



Supplemental Fig.1. Map locations of QTL for seed oil content, silique length, seeds per silique, and seed weight in all environments. For simplicity only the markers in the QTL confidence intervals, along with the QTL-containing chromosomes are show. The color of the lines was used to distinguish the different traits located in the same region, and the red scale bars at the top left of each chromosome are 1 cM.