Figure S1. Profiles of seed fatty acid composition of WT, wri1-1 and wri1-1 expressing HA-EgWRI1 and HA-AtWRI1. Six independent transgenic lines overexpressing HA-EgWRI1 (#1-4, #2-3, #11-1, #13-5, #14-1, and #16-3, respectively; from left to right) and five independent transgenic lines overexpressing HA-AtWRI1 (#1-2,#6-5,#7-2,#8-3, and #9-4, respectively; from left to right) are shown above. Results are means ± SE (n = 3-4).