Additional file 4. Reference sequences included in the phylogenies for comparison and to root the trees. Where RLK subfamilies or generic terms (i.e. Pto-like) are cited, all sequences reported in the source were tested in the phylogenies, except pseudogenes or putative proteins with truncated kinase domains.

I - Reference sequences for phylogenies at the subfamily level				
Proteins	Species	Source		
CrRLK1L	Arabidopsis and rice	[50]		
Pto and relatives	Solanum pimpinellifolium	[39]		
WAK-like	Arabidopsis	[4, 49]		
LRR X	Arabidopsis	[4]		
LRR XII	Arabidopsis	[4]		
Best Pac Blastp matches	Arabidopsis, poplar and rice	NCBI GenBank		

## II - Reference sequences for phylogenetic investigation into the relations among Pto, Pto-like and CrRLK1L

Proteins	Species	Source
Best Pto Blastp matches (i.e. CrRLK1L)	Arabidopsis, Poplar, Castor bean, Grapevine, Rice	NCBI GenBank
Best Blastp matches obtained using wild/cultivated strawberry and bean Pto-like	Fragaria vesca, Phaseolus vulgaris	Phytozome
		F. vesca:
		mrna04250.1-v1.0-hybrid mrna16583.1-v1.0-hybrid
		mrna14010.1-v1.0-hybrid
		mrna23363.1-v1.0-hybrid
		P. vulgaris:
		Phvul.003G239400.1
		Phys. 1 007 C 188 200 1
		Phvul.007G188300.1
Pto and relatives	Solanum pimpinellifolium	[39]
Pto-like	Wild tomato species	[AF4.1]* [76]
Pto-like	Capsicum spp.	NCBI GenBank Acc. Nos: AY316952, AY316954-67
Pto-like	Solanum spp.	[70]
Pto-like	Nicotiana repanda	[75]
Pto-like	Phaseolus vulgaris	[69]
Pto-like	Cultivated and wild strawberry	[74]
Pto-like	Musa acuminata	[73]
Whole CrRLK1L subfamily	Arabidopsis and rice	[50]
Whole CrRLK1L subfamily	Poplar	[5]
Whole CrRLK1L subfamily	Moss	NCBI GenBank, Acc. Nos: XM_001757821, XM_001757822, XM_001760648, XM_001760727, XM_001779604
Whole CrRLK1L subfamily	Tomato	[6]
CrRLK1L-like	Platanus  imes acerifolia	This paper

III - Proteins used to root the trees				
Proteins	Species	Source		
CTR1, Raf kinase family	Arabidopsis	[3], NCBI GenBank		
MPK3, MAP kinase family	Arabidopsis	[3], NCBI GenBank		
CPK7, CDPK family	Arabidopsis	[3], NCBI GenBank		
Several other representatives of eukaryotic protein kinase families (ePKs)	Many species	[3]		
APH(3')III (aminoglycoside 3'-phosphotransferase)	Staphylococcus	[3]		

<sup>\*</sup> References not reported in the text.

AF4.1 Rose LE, Langley CH, Bernal AJ, Michelmore RW: Natural variation in the Pto disease resistance gene within species of wild tomato (Lycopersicon). I. Functional analysis of *Pto* alleles. Population genetics of Pto. *Genetics*. 2005, 171:345-357.