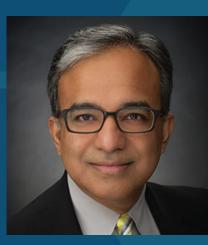
METABOLOMIC AND LIPIDOMIC ANALYSIS OF PATIENTS WITH PRESUMED METABOLIC DYSFUNCTION-ASSOCIATED STEATOHEPATITIS (MASH) TREATED WITH THE SELECTIVE GLUCOCORTICOID RECEPTOR MODULATOR MIRICORILANT



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Summary & Conclusions

- The observed declines in intermediate lipid species suggest that responders (patients with \geq 30% reduction from baseline in LFC following miricorilant treatment), particularly those with rapid LFC clearance accompanied by efficacy-associated ALT elevations (rapid responders [RR]), experienced enhanced metabolic clearance of lipids and FAs
- In RRs, metabolic clearance of acylglycerols continued to improve at week 12 despite discontinuation or interruption of miricorilant after week 6
- Rapid changes in liver metabolism, such as increased FA oxidation, can cause transient hepatic stress. This stress may produce a short-term rise in ALT levels as the liver adjusts to the new metabolic conditions
- Although this may cause transient hepatic stress, it points to improved hepatic lipid metabolism
- Changes in bile metabolism could suggest elevated liver capacity to synthesize bile acids, enhancing lipid emulsification or clearance
- The ongoing phase 2b MONARCH study is assessing 100 mg miricorilant twice weekly, a dose that resulted in improved LFC without ALT elevations in miricorilant's phase 1b study, and a higher dose of 200 mg twice weekly after a 6-week lead-in of 100 mg twice weekly

ALT, alanine aminotransferase; FA, fatty acid; LFC, liver fat content.

The authors want to thank all those who participated in this study: The study patients and their families, the investigators, and the sponsor team.



Background

Aim

Methods

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Results: Demographic Characteristics

Age group, n <30 y 30 to 50 y >50 y

Female, n (% _____ Race, n (%)

White Asian

Hispanic or La _____ BMI, n (%)

30 to <35 kg 35 to <40 kg ≥40 kg/m² Liver stiffness.

8.5 to 12 kPa >12 kPa MRI-PDFF %, n

Hypertensior Diabetes, n (%)

BMI, body mass index; MRI-PDFF, magnetic resonance imaging derived proton density fat fraction; NR, nonresponder; RR, rapid responder; SD, standard deviation; SR, standard responder.

• Cortisol activity has been implicated in metabolic dysfunction-associated steatotic liver disease (MASLD) development and progression^{1,2}

Cortisol binding to the glucocorticoid receptor (GR) increases availability of energy substrates, such as free fatty acids (FFAs), for daytime activities and stress response Cortisol can contribute to increased lipid levels in the liver by increasing uptake and de novo lipogenesis

• Miricorilant (Figure 1) is an orally delivered, nonsteroidal selective glucocorticoid receptor modulator (SGRM) that acts as a mixed GR agonist/ antagonist and modest mineralocorticoid receptor (MR) antagonist (6-fold higher affinity for GR vs MR) and may reduce hepatic steatosis by modulating cortisol activity in the liver³

• In MASLD/MASH preclinical models, miricorilant reversed and prevented hepatic steatosis by preventing lipid accumulation in the liver and reduced inflammation, fibrosis stage, and NAS⁴

• In a phase 1b, multicohort, open-label, dose-finding study (NCT05117489), adults with presumed MASH with fibrosis treated with 30–200 mg miricorilant dosed daily or intermittently for 12 or 24 weeks had notable and rapid liver fat content (LFC) decreases, which was occasionally accompanied by efficacy-associated alanine aminotransferase (ALT) increases⁵

Miricorilant 100 mg twice weekly was safe, welltolerated, and resulted in reduced LFC (-28.2% at week 12), with a corresponding decline in liver

enzymes and improved hepatic, lipid, and glycemic

Miricorilant 100 mg twice weekly is being assessed in the phase 2b MONARCH study (NCT06108219)

• This metabolomic analysis of the phase 1b study data explored miricorilant's mechanism of reducing LFC with the following aims: Test the hypothesis that rapid enhancement in metabolic activity can lead to

transient ALT elevations Characterize metabolomic profiles in plasma from responders vs nonresponders

Figure 1. Miricorilant

following miricorilant treatment

Assess biochemical differences in responders with and without efficacy-associated **ALT** elevations

samples taken at baseline and weeks 6 and 12 from select patients irst 10 study cohorts were stratified into 3 groups based on response ess of miricorilant dose received:

dard responders (SRs): ≥30% reduction from baseline in LFC without ALT tions (n=11)

responders (RRs): \geq 30% reduction from baseline in LFC with treatment acy-associated ALT elevations >3× upper limit of normal (n=9) esponders (NRs): without reduction from baseline in LFC (n=11)

ic profiles in serum were assessed via liquid chromatography/mass

• Metabolite changes within each responder group over time (baseline to week 6, baseline to week 12, week 6 to 12) and between groups at specified time points were analyzed using fold change correlations

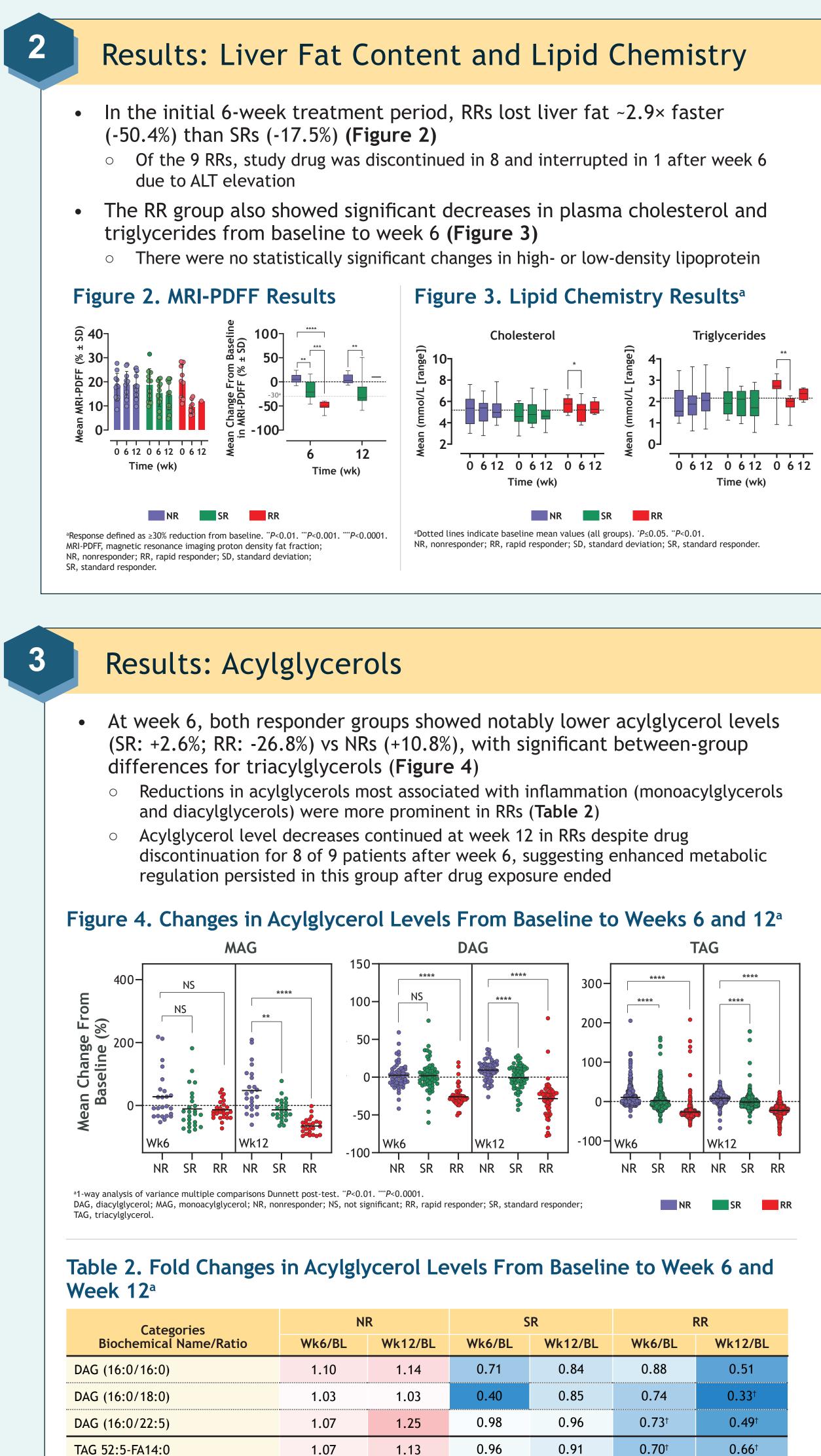
• Results were analyzed by 2-way repeated-measures analysis of variance and matched pairs t-test

NAS, nonalcoholic fatty liver disease activity scor

• NRs were selected to match RRs and SRs based on demographic characteristics (race, sex, and age; Table 1)

Table 1. Baseline Patient Characteristics

	SRs (n=11)	RRs (n=9)	NRs (n=11)	
(%)	0 4 (36.4) 7 (63.6)	0 4 (44.4) 5 (55.6)	2 (18.2) 3 (27.3) 6 (54.5)	
, ,	7 (63.6)	5 (55.6)	6 (54.5)	
	11 (100) 0	8 (88.9) 1 (11.1)	11 (100) 0	
atino ethnicity, n (%)	7 (63.6)	5 (55.6)	6 (54.5)	
/m² /m²	2 (18.2) 4 (36.4) 4 (36.4)	5 (55.6) 1 (11.1) 2 (22.2)	4 (36.4) 4 (36.4) 2 (18.2)	
s, n (%) a	6 (54.5) 5 (45.4)	(n=8) 4 (50.0) 4 (50.0)	10 (90.9) 1 (9.1)	
mean (SD)	18.7 (6.7)	20.3 (6.7)	17.9 (5.6)	
, n (%)	5 (45.4)	5 (55.6)	7 (63.6)	
6)	7 (63.6)	2 (22.2)	3 (27.3)	



References

TAG 52:5-FA22:5

TAG 52:6-FA14:0

TAG 54:7-FA22:5

TAG 56:7-FA22:4

MAG (17:0)

MAG (22:6)

(1) Rahimi L, et al. *Diabetes Metab Syndr Obes*. 2020;13:1133-1145. (2) Rinella ML, et al. *Hepatology*. 2023;77(5):1797-1835. (3) Hunt HJ, et al. Bioorg Med Chem Lett. 2012;22(24):7376-7380.

1.13

1.23

1.21

1.03

1.27

Decreased Trajectory (Fold change <1)

(4) Koorneef LL, et al. Endocrinology. 2018;159(12):3925-3936. (5) Alkhouri N, et al. Presented at: NASH-TAG 2024; January 4-6, 2024; Park City, UT

Acknowledgements

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Presenter Disclosures

NGM, Pfizer, Pliant, and Viking; royalties/licenses from UpToDate; consulting fees from 89bio, Calliditas Therapeutics, CymaBay Gilead, Inipharm, Intercept, Madrigal, Mirum, NGM, and Pliant; payment/honoraria for lectures, presentations speakers bureaus, manuscript writing, or educational events from AbbVie, Gilead, and Intercept.

	S	R	RR		
/BL	Wk6/BL	Wk12/BL	Wk6/BL	Wk12/BL	
4	0.71	0.84	0.88	0.51	
3	0.40	0.85	0.74	0.33†	
5	0.98	0.96	0.7 3 [†]	0.49 [†]	
3	0.96	0.91	0.70 [†]	0.66 [†]	
3	0.94	0.93	0.64*	0.59 [†]	
2	0.97	0.97	0.59*	0.51†	
2	0.93	0.97	0.66 [†]	0.46*	
4	0.85	0.99	0.65*	0.59*	
5	0.47	0.47	0.85	0.06*	
)	0.56	0.72	0.61	0.39 [†]	
	Increased Trajed	ctory (Fold change	>1)		

^aColor scheme based on a 3-color scale with minimum of 0.22, midpoint 1, and maximum 2. ^{*}P≤0.05. [†]0.05<P<0.10 BL, baseline; DAG, diacylglycerol; FA, fatty acid; MAG, monoglyceride; NR, nonresponder; RR, rapid responder; SR, standard responder; TAG, triacylglycerol.

Results: Long-chain Fatty Acids

- Most long-chain FAs (LCFAs), including polyunsaturated FAs (PUFAs) omega-3 and -6, were reduced (fold change <1) at week 6 and week 12 vs baseline in responders regardless of ALT elevation but were elevated (fold change >1) in NRs (Table 3 and Figure 5)
- Significant reductions were observed at week 6 for SRs (-5.3%; P<0.01) and RRs (-16.6%; P<0.0001) but not NRs (+0.6%) vs their respective baselines
- Decreases in LCFAs and PUFAs may suggest improved FA oxidation in responders, which would be expected to reduce circulating FFAs

Figure 5. Changes in LCFA Levels From Baseline to Weeks 6 and 12^a

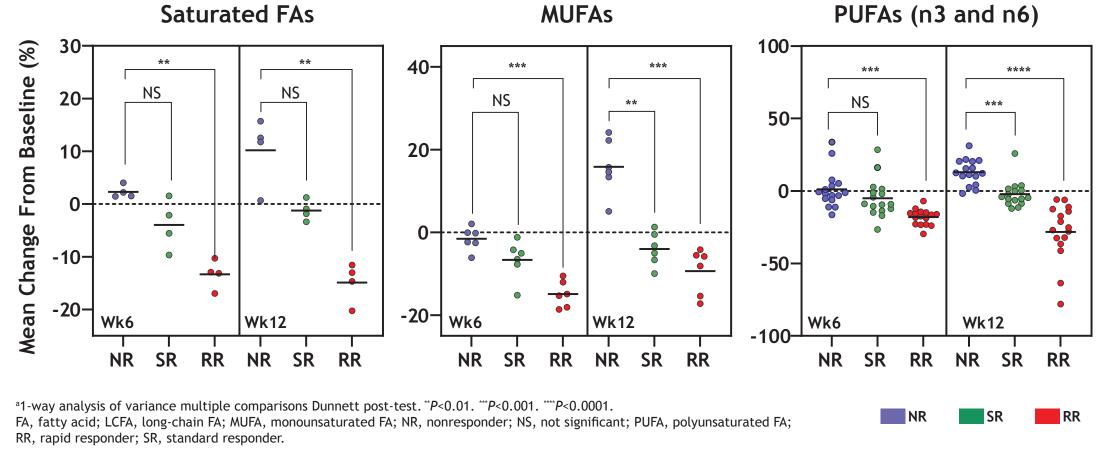


Table 3. Fold Changes in LCFA Levels From Baseline to Weeks 6 and 12^a

	Categories	1	NR	S	R		RR
LUFAS	Biochemical Name/Ratio	Wk6/BL	Wk12/BL	Wk6/BL	Wk12/BL	Wk6/BL	Wk12/BL
	Myristate (14:0)	1.02	1.16	1.02	0.98	0.87	0.88
	Pentadecanoate (15:0)	1.01	1.13	0.94	0.99	0.87	0.87
	Margarate (17:0)	1.02	1.12	0.90	0.97	0.83	0.80
	Myristoleate (14:1n5)	1.00	1.16	0.95	0.93	0.82	0.94
	Palmitoleate (16:1n7)	0.94	1.22	0.94	0.97	0.85	0.96
	10-heptadecenoate (17:1n7)	0.98	1.24	0.85	0.90	0.81	0.85
MUFAS	Oleate/vaccenate (18:1)	1.00	1.13	0.99	1.01	0.89	0.92
	10-nonadecenoate (19:1n9)	0.97	1.05	0.92	0.95	0.85	0.83
	Eicosenoate (20:1)	1.02	1.15	0.96	1.00	0.88	0.94
	Tetradecadienoate (14:2)	0.99	1.04	1.03	1.04	0.82	0.73
	Stearidonate (18:4n3)	1.34	1.21	1.16	0.94	98 0.87 0.87 99 0.87 0.87 97 0.83 0.87 93 0.82 0.97 97 0.85 0.97 97 0.85 0.97 90 0.81 0.87 97 0.85 0.97 90 0.81 0.89 91 0.89 0.97 92 0.85 0.88 91 0.89 0.97 92 0.85 0.88 93 0.77 0.64 94 0.77 0.64 93 0.70 0.27 94 0.77 0.64 93 0.70 0.27 94 0.77 0.64 95 0.77 0.33 96 0.83 0.83 97 0.85 0.94 96 0.84 0.77 97 0.85 0.84 97 0.86 0.84 97 0.86 0.84 97	0.63
LCFAs Biochemical Name/Ratio Wk6/BL Wk6/BL	1.03	0.70	0.22				
	Docosapentaenoate (n3 DPA; 22:5n3)	0.89	K6/BL Wk12/BL Wk6/BL Wk12/BL Wk6/BL 1.02 1.16 1.02 0.98 0.87 1.01 1.13 0.94 0.99 0.87 1.02 1.12 0.90 0.97 0.83 1.00 1.16 0.95 0.93 0.82 0.94 1.22 0.94 0.97 0.85 0.98 1.24 0.85 0.90 0.81 1.00 1.13 0.99 1.01 0.89 0.97 1.05 0.92 0.95 0.85 1.02 1.15 0.96 1.00 0.88 0.99 1.04 1.03 1.04 0.82 1.34 1.21 1.16 0.94 0.77 0.98 1.20 1.01 1.03 0.70 0.89 1.10 0.90 0.95 0.77 0.89 0.98 0.85 0.88 0.77 0.89 0.98 0.80 0.81	0.36			
	Docosahexaenoate (DHA; 22:6n3)	0.89	0.98	0.85	0.88	0.77	0.68
	Docosatrienoate (22:3n3)	0.95	1.22	0.73	0.91	0.85	0.67
ΡΠΕΛς	Hexadecadienoate (16:2n6)	1.03	1.15	0.98	1.00	0.83	0.87
	Linoleate (18:2n6)	1.08	1.20	0.96	1.01	0.85	0.94
n-6)	Linolenate [alpha or gamma; (18:3n3 or 6)]	1.26	1.31	0.99	0.96	0.81	0.94
	Dihomo-linoleate (20:2n6)	1.01	1.13	0.90	0.97	0.85	0.83
	Arachidonate (20:4n6)	0.97	1.10	0.86	0.95	0.84	0.72
	Adrenate (22:4n6)	0.84	1.11	0.83	0.89	0.84	0.75
	Docosapentaenoate (n6 DPA; 22:5n6)	0.94	1.00	0.91	0.95	0.86	0.86
	Docosadienoate (22:2n6)	0.99	1.02	0.91	1.00	0.93	0.89
	Mead acid (20:3n9)	1.05	1.12	1.28	1.26	0.76	0.59 [†]

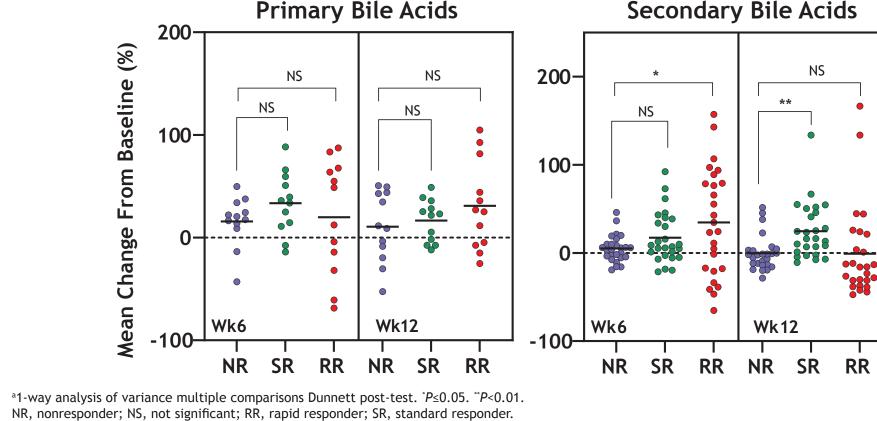
PUFA, polyunsaturated FA; RR, rapid responder; SR, standard responde

Increased Trajectory (Fold change >1

Results: Bile Acids

- At baseline, ~70% of the analyzed primary and secondary bile acids showed elevated levels in RRs compared with SRs and NRs
- For the SR and RR within-group comparisons, most primary bile acid species did not change significantly from baseline to weeks 6 or 12 (**Table 5 and Figure 7**) • For RRs, numerous secondary bile acid species were significantly shifted bidirectionally from baseline
- At week 6, RRs had increasing trajectories (several significant) vs NRs for numerous primary and secondary bile acid species
- SRs trended toward increased bile acid species vs NRs • In SRs, most primary bile acids decreased by ~19% from week 6 to week 12

Figure 7. Changes in Bile Acid Levels From Baseline to Weeks 6 and 12^a



Results: Fatty Acid ß-Oxidation

- Many short- to medium-chain acylcarnitine species were reduced (fold change <1) in RRs vs increased in NRs (fold change >1) (Table 4) Lower levels can reflect improved utilization by the mitochondria at the tissue level
- 3-hydroxy FA species were significantly reduced for RRs at week 6 vs NRs $(P \le 0.05)$, pointing to improved liver FA oxidation (Figure 6)

Figure 6. Changes in Acylcarnitine and 3-hydroxy FA Levels From Baseline to Weeks 6 and 12^a

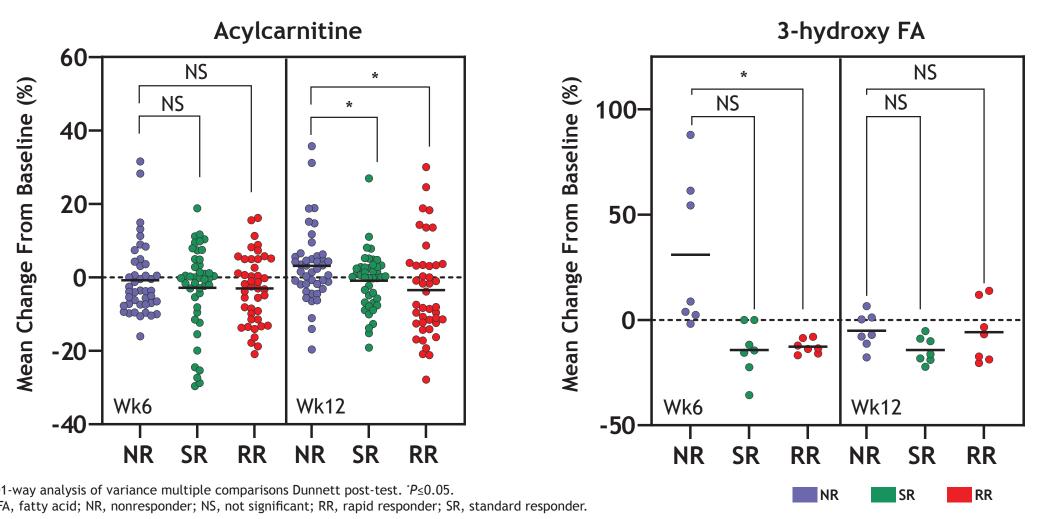


Table 4. Fold Changes in Acylcarnitine and 3-hydroxy FA Levels From Baseline to Weeks 6 and 12^a

Acylcarnitine/	Categories	NR		SR		RR	
3-hydroxy FA	Biochemical Name/Ratio	Wk6/BL	Wk12/BL	Wk6/BL	Wk12/BL	Wk6/BL	Wk12/BL
Short Chain	Acetylcarnitine (C2)	0.90 [†]	0.98	1.00	0.93	0.92	0.99
	Hexanoylcarnitine (C6)	0.90	1.03	1.00	0.95	0.87	0.99
Madium Chain	Octanoylcarnitine (C8)	1.03	1.09	1.01	1.01	Wk6/BL 0.92 0.87 0.81 0.81 1.0.81 0.82 1.00 1.00 0.97 0.97 0.97 0.97 0.97 0.97 0.92 0.93 0.94 0.95 1.07 1.07 0.89 0.91 0.91 0.92 0.83 0.91 0.83 0.84 0.83 0.84 0.85 0.86 0.88 0.86	0.84
Medium Chain	Nonanoylcarnitine (C9)	0.96	0.96	1.03	1.01		0.79 [†]
	Laurylcarnitine (C12)	1.00	1.05	1.11	1.03		0.87
	Myristoylcarnitine (C14)	0.93	0.99	1.05	1.02	1.01	0.91
	Palmitoylcarnitine (C16)	0.94	0.99	1.07	1.02	0.97	0.88
	Margaroylcarnitine (C17)	0.90	0.99	0.98	0.93	0.92	0.89
Long Chain Saturated	Stearoylcarnitine (C18)	0.93	0.89	1.00	0.94	0.97	0.86 [†]
Long Chain Saturated	Arachidoylcarnitine (C20)	1.00	1.00	1.00	1.01	Wk6/BL 0.92 0.87 0.81 0.81 1.03 1.00 1.00 0.97 0.97 0.97 0.97 0.97 0.97 1.00 0.97 0.97 1.097 0.97 0.97 0.97 0.997 0.997 1.009 1.009 0.89 0.91 0.91 0.91 0.93 0.93 0.84 0.92 0.88 0.88 0.88 0.88 0.88 0.86	0.90
	Behenoylcarnitine (C22)	1.09	0.96	0.85	0.87		0.90
	Lignoceroylcarnitine (C24)	0.96	0.93	0.88	1.00		1.03
	Cerotoylcarnitine (C26)	1.01	1.05	1.07	1.03		1.04
	3-hydroxyhexanoate	1.04	0.92	0.86	0.91	0.91	0.93
	3-hydroxyoctanoate	1.61	1.07	0.78	0.78	0.83	1.14
	3-hydroxydecanoate	1.54	1.01	0.88	0.81	Wk6/BL 0.92 0.87 0.87 0.81 0.82 1.00 1.00 0.97 0.91 0.91 0.92 0.93 0.93 0.93 0.93 0.93 0.93 0.93 0.93 0.93 0.93 0.93	0.97
2 huden a Matabalian	3-hydroxysebacate	1.88	1.00	0.64	0.82	0.92	1.12
3-hydroxy Metabolism	3-hydroxylaurate	1.09	0.93	1.00	0.90	 0.87 0.81 0.82 1.00 1.01 0.97 0.97 0.97 0.97 0.89 1.09 1.09 1.07 0.89 0.89 0.81 0.83 0.84 0.92 0.84 0.92 0.84 0.84 0.84 0.84 0.84 0.84 0.88 0.88 0.86 	0.80
	3-hydroxymyristate	0.98	0.89	1.00	0.95		0.81
	3-hydroxystearate	1.02	0.82	0.84	0.84		0.83
	3-hydroxyoleate	0.75*	0.83	0.82	1.20		0.73 [†]

BL, baseline; FA, fatty acid; NR, nonresponder; RR, rapid responder; SR, standard responder.

Increased Trajectory (Fold change >1)

Table 5. Fold Changes in Bile Acid Levels From Baseline to Weeks 6 and 12^a **Bile Acid Biochemical Name/Rational Name** Cholate 1.36 2.55 4.00* 1.55 Glycocholate 1.95 4.21* 2.16 Taurocholate 1.10 0.86 0.78 Chenodeoxycholic acid sulfat cochenodeoxycholate Primary urochenodeoxycholate 2.38 3.35* 1.40 1.12 Cholic acid glucuronide 1.10 1.25 Glycochenodeoxycholate glucuron 1.23 1.81[†] 1.48 Glycochenodeoxycholate 3-sulfat 1.44 Deoxycholate Deoxycholic acid glucuronide 1.34 2.25 1.67 lycodeoxycholate aurodeoxycholate 1.00 0.81 0.71 aurodeoxycholic acid 3-sulfa 1 74 _ithocholic acid sulfate 1.36 1.23 1.94 lycolithocholate sulfat 1.89 1.04 **1.22 2.25[†] 1.84[†]** rsodeoxycholate ycoursodeoxycholate Secondar lycoursodeoxycholic acid sul 0.53 3.59 3.04 0.8 0.66 auroursodeoxycholate auroursodeoxycholic acid sulfa 1.58 6.65 4.20 0.89 1.50 1.69 Hyocholate (gamma-murichola 1.14 2.09 1.84 Glycocholenate sulfate 1.26 2.15* 1.70 aurocholenate sulfate

^aColor scheme based on a 3-color scale with minimum of 0.35, midpoint 1, and maximum 6.65. *P≤0.05. *0.05<P<0.10.

0.98

1.19

1.38 1.09

Bbeta-hydroxy-5-cholenoate

lycodeoxycholate 3-sulfate

NR, nonresponder; RR, rapid responder; SR, standard responde

aurochenodeoxycholic acid 3-sulfate

Decreased Trajectory (Fold change <1 Increased Trajectory (Fold change >1)

0.77 1.47 3.39* 2.30*

0.87

1.21

1.74

1.25 **1.79**[†]

1.23 2.43⁺

0.73[†] 1.03 1.41

1.65 **5.48**^{*} 3.31^{*}