

Transcriptomic, mutational and structural bioinformatics approaches to explore the therapeutic role of FAP in predominant cancer types

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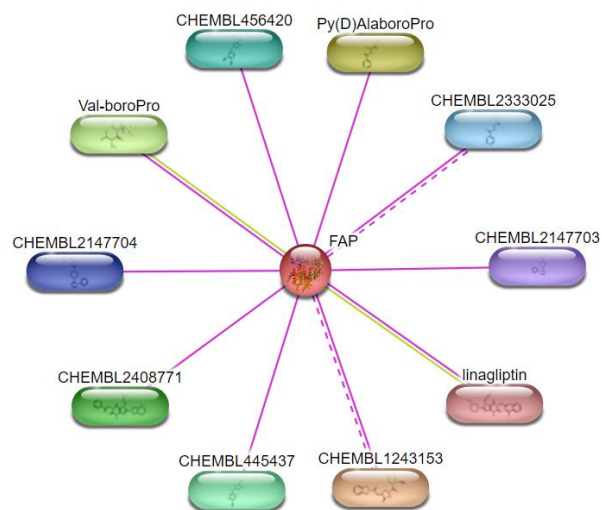
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Online Resource 6 Drug-protein interaction network constructed using STITCH database



Node	Node 2	Drug ID	Experimentally determined interaction	Prediction	Combined score
FAP	CHEMBL1243153	CID046917048	0.898	0.401	0.937
	Py(D)AlaboroPro	CID071655266	0.867	0	0.867
	Val-boroPro	CID006918572	0.827	0	0.842
	CHEMBL2408771	CID066954220	0.838	0	0.838
	CHEMBL445437	CID044590651	0.832	0	0.832
	CHEMBL2333025	CID071655265	0.812	0.105	0.827
	CHEMBL456420	CID044590650	0.814	0	0.814
	CHEMBL2147704	CID067507176	0.81	0	0.81
	CHEMBL2147703	CID067507912	0.807	0	0.807
	Linagliptin	CID010096344	0.708	0	0.804