Supplementary Data Set S1. The list of sequences of the GH16 family and their properties, such as tissue localisation and suggested function.

Organism (plant/yeast)	Gene name	Accession	Tissue localisation	Suggested function	Reference	Entry name used in Figure 2C
Arabidopsis thaliana	XTH3	Q9LJR7	Predominantly expressed in flower buds	XET activity/ CW biogenesis	[121, 184]	Q9LJR7 ARATH
Arabidopsis thaliana	XTH31	P93046	Predominantly expressed in roots	XEH activity/ CW biogenesis	[184 - 187]	P93046 AtXTH31
Hordeum vulgare	HVXEA	P93671	Predominantly expressed in first leaf base	XET activity/ CW biogenesis	[124, 140, 152, 161]	P93671 HvXET3
Hordeum vulgare	HVXEB	P93672	Predominantly expressed in spike	XET activity/ CW biogenesis	[124, 140, 152, 161]	P93672 HvXET4
Hordeum vulgare	EXT	P93668	Protein isolated from 7-days-old seedlings	XET activity/ CW biogenesis	[97, 188]	P93668 HvXET5
Hordeum vulgare	XET6	B1P1S7	Predominantly expressed in flower at anthesis	XET activity/ CW biogenesis	[120, 124]	B1P1S7 HvXET6
Tropaeolum majus	N/A	Q07524	Expressed only in cotyledons	XEH activity/ CW biogenesis	[137, 159, 189]	Q07524 2UWA NXG1
Tropaeolum majus	N/A	2UWC	Expressed only in cotyledons	XEH activity/ CW biogenesis	[137, 189]	2UWC NXG2
Tropaeolum majus	XET1	Q41614	Predominantly expressed in young epicotyls and roots	XET activity/ CW biogenesis	[159]	Q41614 TmXET1
Tropaeolum majus	xet6.3	V5ZEF7	Protein found only in germinating seeds	XET activity/ CW biogenesis	[122]	V5ZEF7 TmXET6.3
Equisitum fluovatile	HTG	A0A0K2WA62	In shoots	HTG activity/ CW biogenesis	[134, 135]	A0A0K2WA62 EfHTG
Equisitum fluovatile	EfXTH-A	MT495433*	N/A	XET activity/ CW biogenesis	[116]	EfXTH-A
Equisitum fluovatile	EfXTH-H	MT495434*	N/A	XET activity/ CW biogenesis	[116]	EfXTH-H
Equisitum fluovatile	EfXTH-I	MT495435*	N/A	XET activity/ CW biogenesis	[116]	EfXTH-I
Populus tremulus x tremuloides	XET16A	Q8GZD5	In vascular tissues	XET activity/ CW biogenesis	[101, 126]	Q8GZD5 PttXET16A
Pinus radiata	XTH1	A0A059SVJ4	In stems (inclined)	XET activity/ CW biogenesis	[155, 190]	A0A059SVJ4 PrXET1
Saccharomyces cerevisiae	CRH1	P53301	Fungal CWs	Transfer of chitin to 1,6-β-glucan / CW biogenesis	[191 - 193]	P53301 CRH1
Saccharomyces cerevisiae	UTR2	P32623	Fungal CWs	Transfer of chitin to 1,6-β-glucan / CW biogenesis	[192 - 194]	P32623 CRH2

*GenBank nucleotide sequence accession number (BankIt2345959)
N/A, not available.

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