

Gln-589 is highlighted in cyan. Note that Ser-462 is located above the α -*genu*, while Ser-528 is located on the side of the thigh domain distal to the α -*genu*.

Fig. 5. Effect of anti- α V β 3 mAbs on fibrinogen binding. A. FITC-fibrinogen binding to cells expressing α V β 3 in the presence of 1 mM Ca^{2+} and 1 mM Mg^{2+} (open column) or in the presence of 2 mM Mg^{2+} and 5 μM EGTA (hatched column) is shown. The ratio of the MFI (FL1) to the MFI (FL2) in the gated cell population was used to normalize the binding with the expression of α V β 3 on the cell surface. B. FITC-fibrinogen binding to cells expressing wild-type α V β 3 in the presence of 2 mM Mg^{2+} and 5 μM EGTA was examined. Binding in the presence of 100 $\mu\text{g}/\text{mL}$ of the indicated mAb is shown in the hatched column. An equivalent volume of PBS, instead of the mAb solution, was included to obtain the control binding. The asterisks indicate statistically different binding abilities, compared with the control (* P <0.01, ** P <0.05).

Fig. 6. Effect of integrin extension on the function of the anti- α V β 3 mAbs. A. SDS-PAGE analysis of α V β 3 expressed on CHO cells. Cell lysates from biotin-labeled cells expressing α V β 3 were immunoprecipitated with anti- α V mAb P2W7. The precipitates were subjected to a 7.5% non-reducing gel and visualized using chemiluminescence. The positions of the molecular weight markers are shown on the left side of panel. Lane 1, parent CHO; lane 2, wild-type α V β 3; lane 3, Q589NAT. Note that the α V carrying the Q589NAT mutation (arrowhead) migrated more slowly than the wild type. B. FITC-fibrinogen binding to cells expressing α V β 3 carrying the Q589NAT mutation was examined as described in Fig. 4A. Binding in the presence of 1 mM Ca^{2+} and 1 mM Mg^{2+} (open column) or in the presence of 2

mM Mg^{2+} and 5 μ M EGTA (solid column) is shown. C. The effect of the anti- $\alpha V\beta 3$ mAbs on fibrinogen binding to cells expressing $\alpha V\beta 3$ carrying the Q589NAT mutation was examined, as described in Fig. 5B. Among the mAbs, only 7E3 significantly inhibited binding.

Fig. 7. Effect of integrin extension on the binding of the anti- $\alpha V\beta 3$ mAbs. Wild-type αV or αV carrying the Q589NAT mutation was transiently expressed together with wild-type $\beta 3$ in CHO cells. The binding of function-blocking mAbs to these cells was examined by FACS. The MFI obtained from the whole cell population with each mAb was normalized by the MFI obtained with an anti- αV mAb 17E6 that represents the αV expression. There was no significant difference in the binding of leg-binding mAbs in cells expressing Q589NAT (solid column) as compared with cells expressing wild-type αV (hatched column).

| | mIgG | 10C4 | 23C6 | LM609 | 17E6 | 69-6-5 | AMF-7 | M9 | P2W7 | P3G8 | SZ21 |
|--------|------|-------|-------|-------|-------|--------|-------|-------|-------|-------|-------|
| CHO | 5.26 | 4.77 | 4.34 | 4.48 | 4.94 | 5.23 | 3.57 | 3.15 | 18.99 | 4.25 | 3.63 |
| β3 | 2.72 | 55.83 | 70.09 | 61.93 | 3.64 | 4.66 | 2.92 | 3.93 | 4.08 | 5.84 | 49.89 |
| αVβ3 | 3.47 | 61.09 | 77.73 | 68.92 | 67.32 | 64.4 | 64.17 | 71.05 | 57.59 | 63.15 | 54.52 |
| αIIbβ3 | 3.92 | 57.69 | 58.25 | 65.47 | 3.66 | 4.25 | 3.07 | 4.08 | 4.61 | 5.56 | 51.73 |
| V/B | 4.18 | 58.73 | 69.11 | 61.9 | 53.13 | 45.19 | 3.08 | 3.78 | 6.00 | 6.36 | 52.16 |
| B/V | 4.24 | 51.6 | 63.63 | 57.3 | 7.11 | 6.44 | 58.39 | 63.47 | 49.97 | 55.63 | 49.16 |

Table 1. **MAB binding to cells expressing tail-swapping chimeras.** MAb binding to cells expressing wild-type human β3 (β3), wild-type human αVβ3 (αVβ3), wild-type human αIIbβ3 (αIIbβ3), tail-swapping mutants (V/B, B/V), and to parent CHO cells (CHO) was examined. The numbers represent the percentage of the cell population stained with each mAb.

| | mIgG | AMF-7 | M9 | P2W7 | P3G8 | 17E6 | SZ21 |
|----------------------|-------|-------|-------|-------|-------|-------|-------|
| CHO | 7.83 | 5.86 | 5.72 | 15.51 | 5.78 | 5.91 | 4.8 |
| α V β 3 | 5.6 | 68.11 | 78.41 | 63.75 | 67.97 | 76.32 | 59.23 |
| VT | 4.72 | 3.37 | 4.09 | 6.91 | 6.44 | 74.37 | 68.6 |
| VC1 | 10.88 | 82.45 | 90.34 | 71.17 | 76.44 | 90.39 | 83.9 |
| VC2 | 4.7 | 64.82 | 81.14 | 69.67 | 69.52 | 78.53 | 74.97 |
| T | 6.4 | 49.93 | 62.12 | 45.45 | 44.62 | 4.45 | 62.75 |
| C1 | 5.87 | 3.8 | 4.03 | 5.82 | 6.37 | 4.05 | 61.79 |
| C2 | 5.36 | 5.53 | 5.16 | 6.57 | 8.25 | 6.8 | 71.52 |

Table 2. **MAB binding to cells expressing domain-swapping chimeras.** The numbers represent the percentage of the cell population stained with each mAb.

| | mIgG | AMF-7 | M9 | P2W7 | P3G8 | 17E6 | SZ21 |
|---------|-------|-------|-------|-------|-------|-------|-------|
| CHO | 3.62 | 2.56 | 2.95 | 13.3 | 3.2 | 4.07 | 1.16 |
| WT | 6.8 | 74.11 | 85.85 | 78.21 | 83.47 | 87.07 | 71.41 |
| I441V | 4.15 | 76.76 | 70.66 | 72.64 | 77.61 | 79.97 | 70.78 |
| T460ICP | 5.15 | 5.84 | 3.26 | 7.01 | 70.72 | 77.52 | 65.15 |
| T460I | 5.88 | 62.11 | 63.27 | 54.8 | 56.43 | 70.81 | 64.01 |
| S462P | 5.52 | 14.66 | 3.75 | 17.17 | 66.41 | 76.18 | 66.88 |
| V486T | 11.93 | 72.65 | 78.35 | 71.75 | 64.14 | 82.23 | 71.1 |
| N492H | 10.04 | 70.78 | 79 | 70.14 | 72.01 | 82.49 | 73.56 |
| E496DV | 14.9 | 73.25 | 76.95 | 67.31 | 75.08 | 75.01 | 58.72 |
| Y515HN | 8.47 | 59.79 | 67.31 | 59.04 | 65.69 | 66.81 | 50.75 |
| S520V | 8.33 | 65.13 | 72.76 | 61.29 | 73.55 | 71.38 | 60.68 |
| N524T | 12.98 | 60.75 | 69.31 | 58.6 | 65.03 | 66.37 | 57.98 |
| I527VF | 9.55 | 74.93 | 78.94 | 59.25 | 9.26 | 84.31 | 81.13 |
| I527V | 3.12 | 75.57 | 78.3 | 64.88 | 77.78 | 83.6 | 67.47 |
| S528F | 5.32 | 80.94 | 82.19 | 76.5 | 10.59 | 85.83 | 78.1 |
| L532Q | 26.68 | 62.2 | 77.22 | 58.2 | 62.38 | 76.84 | 71.26 |
| I539V | 13.13 | 73.98 | 67.01 | 57.12 | 68.84 | 76.24 | 64.92 |
| Y565Q | 11.64 | 70.1 | 77.22 | 64.52 | 71.32 | 80.13 | 72.63 |
| T571A | 27.84 | 68.76 | 69.91 | 54.89 | 59.55 | 75.22 | 75.28 |
| I586V | 9.99 | 70.96 | 78.21 | 61.16 | 70.96 | 68.54 | 76.34 |

Table 3. **MAB binding to cells expressing human-to-mouse α V mutants.** The numbers represent the percentage of the cell population stained with each mAb. Bindings significantly lower than those for 17E6 or SZ21 are marked in red.

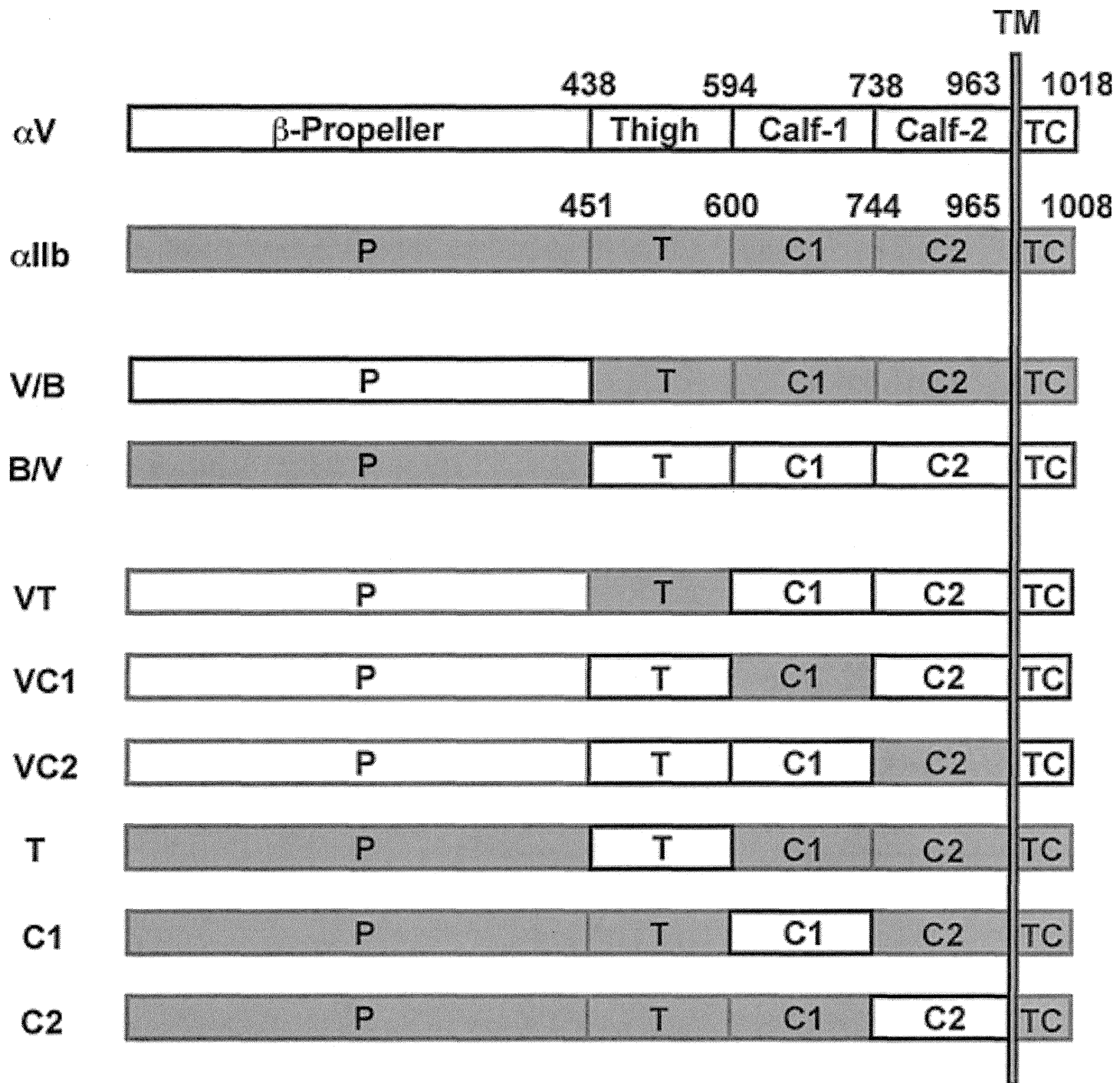
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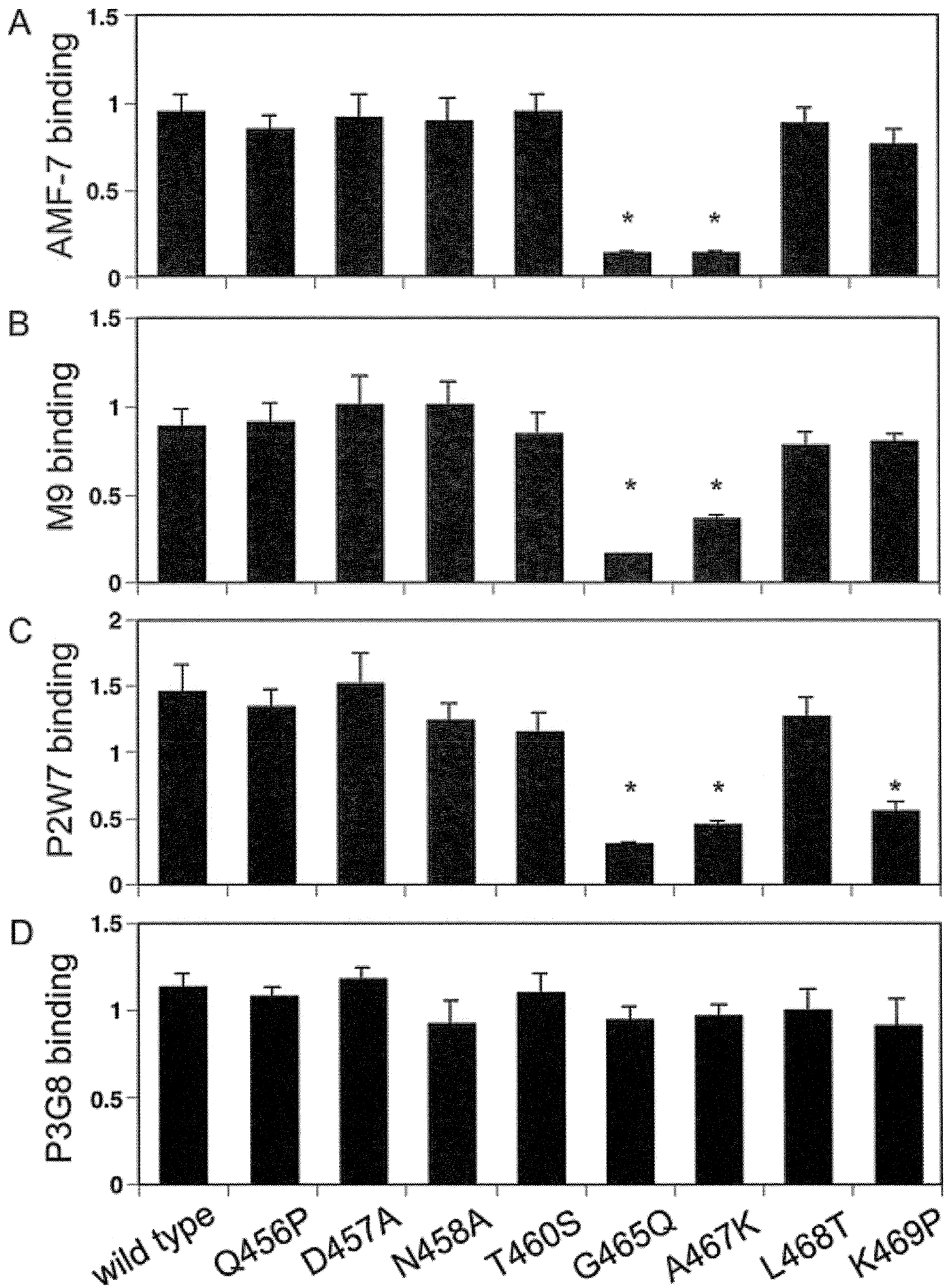
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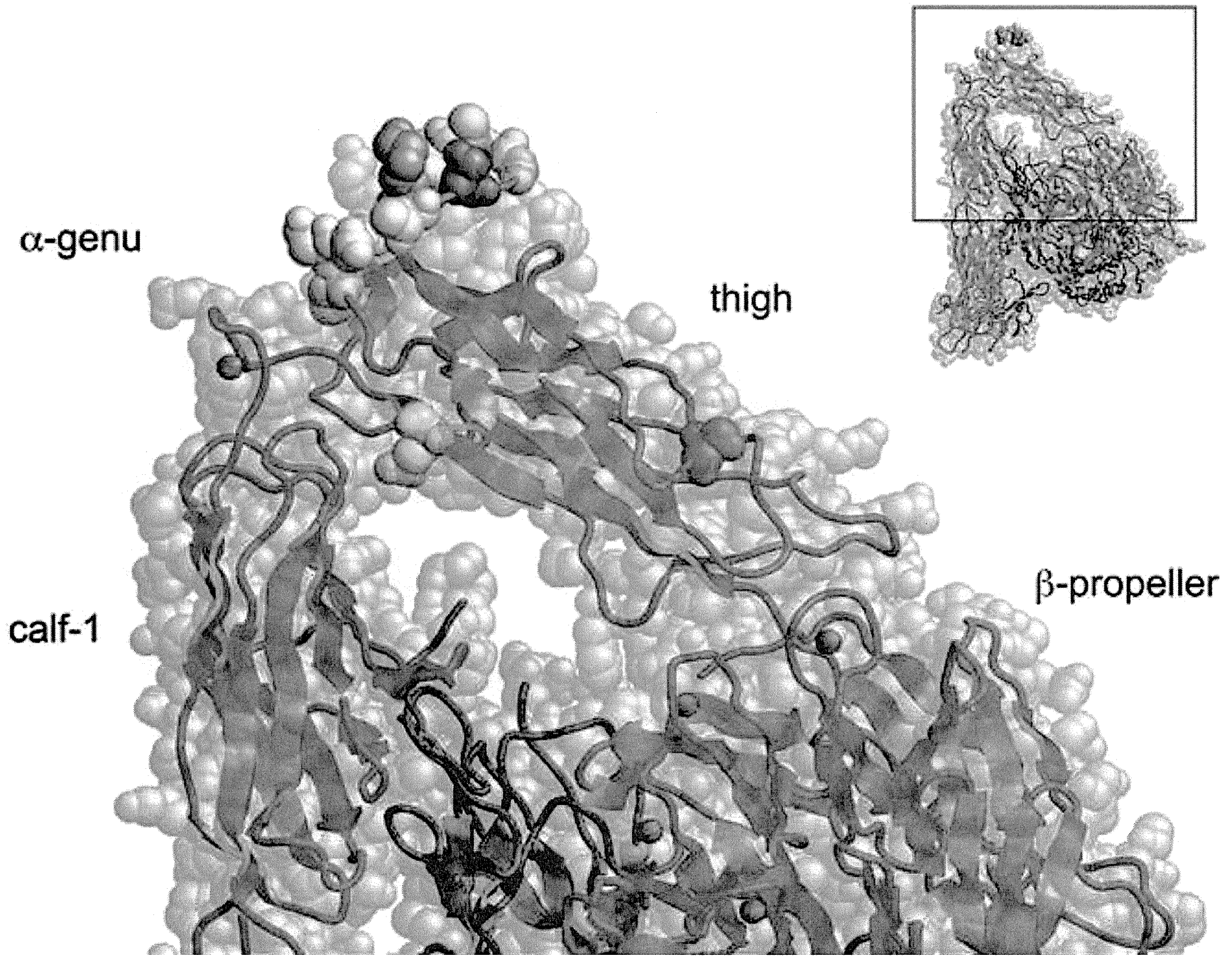
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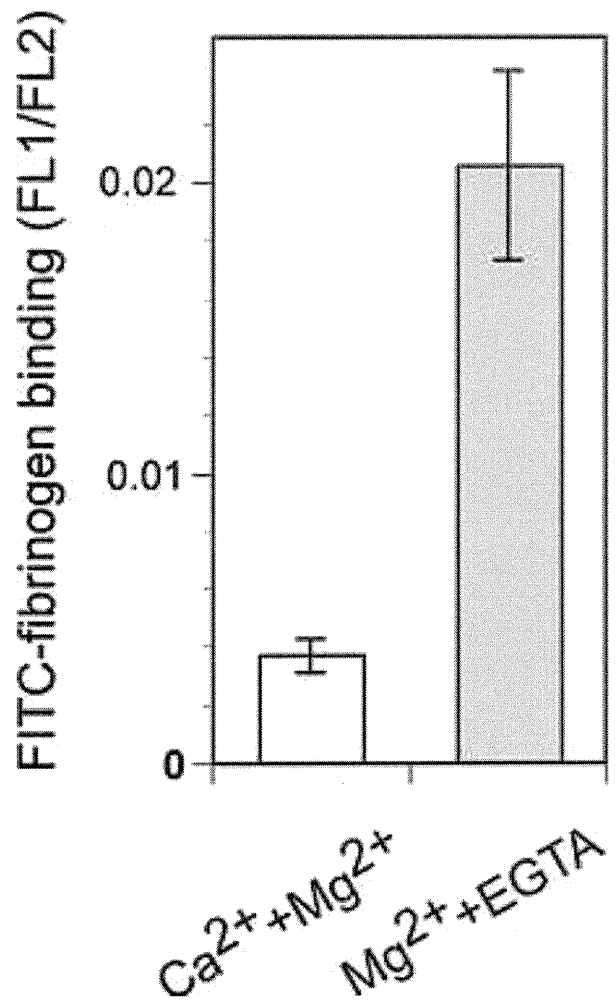


| | | | | | | | |
|-------------------|-----|--|-------------------------------|-------------|-------|----------|-----|
| human α V | 441 | ITVNAGLEVYPSILNQDNKTC | CSLPGTALKVSCFNVRF | CLKADGKGVLP | PRKLN | FQVELLLD | 500 |
| mouse α V | 441 | V.....I | P.....T.....H...DV... | | | | 500 |
| human α Ib | 454 | VKASVQ.L.QD.-..PAV.S.V..Q.KTP.....IQM.VG.T.HNI-.Q..SLNA..Q.. | | | | | 511 |
| human α V | 501 | KLKQKGAIRRALFLYSRSPSHSKNMTISRGG | LMQCEELIAYLRDESEFRDKLTPITIFME | | | | 560 |
| mouse α V | 501 |HN...V...T..VF...Q.....V..... | | | | | 560 |
| human α Ib | 512 | RQ.PRQG-..V.L.G.QQAGTTL.LDLGGKHSP | I.HTTM.F....AD.....S..VLSLN | | | | 571 |
| human α V | 561 | YRLDYRTAADTTGLQPILNQFTPANISRQAHIL | | | | | 593 |
| mouse α V | 561 |Q.....A.....V..... | | | | | 593 |
| human α Ib | 571 | VS.PPTE.----.MA.AVVLHGDTHVQE.TR.V | | | | | 599 |





A



B

