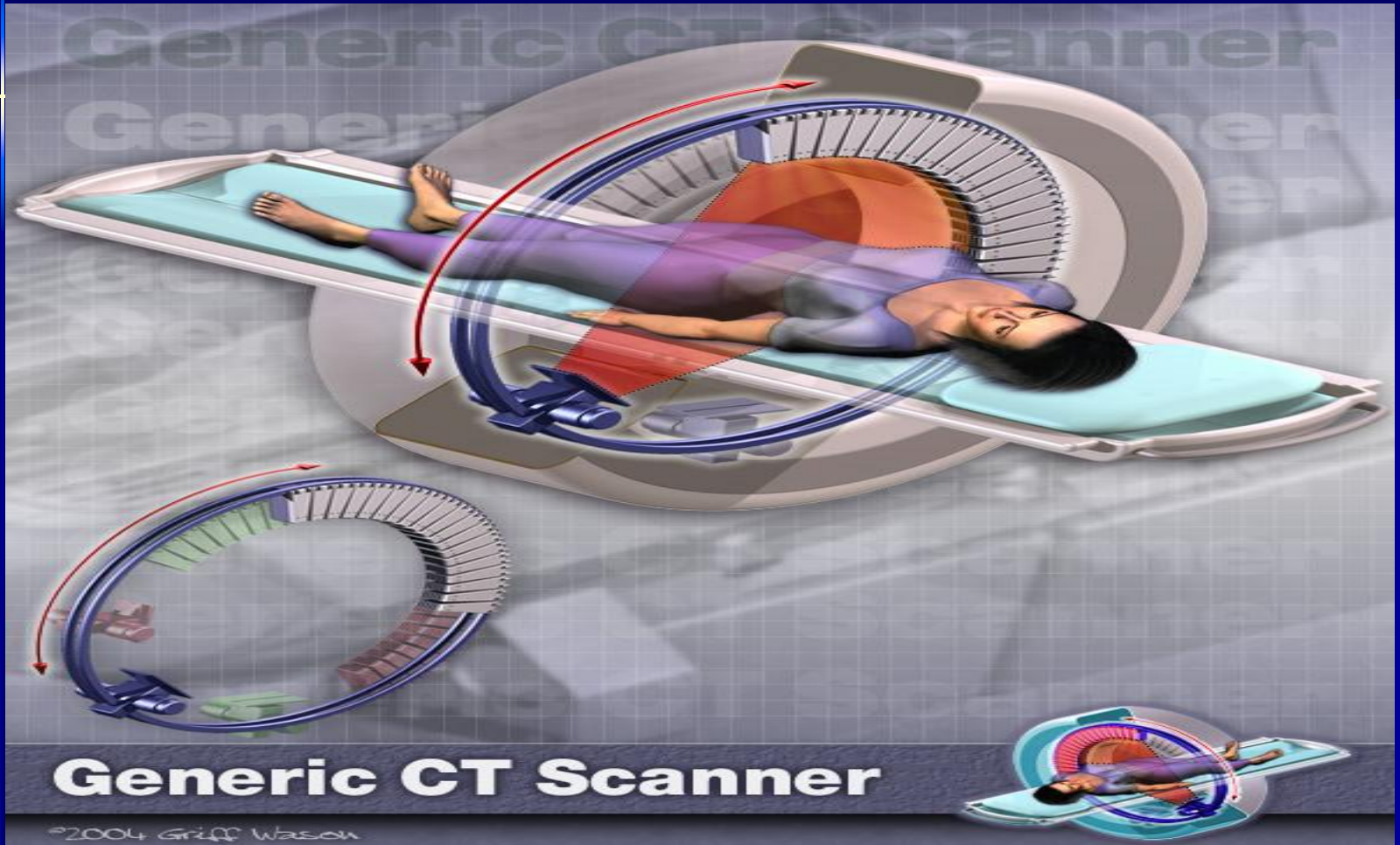


بِسْمِ اللَّهِ الرَّحْمَنِ الرَّحِيمِ

PHYSICAL PRINCIPLES OF COMPUTED TOMOGRAPHY



Presentation: **Mohamad Akbarnejad**
Radiobiology and Radiation Protection MSC

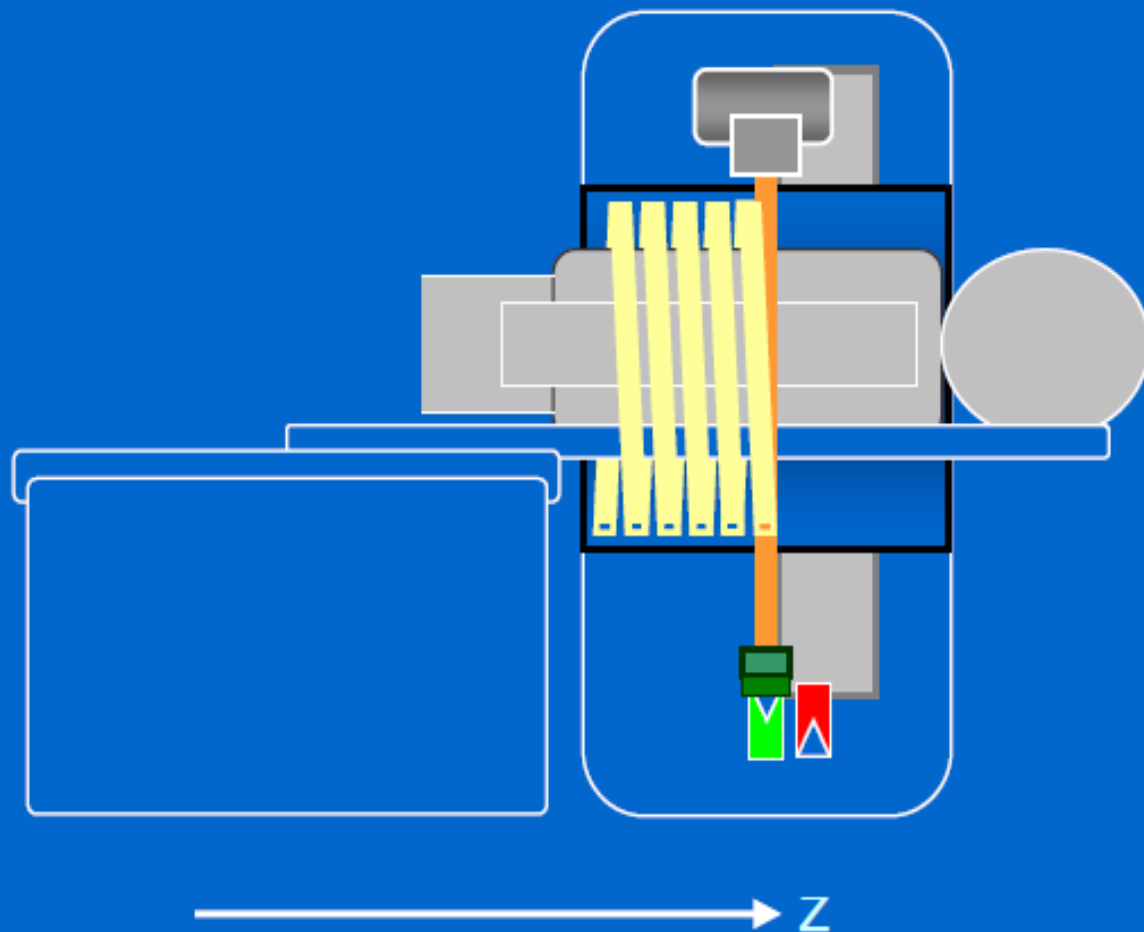
Pitch

The mean of **Pitch** in CT?



Helical (spiral) scanning - pitch

$$\text{Pitch} = \frac{\text{table travel / rotation}}{\text{X-ray beam width}}$$



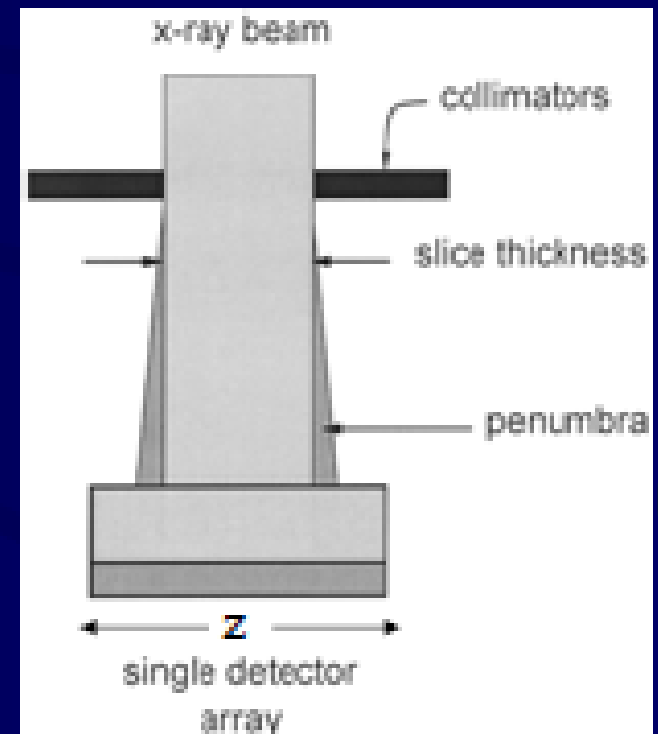
Pitch for Single-Slice CT

- **Image and beam width are same for conventional CT**
- **Pitch = table travel \div beam width**
- **Typical pitch values are 0.7 to 1.5**

Conventional Helical CT Detectors

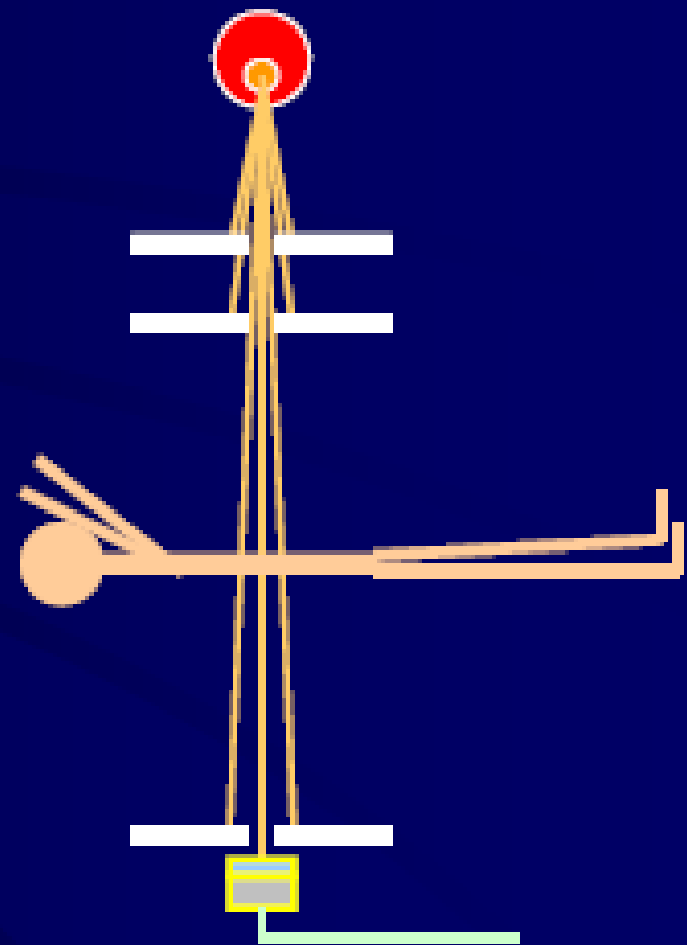
Image width determined by
beam thickness

Pitch = table mm / beam mm

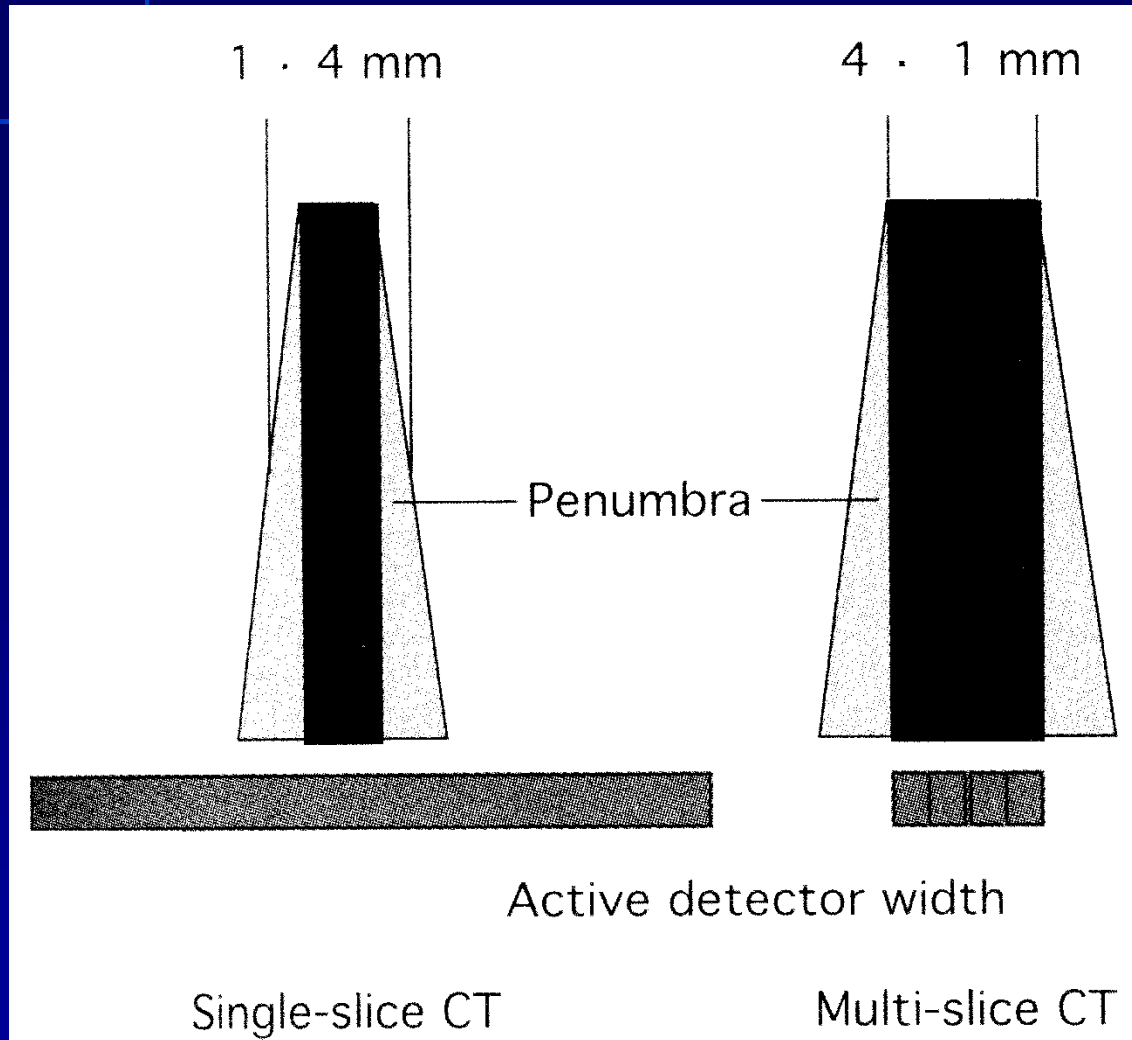


Beam Collimation

- Pre-patient collimators define width of beam in z (all systems)
- “Detector” collimators reduce scatter at detectors (some CTs)

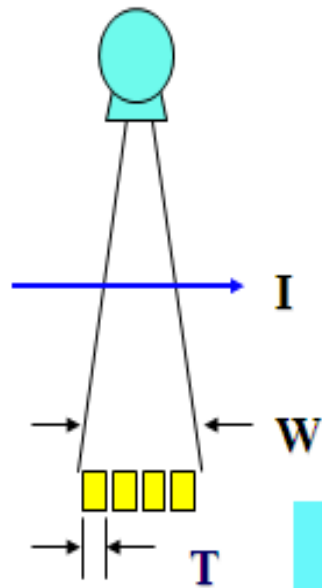


Multi-slice CT



*‘Over beaming’
caused by
wider collimator
settings to avoid
penumbral effects;*

Pitch redefined for MDCT



$$\text{Beam Pitch} = \frac{I}{W}$$

$$\text{Detector Pitch} = \frac{I}{T}$$

$$\text{Beam Pitch} = \frac{\text{Detector Pitch}}{N} = \frac{I}{N * T} = \text{Pitch}$$

I - Table feed (mm/rotation)

W - Beam width (mm)

T - Single DAS channel width (mm)

N - Number of active DAS channels

Pitch factor

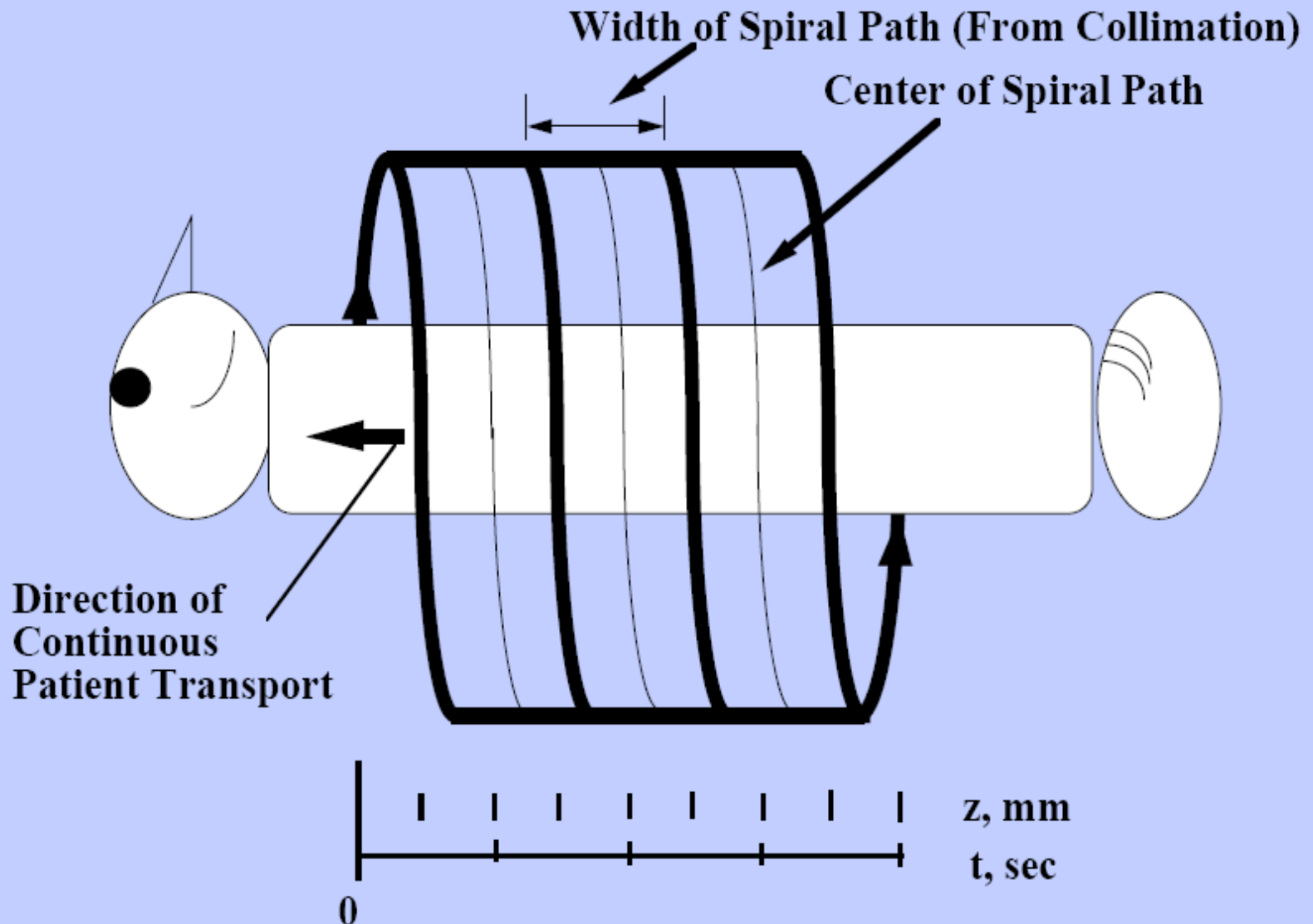
- Inter-slice distance is defined as the couch increment minus nominal slice thickness. In helical CT the pitch factor is the ratio of the couch increment per rotation to the nominal slice thickness at the axis of rotation. In clinical practice the inter-slice distance generally lies in the range between 0 and 10mm, and the pitch factor between 1 and 2.
- The inter-slice distance can be negative for overlapping scans which in helical CT means a pitch < 1 .

Data Acquisition

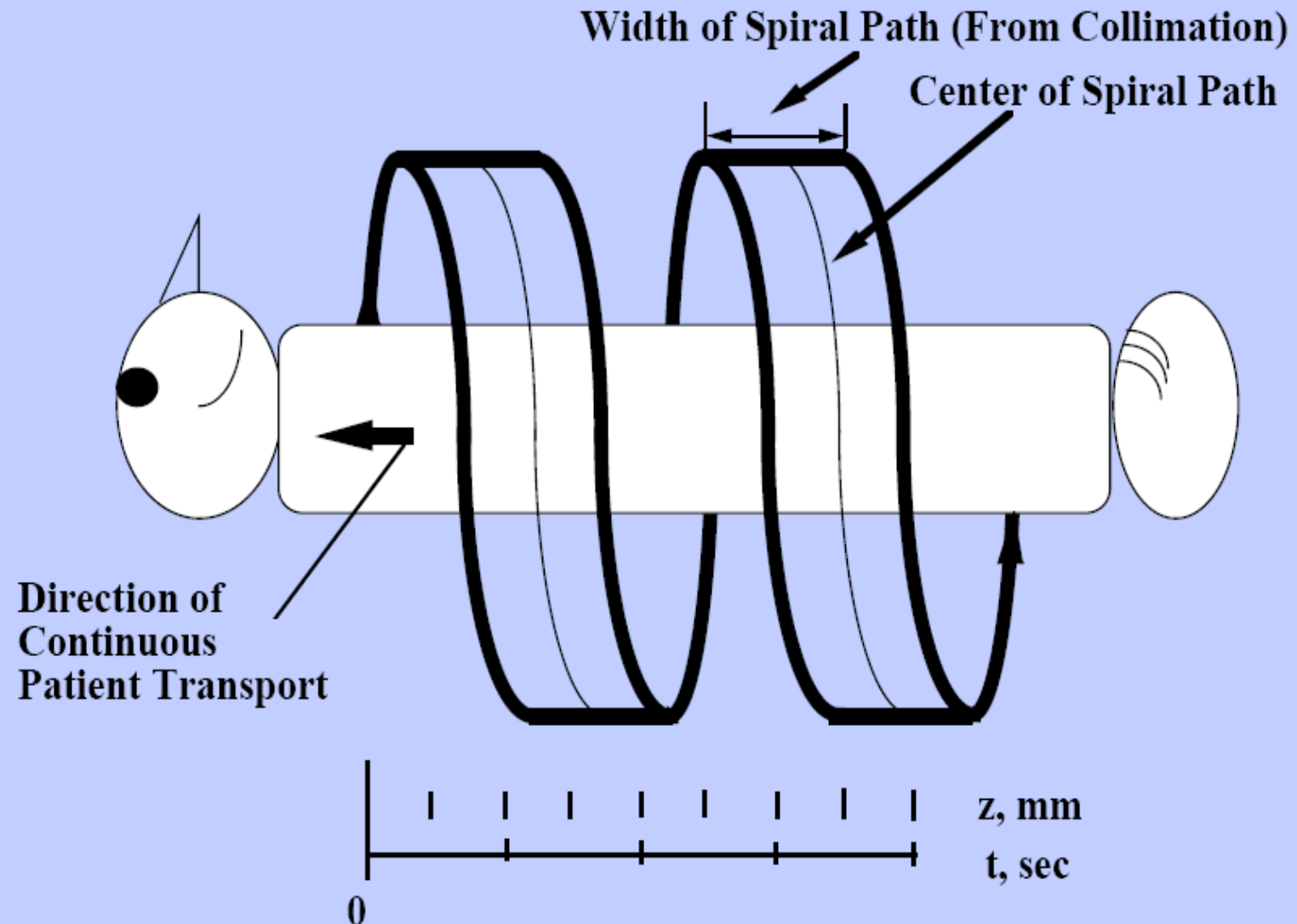
$$Pitch = \frac{\text{Table Movement}}{\text{Collimation}}$$

- Continuous Spiral – Pitch = 1 (10mm/10mm)
- Extended Spiral – Pitch = 2 (20mm / 10mm)
- Overlapping Spiral – Pitch = 1/2 (5mm/10mm)

Pitch=1; Contiguous Spiral



Pitch=2; Extended Spiral

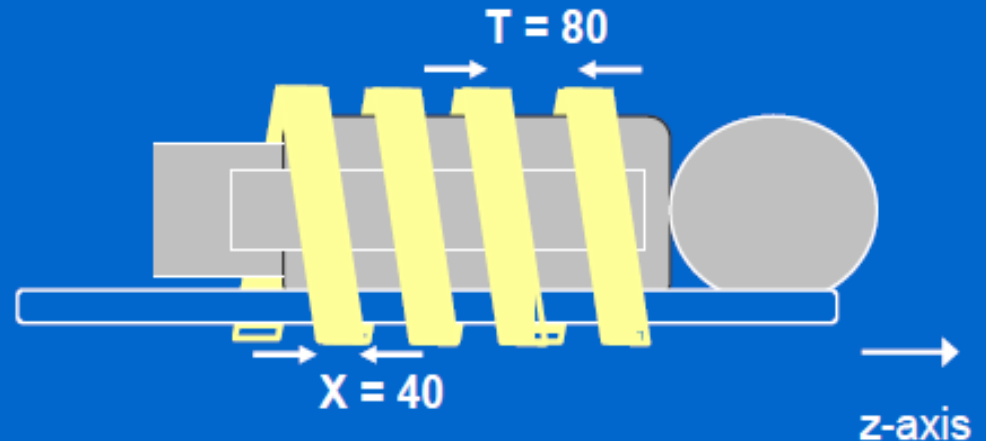


Helical (spiral) scanning - pitch

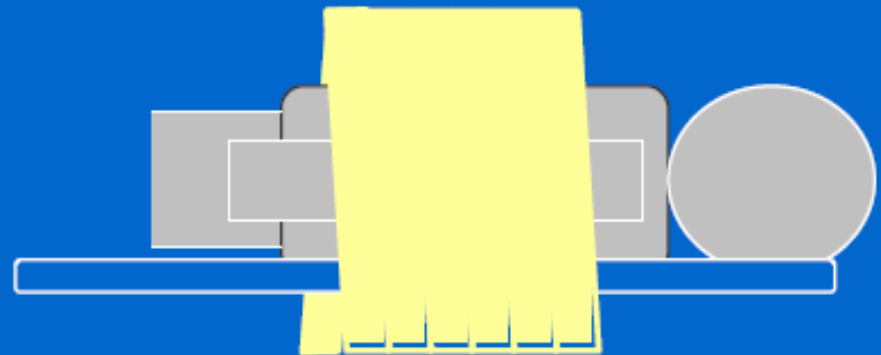
pitch 2

Table travel/rot = 80 mm

Beam width = 40 mm



pitch 1



pitch 0.5



Pitch

Pitch=1
Table Travel = Paint Width
Uniform Paint

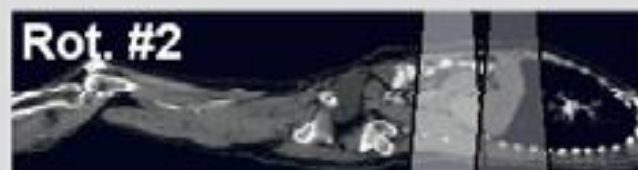
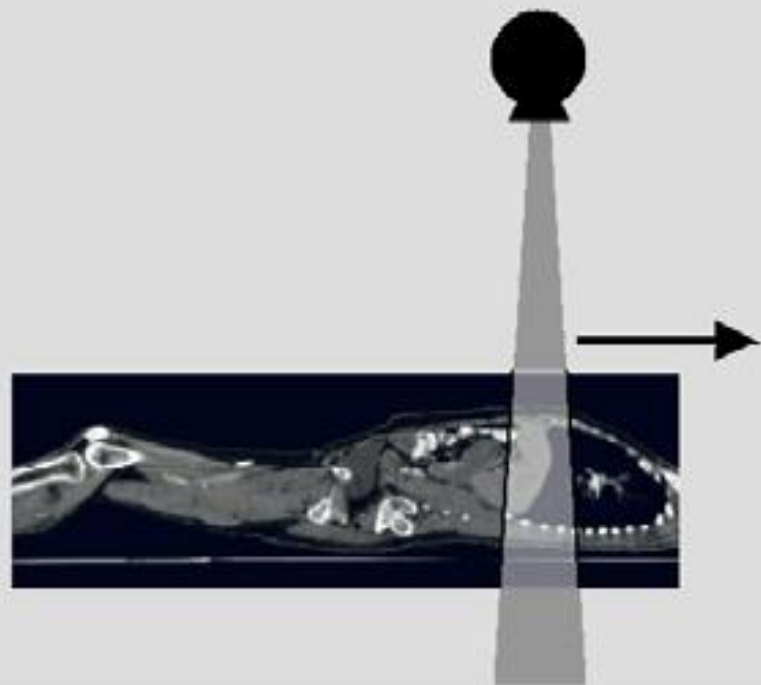


Pitch<1
Table Travel < Paint Width
Uniform + Overlapped Paint

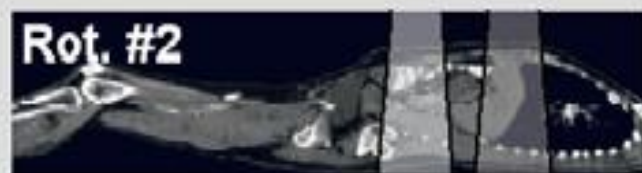


Pitch>1
Table Travel > Paint Width
Candy Cane Stripes

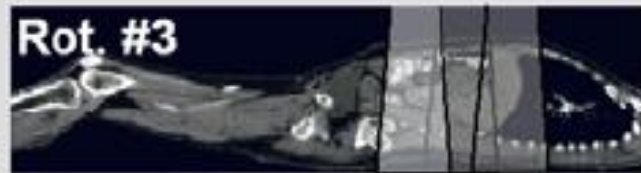
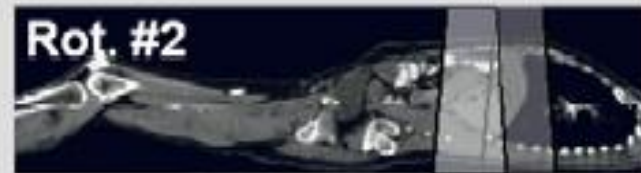




Pitch 1.0



Pitch 1.5



Pitch 0.75

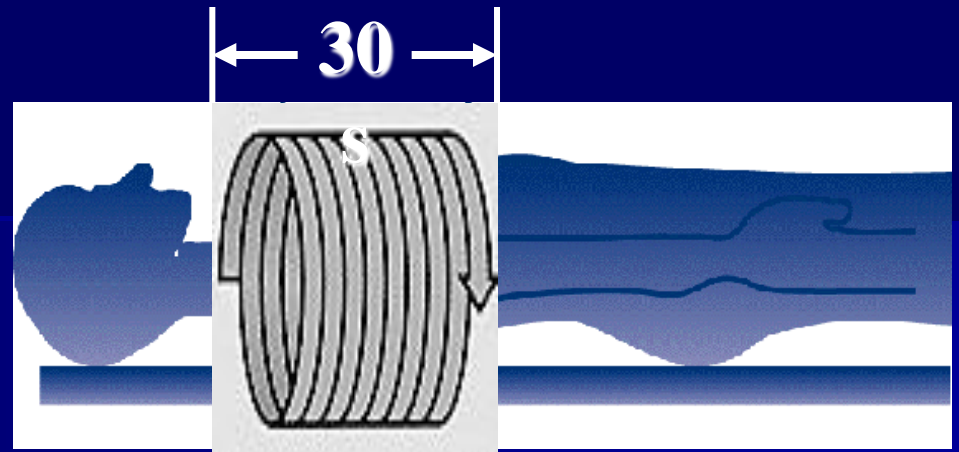
Table Speed & Pitch

Table Speed is defined as distance traveled
in mm
per 360° rotation

$$\text{Pitch} \Rightarrow \frac{\text{Table Feed per rotation}}{\text{Collimation}}$$

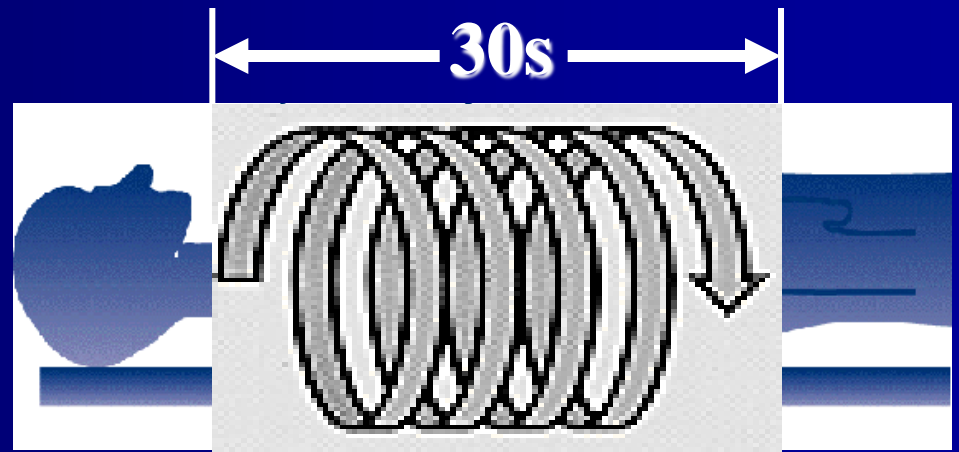
<u>Table Feed</u>	<u>Collimation</u>	<u>Pitch</u>
10 mm/rot	10 mm	1.0
15 mm/rot	10 mm	1.5
20 mm/rot	10 mm	2.0

**Pitch 2 covers
2x
distance as Pitch 1**



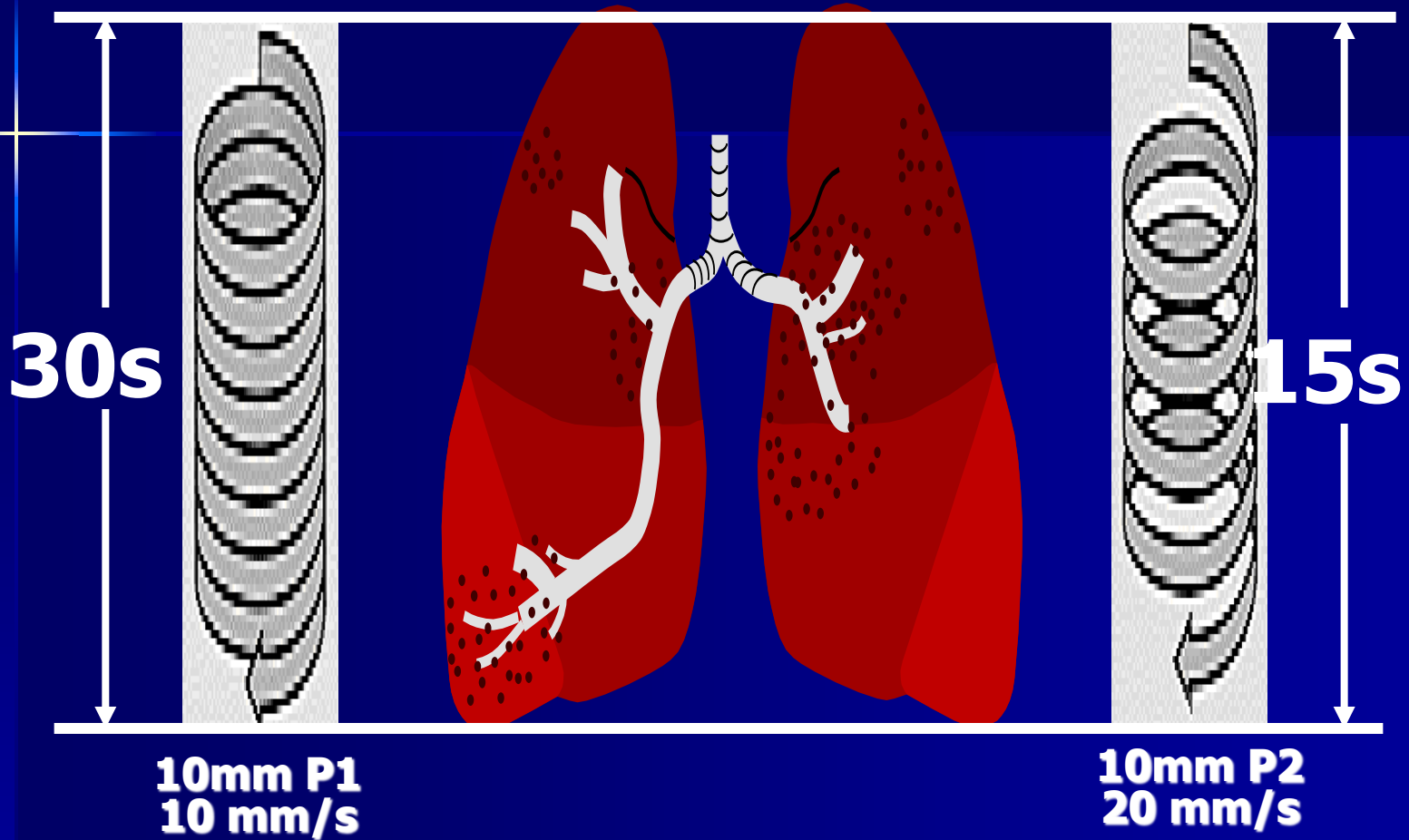
10mm P1

**More Coverage in
the same time with
extended Pitch!!**

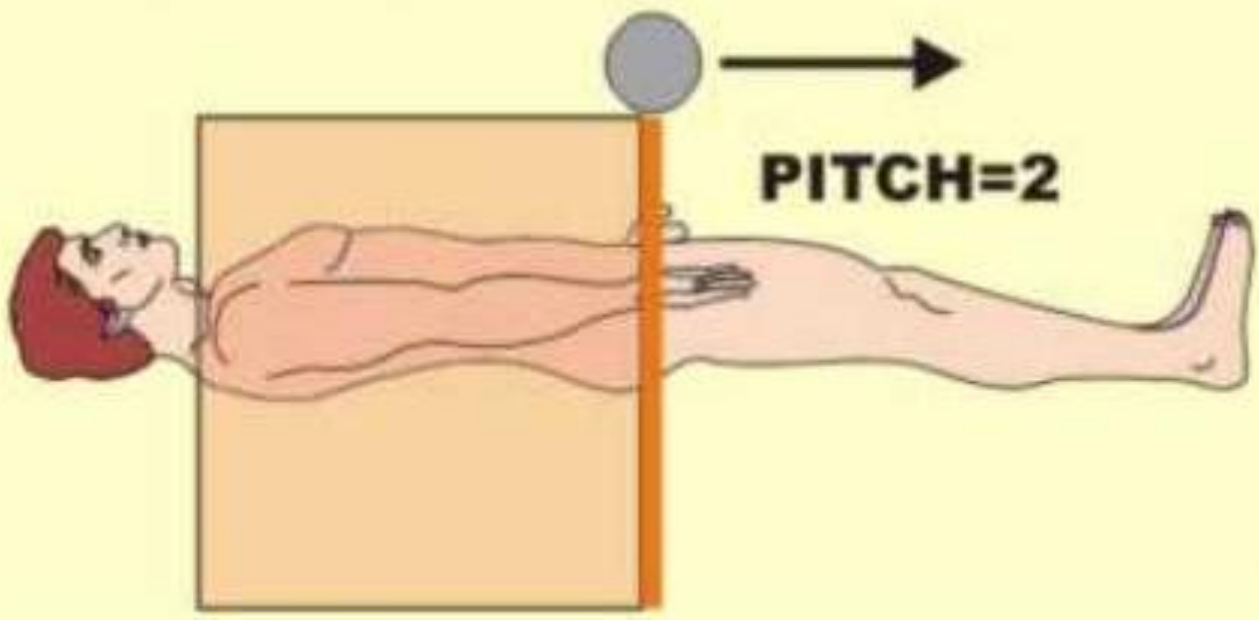
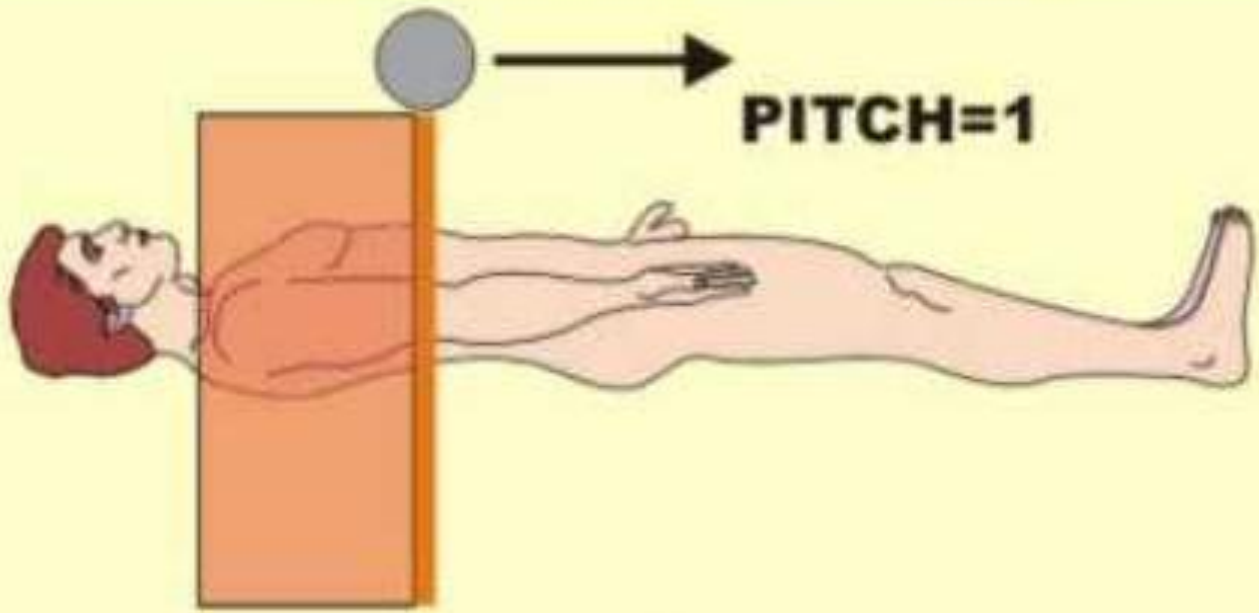


10mm P2

Scan Range = 300mm



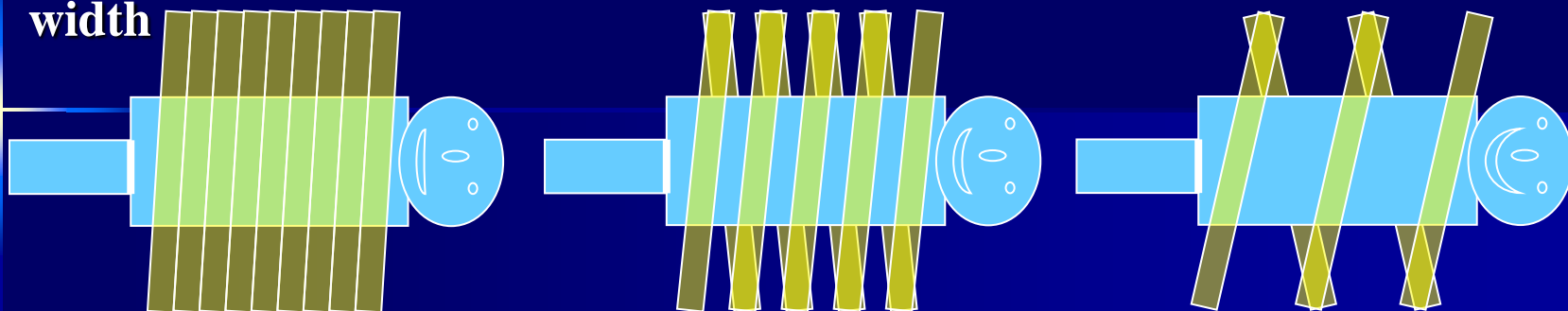
Cover the same volume in shorter time with extended Pitch



Sprawls

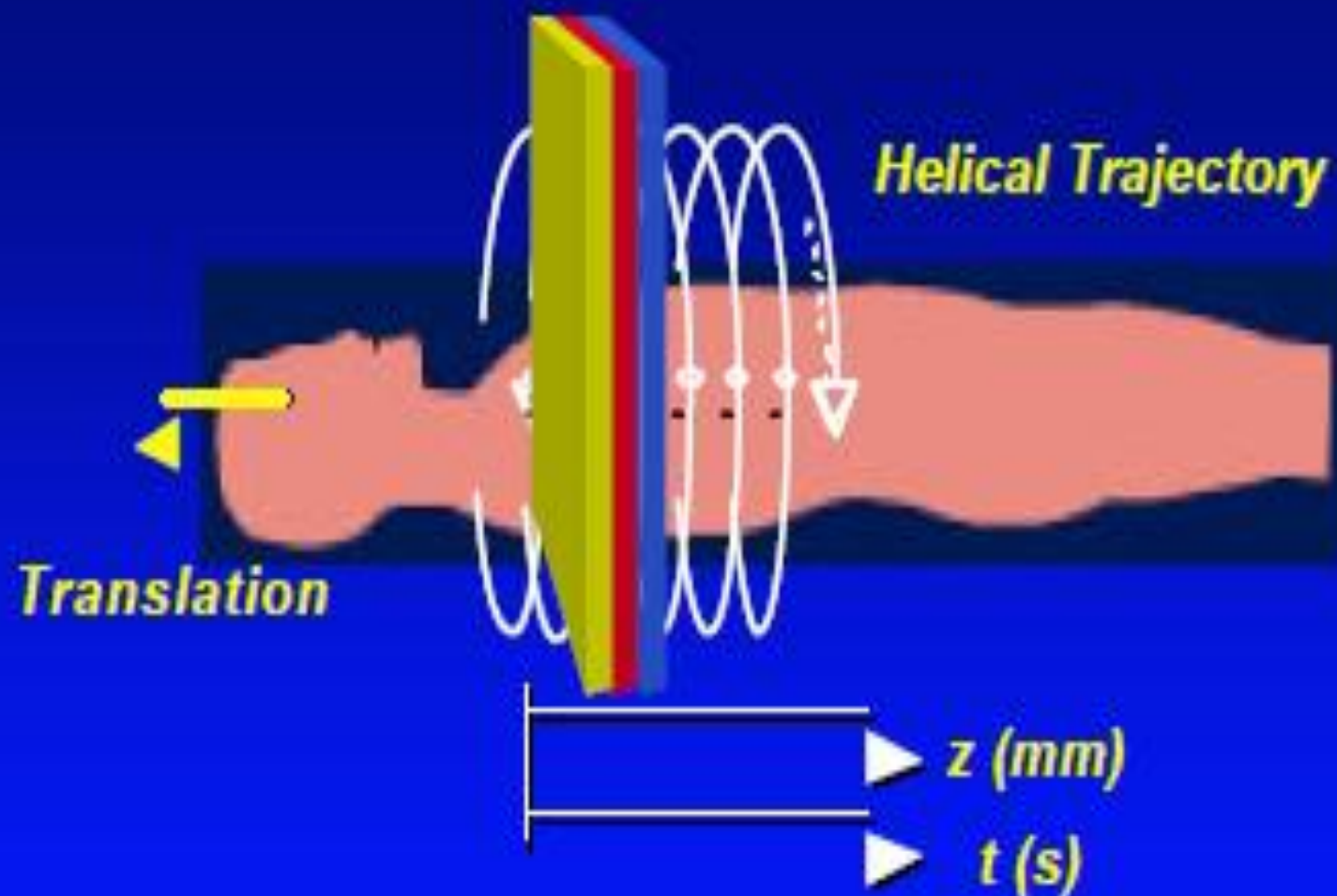
Pitch

ratio of the distance the table travels per rotation to the x-ray beam width



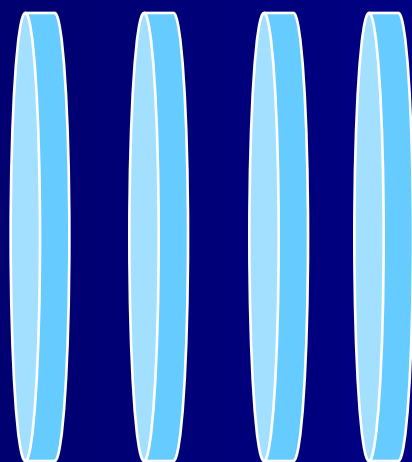
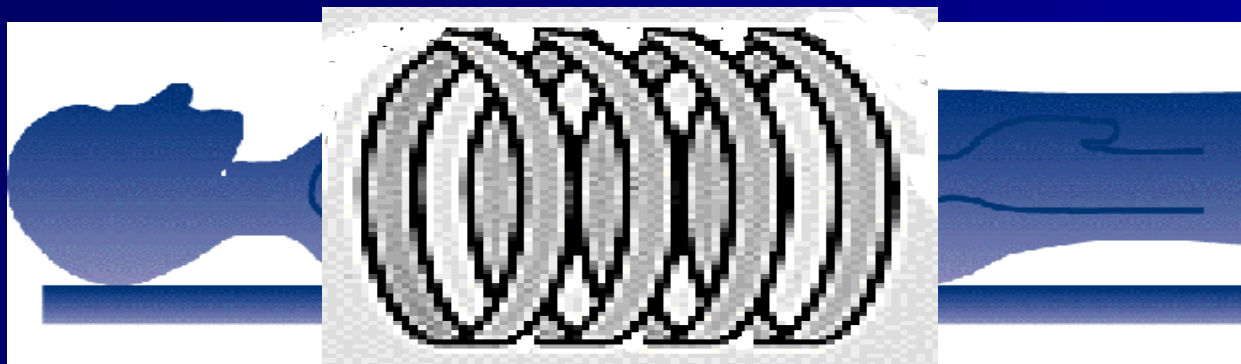
Number rotations	10		5		2.5
Slice thickness	10	10	10	10	10
Table movement per rotation	10	15	20	30	40
Pitch	1	1.5	2	3	4
Dose	10	7.5	5	3.33	2.5

Helical Single-Section Mode



Interpolation using samples from single row detector ring

Conventional

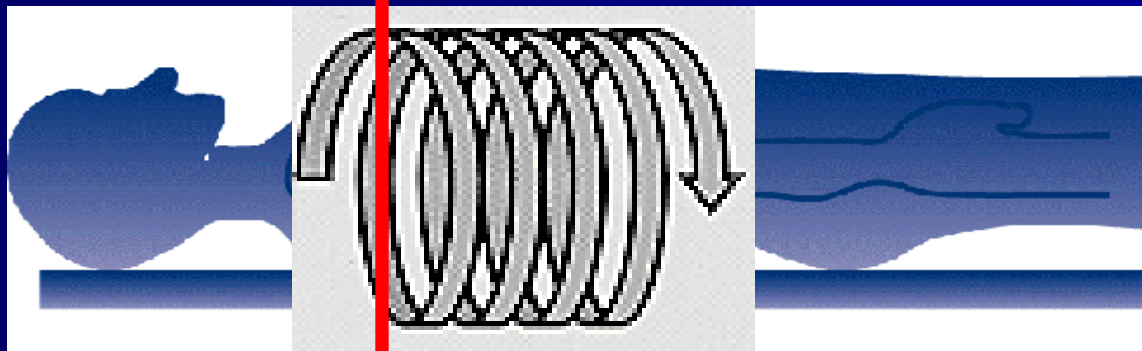


SSP



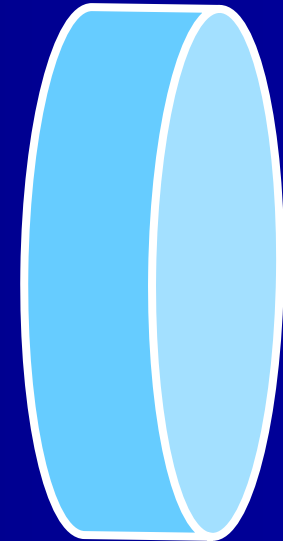
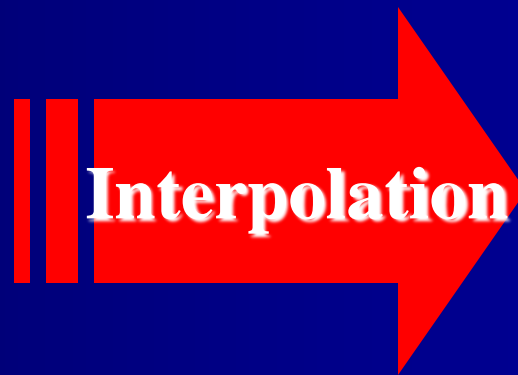
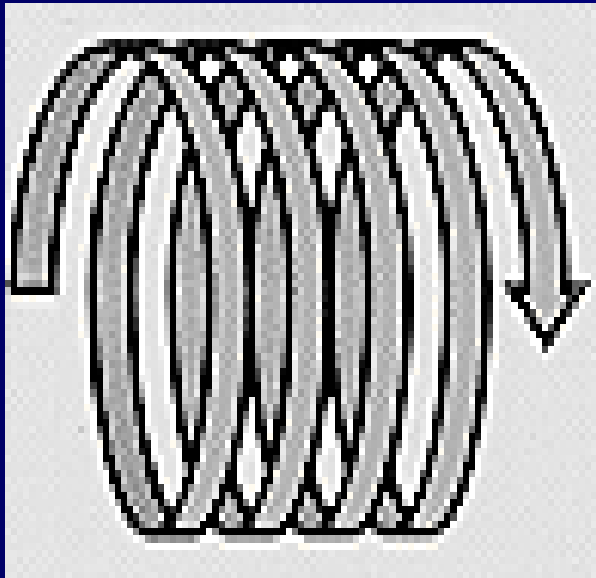
conventional

MSCT



Interpolation Algorithm

Converts volume data into slice 
images



To reduce artifacts due to table motion during spiral scanning, we use a special reconstruction process called *INTERPOLATION*

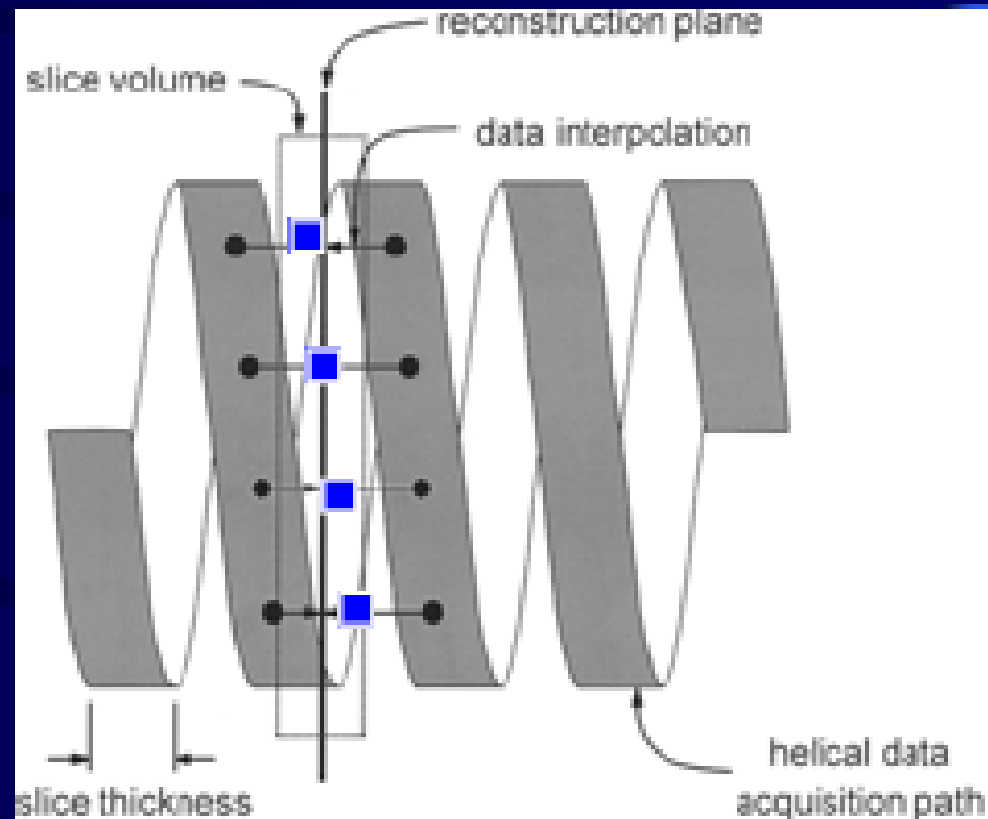
Helical Interpolation

Collect data (black dots)

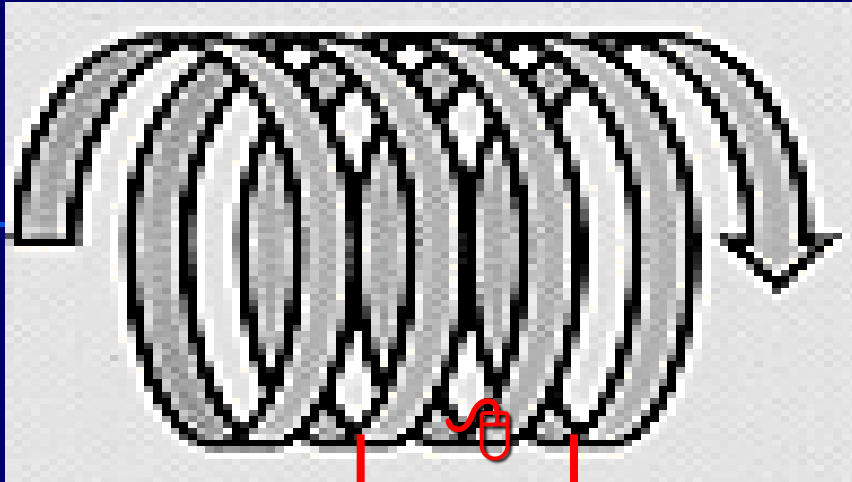
Rebin to estimate the 180° data (blue squares)

Interpolate to estimate image between collected and rebinned data

Helical CT needs fast computers

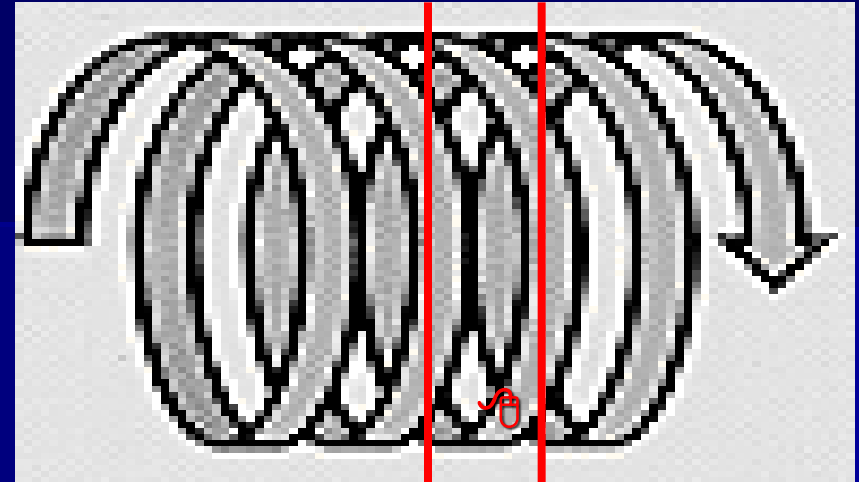


Wide Algorithm



$2 \times 360^\circ$
 $= 720^\circ$
raw data

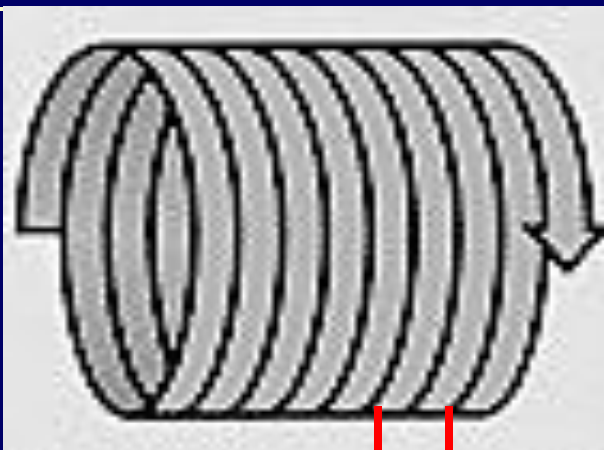
Slim Algorithm



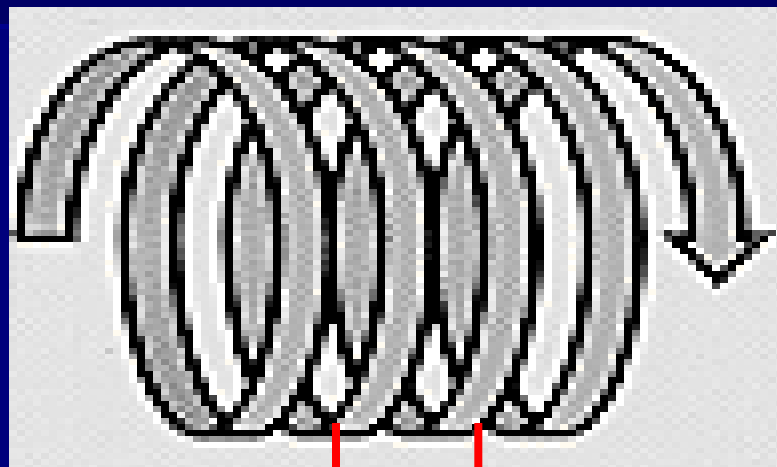
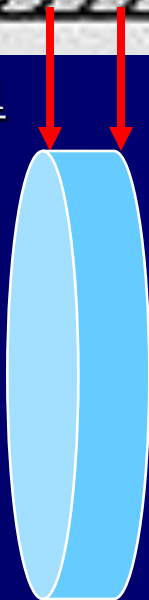
$2 (180+52)$
 $= 464^\circ$
raw data

Wide algorithm produces a broader image thickness
Wide algorithm uses more raw data => less image noise

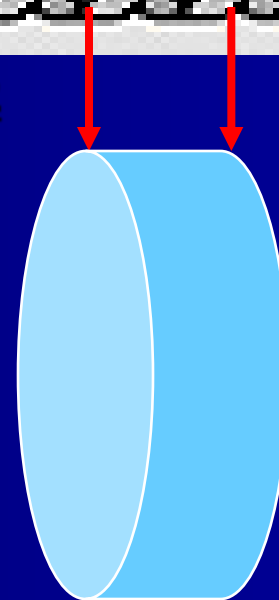
Pitch 2 scanning produces a broader image thickness
Pitch 2 scanning does not increase image noise



PITCH 1



PITCH 2



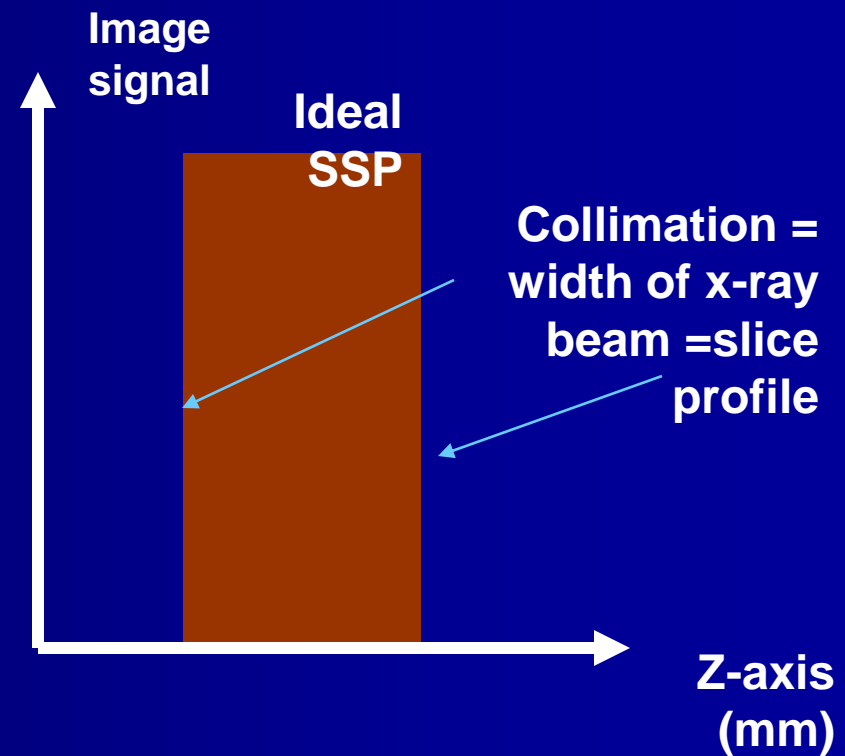
***30% increase in
image thickness
with Pitch 2***

Slice Sensitivity Profile (SSP)

SSP describes the **effective slice thickness** of an image and to what extent anatomy within that slice contribute to the signal



All points within the slice contribute equally & points outside of the slice do not contribute to the image at all .



Slice Profile (SP)

- Effective slice thickness of an image



Resolution



Slice Profile

Factors influencing Slice Profile

- Collimation
- Pitch
- Interpolation algorithm (360° or 180°)

Factors influencing SSP

- Collimator width



Spiral CT

- Table speed or Pitch

- Interpolation Algorithm

=> mathematical process required to reconstruct axial images from the spiral volume data set

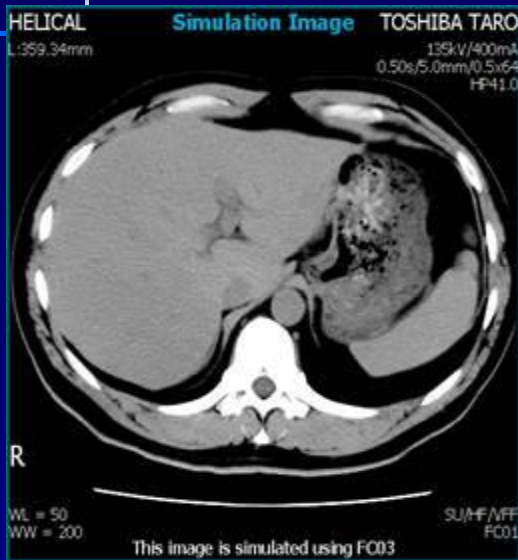


**Smoother
image**

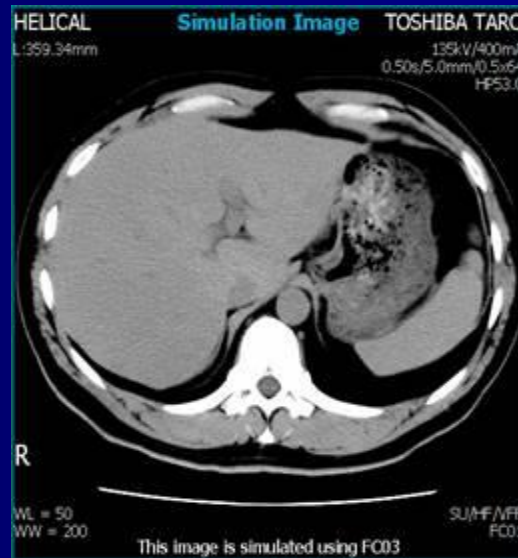


**Noisier
image**

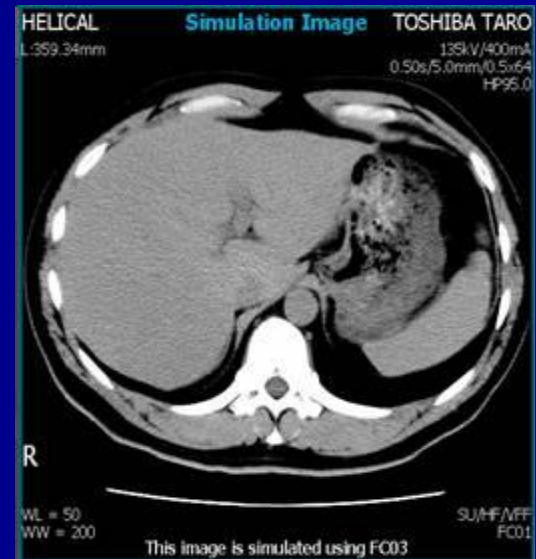
Effect of Pitch on Dose and Image Quality



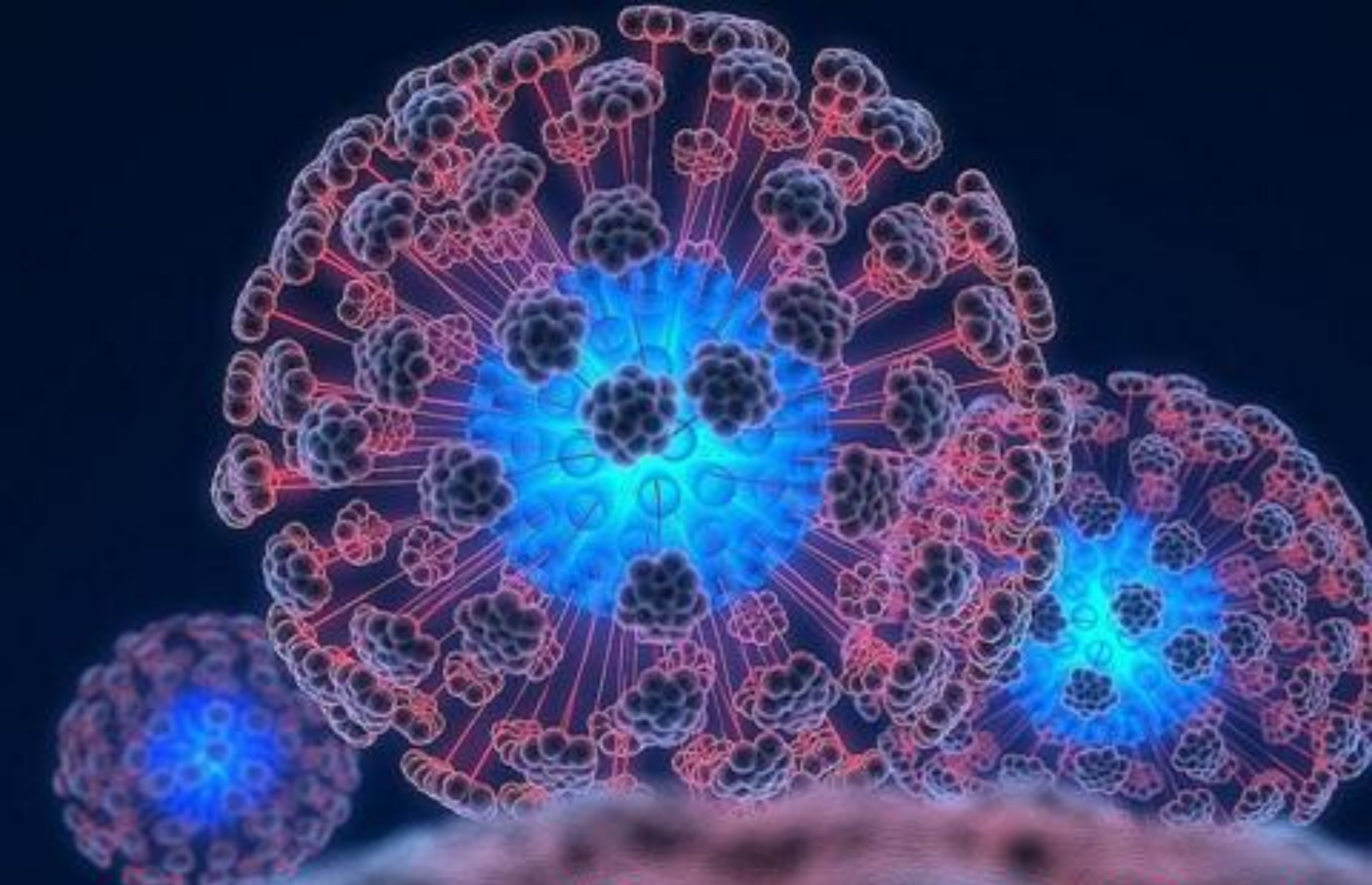
P = 0.64
CTDI = 47.8 mGy
30% higher



P = 0.83
CTDI = 37 mGy



P = 1.48
CTDI = 20.6 mGy
45% lower



Thank you for your attention!