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Discovery of novel sulphonamide hybrids that inhibit LSD1 against bladder cancer cells

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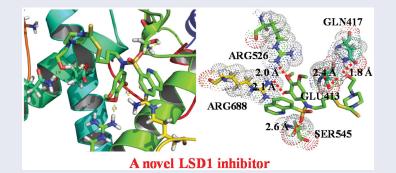
ABSTRACT

Aim:A series of sulphonamide hybrids were designed, synthesised, and identified as potential lysine-specific demethylase 1 (LSD1) inhibitors.

Materials and methods: Bladder cancer cell lines were cultured to evaluate the antiproliferative activity. Inhibitory evaluation of sulphonamide hybrids against LSD1 were performed.

Conclusion: sulphonamide derivative **L8** exhibited the antiproliferative activity against HTB5, HTB3, HT1376, and HTB1 cells with IC_{50} values of 1.87, 0.18, 0.09, and 0.93 μ M, respectively. Compound **L8** as a selective and reversible LSD1 inhibitor could inhibit LSD1 with the IC_{50} value of 60 nM. It effectively inhibited LSD1 by increasing the expression levels of H3K4me1, H3K4me2, and H3K9me2 in HT1376 cells. To the best of our knowledge, this was the first report which showed that sulphonamide–quinoline–dithiocarbamate hybrids potently inhibited LSD1 in bladder cancer cells. Our studies give the potential application of the sulphonamide–based scaffold for developing LSD1 inhibitors to treat bladder cancer.

GRAPHICAL ABSTRACT



HIGHLIGHTS

- 1. L8 exhibits the antiproliferative activity against bladder cancer cells.
- 2. L8 is a selective and reversible LSD1 inhibitor.
- 3. L8 increases the expression levels of H3K4me1, H3K4me2 and H3K9me2.

Introduction

Bladder cancer is the most frequently diagnosed malignancy in the urinary system and has the high morbidity and mortality rates¹. Chemotherapy plays an important role in the treatment of bladder cancer and it is urgent to develop potent anti-bladder cancer drugs^{2,3}. Histone lysine-specific demethylase 1 (LSD1) could catalyse the demethylation of mono and dimethylated H3K4me1/ 2 or H3K9me1/2 and demethylate many other nonhistone substrates⁴. LSD1 is aberrantly expressed in many malignant tumours such as prostate, ovarian, gastric, liver, breast, lung, bladder, neuroblastoma, and blood cancers⁵. The inhibition of LSD1 could prevent tumour cell proliferation, stimulate antitumor immunity, and enhance antitumor efficacy of immune checkpoint blockade⁶. Therefore, LSD1 has been considered as a potential cancer therapeutic target to discover novel anti-bladder cancer agents^{7–9}. LSD1 and MAO-A/-B were belonged to the monoamine oxidase family, and MAO-A/-B shared the similar enzymatic mechanisms and the same cofactor of LSD1 in the cleavage of the inactivated carbon–nitrogen bonds from their substrates¹⁰. Although a variety of LSD1 inhibitors have been reported to date, many of them show insufficient selectivity towards LSD1¹¹.

Sulphonamide has been proven to be an interesting scaffold and many sulphonamide derivatives are designed as potent antitumor agents for cancer therapy¹². Phenylpropanoid-based sulphonamide **1** (Figure 1) induced cell cycle arrest at G1/S transition

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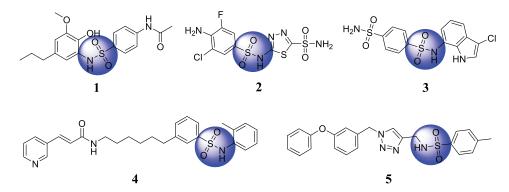


Figure 1. Chemical structures of anticancer sulphonamide derivatives.

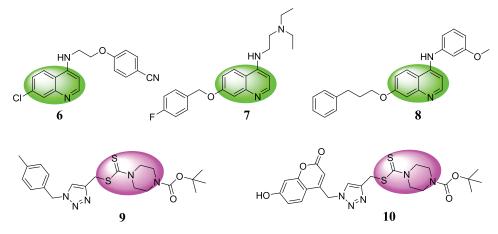


Figure 2. Chemical structures of anticancer quinolines and dithiocarbamate based LSD1 inhibitors.

by reducing the expression levels of cyclin D1 and cyclin E in MCF7 cells¹³. Sulphonamide **2** as a potential antitumor agent was a novel tumour-associated isozyme carbonic anhydrase IX inhibitor¹⁴. Sulphonamide **3** showed a significant antitumor activity against HCT116 human colon carcinoma *in vitro* and *in vivo*¹⁵. *Trans*-3-(pyridin-3-yl)acrylamide-derived sulphamide **4** showed a single-digit nanomolar antiproliferative activity against DU145, Hela, and H1975 cells and inhibited NAMPT with an IC₅₀ value of 5.08 nM¹⁶. Benzenesulphonamide **5** showed the potent inhibitory effects against PC-3 cells with an IC₅₀ value of 4.08 μ M as a potential tubulin polymerisation inhibitor¹⁷.

Quinolines and dithiocarbamates also represent a large group of anticancer agents and there have been many studies on the structural modification based on quinolines or dithiocarbamates^{18,19}. Quinoline **6** (Figure 2) showed the potently inhibitory activity against Raji cells by inducing changes in the cell cycle²⁰. Quinoline derivative **7** triggered p53/Bax-dependent apoptosis by activating p53 transcriptional activity against colorectal cancer HCT-116 cells²¹. Quinoline **8** was a potent antiproliferative agent against HCT-116, RKO, A2780, and Hela cell lines with IC₅₀ values of 2.56, 3.67, 3.46, and 2.71 μ M, respectively²². Dithiocarbamate **9** showed the potent cytotoxicity against MGC-803 and HGC-27 cells by the specific and robust inhibition of LSD1²³. Dithiocarbamate **10** displayed the potent and reversible inhibition against LSD1 with an IC₅₀ value of 0.39 μ M²⁴. So, dithiocarbamate moiety might be a promising fragment to design novel LSD1 inhibitors.

Molecular hybridisation is a useful strategy in anticancer drug design based on the combination of different bioactive scaffolds to produce a new molecular architecture²⁵. Based on these findings, we proposed that the sulphonamide derivatives with

quinoline and dithiocarbamate groups might have the excellent anticancer activity and LSD1 inhibitory activity. Thus, in this study, we designed a series of sulphonamide–quinoline–dithiocarbamate hybrids *via* a molecular hybridisation strategy and evaluated for their antiproliferative activity and inhibitory activity against LSD1.

Experimental

Materials and methods

General procedure for the preparation of J1~J7

A solution of aromatic amine (**11** or **13**, 3.5 mmol) and triethylamine (5.25 mmol) in acetone (10 ml) was cooled to 0 °C. Sulphonyl chloride derivative (**12** or **14**, 5.25 mmol) in acetone (10 ml) was added drop-wise into the solution. After being stirred for 30 min, the reaction mixture was stirred at room temperature for 8 h. The reaction mixture was washed with water (50 ml) and extracted with ethyl acetate (30 ml). The organic extracts were purified by column chromatography on silica (petroleum ether/ ethyl acetate = 10/1) to yield compounds J1~J7.

General procedure for the preparation of K1~K3

To a solution of sulphanilamide **J7** (1.2 mmol) and potassium hydroxide (1.5 mmol) in dichloromethane (6 ml) was added 1,2-dibromoethane or 1,3-dibromopropane or 1,4-dibromobutane (1.5 mmol). The reaction mixture was stirred at $60 \degree C$ for 10 h. The reaction was diluted with brine (45 ml) and extracted with dichloromethane (25 ml). The organic extracts were purified by

column chromatography on silica (petroleum ether/ethyl acetate = 9/1) to yield compounds **K1~K3**.

General procedure for the preparation of L1~L8

A solution of sulphanilamide intermediates **K1~K3** (2 mmol), carbon disulphide (3 mmol), trisodium phosphate dodecahydrate (0.5 mmol), and different piperazine derivatives (3 mmol) in acetone (10 ml) was stirred at room temperature overnight. Solvents were removed and crude products were purified by column chromatography on silica (petroleum ether/ethyl acetate = 8/1) to yield compounds **L1~L8**. The structures of all synthesised compounds **J1~J7**, **K1~K3**, and **L1~L8** were characterised by ¹H NMR, ¹³C NMR, and HRMS. Detailed analytical information was listed in Supporting Information.

3,4-Dimethoxy-N-(quinolin-8-yl)benzenesulphonamide (J1)

White solid, yield:80%, m.p.:1 6 $7 \sim 169$ °C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.84 (s, 1H), 8.88 (dd, J= 4.2, 1.6 Hz, 1H), 8.36 (dd, J= 8.3, 1.6 Hz, 1H), 7.71 (dd, J= 7.6, 1.1 Hz, 1H), 7.68–7.63 (m, 1H), 7.59 (dd, J= 8.3, 4.2 Hz, 1H), 7.56–7.47 (m, 2H), 7.44 (d, J= 2.2 Hz, 1H), 7.00 (d, J= 8.5 Hz, 1H), 3.74 (s, 3H), 3.69 (s, 3H). ¹³C NMR (100 MHz, DMSO-d₆) δ 152.34, 149.30, 148.45, 138.67, 136.51, 133.76, 130.76, 128.05, 126.66, 122.77, 122.31, 120.84, 116.35, 110.91, 109.67, 55.72, 55.68. HRMS (m/z): Calcd. C₁₇H₁₇N₂O₄S, [M + H]⁺ m/z: 345.0909, found: 345.0914.

4-Methoxy-N-(quinolin-8-yl)benzenesulphonamide (J2)

White solid, yield:87%, m.p.:1 6 $1 \sim 163 \,^{\circ}$ C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.79 (s, 1H), 8.87 (dd, J= 4.2, 1.6 Hz, 1H), 8.35 (dd, J= 8.3, 1.6 Hz, 1H), 7.98–7.80 (m, 2H), 7.77–7.41 (m, 4H), 7.09–6.91 (m, 2H), 3.74 (s, 3H). ¹³C NMR (100 MHz, DMSO-d₆) δ 162.62, 149.30, 138.49, 136.51, 133.67, 130.83, 129.17, 128.03, 126.66, 122.67, 122.33, 115.94, 114.27, 55.57. HRMS (m/z): Calcd. C₁₆H₁₅N₂O₃S, [M + H]⁺ m/z: 315.0803, found: 315.0808.

4-(Tert-butyl)-N-(quinolin-8-yl)benzenesulphonamide (J3)

White solid, yield:78%, m.p.:1 4 $2 \sim 144 \,^{\circ}$ C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.91 (s, 1H), 8.86 (dd, J= 4.2, 1.6 Hz, 1H), 8.35 (dd, J= 8.3, 1.6 Hz, 1H), 7.92 – 7.80 (m, 2H), 7.71 (dd, J= 7.6, 1.1 Hz, 1H), 7.65 (dd, J= 8.2, 1.0 Hz, 1H), 7.59 (dd, J= 8.3, 4.2 Hz, 1H), 7.55 – 7.46 (m, 3H), 1.20 (s, 9H). ¹³C NMR (100 MHz, DMSO-d₆) δ 149.34, 139.41, 138.76, 136.48, 133.53, 133.07, 129.10, 128.06, 126.85, 126.62, 123.07, 122.30, 116.77. HRMS (m/z): Calcd. C₁₉H₂₁N₂O₂S, [M + H]⁺ m/z: 341.1324, found: 341.1329.

N-(quinolin-8-yl)benzenesulphonamide (J4)

White solid, yield:89%, m.p.:1 6 $5 \sim 167 \,^{\circ}$ C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.99 (s, 1H), 8.86 (dd, J= 4.2, 1.6 Hz, 1H), 8.35 (dd, J= 8.3, 1.5 Hz, 1H), 8.05–7.86 (m, 2H), 7.79–7.62 (m, 2H), 7.62–7.39 (m, 5H). ¹³C NMR (100 MHz, DMSO-d₆) δ 156.15, 149.28, 138.54, 136.61, 136.49, 133.64, 128.07, 126.78, 126.68, 125.98, 122.75, 122.31, 116.02, 34.78, 30.62. HRMS (m/z): Calcd. C₁₅H₁₃N₂O₂S, [M + H]⁺ m/z: 285.0698, found: 285.0699.

4-Methyl-N-(quinolin-8-yl)benzenesulphonamide (J5)

White solid, yield:84%, m.p.:1 4 $8 \sim 150$ °C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.88 (s, 1H), 8.87 (dd, J= 4.2, 1.5 Hz, 1H), 8.35 (dd, J= 8.3, 1.5 Hz, 1H), 7.81 (d, J= 8.3 Hz, 2H), 7.75–7.43 (m, 4H), 7.28

(d, J= 8.1 Hz, 2H), 2.27 (s, 3H). ¹³C NMR (100 MHz, DMSO-d₆) δ 149.32, 143.56, 138.53, 136.52, 136.46, 133.58, 129.57, 128.04, 126.93, 126.65, 122.80, 122.33, 116.13, 20.86. HRMS (m/z): Calcd. C₁₆H₁₅N₂O₂S, [M + H]⁺ m/z: 299.0854, found: 299.0858.

4-Bromo-N-(quinolin-8-yl)benzenesulphonamide (J6)

White solid, yield:92%, m.p.:1 9 $2 \sim 194 \,^{\circ}$ C. ¹H NMR (400 MHz, DMSO-d₆) δ 10.19 (s, 1H), 8.85 (dd, J= 4.2, 1.6 Hz, 1H), 8.36 (dd, J= 8.3, 1.5 Hz, 1H), 7.83 (d, J= 8.6 Hz, 2H), 7.73–7.67 (m, 4H), 7.63–7.41 (m, 2H). ¹³C NMR (100 MHz, DMSO-d₆) δ 149.41, 139.09, 138.87, 136.49, 133.35, 132.14, 128.90, 128.14, 126.90, 126.63, 123.51, 122.29, 117.68. HRMS (m/z): Calcd. C₁₅H₁₂BrN₂O₂S, [M + H]⁺ m/z: 362.9803, found: 362.9807.

N-(3,4,5-trimethoxyphenyl)quinoline-8-sulphonamide (J7)

White solid, yield:94%, m.p.:1 8 $1 \sim 183$ °C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.87 (s, 1H), 9.16 (dd, J= 4.2, 1.7 Hz, 1H), 8.52 (dd, J= 8.4, 1.7 Hz, 1H), 8.39 (dd, J= 7.3, 1.3 Hz, 1H), 8.27 (dd, J= 8.2, 1.3 Hz, 1H), 7.81–7.60 (m, 2H), 6.35 (s, 2H), 3.51 (s, 6H), 3.47 (s, 3H). ¹³C NMR (100 MHz, DMSO-d₆) δ 152.59, 151.40, 142.72, 136.98, 135.12, 134.23, 133.92, 133.57, 132.25, 128.34, 125.63, 122.58, 98.02, 59.91, 55.56. HRMS (m/z): Calcd. C₁₈H₁₉N₂O₅S, [M + H]⁺ m/z: 375.1015, found: 375.1019.

N-(2-bromoethyl)-N-(3,4,5-trimethoxyphenyl)quinoline-8-sulphonamide (K1)

White solid, yield:72%, m.p.:1 5 8 ~ 160 °C. ¹H NMR (400 MHz, CDCl₃) δ 9.11 (dd, *J*= 4.2, 1.8 Hz, 1H), 8.21 (dd, *J*= 8.6, 1.5 Hz, 2H), 7.94 (dd, *J*= 8.2, 1.3 Hz, 1H), 7.66–7.36 (m, 2H), 6.09 (s, 2H), 4.49 (t, *J*= 7.5 Hz, 2H), 3.67 (s, 3H), 3.48 (t, *J*= 7.5 Hz, 2H), 3.40 (s, 6H). ¹³ C NMR (100 MHz, CDCl₃) δ 152.07, 150.20, 143.11, 136.85, 135.75, 135.63, 133.29, 133.13, 132.51, 127.65, 124.69, 121.11, 105.47, 59.75, 54.85, 53.99, 29.17. HRMS (m/z): Calcd. C₂₀H₂₂BrN₂O₅S, [M + H]⁺ m/z: 481.0433, found: 481.0439.

N-(3-bromopropyl)-*N*-(3,4,5-trimethoxyphenyl)quinoline-8-sulphonamide (K2)

White solid, yield:81%, m.p.:1 2 9 ~ 130 °C. ¹H NMR (400 MHz, CDCl₃) δ 9.09 (dd, *J*= 4.2, 1.7 Hz, 1H), 8.20 (td, *J*= 8.6, 1.5 Hz, 2H), 7.93 (dd, *J*= 8.2, 1.2 Hz, 1H), 7.62–7.32 (m, 2H), 6.12 (s, 2H), 4.22 (t, *J*= 6.8 Hz, 2H), 3.67 (s, 3H), 3.47 (t, *J*= 6.8 Hz, 2H), 3.42 (d, *J*= 8.3 Hz, 6H), 2.09 (p, *J*= 6.8 Hz, 2H). ¹³C NMR (100 MHz, CDCl₃) δ 152.00, 150.12, 143.22, 136.55, 135.78, 135.56, 133.61, 133.14, 132.46, 127.67, 124.63, 121.03, 105.19, 59.77, 54.89, 51.02, 31.75, 29.45. HRMS (m/z): Calcd. C₂₁H₂₄BrN₂O₅S, [M + H]⁺ m/z: 495.0589, found: 495.0593.

N-(4-bromobutyl)-*N*-(3,4,5-trimethoxyphenyl)quinoline-8-sulphonamide (K3)

White solid, yield:96%, m.p.:1 0 $3 \sim 104 \,^{\circ}$ C. ¹H NMR (400 MHz, CDCl₃) δ 9.08 (dd, J= 4.2, 1.8 Hz, 1H), 8.19 (ddd, J= 6.0, 3.9, 1.6 Hz, 2H), 7.92 (dd, J= 8.2, 1.3 Hz, 1H), 7.60–7.31 (m, 2H), 6.08 (s, 2H), 4.12 (t, J= 6.8 Hz, 2H), 3.67 (s, 3H), 3.53–3.33 (m, 8H), 2.01 (dt, J= 14.4, 6.6 Hz, 2H), 1.64 (dt, J= 14.1, 6.9 Hz, 2H). ¹³C NMR (100 MHz, CDCl₃) δ 151.95, 150.08, 143.25, 136.54, 136.01, 135.52, 133.50, 133.06, 132.31, 127.64, 124.63, 120.98, 105.48, 59.76, 54.87, 51.59, 32.67, 28.66, 26.64. HRMS (m/z): Calcd. C₂₂H₂₆BrN₂O₅S, [M + H]⁺ m/z: 509.0746, found: 509.0748.

Tert-butyl-4-(((2-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)ethyl)thio)carbonothioyl)piperazine-1-carboxylate (L1)

White solid, yield:62%, m.p.:1 2 $1 \sim 123 \,^{\circ}$ C. ¹H NMR (400 MHz, CDCl₃) δ 9.07 (dd, *J*= 4.2, 1.8 Hz, 1H), 8.21 (ddd, *J*= 10.1, 7.9, 1.5 Hz, 2H), 7.93 (dd, *J*= 8.2, 1.3 Hz, 1H), 7.58–7.34 (m, 2H), 6.17 (s, 2H), 4.41 (t, *J*= 7.0 Hz, 2H), 4.34–4.06 (m, 2H), 4.04–3.76 (m, 2H), 3.67 (s, 3H), 3.52 (t, *J*= 7.0 Hz, 2H), 3.47 (dd, *J*= 10.0, 5.0 Hz, 4H), 3.44 (s, 6H), 1.41 (s, 9H). ¹³C NMR (100 MHz, CDCl₃) δ 195.89, 153.45, 151.90, 150.18, 143.16, 136.51, 135.93, 135.52, 133.33, 133.03, 132.43, 127.64, 124.62, 121.01, 105.28, 79.58, 59.77, 54.91, 50.77, 35.22, 27.35. HRMS (m/z): Calcd. C₃₀H₃₉N₄O₇S₃, [M + H]⁺ m/z: 663.1981, found: 663.1987.

Tert-butyl-4-(((3-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)propyl)thio)carbonothioyl)piperazine-1-carboxylate (L2)

White solid, yield:91%, m.p.:1 3 0 ~ 132 °C. ¹H NMR (400 MHz, CDCl₃) δ 9.09 (dd, *J*= 4.2, 1.8 Hz, 1H), 8.35–8.06 (m, 2H), 7.92 (dd, *J*= 8.2, 1.3 Hz, 1H), 7.63–7.35 (m, 2H), 6.13 (s, 2H), 4.20 (t, *J*= 6.7 Hz, 4H), 3.91 (s, 2H), 3.67 (s, 3H), 3.51–3.44 (m, 4H), 3.45–3.36 (m, 8H), 2.00–1.85 (m, 2H), 1.40 (s, 9H). ¹³C NMR (100 MHz, CDCl₃) δ 196.61, 153.47, 151.94, 150.17, 143.24, 136.50, 135.97, 135.51, 133.57, 133.07, 132.34, 127.64, 124.61, 121.01, 105.45, 79.56, 59.75, 54.90, 51.54, 33.18, 27.76, 27.35. HRMS (m/z): Calcd. C₃₁H₄₁N₄O₇S₃, [M + H]⁺ m/z: 677.2137, found: 677.2139.

2-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)ethyl-4-(2hydroxyethyl)piperazine-1-carbodithioate (L3)

White solid, yield:89%, m.p.:1 2 $2 \sim 124$ °C. ¹H NMR (400 MHz, DMSO-d₆) δ 9.16 (dd, J= 4.1, 1.5 Hz, 1H), 8.57 (dd, J= 8.4, 1.4 Hz, 1H), 8.29 (d, J= 7.5 Hz, 1H), 8.21 (d, J= 7.3 Hz, 1H), 7.75 (dd, J= 8.3, 4.2 Hz, 1H), 7.65 (t, J= 7.8 Hz, 1H), 6.24 (s, 2H), 4.48 (t, J= 5.3 Hz, 1H), 4.35 (t, J= 6.8 Hz, 2H), 4.19 (s, 2H), 3.86 (s, 2H), 3.55 (d, J= 6.5 Hz, 3H), 3.52 (dd, J= 11.5, 5.9 Hz, 2H), 3.42 (s, 6H), 3.39 (d, J= 6.8 Hz, 2H), 2.50–2.46 (m, 4H), 2.43 (t, J= 6.1 Hz, 2H). ¹³C NMR (100 MHz, DMSO-d₆) δ 194.23, 152.42, 151.54, 143.26, 137.00, 136.75, 136.02, 134.24, 133.97, 133.38, 128.49, 125.67, 122.56, 106.00, 59.95, 59.45, 58.46, 55.55, 52.47, 50.94, 35.54. HRMS (m/z): Calcd. C₂₇H₃₅N₄O₆S₃, [M + H]⁺ m/z: 607.1719, found: 607.1723.

3-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)propyl-4-(2hydroxyethyl)piperazine-1-carbodithioate (L4)

White solid, yield:85%, m.p.:1 0 6 ~ 107 °C. ¹H NMR (400 MHz, CDCl₃) δ 9.10 (d, *J*= 2.1 Hz, 1H), 8.20 (dd, *J*= 6.8, 3.9 Hz, 2H), 7.93 (d, *J*= 8.1 Hz, 1H), 7.65–7.34 (m, 2H), 6.12 (s, 2H), 4.33 (s, 2H), 4.20 (t, *J*= 6.6 Hz, 2H), 3.96 (s, 2H), 3.67 (s, 3H), 3.66–3.57 (m, 2H), 3.42 (s, 8H), 2.73–2.49 (m, 6H), 1.93 (dd, *J*= 13.7, 6.8 Hz, 2H). ¹³C NMR (100 MHz, CDCl₃) δ 196.25, 151.89, 150.19, 143.20, 136.34, 135.86, 135.53, 133.54, 133.08, 132.36, 127.61, 124.62, 121.02, 105.31, 59.76, 58.16, 56.69, 54.85, 51.52, 51.28, 33.21, 27.73. HRMS (m/z): Calcd. C₂₈H₃₇N₄O₆S₃, [M + H]⁺ m/z: 621.1875, found: 621.1879.

4-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)butyl-4-ethylpiperazine-1-carbodithioate (L5)

White solid, yield:85%, m.p.:1 1 $0 \sim 112^{\circ}$ C. ¹H NMR (400 MHz, CDCl₃) δ 9.08 (dd, J= 4.2, 1.8 Hz, 1H), 8.28–8.09 (m, 2H), 7.92 (dd, J= 8.2, 1.3 Hz, 1H), 7.61–7.33 (m, 2H), 6.09 (s, 2H), 4.28 (s, 2H), 4.10 (t, J= 7.0 Hz, 2H), 3.90 (s, 2H), 3.68 (s, 3H), 3.43 (s, 6H), 3.26 (t, J= 7.4 Hz, 2H), 2.60–2.42 (m, 4H), 2.39 (q, J= 7.2 Hz, 2H), 1.79 (dd, J= 14.9, 8.0 Hz, 2H), 1.67–1.59 (m, 2H), 1.04 (t, J= 7.2 Hz, 3H). ¹³C

NMR (100 MHz, CDCl₃) δ 195.96, 151.90, 150.12, 143.25, 136.45, 136.14, 135.46, 133.62, 132.95, 132.25, 127.64, 124.58, 120.97, 105.52, 59.75, 54.88, 52.07, 51.15, 50.89, 35.65, 27.66, 25.04, 10.96. HRMS (m/z): Calcd. $C_{29}H_{39}N_4O_5S_3, \ [M+H]^+ \ m/z: \ 619.2083, found: 619.2087.$

Tert-butyl-4-(((4-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)butyl)thio)carbonothioyl)piperazine-1-carboxylate (L6)

White solid, yield:74%, m.p.:9 8 ~ 100 °C. ¹H NMR (400 MHz, CDCl₃) δ 9.08 (dd, *J*= 4.2, 1.7 Hz, 1H), 8.31–8.13 (m, 2H), 7.92 (dd, *J*= 8.2, 1.2 Hz, 1H), 7.59–7.38 (m, 2H), 6.09 (s, 2H), 4.10 (t, *J*= 7.0 Hz, 6H), 3.67 (s, 3H), 3.54–3.46 (m, 4H), 3.42 (s, 6H), 3.27 (t, *J*= 7.4 Hz, 2H), 1.86–1.75 (m, 2H), 1.66–1.58 (m, 2H), 1.41 (s, 9H). ¹³C NMR (100 MHz, CDCl₃) δ 157.80, 153.47, 151.90, 150.11, 143.23, 136.46, 136.09, 135.47, 133.59, 132.97, 132.27, 127.65, 124.61, 120.97, 105.49, 79.55, 59.75, 54.87, 52.07, 35.65, 27.62, 27.35, 24.95. HRMS (m/z): Calcd. C₃₂H₄₃N₄O₇S₃, [M + H]⁺ m/z: 691.2294, found: 691.2298.

4-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)butyl-4methylpiperazine-1-carbodithioate (L7)

White solid, yield:69%, m.p.:1 3 $5 \sim 137 \,^{\circ}$ C. ¹H NMR (400 MHz, DMSO) δ 9.17 (dd, J= 4.2, 1.7 Hz, 1H), 8.56 (dd, J= 8.4, 1.6 Hz, 1H), 8.23 (ddd, J= 42.3, 7.8, 1.2 Hz, 2H), 7.78-7.59 (m, 2H), 6.17 (s, 2H), 4.23 (s, 2H), 4.08 (t, J= 6.7 Hz, 2H), 3.88 (s, 2H), 3.56 (s, 3H), 3.41 (s, 6H), 3.24 (t, J= 7.3 Hz, 2H), 2.44-2.29 (m, 4H), 2.21 (s, 3H), 1.72 (dd, J= 14.7, 7.7 Hz, 2H), 1.52 (dd, J= 14.5, 6.9 Hz, 2H). ¹³C NMR $(100 \text{ MHz}, \text{ DMSO-d}_6) \delta$ 195.24, 152.43, 151.46, 143.31, 136.92, 136.71, 136.27, 134.27, 134.04, 133.23, 128.48, 125.62, 122.49, 106.15, 59.93, 55.52, 53.94, 52.16, 45.05, 35.70, 27.83, 25.74. HRMS (m/z): Calcd. $C_{28}H_{37}N_4O_5S_3$, $[M + H]^+$ m/z: 605.1926, found: 605.1929.

4-(N-(3,4,5-trimethoxyphenyl)quinoline-8-sulfonamido)butyl-4–(2hydroxyethyl)piperazine-1-carbodithioate (L8)

White solid, yield:73%, m.p.:9 $4 \sim 96$ °C. ¹H NMR (400 MHz, CDCl₃) δ 9.08 (dd, *J*= 4.1, 1.6 Hz, 1H), 8.19 (dd, *J*= 10.7, 3.9 Hz, 2H), 7.92 (d, *J*= 7.3 Hz, 1H), 7.53–7.40 (m, 2H), 6.09 (s, 2H), 4.28 (s, 2H), 4.10 (t, *J*= 7.0 Hz, 2H), 3.88 (d, *J*= 20.8 Hz, 2H), 3.67 (s, 3H), 3.64–3.57 (m, 2H), 3.42 (s, 6H), 3.26 (t, *J*= 7.4 Hz, 2H), 2.53 (dd, *J*= 6.2, 3.8 Hz, 6H), 1.80 (dd, *J*= 14.8, 7.7 Hz, 2H), 1.65–1.55 (m, 2H). ¹³C NMR (100 MHz, CDCl₃) δ 196.27, 151.91, 150.12, 143.25, 136.45, 136.13, 135.47, 133.61, 132.97, 132.25, 127.64, 124.60, 120.97, 105.52, 59.76, 58.04, 56.96, 54.88, 52.08, 51.32, 35.69, 27.63, 25.01. HRMS (m/z): Calcd. C₂₉H₃₉N₄O₆S₃, [M+H]⁺ m/z: 635.2032, found: 635.2037.

Inhibitory evaluation of compounds J1 \sim J7, K1 \sim K3, and L1 \sim L8 against LSD1 and MAO-A/B

LSD1 inhibitor screening assay kit was purchased to evaluate for the enzyme inhibition activity of all synthetic compounds (Cayman Chemical, Fort Annapolis, MI). The MAO-A/B biochemical kit was purchased to explore the inhibitory activity against MAO-A and MAO-B (Promega, Madison, WI). First, compounds **J1~J7**, **K1~K3**, and **L1~L8** were diluted with dimethyl sulphoxide to required concentrations (Innochem, Beijing, China). In a 48-well black plate, a buffer solution of compounds and reagent was prepared and incubated at 37 °C. Then, horseradish peroxidase solution and fluorometric substrate solution were added and incubated at 37 °C. The signal of each well was read by a GloMax Explorer (Beijing Yuanpinghao Biotechnology Co., Ltd, Beijing, China).

Cell culture and cell viability assay

Cell lines (Bel7402, H1299, HCT-15, SGC-7901, HTB3, Jurkat, LX-2, HPAEpiC, HIEC, HTB5, HT1376, and HTB1) were obtained from Shanghai Yuanye Biotechnology Co., Ltd (Shanghai, China). Cell lines were cultured in 89% Roswell Park Memorial Institute 1640 medium supplemented with 10% foetal bovine serum and 1% antibiotic at 37 °C in a humidified CO2 incubator (Shanghai Yuanye Biotechnology Co., Ltd, Shanghai, China). HTB3&shControl, HTB3&shLSD1, HT1376&shControl, and HT1376&shLSD1 cells were supported by the 4th affiliated Hospital of China Medical University. HTB3&shControl, HTB3&shLSD1, HT1376&shControl, and HT1376&shLSD1 cells were cultured in a Roswell Park Memorial Institute 1640 medium with geneticin (Innochem, Beijing, China). Cells were cultured at a 96-well plate for 24 h (Cayman Chemical, Fort Annapolis, MI) and different compounds were added. Thiazolyl blue tetrazolium bromide (Innochem, Beijing, China) was added and cultured for 4 h at 37 °C in a humidified CO₂ incubator. The solution was removed and dimethyl sulphoxide was added. Thirty minutes later, absorption values were read by a Fully Automated Microplate ELISA Analyser (DeTie, NanJing, China).

LSD1 dilution assay

LSD1 recombinant (Cayman Chemical, Fort Annapolis, MI) was incubated with the targeted compound or dimethyl sulphoxide for 60 min. Then, samples were diluted 80 times using a HRP-assay solution containing substrate and coupling reagents. Finally, LSD1 inhibitory activity before and after dilution was examined.

Western blotting

Cells were seeded in four-well plates (Shanghai Yuanye Biotechnology Co., Ltd, Shanghai, China) and exposed to different compounds at different concentrations. Cells were harvested to obtain protein solution by the RIPA Lysis Buffer (DeTie, NanJing, China). The proteins were separated on sodium dodecyl sulphate-polyacrylamide gel electrophoresis (SDS-PAGE) gels and then transferred to poly(vinylidene fluoride (PVDF) membranes (Shanghai Yuanye Biotechnology Co., Ltd, Shanghai, China). It incubated with the primary antibodies and corresponding HRP conjugated secondary antibodies (Servicebio, Wuhan, China). The electrochemiluminescence kit (Servicebio, Wuhan, China) was used to visualise the bands.

Molecular docking analysis

The protein structure of LSD1 was obtained from Protein Data Bank (PDB code: 2v1d). Sulphonamide derivatives **L6~L8** were transferred as PDB files by ChemBio 3 D Ultra 14.0 (CambridgeSoft, Cambridge, MA). Water was removed from LSD1 protein by Pymol. Autodock (The Scripps Research Institute, San Diego, CA) was used to perform antogrid and autodock. The hydrogen-bond interaction between sulphonamide derivatives **L6~L8** and amino acid residues of LSD1 was analyzed^{26,27}.

Statistical methods

All data of biological experiments were expressed as the mean \pm SD. Significant differences between different groups were analysed by GraphPad Prism version 6 (GraphPad Software, La Jolla, CA). Results were considered statistically significant at **p< 0.01 verse control, ***p< 0.001 verse control and ***p< 0.0001 verse control.

Results and discussion

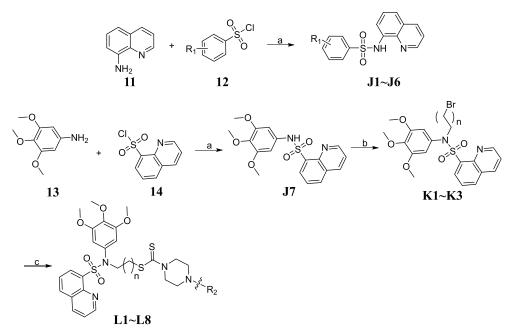
Synthesis of sulphonamide derivatives

Although LSD1 inhibition has been a promising strategy to treat cancer, only few LSD1 inhibitors are currently in clinical trials. Discovery of novel and effective scaffolds to develop LSD1 inhibitors has an important clinical significance. Our aim was to synthesise and identify novel LSD1 inhibitors. Sulphonamide derivatives were synthesised as shown in Scheme 1. Commercially available quinolin-8-amine was treated with different benzenesulfonyl chloride to provide J1~J6 and 3,4,5-trimethoxyaniline was treated with quinoline-8-sulphonyl chloride to afford J7. Compounds $K1\sim K3$ were obtained by the alkylation of sulphonamide J7 with 1,2-dibromoethane or 1,3-dibromopropane or 1,4-dibromobutane. Sulphonamide intermediates $K1\sim K3$ were treated with carbon disulphide and piperazine derivatives in the presence of trisodium phosphate dodecahydrate to obtain sulphonamide-quinoline-dithiocarbamate hybrids L1~L8.

LSD1 inhibition and preliminary structure activity relationship

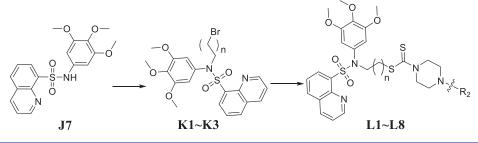
All sulphonamide derivatives in this work were examined *in vitro* for their inhibitory activity against LSD1, MAO-A, and MAO-B. 2-PCPA (tranylcypromine) and ORY-1001 as LSD1 inhibitors were chosen as reference agents²⁸. The inhibitory results of sulphonamides **J1**~**J7** were summarised in Table 1. Sulphonamide derivatives **J1**~**J7** had very weak inhibition effects on MAO-A and MAO-B with IC₅₀ values of >120 μ M. However, all sulphonamide derivatives **J1**~**J7** displayed the moderate activity against LSD1 with IC₅₀ values ranging from 11.57 to 82.46 μ M. Substituent groups on phenyl ring could affect the inhibitory activity against LSD1. Among all these sulphonamides, *N*-(3,4,5-trimethoxyphenyl)quino-line-8-sulphonamide **J7** exhibited the best inhibitory activity, indicating that sulphonamide-quinoline might be a potential scaffold to design LSD1 inhibitors.

Because J7 showed the best inhibitory activity against LSD1, more sulphonamide analogues were designed and synthesised based on compound J7. The in vitro inhibitory activity results of sulphonamides K1~K3 and L1~L8 were listed in Table 2. With the exception of sulphonamide-guinoline-dithiocarbamate hybrids L1~L8, all these sulphonamide analogues bearing a dithiocarbamate fragment exhibit potently inhibitory activity with IC₅₀ values ranging from 0.06 to 4.92 μ M. In comparison with activity results of sulphonamide intermediates K1~K3 without the dithiocarbamate unit, the inhibitory activity of sulphonamide-quinoline-dithiocarbamate hybrids L1~L8 against LSD1 improved obviously, indicating that the dithiocarbamate scaffold play a synergistic role on LSD1 activity. Especially, compound L8 showed the best activity against LSD1, with an IC₅₀ of 0.06 μ M, which is 427 times higher than that of 2-PCPA. During the preliminary structure activity relationship studies, we found that the substituent on the piperazine ring was significant for the LSD1 inhibitory activity showing over 80-fold activity loss, when the hydroxyethyl group was replaced with the ethyl group (compound L8 versus



Scheme 1. Synthesis of the sulphonamide derivatives. Reagents and conditions: (a) triethylamine, acetone, 0°C; (b) 1,2-dibromoethane or 1,3-dibromopropane or 1,4-dibromobutane, dibromobutane, KOH, 60°C; (c) CS₂, piperazine derivatives, and trisodium phosphate dodecahydrate.

Table 1. Inhibitory activity of sulphonamide derivatives J1~J7 against LSD1, MAO-A, and MAO-B.



Compound	R ₁	IC ₅₀ (μM)		
		LSD1	MAO-A	MAO-B
J1	3,4-diOCH ₃	33.78 ± 2.04	>120	>120
J2	4-OCH ₃	45.62 ± 1.12	>120	>120
J3	4-C(CH ₃) ₃	61.90 ± 2.46	>120	>120
J4	Н	82.46 ± 3.38	>120	>120
J5	4-CH ₃	74.52 ± 2.67	>120	>120
J6	4-Br	61.82 ± 5.64	>120	>120
J7	-	11.57 ± 1.92	>120	>120
2-PCPA	-	25.63 ± 2.76	14.57 ± 1.29	6.47 ± 0.62
ORY-1001	-	0.03 ± 0.01	>120	>120

L5). When the substituent on the piperazine ring was tert-butoxycarbonyl group and methyl group, compound **L6** and compound **L7** displayed the potent LSD1 inhibitory activity with IC₅₀ values of 0.23 and 0.73 μ M, respectively. In addition, the length of carbon linker between the sulphonamide and the dithiocarbamate exhibited a significant role in their activities. With the reduction of the carbon tether length, a decrease of LSD1 inhibitory activity was observed (compound **L8** versus **L3**, or compound **L6** versus **L1**). Based on the enzyme inhibitory results, all sulphonamide derivatives **L1~L8** potently inhibited LSD1 with IC₅₀ values ranging from 4.92 to 0.06 μ M. Importantly, they had no significant effects on MAO-A and MAO-B with IC₅₀ values of >120 μ M, indicating that these novel sulphonamide derivatives were selective LSD1 inhibitors. For the preliminary structure activity relationships, dithiocarbamate unit and substituent groups on piperazine of sulphonamide derivatives **L1**~**L8** played significant roles on LSD1 inhibitory activity.

Sulphonamide derivative L8 exhibited good selectivity between cancer cells and normal cells

Based on the LSD1 inhibitory activity results of all sulphonamide derivatives, the most potent LSD1 inhibitor **L8** was prioritised to perform further experiments for evaluating its antiproliferative effects against cancer cell lines and normal cell lines. Bel7402 (human liver cell line), H1299 (human lung cell line), HCT-15 (human colon cell line), SGC-7901 (human gastric cell line), HTB3 (human bladder cell line), Jurkat (human leukaemia cell line), LX-2

(normal human hepatic stellate cell line), HPAEpiC (normal human alveolar epithelial cell line), and HIEC (normal human intestinal epithelial cell line) were selected to do MTT assay with the treatment of compound **L8** for 48 h. The IC₅₀ values of compound **L8** were 0.96, 0.31, 2.17, 2.67, 0.18, and 0.29 μ M against Bel7402, H1299, HCT-15, SGC-7901, HTB3, and Jurkat cells, respectively. However, compound **L8** showed weak antiproliferative activity against normal cell lines (LX-2, HPAEpiC, and HIEC) with IC₅₀ values of >16 μ M. Thus, sulphonamide derivative **L8** exhibited good selectivity between cancer cells and normal cells.

Sulphonamide derivative L8 potently inhibited cell proliferation against bladder cancer in a concentration-dependent and timedependent manner

Among all six cancer cell lines, sulphonamide derivative **L8** displayed the best activity results against bladder cancer HTB3 cells. So, bladder cancer was chosen to investigate the mechanisms of compound **L8**. According to results, sulphonamide derivative **L8** potently inhibited cell proliferation against different bladder cancer cell lines. When these four bladder cancer cell lines were

Table 2. Inhibitory activity of sulphonamide derivatives $K1{\sim}K3$ and $L1{\sim}L8$ against LSD1, MAO-A, and MAO-B.

ĭ,k

				IC ₅₀ (μM)	
Compound	n	R_2	LSD1	MAO-A	MAO-B
J7	-	_	11.57 ± 1.92	>120	>120
K1	1	-	14.79 ± 2.04	>120	>120
K2	2	-	9.63 ± 1.26	>120	>120
K3	3		6.32 ± 0.77	>120	>120
L1	1	Å,	2.73 ± 0.29	>120	>120
L2	2	∕∕ОН	1.39 ± 0.14	>120	>120
L3	1	∕∕ОН	1.85 ± 0.07	>120	>120
L4	2	\sim	0.67 ± 0.10	>120	>120
L5	3	Å.	4.92 ± 0.13	>120	>120
L6	3	/	0.23 ± 0.05	>120	>120
L7	3	∕∕ОН	0.73 ± 0.08	>120	>120
L8	3		0.06 ± 0.01	>120	>120
2-PCPA ORY-1001	- -	-	25.63 ± 2.76 0.03 ± 0.01	14.57 ± 1.29 >120	6.47 ± 0.62 >120

treated with compound **L8**, IC₅₀ values for 48 h were 1.87, 0.18, 0.09, and 0.93 μ M against HTB5, HTB3, HT1376, and HTB1 cells, respectively. In addition, cell viability of bladder cancer decreased obviously in a concentration-dependent and time-dependent manner.

Sulphonamide derivative L8 selectively and reversibly inhibited LSD1 in enzyme-based assays

From the enzyme inhibitory results in Figure 3(A), sulphonamide derivative L8 weakly inhibited MAO-A and MAO-B with the inhibitory rates of only 10.33% and 9.33% at 100 nM, while it showed about 30.33%, 46.00%, and 68.00% of inhibition against LSD1 at 25, 50, and 100 nM, suggesting that compound **L8** selectively inhibited LSD1 in a concentration-dependent. As shown in Figure 3(B), sulphonamide derivative **L8** potently inhibited LSD1 in a time-dependent manner. In the dilution assay, the low concentration of sulphonamide derivative **L8** by the dilution could result in the recovery of LSD1 inhibitory activity. Results in Figure 3(C) showed that sulphonamide derivative **L8** was a reversible LSD1 inhibitor.

Sulphonamide derivative L8 potently inhibited LSD1 in cellbased assays

Before investigating the LSD1 inhibitory mechanisms at the cellular level, we first examined the LSD1 expression levels in four different bladder cancer cell lines (HTB5, HTB1, HTB3, and HT1376). From the results in Figure 4(A), HT1376 cells possessed the highest LSD1 expression levels, followed by HTB3, HTB1, and HTB5. In this work, we used the LSD1 knock-down cells and control cells to investigate the in vitro antiproliferative activity of LSD1 inhibitor L8 (Figure 4(B,C)). Compound L8 inhibited HTB3&shLSD1 cells and HT1376&shLSD1 cells with the IC₅₀ values of 3.07 and 4.36 μ M. However, it significantly inhibited cell proliferation against HTB3&shControl cells and HT1376&shControl cells with the IC₅₀ values of 0.23 and 0.11 μ M, respectively. The great activity discrepancy between LSD1 knock-down cells and control cells showed that the antiproliferative effects of sulphonamide derivative L8 against bladder cancer was dependent on its LSD1 inhibition. Meanwhile, several substrates of LSD1, including H3K4me1, H3K4me2, and H3K9me2 were also investigated to their expression levels with the compound treatment. As shown in Figure 4(D), sulphonamide derivative L8 increased the expression levels of H3K4me1, H3K4me2, and H3K9me2 against HT1376 cells. All these results showed that sulphonamide derivative L8 could inhibit LSD1 in the cellular level.

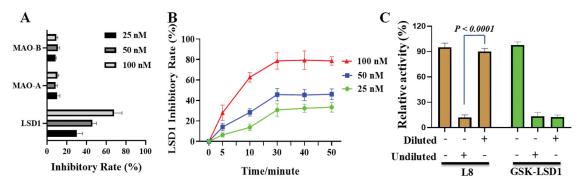


Figure 3. Mechanism studies of sulphonamide derivative L8 on LSD1 inhibitory activity at the enzyme level. (A) Enzyme selectivity of sulphonamide derivative L8. (B) Time dependent curve of LSD1 inhibition by compound L8. (C) Reversible study of compound L8 against LSD1. GSK-LSD1 was a famous LSD1 inhibitor²⁹.

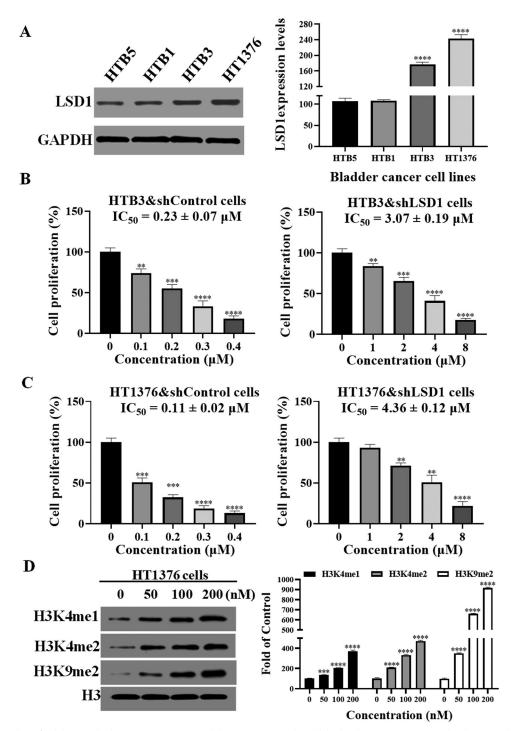


Figure 4. Mechanism studies of sulphonamide derivative L8 on LSD1 inhibitory activity at the cellular level. (A) LSD1expression levels against bladder cancer cell lines. (B) Cell proliferation (%) of HTB3&shControl cells and HTB3&shLSD1 cells. (C) Cell proliferation (%) of HTB376&shControl cells and HTB3&shLSD1 cells. (D) The expression levels of H3K4me1, H3K4me2, and H3K9me2 against HTB376 cells with the treatment of L8. **P < 0.01 verse control, ***p < 0.001 verse control and ****p < 0.001 verse control.

Molecular docking analysis of sulphonamide derivative L8 targeting LSD1

A molecular docking was carried out to predict the binding interaction between sulphonamide derivatives and LSD1. The protein structure of LSD1 was extracted from Protein Data Bank (PDB code: 2V1D, resolution: 3.1 Å) and was prepared by adding hydrogen atoms and removing flavin adenine dinucleotide. The molecular docking results in Figure 5 indicated that sulphonamide derivatives $L6 \sim L8$ showed different binding modes through hydrogen bonds in the active pocket of LSD1. The dithiocarbamate group and amide group of sulphonamide derivative L6formed two hydrogen bonds with LYS14 and GLU559, and the distance was 2.9 and 2.4 Å, respectively. In addition, compound L7also had hydrogen interactions with LYS9 and ASN383 and the distance was 2.2 and 2.5 Å, respectively. Compared with compound L6 and L7, sulphonamide derivative L8 formed more hydrogen bonds with residues of LSD1. Compound L8 makes five

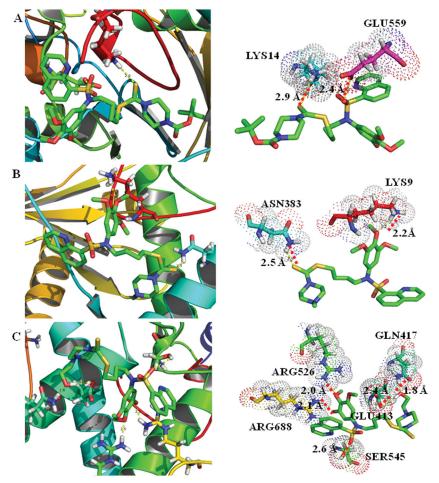


Figure 5. Molecular docking results of sulphonamides $L6 \sim L8$ in binding with LSD1. (A) Binding interactions between sulphonamide L6 and LSD1. (B) Binding interactions between sulphonamide L7 and LSD1. (C) Binding interactions between sulphonamide L8 and LSD1.

hydrogen bonds with amino acid residues GLN417, GLU413, SER545, ARG688, and ARG526. The docking studies of sulphonamide derivatives were in accordance with the results of LSD1 inhibitory activity.

Conclusion

In conclusion, we have designed and synthesised a new class of sulphonamide-quinoline-dithiocarbamate hybrids as LSD1 inhibitors. Novel sulphonamide derivative **L8** selectively and reversibly inhibits LSD1 in a concentration-dependent and time-dependent manner. Importantly, sulphonamide derivative **L8** as a potent antiproliferative agent suppresses the proliferation against bladder cancer cells. To the best of our knowledge, there have been no literature reports regarding sulphonamide-quinoline-dithiocarbamate hybrids as LSD1 inhibitors against bladder cancer cells so far. All these findings provide an effective molecular skeleton for the discovery of LSD1 inhibitors, and sulphonamide based LSD1 inhibitors might be potentially anticancer drugs to treat bladder cancer.

Disclosure statement

The authors declare no competing interests.

Funding

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