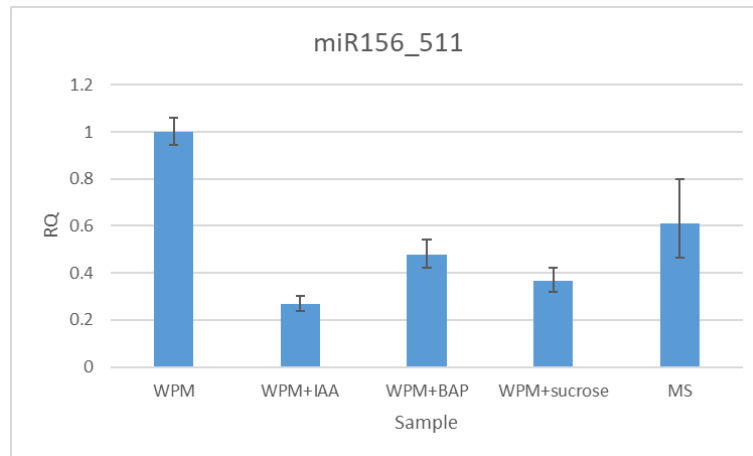
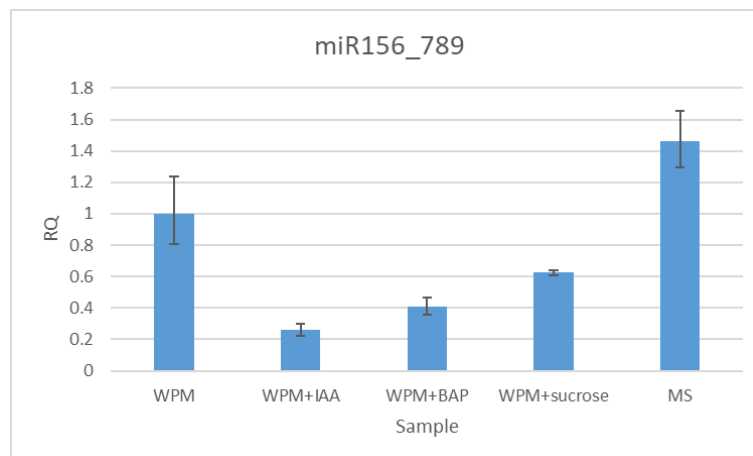


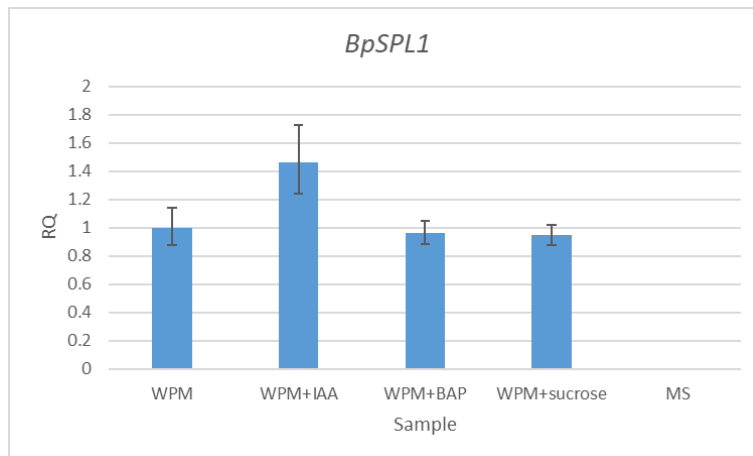
Supplementary file 5. Relative expression (RQ) of selected miRNAs and target genes in samples cultivated on various media. Bars indicate minimum and maximum RQ values. Samples labelled with the same letter are not statistically different ($p>0.05$). nd – expression not detected. The WPM sample was set as the reference sample.



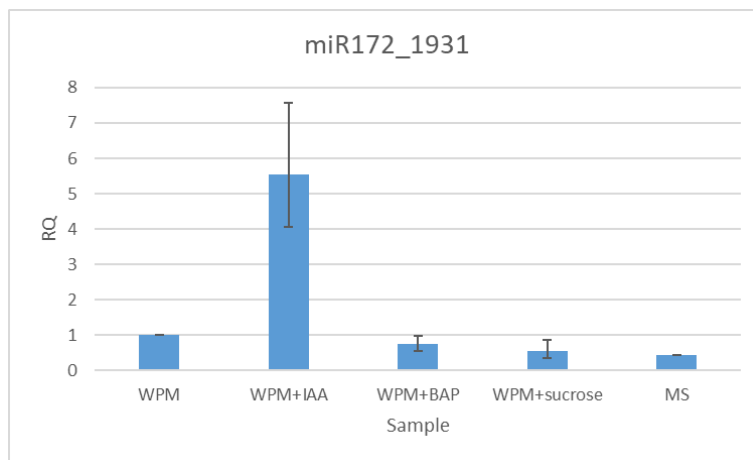
miR156_511			
Sample	RQ	RQ Min	RQ Max
WPM	1	0.944914	1.058298
WPM+IAA	0.269371	0.238948	0.303668
WPM+BAP	0.479791	0.424097	0.542799
WPM+sucrose	0.366425	0.317434	0.422977
MS	0.608901	0.464029	0.799002



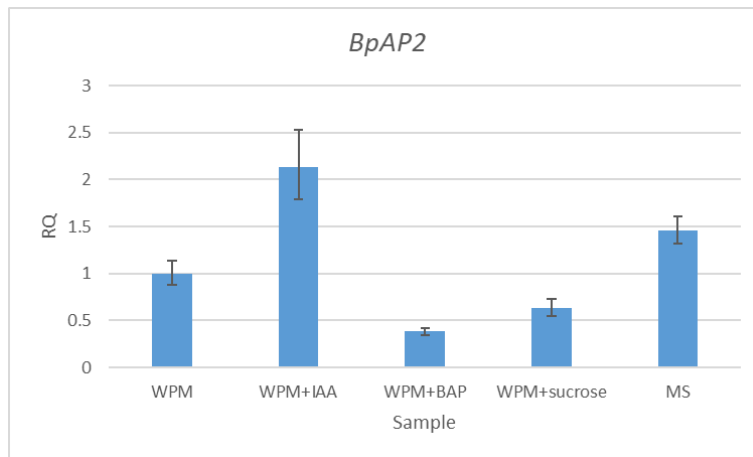
miR156_789			
Sample Name	RQ	RQ Min	RQ Max
WPM	1	0.809474	1.23537
WPM+IAA	0.258103	0.223357	0.298254
WPM+BAP	0.407727	0.357751	0.464685
WPM+sucrose	0.624909	0.610515	0.639643
MS	1.464113	1.296434	1.653478



<i>BpSPL1</i>			
Sample Name	RQ	RQ Min	RQ Max
WPM	1	0.874734	1.143204
WPM+IAA	1.462788	1.238252	1.728041
WPM+BAP	0.963452	0.886002	1.047673
WPM+sucrose	0.946563	0.879723	1.018481
MS	nd	-	-



<i>miR172_1931</i>			
Sample Name	RQ	RQ Min	RQ Max
WPM	1	-	-
WPM+IAA	5.540892	4.051479	7.577845
WPM+BAP	0.739383	0.559748	0.976667
WPM+sucrose	0.553312	0.353219	0.866757
MS	0.428192	-	-



<i>BpAP2</i>			
Sample Name	RQ	RQ Min	RQ Max
WPM	1	0.879362	1.137188
WPM+IAA	2.129686	1.795091	2.526648
WPM+BAP	0.383896	0.348718	0.422623
WPM+sucrose	0.631391	0.546629	0.729296
MS	1.452773	1.316764	1.602831