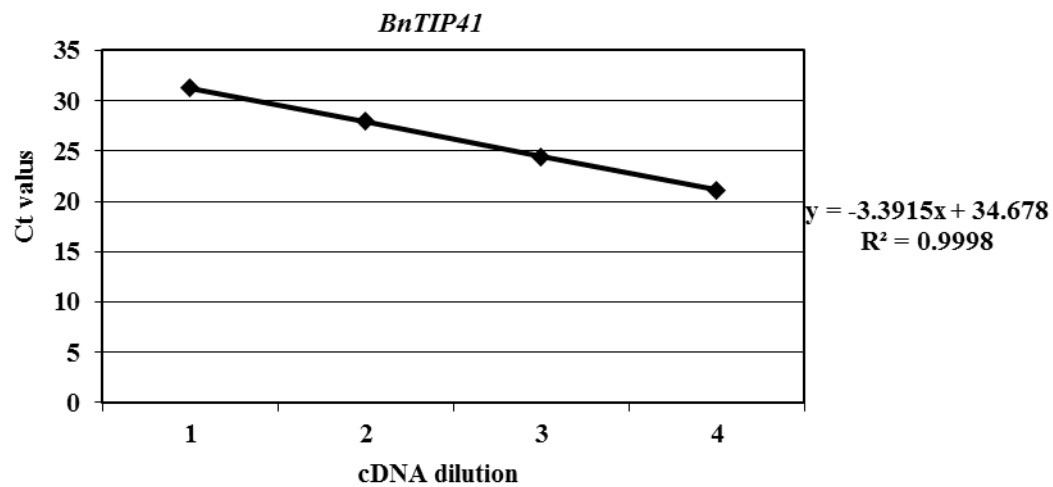


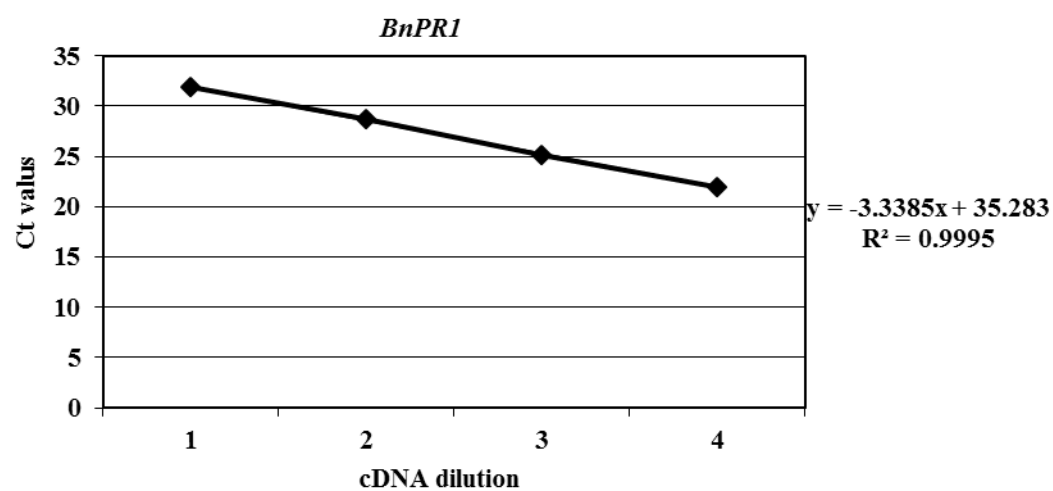
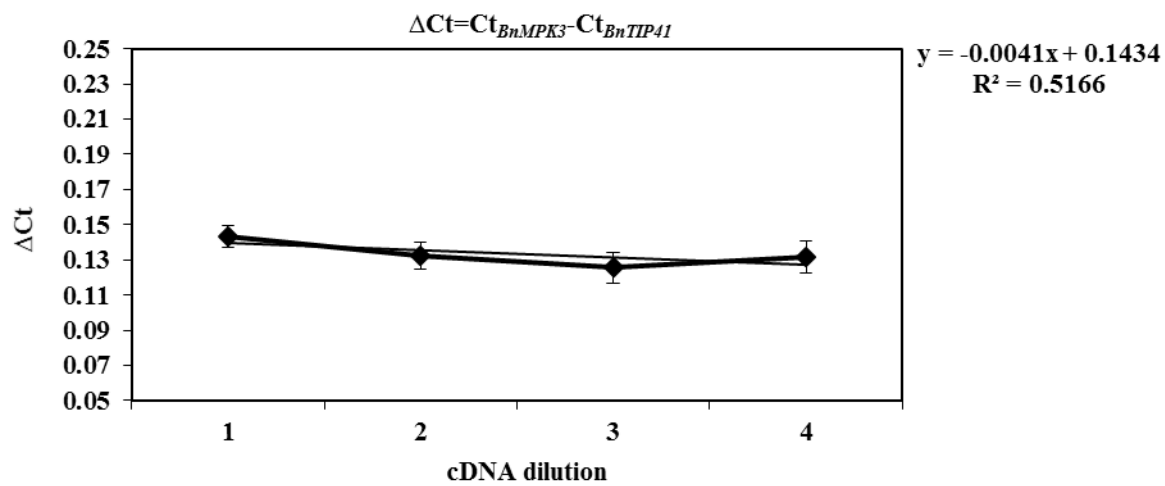
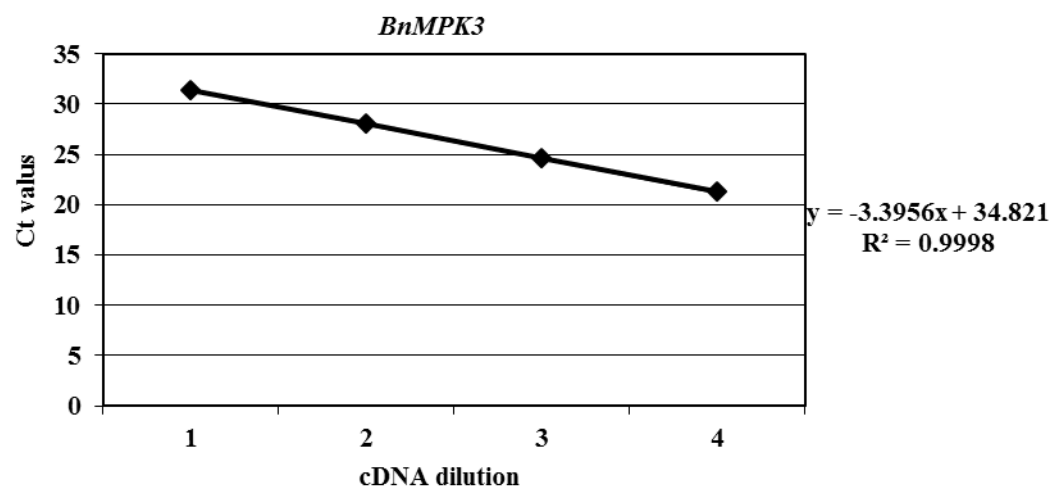
Supplementary File 1

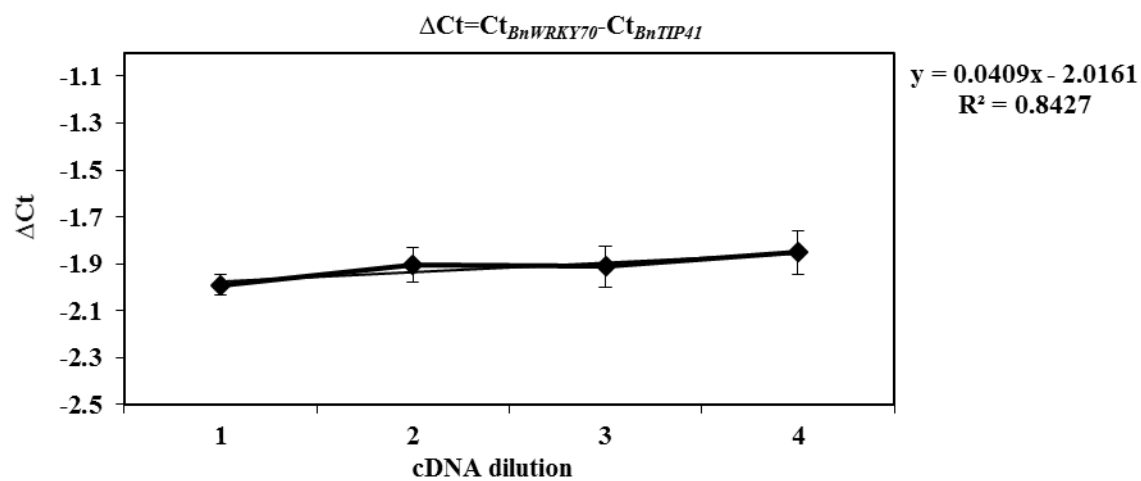
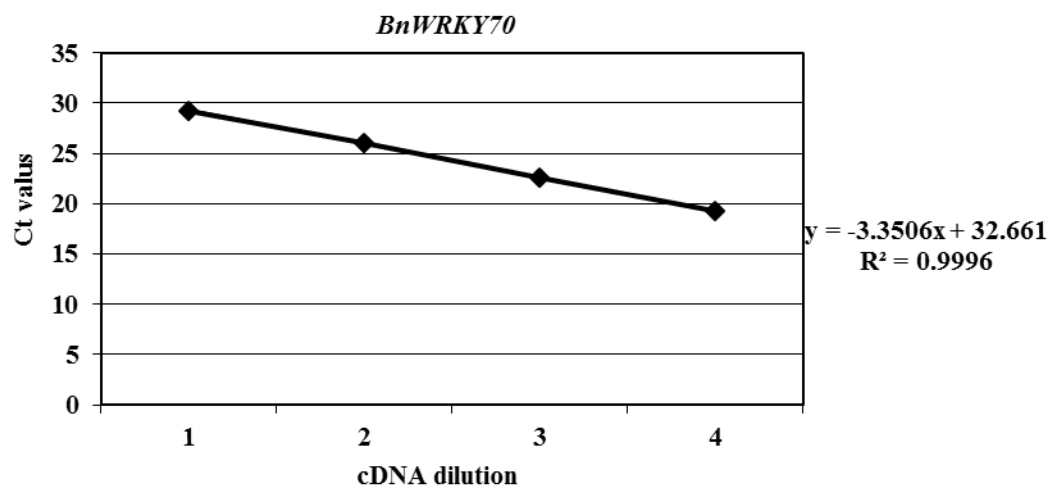
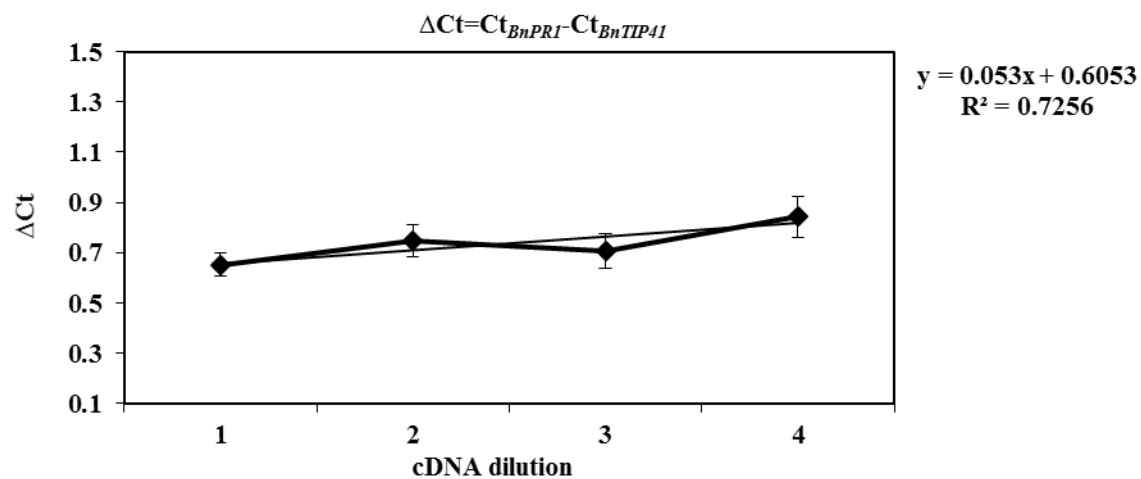
The efficiencies (E) of the primer-specific polymerase chain reaction (PCR) amplifications and variation of ΔCT ($Ct_{Target}-Ct_{TIP41}$) with template dilution

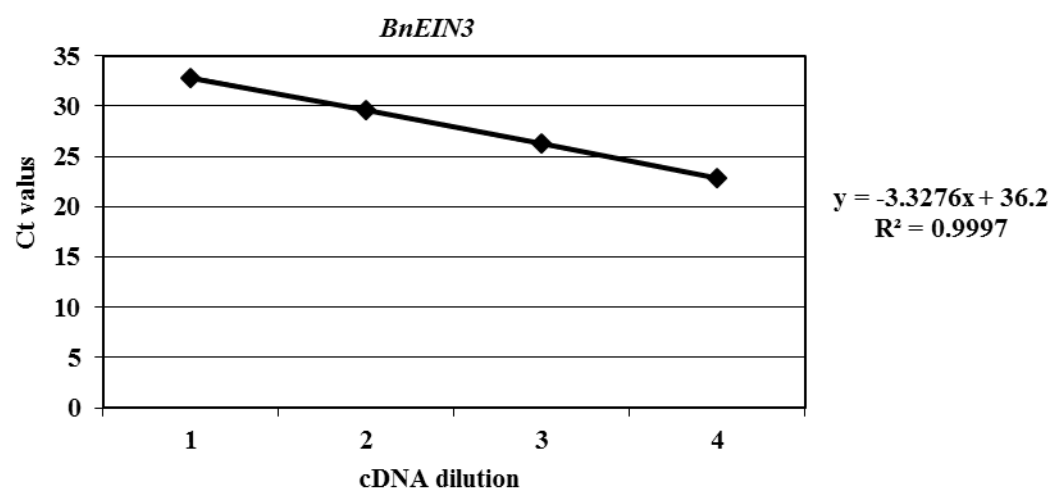
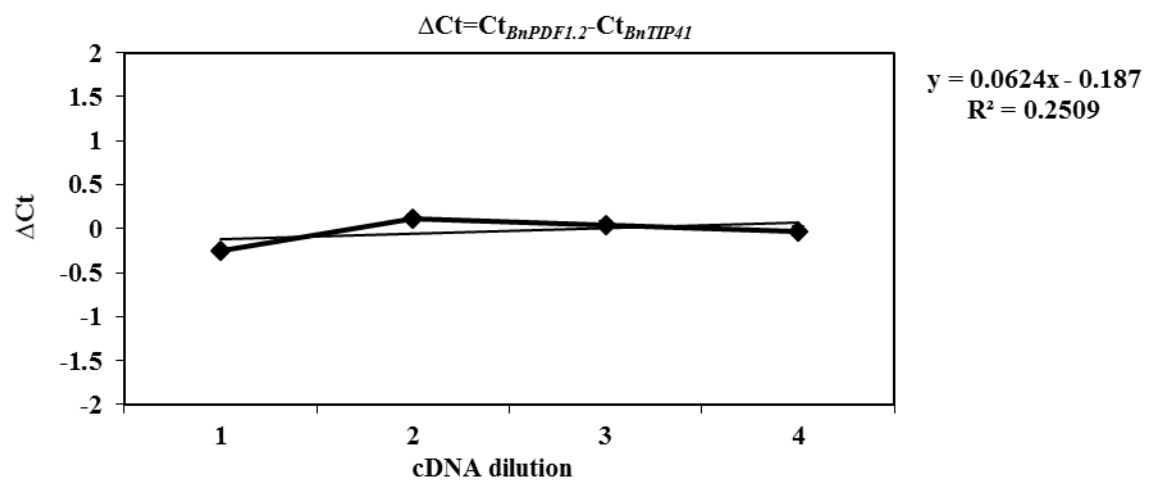
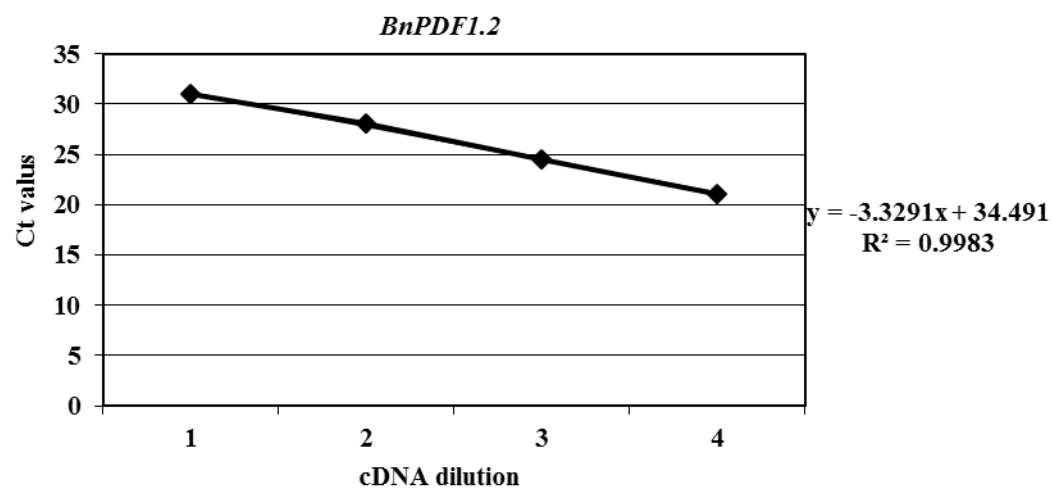
Gene	R ²	Slope	E
<i>BnTIP41</i>	0.9998	-3.3915	97.18%
<i>BnMPK3</i>	0.9998	-3.3956	97.01%
<i>BnPR1</i>	0.9995	-3.3385	99.31%
<i>BnWRKY70</i>	0.9996	-3.3506	98.82%
<i>BnRDF1.2</i>	0.9983	-3.3291	99.70%
<i>BnEIN3</i>	0.9997	-3.3276	99.76%
<i>BnVSP1</i>	0.9997	-3.3514	98.78%
<i>BnVSP2</i>	0.9999	-3.3414	99.19%
<i>BnJR1</i>	1	-3.4384	95.36%
<i>BnPAL</i>	0.9997	-3.3658	98.20%
<i>BnICS1</i>	0.9998	-3.3368	99.38%
<i>BnLOX2</i>	0.9999	-3.3608	98.40%
<i>BnAOS</i>	0.9991	-3.4404	95.28%
<i>BnACS1</i>	0.9997	-3.3708	98.00%
<i>BnACS2</i>	0.9996	-3.3688	98.08%
<i>BnACS6</i>	0.9999	-3.4091	96.49%
<i>BnACS7</i>	0.9995	-3.3591	98.47%
<i>BnACS8</i>	0.9995	-3.3644	98.26%

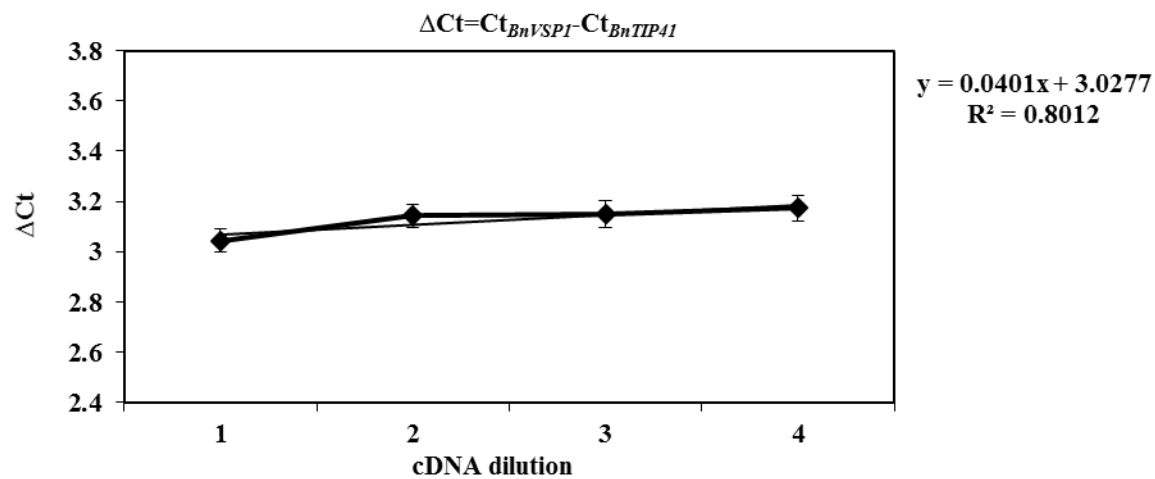
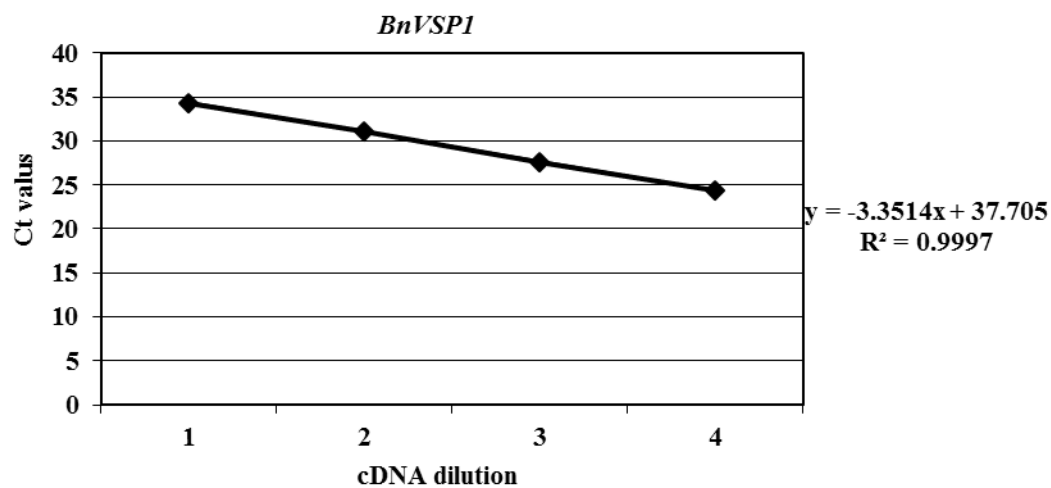
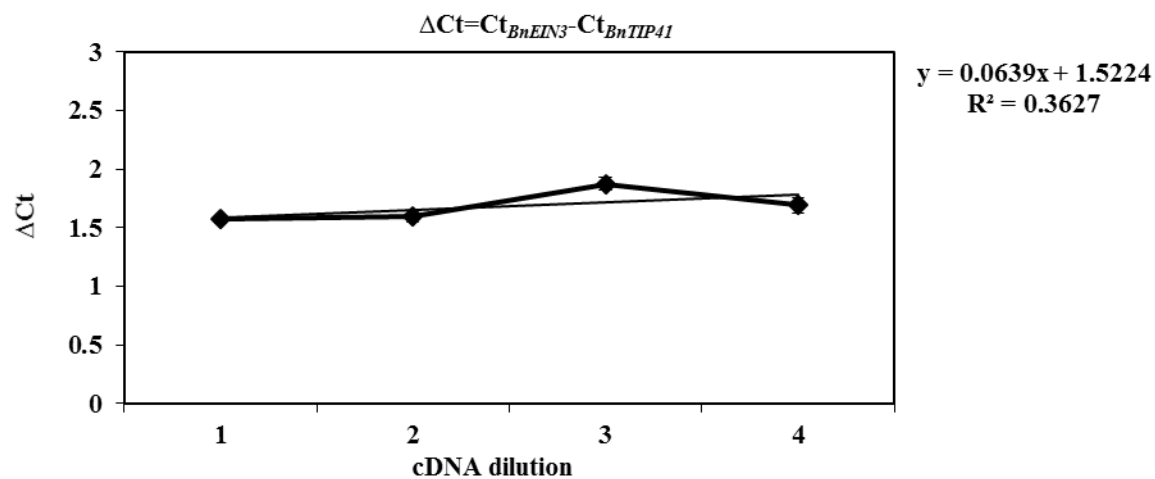
$$E=10^{(1/-\text{slope})}-1$$

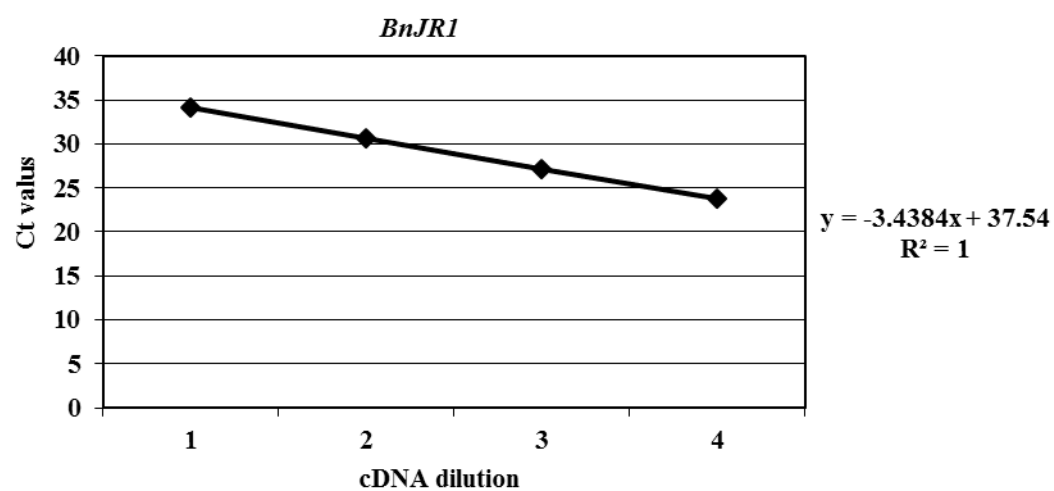
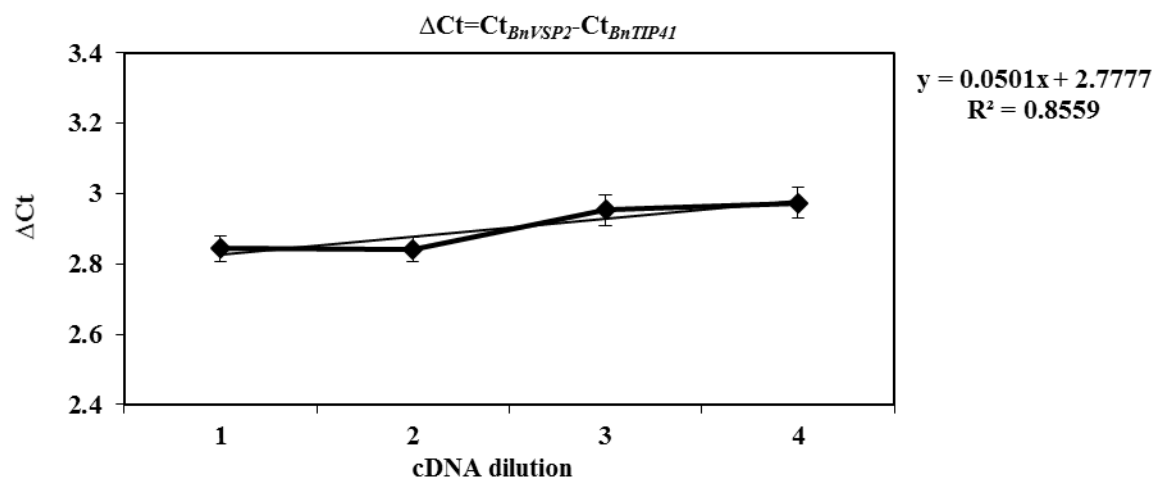
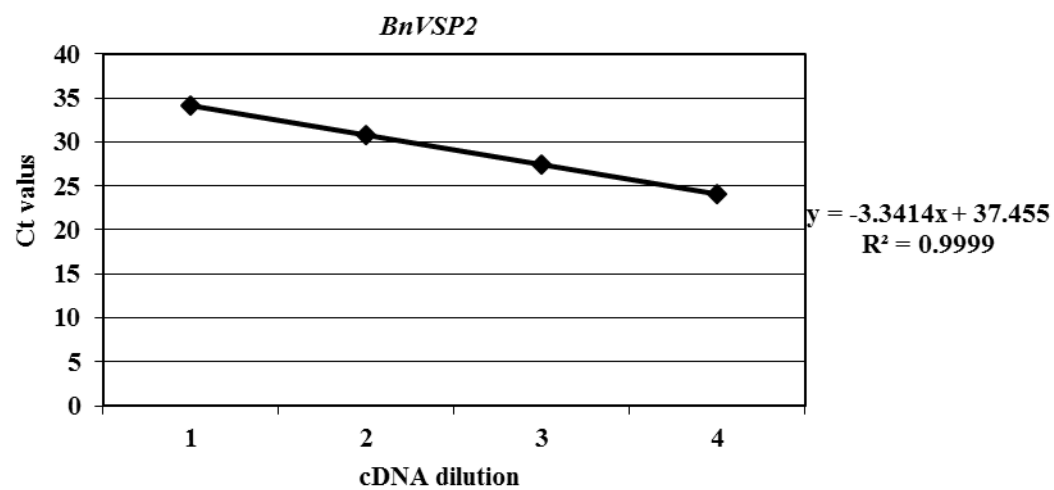


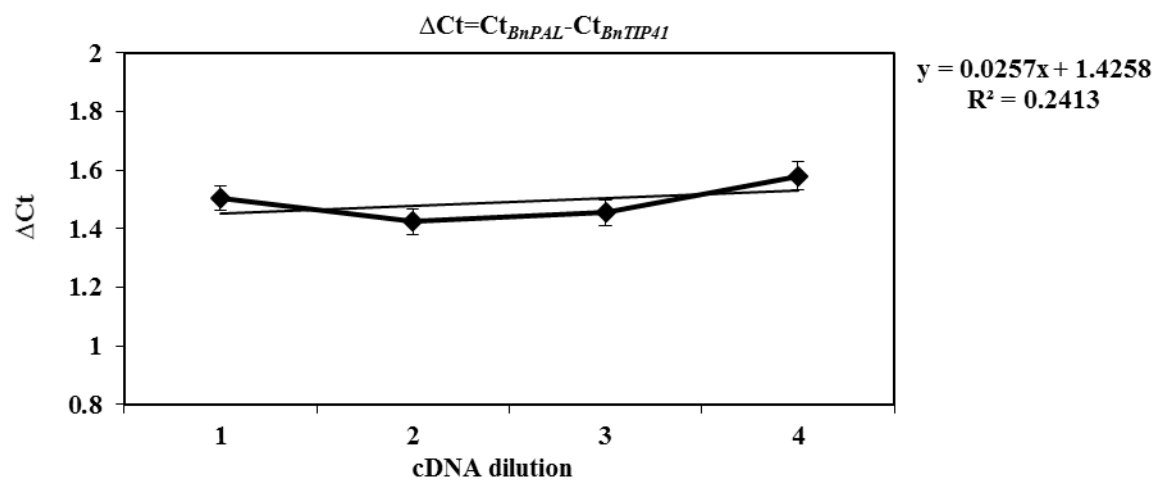
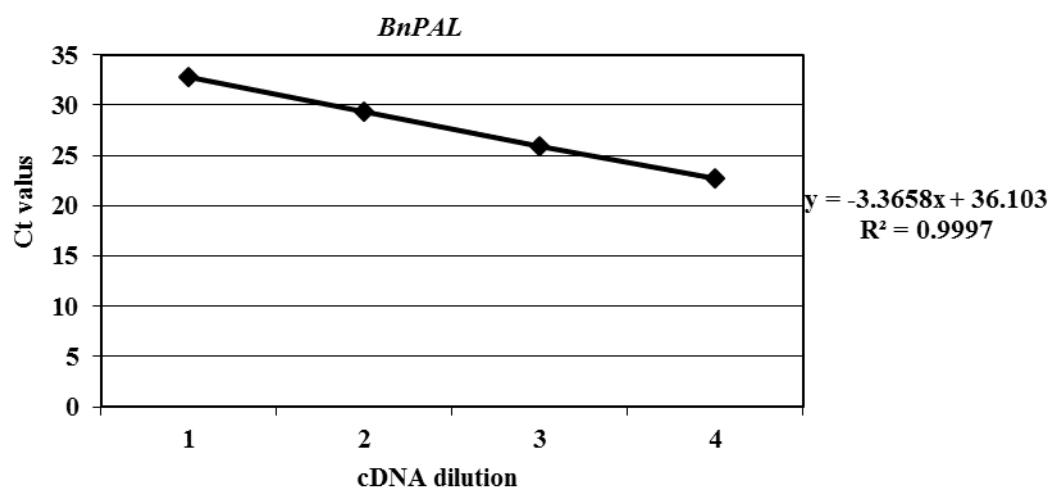
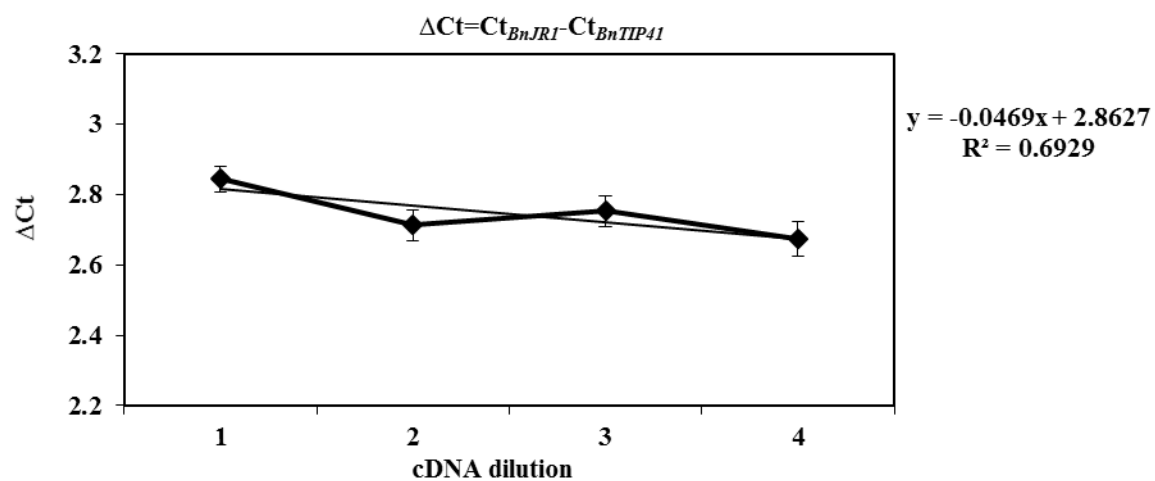


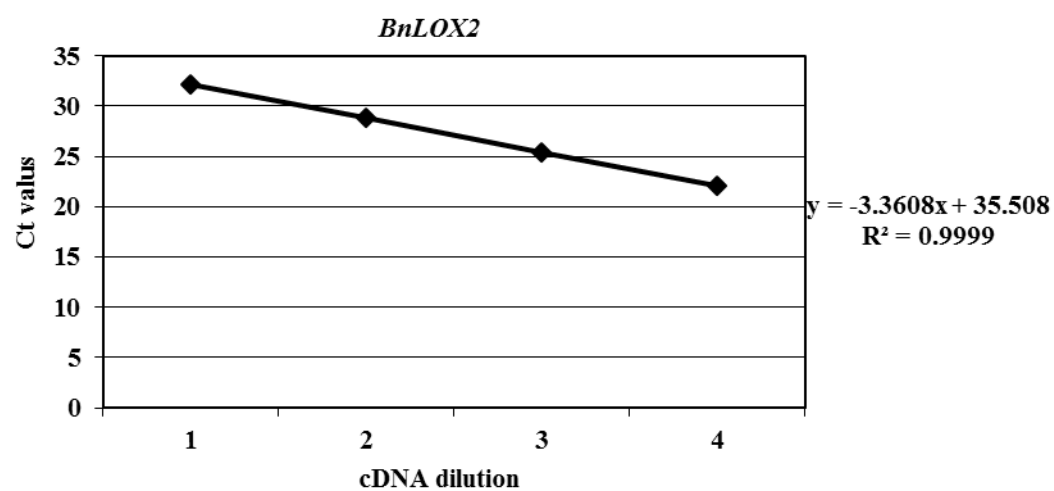
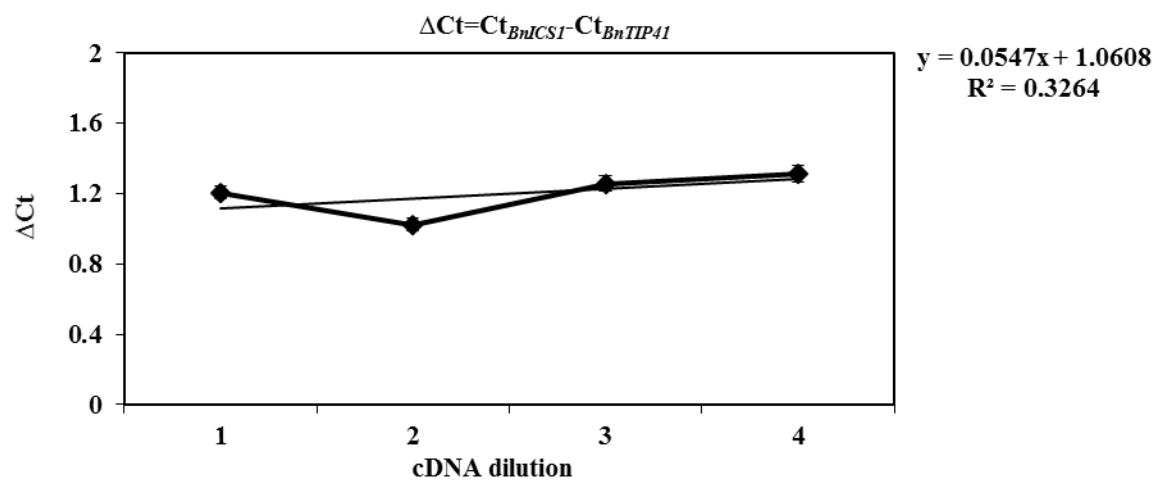
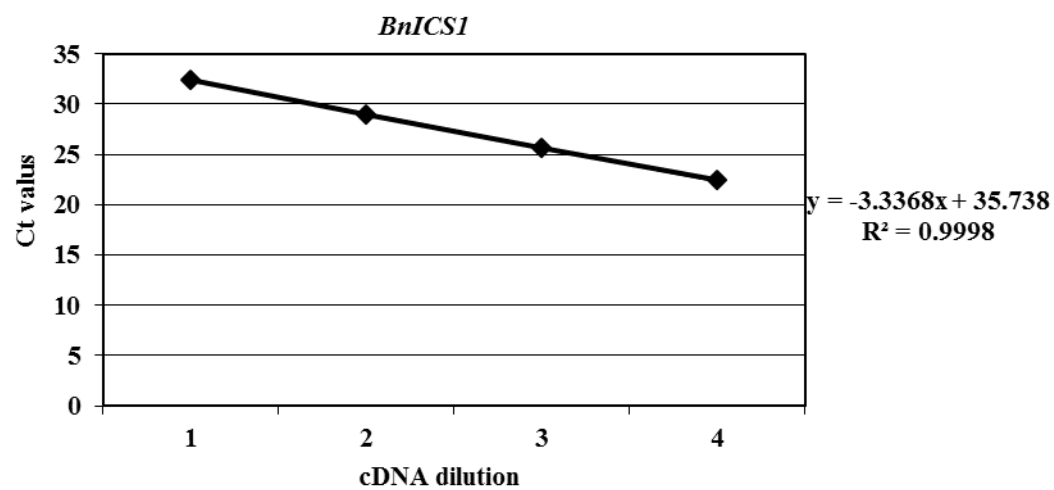


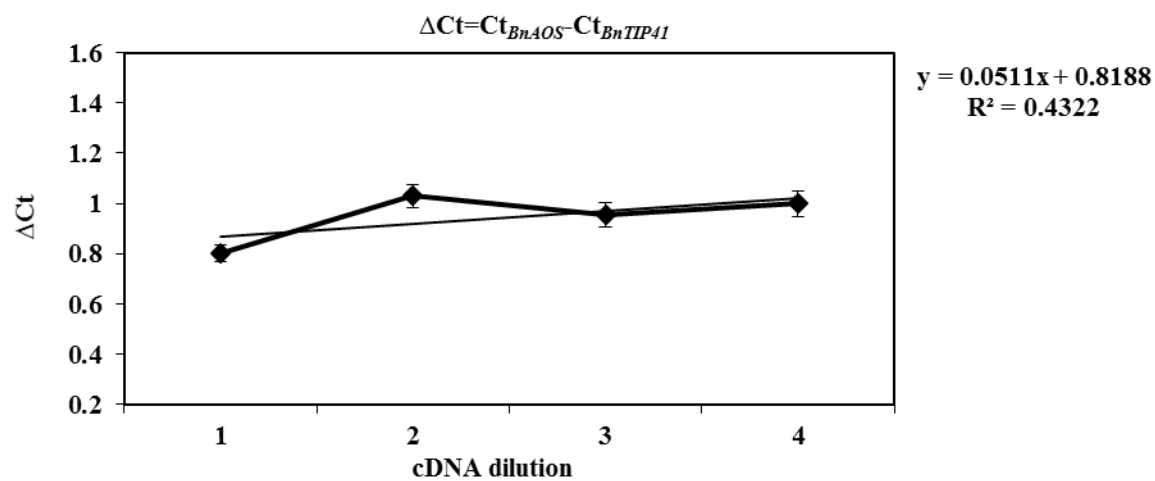
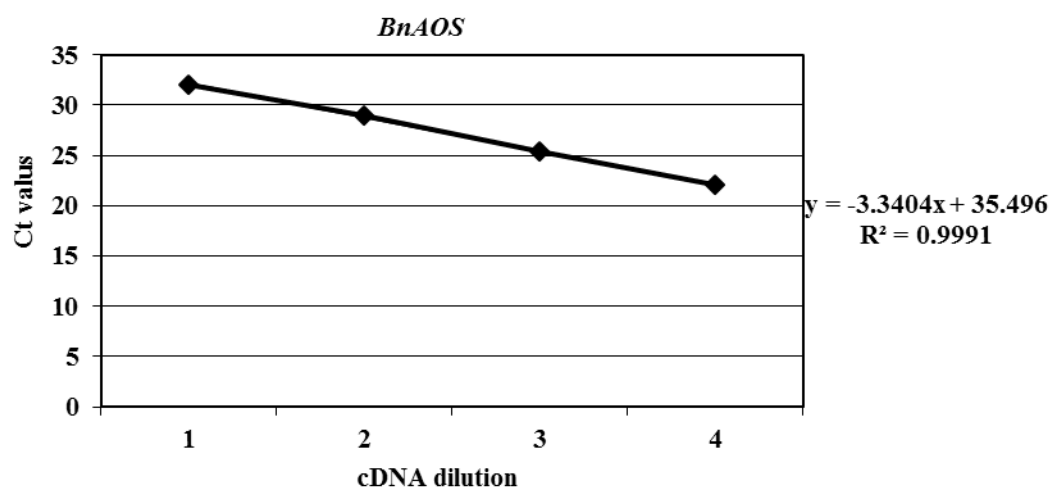
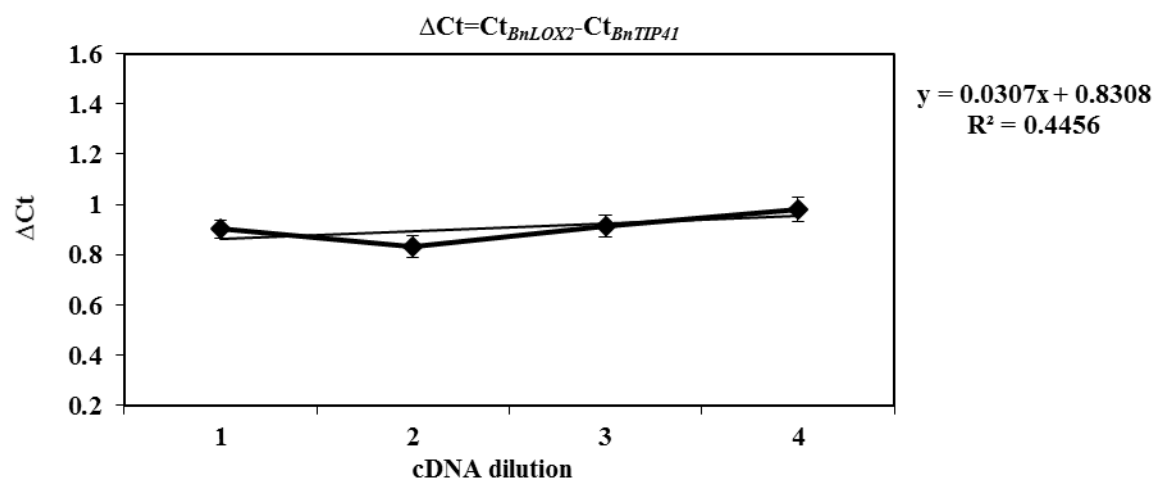


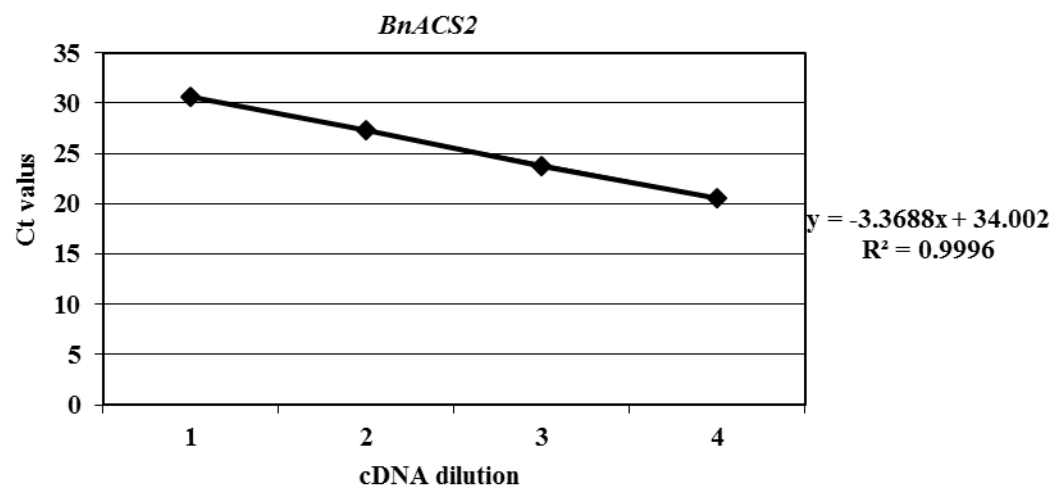
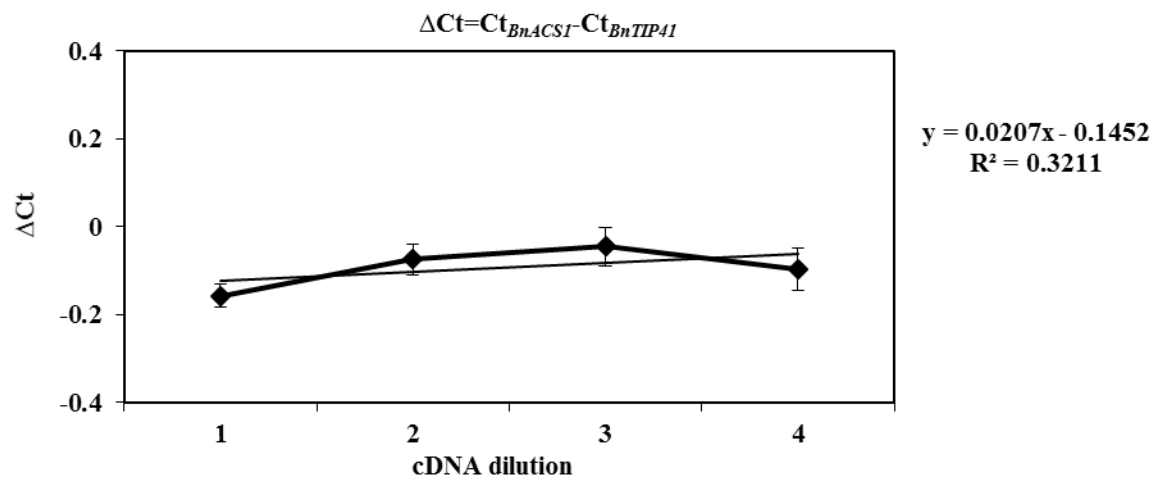
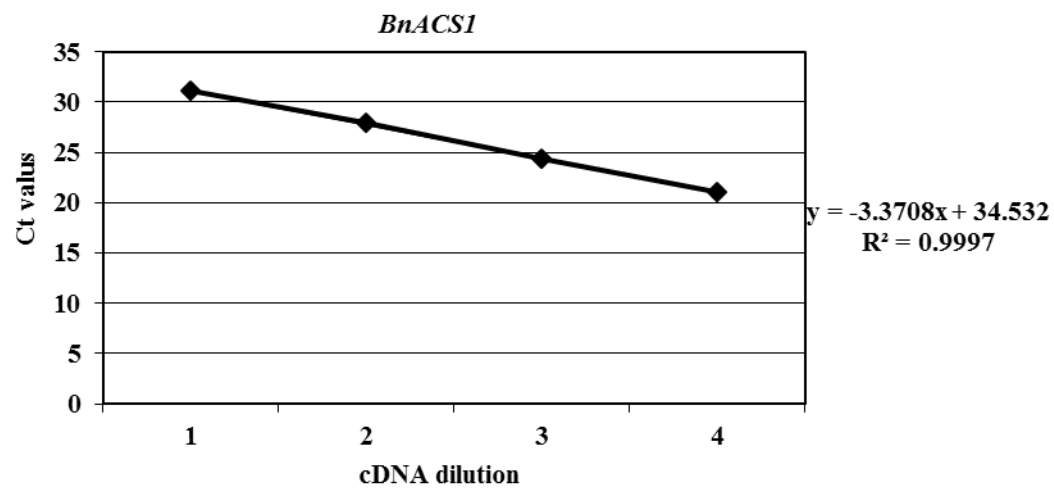


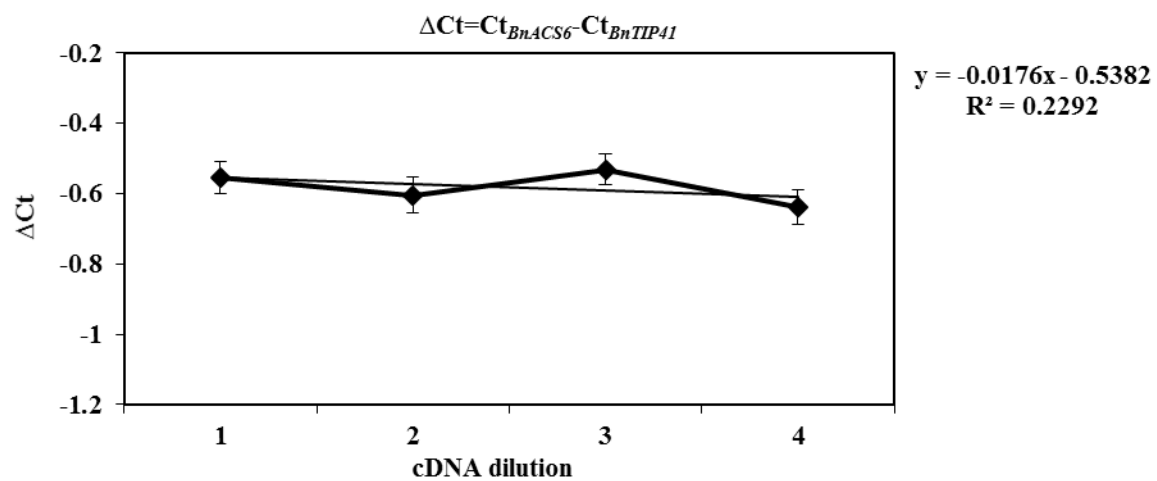
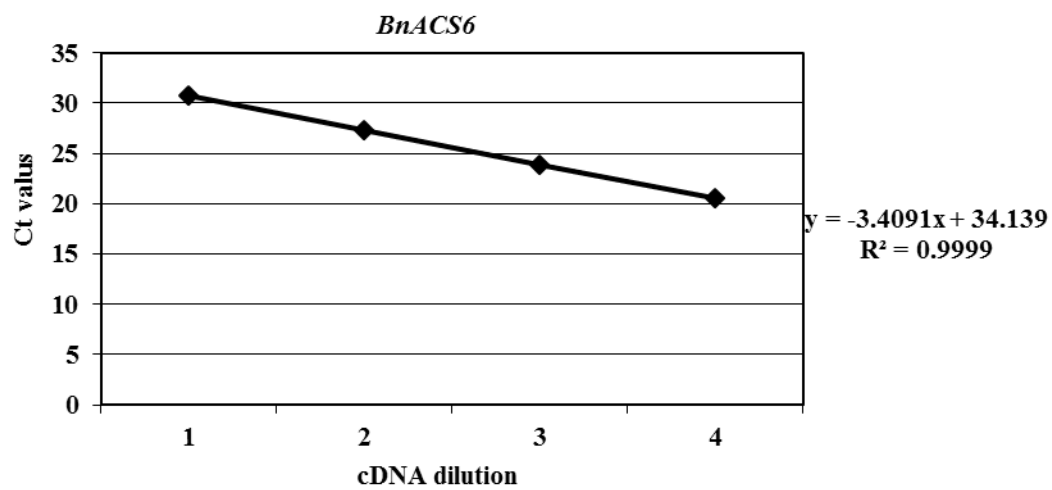
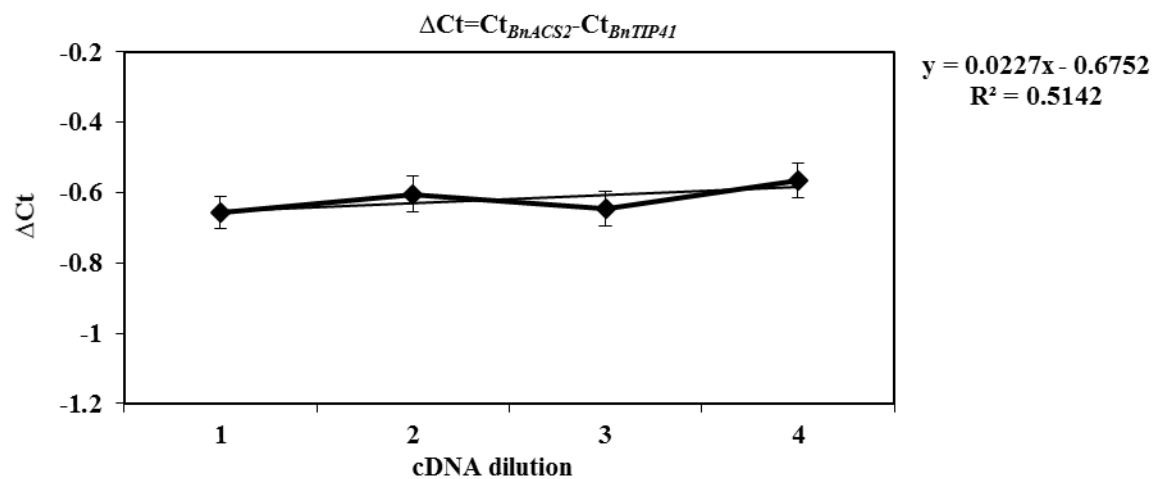


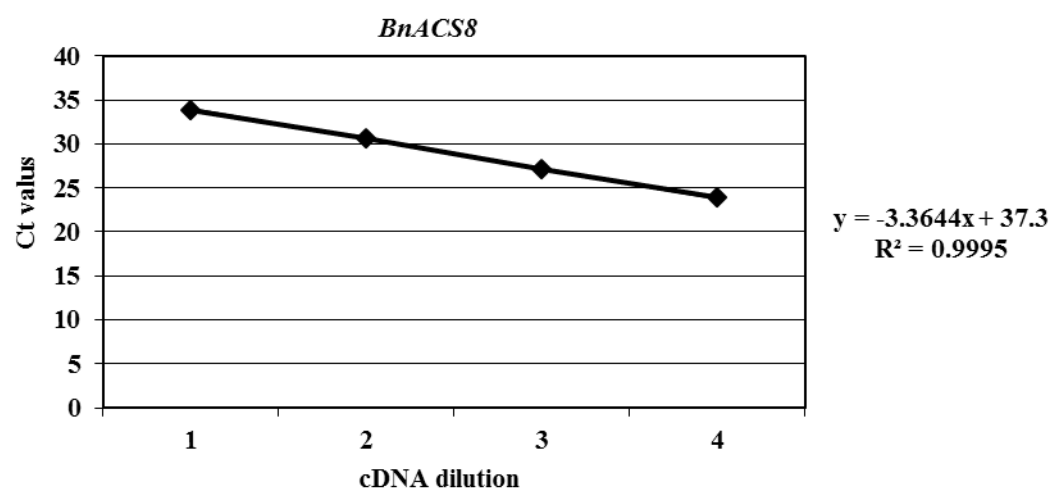
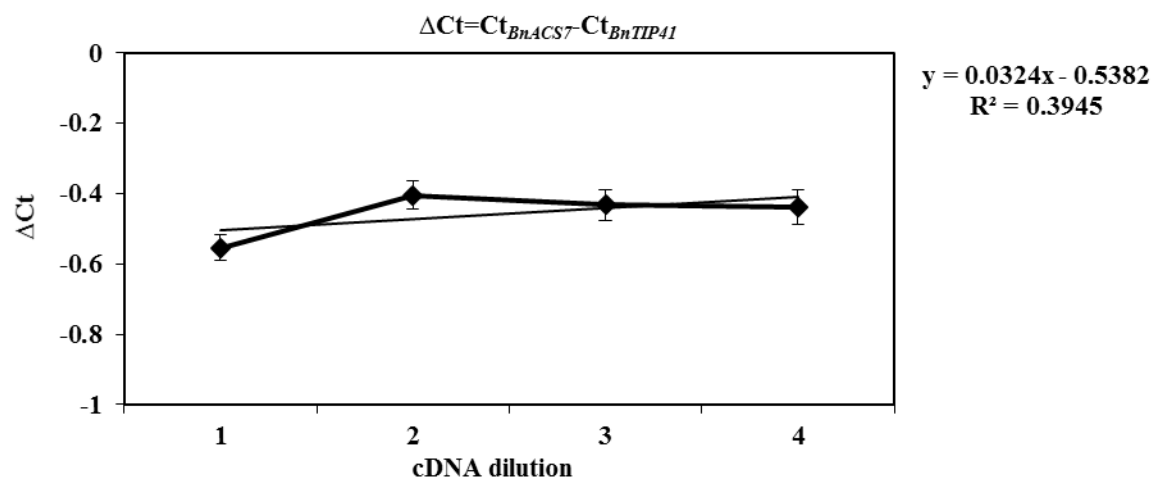
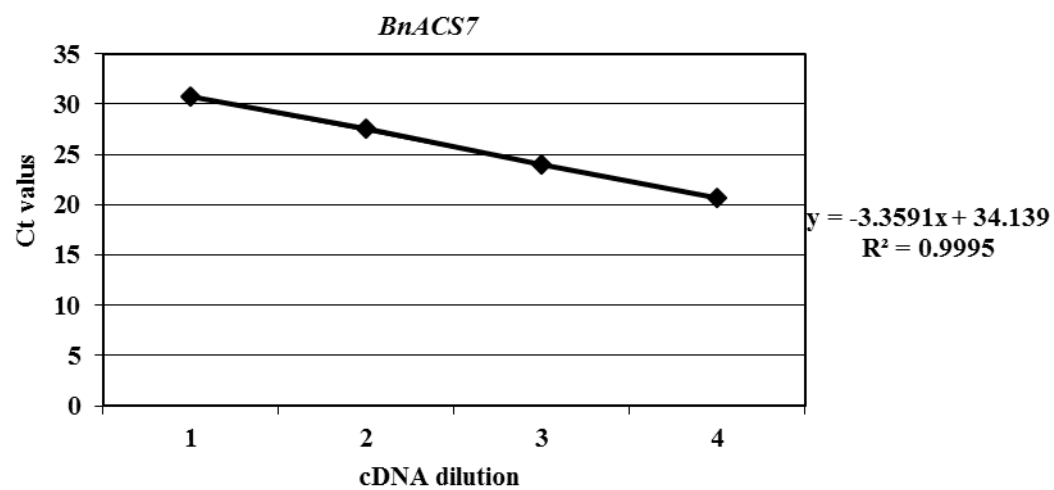


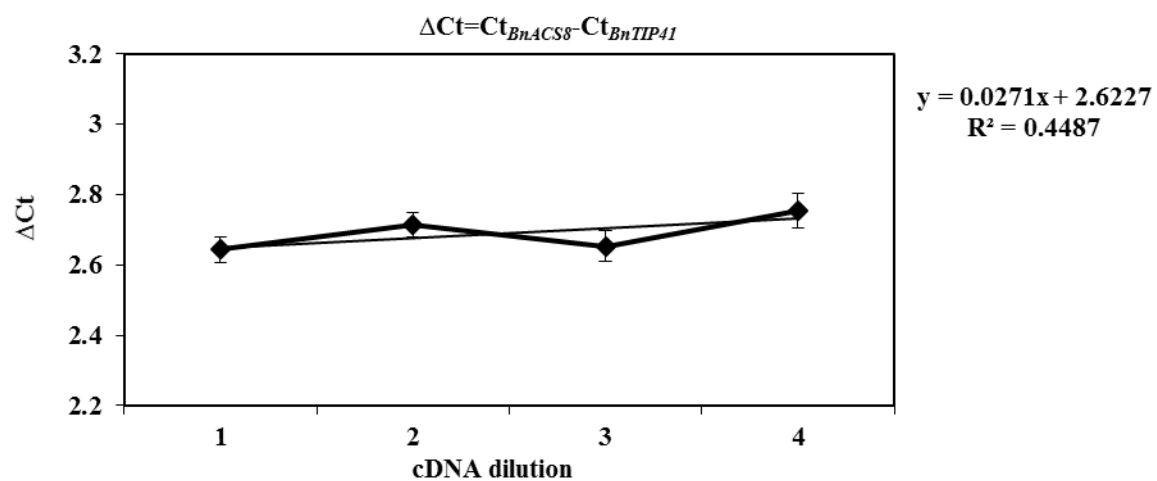












cDNA dilution (1= 0.001; 2= 0.01; 3= 0.1;4=1)