

All samples

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.645	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.855	0.778	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.788	0.934	0.870	-	-	-	-	-	-	-
<i>GAPDH</i>	0.609	0.925	0.841	0.945	-	-	-	-	-	-
<i>PLA2</i>	0.921	0.643	0.924	0.815	0.722	-	-	-	-	-
<i>RP II</i>	0.685	0.899	0.891	0.928	0.931	0.789	-	-	-	-
<i>TUA</i>	0.789	0.789	0.893	0.908	0.849	0.859	0.939	-	-	-
<i>TUB</i>	0.593	0.970	0.761	0.955	0.964	0.657	0.926	0.830	-	-
<i>UBQ10</i>	0.732	0.939	0.836	0.992	0.959	0.784	0.933	0.899	0.972	-
BestKeeper	0.851	0.906	0.939	0.981	0.926	0.893	0.950	0.943	0.912	0.967

Fruit developmental series

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.982	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.933	0.964	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.960	0.990	0.988	-	-	-	-	-	-	-
<i>GAPDH</i>	0.844	0.903	0.980	0.951	-	-	-	-	-	-
<i>PLA2</i>	0.915	0.896	0.939	0.927	0.894	-	-	-	-	-
<i>RP II</i>	0.926	0.962	0.993	0.991	0.977	0.946	-	-	-	-
<i>TUA</i>	0.916	0.937	0.993	0.972	0.977	0.965	0.989	-	-	-
<i>TUB</i>	0.947	0.979	0.985	0.997	0.955	0.938	0.996	0.975	-	-
<i>UBQ10</i>	0.942	0.980	0.992	0.998	0.968	0.924	0.996	0.978	0.997	-
BestKeeper	0.959	0.981	0.993	0.997	0.959	0.949	0.995	0.986	0.995	0.996

Different genotypes

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.988	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.965	0.979	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.965	0.975	1.000	-	-	-	-	-	-	-
<i>GAPDH</i>	0.957	0.981	0.997	0.995	-	-	-	-	-	-
<i>PLA2</i>	0.964	0.914	0.906	0.913	0.879	-	-	-	-	-
<i>RP II</i>	0.976	0.979	0.997	0.998	0.990	0.934	-	-	-	-
<i>TUA</i>	0.993	0.988	0.989	0.989	0.981	0.953	0.996	-	-	-
<i>TUB</i>	0.981	0.982	0.996	0.997	0.989	0.940	1.000	0.998	-	-
<i>UBQ10</i>	0.981	0.993	0.996	0.994	0.995	0.917	0.995	0.995	0.996	-
BestKeeper	0.987	0.990	0.991	0.999	0.989	0.939	0.997	0.994	0.998	0.995

Different storage time series

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.979	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.982	1.000	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.972	0.998	0.997	-	-	-	-	-	-	-
<i>GAPDH</i>	0.952	0.994	0.993	0.996	-	-	-	-	-	-
<i>PLA2</i>	0.972	0.934	0.937	0.937	0.901	-	-	-	-	-
<i>RP II</i>	0.968	0.994	0.993	0.999	0.992	0.945	-	-	-	-
<i>TUA</i>	0.991	0.995	0.996	0.994	0.981	0.966	0.993	-	-	-
<i>TUB</i>	0.964	0.991	0.990	0.998	0.991	0.944	1.000	0.991	-	-
<i>UBQ10</i>	0.964	0.997	0.996	1.000	0.998	0.927	0.998	0.990	0.997	-
BestKeeper	0.984	0.995	0.993	0.998	0.989	0.954	0.997	0.999	0.990	0.992

Different regulator treatments

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.986	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.973	0.998	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.982	1.000	0.999	-	-	-	-	-	-	-
<i>GAPDH</i>	0.968	0.996	0.999	0.998	-	-	-	-	-	-
<i>PLA2</i>	0.991	0.961	0.942	0.955	0.932	-	-	-	-	-
<i>RP II</i>	0.977	0.999	1.000	1.000	0.999	0.947	-	-	-	-
<i>TUA</i>	0.915	0.962	0.977	0.968	0.976	0.883	0.974	-	-	-
<i>TUB</i>	0.985	0.999	0.998	0.999	0.995	0.963	0.998	0.970	-	-
<i>UBQ10</i>	0.989	1.000	0.996	0.999	0.994	0.963	0.997	0.954	0.997	-
BestKeeper	0.991	0.996	0.995	0.998	0.992	0.970	0.999	0.959	0.998	0.997

Different tissues

	<i>18S rRNA</i>	<i>ACT</i>	<i>CYP2</i>	<i>TEF2</i>	<i>GAPDH</i>	<i>PLA2</i>	<i>RP II</i>	<i>TUA</i>	<i>TUB</i>	<i>UBQ10</i>
<i>ACT</i>	0.404	-	-	-	-	-	-	-	-	-
<i>CYP2</i>	0.449	0.863	-	-	-	-	-	-	-	-
<i>TEF2</i>	0.424	0.880	0.999	-	-	-	-	-	-	-
<i>GAPDH</i>	0.089	0.967	0.928	0.937	-	-	-	-	-	-
<i>PLA2</i>	0.789	0.575	0.903	0.891	0.680	-	-	-	-	-
<i>RP II</i>	0.433	0.857	0.998	0.996	0.936	0.893	-	-	-	-
<i>TUA</i>	0.715	0.575	0.905	0.889	0.728	0.963	0.914	-	-	-
<i>TUB</i>	0.156	0.974	0.951	0.961	0.992	0.729	0.951	0.744	-	-
<i>UBQ10</i>	0.405	0.887	0.999	1.000	0.945	0.881	0.997	0.885	0.966	-
BestKeeper	0.504	0.827	0.998	0.995	0.904	0.928	0.996	0.930	0.929	0.993