



The Measurement of Position Resolution of RD53A pixel modules

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RD53A pixel modules





- Pixel detector upgraded for HL-LHC upgrades of ATLAS and CMS
- Demonstrate the suitability of the chosen 65nm CMOS technology
- Only for testing, forms the basis for the production designs

Testbeam setup





• The testbeam determines how devices under test (DUT) respond to a single particle passing through the active area

- Sensors in reference telescopes have better position resolution than DUTs
- Use hits in telescopes to reconstruct the track, and extract position resolution of DUT from the difference between track position and hit position on DUT

Testbeam at SLAC





The Offline testbeam analysis

- EUTelescope v2.0.0 is a very generic and versatile collection of Marlin(Modular Analysis & Reconstruction for the Linear Collider) processors, dedicated to processing of testbeam data
- Track reconstruction algorithm: General Broken Lines(GBL) fitter (https://www.terascale.de/ wiki/generalbrokenlines/)





Effect from tilt angle





Cluster size



• The larger cluster size in tilted direction demonstrates the smaller equivalent pixel width

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Position resolution (Mimosa26)



- residual = hit position track position
- Biased residuals as hits on Mimosa26 are used in track fitting
- $\sigma_{biased}^2 = \sigma_{intrinsic}^2 \sigma_{tracking}^2$



Position resolution (RD53A)



- Unbiased residuals as hits on DUTs are excluded in track fitting $\sigma_{unbiased}^2 = \sigma_{intrinsic}^2 + \sigma_{tracking}^2$
- Fit function = Box(width) convolved with Gaussian (Mean,Sigma)
- $\sigma_{unbiased}^2(DUTs) = width^2/12 + sigma^2$



Tracking resolution on DUTs



- Track-resolution-simulator: https://github.com/simonspa/resolution-simulator/tree/master
- The same geometry of SLAC testbeam setup
- Need intrinsic resolution of Mimosa26 as input



Tracking resolution on DUTs



- The intrinsic resolution of Mimosa26 is measured as 3.85 μm



Results



• The position resolutions of non-tilted RD53A modules are both comparable with $pitch/\sqrt{12}$

pitch: the length or width of a pixel

- RD53A modules with 50×50 μ m² benefit more from 13° tilt angle
- Systematics include uncertainty of material estimation, beam energy, tracking resolution and Z position of DUTs

	RD53A 50μm×50μm non-tilted side(50μm)	RD53A 50μm×50μm tilted side(50μm)	RD53A 100μm×25μm non-tilted side(100μm)	RD53A 100μm×25μm tilted side(25μm)
$pitch/\sqrt{12}$	14.4	14.4	28.8	7.2
Non-tilted	14.51 ± 1.05	14.58 ± 1.04	28.16 ± 0.67	7.92 ± 1.73
13° tilted	14.04 ± 1.07	10.86 ± 1.09	28.54 ± 0.75	6.81 ± 1.82
<u>Non – tilted</u> 13° tilted	0.97 ± 0.10	0.74 ± 0.09	1.01 ± 0.04	0.86 ± 0.30



- The intrinsic position resolution of non-tilted and tilted RD53A modules with 50×50 μm^2 and 100×25 μm^2 pitch are measured using 11 GeV electron beam at SLAC
- The position resolution of 50×50 μm² RD53A reduces by 26% when tilted by 13°, and 14% for 100×25 μm² RD53A
- This information is useful for deciding on the geometry of the pixel layers, which are critical for flavor tagging and other tasks

Thank you!



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Correlation in x direction





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Correlation in y direction





Tilted DUTs





Tilt angle: 13° in XZ plane

After 3 iterations of GBL alignment

<ladder ID="51"

positionX="-1.197375218e+00" positionY="3.239637016e+00" positionZ="5.079932088e+02" rotationXY="-6.855291158e-01" rotationZX="1.257020850e+01" rotationZY="-1.890147900e-01" sizeX="2.000000000e+01" sizeY="9.600000000e+00" thickness="1.000000000e-01" radLength="5.5"

<ladder ID="50"

positionX="5.551223345e+00" positionY="3.128746483e+00" positionZ="5.329198459e+02" rotationXY="-7.342398936e-01" rotationZX="1.246104708e+01" rotationZY="-2.714878546e-01" sizeX="2.000000000e+01" sizeY="9.600000000e+00" thickness="1.000000000e-01" radLength="5.5" .

Cluster size





• Cluster size in tilted direction increase

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21

Position resolution (Mimosa26)







GBL residual at plane 2



Position resolution (Mimosa26)

x resid at plane 5 [µm]

tracks

tracks

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100

y resid at plane 5 [µm]

GBL residual at plane 4 GBL residual at plane 4 tracks 2200F 2000 2000 **Mean = 0.01** μm Mean = 0.00 μ m 1800 1800 **RMS** = 2.44 μ m **RMS** = 2.55 μ **m** 1600 1600 1400 1400 1200 1200 1000 1000 800 800 600 600 400 400 200 200 _100 _100 -80 -60 -40 -20 0 20 40 60 80 100 -80 -60 -20 0 20 40 60 80 -40 y resid at plane 4 [µm] x resid at plane 4 [µm] GBL residual at plane 5 GBL residual at plane 5 tracks 3500 3500 **Mean = 0.01** μm **Mean = 0.01** μm 3000 3000 **RMS** = 1.37 μ **m RMS** = 1.41 μ m 2500 2500 2000 2000 1500 1500 1000 1000 500 500 _100 -100 -20 20 20 -80 -60 -40 0 40 60 80 100 -80 -60 -40 -20 0 40 60 80 100

position resolution (RD53A)



fit function = Box(width) convolved with Gaussian (Mean,Sigma)



Chi^2/d.o.f. comparison





No tilt angle

13° tilt angle

Pull distribution





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Pull distribution

tracks

1007 tracks

0<u></u>5

 $_{-5}^{0L}$





Pull distribution



13345

0.8764

5

13345

0.8926

0.002194



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5

Tracking resolution based on simulator



- Scan intrinsic resolution of Mimosa26
- Simulate track resolution at each Mimosa26 plane
- Minimize the different between measured and calculated unbiased resolution



i are the six mimosa26

Intrinsic resolution

• With pitch of pixels increasing, the residuals should be fit by convolution of box and gaussian function



$$R_{unbiased} = R_{track} + R_{intrinsic}$$

$$\sigma^{2}(unbiased) = \sigma^{2}(track) + \sigma^{2}(intrinsic)$$

$$R_{unbiased} \approx Box(width) + Gauss(sigma)$$

$$\sigma^{2}(unbiased) = width^{2}/12 + sigma^{2}$$

$$(intrinsic) = width^{2}/12 + sigma^{2} - \sigma^{2}(track)$$



Systematics



	RD53A 50μm×50μm non-tilted side(50μm)	RD53A 50μm×50μm tilted side(50μm)	RD53A 100μm×25μm non-tilted side(100μm)	RD53A 100μm×25μm tilted side(25μm)
10% variation of material on DUTs	0.13	0.12	0.03	0.07
track resolution	0.15	0.15	0.3	0.3
10% variation of beam energy	0.02	0.11	0.03	0.07
Z position+/-1 0 mm	1.03	1.02	0.6	1.7
total	1.05	1.04	0.67	1.73