



## Genetic Predisposition and Environment Factors

Bob Mareley\*

Department of Endocrinology, Portland Metropolitan Geriatric Medical Center, Portland

\*Corresponding author: Bob M, Department of Endocrinology, Portland Metropolitan Geriatric Medical Center, Portland; E-mail: bobm123@uw.com

Received date: May 04, 2021; Accepted date: May 19, 2021; Published date: May 26, 2021

### Introduction

In male progeny one-half of the additive genetic variance from sex linked genes is contributed to the dams within sire component and one fourth to the sire component. In female progeny additive genetic variance from sex-linked genes is contributed to the sire component only. Comparison of dam vs. sire component estimates of heritability made separately within male and feminine progeny, and comparison of sire component estimates between male and feminine offspring, may thus indicate the possible involvement of sex-linked genes on the variation within the trait. In both male and feminine offspring the estimate of heritability was larger from the dam within sire component than from the sire component, suggesting the presence of maternal variance and/or no additive, autosomal genetic variance.

The dam within sire component estimate of heritability was substantially larger in male than in female offspring, in accordance with expectation if sex-linked genes influenced variation in the trait. It is our conviction that an entire understanding of the etiology of most psychiatric disorders would require an understanding of the relevant genetic risk factors, the relevant environmental risk factors, and therefore the ways during which these two risk factors interact. Such understanding will only arise from research during which the important environmental variables are measured during a genetically informative design. Such research would require a synthesis of research traditions within psychiatry that have often been at odds with each other within the past.

This interaction between that research tradition has focused on the genetic etiology of psychiatric illness which has emphasized environmental causation will undoubtedly be to the advantage of both. The aim of this study was to research aging-related changes within the contribution of genetic and environmental influences to hand-grip strength in late adulthood. Subjects during this study are 152 intact twin pairs (77 monozygotic and 75 dizygotic pairs) from the National Heart, Lung, and Blood Institute Twin Study assessed repeatedly for hand-grip strength at mean ages of 63 and 73 yr. Structural equation

genetic modeling was used to investigate stability and alter within the genetic and environmental components of variance of hand-grip strength in late adulthood. Average decline in strength over the 10 years of follow-up and was highly significant ( $P = 0.003$ ). The test-retest correlation between baseline and follow-up grip strength was 0.62 ( $P < 0.001$ ). Heritability for weight, height, and BMI was low at birth (between 6.4 and 8.7% for boys, and between 4.8 and 7.9% for girls) but increased over time, accounting for on the brink of half or more of the variance in weight and BMI after 5 months aged in both sexes.

Common environmental influences on all body measures were high at birth (between 74.1%-85.9% altogether measures for boys, and between 74.2 and 87.3% altogether measures for girls) and markedly reduced over time. For body height, the effect of the common environment remained significant for a extended period during infancy (up through 12 years of age). Sex-limitation of genetic and shared environmental effects was observed. Approximately one-third of Chinese obese children are often classified as MHO. Both genetic predisposition and environment factors and their interaction contribute to the prediction of MHO status. This study provides novel insights into the heterogeneity of obesity and has the potential to impact the optimization of the intervention options and regimens within the management of pediatric obesity. Genetic technologies are offering new solutions for disease control, prevention and cure. They are now getting used to diagnose and treat complex diseases like heart condition, asthma, diabetes and cancer. Genetic technologies can also soon allow us to eradicate malaria, a serious health menace in many developing countries. Introduced species aren't always disruptive to an environment, however. Tomás Carlo and Jason Gleditch of Penn State University found that the amount of "invasive" honeysuckle plants within the area correlated with the amount and variety of the birds in the Happy Valley Region of Pennsylvania, suggesting introduced honeysuckle plants and birds formed a interdependent relationship. Presence of introduced honeysuckle was related to higher diversity of the bird populations therein area, demonstrating that introduced species aren't always detrimental to a given environment and it is completely context dependent. For many reasons, children are likely to be more vulnerable than adults to the consequences of environmental contaminants. To better understand the consequences of those exposures on children's health, the NIEHS/EPA Children's Environmental Health and Disease Prevention Research Centers (Children's Centers) were established to explore ways to reduce children's health risks from environmental factors. The program is jointly funded by the National Institute of Environmental Health Sciences (NIEHS) and by EPA through the Science to realize Results (STAR) grants program.