

Supporting Information for Machine Learning-Augmented Docking

1. CYP inhibition prediction

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Code Workflow Procedure

1. Cleaned and combined test and train Pei sets with CYP_clean_files.ipnyb. Sets combined and then clustered to create new train and test sets
2. Dock each ligand 5 times to its respective isoform using FITTED. Docked data can be found here: (to be inserted)
3. Create analogue sets using FITTED. Create max train-test similarity using CYP_TC_DataSets.py
4. Run RF with Feature Importances using max train-test similarity of 0.8 using ML_over_Tanimoto.py which calls CYP_inhibition_functions.py and Do_ML2.py
5. Using these selected features run all ML models on all datasets using ML_over_Tanimoto.py which calls CYP_inhibition_functions.py and Do_ML2.py
6. Use CYP_evaluate_and_ensemble.py which calls CYP_evaluate_and_ensemble_functions.py to make ensembles and evaluate and graph model performance

1 List of input Ligand Features

- Energy
- van der Waals
- Electrostatics
- Hydrogen Bonds
- Hydrogen Bond-like Metal interaction
- Metal Coordination
- Water van der Waals
- Water Electrostatics
- Water Hydrogen Bonds
- Number of Water Molecules
- Number of Rotatable Bonds
- Rotatable Bonds Entropy
- Non-polar Solvation Energy
- Polar Ligand Solvation Energy
- Polar Protein Solvation Energy
- RMSD
- RMSD (MBG)
- Strain Energy
- RankScore
- MatchScore
- Molecular Weight
- Net Charge
- Hydrogen Bond Donors
- Hydrogen Bond Acceptors
- Rotatable Bonds
- Rings
- Ionizable Centers
- Oxygens
- Nitrogens
- Sulfurs
- Heteroatoms
- Heavy Atoms
- Michael Acceptors
- tPSA
- logP
- Fsp3
- Stereogenic Centers Ratio
- logS
- Stereogenic Centers
- Aromatic Rings
- Aromatic Proportion
- Halogens
- McGowan Molecular Volume
- Molecular van der Waals Surface Area
- Polar SASA
- non Polar SASA
- Molecular Density
- Molecular Polarity
- Blood-Brain Barrier
- Molecular Softness
- Molecular Electronegativity
- Negatively Charged Oxygens
- Positively Charged Nitrogen
- Molecular Hardness
- Molecular Polarizability
- 3D-Wiener Index
- Geometric Radius
- Geometric Diameter
- Geometrical Shape Coefficient
- Span
- Radius of Gyration
- Ovality Index
- Globularity Factor

- Dipole moment
- Badapple pscore
- Quantitative Estimate of Druglikeness
- aromatic
- carboxylic acid
- carboxylate
- ester aromatic
- ester aliphatic
- ester conjugated
- ester
- lactone
- aldehyde aromatic
- aldehyde aliphatic
- aldehyde conjugated
- aldehyde
- acyl chloride
- acyl bromide
- amide ter
- amide
- lactame
- anhydride
- ketone
- sulfonamide
- thiol
- thiolate
- carbamate
- primary amine
- secondary amine
- tertiary amine
- quat ammonium
- amine
- aniline
- nitrile
- imine
- oxime
- nitro
- nitroso
- azide
- isocyanate
- hydroxamic acid
- hydroxamate
- alkyl chloride
- alkyl bromide
- alkyl iodide
- aryl chloride
- aryl bromide
- aryl iodide
- boronic acid aromatic
- boronic ester aromatic
- boronic acid aliphatic
- boronic ester aliphatic
- boronic acid vinylic
- boronic ester vinylic
- boronic acid
- sulfonyl chloride
- sulfonyl bromide
- vinyl chloride
- vinyl bromide
- vinyl iodide
- alkene
- ketone O
- formyl
- alcohol
- chloride
- bromide
- fluoride
- iodide
- silicon
- primary amine neutral
- secondary amine neutral
- tertiary amine neutral
- alpha fluoroketone
- alpha chloroketone
- epoxide
- aziridine
- beta-lactam
- primary alcohol
- disulfide
- azo
- isothiocyanate
- hydrazine
- ketene
- peroxide
- silyl fluoride
- silyl bromide
- silyl chloride
- silyl iodide
- thiocarbonyl
- thioester
- betaKeto

Figures and Tables

Table S1: Dataset sizes for 5 CYP P450 isoforms

	1A2	2C19	2C9	2D6	3A4
Before Cleaning Data Size	10072	10769	10050	10851	11436
After Cleaning Data Size	9773	10636	9935	10717	11401
# Actives	3598	4564	2603	1406	3602
# Inactives	6175	6072	7332	9311	7799

Protein structures (PDB IDs)

CYP1A2: 2HI4, CYP2C9: 1R9O, CYP2C19: 4GQS, CYP2D6: 3QM4, CYP3A4: 3NXU

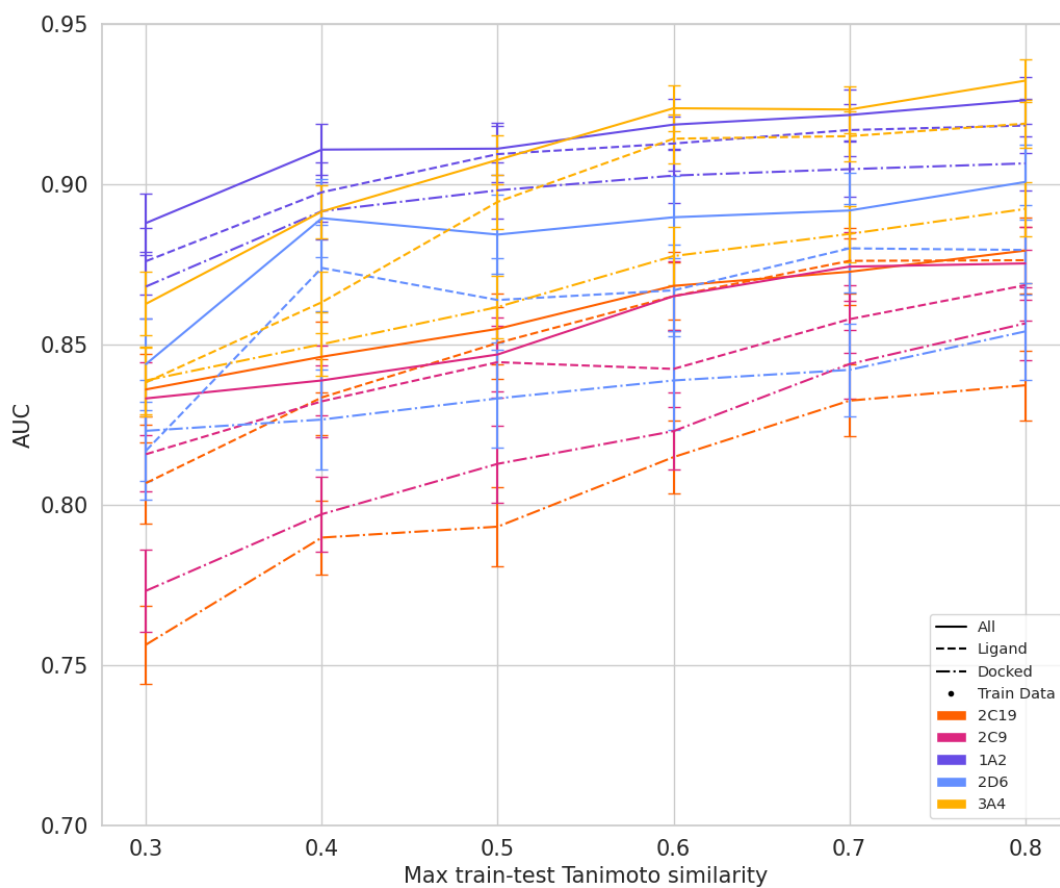


Figure S1: AUC of ML augmented docking using ligand only features, docking features, and all features over max train-test Tanimoto similarity

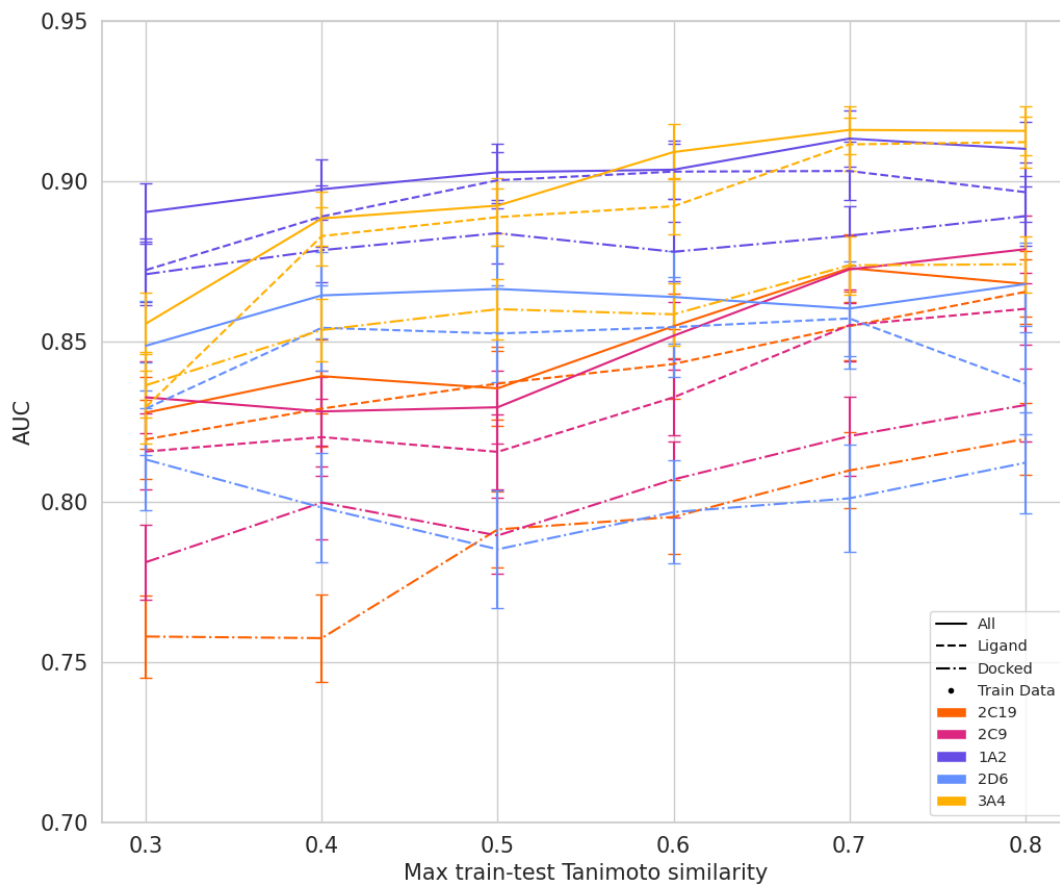


Figure S2: AUC of ML augmented docking using ligand-only features, docking features, and all features over max train-test Tanimoto similarity using same data sizes. Data size capped at data size of 0.3. Data size used; 1A2: 2406, 2C9: 2622, 2C19: 2698, 2D6: 2789, 3A4: 2649. Using all features become more predictive than ligand featurization at low Tanimoto similarity

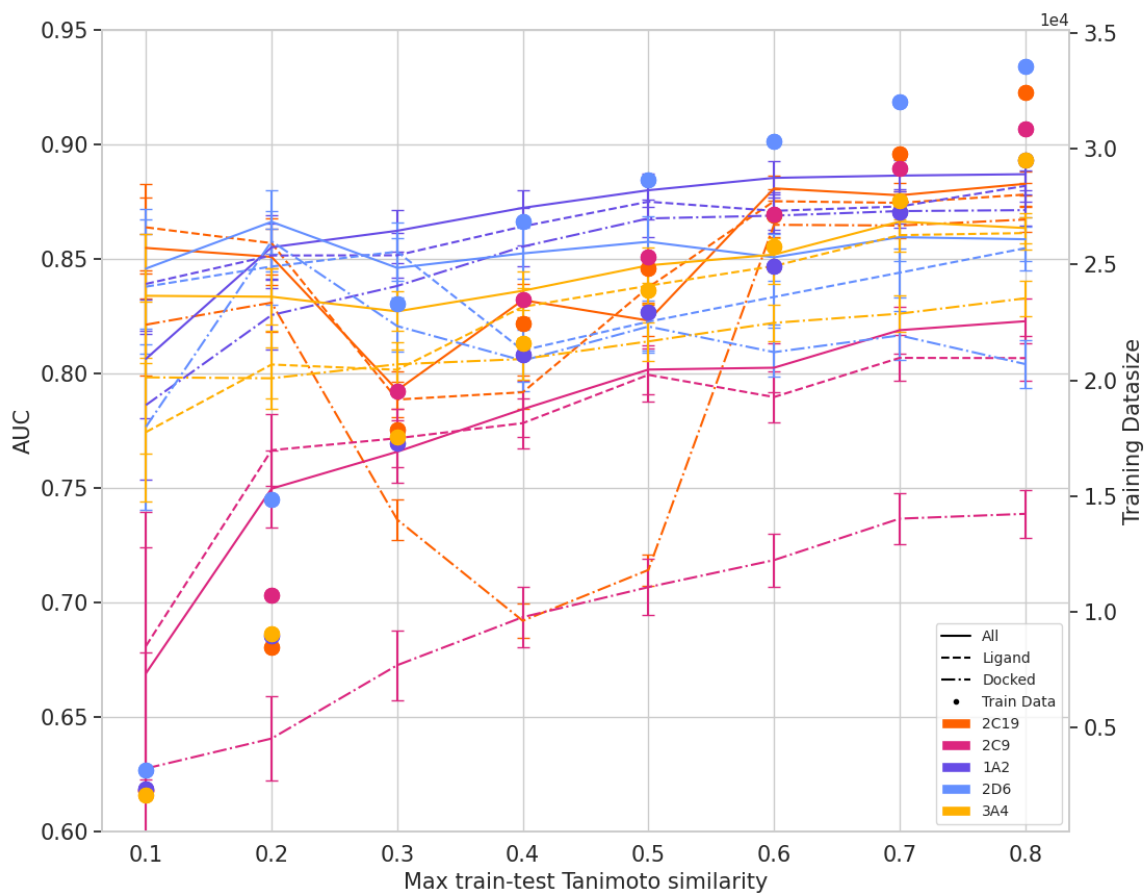


Figure S3: AUC of ML augmented docking using ligand only features, docking features, and all features over max train-test Tanimoto similarity using analog datasets

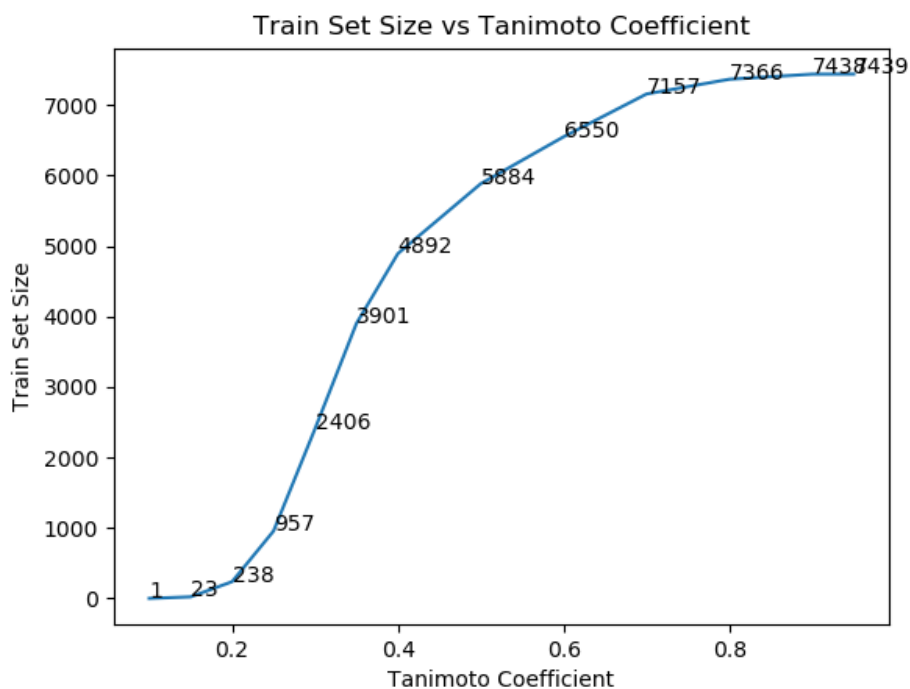


Figure S4: 1A2 CYP inhibition data set size after filtering by max similarity between test and train set using a test set of 100 clusters of 5 molecules.

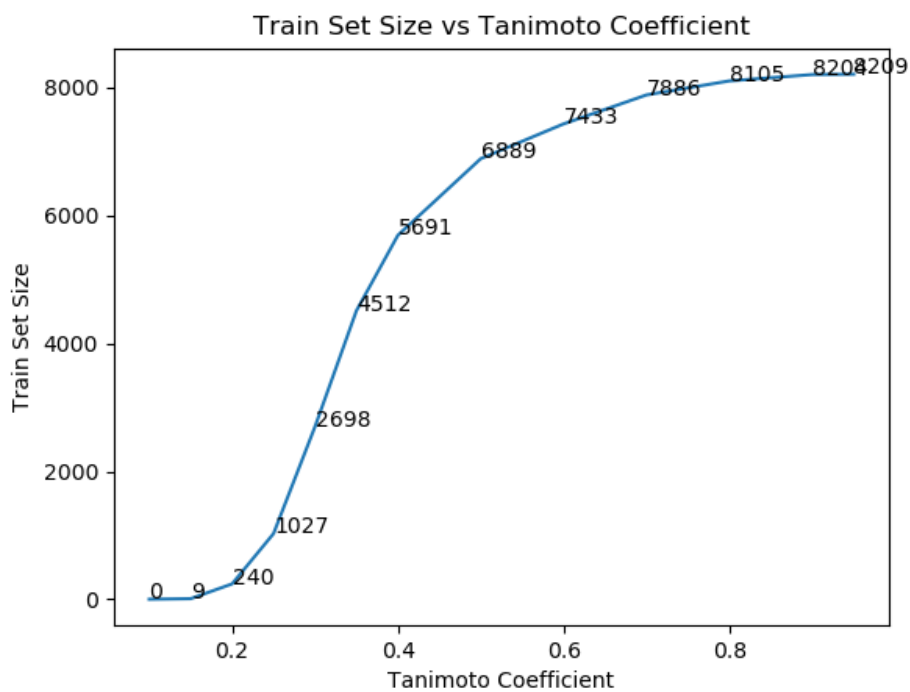


Figure S5: 2C19 CYP inhibition data set size after filtering by max similarity between test and train set using a test set of 100 clusters of 5 molecules.

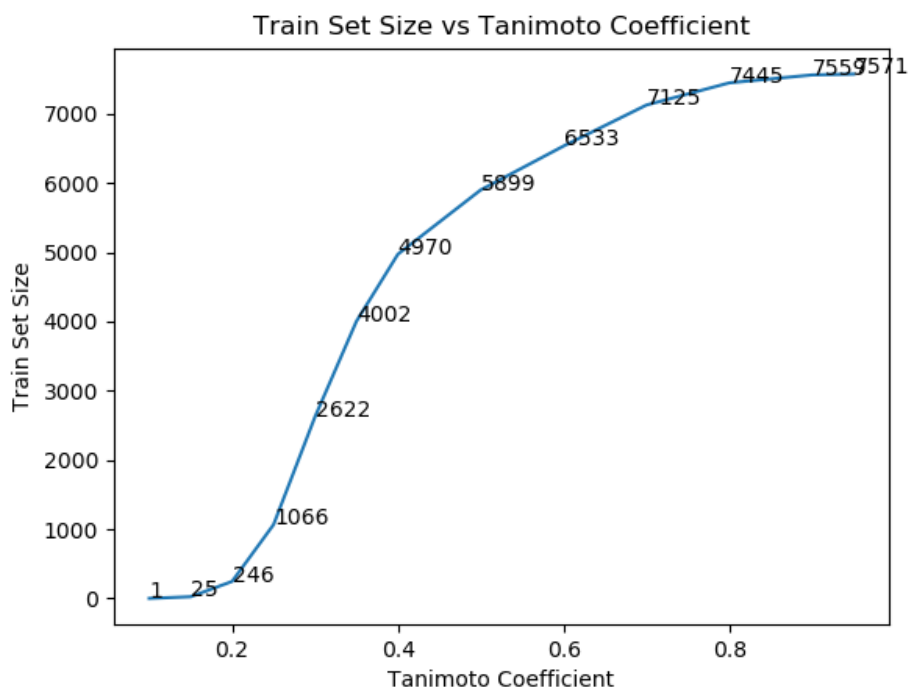


Figure S6: 2C9 CYP inhibition data set size after filtering by max similarity between test and train set using a test set of 100 clusters of 5 molecules.

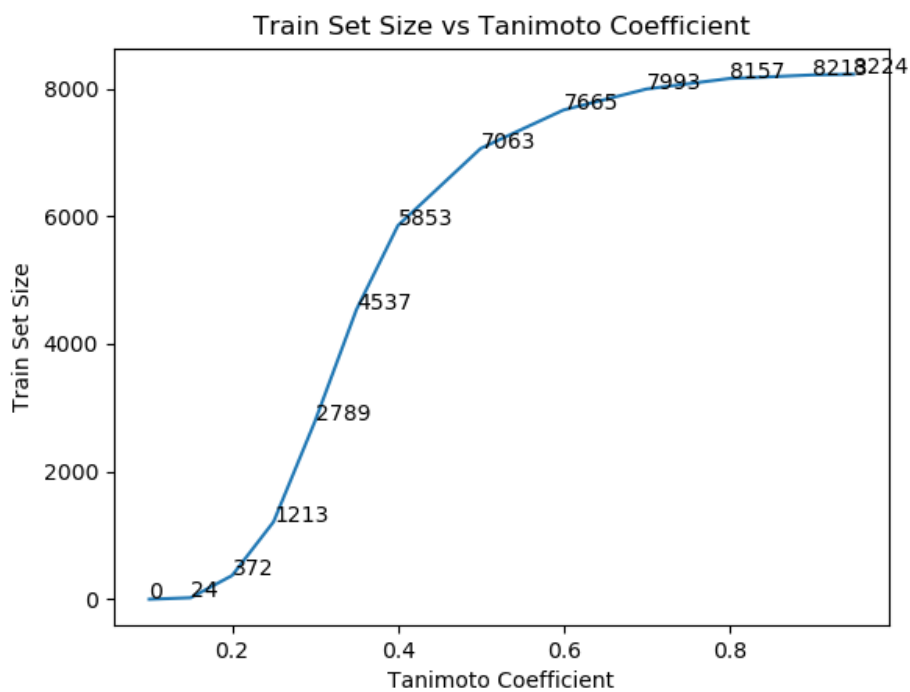


Figure S7: 2D6 CYP inhibition data set size after filtering by max similarity between test and train set using a test set of 100 clusters of 5 molecules.

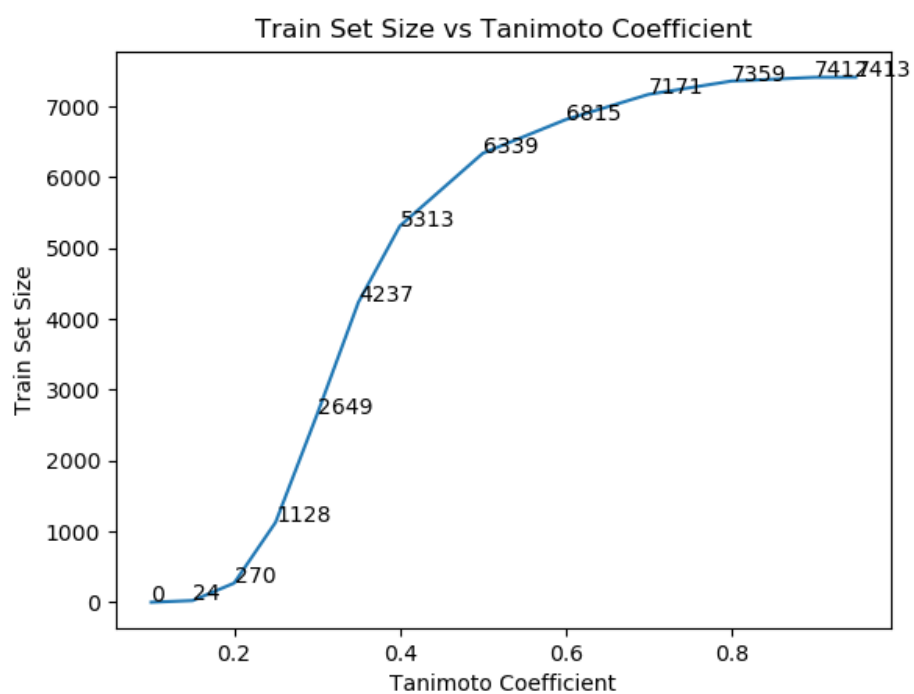


Figure S8: 3A4 CYP inhibition data set size after filtering by max similarity between test and train set using a test set of 100 clusters of 5 molecules.

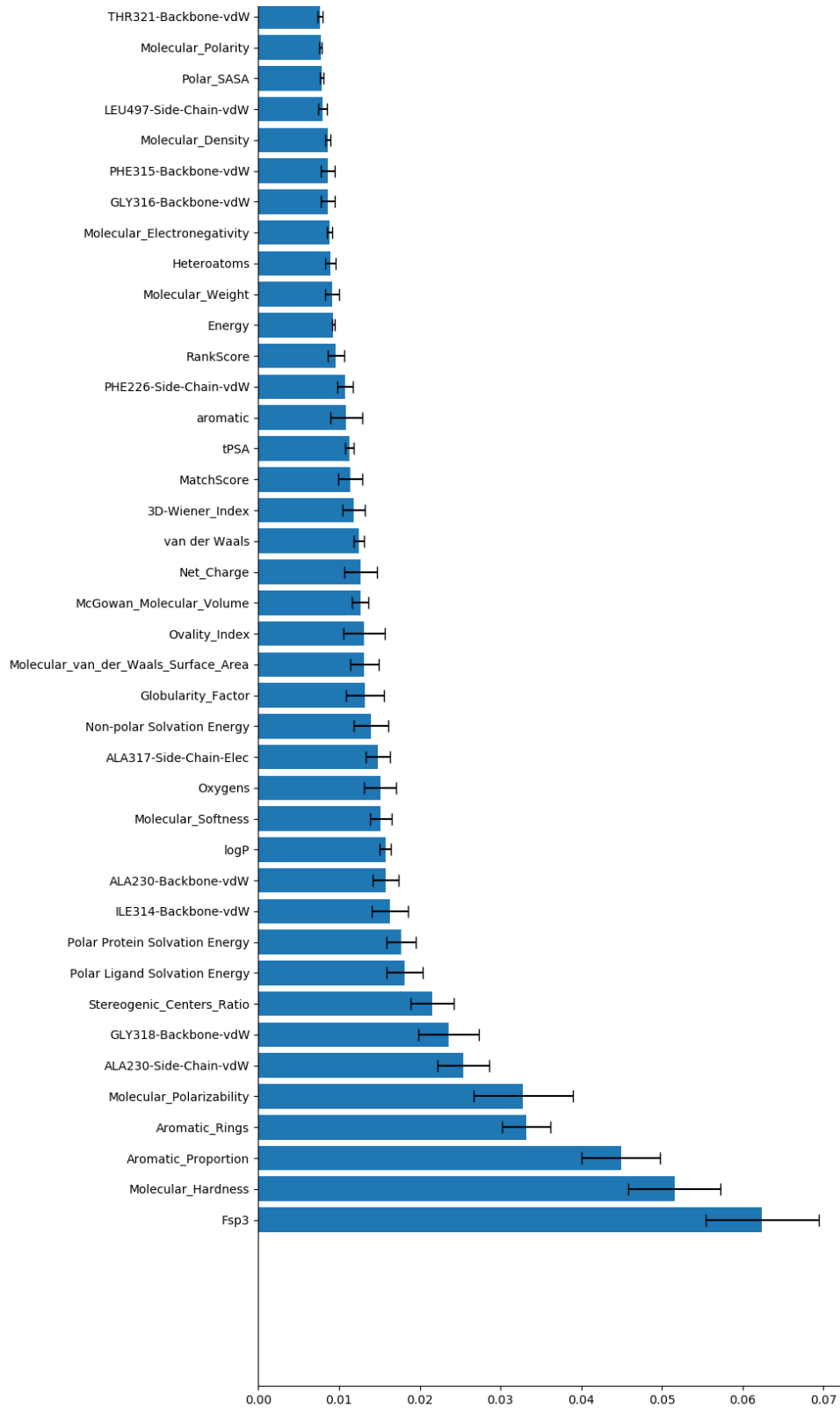


Figure S9: Feature importance of random forest model using selected top 100 features for isoform 1A2.

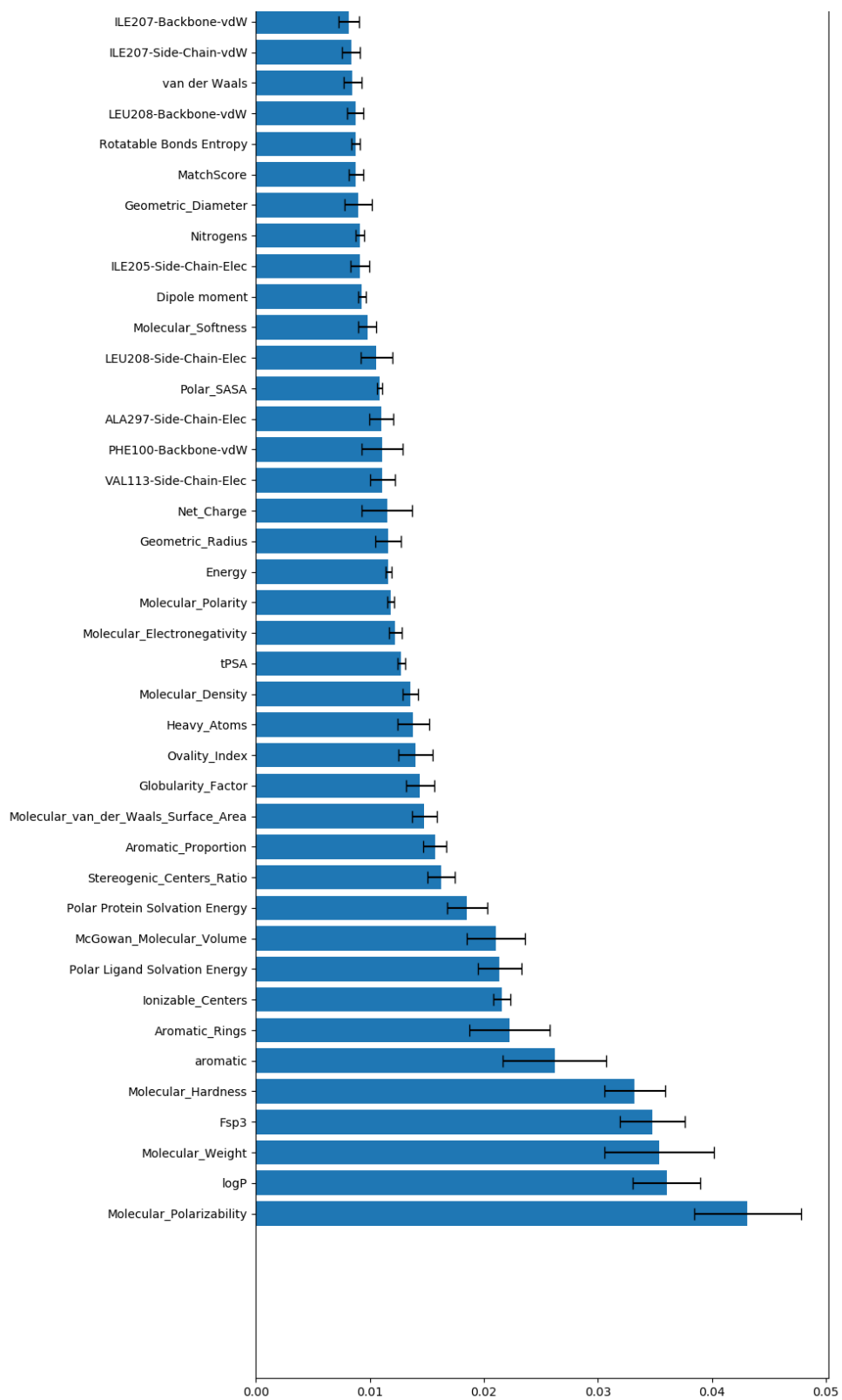


Figure S10: Feature importance of random forest model using selected top 100 features for isoform 2C9.

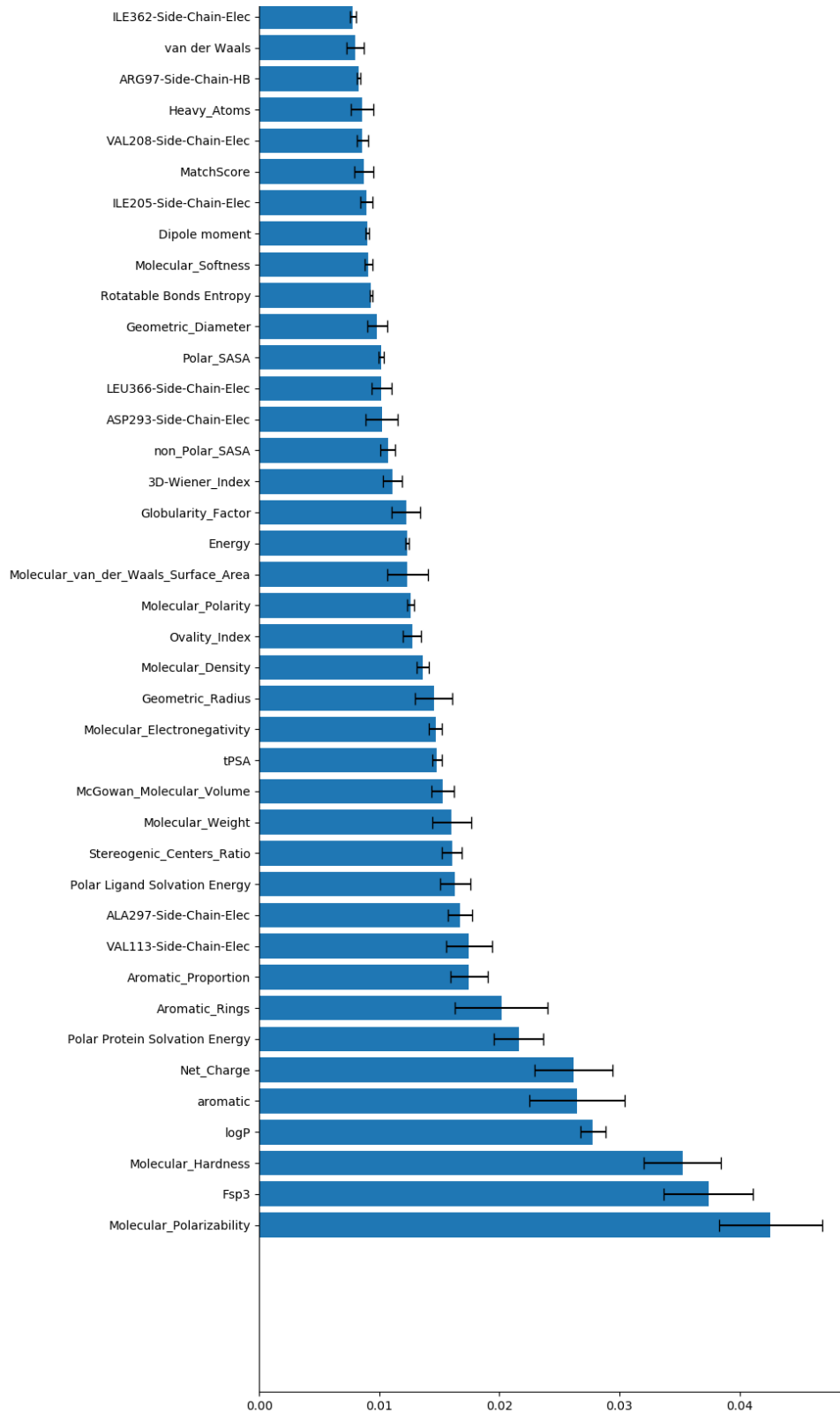


Figure S11: Feature importance of random forest model using selected top 100 features for isoform 2C19.

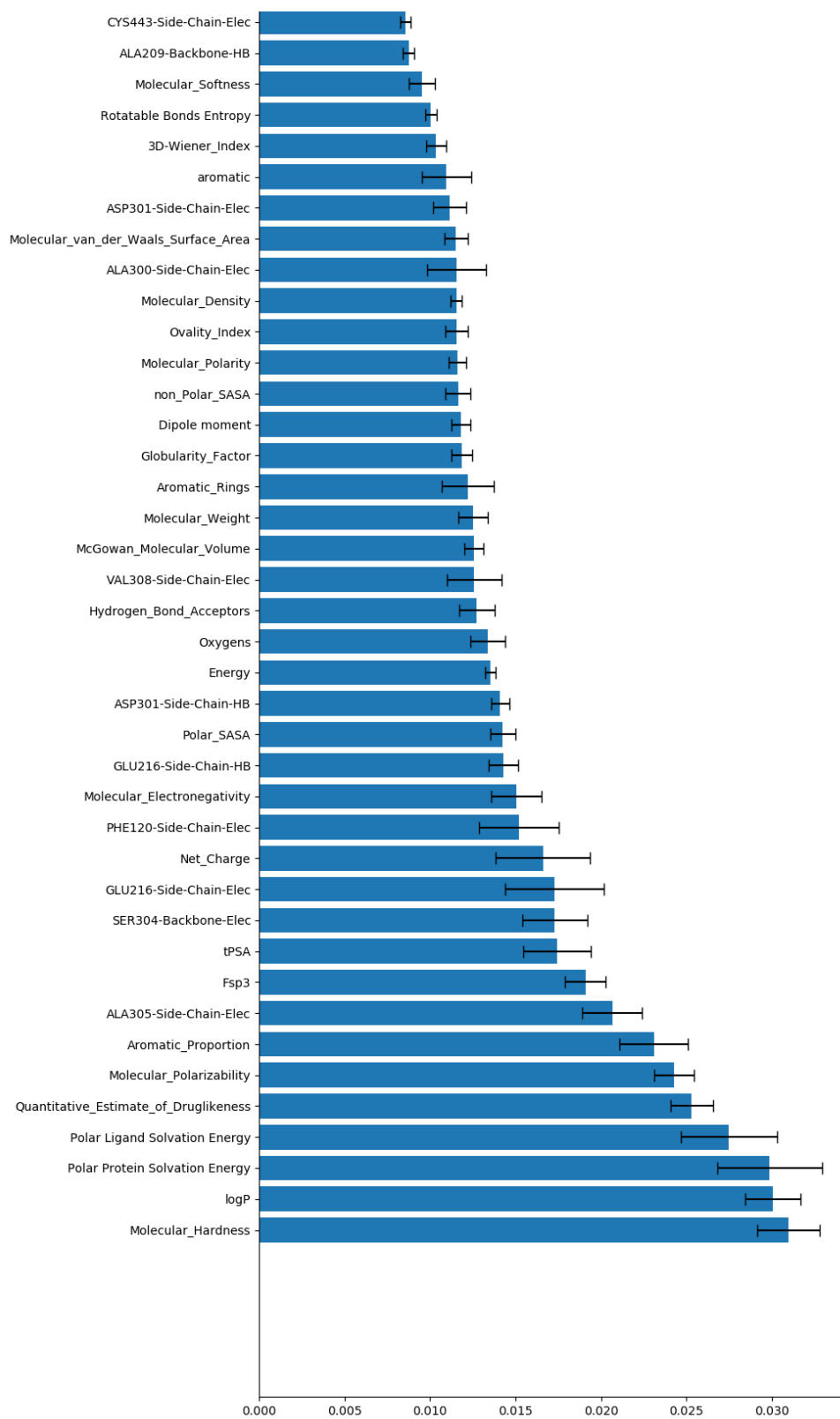


Figure S12: Feature importance of random forest model using selected top 100 features for isoform 2D6.

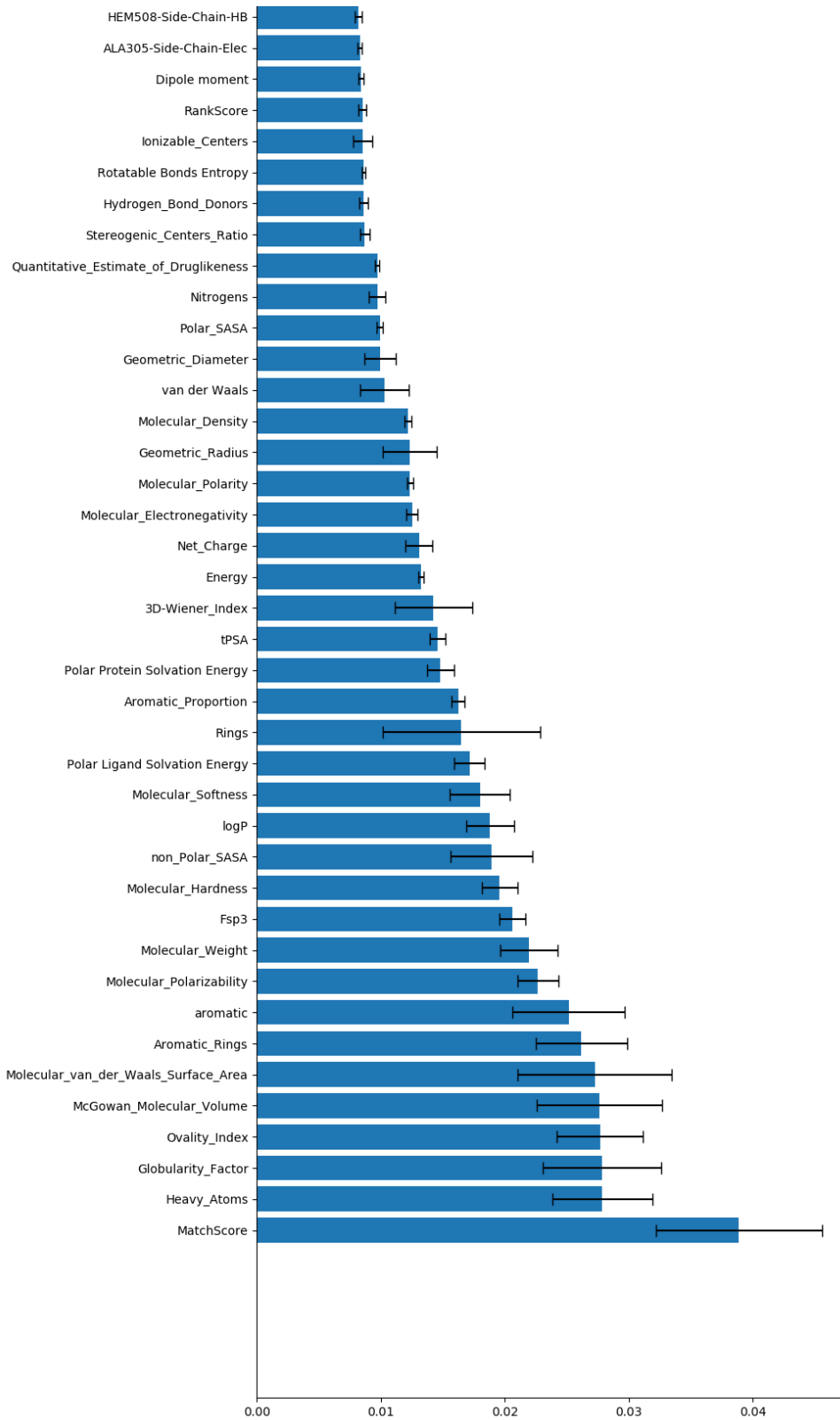


Figure S13: Feature importance of random forest model using selected top 100 features for isoform 3A4.

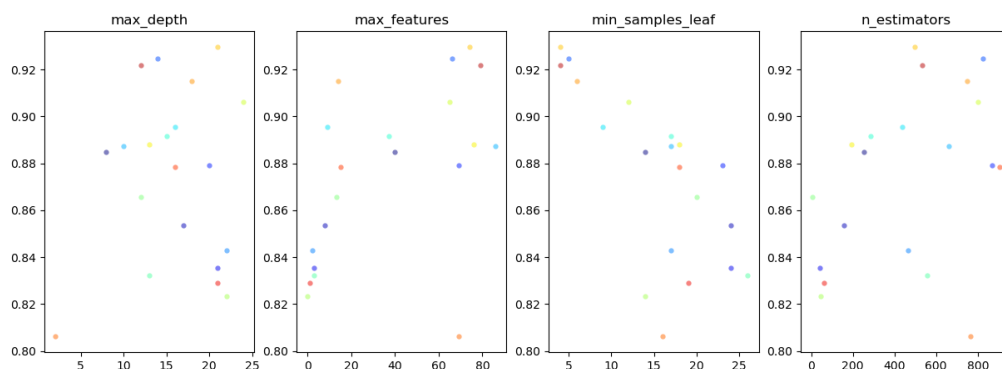


Figure S14: Example of hyperparameter tuning using hyperopt TPE for random forest.

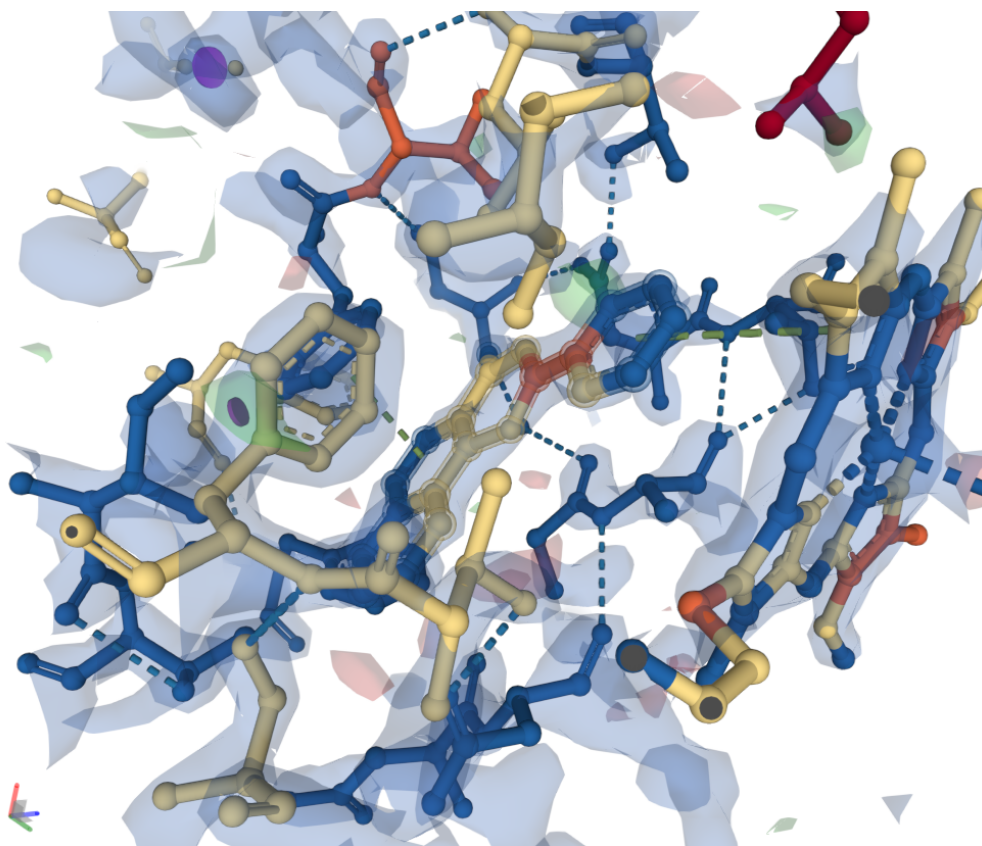


Figure S15: 2HI4: <https://www.rcsb.org/3d-view/2HI4?preset=electronDensityMaps>

Table S2: 1A2 Data

	30	80	70	40	20	50	60
roc_auc_all	0.888	0.926	0.922	0.911	0.849	0.911	0.919
bootstrapped_scores_mean_all	0.888	0.927	0.922	0.911	0.849	0.911	0.918
bootstrapped_scores_std_all	0.009	0.007	0.008	0.008	0.010	0.008	0.008
roc_auc_ligand	0.876	0.918	0.917	0.898	0.830	0.909	0.913
bootstrapped_scores_mean_ligand	0.876	0.918	0.917	0.897	0.831	0.910	0.913
bootstrapped_scores_std_ligand	0.010	0.008	0.008	0.009	0.012	0.009	0.009
roc_auc_docked	0.868	0.907	0.905	0.892	0.783	0.898	0.903
bootstrapped_scores_mean_docked	0.868	0.906	0.905	0.892	0.783	0.898	0.903
bootstrapped_scores_std_docked	0.010	0.008	0.009	0.009	0.012	0.009	0.009
fitted_auc_docked	0.677	0.677	0.677	0.677	0.677	0.677	0.677
auc_improvement_docked	19.120	22.960	22.780	21.470	10.640	22.120	22.580
optimal_threshold_docked	0.505	0.394	0.404	0.374	0.081	0.242	0.343
accuracy_docked	0.786	0.831	0.830	0.813	0.704	0.819	0.821
active_pos_rate_docked	0.783	0.873	0.856	0.876	0.823	0.872	0.875
inactive_neg_rate_docked	0.789	0.798	0.810	0.764	0.609	0.777	0.779
threshold_95_docked	0.152	0.172	0.091	0.182	0.030	0.061	0.162
accuracy_95_docked	0.731	0.769	0.735	0.752	0.639	0.755	0.767
active_acc_95_docked	0.930	0.937	0.945	0.939	0.932	0.939	0.946
inactive_acc_95_docked	0.574	0.637	0.569	0.605	0.408	0.609	0.626
fitted_auc_ligand	0.677	0.677	0.677	0.677	0.677	0.677	0.677
auc_improvement_ligand	19.910	24.140	24.000	22.060	15.350	23.250	23.580
optimal_threshold_ligand	0.192	0.475	0.343	0.455	0.626	0.465	0.495
accuracy_ligand	0.778	0.846	0.839	0.824	0.741	0.840	0.841
active_pos_rate_ligand	0.899	0.864	0.879	0.872	0.836	0.840	0.842
inactive_neg_rate_ligand	0.682	0.832	0.808	0.787	0.667	0.841	0.841
threshold_95_ligand	0.051	0.172	0.071	0.152	0.162	0.061	0.182
accuracy_95_ligand	0.710	0.770	0.761	0.760	0.668	0.757	0.763
active_acc_95_ligand	0.948	0.950	0.948	0.947	0.948	0.945	0.945
inactive_acc_95_ligand	0.522	0.628	0.612	0.613	0.447	0.609	0.619
fitted_auc_all	0.677	0.677	0.677	0.677	0.677	0.677	0.677
auc_improvement_all	21.100	24.930	24.470	23.390	17.170	23.420	24.170
optimal_threshold_all	0.263	0.192	0.303	0.192	0.333	0.485	0.232
accuracy_all	0.796	0.830	0.833	0.798	0.761	0.835	0.816
active_pos_rate_all	0.880	0.915	0.902	0.930	0.842	0.826	0.922
inactive_neg_rate_all	0.729	0.762	0.777	0.694	0.697	0.842	0.732
threshold_95_all	0.101	0.111	0.162	0.182	0.061	0.232	0.172
accuracy_95_all	0.762	0.814	0.798	0.797	0.689	0.799	0.799
active_acc_95_all	0.934	0.943	0.950	0.934	0.964	0.940	0.944
inactive_acc_95_all	0.625	0.713	0.679	0.689	0.471	0.687	0.684

Table S3: 3A4 Data

	60	20	30	80	40	50	70
Roc_auc_all	0.924	0.822	0.863	0.932	0.892	0.908	0.923
Bootstrapped_scores_mean_all	0.924	0.822	0.863	0.932	0.892	0.907	0.923
Bootstrapped_scores_std_all	0.007	0.011	0.010	0.007	0.008	0.008	0.007
Roc_auc_ligand	0.914	0.808	0.838	0.919	0.863	0.895	0.915
Bootstrapped_scores_mean_ligand	0.915	0.808	0.838	0.919	0.863	0.895	0.915
Bootstrapped_scores_std_ligand	0.008	0.012	0.011	0.008	0.010	0.008	0.008
Roc_auc_docked	0.878	0.762	0.839	0.892	0.850	0.862	0.885
Bootstrapped_scores_mean_docked	0.878	0.762	0.839	0.892	0.850	0.862	0.885
Bootstrapped_scores_std_docked	0.009	0.012	0.011	0.008	0.010	0.010	0.009
Fitted_auc_docked	0.742	0.742	0.742	0.742	0.742	0.742	0.742
Auc_improvement_docked	13.610	2.010	9.730	15.080	10.860	12.020	14.300
Optimal_threshold_docked	0.333	0.061	0.384	0.495	0.404	0.273	0.576
Accuracy_docked	0.773	0.634	0.763	0.804	0.776	0.760	0.801
Active_pos_rate_docked	0.862	0.896	0.805	0.790	0.793	0.867	0.737
Inactive_neg_rate_docked	0.714	0.459	0.735	0.813	0.765	0.689	0.844
Threshold_95_docked	0.192	0.040	0.141	0.192	0.152	0.182	0.263
Accuracy_95_docked	0.743	0.622	0.700	0.741	0.713	0.746	0.742
Active_acc_95_docked	0.932	0.925	0.936	0.934	0.939	0.918	0.927
Inactive_acc_95_docked	0.617	0.420	0.542	0.613	0.562	0.631	0.619
Fitted_auc_all	0.742	0.742	0.742	0.742	0.742	0.742	0.742
Auc_improvement_all	18.220	8.090	12.120	19.080	15.000	16.610	18.180
Optimal_threshold_all	0.384	0.404	0.263	0.535	0.222	0.354	0.232
Accuracy_all	0.850	0.730	0.783	0.862	0.817	0.829	0.840
Active_pos_rate_all	0.847	0.755	0.860	0.792	0.857	0.820	0.899
Inactive_neg_rate_all	0.852	0.714	0.732	0.908	0.791	0.836	0.801
Threshold_95_all	0.111	0.051	0.131	0.111	0.081	0.101	0.101
Accuracy_95_all	0.797	0.647	0.734	0.798	0.771	0.799	0.805
Active_acc_95_all	0.938	0.949	0.954	0.948	0.931	0.946	0.947
Inactive_acc_95_all	0.702	0.445	0.587	0.699	0.664	0.701	0.711
Fitted_auc_ligand	0.742	0.742	0.742	0.742	0.742	0.742	0.742
Auc_improvement_ligand	17.270	6.690	9.670	17.740	12.170	15.300	17.350
Optimal_threshold_ligand	0.354	0.212	0.202	0.263	0.111	0.202	0.485
Accuracy_ligand	0.831	0.723	0.753	0.834	0.763	0.803	0.830
Active_pos_rate_ligand	0.883	0.795	0.814	0.877	0.898	0.901	0.792
Inactive_neg_rate_ligand	0.796	0.675	0.713	0.805	0.674	0.738	0.855
Threshold_95_ligand	0.131	0.030	0.051	0.111	0.030	0.101	0.141
Accuracy_95_ligand	0.797	0.680	0.709	0.808	0.713	0.763	0.802
Active_acc_95_ligand	0.939	0.928	0.953	0.933	0.954	0.943	0.936
Inactive_acc_95_ligand	0.702	0.514	0.546	0.724	0.552	0.642	0.712

Table S4: 2C9 Data

	30	40	60	20	50	80	70
Roc_auc_all	0.833	0.839	0.865	0.773	0.847	0.875	0.874
Bootstrapped_scores_mean_all	0.833	0.839	0.865	0.773	0.847	0.875	0.874
Bootstrapped_scores_std_all	0.011	0.011	0.011	0.013	0.012	0.011	0.011
Roc_auc_ligand	0.816	0.832	0.842	0.776	0.845	0.869	0.858
Bootstrapped_scores_mean_ligand	0.816	0.832	0.842	0.775	0.845	0.868	0.858
Bootstrapped_scores_std_ligand	0.012	0.011	0.012	0.011	0.011	0.011	0.011
Roc_auc_docked	0.773	0.797	0.823	0.711	0.813	0.857	0.844
Bootstrapped_scores_mean_docked	0.773	0.797	0.822	0.711	0.813	0.856	0.844
Bootstrapped_scores_std_docked	0.013	0.012	0.012	0.013	0.012	0.011	0.011
Fitted_auc_all	0.687	0.687	0.687	0.687	0.687	0.687	0.687
Auc_improvement_all	14.630	15.190	17.820	8.650	16.000	18.840	18.740
Optimal_threshold_all	0.273	0.646	0.495	0.758	0.636	0.333	0.485
Accuracy_all	0.736	0.786	0.818	0.740	0.810	0.832	0.832
Active_pos_rate_all	0.769	0.632	0.695	0.606	0.612	0.734	0.707
Inactive_neg_rate_all	0.725	0.837	0.860	0.785	0.877	0.865	0.875
Threshold_95_all	0.020	0.040	0.020	0.010	0.030	0.010	0.040
Accuracy_95_all	0.608	0.602	0.613	0.433	0.595	0.649	0.626
Active_acc_95_all	0.946	0.931	0.933	0.952	0.929	0.933	0.937
Inactive_acc_95_all	0.494	0.492	0.507	0.259	0.483	0.554	0.522
Fitted_auc_docked	0.687	0.687	0.687	0.687	0.687	0.687	0.687
Auc_improvement_docked	8.630	11.020	13.610	2.430	12.590	16.970	15.700
Optimal_threshold_docked	0.222	0.394	0.535	0.313	0.414	0.455	0.566
Accuracy_docked	0.631	0.697	0.781	0.689	0.723	0.808	0.806
Active_pos_rate_docked	0.844	0.774	0.646	0.533	0.768	0.717	0.630
Inactive_neg_rate_docked	0.560	0.671	0.827	0.741	0.709	0.838	0.865
Threshold_95_docked	0.061	0.121	0.030	0.030	0.101	0.071	0.061
Accuracy_95_docked	0.516	0.570	0.555	0.465	0.584	0.596	0.584
Active_acc_95_docked	0.944	0.934	0.944	0.923	0.933	0.948	0.941
Inactive_acc_95_docked	0.372	0.449	0.425	0.312	0.468	0.479	0.464
Fitted_auc_ligand	0.687	0.687	0.687	0.687	0.687	0.687	0.687
Auc_improvement_ligand	12.890	14.540	15.550	8.860	15.760	18.160	17.100
Optimal_threshold_ligand	0.586	0.556	0.404	0.869	0.475	0.636	0.253
Accuracy_ligand	0.745	0.779	0.794	0.753	0.786	0.834	0.797
Active_pos_rate_ligand	0.658	0.676	0.738	0.593	0.709	0.646	0.771
Inactive_neg_rate_ligand	0.774	0.813	0.813	0.807	0.812	0.896	0.806
Threshold_95_ligand	0.061	0.020	0.020	0.040	0.051	0.010	0.020
Accuracy_95_ligand	0.581	0.577	0.573	0.423	0.610	0.678	0.623
Active_acc_95_ligand	0.948	0.925	0.939	0.944	0.919	0.911	0.922
Inactive_acc_95_ligand	0.458	0.461	0.450	0.248	0.507	0.601	0.523

Table S5: 2C19

	30	50	70	80	40	60	20
Roc_auc_all	0.836	0.855	0.873	0.879	0.846	0.868	0.720
Bootstrapped_scores_mean_all	0.836	0.855	0.873	0.879	0.846	0.868	0.720
Bootstrapped_scores_std_all	0.011	0.011	0.011	0.010	0.011	0.011	0.012
Roc_auc_ligand	0.807	0.851	0.876	0.876	0.834	0.865	0.730
Bootstrapped_scores_mean_ligand	0.807	0.851	0.877	0.877	0.834	0.866	0.730
Bootstrapped_scores_std_ligand	0.013	0.011	0.010	0.010	0.012	0.011	0.013
Roc_auc_docked	0.756	0.793	0.833	0.837	0.790	0.815	0.659
Bootstrapped_scores_mean_docked	0.756	0.793	0.832	0.838	0.790	0.814	0.658
Bootstrapped_scores_std_docked	0.012	0.012	0.011	0.011	0.012	0.011	0.012
Fitted_auc_ligand	0.616	0.616	0.616	0.616	0.616	0.616	0.616
Auc_improvement_ligand	19.110	23.480	26.040	26.060	21.780	24.940	11.420
Optimal_threshold_ligand	0.364	0.354	0.404	0.434	0.263	0.606	0.323
Accuracy_ligand	0.716	0.757	0.810	0.813	0.745	0.809	0.665
Active_pos_rate_ligand	0.810	0.874	0.838	0.820	0.830	0.727	0.750
Inactive_neg_rate_ligand	0.657	0.684	0.793	0.809	0.693	0.860	0.611
Threshold_95_ligand	0.061	0.141	0.152	0.172	0.030	0.162	0.010
Accuracy_95_ligand	0.628	0.649	0.720	0.715	0.651	0.680	0.537
Active_acc_95_ligand	0.934	0.939	0.944	0.941	0.941	0.941	0.937
Inactive_acc_95_ligand	0.439	0.470	0.581	0.575	0.470	0.517	0.290
Fitted_auc_all	0.616	0.616	0.616	0.616	0.616	0.616	0.616
Auc_improvement_all	22.030	23.920	25.700	26.360	23.050	25.260	10.430
Optimal_threshold_all	0.535	0.475	0.303	0.384	0.293	0.404	0.303
Accuracy_all	0.762	0.785	0.790	0.808	0.744	0.802	0.656
Active_pos_rate_all	0.758	0.805	0.851	0.838	0.868	0.828	0.743
Inactive_neg_rate_all	0.765	0.772	0.752	0.790	0.667	0.785	0.601
Threshold_95_all	0.111	0.111	0.091	0.101	0.101	0.071	0.030
Accuracy_95_all	0.650	0.662	0.711	0.713	0.661	0.704	0.531
Active_acc_95_all	0.937	0.940	0.939	0.946	0.941	0.937	0.944
Inactive_acc_95_all	0.472	0.490	0.569	0.569	0.487	0.559	0.274
Fitted_auc_docked	0.616	0.616	0.616	0.616	0.616	0.616	0.616
Auc_improvement_docked	14.070	17.750	21.680	22.160	17.410	19.920	4.280
Optimal_threshold_docked	0.424	0.364	0.434	0.424	0.414	0.465	0.202
Accuracy_docked	0.661	0.680	0.747	0.756	0.696	0.720	0.580
Active_pos_rate_docked	0.813	0.876	0.807	0.783	0.783	0.778	0.778
Inactive_neg_rate_docked	0.567	0.559	0.709	0.739	0.642	0.684	0.457
Threshold_95_docked	0.202	0.242	0.182	0.162	0.061	0.202	0.040
Accuracy_95_docked	0.578	0.614	0.634	0.665	0.612	0.642	0.482
Active_acc_95_docked	0.944	0.936	0.949	0.939	0.936	0.950	0.938
Inactive_acc_95_docked	0.351	0.413	0.438	0.496	0.411	0.450	0.200

Table S6: 2D6

	30	70	60	80	50	20	40
Roc_auc_all	0.844	0.892	0.890	0.901	0.884	0.692	0.889
Bootstrapped_scores_mean_all	0.844	0.892	0.889	0.901	0.885	0.692	0.890
Bootstrapped_scores_std_all	0.014	0.012	0.013	0.012	0.012	0.015	0.012
Roc_auc_ligand	0.817	0.880	0.867	0.880	0.864	0.715	0.874
Bootstrapped_scores_mean_ligand	0.817	0.880	0.867	0.879	0.864	0.715	0.874
Bootstrapped_scores_std_ligand	0.015	0.014	0.014	0.014	0.013	0.017	0.014
Roc_auc_docked	0.823	0.842	0.839	0.854	0.833	0.624	0.827
Bootstrapped_scores_mean_docked	0.824	0.842	0.838	0.854	0.834	0.624	0.826
Bootstrapped_scores_std_docked	0.016	0.014	0.015	0.015	0.015	0.017	0.016
Fitted_auc_ligand	0.551	0.551	0.551	0.551	0.551	0.551	0.551
Auc_improvement_ligand	26.570	32.880	31.570	32.830	31.270	16.380	32.270
Optimal_threshold_ligand	0.162	0.192	0.313	0.697	0.253	0.051	0.545
Accuracy_ligand	0.847	0.873	0.893	0.915	0.874	0.516	0.899
Active_pos_rate_ligand	0.586	0.711	0.557	0.491	0.601	0.869	0.495
Inactive_neg_rate_ligand	0.886	0.897	0.944	0.980	0.916	0.462	0.961
Threshold_95_ligand	0.010	0.010	0.010	0.010	0.010	0.020	0.020
Accuracy_95_ligand	0.564	0.575	0.801	0.570	0.571	0.381	0.502
Active_acc_95_ligand	0.886	0.938	0.775	0.938	0.944	0.960	0.963
Inactive_acc_95_ligand	0.515	0.519	0.805	0.514	0.514	0.293	0.432
Fitted_auc_docked	0.551	0.551	0.551	0.551	0.551	0.551	0.551
Auc_improvement_docked	27.190	29.090	28.760	30.290	28.200	7.290	27.540
Optimal_threshold_docked	0.202	0.253	0.323	0.212	0.283	0.010	0.293
Accuracy_docked	0.833	0.853	0.874	0.864	0.855	0.727	0.858
Active_pos_rate_docked	0.553	0.590	0.493	0.640	0.530	0.430	0.568
Inactive_neg_rate_docked	0.876	0.893	0.933	0.898	0.905	0.773	0.902
Threshold_95_docked	0.020	0.020	0.010	0.010	0.020	0.010	0.020
Accuracy_95_docked	0.462	0.547	0.558	0.551	0.526	0.727	0.468
Active_acc_95_docked	0.940	0.925	0.919	0.925	0.929	0.430	0.936
Inactive_acc_95_docked	0.389	0.489	0.503	0.494	0.465	0.773	0.397
Fitted_auc_all	0.551	0.551	0.551	0.551	0.551	0.551	0.551
Auc_improvement_all	29.270	34.060	33.850	34.950	33.310	14.110	33.820
Optimal_threshold_all	0.131	0.232	0.323	0.202	0.192	0.121	0.111
Accuracy_all	0.844	0.890	0.899	0.889	0.874	0.698	0.871
Active_pos_rate_all	0.678	0.617	0.565	0.680	0.653	0.526	0.713
Inactive_neg_rate_all	0.869	0.932	0.950	0.921	0.908	0.725	0.896
Threshold_95_all	0.010	0.010	0.010	0.010	0.010	0.010	0.010
Accuracy_95_all	0.593	0.677	0.717	0.624	0.609	0.419	0.694
Active_acc_95_all	0.898	0.909	0.898	0.946	0.940	0.869	0.900
Inactive_acc_95_all	0.547	0.642	0.690	0.575	0.558	0.350	0.662

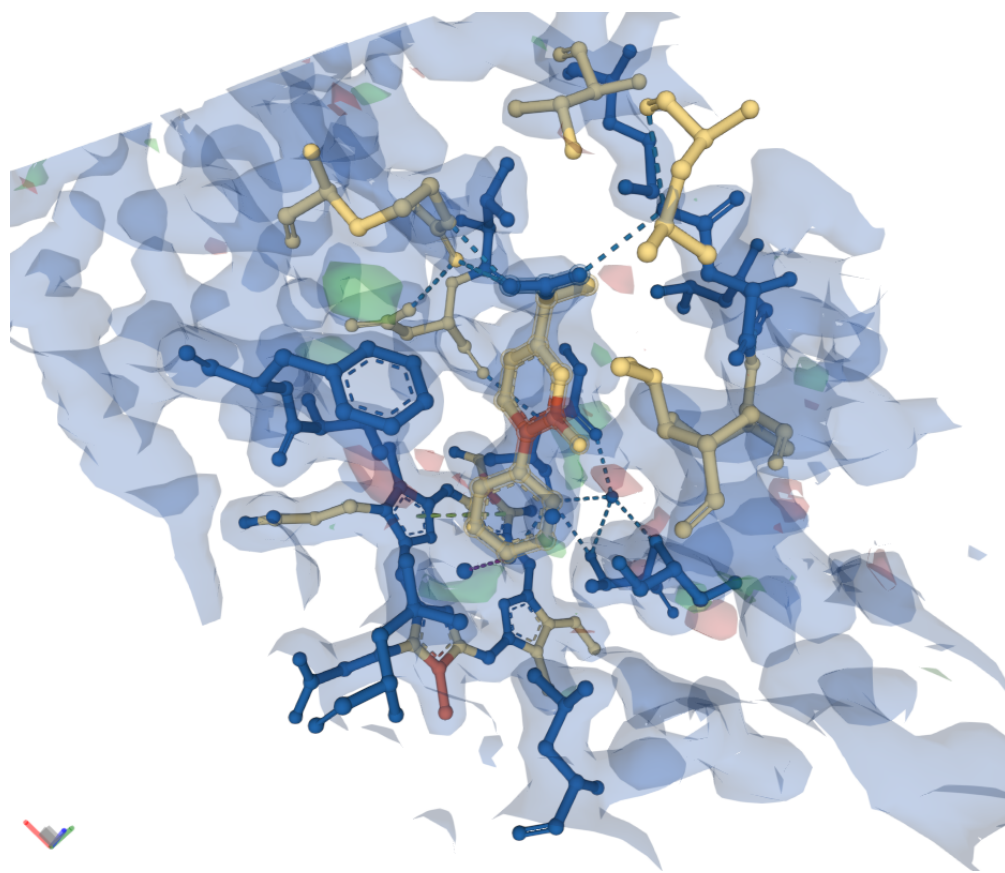


Figure S16: 1R9O: <https://www.rcsb.org/3d-view/1R90?preset=electronDensityMaps>

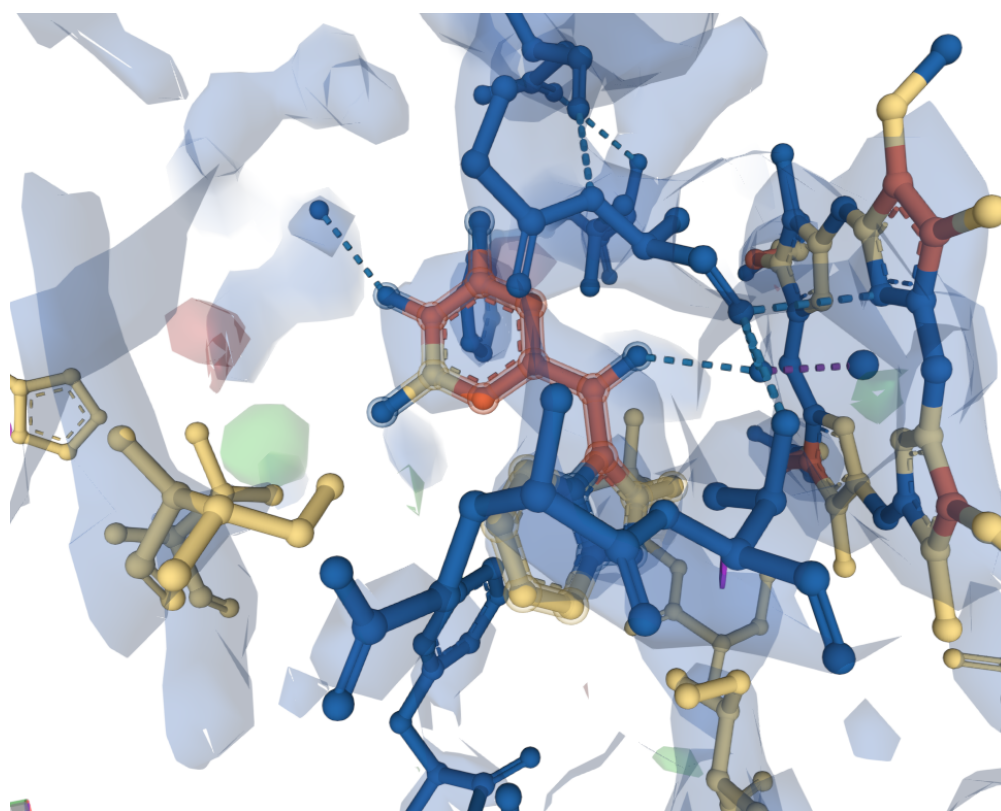


Figure S17: 4GQS: <https://www.rcsb.org/3d-view/4GQS?preset=electronDensityMaps>

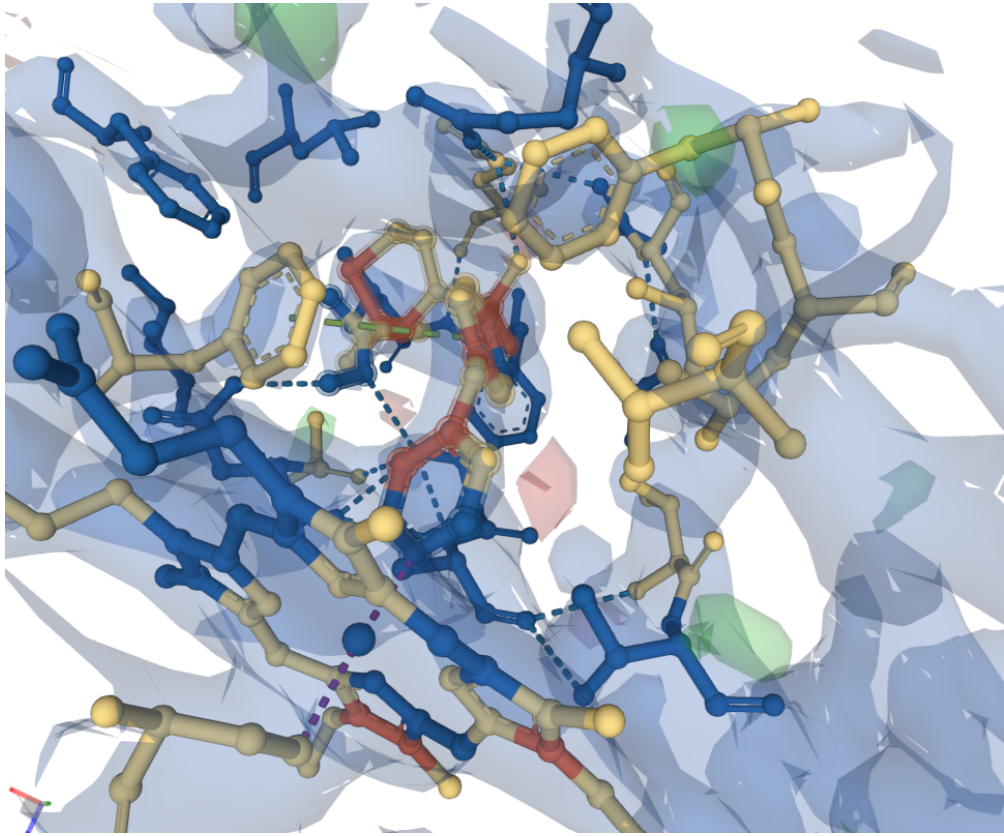


Figure S18: 3QM4: <https://www.rcsb.org/3d-view/3QM4?preset=electronDensityMaps>

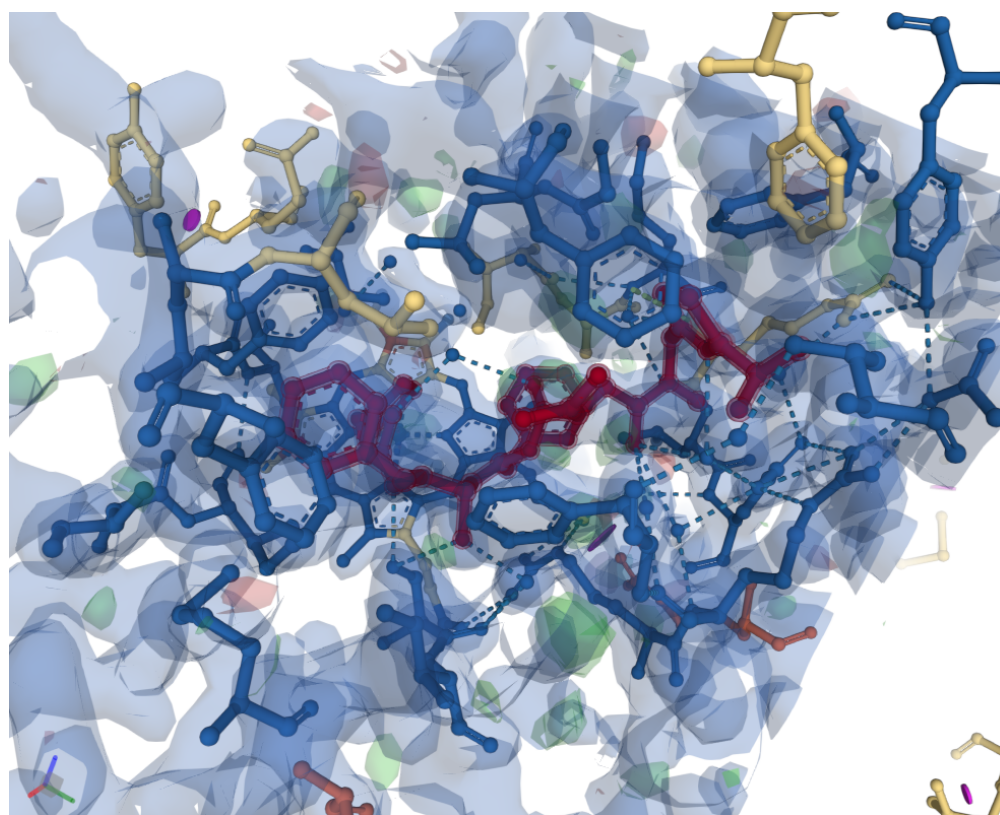


Figure S19: 3NXU: <https://www.rcsb.org/3d-view/3NXU?preset=electronDensityMaps>