

Supplementary Material

Transcriptome analysis showed the metabolic pathway of differentially expressed genes (DEGs) in resistant and susceptible soybean (*Glycine max*) to sclerotinia stem rot (SSR) and candidate gene mining

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Table S1. Phenotypic data of soybean resistance to SSR used for candidate gene based association analysis at two developmental stages.

No.	Stem rot length at seedling stage (cm)	Stem rot length at podding stage (cm)
Line01	1.53	7.70
Line02	3.00	5.83
Line03	5.00	3.80
Line04	6.03	5.03
Line05	5.67	5.17
Line06	1.63	5.47
Line07	4.73	5.37
Line08	5.23	1.97
Line09	5.77	6.83
Line10	1.50	7.53
Line11	5.87	5.37
Line12	2.43	5.87
Line13	2.70	5.53
Line14	8.47	9.40
Line15	4.10	7.13
Line16	1.23	5.73
Line17	2.50	6.37
Line18	6.47	5.10
Line19	6.50	5.20
Line20	4.93	5.57
Line21	6.37	6.50
Line22	3.20	3.03
Line23	3.03	5.27
Line24	1.60	5.73
Line25	6.13	4.57
Line26	4.10	4.80
Line27	6.40	5.37
Line28	6.13	0.00
Line29	5.53	6.03
Line30	3.33	4.40
Line31	1.90	5.10
Line32	5.03	9.17
Line33	2.53	7.60
Line34	3.50	5.90
Line35	3.37	7.57
Line36	2.93	3.97
Line37	7.00	7.17
Line38	9.07	5.77
Line39	4.47	5.20
Line40	10.10	6.60

Line41	6.10	6.17
Line42	4.07	6.97
Line43	4.97	8.60
Line44	4.17	4.93
Line45	1.70	6.50
Line46	7.13	5.00
Line47	1.07	6.83
Line48	1.30	7.40
Line49	6.17	5.80
Line50	6.70	5.83
Line51	4.40	5.37
Line52	2.83	7.47
Line53	5.53	2.83
Line54	0.63	4.03
Line55	2.47	6.23
Line56	5.80	4.90
Line57	1.43	6.77
Line58	4.03	5.50
Line59	3.93	4.30
Line60	1.27	7.67
Line61	7.80	5.77
Line62	3.53	0.00
Line63	3.27	2.60
Line64	4.63	6.50
Line65	6.87	8.17
Line66	1.33	4.60
Line67	1.87	4.80
Line68	1.43	4.77
Line69	4.17	6.80
Line70	3.80	3.90
Line71	7.27	5.83
Line72	4.30	4.03
Line73	5.57	5.43
Line74	7.33	6.37
Line75	1.63	4.43
Line76	4.67	5.17
Line77	7.00	3.63
Line78	1.17	5.03
Line79	3.93	4.83
Line80	3.37	6.23
Line81	5.70	3.10
Line82	2.53	5.97
Line83	1.77	5.07
Line84	1.10	6.67

Line85	1.10	2.63
Line86	5.00	5.53
Line87	1.27	5.70
Line88	6.70	7.80
Line89	7.43	4.07
Line90	7.13	7.40
Line91	5.67	4.60
Line92	5.03	5.97
Line93	4.27	6.83
Line94	4.17	7.57
Line95	1.97	5.23
Line96	8.23	4.70
Line97	6.97	5.50
Line98	5.17	6.13
Line99	4.30	3.97
Line100	7.80	4.07
Line101	4.03	5.10
Line102	3.40	4.30
Line103	1.07	3.50
Line104	7.63	4.73
Line105	1.50	6.60
Line106	6.53	3.37
Line107	1.67	7.67
Line108	6.03	5.17
Line109	3.53	5.73
Line110	4.00	5.40
Line111	3.57	6.47
Line112	5.77	4.60
Line113	7.57	5.87
Line114	4.97	3.33
Line115	3.83	5.70
Line116	3.63	5.03
Line117	2.83	4.07
Line118	2.17	5.57
Line119	2.50	0.00
Line120	5.07	4.97
Line121	2.50	2.27
Line122	7.07	4.27
Line123	1.57	3.50
Line124	3.23	3.70
Line125	1.90	3.97
Line126	4.87	3.90
Line127	5.17	3.30
Line128	4.33	6.60

Line129	5.33	4.20
Line130	9.67	4.23
Line131	8.23	5.43
Line132	4.33	4.70
Line133	3.17	4.70
Line134	5.20	6.03
Line135	2.13	3.83
Line136	5.73	12.10
Line137	4.37	7.10
Line138	3.90	9.63
Line139	5.87	6.07
Line140	1.40	5.67
Line141	6.33	5.67
Line142	3.53	4.23
Line143	2.13	5.40
Line144	9.67	4.70
Line145	2.63	4.63
Line146	2.97	5.77
Line147	4.20	5.43
Line148	3.60	5.13
Line149	5.83	5.33
Line150	4.43	5.87
Line151	5.00	9.83
Line152	6.33	6.53
Line153	4.27	3.80
Line154	3.20	2.40
Line155	4.33	0.00
Line156	6.23	4.67
Line157	5.27	4.63
Line158	7.70	0.00
Line159	2.47	5.33
Line160	3.87	5.03
Line161	4.00	5.13
Line162	6.70	9.07
Line163	7.43	7.77
Line164	1.43	6.03
Line165	4.13	5.50
Line166	3.30	4.50
Line167	6.43	0.00
Line168	8.60	6.60
Line169	6.00	4.23
Line170	7.23	6.67
Line171	8.40	5.50
Line172	3.70	4.50

Line173	1.30	4.03
Line174	5.10	4.37
Line175	5.67	4.97
Line176	6.63	5.33
Line177	5.80	10.93
Line178	7.17	7.13
Line179	5.37	4.80
Line180	3.20	5.87
Line181	5.53	5.20
Line182	3.80	5.43
Line183	6.40	6.40
Line184	3.10	5.93
Line185	3.03	4.27
Line186	5.37	4.27
Line187	5.87	8.60
Line188	3.40	4.73
Line189	7.00	6.27
Line190	5.23	7.10
Line191	1.13	4.40
Line192	4.30	4.10
Line193	3.47	6.40
Line194	1.33	4.73
Line195	4.53	3.77
Line196	1.80	4.20
Line197	2.00	4.87
Line198	2.63	3.80
Line199	5.10	5.53
Line200	6.90	0.00
Line201	2.13	7.00
Line202	1.90	5.13
Line203	5.80	6.67
Line204	3.40	7.60
Line205	0.73	5.33
Line206	2.63	4.83
Line207	4.73	6.80
Line208	3.30	4.43
Line209	8.93	4.43
Line210	4.10	8.60
Line211	1.33	8.27
Line212	1.60	4.93
Line213	1.37	3.67
Line214	2.23	8.60
Line215	1.57	3.63
Line216	2.97	4.90

Line217	1.77	5.80
Line218	4.30	6.47
Line219	1.47	5.60
Line220	2.53	3.33
Line221	3.63	4.60
Line222	2.23	4.87
Line223	1.60	6.40
Line224	6.67	4.50
Line225	1.60	6.50
Line226	1.90	4.10
Line227	1.37	8.03
Line228	1.27	3.07
Line229	6.37	6.03
Line230	7.47	8.83
Line231	1.87	4.80
Line232	5.23	4.40
Line233	2.80	4.90
Line234	7.80	4.47
Line235	6.03	5.20
Line236	6.00	5.13
Line237	3.57	4.93
Line238	4.67	5.00
Line239	6.30	7.40
Line240	6.93	6.83
Line241	4.47	7.83
Line242	4.80	5.63
Line243	7.30	4.63
Line244	7.87	5.40
Line245	7.53	5.13
Line246	1.73	5.47
Line247	2.13	7.47
Line248	1.23	5.87
Line249	3.23	4.33
Line250	1.33	5.33
Line251	4.97	6.07
Line252	6.07	5.17
Line253	5.87	5.57
Line254	1.30	2.87
Line255	3.10	7.53
Line256	5.80	5.80
Line257	4.37	9.67
Line258	3.83	5.27
Line259	6.47	5.27
Line260	5.60	4.93

Line261	3.40	4.07
Line262	2.30	8.77
Line263	4.77	10.10
Line264	8.07	6.03
Line265	4.43	8.23
Line266	1.20	7.97
Line267	3.10	3.83
Line268	4.33	3.87
Line269	7.30	5.20
Line270	6.93	3.10
Line271	1.13	3.63
Line272	6.53	4.73
Line273	6.53	6.67
Line274	4.47	2.77
Line275	3.60	6.37
Line276	7.47	3.07
Line277	7.10	5.43
Line278	2.70	4.00
Line279	1.73	6.17
Line280	3.90	5.43
Line281	5.83	3.07
Line282	9.03	5.33
Line283	5.00	3.07
Line284	1.17	4.40
Line285	6.23	8.10
Line286	0.93	4.90
Line287	7.93	5.07
Line288	6.40	4.97
Line289	4.83	4.80
Line290	3.50	7.50
Line291	6.73	3.50
Line292	7.07	4.53
Line293	6.23	5.30
Line294	4.73	5.90
Line295	6.30	6.60
Line296	4.33	4.67
Line297	2.93	6.00
Line298	8.73	4.47
Line299	6.07	7.67
Line300	6.00	5.67
Line301	8.20	3.87
Line302	7.87	4.03
Line303	4.57	4.00
Line304	3.73	3.30

Line305	4.03	4.27
Line306	6.63	6.03
Line307	1.77	6.10
Line308	8.00	4.43
Line309	9.60	4.80
Line310	7.30	4.07
Line311	4.93	3.43
Line312	4.73	4.87
Line313	4.27	4.77
Line314	4.53	6.73
Line315	6.03	4.27
Line316	8.27	6.77
Line317	8.37	5.40
Line318	4.87	4.60
Line319	2.07	4.80
Line320	3.73	7.10
Line321	3.40	3.33
Line322	3.13	5.70
Line323	4.80	4.23

Table S2. Transcriptome sequencing data and mapping of reference genome of "Maple Arrow " and "Hefeng25".

Soybean line	Sample	Total Clean Reads	Total Mapping Ratio (%)	Uniquely Mapping Ratio (%)	
Maple Arrow	0h-1	37122312	87.98	84.45	
	0h-2	37021526	86.32	82.36	
	0h-3	37065269	88.69	83.12	
	4h-1	51844902	87.57	84.88	
	4h-2	51625487	87.32	83.65	
	4h-3	51938745	87.97	84.96	
	8h-1	54964942	85.97	83.79	
	8h-2	54896325	86.12	82.97	
	8h-3	55036295	85.59	83.2	
	12h-1	34646054	88.48	83.06	
	12h-2	34598623	88.98	82.34	
	12h-3	34789523	88.76	81.98	
	24h-1	54477550	85.18	82.06	
	24h-2	54369852	85.56	81.96	
	24h-3	54489612	85.29	82.77	
	Hefeng25	0h-1	39368826	89.61	85.02
		0h-2	39402698	89.46	85.02
		0h-3	39268579	89.53	85.02
4h-1		43995366	88.39	84.74	
4h-2		43869527	88.98	83.96	
4h-3		43985327	88.72	84.12	
8h-1		43605594	88.16	80.78	
8h-2		43529836	88.96	81.28	
8h-3		44076958	88.42	82.25	
12h-1		46535342	87.76	81.78	
12h-2		46498265	87.32	82.08	
12h-3		46503187	87.87	82.36	
24h-1		44373796	82.12	80.27	
24h-2		44326958	84.03	81.02	
24h-3		44298630	83.42	80.98	