

Clinical Radiology

Can CT measures of tumour heterogeneity stratify risk for nodal metastasis in patients with non-small cell lung cancer?

--Manuscript Draft--

Manuscript Number:	CRAD-D-17-00149
Full Title:	Can CT measures of tumour heterogeneity stratify risk for nodal metastasis in patients with non-small cell lung cancer?
Article Type:	Original Paper
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Abstract:	<p>OBJECTIVES: To undertake a preliminary assessment of the potential for CT measures of tumour heterogeneity to stratify risk of nodal metastasis in patients with non-small cell lung cancer (NSCLC).</p> <p>METHODS: Tumour heterogeneity in CT images from PET/CT examinations in 150 consecutive patients with NSCLC was assessed using CT texture analysis. The short axis diameter of the largest mediastinal node was also measured. 42 patients without distant metastases subsequently had tumour nodal status confirmed by surgery (n=26) or Endobronchial Ultrasound (EBUS); n=16). CTTA parameters and largest nodal diameter were related to nodal status using the rank-correlation and the risk-ratio for each nodal stage (>N0, >N1, >N2) was compared between patients categorised as high and low risk by CTTA or nodal size. The most significant predictor of nodal status was related to overall survival using Kaplan-Meier analysis.</p> <p>RESULTS: N-stage was more significantly correlated with CTTA than nodal diameter (Rs = -0.39, p = 0.011, Rs = -0.45, p=0.0025, Rs = -0.40, p=0.0091 for normalised SD, normalised E and kurtosis respectively; Rs = -0.39, p = 0.042 for nodal diameter). The presence of 2 or more high-risk CTTA values was the greatest risk-factor for mediastinal metastasis (Risk-ratio: 11.0, 95% confidence interval 1.56 - 77.8, p=0.0014) and was associated with significantly poorer overall survival (p=0.016).</p> <p>CONCLUSION: CTTA in NSCLC is related to nodal status in patients without distant metastases and has the potential to inform selection of investigative strategies for the assessment of mediastinal malignancy.</p>

CAN CT MEASURES OF TUMOUR HETEROGENEITY STRATIFY RISK FOR NODAL METASTASIS IN PATIENTS WITH NON-SMALL CELL LUNG CANCER?

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DECLARATION OF INTEREST

Kenneth Miles declares he has a financial interest in Feedback PLC who supply the texture analysis software used in this study.

FUNDING:

This research did not receive any specific grant from funding agencies in the public, commercial or not-for profit sectors.

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AUTHOR CONTRIBUTIONS

1. Guarantor of integrity of entire study – Michelle Craigie
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4. Clinical studies – Michelle Craigie, Julia Squires, Kenneth Miles
5. Experimental studies/data analysis – Michelle Craigie, Julia Squires, Kenneth Miles
6. Statistical analysis – Kenneth Miles
7. Manuscript preparation – Michelle Craigie, Kenneth Miles
8. Manuscript editing – Michelle Craigie, Julia Squires, Kenneth Miles

ABSTRACT

OBJECTIVES: To undertake a preliminary assessment of the potential for CT measures of tumour heterogeneity to stratify risk of nodal metastasis in patients with non-small cell lung cancer (NSCLC).

METHODS: Tumour heterogeneity in CT images from PET/CT examinations in 150 consecutive patients with NSCLC was assessed using CT texture analysis. The short axis diameter of the largest mediastinal node was also measured. 42 patients without distant metastases subsequently had tumour nodal status confirmed by surgery (n=26) or Endobronchial Ultrasound (EBUS); n=16). CTTA parameters and largest nodal diameter were related to nodal status using the rank-correlation and the risk-ratio for each nodal stage (>N0, >N1, >N2) was compared between patients categorised as high and low risk by CTTA or nodal size. The most significant predictor of nodal status was related to overall survival using Kaplan-Meier analysis.

RESULTS: N-stage was more significantly correlated with CTTA than nodal diameter ($R_s = -0.39, p = 0.011, R_s = -0.45, p=0.0025, R_s = -0.40, p=0.0091$ for normalised SD, normalised E and kurtosis respectively; $R_s = -0.39, p = 0.042$ for nodal diameter). The presence of 2 or more high-risk CTTA values was the greatest risk-factor for mediastinal metastasis (Risk-ratio: 11.0, 95% confidence interval 1.56 - 77.8, $p=0.0014$) and was associated with significantly poorer overall survival ($p=0.016$).

CONCLUSION: CTTA in NSCLC is related to nodal status in patients without distant metastases and has the potential to inform selection of investigative strategies for the assessment of mediastinal malignancy.

KEY WORDS

Non-small cell Lung Cancer, Positron-emission tomography, Computed Tomography
Texture Analysis, Nodal metastasis

1 **CAN CT MEASURES OF TUMOUR HETEROGENEITY STRATIFY RISK FOR NODAL**
2 **METASTASIS IN PATIENTS WITH NON-SMALL CELL LUNG CANCER?**

3

4 **INTRODUCTION:**

5 Non-small cell lung cancer (NSCLC) remains one of the leading causes of cancer in
6 the Western World with a poor prognosis. Predictive factors of disease burden are
7 required to help with decisions regarding options for clinical management. These
8 circumstances are illustrated by the current clinical guidance for the management of
9 NSCLC issued by the National Institute of Clinical Health and Excellence (NICE)(1).
10 This guidance recommends different investigative strategies for the assessment of
11 mediastinal disease according to probability of mediastinal malignancy based on
12 nodal size as depicted by CT. For patients with a low probability of mediastinal
13 malignancy (15%; lymph nodes < 10 mm maximum short axis on CT), the optimum
14 strategy was determined to be staging with PET-CT alone. PET-CT, or endobronchial
15 ultrasound (EBUS)-guided transbronchial needle aspiration (TBNA), or endoscopic
16 ultrasound (EUS)-guided fine needle aspiration, or non-ultrasound guided TBNA are
17 recommended for patients with an intermediate probability of mediastinal
18 malignancy (50%; lymph nodes between 10 and 20 mm maximum short axis on CT)
19 whilst neck ultrasound with sampling of visible lymph nodes, or non-ultrasound
20 guided TBNA should be offered to patients with a high probability of mediastinal
21 malignancy (85%; lymph nodes > 20 mm maximum short axis on CT). New methods
22 that can improve the risk stratification for mediastinal disease based on CT therefore

23 have the potential to improve the selection of staging procedures for patients with
24 NSCLC.

25 CT texture analysis (CTTA) is emerging as a technique for derivation of prognostic
26 biomarkers for patients with NSCLC and other tumours (2-5). CT texture analysis
27 evaluates quantitatively the distribution of CT attenuation values within a tumour to
28 determine its heterogeneity. Tumour heterogeneity has been shown to relate to
29 tumour aggression and treatment response, with hypoxia, mutations in EGFR and
30 KRAS genes, and ALK gene re-arrangements having been identified as potential
31 biological correlates for CTTA values in NSCLC (2). Given the prognostic significance
32 of CTTA and its associated biological characteristics, we hypothesize that CTTA can
33 stratify risk for nodal metastatic disease in patients with NSCLC.

34

35 **METHOD:**

36 *Study Design:*

37 A prospective observational study design was adopted, using patient data that was
38 acquired as part of routine clinical care. Our local institutional review board had
39 waived the requirement for individual consent.

40 *Patients:*

41 Imaging data was collected from 150 consecutive patients undergoing PET/CT for
42 staging of NSCLC cancer. The study cohort of 42 patients comprised all those with no
43 distant metastasis detected on PET/CT and subsequent confirmation of nodal status
44 either at surgery (n=26) and or by EBUS with TBNA (n=16). Overall survival was

45 determined from a median clinical follow-up period of 279 days (range: 59-437
46 days).

47 *CT Imaging protocol and data analysis:*

48 Images were acquired using a Siemens mCT PET/CT system 120kV, automated tube
49 modulation (Care dose) with reference tube current set at 80mAs, 5mm slices and
50 collimation 1.2mm(Siemens, Erlangen, Germany). Using TexRAD software
51 (Feedback plc, Cambridge, UK), CTTA was performed on the low dose CT slice that
52 displayed the largest cross sectional area of the tumour on soft tissue windows as
53 described previously (2). Definition of the tumour boundary was assisted with
54 reference to the PET fused images and narrow CT windows (level 40HU, width
55 150HU). Automated segmentation tools were used to optimise consistency in the
56 analysis between operators where possible, for example where the tumour was
57 surrounded by aerated lung. Where the lung tumour was in contact with other
58 tissues such as the chest wall, mediastinum or consolidated lung, manual selection of
59 the region of interest (ROI) was required along that border and the automated
60 segmentation tool could be used on those areas where the tumour bordered
61 aerated lung. Segmentation tools excluded areas of tumour cavitation seen on CT
62 but were not used to exclude areas of necrosis/photopaenia seen on the PET fused
63 images.

64 Based on the filtration-histogram CTTA approach utilised by the CTTA software,
65 tumour heterogeneity at a scale of 4mm was expressed as kurtosis, standard
66 deviation (SD) and entropy (E). SD and E were both log-normalised to the tumour
67 area determined by the number of pixels in the tumour ROI.

68 The short-axis diameter of the largest mediastinal was also measured by a separate
69 operator who was an accredited radiologist with more than 25 years of CT
70 experience.

71 *Statistics*

72 The relationship between CTTA parameters and nodal status were determined using
73 the rank-correlation and compared to the correlation found between N-stage and
74 mediastinal nodal size. If a significant correlation was found, patients were
75 categorised as high or low risk for nodal metastases using the median texture value
76 for the study cohort. The risk for each nodal stage (>N0, >N1, >N2) was compared
77 between high and low risk patients and expressed as the risk ratio (with 95%
78 confidence limits), with comparison against the risk-ratios found for nodal size
79 <10mm versus \geq 10mm, using Fisher's exact test to assess statistical significance. The
80 most significant predictor of nodal status was related to overall survival using
81 Kaplan-Meier analysis.

82

83 **RESULTS:**

84 42 patients NSCLC without distant metastases went on to have nodal status
85 confirmed either at surgery or EBUS with TBNA, the numbers of patients with N-
86 stages 0, 1, 2 and 3 were 26, 4, 8 and 4 respectively. 5 (11.9%) patients died during
87 follow-up.

88 N-stage was shown to correlate significantly with normalised SD, normalised E and
89 kurtosis ($R_s = -0.39$, $p = 0.011$, $R_s = -0.45$, $p=0.0025$, $R_s = 0.40$, $p=0.0091$)

90 respectively). There was a weaker but statistically significant correlation between N-
91 stage and mediastinal nodal diameter ($R_s = -0.39$, $p = 0.042$). CTTA values were
92 categorised as high-risk for nodal metastases if below the median for normalised SD
93 (5.66) and normalised E (0.777), or above the median for kurtosis (-0.295). 21 (50%)
94 patients had 2 or more high-risk CTTA values. Example CT texture results from
95 patients with high and low risk CTTA parameters are given in Figures 1 and 2.

96 The greatest risk factors for node positive disease were log-normalised entropy and
97 the presence of 2 or more high-risk CTTA values, each associated with a 4.3 greater
98 likelihood of nodal metastases ($p=0.0036$, table 1). A mediastinal nodal diameter of \geq
99 10mm was a significant risk-factor for mediastinal malignancy (i.e. N stage 2 or 3) but
100 its risk-ratio of 2.82 was exceeded by that found for all CTTA parameters. The
101 presence of 2 or more high-risk CTTA values was the greatest risk-factor for
102 mediastinal metastases, being associated with an 11-fold risk for mediastinal
103 malignancy ($p=0.0014$) and significantly poorer overall survival ($p=0.016$, Figure 3).
104 The only significant risk factor for N3- disease was mediastinal nodal size ≥ 10 mm.

105

106 **DISCUSSION**

107 In this study we investigated the potential for CT measures of tumour heterogeneity to
108 stratify risk of nodal disease in patients with NSCLC. CTTA parameters within the primary
109 tumour were found to be superior predictors of nodal stage >0 and nodal >1 than the
110 measurement of mediastinal nodal diameter as currently recommended by NICE. However,
111 mediastinal nodal diameter was the best predictor for N-stage >2 . The finding that the
112 presence of two or more high-risk CTTA values is not only associated with a greater risk of

113 nodal stage >1 but also reduced survival, indicates that the presence nodal disease inferred
114 by high-risk CTTA values is of clinical importance.

115 CTTA has the potential to determine the optimum strategy for investigation of nodal disease
116 more accurately than measurements of mediastinal nodal diameter as currently
117 recommended by NICE. For example, the NICE guidance would recommend the nodal status
118 for the patient with N2 disease illustrated in Figure 1 be determined by PET-CT alone. While
119 PET CT is a non-invasive method in assessing for mediastinal nodal metastasis, its sensitivity
120 and specificity of detecting mediastinal metastasis is only around 77% and 86%
121 respectively(6). Furthermore the sensitivity of PET for detecting nodal metastasis is lower
122 when the size of the lymph node is less than 10mm(7). A PET/CT only strategy for this
123 patient would have resulted in underestimation of disease status whereas the presence of
124 high-risk CTTA parameters could feasibly have indicated the need for nodal sampling prior to
125 surgery. Similarly, due to the presence of an enlarged mediastinal node, the NICE guidance
126 would recommend nodal sampling be considered for the patient with N0 disease illustrated
127 in Figure 2 whereas this procedure may not have been considered necessary based on the
128 absence of high-risk CTTA values. As the NICE strategies were optimised on the basis of cost-
129 effectiveness, CTTA therefore also has potential health economic benefits.

130 The observation that tumours with increased metastatic potential exhibit greater genetic
131 instability (8) provides a biological basis for a relationship between CT measures of tumour
132 heterogeneity and nodal status. In the presence of genetic instability, the evolutionary
133 dynamics of tumour development may result in the co-existence of genetically distinct sub-
134 clones within the same tumour (9). For these sub-clones to be detected by CT the genetic
135 status of the sub-clone must correspond to one or more phenotypic features demonstrable
136 by imaging. Furthermore, the spatial separation between sub-clones needs to be sufficiently
137 large relative to the spatial resolution of CT (See Figure 4). The first requirement is shown to

138 be met by radiogenomic studies demonstrating correlations between genomic aberrations
139 and specific CT features in NSCLC and other tumours (5, 10-15). Spatial separation of sub-
140 clones has also been reported and may be sufficiently large to result in sampling error during
141 image-guided biopsy (9). Although plausible, a direct connection between genomic
142 instability, metastatic potential and CT measures of tumour heterogeneity remains to be
143 demonstrated empirically.

144 The main limitation of our study is the small size of our cohort of NSCLC patients without
145 distant metastases for whom nodal status had been confirmed either at surgery or
146 endobronchial biopsy. Larger multi-centre studies are required to confirm our findings. To
147 minimise potential bias, we have adopted median textures as the thresholds for the
148 categorisation of high and low risk for nodal metastases. Larger studies would also allow
149 optimisation of these threshold values through use of separate training and evaluation
150 cohorts. We have also adopted a single slice approach to assess for tumour heterogeneity
151 which has the potential to miss regions of greater heterogeneity and hence under estimate
152 the risk of nodal metastasis (3). The consistency of risk stratification by CTTA may also be
153 potentially improved through automated CT ROI definition based on the PET component of
154 PET/CT examinations(16).

155 The potential prognostic value of CTTA in NSCLC was first reported in 2012 (17) but as yet
156 this finding has not been translated into a widely adopted clinical application. It has been
157 proposed that the prognostic information afforded by CTTA could be used to identify
158 patients with a greater risk for post-surgical recurrence who might benefit most from
159 adjuvant chemotherapy, or to recognise those patients with advanced disease who are
160 unlikely to get sufficient survival benefit to justify the morbidity of chemotherapy in a
161 palliative setting (2). However, the use of CT as a prognostic biomarker represents a novel
162 application for imaging in the management of NSCLC, and in cancer care in general; a fact

163 that may represent a block to the adoption of CTTA into clinical practice. On the other hand,
164 the use of CT markers to assess risk of mediastinal malignancy is embedded within the
165 current clinical guidelines published by NICE. The replacement of measurements of
166 mediastinal nodal size by CTTA measurements of tumour heterogeneity would simply
167 represent a development of an established concept and may therefore be more readily
168 adopted into clinical practice.

169 In summary, CTTA in NSCLC is related to nodal status in patients without distant
170 metastases and has the potential to inform selection of investigative strategies for
171 the assessment of mediastinal malignancy.

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222 in non-small cell lung carcinoma assessed by CT texture analysis: a potential marker of
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224 **FIGURES AND TABLES**

225 Figure 1. A: Axial CT image from a patient with a left upper lobe NSCLC categorised as high
226 risk by CTTA using the filtration-histogram approach. B: Tumour ROI after filtration to
227 highlight image features of radius 4mm. All three CT texture parameters derived from the
228 histogram of values in filtered ROI (C) indicated increased risk for nodal disease: 4.74, 0.73
229 and -0.09 for normalised SD, normalised E and kurtosis respectively. Maximum intensity
230 projections (D) and fused axial images (E) from the patient's PET/CT examination, which
231 showed no nodal disease. N2 disease was confirmed at surgery.

232

233 Figure 2. A: Axial CT image from a patient with a right upper lobe NSCLC categorised as low
234 risk by CTTA using the filtration-histogram approach. B: Tumour ROI after filtration to
235 highlight image features of radius 4mm. All three CT texture parameters derived from the
236 histogram of values in the filtered ROI (C) indicated a reduced risk for nodal disease: 8.61,
237 0.85 and -0.86 for normalised SD, normalised E and kurtosis respectively. The largest

238 mediastinal node measured 11mm on CT (D), indicating a high risk of mediastinal malignancy
239 according to NICE guidance. NO disease was confirmed at surgery.

240

241 Figure 3: The survival curve for patients with less than two high-risk CTTA values (above)
242 showed no deaths in the follow-up period, compared to five deaths for patients with 2 or
243 more high-risk CTTA values (below; $p=0.016$).

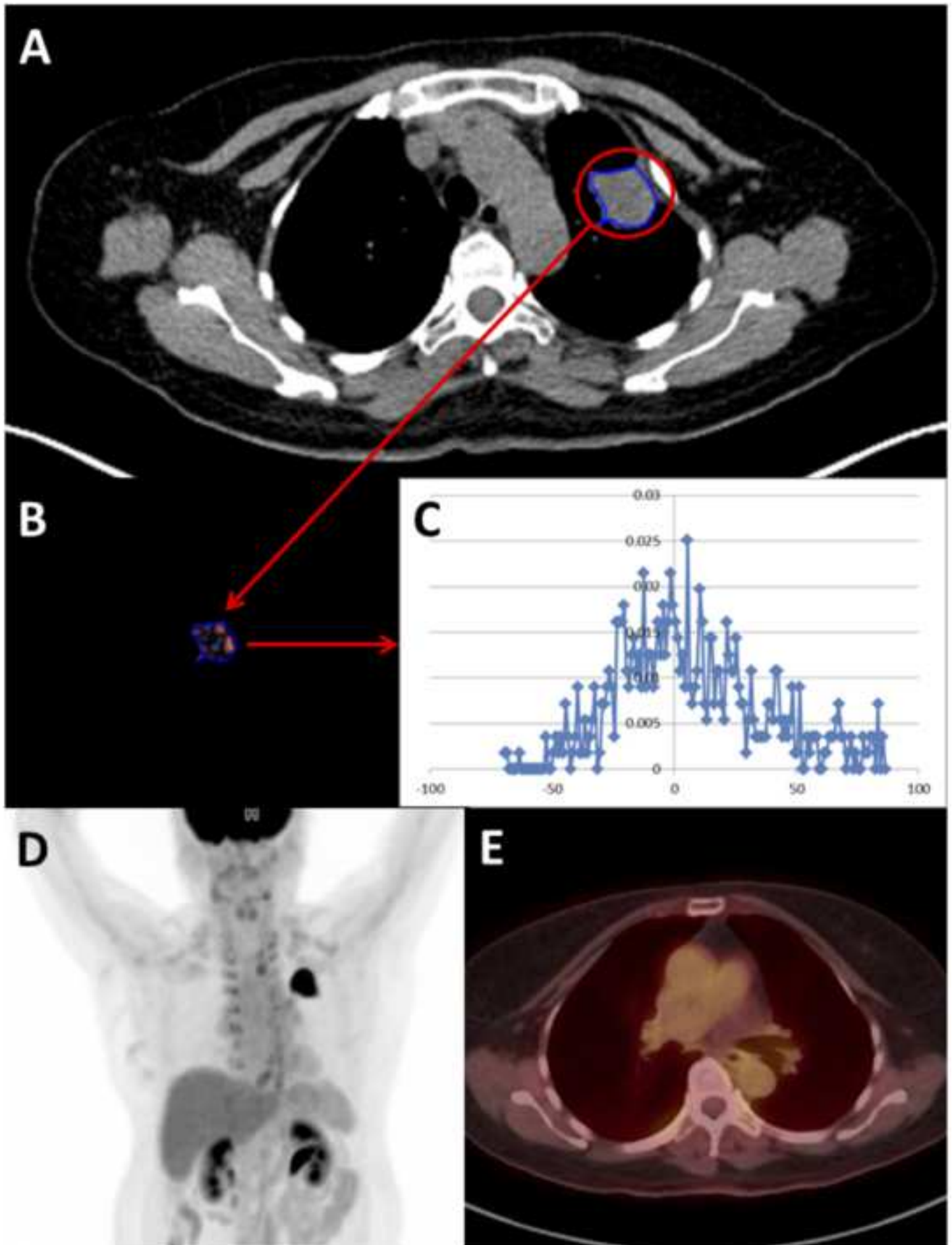
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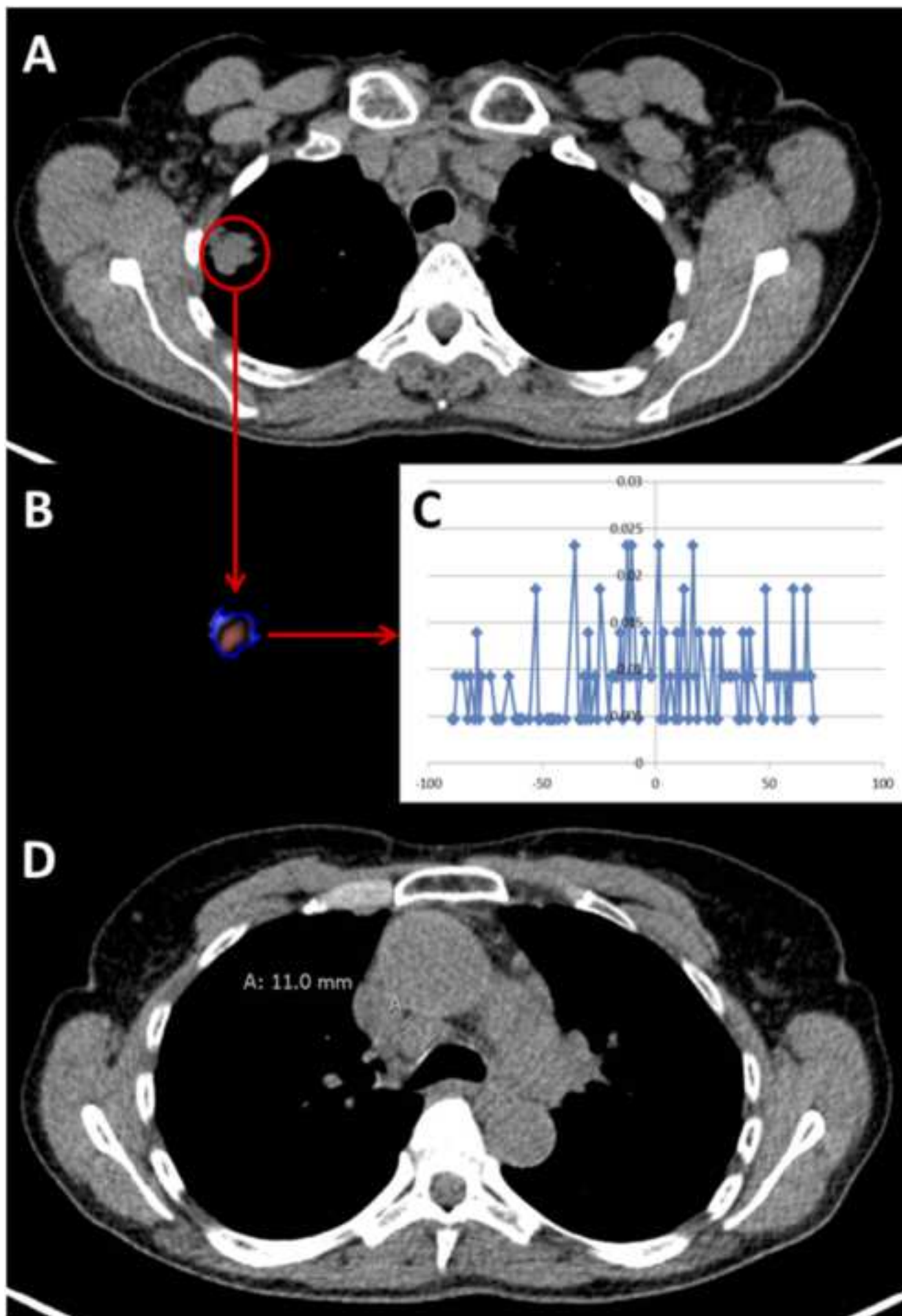
245 Figure 4: Diagrammatic representation of how genetic instability can lead to genetically
246 distinct subclones within a tumour, which express different phenotypes that lead to imaging
247 heterogeneity. This in turn can be assessed quantitatively with CT Texture analysis

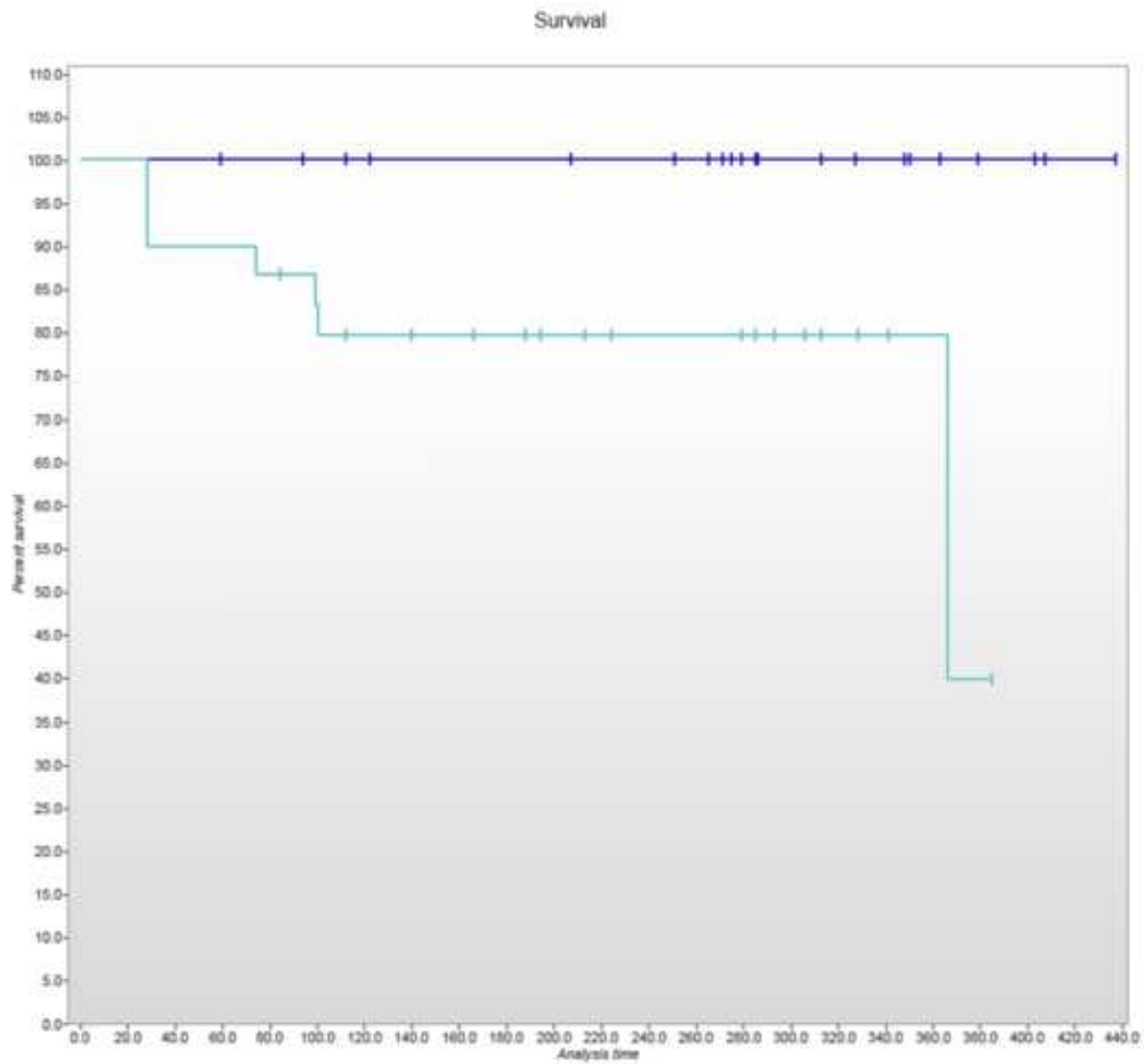
248

249 Table 1. Risk of advancing N-stage for patients categorised by the median texture values
250 with mediastinal nodal size at CT as comparator.

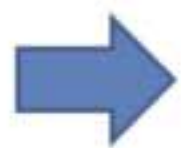
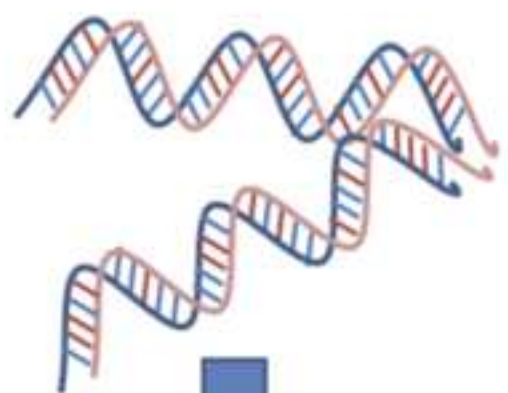
251



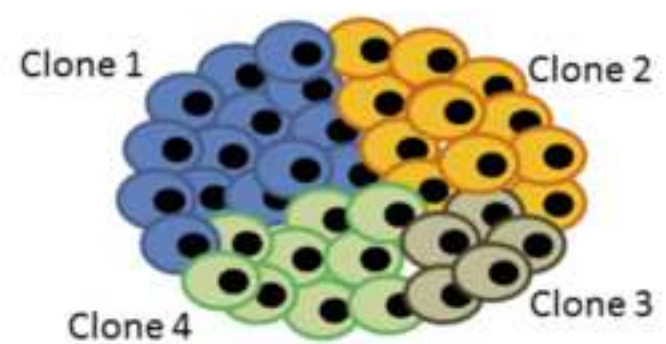




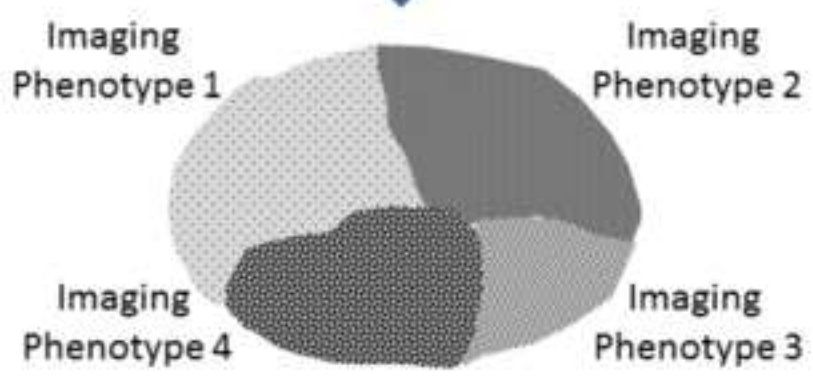
GENETIC INSTABILITY



CLONAL HETEROGENEITY



METASTASIS



IMAGING HETEROGENEITY

Nodal Status	Parameter	High Risk	Low Risk	Risk Ratio (95% CI)	p-value (Fisher's test)
>0	Normalised SD	52.4%	23.8%	2.2 (95% CI 0.92-5.24)	0.11
	Normalised E	61.9%	14.3%	4.3 (95% CI 1.44 -13.0)	0.0036
	Kurtosis	57.1%	19.1%	3.0 (95% CI 1.15-7.80)	0.25
	Any 2 CTTA	61.9%	14.3%	4.3 (95% CI 1.44 -13.0)	0.0036
	Nodal size on CT	54.6%	32.3%	1.69(95% CI 0.80-3.55)	0.28
>1	Normalised SD	42.9%	14.3%	3.0 (95% 0.94-9.55)	0.086
	Normalised E	47.6%	9.5%	5.0 (95% 1.24-20.1)	0.015
	Kurtosis	47.6%	9.5%	5.0 (95% 1.24-20.1)	0.015
	Any 2 CTTA	52.4%	4.76%	11.0 (95% CI 1.56 - 77.8)	0.0014
	Nodal size on CT	54.6%	19.4%	2.82 (95% CI 1.15 – 6.9)	0.049
>2	Normalised SD	19.1%	0%	∞	0.11
	Normalised E	14.3%	4.8%	3.0 (95% 0.34-26.6)	0.61
	Kurtosis	19.1%	0%	∞	0.11
	Any 2 CTTA	19.1%	0%	∞	0.11
	Nodal size on CT	27.3%	3.2%	8.45 (95% CI 0.98 – 73)	0.048

HIGHLIGHTS

- CT texture analysis (CTTA) is emerging as a technique for derivation of prognostic biomarkers for patients with NSCLC
- CT texture analysis evaluates quantitatively tumour heterogeneity, which is linked to tumour aggression.
- CTTA in NSCLC is related to nodal status in patients without distal metastases and has potential to inform selection of investigative strategies for the assessment of mediastinal malignancy.