

# Genetic variants and polygenic risk score associated with the HDL-c response to statin treatment: a GoDARTS study

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Introduction	Results
	Paired Profiles for (B_hdl, H
<ul> <li>Stating mainly act on the reduction of low-</li> </ul>	8 - Mean

Results	
Paired Profiles for (B_hdl, hdl_max_A)	Distribution of Difference: B_hdl - hdl_max_A With 95% Confidence Interval for Mean
8 Mean 8	
	Kernel

## **Discussion and conclusion**

• This study shows statins also helps to improve high-density lipoprotein-cholesterol (HDL-c) levels up to 20% in the study population.

- density lipoprotein-cholesterol (LDL-C) levels.<sup>1</sup>
- Studies have shown that statin therapy also helps in improving high-density lipoproteincholesterol (HDL-c) levels up to 10-15%.<sup>2</sup>
- Inter-individual variation in HDL-c response to statins therapy could be partially explained by genetic variation.
- A recent meta analysis suggested only *CETP* locus for with common genetic variants that influence HDL-C response to statins.<sup>3</sup>
- Global Lipids Genetics Consortium (GLGC)
   2013, has identified 80 genetic variants associated with HDL-c levels.<sup>4</sup>

## **Study Objectives**

• To investigate genetic variants associated



Table 2: Variants associated with HDL-c response (Adjusted for age, sex, dose, Baseline HDL-c, treatment duration)

Parameter	Estimate	Standard Error	t Value	P value
rs247616 ( <i>CETP</i> )	0.011	0.003	3.71	0.0002
rs1532085 ( <i>LIPC</i> )	0.011	0.002	-4.08	<.0001



- Individual genetic variants shows significant positive association with the HDL-c response after adjusting for phenotypic traits.
- Overall effect of PRS with HDL-c response is comparatively less then baseline HDL-c. This suggests that some gene variants differentially contribute to baseline HDL-c levels and HDL-c response.

## Way forward

- Preliminary data suggested that HDL profile between the two population [Scottish  $(1.20\pm0.33)$  and India  $(1.04\pm0.23)$ ] were significantly different (p value <0.001)<sup>5.</sup> Hence, genetic differences needs to be investigated.
- GWAS using Affymetrix, Illumina, and Broad (genetically adjusted) and meta-analysis will be carried out to find the novel loci in MDRF and GoDARTS data.

- with the HDL-c response to statin treatment in GoDARTS cohorts
- To construct and assess the effect of a polygenic risk score (PRS) for HDL-c response in the study population

#### **Study Methodology**

Study population and sample size	<ul> <li>10,633 statin users in GoDARTS cohorts</li> </ul>			
Inclusion criteria	<ul> <li>At least, one off treatment HDL-c level and at least one on- treatment level</li> </ul>			
Exclusion criteria	<ul> <li>Subjects with missing on- or off- treatment measurements</li> </ul>			
Study outcome	<ul> <li>Change in HDL-c (mmol/L) levels</li> </ul>			



#### Graph 2: Normal distribution of Polygenic risk score



Graph 3 (a & b): Regression of Baseline HDL-c (a) and response of HDL-c (b) with Polygenic risk score

Table 3: Effect of PRS on baseline HDL-c (adjusted for age, sex) and HDL-c response (Adjusted for age, sex, dose, baseline HDL- c, treatment duration) (n = 8,271)						
Outcome	parameter	Estimate	S E	P value	Adjusted	

					R square
Baseline HDL-c	PRS	0.137	0.017	<0.001	0.08
HDL-C response	PRS	0.01	0.008	0.0175	0.28

Statin induced elevation in HDL-c levels were (0.27±0.32; t-test p-value <0.001) observed (table 1; graph 1).</li>

- Conditional GWAS will be carried to adjust for variants affecting purely baseline HDL-c levels.
- Discovered novel loci for real pharmacogenetic drug response will be used for polygenic risk score in the study population.

## References

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#### Study predictor

• Polygenic risk score (Top 22 SNPs for HDL-c from GLGC 2013 )

#### Results

Table 1: Difference between before and after HDL-c value (Paired T test )						
Ν	Mean (SD)	Min	Max	t Value	P value	
10,633	-0.27(0.32)	-5.58	1.46	-87.97	<0.0001	

- Among all reported SNPs, rs3764261 (*CETP*), and rs1532085 (*LIPC*) were among few which significantly associated with raise in HDL-c levels (adjusted) (table 2).
- PRS has a significant effect on Baseline HDL-c levels (R square = 0.015) (table 3; graph 3).

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