



Supplemental Figure 3. Plots of the unconditional maximum likelihood point estimates from the 13 mixed stock analyses and the 95% confidence intervals. Point estimates indicate the estimated proportion of individuals in that group that have had source contributions from each of the four wild population genetic clusters. The 13 classes from left to right are: Total, INRA-HA, INRA-RHA, HA-Non-Oil, HA-Oil, RHA-Non-Oil, RHA-Oil, Non-Oil Introgressed, Oil Introgressed, Landrace, OPV, Other Non-Oil, Other Oil. *Note that overlap on error bars is difficult to see, please see Table 3 in the manuscript for detail on confidence interval overlap.*

Genetic diversity and population structure in cultivated sunflower and a comparison to its wild progenitor, *Helianthus annuus* L.

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