Differences in the Metabolic Response to a High-Carbohydrate (HC) Meal Intake between Lean and Overweight/Obese (OW/OB) men - A metabolomics study

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Abstract

The OW/OB individuals exhibited more pronounced changes in the metabolic response to HC meal intake compared to normocarbohyderate (NC) meal consumption. After the HC meal significantly different AUC for metabolites from several classes was observed. Among others, an increase of lysophosphatidylcholines (LPC) (from +35 to +185%, variable importance in the projection (VIP) range 1.1-1.9), lysophosphatidylethanolamines (LPE) (from +133 to +255%, VIP 1.4-2.2), long-chain fatty acids (LCFA) (from +54 to +133%, VIP 1.2-1.7), or leukotrienes (from +84 to 216%, VIP 1.4-1.9) and decrease of sphinganine (SM) (from -60 to -40%, VIP 1.3-1.6) in OW/OB compared to lean individuals was observed.

Background

Since last decade, the dietary recommendations of macronutrient intake have been changing enormously for the obese/overweight and type 2 diabetic patients (T2DM).

Initially, a low-calorie-diet (LC) comprising mainly of fats with a limited restriction on carbohydrates was considered ideal which was, later, changed to the promotion of high-carbohydrate (HC) and low-in-fat-diet consumption after recognizing the role of diabetes in developing the cardiovascular disease (1, 2). However, the consumption of HC-diets with low fat contents also failed to limit the incidences of metabolic syndrome (4, 5). A lot of data published in literature found no considerable difference in postprandial glucose and lipid metabolism in cases of either taking HC or LC-diets (3-5). These conflicting results in literature, prompted us to evaluate the postprandial metabolic response after HC/NC meals intake in normal-weight and overweight/obese men.

Objective

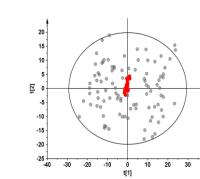
The short-term and immediate postprandial metabolic response to HC meal intake was evaluated and compared between lean and overweight/obese (OW/OB) healthy men.

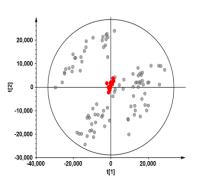
Material and Methods

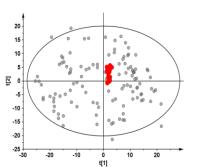
Two groups of non-diabetic male participants (i.e. OW/OB and lean, age: 23-38 years, BMI: from 23.8 to 30.8), each consisting of 12 subjects, underwent a randomized, controlled, and cross-over one-time HC-mealchallenge trial. On the two different visits (with 2-3 weeks of break between them) the participants received standardized HC meal (89% of energy from carbohydrate, 11% protein, and 0% fat) or normalcarbohydrate (NC) meal (45% carbohydrate, 30% protein, 25% fat) Fasting (0 min) and postprandial blood samples were collected at predefined time points (i.e. 30, 60, 120, 180 min) and untargeted plasma metabolomics was performed using LC-QTOF-MS. For each metabolic feature, based on its intensity change in time, the area under the curve (AUC) was calculated. Obtained AUC values were forwarded for multivariate statistical analysis.

This trial was registered at www.clinicaltrials.gov as NCT03792685

Figure 1: PCA models based on postprandial plasma metabolite AUCs obtained after HC-meal intake showing a good separation between obese/overweight and lean men.







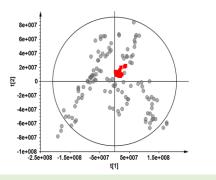
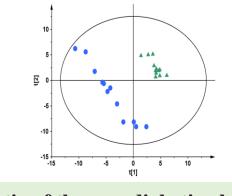


Figure 2: PLS-DA models based on postprandial plasma metabolite AUCs obtained after HC-meal intak showing a good separation between obese/overweight and lean men.



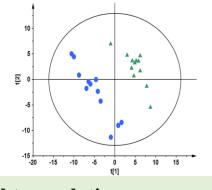


Table 1: Baseline characteristic of the non-diabetic obese/overweight population compared to lean (L) population.

Table 1. Daseline Characteristic of the holf-diabetic obese/o									
Parameters	OB/OW	L	<i>p</i> -value						
Age (years)	38±5.4	23.8±1.6	0.4						
BMI (Kg/m2)	30.8 ± 5.4	23.8±1.6	0.0009						
Body Fat contents (%)	28.0 ± 6.3	17.0 ± 5.3	0.0001						
Fat free mass (%)	69.8±12.2	66.3±6.6	0.4						
Weight height ratio	$0.99.8 \pm 0.06$	0.91 ± 0.06	0.003						
Fasting glucose	87.9 ± 5.8	84.2 ± 8.1	0.2						
concentration (mg/dl)									
Fasting Insulin	12.7 ± 9.3	6.5 ± 1.7	0.06						
concentration									
(IU/mL)									
HOMA-IR	2.8 ± 2.1	1.3 ± 0.3	0.05						
НОМА-β	186.3±121.1	157.5±150.	0.6						
		4							
HbA1c	15.3±0.3	5.2±0.3	0.3						

Note: For quantitative variables with normal distribution, the parametric t-test was used; for the other variables, the nonparametric Mann-Whitney test was applied. The data are represented as the mean ± STD, and p-values< 0.05 were considered significant. * HOMA-IR= Homeostatic Model Assessment of Insulin Resistance; $HOMA-\beta = Homeostatic$ Model Assessment of β-cell function; HbA1c= glycated hemoglobin; WHR= waist-hip ratio

Results

Table 2: Significant metabolites which postprandial change discriminate obese and lean individuals dependently of the meal type

	20 8e+007		RT	Metabolites	High carbohydrate diet		te diet	Normal carbohydrate diet				Remark		
	15 10 5 5 7 7 10 10 10 10 10 10 10 10 10 10 10 10 10	Mass	(min)		Change (%)	VIP	p-corr	p-value	Change (%)	VIP	<i>p</i> -corr	p-value	S	
	∑ 0 -5- -2e+007	370.1814	3.9	Epiandosterone sulfate	114.22	1.3039	-0.514	0.1	-8.0926	0.445	0.0419	0.9	OK	
	-10 -15 -6e+007	304.2397	7.1	Arachidonic acid	89.3829	1.2136	-0.5557	0.1	5.9219	0.635	-0.1622	0.9	OK	
	-20 -25 -30 -20 -10 0 10 20 -1e+008 -2.5e+008 -1.5e+008 -5e+007 1.5e+008	344.2347	5.7	HDOHE	230.9431	2.039	-0.7173	0.02	-22.207	0.491	0.0774	0.6	OK	
	t[1] -2.5e+008 -1.5e+008 -5e+007 - 5e+007 - 1.5e+008 -1.5e+008 -1.	320.2348	5.8	HETE	129.0137	1.7674	-0.6441	0.08	-11.1626	0.465	-0.0092	0.9	OK	
m	a metabolite AUCs obtained after HC-meal intake	300.2659	7.4	Hydroxy stearic acid	53.8667	1.4469	-0.5061	0.07	-17.2622	0.338	0.0774	0.9	Frag	
t and lean men.		213.0097	0.7	Indoxylsulfuric acid	155.3375	2.249	-0.6696	0.07	-25.1716	0.328	0.1112	0.6	OK	
	4	90.03185	0.3	Lactic acid	73.0377	1.7011	-0.5579	0.05	10.4388	0.167	-0.0385	0.6	OK	
	10	467.3369	5.9	LPC 15:0	124.7424	1.6698	-0.6545	0.1	-18.5582	1.163	0.328	0.4	OK	
	5	507.333	5.8	LPC 17:1	34.9017	1.062	-0.5479	0.1	-29.2796	1.177	0.3667	0.2	OK	
	₫ 0	509.348	6.3	LPC 18:0	90.9997	1.6398	-0.5086	0.1	-14.8049	0.314	0.0691	0.8	OK	
	-5-	509.3481	6.2	LPC 18:0	94.4603	1.7454	-0.5447	0.1	-14.1671	0.3	0.0597	0.8	OK	
	-10	551.3587	6	LPC 18:2	92.4192	1.4912	-0.6856	0.05	-22.2853	1.227	0.2816	0.4	OK	
	-15	453.2856	5.6	LPE 16:0	254.733	1.6652	-0.6539	0.1	22.7943	0.331	0.0995	0.6	OK	
	-20 -15 -10 -5 0 5 10 15 t[1]	481.317	6.3	LPE 18:0	164.2111	1.3974	-0.6323	0.1	100.2655	1.479	-0.2972	0.1	OK	
er	weight population compared to lean (L) population.	439.3049	5.8	LPE O-16:0	211.3618	1.7429	-0.6604	0.04	2.3997	0.994	0.3101	0.9	OK	
	Note: Positive/negative value of per cent of change	437.2904	5.8	LPE P-16:0	195.3989	1.7312	-0.6984	0.04	-29.7604	0.921	0.3264	0.2	OK	
	· ·	334.214	4.7	LTB5	83.9859	1.4035	-0.5714	0.1	-7.3469	0.398	0.0464	0.9	OK	
	means higher/lower AUC of postprandial change of	334.2141	4.5	LTB5	216.514	1.9246	-0.6779	0.07	-9.8857 51.4121	0.335	0.0229	0.8	OK	
	metabolite level in obese in comparison to lean	743.5467 686.5363	10.2 8.5	PE 36:2 SM 33:2	59.07282 36.8755	1.2908 1.0984	-0.5767 -0.5048	0.2	51.4131 -24.0824	0.979	-0.2149 0.1741	0.2	OK OK	
	individuals, S - identity of these metabolites was	284.2712	6.3	Stearic acid	125.2366	1.7462	-0.7585	0.03	22.2998	1.064	-0.1803	0.5	OK	
	confirmed by analysis of the standard. The	330.2402	3.8	Trihydroxyoctadecenoic acid	133.0237	1.4716	-0.6404	0.05	6.2266	0.388	-0.1803	0.9	Frag	
	percentage differences in area under curves	168.0282	0.2	Uric acid	144.523	1.4009	-0.6278	0.1	-15.5219	0.304	-0.0718	0.6	OK	
	(AUCs) of postprandial metabolite concentrations in	273.2662	4.1	Hexadecasphinganine	-60.2866	1.5247	0.55948	0.02	-8.2612	0.304	0.0667	0.8	Frag	
	the obese/overweight men compared with the Lean	287.2456	5	Lauroyldiethanolamide	-70.8197	2.436	0.78004	0.002	-3.6148	0.254	0.0843	0.9	Frag	
H		467.3007	5.1	LPC 14:0	185.2085	1.8755	-0.6471	0.002	1.6837	0.234	-0.1421	0.9	ok	
	(L) men are presented, unless otherwise indicated.	495.3317	5.6	LPC 16:0	161.3187	1.6212	-0.7003	0.02	-23.1161	0.181	-0.0415	0.6	ok	
	The <i>P</i> (corr)—predictive loading value and VIP value	523.3631	6.3	LPC 18:0	92.2312	1.3218	-0.6183	0.06	14.276	0.751	-0.2775	0.6	ok	
	were calculated based on respective PLS-DA	507.3684	6	LPC 18:1	180.6202	1.443	-0.5378	0.09	-22.2036	0.665	0.2166	0.5	ok	
	models. Variables with VIP >1.0 and absolute	517.3162	5.2	LPC 18:3	60.05464	1.4165	-0.6009	0.2	120.1516	0.456	-0.0988	0.2	ok	
	P(corr) > 0.4 were considered significant. DG,	549.3789	6.3	LPC 20:1	274.3281	1.8653	-0.5479	0.03	-36.4529	0.335	0.0231	0.6	ok	
	diglyceride; HC, high carbohydrate; HDoHe,	547.3632	6	LPC 20:2	37.03726	1.0398	-0.5139	0.2	53.2641	0.877	-0.3437	0.3	ok	
		503.3008	5.7	LPE 20:3	-71.0096	2.1767	0.5299	0.06	15.201	0.428	0.1244	0.8	ok	
Н	hydroxydocosahexaenoic acid; HETE,	477.3213	5.7	LPE P-19:1	132.7315	1.522	-0.6711	0.03	-22.9944	0.194	-0.0237	0.5	ok	
	hydroxyeicosatetraenoic acid; NC, normo-	271.2509	4.5	Sphinganine	-39.6883	1.3409	0.52921	0.02	11.5367	0.544	0.0982	0.2	ok	
	carbohydrate; LPC, lysophosphatidylcholine; LPE,	287.282	4.2	Sphinganine	-49.0484	1.648	0.50405	0.02	46.6706	0.841	-0.2708	0.8	ok	
	lysophosphoethanolamine; LTB5, leukotrienes B5;	279.2558	5.5	Linoleamide	681.7416	3.60911	-0.77010	0.03	38.4297	0.7851	-0.4016	ESI+	OK	
	PLS-DA, partial least squares discriminant analysis;	297.30246	7.8	Palmitoylethanolamide	69.2	1.67965	-0.501752	0.06	35.8	0.6701	-0.0106	ESI+	Frag	
	RT, retention time; SM, sphingomyelin; VIP, variable	295.25055	5.7	Sphingosine	550.0194	2.5319	-0.54635	0.2	56.6863	0.6979	-0.2706	ESI+	OK	
	importance into projection.	327.31308	7.3	Stearoylethanolamide	42.3067	1.23199	-0.50195	0.06	-47.8736	1.8172	0.4021	ESI+	Frag	
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Conclusion

Obvious postprandial metabolic differences were noted between lean and OW/OB individuals receiving HC meal. Metabolic pathways differentially affected by the HC meal belong to phospholipids and glucose metabolism, sphingomyelin signaling, as purines/pyrimidines and aldosterone catabolism.

Acknowledgement

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