

S2 Fig. HM1 peptide sequence comparison of *hm1* alleles. Sequence comparison of HM1^A with HM1 from the resistant maize inbreds (B73, Va35, W22, and Pr1), maize cultivars (Enano and Pira), HM1 homologs [sorghum (*Sorghum bicolor*), rice (*Oryza sativa*), and barley (*Hordeum vulgare*)], and maize dihydroflavonol 4-reductase (DFR). Among the five amino acid substitutions present in HM1^A (highlighted in red and bold), the L116 residue was a conserved in all *Hm1* alleles and orthologs.

HM1-B73	-----MAEKESNGVRVCVTGGAGFIGS WLVRKLLEKGYTVHATLRNTGDEAKAGLLRR
HM1A	-----MAEKESNGVRVCVTGGAGFIGS WLVRKLLEKGYTVHATLRNTGDEAKAGLLRR
HM1-Va35	-----MAEKESNGVRVCVTGGAGFIGS WLVRKLLEKGYTVHATLRNTGDEAKAGLLRR
HM1-Pr1	-----MAEKESNGVRVCVTGGAGFIGS WLVRKLLEKGYTVHATLRNTGDEAKAGLLRR
HM1-W22	-----MAEKESNGVRVCVTGGAGFIGS WLVRKLLEKGYTVHATLRNTGDEAKAGLLRR
HM1-Enano	-----DEAKAGLLRR
HM1-Pira	-----DEAKAGLLRR
HM1-Sorghum	-----MAENESSGVRCVTGGAGFIGS WLVKLLEKGYTVHATLRNTGDEEKAGLLRR
HM1-Rice	MAEAGR-SGGVAGDGVRCVTGGAGFIASWLVKLLERGCIVHATLRSMGDEEKAGLLRR
HM1-Barley	MAEEGKGTGSNGGRGVACVTGGAGFIGS WLVRKLLEAGYTVHATLRSIGDEGKAGLLRG
DFR-Maize	-----MAGGGATVCVTGAGGFIASWVVKLLLSRGYTVHGTVRHLSDEKT-GHLKR
	S99Y D110Y
HM1-B73	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG S QYKSTAEAVV DA
HM1A	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG Y QYKSTAEAVV YA
HM1-Va35	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG Y QYKSTAEAVV DA
HM1-Pr1	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG Y QYKSTAEAVV DA
HM1-W22	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG S QYKSTAEAVV DA
HM1-Enano	LVPGAA--ERLRLFQADLFYAAATFAPAIAGCQFVFLVATPFGLDSAG S QYKSTAEAVV YA
HM1-Pira	LVPGAA--ERLRLFQADLFDAATFAPAIAGCQFVFLVATPFGLDSAG Y QYKSTAEAVM YA
HM1-Sorghum	LVPGAA--ERLRLFDADLFDAATFAPAIAGCQFVFLVATPFGLQDAGS SKY KSTAEAVV DA
HM1-Rice	LVPGAA--ERLRLFEADLFDAATFAPAIAGCQFVFLIATPYGLEASN SKY KNTADA AVDA
HM1-Barley	LVPGGAPPERLVLFEADIYDAASFAPAIAGCHFVFLVANPSAHE-PAS SKY KTSAA ADA
DFR-Maize	LENA---AGNLRIFKADLLDYDAMAAAVVGCGQGVFHATPVPSEDLT D PELQMLGPAT TG

L116H

HM1-B73 VRAILRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLNVDYPL
 HM1A VRAIHRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLNVDYPL
 HM1-Va35 VRAILRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLKVNYP
 HM1-Pr1 VRAILRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLNVDYPL
 HM1-W22 VHAIRQCEESRTVKRVIHTASVAAASPLLEEE-VPASGVGYRDFIDESCWTSLNVDYPL
 HM1-Enano VRAILRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLNVDYPL
 HM1-Pira VRAILRQCEESRTVKRVIHTASVAAASPLLEEE-VSASGVGYRDFIDESCWTSLNVDYPL
 HM1-Sorghum VRAILRQCEESRTVKRVIHTASVAAASPLKEEE-VS-GVIGYREFISESCWTSPDVDYPL
 HM1-Rice VREILRQCAESKTVKRVIHTASISTASPLIDVPGAGVGAAGYRDFIDESCWTPLDVDYPL
 HM1-Barley VRVILRLCAESRTVKRVIHTASVTAASPLTKSSS--AAAAYGDFISESCWTLLDVNYPL
 DFR-Maize TTNVLKA-ASSANVQRVVVVSSMVAVE-----ISPKDWPPEGKVRDERCWSDREFCRSI

S191N

HM1-B73 RSAHFDKYILSKLRSEQELLSYNGGESPAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1A RSAHFDKYILSKLRSEQELLYNNGGESPAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1-Va35 RSAHFDKYILSKLRSEQELLSYNGGESPAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1-Pr1 RSAHFDKYILSKLRSEQELLSYNGGESPAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1-W22 RSAHFDKYILSKLQSEQELLSYNNNGGESPAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1-Enano RSAHFDKYILSKLRSEQELLYNNGGESPAFEVVTPLGLVAGDTVLGCASETVESAVAPV
 HM1-Pira RSAHFDKYILSKLRSEQELLYNNGSESPEAFEVVTPLGLVAGDTVLGRAPETVESAVAPV
 HM1-Sorghum RSAHFDKYILSKVQSEQELLSYNGGESPAFEVVTLALGLVAGDTVLGRVPETVESAVSPV
 HM1-Rice RSAHFDKYVLSKMMSEKELLGYNDGEGRAFEVVTLPCLVAGDTVLGRAPETLENNAVSPV
 HM1-Barley RSVHFDKYIESKVLSEKELLSYNDGENPAFEVVTPLGLVGGNTVLGYLPETMESVVAPV
 DFR-Maize E----SWYPVAKIISEEAALAYG--QQTGLDVVTINPGLVFGPMLQPTVNTTIQFLI---

L240P

HM1-B73 SRSEPCFGLLRLIQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1A SRSEPCFGLPRILQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1-Va35 SRSEPYFGLPRILQQLLGSLPLVHVDDVCDALVFCMERRRSVAGRFLCAAAYPTIHDVVA
 HM1-Pr1 SRSEPCFGLPRILQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1-W22 SRSEPYFGLLRLIQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1-Enano SRSEPYFGLLRLIQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1-Pira SRSEPCFGLPRILQQLLGSLPLVHVDDVCDALVFCMERRPSVAGRFLCAAAYPTIHDVVA
 HM1-Sorghum SRNEAYFGLPRILQQLLGSLPLVHVDDVCDALIFCMERRHSIAGRFLCAAAYPTIHDVVG
 HM1-Rice SRNEPSFAFLRLLQRLVGSVPLVHADDVCDALVFCMDQ-PSLAGRFLCSAAAYPTIHDIVE
 HM1-Barley TKQEPCFMLPRILQQLLGSLVPLVHVDDVCAALIFCMEQ-PALSGRFLCAAAYPTIHDILD
 DFR-Maize -----YFLKGGPDPVKNLWHIVDVRDVADAMLLYEV-PEATGRHICAPHVISARDLLD

HM1-B73	HYASKFPHLDILKET-EAVATV---RPARDRLGELGFKYKGMEIELDSSVACAARLGSL
HM1A	HYASKFPHLDILKET-EAVATV---RPARDRLGELGFKYKGMEIELDSSVACAARLGSL
HM1-Va35	HYASKFPHLDILKET-EAVATV---RPARDRLGELGFKYKGMEIELDSSVACAARLGSL
HM1-Pr1	HYASKFPHLDILKET-EAVATV---RPARDRLGELGFKYKGMEIELDSSVACAARLGSL
HM1-W22	HYASKFPHLDILKETTEAVATV---RPARDRLGELGFKYKGMEIELDSSVACAARLGSL
HM1-Enano	-----
HM1-Pira	-----
HM1-Sorghum	HYARKFPHLDILKET-EAVARV---QPDGDRLGELGFVYKGIEIELDSSVACAARLGCL
HM1-Rice	HFAAKYPHLDVLKEPEREVARV---QPAADKLGELGFRYKGMEIELDGSVGCAARLGYI
HM1-Barley	HYGSKYPHLDLLRET-DEVARV---QPDKNKLGELGFYKGLEILDESIDCAVRLGSL
DFR-Maize	LLKSMYPDYPCLANE-SILD RDHPAPMTSDKLKKIGWSCR P-LEETIVDTVECCLRAGFL
HM1-B73	DASKLGLQKG-----
HM1A	DASKLGLQKG-----
HM1-Va35	DASKLGLQKG-----
HM1-Pr1	DASKLGLQKA-----
HM1-W22	DGSKLG LQKG-----
HM1-Enano	-----
HM1-Pira	-----
HM1-Sorghum	DATKLSMQKG-----
HM1-Rice	DAAKLRPQE G-----
HM1-Barley	DASKLIVQQG-----
DFR-Maize	DDVGGETSCRFPPPLL NQI