

Table S2. The identity numbers (GB acc. numbers or genomic scaffold numbers) of the sequences used in phylogenetic analyses.

<i>AtMFT</i>	NM101672		<i>JcFT</i>	KF113881	
<i>AtBFT</i>	NM_125597		<i>JcTFL1a</i>	NM_001306038	
<i>ATC</i>	NM_128315		<i>JcTFL1b</i>	NM_001306004	
<i>AtFT</i>	NM_105222		<i>JcTFL1c</i>	NM_001308743	
<i>AtTSF</i>	NM_118156		<i>SmoMFT</i>	XM_002990181	
<i>AtTFL1</i>	ATU77674		<i>SoMFT1</i>	scaff12981	
<i>AtriMFT</i>	XM_006849682		<i>SoMFT2</i>	scaff10102	
<i>AtriMFT1</i>	XM_006841802		<i>SoBFT</i>	scaff14048	
<i>AtriCENlike</i>	XM_006835987		<i>SoCEN1</i>	scaff4602	
<i>AtriHd3a</i>	XM_006858577		<i>SoCEN2</i>	scaff14350	
<i>BvMFT</i>	XM_010687957		<i>SoFTL1-1</i>	scaff16133	
<i>BvMFTlike</i>	HM448918		<i>SoFTL1-2a</i>	scaff8759	
<i>BvBFT</i>	HM448916		<i>SoFTL1-2b</i>	scaff8759	
<i>BvCEN</i>	HM448913		<i>SoFTL2</i>	scaff39498	
<i>BvCENlike</i>	XM_010685445		<i>SoTFL1</i>	scaff68707	scaff48765
<i>BvFT1</i>	HM448910		<i>VvFT</i>	DQ504308	
<i>BvFT2</i>	HM448912		<i>ZmZCN9</i>	EU241900	
<i>BvFTL3</i>	Bvchr6.scaff028	0551.scaff01119	<i>ZmZCN10</i>	NM_001112778	
<i>CrMFT1</i>	KU140602		<i>ZmZCN11</i>	NM_001112793	
<i>CrMFT2</i>	KU140603		<i>ZmZCN8</i>	EU241924	
<i>CrBFT</i>	KU140599		<i>ZmZCN15</i>	EU241930	
<i>CrCEN</i>	KU140601		<i>ZmZCN19</i>	EU241934	
<i>CrFTL1</i>	EU128013		<i>ZmZCN2</i>	EU241918	
<i>CrFTL2</i>	EF445636		<i>ZmZCN3</i>	EU241919	
<i>CrTFL1</i>	KU561656		genomic <i>CrFTL1</i>	KT992793	
<i>CrCAB1</i>	KU140600		genomic <i>CrFTL2</i>	KT992794	
<i>JcMFT</i>	NM_001306009		pseudo <i>CrFTL2</i>	KU140604	
<i>JcMFTlike</i>	NM_001308696		genomic <i>CrFTL3</i>	KT992795	