

Table S5. The Ka/Ks value of segmental duplication gene pairs in cabbage

Gene pairs	Ka	Ks	Ka/Ks
<i>BoMAPK2/BoMAPK1</i>	0.065345207	0.857986718	0.076161094
<i>BoMAPK6.2/BoMAPK6.1</i>	0.025550373	0.512965226	0.049809172
<i>BoMAPK8.2/BoMAPK15</i>	0.111928007	0.808907473	0.138369357
<i>BoMAPK8.2/BoMAPK8.1</i>	0.03885235	0.356228388	0.109065845
<i>BoMAPK16.2/BoMAPK16.1</i>	0.042623594	0.337474565	0.126301649
<i>BoMAPK17.2/BoMAPK17.1</i>	0.017540714	0.288372671	0.060826548
<i>BoMAPK20.2/BoMAPK20.1</i>	0.031080301	0.449822521	0.069094586
<i>BoMAPK18.2/BoMAPK19.1</i>	0.114659333	0.703146879	0.163065978
<i>BoMAPK18.2/BoMAPK18.1</i>	0.112259619	0.742909558	0.151108056
<i>BoMAPK18.1/BoMAPK19.1</i>	0.045305893	0.352802183	0.128417269
<i>BoMAPK19.2/BoMAPK18.2</i>	0.034848154	0.433648526	0.080360366
<i>BoMAPK19.2/BoMAPK19.1</i>	0.104906176	0.706684558	0.148448378
<i>BoMAPK19.2/BoMAPK18.1</i>	0.109052833	0.654146559	0.166710093