

		P-value of association		
		2 clusters	3 clusters	4 clusters
Variable	Age	0.45	0.45	0.59
	Sex	1	0.82	0.8
	Stage (4 levels)	0.56	0.2	0.12
	Stage 3/4 vs. 1/2	0.56	0.16	0.21
	Stage 4 vs. 1/2/3	0.2	0.11	0.17
	HPV (sequencing)	<i>6.20E-08</i> *	<i>1.40E-07</i> *	<i>2.00E-15</i> *
	HPV (clinical)	<i>8.40E-07</i> *	<i>2.80E-06</i> *	<i>5.60E-12</i> *
	Site	<i>3.20E-08</i> *	<i>2.70E-07</i> *	<i>5.00E-09</i> *

Supplemental Table 4. PDX cluster associations with patient demographic and tumor variables.

Correlation of demographic and clinical characteristics of PDXs with cluster membership reveals that HPV status and tumor site are the only variables significantly correlated with hierarchical gene expression-based clustering (when partitioned by either 2, 3, or 4 clusters). Age, sex, and stage were not significantly associated with expression-based clustering in any analyses conducted.