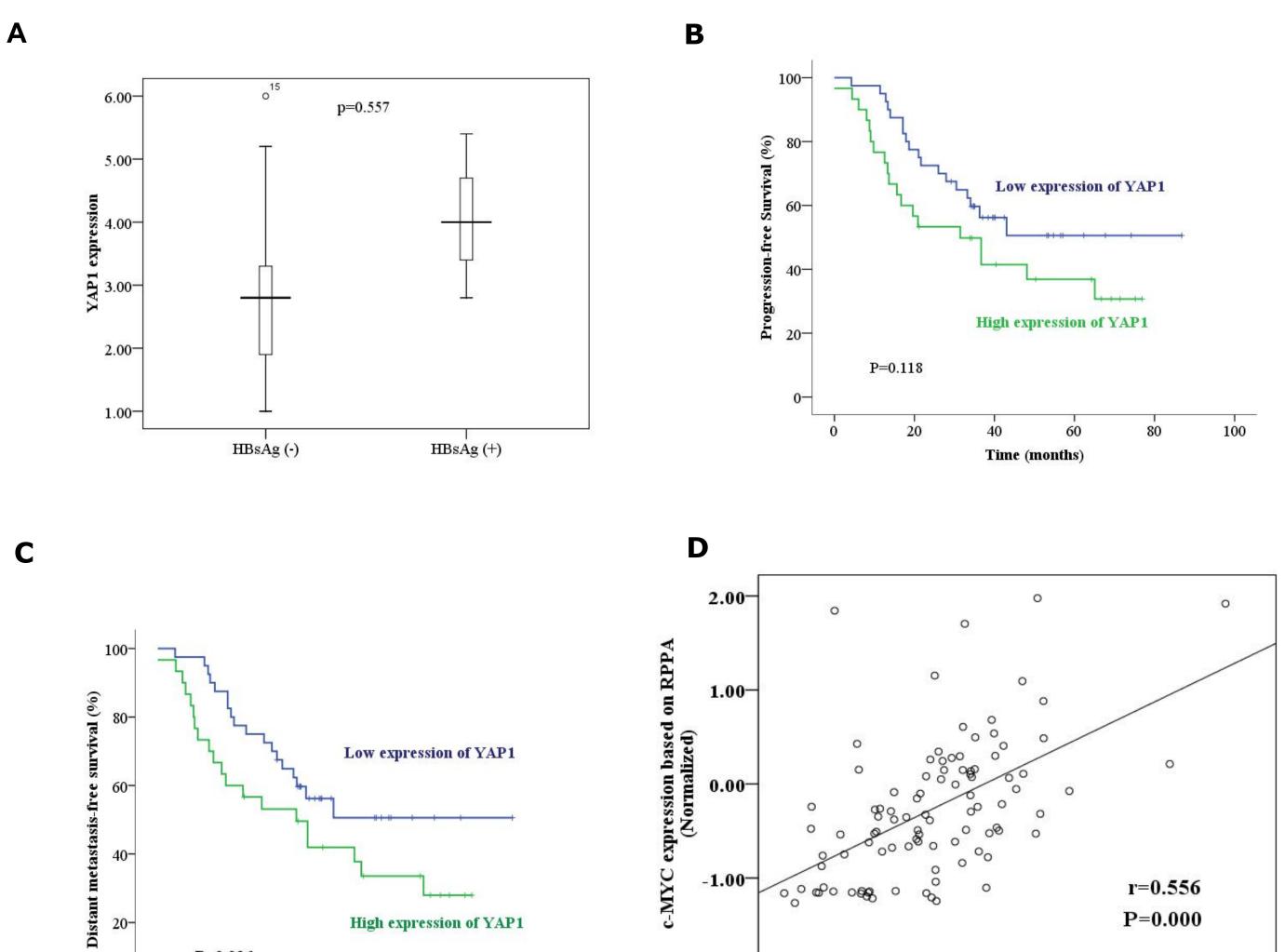
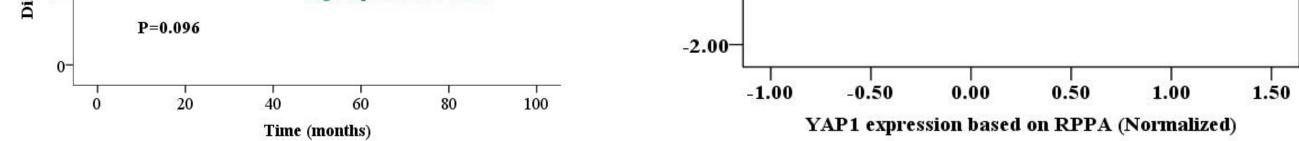
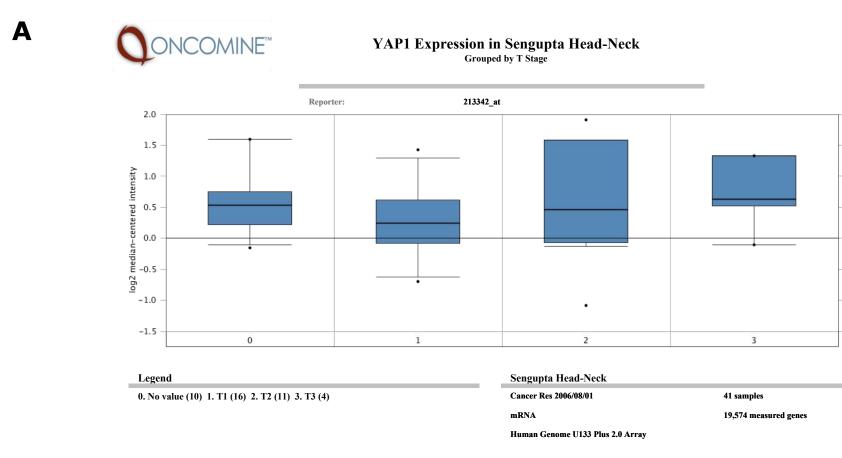
Figure legends

Supplemantary Figure 1. A, YAP1 expressions in HBsAg (+) NPC (n=5) compared to HBsAg (-) NPC (n=65) by IHC analysis (p=0.557).B and C,5-year progression-free survival and distant metastasis-free survival in high YAP1 expression compare to low YAP1 expression groups show no statistic significant (36.9% vs.50.6%, p=0.118,33.6% vs.50.6%, p=0.096, respectively. D, YAP1 expression has positive correlation with c-MYC expression (r=0.556, p<0.001).

Supplemantary Figure 2.The ONCOMINE database (www.oncomine.com/nasophryngeal cancer) was used to analyze yap gene expression with clinical TNM staging. A, yap gene expression with clinical T staging; B, yap gene expression with clinical N staging.



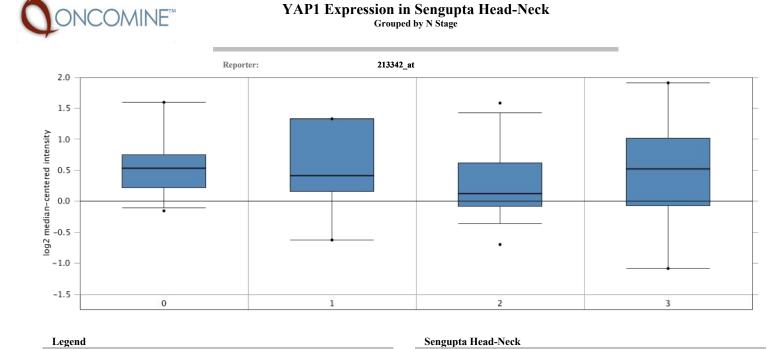




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Oncomine Source:

https://www.oncomine.com/resource/main.html#ac:6301;cso:sizeSmallLarge;cv:detail;d:156636523;dso:geneOverex;dt:dataset;ec:[1];et:over;f:1908396;g:10413;gs:122192584;gt:boxplot;p:200007697; pg:1;pvf:39585[YAP2],800054473[Carcinoma of Nasopharynx];scr:datasets;ss:all;v:18



В

mRNA

19,574 measured genes

41 samples

Human Genome U133 Plus 2.0 Array

Cancer Res 2006/08/01

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Oncomine Source:

0. No value (10) 1. N0 (7) 2. N1 (16) 3. N2 (8)

https://www.oncomine.com/resource/main.html#ac:6301;cso:sizeSmallLarge;cv:detail;d:156636523;dso:geneOverex;dt:dataset;ec:[1];et:over;f:1908396;g:10413;gs:122192584;gt:boxplot;p:200007696; pg:1;pvf:39585[YAP2],800054473[Carcinoma of Nasopharynx];scr:dataset;sss:all;v:18