

# Screening of JAK2 Inhibitors from Natural Curcumin and Its Derivatives as Erythropoiesis Stimulant Agents for Thalassemia Therapy: Computer Aided Molecular Design Approaches



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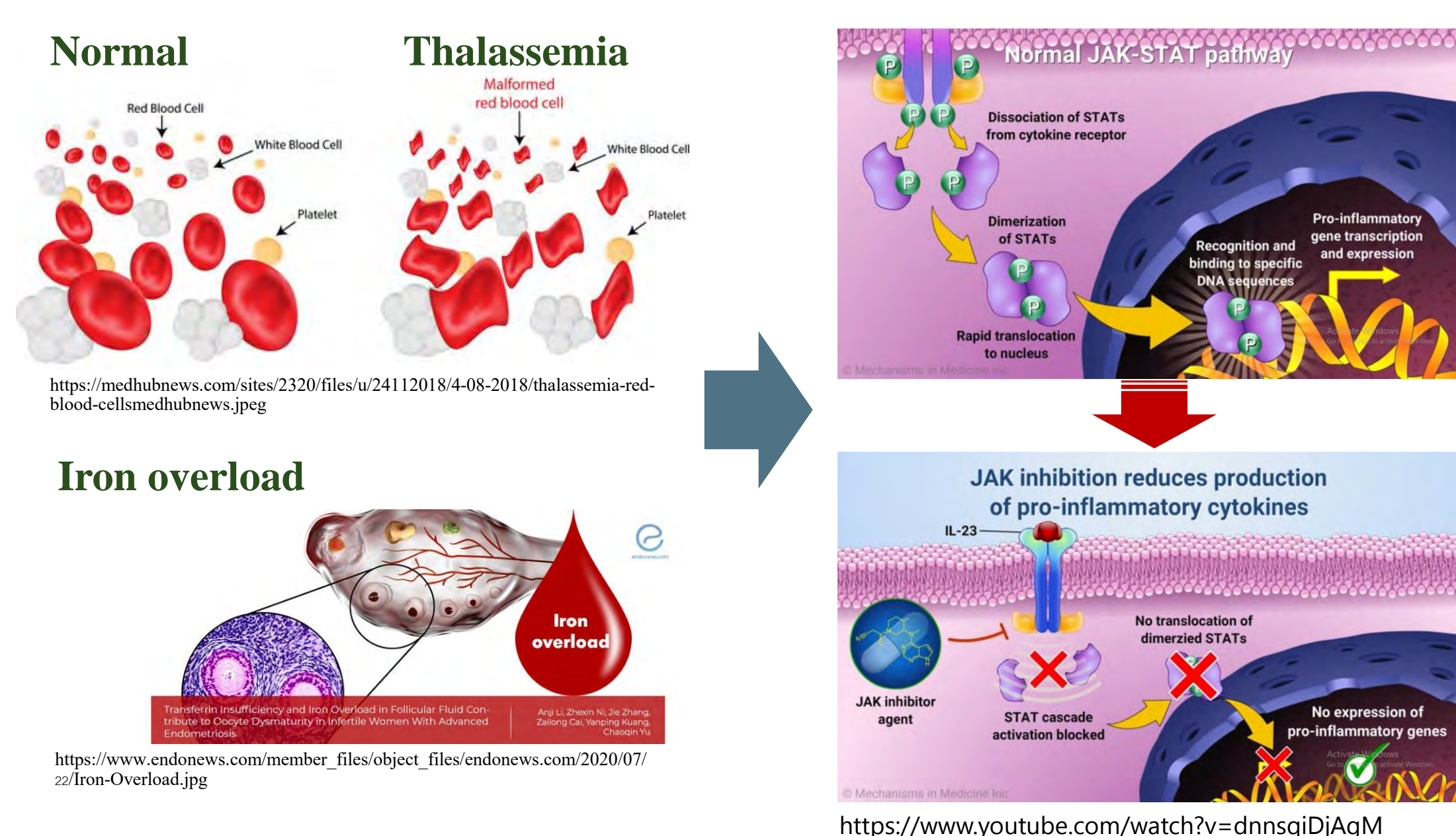
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## Introduction



To developed curcumin and its derivatives as JAK2 inhibitors and erythropoiesis stimulant agents

Computer Aided Molecular Design (CAMD)

- Virtual screening
- Biological predictions
- Pharmacokinetic predictions

Janus kinase 2 (JAK2) is an enzyme responsible for regulating erythropoiesis and has been validated as a novel therapeutic process for thalassemia.

## Materials and Methods

**Curcuminoid analog**  
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**Erythropoiesis stimulant**  
PASS prediction

**Pharmacokinetic properties (ADMET)**  
pkCSM

**Molecular Docking Calculations**  
PDB code: 4D0X  
Glide XP program

**Figure 1** The virtual screening process of natural curcumin and its derivatives to be JAK2 inhibitors

## Conclusions

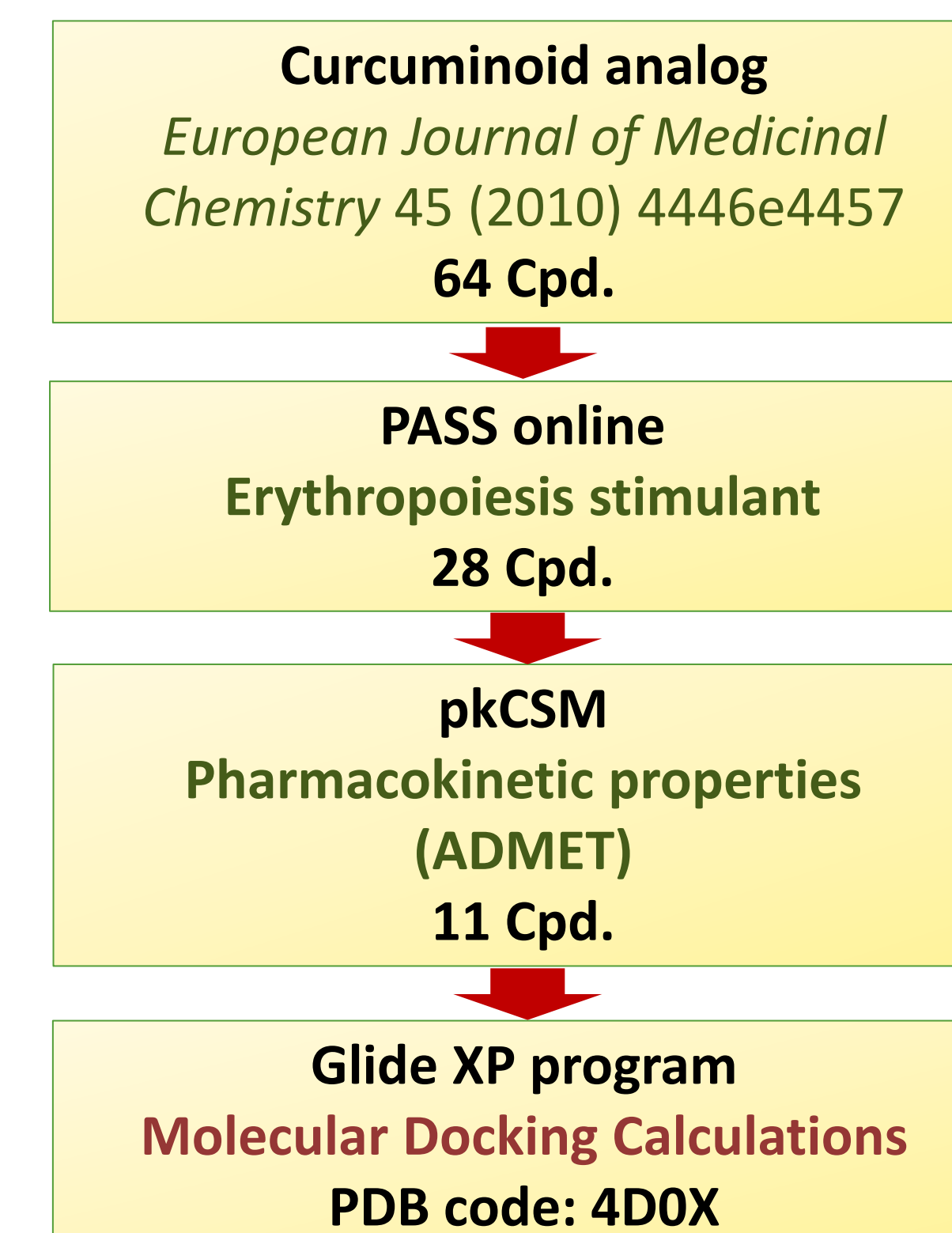
- Eleven curcumin derivatives will be act as JAK2 inhibitors with the high binding affinity in JAK2 binding site docking score ranging from -9.65 to -12.09 kcal/mol.
- H-bond interaction with NH backbone of Leu932 and sigma-pi interaction of aromatic ring on curcumin analog with Leu983 sidechain
- The ADMET predictions suggested that curcumin derivatives were suitable for acting as drug.
- These finding results aid to collect the potential compounds for biological assay evaluations and development as novel drug for thalassemia therapy based on JAK2 inhibition mechanism.

## Results

### Biological activity prediction and molecular docking studies

**Table 1** The results of the erythropoiesis stimulant activity using PASS online prediction and the docking score from molecular docking calculations

Compound	Structures	Erythropoiesis stimulant activity		Docking Score (kcal/mol)
		Pa	Pi	
1		0.307	0.172	-12.086
2		0.332	0.144	-11.364
3		0.273	0.216	-11.326
4		0.273	0.216	-11.001
5		0.334	0.143	-10.634
6		0.298	0.182	-10.618
7		0.274	0.213	-10.510
8		0.267	0.224	-10.449
9		0.523	0.031	-10.084
10		0.279	0.207	-9.948
11		0.353	0.124	-9.650



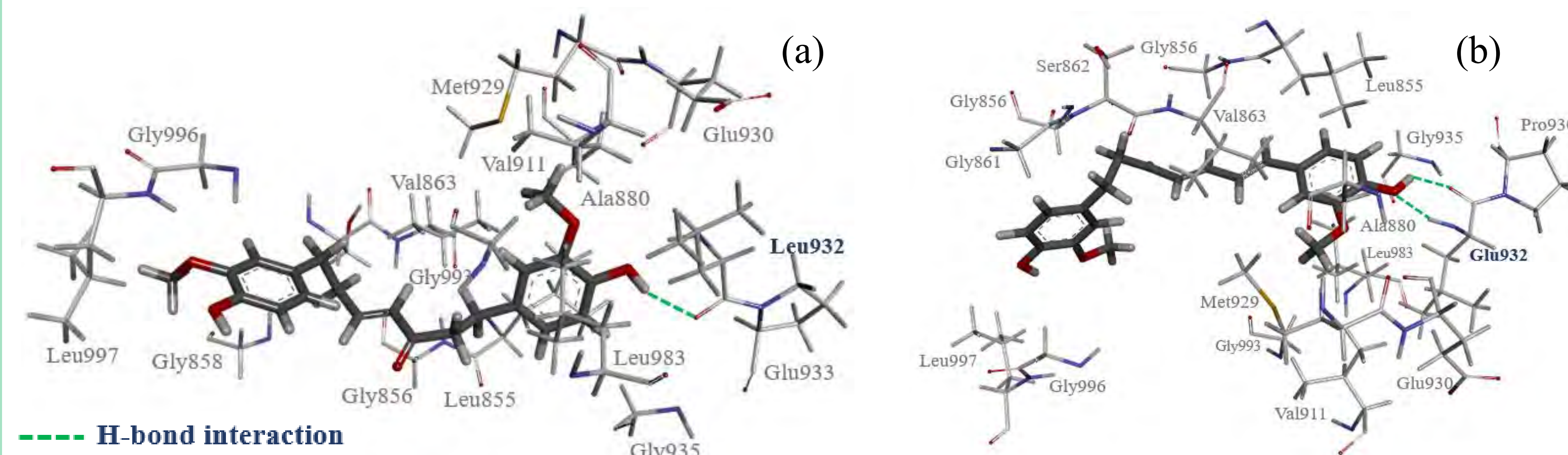
**Figure 2** The virtual screening process of natural curcumin and its derivatives to be JAK2 inhibitors

### The pharmacokinetic properties (ADMET) prediction

**Table 2** The results of ADMET prediction of hit compounds

Compound	1	2	3	4	5	7	8	9	10	11		
Caco2 permeability	1.113	1.079	1.135	1.07	1.096	0.915	0.923	1.103	1.084	1.158	1.152	Caco2 > 0.90
Intestinal absorption (human)	92.9	92.394	92.172	92.37	95.146	89.844	92.74	88.96	92.183	92.523	97.999	high Caco2 permeability
Pgp inhibition	Yes	Yes	Yes	Yes	Yes	Yes	Yes	No	Yes	Yes	Yes	
BBB permeability	-0.314	-0.35	-0.286	-0.285	-0.685	-1.19	-0.973	-0.854	-0.123	-0.259	-0.793	Intestinal absorption (human) < 30% is considered to be poorly adsorbed
CNS permeability	-2.603	-2.655	-2.486	-2.459	-3.011	-3.162	-3.042	-2.295	-2.322	-2.37	-2.971	
CYP2D6 substrate	No	No	No	No	No	No	No	No	No	No	No	
CYP3A4 substrate	Yes	Yes	Yes	Yes	Yes	Yes	Yes	No	No	Yes	Yes	
CYP2C19 inhibitor	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	BBB > 0.3 can readily cross the blood-brain
CYP2C9 inhibitor	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	BBB < -1 poorly distributed to the brain
CYP2D6 inhibitor	No	No	No	No	No	No	No	No	No	No	No	
CYP3A4 inhibitor	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	Yes	
Total Clearance (ml/min/kg)	0.271	0.364	0.244	0.145	0.149	0.431	0.432	0.145	0.044	0.14	0.188	
Renal OCT2 substrate	No	No	No	No	No	No	No	No	No	No	No	
AMES toxicity	No	No	No	No	No	No	No	No	Yes	No	No	CNS > -2 can penetrate the Central Nervous System (CNS)
hERG inhibitor	Yes	Yes	No	Yes	Yes	No	No	No	Yes	No	No	CNS < -3 unable to penetrate the CNS
Oral Rat Acute Toxicity (LD <sub>50</sub> )	2	2.003	2.016	2.064	2.155	2.468	2.396	2.288	1.851	2.032	2.233	
Hepatotoxicity	No	No	No	No	No	No	No	No	No	No	No	

### The binding mode and binding interactions of active compounds in JAK2 binding pocket



**Figure 3** The binding mode of natural curcumin (a) compound 1 and (b) compound 2 in JAK2 binding pocket

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