Table S4 : The level of expression of the 49 commonly quantified proteins in HNSCC patients HPV-16 and HPV negative and t/nt stable.

			T/NT HPV+		T/NT HPV-	
Protein name	**Gene name**	**Protein description**	Ave	SE	Ave	SE
1433B_HUMAN	YWHAB	14-3-3 protein beta/alpha	0.95	0.18	1.40	0.73
1433E_HUMAN	YWHAE	14-3-3 protein epsilon	1.18	0.05	0.86	0.75
ADHX_HUMAN	ADH5	Alcohol dehydrogenase class-3	1.03	0.02	0.94	0.07
ADT3_HUMAN	SLC25A6	ADP/ATP translocase 3	1.16	0.02	1.37	0.18
ALDOA_HUMAN	ALDOA	Fructose-bisphosphate aldolase A	1.10	0.23	0.90	0.43
ANXA2_HUMAN	ANXA2	Annexin A2	0.81	0.05	1.05	0.66
APOA1_HUMAN	APOA1	Apolipoprotein A-I	0.77	0.10	1.22	0.39
ARP2_HUMAN	ACTR2	Actin-related protein 2	1.04	0.13	1.03	0.59
CAPZB_HUMAN	CAPZB	F-actin-capping protein subunit beta	0.96	0.12	1.03	0.41
CBR1_HUMAN	CBR1	Carbonyl reductase [NADPH] 1	0.96	0.10	1.01	0.05
CPNS1_HUMAN	CAPNS1	Calpain small subunit 1	0.79	0.04	0.90	0.17
DDX39_HUMAN	GLUD1	Glutamate dehydrogenase 1, mitochondrial	1.28	0.11	1.04	0.12
EF1A1_HUMAN	EEF1A1	Elongation factor 1-alpha 1	1.10	0.02	1.06	0.42
EF1G_HUMAN	EEF1G	Elongation factor 1-gamma	1.06	0.12	1.05	0.17
EF2_HUMAN	EEF2	Elongation factor 2	0.84	0.13	0.92	0.23
ENOA_HUMAN	ENO1	Alpha-enolase	0.94	0.16	0.96	0.07
ENPL_HUMAN	HSP90B1	Endoplasmin	1.10	0.07	0.88	0.10
ETFB_HUMAN	ETFB	Electron transfer flavoprotein subunit beta	1.11	0.19	0.98	0.05
G3P_HUMAN	GAPDH	Glyceraldehyde-3-phosphate dehydrogenase	0.87	0.02	1.09	0.13
G6PI_HUMAN	GPI	Glucose-6-phosphate isomerase	0.99	0.08	0.89	0.20
GANAB_HUMAN	GANAB	Neutral alpha-glucosidase AB	0.94	0.18	0.99	0.34
GDIB_HUMAN	GDI2	Rab GDP dissociation inhibitor beta	0.89	0.15	0.81	0.06
GRP78_HUMAN	HSPA5	78 kDa glucose-regulated protein	1.26	0.21	0.84	0.11
HNRPU_HUMAN	HNRNPU	Heterogeneous nuclear ribonucleoprotein U	0.95	0.07	0.88	0.14
HSP7C_HUMAN	HSPA8	Heat shock cognate 71 kDa protein	1.10	0.21	1.63	0.47
IF4A1_HUMAN	EIF4A1	Eukaryotic initiation factor 4A-I	0.94	0.09	1.20	0.42
PDIA1_HUMAN	P4HB	Protein disulfide-isomerase	0.93	0.12	0.95	0.45
PDIA4_HUMAN	PDIA4	Protein disulfide-isomerase A4	1.27	0.02	0.98	0.38
PGAM1_HUMAN	PGAM1	Phosphoglycerate mutase 1	1.11	0.27	1.00	0.48

PRDX1_HUMAN	PRDX1	Peroxiredoxin-1	1.10	0.06	0.81	0.49
PRS10_HUMAN	PSMC6	26S protease regulatory subunit 10B	0.87	0.05	0.97	0.49
RL28_HUMAN	RPL28	60S ribosomal protein L28	0.88	0.09	0.85	0.27
RL35A_HUMAN	RPL35A	60S ribosomal protein L35a	0.93	0.17	0.78	0.01
RL5_HUMAN	RPL5	60S ribosomal protein L5	0.92	0.17	0.83	0.17
ROA1_HUMAN	HNRNPA1	Heterogeneous nuclear ribonucleoprotein A1	1.12	0.09	1.10	0.60
ROA2_HUMAN	HNRNPA2B1	Heterogeneous nuclear ribonucleoproteins A2/B1	0.95	0.24	1.20	0.44
RS12_HUMAN	RPS12	40S ribosomal protein S12	0.98	0.04	0.93	0.23
RS15A_HUMAN	RPS15A	40S ribosomal protein S15a	0.91	0.17	0.99	0.35
RS20_HUMAN	RPS20	40S ribosomal protein S20	0.98	0.07	0.96	0.22
RS27A_HUMAN	RPS27A	Ubiquitin-40S ribosomal protein S27a	1.11	0.07	1.08	0.47
RS3_HUMAN	RPS3	40S ribosomal protein S3	1.05	0.10	0.93	0.21
RS3A_HUMAN	RPS3A	40S ribosomal protein S3a	1.07	0.04	0.99	0.21
RU2A_HUMAN	SNRPA1	U2 small nuclear ribonucleoprotein A'	1.05	0.08	1.29	0.22
SET_HUMAN	SET	Protein SET	0.98	0.08	1.44	0.18
TBCB_HUMAN	TBCB	Tubulin-folding cofactor B	1.07	0.00	0.96	0.32
TM109_HUMAN	TMEM109	Transmembrane protein 109	0.90	0.05	0.90	0.17
TPM4_HUMAN	TPM4	Tropomyosin alpha-4 chain	1.26	0.21	0.87	0.41
TRFE_HUMAN	TF	Serotransferrin	0.86	0.01	1.05	0.73
VINC_HUMAN	VCL	Vinculin	1.10	0.23	0.72	0.10