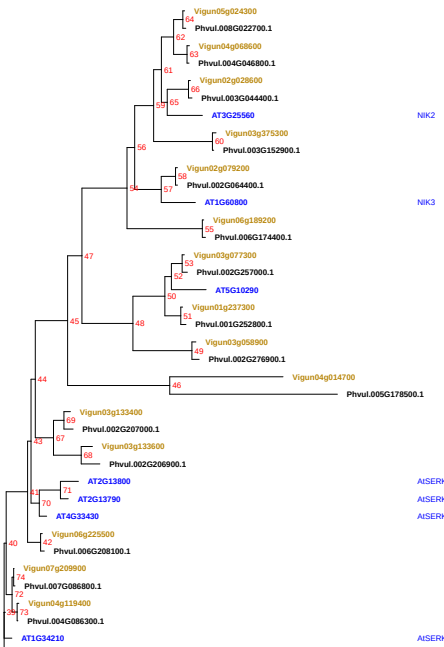

		bjornsen_pamps					Butselaar2024_Col0		Pv_counts					Vu_counts					genome	
		tg2	ef1a	rbl20	vg	cbfa	X2.5ug_pmol	X4.5ug_pmol	Pv_leading	Pvegdry	Pvzim	Pvstem	Pvstrm	Vu_leg	Vu_leg	Vu_leg	Vu_leg	Vu_leg		
Vigun05g024300														2.05	2.93	1.08	3.17	2.7		
Phvul.008G022700.1									8.74	7.66	6.84	8.65	8.69							
Vigun04g068600														1.47	3.42	2.03	3.5	2.53		
Phvul.004G046800.1									6.89	6.02	5.73	6.97	6.72							
Vigun02g028600														5.03	3.81	4.37	3.39	4.73		
Phvul.003G044400.1									8.58	8.05	7.98	8.64	8.71							
AT3G25560	NIK2	−0.24	0.31	0.17	0.24	0.62	8.56	8.22												
Vigun03g375300														5.87	5.81	5.6	5.24	6.16		
Phvul.003G152900.1									6.60	6.28	5.97	6.87	6.74							
Vigun02g079200														7.09	7.48	7.05	7.55	7.73		
Phvul.002G064400.1									9.90	9.34	9.19	9.98	9.99							
AT1G60800	NIK3	0.13	0.11	0.04	−0.12	−0.06	10.02	9.58												
Vigun06g189200														4.77	4.83	4.31	4.82	5.25		
Phvul.006G174400.1									6.17	6.11	6.09	6.37	6.07							
Vigun03g077300														7.45	7.66	7.59	8.53	7.34		
Phvul.002G257000.1									8.52	8.64	8.50	8.69	8.58						a	Pvul218cds.fa
AT5G10290		0.53	0.64	0.35	0.48	0.59	8.35	8.56												
Vigun01g237300														5.16	5.18	5.47	4.91	5.85		
Phvul.001G252800.1									9.10	8.81	8.84	9.12	9.13						a	TAIR10cds.fa
Vigun03g058900														7.15	6.46	6.44	6.64	6.64		
Phvul.002G276900.1									8.81	8.43	8.55	8.89	8.89						a	Vung469cds.fa
Vigun04g014700														7.64	6.73	7.62	6.87	6.5		
Phvul.005G178500.1									9.07	9.15	9.18	9.05	9.25							
Vigun03g133400														7.16	3.46	7.15	3.64	4.24		
Phvul.002G207000.1									2.33	6.35	8.13	2.91	3.44							
Vigun03g133600														6.75	7.17	5.96	6.75	6.09		
Phvul.002G206900.1									8.24	7.85	7.79	8.35	8.52							
AT2G13800	AtSERK5	3.25	3.53	2.45	3.24	2.25	6.70	7.66												
AT2G13790	AtSERK4	3.76	3.49	3.15	3.17	2.46	8.65	11.05												
AT4G33430	AtSERK3	2.85	2.28	2.15	1.60	1.08	7.95	8.72												
Vigun06g225500														6.07	6.32	6.11	6.45	5.44		
Phvul.006G208100.1									9.06	10.03	10.98	8.99	9.06							
Vigun07g209900														6.87	7.8	6.87	7.08	6.77		
Phvul.007G086800.1									7.41	7.21	7.17	7.52	7.59							
Vigun04g119400														1.51	2.87	2.96	2.6	2.5		
Phvul.004G086300.1									6.80	7.61	7.77	6.79	6.89							
AT1G34210	AtSERK2	0.11	−0.03	0.04	0.24	0.11	9.26	8.85												
AT1G71830	AtSERK1	0.03	0.31	0.17	−0.04	0.35	6.93	5.59												