



Main Linac lattice design for TDR (KCS & DKS configurations)

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- The lattices for two layouts
 - KCS - Klystron Cluster Scheme (“4-RFU” and “3-RFU”)
 - DKS - Distributed Klystron Scheme (“3-RFU”)
- Treaty points:
 - T(P/E)RTML2ML &
 - TPML2BDS/TEML2PS
- Details of modified matching procedures
(including optical functions, dispersion minimization and the linac reference orbit following the Earth’s curvature)
- Summary & the present lattice status

ML lattices designed with MAD8 (a special version 51.15.s by M.Woodley) following to the approach.*

** A.Valishev, N.Solyak, M.Woodley, “Status of the ILC Main Linac Lattice Design”, PAC’07, **2007**.*



Changes in TDR layout vs. RDR

- Two configurations:
 - **KCS: new configuration of CryoUnits (CU) vs. DRFS**
 - CU contains long ($4 * \text{RFunit}$) and short ($3 * \text{RFunit}$) cryo-strings
 - Length of CU's are different: from 25 to 52 RF units.
 - Number of RF units are different for Electron (285) and Positron (282) Linacs
 - **DKS:**
 - All cryo-strings are short (3 RF units)
- Treaty points (from RTML and to Undulator/BDS) are modified
- Collimation system migrated from BDS to ML
 - Polarity of the last quad in ELIN and PLIN are different.



KCS version (ver. 6/26/2012, C.Nantista)

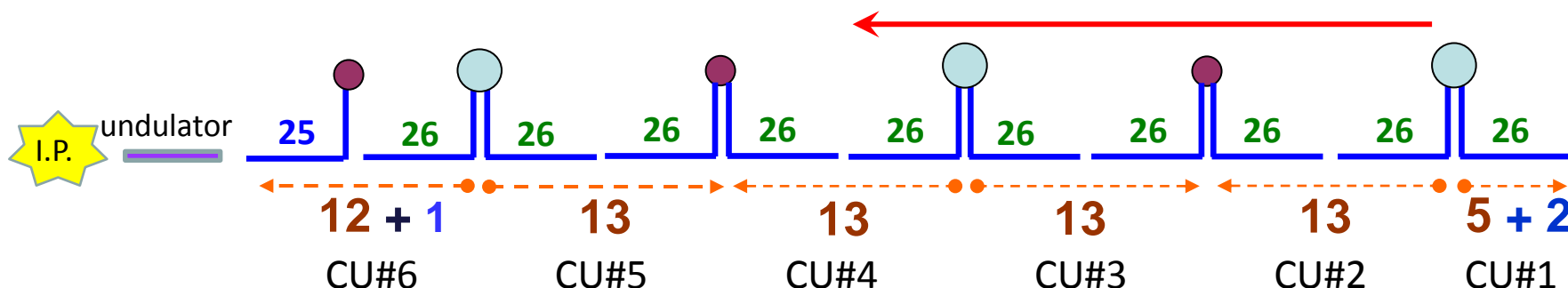
It allows to use most of existing RDR solutions and requires small number of re-matchings

- -- main facilities shaft
- -- additional KCS shaft
- -- cryogenic systems

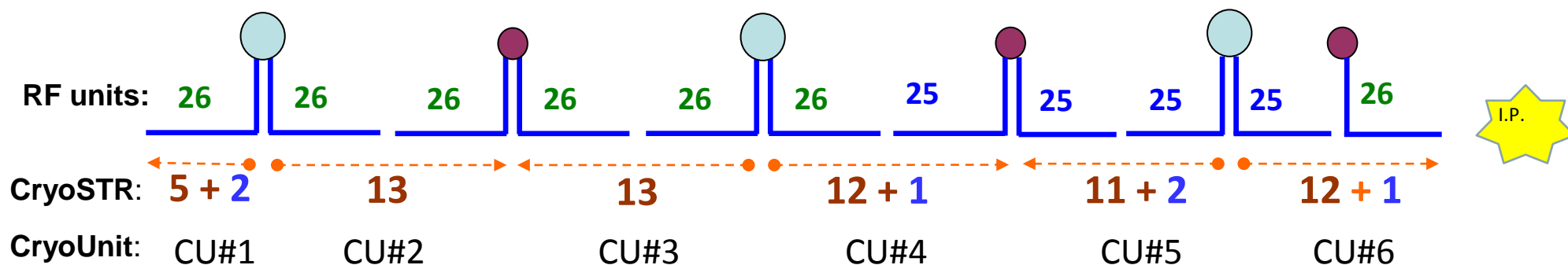
-- 4-rf unit CSTR

-- 3-rf unit CSTR

e^- - beam (ELIN)



e^+ - beam (PLIN)





Basic lattice segmentations in ML

						Length (m)
Name in Lattice	modules	without quad	with quad	without quad	warm section (m)	7.652
RFU#	RF unit (lengths in meters)	12.652	12.652	12.652		
	3 modules					37.956
CSTR#	"4" Long Cryo-String	RF unit 37.956	RF unit 37.956	RF unit 37.956	RF unit 37.956	end-box 2.50
	12 CM's plus string end box					154.324
CSTR#	"3" Short Cryo-String	RF unit 37.956	RF unit 37.956	RF unit 37.956	end-box 2.50	
	9 CM's plus string end box					116.368
CUNIT #	Cryo-Unit	Service end-box 2.500	CSTR	CSTR	CSTR	CSTR
					- - - -	CSTR CSTR



KCS: Layout of Cryo-Units

Electron Main Linac: (72 CSTR = 285 RFunits = 855 CM's)

CUNIT1	7.65	CUNIT2	7.65	CUNIT3	7.65	CUNIT4	7.65	CUNIT5	7.65	CUNIT6
--------	------	--------	------	--------	------	--------	------	--------	------	--------

	CSTR "4"		CSTR "3"		RF units
CUNIT1 =	5		2		26
CUNIT2 =	13				52
CUNIT3 =	13				52
CUNIT4 =	13				52
CUNIT5 =	13				52
CUNIT6 =	12		1		51
Total:	69		3		285

Sbox	01	02	03	04	05	06	07								
Sbox	08	09	10	11	12	13	14	15	16	17	18	19	20		
Sbox	21	22	23	24	25	26	27	28	29	30	31	32	33		
Sbox	34	35	36	37	38	39	40	41	42	43	44	45	46		
Sbox	47	48	49	50	51	52	53	54	55	56	57	58	59		
Sbox	60	61	62	63	64	65	66	67	68	69	70	71	72		

Legend: 7.65 Warm section 7.652m Sbox Service box ## Long (4-RFU) CSTR ## Short (3-RFU) CSTR

Positron Main Linac: (72 CSTR = 282 RFunits = 846 CM's)

CUNIT1	7.65	CUNIT2	7.65	CUNIT3	7.65	CUNIT4	7.65	CUNIT5	7.65	CUNIT6
--------	------	--------	------	--------	------	--------	------	--------	------	--------

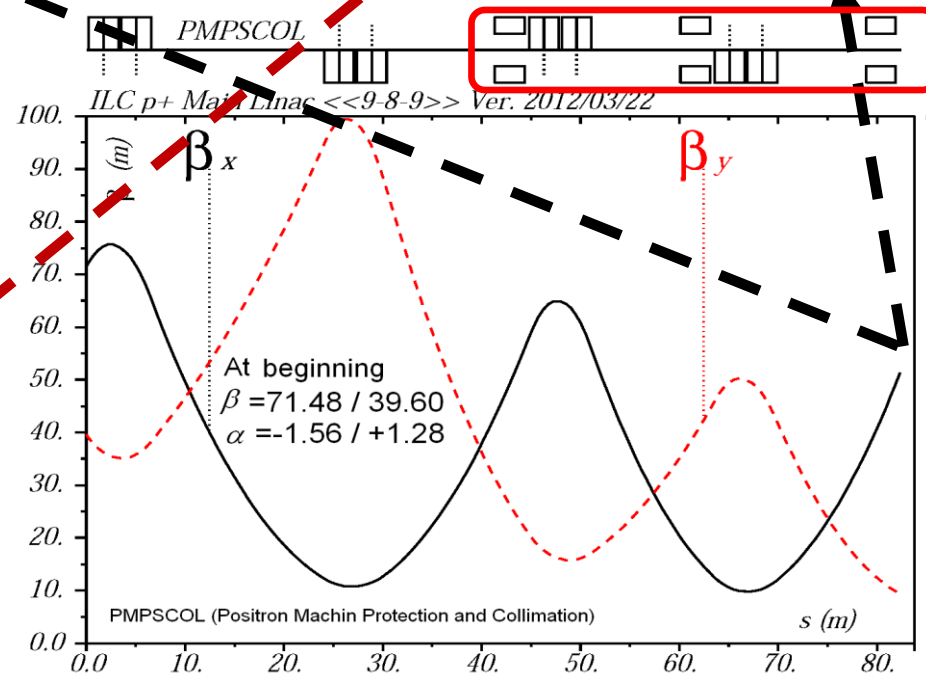
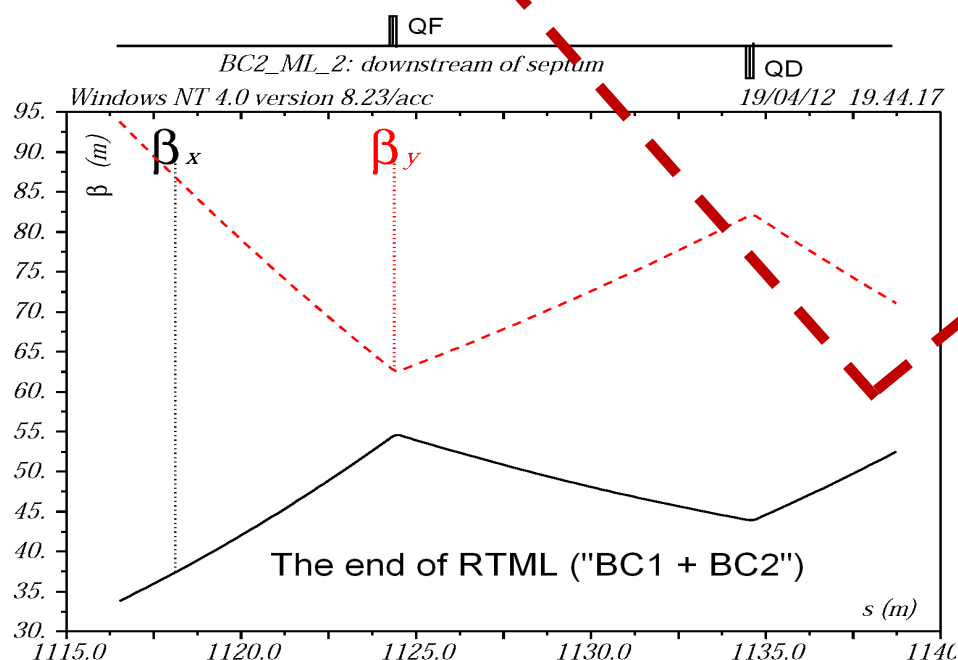
	CSTR "4"		CSTR "3"		RF unit
CUNIT1 =	5		2		26
CUNIT2 =	13				52
CUNIT3 =	13				52
CUNIT4 =	12		1		51
CUNIT5 =	11		2		50
CUNIT6 =	12		1		51
Total:	66		6		282

Sbox	01	02	03	04	05	06	07								
Sbox	08	09	10	11	12	13	14	15	16	17	18	19	20		
Sbox	21	22	23	24	25	26	27	28	29	30	31	32	33		
Sbox	34	35	36	37	38	39	40	41	42	43	44	45	46		
Sbox	47	48	49	50	51	52	53	54	55	56	57	58	59		
Sbox	60	61	62	63	64	65	66	67	68	69	70	71	72		



Treaty points: Optical Functions at ML boundaries

Electron side			Positron side		
Treaty Point	TERTML2ML	TEML2PS	TPS2EBDS	TPRTML2ML	TPML2BDS
	Electron RTML to Main Linac	Electron Main Linac to Positron Source (Undulator Section)	Positron Source (Undulator Section) to Electron BDS	Positron RTML to Main Linac	Positron Main Linac to BDS
Optics Functions					
α_x	-1.142	-2.402	-2.402	-1.142	-2.402
β_x [m]	52.67	51.33	51.33	52.67	51.33
α_y	1.279	0.4888	0.4888	1.279	0.4888
β_y [m]	70.74	9.395	9.395	70.74	9.395



- The RTML ends with defoc. Q => **ML starts with focusing Q**
- ML proper ends at the entry of PMSCOL (p+ machine protection & collimation)

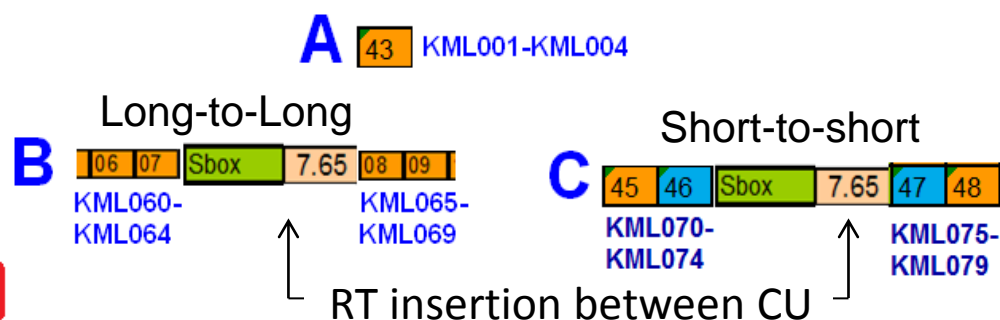


Quadrupoles in e⁻ ML (KCS) cells

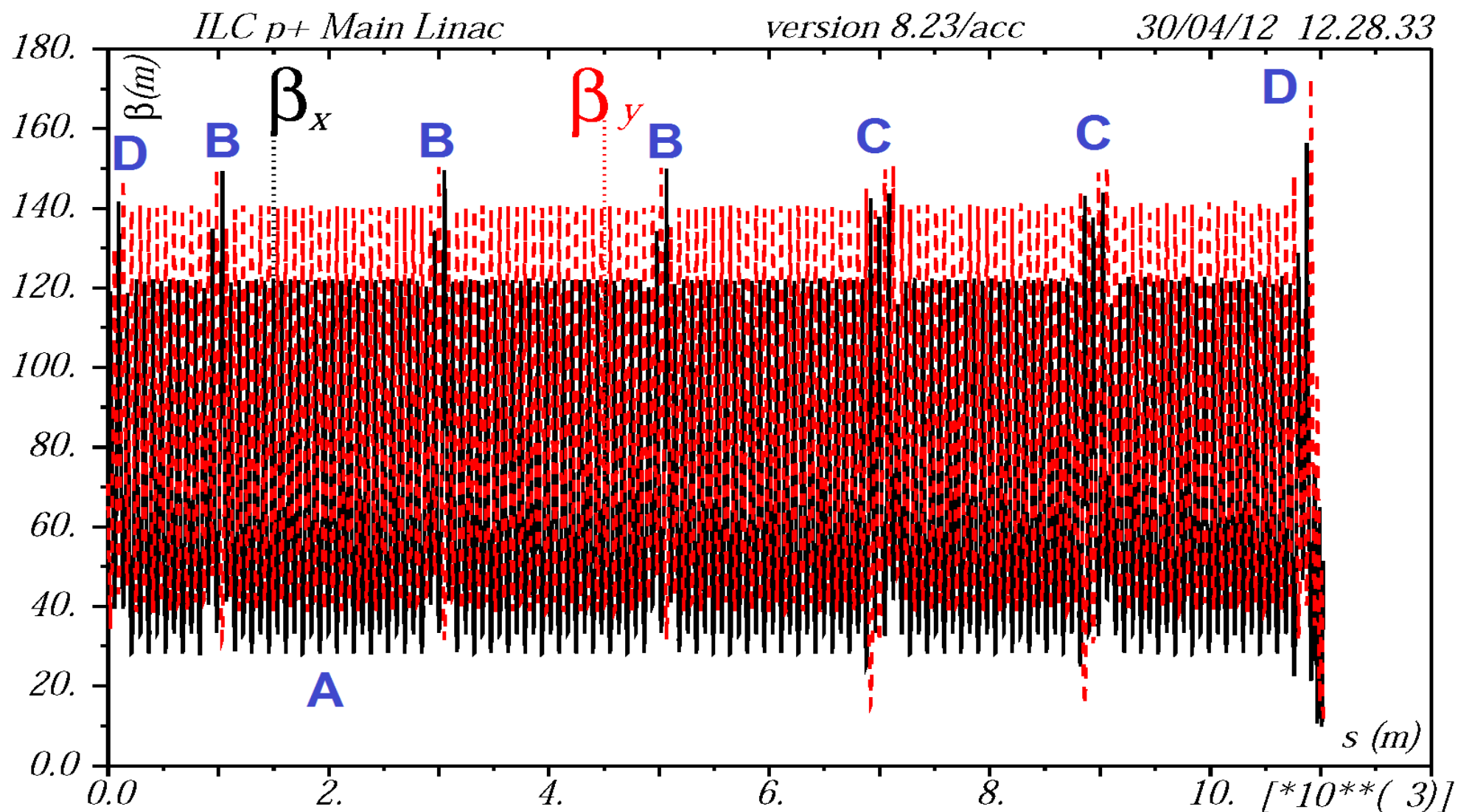
Basic configurations of focusing structure

- A: Quasi-periodical “long” 4-RFU CSTR inside of regular part of CUNITs :
2 FODO quasi-periodical cells (phase advances ~75/60 degrees) =>
4 quads with K1 denoted as K1=KML001, KML002, KML003, KML004
- B: Long 4-RFU CSTR between CUNIT ends separated by warm sections:
“5+5” quad configuration around warm sections with K1 denoted as
KML060-KML064 and KML065-KML069
- C: Two short 3-RFU CSTR at connections of CUNIT#4 with CUNIT#5, and CUNIT#5
and CUNIT#6 (for PLIN only): “5+5” quad configuration around warm sections
with K1 denoted as KML070-KML074 and KML075-KML079
- D: 6 quads at the ML beginning and 6 quads at the ML end are used for matching to
the Twiss parameters b and a at ML boundaries.

	D						Positron ML (KCS)																	
Sbox	01	02	03	04	05	06	07																	
Sbox	08	09	10	11	12	13	14	15	16	17	18	19	20											
Sbox	21	22	23	24	25	26	27	28	29	30	31	32	33											
Sbox	34	35	36	37	38	39	40	41	42	43	44	45	46											
Sbox	47	48	49	50	51	52	53	54	55	56	57	58	59											
Sbox	60	61	62	63	64	65	66	67	68	69	70	71	72	D										



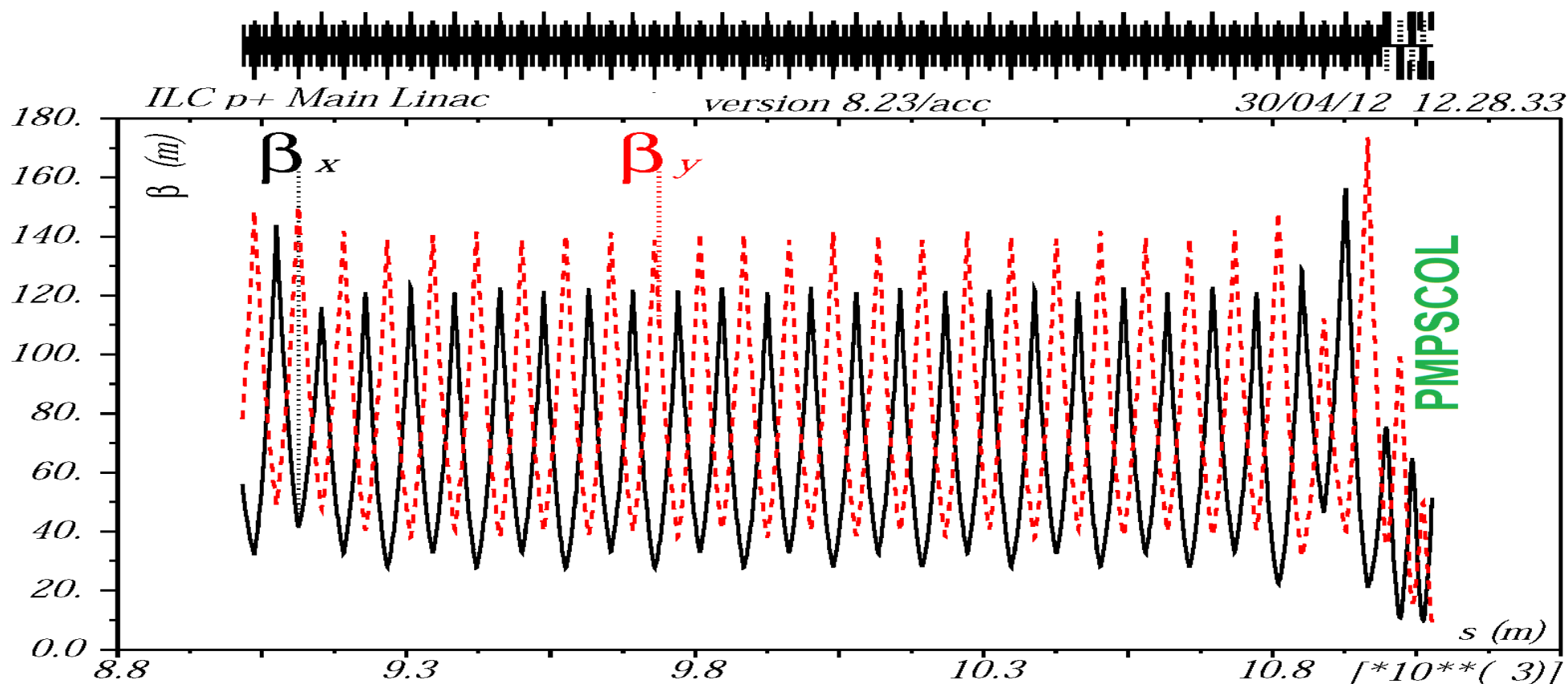
Matched β -functions in PLIN (KCS)



PLIN & PMPSCOL junction

- RTML ends with **defocusing** Quad => ML9 (e⁺&e⁻) starts with **focusing** Quad
- **Positron** ML with 282 Quads (= #RFU) ends with defocusing Quads
- ➔ PMSCOL starts with focusing Quad

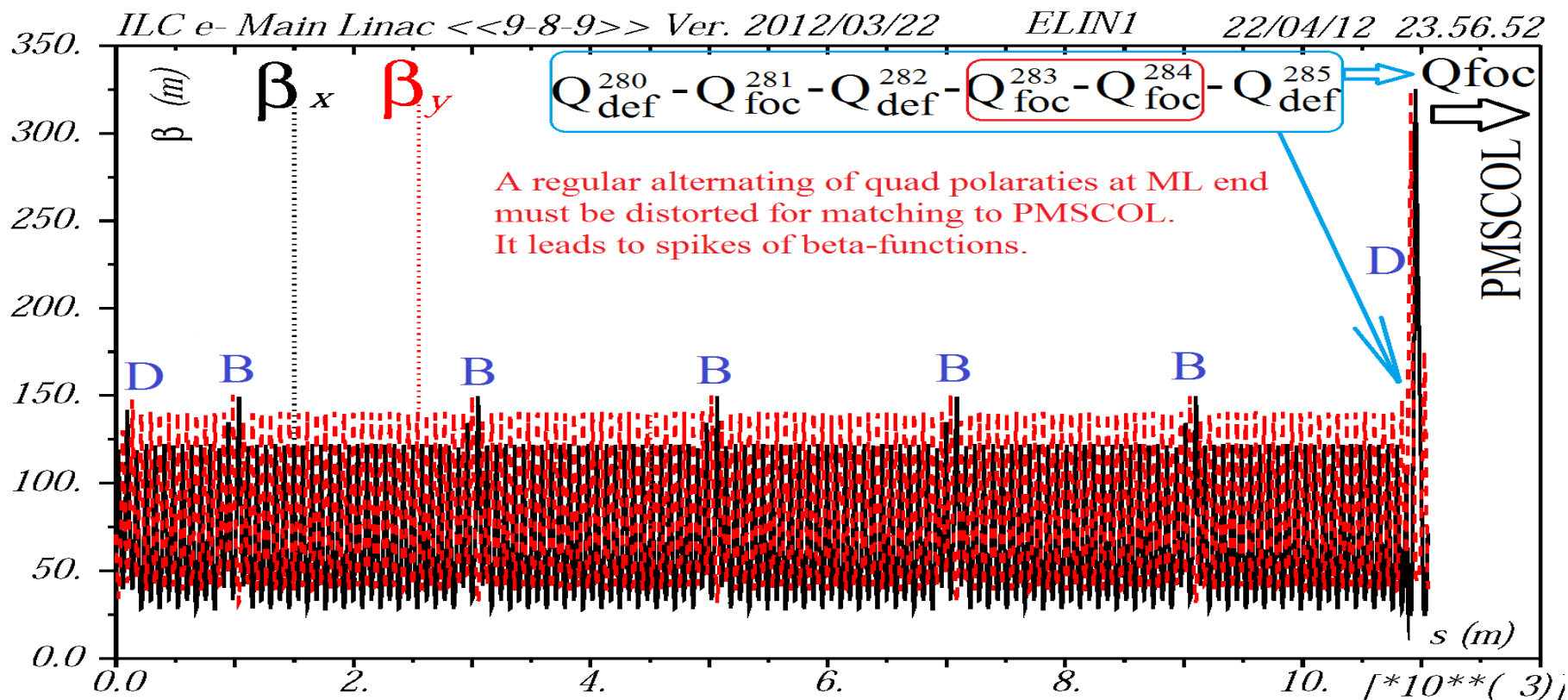
=> Alternative polarity of Quads is kept throughout total Positron ML



Matching features for ELIN

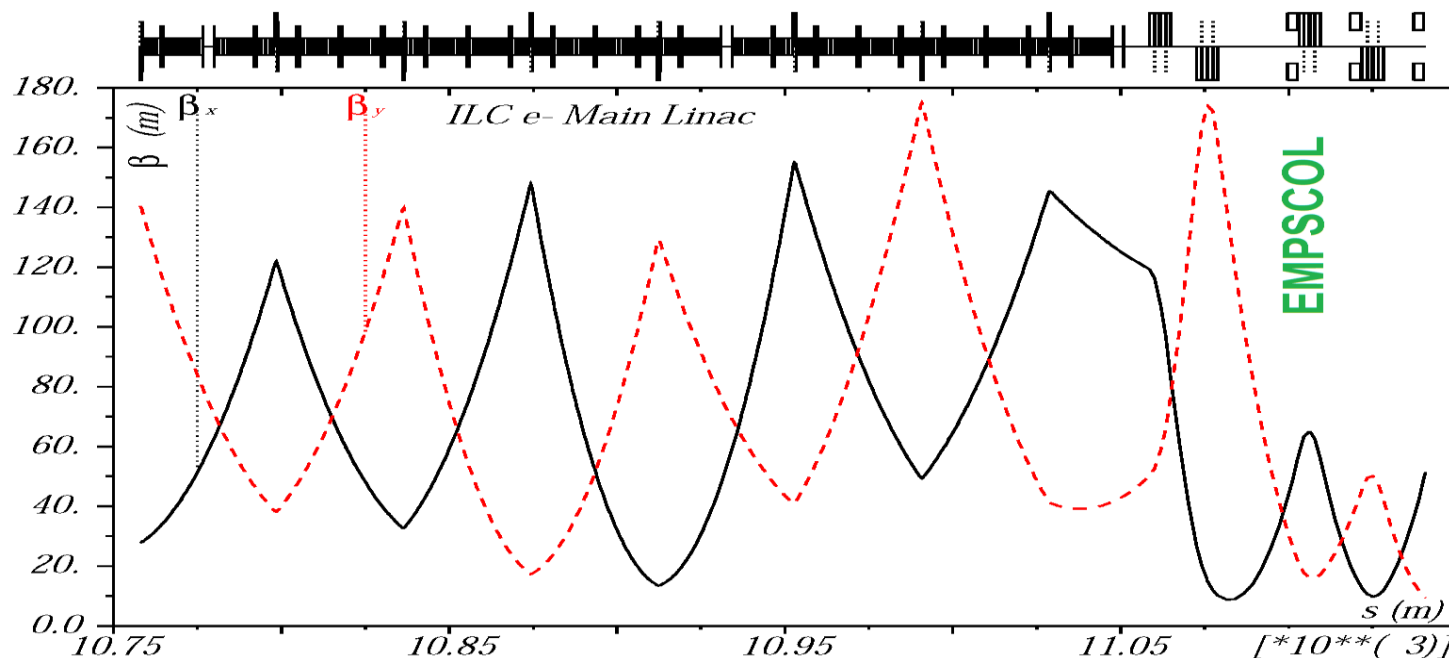
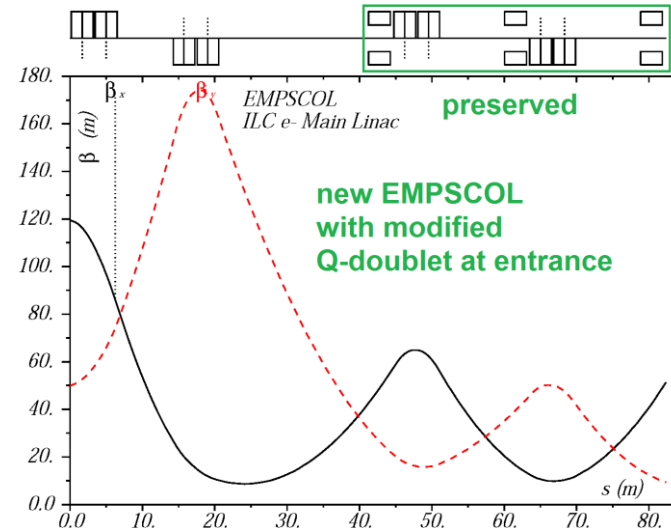
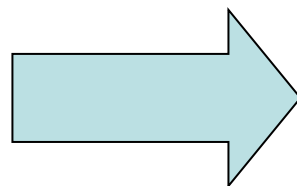
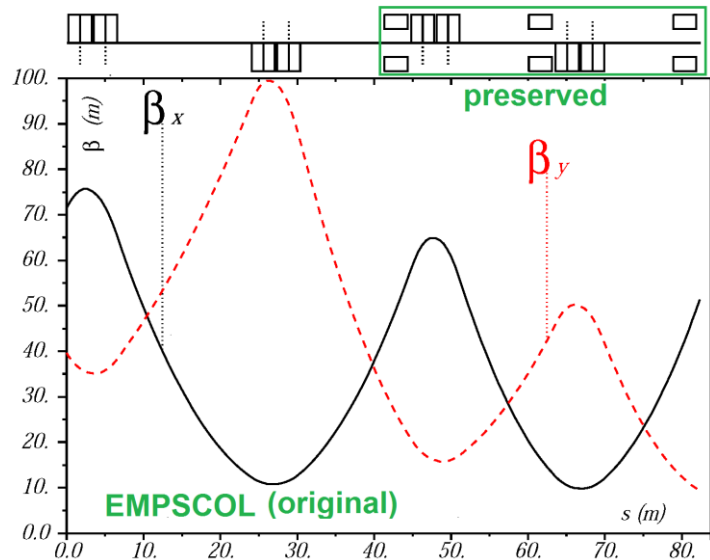
- ELIN has a regular structure inside; no need for “C”- type matching quads
- **Electron** ML with 285 Quads starts and ends with **focusing Quads**
- PMSCOL starts with **Q-foc** => the same Q-polarities at ELIN & COL junction

=> *Difficult matching with COLL structure resulting in large β -functions*



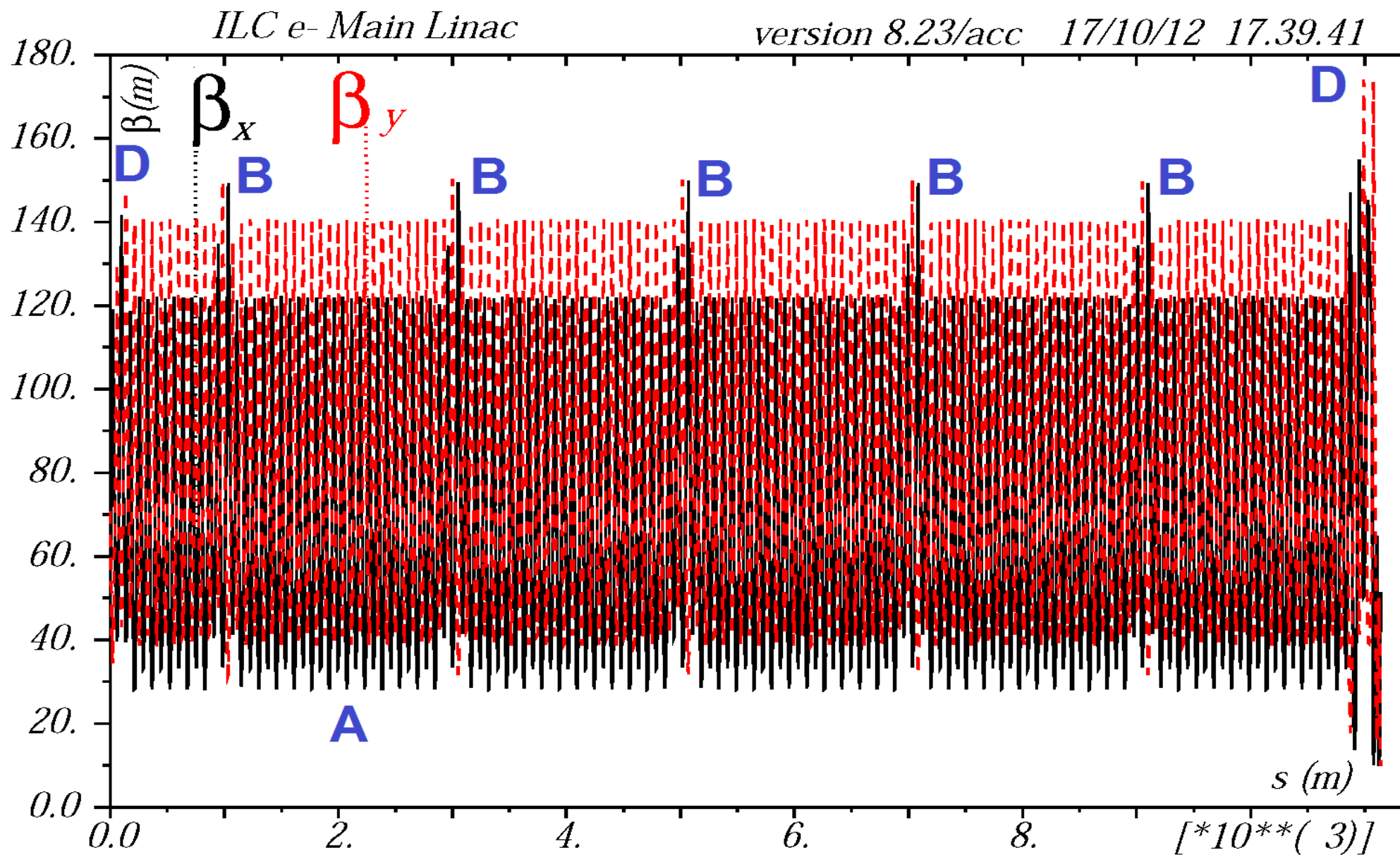
A bad matching at ELIN-end with **original** COL-structure (large β -function)

Improved ELIN & EMPSCOL junction





Matched β -functions in ELIN (KCS)





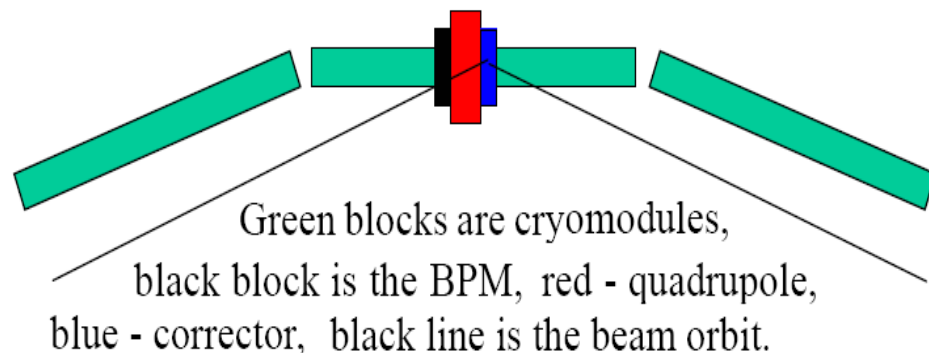
Earth Curvature implementation

- ML follows curvature of the Earth 's surface
- Each CM is straight and aligned along the Earth horizon and the beam-line is kinked at the ends of CMs
- Kinks between CM's are implemented in MAD8 as a thin KML-lines consisting of a dipole (MULT, KOL=p) & a vert. corrector (VKICK):
(The former changes both ref. frame and beam trajectory, the latter cancel the trajectory change)
- In MAD8 curvature (KMLs) are switched ON/OFF by “SET, CURVE, 1” & 0
- KML-lines are set at both ends of every CM. Several types:
 - *KML1 – between CMs inside of RFUs*
 - *KMLQ – at the ends of CM with quads*
 - *KML2 – between CMs at CSTRs ends*
 - *KML4 – between CMs at CUNITs ends*
 - *KML5 – at the end of the last CM (at ML exit)*
 - *KML8 – at the beginning of the first CM (ML entrance)*

Steering to the Earth's curvature

Constraints:

- The beam trajectory is steered by vertical correctors through the centers of quads, i.e. only at every 3rd CM.
- Steering can be switched “ON/OFF” by “SET, STEER,1” or 0



Match corrector strengths along ML:

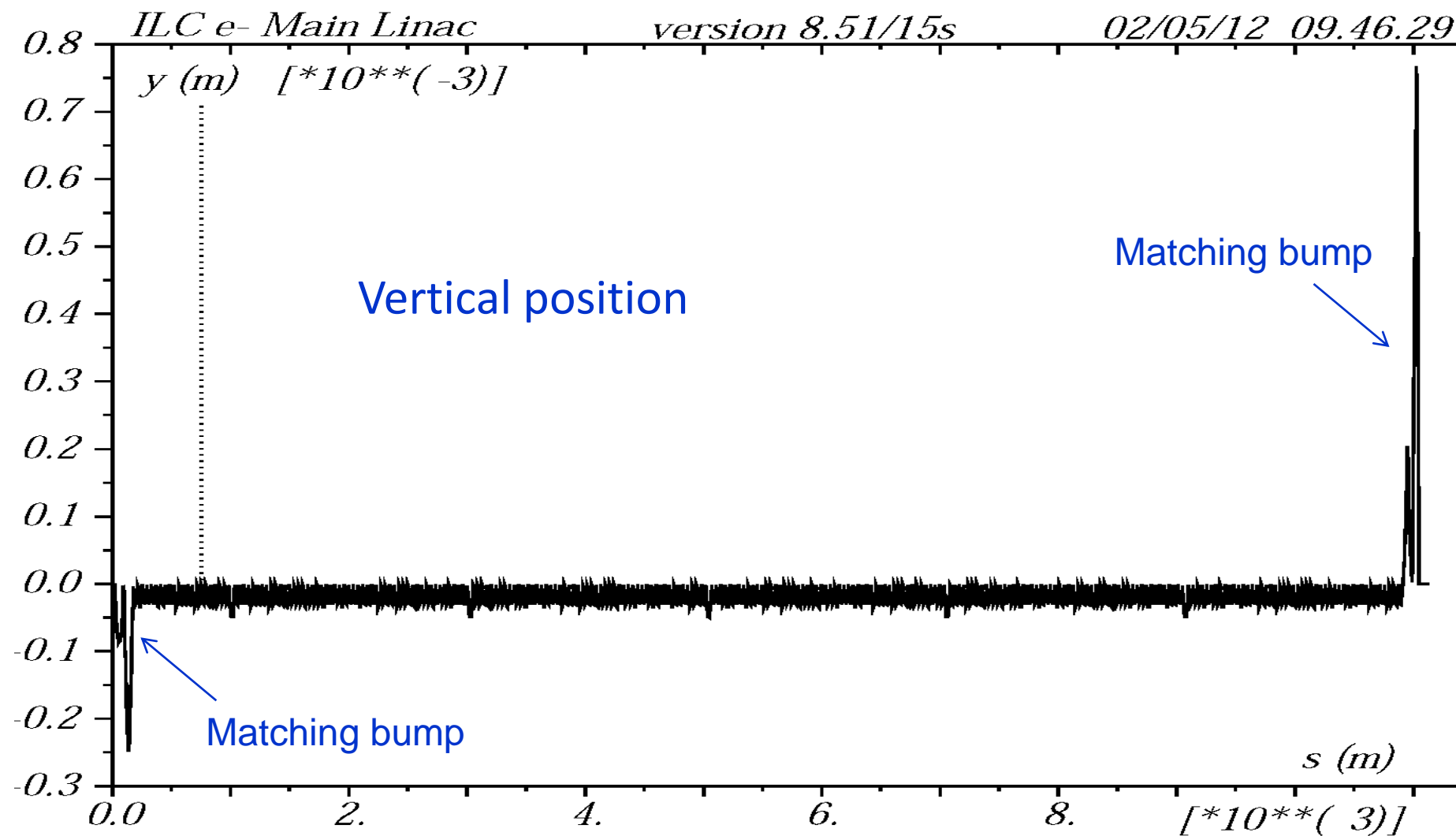
```
MATCH, BETA0=TWSS0
VARY, AMLY# (# = 10, 11,13,15,22,23,25)
.....
CONSTR, PATTERN="YML...", Y=0
LMDIF, TOL=1.E-20,...
MIGRAD, TOL=1.E-20, ...
ENDMATCH
```

Match AML26, AML27 at exit:

```
MATCH, BETA0=TWSS0
VARY, AMLY26, STEP=1.E-9
VARY, AMLY27, STEP=1.E-9
CONSTR, #E, Y=0, PY=0
LMDIF, TOL=1.E-20, CALLS=5000
MIGRAD, TOL=1.E-20, CALLS=5000
ENDMATCH
```

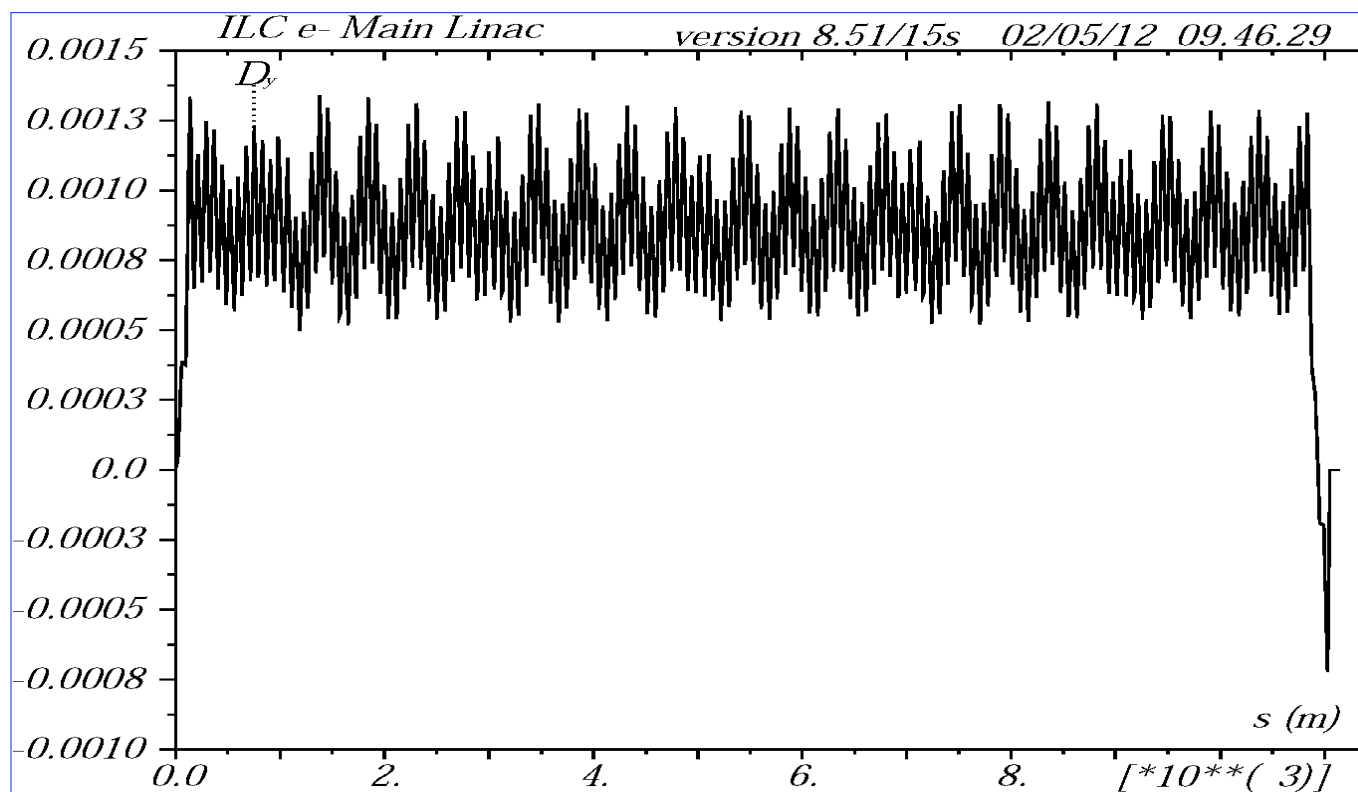
Notice: Another possible constraint with $Y > 0$ (instead of $Y=0$) minimizing wake-field effects (Kubo's proposal) is not realized yet in the present ML lattice.

Beam orbit after steering (KCS)



Dispersion minimization (KCS)

- The beam injected into ML must be matched to the periodic dispersion in curved lattice
- The optimal dispersion at injection (TDY & TDPY) is found by minimizing D_y at every defocusing quad.



! Find TDY & TDPY

SET, CURVE, 1; SET, STEER, 1;

SET, BUMPS, 0; USE, PLIN1

MATCH, BETA0=TWSS0

VARY, TDY; VARY, TDPY

WEIGHT, WX=1.E-9

CONSTR, PATTERN="MQD.*", DY=0

LMDIF, TOL=1.E-20;

ENDMATCH

! Save solution at the 6th RFU

SET, **MDY, TWSS_QML006[DY]**

SET, **MDPY, TWSS_QML006[DPY]**



Matching D_y & ref. orbit at ML entrance

- RTML end with $DY=0$ & w/o curvature is matched into ML beginning with $DY \neq 0$ & $CURVE=>1$
- 5 additional vertical kicks (AMLYi+**AML****DY##i**) for 5 first correctors at ML beginning are switched on by “SET, BUMPS,1”

SET, CURVE, 1; SET, STEER, 1

SET, BUMPS, 1; USE, PLIN1

SAVEBETA, **TWSS1**, **YML003**

SAVEBETA, **TWSS2**, **YML005**

TWISS, BETA0=TWSS0

MATCH, BETA0=TWSS0

VARY, AMLDY11i (12i, 13i, 14i, 15i);

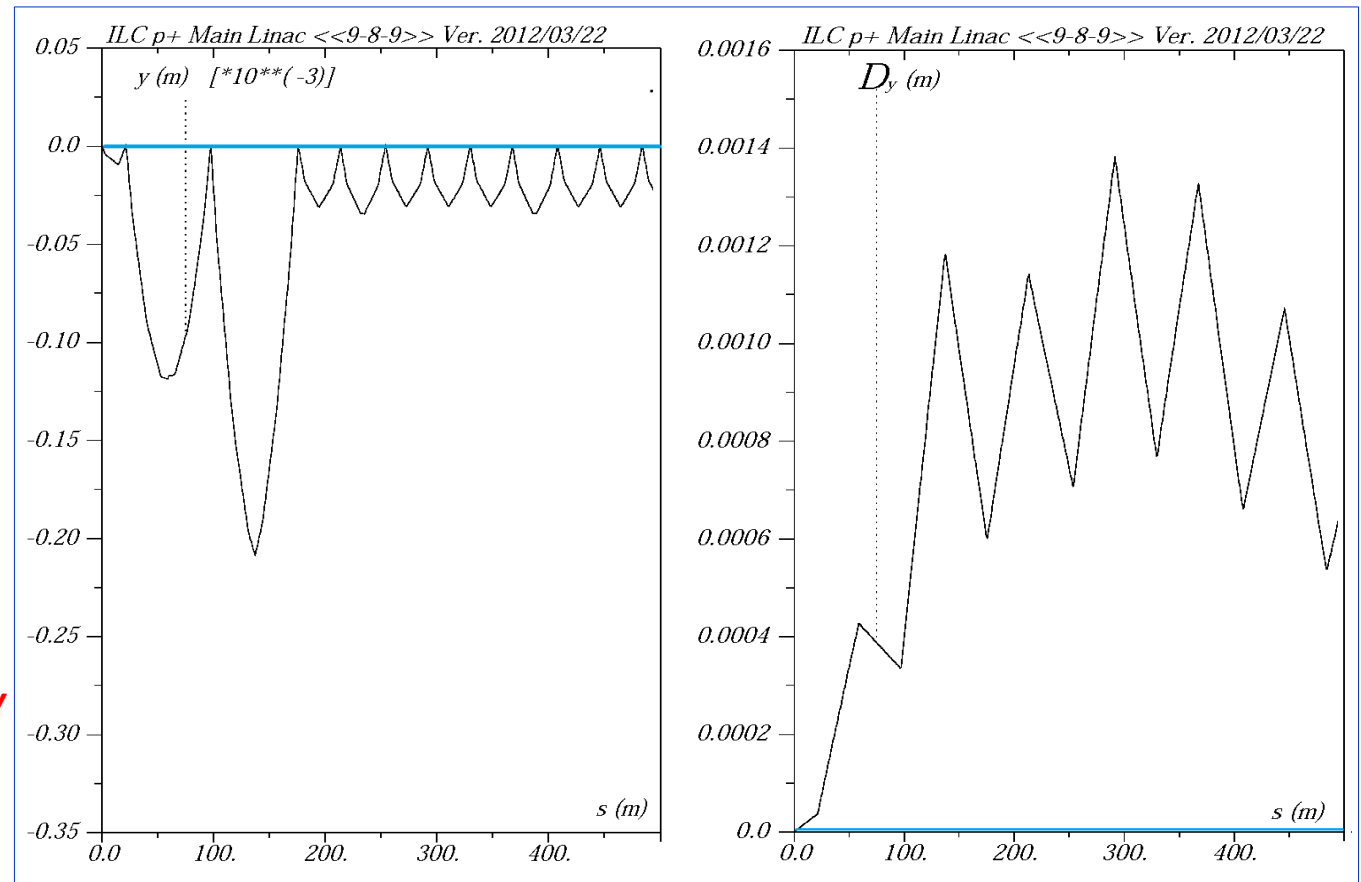
CONSTR, YML003, Y=TWSS1[Y]

CONSTR, YML005, Y=TWSS2[Y],
PY=TWSS2[PY]

CONSTR, QML006[1], DY=MDY, DPY=MDPY

LMDIF (MIGRAD), TOL=1.E-20;

ENDMATCH





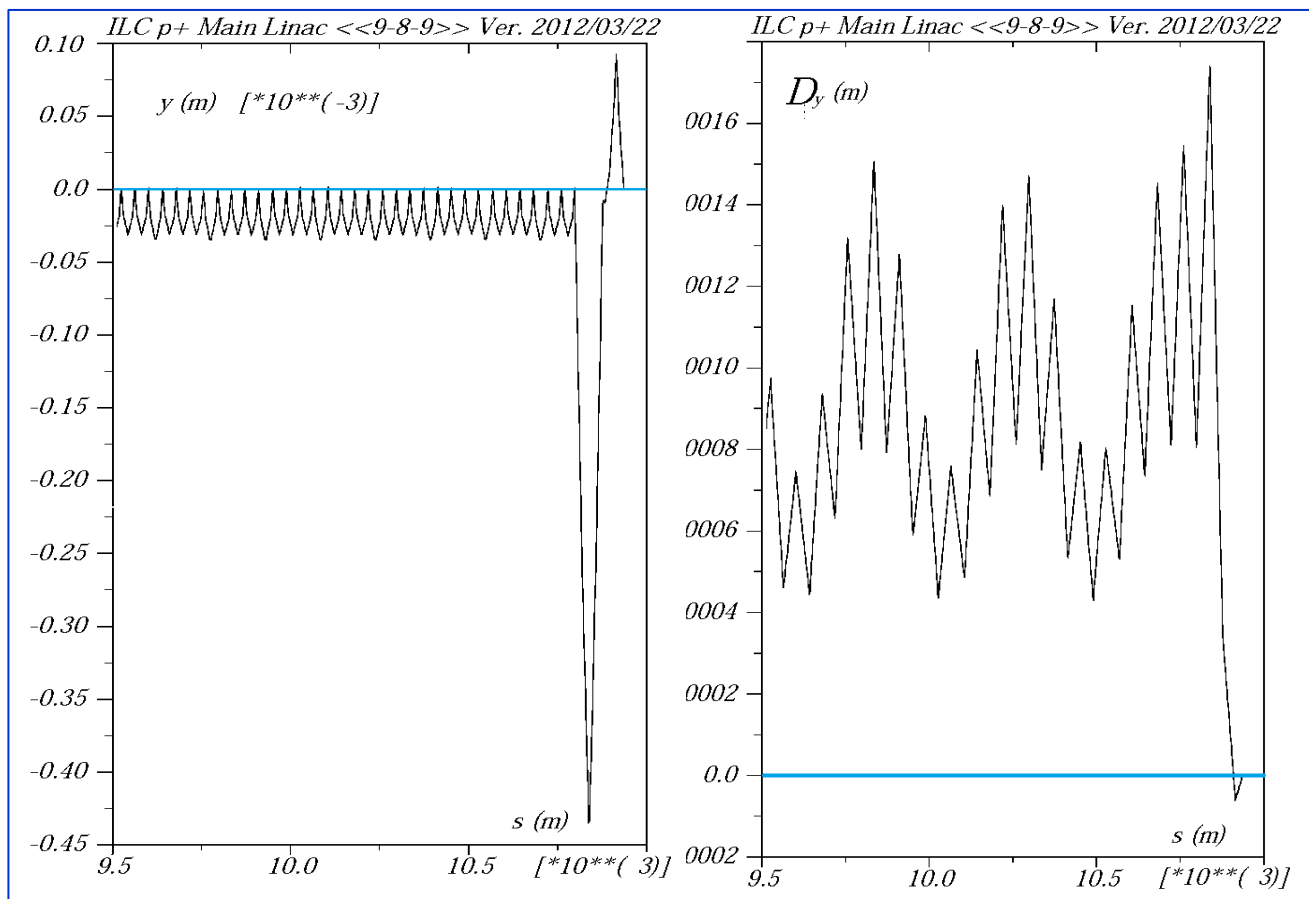
Matching DY & ref. orbit at the ML end

- ML end with $DY \neq 0$ & $CURVE \Rightarrow 1$; is matched PMSCOL end with $DY=0$ & w/o curvature
- 5 additional vertical kicks (AMLYi+ AMLDY##o) for the last correctors at ML end are switched on by “SET, BUMPS,1”

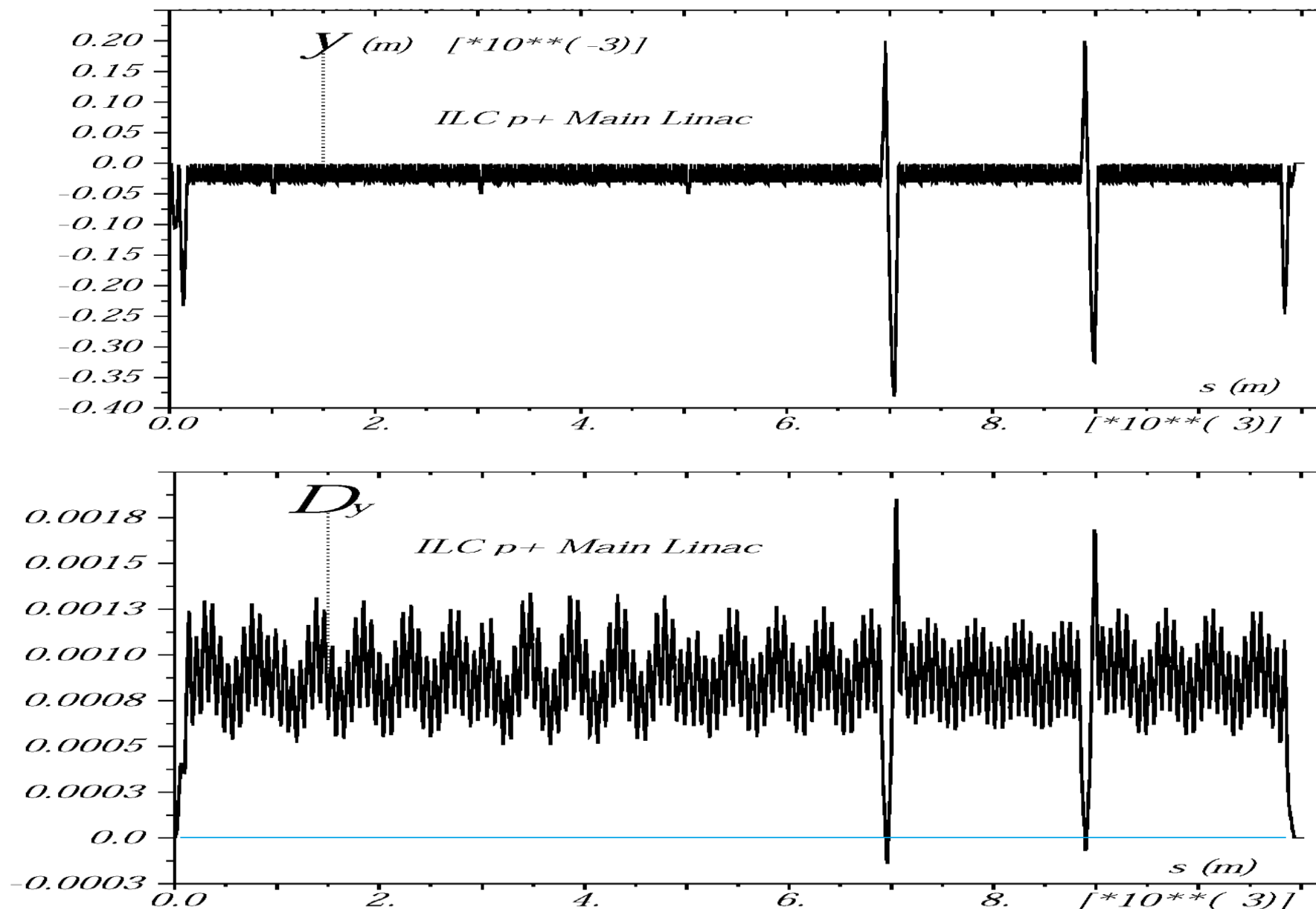
!PLIN example:

```
SET, CURVE, 1; SET, STEER, 1
SET, BUMPS, 1; USE, PLIN1
SAVEBETA, TWSS1_YML281, YML281
!next-to-last
TWISS, BETA0=TWSS0

MATCH, BETA0=TWSS0
VARY, AMLDY21o (22o, 23o, 24o, 25o);
CONSTR, YML281, Y=TWSS1_YML281[Y]
CONSTR, YPLIN2o, Y=0, PY=0, DY=0,
DPY=0
LMDIF (MIGRAD), TOL=1.E-20;
ENDMATCH
```



Matched DY & Y throughout PLIN (KCS)





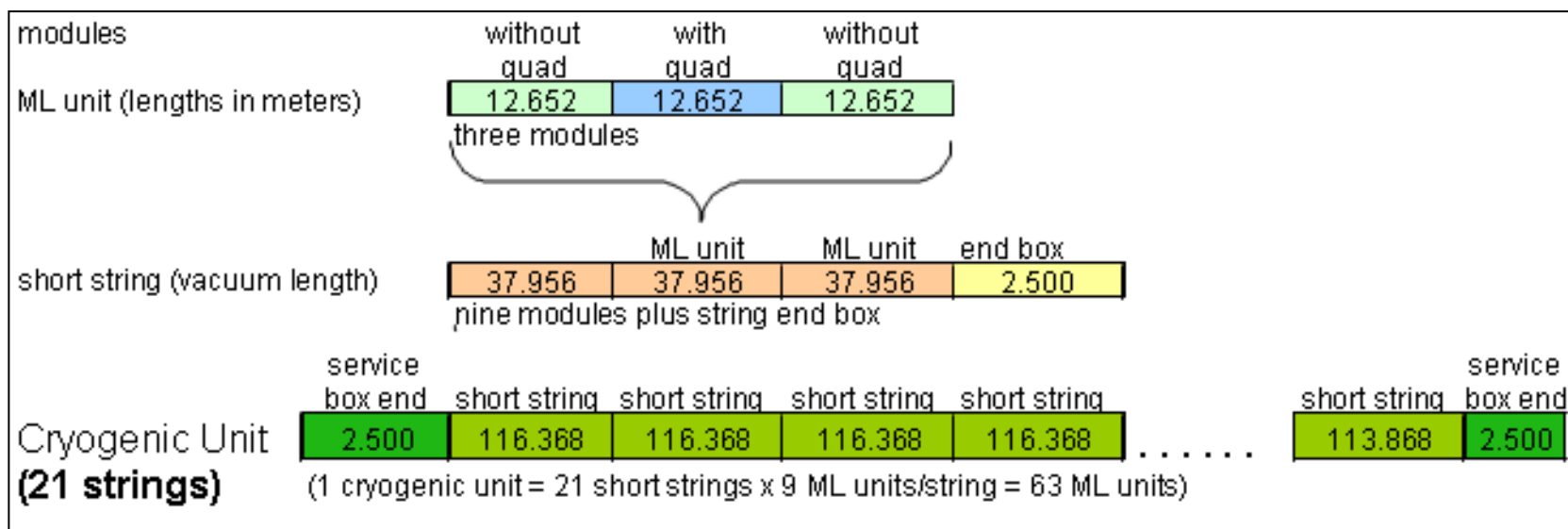
Summary for KCS-lattice status

- Main Linac lattices (9+4Q4+9 configuration) for TDR version have been re-designed, tuned and matched
- Tuning and matching subroutines previously created for RDR in 2007 are checked and adaptively modified for TDR-2012 version
- Presented outlook of lattice tuning is a helpful reference in a future, since the CM length can be slightly changed in the final designs
- ML lattices are ready for a further non-optical “text-information” polishing (like MAD8 “TYPE” statements)
- ML lattices are documented and will be posted at ILC EDMS.



ML DKS version (Ver. 25/6/2012)

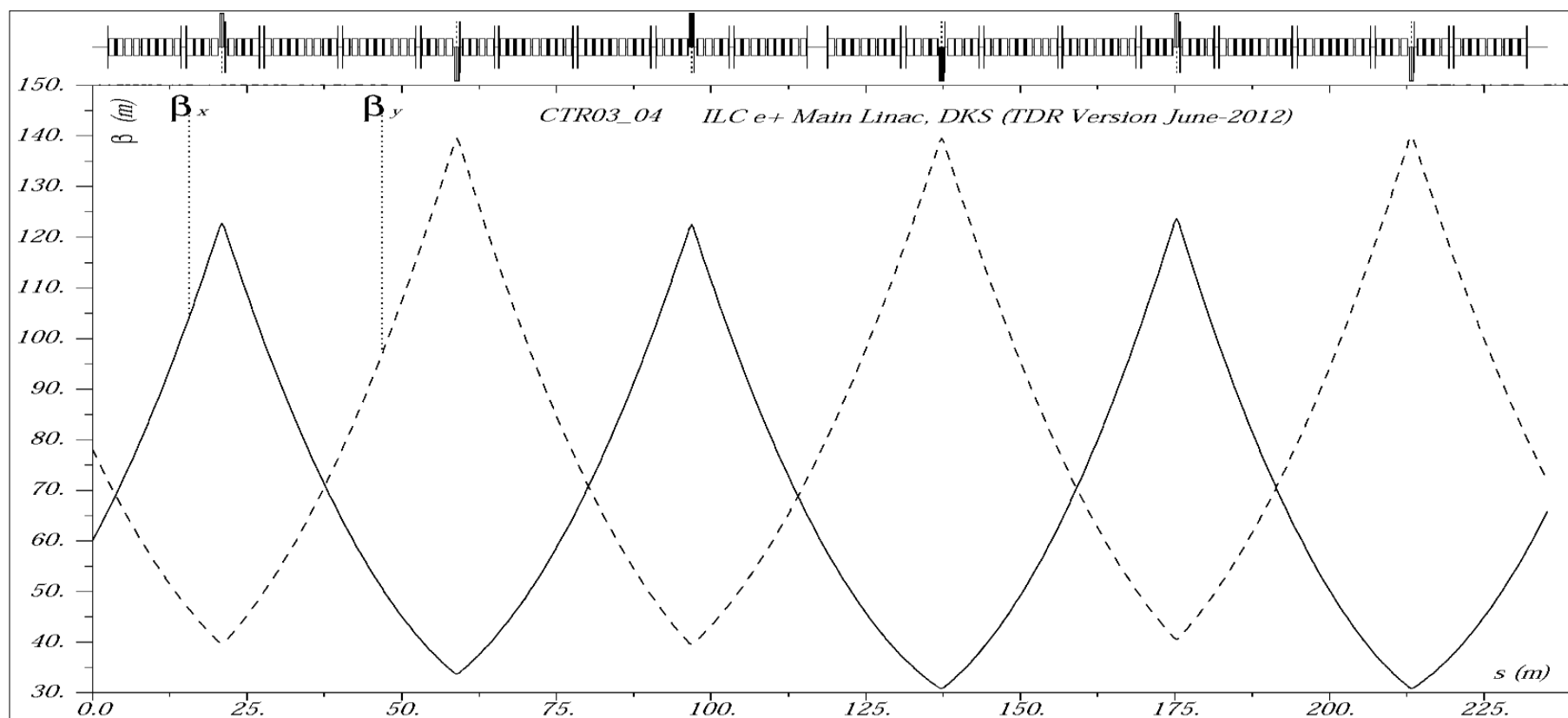
long str 11 short str 33 ML units	warm drift space	long str 21 short str 63 ML units	warm drift space	0 21 short str 63 ML units	warm drift space	long str 21 short str 63 ML units	warm drift space	0 21 short str 63 ML units	warm drift space	long str 21 short str 63 ML units	warm drift space	0 21 short str 63 ML units	collimation section	undulator protection section	66 undulators	Source
1282.5	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	82.4	295.9	318.0	BDS
Electron ML 285 RF units																DR
0 11 short str 33 ML units	warm drift space	0 21 strings 63 ML units	warm drift space	0 21 strings 63 ML units	warm drift space	0 21 strings 63 ML units	warm drift space	0 21 strings 63 ML units	warm drift space	0 20 strings 60 ML units	warm drift space	0 20 strings 60 ML units	collimation section			Source
1282.5	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	7.652	2446.2	82.340			BDS
Positron ML 282 RF units																DR



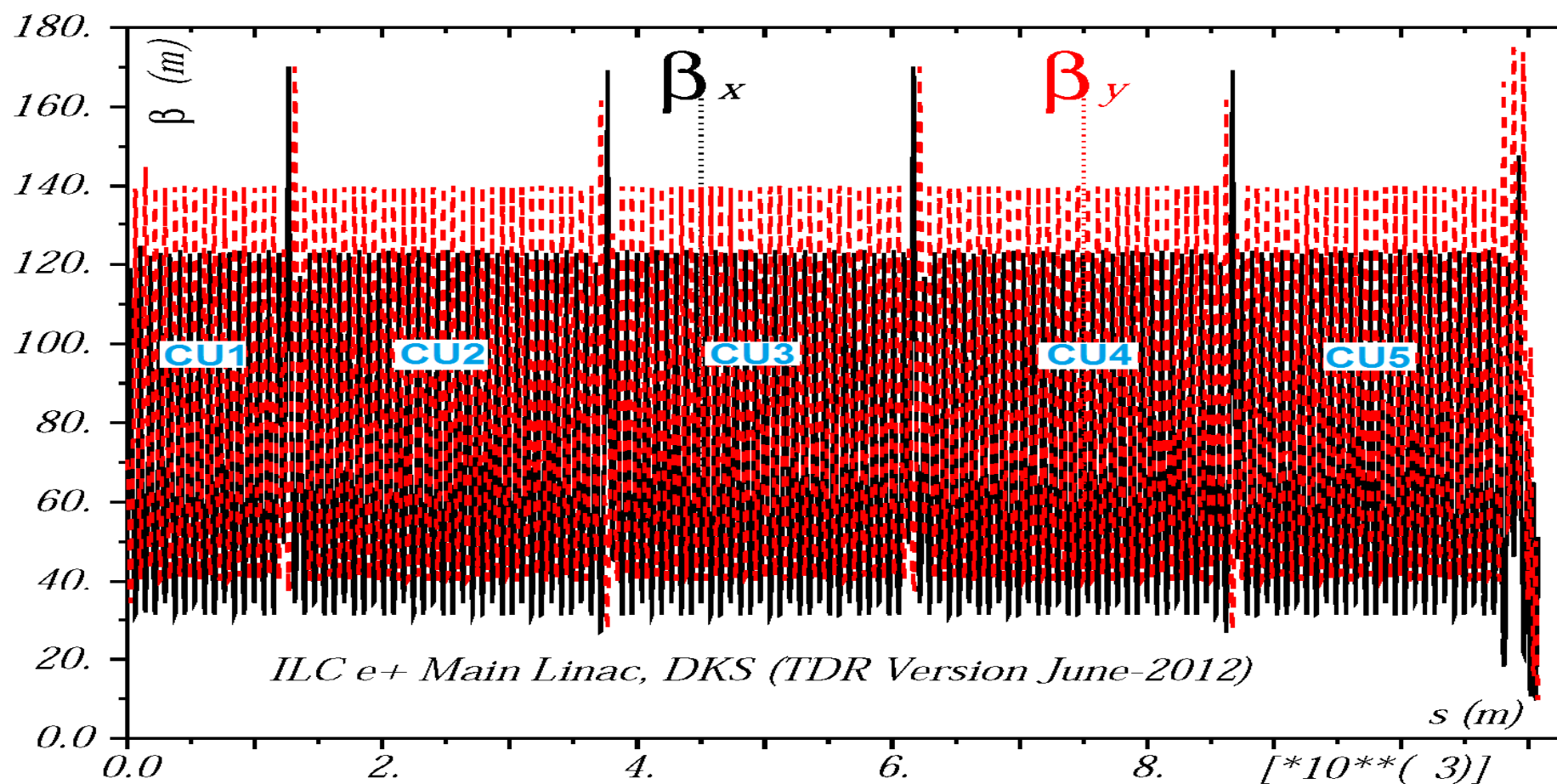


Matched β -functions in PLIN (DKS)

Two strings (6-RFU) forms quasi-periodical 6-Q strong focusing cell
(phase advances $\sim 3 \times (75/60)$ degrees) \Rightarrow
6 quads with K1 denoted as K1=KML001-KML006



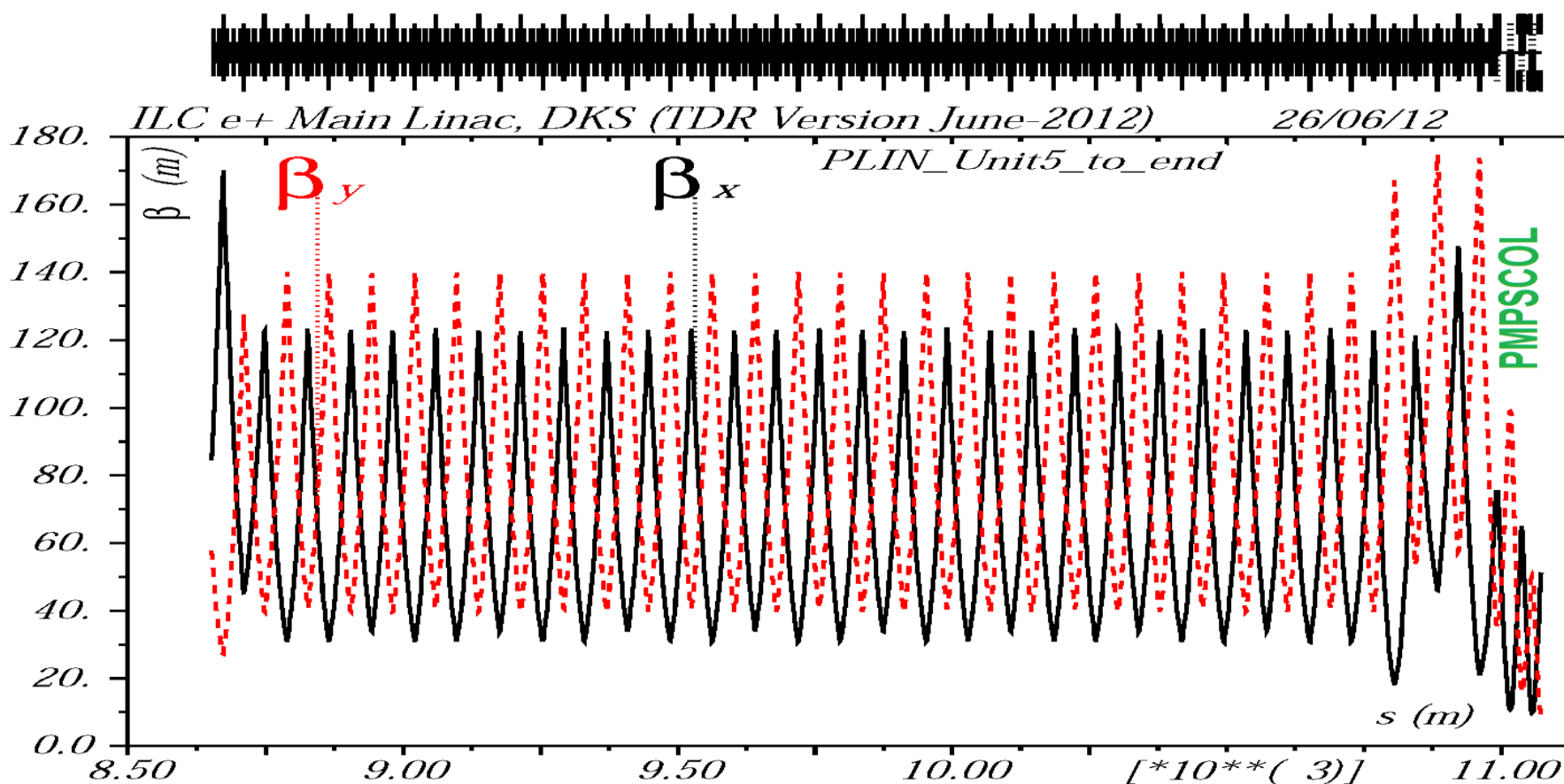
Matched β -functions in PLIN (DKS)



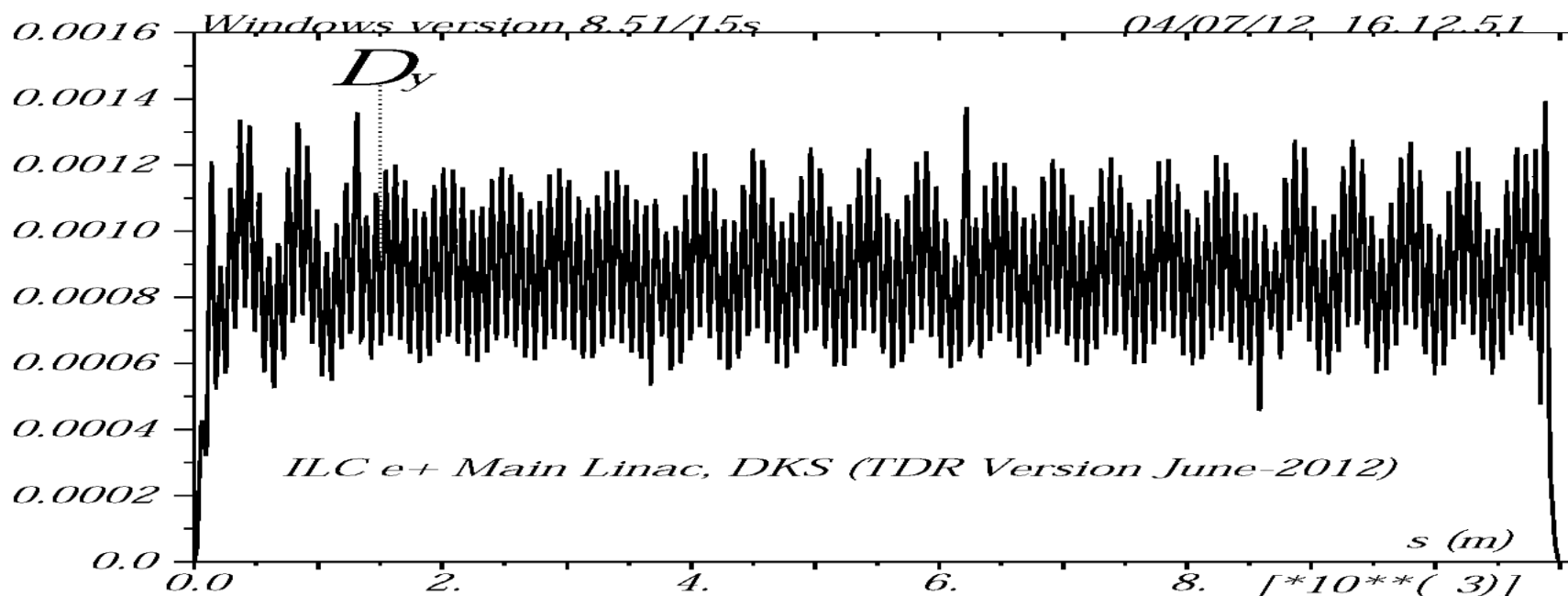
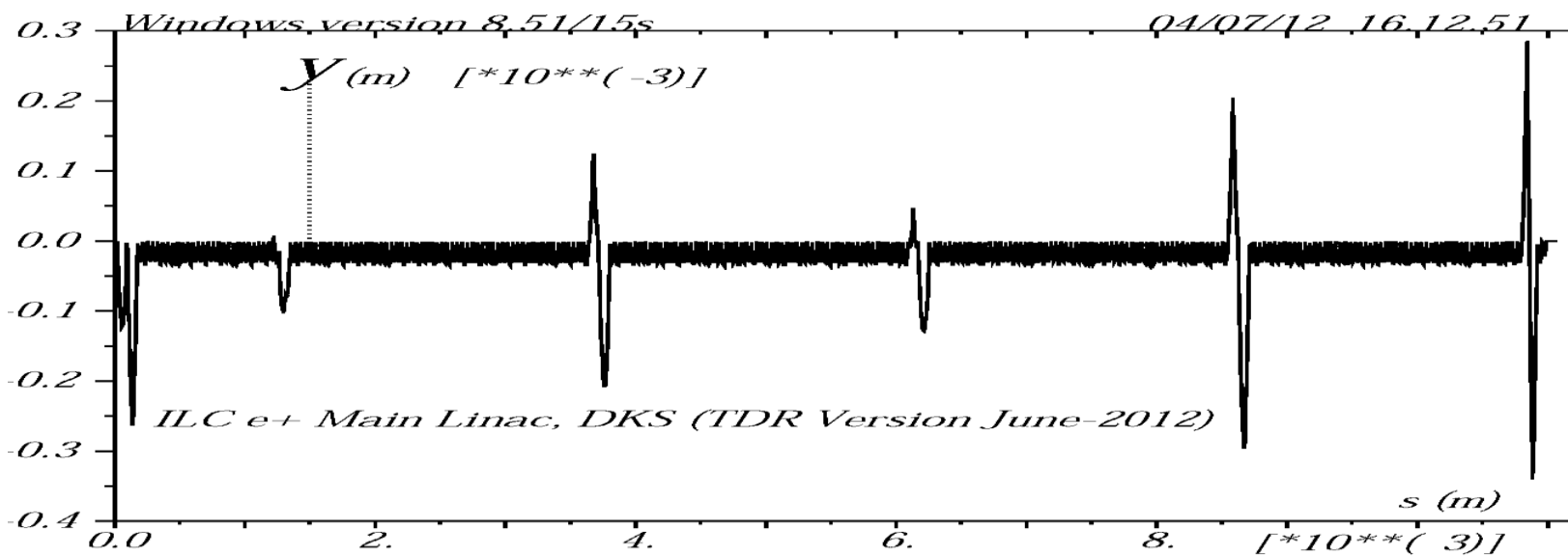
matching at ML entry (after RTML) with 6 quads KML031-KML036;
 matching between CU1 & CU2 and CU3 & CU4 with 6 quads KML051-KML056;
 matching between CU2 & CU3 and CU4 & CU5 with 6 quads KML041-KML045.

PLIN & PMPSCOL junction (DKS)

Alternative polarity of quads is kept throughout total Positron ML
 => easy matching at the junction with PMPSCOL (original)



Matched DY & Y throughout PLIN (DKS)

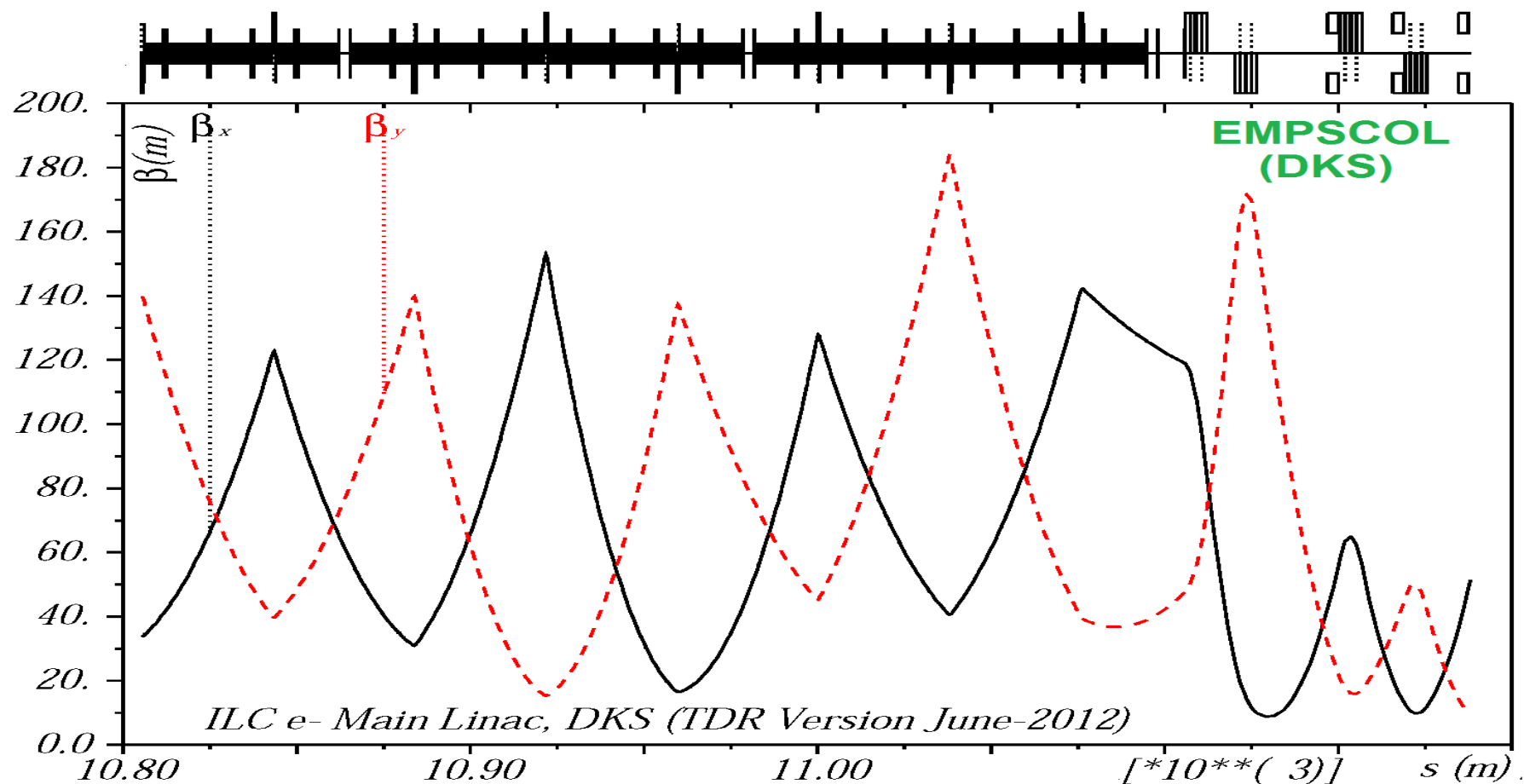


ELIN & EMPSCOL junction (DKS)

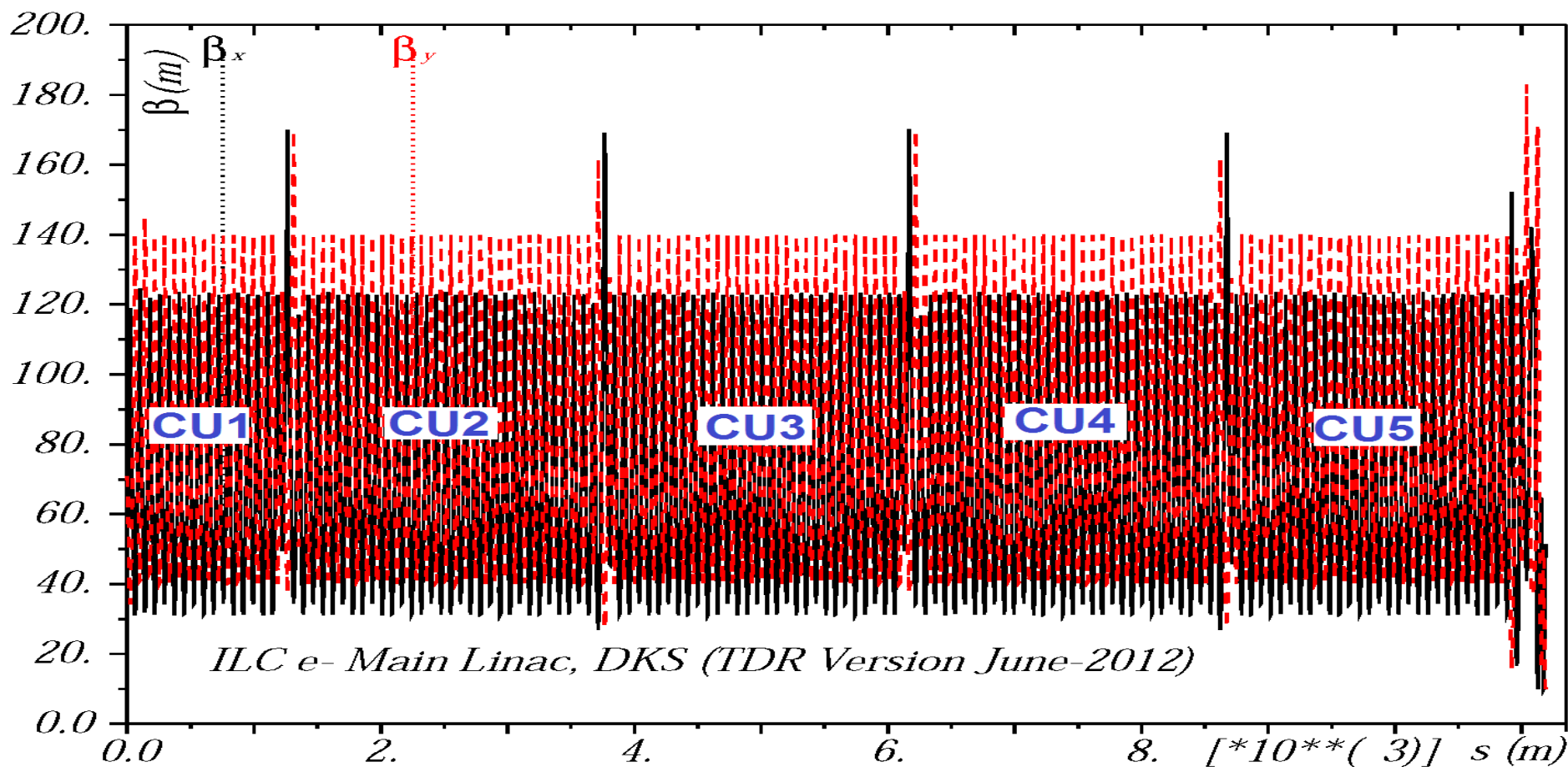
A regular alternating polarity of quads is distorted at the junctions with EMPSCOL.

➔ It leads to large spikes of β -function.

To avoid spikes Q-doublet at the EMPSCOL entrance is modified (similar to KCS)

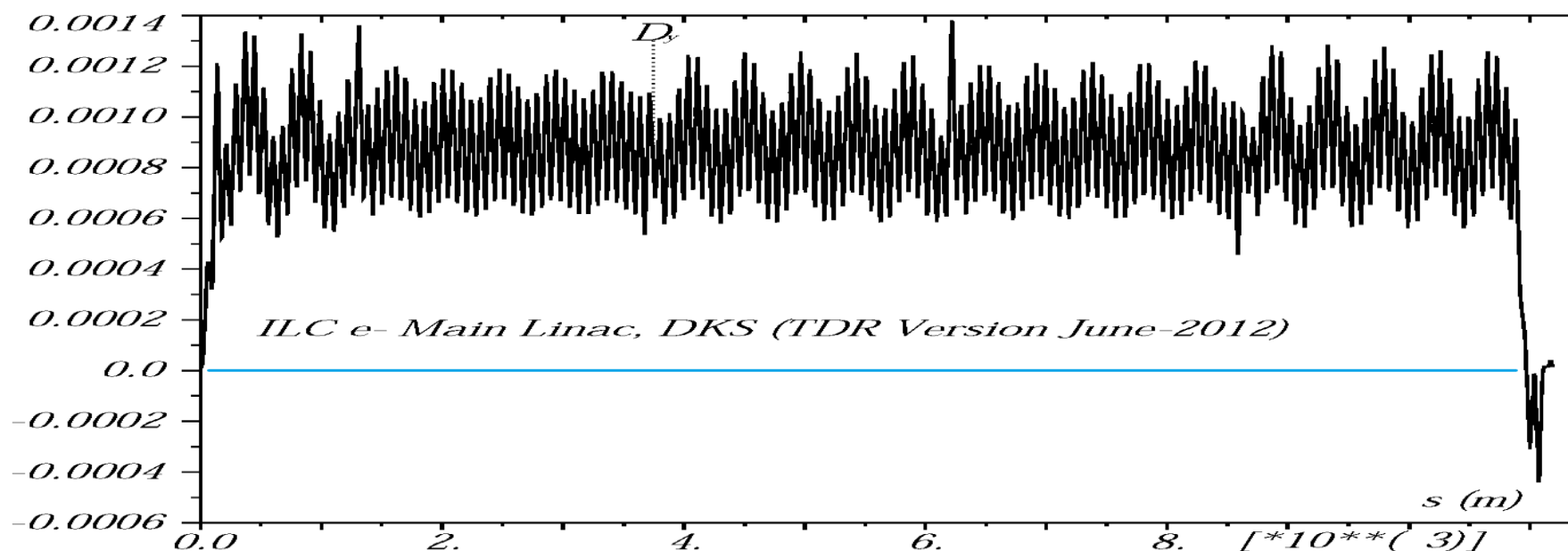
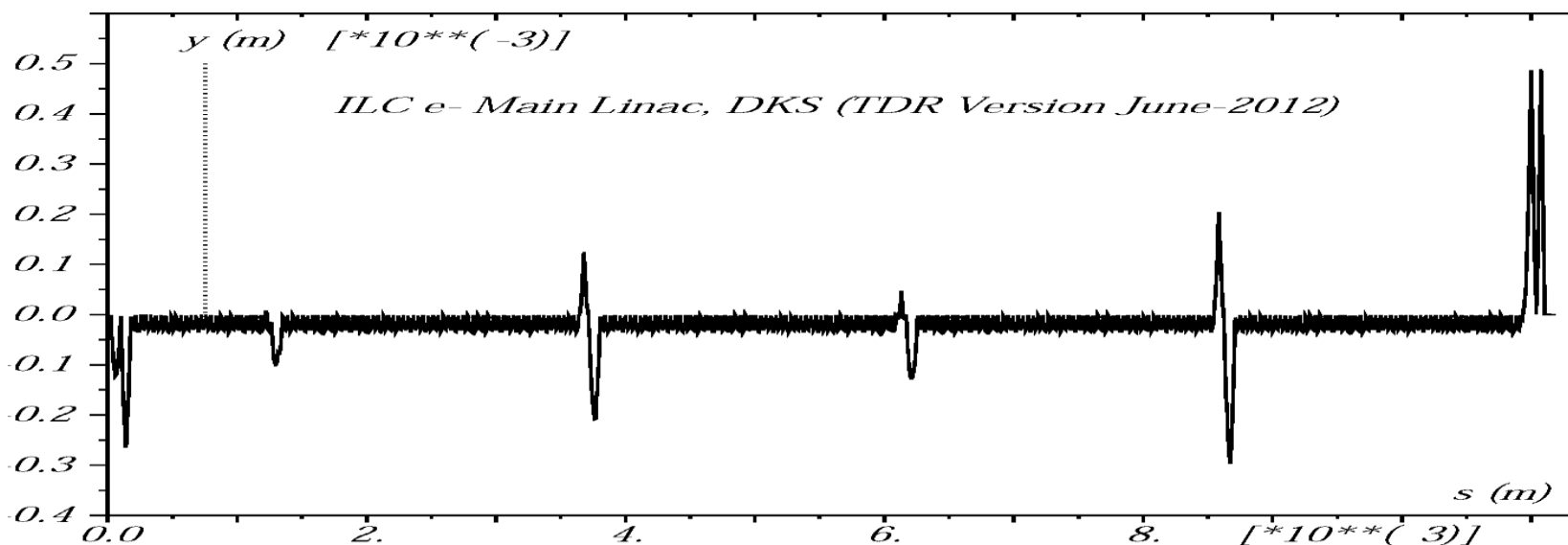


Matched β -functions in ELIN (DKS)



matching at ML entry (after RTML) with 6 quads KML031-KML036;
 matching between CU1 & CU2 and CU3 & CU4 with 6 quads KML051-KML056;
 matching between CU2 & CU3 and CU4 & CU5 with 6 quads KML041-KML045.

Matched DY & Y throughout ELIN (DKS)



Conclusion

- Two Lattices for KCS and DKS Main Linac layouts were designed, based on approach developed for RDR design
- Earth curvature was incorporated in design. Beam reference orbit and dispersion were optimized.
- Tuning and matching subroutines previously created for RDR in 2007 are checked and adaptively modified for TDR-2012 version
- Both ML lattices are documented and will be posted at ILC EDMS.