



Formal Comment

Comment On: Valette et al. Melanocortin-4 Receptor Mutations and Polymorphisms Do Not Affect Weight Loss after Bariatric Surgery. *PLOS ONE* 2012; 7(11):E48221

David Meyre^{1*}, Philippe Froguel^{2,3}, Fritz F. Horber^{4,5}, John G. Kral⁶

1 Department of Clinical Epidemiology and Biostatistics, McMaster University, Hamilton, Ontario, Canada, **2** Centre National de la Recherche Scientifique (CNRS) Unité Mixte de Recherche (UMR) 8199, Lille Pasteur Institute, Lille, Nord, France, **3** Department of Genomics of Common Disease, Imperial College London, London, United Kingdom, **4** Department of Internal Medicine, Landesspital, Vaduz, Liechtenstein, **5** University of Berne, Berne, Switzerland, **6** Department of Surgery, State University of New York Downstate Medical Center, Brooklyn, New York, United States of America

To the Editor

We read with interest the *PLOS ONE* article by Valette et al. in 2012 assessing associations between gene variations at the melanocortin 4 receptor (MC4R) locus and weight loss after bariatric surgery in a French longitudinal cohort [1]. We were the first to describe effects of *MC4R* gene variations on outcomes of bariatric surgery [2] and have several concerns about the current study.

There seems to be a genotyping error of the rs17782313 variant downstream of *MC4R*. As the authors acknowledge, the minor allele frequency (MAF) of this variant was 0.27 in 1443 Swedish bariatric surgery patients [3] which is comparable to the MAF of 0.30 we found in 1274 severely obese similar Swiss patients [4]. In the presence of Hardy-Weinberg equilibrium, MAF of 0.27 for the variant rs17782313 should yield 345 homozygous TT carriers, 256 heterozygous TC carriers and 47 homozygous CC carriers among the 648 patients described by Valette et al. [1]. However, their genotypic distribution is 641 TT, 4 TC and 3 CC carriers implying an error in their genotype count. There are various causes of genotyping artifacts including poor DNA quality, genotyping of different polymorphisms, technical faults, subjective genotype classification, and human errors during genotyping or data transfer [5]. In the absence of information about the precise method used to genotype the rs17782313 polymorphism in the paper it is difficult to assess potential causes of the discordant genotype distribution. The authors failed to describe any quality control procedures to ensure the integrity of their genotyping such as: call rate, Hardy-Weinberg equilibrium test, double-genotyping concordance rate and comparison of the MAF with published databases in comparable ethnic groups [5]. The very significant departure of the genotypic distribution of the variant rs17782313 described by Valette et al from Hardy-Weinberg equilibrium ($P < 10^{-20}$) suggests an error in technique.

We fully acknowledge the value of a matched case control design to longitudinally study responses to bariatric surgery, but such design requires sufficient phenotypic data to interpret outcomes. Valette et al. did not seem to assess associations between *MC4R* genetic variants with preoperative metabolic parameters known to affect surgical outcomes. Potentially, relevant metabolic

phenotypic differences at baseline between carriers and non-carriers of *MC4R* genetic variants might have clarified the findings of their study leading to different conclusions.

The authors mentioned that the analysis of a 'large group of *MC4R* mutation carriers' was one of the strengths of their study. We respectfully disagree with this statement. On the contrary, the small number of subjects in the different *MC4R* genotype groups (e.g. 9 carriers of loss-of-function mutations or 10 carriers of gain-of-function mutations) constitutes a major limitation of this study. The fact that the authors only demonstrated a nominally significant association between the more prevalent *MC4R* genotype group (22 carriers of the -178 A/C polymorphism) and surgical outcomes argues for an overall lack of statistical power in the study.

Lastly, the stated purpose of the paper was to evaluate surgical weight loss in patients with *MC4R* mutations and polymorphisms. Unfortunately the authors chose to do so as early as 12 months postoperatively, before the majority of patients reach weight loss nadir. The rapid weight loss phase after gastric banding and bypass is approximately 9–18 months, with differences in trajectory between the two [6]. By studying patients before they reached a steady, homeostatic state, Valette et al. abrogated their ability to analyze relationships between genetic variants and complications, reoperations, clinically meaningful effects on serious comorbidities or weight regain, altogether undermining the relevance of the study.

Author Contributions

Conceived and designed the experiments: DM PF FH JK. Wrote the paper: DM PF FH JK.

Citation: Meyre D, Froguel P, Horber FF, Kral JG (2014) Comment On: Valette et al. Melanocortin-4 Receptor Mutations and Polymorphisms Do Not Affect Weight Loss after Bariatric Surgery. *PLOS ONE* 2012; 7(11):E48221. PLoS ONE 9(3): e93324. doi:10.1371/journal.pone.0093324

Editor: Franco Folli, University of Texas Health Science Center at San Antonio, United States of America

Received: January 21, 2014; **Accepted:** February 27, 2014; **Published:** March 31, 2014

Copyright: © 2014 Meyre et al. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Funding: No specific funding for this Formal Comment.

Competing Interests: The authors have declared that no competing interests exist.

* E-mail: meyre@mcmaster.ca

Formal Comments are critiques of specific published articles.

References

1. Valette M, Poitou C, Le Beyec J, Bouillot JL, Clement K, et al. (2012) Melanocortin-4 receptor mutations and polymorphisms do not affect weight loss after bariatric surgery. *PLoS one* 7: e48221.
2. Potoczna N, Branson R, Kral JG, Pic G, Steffen R, et al. (2004) Gene variants and binge eating as predictors of comorbidity and outcome of treatment in severe obesity. *J Gastrointest Surg* 8: 971–981; discussion 981–972.
3. Sarzynski MA, Jacobson P, Rankinen T, Carlsson B, Sjostrom L, et al. (2011) Associations of markers in 11 obesity candidate genes with maximal weight loss and weight regain in the SOS bariatric surgery cases. *Int J Obes (Lond)* 35: 676–683.
4. Stutzmann F, Cauchi S, Durand E, Calvacanti-Proenca C, Pigeyre M, et al. (2009) Common genetic variation near MC4R is associated with eating behaviour patterns in European populations. *Int J Obes (Lond)* 33: 373–378.
5. Li A, Meyre D (2013) Challenges in reproducibility of genetic association studies: lessons learned from the obesity field. *Int J Obes (Lond)* 37: 559–567.
6. Kral JG (2006) ABC of obesity. Management: Part III—surgery. *Bmj* 333: 900–903.