

Supplementary Information for:

Fragment-centric Topographic Mapping Method Guides the Understanding of ABCG2-inhibitors Interaction

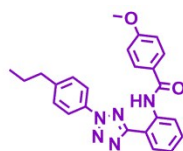
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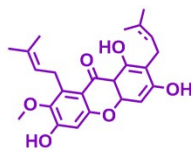
^b College of Physics and Electronics, Shandong Normal University, Jinan 250014, P. R. China

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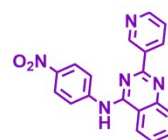
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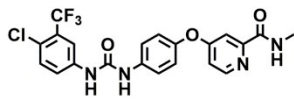
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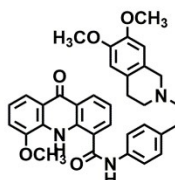
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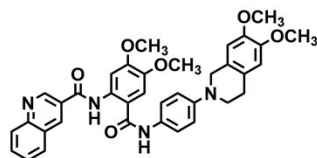
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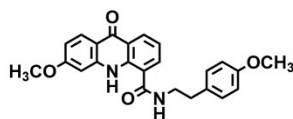
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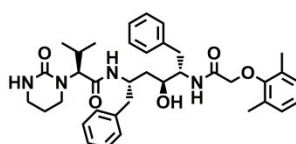
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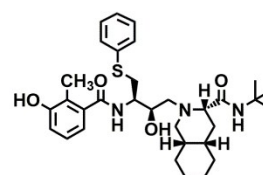
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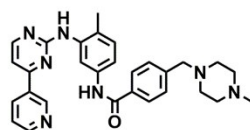
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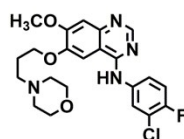
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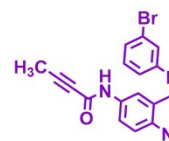
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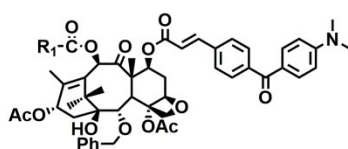
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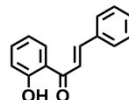
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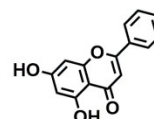
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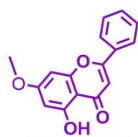
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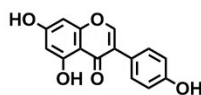
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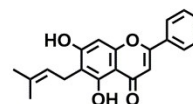
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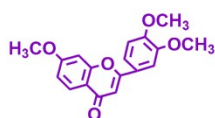
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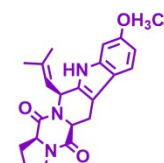
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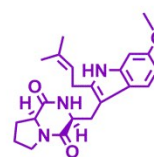
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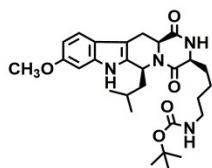
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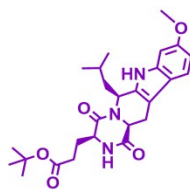
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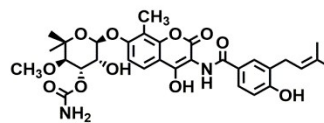
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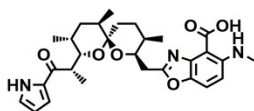
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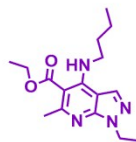
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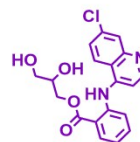
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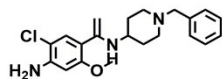
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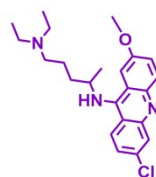
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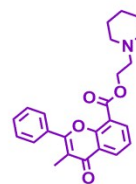
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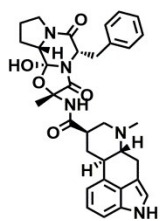
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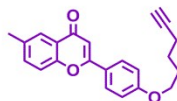
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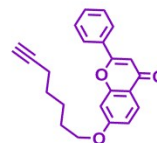
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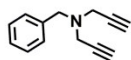
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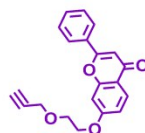
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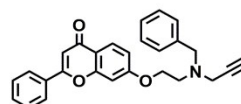
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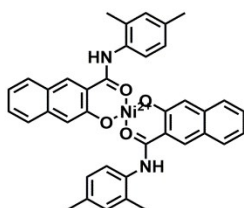
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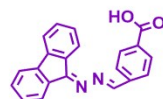
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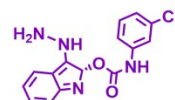
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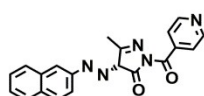
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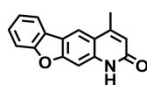
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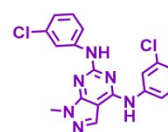
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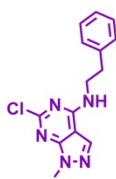
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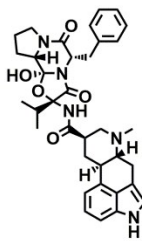
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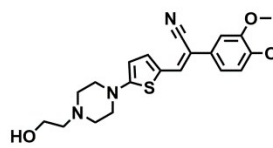
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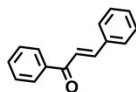
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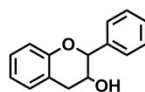
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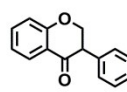
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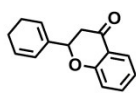
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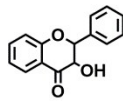
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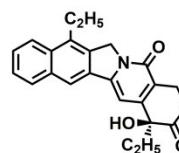
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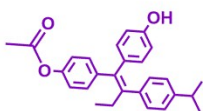
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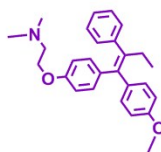
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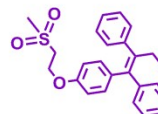
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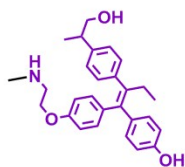
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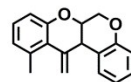
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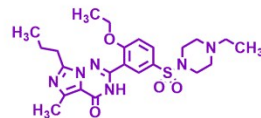
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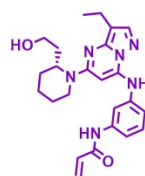
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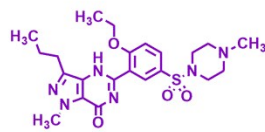
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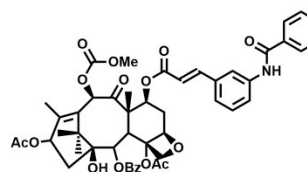
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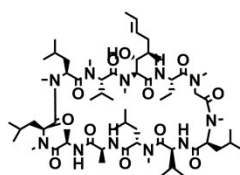
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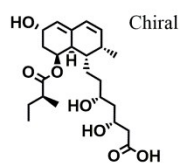
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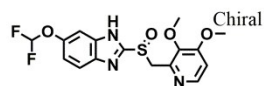
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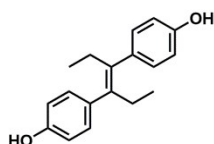
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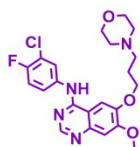
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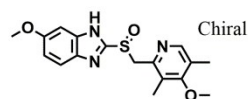
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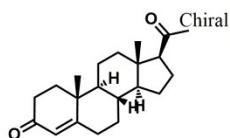
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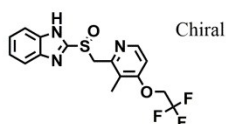
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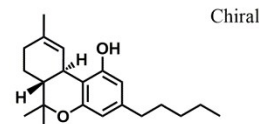
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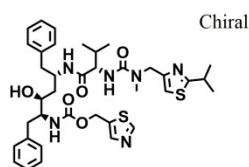
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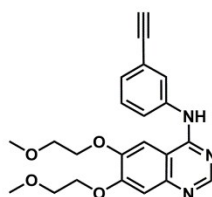
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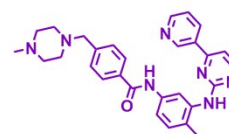
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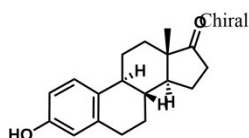
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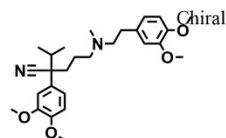
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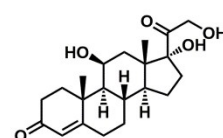
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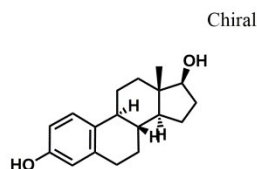
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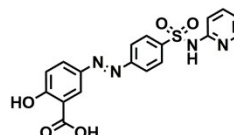
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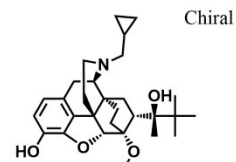
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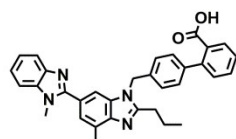
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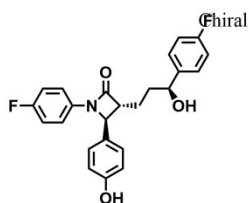
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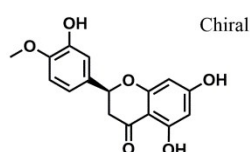
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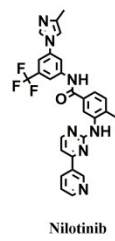
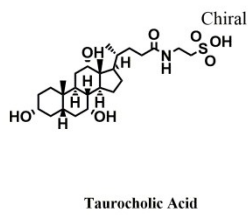
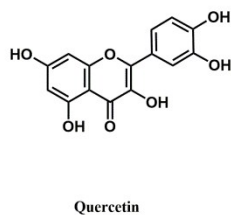
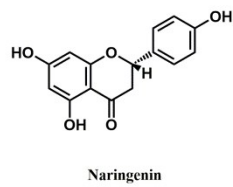
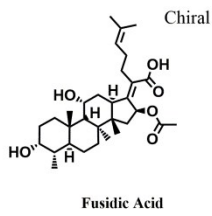
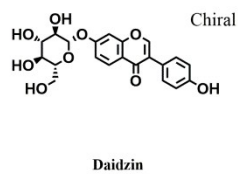
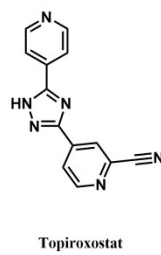
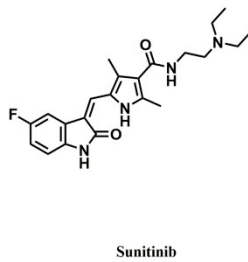
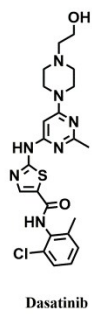
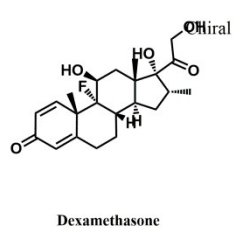
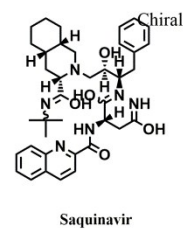
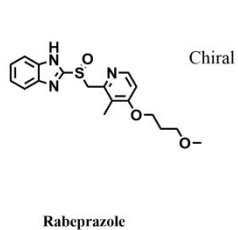
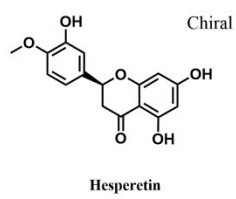
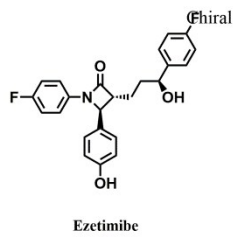
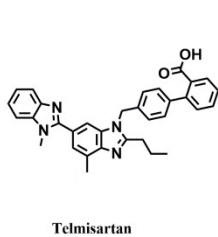
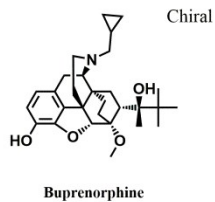
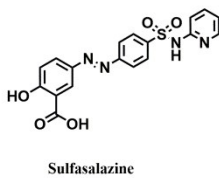
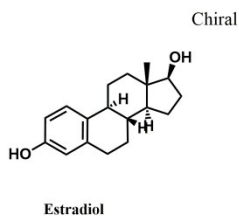
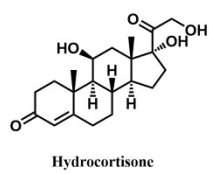
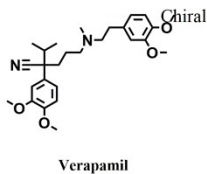
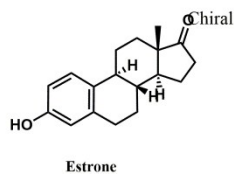
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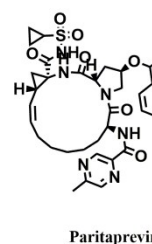
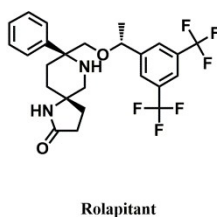
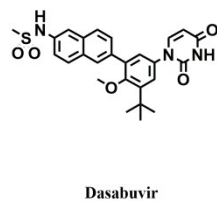
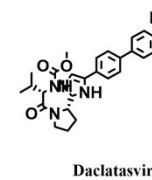
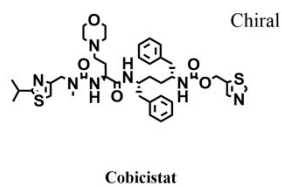
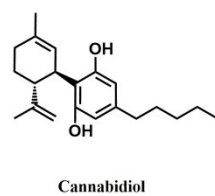
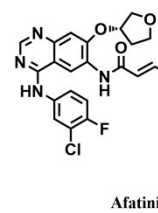
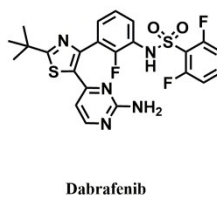
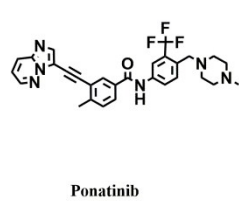
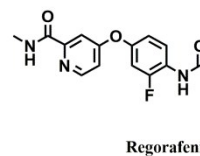
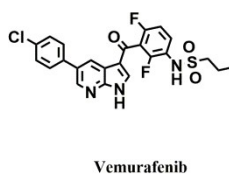
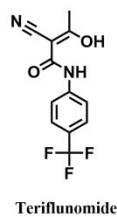
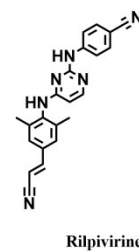
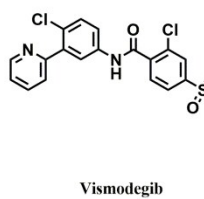
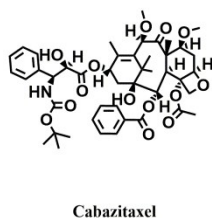
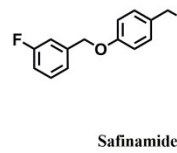
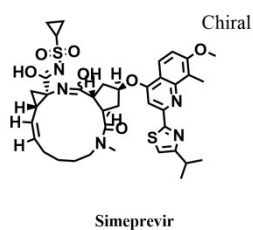
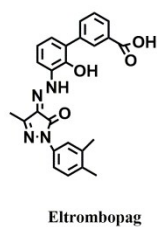
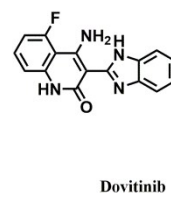
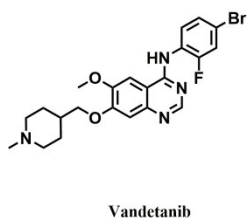
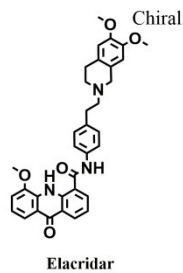


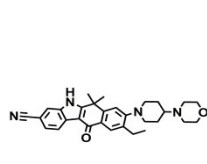
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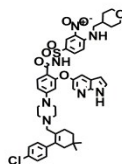
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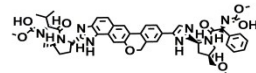




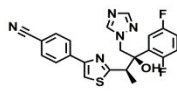
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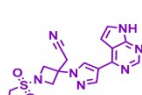
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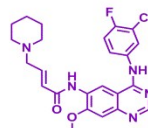
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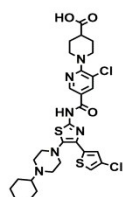
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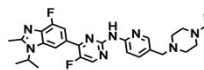
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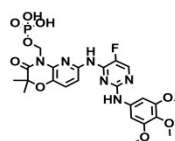
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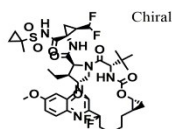
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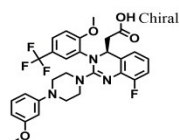
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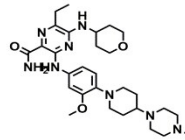
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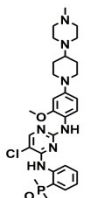
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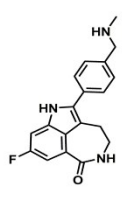
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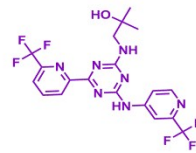
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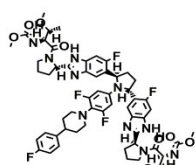
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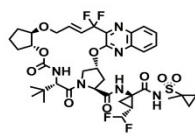
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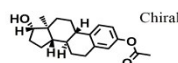
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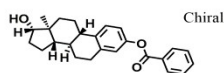
Pibrentasvir



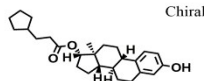
Glecaprevir



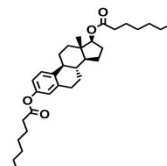
Estradiol acetate



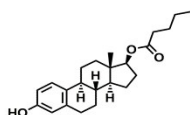
Estradiol benzoate



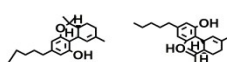
Estradiol cypionate



Estradiol diethanthate



Estradiol valerate



Nabiximols

Inhibitor name	Reference
HM30181 derivatives	1
Alpha-Mangostin	2
4-Anilino-2-pyridylquinazoline	3
Sorafenib	4
GF120918	5
Tariquidar	5
analogue of GF 120918	5
Lopinavir	6
Nelfinavir	6
Gleevec	7
ZD1839	7
EKI-785	7
tRA98006	8
2'-Hydroxychalcone	9-11
Chrysin	9-11
Techtochrysin	9-11
Genistein	9-11
6-Prenylchrysin	9-11
3',4',7-Trimethoxyflavone	9-11
Fumitremorgin C	12
Tryprostatin A	12
Ko132	12
Ko134	12
Novobiocin	13
Calcimycin	14
Tracazolate	14
Glafenine	14
Clebopride	14
Mepacrine	14
Flavoxate	14
Dihydroergotamine	14
Ac5	15
Ac12	15
Ac15	15
Ac13	15
Ac13	15
NSC306698	16
NSC120668	16
NSC320582	16
NSC303769	16
NSC375985	16
NSC11668	16

NSC19139	16
NSC409663	16
EAC 14	17
Chalcones	18
Flavanols	18
Isoflavones	18
Flavanones	18
Flavonols	18
SN-38 analogues	15
TAG-1	19
TAG-2	19
TAG-3	19
TAG-7	19
Rotenoid derivatives	20
Vardenafil	21
E9	22
Sildenafil	23
Paclitaxel analogues	24

Fig. S1 Small molecules library of ABCG2 inhibitors. The inhibitors marked purple are found to meet the criteria. And inhibitors are from DrugBank (from Ciclosporin to Nabiximols)

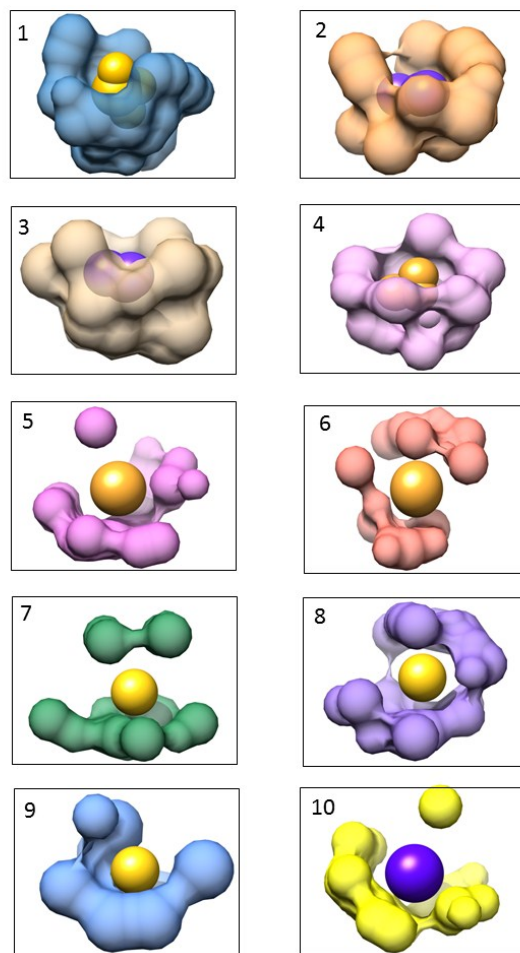


Fig. S2 The shape of ten pockets of central cavity are shown by AlphaSpace. According to the results, the pockets are divided into categories, we can observe that pockets NO.1, NO.2, NO.3, NO.4 belong to deep pocket, which can firmly wrap the atomic group, and compared with these cavities, the others have more planar structure.

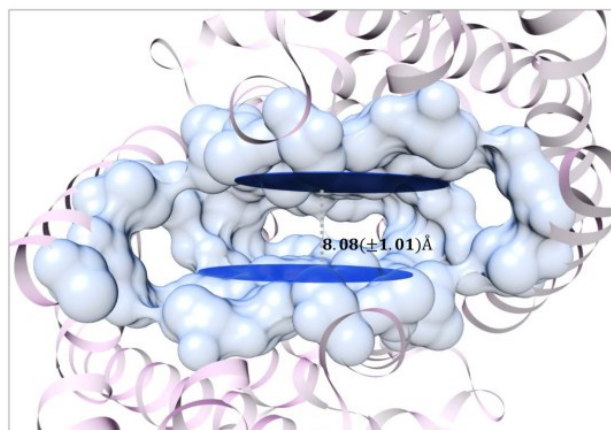


Fig. S3 The height of cavity is 8.08 ± 1.01 Å.

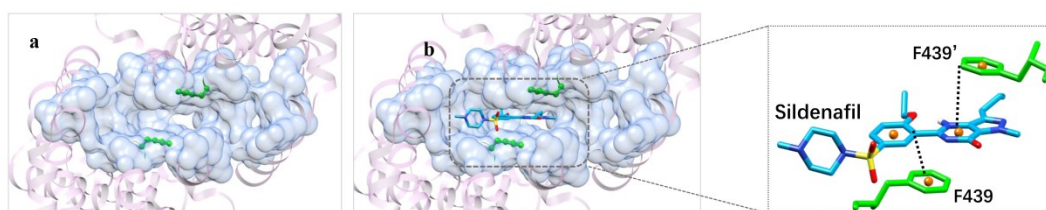


Fig. S4 (a & b) The inhibitor and the residues (F439, F439') located in the upper and lower plane of cavity form π - π staking.



Fig. S5 The docking poses of vardenafil (a) and E9 (b) in pocket-centric topographical map calculated by *AlphaSpace*. And Auto dock scoring parameters are listed.

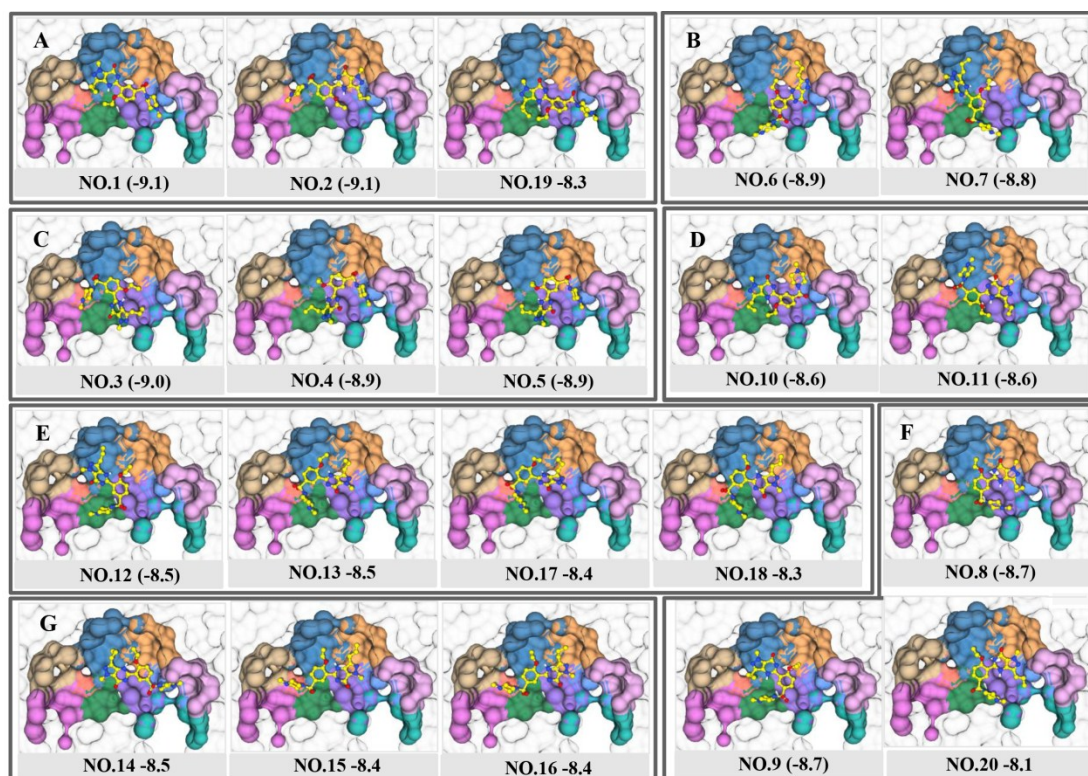


Fig. S6 20 docking poses of sildenafil are divided into 7 categories. And Auto dock scoring parameters are listed.

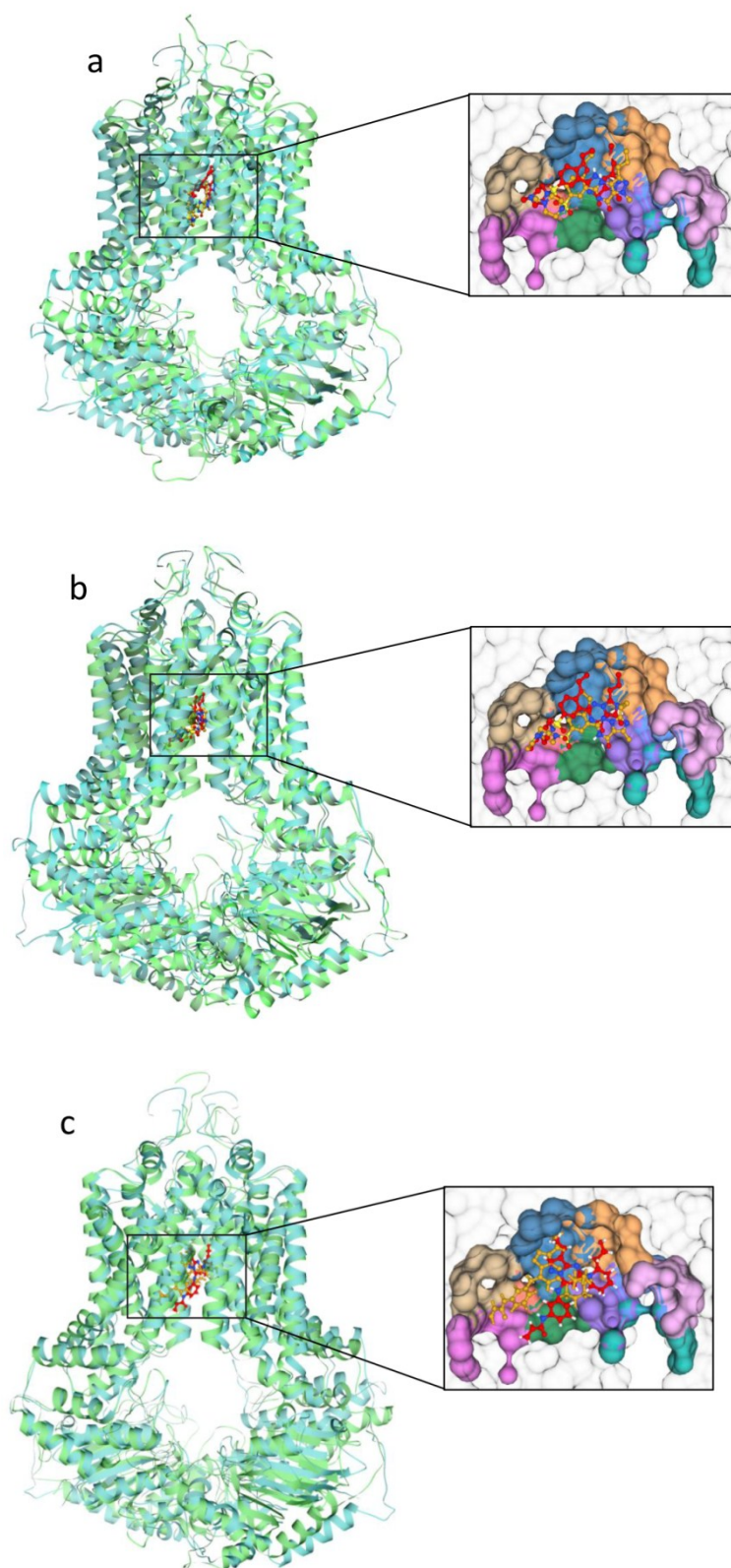


Fig. S7 Docking (protein light sea green, ligand orange) and after MD simulated (protein lime green, ligand red) superpose structures of all the simulated ligands (a Sildenafil, b Vardenafil, and c E9) inside binding pocket.

Table S1 Auto dock energy scoring parameters.

	Sildenafil	Vardenafil	E9
NO.1	-9.1	-9.6	-9.3
NO.2	-9.1	-9.2	-8.9
NO.3	-9.0	-9.0	-8.8
NO.4	-8.9	-9.0	-8.7
NO.5	-8.9	-8.9	-8.5
NO.6	-8.9	-8.8	-8.4
NO.7	-8.8	-8.7	-8.3
NO.8	-8.7	-8.6	-8.2
NO.9	-8.7	-8.6	-8.1
NO.10	-8.6	-8.6	-8.1
NO.11	-8.6	-8.6	-8.0
NO.12	-8.5	-8.5	-8.0
NO.13	-8.5	-8.4	-8.0
NO.14	-8.5	-8.4	-7.9
NO.15	-8.4	-8.3	-7.9
NO.16	-8.4	-8.1	-7.9
NO.17	-8.4	-8.0	-7.9
NO.18	-8.3	-7.8	-7.6
NO.19	-8.3	-7.8	-7.6
NO.20	-8.1	-7.8	-7.5

Table S2 Binding interaction of inhibitors to ABCG2 obtained by MM-PBSA method.

Energy component	Sildenafil (KJ/mol)	Vardenafil (KJ/mol)	E9 (KJ/mol)
ΔE_{ele}	-10.74 ± 3.59	-9.21 ± 1.97	-18.99 ± 3.60
ΔE_{vdw}	-69.70 ± 2.27	-71.93 ± 2.46	-64.46 ± 2.75
ΔG_{pol}	47.94 ± 6.14	50.90 ± 5.68	43.43 ± 6.67
ΔG_{nopol}	-7.20 ± 0.10	-7.39 ± 0.11	-6.93 ± 0.10
ΔH	-39.70 ± 6.06	-37.63 ± 6.40	-46.96 ± 6.72

**Table S3 The high-scoring amino acid residues in each pocket of the three complexes.
(The depth of the color depending on the frequency of amino acid appearance)**

Pocket	Sildenafil	Vardenafil	E9
Pocket 1	F431	F431	F431
	T435	T435	T435
	M549	M549	M549
	M549'	M549'	F432
	-	L555	-
	L555'	L555'	L555'
Pocket 2	F431'	F431'	F431'
	F432'	F432'	F432'
	T435'	T435'	T435'
	V546	V546	V546
	-	-	N436'
	-	-	M549'
	-	-	L555
Pocket 4	A397'	N436'	Q398'
	V401'	S440'	V401'
	L405'	S443'	S440'
	S408'	-	-
	N436'	-	-
	L539	-	-
	I543	-	-
Pocket 7	F439	F439	N436
	F439'	F439'	F439
	T542	F545'	F439'
	T542'	V546'	V546'
Pocket 8	-	L539	T542
	-	-	T542'
Pocket 9	F432	V401'	-
	V546'	T402'	-
	-	L405'	-
	-	I543	-

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