

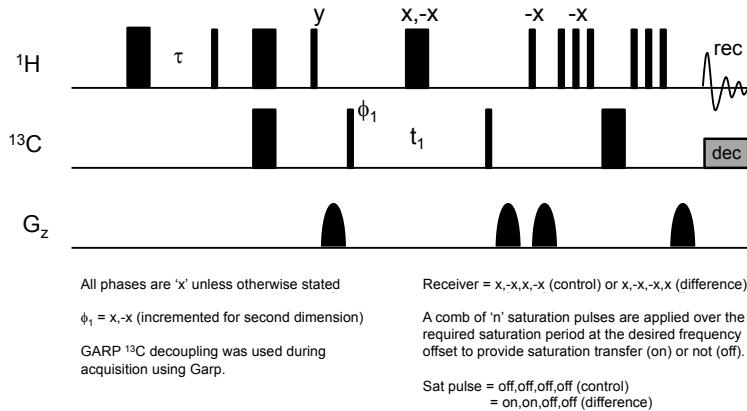
Q2DSTD NMR deciphers epitope-mapping variability for peptide recognition in integrin $\alpha\beta 6$

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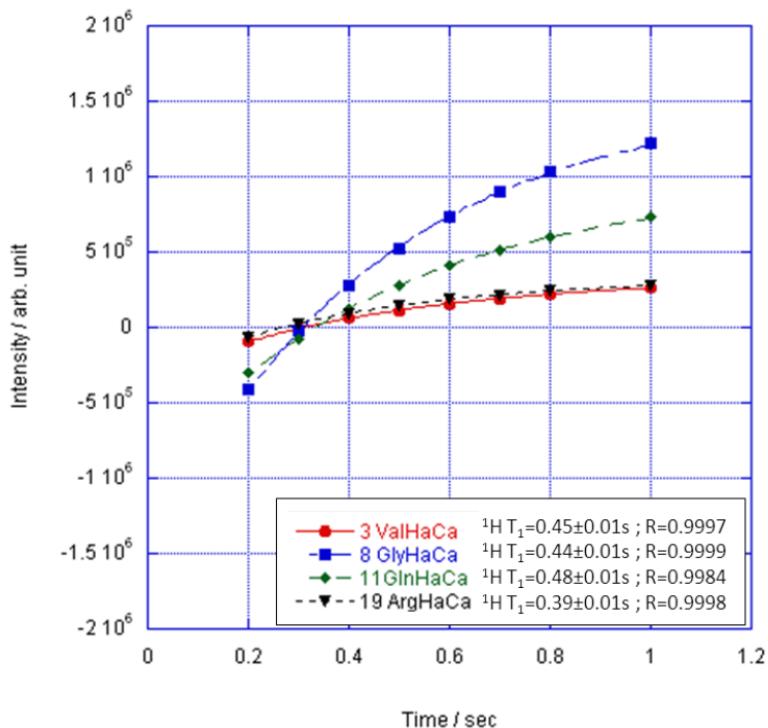
and Mark J. Howard^{*}

SUPPORTING INFORMATION

$^1\text{H}, ^{13}\text{C}$ -HSQC WITH INVERSION RECOVERY



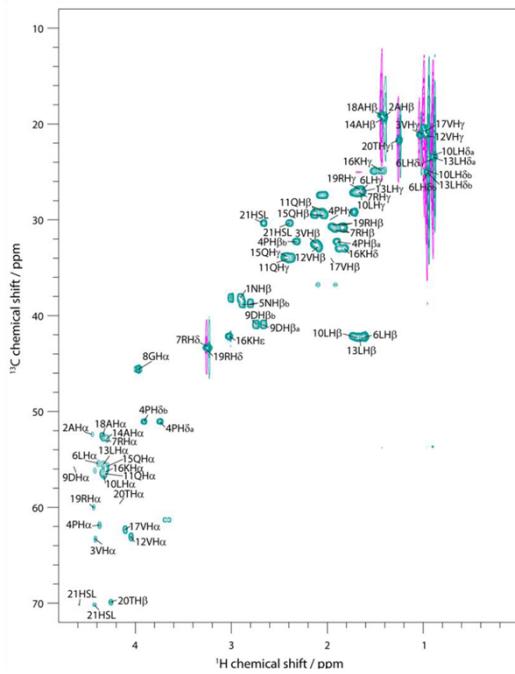
[180- τ -90] unit at the head of the sequence provides the adjustable element to measure T_1 of ^1H attached to ^{13}C . The inter-sequence relaxation delay was set to 5 s and τ values were 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8 and 1.0 s to provide intensities for each ^1H resonance correlated to its ^{13}C in the modified HSQC and example intensity data are shown below for four ^1H - ^{13}C correlations in FMDV2:

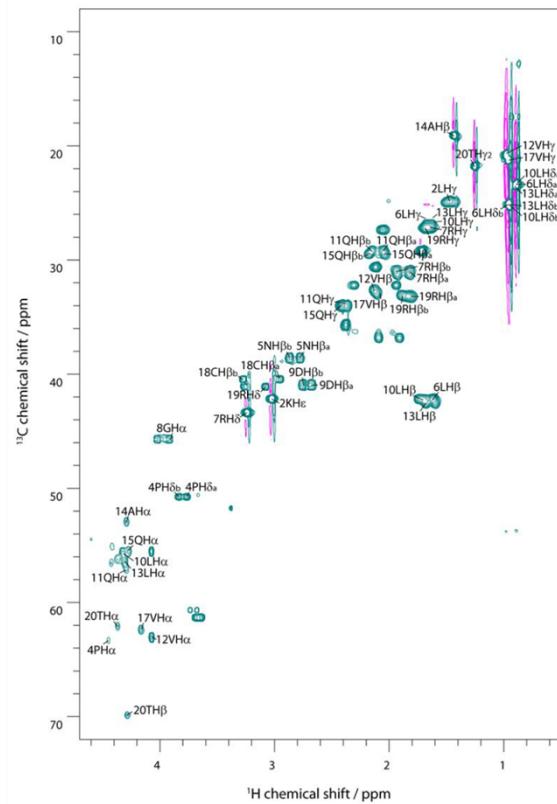


Curves were fitted to: $I_z = I_o [1 - 2e(-t/T_1)]$ where I_z is the measured intensity and t is the tau delay in 180-t-90 delay. Intensities were measured for each 2D dataset obtained with a specific tau value using CCPN Analysis.

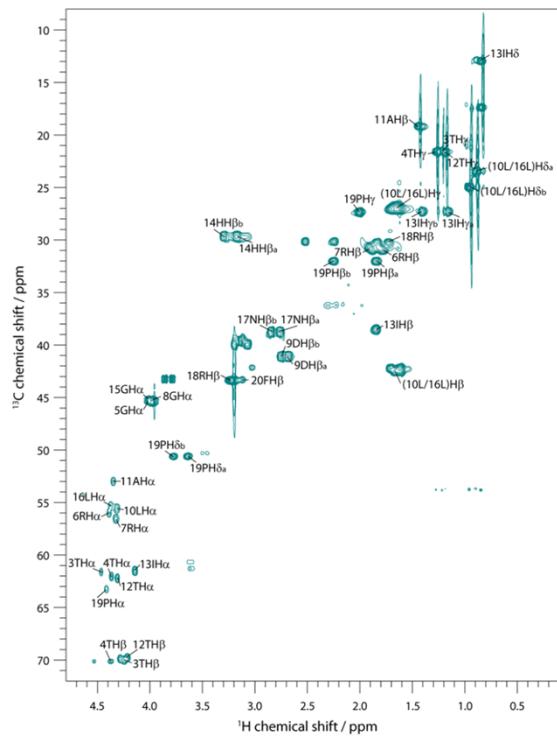
PEPTIDE ^1H , ^{13}C HSQC ASSIGNMENTS

FMDV2





DBD1



LAP2T1

2DSTD, ^1H T_1 and Q2DSTD DATA.

Intensities are in arbitrary units and only for assignable peaks with measurable 2DSTD

| FMDV2 | 2DSTD (arb units) | 2DSTD % | ^1H T_1 (s) | Q2DSTD (arb units) | Q2DSTD % |
|----------|----------------------|---------|-------------------------------|-----------------------|----------|
| 1 AsnHba | | | 0.3358 | | |
| 1 AsnHbb | | | 0.3576 | | |
| 2AlaHa | | | 0.4223 | | |
| 2AlaHb | | | 0.3454 | | |
| 3ValHa | 1.7675 | 28.4 | 0.4503 | 3.9250 | 25.8 |
| 3ValHb | 1.4695 | 23.6 | 0.4070 | 3.6109 | 23.7 |
| 3ValHga | 1.5633 | 25.1 | 0.3826 | 4.0866 | 26.8 |
| 3ValHgb | 1.0405 | 16.7 | 0.3073 | 3.3863 | 22.2 |
| 4ProHa | 0.8140 | 13.1 | 0.4147 | 1.9631 | 12.9 |
| 4ProHb | 0.9490 | 15.3 | 0.3856 | 2.4614 | 16.2 |
| 4ProHb | 0.8269 | 13.3 | 0.3825 | 2.1621 | 14.2 |
| 4ProHd | 0.1112 | 1.8 | 0.3868 | 0.2876 | 1.9 |
| 4ProHd | 2.1186 | 34.1 | 0.3663 | 5.7842 | 38.0 |
| 4ProHg | 1.5850 | 25.5 | 0.3500 | 4.5282 | 29.7 |
| 5AsnHb | 0.7538 | 12.1 | 0.3563 | 2.1155 | 13.9 |
| 5AsnHb | 0.8308 | 13.4 | 0.3505 | 2.3702 | 15.6 |
| 6LeuHa | | | 0.4453 | | |
| 6LeuHb | 2.8360 | 45.6 | 0.3358 | 8.4444 | 55.5 |
| 6LeuHg | 2.5494 | 41.0 | 0.3574 | 7.1334 | 46.8 |
| 6LeuHda | 3.6967 | 59.5 | 0.3800 | 9.7288 | 63.9 |
| 6LeuHdb | 3.1977 | 51.4 | 0.3453 | 9.2612 | 60.8 |
| 7ArgHa | 3.4489 | 55.5 | 0.4720 | 7.3068 | 48.0 |
| 7ArgHb | 1.3923 | 22.4 | 0.3047 | 4.5697 | 30.0 |
| 7ArgHb | 1.4638 | 23.5 | 0.2902 | 5.0436 | 33.1 |
| 7ArgHd | 0.5340 | 8.6 | 0.3183 | 1.6774 | 11.0 |
| 7ArgHg | 1.6649 | 26.8 | 0.3527 | 4.7201 | 31.0 |
| 8GlyHa | 1.7122 | 27.5 | 0.4448 | 3.8491 | 25.3 |
| 9AspHa | | | 0.5460 | | |
| 9AspHb | 1.8762 | 30.2 | 0.3918 | 4.7884 | 31.4 |
| 9AspHb | 2.4276 | 39.0 | 0.3947 | 6.1506 | 40.4 |
| 10LeuHa | 1.7279 | 27.8 | 0.4635 | 3.7277 | 24.5 |
| 10LeuHb | 2.6520 | 42.7 | 0.3266 | 8.1207 | 53.3 |
| 10LeuHg | 2.7561 | 44.3 | 0.3590 | 7.6764 | 50.4 |
| 10LeuHda | 3.6967 | 59.5 | 0.3660 | 10.1013 | 66.3 |
| 10LeuHdb | 3.0376 | 48.9 | 0.3434 | 8.8445 | 58.1 |
| 11GlnHa | 2.0364 | 32.8 | 0.4824 | 4.2218 | 27.7 |
| 11GlnHb | 1.3477 | 21.7 | 0.3206 | 4.2042 | 27.6 |
| 11GlnHb | 1.2536 | 20.2 | 0.3194 | 3.9243 | 25.8 |
| 11GlnHg | 1.6858 | 27.1 | 0.3057 | 5.5147 | 36.2 |
| 12ValHa | 1.4752 | 23.7 | 0.4976 | 2.9644 | 19.5 |
| 12ValHb | 6.2168 | 100.0 | 0.4170 | 14.9084 | 97.9 |
| 12ValHga | 4.4295 | 71.3 | 0.3619 | 12.2400 | 80.4 |

| | | | | | |
|-----------|--------|------|--------|---------|-------|
| 12ValHgb | 3.0588 | 49.2 | 0.3576 | 8.5531 | 56.2 |
| 13LeuHa | 0.8204 | 13.2 | 0.4677 | 1.7543 | 11.5 |
| 13LeuHb | 2.5889 | 41.6 | 0.3393 | 7.6313 | 50.1 |
| 13LeuHg | 5.2711 | 84.8 | 0.3461 | 15.2283 | 100.0 |
| 13LeuHda | 3.6967 | 59.5 | 0.3773 | 9.7969 | 64.3 |
| 13LeuHdb | 3.0376 | 48.9 | 0.3497 | 8.6864 | 57.0 |
| 14AlaHa | 4.1979 | 67.5 | 0.5065 | 8.2879 | 54.4 |
| 14AlaHb | 1.2396 | 19.9 | 0.3277 | 3.7831 | 24.8 |
| 15GlnHa | 3.8847 | 62.5 | 0.5119 | 7.5883 | 49.8 |
| 15GlnHb | 1.1015 | 17.7 | 0.3223 | 3.4176 | 22.4 |
| 15GlnHb | 1.0499 | 16.9 | 0.3166 | 3.3158 | 21.8 |
| 15GlnHg | 0.9749 | 15.7 | 0.3384 | 2.8807 | 18.9 |
| 16LysHa | 0.6526 | 10.5 | 0.4420 | 1.4766 | 9.7 |
| 16LysHd | 2.0962 | 33.7 | 0.3358 | 6.2428 | 41.0 |
| 16LysHe | 0.3440 | 5.5 | 0.3810 | 0.9028 | 5.9 |
| 16LysHg | | | 0.2884 | | |
| 17ValHa | 1.3796 | 22.2 | 0.4532 | 3.0445 | 20.0 |
| 17ValHb | 2.2983 | 37.0 | 0.4014 | 5.7253 | 37.6 |
| 17ValHgb | 1.9290 | 31.0 | 0.3777 | 5.1070 | 33.5 |
| 17ValHga | 2.5163 | 40.5 | 0.3620 | 6.9503 | 45.6 |
| 18AlaHa | 0.4732 | 7.6 | 0.4339 | 1.0907 | 7.2 |
| 18AlaHb | 0.8604 | 13.8 | 0.3330 | 2.5835 | 17.0 |
| 19ArgHa | | | 0.3902 | | |
| 19ArgHb | 2.1983 | 35.4 | 0.3055 | 7.1962 | 47.3 |
| 19ArgHb | 1.2287 | 19.8 | 0.3093 | 3.9724 | 26.1 |
| 19ArgHd | 0.5340 | 8.6 | 0.3025 | 1.7651 | 11.6 |
| 19ArgHg | 2.5494 | 41.0 | 0.3696 | 6.8972 | 45.3 |
| 20ThrHaHa | | | 0.4208 | | |
| 20ThrHb | 1.2814 | 20.6 | 0.4002 | 3.2021 | 21.0 |
| 20ThrHg2 | 0.7073 | 11.4 | 0.3250 | 2.1764 | 14.3 |

| DBD1 | 2DSTD (arb units) | 2DSTD % | ¹ H T ₁ (s) | Q2DSTD (arb units) | Q2DSTD % |
|----------|----------------------|---------|-----------------------------------|-----------------------|----------|
| 2LysHe | 0.6356 | 13.7 | 0.3079 | 2.0641 | 17.2 |
| 2LysHg | 0.5416 | 11.7 | 0.3748 | 1.4451 | 12.0 |
| 4ProHa | 0.5177 | 11.2 | 0.6414 | 0.8072 | 6.7 |
| 4ProHda | 3.6214 | 78.2 | 0.5481 | 6.6072 | 55.0 |
| 4ProHdb | | | 0.5410 | | |
| 5AsnHba | 1.2575 | 27.1 | 0.4269 | 2.9457 | 24.5 |
| 5AsnHbb | 0.4182 | 9.0 | 0.4302 | 0.9721 | 8.1 |
| 6LeuHb | 3.6186 | 78.1 | 0.3861 | 9.3723 | 78.0 |
| 6LeuHda | 3.7871 | 81.7 | 0.3855 | 9.8234 | 81.8 |
| 6LeuHdb | 2.2018 | 47.5 | 0.3491 | 6.3072 | 52.5 |
| 6LeuHg | 2.2122 | 47.7 | 0.4305 | 5.1392 | 42.8 |
| 7ArgHba | 2.0666 | 44.6 | 0.3581 | 5.7711 | 48.0 |
| 7ArgHbb | 1.3528 | 29.2 | 0.3596 | 3.7615 | 31.3 |
| 7ArgHd | 0.8344 | 18.0 | 0.3841 | 2.1724 | 18.1 |
| 7ArgHg | 2.3174 | 50.0 | 0.3275 | 7.0759 | 58.9 |
| 8GlyHa | 3.1384 | 67.7 | 0.5215 | 6.0181 | 50.1 |
| 9AspHba | 2.7270 | 58.8 | 0.4552 | 5.9903 | 49.9 |
| 9AspHbb | 2.5512 | 55.1 | 0.4470 | 5.7081 | 47.5 |
| 10LeuHa | 2.0312 | 43.8 | 0.5454 | 3.7245 | 31.0 |
| 10LeuHb | 4.0946 | 88.4 | 0.3850 | 10.6349 | 88.5 |
| 10LeuHda | 4.6339 | 100.0 | 0.3858 | 12.0126 | 100.0 |
| 10LeuHdb | 3.5080 | 75.7 | 0.3480 | 10.0820 | 83.9 |
| 10LeuHg | 2.4743 | 53.4 | 0.4219 | 5.8654 | 48.8 |
| 11GlnHa | 1.0047 | 21.7 | 0.5429 | 1.8508 | 15.4 |
| 11GlnHba | 2.3655 | 51.0 | 0.3759 | 6.2931 | 52.4 |
| 11GlnHbb | 0.9782 | 21.1 | 0.3756 | 2.6041 | 21.7 |
| 11GlnHg | 1.4832 | 32.0 | 0.3346 | 4.4330 | 36.9 |
| 12ValHa | 4.0880 | 88.2 | 0.5828 | 7.0145 | 58.4 |
| 12ValHb | 3.4079 | 73.5 | 0.4950 | 6.8850 | 57.3 |
| 12ValHga | 3.2130 | 69.3 | 0.3744 | 8.5828 | 71.4 |
| 13LeuHa | 1.4961 | 32.3 | 0.5569 | 2.6866 | 22.4 |
| 13LeuHb | 3.1229 | 67.4 | 0.3925 | 7.9561 | 66.2 |
| 13LeuHda | 3.4468 | 74.4 | 0.3855 | 8.9410 | 74.4 |
| 13LeuHdb | 2.9676 | 64.0 | 0.3488 | 8.5087 | 70.8 |
| 13LeuHg | 2.9533 | 63.7 | 0.4233 | 6.9776 | 58.1 |
| 14AlaHa | 2.1391 | 46.2 | 0.5847 | 3.6581 | 30.5 |
| 14AlaHb | 2.1278 | 45.9 | 0.3455 | 6.1596 | 51.3 |
| 15GlnHa | 1.3481 | 29.1 | 0.4933 | 2.7330 | 22.8 |
| 15GlnHba | 0.4361 | 9.4 | 0.3728 | 1.1697 | 9.7 |
| 15GlnHbb | 1.0740 | 23.2 | 0.3748 | 2.8652 | 23.9 |
| 15GlnHg | 0.8964 | 19.3 | 0.3429 | 2.6143 | 21.8 |
| 17ValHa | 2.7131 | 58.5 | 0.5954 | 4.5566 | 37.9 |
| 17ValHb | | | 0.4754 | | |
| 17ValHga | 2.6573 | 57.3 | 0.3397 | 7.8237 | 65.1 |
| 18CysHba | 2.2176 | 47.9 | 0.5618 | 3.9473 | 32.9 |

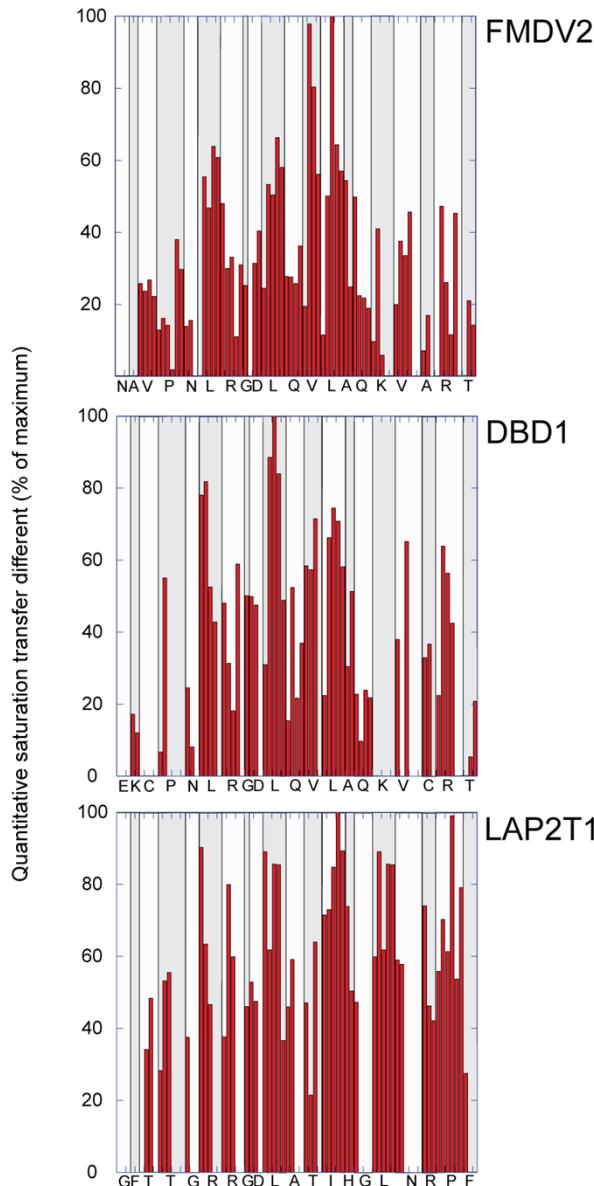
| | | | | | |
|----------|--------|------|--------|--------|------|
| 18CysHbb | 2.5545 | 55.1 | 0.5798 | 4.4059 | 36.7 |
| 19ArgHba | 0.8893 | 19.2 | 0.3307 | 2.6893 | 22.4 |
| 19ArgHbb | 2.6293 | 56.7 | 0.3428 | 7.6704 | 63.9 |
| 19ArgHd | 4.0110 | 86.6 | 0.5928 | 6.7666 | 56.3 |
| 19ArgHg | 1.9662 | 42.4 | 0.3855 | 5.1003 | 42.5 |
| 20ThrHa | | | 0.4708 | | |
| 20ThrHb | 0.2921 | 6.3 | 0.4538 | 0.6438 | 5.4 |
| 20ThrHg2 | 0.8336 | 18.0 | 0.3333 | 2.5008 | 20.8 |

| LAP2T1 | 2DSTD (arb units) | 2DSTD % | ¹ H T ₁ (s) | Q2DSTD (arb units) | Q2DSTD % |
|------------------|----------------------|---------|-----------------------------------|-----------------------|----------|
| 3ThrHa | 0.0000 | | 0.4059 | 0.0000 | |
| 3ThrHb | 1.3159 | 26.9 | 0.3951 | 3.3306 | 34.1 |
| 3ThrHgb | 1.5390 | 31.4 | 0.3261 | 4.7190 | 48.4 |
| 4ThrHa | 1.2231 | 25.0 | 0.4436 | 2.7569 | 28.3 |
| 4ThrHb | 2.1129 | 43.1 | 0.4071 | 5.1902 | 53.2 |
| 4ThrHgb | 1.8330 | 37.4 | 0.3383 | 5.4179 | 55.5 |
| 5GlyHa | 1.3554 | 27.7 | 0.3701 | 3.6619 | 37.5 |
| 6ArgHa | 3.8886 | 79.4 | 0.4416 | 8.8049 | 90.3 |
| 6ArgHb | 1.9230 | 39.3 | 0.3109 | 6.1858 | 63.4 |
| 6ArgHb | 1.4149 | 28.9 | 0.3111 | 4.5487 | 46.6 |
| 7ArgHa | 1.6649 | 34.0 | 0.4534 | 3.6717 | 37.6 |
| 7ArgHb | 2.4377 | 49.8 | 0.3126 | 7.7980 | 79.9 |
| 7ArgHb | 1.8437 | 37.6 | 0.3155 | 5.8431 | 59.9 |
| 8GlyHa | 1.8870 | 38.5 | 0.4200 | 4.4927 | 46.1 |
| 9AspHb | 1.9860 | 40.5 | 0.3851 | 5.1569 | 52.9 |
| 9AspHb | 1.7975 | 36.7 | 0.3879 | 4.6342 | 47.5 |
| (10Leu/16Leu)Hda | 3.3187 | 67.7 | 0.3822 | 8.6839 | 89.0 |
| (10Leu/16Leu)Hg | 1.9889 | 40.6 | 0.3299 | 6.0280 | 61.8 |
| (10Leu/16Leu)Hb | 2.8272 | 57.7 | 0.3385 | 8.3517 | 85.6 |
| (10Leu/16Leu)Hdb | 2.9608 | 60.4 | 0.3552 | 8.3359 | 85.4 |
| 10LeuHa | 1.6649 | 34.0 | 0.4660 | 3.5726 | 36.6 |
| 11AlaHa | 2.2865 | 46.7 | 0.5096 | 4.4866 | 46.0 |
| 11AlaHb | 1.9307 | 39.4 | 0.3346 | 5.7697 | 59.1 |
| 12ThrHa | 2.3102 | 47.2 | 0.5028 | 4.5944 | 47.1 |
| 12ThrHb | 0.9269 | 18.9 | 0.4424 | 2.0953 | 21.5 |
| 12ThrHgb | 2.0515 | 41.9 | 0.3286 | 6.2425 | 64.0 |
| 13IleHa | 3.2749 | 66.8 | 0.4695 | 6.9756 | 71.5 |
| 13IleHb | 3.1528 | 64.4 | 0.4428 | 7.1204 | 73.0 |
| 13IleHda | 4.8993 | 100.0 | 0.5924 | 8.2701 | 84.8 |
| 13IleHga | 3.7093 | 75.7 | 0.3802 | 9.7555 | 100.0 |
| 13IleHgb | 3.2227 | 65.8 | 0.3699 | 8.7135 | 89.3 |
| 14HisHb | 2.7411 | 55.9 | 0.3803 | 7.2072 | 73.9 |
| 14HisHb | 1.9431 | 39.7 | 0.3955 | 4.9132 | 50.4 |
| 15GlyHa | 1.8196 | 37.1 | 0.3945 | 4.6125 | 47.3 |
| 16LeuHa | 2.5773 | 52.6 | 0.4413 | 5.8408 | 59.9 |
| (10Leu/16Leu)Hda | 3.3187 | 67.7 | 0.3822 | 8.6839 | 89.0 |
| (10Leu/16Leu)Hg | 1.9889 | 40.6 | 0.3299 | 6.0280 | 61.8 |
| (10Leu/16Leu)Hb | 2.8272 | 57.7 | 0.3385 | 8.3517 | 85.6 |
| (10Leu/16Leu)Hdb | 2.9608 | 60.4 | 0.3552 | 8.3359 | 85.4 |
| 17AsnHb | 1.9474 | 39.7 | 0.3383 | 5.7558 | 59.0 |
| 17AsnHb | 1.9053 | 38.9 | 0.3380 | 5.6367 | 57.8 |
| 18ArgHb | 2.2601 | 46.1 | 0.3130 | 7.2209 | 74.0 |
| 18ArgHb | 1.4102 | 28.8 | 0.3124 | 4.5148 | 46.3 |
| 18ArgHd | 1.2536 | 25.6 | 0.3052 | 4.1075 | 42.1 |
| 19ProHa | 2.5304 | 51.6 | 0.4645 | 5.4473 | 55.8 |

| | | | | | |
|---------|--------|------|--------|--------|------|
| 19ProHb | 2.7865 | 56.9 | 0.4065 | 6.8546 | 70.3 |
| 19ProHb | 2.5513 | 52.1 | 0.4265 | 5.9817 | 61.3 |
| 19ProHd | 3.7782 | 77.1 | 0.3910 | 9.6628 | 99.0 |
| 19ProHd | 1.9892 | 40.6 | 0.3795 | 5.2413 | 53.7 |
| 19ProHg | 3.5019 | 71.5 | 0.4539 | 7.7156 | 79.1 |
| 20PheHb | 0.7835 | 16.0 | 0.2919 | 2.6840 | 27.5 |

Q2DSTD DATA SHOWN ACROSS EACH PEPTIDE SEQUENCE FOR ALL NUCLEI

Successive residues in sequence shown by alternating white then shaded backgrounds.



Note the general topology of FMDV2 and DBD2 are similar and this is to be expected because they are identical sequences (except for the disulphide cyclisation in DBD2).

However, the cyclisation appears to elevate N-terminal and C-terminal contacts in DBD1 as well as boost RGD-based contacts.

LAP projects significant contacts across the much of the peptide sequence and it is likely these contacts are also used when binding other integrins.

Chemical shifts (ppm) of ¹⁵N/¹³C-A20fmdv2 in PBS

| Residue | N | H ^N | H ^a | Others |
|---------|---------|----------------|----------------|---|
| 1Asn | | | 4.712 | H ^{B2/B3} 2.866, 2.930; C ^B 37.506; H ^{δ21/δ22} 7.046, 7.766; N ^{δ2} 112.751 |
| 2Ala | 118.516 | 8.719 | 4.404 | C ^a 51.388; H ^B 1.427; C ^B 19.381 |
| 3Val | 121.860 | 8.373 | 4.403 | C' 176.451; C ^a 62.460; H ^B 2.105; C ^B 31.634; H ^{γ1/γ2} 0.934, 1.055; C ^{γ1/γ2} 19.239, 20.190 |
| 4Pro | | | | |
| 5Asn | 119.274 | 8.631 | 4.688 | C' 175.560; H ^{B2/B3} 2.813, 2.871; C ^B 37.895; H ^{δ21/δ22} 7.000, 7.708; N ^{δ2} 112.940 |
| 6Leu | 123.657 | 8.402 | 4.363 | C' 174.396; C ^a 54.322; H ^{B2/B3} 1.696; C ^B 41.311; H ^γ 1.594; C ^γ 25.713; H ^{δ1/δ2} 0.892, 0.935; C ^{δ1/δ2} 22.572, 24.085 |
| 7Arg | 121.294 | 8.427 | 4.313 | C' 176.604; C ^a 52.200; H ^{B2/B3} 1.807, 1.931; C ^B 29.964; H ^{γ1/γ2} 1.654; C ^γ 25.790; H ^{δ1/δ2} 3.240; C ^δ 42.295; H ^e 7.471; N ^e 117.442 |
| 8Gly | 109.896 | 8.429 | 3.957 | C' 176.158; C ^a 44.716 |
| 9Asp | 120.457 | 8.392 | 4.584 | C' 173.182; C ^a 53.552; H ^{B2/B3} 2.754; C ^B 39.976 |
| 10Leu | 121.952 | 8.224 | 4.303 | C' 175.904; C ^a 55.541; H ^{B2/B3} 1.706; C ^B 41.303; H ^γ 1.655; C ^γ 26.054; H ^{δ1/δ2} 0.947, 0.996; C ^{δ1/δ2} 22.257, 23.916 |
| 11Gln | 121.142 | 8.374 | 4.289 | C' 176.461; C ^a 55.071; H ^{B2/B3} 2.032, 2.101; C ^B 28.222; H ^{γ1/γ2} 2.375; C ^γ 32.936; H ^{ε21/ε22} 6.937, 7.651; N ^{e2} 112.990 |
| 12Val | 121.838 | 8.146 | 4.024 | C' 175.458; C ^a 62.189; H ^B 2.088; C ^B 31.828; H ^{γ1/γ2} 0.921, 1.014; C ^{γ1/γ2} 19.553, 20.5668 |
| 13Leu | 125.407 | 8.299 | 4.340 | C' 175.770; C ^a 54.228; H ^{B2/B3} 1.682; C ^B 41.279; H ^γ 1.600; C ^γ 25.759; H ^{δ1/δ2} 0.895, 0.959; C ^{δ1/δ2} 22.401, 23.991 |
| 14Ala | 124.435 | 8.256 | 4.282 | C' 176.474; C ^a 51.920; H ^B 1.407; C ^B 18.381 |
| 15Gln | 119.394 | 8.284 | 4.284 | C' 177.137; C ^a 54.693; H ^{B2/B3} 1.997, 2.123; C ^B 28.535; H ^{γ1/γ2} 2.416; C ^γ 32.833; H ^{ε21/ε22} 6.958, 7.631; N ^{e2} 112.926 |
| 16Lys | 123.025 | 8.379 | 4.289 | C' 175.310; C ^a 55.156; H ^{B2/B3} 2.061, 2.365; C ^B 26.397; H ^{γ1/γ2} 1.445; C ^γ 23.877; H ^{δ1/δ2} 1.819; C ^δ 31.972; H ^{e2} 3.032; C ^e 41.159 |
| 17Val | 121.959 | 8.221 | 4.091 | C' 175.666; C ^a 61.405; H ^B 2.084; C ^B 17.969; H ^{γ1/γ2} 0.918, 0.989; C ^{γ1/γ2} 19.558, 20.246 |
| 18Ala | 128.359 | 8.467 | 4.406 | C' 175.041; C ^a 51.471; H ^B 1.412; C ^B 19.252 |
| 19Arg | 121.233 | 8.462 | 4.401 | C' 176.735; C ^a 58.847; H ^{B2/B3} 1.804, 1.919; C ^B 29.617; H ^{γ1/γ2} 1.653; C ^γ 25.306; H ^{δ1/δ2} 3.237; C ^δ 42.605; H ^e 7.243; N ^e 117.559 |
| 20Thr | 115.687 | 8.300 | 4.394 | C' 175.650; C ^a 58.899; H ^B 4.306; C ^B 69.058; H ^{γ1} 1.244; C ^γ 20.678 |
| 21Hsl | 118.587 | 8.745 | 4.735 | C' 173.467; C ^a 69.209; H ^{B2/B3} 2.376, 2.646; C ^B 29.328; H ^{γ1/γ2} 4.411, 4.588 |

Chemical shifts (ppm) of ¹⁵N/¹³C-DBD1 in PBS

| Residue | N | H ^N | H ^a | Others |
|---------|---------|----------------|----------------|--|
| 1Glu | | | | |
| 2Lys | 120.785 | 8.350 | 4.303 | H ^{B2/B3} 1.856, 1.904; H ^{Y1/Y2} 1.452; C ^Y 23.892; H ^{δ1/δ2} 1.716; H ^{ε2} 3.046; C ^ε 41.143 |
| 3Cys | | | | |
| 4Pro | | | 4.461 | H ^{B2/B3} 1.972, 2.325; H ^Y 2.068; H ^{δ1/δ2} 3.786, 3.849; C ^δ 49.656 |
| 5Asn | 117.687 | 8.622 | 4.712 | H ^{B2/B3} 2.796, 2.880; C ^β 37.551; H ^{δ21/δ22} 7.076, 7.723; N ^{δ2} 112.217 |
| 6Leu | 122.229 | 8.321 | 4.438 | C ^α 41.251; H ^{B2/B3} 1.696; C ^β 41.251; H ^Y 1.631; C ^γ 25.384; H ^{δ1/δ2} 0.902, 0.969; C ^{δ1/δ2} 22.357, 24.157 |
| 7Arg | 120.235 | 8.445 | 4.381 | H ^{B2/B3} 1.841, 1.950; C ^β 30.003; H ^{Y1/Y2} 1.688; C ^γ 26.385; H ^{δ1/δ2} 3.271; C ^δ 42.350; H ^ε 7.471; N ^ε 115.791 |
| 8Gly | 109.151 | 8.517 | 3.942 | C ^α 44.612 |
| 9Asp | 119.439 | 8.388 | 4.612 | H ^{B2/B3} 2.693, 2.765; C ^β 39.229 |
| 10Leu | 120.634 | 8.243 | 4.340 | C ^α 54.730; H ^{B2/B3} 1.753; C ^β 41.208; H ^Y 1.638; C ^γ 25.731; H ^{δ1/δ2} 0.903, 0.978; C ^{δ1/δ2} 22.225, 24.090 |
| 11Gln | 119.416 | 8.385 | 4.321 | C ^α 56.098; H ^{B2/B3} 2.066, 2.146; C ^β 28.155; H ^{Y1/Y2} 2.405; H ^{ε21/ε22} 6.964, 7.682; N ^{ε2} 111.927 |
| 12Val | 119.341 | 8.078 | 4.086 | C ^α 62.038; H ^β 2.150; C ^β 31.886; H ^{Y1/Y2} 0.989; C ^{γ1/Y2} 19.847 |
| 13Leu | 123.098 | 8.260 | 4.328 | C ^α 55.135; H ^{B2/B3} 1.726; C ^β 41.338; H ^Y 1.636; C ^γ 25.531; H ^{δ1/δ2} 0.913, 0.982; C ^{δ1/δ2} 22.484, 24.102 |
| 14Ala | 122.829 | 8.207 | 4.306 | C ^α 51.927; H ^β 1.447; C ^β 18.063 |
| 15Gln | 117.523 | 8.275 | 4.306 | C ^α 54.248; H ^{B2/B3} 2.059, 2.179; C ^β 28.552; H ^{Y1/Y2} 2.437; C ^γ 33.222; H ^{ε21/ε22} 7.007, 7.655; N ^{ε2} 111.911 |
| 16Lys | 121.947 | 8.587 | 4.758 | |
| 17Val | 118.487 | 8.124 | 4.176 | C ^α 61.332; H ^β 2.127; C ^β 31.916; H ^{Y1/Y2} 0.984; C ^{γ1} 20.079 |
| 18Cys | 121.045 | 8.899 | 4.786 | H ^{B2/B3} 2.965, 3.282; C ^β 32.195 |
| 19Arg | 122.899 | 8.597 | 4.511 | H ^{B2/B3} 1.842, 1.932; C ^β 32.195; H ^{Y1/Y2} 1.687; C ^γ 26.326 H ^ε 7.307; N ^ε 115.904 |
| 20Thr | 115.075 | 8.383 | 4.380 | C ^α 61.152; H ^β 4.257; C ^β 68.848; H ^{γ1} 1.270; C ^γ 20.750 |
| 21Hsl | 117.642 | 8.768 | 4.726 | H ^{B2/B3} 2.405, 2.267; H ^{Y1/Y2} 4.434, 4.587 |

Chemical shifts (ppm) of ¹⁵N¹³C-A20LAP2T1

| <u>Residue</u> | <u>N</u> | <u>H^N</u> | <u>H^a</u> | <u>Others</u> |
|----------------|----------|----------------------|----------------------|---|
| 1Gly | | | | |
| 2Phe | | | | |
| 3Thr | 115.931 | 8.435 | 4.475 | C ^a 61.606; H ^B 4.255; C ^B 70.151; H ^{γ1} 1.231; C ^γ 21.475 |
| 4Thr | 115.525 | 8.308 | 4.377 | C ^a 62.032; H ^B 4.299; C ^B 69.874; H ^{γ1} 1.282; C ^γ 21.553 |
| 5Gly | 110.541 | 8.561 | 4.022 | C ^a 45.373 |
| 6Arg | 120.226 | 8.394 | 4.398 | C ^a 56.019; H ^{β2/β3} 1.791, 1.888; C ^B 30.986; H ^γ 1.655; H ^{δ1/δ2} 3.230; H ^ε 7.321; N ^ε 116.594 |
| 7Arg | 122.369 | 8.662 | 4.331 | C ^a 56.612; H ^{β2/β3} 1.842, 1.917; C ^B 32.638; H ^{γ1/γ2} 1.693; H ^{δ1/δ2} 3.240; H ^ε 7.405; N ^ε 116.482 |
| 8Gly | 109.453 | 8.603 | 3.976 | C ^a 45.325 |
| 9Asp | 119.548 | 8.324 | 4.637 | H ^{β2/β3} 2.701, 2.755; C ^B 41.067 |
| 10Leu | 121.550 | 8.298 | 4.330 | C ^a 55.472; H ^{β2/β3} 1.707; C ^B 42.416; H ^γ 1.627; C ^γ 26.950; H ^{δ1/δ2} 0.913, 0.971; C ^{δ1/δ2} 23.380, 24.948 |
| 11Ala | 123.034 | 8.374 | 4.359 | C ^a 52.956; H ^B 1.450; C ^B 19.115 |
| 12Thr | 112.477 | 8.108 | 4.319 | C ^a 62.196; H ^B 4.221; C ^B 69.575; H ^{γ1} 1.196; C ^γ 21.643 |
| 13Ile | 121.950 | 8.146 | 4.153 | C ^a 61.460; H ^B 1.862; C ^B 38.495; H ^{γ12/γ13} 1.174, 1.417; C ^γ 27.276; C ^δ 12.907 |
| 14His | 121.910 | 8.524 | 4.730 | H ^{β2/β3} 3.153, 3.266; C ^B 29.621 |
| 15Gly | 109.081 | 8.454 | 3.987 | C ^a 45.102 |
| 16Leu | 120.661 | 8.298 | 4.375 | C ^a 55.259; H ^{β2/β3} 1.700; C ^B 42.416; H ^γ 1.625; C ^γ 26.950; H ^{δ1/δ2} 0.913, 0.971; C ^{δ1/δ2} 23.380, 24.948 |
| 17Asn | 118.394 | 8.614 | 4.715 | H ^{β2/β3} 2.773, 2.852; C ^B 38.726; H ^{δ21/δ22} 7.027, 7.711; N ^{δ2} 118.394 |
| 18Arg | 121.252 | 8.282 | 4.652 | H ^{β2/β3} 1.738, 1.850; C ^B 30.274; H ^{γ1/γ2} 1.668; H ^{δ1/δ2} 3.237; C ^δ 43.334; H ^ε 7.258; N ^ε 116.687 |
| 19Pro | | | 4.425 | C ^a 63.194; H ^{β2/β3} 1.848, 2.267; C ^B 32.020; H ^{γ1/γ2} 2.022; C ^γ 27.324; H ^{δ1/δ2} 3.651, 3.785; C ^δ 50.579 |
| 20Phe | 119.376 | 8.456 | 4.601 | H ^{β2/β3} 3.146; C ^B 43.272 |
| 21Hsl | 117.115 | 8.567 | 4.357 | H ^{β2/β3} 2.264, 2.529; H ^{γ1/γ2} 4.383 |