







## **cIAP1 promoter sequence**

4751 TTACAAAGCCT ATCAGACCTA CATCTGGAAC AGGTCATATA TACTTTGGTG entry score  
 <----- M00076 GATA-2 90.5  
 <----- M00075 GATA-1 89.0  
 -----> M00101 CdxA 87.1  
 <----- M00101 CdxA 86.4  
 <----- M00128 GATA-1 86.3  
 ----- M00050 E2F 86.2 — E2F binding site  
 <----- M00075 GATA-1 85.3  
 <----- M00077 GATA-3 85.0

```

4801 CACAAAGGTA CTTATCAATG TCTGGTTCCCT AACCTCCAAT CCTAACAAAC entry      score
          <----->
          <----->
          ----->
          <----->
          ----->>
          >
          M00203 GATA-X 94.8
          M00101 CdxA   89.3
          M00148 SRY    89.1
          M00100 CdxA   87.2
          M00148 SRY    86.4
          M00050 E2F    86.2 — E2F binding site

```

## **cIAP2 promoter sequence**

5451 GTGTTGGCG GTGTCTCGTG CTATCCAGGC TGGCCTCAAG TTCCGAATGG entry score  
 -----  
 >  
 ----->  
 <-----  
 <-----  
 ----->  
 M00148 SRY 92.7  
 M00100 CdxA 91.0  
 M00050 E2F 88.5  
 M00076 GATA-2 88.1  
 M00123 c-Myc/ 86.8  
 M00240 Nkx-2. 86.0

— E2F binding site

## **Yap1 promoter sequence**

3401 TGAGTTCAG GACAGCCAGG GCTACACAGA GAAACTGTGG CTCGAAAAAC entry score  
 <- M00008 Sp1 86.3  
 <----- M00050 E2F 86.2 — E2F binding site

## **Yap1 intron sequence**



