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Effect of PNPLA3 rs738409 genotype and GDM history on fasting glucagon levels in early NAFLD



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BACKGROUND & AIMS:

The PNPLA3 rs738409 G/G risk genotype is associated to development and progression of NAFLD.

Fasting glucagon levels are higher in both diabetes mellitus NAFLD. There is no data available about the effects of rs73840 genotype or pGDM history on (fasting) glucagon levels.

We assessed the intraheptic lipid content (IHCL) and the pl glucagon levels in a diabetes prone young-middle aged for population in context of the *PNPLA3* genotype and GDM history.

Hypothesis

- GDM history and PNPLA3 rs738409 genotype have an additive on NAFLD development
- GDM history and PNPLA3 rs738409 genotype have an effect on plasma glucagon levels

PATIENTS & METHODS (& STUDY DESIGN):

GDM-genetic association study (2012-2016)*

Follow-up of mothers with information about in the prior pregnancy, incl. routine 75g OGTT $(24-28^{th}gw)$

Targeted enrollment with known PNPLA3 rs738409 C/C or G/G homozygous genotypes

Volunteers: 39 women without using any antidiabetic or lipid lowering drug

- "Routine" clinical laboratory measures (iver enzymes, HbA_{1c}, lipids)
- 75g OGTT: 0'-30'-120', plasma glucose (PG), insulin (using CLIA), 0'-30' glucagon (EURIA Glucagon kit)
- Clinical data and medical history
- Abdominal MR spectroscopy (MRS) & MRI (Multi-echo Dixon methods) measurements for liver&pancreatic proton density fat fraction (PDFF) + routine abdominal MRI (Siemens 3T Prisma MR)
- Patients with elevated liver enzymes were screened for alternative etiology.
- Liver fibrosis scores (NFS, Fib-4) were calculated.
- MW-U, SRO, ANOVA/K-W and post hoc tests were used (Statistica program).

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STUDY POPULATION:

o the	Table 1: Study population's	mean (SD) or
	characteristic at enrollment	n/n
s and)9 risk	Age (yrs)	37.2 (4.8)
	BMI (kg/m²)	28.2 (6.8)
	Time since index pregnancy (yrs) (tot. pop.)	3.5 (1.3)
lasma emale	Time since 1 st GDM pregnancy (yrs) (pGDM group only)	6.1 (4.4)
	pGDM/total	22/39
	rs738409 genotype (CC/GG)	27/12
	DM+prediabetes - at enrollment	
effect	before follow-up 75g OGTT - (IGT,	0/39
	IFG and/or HbA _{1c} criteria) / total	

Table 2: Entire study		
population	n/n or mean	
Clinical data (after study	(SD)	
assesment)		
No of pts with NAFLD	14/39	
(IHCL≥5.5%) / total		
No of pts with HOMA-IR>2.5**	19/39	
/total	13/33	
HOMA-B (%)	154 (81.9)	
DM+prediabetes		
(IGT, IFG and/or HbA1c criteria) /	14/39	
total		
FIB4 ≥1.45 / total	0/38	
NFS	34/5/0	
(<-1.455/≤-1.455<0.675/≥0.675)	54/5/0	

RESULTS:



Ref.: *Rosta K et al. (2017) Association Study with 77 SNPs Confirms the Robust Role for the Mellitus Development, PLOS ONE 12(1): e0169781, https://doi.org/10.1371/iournal.pone.016









CONCLUSION:

- We confirmed the hypothesis that the GDM history and the PNPLA3 rs738409 gene variant have an additive effect on NAFLD development
- The fasting plasma glucagon levels were increased both in the risk and non risk genotype groups, however
- the track of the correlation between the fasting glucagon levels and the IHCL was modified by the rs738409 genotype
- We could not detect significant effect of prior GDM on the fasting glucagon levels with this limited sample size.
- To our knowledge we first report a significant negative correlation between the fasting plasma glucagon and serum HDL levels and
- HDL levels were also inversely correlated with the IHCL values only in patients with rs738409 C/C genotype

