

**Table S2. Results of site models for detection of positively selected sites in plant *GGP* genes.**

Model	Np <sup>a</sup>	lnL <sup>b</sup>	Parameters	Models compared	d.f. <sup>c</sup>	-2ΔlnL <sup>d</sup>	p-value	Positively selected sites
<b>M0</b>	297	-49065.194	ω=0.09302					none
<b>M3</b>	301	-47596.378	p0=0.38716, ω0=0.01357 P1=0.40745, ω1=0.08947 P2=0.20538, ω2=0.27976	M3 vs. M0	4	2937.632***	0.000	none
<b>M1a</b>	298	-48824.857	p0=0.88231, ω0=0.08702 p1=0.11769, ω1=1.0000					Not allowed
<b>M2a</b>	300	-48824.857	p0=0.88232, ω0=0.08702; p1=0.11768 ω1=1.000; p2=0.0, ω2=7.98423	M2a vs. M1a	2	0	1.000	none
<b>M7</b>	298	-47538.644	p=0.66833, q=5.72263					Not allowed
<b>M8</b>	300	-47538.647	p0=0.99999, p=0.66832 q=5.72257 p1=0.00001, ω=2.73695	M8 vs. M7	2	0.006	0.997	none

<sup>a</sup> Np: number of estimated parameters;

<sup>b</sup> lnL: log likelihood score;

<sup>c</sup> df : degrees of freedom;

<sup>d</sup> -2ΔlnL: twice the log-likelihood difference of the model compared.

\*\*\*Significant at  $p < 0.0001$ .

**Table S3. Results of branch-site test by treating each main lineages in the phylogeny as the foreground branch**

Foreground branch	Model	lnL <sup>a</sup>	Parameter estimates	-2ΔlnL <sup>b</sup>	d.f. <sup>c</sup>	p-value	Positively Selected sites
Angiosperms	Model A	-48816.091	p0 = 0.865, p1 = 0.077 p2a = 0.054, p2b = 0.005 background: ω0 = 0.086, ω1 = 1.000 ω2a = 0.086, ω2b = 1.000 foreground: ω0 = 0.086, ω1 = 1.000, ω2a = 1.000, ω2b = 1.000	0	1	1	
	Null model	-48816.091	p0 = 0.865, p1 = 0.077, p2a = 0.054, p2b = 0.005 background: ω0 = 0.086, ω1 = 1.000, ω2a = 0.086, ω2b = 1.000 foreground: ω0 = 0.086, ω1 = 1.000, ω2a = 1.000, ω2b = 1.000				
Angiosperms 1	Model A	-48787.112	p0 = 0.876, p1 = 0.083, p2a = 0.038, p2b = 0.004 background: ω0 = 0.087, ω1 = 1.000, ω2a = 0.087, ω2b = 1.000 foreground: ω0 = 0.087, ω1 = 1.000, ω2a = 1.000, ω2b = 1.000	0	1	1	16L (1.000); 254Q (1.000)
	Null model	-48787.112	p0 = 0.876, p1 = 0.083, p2a = 0.038, p2b = 0.004 background: ω0 = 0.087, ω1 = 1.000, ω2a = 0.087, ω2b = 1.000 foreground: ω0 = 0.087, ω1 = 1.000,				

<b>Angiosperms 2</b>	Model A	-48764.624	$\omega_{2a} = 1.000, \omega_{2b} = 1.000$ $p_0 = 0.839, p_1 = 0.091, p_{2a} = 0.063,$ $p_{2b} = 0.007$ background: $\omega_0 = 0.086, \omega_1 = 1.000,$ $\omega_{2a} = 0.086, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.086, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$	0	1	1	17G (0.997); 28S (0.941);35E (0.955); 45A (0.999);78Q (0.942); 82P (0.999); 107A (1.000)
	Null model	-48764.624	$p_0 = 0.834, p_1 = 0.091, p_{2a} = 0.063,$ $p_{2b} = 0.007$ background: $\omega_0 = 0.086, \omega_1 = 1.000,$ $\omega_{2a} = 0.086, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.086, \omega_1 = 1.000,$ $2a = 1.000, \omega_{2b} = 1.000$				
<b>Eudicots 1</b>	Model A	-48805.662	$p_0 = 0.857, p_1 = 0.105, p_{2a} = 0.034,$ $p_{2b} = 0.004$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$	0	1	1	254Q (1.000)
	Null model	-48805.662	$p_0 = 0.857, p_1 = 0.105, p_{2a} = 0.034,$ $p_{2b} = 0.004$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<b>Eudicots 2</b>	Model A	-48783.609	$p_0 = 0.828, p_1 = 0.100, p_{2a} = 0.064,$	0	1	1	19S (0.996); 45A (1.000);

---

			p2b = 0.008 background: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.086$ , $\omega_{2b} = 1.000$ foreground: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				82P (0.996);107A (0.991); 125Q (0.922);137E (1.000); 145M (1.000);155L (0.938)
	Null model	-48783.609	p0 = 0.828, p1 = 0.100, p2a = 0.064, p2b = 0.008 background: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.086$ , $\omega_{2b} = 1.000$ foreground: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<b>Monocots 1</b>	Model A	-48739.115	p0 = 0.715, p1 = 0.106, p2a = 0.156, p2b = 0.023 background: $\omega_0 = 0.083$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.083$ , $\omega_{2b} = 1.000$ foreground: $\omega_0 = 0.083$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$	0	1	1	1T (1.000); 2V (0.963); 3V (0.997); 5I (1.000); 6N (0.964); 7Q (0.998); 8- (0.999); 9- (0.929);15- (1.000); 18A (0.998); 25D (0.997); 27L(0.988); 28S (0.999); 44T (0.947); 59A (0.996); 60Q (0.992); 75R (0.924); 94D (0.97); 110V(0.999);179P(1.000);194S (1.000); 214N (1.000); 248A(0.998);265V (0.987); 287Y (1.000);288A (0.996); 298E (0.948);299D (0.993)
	Null model	-48739.115	p0 = 0.715, p1 = 0.106, p2a = 0.156, p2b = 0.023 background: $\omega_0 = 0.083$ , $\omega_1 = 1.000$ ,				

---

---

<b>Monocots 2</b>	Model A	-48813.117	$\omega_{2a} = 0.083, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.083, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$ $p_0 = 0.840, p_1 = 0.110, p_{2a} = 0.045,$ $p_{2b} = 0.006$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$	0	1	1	12- (0.999); 28S (1.000); 78Q (0.964)
	Null model	-48813.117	$p_0 = 0.840, p_1 = 0.110, p_{2a} = 0.045,$ $p_{2b} = 0.006$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<b>Gymnosperms</b>	Model A	-48826.351	$p_0 = 0.829, p_1 = 0.128, p_{2a} = 0.037,$ $p_{2b} = 0.006$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$	0	1	1	2V (0.991); 152Y (0.995)
	Null model	-48826.351	$p_0 = 0.829, p_1 = 0.128, p_{2a} = 0.037,$ $p_{2b} = 0.006$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$				

---

---

<b>Chlorophytes</b>	Model A	-48674.379	$\omega_{2a} = 1.000, \omega_{2b} = 1.000$ $p_0 = 0.587, p_1 = 0.088, p_{2a} = 0.283,$ $p_{2b} = 0.042$ background: $\omega_0 = 0.083, \omega_1 = 1.000,$ $\omega_{2a} = 0.083, \omega_{2b} = 1.000$ foreground: $\omega_0 = 0.083, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$	0	1	1	5I (0.998); 6N (0.998); 11- (0.999); 12- (0.999); 18A (0.998); 19S (1.000); 22L (1.000); 47E (1.000); 75R (1.000); 80L (1.000); 96Q (0.991); 101A (1.000); 135C(1.000); 148V(1.000); 190G(0.998); 214N (1.000); 215A (0.999); 216C (0.903); 217I (1.000); 242C (1.000); 245E (0.988); 249R (0.994); 251E (1.000); 277R (1.000); 284T (1.000); 287Y (1.000); 289K (1.000); 294E (1.000); 297S (1.000); 300R (0.938); 304V (0.953); 305K (0.994)
	Null model	-48674.379	$p_0 = 0.587, p_1 = 0.088, p_{2a} = 0.283,$ $p_{2b} = 0.042$ background: $\omega_0 = 0.083, \omega_1 = 1.000,$ $\omega_{2a} = 0.083, \omega_{2b} = 1.000$ foreground; $\omega_0 = 0.083, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<b>Lycophytes</b>	Model A	-48822.348	$p_0 = 0.814, p_1 = 0.128, p_{2a} = 0.050,$ $p_{2b} = 0.008$ background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$	0.592	1	0.442	27L (0.989); 31E (0.958); 112A (0.971); 179P (0.999); 222D (0.956); 300R (0.979)

---

---

			foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.626$ , $\omega_{2b} = 1.626$				
	Null model	-48822.644	$p_0 = 0.801$ , $p_1 = 0.127$ , $p_{2a} = 0.063$ , $p_{2b} = 0.010$				
			background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$				
			foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<b>Bryophytes</b>	Model A	-48829.428	$p_0 = 0.850$ , $p_1 = 0.123$ , $p_{2a} =$ $0.0235$ , $p_{2b} = 0.003$	0	1	1	222D (0.991)
			background: $\omega_0 = 0.088$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.088$ , $\omega_{2b} = 1.000$				
			foreground: $\omega_0 = 0.088$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
	Null model	-48829.428	$p_0 = 0.851$ , $p_1 = 0.123$ , $p_{2a} = 0.023$ , $p_{2b} = 0.003$				
			background: $\omega_0 = 0.088$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.088$ , $\omega_{2b} = 1.000$				
			foreground: $\omega_0 = 0.088$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				

---

<sup>a</sup> lnL: log-likelihood scores.

<sup>b</sup>  $-2\Delta\ln L$ : twice the log-likelihood difference of the models being compared.

<sup>c</sup> d.f.: degree of freedom.

**Table S4. Results of branch-site tests by treating each branch in the phylogeny as the foreground branch.**

Foreground branches	Model	lnL <sup>a</sup>	Parameter estimates	-2ΔlnL <sup>b</sup>	d.f. <sup>c</sup>	p-value	Positively Selected sites
<i>Ananas comosus</i> ( <i>AncGGP-3</i> )	Model A	-48817.404	p0 = 0.784, p1 = 0.125, p2a = 0.079, p2b = 0.013 Background: ω0 = 0.087, ω1 = 1.000, ω2a = 0.087, ω2b = 1.000 Foreground: ω0 = 0.087, ω1 = 1.0000 ω2a = 5.443, ω2b = 5.443	8.041	1	0.0046*	1T (0.987); 7Q (0.987); 59A (0.981); 60Q (0.965); 214N (0.992); 245E (0.984); 265V (0.993); 287Y (0.991); 303E (0.992)
	Null model	-48821.424	p0 = 0.717, p1 = 0.115, p2a = 0.145, p2b = 0.023 Background: ω0 = 0.086, ω1 = 1.000, ω2a = 0.086, ω2b = 1.000 Foreground: ω0 = 0.086, ω1 = 1.000, ω2a = 1.000, ω2b = 1.000				
<i>Arabidopsis thaliana</i> ( <i>AtGGP-1</i> )	Model A	-48830.433	p0 = 0.860, p1 = 0.137, p2a = 0.003, p2b = 0.00046 Background: ω0 = 0.087 ω1 = 1.000, ω2a = 0.087, ω2b = 1.000 Foreground: ω0 = 0.087, ω1 = 1.000, ω2a = 43.808, ω2b = 43.808	5.493	1	0.019*	215A (0.995)
	Null model	-48833.180	p0 = 0.808, p1 = 0.129, p2a = 0.055, p2b = 0.009 Background: ω0 = 0.087 ω1 = 1.000, ω2a = 0.087, ω2b = 1.000 Foreground: ω0 = 0.087, ω1 = 1.000,				



			$\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<i>Brachypodium_stacei</i> (BsGGP-1)	Model A	-48830.372	p0 = 0.861, p1 = 0.137, p2a = 0.002, p2b = 0.00033 Background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 76.292, \omega_{2b} = 76.292$	5.983	1	0.014*	187T (0.995)
	Null model	-48833.364	p0 = 0.82, p1 = 0.13, p2a = 0.043, p2b = 0.007 Background: $\omega_0 = 0.088, \omega_1 = 1.000,$ $\omega_{2a} = 0.088, \omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.088, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<i>Brassica_oleracea</i> (BoGGP-2)	Model A	-48810.375	p0 = 0.838, p1 = 0.134, p2a = 0.025, p2b = 0.004 Background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 23.625, \omega_{2b} = 23.625$	26.261	1	0.0000**	1T (0.975); 5I (1.000); 6N (0.999); 7Q (0.999); 15- (1.000)
	Null model	-48823.506	p0 = 0.715, p1 = 0.114, p2a = 0.148, p2b = 0.024 Background: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 0.087, \omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087, \omega_1 = 1.000,$ $\omega_{2a} = 1.000, \omega_{2b} = 1.000$				
<i>Brassica_rapa</i>	Model A	-48805.884	p0 = 0.816, p1 = 0.13, p2a = 0.047, p2b	20.204	1	0.0000**	1T (0.969); 17G (0.960); 51L (0.969);

<i>(BrGGP-4)</i>			= 0.007 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 45.8$ , $\omega_{2b} = 45.8$				75R (0.969); 107A (0.965); 207E (0.961); 225P (0.966); 251E (0.967); 281D (0.967); 283A (0.998); 284T (1.000); 285E (0.965); 287Y (1.000); 288A (0.965); 289K (0.995)
	Null model	-48815.986	$p_0 = 0.334$ , $p_1 = 0.053$ , $p_{2a} = 0.528$ , $p_{2b} = 0.084$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Chlamydomonas reinhardtii</i> <i>(ChrGGP)</i>	Model A	-48800.092	$p_0 = 0.728$ , $p_1 = 0.112$ , $p_{2a} = 0.139$ , $p_{2b} = 0.021$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 27.048$ , $\omega_{2b} = 27.048$	36.411	1	0.0000*	11- (0.999); 12- (0.999); 22L (0.960); 47E (0.956); 87K (1.000); 148V (0.998); 151P (0.999); 152Y (0.999); 159S (1.000); 203T (0.951); 230I (1.000); 265V (0.953); 284T (0.996); 289K (0.994); 294E (1.000); 297S (0.998)
	Null model	-48818.298	$p_0 = 0.723$ , $p_1 = 0.112$ , $p_{2a} = 0.144$ , $p_{2b} = 0.022$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Coccomyxa subellipsoidea</i> <i>(CosGGP)</i>	Model A	-48808.925	$p_0 = 0.7$ , $p_1 = 0.114$ , $p_{2a} = 0.161$ , $p_{2b} =$ 0.026 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ ,	17.84	1	0.0000**	2V (0.997); 5I (0.992); 6N (0.978); 7Q (1.000); 47E (0.958); 94D (0.999); 131P (0.997); 135C (0.999); 140L (0.995); 190G

			$\omega_{2a} = 0.087$ $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 7.241$ , $\omega_{2b} = 7.241$				(0.975); 220Q (1.000); 241Q (0.990); 245E (0.985); 246R (0.997); 251E (0.984); 260Q (0.985); 287Y (0.993); 296L (0.996); 300R (0.983)
	Null model	-48817.845	$p_0 = 0.713$ , $p_1 = 0.115$ $p_{2a} = 0.148$ $p_{2b} = 0.024$ Background: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.086$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Dunaliella salina</i> (DsGGP)	Model A	-48808.785	$p_0 = 0.724$ , $p_1 = 0.114$ , $p_{2a} = 0.14$ , $p_{2b} = 0.022$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 21.654$ , $\omega_{2b} = 21.654$	29.163	1	0.0000**	4S (0.999); 6N (0.990); 75R (0.996); 80L (0.990); 84D (0.998); 94D (0.953); 134I(0.989); 215A(0.999); 217I(1.000); 226F(0.953); 240I(0.999); 242C(0.968); 246R(0.999); 283A(0.995); 284T(0.992); 297S(0.996)
	Null model	-48823.367	$p_0 = 0.747$ , $p_1 = 0.118$ , $p_{2a} = 0.116$ , $p_{2b} = 0.018$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Eucalyptus grandis</i> (EgGGP-5)	Model A	-48818.108	$p_0 = 0.832$ , $p_1 = 0.133$ , $p_{2a} = 0.031$ , $p_{2b} = 0.005$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$	12.113	1	0.0005*	37G (0.968); 62N (1.000); 63E (1.000); 125Q (0.966); 135C (0.965); 216C (0.968); 264A (0.967); 284T (0.969)

			Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 59.268$ , $\omega_{2b} = 59.268$				
	Null model	-48824.164	$p_0 = 0.379$ , $p_1 = 0.06$ , $p_{2a} = 0.484$ , $p_{2b} = 0.077$  Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$  Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Glycine max</i> ( <i>GmGGP-4</i> )	Model A	-48824.508	$p_0 = 0.858$ , $p_1 = 0.137$ , $p_{2a} = 0.005$ , $p_{2b} = 0.00079$  Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$  Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 73.352$ , $\omega_{2b} = 73.352$	8.194	1	0.004*	304V (0.998); 305K (0.997)
	Null model	-48828.605	$p_0 = 0.71$ , $p_1 = 0.113$ , $p_{2a} = 0.152$ , $p_{2b} = 0.024$  Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$  Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Marchantia polymorpha</i> ( <i>MpGGP</i> )	Model A	-48827.668	$p_0 = 0.838$ , $p_1 = 0.133$ , $p_{2a} = 0.025$ , $p_{2b} = 0.004$  Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$  Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 7.806$ , $\omega_{2b} = 7.806$	8.527	1	0.003*	127K (0.982); 128S (0.955); 203T (0.970)

	Null model	-48831.931	p0 = 0.834, p1 = 0.133, p2a = 0.029, p2b = 0.005 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Micromonas pusilla</i> ( <i>MpGGP</i> )	Model A	-48803.535	p0 = 0.728, p1 = 0.112, p2a = 0.138, p2b = 0.021 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 13.253$ , $\omega_{2b} = 13.253$	26.221	1	0.0000**	15- (0.999); 19S (0.993); 75R (0.998); 123V (1.000); 214N (0.977); 215A (0.977); 241Q (0.999); 242C (1.000); 284T (0.999); 289K (0.989)
	Null model	-48816.645	p0 = 0.675, p1 = 0.105, p2a = 0.19, p2b = 0.03 Background: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.086$ $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Micromonas sp RCC299</i> ( <i>MsGGP</i> )	Model A	-48804.458	p0 = 0.793, p1 = 0.126, p2a = 0.07, p2b = 0.011 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 16.994$ , $\omega_{2b} = 16.994$	34.717	1	0.0000**	4S (0.999); 12- (0.999); 22L (0.986); 47E (0.967); 80L (0.999); 101A (0.965); 107A (0.998); 138T (0.998); 215A (0.998); 287Y (0.998); 289K (0.964); 297S (0.999)
	Null model	-48821.817	p0 = 0.755, p1 = 0.12, p2a = 0.108, p2b = 0.017				

			Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Mimulus guttatus</i> (MgGGP-2)	Mode A	-48831.088	$p_0 = 0.859$ , $p_1 = 0.137$ , $p_{2a} = 0.004$ , $p_{2b} = 0.001$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 17.229$ , $\omega_{2b} = 17.229$	4.794	1	0.029*	129Q (0.953)
	Null model	-48833.485	$p_0 = 0.851$ , $p_1 = 0.136$ , $p_{2a} = 0.012$ , $p_{2b} = 0.002$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Musa acuminata</i> (MaGGP-1)	ModelA	-48831.56	$p_0 = 0.856$ , $p_1 = 0.136$ , $p_{2a} = 0.007$ , $p_{2b} = 0.001$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 9.465$ , $\omega_{2b} = 9.465$	4.01	1	0.045*	29Q (0.983); 190G (0.984)
	Null model	-48833.565	$p_0 = 0.842$ , $p_1 = 0.134$ , $p_{2a} = 0.021$ , $p_{2b} = 0.003$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ $\omega_{2b} = 1.000$				

			Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Ostreococcus lucimarinus</i> ( <i>OIGGP</i> )	Model A	-48801.958	$p_0 = 0.763$ , $p_1 = 0.115$ , $p_{2a} = 0.107$ , $p_{2b} = 0.016$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 9.474$ , $\omega_{2b} = 9.474$	30.123	1	0.0000**	46C (0.997); 47E (0.953); 52P (0.996); 56G (0.992); 63E (0.999); 95Q (0.982); 129Q (0.998); 191C (0.997); 215A (0.955); 216C (0.998); 245E (0.999); 250G (0.990); 277R (0.972); 289K (0.997); 290K (0.980)
	Null model	-48817.019	$p_0 = 0.744$ , $p_1 = 0.117$ , $p_{2a} = 0.12$ , $p_{2b} = 0.019$ Background: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.086$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.086$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Panicum virgatum</i> ( <i>PvGGP-2</i> )	Model A	-48791.388	$p_0 = 0.849$ , $p_1 = 0.136$ , $p_{2a} = 0.013$ , $p_{2b} = 0.002$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 87.418$ , $\omega_{2b} = 87.418$	55.025	1	0.0000**	51 (0.999); 6N (0.999); 7Q (1.000); 8- (1.000); 9- (1.000); 12- (1.000)
	Null model	-48818.900	$p_0 = 0.709$ , $p_1 = 0.113$ , $p_{2a} = 0.154$ , $p_{2b} = 0.025$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				

<i>Picea abies</i> ( <i>PaGGP-3</i> )	Model A	-48799.663	p0 = 0.855, p1 = 0.136, p2a = 0.008, p2b = 0.001 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 969.357$ , $\omega_{2b} = 969.357$	46.735	1	0.0000**	1T (1.000); 2V (0.993); 4S (1.000); 5I (0.999); 6N (1.000); 7Q (0.993)
	Null model	-48823.030	p0 = 0.476, p1 = 0.076, p2a = 0.387, p2b = 0.062 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Selaginella moellendorffii</i> ( <i>SmGGP-2</i> )	Model A	-48824.521	p0 = 0.811, p1 = 0.130, p2a = 0.051, p2b = 0.008 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 4.037$ , $\omega_{2b} = 4.037$	5.590	1	0.018*	109N (0.964); 112A (0.969); 113I (0.984); 151P (0.978); 179P (0.966); 297S (0.992)
	Null model	-48827.316	p0 = 0.782, p1 = 0.125, p2a = 0.08, p2b = 0.013 Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Selaginella moellendorffii</i> ( <i>SmGGP-3</i> )	Model A	-48814.674	p0 = 0.758, p1 = 0.116, p2a = 0.109, p2b = 0.017	10.297	1	0.001*	8- (0.984); 9- (0.971); 34M (0.956); 82P (0.991); 112A (0.955); 138T (0.965);



			Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 3.798$ , $\omega_{2b} = 3.798$				150N (0.953); 190G (0.988); 217I (0.995); 222D (0.974); 231A (0.980); 245E (0.970); 250G (0.983); 253D (0.975); 290K (0.990); 299D (0.989); 300R (0.985)
	Null model	-48819.823	$p_0 = 0.741$ , $p_1 = 0.116$ , $p_{2a} = 0.123$ , $p_{2b} = 0.019$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Volvox carteri</i> (VcGGP)	Model A	-48804.773	$p_0 = 0.755$ , $p_1 = 0.118$ , $p_{2a} = 0.109$ , $p_{2b} = 0.017$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 16.335$ , $\omega_{2b} = 16.335$	32.725	1	0.0000**	8- (0.999); 9- (0.961); 18A (0.965); 22L (0.992); 23Y (0.999); 47E (0.966); 148V (0.990); 184R (0.999); 190G (0.972); 202R (0.998); 251E (0.998); 303E (1.000)
	Null model	-48821.135	$p_0 = 0.745$ , $p_1 = 0.118$ , $p_{2a} = 0.119$ , $p_{2b} = 0.019$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				
<i>Zea mays</i> (ZmGGP-1)	Model A	-48818.760	$p_0 = 0.846$ , $p_1 = 0.135$ , $p_{2a} = 0.016$ , $p_{2b} = 0.003$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$	21.888	1	0.0000**	25D (0.996); 26S (0.996); 27L (0.996); 28S (1.000); 29Q (0.996)

			Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 999.000$ , $\omega_{2b} = 999.000$				
	Null model	-48829.704	$p_0 = 0.449$ , $p_1 = 0.072$ , $p_{2a} = 0.414$ , $p_{2b} = 0.066$ Background: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 0.087$ , $\omega_{2b} = 1.000$ Foreground: $\omega_0 = 0.087$ , $\omega_1 = 1.000$ , $\omega_{2a} = 1.000$ , $\omega_{2b} = 1.000$				

<sup>a</sup> lnL: log-likelihood scores.

<sup>b</sup>  $-2\Delta\ln L$ : twice the log-likelihood difference of the models being compared.

<sup>c</sup> d.f.: degree of freedom.

\*  $p < 0.05$ ; \*\*  $p < 0.0003$  (Bonferroni correction for a significance level of 0.05 over 149 independent tests).

Text S1. Alignment of plant GGP protein sequences and the histidine triad (HIT) motif is marked using box.

	10	20	30	40	50	60	70	80	90	100	110	120	
AcGGP								MLK					
AcGGP-R-1								MLK					
AcGGP-R-2	M						LNT	ATQ-MLT					
AdGGP								MLK					
AeGGP								MLK					
ArGGP								MLK					
AmhGGP-1								MLR					
AmhGGP-2							MGI	LEK-SLT					
AncGGP-3								MLSEA					
AncGGP-1								MLT					
AncGGP-2								MLT					
AqcGGP								MLR					
AhGGP-1								MLK					
AhGGP-2								M-LLK					
AlGGP-1								MLK					
AlGGP-2								M-LLK					
AtGGP-1								MLK					
AtGGP-2								M-LLK					
BosGGP-1								MLK					
BosGGP-2								M-LLK					
BdGGP-1								MEM-KLT					
BdGGP-2								MEV-KLT					
BdGGP-3													
BsGGP-1								MEM-KLT					
BsGGP-2								MEV-KLT					
BsGGP-3													
BoGGP-1								MLK					
BoGGP-2								MLK					
BoGGP-3								M-LLK					
BrGGP-1								MLK					
BrGGP-2								MLK					
BrGGP-3								MLK					
BrGGP-4								M-LLK					
CgGGP-1								MLK					
CgGGP-2								M-LLK					
CrGGP-1								MLK					
CrGGP-2								M-LLK					
ChrGGP								MDSLQ-A	L				
CcGGP-1								M-MLR					
CcGGP-2								MLT					
CsGGP-1								M-MLR					
CsGGP-2								MLT					
CosGGP								MYH					
CusGGP								MLR					
DcGGP-1								MEMLR					
DcGGP-2								MLR					
DsGGP	M			L	QQVTQQQQQL	QQQQQQQQQL	DGNVCNSS	MSR	SNSNHLT				
EgGGP-1								M-MLK					
EgGGP-2								M-MLK					
EgGGP-3								MLT					
EgGGP-4								MLT					
EgGGP-5								MLT					
EsGGP-1								MLS					
EsGGP-2								M-LLK					
FvGGP-1								M-MLK					
FvGGP-2								M-MLK					
FvGGP-3								M-MMK					
GmGGP-1								M-MLK					
GmGGP-2								M-MLR					
GmGGP-3								MLS					
GmGGP-4								MLS					
GnmGGP								MLT					
GrGGP-1								M-MLR					
GrGGP-2								M-MLR					
GrGGP-3								MM-QLS					
KfGGP-1								MGE	STM-MLK				
KfGGP-2								MGE	STM-MLK				
KlGGP-1								MGE	STM-MLK				
KlGGP-2								MGE	STM-MLK				
KlGGP-3								MGE	STM-MLK				
LuGGP	MLVVTLGKVG	CEGGRRVVDI	TAIHGVSRL	LHVRKKEVIV	IQSNPSAHGG	RGALPSEGGS	PSDLLFLAGG	DKM-MLR					
MdGGP								M-MLR					
MeGGP-1	M							S	DNF-MLR				
MeGGP-2	M							S	DNL-MLR				
MeGGP-3									M-LLT				
MpGGP									MLT				
MtGGP-1									M-MLK				
MtGGP-2									MLS				
MpGGP									MAT				
MsGGP									M-SLT				
MgGGP-1									M-MLK				
MgGGP-2									M-MLK				
MaGGP-1									MLT				
MaGGP-2									MLS				
MaGGP-3									MLT				
MaGGP-4													
OtGGP									MEM-KLT				
OsGGP-1									MEM-KLT				
OsGGP-2									MPISPL				
OlgGGP													
PhGGP-1									MEM-KLT				
PhGGP-2													
PvGGP-1									MEM-KLT				
PvGGP-2									MATCSLTRLT	VC			
PvGGP-3													
PhvGGP-1									M-MLK				
PhvGGP-2									MLS				
PpGGP-1									MLT				
PpGGP-2									MLT				
PpGGP-3									MLT				
PaGGP-1									MLT				
PaGGP-2									MVTQKD	HVDKTQTGSL	SLASPPGISA	FSAFPFRPH	DRRILRSRN
PaGGP-3									MPLGR				
PsGGP-1									MLT				
PsGGP-2									MLT				
PsGGP-3									MLT				
PtGGP-1									MLT				
PtGGP-2									MLT				
PtGGP-3									MLT				
PtGGP-4									MLT				
PtGGP-5									MLT				
PotGGP-1									M-MLR				
PotGGP-2									M-MLR				





CsGGP-2	-----TVV	SNYQD	-----	QE	ETA	-----	-----	SENS	-----	E	-----	AGCGNSC	---	---	LGNC	-----
CosGGP	-----SWL	SGSSD	-----	EE	GS	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----
CusGGP	-----TVV	SNYQE	-----	DE	TEN	-----	-----	VSR	-----	S	-----	TGCGRNC	---	---	LNKCC	-----
DcGGP-1	-----TIV	SNYQK	-----	EE	GEE	-----	-----	GACR	-----	-----	-----	VGCGRNC	---	---	LNTCC	-----
DcGGP-2	-----TVV	SNYQK	-----	ED	GEE	-----	-----	ASRQ	-----	T	-----	GGCGRNC	---	---	LNKCC	-----
DsGGP	-----TVV	SMQL	-----	AQ	DDS	-----	-----	-----	-----	-----	-----	GGSGGNAPPS	---	---	NPESISQC	-----
EgGGP-1	-----TVV	SNYQK	-----	EE	VEG	-----	-----	G--R	-----	V	-----	GGCGRNC	---	---	LNKCC	-----
EgGGP-2	-----TVV	SNYQK	-----	EE	AEE	-----	-----	GARR	-----	G	-----	GGCGRNC	---	---	LNKCC	-----
EgGGP-3	-----TVV	SNYQE	-----	EG	-----	-----	-----	GE	-----	A	-----	KGCGRNC	---	---	LGKCC	-----
EgGGP-4	-----TVV	SNYQE	-----	GG	EKD-N	-----	-----	AAVAVEGE	-----	T	-----	KGCGRNC	---	---	LGKCC	-----
EgGGP-5	-----TVV	SNYQE	-----	EG	GEKDN	-----	-----	AAVAVEGE	-----	A	-----	KGCGRNC	---	---	LGKCC	-----
EsGGP-1	-----TVV	SNYQK	-----	DE	AAD	-----	-----	E	-----	S	-----	VGCGRNC	---	---	LGACC	-----
EsGGP-2	-----TVV	SNYQK	-----	DE	TAE	-----	-----	E	-----	G	-----	GGCGRNC	---	---	LSKCC	-----
FvGGP-1	-----TVV	SNYQK	-----	DE	ADE	-----	-----	G-RR	-----	A	-----	GGCGRNC	---	---	LNKCC	-----
FvGGP-2	-----TVV	SNYQK	-----	DE	ADE	-----	-----	G-RR	-----	A	-----	GGCGRNC	---	---	LNKCC	-----
FvGGP-3	-----TVV	SNYQN	-----	DE	ADE	-----	-----	G-RR	-----	V	-----	GGCGRNC	---	---	LNKCC	-----
GmGGP-1	-----TVV	SNFQK	-----	DD	AAD	-----	-----	SPRP	-----	V	-----	GGCGRNC	---	---	LKACC	-----
GmGGP-2	-----TVV	SNYQK	-----	DD	AAD	-----	-----	SPRP	-----	V	-----	GGCGRNC	---	---	LKACC	-----
GmGGP-3	-----TVV	SNYQK	-----	EE	TGE	-----	-----	AA	-----	A	-----	GGCGRNC	---	---	LKSCC	-----
GmGGP-4	-----TVV	SNYQK	-----	EE	TGE	-----	-----	AA-A	-----	A	-----	GGCGRNC	---	---	LKSCC	-----
GnmGGP	-----TIV	SMYHD	-----	-----	-----	-----	-----	GSEA	-----	A	-----	AGCGRNC	---	---	LGQCC	-----
GrGGP-1	-----TVV	SNYQK	-----	DE	ADD	-----	-----	TARR	-----	T	-----	AGCGRNC	---	---	LKSCC	-----
GrGGP-2	-----TVV	SNYQK	-----	DE	TEE	-----	-----	TARR	-----	S	-----	SGCGKNC	---	---	LRSCC	-----
GrGGP-3	-----TVV	SNYQK	-----	DE	AEE	-----	-----	TARR	-----	S	-----	GGCGKNC	---	---	LRSCC	-----
KfGGP-1	-----TVV	SNYQK	-----	DE	ADE	-----	-----	P	-----	-----	-----	KGCGRNC	---	---	LKQCC	-----
KfGGP-2	-----TVL	SNYQK	-----	DE	ADE	-----	-----	P	-----	-----	-----	KGCGRQC	---	---	LKECC	-----
KlGGP-1	-----TVV	SNYQK	-----	DE	ADE	-----	-----	P	-----	-----	-----	KGCGRNC	---	---	LKQCC	-----
KlGGP-2	-----TVL	SNYQK	-----	DE	ADE	-----	-----	P	-----	-----	-----	KGCGRQC	---	---	LKECC	-----
KlGGP-3	-----TVL	SNYQK	-----	DE	ADE	-----	-----	P	-----	-----	-----	KGCGRQC	---	---	LKECC	-----
LuGGP	-----TVV	SNYQK	-----	EE	TED	-----	-----	A	-----	G	-----	GGCGRNC	---	---	LQNC	-----
MdGGP	-----TVV	SNYQK	-----	DE	AEE	-----	-----	GARR	-----	V	-----	EGCGRNC	---	---	LNQCC	-----
MeGGP-1	-----TLV	SNFQK	-----	E	AEE	-----	-----	GASR	-----	S	-----	GGCGRNC	---	---	LKQCC	-----
MeGGP-2	-----TVV	SNFQK	-----	EE	AEE	-----	-----	GARR	-----	S	-----	EGCGRNC	---	---	LRKCC	-----
MeGGP-3	-----TVV	SNYQE	-----	ES	SE	-----	-----	TS	-----	F	-----	EGCGRHC	---	---	LGKCC	-----
MpGGP	-----TVV	SINQV	-----	SG	AAD	-----	-----	SSS	-----	-----	-----	-----	-----	-----	-----	-----
MtGGP-1	-----TVV	SNYQK	-----	EE	VGE	-----	-----	AAPRT	-----	V	-----	GGCGRNC	---	---	LKSCC	-----
MtGGP-2	-----TVV	SNYQK	-----	DE	DAP	-----	-----	-----	-----	V	-----	NGCGRNC	---	---	LKSCC	-----
MpGGP	-----TMI	SMHQM	-----	DD	EDD	-----	-----	PK	-----	-----	-----	-----	---	---	VARVV	-----
MsGGP	-----TVL	SIHQM	-----	DE	EED	-----	-----	PKF	-----	-----	---	---	---	---	VRRC	-----
MgGGP-1	-----TVV	SNYQK	-----	DE	AEE	-----	-----	G	-----	G	-----	AGCGRNC	---	---	LRSCC	-----
MgGGP-2	-----TVV	SNYQK	-----	DE	AEE	-----	-----	GVRP	-----	V	-----	TGCGRNC	---	---	FRSCC	-----
MaGGP-1	-----TVL	SNYQE	-----	DS	-----	-----	-----	GE	-----	S	-----	RGCGRNC	---	---	LGKCC	-----
MaGGP-2	-----TVL	SNYQE	-----	EN	-----	-----	-----	AGE	-----	P	-----	LGCGRSC	---	---	LGKCC	-----
MaGGP-3	-----TVL	SNYQE	-----	DS	-----	-----	-----	GE	-----	P	-----	RGCDRNC	---	---	LSKCC	-----
MaGGP-4	-----M	SVYQP	-----	TQ	GQH	-----	-----	G	-----	E	-----	EKSGRSC	---	---	LLQVH	-----
OtGGP	-----TVV	SNYQE	-----	DS	AAA	-----	-----	A	-----	A	-----	GGCGRNC	---	---	LGDC	-----
OsGGP-1	-----TVV	SNYQE	-----	DA	AAT	-----	-----	AGERP	-----	R	-----	AGCGRDC	---	---	LGDC	-----
OsGGP-2	-----AAF	GA	-----	-----	-----	-----	-----	RP	-----	ARRR	-----	-----	-----	-----	-----	-----
OlGGP	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----
PhGGP-1	-----TVV	SNYQD	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGHC	-----
PhGGP-2	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----
PvGGP-1	-----TVV	SNYQE	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGHC	-----
PvGGP-2	LEPGSRETVD	SGVST	-----	---	---	---	---	---	---	---	---	---	---	---	---	---
PvGGP-3	-----	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---
PhvGGP-1	-----TVV	SNYQK	-----	DG	AAD	-----	-----	TSRP	-----	V	-----	GGCGRNC	---	---	LKACC	-----
PhvGGP-2	-----TVV	SNYQK	-----	EG	SDE	-----	-----	SP	-----	A	-----	EGCGRNC	---	---	LKSCC	-----
PpGGP-1	-----TVV	SMHQ	-----	LE	-----	-----	-----	Q	-----	V	-----	PGCGRDC	---	---	LGTC	-----
PpGGP-2	-----TLV	SINQQ	-----	LD	-----	-----	-----	Q	-----	A	-----	PGCGRDC	---	---	LNKCC	-----
PpGGP-3	-----TVV	SMQQ	-----	LD	-----	-----	-----	Q	-----	G	-----	SGCGRDC	---	---	LNKCC	-----
PaGGP-1	-----TIL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PaGGP-2	-----TVL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRSC	---	---	LGQCC	-----
PaGGP-3	-----FLL	STGEE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PsGGP-1	-----TIL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PsGGP-2	-----TIV	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PsGGP-3	-----TIL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PtGGP-1	-----TIL	SNYQE	-----	GSK	-----	-----	-----	GSK	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PtGGP-2	-----TIL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PtGGP-3	-----TIL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	FGCGRNC	---	---	LGQCC	-----
PtGGP-4	-----TIV	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	VGCGRNC	---	---	LGQCC	-----
PtGGP-5	-----TTL	SNYQE	-----	GSE	-----	-----	-----	GSE	-----	G	-----	LGCGRNC	---	---	LGQCC	-----
PotGGP-1	-----TVV	SNYQK	-----	ED	NE	-----	-----	GSRR	-----	G	-----	GGCGRNC	---	---	LQNC	-----
PotGGP-2	-----TVV	SNYQK	-----	EE	GEE	-----	-----	GSRR	-----	G	-----	GGCGRNC	---	---	LNKCC	-----
PotGGP-3	-----TVV	SNYQE	-----	EN	SE	-----	-----	KG	-----	F	-----	EGCGRNC	---	---	LGKCC	-----
PrpGGP	-----TVV	SNYQK	-----	DE	AEE	-----	-----	GARR	-----	V	-----	GGCGRNC	---	---	LNQCC	-----
RcGGP-1	-----TVV	SNFQK	-----	DE	AED	-----	-----	GGKK	-----	S	-----	GGCGRNC	---	---	LQKCC	-----
RcGGP-2	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----
RcGGP-3	-----TVV	SNYQE	-----	DD	SSD	-----	-----	TS	-----	F	-----	EGCGRDC	---	---	LGNC	-----
SpGGP-1	-----TVV	SNYQK	-----	EE	GEE	-----	-----	GSRR	-----	G	-----	GGCGRNC	---	---	LRNCC	-----
SpGGP-2	-----TVV	SNYQK	-----	ED	GEE	-----	-----	GSRR	-----	G	-----	GGCGRNC	---	---	LNKCC	-----
SpGGP-3	-----TVV	SNYQE	-----	NS	E	-----	-----	KG	-----	F	-----	EGCGRNC	---	---	LGKCC	-----
SmGGP-1	-----TIV	SNYQE	-----	-----	-----	-----	-----	SLES	-----	K	-----	ICCGKNC	---	---	LGSCC	-----
SmGGP-2	-----AIV	SNDRS	-----	-----	-----	-----	-----	ET	-----	N	-----	HGCGRNC	---	---	LGPC	-----
SmGGP-3	-----TVV	SVNQD	-----	-----	-----	-----	-----	-----	-----	C	-----	GSVSRDC	---	---	LQSCF	-----
SiGGP-1	-----TVV	SNYQE	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGDC	-----
SiGGP-2	-----APA	AS	-----	-----	-----	-----	-----	DKP	-----	-----	-----	-----	-----	-----	-----	-----
SvGGP-1	-----TVV	SNYQE	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGDC	-----
SvGGP-2	-----APA	AS	-----	-----	-----	-----	-----	DKP	-----	-----	-----	-----	-----	-----	-----	-----
SlGGP-1	-----TLV	SNFQK	-----	DE	ADE	-----	-----	IAAR	-----	G	-----	AGCGRNC	---	---	LRNCC	-----
SlGGP-2	-----TLV	SNYQE	-----	DV	LEG	-----	-----	N	-----	V	-----	MCGCRKC	---	---	LGKCC	-----
StGGP-1	-----TLV	SNFQK	-----	DE	ADE	-----	-----	IGAR	-----	G	-----	AGCGRNC	---	---	LRNCC	-----
StGGP-2	-----TLV	SNYQE	-----	DV	LEG	-----	-----	N	-----	V	-----	MCGCRKC	---	---	LGKCC	-----
SbGGP-1	-----TVV	SNYQD	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGDC	-----
SbGGP-2	-----APT	AS	-----	-----	-----	-----	-----	DKP	-----	-----	-----	-----	-----	-----	-----	-----
SfGGP-1	-----TLV	SINQE	-----	LE	-----	-----	-----	Q	-----	A	-----	AGCGKDC	---	---	LGSC	-----
SfGGP-2	-----TLV	SFNQE	-----	LE	-----	-----	-----	Q	-----	A	-----	TCCGRDC	---	---	LGSC	-----
SppGGP-1	-----TVV	SNFQE	-----	---	---	---	---	---	---	S	-----	LGCGRSC	---	---	LGKCC	-----
SppGGP-2	-----TVV	SNFQE	-----	---	---	---	---	---	---	A	-----	LGCGRNC	---	---	LGKCC	-----
TcGGP-1	-----TVL	SNYQK	-----	DE	AEE	-----	-----	TARR	-----	S	-----	GGCGKNC	---	---	LRSCC	-----
TcGGP-2	-----TVL	SNYQE	-----	DA	YEK	-----	-----	Q	-----	E	-----	VGCGRNC	---	---	LNKCC	-----
TpGGP	-----TVV	SNYQK	-----	EE	VGD	-----	-----	APRT	-----	V	-----	GGCGRNC	---	---	LKACC	-----
VvGGP-1	-----TVV	SNYQK	-----	ED	SDD	-----	-----	GARQ	-----	V	-----	GGCGRNC	---	---	LKQCC	-----
VvGGP-2	-----TVV	SNYQE	-----	EA	SE	-----	-----	-----	-----	S	-----	LGCGRNC	---	---	LGHC	-----
VcGGP	S-ASSDSTVA	SCLDAYLPIY	IF	-----	---	---	---	---	---	S---	---	---	---	---	---	---
ZmGGP-E-1	-----TVV	SNYQE	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGDC	-----
ZmGGP-E-2	-----AS	PPTAA	SRADE	---	---	---	---	---	---	T	-----	PILGRHCRFLF	---	---	LGKAL	-----
ZmGGP-1	-----TVV	SNYQE	-----	DA	-----	-----	-----	DKP	-----	R	-----	AGCGRNC	---	---	LGDC	-----
ZmGGP-2	-----AS	PPTAA														







MaGGP-4	---GCSL---	Q	---EK-IEA	QSTFLDRL---	---LKEWEDR	KVRGLFHHDV	NAYETKLLSG	DYGFILQLIE	GRDLKKRPT
OtGGP	---SS---	Q	---AD-AAQ	TKPLVNL---	---LTEWENR	LARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
OsGGP-1	---SCSLRKE---	E	---ED-ASN	DEFFVNL---	---LGLWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
OlGGP	---PS---	Q	---EN-GTN	LSPFVRKL---	---FKEWDDR	KARGLFHDDI	SSCETKVL	EHNFVATLIE	GRDQKKRPT
PhGGP-1	---SCSLGM---	E	---DD-SLN	LSRFLDKL---	---ISAWEDR	FAGGLFRYDV	TAVSTKVID	KKKYVAQFNI	GRATNKRPT
PvGGP-1	---PA---	Q	---ED-AAS	TKSLVNL---	---LTEWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
PvGGP-2	---PA---	Q	---ED-AAS	TKSLVNL---	---LAEWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
PvGGP-3	---SCSPGM---	E	---DD-SLN	SSRFLDKL---	---FREWEDR	KARGLFHDDI	SSCETK---	---	---
PhvGGP-1	---D-LALQ---	C	---CG-EPP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFVAQLNE	GRHLKKRPT
PhvGGP-2	---L-PLLG---	C	---KE-HYP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFVAQLNE	GRHLKKRPT
PpGGP-1	---R-AHTG---	S	---VS-PQE	NYFLDELL---	---LAQWEDR	MARGLFRYDV	TACETKML	DCGFIAQLNE	GRHLQKRPT
PpGGP-2	---G-VHTN---	NDDH-CIPGG	---VS-PQE	NYFLDELL---	---LAQWEDR	MARGLFRYDV	TACETKML	DCGFIAQLNE	GRHLQKRPT
PpGGP-3	---S-VHDI---	DESH-CVPPD	---NP-PQE	NYFLDE-L---	---LAQWEDR	MARGLFRYDV	TACETKML	YGFIAQLNE	GRHSQKRPT
PaGGP-1	---YADK---	D	---AE-VAN	IPFLDSVL---	---LGEWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PaGGP-2	---HAVK---	D	---AK-VED	ITLLDTVL---	---LQWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PaGGP-3	---HVDN---	N	---VK-LAD	IPFLDSVL---	---LGLWEER	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PsGGP-1	---YADK---	D	---AE-VAN	IPFLDSVL---	---LGEWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PsGGP-2	---HVDK---	D	---VE-MAD	TPFLDTLL---	---LQWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PsGGP-3	---HVDN---	D	---VK-LAD	IPFLDSVL---	---LGLWEER	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PtGGP-1	---DADK---	D	---AE-VAH	ISFLDSVL---	---LGEWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PtGGP-2	---HVDN---	D	---VK-LTD	VPFLDSL---	---LGLWEER	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PtGGP-3	---DAVK---	D	---SK-VED	ITLLDAVL---	---VQWEDR	MQRGLFRYDV	TTCETKVI	YGFIAQLNE	GRHLKKRPT
PtGGP-4	---HFDK---	D	---VE-MAV	TPFLDSL---	---LQWEDR	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PtGGP-5	---HVDN---	D	---VK-LTD	VPFLDSL---	---LGLWEER	MQRGLFRYDV	TTCETKVI	NYGFIAQLNE	GRHLKKRPT
PotGGP-1	---VG-VFEY---	D	---KG-EPP	VAFLLDSL---	---LGEWEDR	VQRGLFRYDV	TACETKVI	QHGFAQLNE	GRHLKKRPT
PotGGP-2	---D-VFEH---	D	---KS-EPP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TTCETKVI	YGFIAQLNE	GRHLKKRPT
PotGGP-3	---S-VEKS---	S	---EE-QPH	MCFLLHLL---	---LQWEDR	MSRGLFRYDV	TACDTKII	YGFIAQLNE	GRHLKKRPT
PrpGGP	---E-LPGS---	E	---KR-EPP	VDFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
RcGGP-1	---E-VIEH---	E	---NT-EPP	VAFLLDSL---	---LGEWEDR	VQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
RcGGP-2	---E-EIEH---	E	---NI-EPP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
RcGGP-3	---G-EMKF---	S	---KE-QPP	ICFLHLL---	---LQWEDR	MCRGLFRYDV	TACETRIIP	YGFIAQLNE	GRHLKKRPT
SpGGP-1	---VPEH---	G	---KS-EPP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TTCETKVI	SHGFAQLNE	GRHLKKRPT
SpGGP-2	---VG-VSED---	D	---KG-BTT	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
SpGGP-3	---S-VEKS---	S	---ED-QPQ	ICFLHLL---	---LQWEDR	MSRGLFRYDV	TACDTKII	YGFIAQLNE	GRHLKKRPT
SmGGP-1	---GEIQ---	S	---KE-LGQ	NSFLDSAI---	---LQWADK	QABGLFRYDV	TACATKVI	YGFIAQLNE	GRHLKKRPT
SmGGP-2	---	---	---TK-GET	IDFLHSFI---	---LAPWMEK	QKQGLFRYDV	TSCETKIL	DCGFIAQLNE	GRHSKKRPT
SmGGP-3	---EVPS---	S	---PP-TIE	RSFLGTL---	---LSKWEER	ASQGLFRYDV	TACESRVL	NYGFIAQLNE	GRHLKKRPT
SiGGP-1	---PS---	Q	---ED-AAS	TKSLVNL---	---LTEWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
SiGGP-2	---GCSLGM---	E	---ND-SLS	LSPFLNKL---	---FREWEDR	KARGLFHDDI	SSCETKVL	QHNFVATLIE	GRDQKKRPT
SvGGP-1	---PS---	Q	---ED-AAS	TKSLVNL---	---LTEWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
SvGGP-2	---GCSLGM---	E	---ND-SLS	LSPFLNKL---	---FREWEDR	KARGLFHDDI	SSCETKVL	QHNFVATLIE	GRDQKKRPT
SlGGP-1	---SVAD---	E	---TK-ESP	IDFLLESV---	---LGEWEDR	QKGLFRYDV	TACETKVI	YGFVAQLNE	GRHLKKRPT
SlGGP-2	---D-VQTL---	P	---EE-ECQ	MSFLMDLL---	---LGLWEER	MSQGLFRYDV	TTCETKVI	RCGFIAQLNE	GRHLKKRPT
StGGP-1	---S-VAAD---	E	---TK-ESP	IDFLLESV---	---LGEWEDR	QKGLFRYDV	TACETKVI	YGFVAQLNE	GRHLKKRPT
StGGP-2	---D-IQTL---	P	---EE-ECQ	MSFLMDLL---	---LGLWEER	MSQGLFRYDV	TTCETKVI	RCGFIAQLNE	GRHLKKRPT
SbGGP-1	---RS---	Q	---EG-AAS	TKLLVDIL---	---LSEWEDR	MARGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
SbGGP-2	---T---	E	---HE-GSN	LSPFLHRL---	---FNEWDDR	KTRGLFHDDI	SSCETKVL	EHNFVATLIE	GRDQKKRPT
SfGGP-1	---CS-TNGG---	GDGHDTLNNG	---DS-DGD	NGFLDSL---	---LAQWEDR	MDKGLFRYDV	TACETMVL	EHGFIAQLNE	GRHSKKRPT
SfGGP-2	---N-AYSN---	GDGH-KVANG	---NS-EAA	NGFLDSL---	---LAQWEDR	MARGLFRYDV	TACETMVL	EHGFIAQLNE	GRHSKKRPT
SppGGP-1	---KS-LPAA---	P	---DE-DSP	VDFLNLV---	---LQWEDR	MSRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
SppGGP-2	---KK-ALSI---	P	---DE-DPS	IDFLNTLV---	---LQWEDR	LSKGLFRYDV	TACESKVI	YGFIAQLNE	GRHLKKRPT
TcGGP-1	---G-VRGN---	E	---NK-EPP	VAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
TcGGP-2	---G-ELYP---	S	---EE-QPQ	PSFL-NLL---	---LEQWEDR	MRRGLFRYDV	TNCKTKII	YGFIAQLNE	GRHLKKRPT
TpGGP	---L-AIDE---	E	---KE-BLP	LAFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
VvGGP-1	---A-SSGD---	E	---NK-EPP	VDFLLDSL---	---LGEWEDR	MQRGLFRYDV	TACETKVI	YGFIAQLNE	GRHLKKRPT
VvGGP-2	---R-VEVS---	S	---EG-QPP	AAFLLNLL---	---LQWEDR	MSQGLFRYDV	TTCETRIIP	YGFIAQLNE	GRHLKKRPT
VcGGP	---SSYSAFS---	SSIA	---LE-VPS	RSLLSAL---	---MTLWEDR	ADRGLFRYDV	TLCPTRVL	SRGFIAQLNE	GRATNKRPT
ZmGGP-E-1	---GS---	Q	---ED-AAS	TKLLVDIL---	---LSEWEDR	MTRGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
ZmGGP-E-2	---T---	E	---HE-GSN	LSPFLKLL---	---FNEWDDR	KTRGLFHDDI	SSCETKVL	EHNFVATLIE	GRDQKKRPT
ZmGGP-1	---GS---	Q	---ED-AAS	TKLLVLSLV	---CFAPWQEDR	MTRGLFRYDV	TACETKVI	NLGFVAQLNE	GRHLKKRPT
ZmGGP-2	---T---	E	---HE-GSN	LSPFLKLL---	---FNEWDDR	KTRGLFHDDI	SSCETKVL	EHNFVATLIE	GRDQKKRPT
ZomGGP	---GSAV---	A	---GK-NSP	IDFLNTLV---	---LQWEDR	MAQGMFRYDV	TACETKII	NHGFIAQLNE	GRHLQKRPT

AcGGP	FRVDKVLQPF	DESKFNFTKV	QGEVLFQFE	--A--SDD	N--EV--	---QFFP	NAPVDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI
AcGGP-R-1	FRVDKVLQPF	DGSKFNFTKV	QGEVLFQFE	--A--SND	N--EV--	---QFFP	NAPVDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	SILECLPQRI
AcGGP-R-2	FRVDQVLQPF	DENKFNFTKV	QGEVLFQFE	--Q--SND	S--KS--	---HFPT	SSSVNL-D---	---SNSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AdGGP	FRVDKVLQPF	DESKFNFTKV	QGEVLFQFE	--A--SND	N--EV--	---QFFP	NAPVDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	SILECLPQRI
AeGGP	FRVDKVLQPF	DESKFNFTKV	QGEVLFQFE	--A--SVD	N--EV--	---QFFP	NAPVDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI
ArGGP	FRVDKVLQPF	DESKFNFTKV	QGEVLFQFE	--A--SDD	N--EV--	---QFFP	NAPVDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI
AmhGGP-1	FRVDKVLQPF	DGSKFNFTKV	QGEVLFQFE	--A--SED	G--EA--	---YFTP	GAPIDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RIFECPLQRI
AmhGGP-2	FRVDQVLQPF	DDNKFNFTKV	QGEVLFQFE	--P--SND	M--TC--	---YHP	SAPVSL-E---	---SESPN	I----VSP-	IEYGHVLLIP	RIFECPLQRI
AncGGP-3	FRIDEVLQPF	DGDKFNFTKI	QGEVLFQFE	--E--SSN	G--KA--	---GYQP	SAPIMS---	---SSSPN	IIAIN-VSP-	IAHGHVLLIP	RILDCLPQRI
AncGGP-1	FRVDKVLQPF	DPKFNFTKV	QGEVLFQFE	--A--SGS	H--KS--	---SFSE	RAPVDG---	---TKAPN	VLAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AncGGP-2	FRVDQVLQPF	DANKFNFTKV	QGEVLFQFE	--P--ATN	A--VVIN	---TPVDG	---	---TRSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AqGGP	FRVDKVLQPF	DENKFNFTKV	QGEVLFQFE	EKV--NDD	F--GA--	---QFVA	SAPIDT---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	KILEHFPQRI
AhGGP-1	FRVDKVLQSF	DGTFNFTKV	QGEVLFQFE	--A--GED	A--QV--	---QFFP	CMPLDP---	---ENSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AhGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--SNN	DDD--SKI--	---QFLP	SIPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AlGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GED	G--QV--	---QFFP	CMPLDP---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AlGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--SDN	DDD--SQI--	---QFLP	SIPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AtGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GED	A--QV--	---QFFP	CMPLDP---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
AtGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--STN	DDD--SEI--	---QFLA	SMPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BosGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GED	G--QV--	---QFFP	CMPLDP---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BosGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--SND	DDD--SRI--	---QFFP	STPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BdGGP-1	FRVDKVLQSF	DAKFNFTKV	QGEVLFQFE	--N--GGG	D--DS--	---YFLK	SAPTVA---	---DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDRLPQRI
BdGGP-2	FRVDKVLQSF	HSKFNFTKV	QGEVLFQFE	--N--GGG	D--SS--	---YFLA	NAPNTE-S---	---SHPPS	VVAIN-VSP-	IEYGHVLLIP	RVLDRLPQRI
BdGGP-3	FGMNQVLQPF	DSGKFNFTKV	RPEEVIPTFH	--E--TED	E--SD--	---RYFD	GAPPTV---	---LASPS	SILIN-VSP-	IGYCHVLLIP	RILECLPQRI
BsGGP-1	FRVDKVLQSF	DAKFNFTKV	QGEVLFQFE	--N--GGG	D--DS--	---YFLK	SAPTVA---	---DCAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDRLPQRI
BsGGP-2	FRVDKVLQSF	DSKFNFTKV	QGEVLFQFE	--N--GGG	D--RS--	---YFLA	NAPNTE-S---	---SHPPS	VVAIN-VSP-	IEYGHVLLIP	RVLDRLPQRI
BsGGP-3	FGMNQVLQPF	DSGKFNFTKV	RPEEVIPTFH	--E--TEN	E--SD--	---RYFD	GAPPTV---	---LGSPS	SILIN-VSP-	IGYCHVLLIP	RILECLPQRI
BoGGP-1	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--GED	S--EV--	---QFFP	CMPLDA---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BoGGP-2	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GHD	D--EA--	---RFFP	CMPLVA---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BoGGP-3	FRVDQVLQPF	DANKFNFTKV	QGEVLFQFE	--A--GED	C--EV--	---QFFP	CMPLDA---	---ENSPR	VVAIN-VSP-	IEYGHVLLIP	QVFDCLPQRI
BrGGP-1	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--GED	C--EV--	---QFFP	CMPLDS---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BrGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--GED	C--EV--	---QFFP	CMPLDA---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BrGGP-3	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--GED	E--EV--	---QFLP	CMPLLA---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
BrGGP-4	FPVDQVLQPF	DVNKFNFTKV	QGEVLFQFE	--A--GED	D--DLPLEA	---	---	---GNSPS	VVAIN-VSP-	IEYGHVLLIP	QVFDCLPQRI
CgGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GED	D--QV--	---QFFP	CMPLDP---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CgGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--RSD	DDD--SGI--	---QFLP	SIPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CrGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--GED	D--QV--	---QFFP	CMPLDP---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CrGGP-2	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--RSD	DDD--SGI--	---QFFP	SIPLDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
ChrGGP	FSADRVMPQF	DPARFHFNKA	AMGEVLFQFE	--A--DAT	ASA--TSAT	---ATAAPRLLF	SAPMAKSALL	---ASNPSVSGSPN	LVLIN-VSP-	IDHCHVLLIP	RVLDCLPQRI
CcGGP-1	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--SED	G--EV--	---QFFP	SAPIDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CcGGP-2	FRIDQVLQPF	DENKFNFTKI	QGEVLFQFE	--D--	---	---	---	---SKSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CcGGP-3	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--SED	G--EV--	---QFFP	SAPIDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CsGGP-1	FRIDQVLQSF	DENKFNFTKI	QGEVLFQFE	--D--	---	---	---	---SKSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
CosGGP	FCVDQVQRF	DNGKFNFTKA	LQEVLFQFE	--A--ADM	SSK--GS--	---AFLE	LAPV---	---GSGPN	LVLIN-VSP-	IEYGHVLLIP	RALDRNLQRI
CusGGP	FRVDKVLQSF	DGNKFNFTKV	QGEVLFQFE	--A--NEN	G--NT--	---QFIP	NDAIDL---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILDCLPQRI
DcGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--SEE	D--EI--	---HFFP	DASIDV---	---GNSPS	VVAIN-VSP-	IEYGHVLLIP	HVLECLPQRI
DcGGP-2	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--SED	G--EI--	---QFFP	NAPIDV---	---ANSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
DsGGP	VVVDKVCQPF	TEAKFNFKKA	YLKEVLFQFE	--P--RSD	SGA--SGAHINDPSA	---MANKDDEA	---	---SYEPLELL	SAAA	---	---
DgGGP-1	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--SVA	G--EA--	---QFFP	NAPIDV---	---ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI
DgGGP-2	FRVDKVLQSF	DGSKFNFTKV	QGEVLFQFE	--A--SED	G--EV--	---QFFA	KAPIDA---	---DNSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCMPQRI

EgGGP-3	FRVDQVLQDF	DEKKNFNTKV	GQEEVLFMFE	--Q----	SDD	C-----	-----	YS--	-----	HFFP	SAPAIF-K--	-----	PTSPN	VVAIN-VSPS	IQYGHLLLP	RVLDCLPQRI	
EgGGP-4	FRVDQVLQDF	DEKKNFNTKV	GQEEVLFMFE	--Q----	SDD	C-----	-----	YS--	-----	HFFP	SAPAIF-K--	-----	PTSPN	V-----	VSP-	IEYGHVLLIP	RVLDCLPQRI
EgGGP-5	FRVDQVLQDF	DEKKNFNTKV	GQEEVLFMFE	--Q----	SDD	C-----	-----	YS--	-----	HFFP	SAPASF-K-	-----	PTSPN	VVAIN-VSP-	-----	IEYGHVLLIP	HVLDCLPQRI
EsGGP-1	FRVDKVLQSF	DGNKFNFTKV	GQEEVLFQFE	--A----	GED	G-----	-----	EV--	-----	QFFF	CMPLDA--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
EsGGP-2	FRVDKVLQSF	DGNKFNFTKV	GQEEVLFQFE	--A----	DDD	DDD-----	-----	REI-	-----	QFFF	NMPLDA--	-----	DNSPS	VVAIN-VSP-	IEYGHVLLIP	CVLDCLPQRI	
FvGGP-1	FRVDKVLQPF	DGSKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFHF	NAPIDV--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILESCLPQRI	
FvGGP-2	FRVDKVLQPF	DGSKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFHF	NAPIDV--	-----	ENSPN	VVAIN-VSP-	IEYGHVLLIP	RILESCLPQRI	
FvGGP-3	FRVDKVLQPF	DGSKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFHF	NAPIDI--	-----	ATSPS	VVAIN-VSP-	IEYGHVLLIP	RILESCLPQRI	
GmGGP-1	FRVDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	QV--	-----	QFFF	NAPVDV--	-----	DNSPS	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
GmGGP-2	FRVDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	QV--	-----	QFFF	NAPVDV--	-----	DNSPS	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
GmGGP-3	FRVDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--A----	SND	G-----	-----	EV--	-----	QFFF	NAPIDV--	-----	ENSPS	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
GmGGP-4	FRVDKVLQPF	DENKFNFTKV	GQEEVLFQLE	--A----	SND	G-----	-----	EA--	-----	QFFF	NVPIDV--	-----	ENSPS	FVAIN-VSP-	IEYGHVLLIP	QIFECLPQRI	
GnmGGP	FRVDKVLQDF	DPKKNFNTKV	GQEEMLFRFE	--E----	SAA	G-----	-----	IT--	-----	EFLE	KGPV--	-----	LDSPN	VIVIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
GrGGP-1	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFYP	NAPIDV--	-----	ESSPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
GrGGP-2	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFYP	SAPIDV--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
GrGGP-3	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFFF	NAPIDV--	-----	ENHPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
KfGGP-1	FRVDKVLQPF	DEKKNFNTKI	GQEEVVFQFE	--A----	SEC	G-----	-----	EV--	-----	QFFF	EAPIEV--	-----	ENRPN	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
KfGGP-2	FRVDKVLQPF	DEKKNFNTKI	GQEEVVFQFE	--A----	SEC	G-----	-----	EV--	-----	QFFF	EAPIHV--	-----	ENHPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
KlGGP-1	FRVDKVLQPF	DEKKNFNTKI	GQEEVVFQFE	--A----	SEC	G-----	-----	EV--	-----	QFFF	EAPIEV--	-----	ENRPN	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
KlGGP-2	FRVDKVLQPF	DEKKNFNTKI	GQEEVVFQFE	--A----	SEC	G-----	-----	EV--	-----	QFFF	EAPIHV--	-----	ENHPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
KlGGP-3	FRVDKVLQPF	DEKKNFNTKI	GQEEVVFQFE	--A----	SEC	G-----	-----	EV--	-----	QFFF	EAPIHV--	-----	ENHPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
KuGGP	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFD	--A----	SEN	G-----	-----	QV--	-----	EFFP	DAPVDA--	-----	DKSPS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
MeGGP	FRVDKVLQPF	DSSKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	HFFP	SAPIDV--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
MeGGP-1	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--P----	SEN	G-----	-----	EV--	-----	QFFF	SASIDV--	-----	DNSTS	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
MeGGP-2	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--S----	SED	G-----	-----	DV--	-----	QFFF	SAPIDV--	-----	DNSTP	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
MeGGP-3	FRVDKVLQAF	DETKFNFTKI	GQEEVLFQFE	--P----	SND	L-----	-----	KS--	-----	EFVF	TAPPTA-D-	-----	STSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
MpGGP	FRVDQVLQPY	DAKKNFNTKV	DQEEILFQFD	--E----	SSD	H-----	-----	CS--	-----	EYSE	QAFV--	-----	GESPN	VVAIN-VSP-	IEYGHVLLIP	QVKSQIPQRI	
MtGGP-1	FRVDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFYP	NAPIDV--	-----	DNSTP	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
MtGGP-2	FRVDKVLQPF	DETKFNFTKV	GQEEVLFQFK	--S----	SSD	G-----	-----	ET--	-----	QFFF	DAPIDV--	-----	ENSPS	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
MpGGP	FRVDQVQCFE	DAGKFNFTKA	DKAEIFFRFA	--P----	GGV	GQT-----	-----	RS--	-----	EYVA	SAPIEYADAD	RVRGESDAPT	VVFN-VSP-	IEYGHVLLIP	RVTDCLPQRI		
MsGGP	FAVDEVVQAF	DGGKFNFTKA	DKAEIFLFAFE	--R----	GDR	AMK-----	-----	SS--	-----	AYNS	AKTI--	-----	ESSPN	VMLIN-VSP-	IEYGHVLLIP	RVTDCLPQRI	
MgGGP-1	FRVDKVLQPF	DETKFNFTKV	GQEEVLFQFE	--P----	SEN	N-----	-----	HV--	-----	QFFF	NAPIDL--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
MgGGP-2	FRVDKVLQPF	DETKFNFTKV	GQEEVLFQFE	--A----	SND	N-----	-----	EV--	-----	HYFF	NAPIDS--	-----	DNSTP	VVAIN-VSP-	IEYGHVLLIP	RILECLPQRI	
MaGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--A----	GED	D-----	-----	KA--	-----	RFLF	SAAVGE--	-----	TNTPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
MaGGP-2	FRVDKVLQAF	DPKKNFNTKV	GQEEVLFQFE	--A----	AGG	D-----	-----	EA--	-----	HLFE	SDGNEG--	-----	TKSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
MaGGP-3	FRVDKVLQPF	DRKKNFNTKV	GQEEVLFQFE	--P----	GES	G-----	-----	SG--	-----	RLLE	DDAVGD--	-----	ANAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
MaGGP-4	FRIDRVLQPF	DGKKNFNTKI	AQEEMLIRFE	--E----	SEN	D-----	-----	DR--	-----	GFFE	NTPAIDV--	-----	SESPN	VVAIN-VSP-	IEYGHVLLIP	RILDCLPQRI	
OtGGP	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--S----	STG	D-----	-----	DS--	-----	YFLE	SASIPA--	-----	NRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
OsgGGP-1	FRVDKVLQPF	DAKKNFNTKV	GQEEVLFQFE	--N----	GGG	D-----	-----	DS--	-----	PFVE	SSPISV-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
OsgGGP-2	FGMNQVLQPF	DSKFNFTKV	SPEEVLPTFK	--E----	SQN	D-----	-----	SV--	-----	KYFD	NVPHAV--	-----	AAASP	AILIN-VSP-	IEYGHVLLIP	RIQDCLPQRI	
OlGGP	FSVDKVCQDF	DANKFNFTKA	DLKEVLFQFT	--K-----	LAGEAD	NVS-----	-----	RS--	-----	VFPE	SAAV--	-----	GESPT	VVILIN-VSP-	IEYGHVLLIP	RVTDCLPQRI	
PhGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GAG	D-----	-----	DS--	-----	YFLN	NAPVTA-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PhGGP-2	FGMNQVLQPF	DSGKFNFTKV	RPEEVLFRFC	--E----	TDK	D-----	-----	SA--	-----	QYFG	GAPDTI--	-----	SASSS	AILIN-VSP-	IEYGHVLLIP	KIQDCLPQRI	
PvGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GAG	D-----	-----	DS--	-----	YFLN	NAPITA-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	HVLDCLPQRI	
PvGGP-2	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GAG	D-----	-----	DS--	-----	YFLN	NAPITA-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PvGGP-3	-----	PF	DSGKFNFTKV	RPEEVLFRFC	--E----	TDK	D-----	SA--	-----	QYFG	GAPDTI--	-----	SASSS	VILIN-VSP-	IEYGHVLLIP	KIQDCLPQRI	
PhvGGP-1	FRVDKVLQPF	DESKFNFTKV	GQEEVLFQFE	--A----	SDD	G-----	-----	QI--	-----	QFFF	NAPVDV--	-----	DNSTP	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
PhvGGP-2	FRVDKVLQPF	DESKFNFTKV	GQEEVLFQFE	--P----	SDD	G-----	-----	EA--	-----	QFSP	NAPIDV--	-----	DNSTP	FVAIN-VSP-	IEYGHVLLIP	RIFECLPQRI	
PpGGP-1	FRIDQVLQPF	DPKKNFNTKV	GQEEVLFQFM	--P----	NDS	D-----	-----	MS--	-----	EYFE	KATV--	-----	SSSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PpGGP-2	FRVDQVLQAF	DPKKNFNTKV	GQEEVLFQFM	--P----	SED	E-----	-----	VS--	-----	DYFE	KAPV--	-----	LASPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PpGGP-3	FRVDQVLQPF	DSKKNFNTKV	GQEEVLFQFG	--P----	SED	G-----	-----	VS--	-----	EYFE	KALV--	-----	LNSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PaGGP-1	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	N-----	-----	KV--	-----	QYLE	KALV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PaGGP-2	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	N-----	-----	KV--	-----	QYLE	KALV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PaGGP-3	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	N-----	-----	KV--	-----	QYLE	KAPV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PsGGP-1	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SAE	N-----	-----	KV--	-----	QYLE	KALV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PsGGP-2	FRVDKVLQDF	DRSKFNFTKV	GQEEVLFQFE	--E----	SEE	G-----	-----	KV--	-----	QYLE	KAPV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PsGGP-3	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SGE	D-----	-----	KV--	-----	QYLE	KAPV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PtGGP-1	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	N-----	-----	KV--	-----	QYLE	KALV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PtGGP-2	FRVDKVLQDF	DPCKFNFTKV	GQEEVLFQFE	--E----	SEE	D-----	-----	KV--	-----	QYLE	KAPV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PtGGP-3	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	N-----	-----	KI--	-----	QYLE	KAPV--	-----	FDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PtGGP-4	FRVDKVLQDF	DPSKFNFTKV	GQEEVLFQFE	--E----	SEE	D-----	-----	KV--	-----	QYLE	KAPV--	-----	LDSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PtGGP-5	FRVDKVLQDF	DPCKFNFTKV	GQEEVLFQFE	--E----	NEE	G-----	-----	KV--	-----	QYFE	KAPV--	-----	LDSPN	VVAIN-VSP-	VEYGHVLLIP	RVLDCLPQRI	
PotGGP-1	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFG	--A----	SED	G-----	-----	EV--	-----	QFFF	DAPIDP--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PotGGP-2	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFG	--E----	SED	G-----	-----	EV--	-----	KFFF	DATIDA--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
PotGGP-3	FRVDKVLQDF	DETKFNFTKI	GH--FRFE	--K----	SID	H-----	-----	NR--	-----	HFFP	SAPITAD-S	-----	NNSNSS	VVAIN-VSP-	IEYGHVLLIP	QVNLDCLPQRI	
PrpGGP	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	EV--	-----	QFIP	SAPIEP--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILEHLPQRI	
RcGGP-1	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	G-----	-----	DI--	-----	QFFF	SAPIDL--	-----	KNSTP	VVAIN-VSP-	IEYGHVLLIP	RILEHLPQRI	
RcGGP-2	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFE	--A----	SED	D-----	-----	DI--	-----	QFFF	SAPIDV--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RILDCLPQRI	
RcGGP-3	FRVDNVLQSF	DDSKFNFTKI	GQEEVLFQFE	--Q----	GQE	N-----	-----	MS--	-----	HFFP	NPPPSADD-	-----	SSSPS	IVAIN-VSP-	IEYGHVLLIP	RVLDCFPQRI	
SpGGP-1	FRIDKVLQPF	DENKFNFTKV	GQEEVLFQFG	--E----	SED	G-----	-----	EG--	-----	RFFT	DATIDP--	-----	ENSPS	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SpGGP-2	FRVDKVLQPF	DGNKFNFTKV	GQEEVLFQFG	--A----	SEE	G-----	-----	EA--	-----	QFFF	DAPVDP--	-----	ENSTP	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SpGGP-3	FRVDKVLQDF	DETKFNFTKI	GQEEVLFQFE	--K----	SND	H-----	-----	KR--	-----	RFFF	TAPVPTAD-	-----	SDSSR	VVAIN-VSP-	IEYGHVLLIP	QVNLDCLPQRI	
SmGGP-1	FRVDQVLQDF	DHKKFNFTKV	GQEEMLFCFE	--Q----	SNN	D-----	-----	KS--	-----	HYHS	KAYI--	-----	RGSPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCIPQRI	
SmGGP-2	FRVDQVLQDF	DSKFNFTKV	GQEEVLFQFD	--P----	SED	G-----	-----	EV--	-----	QFFF	SAPITAD-S	-----	ENSPF	ATSSNMVSP-	IEYGHVLLIP	RVLDCIPQRI	
SmGGP-3	FRVDQVLQDF	DPKKNFNTKV	NRNEVLFQFG	--E----	GE-	D-----	-----	EA--	-----	GYDD	ITPI--	-----	LSSPD	FVLIN-VSP-	IEYGHVLLIP	RVPDFVQRI	
SlGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GAG	D-----	-----	DS--	-----	YFLN	NAPITV-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SlGGP-2	FGMNQVLQPF	DSGKFNFTKV	RPEEVLFRFC	--E----	ADK	D-----	-----	SA--	-----	QYFN	DAPDTV--	-----	SSSSS	AILIN-VSP-	IEYGHVLLIP	QIQDCLPQRI	
SvGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GAG	D-----	-----	DS--	-----	YFLN	NAPITV-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SvGGP-2	FGMNQVLQPF	DSGKFNFTKV	RPEEVLFRFC	--E----	ADK	D-----	-----	SA--	-----	QYFN	DAPDTI--	-----	SSSSS	AILIN-VSP-	IEYGHVLLIP	QIQDCLPQRI	
SlGGP-1	FRVDKVLQPF	DGSKFNFTKV	GQEEVLFQFE	--A----	SEE	D-----	-----	EV--	-----	QLYP	DAPIDP--	-----	EKSPS	VVAIN-VSP-	IEYGHVLLIP	KVLECLPQRI	
SlGGP-2	FCIDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--P----	STD	Y-----	-----	KA--	-----	HYFS	GMRVNS--	-----	GISPS	IVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
StGGP-1	FRVDKVLQPF	DGSKFNFTKV	GQEEVLFQFE	--A----	SEE	D-----	-----	EV--	-----	QLYP	DAPIDP--	-----	EKSPS	VVAIN-VSP-	IEYGHVLLIP	KVLECLPQRI	
StGGP-2	FCIDKVLQPF	DENKFNFTKV	GQEEVLFQFE	--P----	STD	Y-----	-----	KP--	-----	HYFS	GMRVNG--	-----	GISPS	IVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SbGGP-1	FRVDKVLQPF	DPKKNFNTKV	GQEEVLFQFE	--N----	GDD	D-----	-----	DS--	-----	YFLN	DAPITV-A-	-----	DRAPN	VVAIN-VSP-	IEYGHVLLIP	RVLDCLPQRI	
SbGGP-2	FTMNQVLQPF	VSEKFNFTKV	SPEVLFQFE	--E----	TEK	D-----	-----	SA--	-----	QYFD	GVPDVT--	-----	SASSS	AILIN-VSP-	IEYGHVLLIP	KIQDCLPQRI	
SfGGP-1	FRVDQVLQPF	DGKKNFNTKV	GQEEVLFQFE	--E----	SED	G-----	-----	TS--	-----	QYFE	KAPV--						



















PaGGP-2	P-----	-----	-----	-----	-----	-----
PaGGP-3	-----	-----	-----	-----	-----	-----
PsGGP-1	P-----	-----	-----	-----	-----	-----
PsGGP-2	P-----	-----	-----	-----	-----	-----
PsGGP-3	-----	-----	-----	-----	-----	-----
PtGGP-1	P-----	-----	-----	-----	-----	-----
PtGGP-2	-----	-----	-----	-----	-----	-----
PtGGP-3	P-----	-----	-----	-----	-----	-----
PtGGP-4	P-----	-----	-----	-----	-----	-----
PtGGP-5	-----	-----	-----	-----	-----	-----
PotGGP-1	H-----	-----	-----	-----	-----	-----
PotGGP-2	Q-----	-----	-----	-----	-----	-----
PotGGP-3	Y-----	-----	-----	-----	-----	-----
PrpGGP	Q-----	-----	-----	-----	-----	-----
RcGGP-1	Q-----	-----	-----	-----	-----	-----
RcGGP-2	Q-----	-----	-----	-----	-----	-----
RcGGP-3	H-----	-----	-----	-----	-----	-----
SpGGP-1	Q-----	-----	-----	-----	-----	-----
SpGGP-2	H-----	-----	-----	-----	-----	-----
SpGGP-3	H-----	-----	-----	-----	-----	-----
SmGGP-1	V-----	-----	-----	-----	-----	-----
SmGGP-2	-----	-----	-----	-----	-----	-----
SmGGP-3	-----	-----	-----	-----	-----	-----
SiGGP-1	Q-----	-----	-----	-----	-----	-----
SiGGP-2	-----	-----	-----	-----	-----	-----
SvGGP-1	Q-----	-----	-----	-----	-----	-----
SvGGP-2	-----	-----	-----	-----	-----	-----
SlGGP-1	-----	-----	-----	-----	-----	-----
SlGGP-2	Q-----	-----	-----	-----	-----	-----
StGGP-1	-----	-----	-----	-----	-----	-----
StGGP-2	Q-----	-----	-----	-----	-----	-----
SbGGP-1	Q-----	-----	-----	-----	-----	-----
SbGGP-2	-----	-----L-----	-----	-----	-----	-----
SfGGP-1	QV-----	-----	-----	-----	-----	-----
SfGGP-2	-----	-----	-----	-----	-----	-----
SppGGP-1	-----	-----	-----	-----	-----	-----
SppGGP-2	Q-----	-----	-----	-----	-----	-----
TcGGP-1	Q-----	-----	-----	-----	-----	-----
TcGGP-2	Q-----	-----	-----	-----	-----	-----
TpGGP	Q-----	-----	-----	-----	-----	-----
VvGGP-1	Q-----	-----	-----	-----	-----	-----
VvGGP-2	-----	-----	-----	-----	-----	-----
VcGGP	-----	-----	-----	-----	-----	-----
ZmGGP-E-1	P-----	-----	-----	-----	-----	-----
ZmGGP-E-2	-----	-----LG-----	-----	-----	-----	-----
ZmGGP-1	P-----	-----	-----	-----	-----	-----
ZmGGP-2	-----	-----LG-----	-----	-----	-----	-----
ZomGGP	Q-----	-----	-----	-----	-----	-----