

Supporting Information

Potent Inhibitors of the Hepatitis C Virus NS3 Protease: Design and Synthesis of Macroyclic Substrate-Based b-Strand Mimics

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General

NMR: ^1H NMR spectra of all synthetic intermediates and products were obtained at 23 °C on a 400 MHz NMR spectrometer and the chemical shifts are given in ppm, referenced to the internal deuterated solvent.

Purification, determination of homogeneity and biological testing of ligands: All compounds evaluated *in vitro* biological assays were purified by semi-preparative C₁₈ reversed-phase HPLC on a Whatman Partisil 10 ODS-3 column using a linear gradient from 5% aqueous CH₃CN (containing 0.06% TFA) to 100% CH₃CN (containing 0.06% TFA) to >98% homogeneity based on HPLC ($\lambda = 220$ nm). IC₅₀ determinations were carried out in the *in vitro* enzymatic assays as previously reported using NS3-NS4A heterodimer protein of genotype 1b.¹⁷ The values reported in Scheme 6 are the average of a minimum of three independent determinations.

L-Allylglycine methyl ester hydrochloride (5b) ^1H NMR (DMSO- d₆), δ 2.59 (dd, $J = 7$ Hz, 2H), 3.73 (s, 3H), 4.12 (dd, $J = 7$ Hz, 1H), 5.21-5.15 (m, 2H), 5.82-5.72 (m, 1H), 8.57 (bs, 3 H). ES⁺ MS *m/z* 130 (M+H)⁺

L-S-Allylcysteine methyl ester hydrochloride (6b) ES⁺ MS *m/z* 176 (M+H)⁺; ES⁻ MS *m/z* 174 (M-H)⁻

(2S,4R)-4-(Quinolin-4-yloxy)-pyrrolidine-1,2-dicarboxylic acid 1-*tert*-butyl ester 2-methyl ester (15a) ^1H NMR (DMSO-d₆) δ 1.33 (s, 9H), 2.32-2.42 (m, 1H), 2.62-2.71 (m, 1H), 3.72 (s, 3H), 3.68-3.79 (m, 2H), 4.09-4.18 (m, 1H), 4.44-4.50 (dd, $J = 8.5$ Hz, 1H), 5.35 (bs, 1H), 7.07 (d, $J = 5.1$ Hz, 1H), 7.56 (dd, $J = 7.5$ Hz, 1H), 7.75 (dd, $J = 7.5$ Hz, 1H), 7.95 (d, $J = 8.4$ Hz, 1H), 8.13 (dd, $J = 7.6$ Hz, 1H), 8.73 (d, $J = 5.1$ Hz, 1H). ES⁺ MS *m/z* 359.6 (MH)⁺.

(2S,4R)-4-(7-Methoxy-quinolin-4-yloxy)-pyrrolidine-1,2-dicarboxylic acid 1-*tert*-butyl ester 2-methyl ester (**15b**) ^1H NMR (CDCl_3 , rotamers in ~2:1 ratio): δ (major rotamer) 1.44 (s, 9H), 2.35-2.42 (m, 1H), 2.65-2.74 (m, 1H), 3.78 (s, 3H), 3.95 (s, 3H), 3.89-3.95 (m, 2H), 4.51 (dd, $J = 8$ Hz, 1H), 5.15 (bs, 1H), 6.55 (m, 1H), 7.1 (dd, $J = 9.1, 2.2$ Hz, 1H), 7.36 (d, $J = 2.6$ Hz, 1H), 8.03 (d, $J = 9.1$ Hz, 1H), 8.65 (d, $J = 5.1$ Hz, 1H). ES $^+$ MS m/z 389.4 (MH^+).

(2S)-*N*-Boc-2-amino-6-heptenoic acid (**20a**) ^1H NMR (CDCl_3) δ 1.46 (s, 9H), 1.45-1.57 (m, 2H), 1.62-1.73 (m, 1H), 1.82-1.92 (m, 1H), 2.05-2.12 (m, 2H), 4.32 (bs, 1H), 4.98 (dm, $J = 10.2$ Hz, 1H), 5.10 (dm, $J = 17.0$ Hz, 1H), 5.66-5.84 (tdd, $J = 6.7, 10.2, 17.0$ Hz, 1H). ES $^+$ MS m/z 244.2 ($\text{M}+\text{H}^+$).

Dipeptides **21a,b**. The hydrochloride salts of the *D*-allylglycine methyl ester was coupled to the *L*-Boc-proline fragment **15c** following the standard protocol. In each case, the product was purified by flash column chromatography (using 50% EtOAc in hexane as the eluent) to obtain the corresponding dipeptide as a white solid in ~90-95% yield.

Dipeptide **21a** ^1H NMR ($\text{DMSO}-d_6$), mixture of rotamers in 2:1 ratio: δ (major rotamer) 1.32 (s, 9H), 2.25-2.26 (m, 1H), 2.40-2.30 (m, 1H), 2.43-2.50 (m, 1H), 2.65-2.50 (m, 1H), 3.62 (s, 3H), 3.82-3.70 (m, 2H), 3.92 (s, 3H), 4.48-4.38 (m, 2H), 5.04 (d, $J = 11.0$ Hz, 1H), 5.39 (d, $J = 17.0$ Hz, 1H), 5.53 (br s, 1H), 5.78-5.65 (m, 1H), 7.17 (dd, $J = 3.0, 9.0$ Hz, 1H), 7.39 (d, $J = 2.0$ Hz, 1H), 7.44 (s, 1H), 7.57-7.46 (m, 3H), 7.95-7.92 (m, 1H), 8.32-8.24 (m, 2H), 8.35 (d, $J = 9.0$ Hz, 1H). ES $^+$ MS m/z 576 ($\text{M}+\text{H}^+$); ES $^-$ MS m/z 574 ($\text{M}-\text{H}^-$)

Dipeptide **21b** ^1H NMR ($\text{DMSO}-d_6$), mixture of rotamers in 2:1 ratio: δ (major rotamer) 1.34 (s, 9H), 2.17-2.28 (m, 1H), 2.48-2.36 (m, 2H), 2.61-2.52 (m, 1H), 3.63 (s, 3H), 3.75 (dd, $J = 4.0$,

12.0 Hz, 1H), 3.80 (d, $J = 12$ Hz, 1H), 3.93 (s, 3H), 4.38-4.30 (m, 1H), 4.51-4.40 (m, 1H), 5.05 (d, $J = 10.0$ Hz, 1H), 5.13 (dd, $J = 2.0, 16.0$ Hz, 1H), 5.53 (bs, 1H), 5.84-5.72 (m, 1H), 7.18, (dd, $J = 3.0, 10.0$ Hz, 1H), 7.40 (d, $J = 3$ Hz, 1H), 7.46 (s, 1H), 7.57-7.47 (m, 3H), 7.95 (d, $J = 10.0$ Hz, 1H), 8.27 (d, $J = 7.0$ Hz, 2H), 8.35 (d, $J = 7.0$ Hz, NH). ES⁺ MS m/z 576 (M+H)⁺; ES⁻ MS m/z 574 (M-H)⁻

Dipeptides 22a,b. The hydrochloride salts of **6a** and **6b** were independently coupled to the *L*-Boc-proline fragment **15b** following the standard protocol. In each case, the product was purified by flash column chromatography (using a solvent gradient from 3% to 8% MeOH in EtOAc) to isolate each of the dipeptides **22a** and **22b** as colorless oils (~85% yield).

Dipeptide 22a ¹H NMR (CDCl₃), mixture of rotamers in ~1:5 ratio: d (major rotamer) 1.46 (s, 9H), ~2.4 (br 1H), 2.86-2.91 (dd, $J = 6.0, 14.0$ Hz, 1H), 2.95-3.00 (m, 2H), 3.12-3.18 (m, 2H), 3.75-3.84 (m, 4H), 3.84-3.93 (m, 1H), 3.94 (s, 3H), 4.54-4.57 (m, 1H), 4.80 (bs, 1H), 5.05-5.22 (m, 3H), 5.71-5.81 (m, 1H), 6.64 (bs, 1H), 7.12-7.14 (dd, $J = 2.6, 9.2$ Hz, 1H), 7.36 (d, $J = 2.6$ Hz, 1H), 7.66 (bs, 1H), 7.99 (d, $J = 9.2$ Hz, 1H), 8.66 (d, $J = 5.1$ Hz, 1H). ES⁺ MS m/z 546.3 (M+H)⁺; ES⁻ MS m/z 544.2 (M-H)⁻.

Dipeptide 22b ¹H NMR (CDCl₃), mixture of rotamers in ~1:5 ratio: d (major rotamer) 1.49 (s, 9H), ~2.4 (br 1H), 2.80 (m, 1H), 2.86-2.91 (dd, $J = 6.0, 14.0$ Hz, 1H), 2.95-3.00 (m, 1H), 3.12-3.18 (m, 2H), 3.75-3.84 (m, 5H), 3.94 (s, 3H), ~4.65 (bs, 1H), 4.80 (bs, 1H), 5.05-5.22 (m, 3H), 5.71-5.81 (m, 1H), 6.64 (bs, 1H), 7.12-7.14 (dd, $J = 2.6, 9.2$ Hz, 1H), 7.36 (d, $J = 2.6$ Hz, 1H), 7.66 (bs, 1H), 7.99 (d, $J = 9.2$ Hz, 1H), 8.66 (d, $J = 5.1$ Hz, 1H). ES⁺ MS m/z 546.3 (M+H)⁺; ES⁻ MS m/z 544.2 (M-H)⁻.

Compounds 23b ^1H NMR (CDCl_3) mixture of rotamers (~1:9 ratio): δ (major rotamer) 1.15-1.28 (m, 1H), 1.42 (s, 18H), 1.43-1.75 (m, 4H), 2.26-2.20 (m, 2H), 2.38 (br, 1H), 2.92 (br, 1H), 3.69-3.75 (m, 2H), 3.94 (s, 3H), 4.52 (br, 1H), 4.95 (d, $J = 10.2$ Hz, 1H), 5.00 (d, $J = 17.2$ Hz, 1H), 5.29 (br, 1H), 5.74-5.83 (m, 1H), 7.03 (br, 1H), 7.08 (dd, $J = 2.3, 9.2$ Hz, 1H), 7.42-7.51 (m, 5H), 7.95 (d, $J = 9.2$ Hz, 1H), 8.03 (bd, $J = 7$ Hz, 2H). ES $^+$ MS m/z 658.3 ($\text{M}+\text{H})^+$; ES $^-$ MS m/z 656.2 ($\text{M}-\text{H})^-$.

Dipeptide methyl ester 25b. The *N*-Boc and *t*-butyl ester moieties of compound **23b** were simultaneously hydrolysed using the same procedure as described above for **23a**. In this case, the free acid was directly converted to the corresponding proline hydrochloride salt methyl ester **25b**, which was used directly in the synthesis of tripeptide dienes without purification. **25b**HCl salt ^1H NMR (CDCl_3) mixture of rotamers in (~1:9 ratio): δ (major rotamer) 1.28-1.34 (m, 1H), 1.49-1.66 (m, 4H), 2.02-2.15 (m, 2H), 2.27-2.35 (m, 1H), 2.81-2.89 (m, 1H), 3.46-3.50 (m, 1H), 3.62 (s, 3H), 3.67-3.78 (m, 1H), 3.78-3.88 (m, 1H), 3.99 (s, 3H), 4.49-4.62 (m, 1H), 4.94 (dd, $J = 1.0, 10.2$ Hz, 1H), 5.02 (dd, $J = 1.4, 18.8$ Hz, 1H), 5.80-5.92 (m, 1H), 5.99 (bs, 1H), 7.44 (d, $J = 8.9$ Hz, 1H), 7.65-7.76 (m, 4H), 8.07 (bs, 1H), 8.27 (bd, $J = 7.3$ Hz, 2H), 8.57 (d, $J = 8.9$ Hz, 1H), 9.03 (bs, 1H). ES $^+$ MS m/z 516.2 ($\text{M}+\text{H})^+$

Dipeptide 27 Peptide coupling between the enantiomerically enriched (>98% ee)^{37b} ethyl ester of vinyl ACCA (1*R*, 2*S*)-**9** and the P2 building block **15f**, under the usual peptide coupling conditions, led to the isolation of the corresponding *N*-Boc-protected dipeptide **27**. ^1H NMR (DMSO-d_6) mixture of rotamers in (~1:15 ratio): δ (major rotamer) 1.16 (t, $J = 7.3$ Hz, 3H), 1.28-1.30 (m, 1H), 1.35 (s, 9H), 1.65-1.69 (m, 1H), 2.13-2.18 (m, 1H), 2.25-2.34 (m, 1H), 2.57-

2.65 (m, 1H), 3.80 (br, 2H), 3.93 (s, 3H), 4.00-4.09 (q, $J = 7.3$ Hz, 2H), 4.27-4.33 (m, 1H), 5.11 (d, $J = 8.9$ Hz, 1H), 5.27 (d, $J = 16.2$ Hz, 1H), 5.56 (br, 1H), 5.63-5.73 (m, 1H), 7.18 (dd, $J = 2.3$, 8.9 Hz, 1H), 7.39 (d, $J = 2.3$ Hz, 1H), 7.46 (s, 1H), 7.49-7.56 (m, 3H), 7.94 (d, $J = 8.9$, 1H), 8.27 (d, $J = 7.0$ Hz, 2H), 8.74 (s, 1H). ES⁺ MS m/z 602.3 (M+H)⁺; ES⁻ MS m/z 600.2 (M-H)⁻.

Diene 28a ¹H NMR (CDCl_3) δ 1.40 (s, 9H), 1.25-1.45 (m, 6H), 1.60-1.69 (m, 1H), 1.70-1.80 (m, 1H), 1.95-2.06 (m, 2H), 2.35-2.47 (m, 1H), 2.47-2.55 (m, 1H), 2.55-2.65 (m, 1H), 2.92-3.00 (m, 1H), 3.72 (s, 3H), 3.96 (s, 3H), 3.94-3.98 (m, 1H), 4.31 (bd, $J = 12$ Hz, 1H), 4.45-4.52 (m, 1H), 4.62 (dd, $J = 12$, 7 Hz, 1H), 4.89-4.93 (m, 1H), 4.92 (d, $J = 10$ Hz, 1H), 4.97 (dd, $J = 2$, 17 Hz, 1H), 5.12-5.19 (m, 2H), 5.40 (bs, 1H), 5.63-5.85 (m, 2H), 7.04-7.11 (m, 2H), 7.39 (bd, $J = 8$ Hz, 1H), 7.43-7.54 (m, 5H), 8.02-8.09 (m, 3H). ES⁺ MS m/z 729.5 (M+H)⁺; ES⁻ MS m/z 727.3 (M-H)⁻.

Diene 28b ¹H NMR (CDCl_3), δ 1.39 (s, 9H), 1.22-1.41 (m, 6H), 1.55-1.62 (m, 1H), 1.68-1.75 (m, 1H), 2.00-2.05 (m, 2H), 2.44-2.61 (m, 3H), 2.84-2.92 (m, 1H), 3.74 (s, 3H), 3.96 (s, 3H), 3.96-4.00-(m, 1H), 4.30-4.35 (m, 1H), 4.43-4.50 (m, 1H), 4.61 (dd, $J = 8$ Hz, 1H), 4.80-4.85 (m, 1H), 4.91-5.01- (m, 2H), 5.12-5.18 (m, 2H), 5.41 (bs, 1H), 5.64-5.83 (m, 2H), 7.04-7.11 (m, 3H), 7.43-7.55 (m, 5H), 8.01-8.07 (m, 3H). ES⁺ MS m/z 729.5 (M+H)⁺; ES⁻ MS m/z 727.3 (M-H)⁻.

Diene 29a ¹H NMR (CDCl_3): δ 1.38 (s, 9H), 1.48-1.68 (m, 5H), 1.75-1.85 (m, 1H), 2.05-2.13 (m, 1H), 2.33-2.42 (m, 1H), 2.90-3.00 (m, 2H), 3.15-3.18 (m, 2H), 3.75 (s, 3H), 3.94 (s, 3H), ~3.95 (bs, 1H), 4.26 (d, $J = 11$ Hz, 1H), 4.45-4.52 (m, 1H), 4.70-4.75 (m, 1H), 4.91-5.04 (m, 3H), 5.11-5.20 (m, 3H), 5.28-5.35 (m, 1H), 5.71-5.83 (m, 2H), 6.65 (d, $J = 5.4$ Hz, 1H), 7.12 (dd,

$J = 2.5$, 9.2 Hz, 1H), 7.36 (d, $J = 2.2$ Hz, 1H), 7.56 (d, $J = 6.7$ Hz, 1H), 8.03 (d, $J = 9.2$ Hz, 1H), 8.67 (d, $J = 5.1$ Hz, 1H). ES⁺ MS m/z 671.3 (M+H)⁺; ES⁻ MS m/z 669.3 (M-H)⁻.

Diene 29b ¹H NMR (CDCl₃): δ 1.38 (s, 9H), 1.49-1.64 (m, 5H), 1.75-1.82 (m, 1H), 2.05-2.12 (m, 2H), 2.35-2.45 (m, 1H), 2.92-2.98 (m, 1H), 3.11-3.19 (m, 2H), 3.77 (s, 3H), 3.94 (s, 3H), ~3.98 (bs, 1H), 4.28 (d, $J = 9.5$ Hz, 1H), 4.45-4.52 (m, 1H), 4.70-4.75 (m, 1H), 4.82-4.90 (m, 1H), 4.96-5.05 (m, 2H), 5.12-5.17 (m, 3H), 5.28-5.35 (m, 1H), 5.71-5.83 (m, 2H), 6.64 (d, $J = 5.2$ Hz, 1H), 7.12 (dd, $J = 2.5$, 9.2 Hz, 1H), 7.36 (m, 2H), 8.03 (d, $J = 9.0$ Hz, 1H), 8.67 (d, $J = 5.3$ Hz, 1H). ES⁺ MS m/z 671.3 (M+H)⁺; ES⁻ MS m/z 669.3 (M-H)⁻.

Ligand 29c ¹H NMR (DMSO-*d*₆): δ 1.17 (s, 9H), 1.32-1.64 (m, 4H), 1.96-2.05 (m, 2H), 2.25-2.35 (m, 1H), 2.50-2.54 (m, 2H), 2.85 (dd, $J = 4.8$, 13.6 Hz, 1H), 3.15 (d, $J = 7.0$ Hz, 2H), 3.90 (bd, $J = 10.1$ Hz, 1H), 3.97 (s, 3H), 4.08-4.15 (m, 1H), 4.41-4.46 (m, 1H), 4.51 (d, $J = 12.1$ Hz, 1H), 4.65 (dd, $J = 8.3$ Hz, 1H), 4.94-5.14 (m, 4H), 5.63 (bs, 1H), 5.69-5.83 (m, 2H), 7.12 (d, $J = 6.8$ Hz, 1H), 7.32 (d, $J = 8.9$ Hz, 1H), 7.42 (bs, 2H), 8.26 (d, $J = 8.3$ Hz, 1H), 8.29 (d, $J = 9.1$ Hz, 1H), 9.02 (d, $J = 5.7$ Hz, 1H). FAB HRMS m/z found: 657.295780 (M+H)⁺; calculated for C₃₃H₄₅N₄O₈S: 657.295812

Ligand 29d ¹H NMR (DMSO-*d*₆): δ 1.19 (s, 9H), 1.40-1.65 (m, 4H), 1.95-2.05 (m, 2H), 2.25-2.35 (m, 2H), 2.65-2.82 (m, 2H), 3.15-3.25 (m, 2H), 3.85-3.95 (m, 1H), 3.97 (s, 3H), 4.05-4.15 (m, 1H), 4.37-4.42 (dd, $J = 6.8$, 13.5 Hz, 1H), 4.5 (d, $J = 11$ Hz, 1H), 4.66 (dd, $J = 8.8$ Hz, 1H), 4.95-5.20 (m, 4H), 5.61 (br, 1H), 5.69-5.80 (m, 2H), 7.13 (d, $J = 7.7$ Hz, 1H), 7.28-7.35 (m, 1H), 7.35-7.45 (m, 1H), 7.41 (s, 1H), 8.27 (d, $J = 6.4$ Hz, 1H), 8.29 (d, $J = 9.0$ Hz, 1H), 8.99 (br, 1H),

12.8 (br, 1H). FAB HRMS m/z found: 657.295780 ($M+H$) $^+$; calculated for $C_{33}H_{45}N_4O_8S$: 657.295812

Diene 30c 1H NMR ($CDCl_3$) rotamers in ~10:1 ratio): δ (major rotamer) 1.41 (s, 9H), 1.48-1.60 (m, 5H), 1.60-1.72 (m, 4H), 2.05-2.20 (m, 4H), 2.35-2.45 (m, 1H), 2.95-3.55 (m, 1H), 3.67 (s, 3H), 3.73-3.83 (m, 1H), 3.92-3.98 (m, 1H), 3.96 (s, 3H), 4.25-4.32 (m, 1H) 4.45-4.55 (m, 1H), 4.78-4.85 (m, 1H), 4.93-5.06 (m, 4H), 5.12-5.30 (m, 1H), 5.43 (bs, 1H), 5.73-5.85 (m, 2H), 7.03-7.12 (m, 2H), 7.43-7.54 (m, 6H), 8.01-8.07 (m, 3H). ES $^+$ MS m/z 741.4 ($M+H$) $^+$; ES $^-$ MS m/z 739.3 ($M-H$) $^-$

Dienes 31 1H NMR ($CDCl_3$, mixture of rotamers in ~1:10 ratio): δ (major rotamer) 1.22 (t, $J = 7.3$ Hz, 3H), 1.30-1.50 (m, 7H), 1.41 (s, 9H), 1.56-1.64 (m, 1H), 1.72-1.81 (m, 1H), 1.85-1.91 (m, 1H), 1.98- 2.05 (m, 2H), 2.15 (dd, $J = 8.5, 17.4$ Hz, 1H), 2.36-2.45 (m, 1H), 2.93-3.02 (m, 1H), 3.91-3.95 (m, 1H), 3.96 (s, 3H), 4.16 (q, $J = 7.3$ Hz, 2H), 4.30 (bd, $J = 10.5$ Hz, 1H), 4.41-4.48 (m, 1H), 4.82-4.87 (m, 1H), 4.88-5.00 (m, 2H), 5.07-5.15 (m, 2H), 5.27-5.34 (m, 1H), 5.44 (bs, 1H), 5.70-5.80 (m, 2H), 7.05-7.13 (m, 2H), 7.43-7.58 (m, 5H), 8.00-8.09 (m, 3H). ES $^+$ MS m/z 755.3 ($M+H$) $^+$; ES $^-$ MS m/z 753.3 ($M-H$) $^-$.

Macrocyclic inhibitor 32b 1H NMR ($DMSO-d_6$) δ 1.10 (s, 9H), 1.16-1.26 (m, 1H), 1.42-1.60 (m, 2H), 1.68-1.82 (m, 2H), 2.08-2.15 (m, 1H), 2.43- ~2.5 (m, 3H), 2.67 (dd, $J = 10.5, 13.4$ Hz, 1H), 3.00 (dd, $J = 7.0, 14.1$ Hz, 1H), 3.10 (dd, $J = 8.0, 14.2$ Hz, 1H), 3.87 (bd, $J = 8.9$ Hz, 1H), 3.97 (s, 3H), 3.90-4.03 (m, 1H), 4.28 (ddd, $J = 3.2, 8.6, 10.5$ Hz, 1H), 4.60 (bd, $J = 12.1$ Hz, 1H), 4.75 (dd, $J = 8.0$ Hz, 1H), 5.18-5.25 (m, 1H), 5.48-5.53 (m, 1H), 5.7 (bs, 1H), 7.13 (d, $J =$

5.4 Hz, 1H), 7.32 (dd, $J = 2.7$, 9.2 Hz, 1H), 7.40 (d, $J = 2.7$ Hz, 1H), 7.45 (d, $J = 6.7$ Hz, 1H), 8.29 (d, $J = 9.2$ Hz, 1H), 8.51 (d, $J = 8.9$ Hz, 1H), 9.04 (d, $J = 6.7$ Hz, 1H). FAB HRMS m/z found ($M+H$)⁺ 629.264410; calculated for C₃₁H₄₁N₄O₈S₁ 629.264512

Methyl ester 39a After RCM of diene **28a** under the usual reaction conditions, the mixture of methyl ester olefins obtained (**38**, Z:E ration of 1:1) were reduced under catalytic hydrogenation conditions to give the saturated ester **39a**. ¹H NMR (CDCl₃) δ 1.33 (s, 9H), 1.23-1.41 (m, 7H), 1.55-1.72 (m, 4H), 1.80-2.10 (m, 3H), 2.47-2.57 (m, 1H), 2.92-3.04 (m, 1H), 3.72 (s, 3H), 3.72-3.75 (m, 1H), 4.09 (s, 3H), 4.14-4.25 (m, 1H), 4.33-4.41 (m, 1H), 4.49-4.62 (m, 2H), 5.00 (bs, 1H), 5.17 (d, $J = 8$ Hz, 1H), 5.61-5.74 (m, 1H), 6.80-7.20 (m, 2H), 7.23 (bs, 1H), 7.57 (bs, 4H), 8.11 (d, $J = 9$ Hz, 1H), 8.26 (bs, 2H), 8.97 (s, 1H). ES⁺ MS m/z 703 (M+H)⁺; ES⁻ MS m/z 701 (M-H)⁻.

Methyl ester 41a After RCM of diene **28b** under the usual reaction conditions, the mixture of methyl ester olefins obtained (**40**, Z:E ration of 1:14) were reduced under catalytic hydrogenation conditions to give the saturated ester **41a**. ¹H NMR (CDCl₃) δ 1.23-1.38 (m, 4H), 1.40 (s, 9H), 1.43-1.48 (m, 2H), 1.53-1.62 (m, 1H), 1.90-1.99 (m, 1H), 2.00-2.20 (m, 2H), 2.3-2.43 (m, 2H), 2.63-2.72 (m, 1H), 3.06-3.15 (m, 1H), 3.76 (s, 3H), 3.87-3.94 (m, 1H), 3.95 (s, 3H), 4.28 (bd, $J = 10.0$ Hz, 1H), 4.52-4.64 (m, 2H), 5.02 (dd, $J = 5.1$, 7.3 Hz, 1H), 5.23-5.53 (m, 4H), 7.07 (s, 1H), 7.10 (dd, $J = 2.6$, 8.9 Hz, 1H), 7.44 (d, $J = 2.4$ Hz, 1H), 7.44-7.55 (m, 3H), 7.68 (d, $J = 7.9$ Hz, 1H), 8.00 (d, $J = 8.5$ Hz, 1H), 8.06 (d, $J = 8.0$ Hz, 2H). ES⁺ MS m/z 703.4 (M+H)⁺; ES⁻ MS m/z 701.4 (M-H)⁻

Macrocyclic inhibitor **41b** The saturated macrocyclic methyl ester **41a** was saponified and purified by C₁₈ reversed phase HPLC to afford compound **41b** as a white solid (>99% homogeneity by HPLC). ¹H NMR (DMSO-*d*₆), δ 1.01-1.55 (m, 13H), 1.53 (s, 9H), 1.60-1.65 (m, 1H), 1.70-1.90 (m, 2H), 2.50-2.55 (m, 2H), 3.93-3.96 (m, 1H), 3.96 (s, 3H), 4.12 (m, 1H), 4.42-4.47 (m, 1H), 4.60 (bd, *J*= 13 Hz, 1H), 4.78 (t, *J*= 8 Hz, 1H), 5.77 (br s, 1H), 7.05 (d, *J*= 7 Hz, 1H), 7.23 (bs, 1H), 7.51 (bs, 1H), 7.59-7.70 (m, 4H), 8.15-8.30 (m, 3H), 8.33 (d, *J*= 9 Hz, 1H). FAB HRMS *m/z* found: 689.355260 (M+H)⁺; calculated for C₃₈H₄₉N₄O₈: 689.355040.

Ligand **42** ¹H NMR (DMSO-*d*₆, mixture of rotamers in ~1:10 ratio): δ (major rotamer) 0.86 (2t, *J*= 6.6 , 7.2 Hz, 6H), 1.22 (s, 9H), 1.30-1.40 (m, 9H), 1.40-1.50 (m, 1H), 1.50-1.60 (m, 1H), 1.60-1.69 (m, 1H), 2.20-2.30 (m, 1H), 2.48-2.60 (m, 1H), 3.87 (d, *J*= 9.7 Hz, 1H), 3.96 (s, 3H), 4.14-4.18 (m, 2H), 4.49 (brd, *J*= 11.4 Hz, 1H), 4.61 (dd, *J*= 8.3 Hz, 1H), 5.72 (brs, 1H), 7.07 (d, *J*= 7.0 Hz, 1H), 7.20 (br, 1H), 7.48 (s, 1H), 7.63 (br, 4H), 8.12 (d, *J*= 7.7 Hz, 1H), 8.20 (br, 3H). FAB HRMS *m/z* found (M+H)⁺ 691.371000; calculated for C₃₈H₅₁N₄O₈ 691.370690

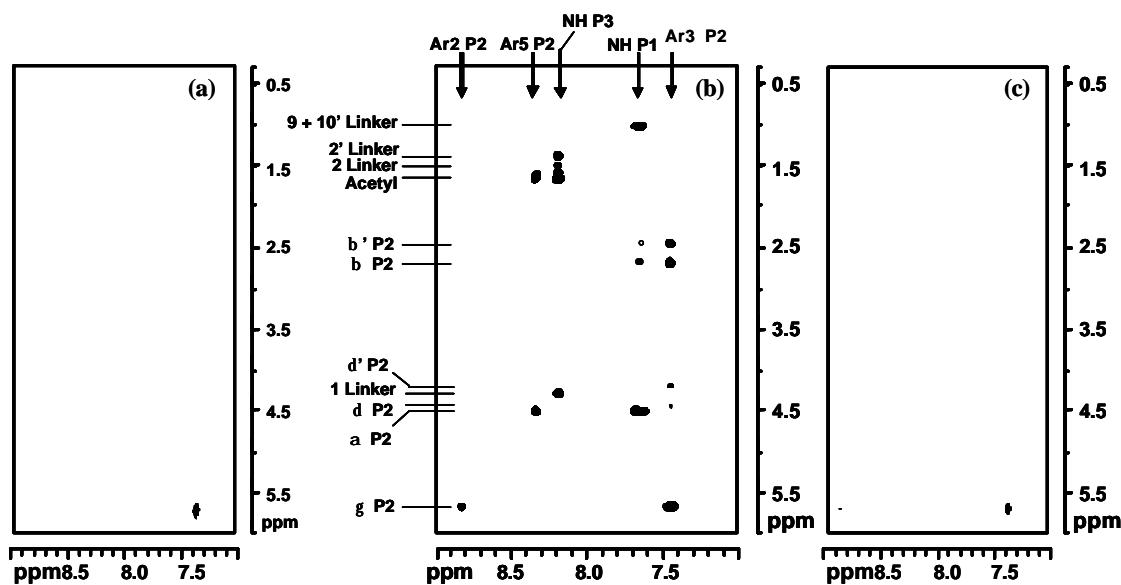
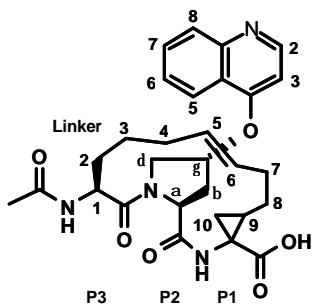
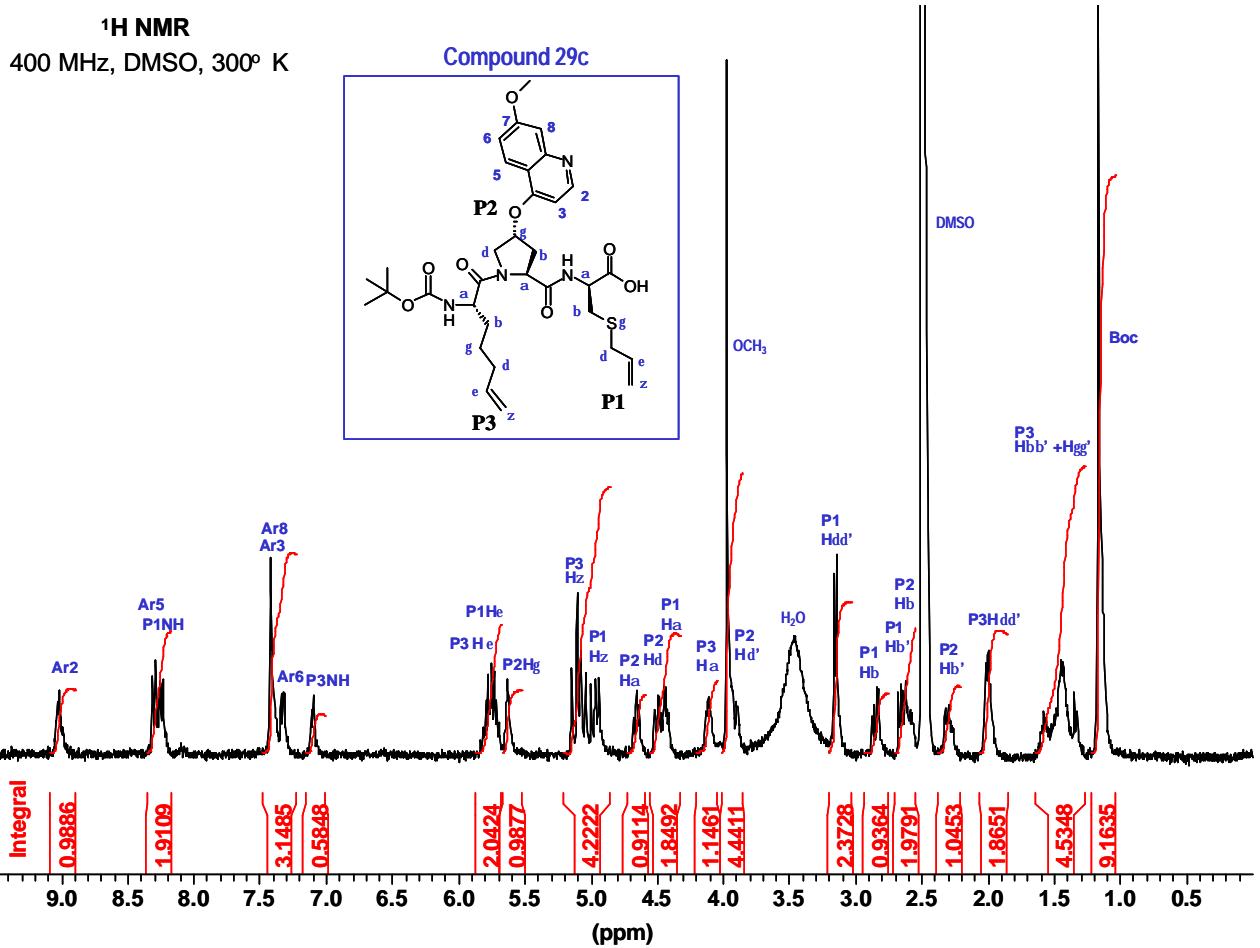
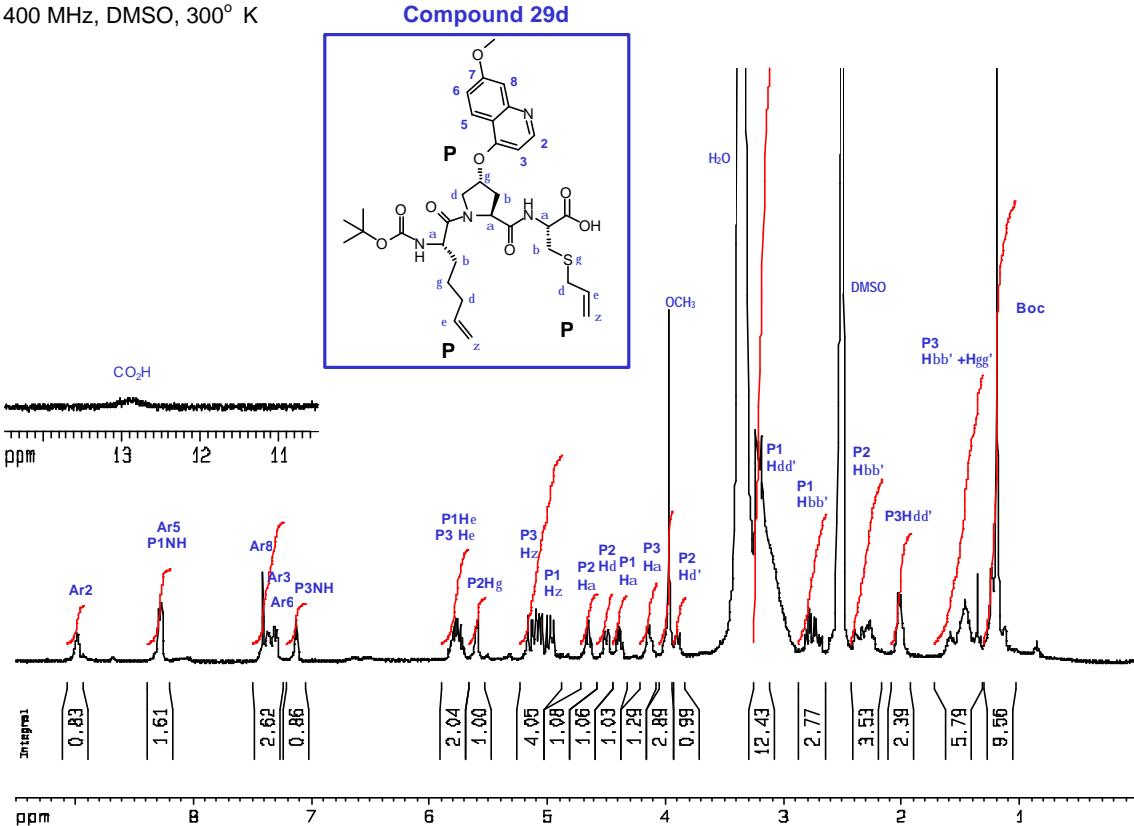


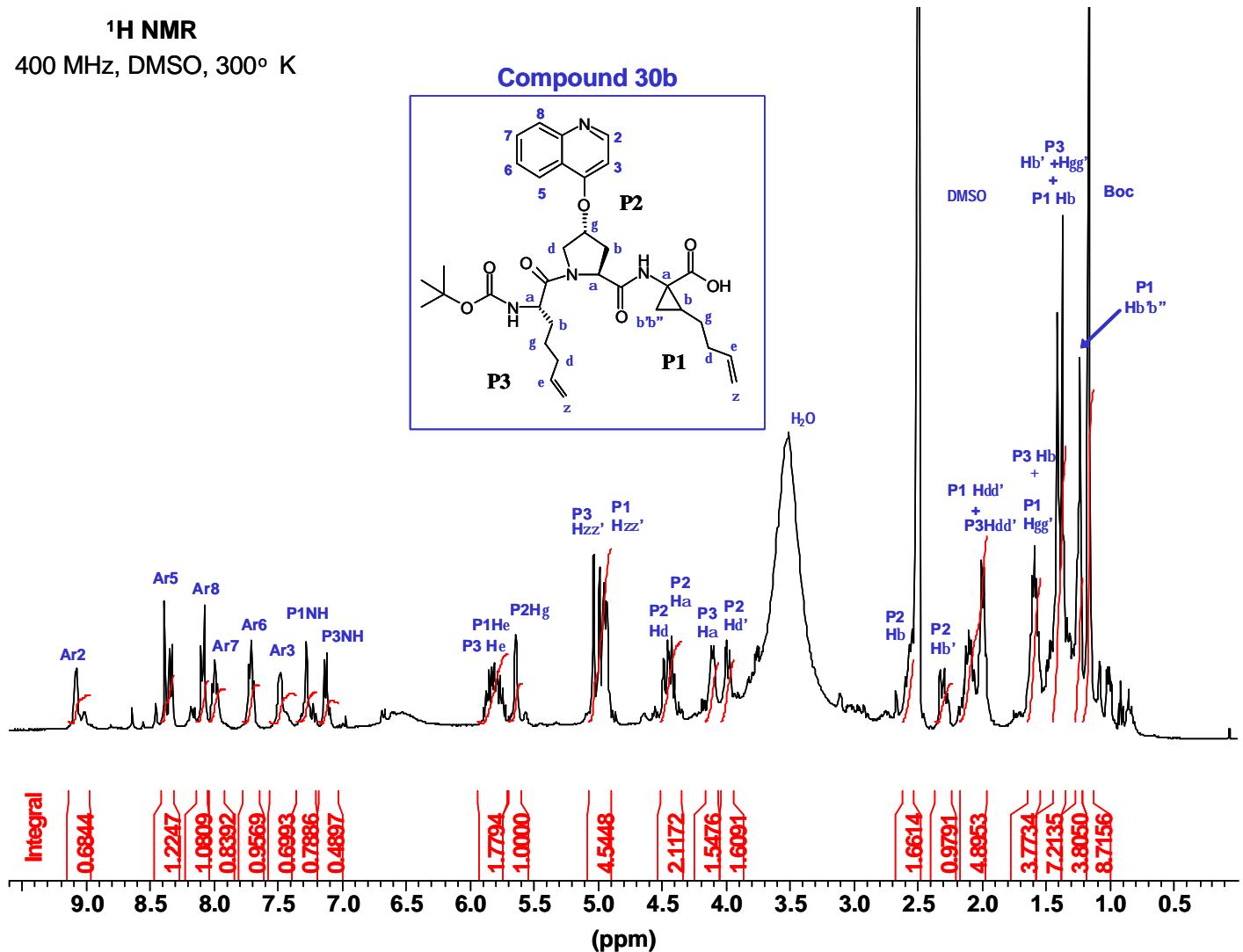
Figure 1. Section of the 2D ^1H NMR spectra of inhibitor **33c**: **a**) NOESY spectrum (200 ms mixing time) of free **33c**; **b**) NOESY spectrum (200 ms mixing time) of **33c** in the presence of NS3 protease BK poly Lys (**33c**:NS3 ratio of 22:1);¹⁴ **c**) NOESY spectrum (200 ms mixing time) of **33c** in the presence of NS3 protease and hexapeptide **43** (**33c**:NS3:**43** ratio of 22:1:2.5). The resonance assignments for all three panels are indicated in panel **(b)**.



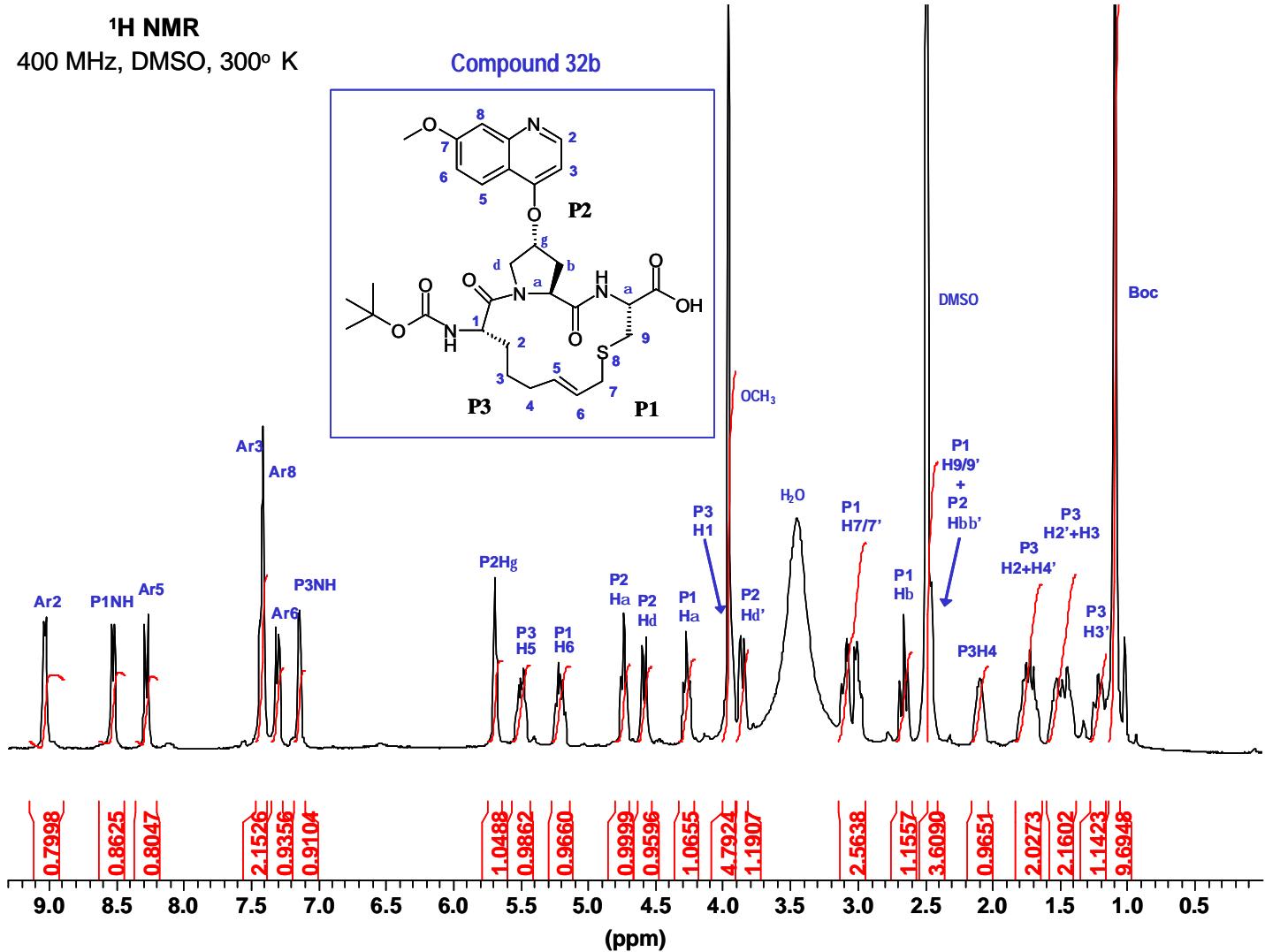
¹H NMR
400 MHz, DMSO, 300° K



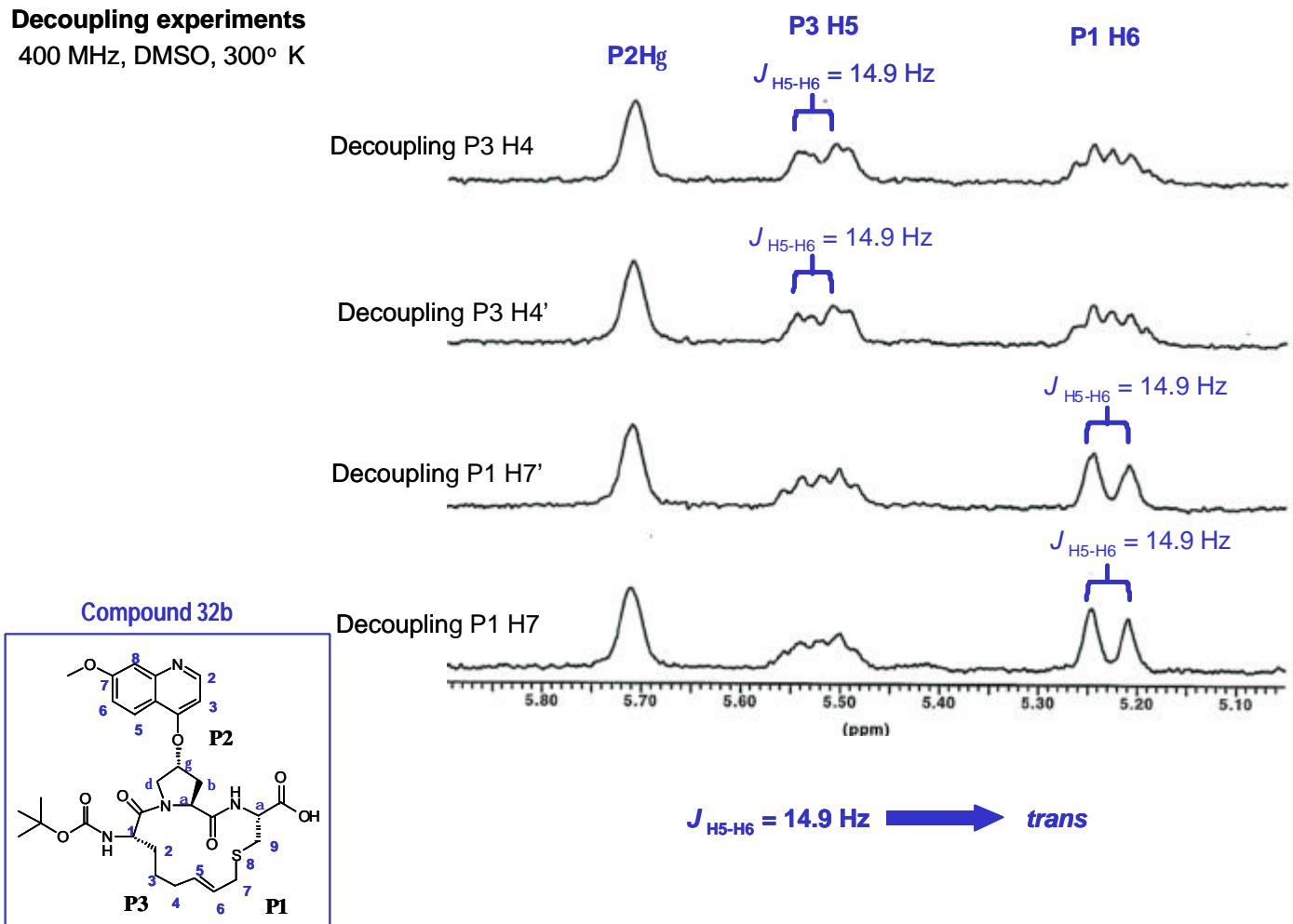
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400 MHz, DMSO, 300° K

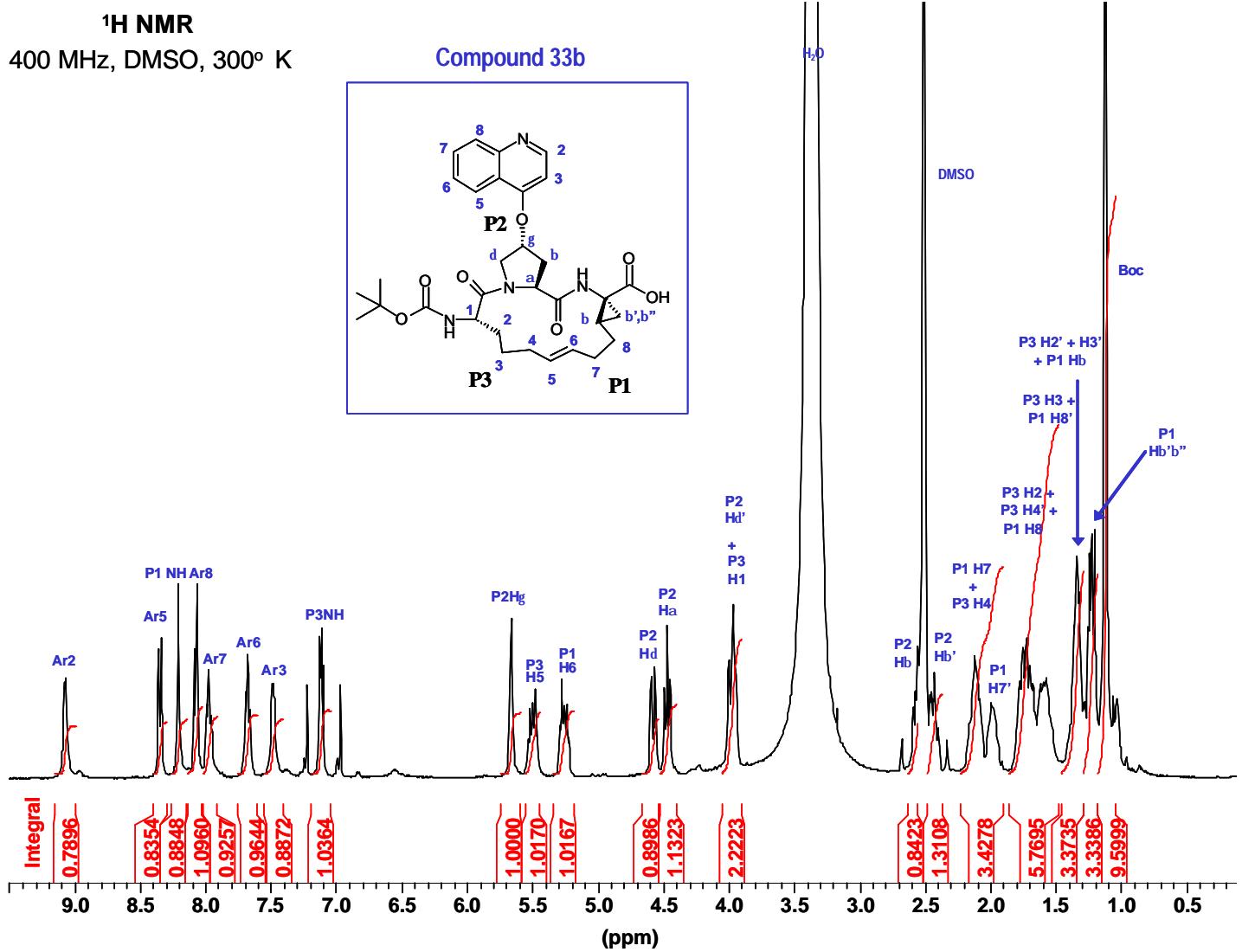


¹H NMR
400 MHz, DMSO, 300° K

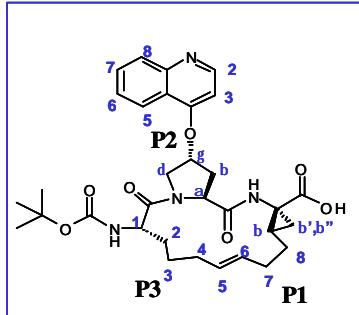
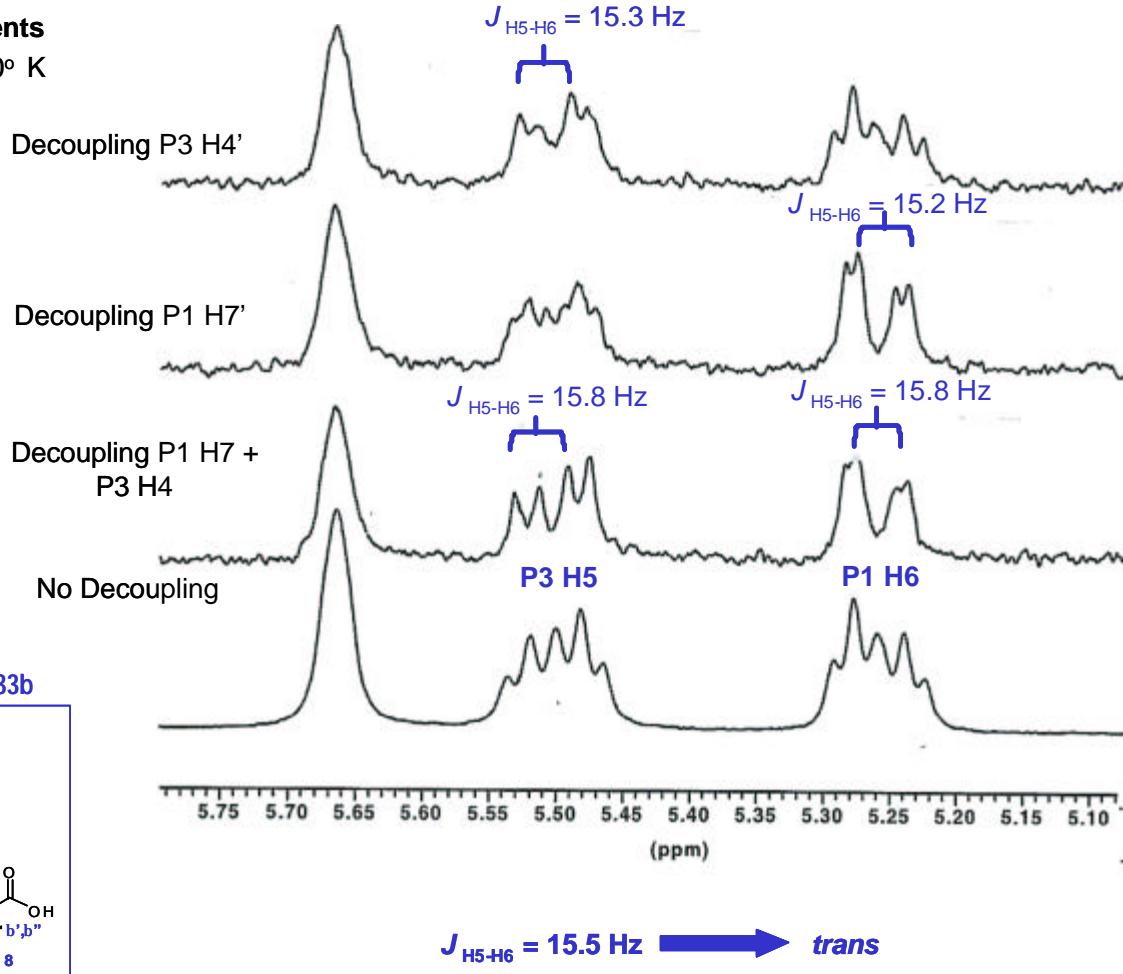


^1H NMR
Decoupling experiments
 400 MHz, DMSO, 300° K



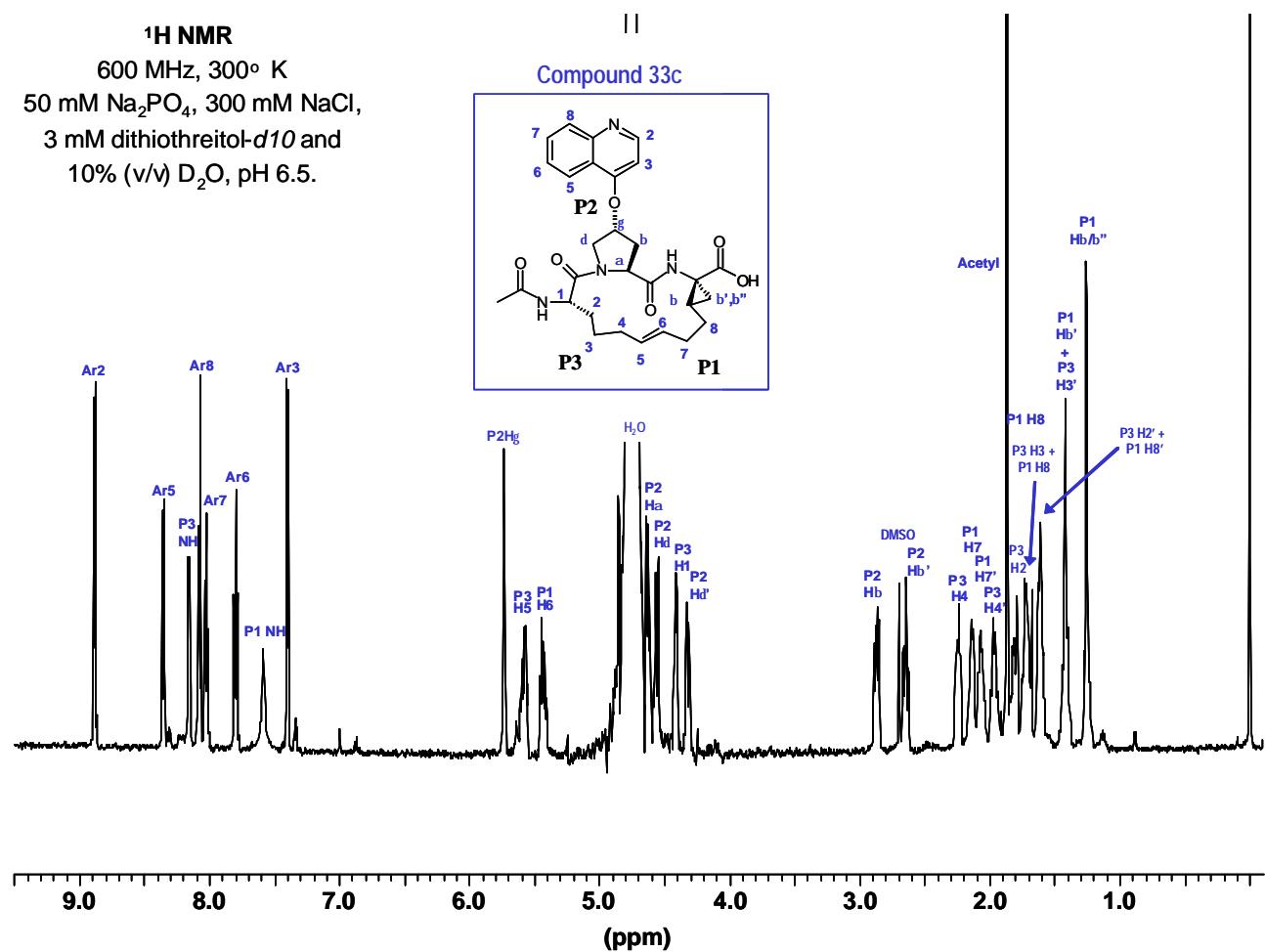


^1H NMR
Decoupling experiments
400 MHz, DMSO, 300° K



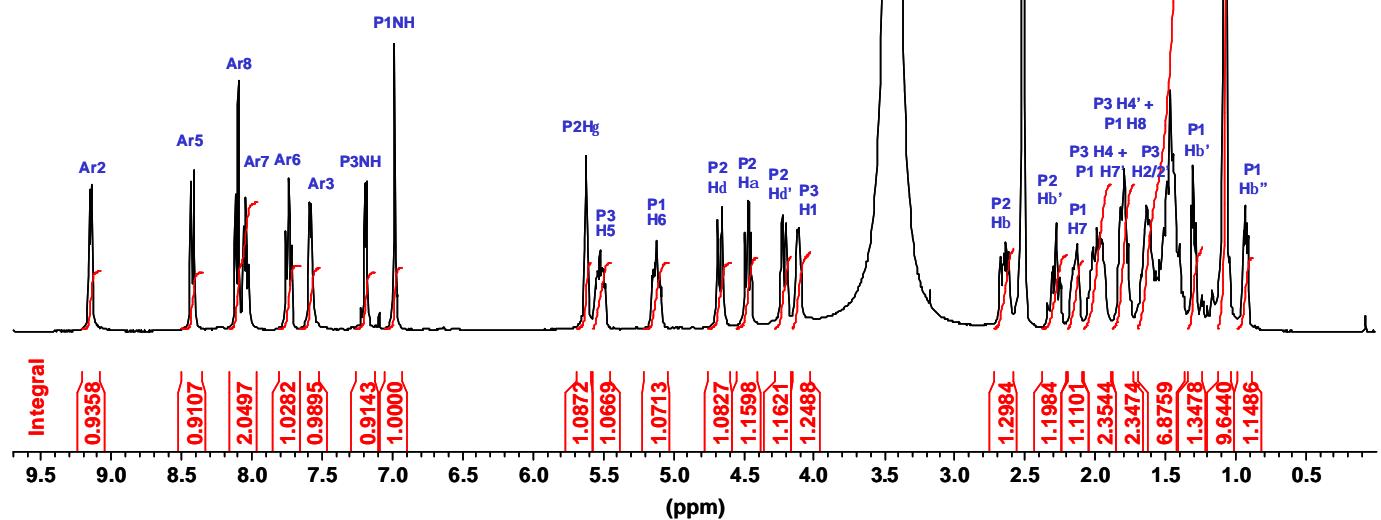
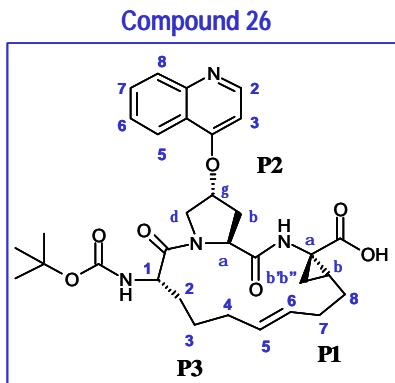
¹H NMR

600 MHz, 300° K
50 mM Na₂PO₄, 300 mM NaCl,
3 mM dithiothreitol-d10 and
10% (v/v) D₂O, pH 6.5.

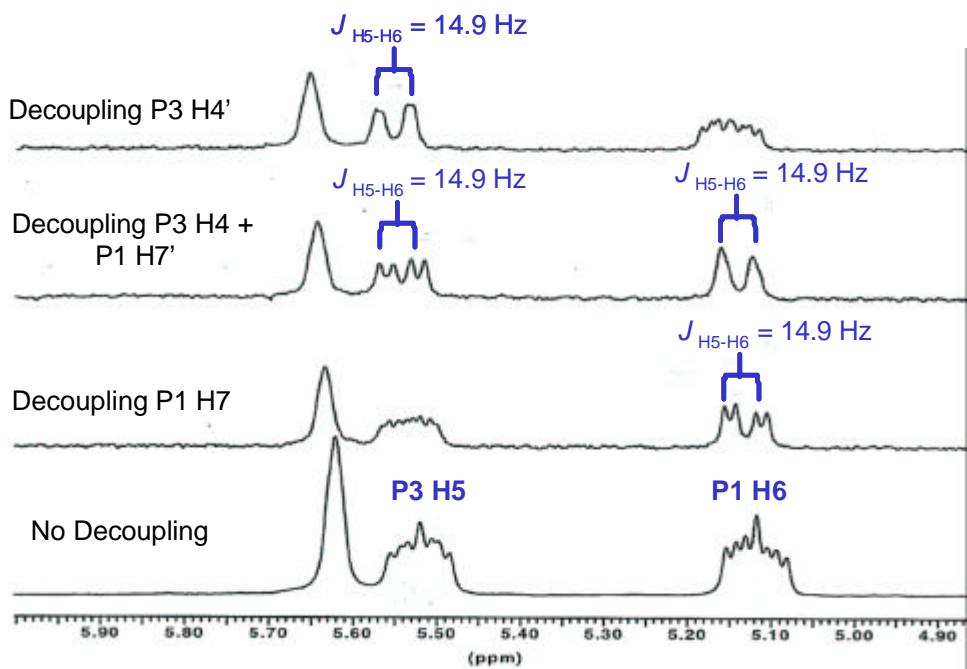


¹H NMR

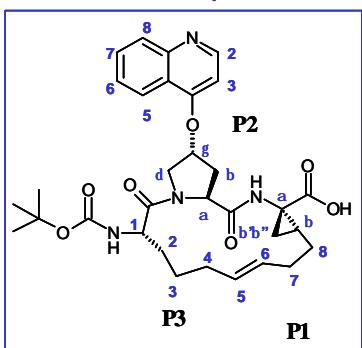
400 MHz, DMSO, 300° K



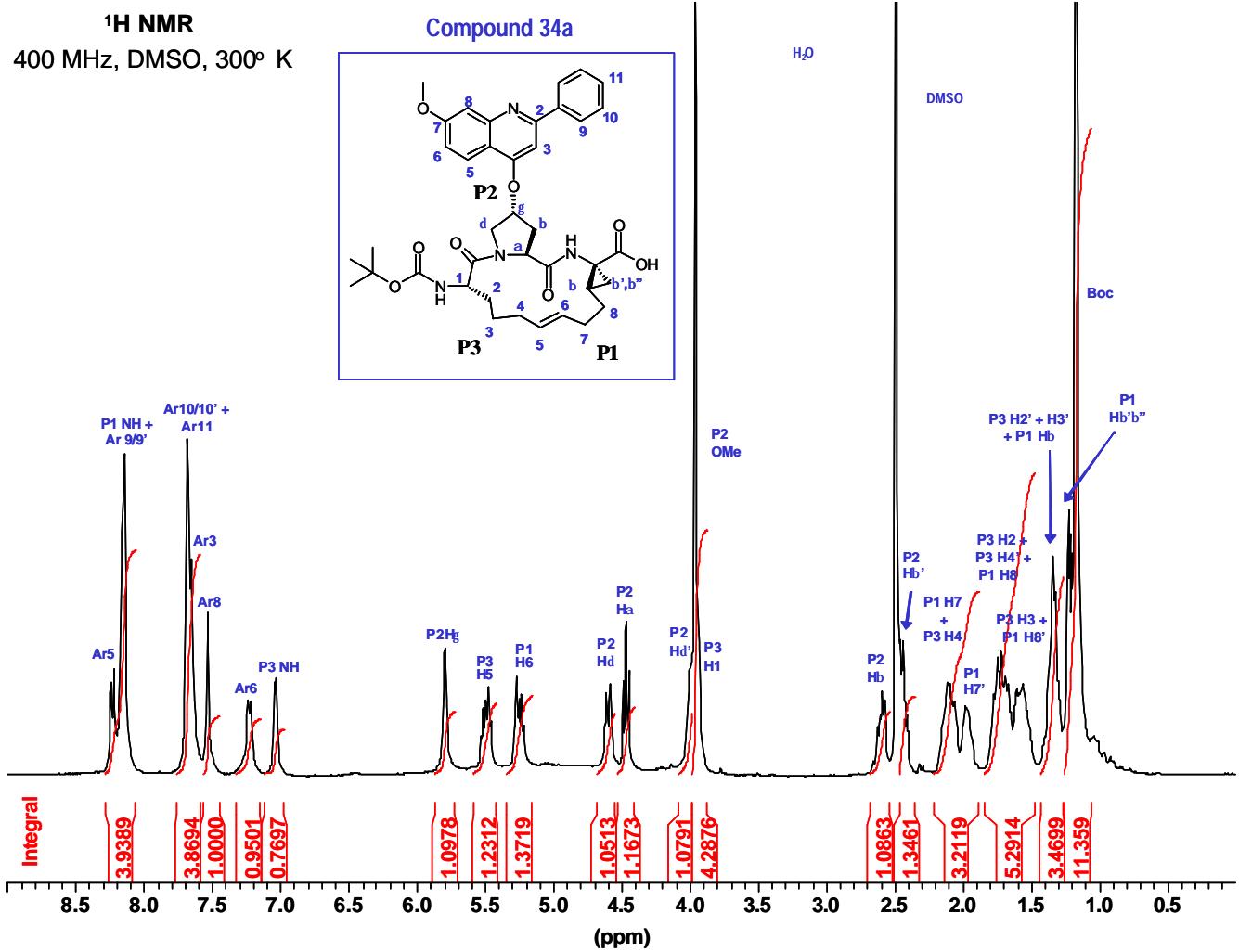
¹H NMR
Decoupling experiments
 400 MHz, DMSO, 300° K



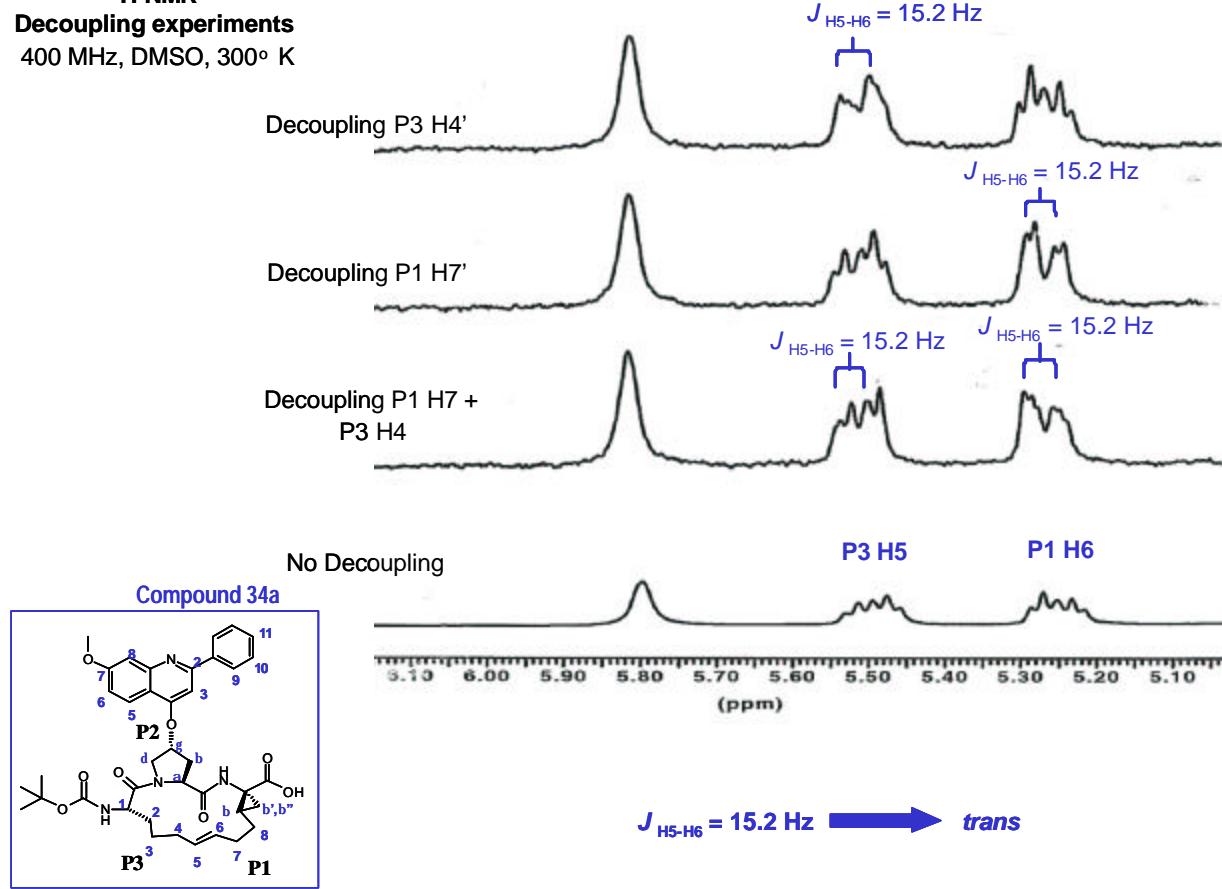
Compound 26

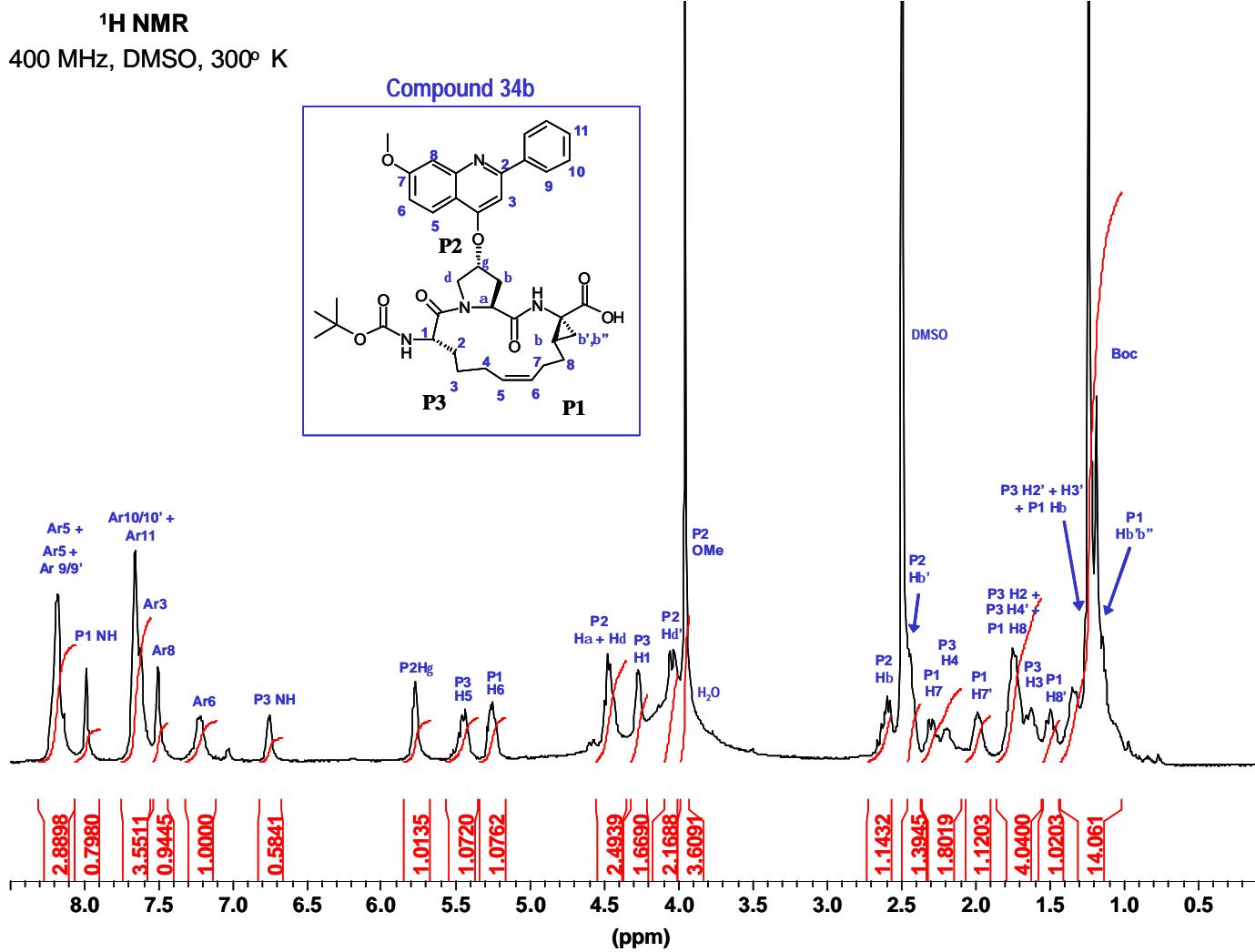


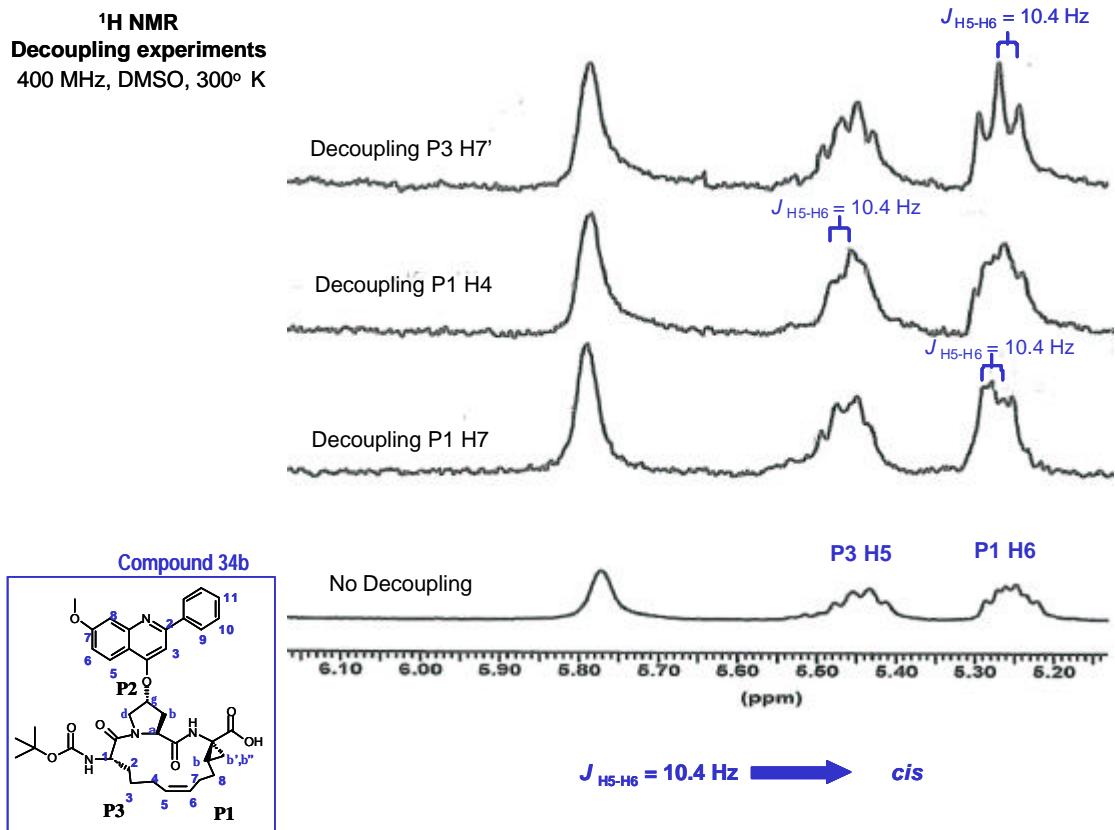
$J_{H5-H6} = 14.9 \text{ Hz}$ → *trans*

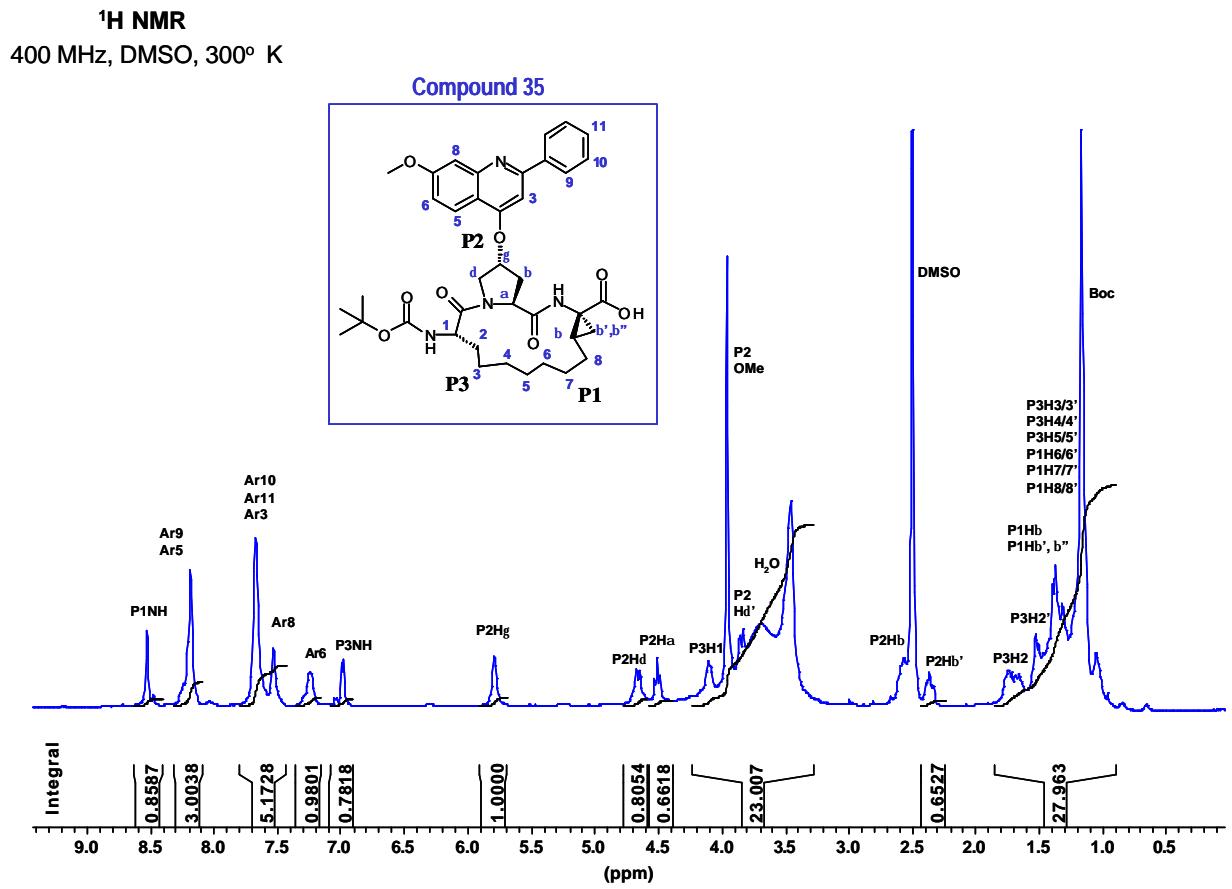


¹H NMR
Decoupling experiments
 400 MHz, DMSO, 300° K

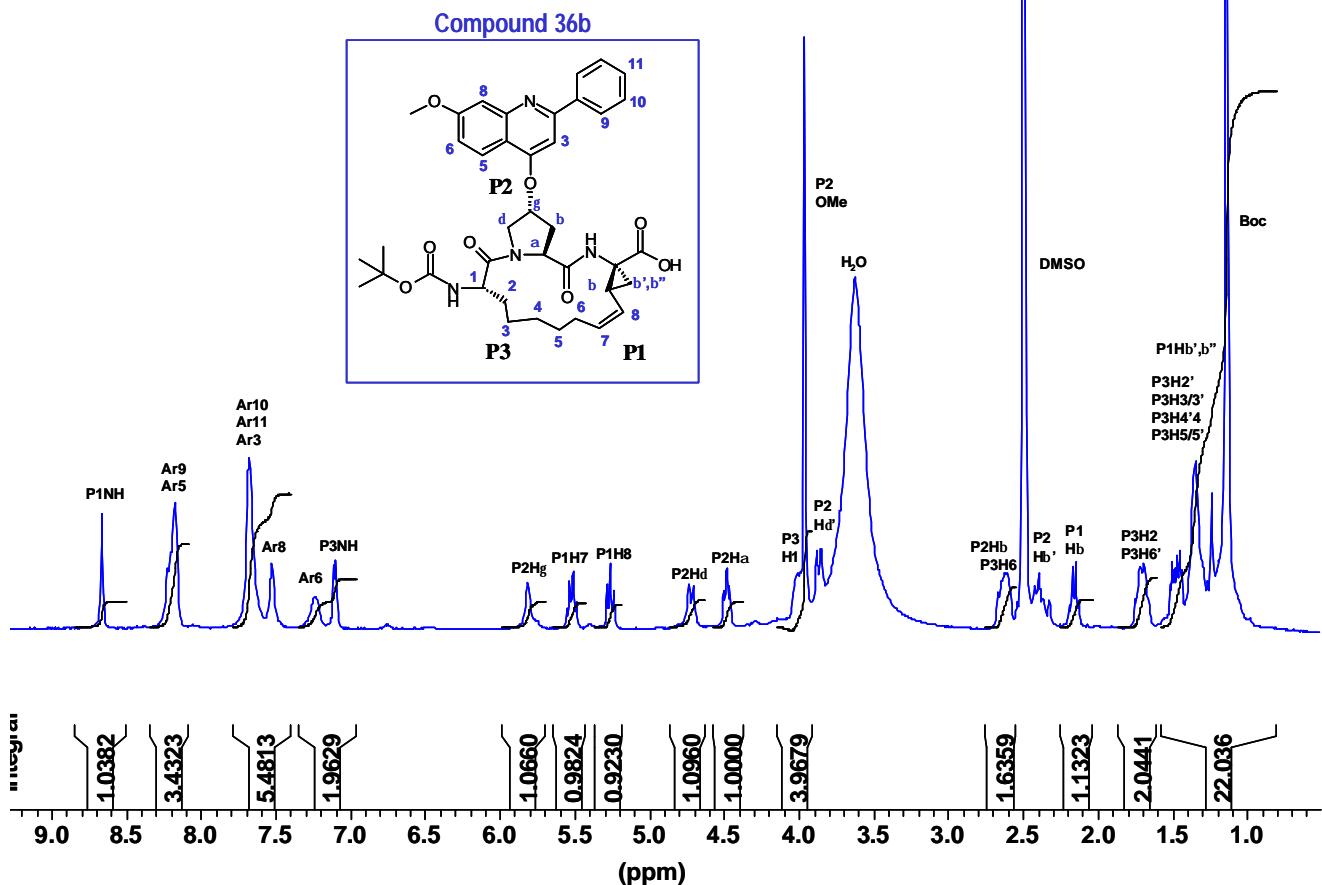




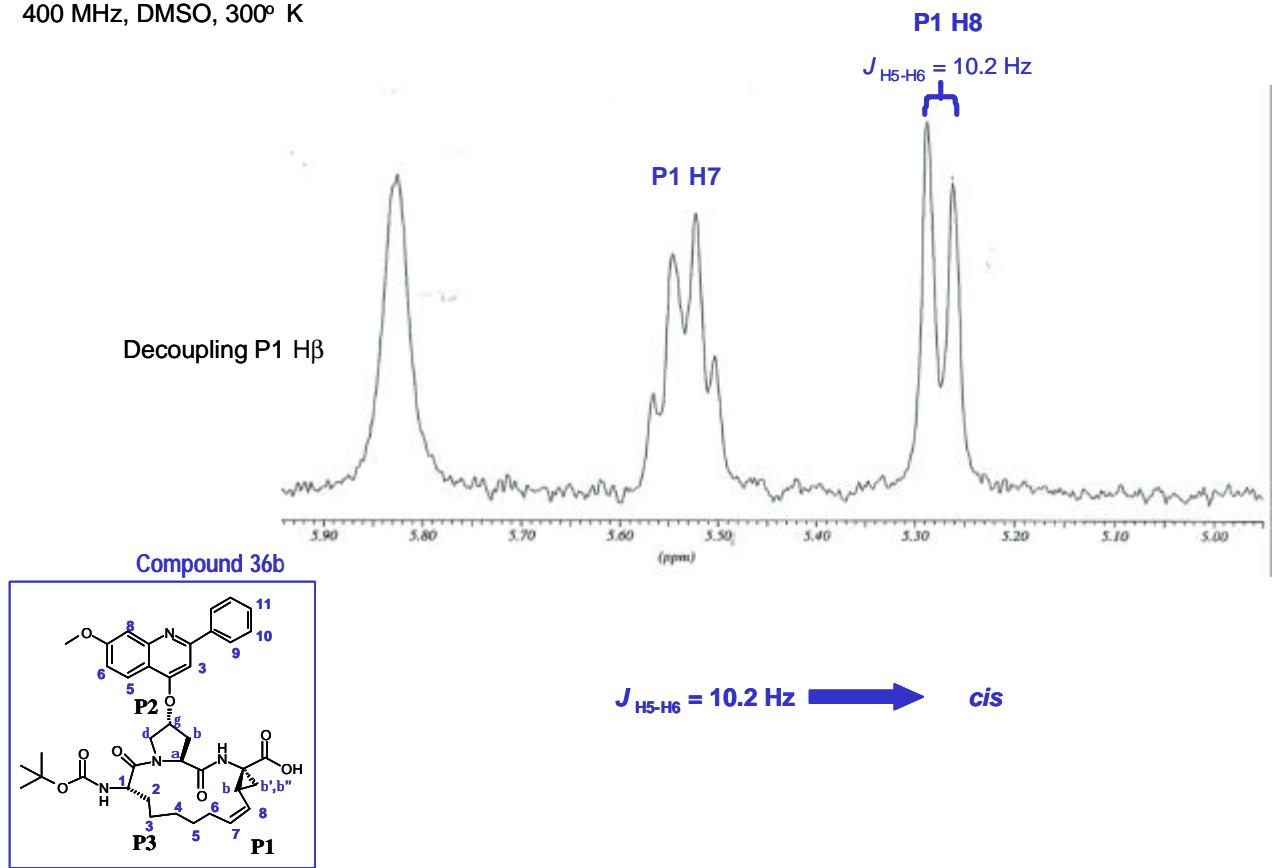




¹H NMR
400 MHz, DMSO, 300° K



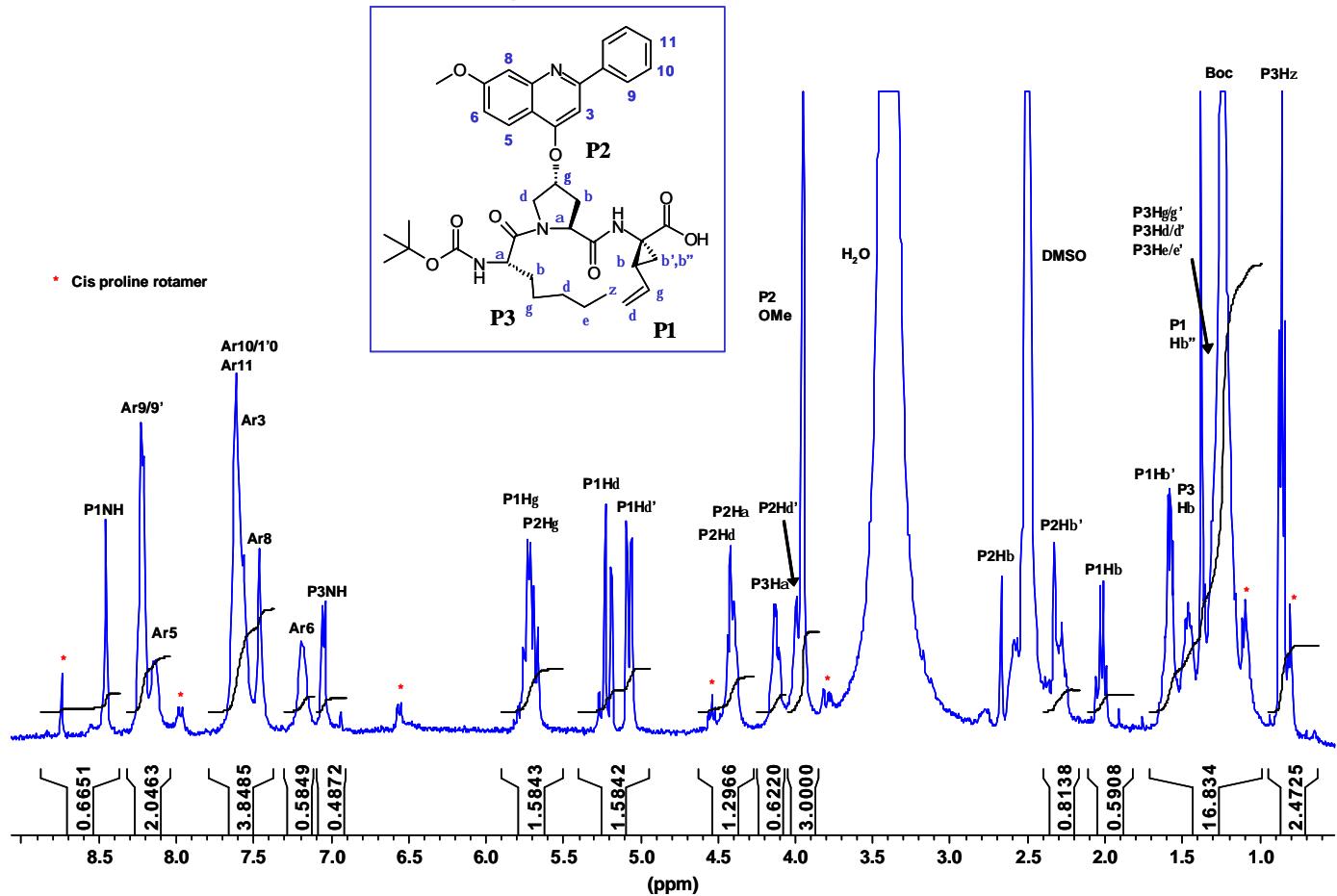
¹H NMR
Decoupling experiments
 400 MHz, DMSO, 300° K

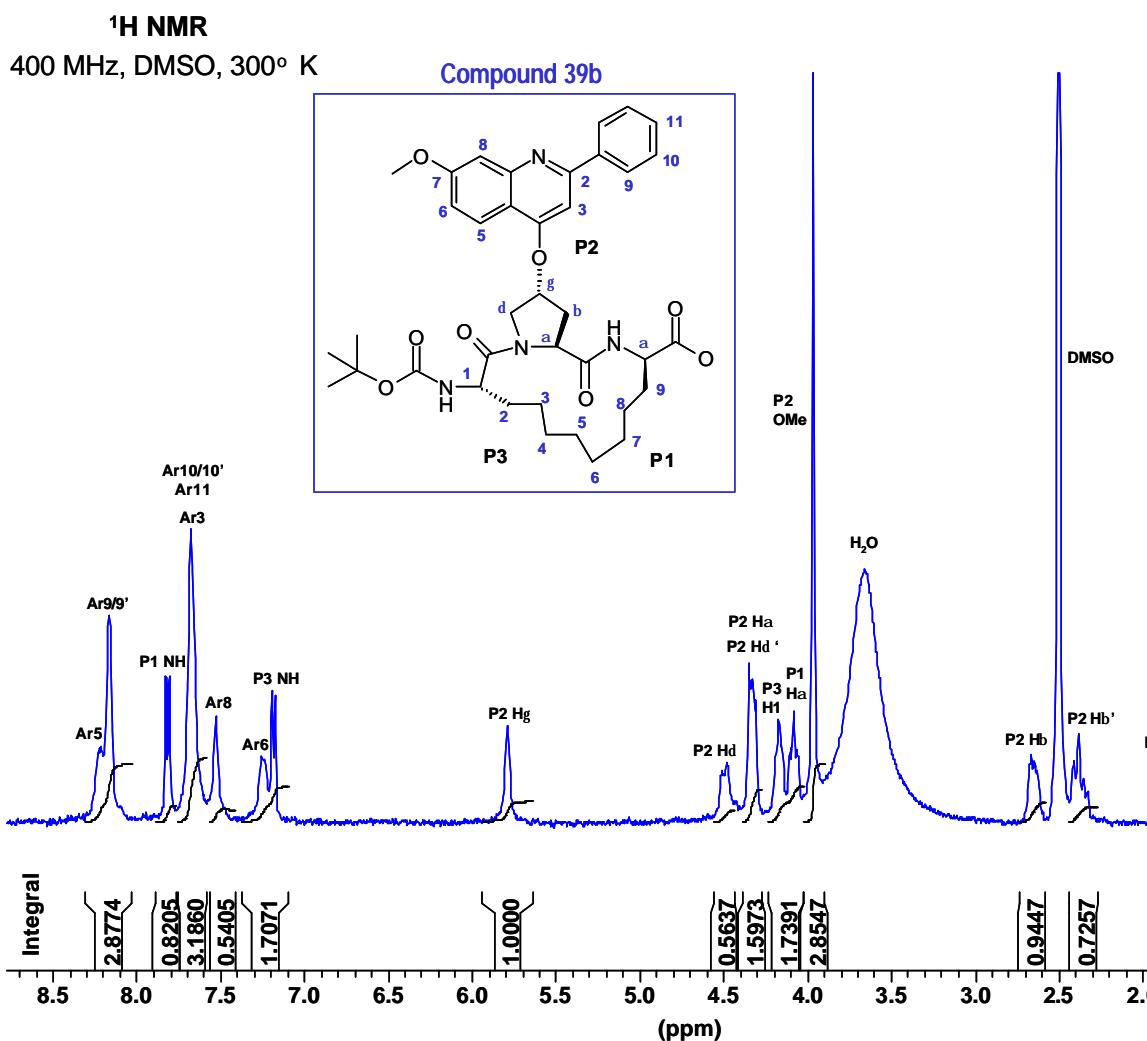


¹H NMR

400 MHz, DMSO, 300° K

Compound 37

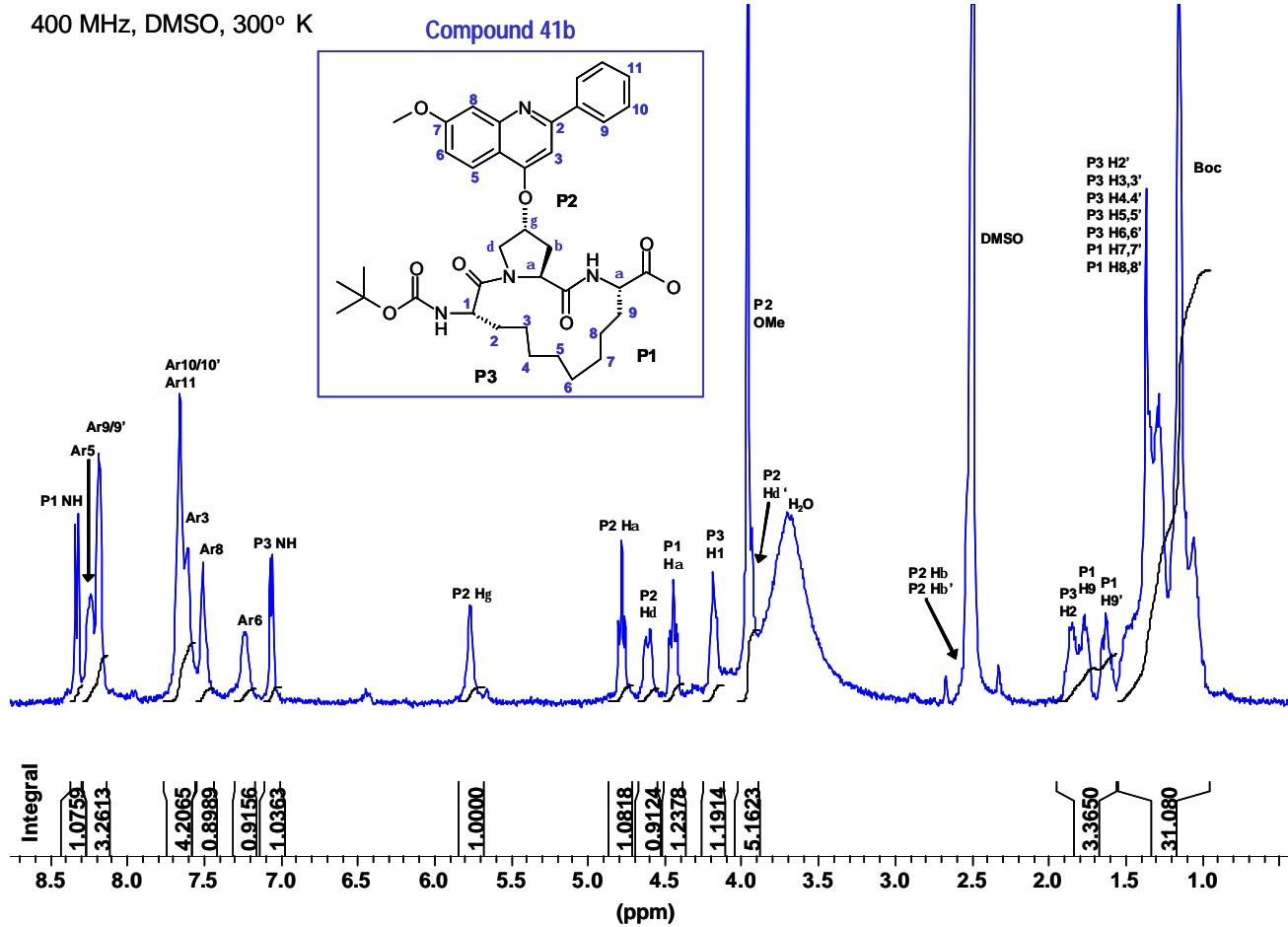




¹H NMR

400 MHz, DMSO, 300° K

Compound 41b



¹H NMR

400 MHz, DMSO, 300° K

