

# Epitope Mapping of Anti-Tiger Podoplanin Monoclonal Antibody PMab-231

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Podoplanin (PDPN) is expressed on podocytes of the kidneys, type I alveolar cells of the lungs, and lymphatic endothelial cells. PDPN comprises three platelet aggregation-stimulating (PLAG) domains (PLAG1, PLAG2, and PLAG3) in the N-terminus and PLAG-like domains in the middle of the PDPN protein. We have previously reported on an anti-tiger PDPN (tigPDPN) monoclonal antibody (mAb), PMab-231, which was developed using the Cell-Based Immunization and Screening (CBIS) method. PMab-231 is very useful in flow cytometry, Western blotting, and immunohistochemical analyses; however, the binding epitope of PMab-231 remains to be elucidated. This study aimed to investigate the epitopes of PMab-231, which was developed by CBIS method, using enzyme-linked immunosorbent assay. The results revealed that the critical epitopes of PMab-231 are Glu29, Asp30, Asp31, Ile32, Met33, Thr34, Pro35, Gly36, and Glu38 of tigPDPN, which is corresponding to PLAG1/2. The findings of our study can be applied to the production of more functional anti-tigPDPN mAbs.

**Keywords:** podoplanin, PDPN, PMab-231, epitope mapping

## Introduction

**P**ODOPLANIN (PDPN) IS A type I transmembrane sialoglycoprotein that is expressed in normal tissues, including renal corpuscles, type I lung alveolar cells, and lymphatic endothelial cells.<sup>(1,2)</sup> PDPN induces platelet aggregation by binding to C-type lectin-like receptor-2 (CLEC-2) on platelets.<sup>(1,3–9)</sup> The interaction between PDPN on lymphatic endothelial cells and CLEC-2 on platelets facilitates the separation of embryonic blood/lymphatic vessels.<sup>(10)</sup> Studies have reported the expression of human PDPN in several malignant tumors, such as oral squamous cell carcinomas,<sup>(11)</sup> malignant mesotheliomas,<sup>(12,13)</sup> lung cancers,<sup>(14)</sup> and malignant brain tumors.<sup>(15–18)</sup> PDPN expression has also been associated with malignant progression and cancer metastasis.<sup>(6,15,19)</sup> Although pathophysiological studies of alveolar cells of the lungs, podocytes of the kidneys, or lymphatic endothelial cells of the colon using anti-tigPDPN monoclonal antibodies (mAbs) are important in many disorders, mAbs against tigPDPN remain to be developed.

In our previous study, we used the Cell-Based Immunization and Screening (CBIS) method to develop specific and sensitive mAbs against the tigPDPN of 163 amino acids to facilitate the immunohistochemical analysis of paraffin-

embedded tissue sections.<sup>(20)</sup> Screening identified strong signals against Chinese hamster ovary (CHO)/tigPDPN cells and weak or no signals against CHO-K1 cells in 18 of 958 wells (1.9%). One of the six clones, PMab-231 (IgG<sub>2a</sub>, kappa), was finally selected using immunohistochemistry against paraffin-embedded feline normal tissue sections because tigPDPN and cat PDPN (cPDPN) shared 99% homology.<sup>(21)</sup> The dissociation constants ( $K_D$ ) of PMab-231 for CHO/tigPDPN and CHO/cPDPN cells were determined to be  $1.2 \times 10^{-8}$  M and  $1.9 \times 10^{-8}$  M, respectively, indicating moderate affinity for CHO/tigPDPN and CHO/cPDPN cells. PMab-231 stained type I alveolar cells of the feline lungs and podocytes of the feline kidneys using immunohistochemistry. This study aimed to determine the binding epitope of PMab-231 to tigPDPN using enzyme-linked immunosorbent assay (ELISA).

## Materials and Methods

### Enzyme-linked immunosorbent assay

Synthesized tigPDPN peptides using PEPSCREEN (Sigma-Aldrich Corp., St. Louis, MO) were immobilized on Nunc Maxisorp 96-well immunoplates (Thermo Fisher Scientific, Inc., Waltham, MA) at 10  $\mu$ g/mL for 30 minutes at 37°C.

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TABLE 1. DETERMINATION OF PMAB-231 EPITOPE BY ENZYME-LINKED IMMUNOSORBENT ASSAY

Peptide	Sequence	PMab-231
24-43	STVRPEDDIMTPGVEDGMVT	++
34-53	TPGVEDGMVTLGAEDNVVTT	-
44-63	LGAEDNVVTTGAGADPEEST	-
54-73	GAGADPEESTGPTGLVPTNT	-
64-83	GPTGLVPTNTESITDFHIED	-
74-93	ESITDFHIEDGPTQESTGHA	-
84-103	GPTQESTGHANEESQSTTTL	-
94-113	NEESQSTTTLNVVTSHSIEK	-
104-123	NVVTSHSIEKIGEDTETTVE	-
114-129	IGEDTETTVEKDGLAT	-

++,  $0.3 \leq OD_{655} < 0.6$ ; -,  $OD_{655} < 0.1$ .

TABLE 2. DETERMINATION OF PMAB-231 EPITOPE BY ENZYME-LINKED IMMUNOSORBENT ASSAY

Mutation	Sequence	PMab-231
S24A	ATVRPEDDIMTPGVEDGMVT	+
T25A	SAVRPEDDIMTPGVEDGMVT	+
V26A	STARPEDDIMTPGVEDGMVT	+
R27A	STVAPEDDIMTPGVEDGMVT	+
P28A	STVRAEDDIMTPGVEDGMVT	+
E29A	STVRPADDIMTPGVEDGMVT	-
D30A	STVRPEADIMTPGVEDGMVT	-
D31A	STVRPEDAIMTPGVEDGMVT	-
I32A	STVRPEDDAMTPGVEDGMVT	-
M33A	STVRPEDDIATPGVEDGMVT	-
T34A	STVRPEDDIMAPGVEDGMVT	-
P35A	STVRPEDDIMTAGVEDGMVT	-
G36A	STVRPEDDIMTPAVEDGMVT	-
V37A	STVRPEDDIMTPGAEDGMVT	+
E38A	STVRPEDDIMTPGVADGMVT	-
D39A	STVRPEDDIMTPGVEAGMVT	+
G40A	STVRPEDDIMTPGVEDAMVT	+
M41A	STVRPEDDIMTPGVEDGAVT	+
V42A	STVRPEDDIMTPGVEDGMAT	+
T43A	STVRPEDDIMTPGVEDGMVA	+

+,  $0.1 \leq OD_{655} < 0.3$ ; -,  $OD_{655} < 0.1$ .

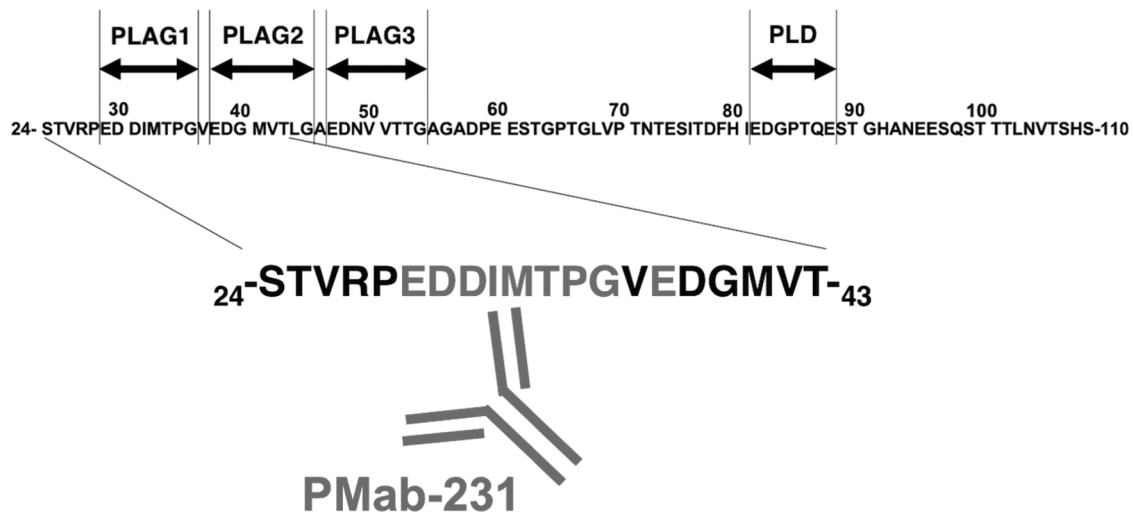
After blocking with SuperBlock T20 (PBS) Blocking Buffer (Thermo Fisher Scientific, Inc.), the plates were incubated with purified PMab-231 ( $1 \mu\text{g/mL}$ ), followed by a peroxidase-conjugated anti-mouse IgG (Agilent Technologies, Inc., Santa Clara, CA) dilution of 1:2000. Enzymatic reactions were performed using 1-Step Ultra TMB-ELISA (Thermo Fisher Scientific, Inc.). Optical density was measured at 655 nm using an iMark microplate reader (Bio-Rad Laboratories, Inc., Berkeley, CA). These reactions were performed at  $37^\circ\text{C}$  using a total sample volume of 50–100  $\mu\text{L}$ .

## Results and Discussion

We have previously produced mAbs against HER2<sup>(22)</sup> or EGFR<sup>(23)</sup> using CasMab technology and mAbs against CD44, PD-L1, or CD133 using the CBIS method.<sup>(24–28)</sup> However, the binding epitopes of these mAbs (H<sub>2</sub>Mab-77 against HER2, EMab-51 against EGFR, C<sub>44</sub>Mab-5 against CD44, L<sub>1</sub>Mab-13 against PD-L1, and CMab-43 against CD133) remain to be determined because these mAbs did not react with synthesized peptides. Conversely, the binding epitopes of mAbs produced by immunizing with synthesized peptides have been easily determined using deletion mutants and point mutants of synthesized peptides.<sup>(29,30)</sup>

We first synthesized a series of tigPDPN peptides, which are summarized in Table 1. Using ELISA, PMab-231 detected 24–43 peptides corresponding to the 24th–43rd amino acids (aas) of tigPDPN. We then synthesized point mutants of 24–43 peptides (Table 2). Using ELISA, PMab-231 detected S24A, T25A, V26A, R27A, P28A, V37A, D39A, G40A, M41A, V42A, and T43A. Conversely, it did not detect E29A, D30A, D31A, I32A, M33A, T34A, P35A, G36A, and E38A, indicating that Glu29, Asp30, Asp31, Ile32, Met33, Thr34, Pro35, Gly36, and Glu38 are included in the critical epitope of PMab-231. Although PMab-231 was produced using the CBIS method, the epitope of PMab-231 was fortunately determined by ELISA.

We previously developed mAbs against human,<sup>(31)</sup> mouse,<sup>(31)</sup> rat,<sup>(32)</sup> rabbit,<sup>(33)</sup> dog,<sup>(34)</sup> cat,<sup>(27)</sup> bovine,<sup>(35)</sup> horse,<sup>(36–38)</sup> and pig<sup>(39)</sup> PDPNs. PDPN comprises three tandem-repeat of the “EDxxVTPG” sequences, which were defined to be platelet aggregation-stimulating (PLAG) domains (PLAG1,



**FIG. 1.** Schematic illustration of the epitope recognized by PMab-231. The critical epitope of PMab-231 are shown in gray letters. PLAG, platelet aggregation-stimulating; PLD, PLAG-like domain.

PLAG2, and PLAG3) in the N-terminus of the PDPN protein.<sup>(4)</sup> There are several PLAG-like domains (PLDs) of the “E(D/E)xx(T/S)xx” sequence in the middle of the PDPN protein. PLDs are reportedly important for PDPN–CLEC-2 interactions.<sup>(40)</sup> Almost all mAbs against PDPNs reportedly react with PLAG domains or PLDs.<sup>(40,41)</sup> As summarized in Figure 1, PMab-231 also detected PLAG1 (Glu29, Asp30, Asp31, Ile32, Met33, Thr34, Pro35, and Gly36) and a part of PLAG2 (Glu38) of tigPDPN. Therefore, PLAG1 of tigPDPN was clarified to be the advantageous epitope for several applications, such as flow cytometry, Western blotting, and immunohistochemical analyses. These findings can be applied to the production of more functional anti-tigPDPN mAbs.

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