

**Table S7. Bisulfite sequencing of the promoter DNA
hypermethylation detection in primary tissues from HNSCC
and non-cancerous patients of the validation cohort**

	Percentage of patients with hemi or fully methylated promoters, detected in primary tumor tissues	Percentage of patients with hemi or fully methylated promoters, detected in normal tissues	Fisher's exact test
	%	%	p-value
ZNF14	43.8	0	<0.0001
ZNF141	9.4	78.6	<0.0001
ZNF160	46.9	14.3	0.0011
ZNF211	0.0	0	ND
ZNF420	25.0	0	0.0018
ZNF585B	34.4	0	<0.0001
ZNF71	37.5	0	<0.0001