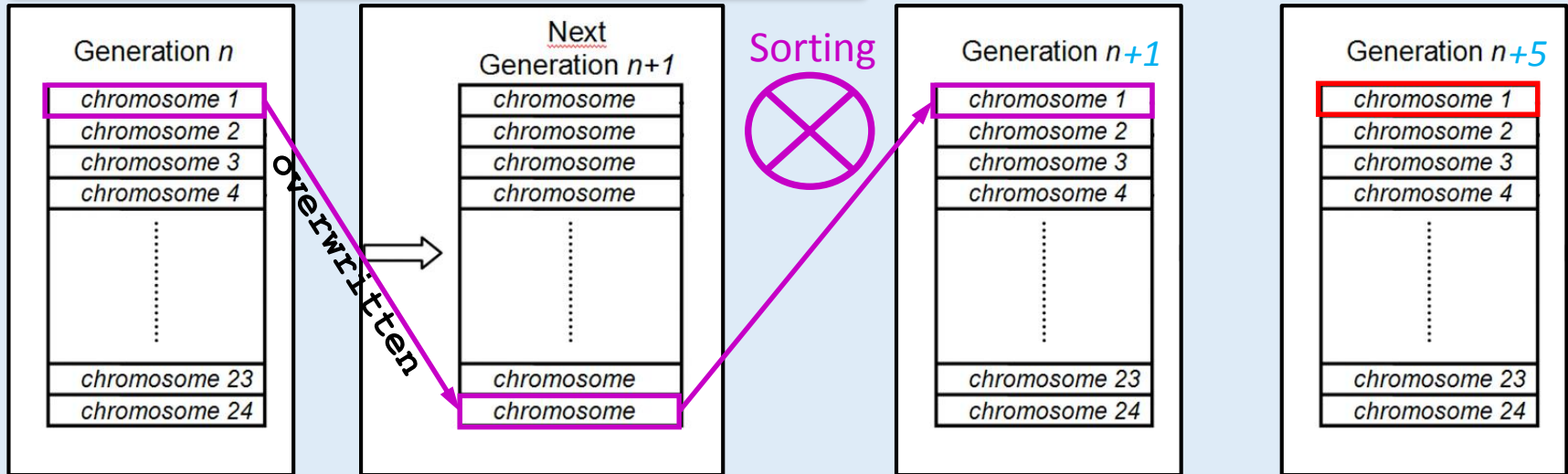


2) Binary Coding



## ➤ Following Genetic Laws: Elitism

best chromosome is kept in next generations unless it improves



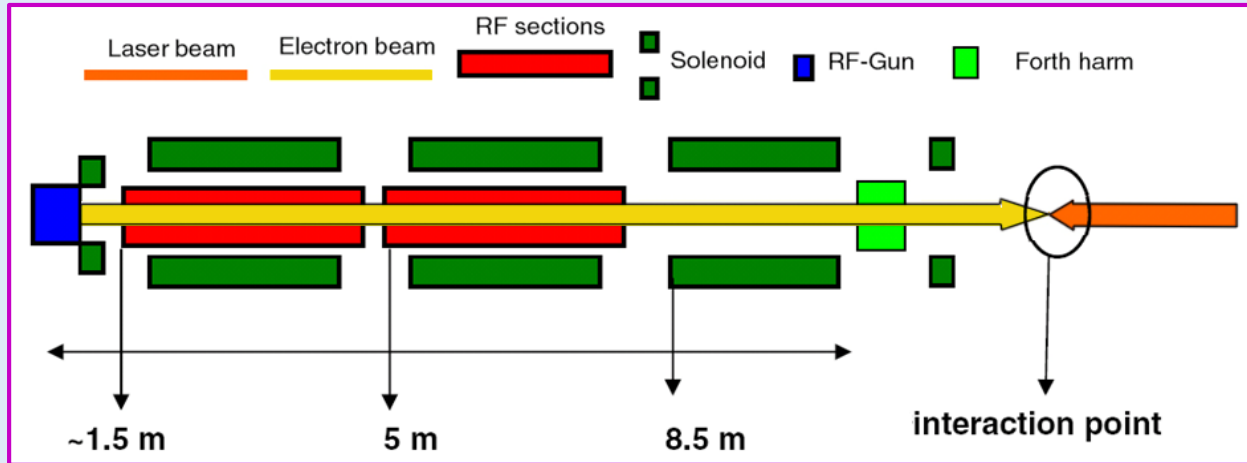
“quasi-classical” optimization techniques:

- elitism
- advanced mutation operators
- hill climbing
- regeneration from best solutions
- ... ..

parallelization is-mandatory

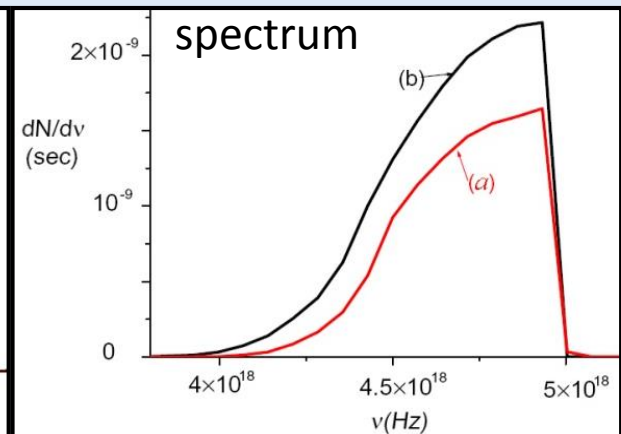
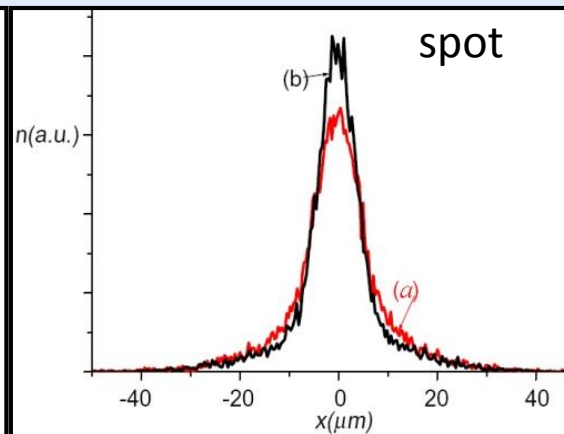
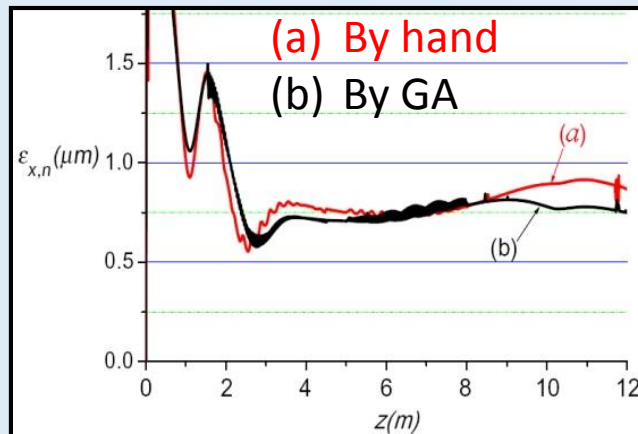


# ➤ SPARC beamline optimization in Thomson case



Old Kind of Fitness Function

$$F_{fitness} = \frac{1}{\epsilon_{x,n} \cdot \frac{\Delta\gamma}{\gamma}}$$



**“MAXIMIZING THE BRIGHTNESS OF AB ELECTRON BEAM BY MEANS OF A GENETIC ALGORITHM”**

A. Bacci, C. Maroli, V. Petrillo, A. Rossi, L. Serafini - NIMB 263 (2007) 488-496

The  
GIOTTO  
code

# GIOTTO: Genetic Interface for OpTimising Tracking with Optics

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- ❖ The code was born in 2008;      Language: Fortran 90/95
- ❖ ... born for optimization of beamline parameters & statistical (Jitters) analysis
- ❖ INPUTS based on NameList & two internal DataBase
- ❖ CAN easily Drive different codes (NameList natively):  
Now: ASTRA's Generator,      ASTRA,      QFluid (Plasma acceleration, A. Rossi modifications)
- ❖ Current Version (Ver. 18.4):  
Linux & Windows 64 bit - (compilers gfortran or INTEL fortran compiled). Parallelized: OPEN-MPI (Linux), MS-MPI or INTEL MPI (Windows)  
Tested @ PSI (S. Bettoni), @ Desy-Pitz (Martin), LAL (Luca, Harsh)
- ❖ Code and Documentations:  
URL: <http://pcfasci.fisica.unimi.it/Pagine/GIOTTO/GIOTTO.htm>  
(server down, pardon!)  
Exist an User manual for version 8.5 2012 (needs updates)

# GIOTTO – Genetic Interface for Optimising Tracking with Optics

From 2008 up to day, the code is grew in power and capability

Optimization techniques: elitism; advanced mutation operators; hill climbing; ant colony; regeneration from best solutions  
Targets: bunch PosZ, bunch Time, En, En<sub>spread</sub>, SigZ, Xemit, sigX, divergX, Yemit, ...

## Important GIOTTO's features:

Every NameList 's variables can be used as a GENE (optimizable) & Any code working with NameList is directly importable in GIOTTO.

ASTRA: Phi(1)...Phi(50), MaxE(1:50), MaxB(1:50), sig\_x (laser cathode), sig\_clock (Laser @ cathode), ..., ... (no limit on the number)

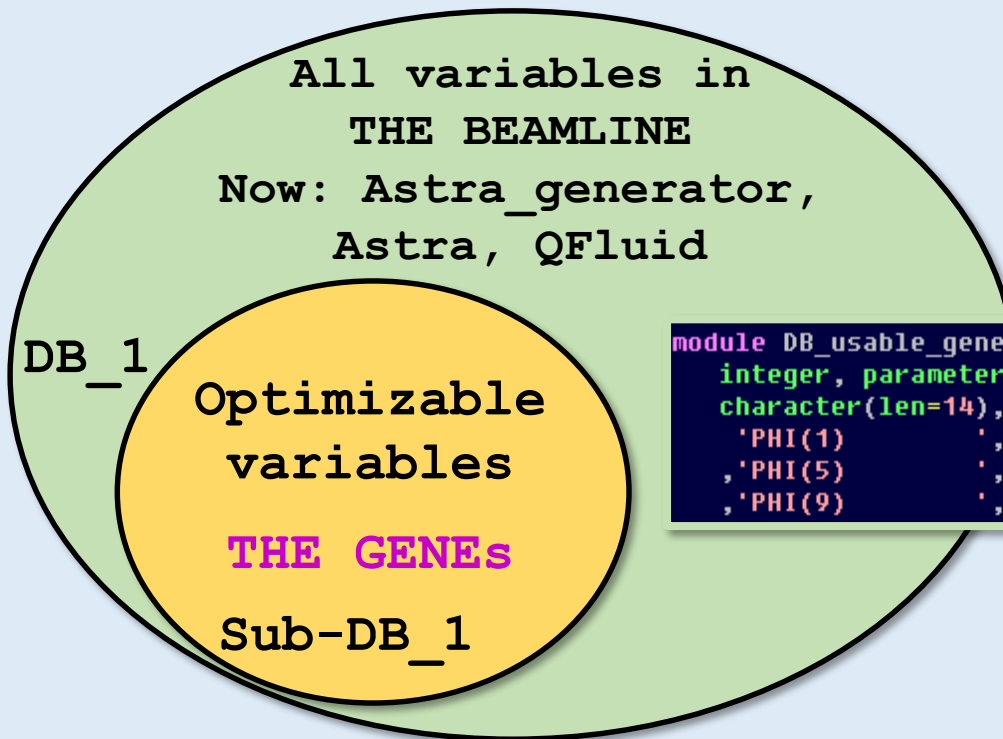
Constraints freely defined by the user (under test)

switches from Optimizations to Statistical analysis

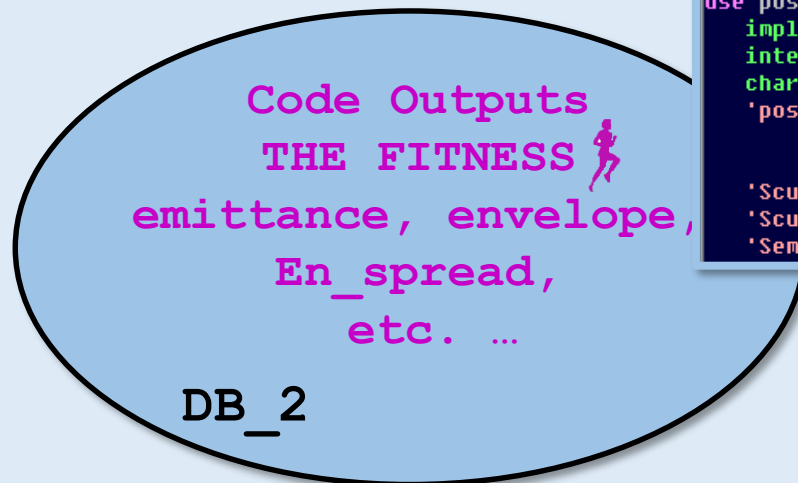
Jitters sampling interval Uniform or Gaussian



# ➤ GIOTTO's Data-Bases



```
module DB_usable_gene
  integer, parameter :: DB_extension=269
  character(len=14), dimension(DB_extension) :: DB_genes=(/ &
    'PHI(1)      ', 'PHI(2)      ', 'PHI(3)      ', 'PHI(4)      ', '&
    'PHI(5)      ', 'PHI(6)      ', 'PHI(7)      ', 'PHI(8)      ', '&
    'PHI(9)      ', 'PHI(10)     ', 'PHI(11)     ', 'PHI(12)     ', '&
```



```
module charge_rpn_idon      !rpn reverse polish notation
  use post_processors
  implicit none
  integer, parameter, private :: DB_out_dim=107
  character(len=10), dimension(DB_out_dim), private :: DB_out=(/ &
    'pos2      ', 'time      ', 'En      ', 'sigZ      ', 'DEn      ', 'emit2      ',
    'Xmed      ', 'sigX      ', 'divergX    ', 'emitX      ',
    'Ymed      ', 'sigY      ', 'divergY    ', 'emitY      ',
    'Scurr(1)   ', 'Scurr(2)   ', 'Scurr(3)   ', 'Scurr(4)   ', 'Scurr(5)   ', '&
    'Scurr(6)   ', 'Scurr(7)   ', 'Scurr(8)   ', 'Scurr(9)   ', 'Scurr(10)  ', '&
    'SemitX(1)  ', 'SemitX(2)  ', 'SemitX(3)  ', 'SemitX(4)  ', 'SemitX(5)  ', '&
```

## ➤ GINxx.xx.ini GIOTTO's INPUT FILE

It is divided in two parts:

- 1) A **NameList** (&GA) giving all the directive to GIOTTO
- 2) Three **keyNames** defining: **CONSTRAINTS**, **FITNESS** and **GENES**

```
1) {
    &GA
    Astra_in='Astra_23_Jan2014.exe','pls-start.in'
    ....
    .....
    /

    =====From here Key_Names=====
    =====(lines after one blank-record are comments)=====
    =====

    [constr01]
    sigZ En * sqr emitZ sqr sigX * / 5300 * 1 >

    [constr02]
    sigZ En * sqr emitZ sqr sigX * / 6300 * 10 >

    [idoneity] !must be used the Reverse Polish Notation
    emitX 2.5 / sqr -1. * exp 50 *
    sigZ 0.150 / sqr -1. * exp 50 * +
    sigX 2.5 / sqr -1. * exp 50 * +

    !Gene(i)___Delta___JoinGenes(i:i+N)___u-uniform___g-gaussian___JoingSign
    [genes]
    sig_clock    0.2E-3    1      u      1    0.0 0
    SIG_x        0.02     1      u      1    0.0 0
    MaxB(1)      0.08     1      u      1    0.0 0
    MaxB(2)      0.08     1      u      1    0.0 0
}
```

## ➤ INPUT FILE: &GA NameList

```
&GA
Astra_in='Astra_23_Jan2014.exe','pls-start.in'
Gener_in='generator Mar2013.exe','generator-start1.in'
Genes_number=7 !max value is 40
pop_size=8 !must to be a multiple of the nodes numbers
generations_number=410

keepADistribution=.false. !".f." start from "Gener_in", ".t." from Distribution
keepInParameters=.true.
!".t." the chrom in the first gen;".f." All rnd, the chrom as central values

!*****turn on and Control the constraints
constraints=.false. !if "true" turn-on constraints-----
minimum2=1.0 !this is valid for all constraints
constr_number=2 !max number of constraints is 10
constr_name='[constr01]','[constr02]'
lower_Zbound=1.6,5.0,16.0 !intervals where the constraint
upper_Zbound=25.0,25.0,25.0

!*****Post processor for COMB bunch distributions*****
LaserComb=.false. !if true Gener_in is a generator input with SPIKES namelist--
!if "true" uses "ID_bunch_LC" for the spikes analysis and compute the Fitness--
LaserComb_PostPro=.false.
PostPro_in_index=0012

!*****Forced Optimization Process*****
step_OP_forced=25
variation_x100=5.0 !Best Cromosom Variation %. Every "step_OP_f
!-----variation in % is halved----untill "step_OP_forced" X "Step2start_x100"
Step2start_x100=4

!*****Statistical Analysis*****
statistic=.false.
Runs_Number=360 !must be a multiple of the pop_size
/
```

Usually  
few  
variables  
are used

Under  
developing (it  
slows down  
heavily GIOTTO)

Optimization  
Rarely needs  
changes

Fitness  
&  
Genes

# ➤ INPUT FILE: **Key\_Names**

```
=====From here Key_Names=====
===== (lines after one blank-record are comments)=====

!*****Equation in Reverse Polish Notation*****
!-----Variables that can be used are:-----
!-----PosZ time En DEn sigZ emitZ Xmed sigX divergX emitX Ymed sigY-----
!-----divergY emitY-----
!*****

[constr01]
sigZ En * sqr emitZ sqr sigX * / 5300 * 1 >

[constr02]
sigZ En * sqr emitZ sqr sigX * / 6300 * 10 >

[idoneity] !must be used the Reverse Polish Notation
emitX 2.5 / sqr -1. * exp 50 *
sigZ 0.150 / sqr -1. * exp 50 * +
sigX 2.5 / sqr -1. * exp 50 * +

emitX 0.4 - 0.05 / sqr -1. * exp 50 * sigZ 0.280 - 0.01 / sqr -1. * exp 50 * +
sigX sigY - 1.0 / sqr -1. * exp 50 * + Xmed 1000 - 10.0 / sqr -1. * exp 50 * +
SemitX(1) 4 / sqr -1 * exp SemitX(2) 4 / sqr -1 * exp
Sdist(1) 380E-6 - 100E-6 / sqr -1 * exp 50 * +

!Gene(i)___Delta___JoinGenes(i:i+N)___u-uniform___g-gaussian___JoinSign
[genes]
sig_clock    0.2E-3    1      u      1    0.0 0
SIG_x        0.02     1      u      1    0.0 0
MaxB(1)      0.08     1      u      1    0.0 0
MaxB(2)      0.08     1      u      1    0.0 0
Phi(1)       2.0      1      u      1    0.0 0
Phi(2)       2.0      1      u      1    0.0 0
Phi(3)       10.0     1      u      1    0.0 0

MaxE(5)      5.0      1      u      1    0.0 0
c_pos(5)     0.05     1      u      1    0.0 0
Q_grad(4)    0.002    1      u      1    0.01 Q_grad(6) 0
Zstop        0.05     1      u      1    0.0 0
```

Rarely  
needed

Comments

Comments

Definition

GENES  
Definition

# ➤ GIOTTO: FITNESS FUNCTION

```
[idoneity] !must be used the Reverse Polish Notation
emitX 2.5 / sqr -1. * exp 50 *
sigZ 0.150 / sqr -1. * exp 50 * +
sigX 2.5 / sqr -1. * exp 50 * +
```

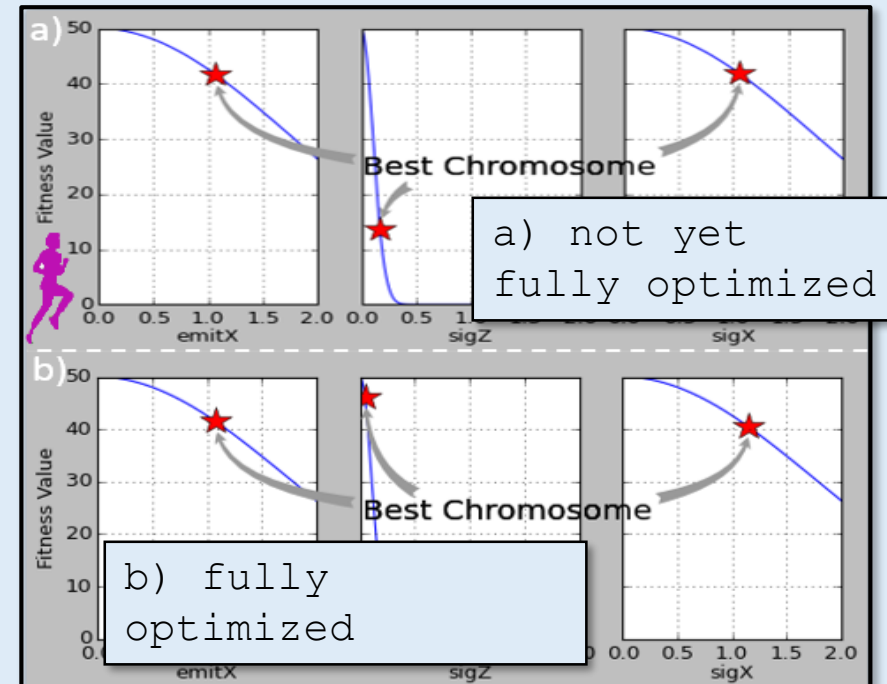
## Reverse Polish Notation:

Operands Follow Operations



- 3 4 + = 7
- Stack based operation
- Does not need brackets



$$50 \cdot e^{-\left(\frac{\text{emitX}}{2.5}\right)^2} + 50 \cdot e^{-\left(\frac{\text{sigZ}}{0.15}\right)^2} + 50 \cdot e^{-\left(\frac{\text{sigX}}{2.5}\right)^2}$$



## Strategy to Cope with Multi Objectives Problems (MO):

- o One Single Criterium per Equation piece (Objectives Wights)
- o Close to the Goal mean close to the Gaussian Curve Top
- o The 'Far region' (referring to optimization) has to be on the maximum Gaussian slope
- o Change the  in real time (under implementation)
- o Lorentzian  dist.: very powerful if starting from scratch

## ➤ FITNESS FUNCTION if starting from scratch

Lorentzian much better than Gaussian

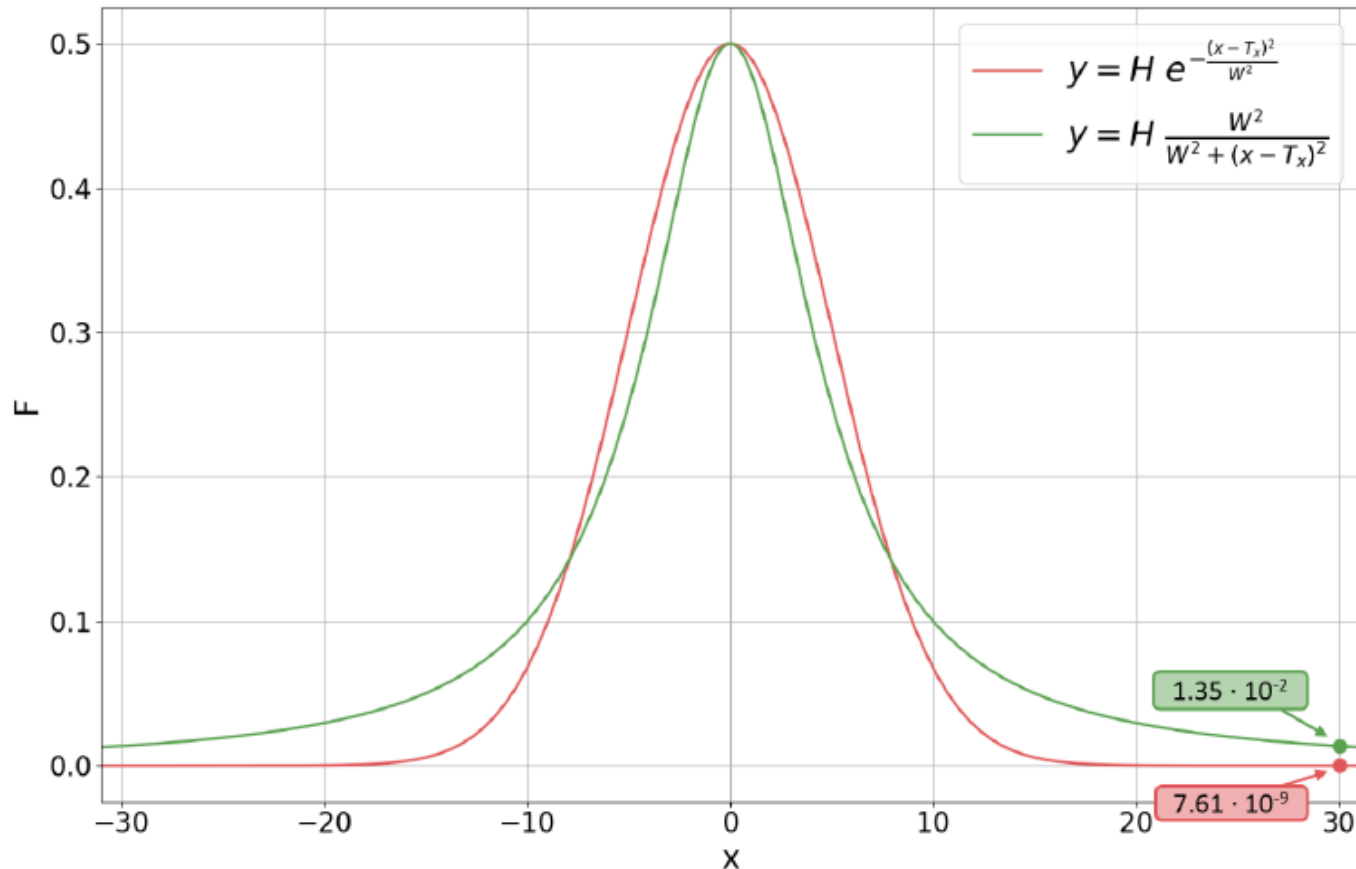


Figure 3.4: The figure shows the comparison between a Gaussian and a Lorentzian. The height, width and target value parameters are the same in both the functions ( $H = 0.5$ ,  $W = 5$ ,  $T_x = 0$ ). Both functions are evaluated in the point  $x = 30 = 6W$ .

# GIOTTO "RISULTATI.TXT" output file



```
[idoneity] !must be used the Reverse
emitX 1.2 / sqr -1. * exp 100 *
sigZ 0.016 / sqr -1. * exp 400 * +
DEn 200.0 / sqr -1. * exp 50 * +
```

GENES

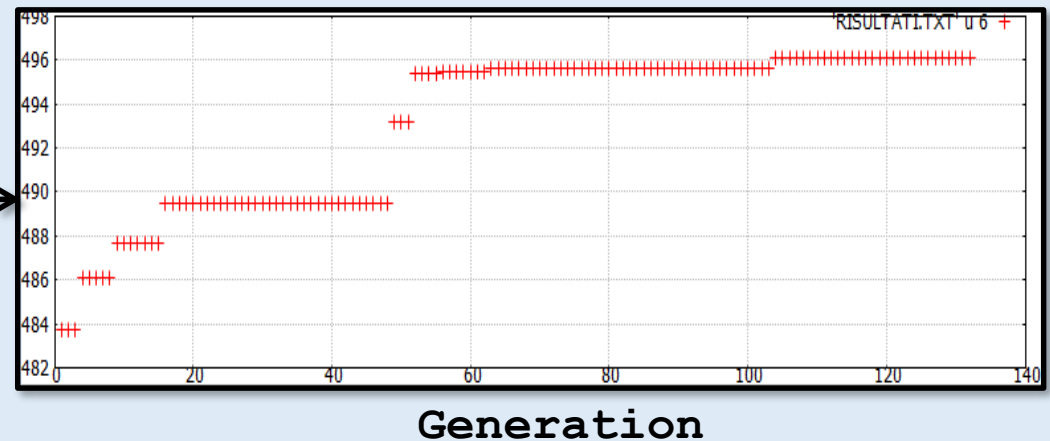
```
[genes]
MaxB(1) 0.003 1 u 1 0.0 0
MaxB(2) 0.01 1 u 1 0.0 0
Phi(1) 1.2 1 u 1 0.0 0
Phi(2) 1.5 1 u 1 0.0 0
Phi(3) 1.5 1 u 1 0.0 0
```

## RISULTATI.TXT

	MAXB(1)	MAXB(2)	PHI(1)	PHI(2)	PHI(3)	***id best***	***id_worst***	Emit	sigZ	$\Delta E$
1	MAXB(1)	MAXB(2)	PHI(1)	PHI(2)	PHI(3)	***id best***	***id_worst***	emitX	sigZ	DEn
2	3.33234E-01	5.76714E-02	2.34351E+00	1.91001E+02	-8.48726E+01	4.83694E+02	7.83022E+01	5.79140E-01	5.08770E-03	7.80340E+01
3	3.33234E-01	5.76714E-02	2.34351E+00	1.91001E+02	-8.48726E+01	4.83694E+02	1.72762E+02	5.79140E-01	5.08770E-03	7.80340E+01
4	3.33234E-01	5.76714E-02	2.34351E+00	1.91001E+02	-8.48726E+01	4.83694E+02	3.32879E+02	5.79140E-01	5.08770E-03	7.80340E+01
5	3.33234E-01	5.76714E-02	2.18337E+00	1.90543E+02	-8.50834E+01	4.86096E+02	3.06747E+02	6.46100E-01	4.62500E-03	7.56980E+01
6	3.33234E-01	5.76714E-02	2.18337E+00	1.90543E+02	-8.50834E+01	4.86096E+02	3.27456E+02	6.46100E-01	4.62500E-03	7.56980E+01
7	3.33234E-01	5.76714E-02	2.18337E+00	1.90543E+02	-8.50834E+01	4.86096E+02	2.90264E+02	6.46100E-01	4.62500E-03	7.56980E+01
8	3.33234E-01	5.76714E-02	2.18337E+00	1.90543E+02	-8.50834E+01	4.86096E+02	2.92544E+02	6.46100E-01	4.62500E-03	7.56980E+01
9	3.33234E-01	5.76714E-02	2.18337E+00	1.90543E+02	-8.50834E+01	4.86096E+02	3.01542E+02	6.46100E-01	4.62500E-03	7.56980E+01
10	3.33234E-01	5.76714E-02	2.34351E+00	1.90543E+02	-8.50834E+01	4.87645E+02	4.29286E+02	6.16760E-01	4.63930E-03	7.69790E+01
11	3.33234E-01	5.76714E-02	2.34351E+00	1.90543E+02	-8.50834E+01	4.87645E+02	4.31203E+02	6.16760E-01	4.63930E-03	7.69790E+01
12	3.33234E-01	5.76714E-02	2.34351E+00	1.90543E+02	-8.50834E+01	4.87645E+02	4.40076E+02	6.16760E-01	4.63930E-03	7.69790E+01
13	3.33234E-01	5.76714E-02	2.34351E+00	1.90543E+02	-8.50834E+01	4.87645E+02	4.40076E+02	6.16760E-01	4.63930E-03	7.69790E+01

Best Chromosome  
of the  
Generation N.5

Id value

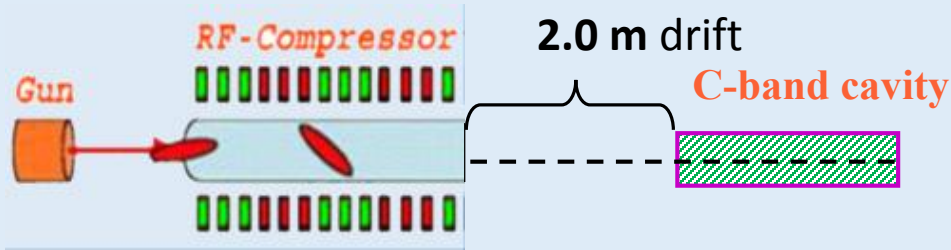


GIOTTO

Optimization  
&  
Statistical analysis

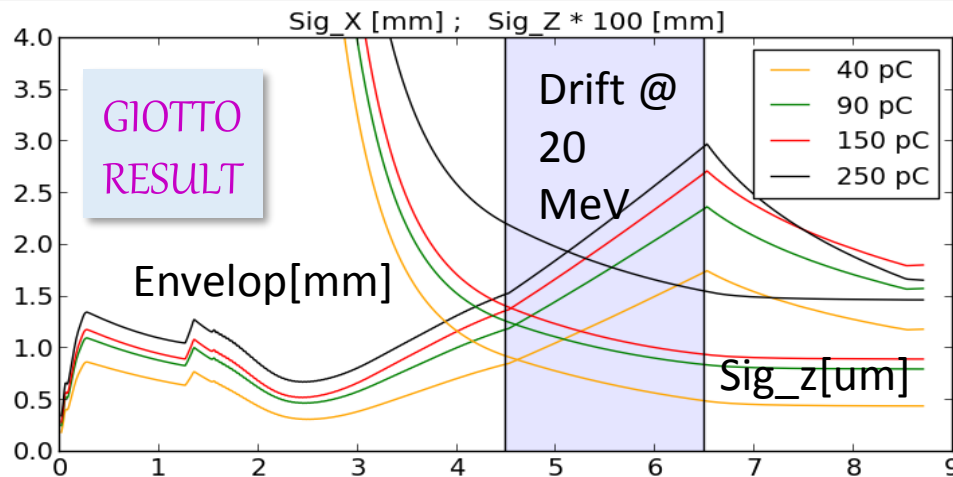


# Beamline Optimization for: ultra short, ultra cold, High brightness bunches



## GOALS :

- Low Emittance
- Low Energy Spread
- Sig\_Z hundred of nm

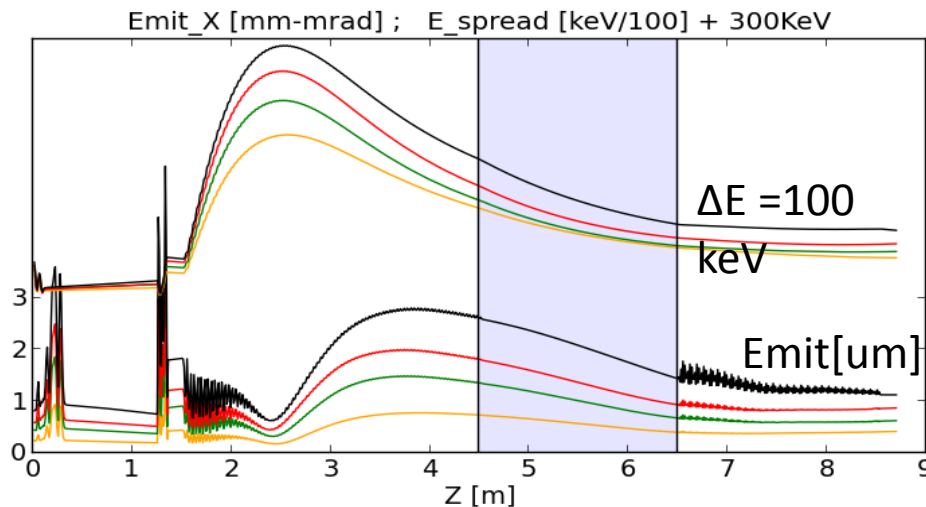


## A Beam-Line studied with:

- Experience
- An Ad hoc GIOTTO use

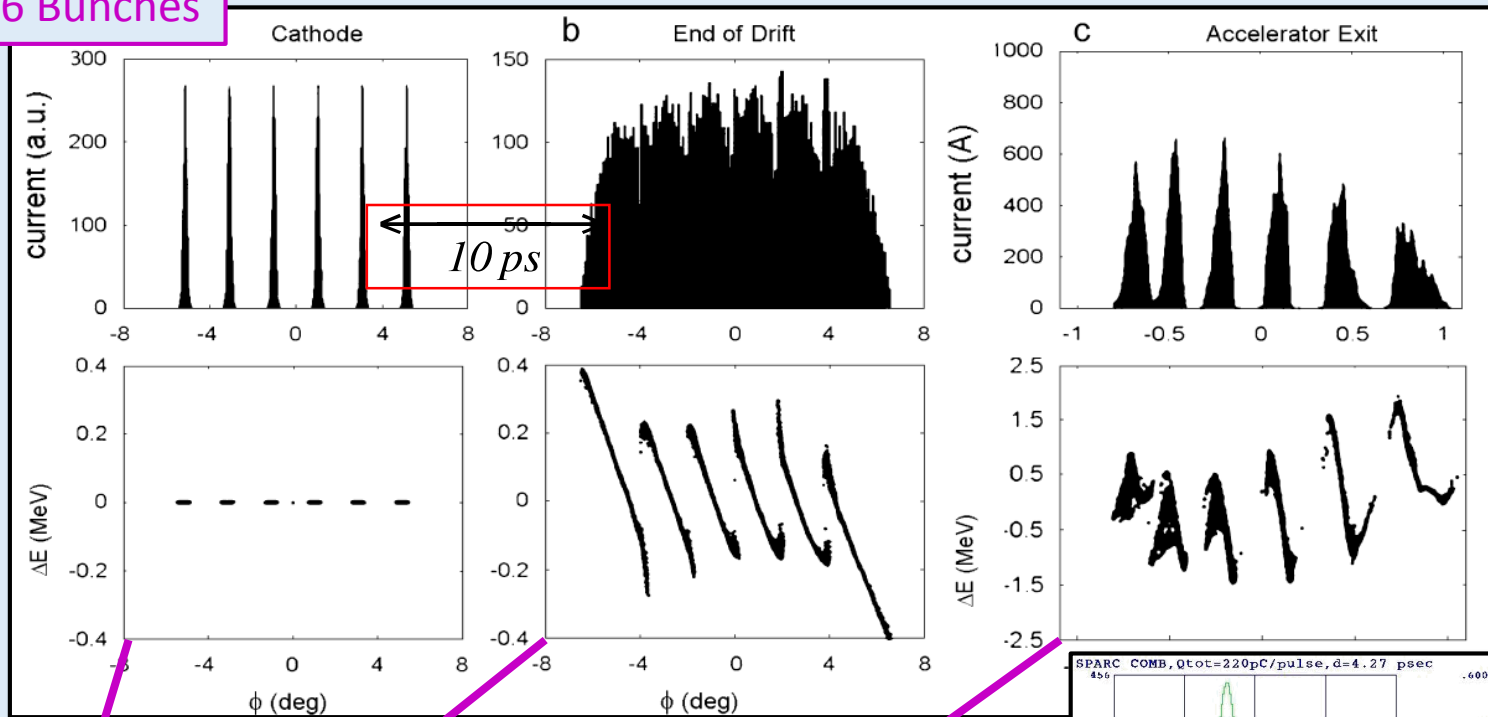
## GENES in the Optimization:

- Gun:
  - (1) Phase & (2) Solenoid ( $B_z$ )
- TW cavity (RF- Compressor):
  - (3) Phase & (4) Solenoids
- C-band cavity: (5) Phase

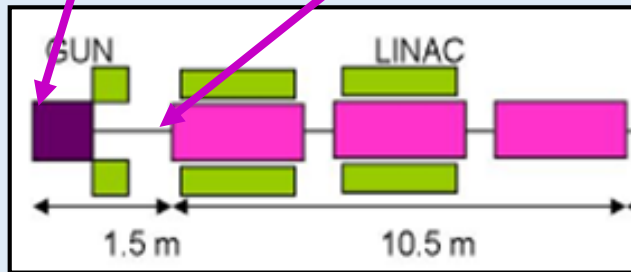
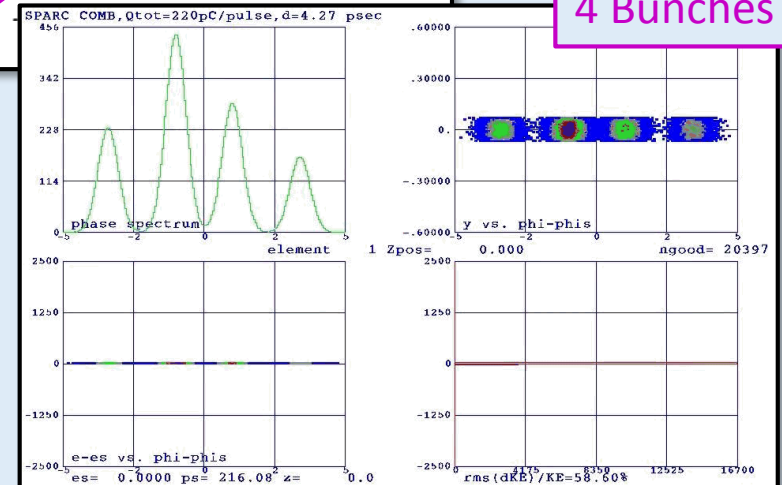


# Beam-Line STATISTIC for Laser Comb (ECHO Bunch Generations)

6 Bunches



4 Bunches

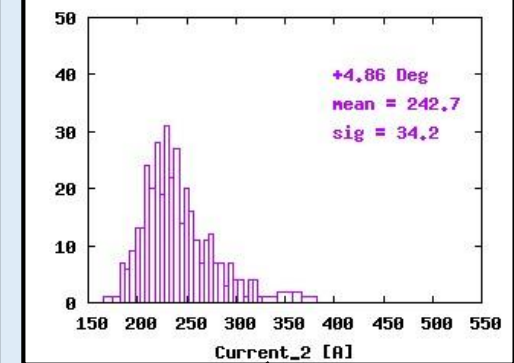
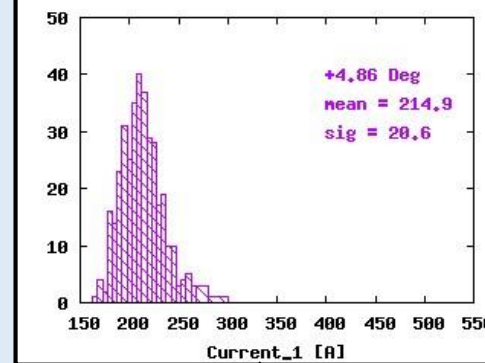


- P.O.Shea et al., Proc. of 2001 IEEE PAC, Chicago, USA (2001) p.704.
- M. Ferrario. M. Boscolo et al., Int. J. of Mod. Phys. B, 2006 (Taipei 05 Workshop)

# Beamline STATISTIC Laser Comb two **bunches case**

```
!*****Statistical Analysis*****
statistic=.true.
Runs_Number=360      !must be a multiple of the pop_size
!*****
```

```
[genes]
phi(1)    0.9      1      g      1
phi(2)    0.9      1      g      1
```



360 cases

bacci@pcsimul2:~/Doc_Lavoro/08_SPARC/03_LaserComb/02_Simulazioni/05_Giotto/02b_Titti_comparison/02_result_p5										
5	posZ	time	En	sigZ	DEn	emitZ	zXEpMed	Xmed		
6	12.0000000	40.1020012	127.5899963	0.1931200	315.2000122	49.8079987	181.1999969	-0.0012153		
7	12.0000000	40.0989990	127.5899963	0.1988300	287.4100037	47.6329994	158.7799988	-0.0011859		
8	12.0000000	40.1010017	127.5999985	0.1907200	329.6600037	51.0439987	192.4600067	-0.0012634		
9	12.0000000	40.0989990	127.5899963	0.2004300	282.3500061	47.5709991	152.9400024	-0.0012552		
10	12.0000000	40.0999985	127.5899963	0.2022500	275.6199951	47.2900009	145.9199982	-0.0012090		
11	12.0000000	40.0970001	127.6399994	0.1682500	445.8599854	58.4290009	279.6300049	-0.0011903		

sigX	divergX	emitX	xXxpMed	Ymed	sigY	divergY	emitY
0.1552100	0.0777850	1.6316000	0.0655140	0.0003971	0.1546800	0.0778750	1.6328000
0.1575600	0.0799780	1.5186000	0.0701300	0.0004734	0.1574000	0.0801450	1.5223000
0.1577300	0.0770300	1.7099000	0.0637480	0.0002847	0.1572100	0.0771520	1.7097000
0.1560300	0.0800730	1.4665999	0.0707510	0.0005361	0.1559900	0.0802450	1.4737000
0.1585100	0.0807810	1.4794000	0.0716900	0.0005785	0.1585100	0.0810230	1.4844000
0.1906800	0.0691680	2.5023999	0.0452270	0.0004415	0.1896900	0.0691130	2.5088999
0.1604000	0.0805670	1.5132999	0.0712350	0.0005018	0.1602400	0.0807310	1.5175000

yXypMed	Scurr01	Scurr02	SemtX01	SemtX02	SemtY01	SemtY02	Sdist01	Sdist02
0.0655090	216.6100006	234.6699982	1.0002000	1.6055000	1.0131000	1.5854000	0.0003813	0.0000000
0.0702470	227.5399933	218.2200012	1.0061001	1.6256000	1.0204000	1.6002001	0.0003928	0.0000000
0.0638020	209.6300049	244.7700043	1.0125999	1.7193000	1.0254000	1.7012000	0.0003765	0.0000000
0.0708450	230.9100037	215.6799927	0.9868600	1.6118000	1.0041000	1.5850000	0.0003960	0.0000000
0.0718970	234.2200012	212.6699982	1.0067000	1.6318001	1.0206000	1.6045001	0.0003997	0.0000000
0.0446640	170.4299927	359.5400085	1.0303000	1.6545000	1.0482000	1.6513000	0.0003305	0.0000000

# Quads matching lines for plasma acc. beams – from scratch

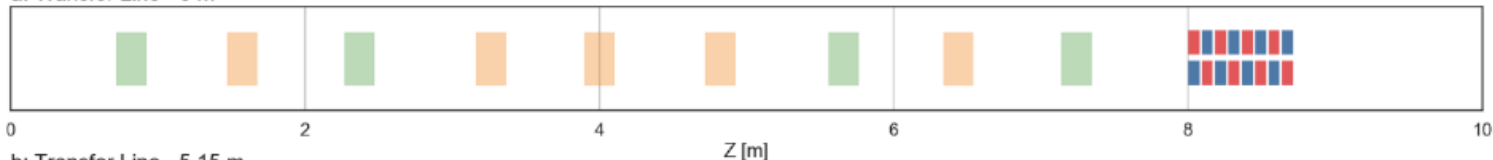
EUPRAXIA project  
From  
Marcello Thesis

**Table 3.3:** Magnetic elements in the 4 different TLs shown in Fig. 3.10

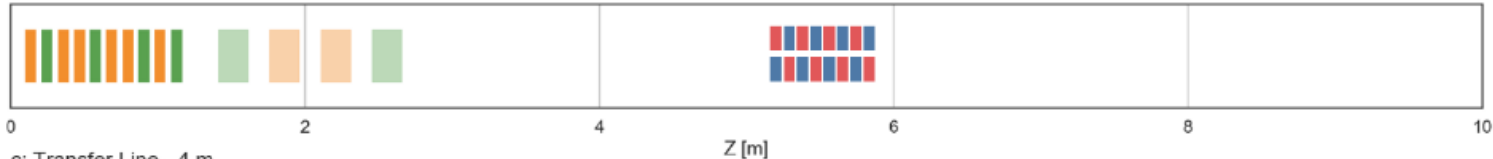
	Len.	EMQ	EMQ len.	EMQ bore	PMQ	PMQ len.	PMQ bore
A	8 m	9	20 cm	1 cm			
B	5.15 m	4	20 cm	1.5 cm	10	7 cm	1 cm
C	4 m	4	20 cm	1.5 cm	6	7 cm	1 cm
D	3 m	4	20 cm	1.5 cm	6	7 cm	1 cm

GIOTTO starts with all K=0 and matched all the lines

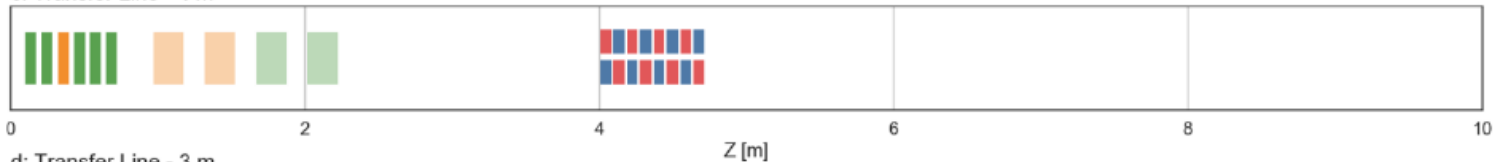
a: Transfer Line - 8 m



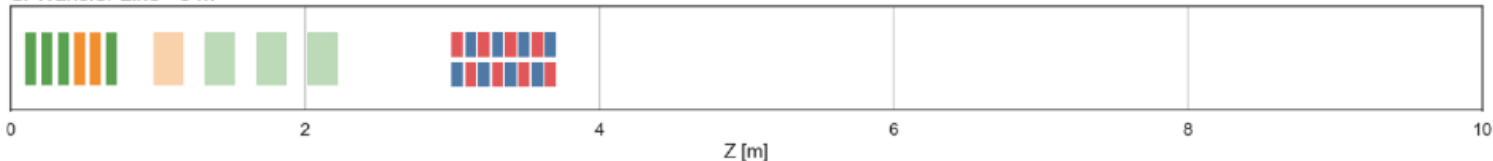
b: Transfer Line - 5.15 m



c: Transfer Line - 4 m

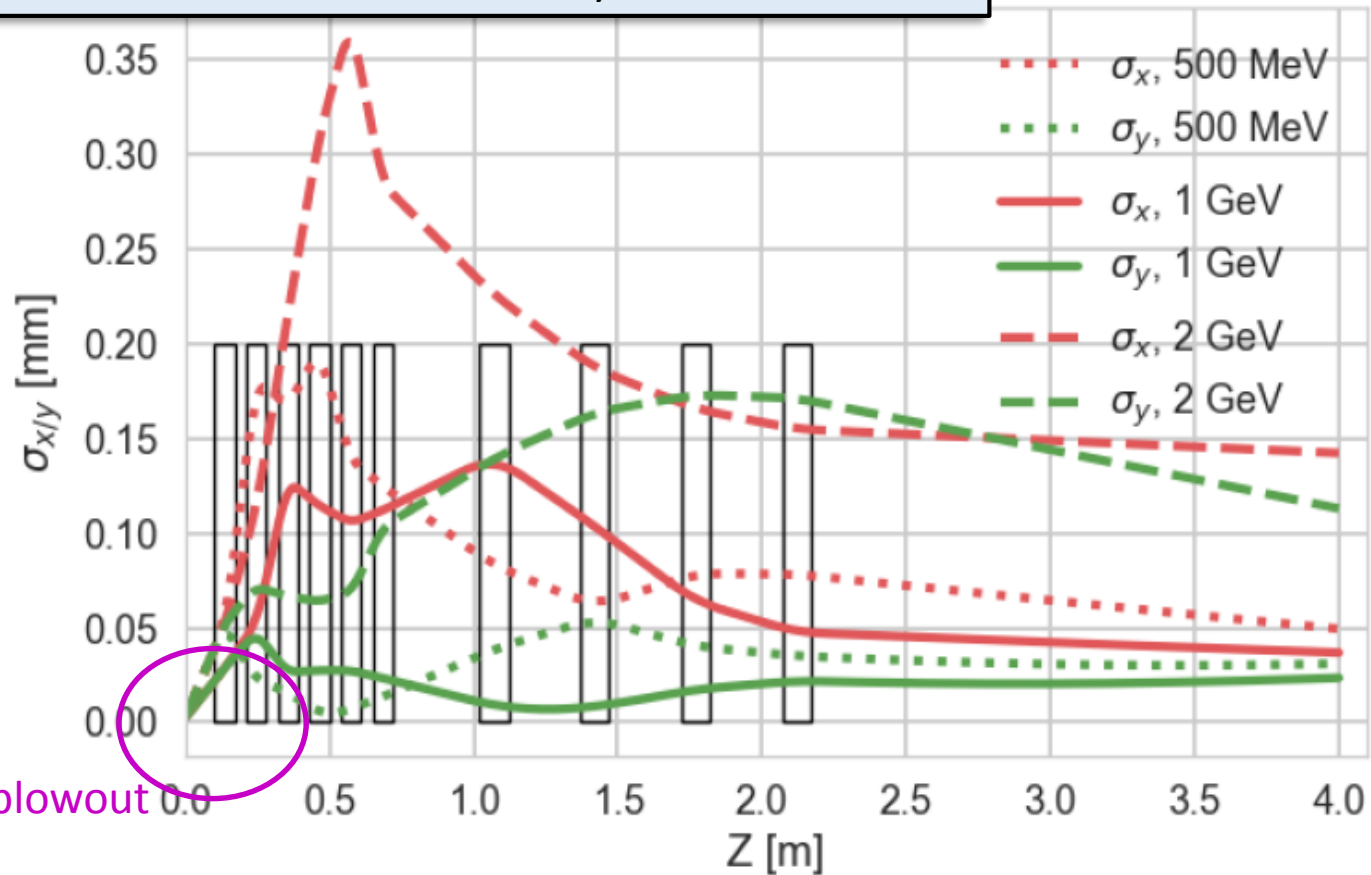


d: Transfer Line - 3 m



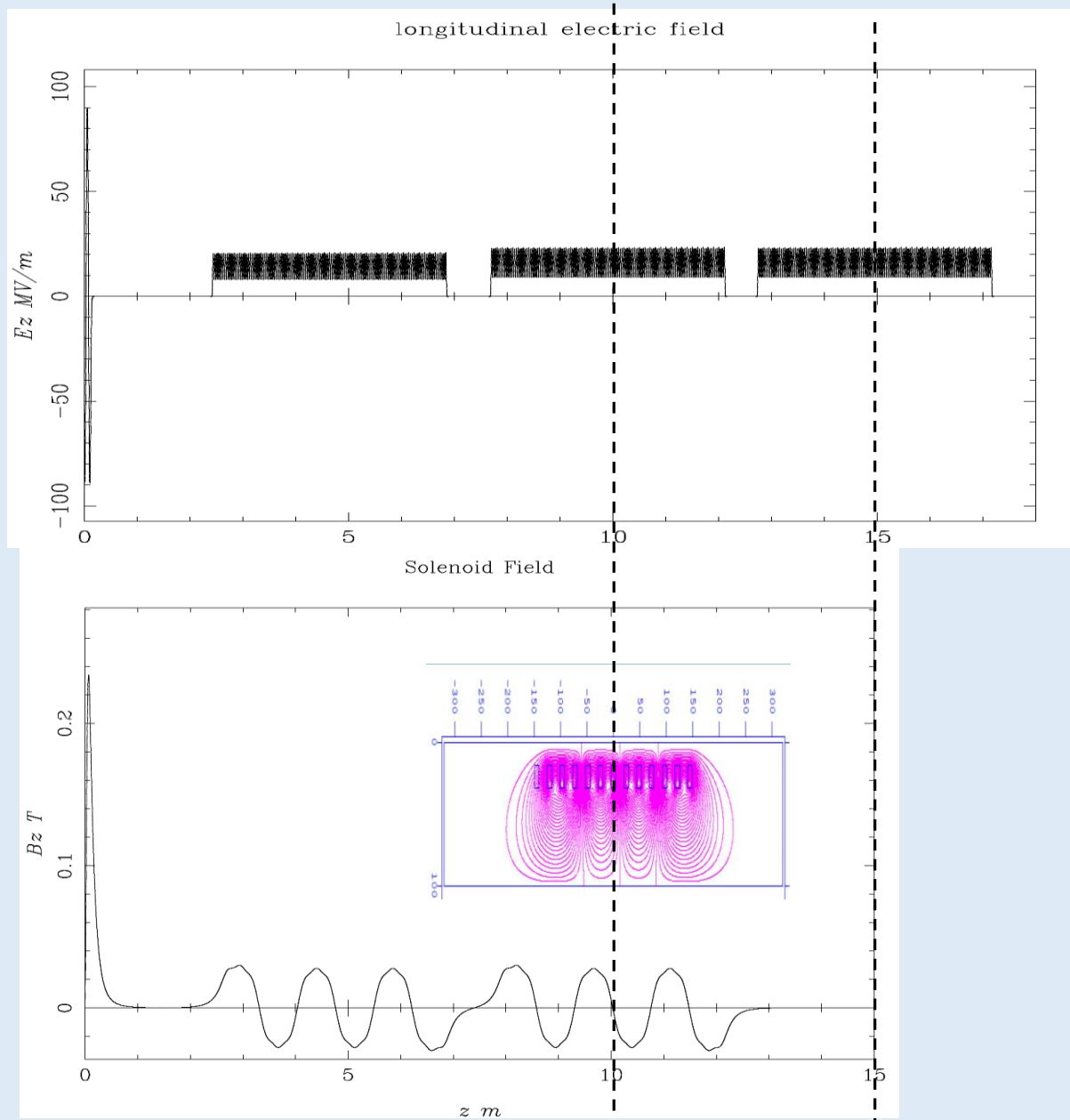
matching lines for plasma acc. Beams (same line diff. energy)

Isseu: the emittance conservation by chromatic effects



**Figure 3.11:** Comparison between the beams envelopes for matched bunches of significantly different energies. The position and the dimension of the elements of the line is the same in the three cases (rectangles), only the quads strengths are changed.

# Velocity bunching test on CLEAR machine (1)



# Velocity bunching test on CLEAR machine (2)

```
&NEWRUN
  RUN=2
  Distribution = 'Q_80_FD_300_AB.ini'
  TRACK_ALL=T, AUTO_PHASE=T, PHASE_SCAN=T
  check_ref_part=F,
  RefS=T,
  EmitS=T,
  PhaseS=T,
  Lmagnetized=F,
  H_max=0.002
  ZSTART=0.00, ZSTOP=18.00,
  Zemit=1800,
  Zphase=1,
/
&CHARGE
  LSPCH=T,
  Nrad=5, Nlong_in=10,
  Cell_var=2.0,
  ?min_grid=0.40-6,
  min_grid=0.0,           ?AB
  Max_scale=0.1,
  Max_count=20           ?AB
  Lmirror=T
/
&CAVITY
  LEfield=T,
  File_Efield(1)= 'sonde_ideal_SF_100.txt', phi(1)=0.0, maxE(1)=90.0, nue(1)=2.99855, c_pos(1)=0.0
  File_Efield(2)= 'TWS_LIL.txt', phi(2)=-40, maxE(2)=20.5, nue(2)=2.99855, c_numb(2)=132, c_pos(2)=2.3904
  File_Efield(3)= 'TWS_LIL.txt', phi(3)=0, maxE(3)=23.0, nue(3)=2.99855, c_numb(3)=132, c_pos(3)=7.6592
  File_Efield(4)= 'TWS_LIL.txt', phi(4)=0, maxE(4)=23.0, nue(4)=2.99855, c_numb(4)=132, c_pos(4)=12.7079
/
&SOLENOID
  LBfield=T,
  File_Bfield(1) = 'profile.txt',           MaxB(1)=0.234153, S_pos(1)=0.00
  File_Bfield(2) = 'TW_LIL_Solen_4VB.dat', MaxB(2)=0.03, S_pos(2)=4.640
  File_Bfield(3) = 'TW_LIL_Solen_4VB.dat', MaxB(3)=0.03, S_pos(3)=9.9092
/
```

really negligible Velocity Bunching.  
compression starts @  $\phi = -80^\circ$   
don't know machine behaviour I  
don't want to lose the beam

# Velocity bunching test on CLEAR machine (3)

```
&&GA
Astra_in='Astra_Mar2018.exe','phi_3LIL_Clear.in'
Gener_in='', ''
Genes_number=6    !max value is 40
pop_size=32       !must to be a multiple of the nodes numbers
generations_number=1600

use_generator=.false.
del_NORRAN=.false.    !if .true. the Generator starts from a static seed
keepInParameters=.false.
!.true.: the starting values are used as a cromosom,
!.false.: " ...." are used only as sampling intervals central values

!***
!--
!--
!--
!--
!--
!--
!***
[ide]
2 sqr emitX  sqr 2 sqr + / 50 *
1 sqr sig2  sqr 1 sqr + / 50 * +

5 sqr Den 1 -  sqr 5 sqr + / 60 * +
0.4 sqr emitY 0.8 -  sqr 0.4 sqr + / 150 * +

!Gene(i)___Delta___JoinGenes(i:i+N)___u-uniform___g-gaussian___JoingSign
[genes]
Phi(1)      3.0      1      u      1  0.0 0 !buncher
Phi(2)     15.0      1      u      1  0.0 0 !booster
Phi(3)     30.0      1      u      1  0.0 0 !booster
MaxB(1)     0.015     1      u      1  0.0 0 !solenoid peak field
MaxB(2)     0.02      1      u      1  0.0 0 !solenoid peak field
MaxB(3)     0.03      1      u      1  0.0 0 !solenoid peak field
```

- The optimization after few runs:  
bunch\_length= 2ps, emit: 1.2 mm-mrad
- After 4 hours (16 core working station -2012 machine-):  
bunch\_length=0.5ps, emit:0.9 mm-mrad



## Concluding: some publications to deepen the topics discussed

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Daniel S. Weile and Eric Michielssen “**Genetic Alforithm Optimization Applied to Electromagnetics: A Review**”, IEEE Trans. On Antennas and Propagation, Vol 45, NO. 3, March 1997

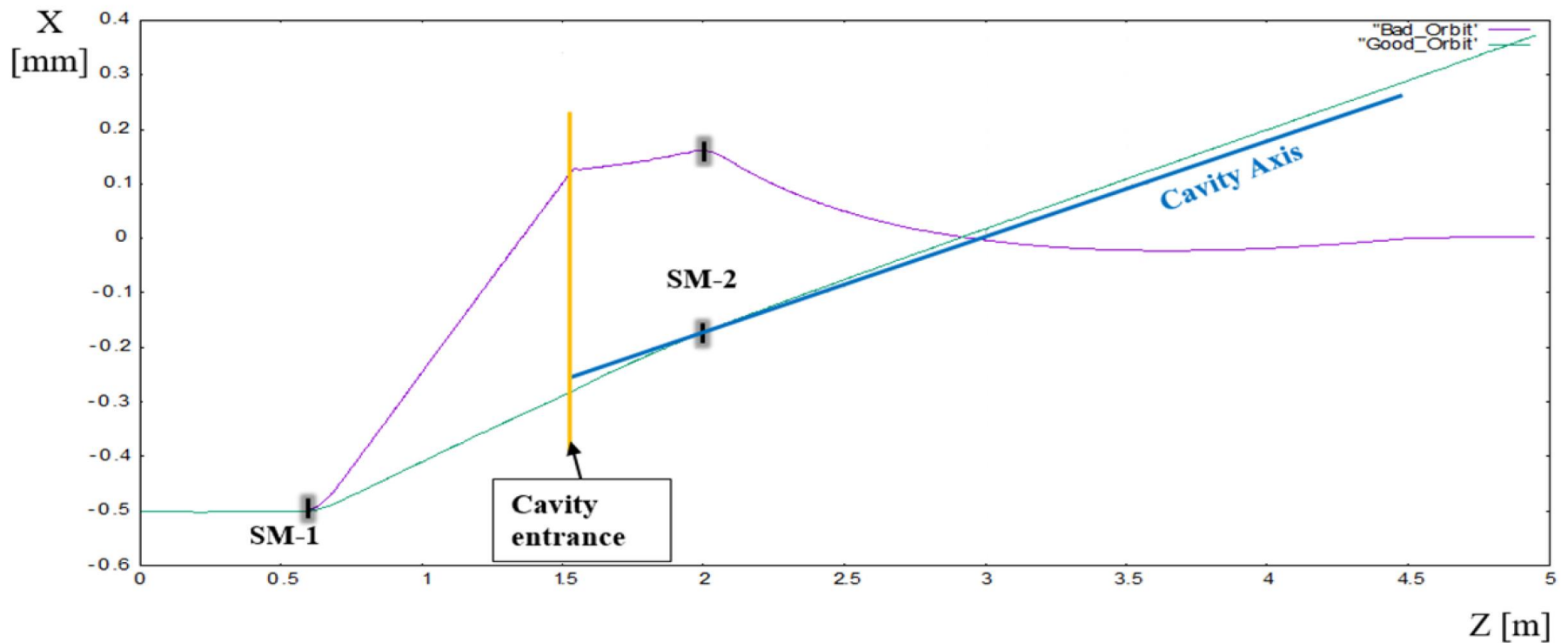
A. Bacci, et al. “**GIOTTO: A Genetic Code for Demanding Beam-dynamics Optimizations**”, doi: 10.18429/JACoW-IPAC2016-WEPOY03

M. Rossetti Conti, A. Bacci “**Beam based alignment methods for cavities and solenoids in photo-injectors**”, Proceedings of IPAC2016, Busan, Korea THPMB011

A. Bacci, S. Gallo, et al., “**STUDY OF A C-BAND HARMONIC RF SYSTEM TO OPTIMIZE THE RF BUNCH COMPRESSION PROCESS OF THE SPARC BEAM**” IPAC 2015  
doi:10.18429/JACoW-IPAC2015-TUPW

Thanks for your attention

M. Rossetti Conti, A. Bacci “Beam based alignment methods for cavities and solenoids in photo-injectors”, Proceedings of IPAC2016, Busan, Korea THPMB011



# A Radio Frequency (RF) Knob

Catching beams @ different envelope sizes MEANS:

a strong control on Focal Lengths of the RF lens, MEANS:

a KNOB to focus Adiabatically and in Full (x,y) Symmetry

NO Quadrupoles

→ Symmetric Beams

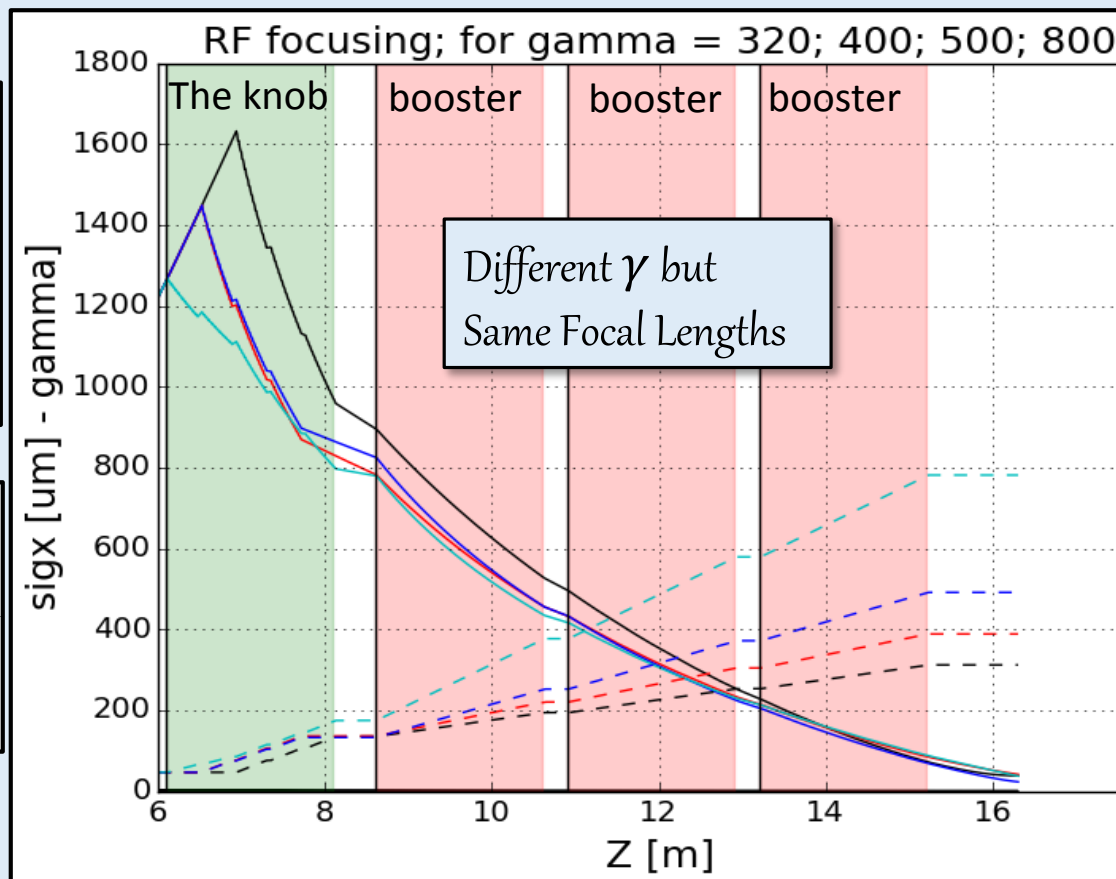
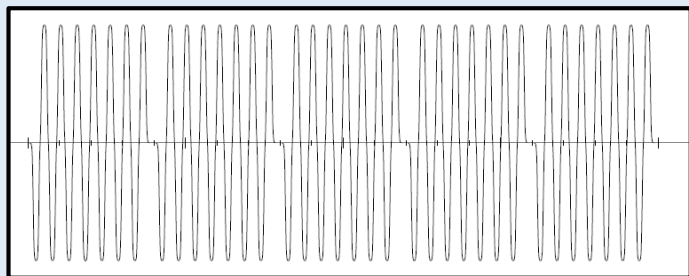
→ Very Good for BD

## The knob

C-band five sectors Standing Wave (SW) cavity. Why C-band and SW:

- Irises large enough
- One feeder for sector
- A bit stronger focusing effect

## The knob - field

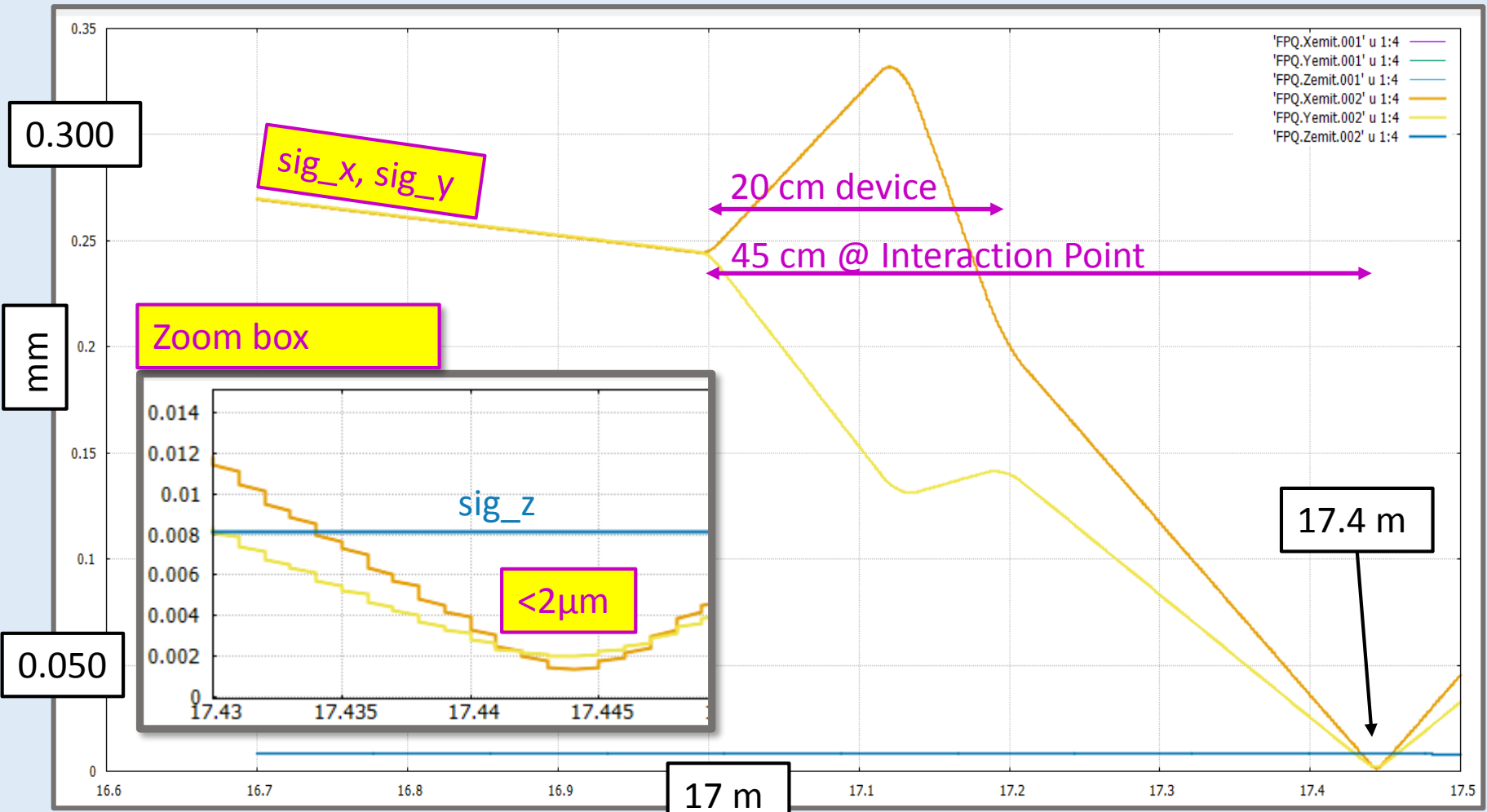


# A Radio Frequency (RF) Knob – Final focus

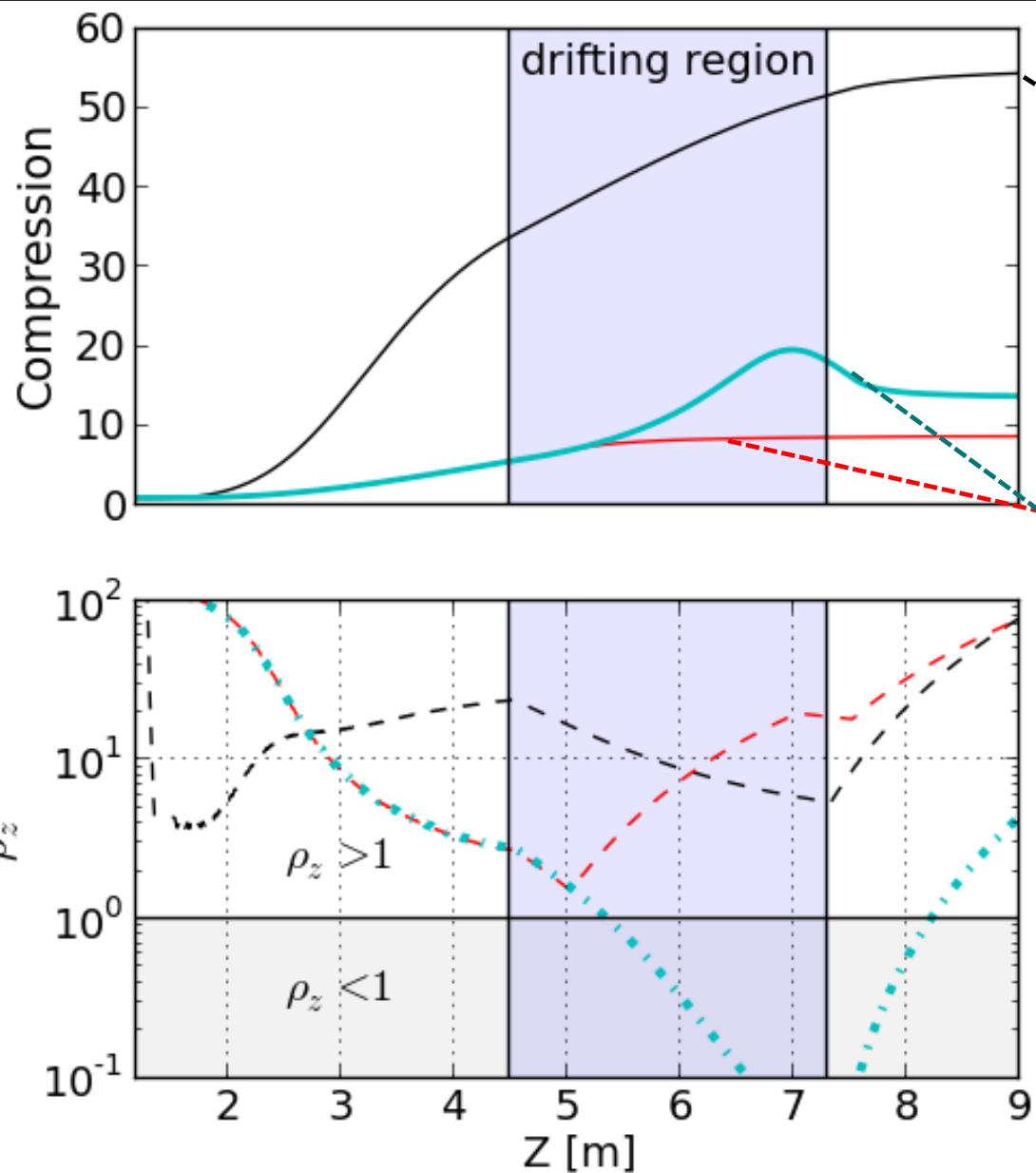
An IMPORTANT question:

Can we focus the bunch at Linac exit after RF focusing down to final focus specs?

Final focusing channel by 5 Permanet Quads (20 cm)



## $\rho_z$ Laminar parameter for LB & VB

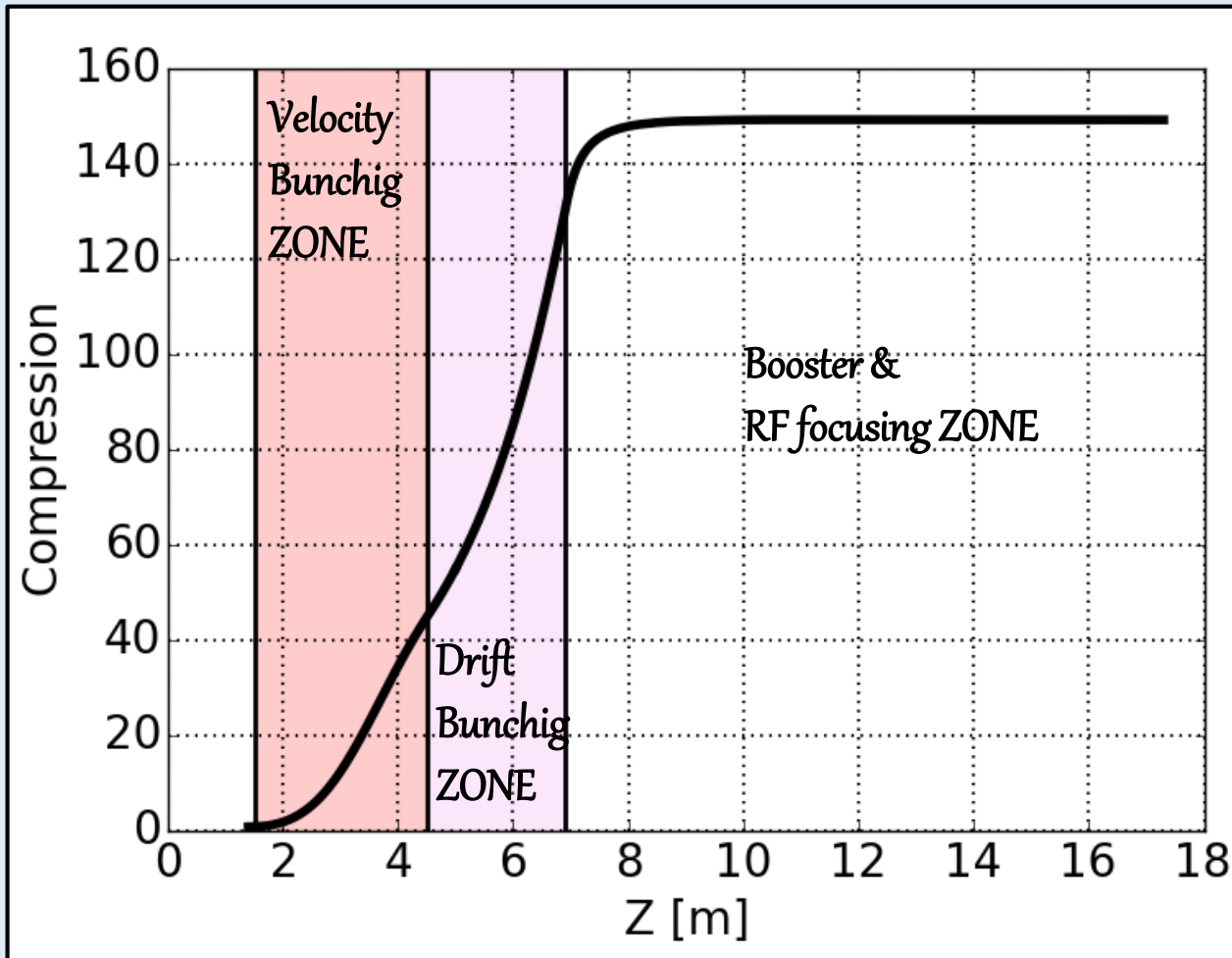


Laminar Bunching  
 $\rho_z$  is low but  $> 1$   
A soft bunch for  
the compression

Velocity Bunching  
NO DRIFT (STANDARD  
case)  
 $\rho_z$  starts  $\gg 1$   
A bunch stiff to be  
compressed

Velocity Bunching  
SI DRIFT (TEST  
case)  
Laminarity is lost  
 $\rho_z < 1$

## compression factor 40 pC case



@ booster end  
= 17 m (0.5 GeV)

$$Q_b = 40 \text{ pC}$$

$$\sigma_z = 4.2 \text{ } \mu\text{m}$$

$$I_{peak} > 2.5 \text{ kA}$$

$$\epsilon_{peak} = 0.3 \text{ } \mu\text{m}$$

$$B_{peak} = 3 \cdot 10^{16}$$

$$(I_p = 1.5 \text{ kA})$$

$$(\Delta\gamma/\gamma)_{@I_{peak}} \cong 8 \cdot 10^{-5}$$

