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DOI: 10.1016/j.placenta.2021.02.015

Document Version Peer reviewed version

Link to publication record in King's Research Portal

Citation for published version (APA):

Steinweg, J. K., Hui, G. T. Y., Pietsch, M., Ho, A., van Poppel, M., F A Lloyd, D., Colford, K., Simpson, J., Razavi, R., Pushparajah, K., Rutherford, M., & Hutter, J. (2021). T2\* placental MRI in pregnancies complicated with fetal congenital heart disease. *Placenta*, *108*, 23-31. https://doi.org/10.1016/j.placenta.2021.02.015

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### 1 Title

2 T2\* placental MRI in pregnancies complicated with fetal congenital heart disease

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28	Abstract					
29	Background					
30	Congenital heart disease (CHD) is one of the most important and common group of					
31	congenital malformations in humans. Concurrent development and close functional					
32	links between the fetal heart and placenta emphasise the importance of understanding					
33	placental function and its influence in pregnancy outcomes. The aim of this study was					
34	to evaluate placental oxygenation by relaxometry (T2*) to assess differences in					
35	placental phenotype and function in CHD.					
36	Methods					
37	In this prospective cross-sectional observational study, 69 women with a fetus affected					
38	with CHD and 37 controls, whole placental T2* was acquired using a 1.5-Tesla MRI					
39	scanner. Gaussian Process Regression was used to assess differences in placental					
40	phenotype in CHD cohorts compared to our controls.					
41	Results					
42	Placental T2* maps demonstrated significant differences in CHD compared to controls					
43	at equivalent gestational age. Mean T2* values over the entire placental volume were					
44	lowest compared to predicted normal in right sided obstructive lesions (RSOL) (Z-					
45	Score 2.30). This cohort also showed highest lacunarity indices (Z-score -1.7), as a					
46	marker of lobule size. Distribution patterns of T2* values over the entire placental					
47	volume were positively skewed in RSOL (Z-score -4.69) and suspected, not confirmed					
48	coarctation of the aorta (CoA-) (Z-score -3.83). Deviations were also reflected in					
49	positive kurtosis in RSOL (Z-score -3.47) and CoA- (Z-score -2.86).					
50	Conclusion					
51	Placental structure and function appear to deviate from normal development in					
52	pregnancies with fetal CHD. Specific patterns of altered placental function assessed by					
53	T2* deliver crucial complementary information to antenatal assessments in the					
54	presence of fetal CHD.					
55	Keywords					
56	Congenital Heart disease (CHD), Placenta, Magnetic resonance imaging (MRI), T2*					
57	Mapping, Gaussian Process Regression (GPR), Machine learning					
58						

### Introduction

59

60 Congenital heart disease (CHD) is a common group of congenital malformations with a 61 prevalence of up to 1% of all live births, a leading cause of neonatal and infant death 62 and a global burden in child health [1,2]. Underlying aetiology is considered 63 multifactorial combining genetic, epigenetic and environmental causes [3-5]. Recently, 64 placental vascular malperfusion has been reported at pathological examination in 65 association with CHD [6,7] and it has been postulated that this may contribute to the 66 neurodevelopmental abnormalities observed in children with CHD [8]. 67 The placenta is the only connection between the fetal and maternal circulation. It plays 68 a pivotal role in fetal development due to its responsibility for all fetomaternal 69 exchange, including oxygen, carbon dioxide and nutrients, and essential immunological 70 and homeostatic functions. The placenta and fetal heart develop concurrently and 71 share similar developmental pathways – amplifying its unique role and vulnerability to 72 disturbances especially in the presence of fetal CHD [9,10]. A sophisticated in vivo 73 assessment of placental structure and function may inform understanding of complex 74 antenatal pathophysiology in pregnancies with fetal CHD and help to identify fetuses at 75 greater risk of adverse long term neurodevelopmental outcomes. 76 A common MR contrast mechanism employed to study tissue oxygen concentration is 77 produced by the blood oxygen level dependency (BOLD) effect, exploited in 78 quantitative T2 relaxometry (T2\*). The parametric properties of deoxyhaemoglobin 79 allow for a faster T2\* decay than its oxygenated counterpart [11,12]. Widely used in 80 functional brain MRI, it has also recently found application in placental MRI, allowing in-81 vivo insights into the tissue properties without the use of exogenous contrast agents 82 [13-15]. 83 Previous studies in humans have shown a linear negative correlation of placental mean 84 T2\* values over gestational age (GA) and found it to be predictive for low birth weight 85 [15,16]. An increase in placental mean T2\* was previously reported in response to 86 maternal oxygen administration [17,18]. Decreased mean T2\* values can be induced by vasoconstrictive agents in animal models and have been demonstrated in 87 88 pregnancies with fetal growth restriction in both animals and humans [19-21]. Reduced 89 whole placental mean T2\* values for GA were recently also associated with 90 preeclampsia [22]. 91 Mean T2\* value calculated over the placental parenchyma, either assessed in selected 92 slices or the entire placental volume have been used for quantification in previous

93	studies. However, novel advances extend this to include spatial analysis using
94	histograms and texture-based measurements, first shown in rhesus monkeys [23,24].
95	This study aimed to evaluate whether CHD is associated with altered placental
96	phenotype and function, including signs of lower oxygenation and altered tissue
97	morphology, as estimated by high-resolution whole placental MR relaxometry (T2*) and
98	to establish methods to evaluate deviations from the expected values over GA.

## **Methods**

100	In this prospective cross-sectional observational study, we recruited pregnant women
101	carrying a fetus diagnosed with congenital heart disease (CHD) from a tertiary fetal
102	cardiology service for CHD (St Thomas' Hospital, London, UK) alongside a control
103	group of women with uncomplicated pregnancies during the second and third trimester
104	Inclusion criteria for both cohorts were singleton pregnancy, maternal age over 18
105	years and ability to consent in English language. Exclusion criteria for both cohorts
106	were multiple pregnancies, major maternal health issues, any treatment for
107	hypertension at time of scan as well as contraindications for MRI such as metallic
108	implants and claustrophobia. Exclusion criteria for controls also included diagnosis of
109	fetal or intrauterine growth restriction (FGR/IUGR) and low birth weight (SGA), signs of
110	preeclampsia or (gestational) diabetes at time of scan.
111	Prospectively specified data collection included demographic characteristics and
112	maternal pregnancy data at time of scan, as well as delivery notes and neonatal
113	outcomes. Maternal data included age and BMI at scan, parity, gravida, medical history
114	of smoking, (gestational) diabetes, hypertension, preeclampsia, HELLP syndrome,
115	thyroid disease and anxiety or depression as noted at time of scan and in their delivery
116	notes. Neonatal data included fetal and neonatal cardiac diagnosis, sex, GA at scan
117	and birth, birth weight, 5 min APGAR, neonatal outcome, genetic testing, suspicion of
118	genetic abnormality and place of birth as noted on the delivery or (cardiology)
119	discharge notes.
120	Centiles and standard scores for birth weight were calculated following the
121	INTERGROWTH-21 <sup>st</sup> project [25]. SGA was defined as birthweight less than 10 <sup>th</sup>
122	centile in keeping with common clinical practice. Our CHD cohort was classified into 7
123	groups based on the main antenatal diagnosis (Figure 1, Table 1). We divided our
124	cohort into left sided obstructive lesions, namely hypoplastic left heart syndrome
125	(HLHS) and coarctation of the aorta (CoA), right-sided obstructive lesions (RSOL)
126	defined as structural or functional obstruction of the right ventricular outflow tract,
127	disorders of mixing (e.g., transposition of the great arteries (TGA)), suspected vascular
128	rings (VR) and other major lesions comprising of common arterial trunk (CAT), partial
129	anomalous pulmonary venous drainage (PAPVD) and cardiac rhabdomyomas (CR).
130	The group of suspected CoA was further divided by those with postnatal confirmation
131	requiring surgery within the neonatal period (CoA+) and without postnatal CoA during
132	follow up (CoA-). All our infants are therefore followed up by our team at least until 1
133	year of age.

- 134 Ethics:
- 135 This study was approved by London Research Ethics Committees of the Health
- 136 Research Authority (HRA) of the Department of Health in the United Kingdom,
- "Quantification of fetal brain growth and development using MRI" (REC:
- 138 07/H0707/105), "Fetal Imaging with Maternal Oxygen" (REC: 17/LO/0282) and "iFIND-
- 2. Further Imaging" (REC: 14/LO/1806). Controls were also included from the
- 140 "Placenta Imaging Project" (REC: 16/LO/1573).
- 141 MRI Acquisition:
- 142 All women were scanned using a clinical 1.5-Tesla Philips Achieva MRI with a 28-
- 143 channel torso coil under clinical monitoring and medical cover during the entire scan.
- 144 All mothers carrying a fetus with CHD were scanned in left lateral tilt. A multi-echo
- 145 gradient echo sequence covering the entire uterus in coronal orientation was acquired
- with a resolution of 2.5mm isotropic, FOV=360x360, 50-88 slices, no SENSE, no half-
- 147 scan, TR=14s, TE=11.376ms / 57.313ms / 103.249ms / 149.186ms / 195.122ms, free
- 148 breathing, TA=1min.
- 149 MRI data processing:
- 150 T2\* maps were calculated using an in-house monoexponential fitting in MATLAB (The
- 151 MathWorks Inc, USA). The whole placenta was manually segmented by experienced
- observers (JKS, GH, AH, JH) using ITK-SNAP [26], with exactly matched instructions
- to segment the placenta conservatively, avoiding inclusion of both amniotic fluid and
- maternal vasculature. Reproducibility of manual segmentations between observers
- was assessed using the Sørensen–Dice coefficient [27].
- 156 The masks were automatically refined by excluding non-physiological values. Mean
- 157 T2\*, skewness, kurtosis and lacunarity values were obtained using a purpose-build
- python script [13,28]. Lacunarity values thereby reflect the spatial distribution of gaps of
- a specific size within lobules [29] and a box-size matched to typical placental lobule
- 160 size.
- To estimate placental development over GA we used Gaussian Process Regression
- 162 (GPR), a Bayesian non-parametric regression [30]. Clinical cohorts are inherently
- heterogenous necessitating an estimation of uncertainty of probable distributions.
- Accounting for this and a covariance between the data points GPR provides normative
- group mean function, allowing point estimates and predictive confidence for each
- observation. This allows the calculation of standard (Z-) scores for all measured test
- data points, describing the distance to expected value following the normative group
- mean function [31].

169 The control cohort was split into training and test data (0.7 to 0.3 ratio) and the test 170 subjects were used to train the model. The kernel function used was an additive 171 combination of Constant function, Radial Basis Functions and Noise kernel. The data 172 was scaled pre-training. Predictive posterior distributions were obtained for mean T2\*, 173 lacunarity, skewness and kurtosis separately. Z-scores for the observed values were 174 derived for all CHD cohorts by estimation of mean deviation and median deviation from 175 GPR predicted value normalised by the prediction uncertainty. 176 For comparison of clinical parameters controls were restricted to GA at scan between 177 28 and 37 weeks to match GA for CHD cohort, categorical variables were compared 178 using Fisher's exact test, continuous variables were evaluated for normality by Shapiro 179 compared using Independent T-test (normal distribution), 2-tailed Mann-Whitney U test 180 (non-normal distribution). Dependence of results from clinical parameters was 181 evaluated by coefficient of determination. All statistical analysis and visualisation were 182 performed using SPSS Statistics v27 (IBM) and Jupyter Notebook, python3.

184	Results
185	Maternal and neonatal demographics are demonstrated in Table 1. Overall, 119
186	participants were enrolled in this study. Eight subjects from our CHD cohort were
187	excluded: for twin pregnancy (n=1), insufficient scan data (n=3) and insufficient
188	outcome data at time of analysis (n=4). Five controls were excluded for small for GA at
189	birth (n=3) and insufficient scan data (n=2). In total 69 women with pregnancies with
190	fetal diagnosis of CHD and 37 women with uncomplicated pregnancies were included
191	in the analysis (Figure 1). The median GA at scan was 31.3 weeks (IQR 2.21) and 31.2
192	weeks (IQR 8.11) respectively. Results of genetic testing from antenatal invasive
193	procedures or postnatal blood sample using at least array comparative genomic
194	hybridization (aCGH) were available for 46 (67%) subjects in our CHD cohort. 5/46
195	identified an abnormal result (two with Chr 2q31 deletion, one with a Chr 9 deletion and
196	gene copy number on Chr 2, one with TSC1 gene mutation, and one with mosaic
197	Monosomy X). All remaining patients caused no phenotypic suspicion for a genetic
198	abnormality after birth.
199	Seven newborns with CHD from our cohort died due to necrotising enterocolitis (1),
200	hypoxic ischemic encephalopathy (1), congenital diaphragmatic hernia (1) and cardiac
201	collapse or palliation (4). The CHD cohort had a higher incidence of low APGAR score
202	(<7) at 5 minutes (p=0.024) compared to our control cohort. GA at birth (p<0.001) and
203	birth weight (p<0.001) and birth head circumference (p=0.005) were significantly lower
204	in the CHD cohort, whose mothers were also younger (p=0.017). None of the other
205	clinical collected parameters achieved clinical significance.
206	We have reviewed medical records at time of scan and delivery. Our CHD cohort
207	included three women diagnosed with preeclampsia (4.3%). Two needed delivery at $32$
208	weeks of gestation, one for severe HELLP and one for uncontrollable hypertension.
209	One woman delivered after induction early term at 37 weeks of GA.
210	In our CHD cohort 18 (26%) newborns were born <10 <sup>th</sup> birth weight centile (SGA), 25
211	(36%) newborns were born with low head circumference (HC) <10 <sup>th</sup> centile, 13
212	newborns (19%) were SGA and had low HC. Only one of these 13, would be defined
213	as growth restricted following the consensus-based definition in the newborn by Beune
214	et al. [32].
215	Placental histology was available in 12 pregnancies complicated by fetal CHD on
216	special clinical request mostly due to maternal pyrexia during delivery (Supplemental
217	Table 1). From the two placentas demonstrating signs of maternal vascular
218	malperfusion (MVM), one placenta showed also acute subchorionitis and maternal

- 219 inflammatory response (MIR) in keeping with ascending intra-uterine infection showing
- infarcts of variable age, but no thrombi. The other placenta was from a patient
- 221 diagnosed with severe preeclampsia and HELLP syndrome leading to early delivery.
- 222 Placental histology was also available for 15 control subjects as part of other study
- 223 protocols (Supplemental Table 1).

### 224 Qualitative assessments

- 225 Placental T2\* maps in our cohort of RSOL showed most marked differences compared
- 226 to age-matched controls as depicted in mid-parenchymal slices (Figure 2). Specifically,
- short T2\* values were noted in the entire placenta with additional and faster decay from
- the centre to the periphery of the lobules. Furthermore, increased heterogeneity could
- be observed in RSOL. Our cohorts of left sided obstructive lesions, disorders of mixing
- 230 and other lesions appeared only moderately different to controls at similar GA (Figure
- 231 2). Overall, our CHD cohort appeared to have generally lower signal intensity
- throughout the placenta, advanced lobularity and higher granularity within the lobules
- at a given GA compared to our control cohort.

### **Quantitative assessments**

- 235 Interobserver variability of manually segmented placental masks showed good
- correlation in 10 randomly selected with a Sørensen–Dice coefficient of 0.87.
- 237 Quantitative results from the control cohort illustrate decay in mean T2\* with increasing
- 238 GA, in keeping with previous literature [15,29]. Lacunarity, kurtosis and skewness tend
- to increase over GA in all our cohorts as previously shown in controls [28]. The
- obtained posterior mean of the Gaussian process is given for all quantitative values in
- 241 Figure 3.

- 242 Mean T2\* values over the entire placental volume were lowest compared to predicted
- 243 normal in RSOL (Z-Score 2.30) and our cohort with other major lesions [CAT, PAPVD,
- 244 CR] (Z-Score 2.31). Our CoA- cohort had a larger deviation from expected values (Z-
- 245 Score 1.39) than CoA+ (Z-score 0.24). Mean T2\* values for HLHS (Z-score 0.63), VR
- 246 (0.09) and TGA (Z-score -0.11) were within one standard deviation (SD) from expected
- 247 results.
- 248 RSOL (Z-score -1.7), our group of other major lesions (Z-score -1.26) and CoA- (Z-
- 249 score -1.02) showed significantly higher lacunarity compared to expected results at GA
- equivalent. HLHS (Z-score -0.18), TGA (Z-score -0.01) and suspected VR (Z-score
- 251 0.06) were similar to expected controls. CoA+ showed slightly lower lacunarity (Z-score
- 252 0.26).

- 253 Distribution of T2\* values over the entire placental volume was positively skewed in
- 254 RSOL (Z-score -4.69) and CoA- (Z-score -3.83), followed by our group of other major
- 255 lesions (Z-score -1.75) and HLHS (Z-score -1.12). Suspected VR (Z-score -0.8), CoA+
- 256 (Z-score -0.66) distributions were positively skewed within one SD. Our TGA cohort
- was closest to expected skewness (Z-score 0.02).
- We found positive kurtosis of distribution most significantly in RSOL with highest mean
- deviation (Z-score -3.47), followed by CoA- (Z-score -2.86) and our group of other
- 260 major lesions (Z-score -1.97). HLHS and CoA+ showed positive kurtosis with almost
- one SD from expected results (Z-Scores -0.98), while suspected VR (Z-score -0.73)
- and TGA (Z-Score -0.27) showed similar kurtosis of distribution compared to our
- 263 control cohort.
- 264 All placentas (19/69, 28%) with individual z-scores > ± 3 for any of calculated results
- were individually reviewed for correlation with clinical confounders, while we found
- 266 MVM in two placentas and SGA complicating one pregnancy, overall numbers did not
- reach statistical significance.
- Mean and median deviation form GP (Z-score) for all cohorts and measured values are
- 269 listed in Table 2. Individual heterogeneity is demonstrated in histograms of occurrence
- 270 fraction of T2\* values in all voxels within the individual placenta from all participants in
- Figure 4. These histograms are color-coded by results of genetic testing in
- 272 Supplemental Figure 1.
- Maternal BMI at scan did not have an effect on mean  $T2^*$  values ( $R^2 = 0.004$ ). Maternal
- lie supine or left lateral, 30 and 7 scans respectively, had no significant effect on mean
- 275 T2\* values in our control group (p = 0.44). Placental position dichotomised in mostly
- anterior or posterior was also not associated with mean T2\* values (p = 0.98). There
- was no linear correlation with mean  $T2^*$  Z-scores at scan and weight ( $R^2 = 0.13$ ) or
- 278 head circumference at birth ( $R^2 = 0.11$ ).

279	DISCUSSION
280	In this study we have shown for the first time a comprehensive approach to placental
281	tissue characterisation in CHD. Using T2 relaxometry (T2*) and Gaussian Process
282	Regression (GPR) we were able to provide standard deviations for a range of placenta
283	metrics across various CHD groups from predicted values derived by our model trained
284	on a control group. Mean T2* values over the entire placenta may not represent
285	regional differences adequately and can lead to misinterpretation of imaging findings.
286	Therefore, we also used histograms and evaluated skewness and kurtosis to show
287	specific pattern of distribution depending on CHD subtype, which in turn may represent
288	pathophysiological substrates.
289	Requiring minimal acquisition time (<1min) and minor modification to clinical scan
290	protocols, these baseline placental assessments have been included into all our clinical
291	fetal MRI scans.
292	While mean T2* deviation for RSOL (Z-score 2.30) is comparable with our group of
293	other cardiovascular lesions (Z-score 2.31), the latter shows much less deviation from
294	calculated normal distribution as described in skewness and kurtosis (Z-scores -1.75
295	and -1.97, respectively). Strikingly, right-sided obstructive lesions (RSOL) show highest
296	deviation from expected normal distribution represented in skewness (Z-score -4.69)
297	and kurtosis (Z-score -3.47).
298	Our findings of abnormal placental imaging appear CHD lesions specific. One might
299	speculate that common intrinsic developmental pathways of placenta and fetal heart
300	may play a larger role, warranting further investigations towards understanding the
301	pathophysiology. Future investigations in conjunction with assessments of fetal
302	circulation in the presence of CHD, a flourishing field in both ultrasound and MRI
303	research [33,34], may allow a more detailed interpretation of our results.
304	Given the severe implications after birth, the cohort of HLHS appears to have
305	surprisingly limited deviations from control in placental structure and function. This
306	suggests less co-dependence between the development and or antenatal effects of
307	this anomaly and the development and function of the placenta also requiring further
308	research.
309	Our findings are in keeping with data from a very large Danish cohort from Matthiesen
310	et al. also depicting an association of RSOL, but not left outflow tract obstructions
311	including HLHS, aortic valve stenosis, and coarctation of the aorta, with lower placenta
312	weight [35]. Moreover Llurba et al. have suggested an imbalance in maternal and fetal

313	angiogenic factors may contribute to CHD and placental dysfunction most marked in
314	cohorts other than left sided lesions [36].
315	Fetuses with antenatal suspicion of CoA without confirmation after birth (CoA-) were
316	purposely not classified as controls in the design of this study. Firstly, presentation of
317	CoA may present several weeks (up to a year) after closure of the arterial duct [37].
318	None of the infants included here showed signs of coarctation at the time of manuscript
319	preparation. Secondly, there is a growing evidence that the observed ventricular
320	asymmetry is a feature of abnormal loading conditions leading to abnormal myocardial
321	deformation in fetuses with suspected coarctation [38]. In turn our CoA cohort spreads
322	relatively widely through the range of values provided by our placental assessments
323	(Supplemental Figure 2), which may reflect the wide clinical spectrum of CoA+
324	observed after birth, but also may indicate previously underexplored CHD entities
325	some of which do not require an extensive surgical arch repair within the first year of
326	life.
327	Studies of placental dysfunction and vascular malperfusion have shown altered
328	baseline conditions on assessment with T2 relaxometry accounting for a higher relative
329	response to maternal hyperoxygenation [15]. In our study we show similar altered
330	placental baseline conditions in case of fetal CHD. This is a critical insight to
331	understand the reported effects of short term maternal hyperoxygenation in CHD,
332	which are being explored to improve fetal oxygenation in CHD [39].
333	Altered placental baseline functions are of particular importance to understand the
334	complex intertwined fetomaternal environment in CHD and will also be essential to
335	understand further approaches on antenatal intervention such as maternal
336	hyperoxygenation [40].
337	Furthermore, structural assessments based on T2-weighted, and more recently also T1
338	imaging, depict changes over GA encompassing advanced lobulation with varied
339	lobule sizes, higher granularity, and substantial areas of low-signal intensity and an
340	increasing microstructural heterogeneity [41]. These changes are pronounced in FGR
341	or PE compared to healthy controls [22,42]. Although we did not report weighted MR
342	imaging in this study, observations from our T2* maps of increasing lacunarity over GA
343	is in keeping with the literature for our control and CHD cohort. Moreover, our CHD
344	cohorts also present higher lacunarity compared to healthy controls, most pronounced
345	in RSOL (Z-score -1.70).
346	As previously reported factors such as maternal age, BMI, fetal sex, parity, mode of
347	delivery and placental location were not correlated with T2* once corrected for GA [28].

348	Limitations of this study
349	Although we are based at a major referral centre for congenital heart disease in the
350	UK, this was a single centre study, with some patients delivering outside of our
351	hospital, having been referred for antenatal imaging. Our CHD groups are of moderate
352	sample size and may influence statistical significance.
353	Our placental scan protocol consistent of one single timepoint, therefore temporal
354	variance in placental T2* measurements, as recently suggested by other groups, might
355	not be accounted for in our data [43]. Recent literature describes also the possible
356	influence of maternal or placental position on T2* signals in higher field MRI [44]. While
357	we have scanned all our CHD patients in left lateral position, controls were also
358	scanned supine, resulting in higher statistical variance, 371.6 vs 776.5 respectively, but
359	not reaching statistical significance.
360	While in-utero MRI provides an excellent opportunity to provide early, in-vivo evidence
361	for specific placental phenotypes associated with CHD and hence a window of
362	observation into the intertwined relation between developing heart and placenta, it
363	cannot directly answer the question on causality and order of events.
364	Despite recent attempts to standardize definitions for FGR/IUGR, there is currently no
365	universal definition, which would include fetuses with congenital malformations such as
366	CHD [45]. Furthermore, the recent consensus-based definition of growth restriction on
367	the newborn excludes congenital and chromosomal abnormalities specifically, although
368	stating it may be applicable for this group due to the lack of any other option [32].
369	Previous literature, including a large population of 924,422 cohorts with over 5500 CHE
370	cases from Denmark also suggests an association of some CHD subtypes with lower
371	birth weight and head circumference [46]. Therefore, we did not exclude any
372	participants in our CHD cohort solely due diagnosis for FGR/IUGR or SGA.
373	Placental histopathology results were only available in 25% of all our assessments and
374	in CHD patients only with a specific clinical question at time of birth, such as maternal
375	pyrexia. Statistical analysis of features of MVM or fetal vascular malperfusion (FVM) or
376	Chorioamnionitis could therefore not be included. Similarly, only two placentas showed
377	infarcts in histopathology. These could be identified on the in-vivo imaging but did not
378	significantly alter the observed whole placental mean T2* results and have therefore
379	not been factored in the statistical analysis. Previous literature from postnatal placental
380	histopathology describe increased findings of FVM and MVM in the presence of
381	preeclampsia as well as CHD supporting the hypothesis of similar etiopathogenetic
382	factors contributing to the development of placental malformation and CHD [6].

383 While T2\* values are an indicator of oxygen concentration, direct measurement was 384 not possible in our setting. Oxygen-haemoglobin dissociation curves derived from MRI 385 in animal models might be available for future studies to allow close estimation of 386 actual oxygen content in the fetoplacental circulation [47,48]. 387 Including only five pregnancies with confirmed fetal chromosomal abnormalities in our 388 cohort did not allow for further statistical analysis of potential associations. This study 389 did not include volumetric assessments of the placenta given echo planar imaging is 390 associated with geometric distortions. 391 Future studies applied to CHD cohorts with more complex and time-consuming scan 392 protocols, including diffusion-weighted imaging or texture analysis using T1 and T2 393 weighted methods, may allow modelling of fetal and maternal peculiarities in circulation 394 and disentangle tissue characteristics and flow pathophysiology, as recently in CHD 395 cohorts [41,49-51]. Slator et al. have recently published a study with a focused 396 extensive research protocol on the microenvironments within the placenta depicting 397 heterogenous compartments from maternal and fetal side [52]. Using Velocity-398 Selective Arterial Spin Labelled MRI Zun et al. also reported that global placental 399 perfusion significantly decreased and regional variation of placental perfusion 400 significantly increased over GA in fetuses with CHD [53,54].

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### Conclusion

This study describes in vivo differences in placental tissue phenotypes in healthy controls and fetuses with antenatal diagnosis of congenital heart disease based on T2 relaxometry. Using machine learning we depict unique features of T2\* value distribution and their standard score from our normal cohort for a wide range of major cardiac lesions, providing information on placental dysfunction complementary in the antenatal assessment of CHD.

409	Acknowledgements
410	The authors acknowledge all clinical and scanning team involved in these projects and
411	we are immensely grateful to the patients and volunteers recruited for this project.
412	Funding
413	This work was supported by the UKRI Future Leaders Fellowship [MR/T018119/1], the
414	Wellcome/EPSRC Centre for Medical Engineering [WT 203148/Z/16/Z], the Wellcome
415	Trust IEH Award 102431 (iFIND project), the NIH Human Placenta Project grant
416	1U01HD087202-01 (Placenta Imaging Project (PIP)), the Wellcome Trust Sir Henry
417	Wellcome Fellowship [201374/Z/16/Z] and by the National Institute for Health Research
418	(NIHR) Biomedical Research Centre based at Guy's and St Thomas' NHS Foundation
419	Trust and King's College London. The views expressed are those of the authors and
420	not necessarily those of the NHS, the NIHR or the Department of Health.
	Declaration of commeting interests
421	Declaration of competing interests
422	The authors declare that they have no competing interest.
	Marriage
423	Keywords
424	Congenital Heart disease (CHD), Placenta, Magnetic resonance imaging (MRI), T2*
425	Mapping, Gaussian Process Regression (GPR), Bayesian non-parametric regression,
426	Machine learning

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# 669 Tables/figures

	Median	IQR	Min	Max	р
Control cohort (n=37)					
Maternal age at scan (a)	34.65	4.65	27.69	44.03	
Maternal BMI at scan (kg/m²)	30.03	8.67	18.75	42.59	
Gestational age at scan (w)	30.86	8.36	19.57	37.86	
Fetal sex (female/total)	0.62				
Gestational age at birth (w)	40.14	2.14	38.14	42.14	
Birth weight (g)	3583	534	2780	4450	
BW Z-score	0.9558	1.2147	-1.0550	2.1966	
BW Centile	83.04	39.16	14.57	98.60	
CHD cohort (n=69)					
Maternal age at scan (a)	32.63	8.39	18.60	40.79	0.002†
Maternal BMI at scan (kg/m²)	28.39	5.63	18.81	40.35	0.094†
Gestational age at scan (w)	31.57	2.64	28.43	36.29	0.069†
Fetal sex (female/total)	0.58				0.836
Gestational age at birth (w)	38.71	1.57	32.71	41.71	<0.001
Birth weight (g)	3080	964	1610	5150	<0.001
BW Z-score	-0.1230	1.7762	-2.9547	3.7986	0.003
BW Centile	48.20	62.20	0.16	99.99	0.003
Diagnosis	N	%			
Left-sided obstructive lesions	10	14.5	HLHS	HLHS	
	13	18.8	COA- (susp	COA- (suspected, not confirmed)	
	8	11.6	COA+ (con	COA+ (confirmed neonatal)	
Right-sided obstructive lesions	10	14.5	TOF, PA, PS, TA, Ebstein anomaly		
Disorders of mixing	5	7.2	TGA		
Suspected vascular rings	17	24.6	DAA, RAA (+/- ALSA)		
Other major lesions	Other major lesions 6 8.7 CAT, PAPVD, CR				

Table 1: Maternal and pregnancy characteristics.

 $(\dagger)$  For statistical analysis controls were restrained to GA range at scan (28 – 37

672 weeks, N=23).

Mean deviation from GP [Z-score] +/- SD of Z-scores within group									
	Control	CHD	HLHS	COA-	COA+	RSOL	TGA	Other	VR
N	9	69	10	13	8	10	5	6	17
Mean T2*	-0.56	0.93*	0.64	1.39*	0.24	2.30***	-0.11	2.31**	0.09
	+/-1.34	+/-2.26	+/-2.03	+/-2.91	+/-1.34	+/-2.07	+/-1.86	+/-2.01	+/-2.06
Skewness	-0.08	-1.99*	-1.12	-3.83	-0.66	-4.69*	0.02	-1.75**	-0.80
	+/-0.59	+/-4.27	+/-1.58	+/-7.18	+/-0.73	+/-5.56	+/-0.73	+/-1.25	+/-2.70
Kurtosis	-0.53	-1.67	-0.98	-2.86*	-0.98	-3.47**	-0.27	-1.97**	-0.73
	+/-1.01	+/-2.34	+/-1.39	+/-3.25	+/-0.67	+/-3.02	+/-1.50	+/-0.93	+/-1.66
Lacunarity	-0.15	-0.54	-0.18	-1.02	0.26	-1.70*	-0.01	-1.26	0.06
	+/-1.27	+/-1.55	+/-1.21	+/-1.69	+/-0.53	+/-1.99	+/-0.98	+/-2.35	+/-0.93
Median devi	Median deviation from GP [Z-score] (IQR within group)								
	Control	CHD	HLHS	COA-	COA+	RSOL	TGA	Other	VR
N	9	69	10	13	8	10	5	6	17
Mean T2*	-0.78	0.70*	0.57	0.89*	0.25	2.36***	-0.41	2.10**	-0.59
	(2.05)	(2.93)	(2.37)	(4.05)	(2.06)	(3.32)	(3.68)	(3.77)	(2.76)
Skewness	0.01	-0.47*	-0.49	-0.96	-0.62	-2.12*	0.73	-1.89**	0.02
	(0.71)	(2.19)	(2.94)	(4.38)	(0.98)	(10.23)	(2.32)	(2.40)	(0.83)
Kurtosis	-0.80	-1.10	-0.75	-1.72*	-0.86	-2.15**	0.41	-1.83**	-0.63
	(1.67)	(1.95)	(2.07)	(3.35)	(1.04)	(5.64)	(2.65)	(1.83)	(1.23)
Lacunarity	0.01	-0.18	-0.09	-0.68	0.17	-1.43*	0.29	-0.37	0.05
	(1.95)	(1.67)	(1.28)	(2.49)	(0.82)	(2.22)	(1.84)	(3.84)	(1.35)

673 Table 2: Results Gaussian process Regression, groupwise.

Entire CHD cohort (CHD), Hypoplastic left heart syndrome (HLHS), Coarctation suspected not confirmed (CoA-), Coarctation confirmed operated (CoA+), Right-sided obstructive lesions (RSOL), Disorders of mixing (TGA), Other lesions, Suspected vascular rings (VR). Significance calculated by Independent t-test (Mean T2\*) or Mann-Whitney U (Skewness, Kurtosis, Lacunarity) in comparison to test control cohort, \*  $p \le 0.05$ , \*\*  $p \le 0.01$ , \*\*\*  $p \le 0.001$ .

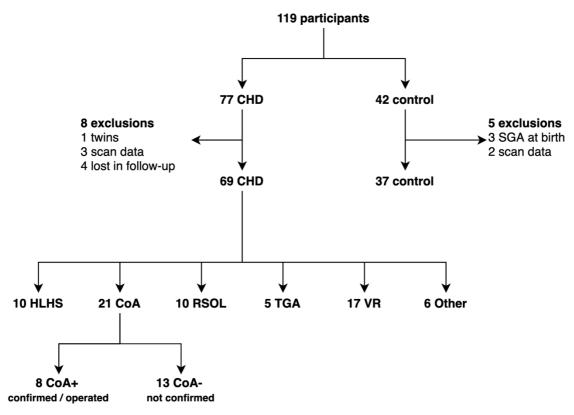


Figure 1 - Study cohort.

CHD subtypes classified by main antenatal diagnosis: Hypoplastic left heart (HLHS), coarctation of the aorta (CoA), right-sided obstructive lesions (RSOL), transposition of the great arteries (TGA), suspected vascular ring (VR) and other complex lesions. Controls were excluded for low birth weight (SGA) and insufficient scan data, pregnancies complicated by fetal CHD had to be excluded for twin pregnancy and loss in follow-up at time of analysis.

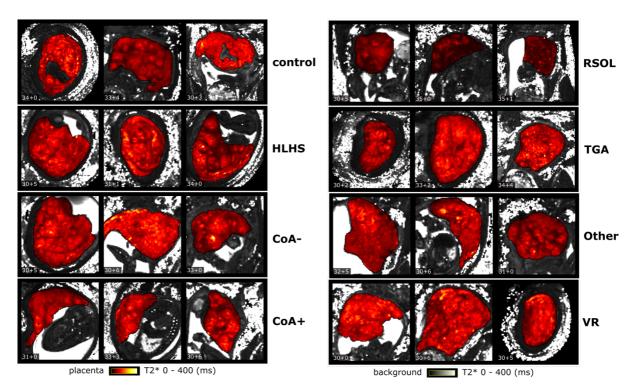


Figure 2: Qualitative Imaging

Illustration of mid-stack slice in coronal orientation from T2\* maps from 3 individual placentas for all included cohorts. Segmented placental parenchyma is highlighted in red/yellow scales.

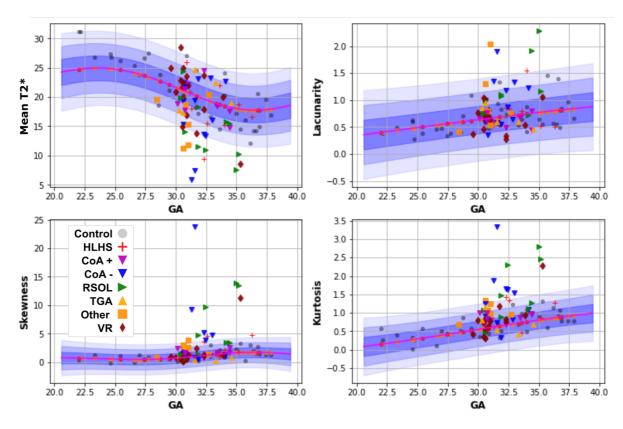


Figure 3 – Gaussian Process Regression results

The posterior probability for Mean T2\*, Lacunarity, Skewness and Kurtosis over GA for all considered controls in red, the corridors corresponding to Z-scores of 1, 2 and 3 are illustrated in blue. Cohorts: Control (test subset), Hypoplastic left heart syndrome (HLHS), Coarctation of the aorta confirmed (COA+) and not confirmed (COA-), right-sided obstructive lesions (RSOL), disorders of mixing (TGA), other major lesions and suspected vascular rings (VR).

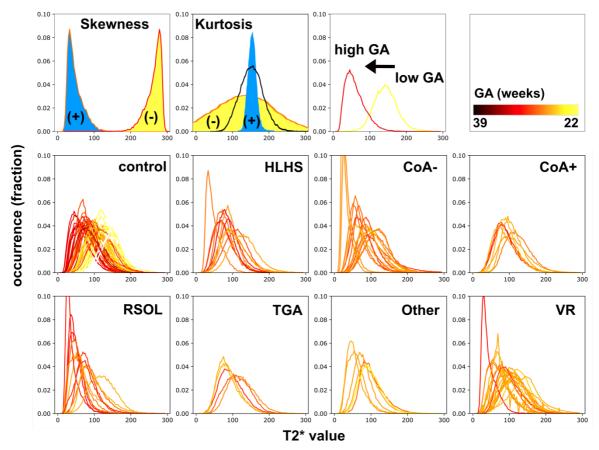
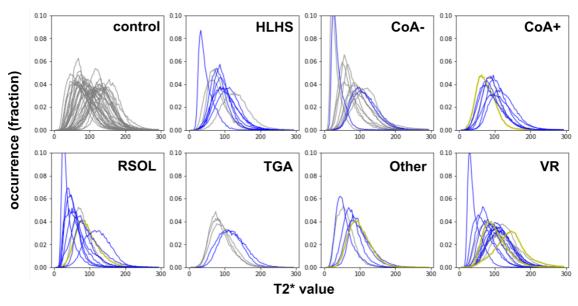


Figure 4 - Individual Histograms

Individual Histograms, derived from about 70000 voxels per subject binned in 100, depicting occurrence fraction of T2\* values over the entire placental volume from all participants, coloured by GA at scan. Groups from top left to bottom right. Control, Hypoplastic left heart syndrome (HLHS), coarctation of the aorta not confirmed (CoA-) and confirmed (COA+), right-sided obstructive lesion (RSOL), Transposition of the great arteries (TGA), Other lesions and suspected vascular ring (VR).



#### Supplemental Figure 1 – Histograms with genetic test results

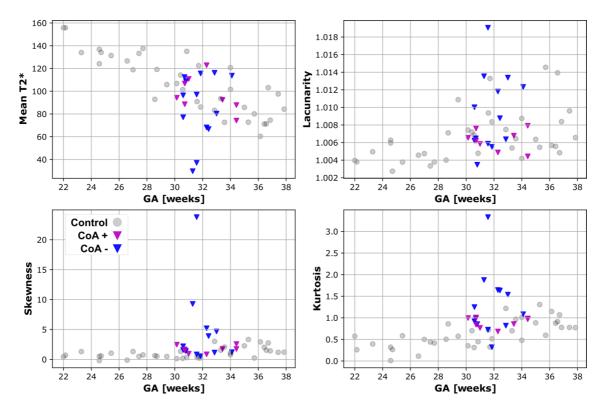
Individual Histograms, as shown in Figure 4. Color coded by results of genetic testing. Abnormal genetics (yellow), normal genetics (blue), not tested (grey), Hypoplastic left heart syndrome (HLHS), coarctation of the aorta not confirmed (CoA-) and confirmed (COA+), right-sided obstructive lesion (RSOL), Transposition of the great arteries (TGA), Other lesions and suspected vascular ring (VR).

7	1	8	

N (%)	Control	CHD
Available placental histology	15/37 (40%)	12/69 (17%)
MVM	1	2
With infarct	1	2
(Sub-)Chorionitis	8	7
With MIR	4	3
With FIR	5	4

### Supplemental Table 1 – Placental Histology

Table shows available placental histology results. Maternal vascular malperfusion (MVM), maternal inflammatory response (MIR), fetal inflammatory response (FIR).



Supplemental Figure 2 – Coarctation of the aorta cohort and controls

Mean T2\*, Lacunarity, Skewness and Kurtosis over GA for all included controls in grey,
confirmed Coarctation of the aorta (CoA+) in pink and not confirmed (CoA-) in blue.