Supplementary Information

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Enhancing the Productivity and Proliferation of CHO-K1 Cells by Oncoprotein YAP (Yes-Associated Protein)

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1 MEPAQQPPPQPAPQGPAPP------SVSPAGTPAAPPAPPAGHQVVHVRGD 45
mYAP
            1 MEPAQQPPPQPAPQGPVPPSA------PAPPSGTPAAPPAPPAGHQVVHVRGD 47
chYAP
            1 MDPGQQPPPQPAPQGQGQPPSQPPQGQGPPSGPGQPAPAATQAAPQAPPAGHQIV<mark>HVRGD</mark> 60
hYAP
                                            .*:.* *** *****
              *:*.******
mYAP
           46 SETDLEALFNAVMNPKTANVPQTVPMRLRKLPDSFFKPPEPKSHSRQASTDAGTAGALTP 105
           48 SETDLEALFNAVMNPKTANVPQTVPMRLRKLPDSFFKPPEPKSHSRQASTDAGTAGALTP 107
chYAP
           61 SETDLEALFNAVMNPKTANVPQTVPMRLRKLPDSFFKPPEPKSHSRQASTDAGTAGALTP 120
hYAP
               mYAP
          106 QHVRAHSSPASLQLGAVSPGTLTASGVVSGPAAAPAAQHLRQSSFEIPDDVPLPAGWEMA 165
           108 QHVRAHSSPASLQLGAVSPGTLTPT-VVSGPAATPAAQHLRQSSFEIPDDVPLPAGWEMA 166
chYAP
          121 QHVRAHSSPASLQLGAVSPGTLTPTGVVSGPAATPTAQHLRQSSFEIPDDVPLPAGWEMA 180
hYAP
              *******************
mYAP
          166 KTSSGQRYFLNHNDQTTTWQDPRKAMLSQLNVPAPASPAVPQTLMNSASGPLPDGWEQAM 225
chYAP
           167 KTSSGQRYFLNHIDQTTTWQDPRKAMLSQLNVPTPASTAVPQTLMNSASGPLPDGWEQAM 226
          181 KTSSGQRYFLNHIDQTTTWQDPRKAMLSQMNVTAPTSPPVQQNMMNSASGPLPDGWEQAM 240
hYAP
              ******** ***************************
          226 TQDGEVYYINHKNKTTSWLDPRLDPRFAMNQRITQSAPVKQPPPLAPQSPQGGVLGGGSS 285
mYAP
chYAP
           227 TQDGDVYYINHKNKTTSWLDPRLDPRFAMNQRITQSAPVKQPPPLAPQSPQGGVLGGGNS 286
           241 TQDGEIYYINHKNKTTSWLDPRLDPRFAMNQRISQSAPVKQPPPLAPQSPQGGVMGGSNS 300
hYAP
              286 NQQQQIQLQQLQMEKERLRLKQQELFRQELALRSQLPTLEQDGGTPNAVSSPGMSQELRT 345
mYAP
chYAP
           287 SQQQQMQLQQLQMEKERLRLKQQELLRQELALRSQLPTLEQDGGTQNAVSSPGMSQELRT 346
hYAP
           301 NQQQMRLQQLQMEKERLRLKQQELLRQELALRSQLPTLEQDGGTQNPVSSPGMSQELRT 360
              346 MTTNSSDPFLNSGTY<mark>HSRDES</mark>TDSGLSMSSYSIPRTPDDFLNSVDEMDTGDTISQSTLPS 405
mYAP
           347 MTTNSSDPFLNSGTY<mark>HSRDES</mark>TDSGLSMSSYSVPRTPDDFLNSVDEMDTGDTINQSTLPS 406
chYAP
           361 MTTNSSDPFLNSGTY<mark>HSRDES</mark>TDSGLSMSSYSVPRTPDDFLNSVDEMDTGDTINQSTLPS 420
hYAP
              mYAP
          406 QQSRFPDYLEALPGTNVDLGTLEGDAMNIEGEELMPSLQEALSSEIL-DVESVLAATKLD 464
chYAP
           407 QQSRFPDYLEAIPGTNVDLGTLEGDAMNIEGEELMPSLQEALSSDILNDMESVLAATKLD 466
hYAP
          421 QQNRFPDYLEAIPGTNVDLGTLEGDGMNIEGEELMPSLQEALSSDILNDMESVLAATKLD 480
              mYAP
          465 KESFLTWL 472
          467 KESFLTWL 474
chYAP
          481 KESFLTWL 488
hYAP
             *****
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Fig. S1 Amino acid sequence alignment of YAP from various organisms, including human, mouse, and Chinese hamster, for the identification of the consensus phosphorylation sequence for LATS. The LATS consensus phosphorylation sites are highlighted.

ATGGAGCCCGCGCAGCAGCCGCCCCCAGCCGGCCCGCAAGGCCCCGTGCCGCCGCTCCGCGCCCGCGC C<mark>GCG</mark>GAGACCGACCTGGAGGCGCTCTTCAACGCCGTCATGAACCCCAAGACGGCCAACGTGCCGCAGACC GTGCCCATGCGGCTGCGCAAGCTGCCCGACTCCTTCTTCAAGCCGCCCGAGCCCAAGTCCCACTCGCGAC AGGCT<mark>GCT</mark>ACTGATGCAGGCACTGCAGGAGCTCTGACTCCACAGCACGTTCGGGCTCAC<mark>GCCGCT</mark>CCAGC AGCCCTGCAGCTGGGAGCCGTTTCTCCTGGGACTCTCACCCCTACTGTGGTGTCTTGGTCCTGCAGCCACC CCTGCAGCACAGCATCTCCGACAG<mark>GCTGCC</mark>TTTGAGATCCCTGATGATGTACCCCTGCCTGCAGGCTGGG AGATGGCCAAGACATCGTCTGGTCAGAGATACTTCTTAAATCACATCGATCAGACAACAACATGGCAGGA CCCCAGGAAGGCCATGCTCTCCCAACTGAATGTCCCTACGCCTGCCAGCACAGCAGTGCCTCAGACCCTG ATGAACTCCGCCTCAGGACCTCTTCCTGATGGATGGAGCCATGACGCAGGATGGAGATGTTACT ACATAAACCATAAGAACAAGACTACGTCCTGGCTGGACCCAAGGCTCGACCCTCGTTTTGCTATGAACCA GAGAATCACTCAGAGCGCTCCAGTGAAGCAGCCGCCACCCTTGGCTCCCCAGAGCCCACAGGGAGGTGTC CTGGGTGGTGGCAACTCCAGCCAGCAGCAGCAGATGCAACTGCAGCAGCTCCAGATGGAGAAGGAGAGAC TGCGGTTGAAACAGCAAGAATTACTGCGGCAGGAATTAGCTCTCCGAAGCCAGTTGCCCACACTGGAGCA GGACGGAGGAACTCAGAATGCTGTGTCTTCCCCCGGGATGTCTCAGGAATTGAGGACAATGACAACCAAT AGTTCAGATCCCTTTCTTAACAGTGGCACCTATCACTCTAGAGATGAGGCC ACGGACAGTGGCCTCAGCATGAGTAGCTACAGTGTCCCTCGGACCCCGGATGACTTCCTGAACAGTGTTGATGAAATGGATACAGGTGA CACTATCAACCAAAGCACCCTGCCTTCACAGCAGAGCCGTTTCCCAGATTACCTTGAAGCCATCCCTGGG ACAAATGTGGACCTTGGAACACTGGAAGGAGATGCGATGAACATAGAAGGAGGAGCTGATGCCCAGCC TGCAGGAAGCACTGAGTTCCGACATCCTTAATGACATGGAGTCTGTGTTTGGCTGCCACCAAGCTAGATAA AGAAAGCTTTCTCACATGGTTATAG

Fig. S2 Nucleotide sequence encoding the YAP-5SA which is inserted within the PB513B-1 plasmid. Letters in red indicate bases changed to create the codon for alanine, resulting in the substitution of the serine codon with alanine, shown in highlighted color.