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DOI:
[10.1016/j.placenta.2016.05.012](https://doi.org/10.1016/j.placenta.2016.05.012)

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Document Version
Peer reviewed version

Citation for published version (Harvard):
Dunn, W, Allwood, JW, Van Mieghem, T, Morris, RK, Mackie, F, Fox, C, Kilby, M & MacKie, F 2016, 'Carbohydrate and fatty acid perturbations in the amniotic fluid of the recipient twin of pregnancies complicated by twin-twin transfusion syndrome in relation to treatment and fetal cardiovascular risk', *Placenta*, vol. 44, pp. 6-12. <https://doi.org/10.1016/j.placenta.2016.05.012>

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Checked 1/6/2016

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Accepted Manuscript

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PII: S0143-4004(16)30107-2

DOI: [10.1016/j.placenta.2016.05.012](https://doi.org/10.1016/j.placenta.2016.05.012)

Reference: YPLAC 3420

To appear in: *Placenta*

Received Date: 5 May 2016

Revised Date: 23 May 2016

Accepted Date: 26 May 2016

Please cite this article as: Dunn WB, Allwood JW, Van Mieghem T, Morris RK, Mackie FL, Fox CE, Kilby MD, Carbohydrate and fatty acid perturbations in the amniotic fluid of the recipient twin of pregnancies complicated by twin-twin transfusion syndrome in relation to treatment and fetal cardiovascular risk, *Placenta* (2016), doi: [10.1016/j.placenta.2016.05.012](https://doi.org/10.1016/j.placenta.2016.05.012).

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Carbohydrate and fatty acid perturbations in the amniotic fluid of the recipient twin of pregnancies complicated by twin-twin transfusion syndrome in relation to treatment and fetal cardiovascular risk

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ACCEPTED MANUSCRIPT

Abstract

Introduction: Twin-twin transfusion syndrome (TTTS) complicates 15% of monochorionic twin pregnancies, often being associated with recipient cardiac dysfunction. Untreated, it has a fetal mortality rate of at least 90%; although treatment by fetoscopic laser coagulation significantly improves prognosis. Measurement of recipient amniotic fluid metabolites, such as cardiac Troponin T and atrial natriuretic polypeptide, correlate with cardiac function in this fetus. The aim of this study is to describe the amniotic fluid metabolomic profile in TTTS, relate this to fetal recipient cardiac function and assess the metabolomic changes induced by fetoscopic laser coagulation.

Methods: Prospective single centre cohort study. The metabolomics profile of the amniotic fluid from the recipient sac of TTTS pregnancies was assessed using ultra high performance liquid chromatography-mass spectrometry. Profiles were compared pre- and post-laser coagulation and related to fetal recipient cardiac function, as assessed using Doppler ultrasound within 6 hours of treatment.

RESULTS: Eleven metabolites had significant associations with recipient fetal right and left ventricular myocardial performance index pre-laser. 200 metabolites in recipient amniotic fluid demonstrated a change in relative concentrations when comparing pre- and post-laser coagulation ($p < 0.005$). The most prominent change is in the balance of carbohydrate and fatty acid metabolic profile contributing to fetal or placental energy metabolism. These changes were also associated with the echocardiographic measures of recipient cardiac function.

Discussion: Changes in carbohydrate and fatty acid metabolic profiles are noted in recipients with cardiac dysfunction, and further changes are noted after treatment.

Validation and investigation may identify targets for potential pharmacological treatment.

Key words: Echocardiography, fetoscopy, laser coagulation, metabolomics, monochorionic twins, twin-twin transfusion syndrome

Abbreviations:

ANF – atrial natriuretic factor

AVA – arteriovenous anastomoses

BNP – brain-type natriuretic factor

DV – ductus venosus

E/A – early passive/atrial

FLC- fetoscopic laser coagulation

IUGR – intrauterine growth restriction

LV – left ventricle

MCDA – monochorionic diamniotic

MPI – myocardial performance index

QC – quality control

RV – right ventricle

TTTS – twin-twin transfusion syndrome

UHPLC-MS - Ultra High Performance Liquid Chromatography-Mass Spectrometry

1 Introduction

2 Approximately 15% of monochorionic dichorionic (MCDA) twins are complicated by
3 twin-twin transfusion syndrome (TTTS), a condition associated with unidirectional intertwin
4 blood flow through placental arteriovenous anastomoses (AVA) and high perinatal mortality.
5 This leads to the severe haemodynamic imbalances seen within the fetal circulations of this
6 condition with a hypertensive circulation in the “recipient” twin and subsequent cardiac
7 dysfunction (1), noted in up to 70% of pregnancies. There is corresponding dysregulation of
8 fetal endocrine systems, including the renin-angiotensin-aldosterone (2), atrial natriuretic
9 factor (ANF) (3) and Endothelin-1 systems (4). Such changes have been measured in the
10 amniotic fluid of the recipient twin, and related to cardiac dysfunction and overall fetal
11 prognosis (3).

12 Untreated, fetal mortality is at least 90%, for each fetus (5). Treatment by fetoscopic
13 laser coagulation (FLC) is effective with survival of two babies approaching 60% (6). FLC
14 also reduces the risk of neurodevelopmental morbidity compared to other treatments (7).
15 Improvement in recipient cardiac function is noted within 48 hours of FLC in approximately
16 half of recipients (8), which is associated with improved fetal outcome (9).

17 Metabolomics is the holistic untargeted study of metabolism (10) and investigates the
18 final downstream product of genotype-environment interactions. It provides the identification
19 of a dynamic and sensitive phenotypic signature associated with human health ageing and
20 disease molecular pathophysiology (10). In human pregnancy, this technique has been
21 used to investigate intrauterine growth restriction (IUGR) (11-14), pre-eclampsia (15-17)
22 and gestational diabetes (18, 19). In non-pregnant adults, it has been used to profile
23 functional and metabolic changes associated with heart failure (20, 21).

24 In MCDA pregnancies affected by TTTS, a small number of studies, targeting
25 specific areas of metabolism have been reported (22, 23). Molecular patho-mechanistic
26 changes have been observed in angiogenic growth factors (24), cytokine levels (25) and
27 gene transcripts in amniotic fluid and maternal plasma which appear to predict fetal
28 outcome (26, 27).

29 We describe the metabolomic profiles in amniotic fluid from the recipient sac of
30 MCDA twin pregnancies complicated by severe TTTS and note associations between fetal
31 echocardiographic recipient right ventricular (RV) and left ventricular (LV) function. In
32 addition, the effects of FLC on the metabolomic profile signatures are described.

33 **Methods**

34 This study had ethical approval from Birmingham Black Country Local Research Ethics
35 Committee (No: 06/Q2702/71 accepted in 2006) with written consent obtained from all
36 subjects.

37 ***Patient selection***

38 The cohort consisted of MCDA twins complicated by severe TTTS (presenting before 24
39 weeks) treated between August 2011 – June 2012 and TTTS was defined as
40 polyhydramnios (>8cm in the deepest vertical pocket of the recipient at <20 weeks of
41 gestation or >10cm from 20 weeks of gestation onwards) in combination with
42 oligohydramnios in the donor (<2cm deepest vertical pool depth). All cases were
43 prospectively staged using the Quintero system (28).

44 ***Cardiac function assessment***

45 High-resolution fetal ultrasound and echocardiography were performed in the recipient with
46 curvilinear array transducers (7–3.5 MHz) on a Siemens S3000 ultrasound machine

47 (Siemens Ltd, Erlangen, Germany) by a single operator (MDK) and the myocardial
48 performance index (MPI) calculated for each ventricle (1, 29) as previously described.
49 Cardiac dysfunction was indicated by the presence of tricuspid regurgitation, reversed flow
50 in the DV during atrial contraction, and a tricuspid early passive/atrial contraction (E/A) ratio
51 of >95% CI outside the normal limits. This was performed within 4 hours of starting the laser
52 procedure and repeated with 6 hours post-FLC.

53 Fetal cardiac function was assessed 24 hours prior to FLC and repeated within 4 hours
54 post-FLC.

55 ***Fetoscopic laser coagulation (FLC)***

56 FLC was performed using local anaesthesia (1% lignocaine skin/myometrial infiltration) and
57 maternal Remifentanil sedation as previously described (30). A selective sequential FLC
58 technique was used, with an additional “Solomon” procedure in some cases (31). No
59 amnio-infusion was performed.

60 ***Non-targeted UHPLC-MS metabolomics analysis***

61 All solvents and chemicals applied were of HPLC analytical grade (J.T. Baker, U.K.).

62 ***Sample collection and preparation.*** A 10ml sample of amniotic fluid was taken at
63 insertion of the fetoscope into the recipient amniotic sac and then a further 10ml sample
64 withdrawn at the end of the laser coagulation treatment. The median duration of the laser
65 procedure and amniodrainage the median duration was 34 (range 21 - 45) minutes.
66 Samples were stored at -80°C before preparation and analysis. All samples were
67 randomised to ensure no correlation between order of preparation and subject, disease
68 grade or date of sample collection. Deproteinisation was performed as described below.
69 250µL of amniotic fluid was vortex-mixed with 1000µL of methanol for 15 seconds to
70 precipitate proteins and DNA followed by centrifugation (15 minutes, 13,000 g) drying to

71 induce metabolite stability and then stored at -80°C prior to analysis. A pooled quality
72 control (QC) sample was prepared by combining $80\mu\text{L}$ aliquots of each of the 38 samples
73 (32).

74 **Ultra High Performance Liquid Chromatography-Mass Spectrometry (UHPLC-MS)**

75 **analysis.** UHPLC-MS analysis of amniotic fluid extracts and QC samples was performed
76 applying a Dionex U3000 coupled to an electrospray LTQ-FT-MS Ultra mass spectrometer
77 (Thermo Scientific Ltd. UK). Samples were reconstituted in $100\mu\text{L}$ of 50:50 methanol:water,
78 vortex-mixed for 15 seconds, centrifuged (15 minutes, $13,000\text{ g}$) and transferred to vials
79 with $200\mu\text{L}$ fixed inserts (Thermo-Fisher Ltd. U.K.). All samples were stored in the
80 autosampler at 5°C and analysed separately in negative and positive electrospray
81 ionisation (ESI) modes within 72 h of reconstitution. UHPLC separations were performed
82 applying a Hypersil Gold C_{18} reversed phase column ($100 \times 2.1\text{mm}$, $1.9\mu\text{m}$) at a flow rate of
83 $400\mu\text{L}\cdot\text{min}^{-1}$, column temperature of 40°C and with two solvents: solvent A (HPLC grade
84 water + 0.1% formic acid) and solvent B (HPLC grade methanol + 0.1% formic acid). A
85 gradient elution was performed as follows: hold 100% A 0-1.5 min, 100% A - 100% B 1.5-6
86 min curve 3, hold 100% B 6-12 min, 100% B – 100% A 12-13 min curve 3, hold 100% A 13-
87 15 min. Injection volume was $5\mu\text{L}$. UHPLC eluent was introduced directly in to the
88 electrospray LTQ-FT Ultra mass spectrometer with source conditions as follows: spray
89 voltage -4.5 kV (ESI-) and $+5\text{ kV}$ (ESI+), sheath gas 30 arbitrary units, aux gas 15 arbitrary
90 units, capillary voltage 35 V , tube lens voltage -100 V (ESI-) and $+90\text{ V}$ (ESI+), capillary
91 temperature 280°C , ESI heater temperature 300°C . Data were acquired in the FT mass
92 spectrometer in the m/z range 100-1000 at a mass resolution of 50,000 (FWHM defined at
93 m/z 400), with a scan speed of 0.4 sec and an AGC setting of 1×10^6 . Analysis order was
94 composed of 10 QC sample injections for system conditioning followed by a QC sample
95 injection every 6th injection with two QC sample injections at the end of the analytical run.

96 Amniotic fluid extracts for each subject were analysed in a random order; the two samples
97 for each subject were analysed sequentially.

98 **Data pre-processing.** UHPLC-MS raw data profiles were first converted into a NetCDF
99 format within the Xcalibur software's File Converter program. Each NetCDF based three-
100 dimensional data matrix (intensity \times m/z \times retention time – one per sample) was converted
101 into a vector of peak responses, using the freely available XCMS software as described
102 previously (33). Data were exported from XCMS as a .csv file for further data analysis.
103 Metabolite annotation was performed applying the PUTMEDID_LCMS workflow as
104 previously described (34). All metabolite annotations are reported at level 2 (putatively
105 annotated compounds) according to MSI reporting standards (35). In cases where a single
106 metabolite is detected as multiple metabolite features (as described previously (36)), only a
107 single feature is reported chosen as having a p-value nearest to 0.05).

108 **Univariate and multivariate analysis.** Intergroup comparisons for continuous variables
109 with a non-parametric distribution were made using the Mann-Whitney U test to determine
110 significant differences between the data sets. Median values and 95% confidence intervals
111 (CI) are described. Categorical data were analysed using Fisher's exact test and odds
112 ratios (OR) and 95%CI. Significance was taken as $p < 0.05$. Metabolomics processed data
113 were analysed in 'R' applying the unsupervised multivariate principal components analysis
114 (PCA), supervised multivariate Partial Least Squares-Discriminant Analysis (PLS-DA),
115 univariate non-parametric Wilcoxon Signed Rank test and Spearman rank correlation
116 analysis. The fold change (median peak area before treatment/median peak area after
117 treatment) was calculated including 95%CI. Metabolites were manually clustered into
118 classes defining similar chemical structure or metabolic pathway to identify biologically
119 relevant and robust metabolic changes.

120 **Results**

121 **Cohort of monochorionic twins complicated by TTTS and the effects of treatment**

122 Table 1 summarises the baseline demographic data of the whole cohort (n=19) at
123 the time of diagnosis. In the total cohort of MCDA twins complicated by TTTS, 5.2% (1/19)
124 had Quintero Stage-I, 15.8% (3/19) Stage-II, and 79% (15/19) Stage-III disease. The
125 measured RV and LV MPI was elevated (>95%CI for gestation) in 89.4% (17/19) and
126 73.7% (14/19) of recipients, respectively (Figure 1a and b). Of the two cases which had a
127 RV MPI within 95%CI for gestation, one twin set had Quintero stage-I and one Quintero
128 stage-III. In the 5 recipient fetuses with LV MPI within 95%CI, all had Quintero stage >II. In
129 the recipient twin, additional features of cardiac dysfunction are described in Table 1.

130 FLC was performed at a median gestation of 20 weeks and 2 days (142 days)
131 (95%CI 137.7 – 148.3). Recipient RV MPI (p=0.02) and LV MPI (p=0.03) decreased
132 significantly after FLC (Figure 2a and b).

133 **Amniotic fluid metabolome of the recipient amniotic fluid**

134 38 paired amniotic fluid samples were collected before and after FLC. Following
135 quality assurance of the data, 2694 and 1510 metabolite features remained in positive and
136 negative ion modes, respectively.

137

138 **Correlation analysis of metabolic profiles and recipient cardiac function in TTTS**

139 Spearman rank correlation analysis was performed to identify associations between
140 recipient RV and LV MPI at the time of diagnosis, and the concentration of each metabolite.
141 A summary of the results demonstrating associations between recipient RV or LV MPI and
142 amniotic fluid metabolites are shown in Table 2 and the full data are shown in

143 Supplementary Tables 2 and 3. The 118 and 102 metabolites identified as having an
144 association with LV MPI or RV MPI each showed a positive correlation (denoted as '+'
145 showing that as LV MPI or RV MPI increase so does the metabolite concentration) or a
146 negative correlation (denoted as '-' showing that as LV MPI or RV MPI increase the
147 metabolite concentration decreases). 15 metabolites showed an association with LV MPI
148 and RV MPI. Acyl carnitines, acyl glycerides, fatty acids and oxidised fatty acids were all
149 negatively correlated with cardiac function whereas carbohydrates were positively
150 correlated with cardiac function. Ceramides, sphingolipids and glycerophospholipids were
151 also negatively correlated with cardiac function and may be related to changes in cell
152 membranes. Hormones were negatively correlated with cardiac function. Finally, for LV
153 MPI, two oxidative phosphorylation metabolites were negatively correlated with cardiac
154 function but no metabolites were correlated with the RV MPI data.

155

156 **Comparison of metabolic changes before and after laser treatment independent of** 157 **type of laser treatment**

158 200 metabolites showed a statistically significant change ($p < 0.005$) in relative
159 concentrations when comparing amniotic fluid samples taken before laser, and after laser
160 treatment; this was unaffected by whether or not the Solomon technique was applied.
161 Supplementary Table 1 lists the significantly altered metabolomic profiles (all metabolites
162 are grouped into classes of chemical structure or metabolic pathway). Figure 3
163 demonstrates these data as a "heat map" of relative concentration changes for all
164 metabolite classes containing three or more metabolites. There were 13 metabolite
165 "classes" consisting of ≥ 3 metabolites which demonstrated significant fold changes. These
166 include acyl carnitines, acyl glycerides, amino acid metabolism, carbohydrate metabolism,

167 cholesterol esters, ceramides and sphingolipids, fatty acid metabolism,
168 glycerophospholipids, haem metabolism, nucleosides, oxidised fatty acids, oxidative
169 phosphorylation (electron transport chain) and thyroid/steroid hormone metabolism. These
170 suggest a change in fetal or placental energy metabolism, specifically between
171 carbohydrates (shown by higher levels of carbohydrates before treatment) and fatty acids
172 (shown by higher levels of acyl carnitines, acyl glycerides, fatty acids, oxidised fatty acids
173 and TCA/oxidative phosphorylation metabolites) in amniotic fluid post-FLC. Amniotic fluid
174 oxidised fatty acid concentration shows a 50-fold change (higher concentration post-FLC).
175 This is potentially a prostaglandin derivative known to have vasodilatory roles within the
176 endothelium (37). Of five acyl carnitines, it is the medium chain derivatives (hexanoyl,
177 octanoyl and decanoyl) that predominate. Also changes related to oxidative
178 phosphorylation were noted (Supplementary Table 1). Concentrations of thyroid and steroid
179 hormones were decreased post-FLC potentially related to an attenuated “stress” response
180 post-FLC treatment.

181 These observed changes may be secondary to placental destruction caused by
182 coagulation (38), changes in recipient cardiovascular function post-laser coagulation or a
183 combination of these events. Finally, uridine/pseudouridine was observed to be 20% higher
184 before treatment and these metabolites have been observed to be increased in heart failure
185 in adults (39).

186

187 **Discussion**

188 This cohort study demonstrates that the metabolomic profile in the amniotic fluid of
189 recipient monozygotic twins complicated by TTTS is different when there is significant
190 cardiac dysfunction, and that amniotic fluid metabolic profiles change in response to FLC.

191 Following FLC treatment, a wide range of different metabolite classes are perturbed in
192 amniotic fluid in the recipient sac. The balance between carbohydrates and fatty acid for
193 energy production appears to change following treatment to be preferential for fatty acids.
194 The balance between carbohydrate and fatty acid metabolism appears to be related to
195 recipient cardiac function (as measured by LV and RV MPI) as fatty acids are negatively
196 correlated and carbohydrates are positively correlated with cardiac function. However, it is
197 difficult to delineate whether the source of the metabolites is fetal or placental, as amniotic
198 fluid is a composite of placental metabolite secretion and fetal metabolite secretion (e.g. as
199 urine). As amniotic fluid sampling was repeated after a median time of 34 minutes, it is
200 more probable that these changes are secondary to a combination of trophoblast/vessel
201 destruction (as previously noted by an elevation in alpha-fetoprotein and human chorionic
202 gonadotrophin) (38) and fetal cardiovascular change.

203 In this relatively small cohort study a significant proportion of the recipient fetuses
204 had evidence of cardiac dysfunction and an elevated LV and RV MPI. Such data are
205 consistent with those previously described in the literature (3) and that recipient RV and LV
206 MPI rapidly alters post-FLC (40). Much of the recipient cardiac effects are due to RV and LV
207 diastolic dysfunction or hypertension (3).

208 In adults with cardiac failure, prediction of survival is optimal when cardiac ultrasound
209 measurements are combined with serum-derived biomarkers (41). Classically, the
210 combination of cardiac Troponin T, atrial natriuretic polypeptide (ANP) and BNP is used as
211 a marker to reflect cardiac dysfunction and a combination is better than one peptide alone.
212 In recent years there has been a move to a metabolomic approach for profiling functional
213 and metabolic changes in adults with heart failure (21).

214 In the fetal setting, determination of these peptides within fetal plasma would require
215 fetal blood sampling which would carry significant procedure-related complications.
216 However, therapy for TTTS by FLC involves the removal of redundant amniotic fluid from
217 the recipient's sac after completion of the procedure. At this gestation, amniotic fluid is
218 composed mainly of fetal urine and is therefore a potential source for fetal cardiac
219 biomarkers. The presence of cardiac Troponin T in amniotic fluid has previously been
220 described in severely growth restricted fetuses (42) and also in recipient fetuses with fetal
221 cardiac dysfunction in TTTS (3). Also, in pregnancies complicated by TTTS, ANP and brain-
222 type natriuretic peptide (BNP) have been identified in the amniotic fluid (4, 43) and have
223 also been noted to be associated with fetal recipient cardiac dysfunction in TTTS (3).

224 15 metabolites in the amniotic fluid of the recipient sac prior to laser coagulation
225 demonstrated an association with LV MPI and RV MPI. Acyl carnitines, acyl glycerides, fatty
226 acids and oxidised fatty acids were all negatively correlated with cardiac function whereas
227 carbohydrates were positively correlated with cardiac function. This implies that the balance
228 between using carbohydrates and fatty acids as substrates for energy metabolism has an
229 influence on fetal cardiac function, though causality still requires testing. This balance has
230 been shown to be important in cardiac diseases including hypertrophic cardiomyopathy
231 (44). Ceramides, sphingolipids and glycerophospholipids were also negatively correlated
232 with cardiac function and may be related to changes in cell membranes. Hormones were
233 negatively correlated with cardiac function. Finally, for LV MPI, two oxidative
234 phosphorylation metabolites were negatively correlated with cardiac function but no
235 metabolites were for the RV MPI data. Of interest and worthy of further study also are the
236 changes in N,N-dimethylarginine and the structurally similar N,N-diacetylspermine. N,N-
237 dimethylarginine is a known inhibitor of NO synthesis from arginine and therefore reduces

238 vasodilation. Symmetrical and asymmetrical N,N-dimethylarginine are implicated in cardiac
239 function and cardiovascular health (45, 46).

240 There were also 13 metabolite “classes” consisting of ≥ 3 metabolites which
241 demonstrated significant fold changes in amniotic fluid after FLC. These include acyl
242 carnitines, acyl glycerides, amino acid metabolism, carbohydrate metabolism, cholesterol
243 esters, ceramides and sphingolipids, fatty acid metabolism, glycerophospholipids, haem
244 metabolism, nucleosides, oxidised fatty acids, oxidative phosphorylation (electron transport
245 chain) and thyroid/steroid hormone metabolism. These suggest a change in fetal or
246 placental energy metabolism, specifically between carbohydrates (shown by higher levels of
247 carbohydrates before treatment) and fatty acids (shown by higher levels of acyl carnitines,
248 acyl glycerides, fatty acids, oxidised fatty acids and TCA/oxidative phosphorylation
249 metabolites) in amniotic fluid post-FLC. Similar changes in the balance of carbohydrate and
250 fatty acid usage for energy production in hypertrophic cardiomyopathy have been
251 previously reported (44). In TTTS, amniotic fluid levels of BNP appear to correlate with the
252 severity of recipient cardiac dysfunction (3). The higher levels of fatty acids and lower levels
253 of carbohydrates post-FLC suggest that there is a switch from fetal or placental use of fatty
254 acids as precursors for energy metabolism to carbohydrates.

255 Amniotic fluid oxidised fatty acid concentration shows a 50-fold change (higher
256 concentration post-FLC). This is potentially a prostaglandin derivative known to have
257 vasodilatory roles within the endothelium (37). Of five acyl carnitines, it is the medium chain
258 derivatives (hexanoyl, octanoyl and decanoyl) that predominated suggesting a specific
259 perturbation in medium chain fatty acid oxidation.

260 Two published studies (from the same group) in singleton pregnancies (47, 48) have
261 investigated the amniotic fluid metabolome. These data noted 70 metabolomic compounds

262 using $(1)H$ NMR. Pregnancies complicated by a heterogeneous group of fetal
263 malformations demonstrated “variations in glucose, some amino acids and organic acids
264 and proteins”. However, it is recognised that this group is heterogeneous for malformations
265 and thus it was difficult to draw conclusions. The amniotic fluid samples were taken by
266 amniocentesis at a wide gestational range (13 – 42 weeks) in these singleton pregnancies.
267 Subjects with gestational diabetes showed an average increase in glucose and small
268 decreases in several amino acids along with acetate, formate, creatinine, and
269 glycerophosphocholine. Small metabolite changes were also observed in the amniotic fluid
270 of singleton pregnancies which eventually underwent preterm delivery and premature
271 rupture of membranes. It is difficult to draw comparisons to our cohort, as these
272 pregnancies have taken amniotic fluid from a recipient sac in complex monochorionic
273 multiple pregnancies (in a narrow gestational age range).

274 This is the first study of energy metabolism in TTTS and FLC. We have reported the
275 correlation of the balance between fatty acid and carbohydrate use in energy metabolism
276 and their associations with measures of global recipient cardiac dysfunction and the effects
277 of laser ablative treatment. Additional research is required to delineate the origin of these
278 metabolomic changes, although it would be difficult to obtain ethical approval to perform
279 repeat amniocenteses in these high-risk pregnancies which we would enable us to explore
280 these changes. Further targeted metabolomics studies in different biofluids and tissues are
281 now required to identify potential prognostic ‘biomarkers’ to improve outcome in
282 monochorionic twin pregnancies complicated by TTTS.

283

284 **Contributors:** WBD and JWA analysed samples, interpreted data and wrote manuscript.
285 TVM helped with calculation of the MPI and normalising these data as gestationally related

286 z-scores, as well as contributing to writing the manuscript. RKM, FLM, and CEF helped with
287 procedures, collecting samples, interpreting results and writing manuscript. MDK performed
288 the procedures, interpreted the results and wrote the manuscript. MDK conceived idea. All
289 authors contributed to writing the manuscript and have seen and approved the final version.

290

291 **Acknowledgements:** JWA and WD wish to acknowledge the Systems Science for Health
292 Initiative at the University of Birmingham for financial support. This work was supported by
293 Wellbeing of Women (through an Entry Level Scholarship to CEF) and the Wiseman Trust
294 (supporting FLM).

295

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432

433 **Tables and Figures Legend**

434 **Table 1: Demographic and clinical data for all participating subjects (n=19 twin**
435 **pregnancies; n=38 fetuses)**

436 **Table 2: Correlation of metabolic profiles and cardiac function (Left Ventricle (LV)**
437 **and Right Ventricle (RV) Myocardial Performance Index (MPI)) according to**
438 **metabolite class**

439 **Figure 1: (a) Right ventricular myocardial performance index (RV-MPI) (b) Left**
440 **ventricular myocardial performance index (LV-MPI) in recipient fetuses**

441 The graph demonstrates individual fetal values against gestational age (and with reference
442 to the 95%CI).

443 **Figure 2: Changes in: a) Right Ventricular Myocardial Performance Index (RVMPI)**
444 **and b) Left Ventricular Myocardial Performance Index (LVMPI) after pre- and post-**
445 **fetoscopic laser coagulation in recipient twins**

446 RV and LV MPI z-scores before and immediately post- fetoscopic laser coagulation
447 (individual data shown).

448 **Figure 3: Heat map showing the distribution of concentrations for individual**
449 **metabolites (rows) for samples collected before and after fetoscopic laser**
450 **coagulation (columns).** Green shows a low concentration whereas red shows a high
451 concentration in the range of concentrations for each metabolite. Abbreviations are amino
452 acid metabolism (AA), acyl carnitines (AC), acyl glycerides (AG), carbohydrates (C),
453 cholesterol esters (CE), ceramides and sphingolipids (CS), fatty acids (FA),
454 glycerophospholipids (GPL), haem metabolism (H), nucleotides (N), oxidised fatty acids
455 (OFA), oxidative phosphorylation (OP) and thyroid/steroid hormones (TSH).

456

457 **Supplementary Tables**

458 **Supplementary Table 1: Metabolites showing a statistically significant ($p < 0.005$)**
459 **change in relative metabolite concentrations before and after fetoscopic laser**
460 **coagulation.** All metabolites are grouped into classes of chemical structure or metabolic
461 pathway. Fold change is calculated as the median (before treatment)/median (after
462 treatment) and the 95% confidence intervals are included in brackets.

463 **Supplementary Table 2: Metabolites showing an association between their**
464 **concentration and the Right Ventricle Myocardial Performance Index (MPI) at**
465 **diagnosis and before fetoscopic laser coagulation.** All metabolites listed demonstrated
466 a Spearman Rank correlation between -1.0 to -0.3 or +0.3 to +1.0 and are grouped into
467 classes of chemical structure or metabolic pathway.

468 **Supplementary Table 3: Metabolites showing an association between their**
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472 classes of chemical structure or metabolic pathway.

473

474

475 **Table 1: Demographic and clinical data for all participating subjects (n=19 twin**
 476 **pregnancies; n=38 fetuses)**

Patient characteristics	Median (95%CI)
Maternal age (years)	29 (26.1 – 31.3)
Maternal BMI	24 (22.8 – 27.7)
Gestational age at diagnosis and FLC (days)	142 (137.7 – 148.3)
% difference in EFW	25.2 (21.9 – 31.9)
Recipient fetal cardiac measurements	
Recipient RV MPI (z-score) Median (95%CI)	4.96 (3.62 – 5.50)
Recipient absent/reversed “A-wave” in Ductus Venosus <i>n</i> (%)	7/19 (36.8)
Recipient tricuspid regurgitation <i>n</i> (%)	15/19 (78.9)
Recipient RV E/A ratio > 95%CI for gestation <i>n</i> (%)	11/19 (57.9)
Recipient LV MPI (z-score) Median (95%CI)	2.71 (2.07 – 3.63)
Fetoscopic laser coagulation (FLC) variables	
Duration of FLC and amniodrainage (minutes) Median (range)	34 (21 - 45)
Duration of FLC (minutes) Median (range)	19 (9 – 28)
Number of arteriovenous anastomoses coagulated Median (95%CI)	8 (7.4 – 8.3)
Amniodrainage post-laser coagulation (ml) Median (95%CI)	2600 (1800 – 4000)
Pregnancy outcomes	
Gestational age at delivery (days) Median (95%CI)	227 (207.1 – 232.7)

Perinatal survival of all fetuses* (at 28 days) <i>n</i> (%)	30/38 (78.9)
At least one survivor in pregnancy <i>n</i> (%)	18/19 (94.7)
Two survivors <i>n</i> (%)	12/19 (63.2)
One survivor <i>n</i> (%)	6/19 (31.5)
No survivors <i>n</i> (%)	1/19 (5.3)

477

478 Key: BMI: body mass index, EFW: estimated fetal weight, MPI: myocardial performance
479 index, RV: right ventricle.

480 (†Intergroup comparisons for continuous variables with a non-parametric distribution were
481 made using the Mann-Whitney U test to determine significant differences between the data
482 sets. For such data, median values and 95%CI are described. Categorical data were
483 analysed using Fisher's exact test and relative risk ratios and 95% confidence intervals.
484 Significance was taken as $P < 0.05$ unless otherwise stated)

485 *Perinatal mortality defined as total number of survivors (all fetuses) who survived until at
486 least 28 days of age.

487

Table 2: Correlation of metabolic profiles and cardiac function (Left Ventricle (LV) and Right Ventricle (RV) Myocardial Performance Index (MPI)) according to metabolite class

Metabolite Class	Left Ventricle (LV) MPI		Right Ventricle (RV) MPI	
	Number of Metabolites	Correlation coefficient range	Number of Metabolites	Correlation coefficient range
Acyl amino acids	2	+0.34 to +0.39	2	+0.30 to +0.37
Acyl carnitine	5	-0.46 to +0.41	6	-0.43 to -0.31
Acyl glycerides	12	-0.52 to +0.54	9	-0.49 to +0.60
Bile acid metabolism	3	-0.38 to +0.44	3	+0.31 to +0.36
Carbohydrates	3	-0.57 to +0.65	4	-0.33 to +0.45
Ceramides and sphingolipids	8	-0.43 to -0.30	9	-0.49 to +0.42
CoA metabolism	2	-0.34 to +0.36	2	-0.40 to -0.30
Fatty acid metabolism	10	-0.67 to +0.54	13	-0.52 to +0.46
Glycerophospholipids	41	-0.53 to +0.50	28	-0.64 to +0.51
Nucleoside	3	-0.48 to +0.34	4	-0.36 to +0.54
Oxidised fatty acids	4	-0.31 to -0.30	7	-0.48 to +0.57

The metabolites identified as having an association with LV MPI or RV MPI (denoted as '+' correlation coefficient shows that as LV MPI or RV MPI increase so does the metabolite concentration) or a negative correlation (denoted as '-' shows that as LV MPI or RV MPI increase the metabolite concentration decreases).

Figure 1: (a) Right ventricular myocardial performance index (RV-MPI) (b) Left ventricular myocardial performance index (LV-MPI) in recipient fetuses

The graph demonstrates individual fetal values against gestational age (and with reference to the 95%CI).

Figure 2: Changes in: a) Right Ventricular Myocardial Performance Index (RVMPI) and b) Left Ventricular Myocardial Performance Index (LVMPI) after pre- and post-fetoscopic laser coagulation in recipient twins

RV and LV MPI z-scores before and immediately post- fetoscopic laser coagulation (individual data shown).

Figure 3: Heat map showing the distribution of concentrations for individual metabolites (rows) for samples collected before and after fetoscopic laser coagulation (columns). Green shows a low concentration whereas red shows a high concentration in the range of concentrations for each metabolite. Abbreviations are amino acid metabolism (AA), acyl carnitines (AC), acyl glycerides (AG), carbohydrates (C), cholesterol esters (CE), ceramides and sphingolipids (CS), fatty acids (FA), glycerophospholipids (GPL), haem metabolism (H), nucleotides (N), oxidised fatty acids (OFA), oxidative phosphorylation (OP) and thyroid/steroid hormones (TSH).

Supplementary Tables

Supplementary Table 1: Metabolites showing a statistically significant ($p < 0.005$) change in relative metabolite concentrations before and after fetoscopic laser coagulation. All metabolites are grouped into classes of chemical structure or metabolic pathway. Fold change is calculated as the median (before treatment)/median (after treatment) and the 95% confidence intervals are included in brackets.

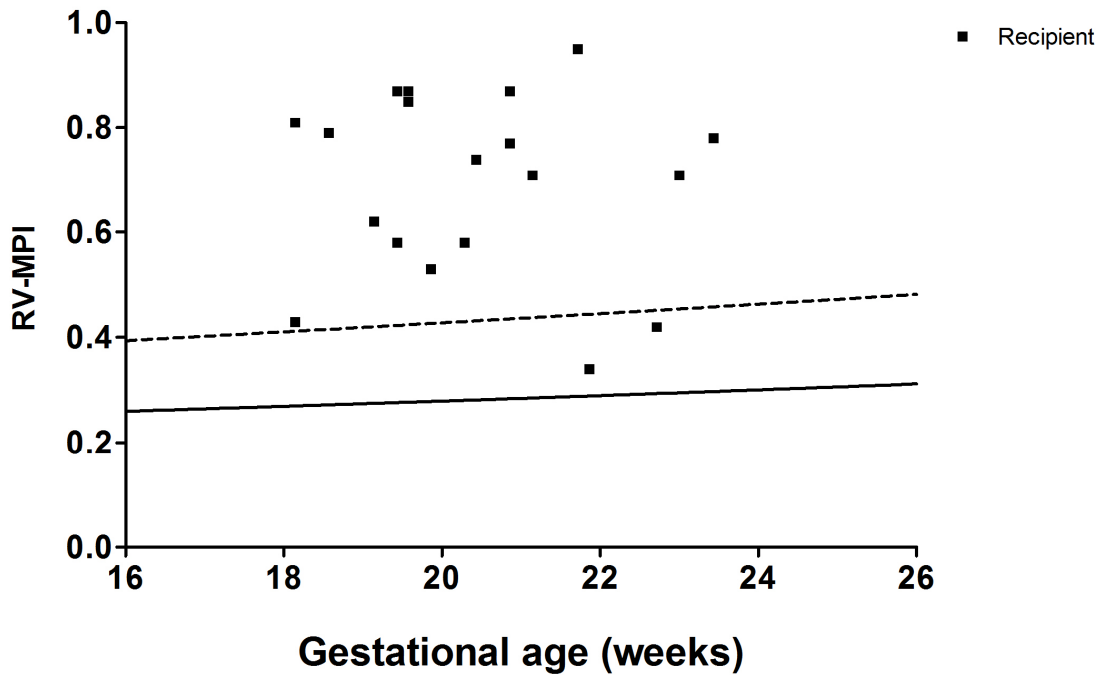
Supplementary Table 2: Metabolites showing an association between their concentration and the Right Ventricle Myocardial Performance Index (MPI) at diagnosis and before fetoscopic laser coagulation. All metabolites listed demonstrated a Spearman Rank correlation between -1.0 to -0.3 or +0.3 to +1.0 and are grouped into classes of chemical structure or metabolic pathway.

Supplementary Table 3: Metabolites showing an association between their concentration and the Left Ventricle Myocardial Performance Index (MPI) at diagnosis and before fetoscopic laser coagulation. All metabolites listed demonstrated a Spearman Rank correlation between -1.0 to -0.3 or +0.3 to +1.0 and are grouped into classes of chemical structure or metabolic pathway.

Figure 1: (a) Right ventricular myocardial performance index (RV-MPI) (b) Left ventricular myocardial performance index (LV-MPI) in recipient fetuses

The graph demonstrates individual fetal values against gestational age (and with reference to the 95%CI).

(a)



(b)

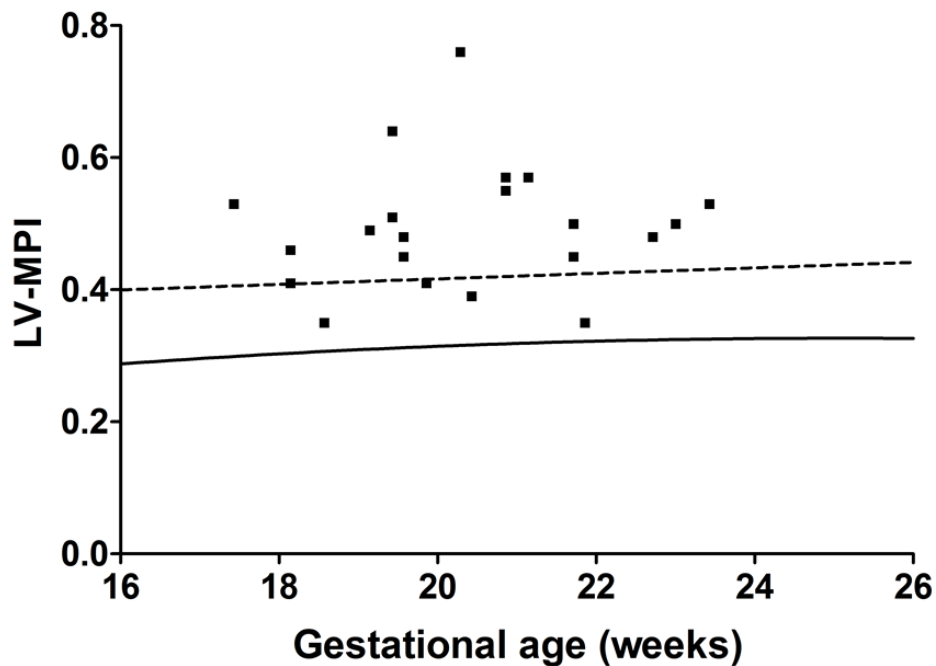
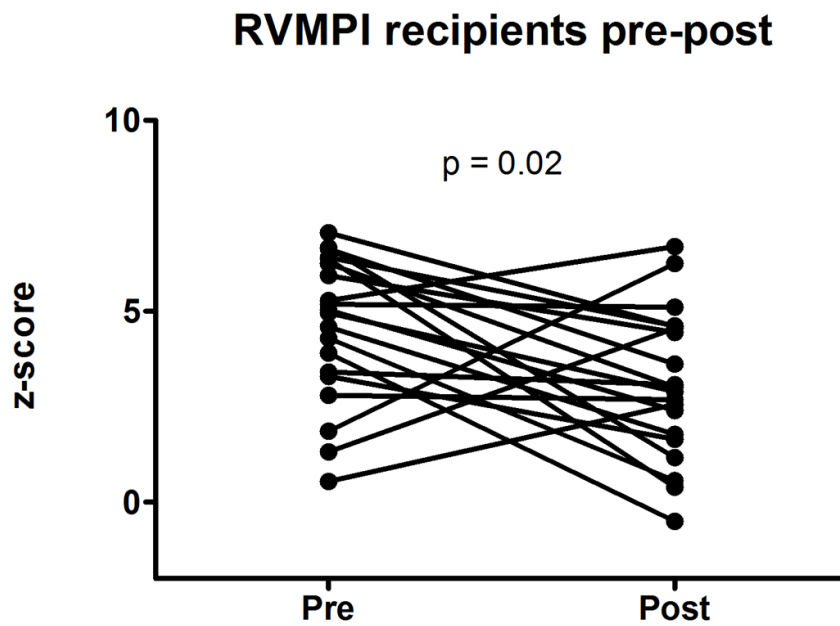


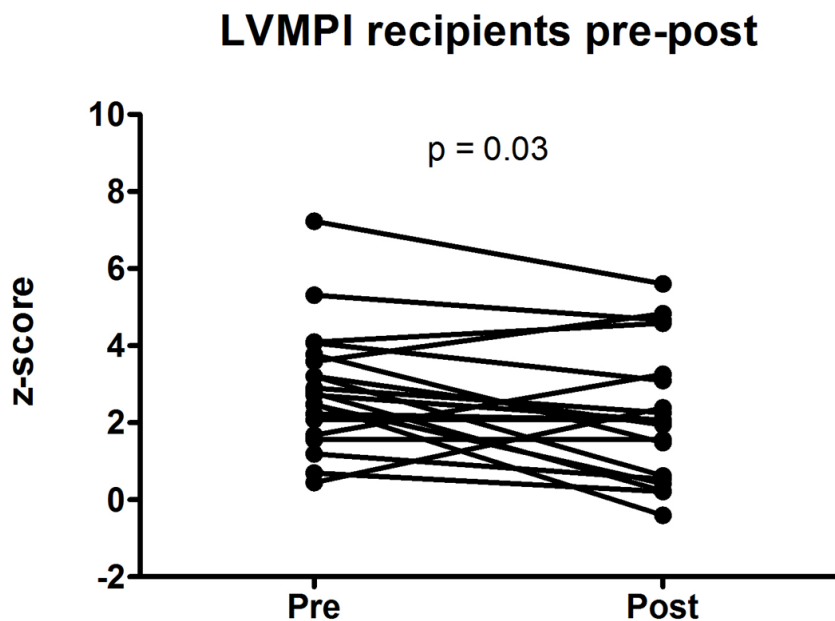
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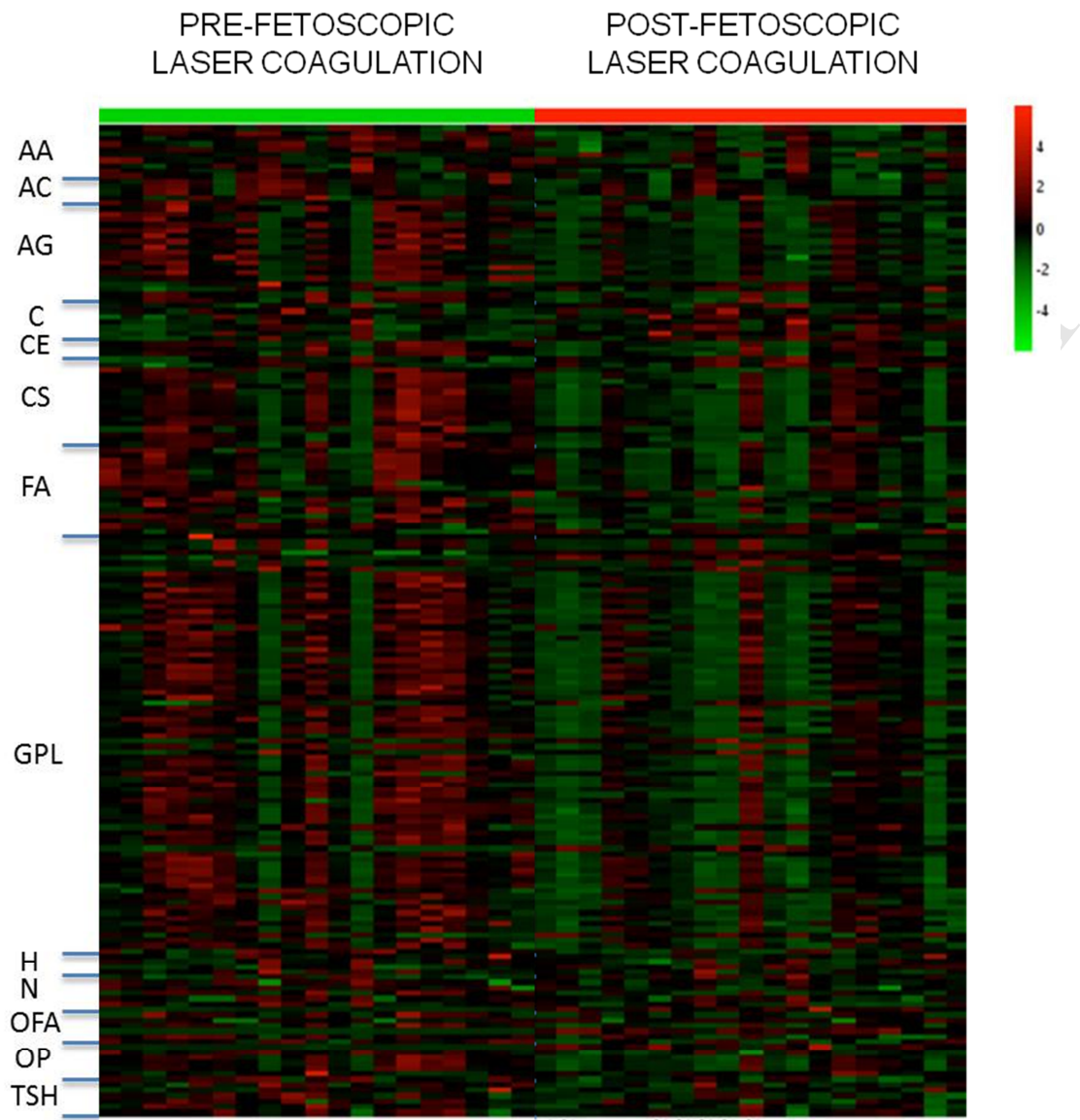
RV and LV MPI z-scores before and 6 hours post- fetoscopic laser coagulation (individual data shown).

a)



b)





Highlights

1. Metabolomic changes are seen after fetoscopic laser coagulation (FLC)
2. Carbohydrate and fatty acid metabolism appears altered following FLC
3. These findings are in keeping with changes seen in adults with cardiomyopathy
4. Metabolomics may provide new biomarkers for twin twin transfusion syndrome