

Original Article

A novel ultrasound-based approach to investigate extramedullary haematopoiesis in foetal spleen

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Abstract: Foetal spleen is described as a transient focus of haematopoiesis between the 3rd and 5th month of gestation: this function is however entirely replaced by the bone marrow before the end of pregnancy. This study identifies haematopoiesis in foetal spleen by exploring changes of echogenicity during its development throughout gestation. Two intervals of pregnancy were studied: Mid-Pregnancy (Mid-P, 19-23 weeks) and End-Pregnancy (End-P, 37-41 weeks). The foetal spleen was investigated in 80 pregnant women (41 vs 39). Due to quality criteria the comparison was made between 60 images (30 Mid-P vs 30 End-P). The acquisition of splenic parenchyma was followed by clustering segmentation. We identified two new parameters resulted from the clustering segmentation: Dark Ratio (DR) and Light Ratio (LR). These are related to splenic echogenicity expressing the percentage of dark and light signal in the clustered image, influenced by blood cellularity. The mean of DR value was different among the 2 groups (0.0631 vs 0.0483, $P = 0.014$), while LR did not show any significant differences. We conclude that DR may represent a reliable radiomic parameter in the determination of extramedullary haematopoiesis in the spleen.

Keywords: Foetal spleen, extramedullary haematopoiesis, ultrasound, radiomics, clustering segmentation

Introduction

Spleen is the largest lymphoid organ in human body. It appears at 5th-6th week of gestational age (GA) on the left side of the dorsal mesogastrium [1]. From the 18th week of GA, two functionally distinct compartments arise: the red and white pulp [2]. Haematopoiesis can be detected in foetal spleen from the third month of pregnancy [3-6]. This function is transitional and tends to be replaced by the bone marrow (BM) at the sixth-seventh month [7]. After delivery, spleen acquires the germinal centres which give the definitive lymphoid structure [2].

The role of foetal spleen in extramedullary haematopoiesis (EMH) is still controversial, however, many studies support its function as mid-gestation haematopoietic organ [4, 7, 8]. The assumption of haematopoietic activity in foetal

spleen is historically based on murine models [2, 9]: haematopoietic development of spleen in humans parallels that of mouse [10].

Increasing evidence [6, 8, 11] reveals that haematopoietic stem cells (HSCs) can be collected from foetal spleen: the migration into the organ may occur during the first trimester of pregnancy. Electron microscopy studies [12] confirm presence of resident splenic immature HSCs at the first and second trimester, possibly as result of their entrapment from foetal blood, rich of HSCs [13, 14]. However, this finding seems to be a transient phenomenon: HSCs are not observed in spleen at the ninth month [11], when the blood still contains an amount of CD34+ three times that of adult BM [5, 15].

An increase of absolute number of CD34+ cells in human foetal spleen is detectable: 1×10^6 at

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16-17 weeks of GA to 11×10^6 at 21-22 weeks of GA; notably the spleen exhibits the higher percentage increase of CD34+ compared to BM and foetal liver [16].

In adult life, spleen may reacquire its role in EMH under certain circumstances, such as infections, inflammations and malignancy [17]. A typical example is what happens in the setting of myelofibrosis, a disease in which BM becomes inhabitable [18]. The haematopoietic stem and progenitor cells migrate and seed in the spleen: these cells initiate splenic EMH. However, this new microenvironment is not suitable as the BM, therefore this EMH is ineffective [19].

Spleen may be investigated through ultrasound (US) only from the 18th week of GA [20]. Previous studies [20, 21] generated US splenic nomograms to characterize the growth of foetal spleen during pregnancy. However, the sonographic appearance of parenchyma was not considered.

US represents the technique of choice for regular monitoring of pregnancy [22]; the extension of radiomic analysis may constitute a promising approach in this field [23-25]. The spread of this novel technique is transforming the conception of diagnosis and prognosis in medical imaging [23]. However, there are no published data on the use of radiomic analysis investigating the spleen by using US, but only on computed tomography technique [24].

The aim of this work is to assess the echogenicity of foetal spleen during pregnancy by using US imaging, in order to identify splenic EMH. Our study constitutes the first assessment of splenic echogenicity through two radiomic parameters, which resulted from clustering segmentation of the images [25, 26].

To evaluate any significant differences, two groups were studied: Mid-Pregnancy (Mid-P, 19-23 weeks of GA) and End-Pregnancy (End-P, 37-41 weeks of GA).

Material and methods

Patient population

The study was conducted at “Antonio Cardarelli” hospital in Campobasso, Molise, Italy. A to-

tal number of 80 (41 Mid-P vs 39 End-P) different healthy pregnant women were retrospectively enrolled in this study. They had given their written consent to collect and publish the US images of foetal spleen and their own information, anonymously. Due to the spread of COVID-19 pandemic, our group of study decided to stop further recruitment because of a slower-than-anticipated enrolment rate, that rendered the number of participants unlikely to reach the projected target within a reasonable time frame. Data related to hypertension, dyslipidaemia, diabetes, smoking history, alcohol, drug use, age, weight, height and ethnicity group were collected. Patients characteristics are summarized in **Table 1**. This work received the approval by the “Comitato Etico Università Federico II” ethics committee of University of Naples Federico II (n. 400/20).

Ultrasound examination

The US scans were performed using the 2D Esaote MyLab™ Seven eHD Crystaline, AC2541 1-8 MHz convex US probe, on “obstetrics” machine setting. All the scans were acquired by the same operator, who was blind to the study. After each US examination, the acquired scans were stored in the digital diagnostic imaging archive of Obstetrics and Gynaecology Operating Unit. GA was estimated by US foetal biometry [20]. The spleen was best selected and visualized on axial abdominal scan, in absence of acoustic shadows involving the organ, when the foetus lied in an occipito-posterior position [27]: gastric bubble and first portion of umbilical vein were chosen as reference points to select comparable scans (**Figure 1**). The dimensional parameters (*length*, *width*, *circumference* and *area*) were digitally measured and registered as the average of three different measurement: resulting values were aligned with splenic nomograms of previous studies [20, 21]. Presence of maternal (gestational diabetes, pre-eclampsia, infections), foetal (small for gestational age, large for gestational age), placental (placenta previa, accreta, abruptio placentae) and/or amniotic (oligo- or polyhydramnios) pathological conditions and/or multiple gestations were defined as exclusion criteria. Only two pregnant women in the End-P group were under anti-hypertensive agents. Inclusion criteria are described as follows: in Mid-P group were included healthy foetuses at

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Table 1. Features of the pregnant women (*Mean ± SD*)

	<i>n</i>	<i>GA (weeks)</i>	<i>Age (years)</i>	<i>Weight (kg)</i>	<i>Height (m)</i>	<i>BMI</i>	<i>Hypertension</i>		<i>Diabetes</i>		<i>Dyslipidaemia</i>		<i>Smoke</i>		<i>Ethnicity</i>	
							<i>no</i>	<i>yes</i>	<i>no</i>	<i>yes</i>	<i>no</i>	<i>yes</i>	<i>no</i>	<i>yes</i>	<i>white</i>	<i>black</i>
<i>Group Mid-P</i>	41	20.39 ± .74	29.78 ± 6.00	65.09 ± 8.81	1.64 ± .06	24.16 ± 2.48	41	0	41	0	39	2	31	10	36	5
<i>End-P</i>	39	38.62 ± 1.44	34.18 ± 4.96	77.97 ± 14.00	1.66 ± .06	28.10 ± 4.51	36	3	38	1	37	2	35	4	36	3

Abbreviations: Mid-P (Mid-Pregnancy), End-P (End-Pregnancy), n (number), GA (Gestational Age), BMI (Body Mass Index), SD (Standard Deviation).

RADIOMIC MULTISTEP PROCESS

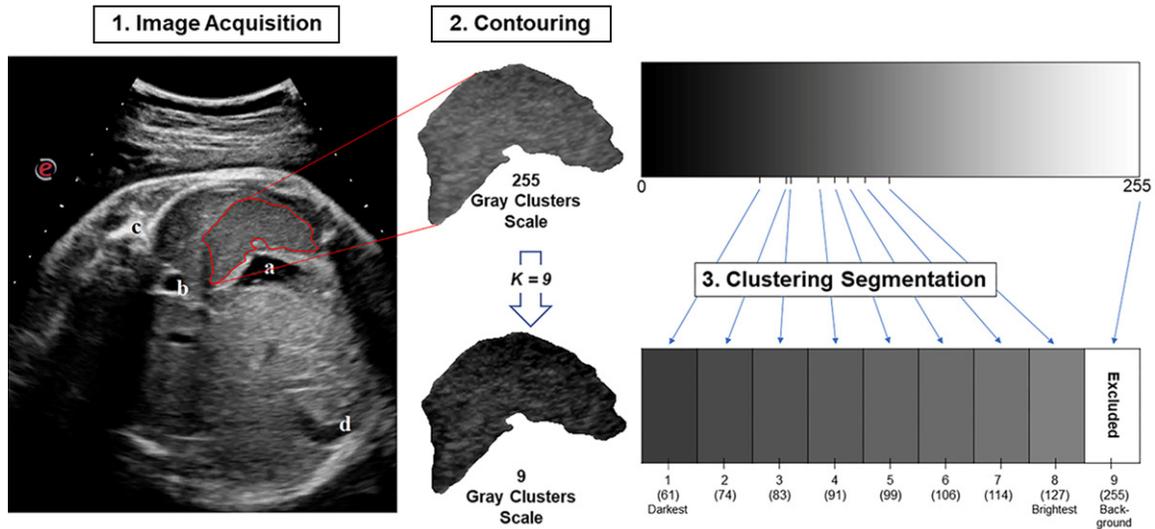


Figure 1. Schematic representation of radiomic workflow ($k = 9$ in k-means algorithm). 1. Detection of foetal spleen in the abdominal scan (red trace): a - gastric bubble, b - aorta, c - vertebra, d - umbilical vein. 2. Computed isolation of spleen parenchyma (“region of interest”) from adjacent structures. 3. Image conversion from a 0-255 to 1-9 greyscale clusters. As neutral background of clustered image, white cluster is not considered the brightest cluster.

19-23 weeks of GA; in End-P group were included healthy foetuses at 37-41 weeks of GA. Spleen images were considered for inclusion only if abdominal scans (axial view), with a clearly visible spleen, without acoustic shadows involving the region of interest and in presence of the aforementioned reference points.

From February 2019 to February 2020, a total of 80 images were collected (41 vs 39). Twenty images were excluded (13 malposition, 2 small for gestational age, 1 gestational diabetes, 2 pre-eclampsia, 2 oligohydramnios). The remaining 60 images, 30 Mid-P and 30 End-P were analysed, thus underwent clustering segmentation. We digitally isolated the splenic parenchyma by photo editing software (Adobe Photoshop® CC 2017): this tool allowed the accurate isolation of spleen parenchyma, preserving images and informative content quality (Figure 1).

Computer processing

Radiomics is a new challenging approach that transforms digitally encoded medical images into mineable high-dimensional data [28]. The first step consists of image acquisition, followed by the segmentation of the “region-of-interest” [25]. In this study the algorithm k-means was used as part of clustering segmentation [26]. The K-Means algorithm oper-

ates finding and grouping datapoints in classes that have high similarity between them. This similarity is understood by the system as the opposite of the distance between datapoints. The closer the data points are, the more similar and more likely to belong to the same cluster (see below) they will be [25]. The distance between the datapoints is calculated by Euclidean distance.

The algorithm works through several steps: a) selection of the “k” number of clusters in which the datapoints need to be grouped; b) first random selection of the cluster centroids; c) assignment of each datapoint to the closest centroid; d) calculation of within-the-cluster-sum-of-squares value; e) final definition of the “new” centroids by calculation of the minimum quadratic error between datapoints and the centre of each cluster; f) re-assignment of the datapoints to the definitive centroids. This process can be repeated multiple times.

In this study, the datapoints represent the pixels within the grayscale image (single channel grayscale map) of the isolated spleen parenchyma (the “region of interest”): every pixel in the image is defined by one grayscale value, so similarities can be detected from the system.

Here the k-means assigns each pixel of the grayscale scan (0 to 255 shades of grey; 0 -

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Table 2. Differences in foetal splenic parameters of Mid-P and End-P groups. DR9 and LR9 values are shown (Mean \pm SD)

	Length (mm)	Width (mm)	Circumference (mm)	Area (mm ²)	DR9	LR9
Group Mid-P	14.99 \pm 2.50	8.28 \pm 1.41	42.42 \pm 5.65	.95 \pm 0.23	.063 \pm .026	.039 \pm .025
End-P	40.42 \pm 6.31	16.79 \pm 3.21	113.12 \pm 17.12	5.68 \pm 1.67	.048 \pm .018	.036 \pm .020

Abbreviations: Mid-P (Mid-Pregnancy), End-P (End-Pregnancy), DR9 (Dark Ratio, when k = 9 in k-means algorithm), LR9 (Light Ratio, when k = 9 in k-means algorithm), SD (Standard Deviation).

Table 3. Pearson rank correlation analysis

Variable	GA	Weight	Height	BMI	DR9	LR9
GA	P = 1	0.504	0.219	0.495	-0.298	-0.065
	p-value = 1	< 0.001	0.051	< 0.001	0.021	0.619
Weight		P = 1	0.572	0.928	-0.172	-0.083
		p-value = 1	< 0.001	< 0.001	0.188	0.528
Height			P = 1	0.229	-0.023	0.081
			p-value = 1	0.041	0.862	0.539
BMI				P = 1	-0.181	-0.125
				p-value = 1	0.166	0.341
DR9					P = 1	-1.48
					p-value = 1	0.258
LR9						P = 1
						p-value = 1

Abbreviations: GA (Gestational Age), BMI (Body Mass Index), DR9 (Dark Ratio, when k = 9 in k-means algorithm), LR9 (Light Ratio, when k = 9 in k-means algorithm).

black, 255 - white) into a “k” number of clusters. Furthermore, the algorithm calculates the absolute number of pixels within the clusters with the darkest (Dark Cluster-DC) and brightest (Light Cluster-LC) centroids. Also, the system calculates the percentage of pixels attributed to the DC and to the LC in relation to the total number of pixels in the region of interest: Dark Ratio (DR) and Light Ratio (LR), respectively. After the k-means algorithm run, the isolated region of interest is clustered in a “k-1” number of clusters: one cluster is always excluded because it constitutes the white background surrounding the image. **Figure 1** shows a visual representation of the segmentation algorithm, run with k = 9: each pixel of the image is assigned to one of the 8 clusters, and is represented using the greyscale colour that corresponds to the value of the centroid assigned to its own cluster.

K-means algorithm was run with a convergence tolerance value of 0.0001, using the k-means++ cluster initialization method [29] and each run was executed 4 times, selecting the best one on the basis of the within-the-cluster-sum-of-square score. The clustering seg-

mentation was carried out in six different number of clusters (k = 4, 5, 6, 7, 8, 9); values of DR and LR for each clustering were calculated (DR4-LR4, DR5-LR5, DR6-LR6, DR7-LR7, DR8-LR8, DR9-LR9). The “silhouette score” [26] was determined as a cluster validity method for each number of clusters: k = 9 exhibited the higher value (0.7774). Therefore, DR9 and LR9 were used to match the two groups. Splenic parameters are summarized in **Table 2**.

Statistical analysis

Shapiro-Wilk test was used to investigate the normal distribution of DR9 and LR9 parameters [30]. Because of the normal distribution of variables, parametric tests were used. Pearson rank correlation analysis was performed to investigate the relation between DR9 and LR9 with the weeks of gestations, weight, height and body mass index of pregnant women. The difference in DR9 and LR9 between the groups was further investigated through the independent samples t-test and ANOVA one-way test. Since DR9 showed a normal distribution in Mid-P group and non-normal in End-P group, we performed both the independent samples

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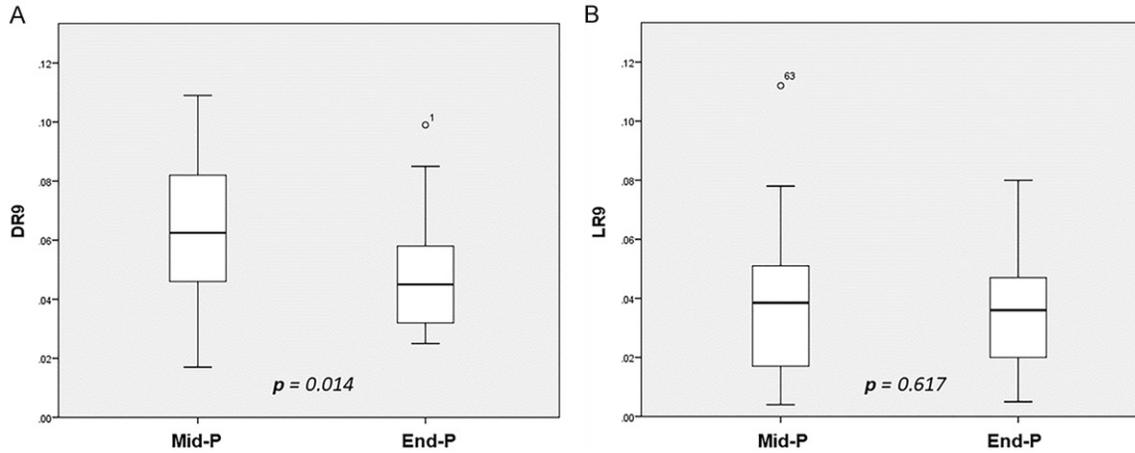


Figure 2. DR9 values in Mid-P and End-P groups (A). LR9 values in Mid-P and End-P groups (B). *P*-values are shown in the graphs. Abbreviations: Mid-P (Mid-Pregnancy), End-P (End-Pregnancy), DR9 (Dark Ratio, when $k = 9$ in k -means algorithm), LR9 (Light Ratio, when $k = 9$ in k -means algorithm).

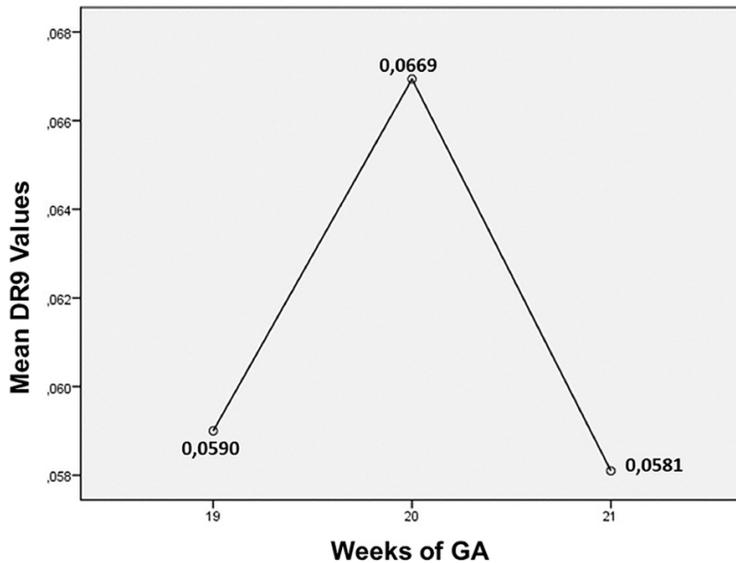


Figure 3. Mean DR9 values in Mid-P group at 19-21 weeks of GA. Abbreviations: Mid-P (Mid-Pregnancy), GA (Gestational Age), DR9 (Dark Ratio, when $k = 9$ in k -means algorithm).

t-test (parametric) and Mann-Whitney test (non-parametric) to investigate the differences of DR9 in these two groups. All tests were performed by using IBM SPSS® Statistics 21.0 and only $p < 0.05$ results were considered statistically significant.

Results

Shapiro-Wilk normality test showed a normal distribution of variables except for a non-normal distribution of DR9 in the End-P group. Pearson rank test showed a significant correla-

tion between DR9 and weeks of gestation ($r = -0.298$, p -value = 0.021). Height, weight, body mass index and age of the patients did not correlate to the DR9 or LR9. Results from Pearson correlation test are shown in **Table 3**. Independent samples t-test showed a difference between the DR9 in Mid-P versus End-P (0.0631 and 0.0483 respectively - $p = 0.014$). Because of the non-normal distribution of DR9 in End-P group, we performed Mann-Whitney test, which showed comparable results ($p = 0.023$). LR9 did not show any difference between groups: $p = 0.617$ (**Figure 2**).

Discussion and conclusion

Previous applications of US in foetal spleen evaluation were merely focused on morphological description (shape, dimensions) [20, 31, 32] or in assessing parenchymal lesions [33]. Currently, no data are available on US-definition of normal/healthy foetal spleen parenchyma: its echogenicity was simply defined as similar to that of liver and kidneys [21]. This study represents the first US-based assessment of healthy spleen parenchyma in two GAs (Mid-P vs End-P), during which this organ seems to play different roles (haematopoietic vs non-haematopoietic) [3, 4, 6]. The variation

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of splenic echogenicity is hard to appreciate without an image processing computer system, which generates radiomic parameters related to echogenicity (DR and LR). The application of radiomics to US technique and the expansion of this new challenging field to prenatal US screenings seems to be a rapidly expanding phenomenon [34-36].

Our results show that sonographic appearance of spleen may change during the course of pregnancy. This may reflect a different activity of spleen between the fifth and ninth month. The difference is significant in DR9 ($p = 0.021$); we speculate that the difference in anechoic/dark signals during these GAs may depend on different cell populations, rather than vascular structure. The slight difference between the two groups is non-significant ($p = 0.617$) in LR9: this may hypothetically reflect a similar architecture of hyperechoic regions.

In fact, considering the equivalent vascular development of foetal spleen at 22nd and 38th GA [11], we believe that darker sonographic signal in Mid-P group correlates with higher density of HSCs and this may impact on the echogenicity of spleen parenchyma.

HSCs can be collected from the spleen during 13th to 36th weeks of GA [11], but these are undetected at ninth month, when splenic haematopoiesis is described to stop [6]. This finding could support the hypothesis that spleen may contribute as a minor and transient haematopoietic organ in foetal life.

Interestingly, based on DR values progression (Figure 3) in Mid-P group, the spleen exhibits its darker appearance at the 20th week of GA. This event could represent the peak of splenic HSCs due to a greater haematopoietic or trapping activity [9]. The difference between DR9 mean at 20th week of GA versus End-P group was statistically significant ($P = 0.008$).

This study represents the first attempt to examine an historical controversy using US, a non-invasive and widespread technique. Considering that histology of spleen cannot be investigated in living foetuses, US-based radiomic analysis could represent a valid approach to evaluate splenic structure and activity. Ideally, the sonographic appearance of spleen should be followed throughout the whole gestation and in a larger cohort of pregnant women.

We also believe this novel strategy could be employed as modern imaging approach to explore EMH in adult pathological conditions, such as myeloproliferative neoplasms. The characterization of this phenomenon could be the first step for a future non-invasive diagnostic tool and provide useful information regarding prognosis and response-to-treatment of these conditions.

Disclosure of conflict of interest

None.

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