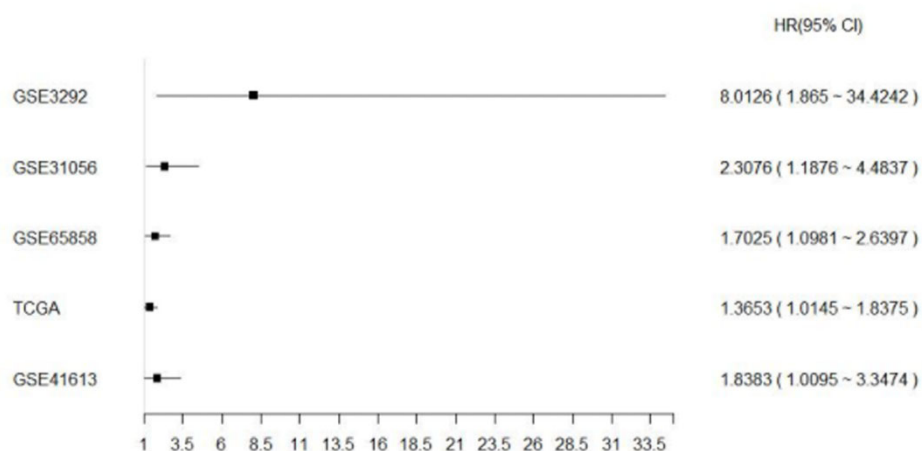


# OShnscc

OShnscc ( Online consensus Survival for Head and Neck Squamous Cell Carcinoma) integrates 1366 unique HNSCC cases with available gene expression profiles and the corresponding clinical information from 9 independent cohorts, and provides 6 types of prognosis endpoints for survival analysis.

Gene symbol:  [Survival analysis](#)  
Data source:  Survival:

Forest plot reporting the correlation of IL11 with OS



Univariate Cox regression analysis of IL11 expression with OS in each dataset of OShnscc					
Dataset	P value	HR	95%CI	Prognostic	
GSE3292	0.0051	8.0126	1.8650 ~ 34.4242	Poor	<a href="#">Go</a>
GSE31056	0.0136	2.3075	1.1876 ~ 4.4837	Poor	<a href="#">Go</a>
GSE65858	0.0174	1.7025	1.0981 ~ 2.6397	Poor	<a href="#">Go</a>
TCGA	0.0399	1.3653	1.0145 ~ 1.8375	Poor	<a href="#">Go</a>
GSE41613	0.0465	1.8383	1.0095 ~ 3.3474	Poor	<a href="#">Go</a>

Note: cutoff: upper 25% vs other 75%

[Click here](#) to see the definitions of Survival terms including overall survival (OS), disease specific survival (DSS), disease free survival (DFS), progression free survival (PFS), disease free interval (DFI), and progression free interval (PFI).

Fig. S1 Summary of prognostic analysis of IL11 expression in OShnscc using default parameters.