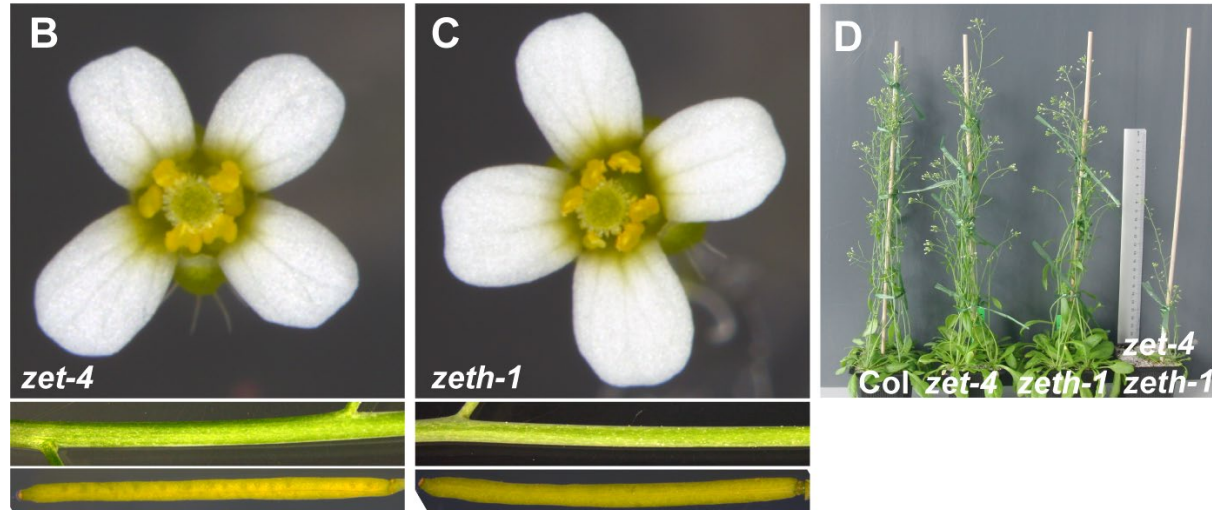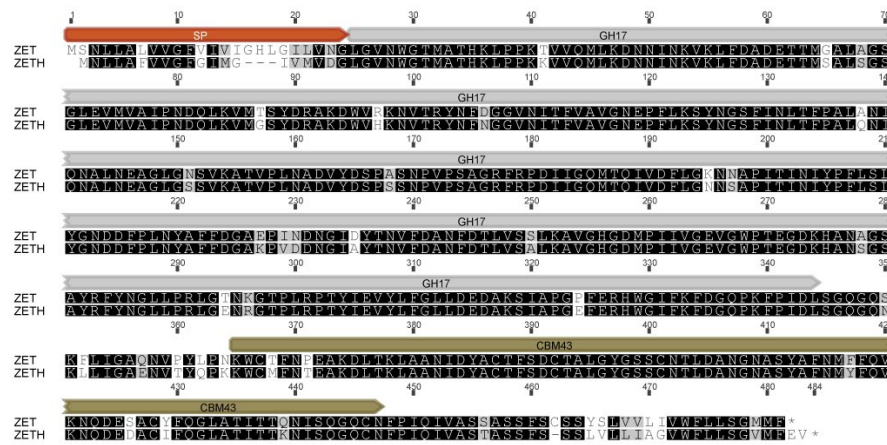1   **Asymmetric redundancy of *ZERZAUST* and *ZERZAUST HOMOLOG* in different**

2   **accessions of *Arabidopsis thaliana***

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4   **Supplementary data**

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7   **Fig. S1.** A) Alignment of ZET and ZETH amino acid sequences. B) *zet-4*. C) *zeth-1*. Flower,

8   stem and silique morphology is essentially normal. (D) Whole-plant appearance. Genotypes

9   are indicated.

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Fig. S2. Evolutionary history of ZET (AT1G64760) and ZETH (AT2G19440).

Phylogenetic tree showing the evolution of the ZET and its closest homolog (ZETH). The

tree shows that duplication is specific to Arabidopsis and it is not present in any other

Brassicaceae members. This phylogenetic analysis is adapted from the source:

http://ensembl.gramene.org. Full version of the tree is available at https://bit.ly/2CxlEWn.

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18   **Supplementary Table S1. Primers used in this study.**

Primer Name	Sequence (5'-3')
ZETHCol_F	ATCCCGGGCTAGGTGTCAACTGGGGAACAATG
ZETHCol_R	TAGGATCCCATATTATGCAATTTAATACATGG
ZETHL*er*_F/	ATCCCGGGCTAGGTGTGAACTGGGGAACAATG
ZETHL*er*_R	TAGGATCCATCACGTACATTAACTGCCGTTAG
ZETqRT_F	TCAAGATGAGAGTGCTTGCTATTTT
ZETqRT_R	CACAACCAAAGAATAAGACGAACAAGA
ZETHqRT_F	CCAGGATGAGGATGCTTGTATCTTC
ZETH qRT_R	TCCTGATAAGAGAAACCAAACTCCAGC
DT1-F0_ZETH	TGTAATGATCAGCTTAAGGTTGTTTTAGAGCTAGAAATAGC
DT2-R0_ZETH	AACAACCTTAAGCTGATCATTACAATCTCTTAGTCGACTCTAC
DT1-BsF_ZETH	ATATATGGTCTCGATTGTAATGATCAGCTTAAGGTTGTT
DT2-BsR_ZETH	ATTATTGGTCTCGAAACAACCTTAAGCTGATCATTACAA

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