

In the article "Structure of the human B lymphocyte receptor for C3d and the Epstein-Barr virus and relatedness to other membranes of the family of C3/C4 binding proteins" by Janis J. Weis, Lorraine E. Toothaker, John A. Smith, John H. Weis, and Douglas T. Fearon (March 1988, 167:1047), the printer omitted a line from the nucleotide sequence in Fig. 1 *B*. The corrected figure and legend appear below.

B		+1
-55	CCCGACGCTCGCGGGTCTCGGAACGCATCCCGCCGCGGGGGCTTCGGCCGTGGCATGGG	5
6	CGCCGCGGGGCTGCTCGGGGTTTCTTGGCTCTCGTCGCACCGGGGGTCTCGGGATTTC	65
66	TTGTGGCTCTCTCCGCTATCCTAAATGGCCGATTAGTTATTATTCTACCCCATTCG	125
126	TGTTGGTACCGTATAAGGTACAGTTGTTCAAGTACCTTCGCGCTCATTGGAGAAAAAG	185
186	TCTATTATGCATAACTAAAGACAAAGTGGATGGAACTGGGATAAACCTGCTCCTAAATG	245
246	TGAATATTTCATAAATATTCTTCTTGCCCTGAGCCCATAGTACCAGGAGGATACAAAAT	305
306	TAGAGGCTCTACACCTACAGACATGGTGATTCTGTGACATTTGCCTGTAAACCAACTT	365
366	CTCCATGAACGGAACAAGTCTGTTTGGTGTCAAGCAAATAATATGTGGGGGCCGACAG	425
426	ACTACCAACCTGTGTAAGTGTTTCCCTCTCGAGTGTCCAGCACTTCTATGATCCACAA	485
486	TGGACATCACACAAGTGAGAATGTTGGCTCCATTGTCTCCAGGATTGTCTGTGACTTACAG	545
546	CTGTGAATCTGGTTACTTGTCTTGGAGAAAAGATCATTAACTGTTTGTCTTCGGGAAA	605
606	ATGGAGTGTCTCCCCCCACATGTGAAGAGGCACGCTGTAAATCTCTAGGACGATTTCC	665
666	CAATGGGAAGGTAAAGGAGCCTCCAATTCTCCGGTGTGGTGAAGTCAAACTTTTCTG	725
726	TGATGAAGGTATCGACTGCAAGGCCACCTTCTAGTCGGTGTGAATTGCTGGACAGGG	785
786	AGTTGCTTGGACCAAAATGCCAGTATGTGAAGAAATTTTTTGCCCATCACCTCCCTTAT	845
846	TCTCAATGGAAGACATATAGGCAACTCACTAGCAAAATGTCTCATATGGAAGCATAGTCAC	905
906	TTACACTTGTGACCCGACCCAGAGGAAGGAGTGAACCTCATCCTTATTGGAGAGAGCAC	965
966	TCTCGTTGTACAGTTGATAGTCAGAAGACTGGGACCTGGAGTGGCCCTGCCCCACGCTG	1025
1026	TGAACCTTCTACTTCTCGGGTTCAGTGTCCACATCCCAGATCCTAAGAGGCCGAATGGT	1085
1086	ATCTGGGCAGAAAGATCGATATACCTATAACGACACTGTGATATTGTCTGCATGTTTGG	1145
1146	CTTCACCTTGAAGGCGAGCAAGCAAAATCCGATGCAATGCCCAAGGCACATGGGAGCCATC	1205
1206	TGACACAGTCTGTGAAAGGAATGCCAGGCCCTCTAACATCCTCAATGGGCAAAAGGA	1265
1266	AGATAGACACATGTCGCTTTGACCCCTGGAACATCTATAAAATATAGCTGTAAACCTGG	1325
1326	CTATGTGCTGGTGGGAGAAGAATCCATACAGTGTACCTCTGAGGTGTGGACACCCCTGT	1385
1386	ACCCCAATGCAAAAGTGGCAGCGTGTGAAGCTACAGGAAGGCAACTCTTGACAAAACCCCA	1445
1446	GCACCAATTGTAGACAGATGTCAACTCTTCTTGTGGTGAAGGTACAAGTTAAGTGG	1505
1506	GAGTGTTCATCAGGAGTGTCAAGGCACAATTCCTTGGTTTATGGAGATTCGTCTTTGTAA	1565
1566	AGAAATCACCTGCCACCAACCCCTGTTATCTACAATGGGGCACACACCGGGAGTTCCTT	1625

FIGURE 1. Nucleotide sequence of the human CR2 cDNA. Nucleotide sequence analysis was performed on three overlapping cDNA clones isolated from a human tonsillar cDNA library in λ gt11. The composite cDNA is represented by the top line of *A*, with the positions of the following restriction endonuclease sites indicated: SSp I (S), Pst I (P), Bbv I (B), Eco RI (R), and Xba I (X). The placement of the overlapping cDNA clones is indicated below. The insert above λ 6.11, labeled 10a, indicates the position of the additional sequence present in this clone but not in λ 4.11 or λ 6.21. The position and direction of the M13 subclones which were sequenced are indicated by the arrows in the lower portion of *A*. The nucleotide sequence of the human CR2 cDNA is shown in *B*. The sequences corresponding to the predicted signal peptide (+1-60) and the transmembrane region (2,923-2,994) are underlined. The position of the termination codon (3,150) is double underlined and the recognition sequence for

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1626 AGAAGATTTTCCATATGGAACACGCTCACTTACACATGTAACCTGGGCCAGAAAAGAGG 1685
1686 AGTGGAATTCAGCCTCATTGGAGAGAGCACCATCCGTTGTACAAGCAATGATCAAGAAAG 1745
1746 AGGCACCTGGAGTGGCCCTGCTCCCTGTGTAAACTTTCCCTCCTTGCTGTCCAGTGCTC 1805
1806 ACATGTCCATATTGCAATGGATACAAGATATCTGGCAAGGAAGCCCATATTTCTACAA 1865
1866 TGACACTGTGACATTCAAGTGTTATAGTGGATTTACTTTGAAGGGCAGTAGTCAGATTCG 1925
1926 TTGCAAAAGCTGATAACACCTGGGATCCTGAAATACCAGTTTGTGAAAAGAAAACATGCCA 1985
1986 GCATGTGAGACAGAGTCTTCAAGAACTTCCAGCTGGTTACCGTGTGGAGCTAGTTAATAC 2045
2046 GTCTGCCAAGATGGGTACCAGTTGACTGGACATGCTTATCAGATGTGTCAAGATGCTGA 2105
2106 AAATGGAATTTGGTTCAAAAAGATTCCACTTTGTAAAGTTATTCACTGTACCCCTCCACC 2165
2166 AGTGATTGTCAATGGGAAGCACACAGGCATGATGGCAGAAAACCTTCTATATGGAATGA 2225
2226 AGTCTCTTATGAATGTGACCAAGGATTCTATCTCCTGGGAGAGAAAAAATTGCAGTGCAG 2285
2286 AAGTGATTCTAAAGGACATGGATCTTGGAGCGGGCCTTCCCCACAGTGCTTACGATCTCC 2345
2346 TCCTGTGACTCGCTGCCCTAATCCAGAAGTCAAACATGGGTACAAGCTCAATAAAACACA 2405
2406 TTCTGCATATCCCACAATGACATAGTGTATGTTGACTGCAATCCTGGCTTCATCATGAA 2465
2466 TGGTAGTCGCGTGATTAGGTGTCTACTGATAACACATGGGTGCCAGGTGTGCCAACTTG 2525
2526 TATCAAAAAGCCTTCATAGGGTGTCCACCTCCGCCTAAGACCCCTAACGGGAACCATAC 2585
2586 TGGTGGAACATAGCTCGATTTTCTCCTGGAATGTCAATCCTGTACAGCTGTGACCAAGG 2645
2646 CTACCTGCTGGTGGGAGAGGCACTCCTTCTTTGCACACATGAGGGAACCTGGAGGCCAAC 2705
2706 TGCCCTCATTTGTAAGAGGTAAACTGTAGCTCACCAGCAGATATGGATGGAATCCAGAA 2765
2766 AGGGCTGGAACCAAGGAAAAATGATCAGTATGGAGCTGTTGTAACCTCGGAGTGTGAAGA 2825
2826 TGGGTATATGCTGGAAGGCAGTCCCAGAGCCAGTGCCCAATCGGATCACCAATGGAACCC 2885
2886 TCCCTGGCGGTTTGCAGATCCCGTTCACTTGCTCCTGTCCTTTGTGGTATTGCTGCAGG 2945
2946 TTTGATACTTCTTACCTTCTTGATTGTCGTTACCTTATACGTGATATCAAAACACAGAGC 3005
3006 ACGCAATATTATACAGATACAAGCCAGAAAGAAGCTTTTCATTAGAAAGCAGCGAAGT 3065
3066 ATATTCTGTTGATCCATACAACCCAGCCAGCTGATCAGAAGACAACTGGTGTGTGCCTC 3125
3126 ATTGCTTGAATTCAGCGGAATATTGATTAGAAAGAACTGCTCTAATATCAGCAAGTCT 3185
3186 CTTTATATGGCCTCAAGATCAATGAAATGATGTCTAAGCGGATCACTTCTATATGCACT 3245
3246 TATTCTCAAGAAGAACATCTTTATGGTAAAGATGGGAGCCAGTTTCACTGCCATATACT 3305
3306 CTTCAAGGACTTTCTGAAGCCTCACTTATGAGATGCCTGAAGCCAGGCCATGGCTATAAA 3365
3366 CAATTACATGGCTCTAAAAAGTTTGGCCCTTTTAAGGAAGGCACTAAAAAGAGCTGTCC 3425
3426 TGGTATCTAGACCATCTCTTTTGAATCAGCATACTCAATGTTACTATCTGCTTTTG 3485
3486 GTTATAATGTGTTTTTAATTATCTAAAGTATGAAGCATTTCGGGGTTATGATGGCCTT 3545
3546 ACCTTTATTAGGAAGTATGGTTTTATTGATAGTAGCTTCCTCCTCTGGTTGGTGTAA 3605
3606 TCATTTCATTTTACCCCTTACTTGGTTGAGTTTCTCTACATTACTGTATATACTTTGCC 3665
3666 TTTCCATAATCACTCAGTGATTGCAATTGCAAGTTTTTTTAAATTATGGGAATCAAG 3725
3726 ATTTAATCTAGAGATTGGTGTACAATTGAGGCTTTGGATGTTTCTTTAGCAGTTTGT 3785
3786 GATAAGTTCTAGTTGCTTGTAAAAATTCACTTAATAATGTGTACATTAGTCATTCAATAA 3845
3846 ATTGTAATTGTAAGAAAACAAA 3868

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polyadenylation (3,841) is in bold. The arrow between positions 1,974 and 1,975 indicates the site of insertion of the additional 177 nucleotides in $\lambda 6.11$. *C* contains the nucleotide sequence of this region of $\lambda 6.11$. These sequence data have been submitted to EMBL/Gen Bank Libraries under the accession number Y00649.