

iScience, Volume 27

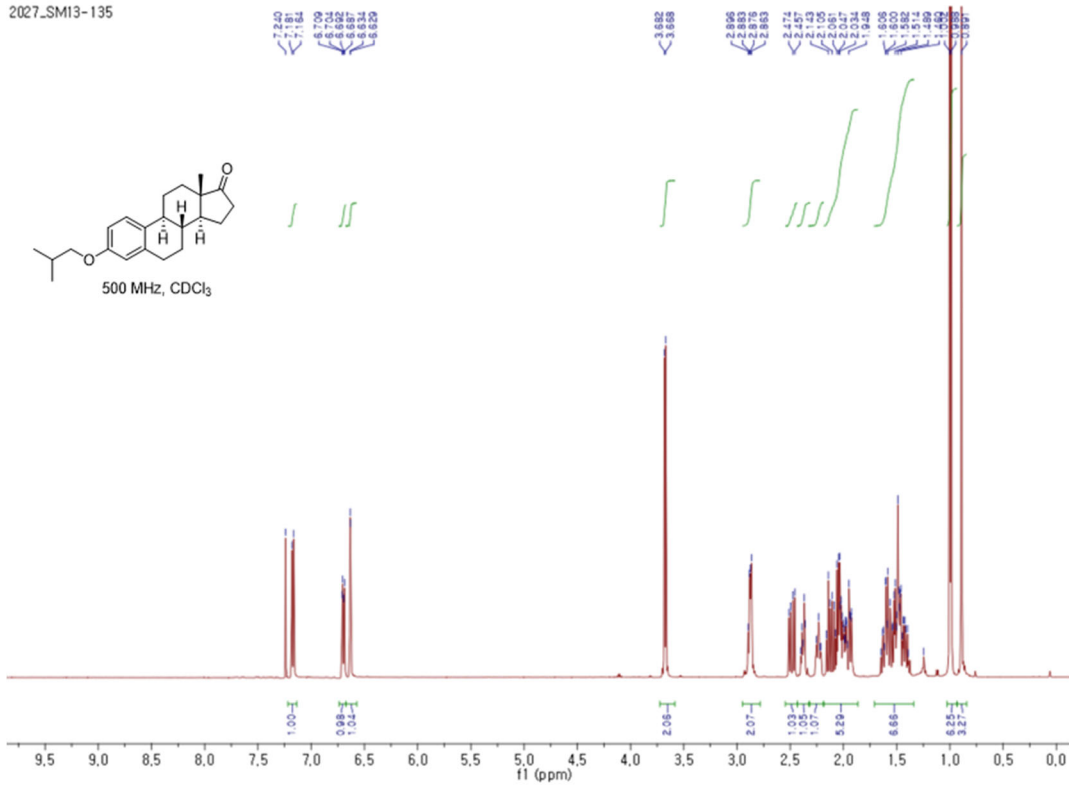
Supplemental information

Dual inhibition of aminoacyl-tRNA synthetase interacting multifunctional protein-2 and α -synuclein by steroid derivative is neuroprotective in Parkinson's model

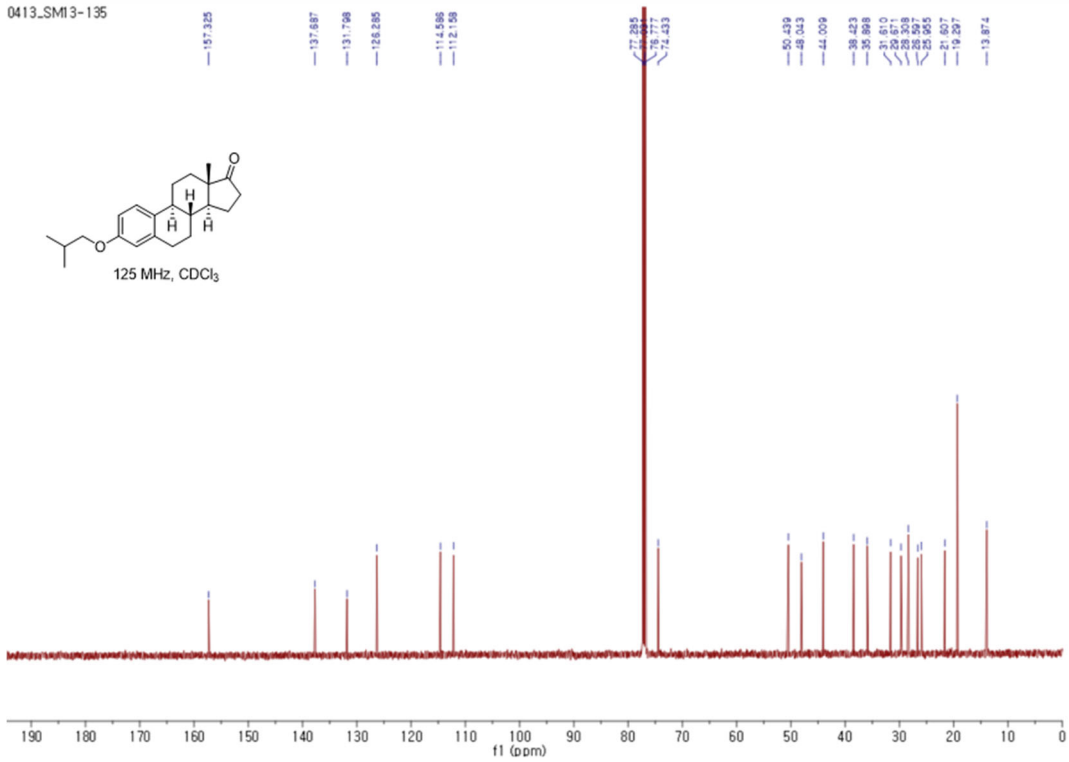
Jeong-Yong Shin, Min Woo Ha, Ji Hun Kim, Jiwon Cheon, Gum Hwa Lee, Seung-Mann Paek, and Yunjong Lee

<Spectral data of SG13-135>

2027_SMI3-135



0413_SMI3-135



Single Mass Analysis

Tolerance = 20.0 PPM | DBE: min = -3.0, max = 100.0

Element prediction: Off

Number of isotope peaks used for iFIT = 9

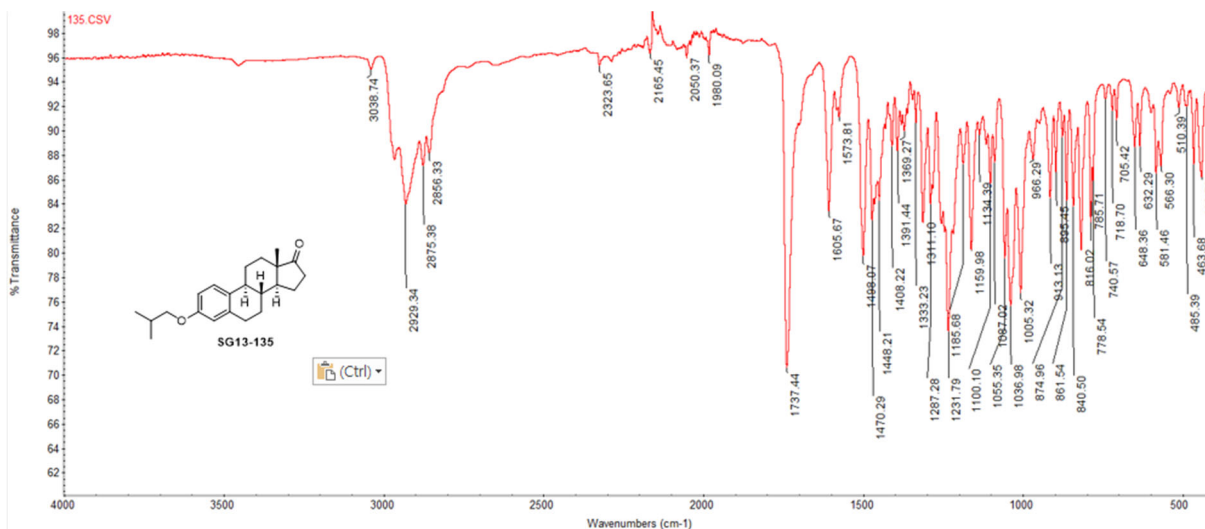
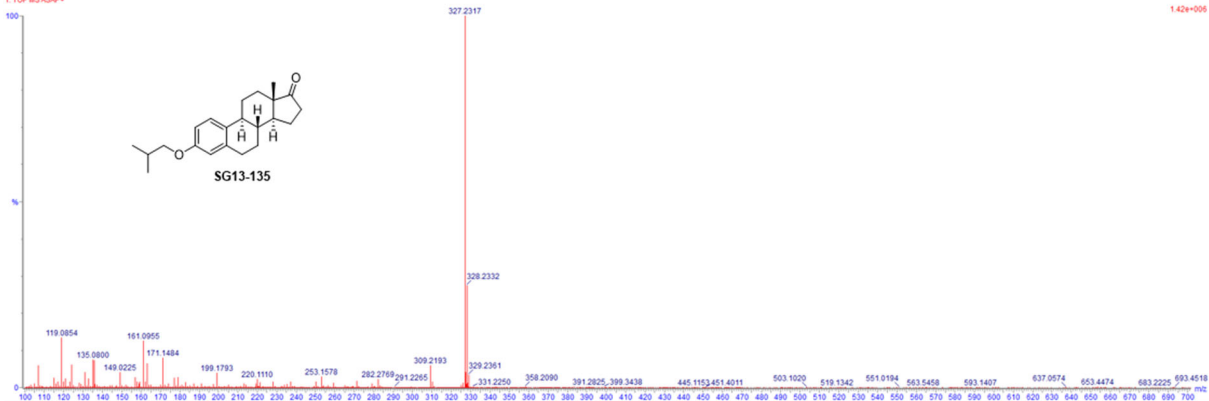
Monoisotopic Mass, Even Electron Ions

22 formulae evaluated with 1 result within limits (up to 10 closest results for each mass)

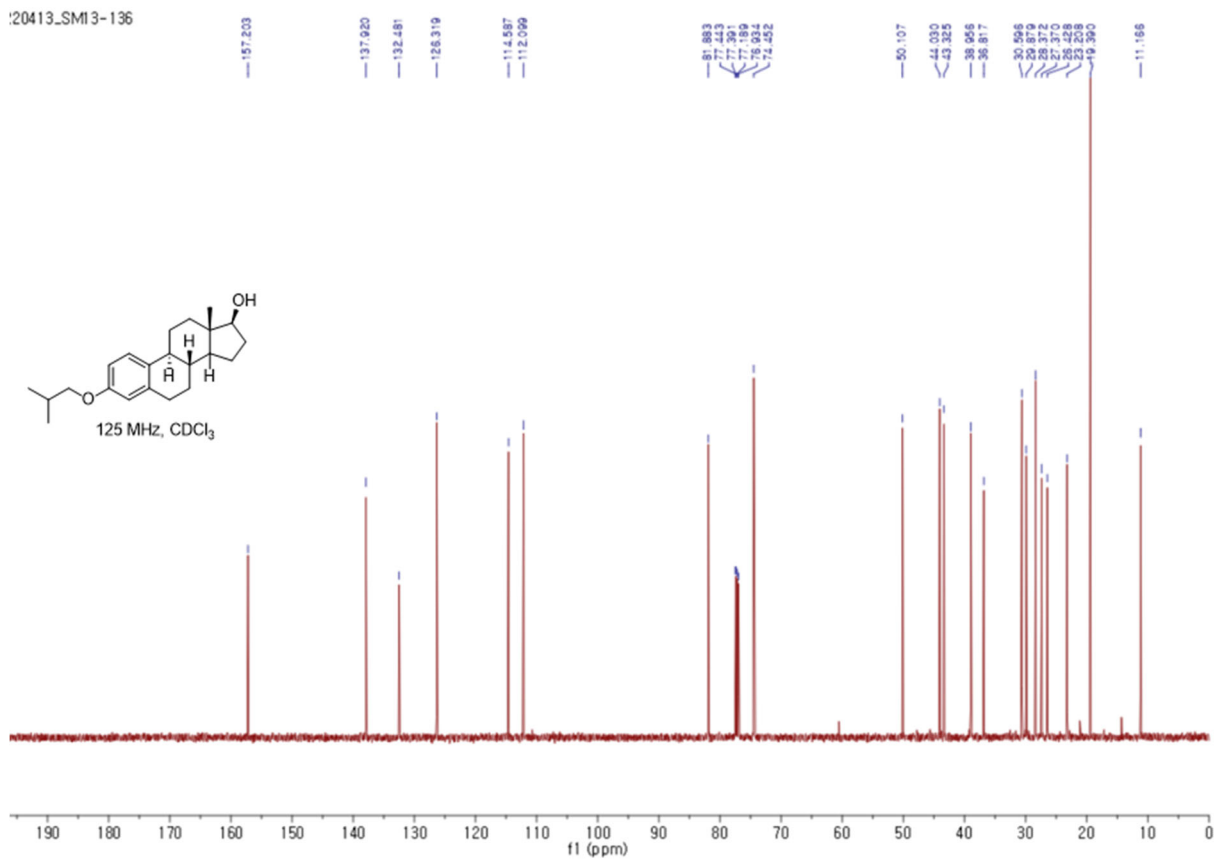
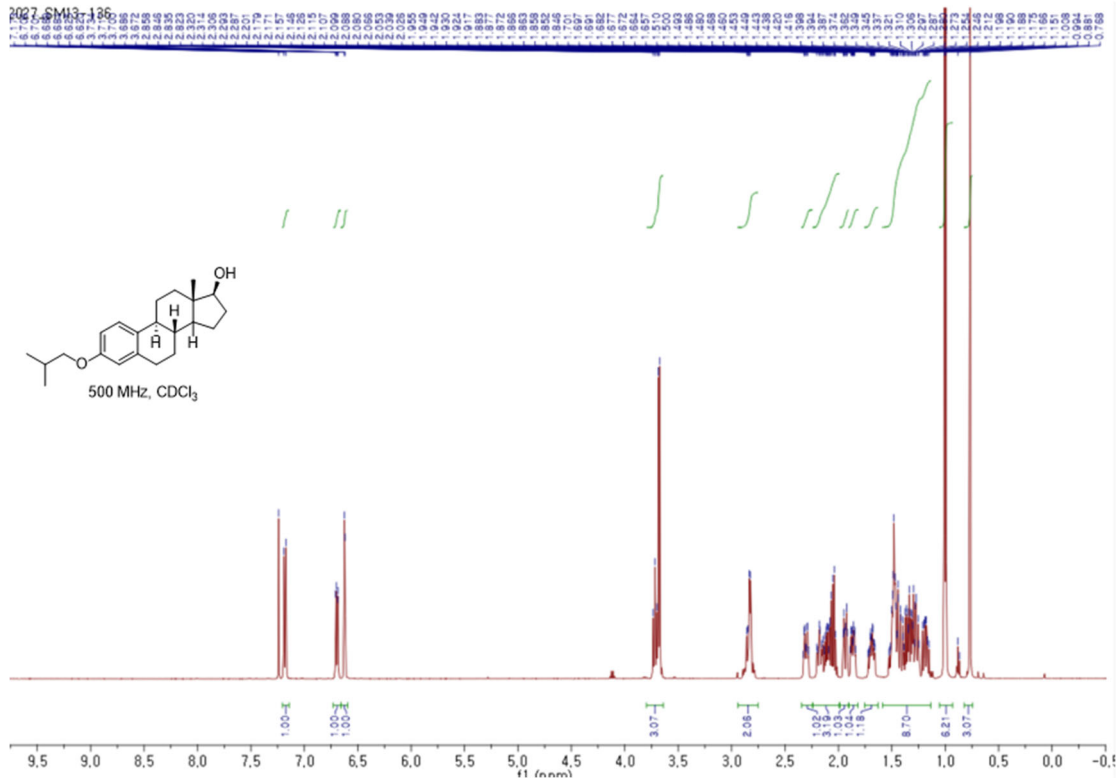
Elements Used

Mass	Calc. Mass	mDa	PPM	DBE	Formula	iFIT	iFIT Norm	Fit Conf %	C	H	O
327.2317	327.2324	-0.7	-2.1	7.5	C ₂₂ H ₃₁ O ₂	1741.6	n/a	n/a	22	31	2

20220413_SM13-135_ASAP_1415 (3.848)
1. TOP MS ASAP+



<Spectral data of SG13-136>



Single Mass Analysis

Tolerance = 30.0 PPM / DBE: min = -3.0, max = 100.0

Element prediction: C#F

Number of isotopic peaks used for i-FT = 9

Monoisotopic Mass, Even Electron Ions

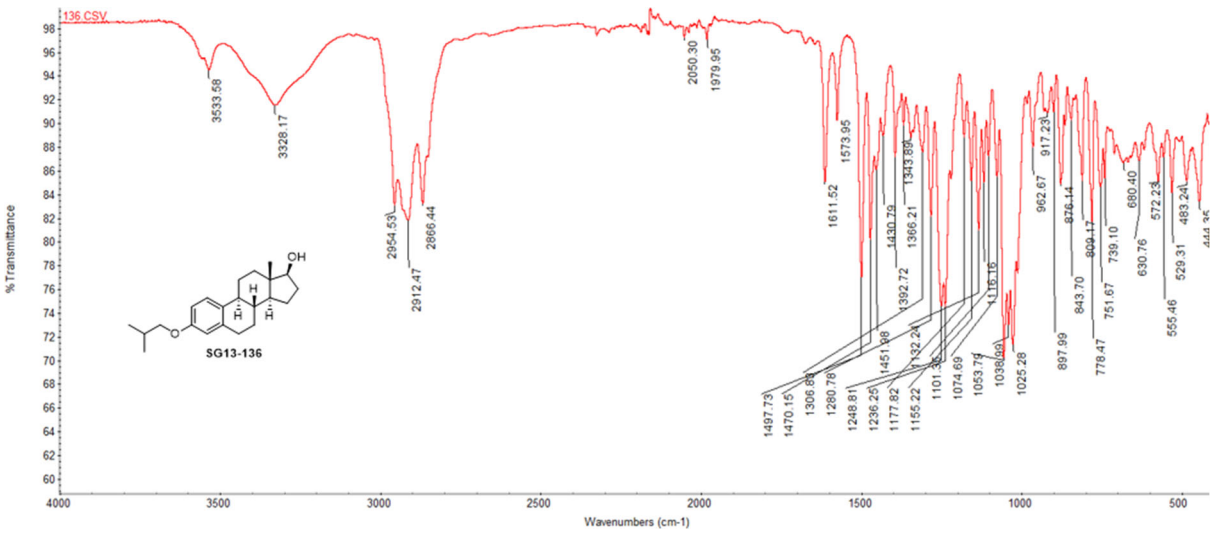
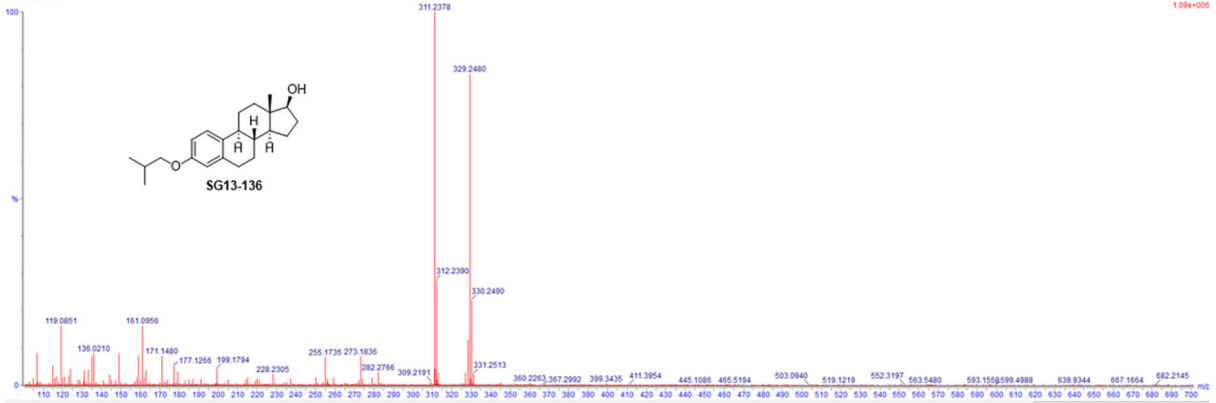
23 formula(s) evaluated with 1 result within limits (up to 10 closest results for each mass)

Elements Used

Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FT	i-FT Norm	Fit Conf %	C	H	O
329.2480	329.2481	-0.1	-0.3	6.5	C22 H33 O2	1000.0	1000.0	100	22	33	2

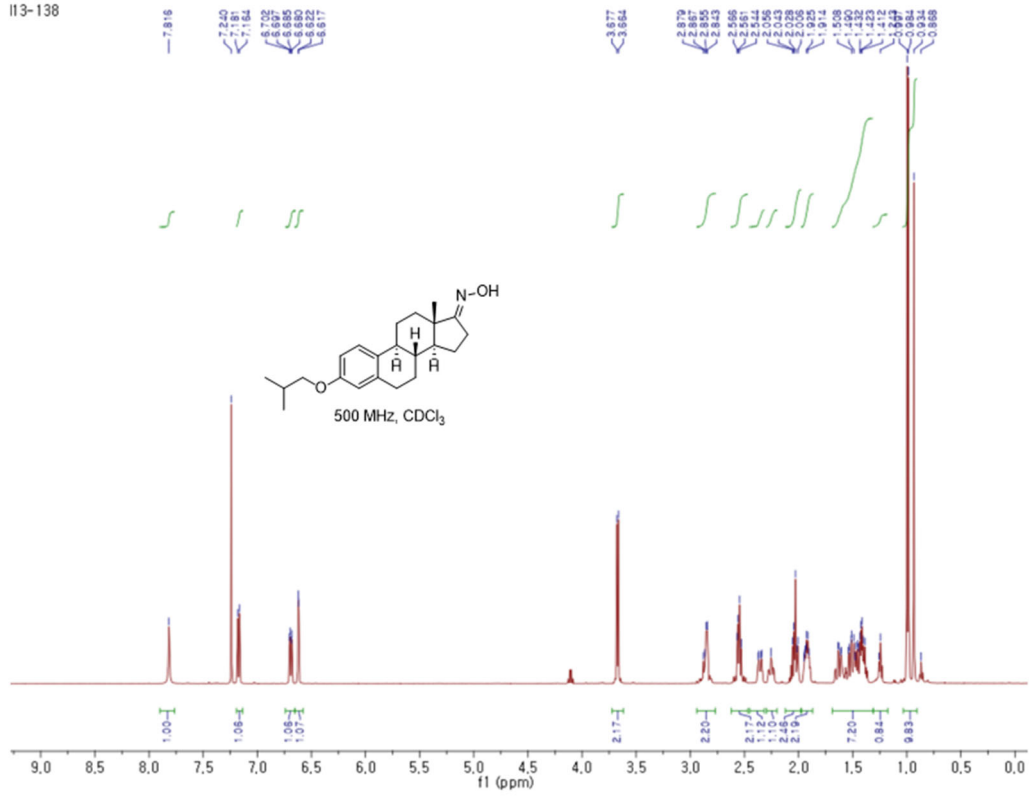
20220413_0M13-136_ADAP 523 (4.843)

1: TOF MS ADAP+

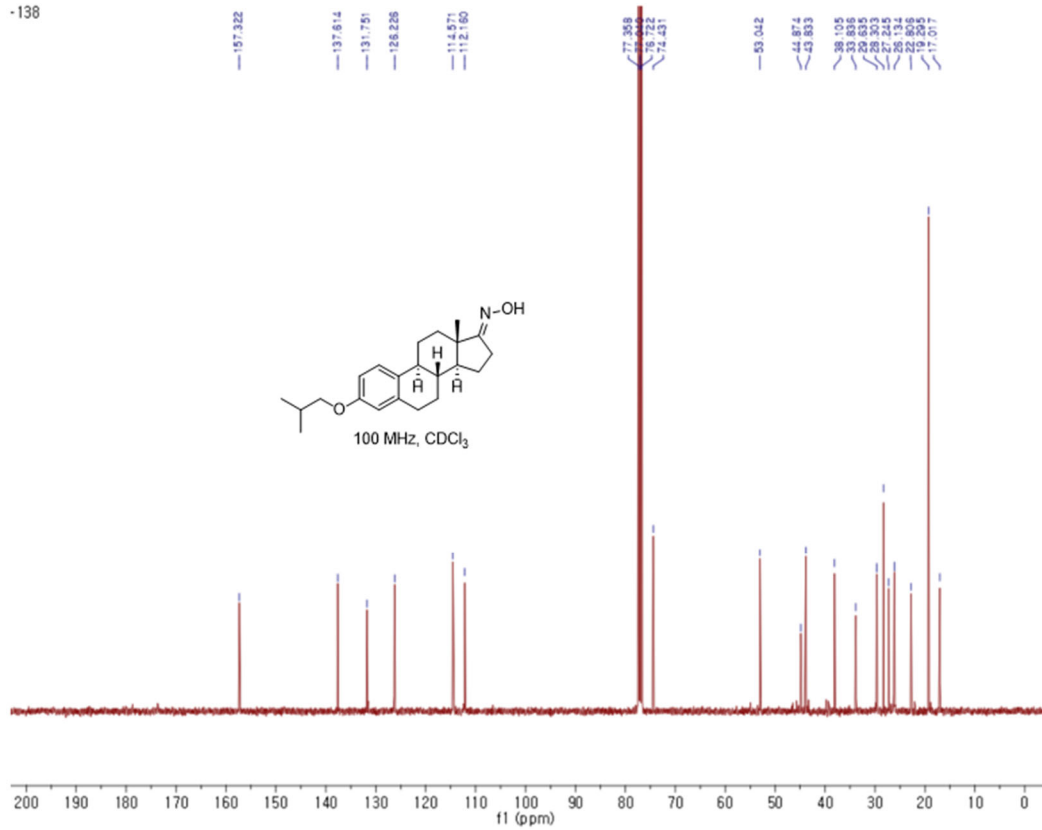


<Spectral data of SG13-138>

113-138

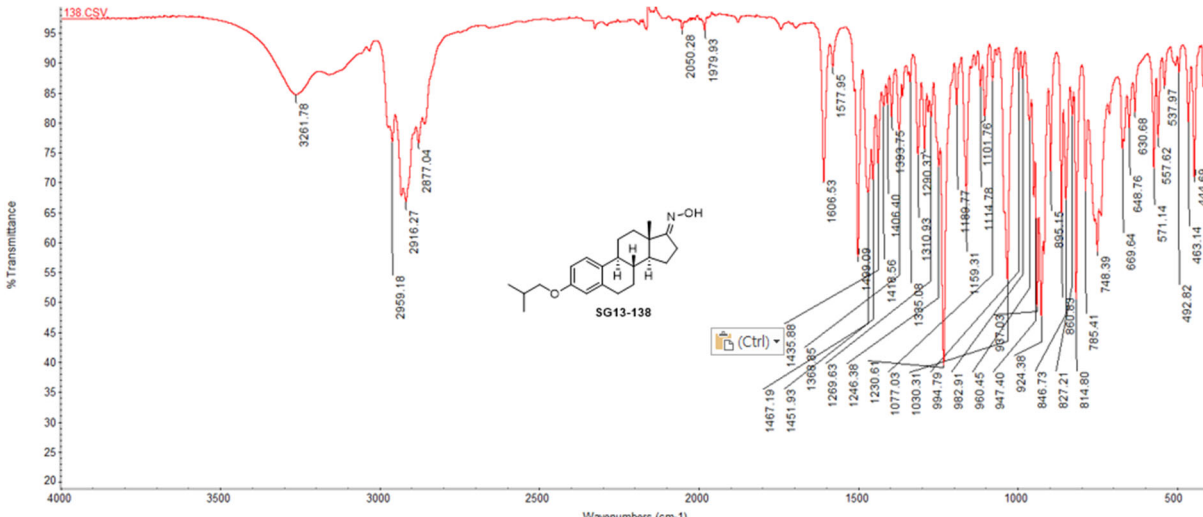
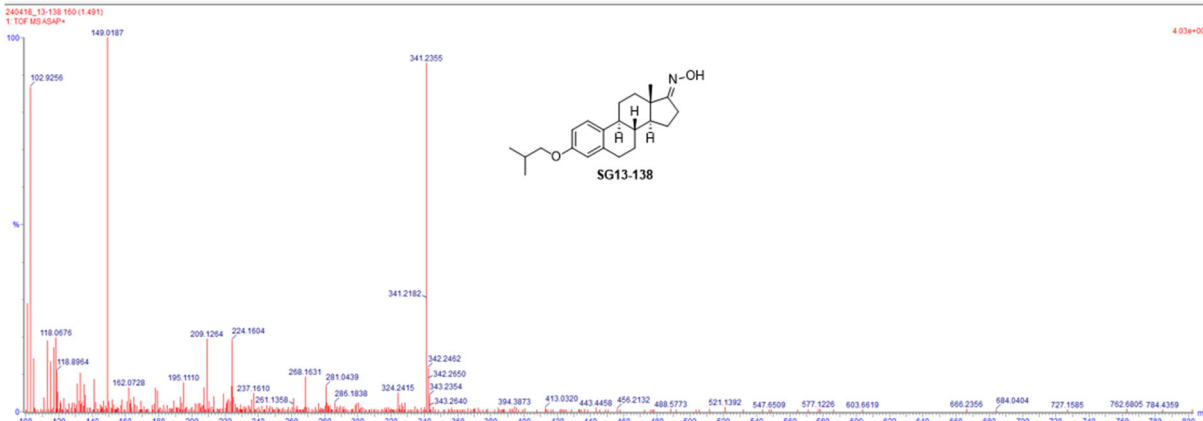


-138



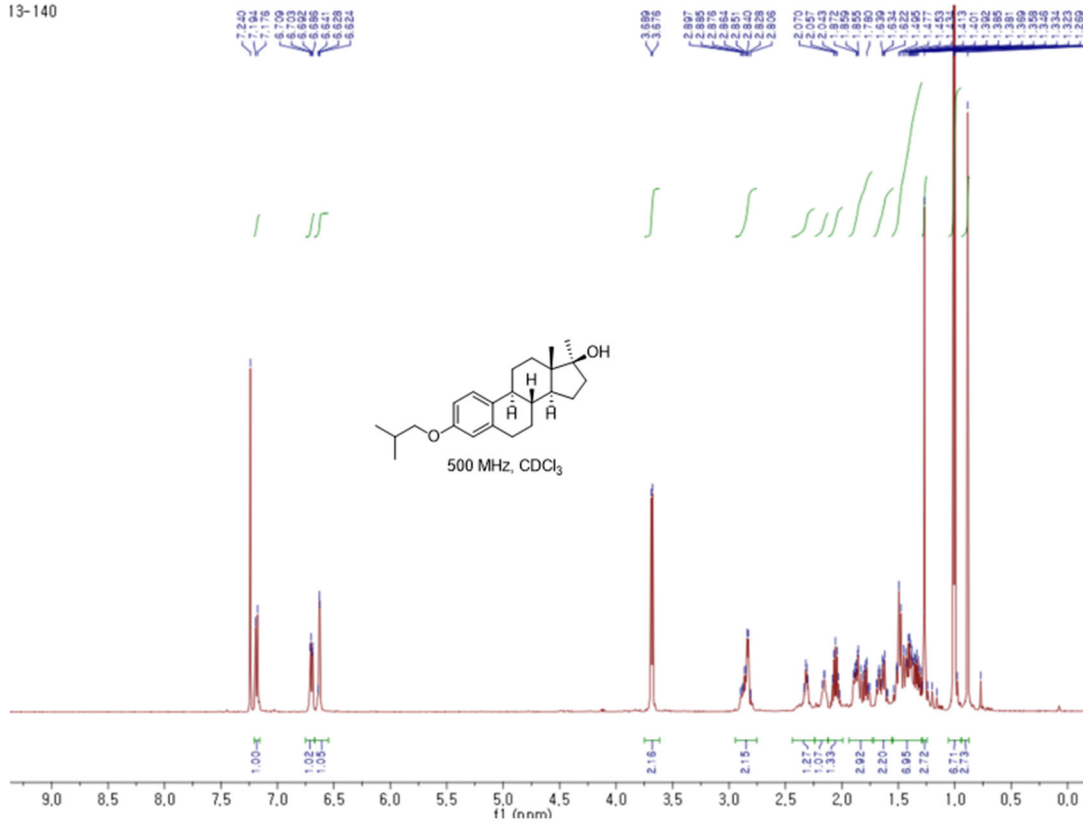
Single Mass Analysis
Tolerance = 20.0 PPM / DBE: min = -30.0, max = 80.0
Element prediction: C#F
Number of isotope peaks used for iFIT = 5
Monoisotopic Mass, Odd and Even Electron Ions
102 formula(s) evaluated with 1 result within limits (up to 5 closest results for each mass)
Elements Used:

Mass	Calc. Mass	mDa	PPM	DBE	Formula	iFIT	iFIT Norm	Fit Conf %	C	H	N	O
341.2355	341.2355	0.0	0.0	8.0	C22 H31 N O2	102.7	n/a	n/a	22	31	1	2

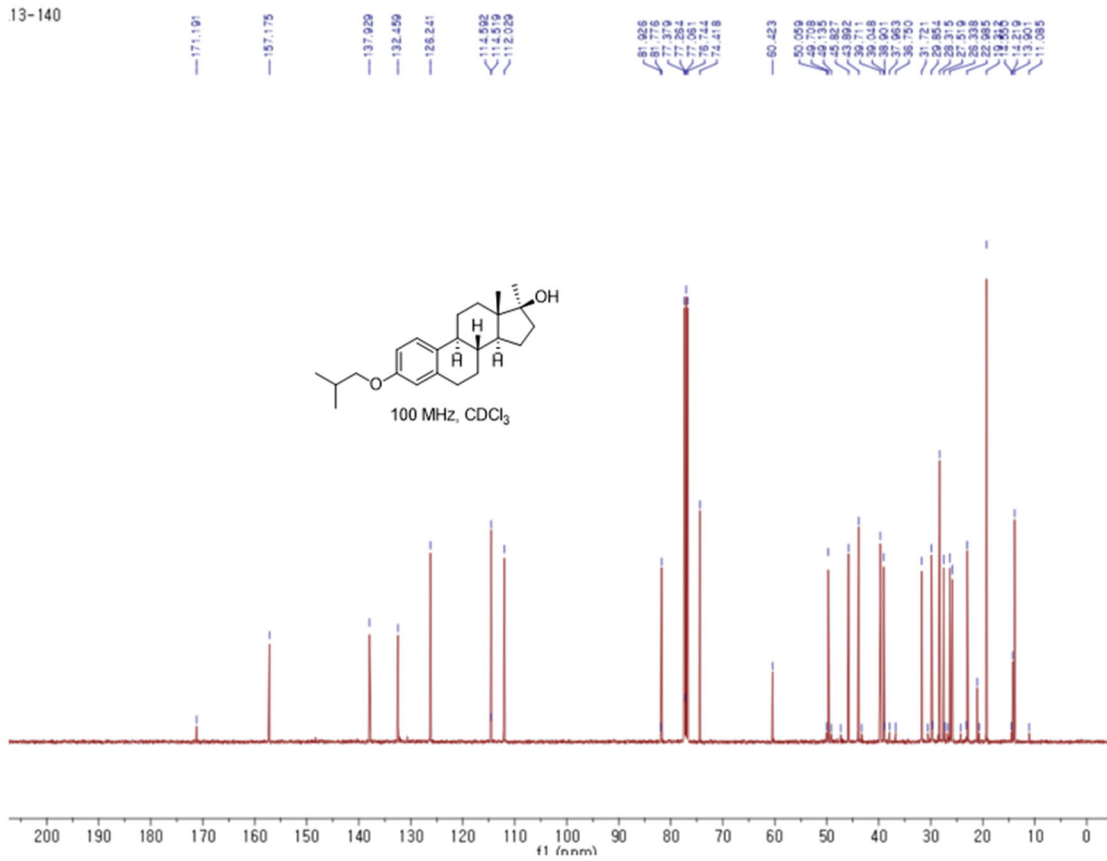


<Spectral data of SG13-140>

13-140



13-140

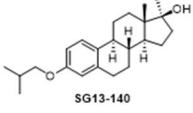
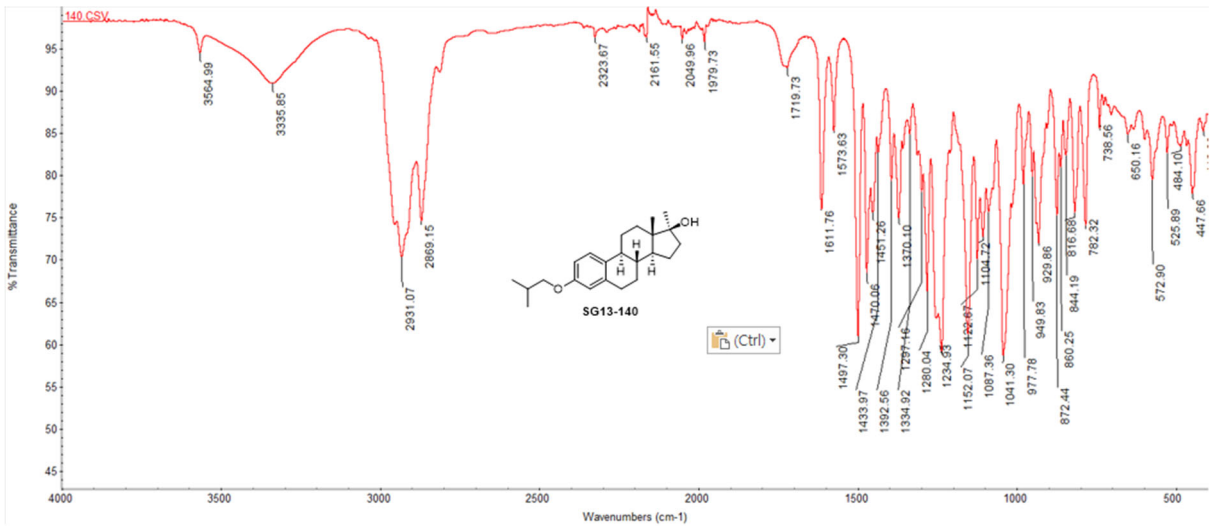
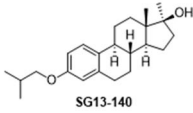
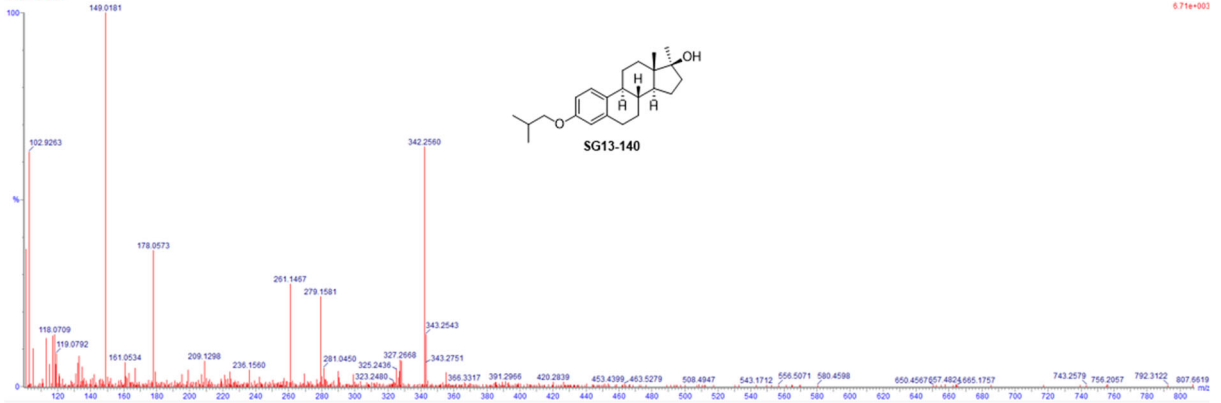


Single Mass Analysis
 Tolerance = 20.0 PPM / DBE: min = -30.0, max = 80.0
 Element prediction: Off
 Number of isotope peaks used for i-FT = 5
 Monoisotopic Mass, Odd and Even Electron Ions
 25 formula(s) evaluated with 1 result within limits (up to 5 closest results for each mass)
 Elements Used

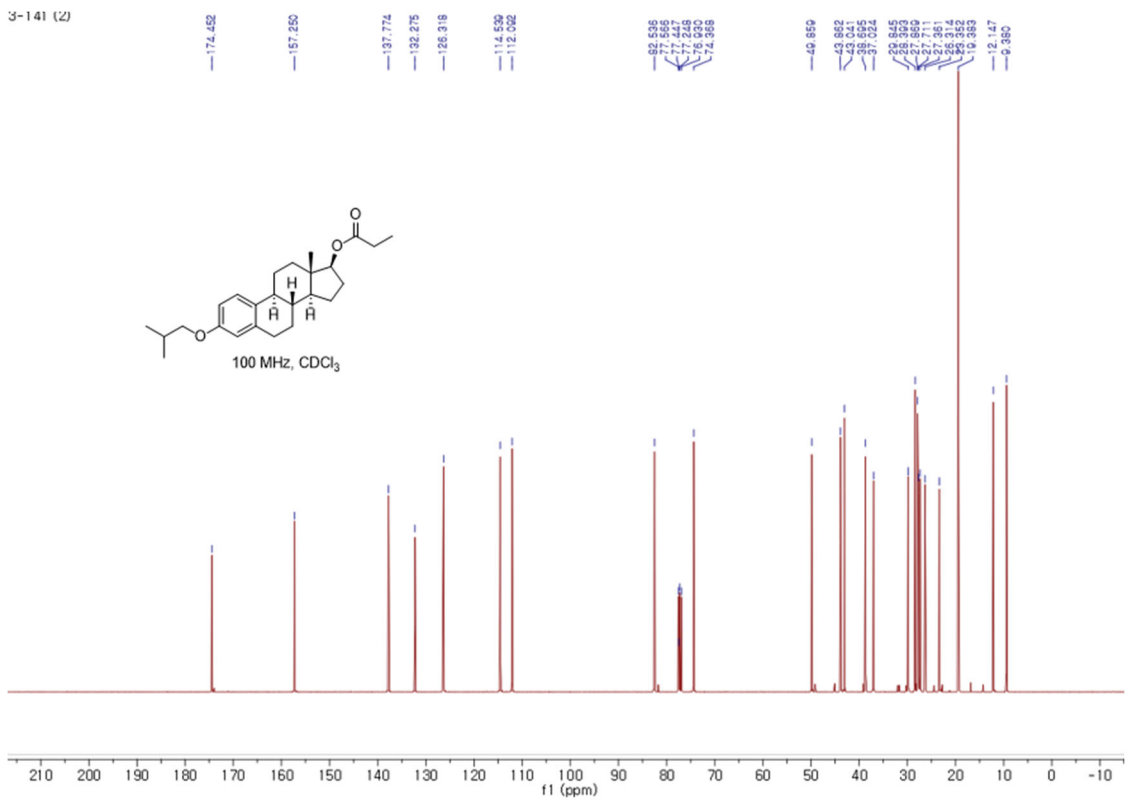
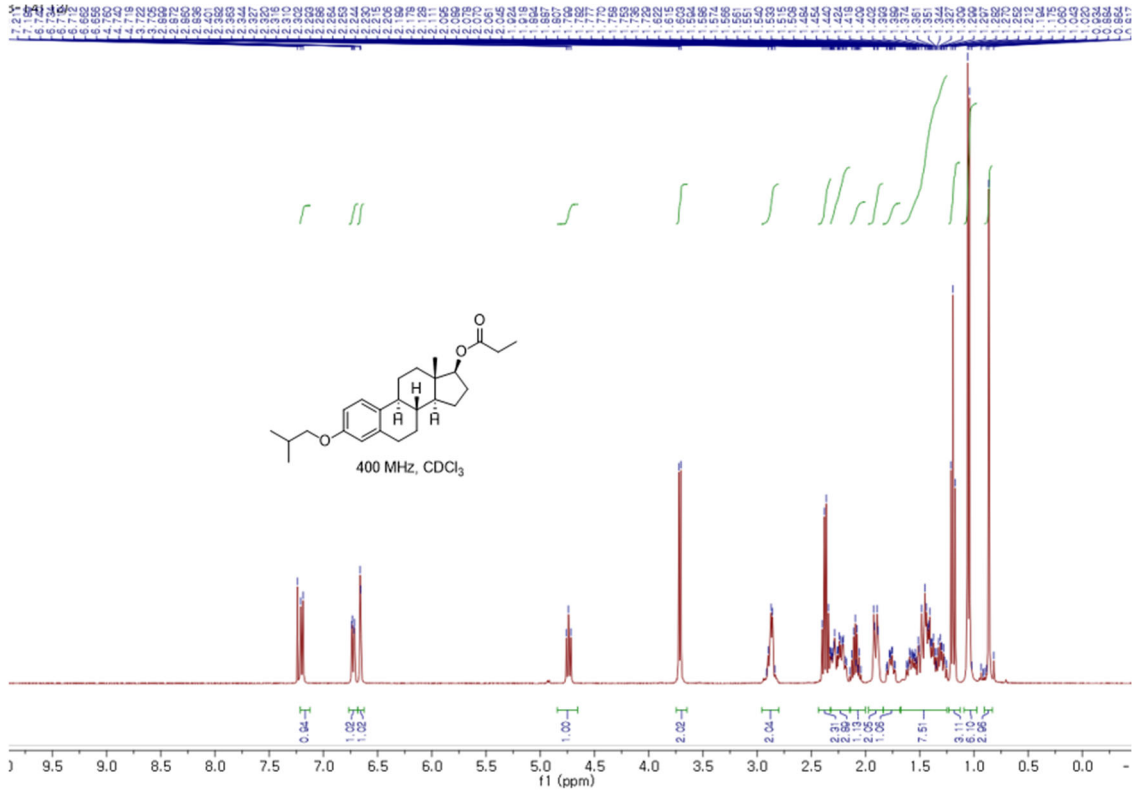
Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FT	i-FT Norm	Fit Conf %	C	H	O
342.260	342.259	0.1	0.3	7.0	C ₂₃ H ₃₄ O ₂	93.8	n/a	n/a	23	34	2

240418_13-140.00 (0.645)

1: TDF.8045AP



<Spectral data of SG13-141>



Single Mass Analysis

Tolerance = 20.0 PPM | DGE: min = -30.0, max = 80.0

Element prediction: O#

Number of isotope peaks used for iFIT = 5

Monoisotopic Mass, Even Electron Ions

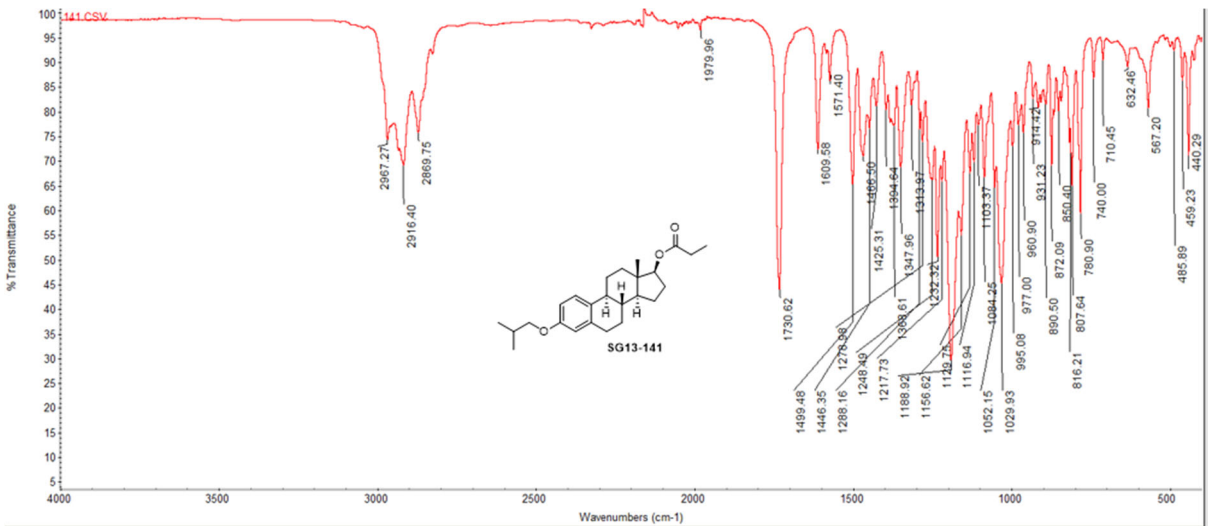
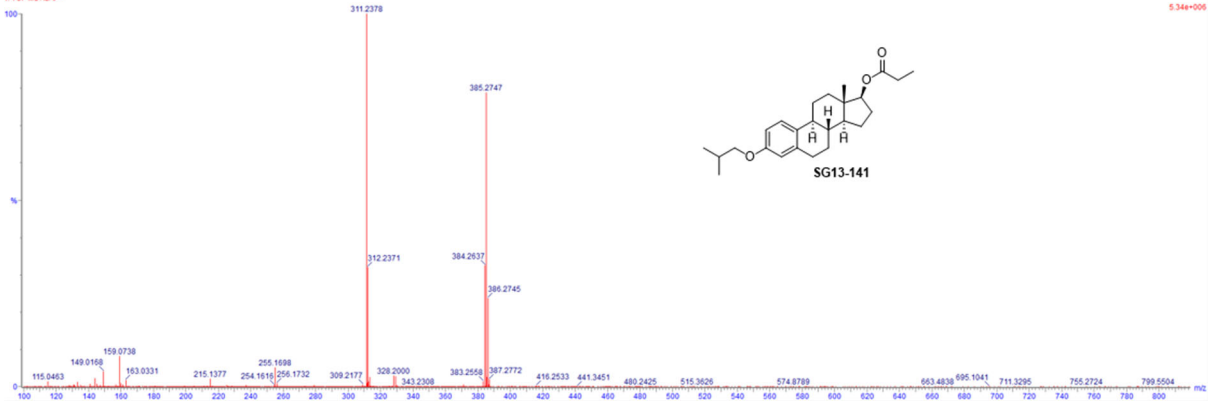
42 formula(s) evaluated with 1 result within limits (up to 5 closest results for each mass)

Elements Used

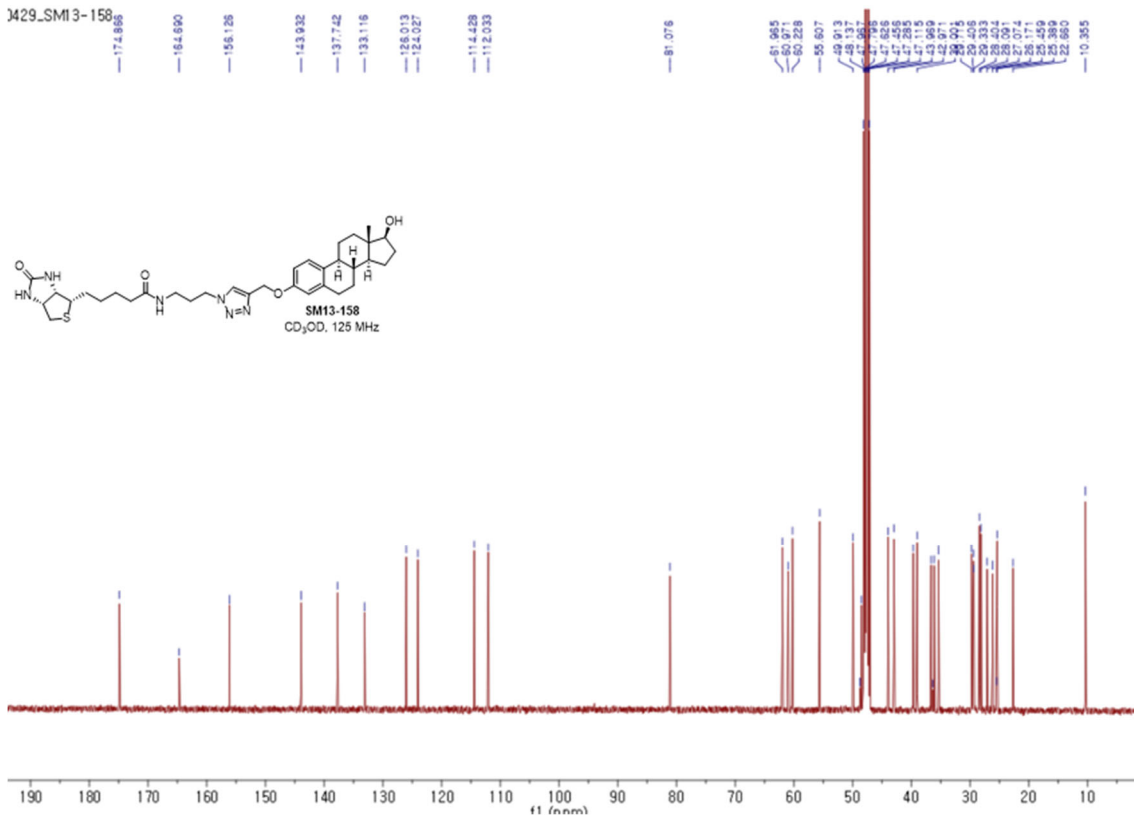
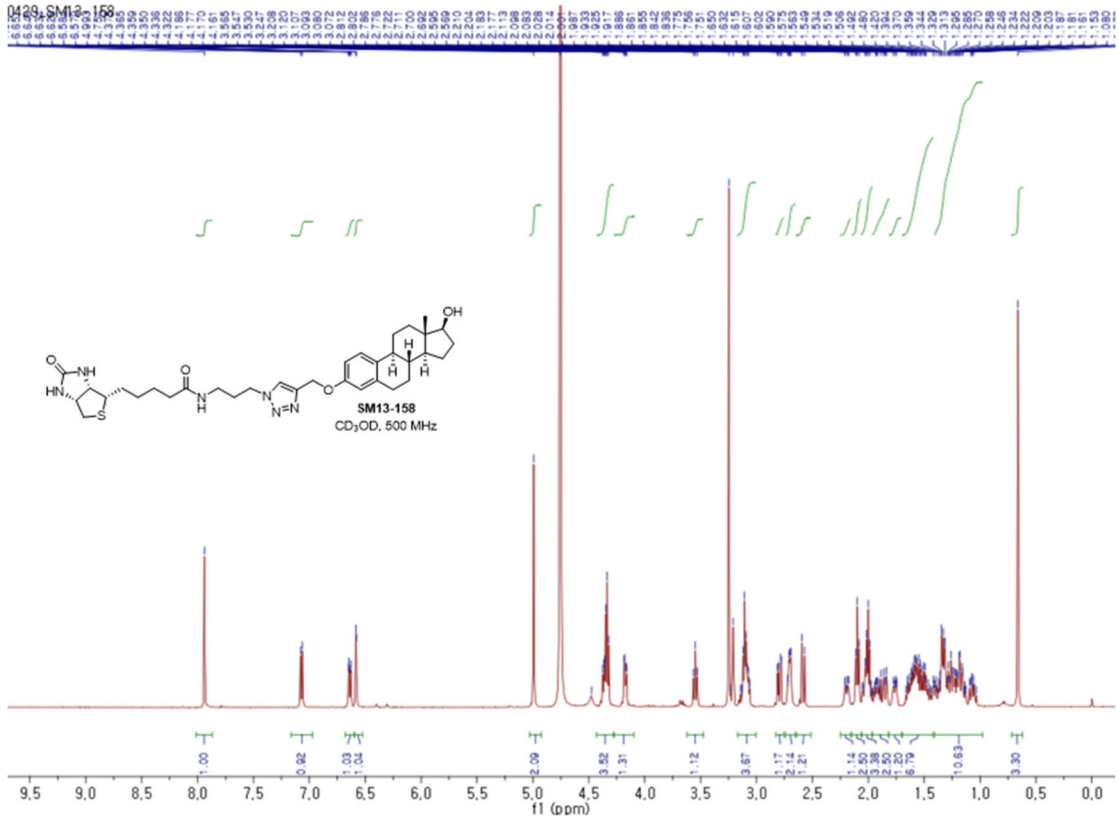
Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FIT	i-FIT Norm	Fit Conf %	C	H	O
385.2747	385.2743	0.4	1.0	7.5	C ₂₅ H ₃₇ O ₃	588.5	n/a	n/a	25	37	3

242425_13-141_356 (3.297)

1. TOP MS ASAP



<Spectral data of SG13-158>



Single Mass Analysis

Tolerance = 20.0 PPM / DBE: min = -30.0, max = 80.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Odd and Even Electron Ions

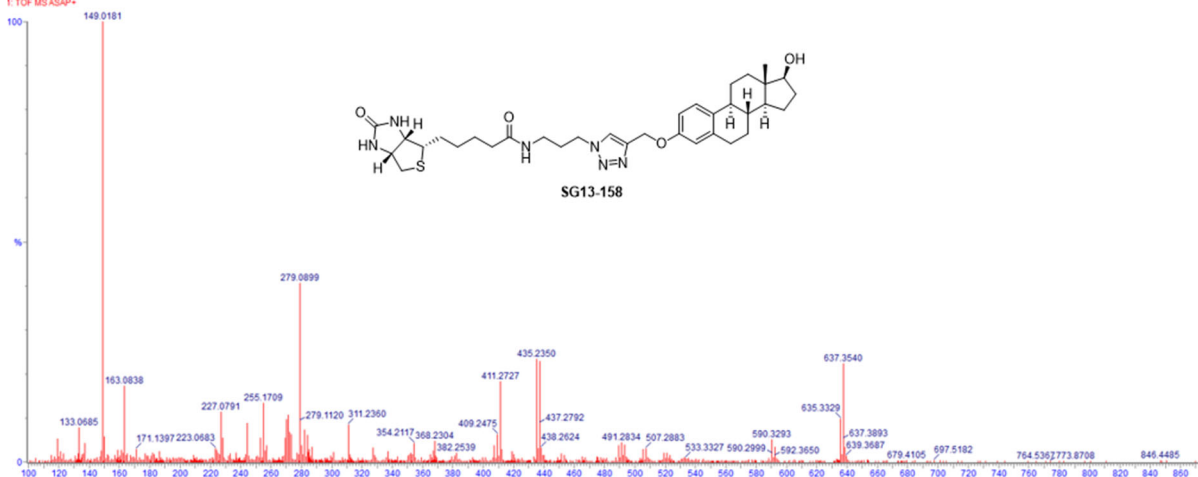
416 formula(e) evaluated with 13 results within limits (up to 5 closest results for each mass)

Elements Used:

Mass	Calc. Mass	mDa	PPM	DBE	Formula	i-FIT	i-FIT Norm	Fit Coef %	C	H	N	O	S
637.3540	637.3536	0.4	0.6	13.5	C34 H49 N6 O4 S	96.7	1.969	13.07	34	49	6	4	1
	637.3549	-0.9	-1.4	13.0	C36 H51 N3 O5 S	96.4	1.707	18.14	36	51	3	5	1
	637.3523	1.7	2.7	14.0	C32 H47 N9 O3 S	96.9	2.218	10.88	32	47	9	3	1
	637.3563	-2.3	-3.6	18.0	C37 H47 N7 O 5	96.1	1.409	24.44	37	47	7	1	1
	637.3576	-1.6	-5.6	17.5	C39 H49 N4 O2 S	95.8	1.122	32.57	39	49	4	2	1

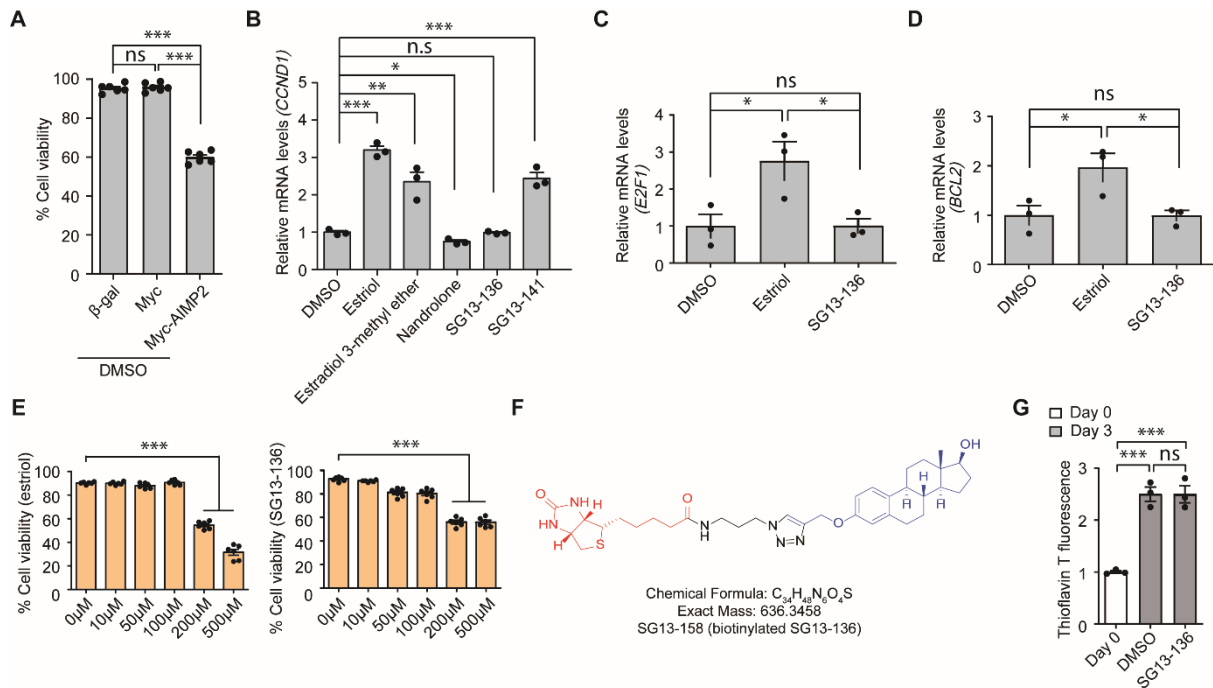
140418_13-158 505 (5.505)

1: TOF MSABAP+



Data S1. Related to Figure 1. Chemical structures and ^1H -NMR spectra of newly synthesized steroid derivatives.

The ^1H -NMR spectra illustrate the chemical structures of designated compounds (SG13-135, SG-13-136, SG13-138, SG13-140, SG13-141, and SG13-158).



Supplementary Figure 1. Related to Figure 2 and 3. Reduction of cytotoxicity, absence of estrogenic activity, and no inhibition of Tau aggregation of SG13-136.

(A) Trypan blue exclusion cell viability assessment conducted in SH-SY5Y cells transfected with β -gal, Myc empty vector control, or Myc-AIMP2 (72 h) (n = 6 per group).

(B) Quantification of the estrogen responsive gene, *CCND1* messenger RNA levels in a human breast tumor cell line, MCF7, treated with the specified compounds (10 μ M, 48 h), monitored by real-time quantitative PCR (n = 3 separate experiments per group). *GAPDH* served as an internal loading control for normalization.

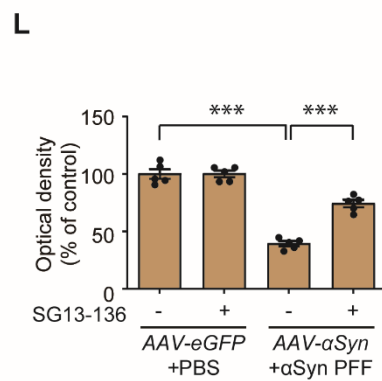
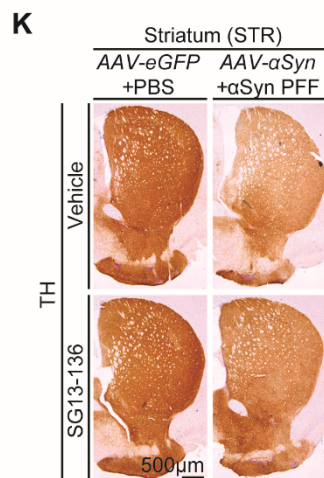
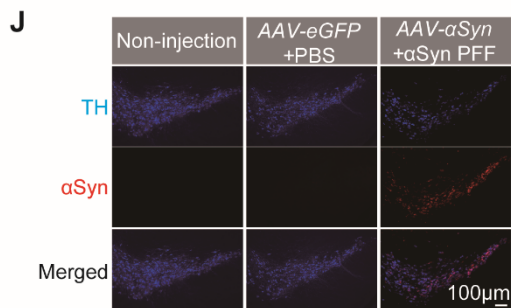
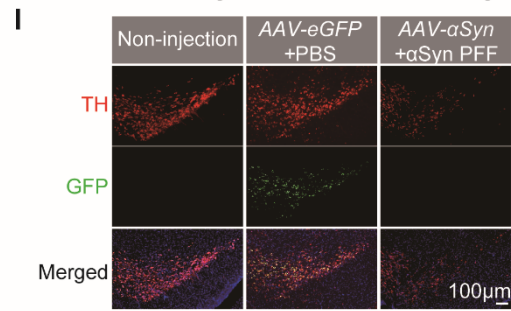
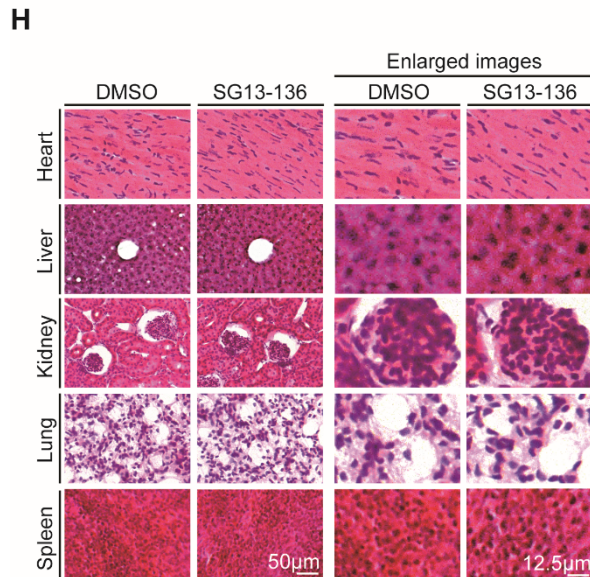
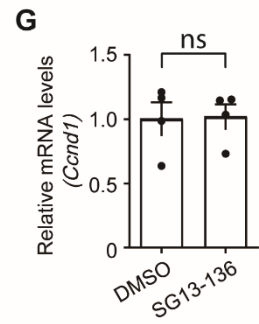
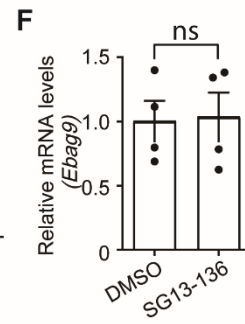
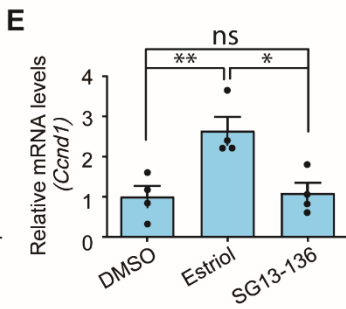
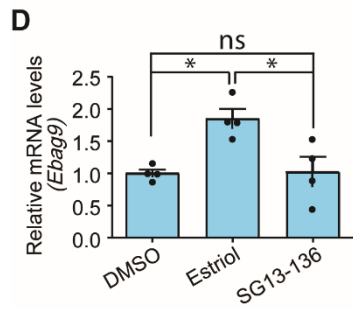
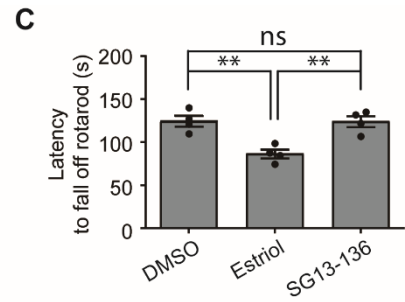
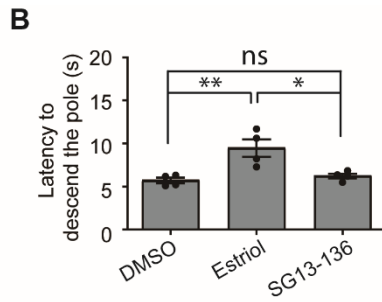
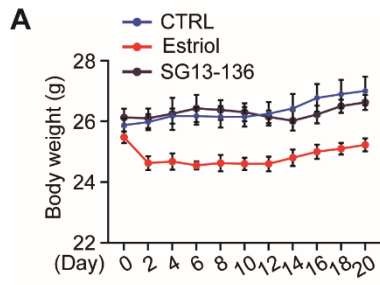
(C, D) Quantification of estrogen responsive genes, *E2F1*, *BCL2* messenger RNA levels in the human breast tumor cell line, MCF7 treated with the indicated compounds (10 μ M, 48 h) (n = 3 separate experiments per group). *GAPDH* served as internal loading control for normalization.

(E) Trypan blue exclusion cell viability assessment conducted in SH-SY5Y cells treated with increasing concentrations of estriol or SG13-136 (0, 10, 50, 100, 200, 500 μ M, 48 h) (n = 6 separate experiments per group).

(F) Chemical structure of biotin-conjugated SG13-136 (also named SG13-158), along with chemical formula and mass.

(G) Amyloid-like aggregation of recombinant tau incubated for three days (20 μ M in PBS with 5 μ M heparin) in the presence of indicated compounds (50 μ M), monitored by thioflavin T fluorescence assay (n = 3 per group). Thioflavin T measurement of recombinant tau at day 0 served as baseline, with DMSO as vehicle control.

Data in all panels represent mean \pm standard error of the mean. * P < 0.05, ** P < 0.01, and *** P < 0.001, determined by one-way (A, B, C, D, and E) or two-way (G) analysis of variance (ANOVA) followed by Tukey's post hoc analysis.



Supplementary Figure 2. Related to Figure 5. The steroid derivative, SG13-136, devoid of estrogenic activity, showed no organ toxicity and conferred protection to dopaminergic axon in a α Syn PFF/AAV- α Syn injection PD mouse model.

(A) Measurement of mouse body weights (every two days) with daily administrations of vehicle, estriol, or SG13-136 (0.5 mg/kg/d, p.o. for 20 days) (n = 4 mice per group).

(B) Assessment of bradykinesia in wild-type mice administered with vehicle DMSO, estriol, or SG13-136 (0.5 mg/kg, p.o. once daily for three weeks) or vehicle monitored by pole test (n = 4 mice per group).

(C) Assessment of motor coordination in wild-type mice administered with vehicle DMSO, estriol, or SG13-136 (0.5 mg/kg, p.o. once daily for three weeks) or vehicle monitored by pole test (n = 4 mice per group).

(D) Quantification of the estrogen responsive gene, *Ebag9* messenger RNA levels in brains from the mice with the indicated compound administration (0.5 mg/kg/d estriol, or SG13-136, p.o., for three weeks) monitored by real-time quantitative PCR and normalized by *Gapdh* internal loading control (n = 4 mice per group).

(E) Quantification of the estrogen responsive gene, *Ccnd1* messenger RNA levels in brains from the mice with the indicated compound administration (0.5 mg/kg/d estriol, or SG13-136, p.o., for three weeks) monitored by real-time quantitative PCR and normalized by *Gapdh* internal loading control (n = 4 mice per group).

(F, G) Quantification of the estrogen responsive genes, *Ebag9*, *Ccnd1* messenger RNA levels in liver tissues from the mice with the administration of SG13-136 (0.5 mg/kg/d, p.o., for three weeks) monitored by real-time quantitative PCR and normalized by *Gapdh* internal loading control (n = 4 mice per group).

(H) Representative H&E staining images of tissue sections (heart, liver, kidney, lung, and spleen) from the mice administered with vehicle or SG13-136 (0.5 mg/kg/d, p.o., for three weeks). Enlarged images were also presented in the right panel. Scale bar = 50, and 12.5 μm , respectively.

(I, J) Representative co-immunofluorescence images showing the expression and distribution of eGFP and α -synuclein in dopaminergic neurons labeled with TH on coronal ventral midbrain sections from each experimental mouse groups. Scale bar = 100 μm .

(K) Representative anti-TH immunohistochemistry images of the striatum coronal sections from each mouse group with the indicated nigral injections and SG13-136 p.o. administration. Scale bar = 500 μm .

(L) Quantification of relative TH-stained dopaminergic axon terminal densities in the striatum of the indicated experimental groups (n = 5 mice per group).

Data in all panels represent mean \pm standard error of the mean. * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$, determined by unpaired two-tailed Student's t-test (F, and G), one-way (B, C, D, and E) or two-way (L) analysis of variance (ANOVA) followed by Tukey's post hoc analysis. n.s., non-significant.