

Figure S1: Genevestigator Arabidopsis XTH gene expression in response to light quality manipulations. The heat map was constructed using data compiled from relevant microarray experiments as reported by Genevestigator. Expression values are depicted as log2 ratios indicating fold changes and colour coded to indicate up (red) and down (green) regulation relative to controls.

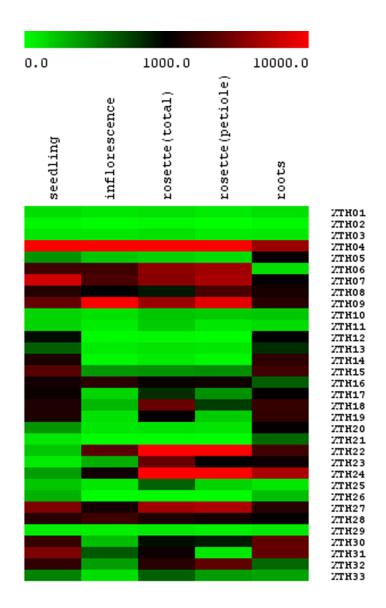


Figure S2: Genevestigator XTH organ-specific expression. Expression data was compiled from relevant microarray experiments as reported by Genevestigator. Data are depicted as absolute expression values and colour coded to indicate high (red) and low (green) expression values.

Table S1: *XTH* genes for which *XTHpro*::*GUS* reporter lines were available were screened for shade-induced differential petiole growth.

Gene name	AGI ID	GUS staining in petioles**
AtXTH1	At4g13080	None
AtXTH5	At5g13870	None
AtXTH8	At1g11545	None
AtXTH9	At4g03210	+
AtXTH12	At5g57530	None
AtXTH14	At4g25820	None
AtXTH15	At4g14130	+
AtXTH18	At4g30280	None
AtXTH19	At4g30290	+
AtXTH20	At5g48070	None
AtXTH21	At2g18800	None
AtXTH23	At4g25810	None
AtXTH24	At4g30270	None
AtXTH25	At5g57550	None
AtXTH27	At2g01850	None
AtXTH28	At1g14720	None
AtXTH29	At4g18990	None
AtXTH32	At2g36870	None
AtXTH33	At1g10550	None

<sup>\*\*</sup> Based on staining in petioles from plants treated for 24 h in shade relative to petioles from plants treated for 24 h in white light