Synthesis of Rigidified Shikimic Acid Derivatives by Ring-Closing Metathesis to Imprint Inhibitor Efficacy against Shikimate Kinase Enzyme

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1. PREPARATION OF ALCOHOLS 18, 20 AND 21



Preparation of 2-(cyclopropylmethyl)prop-2-en-1-ol (18) from diethyl malonate (22)

(a) Diethyl 2-(cyclopropylmethyl)malonate. A solution of diethyl malonate (22) (0.29 mL, 1.71 mmol) in dry THF (3 mL), at 0 °C and under argon, was treated with NaH (75 mg, 1.87 mmol, *ca* 60% in mineral oil). The resultant mixture was stirred at room temperature for 30 min, and then cooled to 0 °C. Cyclopropylmethyl bromide (0.21 mL, 2.20 mmol) was then added and the reaction mixture was first stirred at room temperature for 1 h and then heated under reflux for 6 h. After cooling to room temperature, the solvent was removed under reduced pressure and the crude was dissolved in a mixture of diethyl ether and saturated NH₄Cl. The organic layer was separated and the aqueous layer was extracted with diethyl ether (×2). The combined organic extracts were washed with brine, dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure. The resulting residue was purified by flash chromatography, eluting with (5:95) ethyl acetate/hexane, to give diethyl 2-(cyclopropylmethyl)malonate (245 mg, 67%) as a colorless oil. ¹H NMR (300 MHz, CDCl₃) δ : 4.14 (q, *J* = 6.6 Hz, 4H, 2×CH₂), 3.38 (t, *J* = 7.7 Hz, 1H, CH), 1.74 (t, *J* = 7.2 Hz, 2H, CH₂), 1.21 (t, *J* = 7.1 Hz, 6H, 2×CH₃), 0.73–0.61 (m, 1H, CH₂CH(CH₂)₂), 0.42–0.36 (m, 2H, cPr) and 0.06–0.01 (m, 2H, cPr) ppm.

(b) 2-(Cyclopropylmethyl)malonic acid. A solution of diethyl 2-(cyclopropylmethyl)malonate (1.357 g, 6.34 mmol) in NaOH (8.5 mL, 16.92 mmol, 2 M) was heated under reflux for 2 h. After cooling to room temperature, the reaction mixture was washed with hexane (x2), acidified with HCl (1 M) until pH 1 and then extracted with ethyl acetate (x3). The combined organic extracts were washed with brine, dried (anh. MgSO₄), filtered and concentrated under reduced pressure to afford the corresponding 2-(cyclopropylmethyl)malonic acid (898 mg, 90%) as a white solid. ¹H NMR (300 MHz,DMSO) δ : 12.68 (br s, 2H, 2×OH), 3.26 (t, *J* = 7.5 Hz, 1H, CH), 1.61 (t, *J* = 7.2 Hz, 2H, CH₂), 0.74–0.61 (m, 1H, CH₂CH(CH₂)₂), 0.40–0.35 (m, 2H, *c*Pr) and 0.08–0.04 (m, 2H, *c*Pr) ppm.

(c) 2-(Cyclopropylmethyl)acrylic acid (23). A solution of 2-(cyclopropylmethyl)malonic acid (878 mg, 5.55 mmol), piperidine (0.66 mL, 6.66 mmol) and HCHO (2 mL, 27.75 mmol, *ca* 37% wt) in ethanol (22 mL) was heated at 80 °C for 12 h. After cooling to room temperature, the organic solvent was removed under reduced pressure. The residue was diluted with diethyl ether and saturated NaHCO₃. The aqueous phase was separated and the organic layer was extracted with saturated NaHCO₃. The combined aqueous phases were washed with diethyl ether and then acidified with HCl (conc.) until pH 1. The solution was extracted with diethyl ether (x3). The combined organic extracts were washed with brine, dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure to afford 2-(cyclopropylmethyl)acrylic acid (23) (663 mg, 95%) as a colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 9.96 (br s, 1H, OH), 6.30 (s, 1H, C=CHH), 5.80 (m, 1H, C=CHH), 2.18 (d, *J* = 6.8 Hz, 2H, CH₂CH(CH₂)₂), 0.94–0.81 (m, 1H, CH₂CH(CH₂)₂), 0.52–0.46 (m, 2H, CPr) and 0.08 (m, 2H, *c*Pr) ppm.

(e) 2-(Cyclopropylmethyl)prop-2-en-1-ol (18). A solution of 2-(cyclopropylmethyl)acrylic acid (23) (651 mg, 5.16 mmol) in dry THF (52 mL), at 0 °C and under argon, was treated with BH₃.Me₂S complex (3 mL, 6.20 mmol, 2 M). The reaction mixture was stirred at room temperature for 5 h and then diluted with ethyl acetate and water (1:1). The organic layer was separated and the aqueous layer was extracted with ethyl acetate (×3). The combined organic extracts were dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure to afford 2-(cyclopropylmethyl)prop-2-en-1-ol (18) (350 mg, 60%) as a colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.02 (s, 2H, C=CH₂), 4.13 (s, 2H, CH₂OH), 1.97 (d, *J* = 6.8 Hz, 2H, CH₂CH(CH₂)₂), 0.86–0.76 (m, 1H, CH₂CH(CH₂)₂), 0.53–0.47 (m, 2H, *c*Pr) and 0.09 (m, 2H, *c*Pr) ppm.



2-(Ethoxymethyl)prop-2-en-1-ol (20). A solution of 2-methylene-1,3-propanodiol (**19**) (50 mg, 0.57 mmol) in dry DMF (1.1 mL), at 0 °C and under argon, was treated with NaH (34 mg, 0.86 mmol, *ca* 60% w/w in mineral oil). After 30 min stirring, ethyl bromide (0.1 mL, 1.14 mmol) was added, the ice bath was removed and the reacion mixture was stirred for 1.5 h at room temperature. Ethyl acetate and water (4:1) were added, the aqueous fase was extracted with ethyl acetate (×3). The combined organic extracts were dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure. The resulting residue was purified by flash chromatography, eluting with (60:40) ethyl acetate/hexane, to give compound **20** (29 mg, 44%) as a colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.15 (s, 1H, CHH), 5.10 (s, 1H, CHH), 4.15 (s, 2H, OCH₂), 4.03 (s, 2H, OCH₂), 3.48 (q, *J* = 7.0 Hz, 2H, OCH₂CH₃), 2.17 (s, 1H, OH) and 1.20 (t, *J* = 7.0 Hz, 3H, CH₃) ppm. IR (film): 3399 (OH) cm⁻¹. MS (ESI) *m/z* = 117 (MH⁺). HRMS calcd for C₆H₁₃O₂ (MH⁺): 117.0910; found, 117.0909.



2-(Benzyloxymethyl)prop-2-en-1-ol (21). A solution of 2-methylene-1,3-propanodiol (**19**) (50 mg, 0.57 mmol) in dry THF (1.9 mL), at 0 °C and under argon, was treated with NaH (34.2 mg, 0.86 mmol, *ca* 60% w/w in mineral oil). After 30 min stirring, benzyl bromide (0.1 mL, 0.9 mmol) was added, the ice bath was removed and the reacion mixture was stirred for 3 h at room temperature. Saturated aqueous solution of NH₄Cl was added, the organic solvent was removed under reduced pressure and the aqueous extract was extracted with ethyl acetate (×3). The combined organic extracts were dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure. The resulting residue was purified by flash chromatography, eluting with (60:40) ethyl acetate/hexane, to give compound **21** (28 mg, 28%) as a colorless oil. ¹H NMR (300 MHz, CDCl₃) δ : 7.39–7.26 (m, 5H, 5×ArH), 5.21 (s, 1H, C=CHH), 5.16 (s, 1H, C=CHH), 4.52 (s, 2H, OCH₂Ph), 4.17 (s, 2H, CH₂OH), 4.09 (s, 2H, CH₂OBn) and 2.55 (br s, 1H, OH) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 145.1 (C), 138.0 (C), 128.5 (2×CH), 127.8 (3×CH), 113.4 (CH₂), 72.3 (CH₂), 71.6 (CH₂) and 64.2 (CH₂) ppm. IR (film): 3372 (OH) cm⁻¹. MS (ESI) *m/z* = 201 (MNa⁺). HRMS calcd for C₁₁H₁₄O₂Na (MNa⁺): 201.0886; found, 201.0888.

2. PREPARATION OF CARBONATES 8

Methyl-2-methylallyl carbonate (**8a**) and methyl (2-methylenebutyl) carbonate (**8b**) were prepared as reported previously described by Dai *et al.*¹⁷ from commercially available 2-methylprop-2-en-1-ol (**14**) and 2-methylenebutan-1-ol (**15**), which was prepared from commercially available ketone **11** as reported Sawayama *et. al.*¹

General procedure for synthesis of carbonates 8a-h. A solution of the alcohols **14–21** (1 mmol) and dry pyridine (3 mmol) in dry dichloromethane (0.4 mL/mmol of alcohol), at 0 °C and under argon, was treated with methyl chloroformate (2 mmol) and the resultant solution was stirred at room temperature for 2–2.5 h. The reaction mixture was washed with brine and the organic layer was extracted with diethyl ether (×3). The combined organic extracts, were washed with HCl (1 M), dried (anh. MgSO₄), filtered and concentrated under reduced pressure. The reaction crude was purified by flash chromatography to give the carbonates **8a-h**.



Methyl (2-methylenepentyl) carbonate (8c). It was prepared following the general procedure using 2-methylenepentan-1ol (**16**)² (627 mg, 6.27 mmol), pyridine (1.5 mL), methyl chloroformate (1.0 mL, 12.54 mmol), dichloromethane (16 mL). Reaction time = 2.5 h. Eluent for chromatography = (75:25) diethyl ether/hexanes. Yield = 78% (919 mg). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.03 (br s, 1H, C=CHH), 4.92 (br s, 1H, C=CHH), 4.54 (s, 2H, OCH₂), 3.76 (s, 3H, OCH₃), 2.02 (t, *J* = 7.4 Hz, 2H, CH₂CH₂CH₃), 1.46 (sext, *J* = 7.4 Hz, 2H, CH₂CH₂CH₃) and 0.89 (t, *J* = 7.3 Hz, 3H, CH₂CH₂CH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.7 (C), 143.3 (C), 112.8 (CH₂), 70.3 (CH₂), 54.8 (OCH₃), 35.1 (CH₂), 20.7 (CH₂) and 13.8 (CH₃) ppm. FTIR (film): 1750 (CO) cm⁻¹. MS (APCI) *m/z* = 159 (MH⁺). HMRS calcd for C₈H₁₅O₃ (MH⁺): 159.1016; found, 159.1014.



Methyl 2-(methylenehexyl) carbonate (8d). It was prepared following the general procedure using 2-methylenehexyl-1-ol $(17)^2$ (3.16 g, 31.59 mmol), pyridine (5 mL), methyl chloroformate (7 mL, 94.77 mmol), dichloromethane (63 mL). Reaction time = 2.5 h. Yield = 94% (5.12 g). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ: 4.89 (br s, 1H, C=CHH), 4.78 (br s, 1H, C=CHH), 4.41 (s, 2H, OCH₂), 3.61 (s, 3H, OCH₃), 1.92 (t, *J* = 7.3 Hz, 2H, CH₂(CH₂)₂CH₃), 1.30 (m, 2H, CH₂CH₂CH₂CH₃), 1.17 (m, 2H, CH₂CH₂CH₃) and 0.76 (t, *J* = 7.2 Hz, 3H, CH₂(CH₂)₂CH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ: 155.3 (C), 143.4 (C), 111.9 (CH₂), 69.7 (CH₂), 54.2 (OCH₃), 32.4 (CH₂), 29.4 (CH₂), 22.1 (CH₂) and 13.5 (CH₃) ppm. FTIR (film): 1750 (CO) cm⁻¹. MS (APCI) *m/z* = 173 (MH⁺). HRMS calcd for C₉H₁₇O₃ (MH⁺): 173.1172; found, 173.1171.

¹ Sawayama, A. M.; Tanaka,H.; Wandless, T. J. Total synthesis of ustiloxin D and considerations on the origin of selectivity of the asymmetric allylic alkylation. *J. Org. Chem.* **2004**, *69*, 8810–8820.

² Ma, W.; Fang, J.; Ren, J.; Wang, Z. Lewis acid catalyzed formal intramolecular [3 + 3] cross-cycloaddition of cyclopropane 1,1-diesters for construction of benzobicyclo[2.2.2]octane skeletons. Org. Lett. **2015**, *17*, 4180–4183.



8e

2-(Cyclopropylmethyl)allyl methyl carbonate (8e). It was prepared following the general procedure using 2-(cyclopropylmethyl)prop-2-en-1-ol (**18**) (321 mg, 2.86 mmol), pyridine (0.7 mL), methyl chloroformate (0.4 mL), dichloromethane (7.2 mL). Reaction time = 2.5 h. Eluent for chromatography = (10:90) ethyl acetate/hexane. Yield = 72% (350 mg). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.08 (br s, 1H, C=CHH), 5.04 (br s, 1H, C=CHH), 4.61 (br s, 2H, OCH₂), 3.76 (s, 3H, OCH₃), 1.95 (d, *J* = 6.9 Hz, 2H, CH₂CH(CH₂)₂), 0.80–0.73 (m, 1H, CH₂CH(CH₂)₂), 0.50–0.44 (m, 2H, CH₂CHCH₂CH₂) and 0.06 (m, 2H, CH₂CHCH₂CH₂) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.8 (C), 143.6 (C), 112.9 (CH₂), 70.5 (CH₂), 54.8 (OCH₃), 38.1 (CH₂), 9.1 (CH) and 4.7 (2×CH₂) ppm. IR (film): 1744 (CO) cm⁻¹. MS (ESI) *m/z* = 171 (MH⁺). HRMS calcd for C₉H₁₅O₃ (MH⁺): 171.1016; found, 171.1020.



2-(Ethoxymethyl)allyl methyl carbonate (8f). It was prepared following the general procedure using 2-(ethoxymethyl)prop-2-en-1-ol (**19**) (251 mg, 1.16 mmol), pyridine (0.3 mL), methyl chloroformate (0.18 mL, 2.27 mmol), solution of methyl chloroformate in dry dichloromethane (2.4 mL, 1.0 M), dichloromethane (2.1 mL). Reaction time = 2 h. Eluent for chromatography = (30:70) ethyl acetate/hexane. Yield = 78% (240 mg). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.16 (s, 2H, C=CH₂), 4.58 (s, 2H, OCH₂), 3.91 (s, 2H, CH₂), 3.70 (s, 3H, OCH₃), 3.40 (q, *J* = 7.0 Hz, 2H, OCH₂CH₃) and 1.12 (d, *J* = 7.0 Hz, 3H, CH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.5 (C), 140.5 (C), 115.2 (CH₂), 71.0 (CH₂), 68.0 (CH₂), 65.7 (CH₂), 54.7 (OCH₃) and 15.0 (CH₃) ppm. IR (film): 1751 (CO) cm⁻¹. MS (ESI) *m/z* = 175 (MH⁺). HRMS calcd for C₈H₁₅O₄ (MH⁺): 175.0965; found, 175.0964.



2-(Benzyloxymethyl)allyl methyl carbonate (8g). It was prepared following the general procedure using 2-(benzyloxymethyl)prop-2-en-1-ol (**20**) (207 mg, 1.16 mmol), pyridine (0.3 mL), methyl chloroformate (0.2 mL), dichloromethane (2.9 mL). Reaction time = 2.5 h. Eluent for chromatography = (20:80) ethyl acetate/hexane. Yield = 94% (257 mg). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 7.34–7.26 (m, 5H, 5×ArH), 5.30 (s, 2H, C=CH₂), 4.71 (s, 2H, OCH₂), 4.51 (s, 2H, OCH₂Ph), 4.07 (s, 2H, CH₂OBn) and 3.78 (s, 3H, CH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.6 (C), 140.2 (C), 138.0 (C), 128.4 (2×CH), 127.7 (2×CH), 127.7 (CH), 115.9 (CH₂), 72.2 (CH₂), 70.6 (CH₂), 68.1 (CH₂) and 54.8 (CH₃) ppm. IR (film): 1744 (CO) cm⁻¹. MS (ESI) *m/z* = 259 (MNa⁺). HRMS calcd for C₁₃H₁₆O₄Na (MNa⁺): 259.0941; found, 259.0941.



8ŀ

2-(Hydroxymethyl)allyl methyl carbonate (8h). It was prepared following the general procedure using 2-methylene-1,3-propanediol (**19**) (205 mg, 2.33 mmol), pyridine (0.4 mL), methyl chloroformate (0.2 mL), dichloromethane (2.3 mL). Reaction time = 2.5 h. Eluent for chromatography = 40:60) ethyl acetate/hexane. Yield = 55% (188 mg). Colorless liquid. ¹H NMR (300 MHz, CDCl₃) δ : 5.19 (m, 1H, C=CHH), 5.14 (s, 1H, C=CHH), 4.61 (s, 2H, OCH₂), 4.08 (s, 1H, CHHOH), 4.07 (s, 1H, CHHOH) and 3.72 (s, 3H, OCH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.7 (C), 142.8 (C), 114.4 (CH₂), 68.1 (CH₂), 63.2 (CH₂) and 54.9 (OCH₃) ppm. IR (film) u: 3390 (OH) and 1747 (CO) cm⁻¹.



2-((tert-Butyldimethylsilyloxy)methyl)allyl methyl carbonate (8i). A solution of 2-(hydroxymethyl)allyl methyl carbonate **(8h)** (180 mg, 1.23 mmol), DMAP (42 mg, 0.35 mmol), TBAI (46 mg, 0.12 mmol) and dry triethylamine (0.2 mL, 1.35 mmol) in dry DMF (2.0 mL), at 0 °C and under argon, was treated with *tert*-butyldimethylsilyl chloride (204 mg, 1.35 mmol). The reaction mixture was stirred at 0 °C for 1 h and at room temperature for 3 h. The resultant suspension was diluted with

ethyl acetate and filtered over a plug of Celite[®]. The filtrate and the washings were washed with HCl (1 M, ×3) followed by brine (×3). The organic extract was dried (anh. Na₂SO₄), filtered and concentrated under reduced pressure. The resulting residue was purified by flash chromatography, eluting with (7:93) ethyl acetate/hexane, to give the silyl ether **22** (297 mg, 93%) as a colorless oil. ¹H NMR (300 MHz, CDCl₃) δ : 5.21 (m, 1H, C=CHH), 5.12 (m, 1H, C=CHH), 4.61 (s, 2H, OCH₂), 4.14 (br s, 2H, CH₂OTBS), 3.74 (s, 3H, OCH₃), 0.87 (s, 9H, C(CH₃)₃) and 0.03 (s, 6H, 2×CH₃) ppm. ¹³C NMR (75 MHz, CDCl₃) δ : 155.7 (C), 142.8 (C), 113.5 (CH₂), 68.1 (CH₂), 63.7 (CH₂), 54.8 (OCH₃), 25.9 (C(CH₃)₃), 18.4 (*C*(CH₃)₃) and 5.4 (2×CH₃) ppm. IR (film): 1751 (CO) cm⁻¹. MS (ESI) *m/z* = 283 (MNa⁺). HRMS calcd for C₁₂H₂₄O₄SiNa (MNa⁺): 283.1336; found, 283.1336.

3. SHIKIMATE KINASE ASSAY

Both enzymes were purified as described previously.^{3,18} Concentrated solutions of Hp-SK (7.4 mg mL⁻¹) or Mt-SK (1.5 mg mL⁻¹) were stored in potassium phosphate buffer (50 mM, pH 7.2), DTT (1 mM) and NaCl (150 mM). When required for assays, aliquots of the enzyme stock solutions were diluted in water and buffer and stored on ice. Enzyme activity was measured by monitoring the decrease in absorbance at 340 nm in the UV spectrum due to the absorbance of NADH (ϵ/M^{-1} cm⁻¹ 6 220) in a coupled assay format wherein ADP formed after the formation of shikimate-3-phosphate was detected using pyruvate kinase (PK) and lactate dehydrogenase (LDH). Oxidation of NADH to NAD during PK-LDH activity was monitored at 340 nm. Standard assay conditions for shikimate kinase were 100 mM Tris.HCl pH 7.7, 100 mM KCl, 5 mM MgCl₂, 2.5 mM ATP, 1 mM shikimic acid, 1 mM PEP, 0.20 mM NADH, ~2.8 units of PK-LDH and 0.03 unit of SK protein. One unit of enzyme is defined as the amount of enzyme required to convert 1 µmol of substrate to product in 1 min. Each assay was initiated by addition of shikimic acid. Solutions of ATP (ϵ/M^{-1} cm⁻¹ 15 400) and NADH (ϵ/M^{-1} cm⁻¹ 6220) were calibrated by equilibration with SK and measurement of the change in the UV absorbance at 340 nm due to the disappearance of NADH. The K_i values of compounds **2–5** against SK were obtained from Dixon plots (1/v vs [I]) of assay data. The initial rates at fixed enzyme and substrate concentrations (0.25–1.2 K_m) were measured in the absence and in the presence of various inhibitor concentrations.



³ Krell, T.; Maclean, J.; Boam, D. J.; Cooper, A.; Resmini, M.; Brocklehurst, K.; Kelly, S. M.; Price, N. C.; Lapthorn, A. J.; Coggins, J. R. Biochemical and Xray crystallographic studies on shikimate kinase: The important structural role of the P-loop lysine. *Protein Science* **2001**, *10*, 1137–1149.



Figure S1. Conformation of minimum energy of **6** (R = H) obtained by computational calculations (Gaussian 09, AM1).



Figure S2. The rmsd plots for the protein backbone (C α , C, N and O atoms) (orange line) and ligand (blue line) calculated for the *Hp*-SK/ATP/Mg⁺²/ligand complex obtained from MD simulation studies. A. *Hp*-SK/ATP/Mg⁺²/**4a** complex. B. *Hp*-SK/ATP/Mg⁺²/**4b** complex. C. *Hp*-SK/ATP/Mg⁺²/**4c** complex. D. *Hp*-SK/ATP/Mg⁺²/**4d** complex. E. *Hp*-SK/ATP/Mg⁺²/**4f** complex.



Figure S3. The rmsd plots for the protein backbone (C α , C, N and O atoms) (orange line) and ligand (blue line) calculated for the *Mt*-SK/ATP/Mg⁺²/ligand complex obtained from MD simulation studies. A. *Mt*-SK/ATP/Mg⁺²/**4a** complex. B. *Mt*-SK/ATP/Mg⁺²/**4b** complex. C. *Mt*-SK/ATP/Mg⁺²/**4c** complex. D. *Mt*-SK/ATP/Mg⁺²/**4d** complex. E. *Mt*-SK/ATP/Mg⁺²/**4f** complex.



Figure S4. Detailed view of the binding mode of the most potent compounds against *Hp*-SK (A, compound **4f**) and *Mt*-SK (B, compound **4a**) in *Hp*-SK/ATP/Mg⁺²/**4f** and *Mt*-SK/ATP/Mg⁺²/**4a** enzyme complexes, respectively. Two perspectives are provided. Snapshots after 80 ns of simulation are shown. Relevant side chain residues are shown and labelled. Polar contacts are shown as blue dashed lines. The indirect interaction of the oxygen atom of the substituent in the ether bridge in **4f** with the Val44 side chain through a water network is highlighted with blue shading. The proximity of the methyl group in **4a** to the carbon chain of the catalytic residue Arg117 is shown as yellow shading.



Time (ns)





Time (ns)



Figure S5. Variation of the dihedral angle shown in red in ligands **4b** (A), **4c** (B), **4d** (C) and **4f** (D) calculated for the *Hp*-SK/ATP/Mg⁺²/ligand complex obtained from MD simulation studies.









Figure S6. Variation of the dihedral angle shown in red in ligands **4b** (A), **4c** (B), **4d** (C) and **4f** (D) calculated for the *Mt*-SK/ATP/Mg⁺²/ligand complex obtained from MD simulation studies.

$\leftarrow \textbf{P} \quad \textbf{Loop} \rightarrow$

				← SB	domain \rightarrow		
	10) 20) 30) 4	10 50		
Mt-SK	-MAPKAV <mark>L</mark>	V <mark>G</mark> LP <mark>G</mark> S <mark>GKS</mark> T	IGRRLAKALG	VGLL <mark>D</mark> T <mark>D</mark> VA <mark>I</mark>	EQRTGRSIAD		
Hp-SK	MQHLV <mark>L</mark>	I <mark>G</mark> FM <mark>G</mark> S <mark>GKS</mark> S	LAQELGLALK	LEVL <mark>D</mark> T <mark>D</mark> MI <mark>I</mark>	SERVGLSVRE		
Hp-SK*	MQHLV <mark>L</mark>	I <mark>G</mark> FM <mark>G</mark> S <mark>GKS</mark> S	LAQELGLALK	levl <mark>d</mark> t <mark>d</mark> mi	SERVGLSVRE		
Ec-SK	-MAEKRNIF <mark>l</mark>	V <mark>G</mark> PM <mark>G</mark> AGKST	IGRQLAQQLN	MEFY <mark>D</mark> S <mark>D</mark> QE <mark>I</mark>	EKRTGADVGW		
Sa-SK	MNHDKSPII <mark>l</mark>	I <mark>G</mark> FM <mark>G</mark> TGKST	IGKYVADEQN	LSFI <mark>D</mark> I <mark>D</mark> SY <mark>I</mark>	EEKYKLTIPE		
St-SK	-MAEKRNIF <mark>L</mark>	V <mark>G</mark> PM <mark>G</mark> A <mark>GKS</mark> T	IGRQLAQQLN	MEFY <mark>D</mark> S <mark>D</mark> QE <mark>1</mark>	EKRTGADVGW		
	<i>←SB domain→</i>						
•	60) 7() 80)	0 100		
Mt-SK	I FATD <mark>GE</mark> QEF	RRIEEDVVRA	ALADHD-GVL	SL <mark>GGG</mark> AVTSI	? GVRAA <mark>L</mark> AGH-		
Hp-SK	IFEEL <mark>GE</mark> DNF	RMFEKNLIDE	LKTLKTPHVI	ST <mark>GGG</mark> IVMHE	E N <mark>L</mark> KGLG		
Hp-SK*	IFEEL <mark>GE</mark> DNF	RMFEKNLIDE	LKTLKTPHVI	ST <mark>GGG</mark> IVMHE	E N <mark>L</mark> KGLG		
Ec-SK	VFDLEGEEGF	RDREEKVINE	LTEKQG-IVL	ATGGGSVKSF	R ETRNR <mark>L</mark> SARG		
Sa-SK	IFCKHGEQYF	RNLEFTCLQE	CINTADII	ATGGGIIESE	E EAFNF <mark>L</mark> KNQK		
St-SK	v <mark>f</mark> dve <mark>ge</mark> dg <mark>f</mark>	RNREEKVINE	L'I'EKQG-IVL	AT <mark>GGG</mark> SVKSI	R ETRNR <mark>L</mark> SARG		
	$\leftarrow \textit{LID domain} \rightarrow $						
	 11() 120) 130	···· ····) 14	···· ···· 10 150		
Mt-SK	TVVY <mark>l</mark> eisaa	EGVR <mark>R</mark> TGGNT	V <mark>RP</mark> LLAGP	DR-AEKYRAI	l makrapl <mark>y</mark> rr		
Hp-SK	TTFY <mark>L</mark> KMDFE	TLIK <mark>R</mark> LNQKE	REK <mark>RP</mark> LLNNL	TQAKELFE	KRQAL <mark>Y</mark> EK		
Hp-SK*	TTFY <mark>L</mark> K <mark>I</mark> DFE	TLIK <mark>R</mark> LNQKE	REK <mark>RP</mark> LLNNL	AQAKELFE	KRQAL <mark>Y</mark> EK		
Ec-SK	VVVY <mark>L</mark> ETTIE	KQLA <mark>R</mark> TQRDK	K <mark>RP</mark> LLHVE	TPPREVLEAI	L ANERNPL <mark>Y</mark> EE		
Sa-SK	NIIW <mark>L</mark> DCNID	IIYS <mark>R</mark> INDDP	H <mark>RP</mark> NAN	NKTIKQLNDI	L YCSRNLR <mark>Y</mark> NE		
St-SK	VVVY <mark>L</mark> ETTIE	KQLA <mark>R</mark> TQRDK	K <mark>RP</mark> LLQVE	APPREVLEAI	l anernpl <mark>y</mark> ee		
	160) 170) 180)			
Mt-SK	V <mark>A</mark> TMRVDTNR	RNPGAVVRHI	LSRLQVPSPS	EAAT			
Hp-SK	N <mark>A</mark> SFIIDARG	-GLNNSLKQV	LQFIA				
Hp-SK*	N <mark>A</mark> SFIIDARG	-GLNNSLKQV	LQFIA				
Ec-SK	I <mark>A</mark> DVTIRTDD	QSAKVVANQI	IHMLESN				
Sa-SK	I <mark>A</mark> FKKFDSHL	LSISEIYYEL	LNLIKASDQY				
St-SK	I <mark>A</mark> DVTIRTDD	QSAKVVANQI	IHMLESN				

Figure S7. Amino acid sequence alignments for *M. tuberculosis, H. pylori, E. coli, S. aureus* and *S. typhi* shikimate kinase. For the *H. pylori* enzyme, two strains are shown: AROK_HELPJ, Q9ZMS3 (strain J99 / ATCC 700824) and clinically isolated strain from Guadalajara Hospital (Spain), which was employed in these studies. Protein sequences were aligned using the CLUSTAL Omega multiple sequence alignment (<u>http://www.ebi.ac.uk/Tools/msa/clustalo/</u>). Changed residues in the clinically isolated strain used in these studies are highlighted in red. Conserved residues are highlighted in yellow. The P loop, SB and LID domains are also indicated.



Figure S8. Superposition of several snapshots of Hp-SK/ATP/Mg⁺²/**5S** (A) and Mt-SK/ATP/Mg⁺²/**5S** (B) complexes. Note how the methyl group would interact with the side chain of apolar residues of the lid and P-loop (Leu118 and Met10 for Hp-SK and Leu119, Leu120 and Pro11 for Mt-SK). No interaction with the carbon chain of the essential arginine was identified.





























Figure S9. Dixon plots of compounds 2–5 against *Hp*-SK.



Figure S10. Dixon plots of compounds 4–5 against *Mt*-SK.