Table S2. Similarity between intact copies of the HELPO1 and HELPO2 families in PC15 genome.

helpo1.3_scaffold_08_RC	100.00	96.58	96.27	96.01	96.63	96.81	96.67	47.45	47.56	47.56	47.56	47.56	40.72	40.65	37.84	37.85
helpo1.3_scaffold_07_2	96.58	100.00	98.63	98.50	99.06	99.02	98.89	46.94	47.05	47.05	47.05	47.05	41.21	41.15	39.33	38.42
helpo1.3_scaffold_02_RC	96.27	98.63	100.00	98.19	98.59	98.83	98.70	47.11	47.22	47.22	47.22	47.22	40.55	40.49	39.24	38.34
helpo1.3_scaffold_01_RC	96.01	98.50	98.19	100.00	98.46	98.58	98.45	47.01	47.11	47.11	47.11	47.11	40.94	40.88	39.15	38.59
helpo1.3_scaffold_08_42_RC	96.63	99.06	98.59	98.46	100.00	98.93	98.86	47.23	47.33	47.33	47.33	47.33	40.74	40.67	39.00	38.51
helpo1.3_scaffold_07_1	96.81	99.02	98.83	98.58	98.93	100.00	99.87	47.23	47.33	47.33	47.33	47.33	40.87	40.81	39.32	38.51
helpo1.3_scaffold_09_RC	96.67	98.89	98.70	98.45	98.86	99.87	100.00	47.23	47.33	47.33	47.33	47.33	40.87	40.81	39.24	38.51
HELPO2_scaffold_01	47.45	46.94	47.11	47.01	47.23	47.23	47.23	100.00	100.00	100.00	100.00	100.00	50.20	50.25	50.18	49.01
HELPO2_scaffold_06_RC	47.56	47.05	47.22	47.11	47.33	47.33	47.33	100.00	100.00	100.00	100.00	100.00	50.19	50.25	50.19	49.00
HELPO2_scaffold_07_RC	47.56	47.05	47.22	47.11	47.33	47.33	47.33	100.00	100.00	100.00	100.00	100.00	50.19	50.25	50.19	49.00
HELPO2_scaffold_08	47.56	47.05	47.22	47.11	47.33	47.33	47.33	100.00	100.00	100.00	100.00	100.00	50.19	50.25	50.19	49.00
HELP02_scaffold_08_2_RC	47.56	47.05	47.22	47.11	47.33	47.33	47.33	100.00	100.00	100.00	100.00	100.00	50.19	50.25	50.19	49.00
HELP01.1_scaffold_07_1	40.72	41.21	40.55	40.94	40.74	40.87	40.87	50.20	50.19	50.19	50.19	50.19	100.00	99.74	56.93	56.69
HELP01.1_scaffold_07_2	40.65	41.15	40.49	40.88	40.67	40.81	40.81	50.25	50.25	50.25	50.25	50.25	99.74	100.00	56.91	56.71
HELP01.1_scaffold_05_RC	37.84	39.33	39.24	39.15	39.00	39.32	39.24	50.18	50.19	50.19	50.19	50.19	56.93	56.91	100.00	60.05
HELPO1.2 scaffold 11	37.85	38.42	38.34	38.59	38.51	38.51	38.51	49.01	49.00	49.00	49.00	49.00	56.69	56.71	60.05	100.00