Author Correction: Exome sequencing of Finnish isolates enhances rare-variant association power

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In this Article, several errors have been drawn to our attention. In the author list, some of the affiliation numbers were incorrect (see PDF version of the original Article for affiliation numbering): Robert S. Fulton's affiliations should be 2,13 rather than 2; Hannele Laivuori's affiliations should be 8,35,36 rather than 7,35,36; Chiara Sabatti's affiliations should be 10,39 rather than 9,39; Aarno Palotie's affiliations should be 8,40,41 rather than 7,40,41; and Samuli Ripatti's affiliations should be 8,11,41 rather than 7,11,41. In addition, the faculty of affiliation 36 should read 'Faculty of Medicine and Health Technology' rather than 'Faculty of Medicine and Life Sciences'. In the main text on page 326, the protein consequence for the associated *KRT40* variant, listed as 'Ser32Pro', should be 'Ser328Pro'. In the legend for Extended Data Fig. 7, the definitions of SuK and Nfi were missing and have been added. To clarify our description of the Finnish map in Fig. 3, the last line of the legend should read 'Birthplaces of carrier and non-carrier individuals were plotted on a map of Finland, including regions of Finland, later ceded, as they existed before the Second World War.' rather than 'Birthplaces of carrier and non-carrier individuals were plotted on a map of Finland, including regions that were ceded before the Second World War.'. In the 'Apolipoprotein B' section of the Supplementary Information, the protein consequence for the associated *APIM2* variant, listed as '418Asn', should be 'Tyr418Asn'. In the 'Multiplicity adjustment procedure' section of the Supplementary Information, we corrected the annotation in step II. (a) to correctly identify the ordered *P* values (*m*) for all 64 traits at each variant. These errors have all been corrected online.