

# The sequence of TGF- $\beta$ 2 from *Xenopus laevis*

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TGF- $\beta$ 2 is one of the five known isoforms of transforming growth factor- $\beta$  (1). While TGF- $\beta$ 2 is equipotent to TGF- $\beta$ 1 in several assays involving different mammalian cell lines (2), it is differentially active in inducing mesoderm differentiation in ectodermal explants from *Xenopus laevis* embryos (3). We have shown that the *Xenopus* cell line XTC produces TGF- $\beta$ 2 (2). Here we report the sequence of *X. laevis* TGF- $\beta$ 2 based on cDNA clones isolated from an XTC cell library. The sequence is highly

homologous to that of mammalian TGF- $\beta$ 2, with 95% identity in the region of mature TGF- $\beta$ 2, and 82% in the precursor region.

## REFERENCES

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CCACATAGGTCTATTACATTTTTTGATATACTTTCAATTGGACTTGTGGATCATTAGCAAAAGAAGAAAATGCACTACTATGTCCTGTTAACCTTC	105
CTGACCCCTGGACCTGGCCCCCTGGCCCTCAGTCTGTCACCTGCAGTGCCTCGACATGGATCAATTCTATGCGCAAGAGGATCGAGGCCATTAGGGGGCAGATC	210
L T L D L A P V A L S L S T C S A L D N D O F M R K R I E A I R G Q I	
CTGAGCAAACCTGAATAACAGCCCCCTGAGGATTACCCGGAGACCGGGAGGGAGTTCCAGGATGTCATTCTCATCACAGCAGAGACCTCTTCAG	315
L S K L K L N S P D E Y P E P G E V S O D V I S I Y N S T R D L L Q	
GAGAAGGCCAACGAGAGAGCTACCTCTGTGAGAGGGAGAGGAGTGAAGATGAGTATTATGCCAAAGAGGTTTACAAAATGACATGCTACCCATTACACCTCT	420
E R A N E R A T S C E R E R E S D E Y Y K A E V Y R I D M L P Y Y T S	
GAAAATGTCATCCACCAAGTTATACACCCCATACTTCCGAATCTGATCTGAGTTGATCTCTTCATGAGAAAATCTCCAACTCTCTGAAGGCCAGAATT	525
E N V I P P S Y T T P Y F R I V R F D V S S M E K N A S N L V K A E F	
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R V F R L M N T K A R V S E Q R I E L Y Q I P D V S A L S P T Q R Y	
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I D S K V V K T R A E G E W L S F D V T E A V N E W L H H K D R N L G	
TTTAAGATAAGTTGCACTGTCCTGCTGCACCTTATTCCCTCTAAATACTACATCATCCAAAACAAAAGTGAAGAATTGAAACCAGTTGCAAGGTATTGAT	840
F K I S L H P C P C T F I P S N N Y I P N K S E E L E T R F A G I D	
GATGCTCTACATCTGATGTCGTTGGGATTCCAAGAGCAAGACTGGCGGAAGAACGACACGGCAGAACCTCCCACCTCTGCTCATGTTTGCCCTCATATAGG	945
D A Y M Y A G G D S K S K T G R K K H T G R T P H L L L M L L P S Y R	
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L E S Q O S R R K K R A L D A Y C F R N V Q D N C C L R P L Y I D	
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N K P K I E O L S N M I V K S C K C S *	
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AAATGTCCTG	2730
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CACTAGCAAAATAGTCGTTCTG	3150
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CCAATTTCGCTTGTGTTAGTAATGTCGCTG	3507

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