

Table S2. Summarized data of HPV16 E6 splice isoform-specific real-time RT-PCR.

Sample	Ct value				Log2(Ct value)				Relative copy number				Relative abundance			
	Unspliced	226^409	226^526	226^742	Unspliced	226^409	226^526	226^742	Unspliced	226^409	226^526	226^742	Unspliced	226^409	226^526	226^742
CaSki	27.544	24.052	27.951	33.368	14.741	16.744	13.460	9.493	27,376	109,724	11,267	721	18.40%	73.60%	7.60%	0.50%
SiHa	27.720	26.817	30.265	35.910	14.581	14.132	11.306	7.221	24,510	17,955	2,532	149	54.30%	39.80%	5.60%	0.30%
20861	24.954	20.954	25.143	31.365	17.075	19.671	16.073	11.284	138,054	834,296	68,941	2,493	13.20%	79.90%	6.60%	0.20%
20863	26.082	24.177	28.456	33.458	16.058	16.626	12.990	9.413	68,203	101,139	8,137	682	38.30%	56.80%	4.60%	0.40%
T1	38.533	35.341	36.656	38.445	4.835	6.081	5.360	4.955	29	68	41	31	17.00%	40.20%	24.40%	18.40%
T2	35.742	32.932	35.647	41.359	7.351	8.357	6.299	2.350	163	328	79	5	28.40%	57.00%	13.70%	0.90%
T3	26.131	25.421	28.384	34.839	16.014	15.451	13.056	8.178	66,155	44,792	8,520	290	55.20%	37.40%	7.10%	0.20%
T4	35.776	32.388	33.962	40.621	7.320	8.870	7.867	3.010	160	468	233	8	18.40%	53.80%	26.90%	0.90%
T5	31.456	28.604	29.474	38.750	11.215	12.444	12.043	4.682	2,376	5572	4220	26	19.50%	45.70%	34.60%	0.20%
T6	27.779	26.266	28.802	34.409	14.529	14.653	12.668	8.563	23,632	25,761	6,509	378	42.00%	45.80%	11.60%	0.70%
T7	25.479	24.999	27.176	33.446	16.602	15.850	14.180	9.424	99,435	59,033	18,568	687	55.90%	33.20%	10.40%	0.40%
T8	35.620	33.577	33.925	36.588	7.461	7.747	7.901	6.615	176	215	239	98	24.20%	29.50%	32.80%	13.50%
T9	32.204	28.905	31.420	36.399	10.540	12.160	10.232	6.783	1,489	4,577	1,203	110	20.20%	62.00%	16.30%	1.50%