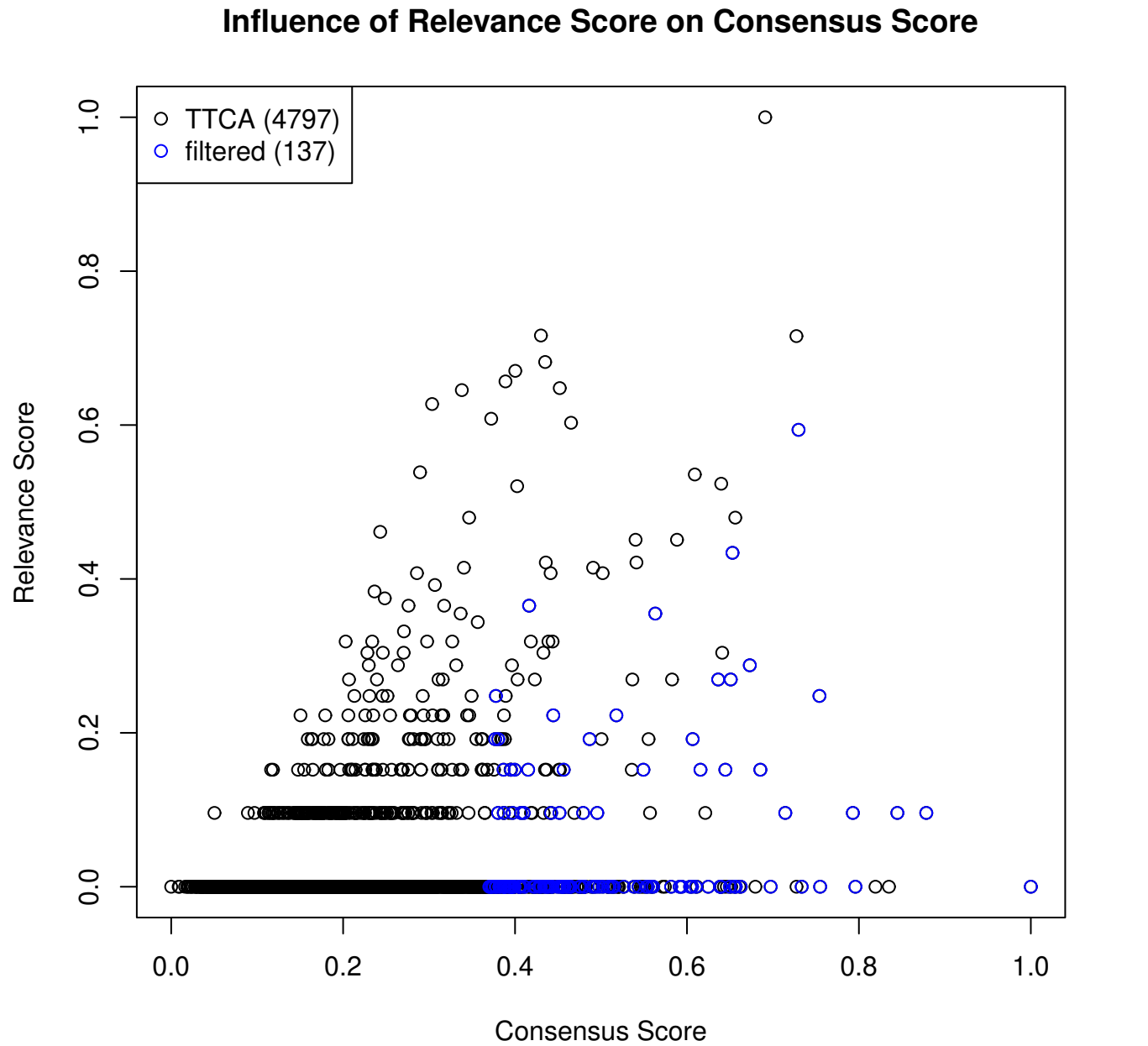

Supplemental Figure 2: The correlation between relevance and consensus score. Among the 137 filtered genes (blue) with high consensus scores, some have high relevance score, meaning that publications in PubMed already associate the gene with the term 'fat distribution'. Other genes have no co-occurences with the term 'fat distribution' and thus represent novel gene candidates. The pearson correlation coefficient for the filtered genes is 0.23.