

**Table 1.** Cross validation on the 1-step reaction sequence-likeness (enzymatic-reaction-likeness).

Chemical fingerprints/descriptors	Diff-common L1SVM		Diff-only L1SVM		Baseline		Random	
	AUC	AUPR	AUC	AUPR	AUC	AUPR	AUC	AUPR
CDK extended	0.6917	0.0603	0.6742	0.0468	0.6161	0.0289	0.5000	0.0199
CDK graph-only	0.7161	0.0602	0.6960	0.0432	0.6255	0.0228	0.5000	0.0199
CDK hybridization	0.7003	0.0489	0.6761	0.0403	0.6023	0.0272	0.5000	0.0199
E-state	0.6088	0.0371	0.5567	0.0297	0.5085	0.0169	0.5000	0.0199
KlekotaRoth	0.6031	0.0347	0.5683	0.0304	0.5203	0.0042	0.5000	0.0199
MACCS	0.6837	0.0489	0.6582	0.0342	0.5914	0.0189	0.5000	0.0199
PubChem	0.7170	0.0531	0.7026	0.0422	0.6752	0.0307	0.5000	0.0199
NS-descriptor	0.8858	0.2134	0.8429	0.0968	0.6566	0.0446	0.5000	0.0199
KCF-S descriptor	0.9659	0.3943	0.9610	0.2801	0.6945	0.0755	0.5000	0.0199

**Table 2.** Cross validation on the 2-step reaction sequence-likeness prediction (with 1 intermediate compound).

Chemical fingerprints/descriptors	Diff-common L1SVM		Diff-only L1SVM		Baseline		Random	
	AUC	AUPR	AUC	AUPR	AUC	AUPR	AUC	AUPR
CDK extended	0.7747	0.1730	0.7178	0.1352	0.4815	0.0576	0.5000	0.0665
CDK graph-only	0.7429	0.1711	0.7510	0.1626	0.4969	0.0567	0.5000	0.0665
CDK hybridization	0.7871	0.1879	0.6971	0.1271	0.4631	0.0554	0.5000	0.0665
E-state	0.6318	0.0900	0.5389	0.0563	0.3941	0.0407	0.5000	0.0665
KlekotaRoth	0.6462	0.1080	0.5309	0.0807	0.4949	0.0181	0.5000	0.0665
MACCS	0.7474	0.1418	0.6634	0.1152	0.4465	0.0502	0.5000	0.0665
PubChem	0.7674	0.1589	0.7270	0.1357	0.5732	0.0710	0.5000	0.0665
NS-descriptor	0.8898	0.2937	0.8673	0.2651	0.6187	0.0937	0.5000	0.0665
KCF-S descriptor	0.9411	0.4493	0.9419	0.4473	0.6621	0.0635	0.5000	0.0665

**Table 3.** Cross validation on the 3-step reaction sequence-likeness prediction (with 2 intermediate compounds).

Chemical fingerprints/descriptors	Diff-common L1SVM		Diff-only L1SVM		Baseline		Random	
	AUC	AUPR	AUC	AUPR	AUC	AUPR	AUC	AUPR
CDK extended	0.8103	0.1436	0.7542	0.0959	0.5474	0.0368	0.5000	0.0367
CDK graph-only	0.8363	0.1508	0.7785	0.0949	0.5621	0.0369	0.5000	0.0367
CDK hybridization	0.8293	0.1558	0.7631	0.0973	0.5226	0.0345	0.5000	0.0367
E-state	0.6192	0.0515	0.5450	0.0495	0.4125	0.0225	0.5000	0.0367
KlekotaRoth	0.6395	0.0605	0.5463	0.0539	0.5165	0.0116	0.5000	0.0367
MACCS	0.7608	0.0986	0.6770	0.0713	0.4959	0.0309	0.5000	0.0367
PubChem	0.8097	0.1239	0.7656	0.0910	0.6365	0.0489	0.5000	0.0367
NS-descriptor	0.9284	0.2638	0.9028	0.1989	0.7069	0.0807	0.5000	0.0367
KCF-S descriptor	0.9624	0.4232	0.9585	0.4094	0.6621	0.0635	0.5000	0.0367

**Table 4.** Cross validation on the 4-step reaction sequence-likeness prediction (with 3 intermediate compounds).

Chemical fingerprints/descriptors	Diff-common L1SVM		Diff-only L1SVM		Baseline		Random	
	AUC	AUPR	AUC	AUPR	AUC	AUPR	AUC	AUPR
CDK extended	0.8577	0.1062	0.7867	0.0649	0.5863	0.0172	0.5000	0.0156
CDK graph-only	0.8611	0.0988	0.7957	0.0482	0.5961	0.0175	0.5000	0.0156
CDK hybridization	0.8581	0.1083	0.7869	0.0583	0.5490	0.0157	0.5000	0.0156
E-state	0.6053	0.0235	0.5452	0.0230	0.4336	0.0097	0.5000	0.0156
KlekotaRoth	0.6270	0.0273	0.5519	0.0248	0.5104	0.0057	0.5000	0.0156
MACCS	0.7663	0.0582	0.6898	0.0351	0.5187	0.0141	0.5000	0.0156
PubChem	0.8536	0.0818	0.7962	0.0481	0.6590	0.0234	0.5000	0.0156
NS-descriptor	0.9535	0.2058	0.9304	0.1341	0.7521	0.0436	0.5000	0.0156
KCF-S descriptor	0.9772	0.3283	0.9837	0.3202	0.7039	0.0315	0.5000	0.0156