

Immunodominant T Cell Epitope from Signal Sequence

R. A. Henderson *et al.* (1) suggest that signal peptides or fragments thereof may be presented by class I major histocompatibility complex (MHC) antigens as T cell epitopes that are biologically relevant. This possibility is illustrated by the fact that a dominant target peptide presented by D^b and recognized by cytotoxic T cells specific for lymphocytic choriomeningitis virus (LCMV) in H-2^b mice is derived from the signal sequence of the viral glycoprotein precursor GP-C. The dominant immunogenic peptide is gp-a, which is

contained within amino acids 32 through 42 (2). This peptide is part of the leader sequence (amino acids 1 to 58) of the GP-C protein, the signal peptide of which was deduced by the predictive algorithm of G. von Heijne (3) and confirmed directly by NH₂-terminal amino acid sequencing of the GP-1 glycoprotein (4). Moreover, investigation of the pathway of signal peptide epitope presentation may also yield information of general biological interest about the fate of cleaved signal peptides in eukaryotic cells.

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Plakoglobin and β -Catenin: Distinct But Closely Related

Catenins are cytoplasmic proteins that were originally identified in association with the cytoplasmic domain of the cell adhesion molecule uvomorulin (E-cadherin), but they also form complexes with other proteins of the cadherin family (1, 2). P. D. McCrea *et al.* (3) isolated cDNA for the *Xenopus* β -catenin and performed a sequence comparison that revealed it to be homologous to mammalian plakoglobin and to the product of the segment polarity gene *armadillo* of *Drosophila*. They also found that a monoclonal antibody to bovine plakoglobin recognized canine β -catenin in Madin-Darby canine kidney (MDCK) cells, which raises the possibility that β -catenin may be identical to plakoglobin. However, we had found previously that antibodies to plakoglobin did not recognize components of the isolated uvomorulin-catenin complex (1).

To resolve this apparent contradiction, we used a cDNA probe of human plakoglobin (4) to screen a mouse cDNA library prepared from embryonal carcinoma cells, PCC4. We used intermediate hybridization and washing conditions to detect different degrees of homology (5). Phage clones with strong and weaker signals were plaque-purified. Restriction sites in the largest insert found in each group, 3.4 and 2.8 kb, respectively, were mapped and the insert was sequenced (Fig. 1).

We found the primary structure encoded by the cDNA with strong hybridization signals to have a high homology (99% identity) to human plakoglobin. In contrast, the deduced amino acid sequence of the cDNA with the weaker hybridization signal showed a strong homology (97% identity) to *Xenopus* β -catenin and a weak homology (68%) to human plakoglobin.

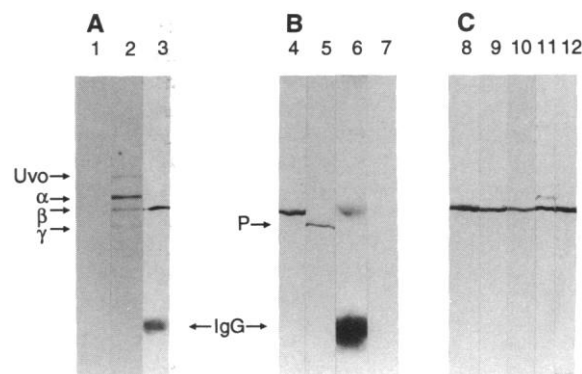
The cDNAs each detected a single copy gene in Southern blot (DNA) analysis of mouse genomic DNA (6).

These results indicate that mouse plakoglobin and β -catenin are distinct but related proteins. Out of 781 amino acids, β -catenin from mouse and from *Xenopus* differed only in 21 residues. Such evolutionary conservation indicates that β -catenin manifests basic cellular functions. To characterize β -catenin further and to demonstrate that

it forms a complex with uvomorulin, we produced rabbit antibodies to a synthetic peptide specific for β -catenin (Fig. 1). These antibodies specifically recognized β -catenin from uvomorulin-catenin complexes collected by antibodies to uvomorulin (Fig. 2A, lane 3). The antibodies specific for β -catenin cross-reacted on immunoblots (8) with the homologous protein from human, bovine, mouse, chicken, and *Xenopus* cell lysates (Fig. 2C). When immunoblots of whole-cell lysates from different species were stained with antibodies to β -catenin and to plakoglobin, these proteins exhibited different electrophoretic mobilities (Fig. 2B, lanes 4 and 5); the

Fig. 1 (right). Alignment of the mouse β -catenin (Mouse-Beta) and mouse plakoglobin (Mouse-Plako) protein sequences with the corresponding sequences of *Xenopus* β -catenin (Xenopus-Beta) and human plakoglobin (Human-Plako). The amino terminus of mouse plakoglobin (around 124 residues) is missing, so the residue number is provisional. Amino acids identical in three or four protein sequences are boxed; dots represent gaps. A synthetic peptide of the amino acid sequence, denoted by an overline, was used for antibody production. Abbreviations for the amino acid residues are as follows. A, Ala; C, Cys; D, Asp; E, Glu; F, Phe; G, Gly; H, His; I, Ile; K, Lys; L, Leu; M, Met; N, Asn; P, Pro; Q, Gln; R, Arg; S, Ser; T, Thr; V, Val; W, Trp; and Y, Tyr.

Fig. 2. Detection of β -catenin in uvomorulin-catenin complexes by antibodies to β -catenin and recognition of the homologous proteins in human, bovine, canine, chicken, and *Xenopus* cell lysates by SDS-PAGE analysis. (A) Uvomorulin-catenin complexes immunoprecipitated by antibodies to uvomorulin (lane 2) from cell lysates of mouse epithelial cells labeled with ³⁵S-methionine, CMT. Lane 1 is a control with an unrelated antibody. Lane 3 is an immunoblot of lane 2 stained with antibodies to β -catenin. (B) Whole-cell lysates (lanes 4 and 5) from MDCK cells, or immunoprecipitates collected with antibody to uvomorulin (lanes 6 and 7), were probed in immunoblots with antibodies to β -catenin (lanes 4 and 6) or to plakoglobin (lanes 5 and 7). (C) Whole-cell lysates from mouse embryonal carcinoma cells, F9 (lane 8), bovine MDBK cells (lane 9), human JAR cells (lane 10), chicken fibroblast cells (lane 11), and *Xenopus* epithelial cells, A6 (lane 12) were probed with antibodies to β -catenin.



antibodies reacted only with the respective proteins, which allowed us to investigate

the reactivity of both antibodies to components of the uvomorulin-catenin complex.

We investigated the possible association of plakoglobin with the uvomorulin-catenin complex in bovine and canine cells because the monoclonal antibody to bovine plakoglobin is of mouse origin. Immuno-complexes collected from MDCK and Madin-Darby bovine kidney (MDBK) cell lysates with rabbit antibodies to uvomorulin were blotted and stained with antibodies to either β -catenin or to plakoglobin. Where-as antibodies to plakoglobin were negative, β -catenin was easily detectable in the complex (Fig. 2B, lanes 6 and 7). Therefore, in spite of the high degree of homology, plakoglobin and β -catenin exhibited distinct functional properties. These results are in agreement with pulse-chase experiments in which β -catenin was found to associate with the uvomorulin precursor polypeptide (9), which suggests that only β -catenin can interact directly with uvomorulin. Nevertheless, a possible association of plakoglobin with the uvomorulin-catenin complex cannot be ruled out. Different solubilization procedures, with other detergents or ionic strengths, could result in soluble protein complexes of various sizes. It seems possible that under certain solubilization conditions plakoglobin could interact with the uvomorulin-catenin complex. In immunofluorescence tests on permeabilized canine MDCK and bovine MDBK cells, both proteins localized at the plasma membrane. Moreover, in epithelial cells, plakoglobin is a component of the desmosomal dense plaque, but it is also localized in the adherens-type junctions, where uvomorulin and catenins are found (10). These findings suggest a close colocalization and, together with the structural homology, indicate that both proteins may be involved in similar cellular processes.

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5. To obtain cDNA clones coding for mouse plakoglobin or mouse β -catenin, we screened 5×10^5 clones of a mouse λ gt11 cDNA library prepared from embryonal carcinoma cells PCC4 and probed with radioactively labeled, full-length cDNA for human plakoglobin (4) under intermediate conditions ($1 \times$ standard sodium citrate; 0.1% sodium dodecyl sulfate at 65°C). The inserts from plaques with strong and weak signals were sized, the bacteriophages with the largest insert

Mouse-Beta	MATQADLMEL	DMAMEPDRKA	AVSHNQQQSY	LDSGIHSGAT	40
Xenopus-Beta	MATQADLMEL	DMAMEPDRKA	AVSHNQQQSY	LDSGIHSGAT	40
Human-Plako		EVMMNLMEQPI	KVTEHQQQSY	YDSGIHSGAN	31
Mouse-Plako					
Mouse-Beta	TTAPSLGK	NPEEDVDVTS	QVLYEWEQGF	SSFTQEQVA	80
Xenopus-Beta	TTAPSLGK	NPEEDVDVTS	QVLYEWEQGF	SSFTQEQVA	80
Human-Plako	TCVPSVSG	IMEDEACGR	QYTLKKTTTY	TSGVFPSSGD	71
Mouse-Plako					
Mouse-Beta	DIDGQYAMTE	QQRVFAAMFF	ETLDEGMOQIP	STQFDAAHPT	120
Xenopus-Beta	DIDGQYAMTE	QQRVFAAMFF	ETLDEGMOQIP	STQFDAAHPT	120
Human-Plako	LEYQMSTTAR	AKRVFAAMCF	GVSGEGQQLAL	LATQVEGQAT	111
Mouse-Plako					
Mouse-Beta	RVORLAEPSC	MLPHHAVNLI	HYODDAELAT	RAIFELTILL	160
Xenopus-Beta	RVORLAEPSC	MLPHHAVNLI	HYODDAELAT	RAIFELTILL	160
Human-Plako	NLORLAEPSC	MLPHHAVNLI	HYODDAELAT	RAIFELTILL	151
Mouse-Plako					27
Mouse-Beta	NDELQVVVN	AAVMHQLSL	FEASRRAIMR	SPQMVAIVR	200
Xenopus-Beta	NDELQVVVN	AAVMHQLSL	FEASRRAIMR	SPQMVAIVR	200
Human-Plako	NDEDPVVVT	AAVMHQLSL	FEASRRAIMR	SPQMVAIVR	191
Mouse-Plako	NDEDPVVVT	AAVMHQLSL	FEASRRAIMR	SPQMVAIVR	67
Mouse-Beta	THONTNVEET	ARCTAGTLLH	LSHHREGGLLA	IFKSGGIPAL	240
Xenopus-Beta	THONTNVEET	ARCTAGTLLH	LSHHREGGLLA	IFKSGGIPAL	240
Human-Plako	THONTNVEET	ARCTAGTLLH	LSHHREGGLLA	IFKSGGIPAL	231
Mouse-Plako	THONTNVEET	ARCTAGTLLH	LSHHREGGLLA	IFKSGGIPAL	107
Mouse-Beta	VKMLGSPVD	VLFFAATTLLH	NLLHVOEGAF	MAVRLAGSLO	280
Xenopus-Beta	VKMLGSPVD	VLFFAATTLLH	NLLHVOEGAF	MAVRLAGSLO	280
Human-Plako	VRMLGSPVD	VLFFAATTLLH	NLLHVOEGAF	MAVRLAGSLO	271
Mouse-Plako	VRMLGSPVD	VLFFAATTLLH	NLLHVOEGAF	MAVRLAGSLO	147
Mouse-Beta	EMVALLNTH	VFPLAITTDC	LCLLAYGHQ	CELILLANS	320
Xenopus-Beta	EMVALLNTH	VFPLAITTDC	LCLLAYGHQ	CELILLANS	320
Human-Plako	EMVALLNTH	VFPLAITTDC	LCLLAYGHQ	CELILLANS	311
Mouse-Plako	EMVALLNTH	VFPLAITTDC	LCLLAYGHQ	CELILLANS	187
Mouse-Beta	FOALNIMRT	ATFERLLTTH	SRVLEVLVSV	SENFPAIVEA	360
Xenopus-Beta	FOALNIMRT	ATFERLLTTH	SRVLEVLVSV	SENFPAIVEA	360
Human-Plako	FOALNIMRT	ATFERLLTTH	SRVLEVLVSV	SENFPAIVEA	351
Mouse-Plako	FOALNIMRT	ATFERLLTTH	SRVLEVLVSV	SENFPAIVEA	227
Mouse-Beta	SGHOALGRL	TDPSQELVQ	LSLTLERLST	ATFQEGMES	400
Xenopus-Beta	SGHOALGRL	TDPSQELVQ	LSLTLERLST	ATFQEGMES	400
Human-Plako	SGHOALGRL	TDPSQELVQ	LSLTLERLST	ATFQEGMES	391
Mouse-Plako	SGHOALGRL	TDPSQELVQ	LSLTLERLST	ATFQEGMES	267
Mouse-Beta	LLGTLVQLG	SDIINAVTCA	ATLSHLTCH	YRHHMMYCO	440
Xenopus-Beta	LLGTLVQLG	SDIINAVTCA	ATLSHLTCH	YRHHMMYCO	440
Human-Plako	VLKILVNQCS	VDIIVNLTCA	ATLSHLTCH	YRHHMMYCO	431
Mouse-Plako	VLKILVNQCS	VDIIVNLTCA	ATLSHLTCH	YRHHMMYCO	307
Mouse-Beta	VGIIEALVRT	VLRAGIREDO	TEPAICALLR	LTSRHPQEAEM	480
Xenopus-Beta	VGIIEALVRT	VLRAGIREDO	TEPAICALLR	LTSRHPQEAEM	480
Human-Plako	NSGIEALVTH	VLRAGIREDO	TEPAICALLR	LTSRHPQEAEM	471
Mouse-Plako	NSGIEALVTH	VLRAGIREDO	TEPAICALLR	LTSRHPQEAEM	347
Mouse-Beta	AQAARREHYG	LIVVFLHP	ISHIIPAT	VRLIHLALC	520
Xenopus-Beta	AQAARREHYG	LIVVFLHP	ISHIIPAT	VRLIHLALC	520
Human-Plako	AQAARREHYG	LIVVFLHP	ISHIIPAT	VRLIHLALC	511
Mouse-Plako	AQAARREHYG	LIVVFLHP	ISHIIPAT	VRLIHLALC	387
Mouse-Beta	PAHHAPRERQ	GATIELLYQL	RAHOITQVR	TSMGSTOCQF	560
Xenopus-Beta	PAHHAPRERQ	GATIELLYQL	RAHOITQVR	TSMGSTOCQF	560
Human-Plako	PAHHAPRERQ	GATIELLYQL	RAHOITQVR	TSMGSTOCQF	550
Mouse-Plako	PAHHAPRERQ	GATIELLYQL	RAHOITQVR	TSMGSTOCQF	426
Mouse-Beta	VEVHEEIA	EACTGALHII	ARIHHPFIVI	RGHTIHLFPA	600
Xenopus-Beta	VEVHEEIA	EACTGALHII	ARIHHPFIVI	RGHTIHLFPA	600
Human-Plako	TDVHEEIA	EACTGALHII	ARIHHPFIVI	RGHTIHLFPA	590
Mouse-Plako	TDVHEEIA	EACTGALHII	ARIHHPFIVI	RGHTIHLFPA	466
Mouse-Beta	LLLYPIENT	EPAAALCL	LAVIDEAAES	ENEGATAPL	640
Xenopus-Beta	LLLYPIENT	EPAAALCL	LAVIDEAAES	ENEGATAPL	640
Human-Plako	LLLYSVENI	EPAAALCL	LAVIDEAAES	ENEGATAPL	630
Mouse-Plako	LLLYSVENI	EPAAALCL	LAVIDEAAES	ENEGATAPL	506
Mouse-Beta	TELLHFNHG	VNTAAALF	IMEDFPQD	KHLLVELTS	680
Xenopus-Beta	TELLHFNHG	VNTAAALF	IMEDFPQD	KHLLVELTS	680
Human-Plako	MELLHFNHG	VNTAAALF	IMEDFPQD	KHLLVELTS	670
Mouse-Plako	MELLHFNHG	VNTAAALF	IMEDFPQD	KHLLVELTS	546
Mouse-Beta	LFRTEMP	NETADLGLDI	GAQGEALGYR	QDPSISFPH	720
Xenopus-Beta	LFRTEMP	NETADLGLDI	GAQGEALGYR	QDPSISFPH	720
Human-Plako	LFRTEMP	NETADLGLDI	GAQGEALGYR	QDPSISFPH	705
Mouse-Plako	LFRTEMP	NETADLGLDI	GAQGEALGYR	QDPSISFPH	581
Mouse-Beta	GGYGQAG	DPMMEHG	HHGADPV	DGLDLGHQ	760
Xenopus-Beta	GGYGQAG	DPMMEHG	HHGADPV	DGLDLGHQ	760
Human-Plako	SDVPLPE	DPMMEHG	HHGADPV	DGLDLGHQ	741
Mouse-Plako	SDVPLPE	DPMMEHG	HHGADPV	DGLDLGHQ	617
Mouse-Beta	DLMDGLFPGD	SNQLAWFDTD	L781		
Xenopus-Beta	DLMDGLFPGD	SNQLAWFDTD	L781		
Human-Plako	HMLA		745		
Mouse-Plako	HMLA		621		

in each class, designated λ W6 and λ H8, were cloned into pBluescript I SK+ vector, and both strands were sequenced (Sequenase Version 2.0, protocol U.S. Biochemicals, Cleveland, OH). The 3.4 kb fragment of λ H8 represents the full-length cDNA of mouse β -catenin, whereas the 2.8 kb fragment of λ W6 encodes part of mouse plakoglobin, beginning at amino acid 124. The deduced amino acid sequence of mouse β -catenin has a calculated molecular size of 85.5 kD, which agrees with 88 kD, the molecular size of uvomorulin-complexed β -catenin. Protein sequence alignment was done with a Genetics Computer Group program for the VAX. The GenBank accession number for mouse β -catenin is M90364 and for mouse plakoglobin, M90365.

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7. A peptide containing the amino acid sequence overlined in Fig. 1, which represents the carboxyl terminus of β -catenin, was coupled by glutaraldehyde to keyhole limpet hemocyanin (Sigma, St. Louis, MO). After three subcutaneous immunizations at 3-week intervals, specific antibodies were isolated on a peptide- ϵ -amino-hexanoyl (EAH)-Sepharose (Pharmacia, Fairfield, NJ), column (5 mg of peptide coupled to 1 ml of EAH-Sepharose 4B, as described by the manufacturer).
8. Immunoprecipitation and immunoblot experiments were performed as described in (1). Immunocomplexes were collected from lysates of 4×10^6 cells with 4 μ g of antibodies to uvomorulin and Protein A-Sepharose (Pharmacia). Proteins were separated by 8% SDS-PAGE under reducing conditions. Antibodies to β -catenin (5 μ g per milliliter) and to plakoglobin (1 μ g per milliliter) were diluted in phosphate-buffered saline in immunoblots. Mouse monoclonal antibody to bovine plakoglobin was purchased from Progen (Heidelberg, Federal Republic of Germany).
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Response: The work presented by Butz *et al.* appears to be in agreement with the principal conclusion of our earlier work (1), which demonstrated that β -catenin is highly homologous to the *Drosophila* gene product *armadillo* and to human plakoglobin. At that time, we erred in favoring the interpretation that plakoglobin and β -catenin were the same protein because we did not notice the small difference in their gel mobilities and did not yet have antibodies to β -catenin (or to *armadillo*) that would have made that difference more obvious. Nonetheless, the greater degree of sequence conservation maintained over evolutionary time between *Xenopus* β -catenin and *Drosophila armadillo*, compared with that between *Xenopus* β -catenin and human plakoglobin (both from vertebrates), led us to discuss the possibility that β -catenin and plakoglobin might be distinct members of a gene family.

Butz *et al.* present excellent and interesting evidence that β -catenin and plakoglobin are distinct, although closely related; proteins within the same cell. We are in complete agreement concerning this issue; on the basis of recent immunological evidence from experiments with MDCK cells, we have, in a collaborative effort, reached the same conclusion (2).

Our findings in (2) and that of Butz *et al.* differ on one point. We find that plakoglo-

bin is a component of the E-cadherin-catenin complex. Although it is more weakly associated with the complex than is β -catenin [as determined by the ease with which it is removed from the complex with detergents and other washes (3)], it specifically coimmunoprecipitates with E-cadherin from MDCK cells (1, 2). As Butz *et al.* point out, it is conceivable (we think likely) that differences in the composition of the solutions used for cell extraction, immunoprecipitate washing, and so forth may explain why we more readily observe plakoglobin in E-cadherin immunoprecipitates. It remains to be determined whether plakoglobin is the γ -catenin polypeptide present in E-cadherin immunoprecipitates.

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