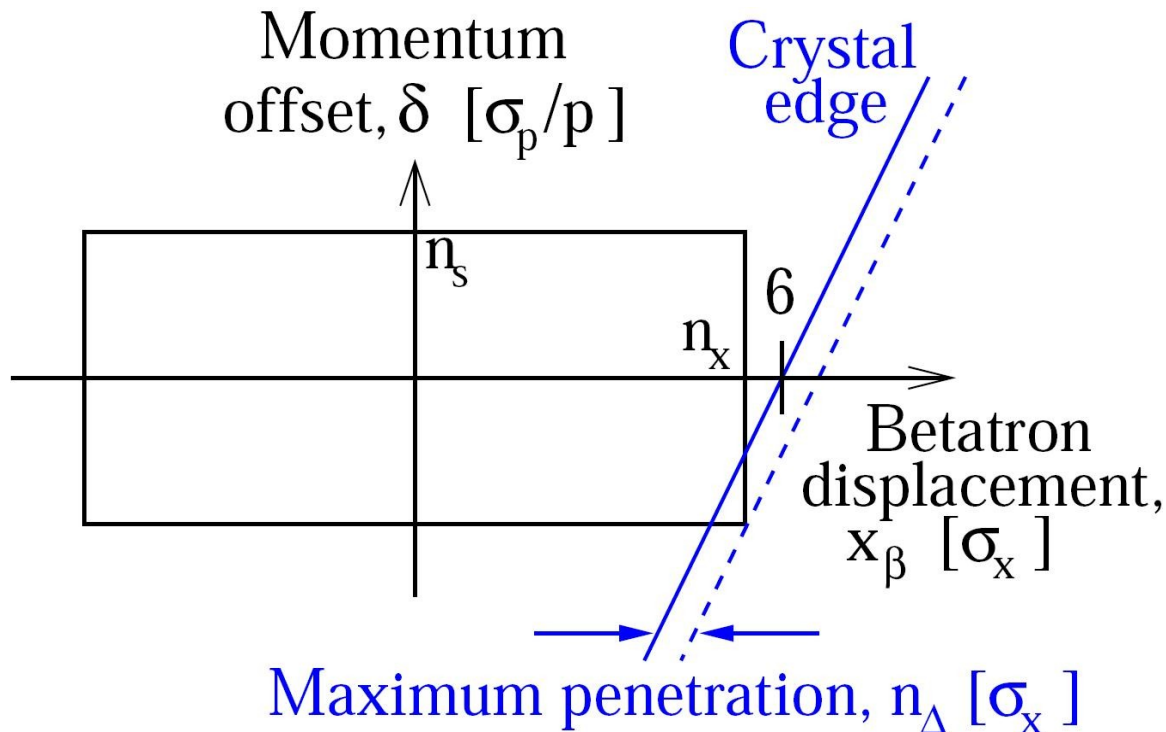


The grazing function g

S. Peggs (BNL) & V. Previtalli (CERN)

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A “grazing particle” just touches the crystal during simultaneous betatron & synchrotron oscillation extrema.

$$a_x + |\eta| a_s = |x_c|$$

Thus the grazing angle depends linearly on the synchrotron amplitude a_s according to

$$x'_G = -\frac{\alpha}{\beta}x_c + \text{sgn}(x_c) \text{sgn}(\eta) g a_s \quad (15)$$

where the linear slope of grazing angle with respect to synchrotron amplitude is

$$\frac{dx'_G}{da_s} = \text{sgn}(x_c) \text{sgn}(\eta) g \quad (16)$$

The *grazing function* g that enters these equations is an optical quantity defined as

$$g \equiv \left(\frac{\alpha}{\beta} \eta + \eta' \right) \quad (17)$$

The grazing function is thus revealed to be just

$$g = \sqrt{\beta} \eta'_N$$

and the rigorous general synchrotron condition $g = 0$ is just

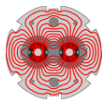
$$\eta'_N = 0$$

Efficient collimation: the grazing angle must be within the crystal acceptance angle for “all” synchrotron amplitudes.

$$|g| < \frac{\sigma'_A}{n_{max} (\sigma_p/p)}$$

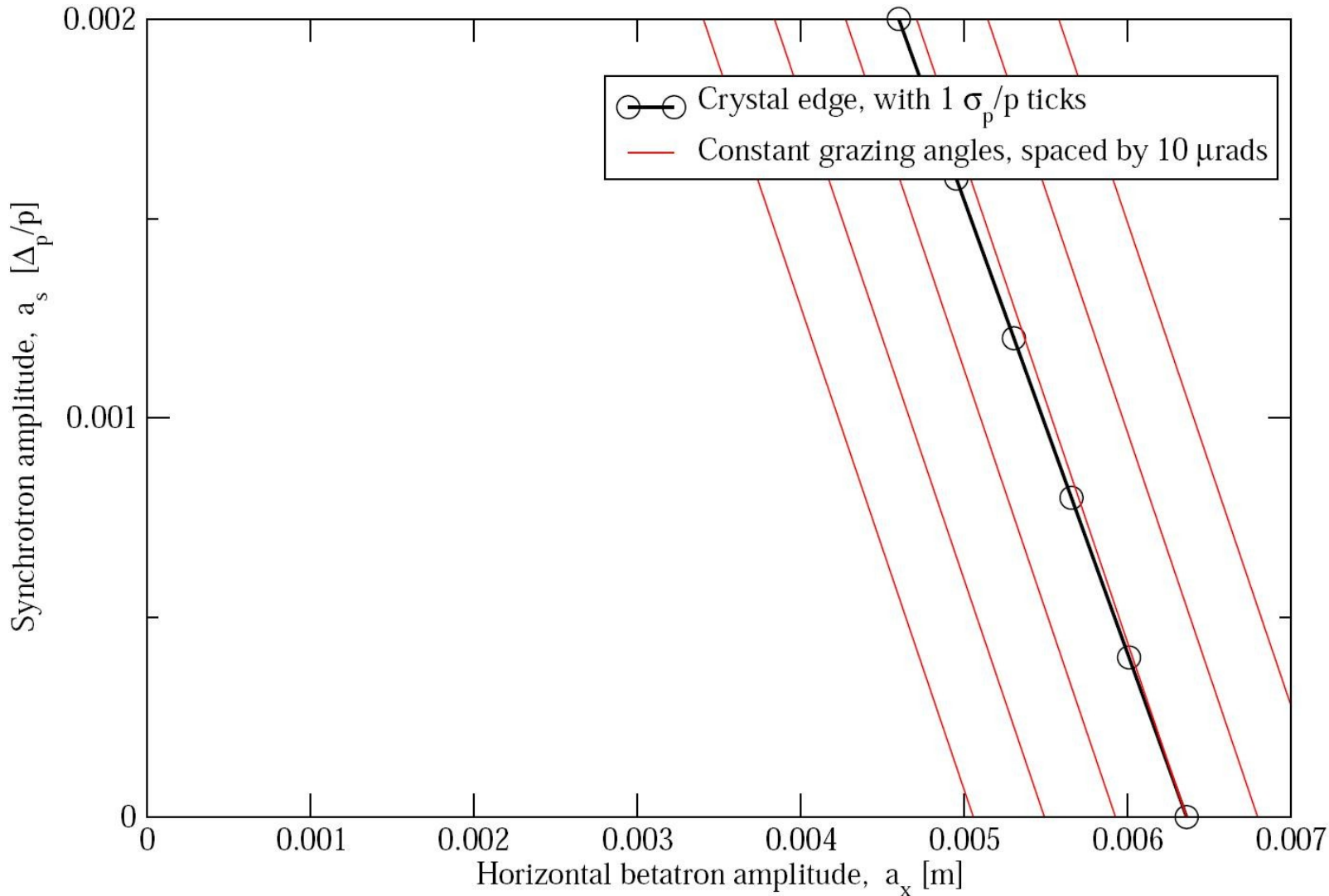
σ'_A [μrad] \sim 2 channeling at 7 TeV
 \sim 10 channeling at 0.1 TeV
 \sim 100 volume reflection at any energy

	α	β [m]	η [m]	η' [10^{-3}]	g [10^{-3}]	E [TeV]	σ_p/p [10^{-3}]	σ'_G [μ rad]
RHIC	-26.5	1155.0	-0.864	-16.2	3.6	0.10	0.50	1.81
SPS	-2.21	96.1	-0.880	-19.0	1.2	0.12	0.40	0.48
Tevatron	-0.425	67.5	1.925	15.0	2.9	0.98	0.14	0.41
LHC	1.94	137.6	0.559	-8.9	-1.0	0.45	0.31	0.31
						7.0	0.11	0.11



LARP

SPS UA9 expectations

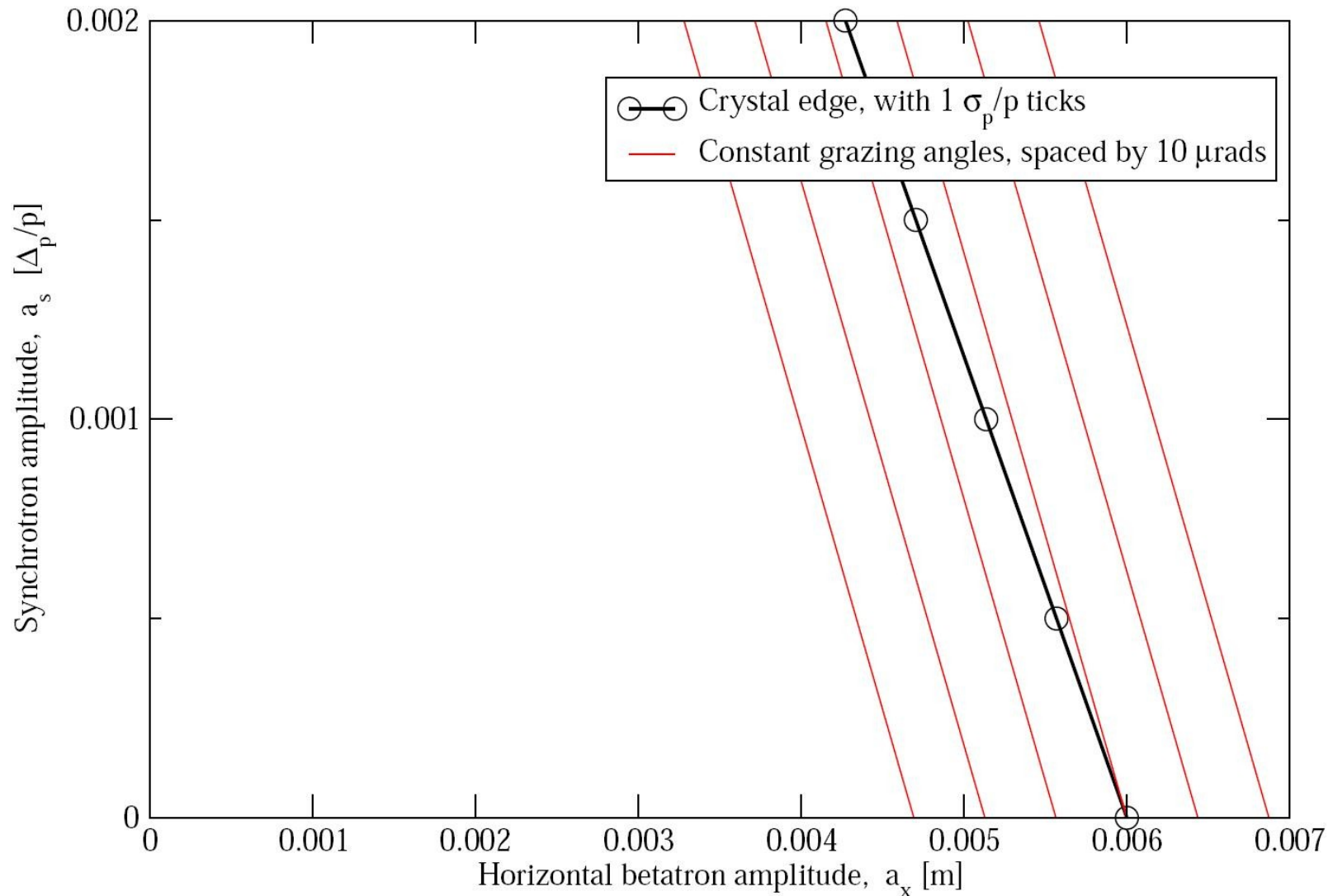


Black line:
Crystal edge

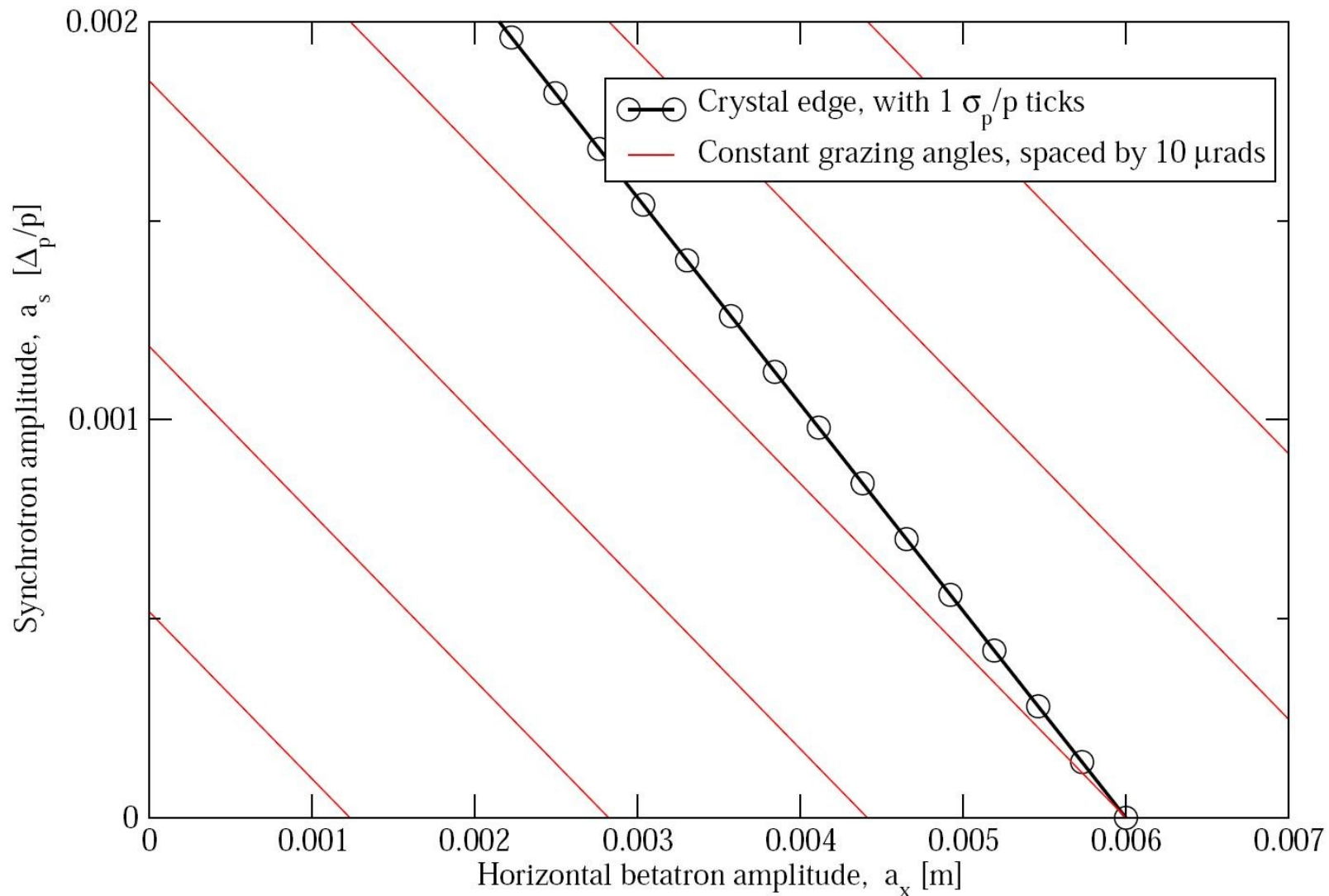
Circles:
Synchrotron amplitudes spaced by $1 \sigma_p/p$

Red lines:
Constant grazing angles spaced by $10 \mu\text{rad}$

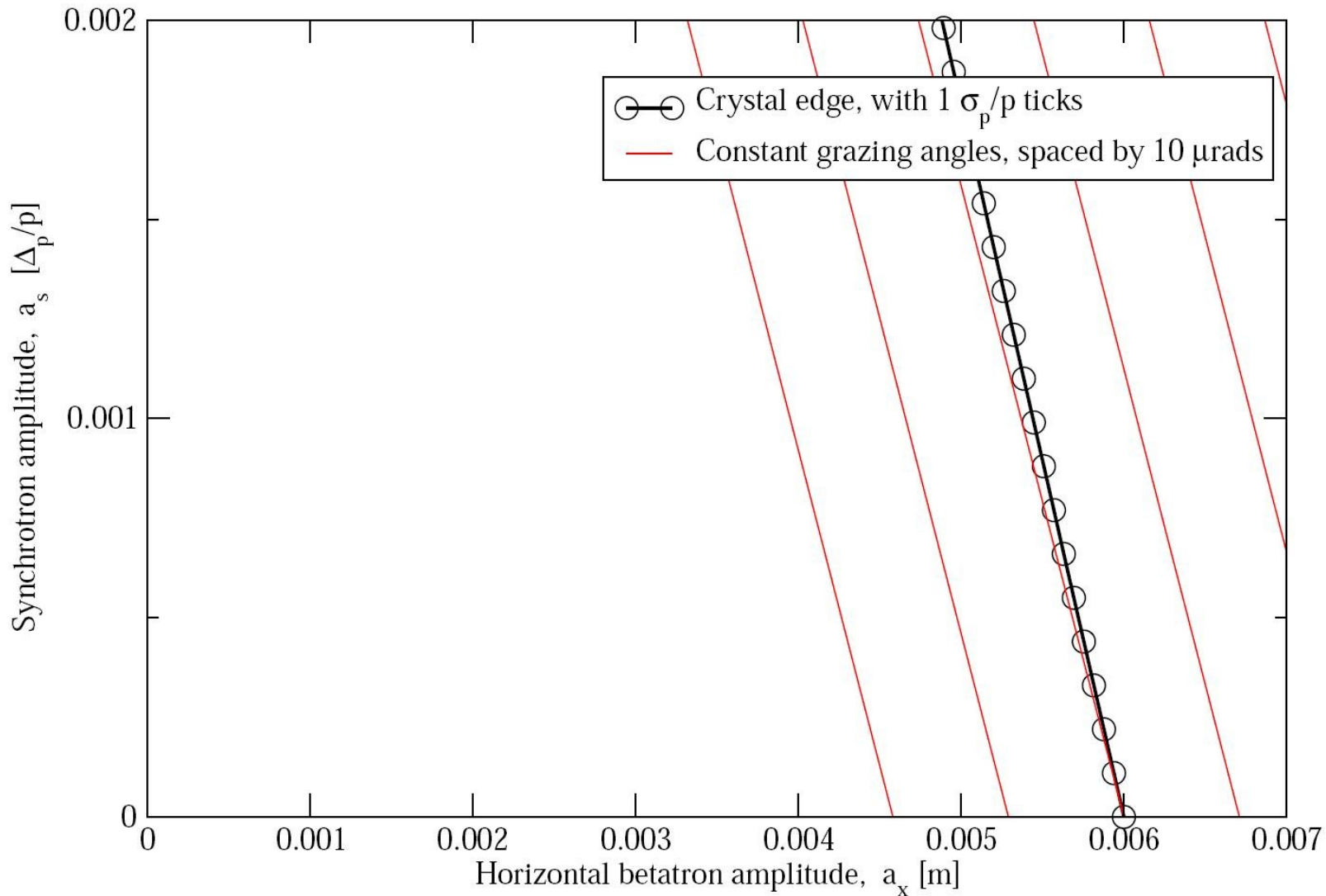
UA9 looks ok: the grazing angle changes by only $0.48 \mu\text{rad}$ per σ_p/p , despite large negative dispersion at the crystal.



Large σ_p/p : grazing angle changes by $5.4 \mu\text{rad}$ over $3 \sigma_p/p$.
 Helps explain disappointing performance?



Flattened slopes: grazing angle changes by $0.41 \mu\text{rad}$ per $3 \sigma_p/p$. Measured optics values also available.



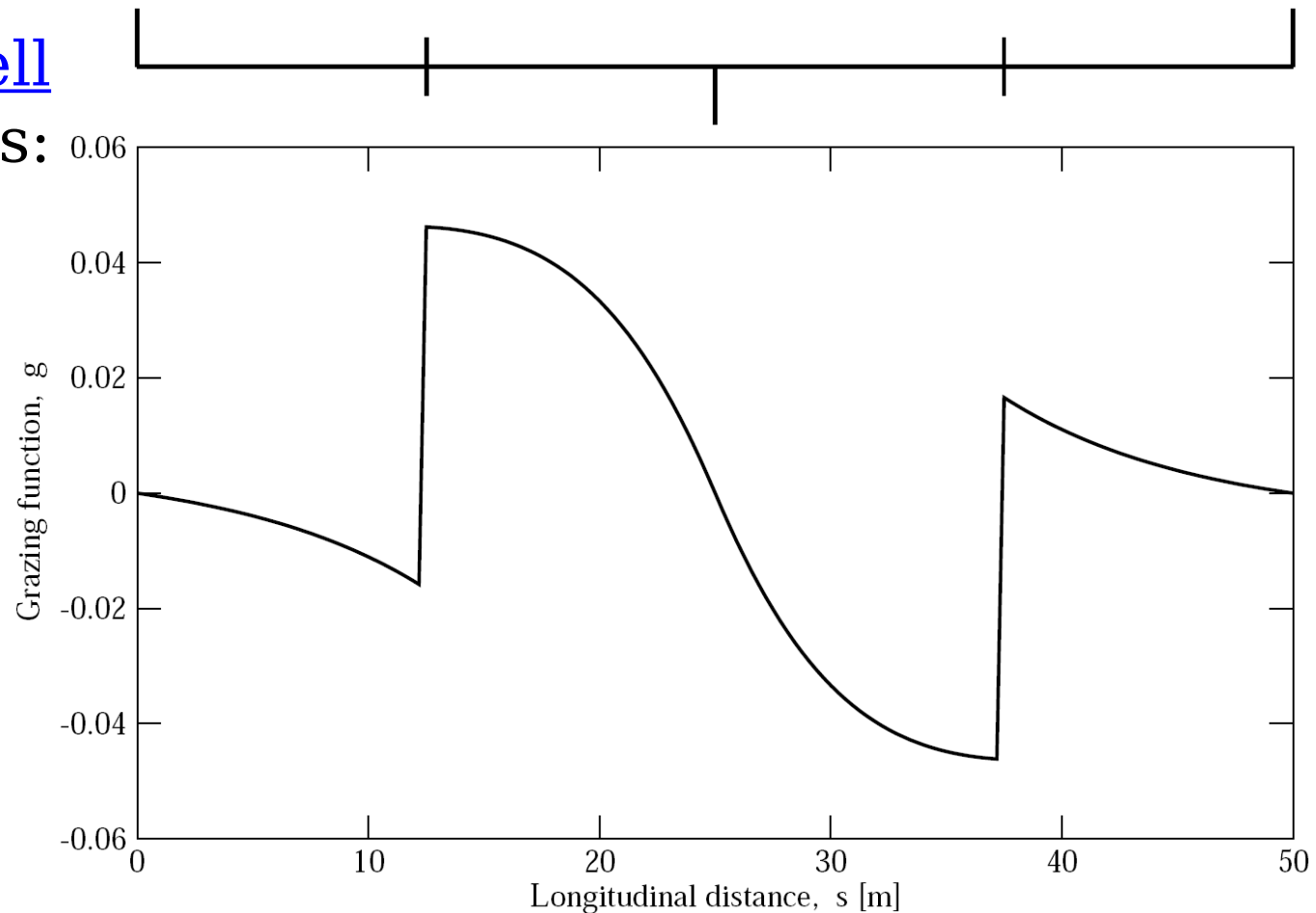
Small σ_p/p , small channeling acceptance: grazing angle changes by only 0.11 (0.31) μ rad per σ_p/p at 7 (0.45) TeV.

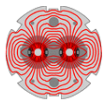
How does g propagate?

A thin dipole: $\Delta g = \left(\Delta\eta' - \frac{\eta \Delta b'}{b} \right) = \Delta\theta$

A thin quad: $\Delta g = \eta \left(\frac{\Delta\eta'}{\eta} - \frac{\Delta b'}{b} \right) = 0$

A matched FODO cell
thin dipoles & quads:





Propagation - 2

LARP



A matched FODO cell

short dipoles
thick quads:

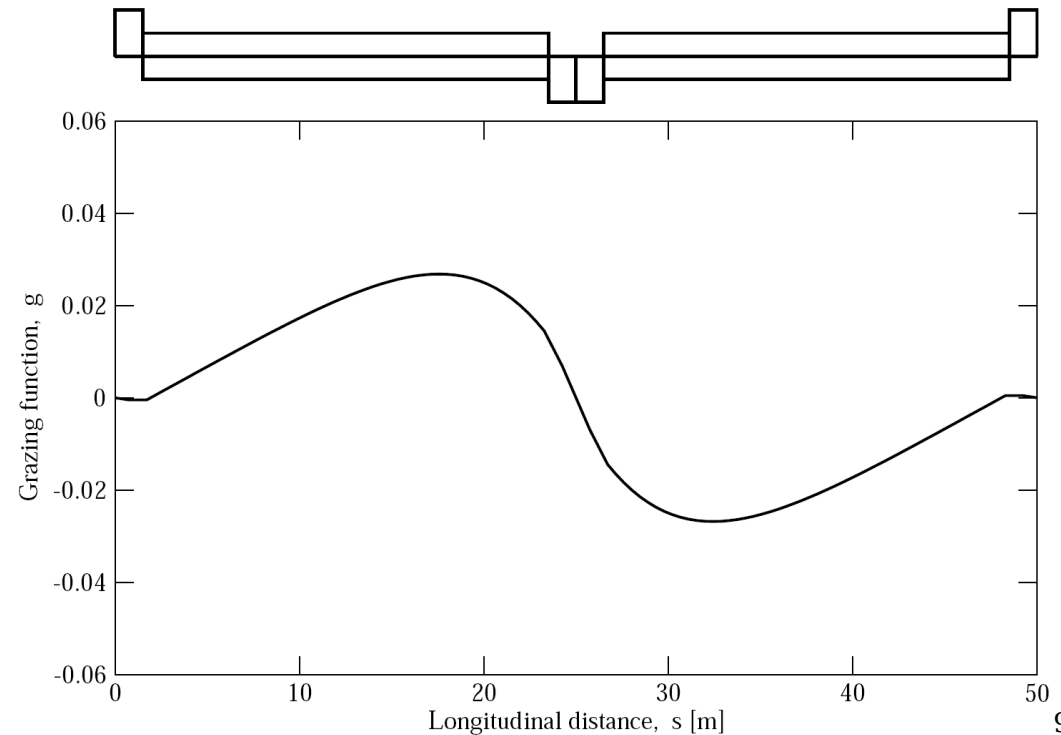
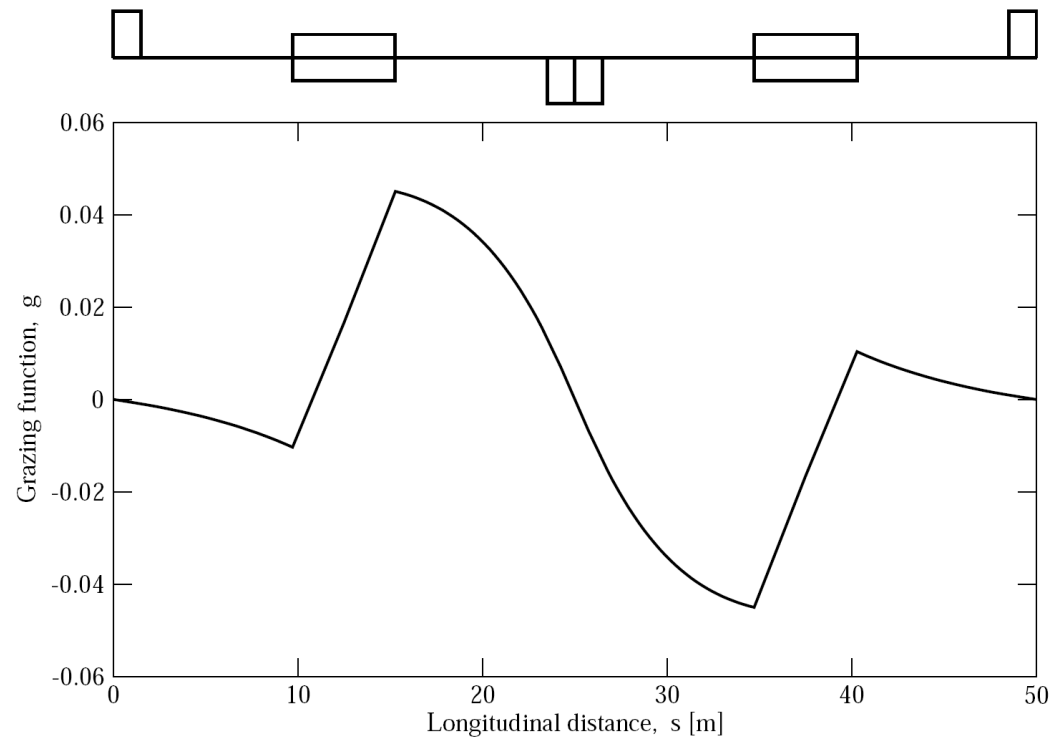
A matched FODO cell

long dipoles
thick quads:

g is small when the normalized dispersion is almost constant

$$g = \sqrt{\beta} \eta'_N$$

LCU section mtg, CERN, Dec 2 2008



Simple prediction vs numerical testing

Predict $|g|_{max} \approx \theta \frac{3}{4\sqrt{2}} \frac{\sqrt{1+C^2}}{S^2C} \left((2-S)\sqrt{1+S} - (2+S)\sqrt{1-S} \right)$

where $S \equiv \sin \phi/2$, $C \equiv \cos \phi/2$

for a matched FODO cell, phase advance per cell ϕ , half cell length L , half cell bend angle θ

In reasonable agreement with numerical testing:

$$g_{max} \approx 0.427 L^0 \theta^1 \quad (51)$$

when the phase advance per full-cell is 90 degrees. There is no dependence on the cell length! A fair rule of thumb is that

$$g_{max} \approx \theta/2 \quad (52)$$

CAUTION: these best case results assume **matched optics**.

Conclusions

- 1) The grazing function g parameterizes the rate of change of total angle with synchrotron amplitude.
- 2) A pure optics function, it is related to the slope of the normalized dispersion. Ideal crystal value $g = 0$
- 3) g should be small enough that “all” synchrotron amplitudes are within the crystal acceptance angle.
- 4) This appears to be reasonable to achieve in practice, especially in VR mode, and at lower energies.
- 5) Place crystal away from betatron & dispersion waves, since they may increase g by an order of magnitude.
- 6) Planning for future crystals should include a grazing function analysis, both in design and in error analysis.