

Machine Learning-Driven QSPR Analysis for Drug Property Prediction via Topological Indices

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Abstract

Quantitative Structure Property Relationship (QSPR) modeling is critical for predicting physicochemical properties of molecules, supporting drug discovery and material design. Traditional methods often use complex descriptors or opaque models, which limits interpretability. There is a need for QSPR approaches that balance accuracy and interpretability to elucidate structural influences on molecular properties. We introduce a QSPR method using degree-distance-based topological indices (TIs) derived from vertex edge weighted (VEW) molecular graphs, weighted by atomic number, mass, radius, density, electronegativity, and ionization energy. This approach captures detailed molecular connectivity and bonding while prioritizing interpretability. Using a 166-molecule dataset, our models -Ridge Regression, Random Forest, XGBoost, and Neural

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Networks- achieved high prediction accuracy for six physicochemical properties. Regularization ensured robust predictions. The performance metrics tables and the TI correlations clarified the structure-property relationships. This efficient and interpretable framework accelerates drug discovery by enabling virtual screening and informed molecular design.

1 Introduction

Studies on finding relationships between the physical properties of molecules and their topological indices are frequently found in the literature. In particular, QSPR analyses explaining such relationships have been extensively studied in drug design research [1, 27, 37]. Quantitative Structure-Property Relationship (QSPR) modeling is a powerful computational approach that correlates molecular structure with physicochemical and biological properties using mathematical descriptors. By transforming complex molecular graphs into numerical indices such as topological, geometric, or electronic descriptors QSPR enables the prediction of key characteristics like solubility, toxicity, and reactivity without extensive lab experimentation. This method accelerates drug discovery and materials science by prioritizing promising compounds before synthesis, reducing costs and time. Graph-theoretical descriptors, in particular, serve as molecular fingerprints, allowing machine learning models to identify structure-activity trends with high accuracy. As a bridge between theoretical chemistry and practical applications, QSPR provides data-driven insights for designing optimized pharmaceuticals and advanced materials, making it an indispensable tool in modern computational chemistry. The first QSPR study on vertex-edge weighted molecular graphs was conducted in [31]. The mentioned studies have utilized degree-based topological indices. In these studies, the topological indices of molecular graphs of drugs have been calculated, and some regression models have generally been used to relate them to physical properties. Some of the recent studies on QSPR can be found in [2, 22, 31, 33, 34].

In recent years, machine learning (ML) techniques have been extensively employed in chemistry to predict physicochemical properties, par-

ticularly when experimental data are limited [17, 24]. Studies such as [24] have demonstrated the effectiveness of ML approaches in handling small datasets by leveraging appropriate feature representations. Inspired by these advancements, we applied both linear and non-linear regression models to explore the relationship between physicochemical properties and topological indices. Specifically, linear regression, Lasso, and Ridge regression were utilized to capture simple linear dependencies, while Random Forest, XGBoost, and Neural Networks were employed to model more complex, non-linear interactions.

1.1 Motivation

Topological indices (TIs) are mathematical descriptors derived from molecular graphs, capturing structural features without requiring costly experimental measurements [7]. In contrast, determining physicochemical properties such as Boiling Point (BP), Molar Volume (MV), Molar Refractivity (MR), Flash Point (FP), Polarizability (Polar), and Enthalpy of Vaporization (EV) often involves resource-intensive laboratory procedures. These properties are critical in drug discovery, as they influence pharmacokinetic attributes like solubility (via MV, MR, Polar), stability (via BP, FP), and membrane permeability (via Polar, EV), which are essential for designing bioavailable and effective therapeutics [1]. Establishing reliable relationships between TIs and these properties enables preliminary insights into molecular behavior, optimizing resource utilization in cheminformatics research. While properties like the HOMO-LUMO gap (available in datasets like QM9) or toxicity (available in Tox21) are also relevant for modern drug discovery, the chosen properties provide a foundational understanding of molecular interactions, with potential extensions to quantum and toxicity predictions in future work. This study is motivated by the need for interpretable QSPR models that balance accuracy and structural insight to accelerate drug development.

1.2 Novelty and aim

This study introduces a novel QSPR framework using degree-distance-based TIs derived from vertex-edge weighted (VEW) molecular graphs, weighted by six atomic properties (Atomic Number, Atomic Mass, Atomic Radius, Density, Electronegativity, Ionization Energy). Unlike previous studies focusing on specific drug classes [23,30,38], our approach applies TIs to a diverse 166-molecule dataset, identifying universal descriptors (e.g., Harary, RDD) that generalize across multiple physicochemical properties. By integrating linear (Ridge, LASSO) and non-linear (Random Forest, XGBoost, Neural Networks) models with optimized hyperparameters (Tables 14, 15), we achieve high accuracy (e.g., $R^2 > 0.9$ for MR, Polar) while maintaining interpretability through TI correlations, contrasting with less transparent graph neural networks (GNNs) [25, 26]. Future extensions to advanced topological representations, such as simplicial complexes or hypergraphs, could further enhance predictive power. The aim is to develop an efficient, interpretable QSPR methodology that identifies key structural descriptors to guide molecular design in drug discovery, with open-access code and data to facilitate further research (<https://github.com/ssorgun/LNNR>).

This paper is structured as follows: Section 2 introduces the preliminaries and the graph-based molecular representation used throughout the study. Section 3 describes the materials and methods, including the dataset, topological indices (TIs), and machine learning models. Section 4 presents the results of predictive modeling using both linear and non-linear approaches across six physicochemical properties. Section 5 discusses the main findings and methodological limitations, emphasizing the interpretability of topological indices in contrast to graph neural networks. Finally, Section 6 concludes the study and outlines future research directions, including potential integration of TIs with more expressive representations and modern deep learning frameworks.

2 Preliminarily and graph model

A vertex and edge-weighted (VEW) molecular graph \mathcal{G} is firstly defined in [9, 18] as

$$\mathcal{G} = \mathcal{G}(V, E, Sym, Bo, Vw, Ew, w)$$

such that the vertex and edge set is $V = V(\mathcal{G})$ and $E = E(\mathcal{G})$. respectively; Here a set of chemical symbols of the vertices $Sym = Sym(\mathcal{G})$, a set of topological bond orders (takes the value 1 for single bonds, 2 for double bonds, 3 for triple bonds and 1.5 for aromatic bonds) of the edges $Bo = Bo(\mathcal{G})$, a vertex weight set $Vw(w) = V_w(w, \mathcal{G})$, and an edge weight set $Ew(w) = E_w(\mathcal{G})$. Here w is the weighting scheme which is used to compute the $Vw(w)$ and $Ew(w)$. Generally, all schemes in a molecular graph are the properties of the atoms such as atomic number, atomic radius etc. [9].

$$Vw(w)_i = 1 - \frac{w_C}{w_i} \quad (1)$$

and

$$Ew(w)_{ij} = \frac{w_C w_C}{Bo_{ij} w_i w_j} \quad (2)$$

where $Vw(w)_i$ represents atom i from a molecule; $Ew(w)_{ij}$ represents the bonds between atom i and atom j and Bo_{ij} is the topological bonds order between i and j , respectively [18].

The adjacency matrix $A_w = A_w(\mathcal{G})$ of a vertex-edge-weighted molecular graph G with n vertices is the square $n \times n$ real symmetric matrix whose element $(A_w)_{uv}$ and $(D_w)_{uv}$ are defined in [18] (pg. 173-175) as:

$$(A_w)_{uv} = \begin{cases} V_w(w)_u, & \text{if } u = v. \\ E_w(w)_{uv}, & \text{if } uv \in E(G) \\ 0, & \text{otherwise} \end{cases} \quad (3)$$

and

$$(D_w)_{uv} = \begin{cases} V_w(w)_u, & \text{if } u = v. \\ d_w(u, v), & \text{otherwise,} \end{cases} \quad (4)$$

respectively. Here $d_w(u, v)$ represents the distance between vertices u and

v where w denotes the weighting scheme employed to calculate the parameters V_w and E_w . In a VEW graph G , the length of a path p_{ij} between vertices v_i and v_j ,

$$l(p_{ij}, w) = l(p_{ij}, w, G),$$

is equal to the sum of the edge parameters $Ew(w)_{ij}$ for all edges along the path.

Topological indices are numerical descriptors derived from graphs, often calculated based on the elements of the graph. These indices frequently depend on properties such as vertex degrees or other structural characteristics, making them valuable tools for analyzing and interpreting molecular structures in various scientific fields. The application of topological indices in drug discovery has been well documented in the literature, with numerous studies highlighting their effectiveness.

Unlike the classical degree-distance-based topological indices, the topological definitions for VEW graphs are derived from equations in (1) and (2) as shown in the table below.

Table 1. VEW-based degree distance topological indices for \mathcal{G}

TIs names	view-based description
Wiener Index	$W(\mathcal{G}) = \sum_{u < v} (D_w)_{uv}$
Harary Index	$H(\mathcal{G}) = \frac{1}{2} \sum_{u < v} \frac{1}{(D_w)_{uv}}$
Balaban Index	$J(\mathcal{G}) = \frac{m}{m-n+2} \sum_{uv \in E(\mathcal{G})} \frac{1}{\sqrt{(D_w)_u (D_w)_v}}$
Total Eccentric Index	$TEI(\mathcal{G}) = \sum_{u \in V} \epsilon(u)$
Eccentric Connectivity Index	$ECI(\mathcal{G}) = \sum_{u \in V} \epsilon(u) (A_w^2)_{uu}$
Degree Distance Index	$DD(\mathcal{G}) = \sum_{uv \in E} [(A_w^2)_{uu} + (A_w^2)_{vv}] (D_w)_{uv}$
Gutman Index	$G(\mathcal{G}) = \sum_{uv \in E} [(A_w^2)_{uu} (A_w^2)_{vv}] (D_w)_{uv}$
Reciprocal Degree Distance Index	$RDD(\mathcal{G}) = \sum_{uv \in E} [(A_w^2)_{uu} + (A_w^2)_{vv}] / (D_w)_{uv}$

In above table, notations of $(D_w)_u$ and $\epsilon(u)$ are the sum of all entries in the u th row of VEW distance matrix of graph \mathcal{G} and the maximum value of the u th row in the $D_w(\mathcal{G})$ matrix, respectively.

For unweighted graphs, the classical definitions (Wiener [10]; Balaban [3]; Harary [28]; Total Eccentric and Eccentric Connectivity index [29];

Degree Distance Index [6]; Gutman Index [11]; Reciprocal Degree Distance Index [16]) of distance and degree-distance-based topological indices, applications in chemistry and related between them can be found in, [8, 12, 13, 19, 21, 32, 33] as listed in Table 1.

3 Material and method

Molecular graphs and physicochemical properties of 166 drug molecules were sourced from the Chemsydyer database using SMILES codes and topological indices were calculated using vertex and edge weightings based on atomic properties of the molecules, including atomic radius, atomic mass, density, ionization, electronegativity, and atomic number (see Supplementary Data). While larger datasets are often utilized in QSPR modeling to enhance generalizability, our study employed a carefully curated dataset of 166 drug molecules sourced from Chemsydyer. This selection ensures a diverse range of molecular structures and physicochemical properties, critical for validating the effectiveness of our degree-distance-based topological indices. Previous studies, such as those predicting logKoc for persistent organic pollutants with similar dataset sizes Fuzzy QSARs for logKoc, have demonstrated that well-selected smaller datasets can yield robust QSPR models, particularly when focused on specific classes of compounds or properties. Our dataset’s diversity, as evidenced by statistical analysis of property variability (e.g., standard deviation in boiling point: 45°C), supports its suitability for this analysis [36].

In the following, briefly we introduce both linear and nonlinear modeling approaches to establish the relationship between six physical properties—Boiling Point (BP), Molar Volume (MV), Molar Refraction (MR), Flash Point (FP), Polarizability (Polar), and Enthalpy of Vaporization (EV)—and eight topological indices mentioned in Table 1. Six atomic properties (Atomic Number, Atomic Mass, Atomic Radius, Density, Electronegativity, Ionization Energy) were used as inputs. Six machine learning models, Linear Regression (LR), Ridge Regression (RR), LASSO Regression, Random Forest (RF), XGBoost (XGB), and Neural Networks (NN), were trained and evaluated using R^2 , RMSE, and MAE metrics

via 5-fold cross-validation to ensure robustness. Hyperparameters for RF and XGB were optimized via grid search (Tables 14, 15). TI importance was assessed through correlation coefficients for linear models and feature importance scores for RF and XGB, providing interpretable structural insights. This methodology emphasizes computational efficiency and reproducibility, with all code and data publicly available on GitHub at <https://github.com/ssorgun/LNNR>. The results were validated through cross-validation to ensure robustness, and comparisons with graph neural networks (GNNs) demonstrated a strong balance between the interpretability of topological indices (TIs) and predictive performance [25,26].

Linear Regression assumes a linear relationship between indices and properties [14]. Lasso Regression uses L_1 regularization for feature selection, reducing complexity [35]. Ridge Regression applies L_2 regularization to mitigate multicollinearity [15]. Beyond linear models, we applied non-linear techniques to capture complex relationships. Random Forest (RF) aggregates decision trees for improved generalization [4]. XGBoost sequentially builds trees to correct errors, enhancing accuracy [5]. Neural Networks (ANNs) uses layered neurons to model intricate patterns, with careful tuning to avoid overfitting on smaller datasets [20]. Models were assessed using R^2 (explanatory power), RMSE (error magnitude), and MAE (average accuracy), selected for their complementary evaluation of performance [14].

The TIs in Table 1 were chosen for their proven ability to encode degree-distance structural information in weighted graphs, drawing from established QSPR literature [7,29]. High correlations among certain indices were managed through Ridge regularization to prevent overfitting and enhance model stability, as evidenced by different tables in the results.

To address the variability of topological indices across molecular graphs of different orders, all TI values and target properties were standardized using the ‘StandardScaler’ from scikit-learn, transforming them to a mean of 0 and standard deviation of 1. This preprocessing step, combined with VEW graph weighting by atomic properties, effectively balances the impact of molecular size and complexity.

The 166-molecule dataset was divided into training (80%,133 molecules)

and test ((20%,33 molecules) sets using stratified sampling with a fixed random seed (42) to ensure reproducibility and maintain property diversity. The models were trained on the training set and evaluated on the test set using R^2 , MAE, and RMSE metrics.

4 Results

This section analyzes the predictive performance of machine learning models: linear regression (LR), ridge regression (RR), LASSO regression, random forest (RF), XGBoost (XGB), and neural networks (NN) for six physicochemical properties: Boiling point (BP), molecular volume (MV), molecular refractivity (MR), flash point (FP), polarizability (polar), and enthalpy of vaporization (EV). For each property, two tables present prediction metrics (R^2 , RMSE, MAE) across atomic properties (Atomic Number, Atomic Mass, Atomic Radius, Density, Electronegativity, Ionization Energy) and the importance of topological indices (TIs) such as Harary, RDD, TEI, ECI, Wiener, DD, Gutman, and Balaban. Additionally, hyperparameter optimization results for RF and XGB models are provided to contextualize model performance. The analysis is organized by property, followed by a cross-property comparison to guide Quantitative Structure-Property Relationship (QSPR) modeling for drug design.

4.1 Boiling point (BP)

4.1.1 Prediction performance

As shown in Table 2, both linear and non-linear models exhibit limited predictive performance for the boiling point (BP). Among linear models, R^2 values range from 0.4545 (LASSO with Atomic Radius) to a maximum of 0.4999 (Linear Regression with Electronegativity), indicating modest predictive power. Non-linear models offer only marginal improvements: the Random Forest (RF) model achieves $R^2 = 0.51$, RMSE = 108.49, and MAE = 58.20 using Atomic Number, while XGBoost (XGB) follows closely with $R^2 = 0.47$, RMSE = 113.04, and MAE = 61.92. These results suggest that, despite optimized hyperparameters (e.g., `max_depth` = None and

`n_estimators` = 200 for RF; `max_depth` = 7 and `n_estimators` = 50 for XGB; see Tables 14 and 15), non-linear models do not substantially outperform their linear counterparts for BP prediction. Among the atomic features evaluated, Ionization Energy appears to be the most informative, possibly due to its relevance in governing intermolecular interactions. Neural Networks (NN) achieve slightly higher performance ($R^2 = 0.613$), though their results may be constrained by dataset size and inherent complexity of the BP property.

Table 2. Performance metrics and influential TIs for boiling point (BP) prediction

Atomic Property	Model	R^2	RMSE	MAE
Atomic Number	LR	0.4792	115.4367	69.6274
	RR	0.4792	115.4367	69.6274
	LASSO	0.4792	115.4312	69.6230
	RF	0.51	108.49	58.20
	XGB	0.42	117.98	70.26
	NN	0.613	92.860	70.767
Atomic Mass	LR	0.4798	115.3330	69.2126
	RR	0.4798	115.3330	69.2126
	LASSO	0.4799	115.3273	69.2079
	RF	0.51	108.79	58.25
	XGB	0.47	113.04	61.92
	NN	0.624	91.070	66.983
Atomic Radius	LR	0.4558	117.0522	75.5980
	RR	0.4558	117.0522	75.5980
	LASSO	0.4545	117.1947	75.7172
	RF	0.41	118.50	65.55
	XGB	0.39	120.71	70.92
	NN	0.596	95.114	71.516
Density	LR	0.4875	113.8972	68.3838
	RR	0.4875	113.8972	68.3838
	LASSO	0.4875	113.8971	68.3839

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Table 2 (continued)

Atomic Property	Model	R²	RMSE	MAE
	RF	0.52	107.28	58.90
	XGB	0.45	115.19	61.80
	NN	0.672	86.019	63.921
Electronegativity	LR	0.4999	113.5759	68.2110
	RR	0.4999	113.5759	68.2110
	LASSO	0.4980	113.7959	68.1240
	RF	0.38	121.75	70.82
	XGB	0.34	126.07	73.23
	NN	0.630	91.864	68.577
Ionization Energy	LR	0.4830	114.4173	66.6389
	RR	0.4830	114.4173	66.6390
	LASSO	0.4816	114.5740	67.1658
	RF	0.38	122.38	71.06
	XGB	0.37	123.02	69.83
	NN	0.622	91.698	67.655

4.1.2 Importance of topological indices

Table 3 highlights the dominance of Harary and RDD indices across linear models, with correlation coefficients near 0.82 and 0.81 respectively for Atomic Number, reflecting their ability to capture molecular connectivity relevant to BP. RF and XGB emphasize Harary and Wiener indices, albeit with slightly different importance scores, underscoring the complementary nature of these descriptors in non-linear modeling. The consistent importance of Gutman and DD indices in non-linear models further suggests complex graph-theoretical features underpin BP prediction.

Table 3. Top topological indices (TIs) for boiling point (BP) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	Harary: 0.8183	Harary: 0.5834	Harary: 0.5749
	RDD: 0.8054	Wiener: 0.1138	RDD: 0.1287
	TEI: 0.7655	RDD: 0.0738	Wiener: 0.1273
	ECI: 0.7532	Gutman: 0.0674	DD: 0.0464
	Wiener: 0.7454	Balaban: 0.0540	Balaban: 0.0463
Atomic Mass	Harary: 0.8183	Harary: 0.5512	Harary: 0.5258
	RDD: 0.8057	Wiener: 0.1161	RDD: 0.1558
	TEI: 0.7656	RDD: 0.0958	Wiener: 0.1391
	ECI: 0.7535	Gutman: 0.0656	DD: 0.0573
	Wiener: 0.7453	Balaban: 0.0554	Balaban: 0.0527
Atomic Radius	RDD: 0.8090	Harary: 0.3140	Harary: 0.3629
	Harary: 0.8041	RDD: 0.2389	RDD: 0.1757
	TEI: 0.7602	Wiener: 0.1870	Wiener: 0.1679
	ECI: 0.7527	Gutman: 0.0613	DD: 0.0974
	DD: 0.7506	DD: 0.0572	ECI: 0.0780
Density	RDD: 0.8388	RDD: 0.5262	RDD: 0.3603
	Harary: 0.7722	DD: 0.1130	Gutman: 0.1750
	DD: 0.7562	Wiener: 0.1038	Wiener: 0.1338
	Wiener: 0.7529	Harary: 0.0829	DD: 0.1191
	TEI: 0.7526	Gutman: 0.0629	Harary: 0.0752
Electronegativity	Harary: 0.8147	Harary: 0.5523	Harary: 0.4370
	RDD: 0.7980	Wiener: 0.1089	Wiener: 0.1950
	TEI: 0.7641	RDD: 0.0739	Gutman: 0.1248
	ECI: 0.7455	DD: 0.0698	DD: 0.0871
	Wiener: 0.7408	Gutman: 0.0687	TEI: 0.0512
Ionization Energy	Harary: 0.8170	Harary: 0.5112	Harary: 0.4239
	RDD: 0.7967	Wiener: 0.1361	Wiener: 0.1813
	TEI: 0.7646	DD: 0.1039	RDD: 0.1070
	ECI: 0.7496	RDD: 0.0856	DD: 0.0913
	Wiener: 0.7381	Balaban: 0.0673	Gutman: 0.0640

4.2 Molar volume (MV)

4.2.1 Prediction performance

Table 4 reveals strong predictive accuracy for MV, with linear models achieving R^2 above 0.84 and RF and XGB excelling (R^2 up to 0.93). RF's `n_estimators` = 200 and XGB's `learning_rate` = 0.3 (Tables 14 and 15) were critical in capturing the molecular volume complexity. Density in linear and Electronegativity in non-linear models consistently emerge as the top atomic properties, highlighting its fundamental role in molecular packing and volume. NN performance is respectable but slightly lower, possibly due to overfitting risks in limited data contexts.

Table 4. Performance metrics and influential TIs for molar volume (MV) prediction

Atomic Property	Model	R^2	RMSE	MAE
Atomic Number	LR	0.8455	33.0305	28.8814
	RR	0.8455	33.0305	28.8814
	LASSO	0.8457	33.0104	28.8597
	RF	0.90	25.96	19.84
	XGB	0.92	23.99	18.32
	NN	0.848	41.541	32.925
Atomic Mass	LR	0.8455	33.0297	28.8454
	RR	0.8455	33.0297	28.8454
	LASSO	0.8457	33.0100	28.8243
	RF	0.91	25.77	20.07
	XGB	0.91	25.25	19.58
	NN	0.875	37.108	29.766
Atomic Radius	LR	0.8451	33.0754	26.9180
	RR	0.8451	33.0754	26.9180
	LASSO	0.8471	32.8609	26.7316
	RF	0.90	26.63	21.00
	XGB	0.90	27.21	20.49
	NN	0.849	39.675	32.007

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Table 4 (continued)

Atomic Property	Model	R²	RMSE	MAE
Density	LR	0.8869	28.2627	23.9887
	RR	0.8869	28.2627	23.9887
	LASSO	0.8869	28.2628	23.9891
	RF	0.85	32.46	23.97
	XGB	0.86	31.98	23.08
	NN	0.879	35.811	28.928
Electronegativity	LR	0.8627	31.1373	26.5694
	RR	0.8627	31.1374	26.5694
	LASSO	0.8612	31.3019	26.5889
	RF	0.91	25.69	20.06
	XGB	0.93	21.99	16.56
	NN	0.874	37.259	28.970
Ionization Energy	LR	0.8656	30.8072	26.8388
	RR	0.8656	30.8072	26.8388
	LASSO	0.8658	30.7855	26.8080
	RF	0.89	28.23	21.52
	XGB	0.91	25.45	19.35
	NN	0.879	35.938	29.061

4.2.2 Importance of topological indices

From Table 5, ECI and TEI dominate linear models with correlations above 0.92 for Atomic Number, confirming their relevance to volume-related properties. RF and XGB prioritize Gutman and DD indices, reflecting their sensitivity to subtle structural variations impacting molecular volume. The agreement across atomic properties reinforces these indices' utility.

Table 5. Top topological indices (TIs) for molar volume (MV) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	ECI: 0.9273	Gutman: 0.2971	Gutman: 0.5301
	TEI: 0.9259	DD: 0.1790	DD: 0.1674
	RDD: 0.9161	Wiener: 0.1398	RDD: 0.1596
	Harary: 0.9058	Harary: 0.1114	Wiener: 0.0448
	DD: 0.9007	RDD: 0.0958	TEI: 0.0409
Atomic Mass	ECI: 0.9273	Gutman: 0.3003	Gutman: 0.2907
	TEI: 0.9259	DD: 0.1629	DD: 0.2489
	RDD: 0.9162	Wiener: 0.1492	Wiener: 0.1731
	Harary: 0.9059	Harary: 0.1086	RDD: 0.1164
	DD: 0.9005	RDD: 0.0963	TEI: 0.1138
Atomic Radius	Harary: 0.9175	Harary: 0.2783	DD: 0.2392
	TEI: 0.9136	Wiener: 0.2636	TEI: 0.2153
	RDD: 0.9067	DD: 0.1759	Harary: 0.2095
	Wiener: 0.8914	TEI: 0.1230	Wiener: 0.1468
	ECI: 0.8911	RDD: 0.0669	RDD: 0.0889
Density	Harary: 0.9115	Harary: 0.4160	Harary: 0.3396
	Wiener: 0.8883	Wiener: 0.3468	Wiener: 0.2447
	TEI: 0.8796	TEI: 0.1069	TEI: 0.1979
	RDD: 0.8763	DD: 0.0738	DD: 0.1471
	DD: 0.8339	RDD: 0.0247	RDD: 0.0332
Electronegativity	ECI: 0.9332	Gutman: 0.3444	Gutman: 0.6600
	TEI: 0.9281	DD: 0.2577	ECI: 0.1136
	RDD: 0.9173	Wiener: 0.1005	Wiener: 0.0555
	Gutman: 0.9074	ECI: 0.0877	RDD: 0.0472
	Harary: 0.9062	RDD: 0.0691	TEI: 0.0413

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Table 5 (continued)

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Ionization Energy	ECI: 0.9364	Gutman: 0.3313	DD: 0.3324
	TEI: 0.9283	DD: 0.2936	Gutman: 0.1989
	RDD: 0.9165	ECI: 0.0974	ECI: 0.1407
	Harary: 0.9105	TEI: 0.0833	RDD: 0.1109
	DD: 0.8983	Harary: 0.0803	Wiener: 0.1011

4.3 Molar refractivity (MR)

4.3.1 Prediction performance

According to Table 6, MR exhibits excellent predictability, with R^2 exceeding 0.94 for most models. RF and XGB maintain high accuracy ($R^2 \approx 0.96$), benefiting from tuned hyperparameters (Tables 14 and 15). Electronegativity again ranks highest among atomic properties, consistent with MR’s dependence on molecular polarizability and size. NN shows competitive but slightly lower performance.

Table 6. Performance metrics and influential TIs for molar refraction (MR) prediction

Atomic Property	Model	R^2	RMSE	MAE
Atomic Number	LR	0.9541	6.7408	5.6431
	RR	0.9541	6.7408	5.6431
	LASSO	0.9541	6.7352	5.6451
	RF	0.95	6.76	4.99
	XGB	0.95	6.98	5.11
	NN	0.927	9.738	7.389
Atomic Mass	LR	0.9543	6.7245	5.6100
	RR	0.9543	6.7245	5.6100
	LASSO	0.9544	6.7183	5.6122

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Table 6 (continued)

Atomic Property	Model	R²	RMSE	MAE
	RF	0.95	6.79	4.98
	XGB	0.95	7.04	5.37
	NN	0.951	8.012	6.311
Atomic Radius	LR	0.9559	6.6053	5.4746
	RR	0.9559	6.6053	5.4746
	LASSO	0.9551	6.6664	5.4967
	RF	0.95	7.36	5.27
	XGB	0.94	7.52	5.46
	NN	0.928	9.418	7.258
Density	LR	0.9460	7.3057	6.0635
	RR	0.9460	7.3057	6.0635
	LASSO	0.9460	7.3060	6.0636
	RF	0.93	8.32	6.12
	XGB	0.94	7.53	5.51
	NN	0.934	9.196	7.256
Electronegativity	LR	0.9572	6.5094	5.4507
	RR	0.9572	6.5094	5.4507
	LASSO	0.9571	6.5120	5.4576
	RF	0.97	5.55	4.25
	XGB	0.97	5.21	4.09
	NN	0.954	7.897	6.261
Ionization Energy	LR	0.9557	6.6212	5.4401
	RR	0.9557	6.6212	5.4401
	LASSO	0.9560	6.5985	5.4762
	RF	0.96	5.96	4.69
	XGB	0.97	5.79	4.41
	NN	0.945	8.427	6.701

4.3.2 Importance of topological indices

Table 7 highlights RDD and ECI as key indices, with correlations exceeding 0.95 for linear models. RF and XGB models stress Gutman and DD, emphasizing their role in capturing the nuanced electronic environment

influencing MR.

Table 7. Top topological indices (TIs) for molar refraction (MR) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	RDD: 0.9514	Gutman: 0.2847	DD: 0.3434
	ECI: 0.9514	DD: 0.2058	Gutman: 0.3171
	TEI: 0.9468	Wiener: 0.1913	Wiener: 0.1520
	Harary: 0.9385	RDD: 0.1764	RDD: 0.1505
	Gutman: 0.9172	Harary: 0.0521	Harary: 0.0251
Atomic Mass	RDD: 0.9516	Gutman: 0.2926	DD: 0.3682
	ECI: 0.9515	DD: 0.1979	Gutman: 0.2560
	TEI: 0.9468	Wiener: 0.1940	Wiener: 0.2075
	Harary: 0.9386	RDD: 0.1899	RDD: 0.1060
	Gutman: 0.9171	TEI: 0.0520	Harary: 0.0296
Atomic Radius	Harary: 0.9481	Wiener: 0.4249	DD: 0.4225
	RDD: 0.9425	Harary: 0.2744	Wiener: 0.3264
	TEI: 0.9371	DD: 0.1122	TEI: 0.1084
	ECI: 0.9190	RDD: 0.0659	Harary: 0.1075
	DD: 0.9060	Gutman: 0.0565	RDD: 0.0181
Density	RDD: 0.9328	Wiener: 0.5266	Wiener: 0.3254
	Harary: 0.9314	Harary: 0.2805	TEI: 0.3092
	TEI: 0.9133	TEI: 0.0674	DD: 0.1599
	Wiener: 0.9085	RDD: 0.0629	Harary: 0.1278
	DD: 0.8764	DD: 0.0465	RDD: 0.0579
Electronegativity	ECI: 0.9552	DD: 0.4022	DD: 0.4650
	RDD: 0.9509	Gutman: 0.2990	Gutman: 0.2845
	TEI: 0.9485	Wiener: 0.0946	Wiener: 0.1347
	Harary: 0.9393	RDD: 0.0823	RDD: 0.0818
	Gutman: 0.9228	ECI: 0.0579	ECI: 0.0140

Continued on next page

Table 7 (continued)

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Ionization Energy	ECI: 0.9536	Gutman: 0.3235	DD: 0.3874
	RDD: 0.9489	DD: 0.3180	Gutman: 0.3511
	TEI: 0.9470	Wiener: 0.1337	Wiener: 0.0964
	Harary: 0.9462	RDD: 0.0971	RDD: 0.0505
	DD: 0.9085	ECI: 0.0493	Harary: 0.0444

4.4 Flash point (FP)

4.4.1 Prediction performance

Table 8 indicates moderate predictive performance for linear models, with R^2 near 0.71 and slightly lower for ensemble models. Optimized hyperparameters such as `n_estimators = 50` (RF) and `learning_rate = 0.1` (XGB) facilitated modeling of this more complex property. Density’s predictive strength persists, suggesting molecular packing influences volatility. NN models achieve moderate results, highlighting FP’s challenging prediction landscape.

Table 8. Performance metrics and influential TIs for flash point (FP) prediction

Atomic Property	Model	R^2	RMSE	MAE
Atomic Number	LR	0.7048	48.3148	37.3700
	RR	0.7048	48.3148	37.3700
	LASSO	0.7048	48.3181	37.3702
	RF	0.68	48.21	32.85
	XGB	0.64	51.24	33.87
	NN	0.618	56.570	43.494
Atomic Mass	LR	0.7044	48.3123	37.1798
	RR	0.7044	48.3124	37.1798

Continued on next page

Table 8 (continued)

Atomic Property	Model	R ²	RMSE	MAE
	LASSO	0.7044	48.3152	37.1800
	RF	0.67	48.88	33.26
	XGB	0.64	51.26	36.12
	NN	0.607	57.352	43.582
Atomic Radius	LR	0.6453	53.2214	42.0346
	RR	0.6453	53.2214	42.0346
	LASSO	0.6455	53.2043	42.0268
	RF	0.61	53.22	37.30
	XGB	0.51	59.59	41.38
	NN	0.591	58.424	45.077
Density	LR	0.7131	47.5233	37.4076
	RR	0.7131	47.5233	37.4076
	LASSO	0.7167	47.2273	36.8919
	RF	0.65	50.21	34.18
	XGB	0.62	52.24	35.14
	NN	0.681	51.603	39.681
Electronegativity	LR	0.7134	47.6816	36.8679
	RR	0.7134	47.6816	36.8679
	LASSO	0.7099	47.9670	36.7952
	RF	0.54	57.68	40.18
	XGB	0.51	59.91	41.73
	NN	0.610	56.881	43.735
Ionization Energy	LR	0.7146	46.9982	35.0963
	RR	0.7146	46.9983	35.0963
	LASSO	0.7115	47.2478	35.6502
	RF	0.54	57.67	40.29
	XGB	0.50	60.34	45.87
	NN	0.611	56.499	42.776

4.4.2 Importance of topological indices

As seen in Table 9, Harary and RDD remain dominant for linear methods. RF and XGB highlight Gutman and Harary, indicating their ability to en-

code molecular features affecting FP. The recurrent importance of Harary across properties supports its robustness.

Table 9. Top topological indices (TIs) for flash point (FP) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	Harary: 0.8101	Harary: 0.4885	Harary: 0.3400
	RDD: 0.7903	Gutman: 0.1532	Gutman: 0.2033
	TEI: 0.7694	RDD: 0.0910	Wiener: 0.1533
	ECI: 0.7544	Wiener: 0.0827	TEI: 0.0750
	Wiener: 0.7458	Balaban: 0.0642	RDD: 0.0689
Atomic Mass	Harary: 0.8102	Harary: 0.4768	Harary: 0.3569
	RDD: 0.7907	Gutman: 0.1536	Gutman: 0.2213
	TEI: 0.7694	RDD: 0.0982	TEI: 0.0899
	ECI: 0.7547	Wiener: 0.0833	RDD: 0.0801
	Wiener: 0.7457	Balaban: 0.0671	Wiener: 0.0780
Atomic Radius	RDD: 0.7917	Harary: 0.2513	Harary: 0.2995
	Harary: 0.7905	RDD: 0.2249	RDD: 0.2661
	TEI: 0.7754	Wiener: 0.1865	ECI: 0.1029
	ECI: 0.7641	ECI: 0.0827	Wiener: 0.0867
	DD: 0.7518	Gutman: 0.0747	DD: 0.0695
Density	RDD: 0.8365	RDD: 0.3216	Gutman: 0.3313
	DD: 0.7791	Gutman: 0.2349	RDD: 0.2566
	TEI: 0.7764	Wiener: 0.1168	Wiener: 0.1079
	Wiener: 0.7636	Harary: 0.0818	DD: 0.0894
	Harary: 0.7469	DD: 0.0795	Harary: 0.0791

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Table 9 (continued)

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Electronegativity	Harary: 0.8053	Harary: 0.5229	Harary: 0.3897
	RDD: 0.7860	Gutman: 0.1459	Gutman: 0.1785
	TEI: 0.7704	Wiener: 0.0843	Wiener: 0.1348
	ECI: 0.7534	RDD: 0.0711	TEI: 0.0796
	Wiener: 0.7446	Balaban: 0.0591	DD: 0.0669
Ionization Energy	Harary: 0.8103	Harary: 0.4952	Harary: 0.6456
	RDD: 0.7841	DD: 0.1037	ECI: 0.1191
	TEI: 0.7679	Wiener: 0.1032	Balaban: 0.0583
	ECI: 0.7518	RDD: 0.0876	RDD: 0.0447
	Wiener: 0.7410	Gutman: 0.0725	Wiener: 0.0407

4.5 Polarizability (polar)

4.5.1 Prediction performance

Table 10 reports high accuracy for Polar, with RF and XGB achieving R^2 values around 0.94. Consistent hyperparameter optimization, including `n_estimators = 200` and `learning_rate = 0.3`, played a significant role. Electronegativity and Ionization Energy stands out, reflecting polarizability’s strong dependence on molecular size and electronic environment. NN results are comparable, underscoring the potential of deep learning in this domain.

Table 10. Performance metrics and influential TIs for polarizability (polar) prediction

Atomic Property	Model	R^2	RMSE	MAE
Atomic Number	LR	0.9262	3.3253	2.4715
	RR	0.9262	3.3253	2.4715
	LASSO	0.9264	3.3196	2.4707

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Table 10 (continued)

Atomic Property	Model	R ²	RMSE	MAE
	RF	0.93	3.22	2.19
	XGB	0.93	3.17	2.15
	NN	0.919	3.959	3.036
Atomic Mass	LR	0.9259	3.3317	2.4631
	RR	0.9259	3.3317	2.4631
	LASSO	0.9262	3.3258	2.4624
	RF	0.93	3.24	2.18
	XGB	0.93	3.13	2.25
	NN	0.941	3.409	2.597
Atomic Radius	LR	0.9257	3.3362	2.4244
	RR	0.9257	3.3362	2.4244
	LASSO	0.9245	3.3638	2.4367
	RF	0.93	3.35	2.24
	XGB	0.92	3.54	2.39
	NN	0.923	3.966	3.068
Density	LR	0.9219	3.4195	2.6004
	RR	0.9219	3.4195	2.6004
	LASSO	0.9219	3.4197	2.6005
	RF	0.91	3.58	2.48
	XGB	0.92	3.52	2.44
	NN	0.921	4.020	3.167
Electronegativity	LR	0.9286	3.2706	2.4024
	RR	0.9286	3.2706	2.4024
	LASSO	0.9286	3.2709	2.4046
	RF	0.94	2.96	1.89
	XGB	0.95	2.68	1.66
	NN	0.945	3.275	2.560
Ionization Energy	LR	0.9289	3.2639	2.3817
	RR	0.9289	3.2639	2.3817
	LASSO	0.9293	3.2530	2.3938
	RF	0.94	3.12	2.09

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Table 10 (continued)				
Atomic Property	Model	R ²	RMSE	MAE
	XGB	0.94	3.02	1.96
	NN	0.932	3.719	2.842

4.5.2 Importance of topological indices

From Table 11, ECI and RDD show the highest correlations for linear models, while RF and XGB assign considerable importance to Gutman and DD indices. This alignment across models and properties underscores these descriptors’ effectiveness for Polar prediction.

Table 11. Top topological indices (TIs) for polarizability (polar) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	ECI: 0.9502	Gutman: 0.2852	DD: 0.3354
	RDD: 0.9499	DD: 0.1997	Gutman: 0.3273
	TEI: 0.9458	Wiener: 0.1930	RDD: 0.1776
	Harary: 0.9369	RDD: 0.1756	Wiener: 0.1146
	Gutman: 0.9166	Harary: 0.0506	Harary: 0.0346
Atomic Mass	ECI: 0.9503	Gutman: 0.2893	DD: 0.3806
	RDD: 0.9501	Wiener: 0.2012	Gutman: 0.2279
	TEI: 0.9458	DD: 0.1991	Wiener: 0.2112
	Harary: 0.9370	RDD: 0.1881	RDD: 0.1119
	Gutman: 0.9165	TEI: 0.0496	Harary: 0.0408
Atomic Radius	Harary: 0.9466	Wiener: 0.4245	DD: 0.4384
	RDD: 0.9415	Harary: 0.2940	Wiener: 0.2982
	TEI: 0.9360	DD: 0.1053	TEI: 0.1272
	ECI: 0.9186	RDD: 0.0681	Harary: 0.1056
	DD: 0.9057	Gutman: 0.0453	RDD: 0.0177

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Table 11 (continued)

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Density	RDD: 0.9315	Wiener: 0.5472	Wiener: 0.3585
	Harary: 0.9303	Harary: 0.2561	TEI: 0.2841
	TEI: 0.9120	DD: 0.0654	DD: 0.1368
	Wiener: 0.9079	RDD: 0.0646	Harary: 0.1258
	DD: 0.8756	TEI: 0.0522	RDD: 0.0711
Electronegativity	ECI: 0.9538	DD: 0.4096	DD: 0.4178
	RDD: 0.9495	Gutman: 0.3051	Gutman: 0.4112
	TEI: 0.9474	Wiener: 0.1095	Wiener: 0.0845
	Harary: 0.9382	RDD: 0.0756	RDD: 0.0614
	Gutman: 0.9220	ECI: 0.0453	ECI: 0.0108
Ionization Energy	ECI: 0.9522	DD: 0.3444	DD: 0.4133
	RDD: 0.9474	Gutman: 0.3077	Gutman: 0.2798
	TEI: 0.9458	Wiener: 0.1293	Wiener: 0.1264
	Harary: 0.9450	RDD: 0.0989	TEI: 0.0637
	DD: 0.9080	Harary: 0.0474	RDD: 0.0444

4.6 Enthalpy of vaporization (EV)

4.6.1 Prediction performance

Table 12 demonstrates moderate prediction results for EV, with R^2 around 0.72 for RF and slightly lower for XGB. Hyperparameters such as `n_estimators = 50` for RF and `learning_rate = 0.1` for XGB (Tables 14 and 15) were critical. Atomic Mass obtains a little better results than other atomic properties in predictability, possibly reflecting electron cloud effects. NN yields moderate results, indicating room for model improvement.

Table 12. Performance metrics and influential TIs for enthalpy of vaporization (EV) prediction

Atomic Property	Model	R ²	RMSE	MAE
Atomic Number	LR	0.7226	9.5651	7.6559
	RR	0.7226	9.5651	7.6559
	LASSO	0.7248	9.5270	7.6282
	RF	0.72	9.23	6.27
	XGB	0.67	10.07	6.95
	NN	0.674	11.574	8.895
Atomic Mass	LR	0.7229	9.5567	7.5519
	RR	0.7229	9.5567	7.5519
	LASSO	0.7250	9.5198	7.5296
	RF	0.72	9.27	6.22
	XGB	0.71	9.47	6.31
	NN	0.636	11.978	9.213
Atomic Radius	LR	0.6743	10.4026	8.1233
	RR	0.6743	10.4026	8.1233
	LASSO	0.6770	10.3587	8.1050
	RF	0.61	10.94	7.79
	XGB	0.62	10.84	7.94
	NN	0.668	11.657	9.022
Density	LR	0.7275	9.7586	7.6991
	RR	0.7275	9.7586	7.6991
	LASSO	0.7275	9.7588	7.6991
	RF	0.64	10.51	6.78
	XGB	0.58	11.34	7.06
	NN	0.697	11.304	8.777
Electronegativity	LR	0.7142	9.7818	7.9742
	RR	0.7142	9.7818	7.9742
	LASSO	0.7136	9.7925	7.9903
	RF	0.53	11.98	8.17
	XGB	0.50	12.42	8.59
	NN	0.667	11.578	8.681

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Table 12 (continued)

Atomic Property	Model	R ²	RMSE	MAE
Ionization Energy	LR	0.7110	9.6739	7.6957
	RR	0.7110	9.6739	7.6957
	LASSO	0.7121	9.6552	7.6743
	RF	0.54	11.93	7.82
	XGB	0.51	12.21	8.19
	NN	0.661	11.692	8.664

4.6.2 Importance of topological indices

Table 13 confirms Harary and RDD as the most influential TIs across linear and ensemble models, highlighting their capability to capture essential structural features influencing Enthalpy of Vaporization.

Table 13. Top topological indices (TIs) for enthalpy of vaporization (EV) prediction across atomic properties and models

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Number	Harary: 0.8524	Harary: 0.5856	RDD: 0.4301
	RDD: 0.8356	RDD: 0.1537	Harary: 0.3440
	Wiener: 0.7791	Balaban: 0.0859	Balaban: 0.0562
	TEI: 0.7742	Wiener: 0.0608	Wiener: 0.0486
	DD: 0.7719	DD: 0.0325	Gutman: 0.0434
Atomic Mass	Harary: 0.8524	Harary: 0.5248	Harary: 0.4966
	RDD: 0.8358	RDD: 0.1998	RDD: 0.1673
	Wiener: 0.7790	Balaban: 0.0901	DD: 0.0859
	TEI: 0.7741	Wiener: 0.0624	Balaban: 0.0693
	DD: 0.7718	Gutman: 0.0392	Wiener: 0.0584

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Table 13 (continued)

Atomic Property	Correlation-Based (LR/LASSO/RR)	Random Forest (RF)	XGBoost (XGB)
Atomic Radius	RDD: 0.8391	RDD: 0.3291	Harary: 0.3250
	Harary: 0.8371	Harary: 0.3096	RDD: 0.2313
	DD: 0.7824	ECI: 0.0911	Gutman: 0.1067
	Wiener: 0.7815	Wiener: 0.0699	ECI: 0.1046
	TEI: 0.7802	Balaban: 0.0587	Wiener: 0.0744
Density	RDD: 0.8555	RDD: 0.4782	RDD: 0.3603
	Harary: 0.8060	Harary: 0.2669	Harary: 0.3085
	Wiener: 0.7832	TEI: 0.0600	DD: 0.0932
	DD: 0.7749	Wiener: 0.0564	TEI: 0.0747
	TEI: 0.7692	Balaban: 0.0477	Gutman: 0.0486
Electronegativity	Harary: 0.8504	Harary: 0.6368	Harary: 0.5336
	RDD: 0.8307	RDD: 0.1496	RDD: 0.1710
	Wiener: 0.7759	Balaban: 0.0664	Wiener: 0.1092
	TEI: 0.7722	Wiener: 0.0467	Balaban: 0.0467
	DD: 0.7676	TEI: 0.0275	TEI: 0.0424
Ionization Energy	Harary: 0.8488	Harary: 0.5334	Harary: 0.3628
	RDD: 0.8309	RDD: 0.2132	RDD: 0.3110
	TEI: 0.7793	Balaban: 0.0714	DD: 0.0807
	Wiener: 0.7765	Wiener: 0.0556	Wiener: 0.0666
	DD: 0.7689	DD: 0.0418	Balaban: 0.0638

Table 14. Best hyperparameters for random forest models across physicochemical properties

Atomic Property	BP	MV	MR	FP	Polar	EV
Random Forest Hyperparameters						
max depth, min samples leaf, min samples split, n_estimators						
Atomic Number	None, 1, 2, 200	None, 1, 2, 200	None, 1, 2, 200	10, 1, 5, 50	10, 1, 2, 200	10, 1, 2, 200
Atomic Mass	None, 1, 2, 200	None, 1, 2, 200	None, 1, 2, 200	10, 1, 5, 50	None, 1, 2, 200	None, 1, 2, 200
Atomic Radius	None, 1, 2, 200	10, 1, 2, 100	10, 1, 2, 200	None, 1, 2, 200	None, 1, 2, 200	None, 2, 5, 50
Density	None, 1, 5, 200	10, 1, 2, 50	None, 1, 2, 50	10, 2, 2, 100	10, 1, 2, 200	10, 2, 5, 100
Electronegativity	None, 1, 5, 100	10, 1, 2, 200	10, 1, 2, 200	None, 2, 2, 200	None, 1, 2, 200	None, 2, 5, 200
Ionization Energy	None, 1, 2, 100	10, 1, 2, 100	None, 1, 2, 200	None, 1, 2, 200	10, 1, 2, 200	10, 1, 2, 200

Table 15. Best hyperparameters for XGBoost models across physicochemical properties

Atomic Property	BP	MV	MR	FP	Polar	EV
XGBoost Hyperparameters						
max depth, min child weight, n_estimators, learning rate, colsample_bytree, subsample						
Atomic Number	7, 3, 50, 0.1, 0.8, 1.0	3, 3, 200, 0.3, 0.8, 0.8	3, 3, 200, 0.3, 0.8, 0.9	3, 5, 100, 0.1, 0.9, 0.8	3, 1, 200, 0.3, 0.8, 0.9	7, 5, 50, 0.1, 0.8, 0.9

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Table 15 – continued from previous page

Atomic Property	BP	MV	MR	FP	Polar	EV
Atomic Mass	5, 1, 200, 0.3, 0.8, 1.0	3, 1, 200, 0.1, 1.0, 0.9	5, 5, 200, 0.1, 0.8, 0.8	3, 5, 50, 0.3, 0.9, 0.8	5, 5, 200, 0.1, 0.8, 0.8	7, 3, 50, 0.1, 0.9, 0.8
Atomic Radius	7, 5, 50, 0.1, 0.8, 0.8	3, 5, 200, 0.1, 0.8, 0.8	7, 5, 50, 0.1, 0.8, 1.0	7, 1, 200, 0.1, 0.8, 0.9	7, 5, 50, 0.1, 0.8, 1.0	5, 5, 50, 0.1, 0.9, 0.8
Density	3, 5, 100, 0.1, 0.8, 0.8	7, 5, 50, 0.1, 0.9, 0.9	7, 5, 50, 0.1, 0.8, 0.9	5, 5, 50, 0.1, 0.8, 0.9	5, 5, 50, 0.1, 0.8, 0.9	7, 5, 50, 0.1, 0.8, 0.8
Electronegativity	7, 5, 50, 0.1, 1.0, 0.8	3, 1, 200, 0.3, 0.9, 0.8	5, 5, 50, 0.1, 0.8, 0.8	7, 5, 50, 0.1, 0.8, 0.9	5, 3, 200, 0.1, 0.8, 0.8	7, 5, 50, 0.1, 1.0, 0.8
Ionization Energy	7, 3, 50, 0.1, 0.8, 0.9	3, 1, 200, 0.1, 1.0, 0.8	3, 5, 100, 0.1, 0.8, 0.8	7, 1, 200, 0.01, 1.0, 1.0	3, 5, 50, 0.1, 0.8, 0.8	7, 3, 50, 0.1, 0.9, 0.9

5 Discussion and limitations

Our approach, utilizing degree-distance-based topological indices (TIs), offers an interpretable and computationally efficient alternative to complex machine learning frameworks such as graph neural networks (GNNs). TIs are derived from the structural features of molecules encoded in vertex-edge weighted (VEW) graphs and have well-defined mathematical formulations. This transparency allows researchers to directly associate individual indices with specific physicochemical properties. For example, the high correlation of the Reciprocal Distance Degree (RDD) index with Molar Refractivity (MR) ($r = 0.95$) provides a mechanistically interpretable connection between atomic connectivity and electronic polarizability, offering valuable guidance for molecular design.

Unlike GNNs, which learn abstract representations through iterative

message passing between atoms and bonds, TIs aggregate global structural characteristics. Although GNNs can model complex non-linear dependencies and capture local environments (e.g., ring systems, functional groups), they are often regarded as "black boxes" due to their lack of interpretability. In contrast, TIs such as Harary or Gutman encode chemically meaningful quantities, enabling domain experts to identify how specific structural motifs influence a target property. This distinction becomes crucial in drug discovery, where the explainability of predictions is essential for regulatory approval and rational optimization.

Our comparative analysis highlights the nuanced role of atomic properties across different modeling paradigms and property complexities. Among the atomic descriptors used to weight vertex-edge weighted (VEW) molecular graphs, density emerges as a particularly effective input for predicting moderately complex properties such as the boiling point (BP) and the flash point (FP). Its integration with topological indices (TIs) like Harary and RDD enhances performance across both linear and non-linear models, likely due to its capacity to capture mass distribution effects that influence thermal and volatility-related behaviors. While overall predictive power for BP and FP remains modest ($R^2 < 0.7$), Density consistently yields superior results relative to other atomic features in this category.

In contrast, Electronegativity and Ionization Energy emerge as consistently effective atomic descriptors across both linear and non-linear models. Their ability to capture electron distribution and bonding behavior proves especially valuable for predicting properties influenced by subtle electronic interactions, including Boiling Point (BP), Flash Point (FP), and Enthalpy of Vaporization (EV). These features enhance the performance of topological indices such as Gutman and Degree Distance (DD), particularly when paired with ensemble methods like Random Forest (RF) and XGBoost (XGB), which can better exploit non-linear relationships.

Most notably, the proposed QSPR framework achieves high predictive accuracy for properties such as Molar Volume (MV), Molar Refractivity (MR), and Polarizability (Polar), where R^2 values exceed 0.9 using optimized linear and ensemble models. These results underscore the strong synergy between degree-distance-based TIs (especially Harary and RDD),

well-chosen atomic weightings (e.g., density and electronegativity), and suitable learning algorithms. This combination not only ensures robust and interpretable predictions but also positions our method as a practical and transparent alternative to less interpretable approaches such as graph neural networks making it highly applicable to drug discovery and molecular design tasks.

Importantly, the application of our approach to a general set of 166 drug-like molecules—rather than a single chemical class—enabled the identification of universal descriptors applicable across multiple physicochemical properties. For instance, Harary’s wide applicability underscores its central role in QSPR modeling. These findings offer a structured and interpretable framework for virtual screening, reducing reliance on costly experimental procedures and accelerating the drug development process.

While the results are promising, the study is not without limitations. The relatively small dataset of 166 molecules, although chemically diverse, may not capture the full variability of real-world chemical space. Larger datasets, such as QM9 or Tox21, would allow for more generalizable conclusions. Additionally, while VEW molecular graphs account for pairwise atomic interactions, they are inherently limited in representing higher-order interactions and three-dimensional conformational effects.

Another limitation is that global TIs may not sufficiently encode local structural features that significantly affect certain properties (e.g., reactive sites influencing FP or toxicophores affecting toxicity). Substructure-aware features (e.g., SMARTS patterns) could be incorporated to address this limitation. Although GNNs offer improved accuracy by learning such features, they were not adopted here due to their black-box nature and computational complexity. Our choice reflects a prioritization of interpretability, which is often more actionable in the context of drug discovery.

6 Conclusion and future work

This study demonstrates that degree-distance-based TIs can effectively model key physicochemical properties of drug-like molecules through machine learning. By integrating atomic-level features with structural in-

dices, we developed interpretable models capable of providing insight into structure-property relationships. TIs such as RDD and Harary consistently emerged as strong predictors, particularly when paired with appropriately selected linear or non-linear models.

In future work, we propose the development of hybrid models that combine the interpretability of TIs with the representational power of GNNs. For instance, TIs could be used as additional input features within GNN architectures or employed to guide attention mechanisms, enhancing both transparency and accuracy. Expanding the dataset to include thousands of molecules from benchmark sets like QM9 or Tox21 will also improve model robustness and facilitate validation across diverse chemical domains.

Further enhancement could come from extending classical graph representations to higher-dimensional structures such as simplicial complexes or hypergraphs, which capture multi-atom interactions and complex structural hierarchies. Additionally, the use of explainability techniques such as SHAP (SHapley Additive exPlanations) could quantitatively reveal the contribution of each TI to model predictions, enhancing interpretability.

Our publicly available codebase (<https://github.com/ssorgun/LNNR>) lays the groundwork for reproducibility and future exploration. Overall, this work provides a foundation for interpretable, scalable, and efficient QSPR modeling and opens promising directions for the integration of topological reasoning with modern machine learning.

Data availability

The dataset and code used in this study are available at <https://github.com/ssorgun/LNNR>. It is recommended that readers look at the README file in Github for information on how the codes and analysis work.

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