

Impact of crab cavity bump nonclosure on DA during collisions

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<u>Overview</u>

- Simulation parameters
- Sanity checks
- Impact of CC bump non-closure on orbit, tune diffusion & DA

185th Hi-Lumi WP2 meeting 24/11/2020

Simulation parameters

DA studies with beam-beam at the end of β^* -leveling

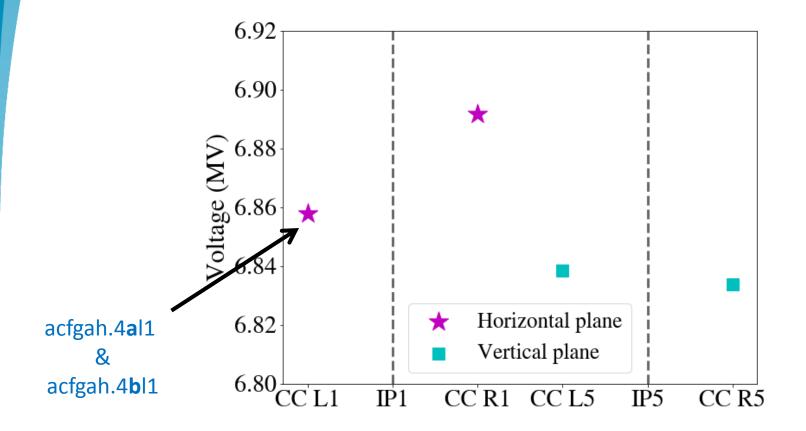
HL-LHC V1.5*						
7 TeV	β [*] =15 cm	1.2e11 ppb	ε _n =2.5 μm	C⁻=1e-3	I _{oct} =+300 A	Q'=15
N _{IP1} , N _{IP2} , N _{IP8} = 2748, 2494,2572		φ/2 _{ιΡ1} = 250 μrad	φ/2 _{IP8} = -200 μrad Positive polarity	on_crab= -190 µrad	(Q _x , Q _y) =(0.31, 0.315)	V _{RF} =16 MV

<u>*https://github.com/lhcopt/lhcmask/tree/master/python_examples/hl_lhc_collisions</u> <u>_python</u>



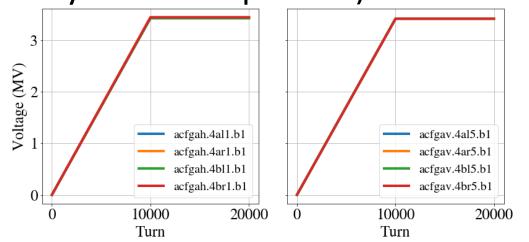
https://gitlab.cern.ch/skostogl/crab-cavities-da-studies

For on_crab1=on_crab5 =-190 µrad:





Simulation parameters ➤ CC voltages are ramped adiabatically during 10 K turns (~18 synchrotron periods).



Output of DYNK block SixTrack



Scc voltages are ramped adiabatically during 10 K turns (~18 synchrotron periods). Output of DYNK block SixTrack

≻δp/p=27e-5 (3/4 bucket height) in DA simulations→ z~200 mm. Studies also with z=75 mm.

20000

0

acfgav.4bl5.b1

acfgav.4br5.b1

20000

10000

Turn

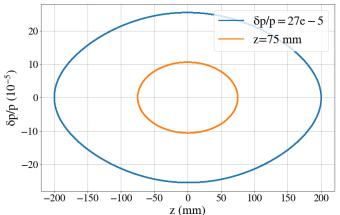
acfgah.4bl1.b1

acfgah.4br1.b1

10000

Turn

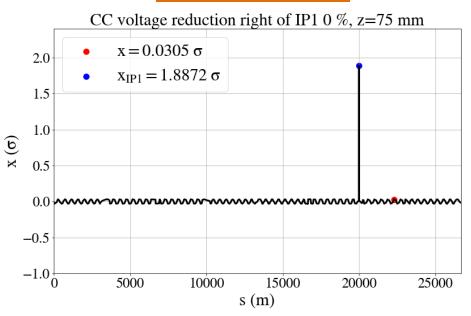
0





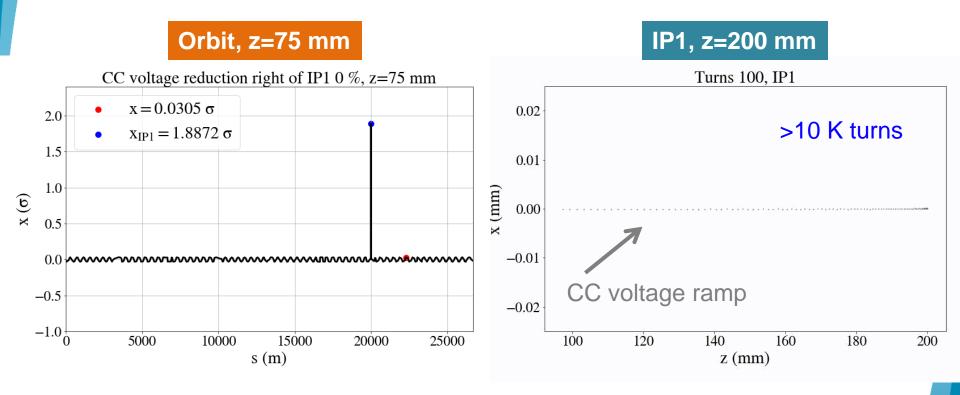
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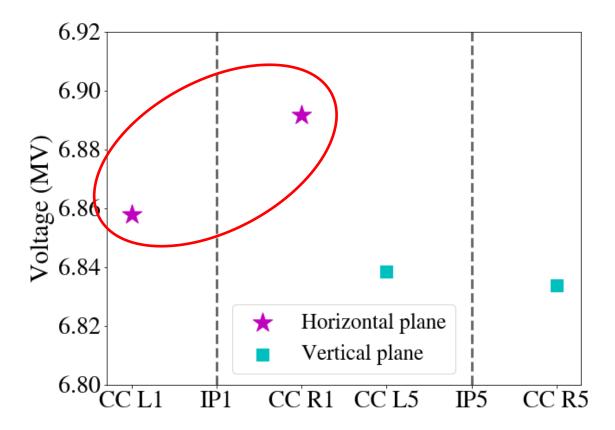








For on_crab1=on_crab5 =-**190 µrad**:

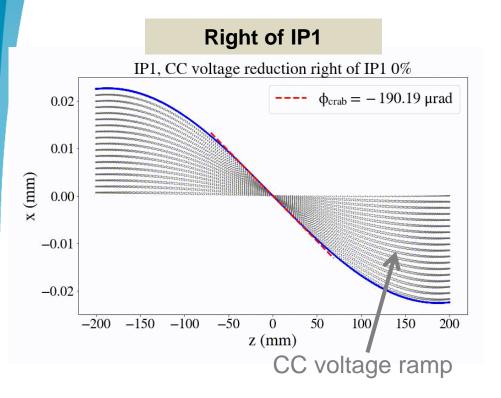


Modify the CC voltage right (or/and) left of IP1 with the formula: $V_{err} = V_{closed} \times (1-error)$



CC bump non-closure: Impact on crabbing angle



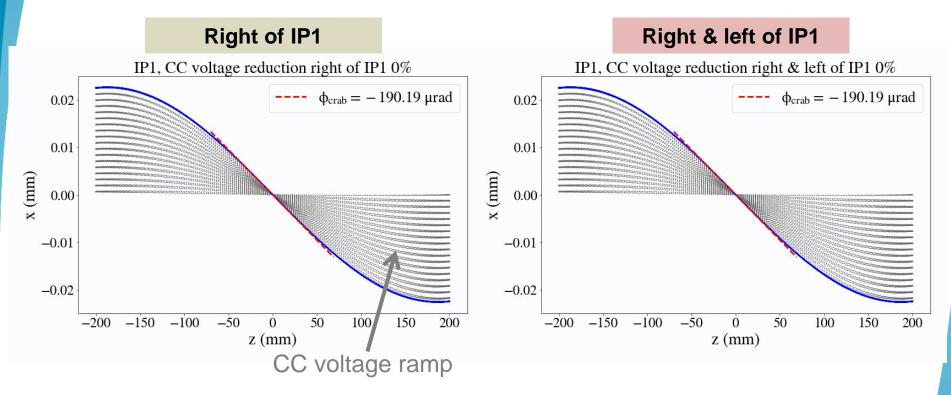


Minimum crabbing angle for **200%** reduction of voltage right of IP1



CC bump non-closure: Impact on crabbing angle

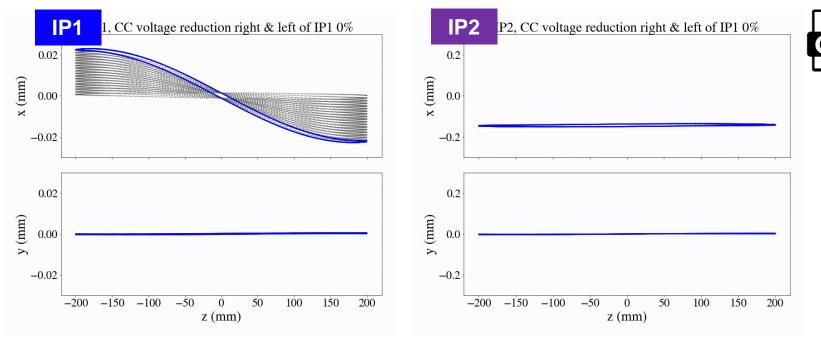




Minimum crabbing angle for **200%** reduction of voltage right of IP1 Minimum crabbing angle at 100% reduction of voltage right & left of IP1

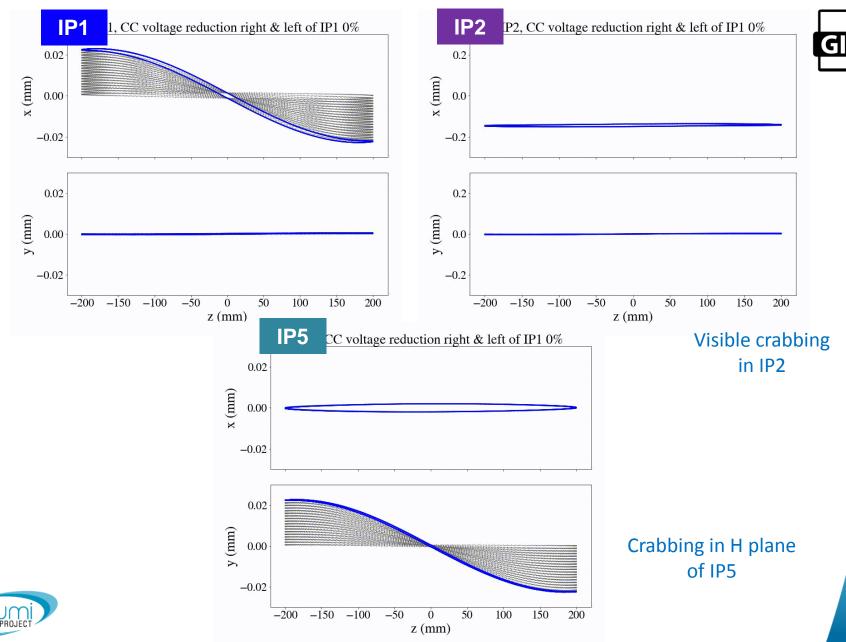


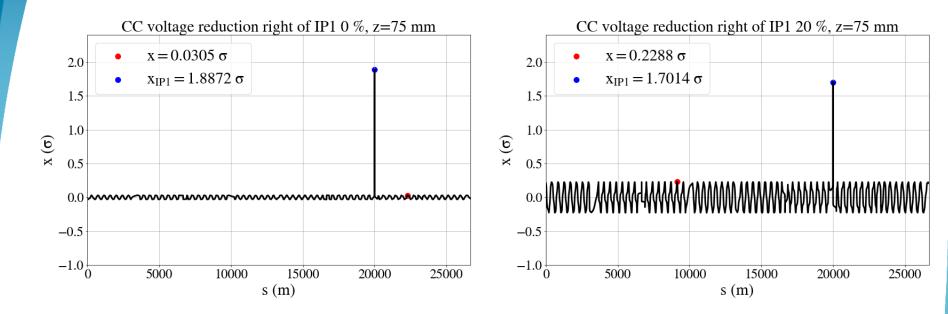
CC bump non-closure: Impact on other IPs



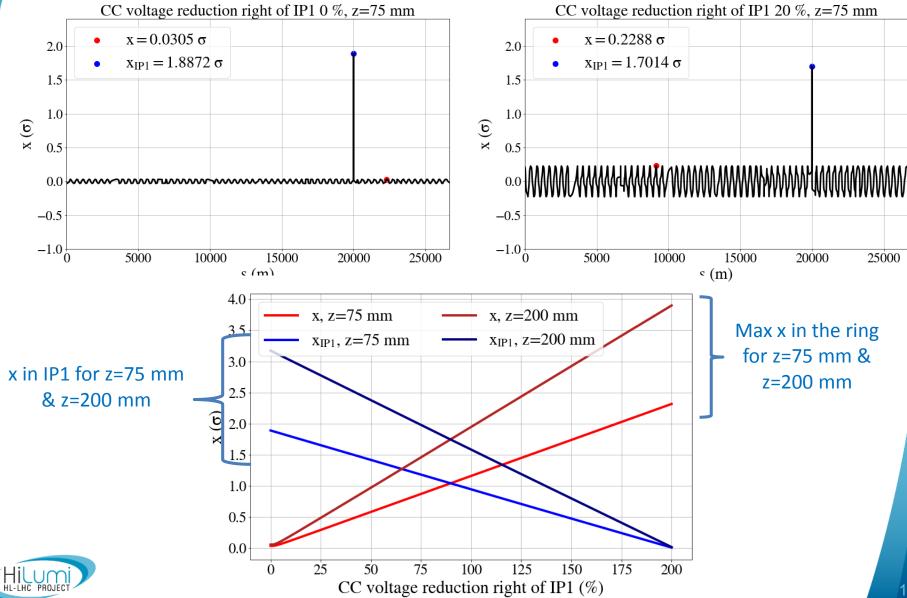


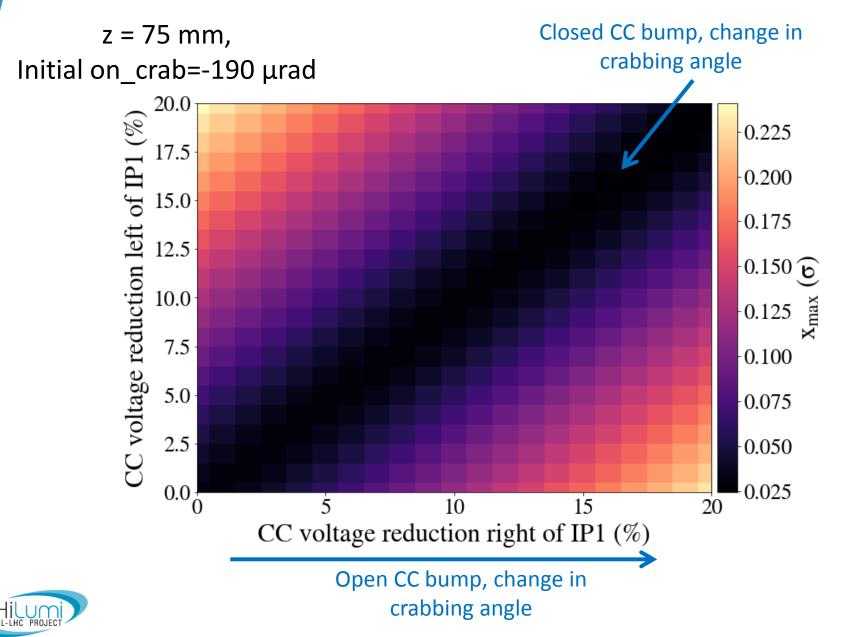
CC bump non-closure: Impact on other IPs

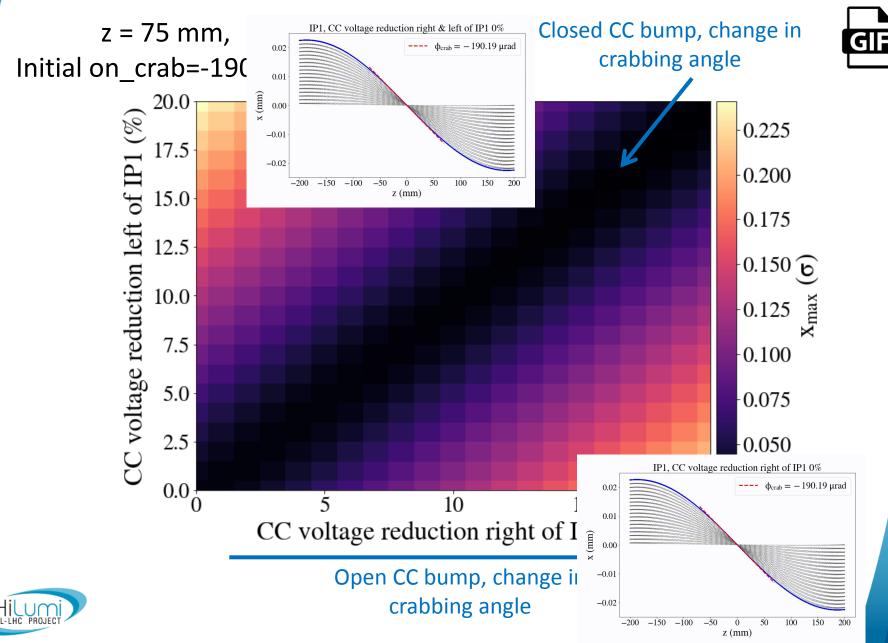






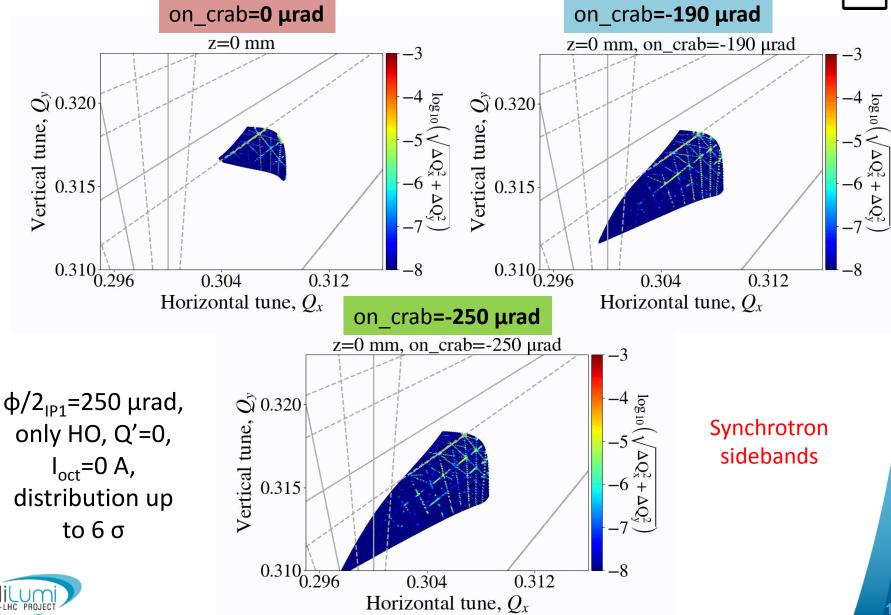






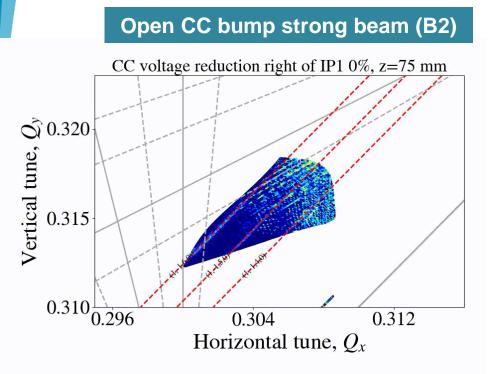
6D FMAs: Scan in z





6D FMAs: CC bump non-closure

z = 75 mm, Initial on_crab=-190 μrad

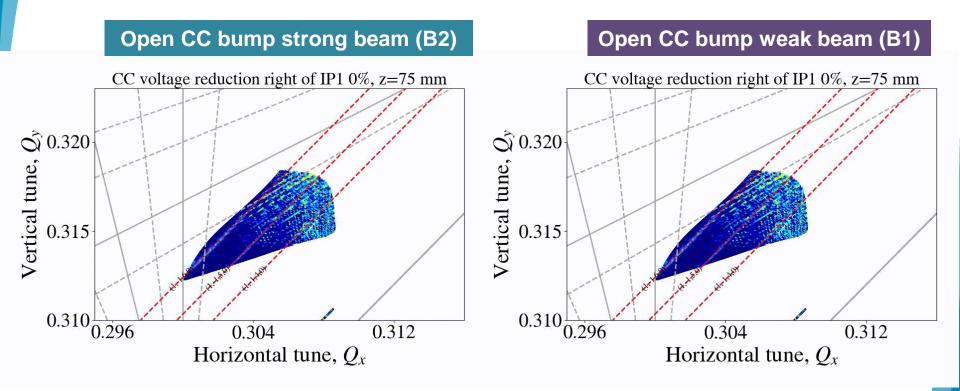




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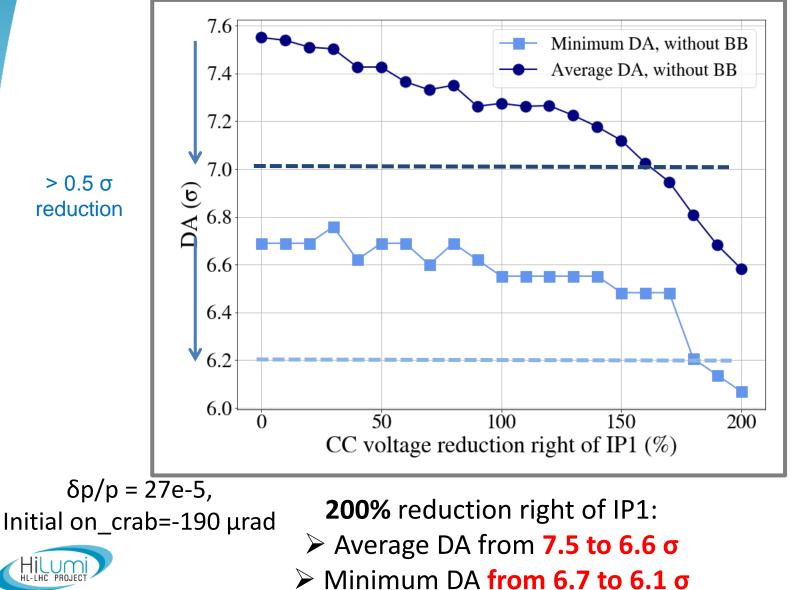




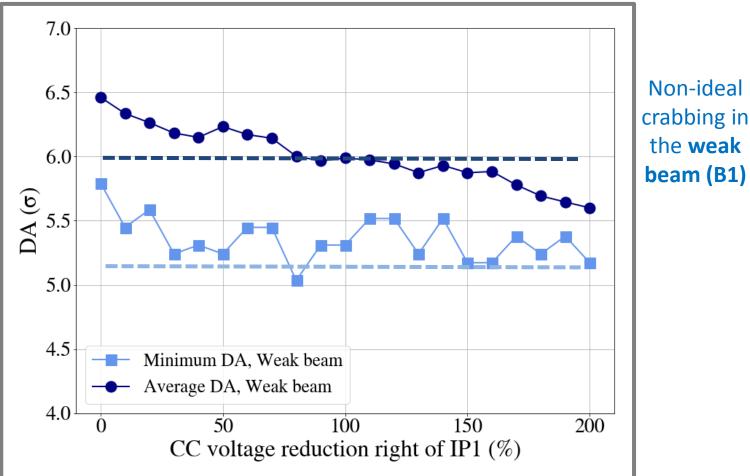
When including imbalance in both beams, main contributor is the weak beam (see also appendix).



Impact of CC bump non-closure on DA without BB



Impact of CC bump non-closure on DA with BB

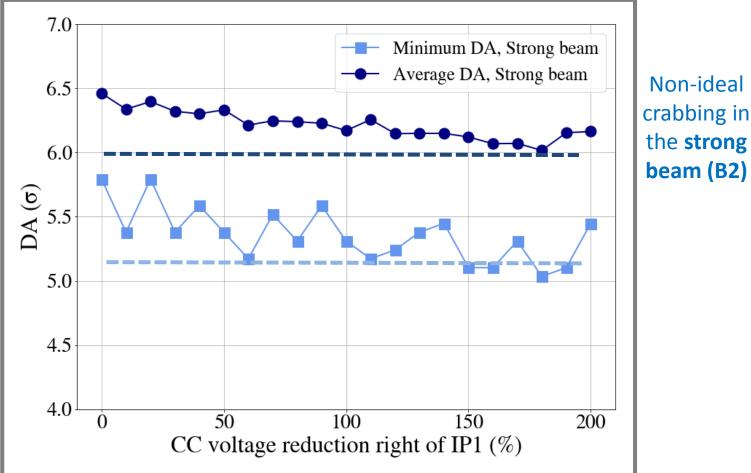


200% reduction right of IP1:

Average DA from 6.5 to 5.6 σ



Impact of CC bump non-closure on DA with BB

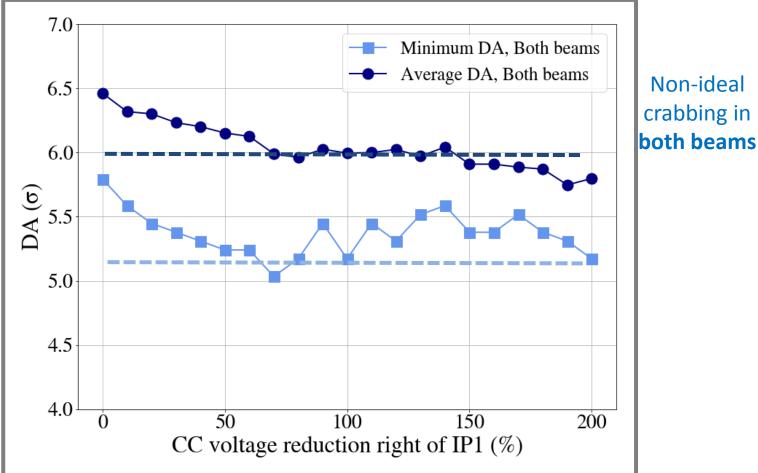


200% reduction right of IP1:

Average DA from 6.5 to 6.2 σ



Impact of CC bump non-closure on DA with BB



200% reduction right of IP1:

Average DA from 6.5 to 5.8 σ



- Investigated impact of CC bump non-closure on tune diffusion & DA at the end of the β*-leveling for HL-LHC v1.5.
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 - Without beam-beam (& up to 200% CC voltage reduction right of IP1):
 - From 7.5 to 6.6 σ average DA and from 6.7 to 6.1 σ minimum DA.



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Without beam-beam (& up to 200% CC voltage reduction right of IP1):

- From 7.5 to 6.6 σ average DA and from 6.7 to 6.1 σ minimum DA. With beam-beam (& up to 200% CC voltage reduction right of IP1):
- Considered non-perfect crabbing for weak and/or strong beam.
- Weak beam (B1): from 6.5 to 5.6 σ average DA (6D FMAs: HO tune spread ↓ as crabbing angle ↓ & ↑ of tune diffusion due to synchrobetatron sidebands)
- Strong beam (B2): from 6.5 to 6.2 σ average DA (6D FMAs: HO tune spread \downarrow as crabbing angle \downarrow , no impact on tune diffusion)
- Both beams: from 6.5 to 5.8σ average DA.



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- Both beams: from 6.5 to 5.8 σ average DA.

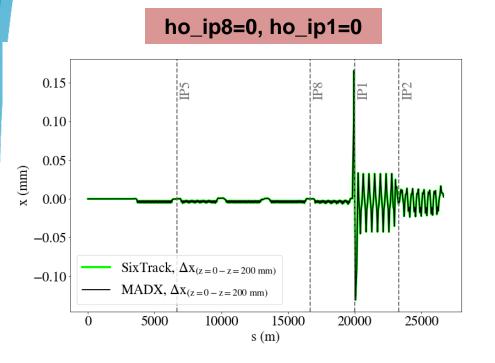
Next steps: Include CC multipolar components.



Backup



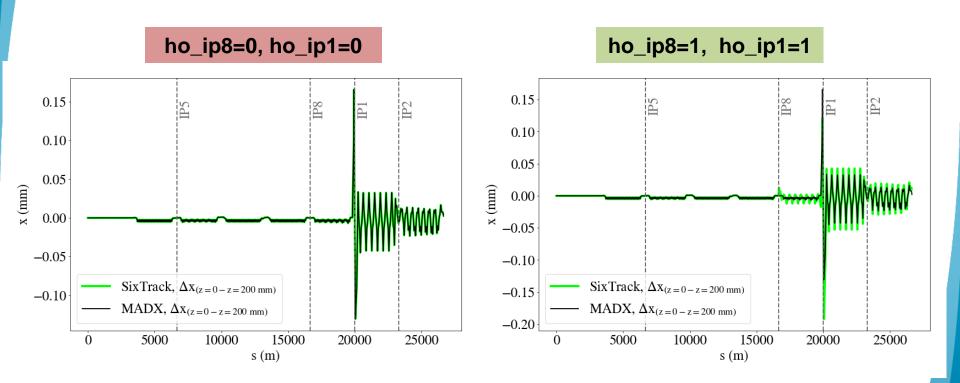
Sanity check MADX-SixTrack:



Initial conditions $(x, p_x, y, p_y)=(0,0,0,0), z=0 \text{ and } z=200 \text{ mm.}$



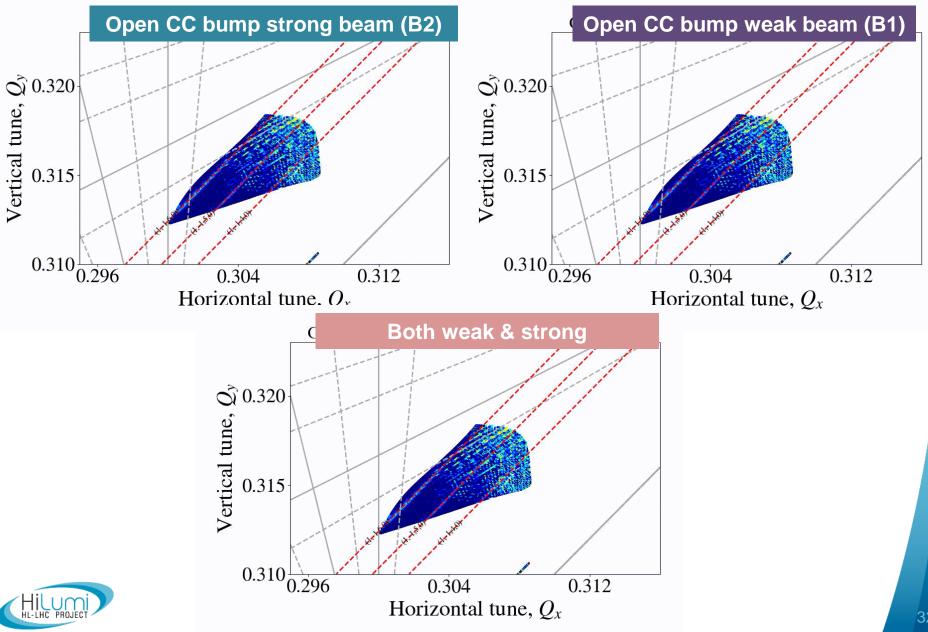
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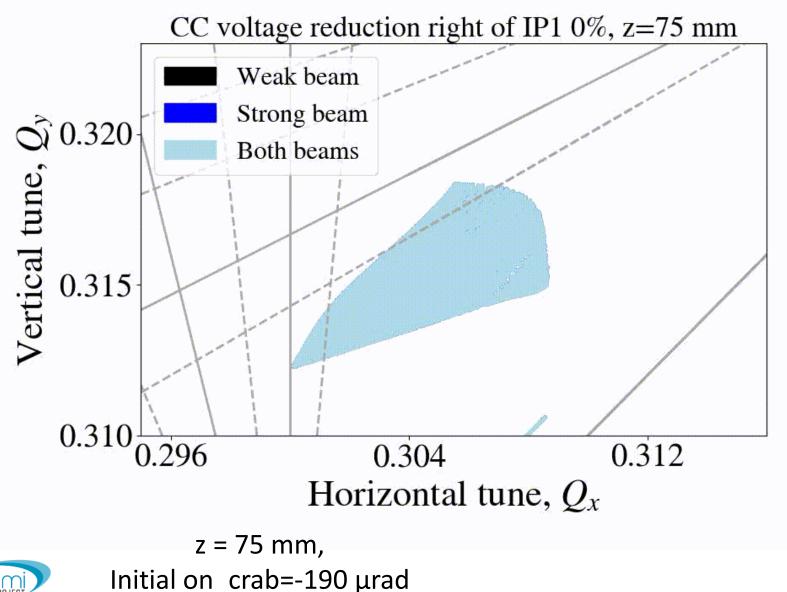
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6D FMAs: CC bump non-closure



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