

Supplementary Information

Interaction of *Jania rubens* Polyphenolic Extract as an Antidiabetic Agent with α -Amylase, Lipase, and Trypsin: In Vitro Evaluations and In Silico Studies

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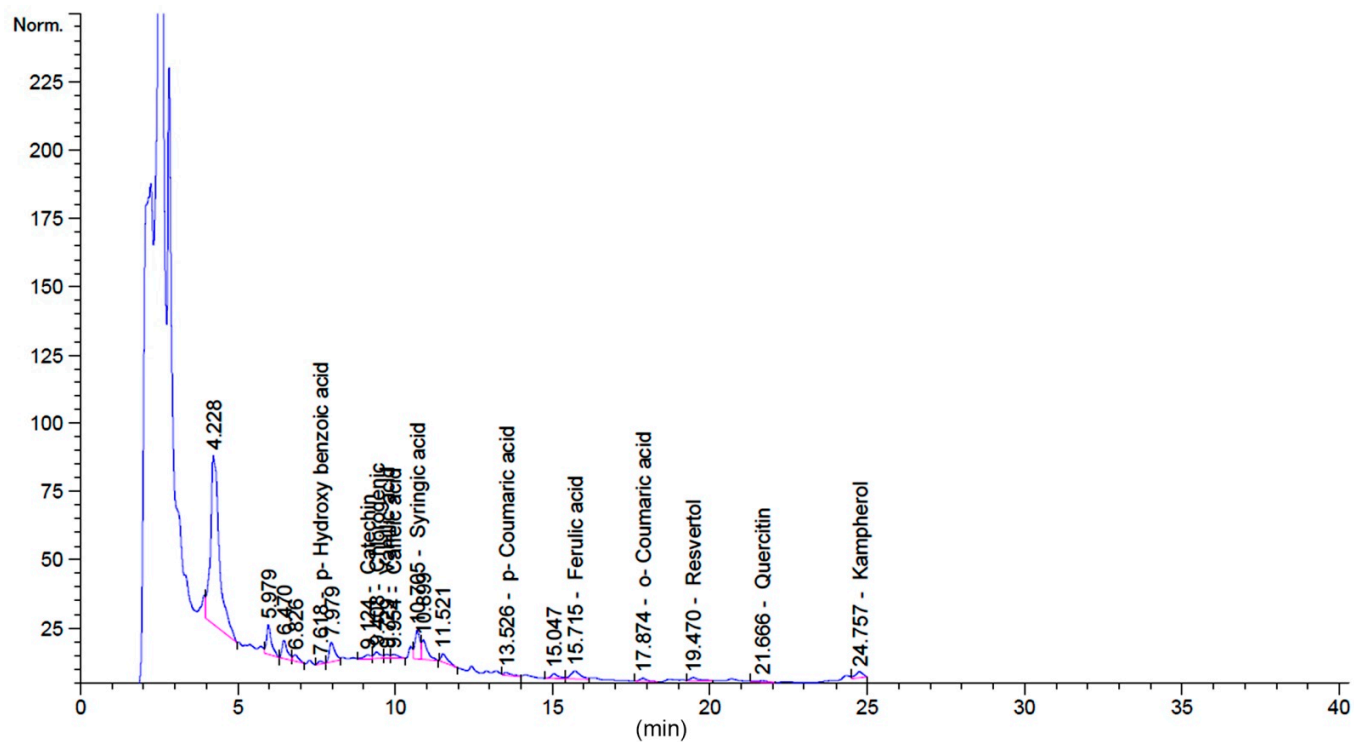
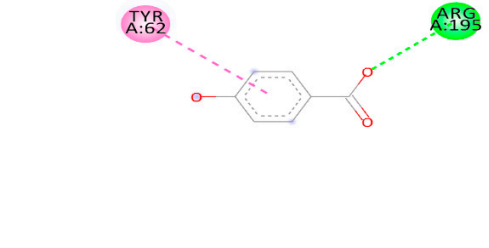
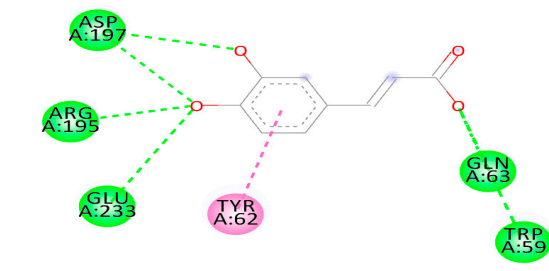
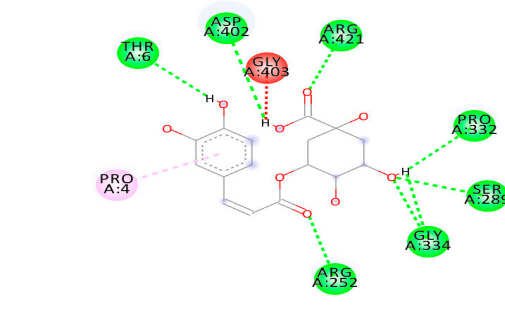
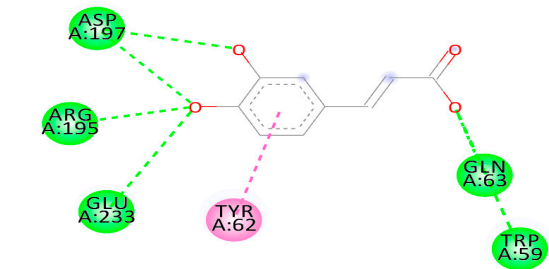


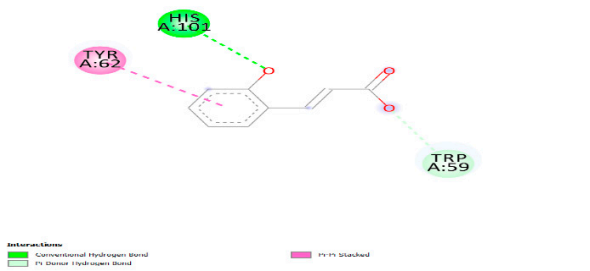
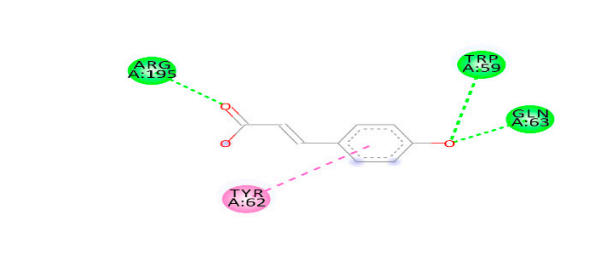
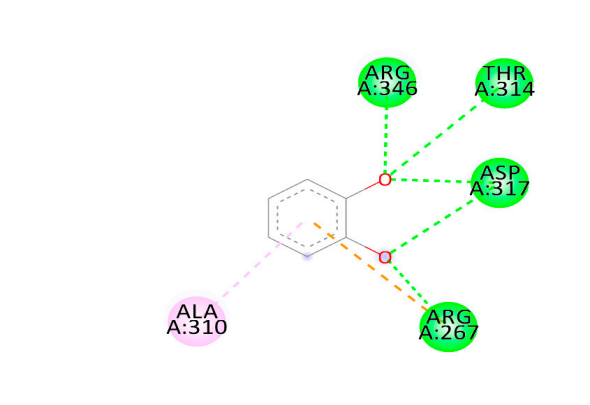
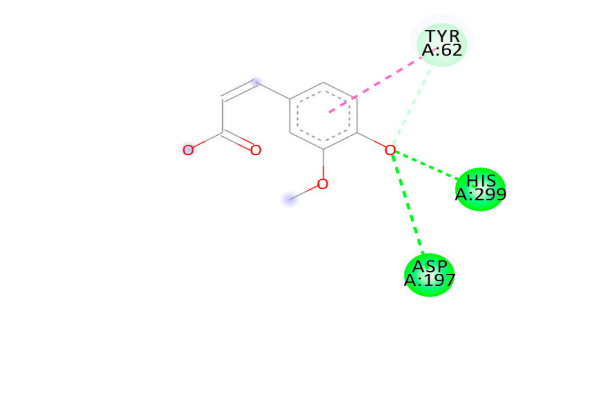
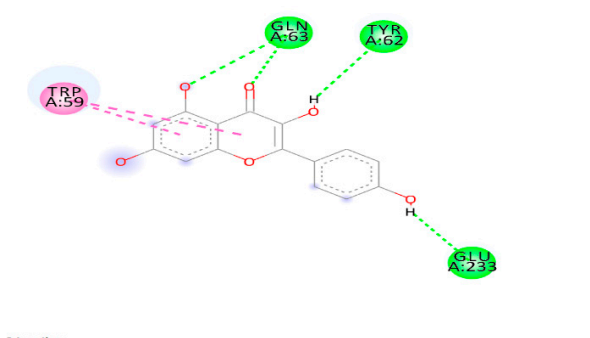
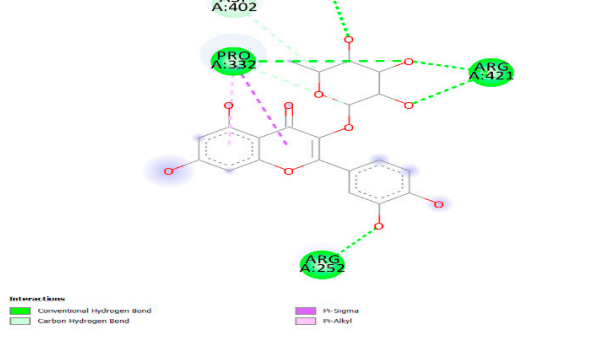
Figure S1. HPLC chromatogram for the JRPE shows different polyphenolic compounds (phenolic and flavonoids).

Table S1. The docking affinity scores of the different polyphenolics' compounds in the JRPE.

Compounds	α -Amylase Affinity	a.a residue	Lipase Affinity	a.a residue	Trypsin Affinity	a.a residue
<i>P</i> - Hydroxybenzoic acid	-5.3	TYR62 ,ARG195	-4.9	SER195 ,CYS220 ,GLY226	-5	SER195, CYS220 ,GLY226
Caffeic acid	-5.7	TRP59 ,TYR62 ,GLN63 ,ARG195 ,ASP197 ,GLU233	-4.7	CYS42 ,HIS57 ,TYR59 ,GLY193	-4.7	CYS42, HIS57 ,TYR59 ,GLY193
Catechin	-4.5	ARG267 ,ALA310 ,THR314 ,ASP317 ,ARG346	-4.3	SER195	-4.3	SER195
Chlorogenic	-8	PRO4 ,THR6 ,ARG252 ,SER289 ,PRO332 ,GLY334 ,ASP402 ,GLY403 ,ARG421	-7.1	LEU18 ,HIS30 ,ILE33 ,SER35 ,SER37 ,TYR55 ,ILE79 ,THR80	-7.1	PHE41 ,TYR151 ,GLY193 SER195 ,GLY219
Ferulic acid	-6.1	TYR62 ,ASP197 ,HIS299	-5.3	HIS57 ,SER190 ,CYS191 SER195 ,CYS220	-5.2	HIS57 ,SER190 ,CYS191 ,SER195 ,CYS220
Kaempferol	-8.4	TRP59, TYR62 ,GLN63 ,GLU233	-7.1	HIS57 ,ASP189 ,SER190 ,GLN192 ,GLY193 ,TRP215 ,GLY216	-7.2	HIS57 ,ASP189 ,SER190 ,GLN192 ,GLY193 ,TRP215 ,GLY216
<i>o</i> -Coumaric acid	-5.8	TRP59, TYR62 ,HIS101	-4.7	TYR59 ,THR90	-4.6	TYR59 ,THR90
<i>p</i> -Coumaric acid	-5.7	TRP59, TYR62 ,GLN63 ,ARG195	-4.8	ASN34 ,ARG62 ,GLN64 ,ARG66	-4.3	ASN34 ,ARG62, GLN64 ,ARG66
Quercetin	-8.8	ARG252 ,PRO332 ,ARG398 ,ASP402 ,ARG421	-7.4	HIS57 ,TYR151 ,CYS191 ,GLY193 ,ASP194 ,SER195	-7.2	HIS57 ,TYR151 ,CYS191 ,GLY193 ,ASP194 ,SER195
Resveratrol	-7.4	TYR62 ,THR163 ,ASP197 ,HIS299	-6	HIS57 ,LEU99 ,SER214 ,TRP215 ,GLY216	-5.8	HIS57 ,LEU99 ,SER214 ,TRP215 ,GLY216
Syringic acid	-5.8	ARG252 ,SER289 ,GLY334 ,PHE335 ,ARG398	-4.9	SER190 ,CYS191 ,GLY193 ,SER195 ,GLY216 ,GLY219	-5.5	SER190 ,CYS191 ,GLY193 ,SER195 ,GLY216 ,GLY219
Vanillic acid	-5.4	ARG252 ,SER289 ,ASP290 ,PRO332 ,GLY334 ,ARG398	-4.8	LYS60 ,SER61 ,ARG62	-4.1	LYS60 ,SER61 ,ARG62

Table S2. Affinity scores for docking analysis of α -amylase (PDB ID: 4W93) using twelve polyphenolic compounds identified from JRPE.

 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked
<p><i>P</i>-Hydroxybenzoic acid</p>	<p>Caffeic acid</p>
 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Unfavorable Donor-Donor Pi-Allyl 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked
<p>Chlorogenic</p>	<p>Vanillic acid</p>

 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Donor Hydrogen Bond Pi-Pi Stacked 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked
<p>O-Coumaric acid</p>	<p>P-Coumaric acid</p>
 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Cation Pi-Alkyl 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Donor Hydrogen Bond Pi-Pi Stacked
<p>Catechin</p>	<p>Ferulic acid</p>
 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Carbon Hydrogen Bond Pi-Sigma Pi-Alkyl
<p>Kaempferol</p>	<p>Quercetin</p>

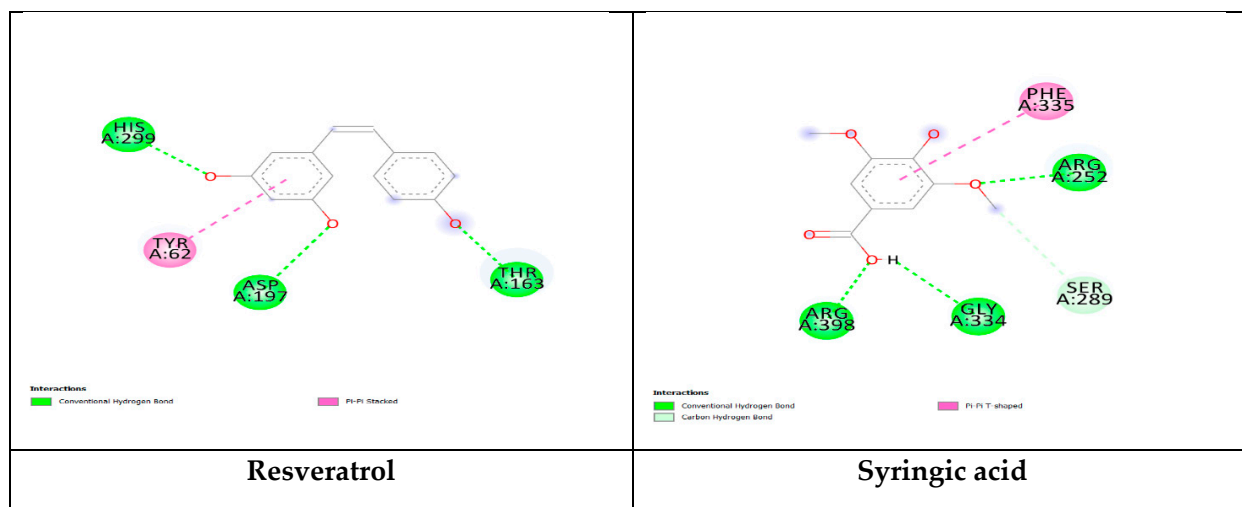
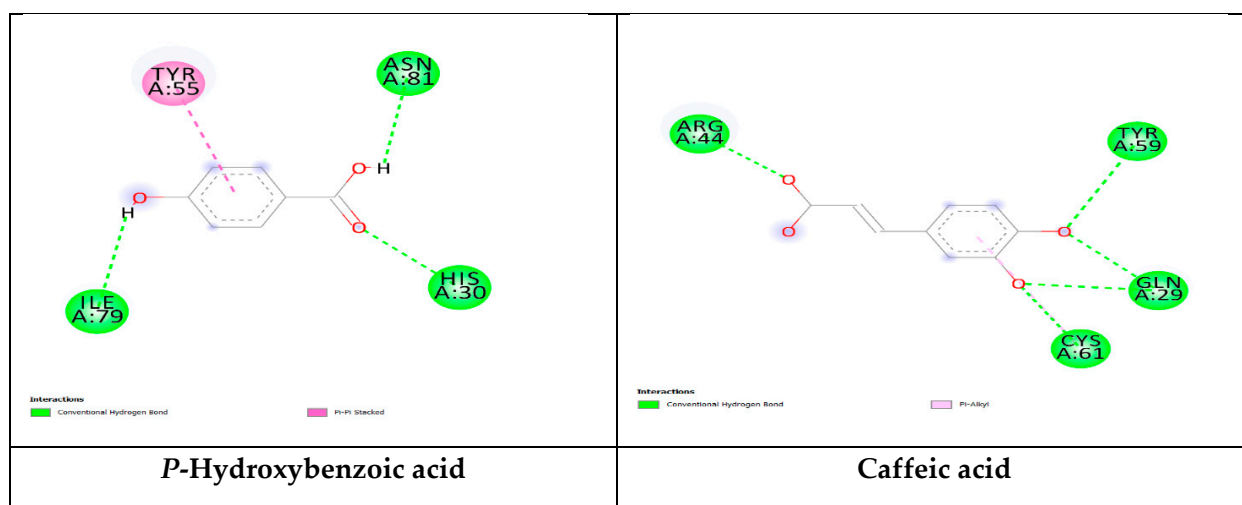
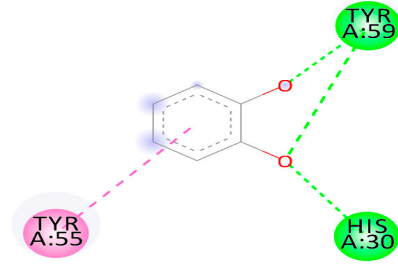
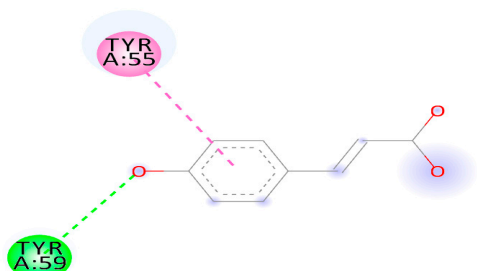
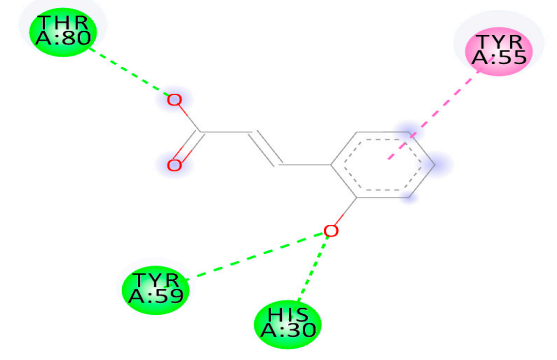
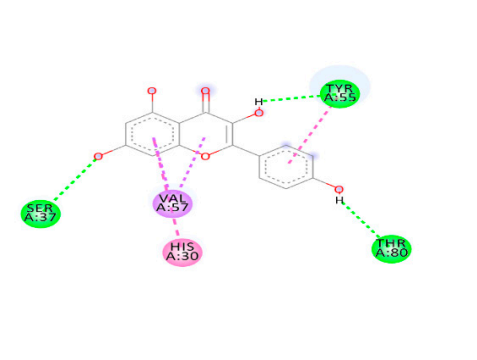
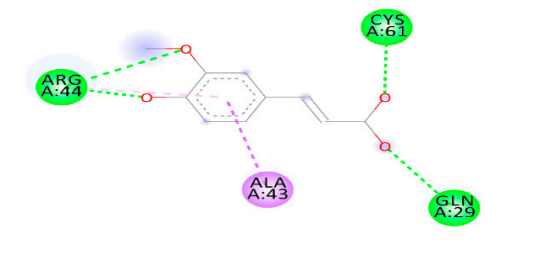
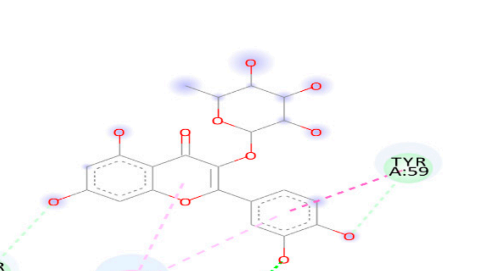


Table S3. Affinity scores for docking analysis of lipase (PDB ID: 1LPB) using twelve polyphenolic compounds identified from JRPE.



 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked
<p>Catechin</p>	<p>P-coumaric acid</p>
 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Sigma Pi-Pi Stacked Pi-Pi T-shaped
<p>o-Coumaric</p>	<p>Kaempferol</p>
 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Sigma Pi-Alkyl 	 <p>Interactions</p> <ul style="list-style-type: none"> Conventional Hydrogen Bond Pi-Pi Stacked Pi-Alkyl Pi-Donor Hydrogen Bond
<p>Ferulic acid</p>	<p>Quercetin</p>

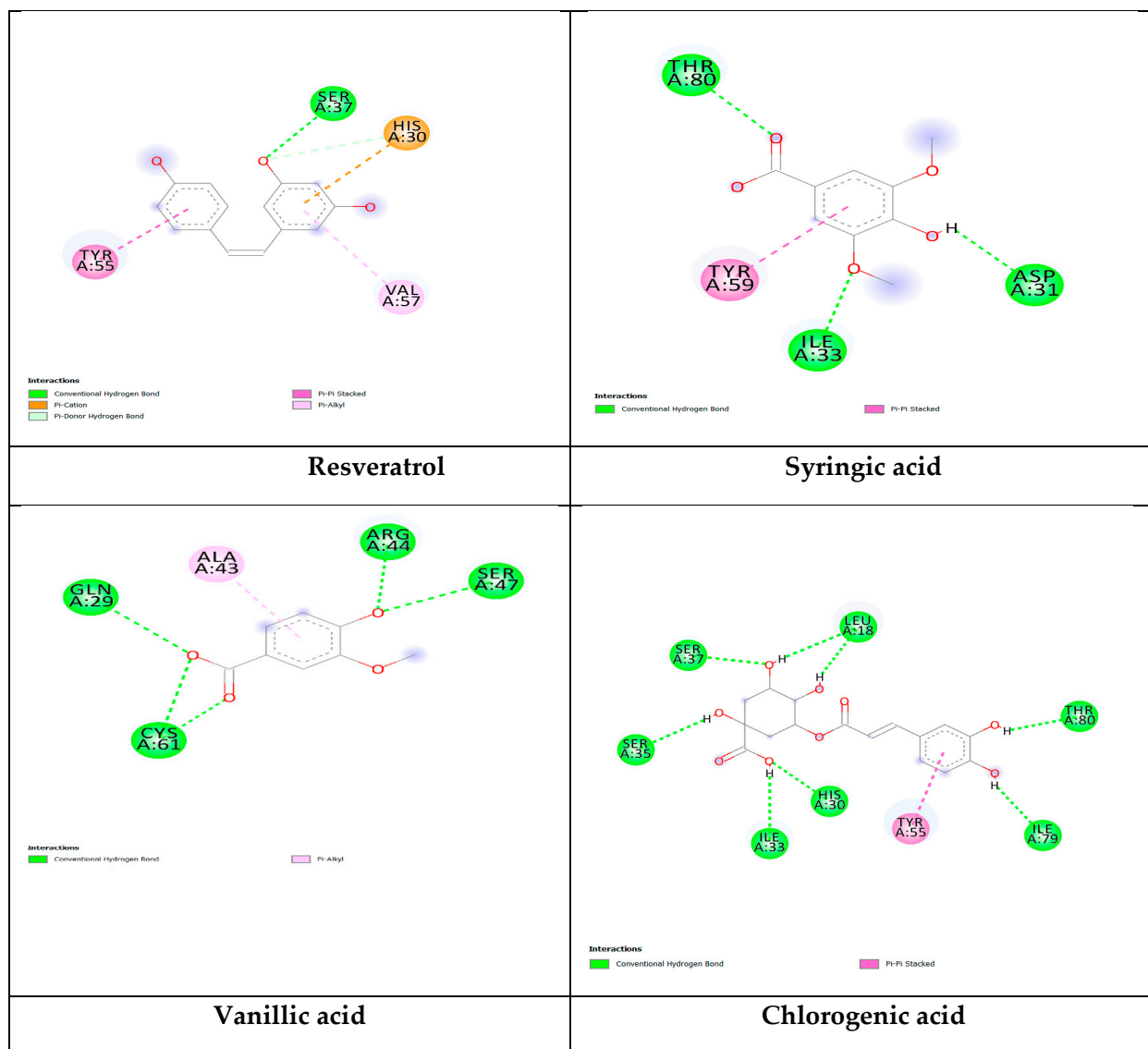
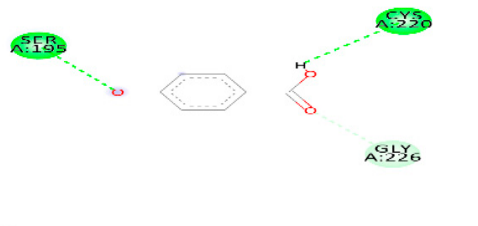
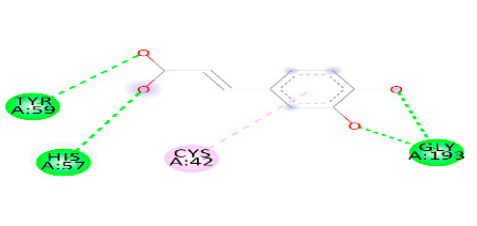
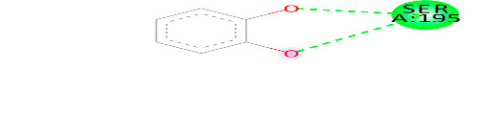
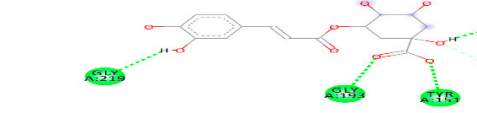
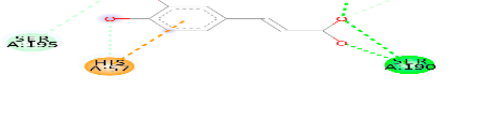
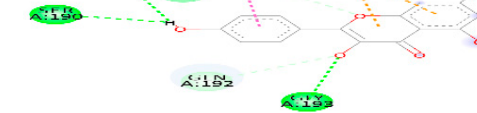
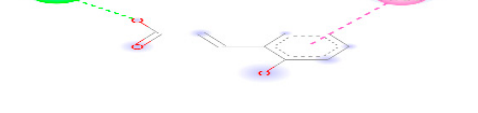
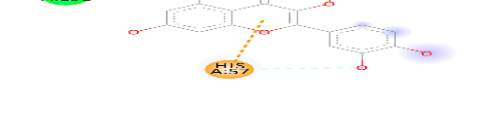


Table S4. Affinity scores for docking analysis of trypsin (PDB ID: 4DOQ) using twelve polyphenolic compounds identified from JRPE.

 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond 	 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond
<p><i>p</i>-Hydroxybenzoic acid</p>	<p>Caffeic acid</p>
 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond 	 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond
<p>Catechin</p>	<p>Chlorogenic acid</p>
 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond 	 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond
<p>Ferulic</p>	<p>Kaempferol</p>
 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond 	 <p>Active site:</p> <ul style="list-style-type: none"> SER A:19 GLY A:226 <p>Interactions:</p> <ul style="list-style-type: none"> Conventional hydrogen bond Carbon hydrogen bond
<p><i>o</i>-coumaric acid</p>	<p>Quercetin</p>

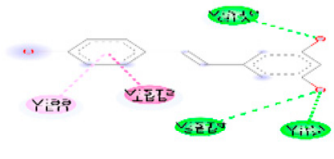
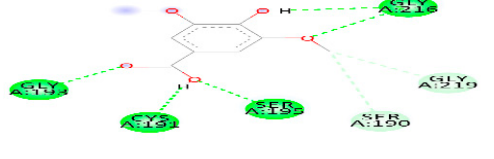
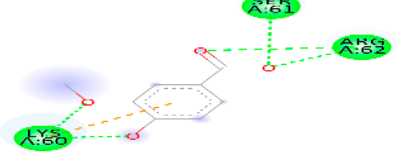
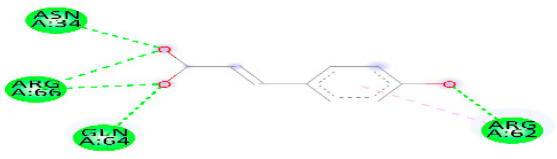
	
<p>Resveratrol</p>	<p>Syringic acid</p>
	
<p>Vanillic acid</p>	<p><i>p</i>-Coumaric acid</p>

Table S5. Pharmacodynamics and pharmacokinetics analysis of twelve polyphenolic compounds identified from JRPE.

Compounds	Pharmacokinetics	Medicinal Chemistry	Drug likeness	Physicochemical Properties	Lipophilicity	Water Solubility
Resveratrol	GI absorption High BBB permeant Yes P-gp substrate No CYP1A2 inhibitor Yes CYP2C19 inhibitor No CYP2C9 inhibitor Yes CYP2D6 inhibitor No CYP3A4 inhibitor Yes Log Kp (skin permeation) -5.47 cm/s	PAINS 0 alert Brenk 1 alert: stilbene Leadlikeness No; 1 violation: MW<250 Synthetic accessibility 2.02	Lipinski Yes; 0 violation Ghose Yes Veber Yes Egan Yes Muegge Yes Bioavailability Score 0.55	Formula C ₁₄ H ₁₂ O ₃ Molecular weight 228.24 g/mol Num. heavy atoms 17 Num. arom. heavy atoms 12 Fraction Csp ³ 0.00 Num. rotatable bonds 2 Num. H-bond acceptors 3 Num. H-bond donors 3 Molar Refractivity 67.88 TPSA 60.69 Å ²	Log Po/w (iLOGP) 1.71 Log Po/w (XLOGP3) 3.13 Log Po/w (WLOGP) 2.76 Log Po/w (MLOGP) 2.26 Log Po/w (SILICOS-IT) 2.57 Consensus Log Po/w 2.48	Log S (ESOL) -3.62 Solubility 5.51e-02 mg/ml ; 2.41e-04 mol/l Class Soluble Log S (Ali) -4.07 Solubility 1.93e-02 mg/ml ; 8.44e-05 mol/l Class Moderately soluble Log S (SILICOS-IT) -3.29 Solubility 1.18e-01 mg/ml ; 5.16e-04 mol/l Class Soluble
P-Coumarate	GI absorption High BBB permeant Yes P-gp substrate No CYP1A2 inhibitor No CYP2C19 inhibitor No CYP2C9 inhibitor No CYP2D6 inhibitor No CYP3A4 inhibitor No Log Kp (skin permeation) -6.26 cm/s	PAINS 0 alert Brenk 1 alert: michael_acceptor_1 Leadlikeness No; 1 violation: MW<250 Synthetic accessibility 1.61	Lipinski Yes; 0 violation Ghose Yes Veber Yes Egan Yes Muegge No; 1 violation: MW<200 Bioavailability Score 0.85	Formula C ₉ H ₈ O ₃ Molecular weight 164.16 g/mol Num. heavy atoms 12 Num. arom. heavy atoms 6 Fraction Csp ³ 0.00 Num. rotatable bonds 2 Num. H-bond acceptors 3 Num. H-bond donors 2 Molar Refractivity 45.13 TPSA 57.53 Å ²	Log Po/w (iLOGP) 0.95 Log Po/w (XLOGP3) 1.46 Log Po/w (WLOGP) 1.38 Log Po/w (MLOGP) 1.28 Log Po/w (SILICOS-IT) 1.22 Consensus Log Po/w 1.26	Log S (ESOL) -2.02 Solubility 1.58e+00 mg/ml ; 9.65e-03 mol/l Class Soluble Log S (Ali) -2.27 Solubility 8.73e-01 mg/ml ; 5.32e-03 mol/l Class Soluble Log S (SILICOS-IT) -1.28 Solubility 8.67e+00 mg/ml ; 5.28e-02 mol/l Class Soluble
Caffeic Acid	GI absorption High	PAINS 1 alert:	Lipinski Yes; 0 violation	Formula C ₉ H ₈ O ₄	Log Po/w (iLOGP) 0.97	Log S (ESOL) -1.89

	BBB permeant	No	catechol_A	Ghose	Yes	Molecular weight	180.16 g/mol	Log Po/w (XLOGP3)	1.15	Solubility	2.32e+00 mg/ml ; 1.29e-02 mol/l
	P-gp substrate	No	Brenk 2 alerts: catechol, michael_acceptor_1	Veber	Yes	Num. heavy atoms	13	Log Po/w (WLOGP)	1.09	Class	Very soluble
	CYP1A2 inhibitor	No		Egan	Yes	Num. arom. heavy atoms	6	Log Po/w (MLOGP)	0.70	Log S (Ali)	-2.38
	CYP2C19 inhibitor	No	Lead likeness No; 1 violation: MW<250	Muegge	No; 1 violation: MW<200	Fraction Csp3	0.00	Log Po/w (SILICOS-IT)	0.75	Solubility	7.55e-01 mg/ml ; 4.19e-03 mol/l
	CYP2C9 inhibitor	No				Num. rotatable bonds	2	Consensus Log Po/w	0.93	Class	Soluble
	CYP2D6 inhibitor	No	Synthetic accessibility 1.81	Bioavailability Score	0.56	Num. H-bond acceptors	4			Log S (SILICOS-IT)	-0.71
	CYP3A4 inhibitor	No				Num. H-bond donors	3			Solubility	3.51e+01 mg/ml ; 1.95e-01 mol/l
	Log Kp (skin permeation)	-6.58 cm/s				Molar Refractivity	47.16			Class	Soluble
						TPSA	77.76 Å²				
Catechin	GI absorption	High	PAINS 1 alert: catechol_A	Lipinski	Yes; 0 violation	Formula	C15H14O6	Log Po/w (iLOGP)	1.47	Log S (ESOL)	-2.22
	BBB permeant	No	Brenk 1 alert: catechol	Ghose	Yes	Molecular weight	290.27 g/mol	Log Po/w (XLOGP3)	0.36	Solubility	1.74e+00 mg/ml ; 5.98e-03 mol/l
	P-gp substrate	Yes	Leadlikeness Yes	Veber	Yes	Num. heavy atoms	21	Log Po/w (WLOGP)	1.22	Class	Soluble
	CYP1A2 inhibitor	No		Egan	Yes	Num. arom. heavy atoms	12	Log Po/w (MLOGP)	0.24	Log S (Ali)	-2.24
	CYP2C19 inhibitor	No	Synthetic accessibility 3.50	Muegge	Yes	Fraction Csp3	0.20	Log Po/w (SILICOS-IT)	0.98	Solubility	1.66e+00 mg/ml ; 5.72e-03 mol/l
	CYP2C9 inhibitor	No		Bioavailability Score	0.55	Num. rotatable bonds	1	Consensus Log Po/w	0.85	Class	Soluble
	CYP2D6 inhibitor	No				Num. H-bond acceptors	6			Log S (SILICOS-IT)	-2.14
	CYP3A4 inhibitor	No				Num. H-bond donors	5			Solubility	2.09e+00 mg/ml ; 7.19e-03 mol/l
	Log Kp (skin permeation)	-7.82 cm/s				Molar Refractivity	74.33			Class	Soluble
					TPSA	110.38 Å²					
Ferulate	GI absorption	High	PAINS 0 alert	Lipinski	Yes; 0 violation	Formula	C10H10O4	Log Po/w (iLOGP)	1.62	Log S (ESOL)	-2.11
	BBB permeant	Yes	Brenk 1 alert: michael_acceptor_1	Ghose	Yes	Molecular weight	194.18 g/mol	Log Po/w (XLOGP3)	1.51	Solubility	1.49e+00 mg/ml ; 7.68e-03 mol/l
	P-gp substrate	No	Leadlikeness No; 1 violation: MW<250	Veber	Yes	Num. heavy atoms	14	Log Po/w (WLOGP)	1.39	Class	Soluble
				Egan	Yes	Num. arom. heavy atoms		Log Po/w (MLOGP)	1.00		

	CYP1A2 inhibitor	No	Synthetic accessibility	Muegge	No; 1 violation:	6	Log Po/w (SILICOS-IT)	Log S (Ali) -2.52
	CYP2C19 inhibitor	No	1.93	MW<200		Fraction Csp3	SILICOS-IT: Hybrid	Solubility
	CYP2C9 inhibitor	No		Bioavailability Score	0.85	Num. rotatable bonds	fragmental/topological	mol/l
	CYP2D6 inhibitor	No				Num. H-bond acceptors	method calculated by	Class
	CYP3A4 inhibitor	No				4	FILTER-IT program, version	Soluble
	Log Kp (skin permeation)	-6.41 cm/s				Num. H-bond donors	1.0.2, courtesy of SILICOS-IT, http://www.silicos-it.com	Log S (SILICOS-IT)
						Molar Refractivity	1.26	-1.42
						TPSA	Consensus Log Po/w	Solubility
							1.36	mol/l
								Class
								Soluble
Kaempferol	GI absorption	High	PAINS	0 alert	Lipinski	Yes; 0 violation	Formula	Log S (ESOL)
	BBB permeant	No	Brenk	0 alert	Ghose	Yes	Molecular weight	-3.31
	P-gp substrate	No	Leadlikeness	Yes	Veber	Yes	g/mol	Solubility
	CYP1A2 inhibitor	Yes	Synthetic accessibility	3.14	Egan	Yes	Num. heavy atoms	mol/l
	CYP2C19 inhibitor	No			Muegge	Yes	16	Class
	CYP2C9 inhibitor	No			Bioavailability Score	0.55	Num. arom. heavy atoms	Soluble
	CYP2D6 inhibitor	Yes					16	Log S (Ali)
	CYP3A4 inhibitor	Yes					Fraction Csp3	-3.86
	Log Kp (skin permeation)	-6.70 cm/s					0.00	Solubility
							Num. rotatable bonds	mol/l
							1	Class
							Num. H-bond acceptors	Soluble
							6	Log S (SILICOS-IT)
							Num. H-bond donors	-3.82
							4	Solubility
							Molar Refractivity	mol/l
							76.01	Class
							TPSA	Soluble
							111.13 Å²	
Quercetin	GI absorption	Low	PAINS	1 alert: catechol_A	Lipinski	No; 2 violations: NorO>10, NHorOH>5	Formula	Log S (ESOL)
	BBB permeant	No	Brenk	1 alert: catechol	Ghose	Yes	Molecular weight	-3.33
	P-gp substrate	No	Leadlikeness	No;	Veber	No; 1 violation: TPSA>140	g/mol	Solubility
	CYP1A2 inhibitor	No	1 violation: MW>350		Egan	No; 1 violation: TPSA>131.6	448.38	mol/l
	CYP2C19 inhibitor	No	Synthetic accessibility	5.28	Muegge	No; 3 violations:	32	Class
	CYP2C9 inhibitor	No					Num. arom. heavy atoms	Soluble
							16	Log S (Ali)
							Fraction Csp3	-4.44
							0.29	Solubility
								mol/l

	CYP2D6 inhibitor No CYP3A4 inhibitor No Log Kp (skin permeation) -8.42 cm/s		TPSA>150, H-acc>10, H-don>5 Bioavailability Score 0.17	Num. rotatable bonds 3 Num. H-bond acceptors 11 Num. H-bond donors 7 Molar Refractivity 109.00 TPSA 190.28 Å ²	Consensus Log Po/w 0.22	Class Moderately soluble Log S (SILICOS-IT) -2.08 Solubility 3.69e+00 mg/ml ; 8.22e-03 mol/l Class Soluble
Chlorogenic Acid	GI absorption Low BBB permeant No P-gp substrate No CYP1A2 inhibitor No CYP2C19 inhibitor No CYP2C9 inhibitor No CYP2D6 inhibitor No CYP3A4 inhibitor No Log Kp (skin permeation) -8.62 cm/s	PAINS 1 alert: catechol_A Brenk 3 alerts: catechol, michael_acceptor_1, more_than_2_esters Leadlikeness No; 1 violation: MW>350 Synthetic accessibility 4.27	Lipinski Yes; 0 violation Ghose No; 1 violation: WLOGP<-0.4 Veber No; 1 violation: TPSA>140 Egan No; 1 violation: TPSA>131.6 Muegge No; 1 violation: TPSA>150 Bioavailability Score 0.55	Formula C17H20O9 Molecular weight 368.34 g/mol Num. heavy atoms 26 Num. arom. heavy atoms 6 Fraction Csp3 0.41 Num. rotatable bonds 6 Num. H-bond acceptors 9 Num. H-bond donors 5 Molar Refractivity 87.82 TPSA 153.75 Å ²	Log Po/w (iLOGP) 1.64 Log Po/w (XLOGP3) -0.10 Log Po/w (WLOGP) -0.67 Log Po/w (MLOGP) -0.81 Log Po/w (SILICOS-IT) -0.07 Consensus Log Po/w -0.00	Log S (ESOL) -1.84 Solubility 5.38e+00 mg/ml ; 1.46e-02 mol/l Class Very soluble Log S (Ali) -2.68 Solubility 7.76e-01 mg/ml ; 2.11e-03 mol/l Class Soluble Log S (SILICOS-IT) -0.29 Solubility 1.89e+02 mg/ml ; 5.13e-01 mol/l Class Soluble
4-Hydroxybenzoic acid	GI absorption High BBB permeant Yes P-gp substrate No CYP1A2 inhibitor No CYP2C19 inhibitor No CYP2C9 inhibitor No CYP2D6 inhibitor No CYP3A4 inhibitor No	PAINS 0 alert Brenk 0 alert Leadlikeness No; 1 violation: MW<250 Synthetic accessibility 1.00	Lipinski Yes; 0 violation Ghose No; 3 violations: MW<160, MR<40, #atoms<20 Veber Yes Egan Yes Muegge No; 1 violation: MW<200 Bioavailability Score 0.85	Formula C7H6O3 Molecular weight 138.12 g/mol Num. heavy atoms 10 Num. arom. heavy atoms 6 Fraction Csp3 0.00 Num. rotatable bonds 1 Num. H-bond acceptors	Log Po/w (iLOGP) 0.85 Log Po/w (XLOGP3) 1.58 Log Po/w (WLOGP) 1.09 Log Po/w (MLOGP) 0.99 Log Po/w (SILICOS-IT) 0.74 Consensus Log Po/w 1.05	Log S (ESOL) -2.07 Solubility 1.18e+00 mg/ml ; 8.52e-03 mol/l Class Soluble Log S (Ali) -2.40 Solubility 5.51e-01 mg/ml ; 3.99e-03 mol/l Class Soluble

	Log Kp (skin permeation) -6.02 cm/s				3 Num. H-bond donors 2 Molar Refractivity 35.42 TPSA 57.53 Å²		Log S (SILICOS-IT) -1.17 Solubility 9.40e+00 mg/ml ; 6.81e-02 mol/l Class Soluble
o-Coumaric acid	GI absorption	High	PAINS 0 alert	Lipinski Yes; 0 violation	Formula C9H8O3	Log Po/w (iLOGP) 1.09	Log S (ESOL) -2.37
	BBB permeant	Yes	Brenk 1 alert: michael_acceptor_1	Ghose Yes	Molecular weight 164.16 g/mol	Log Po/w (XLOGP3) 2.03	Solubility 6.93e-01 mg/ml ; 4.22e-03 mol/l
	P-gp substrate	No	Leadlikeness No; 1 violation: MW<250	Veber Yes	Num. heavy atoms 12	Log Po/w (WLOGP) 1.38	Class Soluble
	CYP1A2 inhibitor	No	Synthetic accessibility 1.85	Egan Yes	Num. arom. heavy atoms 6	Log Po/w (MLOGP) 1.28	Log S (Ali) -2.87
	CYP2C19 inhibitor	No		Muegge No; 1 violation: MW<200	Fraction Csp3 0.00	Log Po/w (SILICOS-IT) 1.22	Solubility 2.24e-01 mg/ml ; 1.36e-03 mol/l
	CYP2C9 inhibitor	No		Bioavailability Score 0.85	Num. rotatable bonds 2	Consensus Log Po/w 1.40	Class Soluble
	CYP2D6 inhibitor	No			Num. H-bond acceptors 3		Log S (SILICOS-IT) -1.28
	CYP3A4 inhibitor	No			Num. H-bond donors 2		Solubility 8.67e+00 mg/ml ; 5.28e-02 mol/l
	Log Kp (skin permeation) -5.86 cm/s				Molar Refractivity 45.13		Class Soluble
Vanillic acid	GI absorption	High	PAINS 0 alert	Lipinski Yes; 0 violation	Formula C8H8O4	Log Po/w (iLOGP) 1.40	Log S (ESOL) -2.02
	BBB permeant	No	Brenk 0 alert	Ghose Yes	Molecular weight 168.15 g/mol	Log Po/w (XLOGP3) 1.43	Solubility 1.60e+00 mg/ml ; 9.52e-03 mol/l
	P-gp substrate	No	Leadlikeness No; 1 violation: MW<250	Veber Yes	Num. heavy atoms 12	Log Po/w (WLOGP) 1.10	Class Soluble
	CYP1A2 inhibitor	No	Synthetic accessibility 1.42	Egan Yes	Num. arom. heavy atoms 6	Log Po/w (MLOGP) 0.74	Log S (Ali) -2.44
	CYP2C19 inhibitor	No		Muegge No; 1 violation: MW<200	Fraction Csp3 0.12	Log Po/w (SILICOS-IT) 0.73	Solubility 6.15e-01 mg/ml ; 3.66e-03 mol/l
	CYP2C9 inhibitor	No		Bioavailability Score 0.85	Num. rotatable bonds 2	Consensus Log Po/w 1.08	Class Soluble
	CYP2D6 inhibitor	No			Num. H-bond acceptors 4		Log S (SILICOS-IT) -1.32
	CYP3A4 inhibitor	No			Num. H-bond donors 2		Solubility 8.10e+00 mg/ml ; 4.82e-02 mol/l
	Log Kp (skin permeation) -6.31 cm/s				Molar Refractivity 41.92		Class Soluble

				TPSA 66.76 Å ²		
Syringic acid	GI absorption High	PAINS 0 alert	Lipinski Yes; 0 violation	Formula C ₉ H ₁₀ O ₅	Log Po/w (iLOGP) 1.54	Log S (ESOL) -1.84
	BBB permeant No	Brenk 0 alert	Ghose Yes	Molecular weight 198.17 g/mol	Log Po/w (XLOGP3) 1.04	Solubility 2.84e+00 mg/ml ; 1.44e-02 mol/l
	P-gp substrate No	Leadlikeness No; 1 violation: MW<250	Veber Yes	Num. heavy atoms 14	Log Po/w (WLOGP) 1.11	Class Very soluble
	CYP1A2 inhibitor No	Synthetic accessibility 1.70	Egan Yes	Num. arom. heavy atoms 6	Log Po/w (MLOGP) 0.49	Log S (Ali) -2.23
	CYP2C19 inhibitor No		Muegge No; 1 violation: MW<200	Fraction Csp3 0.22	Log Po/w (SILICOS-IT) 0.77	Solubility 1.18e+00 mg/ml ; 5.94e-03 mol/l
	CYP2C9 inhibitor No		Bioavailability Score 0.56	Num. rotatable bonds 3	Consensus Log Po/w 0.99	Class Soluble
	CYP2D6 inhibitor No			Num. H-bond acceptors 5		Log S (SILICOS-IT) -1.46
	CYP3A4 inhibitor No			Num. H-bond donors 2		Solubility 6.93e+00 mg/ml ; 3.50e-02 mol/l
	Log Kp (skin permeation) -6.77 cm/s			Molar Refractivity 48.41		Class Soluble
				TPSA 75.99 Å ²		