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Supplemental information

An integrative framework to prioritize genes

in more than 500 loci associated

with body mass index

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536 BMI-associated variants + proxies in LD $r^2 \ge 0.8$, locus distance $\pm 1 \text{ MB}$

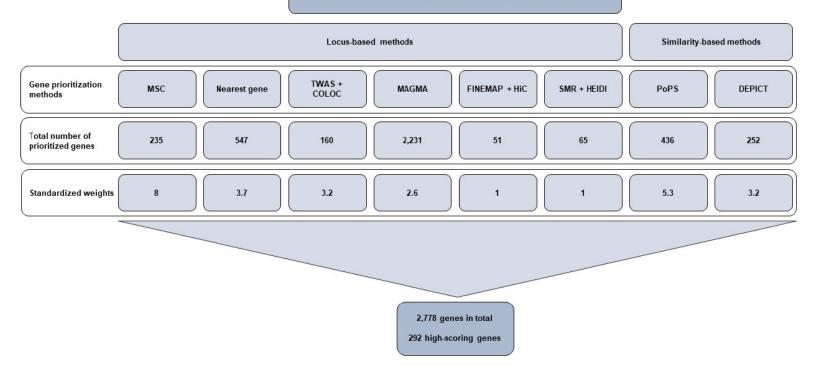


Figure S1. Flowchart of pipeline, number of genes prioritized and contribution (weights) of each method to the overall scoring of prioritized genes.