

```
In [64]: import numpy as np
import pandas as pd
from rdkit import Chem
from rdkit.Chem import Descriptors
from rdkit.Chem.MolStandardize import rdMolStandardize
from rdkit.ML.Descriptors import MoleculeDescriptors as md
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, f1_score, roc_auc_score
import xgboost as xgb
import matplotlib.pyplot as plt
import pandas as pd
import xgboost as xgb
from sklearn.model_selection import train_test_split
from sklearn.metrics import (
    accuracy_score, f1_score, precision_score, recall_score,
    roc_auc_score, confusion_matrix, classification_report, roc_curve
)
import matplotlib.pyplot as plt
import seaborn as sns
```

```
In [65]: def molecule_from_smiles(smiles):
    try:
        # Extract molecule
        molecule = Chem.MolFromSmiles(smiles, sanitize=True)
        if molecule is None:
            return None, "failed"

        # Remove salts
        clean_molecule = rdMolStandardize.LargestFragmentChooser()
        molecule = clean_molecule.choose(molecule)

        # Sanitize molecule again to reflect changes
        Chem.SanitizeMol(molecule)
        return molecule, "succeed"
    except Exception as e:
        return None, f"error: {e}"

def calculate_descriptors(molecule):
    # Get all descriptors (1D/2D)
    descriptor_names = []
    for descriptor, _ in Descriptors._descList:
        descriptor_names.append(descriptor)

    # Use descriptors to calculate values
    calculator = md.MolecularDescriptorCalculator(descriptor_names)
    descriptor_values = calculator.CalcDescriptors(molecule)

    # Create dictionary
    descriptors = dict(zip(descriptor_names, descriptor_values))
    return descriptors
```

```
In [66]: dataset = pd.read_excel("in_chemico_dataset.xlsx", engine="openpyxl", skiprows=1)

descriptor_rows = []
state_molecules = []
molecules = []
```

```
for smiles in dataset["SMILES code"].astype(str):
    # Get molecule
    molecule, state = molecule_from_smiles(smiles)
    state_molecules.append(state)
    molecules.append(molecule)

    # Calculate
    if molecule is None:
        descriptor_rows.append({})
        continue

    descriptors = calculate_descriptors(molecule)
    descriptor_rows.append(descriptors)

descriptor_data = pd.DataFrame(descriptor_rows)

numeric_columns = []
for col in descriptor_data.columns:
    if pd.api.types.is_numeric_dtype(descriptor_data[col]):
        numeric_columns.append(col)

clean_desc = descriptor_data[numeric_columns].copy()

for col in clean_desc.columns:
    clean_desc[col] = clean_desc[col].replace([np.inf, -np.inf], np.nan)

# Drop columns with any NaN
cols_with_nan = []
for col in list(clean_desc.columns):
    if clean_desc[col].isna().any():
        cols_with_nan.append(col)

if len(cols_with_nan) > 0:
    clean_desc = clean_desc.drop(columns=cols_with_nan)

constant_cols = []
for col in list(clean_desc.columns):
    if clean_desc[col].nunique(dropna=False) <= 1:
        constant_cols.append(col)

if len(constant_cols) > 0:
    clean_desc = clean_desc.drop(columns=constant_cols)

output = pd.concat([dataset.reset_index(drop=True), descriptor_data.reset_index()])
output["MoleculeStatus"] = state_molecules

# Save
with pd.ExcelWriter("in_chemico_dataset_processed.xlsx", engine="openpyxl"):
    # Full report with raw 1D/2D descriptors
    output.to_excel(writer, index=False, sheet_name="Raw_Descriptors")

    # Clean X for XGBoost
    clean_desc.to_excel(writer, index=False, sheet_name="X_1D2D_clean")

clean_desc.to_csv("X_1D2D_descriptors.csv", index=False)

print(f"Rows: {len(output)}/Columns in raw: {output.shape[1]}")
print(f"Columns in X (clean): {clean_desc.shape[1]}")
print("First rows of X:")
```

```
print(clean_desc.head().to_string(index=False))
```

```
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: C=C[C@H]1CN2CC[C@H]1C[C@H]2[C@H](0)c1ccnc2ccc(OC)cc12
[02:34:29] New largest fragment: C=C[C@H]1CN2CC[C@H]1C[C@H]2[C@H](0)c1ccnc2ccc(OC)cc12 (48)
[02:34:29] Fragment: Cl
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: CN(C)CCCN1c2cccc2Sc2ccc(Cl)cc21
[02:34:29] New largest fragment: CN(C)CCCN1c2cccc2Sc2ccc(Cl)cc21 (40)
[02:34:29] Fragment: Cl
[02:34:29] Running LargestFragmentChooser
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: CC1c2cccc(0)c2C(0)=C2C(=0)C3(0)C(0)=C(C(N)=0)C(=0)C(N(C))C3C(0)C21
[02:34:29] New largest fragment: CC1c2cccc(0)c2C(0)=C2C(=0)C3(0)C(0)=C(C(N)=0)C(N(C))C3C(0)C21 (56)
[02:34:29] Fragment: Cl
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: [0-]n1cccc1=S
[02:34:29] New largest fragment: [0-]n1cccc1=S (12)
[02:34:29] Fragment: [Na+]
[02:34:29] Running LargestFragmentChooser
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: CN1CCN(CCCN2c3cccc3Sc3ccc(Cl)cc32)CC1
[02:34:29] New largest fragment: CN1CCN(CCCN2c3cccc3Sc3ccc(Cl)cc32)CC1 (49)
[02:34:29] Fragment: O=C(0)/C=C\C(=O)O
[02:34:29] Fragment: O=C(0)/C=C\C(=O)O
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: CC(CN1c2cccc2Sc2cccc21)N(C)C
[02:34:29] New largest fragment: CC(CN1c2cccc2Sc2cccc21)N(C)C (40)
[02:34:29] Fragment: Cl
[02:34:29] Running LargestFragmentChooser
[02:34:29] Fragment: Cc1ncc(C0)c(C0)c10
[02:34:29] New largest fragment: Cc1ncc(C0)c(C0)c10 (23)
[02:34:29] Fragment: Cl
[02:34:29] Running LargestFragmentChooser
[02:34:29] Running LargestFragmentChooser
```



```
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Fragment: CCOC(=0)C1=C(COCCN)NC(C)=C(C(=0)OC)C1c1cccc1Cl  
[02:34:30] New largest fragment: CCOC(=0)C1=C(COCCN)NC(C)=C(C(=0)OC)C1c1cc  
ccc1Cl (53)  
[02:34:30] Fragment: O=S(=0)(0)c1cccc1  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Fragment: COC(=0)C1=C(C)NC(C)=C(C(=0)O[C@H]2CCCN(Cc3cccc3)C2)  
[C@H]1c1cccc([N+](=0)[0-])c1  
[02:34:30] New largest fragment: COC(=0)C1=C(C)NC(C)=C(C(=0)O[C@H]2CCCN(C  
c3cccc3)C2)[C@H]1c1cccc([N+](=0)[0-])c1 (68)  
[02:34:30] Fragment: Cl  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Fragment: OCCN1CCN(CCCN2c3cccc3Sc3ccc(C(F)(F)F)cc32)CC1  
[02:34:30] New largest fragment: OCCN1CCN(CCCN2c3cccc3Sc3ccc(C(F)(F)F)cc3  
2)CC1 (56)  
[02:34:30] Fragment: Cl  
[02:34:30] Fragment: Cl  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Fragment: COC(=0)C1=C(C)NC(C)=C(C(=0)OCCN(C)Cc2cccc2)C1c1cccc  
([N+](=0)[0-])c1  
[02:34:30] New largest fragment: COC(=0)C1=C(C)NC(C)=C(C(=0)OCCN(C)Cc2cccc  
c2)C1c1cccc([N+](=0)[0-])c1 (64)  
[02:34:30] Fragment: Cl  
[02:34:30] Running LargestFragmentChooser  
[02:34:30] Fragment: CC(=0)CC(c1cccc1)c1c([0-])c2cccc2oc1=0  
[02:34:30] New largest fragment: CC(=0)CC(c1cccc1)c1c([0-])c2cccc2oc1=0  
(38)  
[02:34:30] Fragment: [Na+]  
[02:34:30] Running LargestFragmentChooser
```

Rows: 162/Columns in raw: 230
 Columns in X (clean): 191
 First rows of X:

	MaxAbsEStateIndex	MaxEStateIndex	MinAbsEStateIndex	MinEStateIndex																																																																		
qed	SPS	MolWt	HeavyAtomMolWt	ExactMolWt	NumValenceElectrons	Ma																																																																
xPartialCharge	MinPartialCharge	MaxAbsPartialCharge	MinAbsPartialCharge	FpDensityMorgan1	FpDensityMorgan2	FpDensityMorgan3	BCUT2D_MWHI	BCUT2D_M																																																														
MWLOW	BCUT2D_CHGHI	BCUT2D_CHGL0	BCUT2D_LOGPHI	BCUT2D_LOGPL0W	BCUT2D_M	RHI	BCUT2D_MRLOW	AvgIpc	BalabanJ	BertzCT	Chi0	Chi0n	Ch																																																									
i0v	Chi1	Chi1n	Chi1v	Chi2n	Chi2v	Chi3n	Chi3v	Chi	4n	Chi4v	HallKierAlpha	Ipc	Kappa1	Kappa2	Kappa3	Labu																																																						
teASA	PEOE_VSA1	PEOE_VSA10	PEOE_VSA11	PEOE_VSA12	PEOE_VSA13	PEOE_VSA	14	PEOE_VSA2	PEOE_VSA3	PEOE_VSA4	PEOE_VSA5	PEOE_VSA6	PEOE_VSA7	PEOE_VSA8	PEOE_VSA9	SMR_VSA1	SMR_VSA10	SMR_VSA3	SMR_VSA4	SMR_VSA5	SMR_V																																																	
SA6	SMR_VSA7	SMR_VSA9	SlogP_VSA1	SlogP_VSA10	SlogP_VSA11	SlogP_VSA12	SlogP_VSA2	SlogP_VSA3	SlogP_VSA4	SlogP_VSA5	SlogP_VSA6	SlogP_VSA7	SlogP_VSA8	TPSA	EState_VSA1	EState_VSA10	EState_VSA11	EState_VSA2	ESt	ate_VSA3	EState_VSA4	EState_VSA5	EState_VSA6	EState_VSA7	EState_VSA8																																													
EState_VSA9	VSA_EState1	VSA_EState10	VSA_EState2	VSA_EState3	VSA_ESta	te4	VSA_EState5	VSA_EState6	VSA_EState7	VSA_EState8	VSA_EState9	Frac	tionCSP3	HeavyAtomCount	NHOHCount	NOCount	NumAliphaticCarbocycles	Num	AliphaticHeterocycles	NumAliphaticRings	NumAmideBonds	NumAromaticCarbo	cycles	NumAromaticHeterocycles	NumAromaticRings	NumAtomStereoCenters	Nu	mBridgeheadAtoms	NumHAcceptors	NumHDonors	NumHeteroatoms	NumHeterocycl	es	NumRotatableBonds	NumSaturatedCarbocycles	NumSaturatedHeterocycles	NumSaturatedRings	NumSpiroAtoms	NumUnspecifiedAtomStereoCenters	Phi																														
RingCount	MolLogP	MolMR	fr_Al_COO	fr_Al_OH	fr_Al_OH_noTert	fr_ArN	fr_Ar_COO	fr_Ar_N	fr_Ar_NH	fr_Ar_OH	fr_COO	fr_COO2	fr_C_0	fr_C_0_no	COO	fr_HOCCN	fr_Imine	fr_NH0	fr_NH1	fr_NH2	fr_N_0	fr_Ndealkylation1	fr_Ndealkylation2	fr_Nhpyrrole	fr_alkyl_halide	fr_allylic_oxid	fr_amid	e	fr_amidine	fr_aniline	fr_aryl_methyl	fr_azo	fr_benzene	fr_bicyclic	fr_dihydropyridine	fr_ester	fr_ether	fr_furan	fr_guanido	fr_halogen	fr_hdrzine	fr_hdrzone	fr_imidazole	fr_imide	fr_ketone	fr_ketone_Topl	iss	fr_lactam	fr_lactone	fr_methoxy	fr_nitro	fr_nitro_arom	fr_nitro_a	rom_nonortho	fr_para_hydroxylation	fr_phenol	fr_phenol_noOrthoHbond	fr_piperidine	fr_piperazine	fr_priamide	fr_pyridine	fr_sulfide	fr_sulfona	md	fr_sulfone	fr_tetrazole	fr_thiazole	fr_thiophene	fr_unbrch_alkane	fr_urea
11.151855	11.151855	0.177547	-0.503545	0.87																																																																		
7602	30.250000	324.424	300.232	324.183778	126																																																																	
0.119124	-0.496743	0.496743	0.119124																																																																			
1.416667	2.291667	3.000000	16.465327	9.733492																																																																		
2.417618	-2.502700	2.417007	-2.544133	5.834889																																																																		
-0.044441	2.661658	1.687085	760.387301	16.681434	14.058926	14.058926	11.7																																																															
07040	8.682990	8.682990	6.974394	6.974394	5.819484	5.819484	4.444306	4.444																																																														
306	-1.91	512499.659037	15.608011	6.187608	2.502406	142.313426																																																																
9.843390	5.749512	0.000000	0.0	0.000000	0.000000	9.8																																																																
83888	0.000000	0.000000	6.578936	6.076020	61.051132	24.169665	1																																																															
8.730465	9.843390	10.902925	9.883888	11.835812	24.987450	20.199310	48.6																																																															
80719	5.749512	4.736863	0.0	5.749512	0.0	41.23																																																																
1567	0.000000	11.835812	24.509061	43.117268	0.0	10.9029																																																																
25	45.59	6.103966	5.106527	0.0	6.041841	11.8358																																																																
12	41.726223	6.420822	13.306641	24.265468	22.538844	4.																																																																
736863	5.349572	0.0	6.858527	12.130807	1.846194																																																																	
2.006859	7.957657	5.620375	6.068442	1.661567	0.450000																																																																	
24	1	4	0		3																																																																	
3	0		1		1																																																																	
2		5	2		1																																																																	
4	4		4	0																																																																		
3	3	0		1	4.024																																																																	

010	5	3.17320	95.0268	0	1	1
0	0	1	0	0	0	0
0	0	0	2	0	0	0
0	0	0	0	0	0	0
0	0	0	0	1	4	0
0	1	0	0	0	0	0
0	0	0	0	0	0	0
1	0	0	0	0	0	0
0	0	0	3	0	0	0
1	0	0	0	0	0	0
0	0	0	0	0	0	0
	12.268665	12.268665	0.241088	-4.588877	0.66	
0374	11.142857	308.311	296.215	308.035459		110
0.297751	-0.507039		0.507039		0.297751	
1.142857	1.761905		2.285714	32.239784	10.022356	
2.228520	-2.100377	2.321408	-2.067037	7.855380	0.	
103128	2.454894	2.629387	780.611574	15.620956	10.977121	11.793617
54	5.825519	7.265583	4.190432	5.580391	2.840472	3.715321
-2.38	42155.432869	15.017356	5.547507	3.037599	120.916319	9.843390
.394507	5.783245	0.0	10.118127	0.000000	9.347287	0.000
000	8.417797	0.000000	30.331835	6.066367	11.629819	12.673249
08474	15.901372	0.000000	0.000000	4.895483	7.109798	53.591472
24	4.736863	0.0	11.499024	0.0	30.970117	10.118
127	0.000000	15.921440	47.360053	0.0	0.000000	100.90
26.546367	18.318862	0.0	16.876415	0.000000		12.1327
34	19.242532	18.199101	0.000000	0.000000	9.289613	36.
586252	0.0	11.687259	9.867079	0.036000	-1.256141	
9.902413	0.000000	0.000000	-3.406195	0.071429		
21	2	6		0		0
0	0		2		0	
2	0		0		5	2
7	0		4		0	
0	0		0		0	3.967
090	2	1.87850	74.3479	0	0	0
0	0	0	0	1	0	1
1	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	2	0	0	0
0	1	0	0	0	0	0
0	0	1		0	0	0
1	0	0		0	0	0
1		1	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
	11.165950	11.165950	0.325325	-0.325325	0.70	
5368	10.533333	204.225	192.129	204.078644		78
0.335962	-0.493745		0.493745		0.335962	
1.333333	2.133333		2.866667	16.476922	10.129957	
2.026932	-2.042365	2.213955	-1.980500	5.806853	0.	
339807	2.167858	2.653374	540.048796	10.836499	8.741253	8.741253
09	4.877663	4.877663	3.339426	3.339426	2.233829	2.233829
-1.77	2829.054631	9.772439	3.768723	1.860292	87.287778	9.154014
.332532	0.000000	0.0	0.000000	5.625586	0.000000	4.794
537	0.000000	0.000000	0.000000	31.543660	17.518958	6.606882
54014	10.969244	0.000000	0.000000	13.847474	6.606882	40.249043
12	10.362449	0.0	5.749512	0.0	6.606882	0.000
000	6.923737	12.487189	33.477156	0.0	10.969244	39.44
0.000000	4.794537	0.0	5.625586	12.189902	16.69918	
8	6.066367	6.066367	25.980209	0.000000	9.154014	10.4
25547	0.0	11.165950	0.942642	1.170275	0.721065	

7.007289	0.000000	4.400566	0.000000	0.250000		
15	0	3	0	0	0	0
0	0		1	0	1	0
2		0	0	3	0	0
3		1	2		0	0
0		0	0		0	2.455
308	2	2.50012	58.3900	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	1	0	1	1	0	0
0	1	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
	10.801691	10.801691	0.299673	-0.299673	0.55	
2257	10.416667	160.172	152.108	160.052429		60
0.335690		-0.422729		0.422729		0.335690
1.166667		2.000000	2.833333	16.391868		10.153925
1.946105	-1.951119	2.120754	-1.934028	5.762922		0.
560734	2.053587	2.873950	468.467428	8.552042	6.703248	6.703248
20	3.761090	3.761090	2.795711	2.795711	1.774814	1.774814
-1.57	692.058854	7.099295	2.482514	1.114984	69.444349	4.417151
.583020	0.000000	0.0	0.000000	5.625586	0.000000	4.794
537	0.000000	0.000000	11.629819	25.122838	11.452591	0.000000
17151	10.969244	0.000000	0.000000	6.923737	0.000000	46.315410
00	5.625586	0.0	0.000000		0.0	0.000000
000	6.923737	5.563451	39.543523		0.0	10.969244
0.000000	4.794537		0.0	5.625586	5.583020	10.94967
6	6.066367	6.066367	25.122838	0.000000	4.417151	4.9
61065	0.0	10.801691	0.967222	1.510002		0.000000
8.922176	0.000000	2.004511	0.000000	0.100000		
12	0	2		0		0
0	0		1	0	1	0
2		0	0	2	0	0
2		1	0		0	0
0		0	0			0 1.468
675	2	2.10142	47.2210	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	1	0	1	1	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
0	0	0	0	0	0	0
	10.875576	10.875576	0.346525	-0.346525	0.62	
1753	10.230769	176.171	168.107	176.047344		66
0.335693		-0.496624		0.496624		0.335693
1.230769		2.000000	2.769231	16.466276		10.208834
1.975995	-1.992579	2.179711	-1.938475	5.771611		0.
413783	2.163185	2.746434	484.738995	9.259149	7.111496	7.111496
25	3.873462	3.873462	2.658188	2.658188	1.830255	1.830255
-1.77	1243.831488	7.857386	2.935678	1.402156	74.557894	9.154014
.332532	0.000000	0.0	0.000000	5.625586	0.000000	4.794
537	0.000000	0.000000	0.000000	18.199101	17.518958	7.109798
						9.1

```
54014 10.969244 0.000000 0.000000 0.000000 7.109798 40.751959 5.7495
12 10.362449 0.0 5.749512 0.0 7.109798 0.000
000 0.000000 0.000000 39.543523 0.0 10.969244 39.44
0.000000 4.794537 0.0 5.625586 11.332532 5.38622
4 6.066367 19.242532 12.132734 0.000000 9.154014 9.9
68674 0.0 10.875576 0.889259 0.201484 0.682269
8.477764 0.000000 0.000000 1.571641 0.100000
13 0 3 0 0 1 0 1 0
0 0 0 1 0 3 0 0
2 0 0 0 0 0 0 0 0
3 1 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0 1.774
366 2 1.80160 49.0360 0 0 0 0 0
0 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
0 0 0 0 1 0 0 0 0
0 1 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
1 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
0 0 0 0 0 0 0 0 0
```

```
In [67]: DESC_CSV = "X_1D2D_descriptors.csv"
ORIG_XLSX = "in_chemico_dataset.xlsx"
TARGET_COL = "Phototoxicity"
EXCEL_SKIPROWS = 1

# Output
TOP_FEATURES_CSV = "top_features_best.csv"

# Split train-test
X = pd.read_csv(DESC_CSV)
y = pd.read_excel(ORIG_XLSX, engine="openpyxl", skiprows=EXCEL_SKIPROWS)

X_train, X_test, y_train, y_test = train_test_split(
    X, y, test_size=0.2, random_state=42, stratify=y
)

# Initial model
base_model = xgb.XGBClassifier(
    n_estimators=400,
    learning_rate=0.05,
    max_depth=6,
    subsample=0.8,
    colsample_bytree=0.8,
    random_state=42,
    n_jobs=-1,
    tree_method="hist",
    eval_metric="logloss",
)
base_model.fit(X_train, y_train)

# Get features
importances = base_model.feature_importances_
feat_imp = pd.DataFrame({
    "feature": X.columns,
    "importance": importances
}).sort_values("importance", ascending=False)
```

```
print("\nTop 10 features:")
print(feat_imp.head(10))

y_pred = base_model.predict(X_test)
y_prob = base_model.predict_proba(X_test)[:, 1]

# Metrics
acc = accuracy_score(y_test, y_pred)
prec = precision_score(y_test, y_pred)
rec = recall_score(y_test, y_pred)
f1 = f1_score(y_test, y_pred)
auc = roc_auc_score(y_test, y_prob)

print("\nModel metrics")
print(f"Accuracy: {acc:.4f}")
print(f"Precision: {prec:.4f}")
print(f"Recall: {rec:.4f}")
print(f"F1 Score: {f1:.4f}")
print(f"ROC AUC: {auc:.4f}")

# Classification report
print("\nClassification report:")
print(classification_report(y_test, y_pred, digits=3))

# Confusion Matrix
cm = confusion_matrix(y_test, y_pred)
plt.figure(figsize=(5, 4))
sns.heatmap(cm, annot=True, fmt="d", cmap="Blues", cbar=False)
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Confusion matrix")
plt.tight_layout()
plt.show()

# ROC Curve
fpr, tpr, thresholds = roc_curve(y_test, y_prob)

plt.figure(figsize=(6, 5))
plt.plot(fpr, tpr, color="blue", label=f"ROC Curve (AUC = {auc:.3f})")
plt.plot([0, 1], [0, 1], linestyle="--", color="gray", label="Random Guess")
plt.xlabel("False positive rate")
plt.ylabel("True positive rate")
plt.title("Receiver operating characteristic (ROC) curve")
plt.legend(loc="lower right")
plt.tight_layout()
plt.show()
```

Top 10 features:

	feature	importance
58	PEOE_VSA8	0.031478
104	NOCount	0.029348
62	SMR_VSA3	0.029012
108	NumAmideBonds	0.024193
30	Chi0v	0.023395
128	fr_Al_COO	0.017744
78	SlogP_VSA8	0.017709
39	Chi4v	0.016871
1	MaxEStateIndex	0.016812
92	VSA_EState10	0.016808

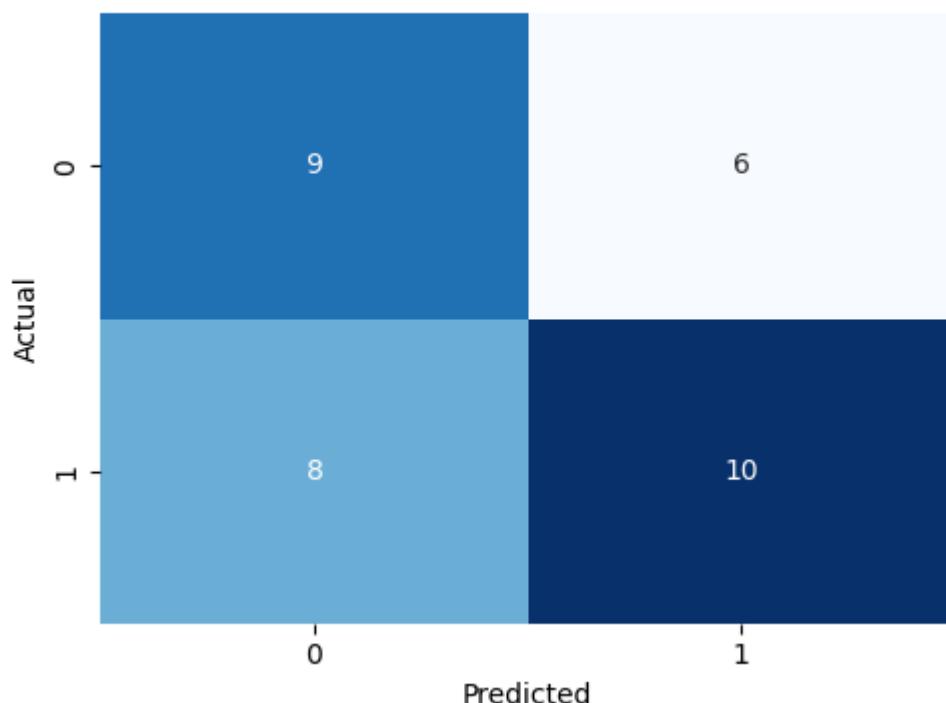
Model metrics

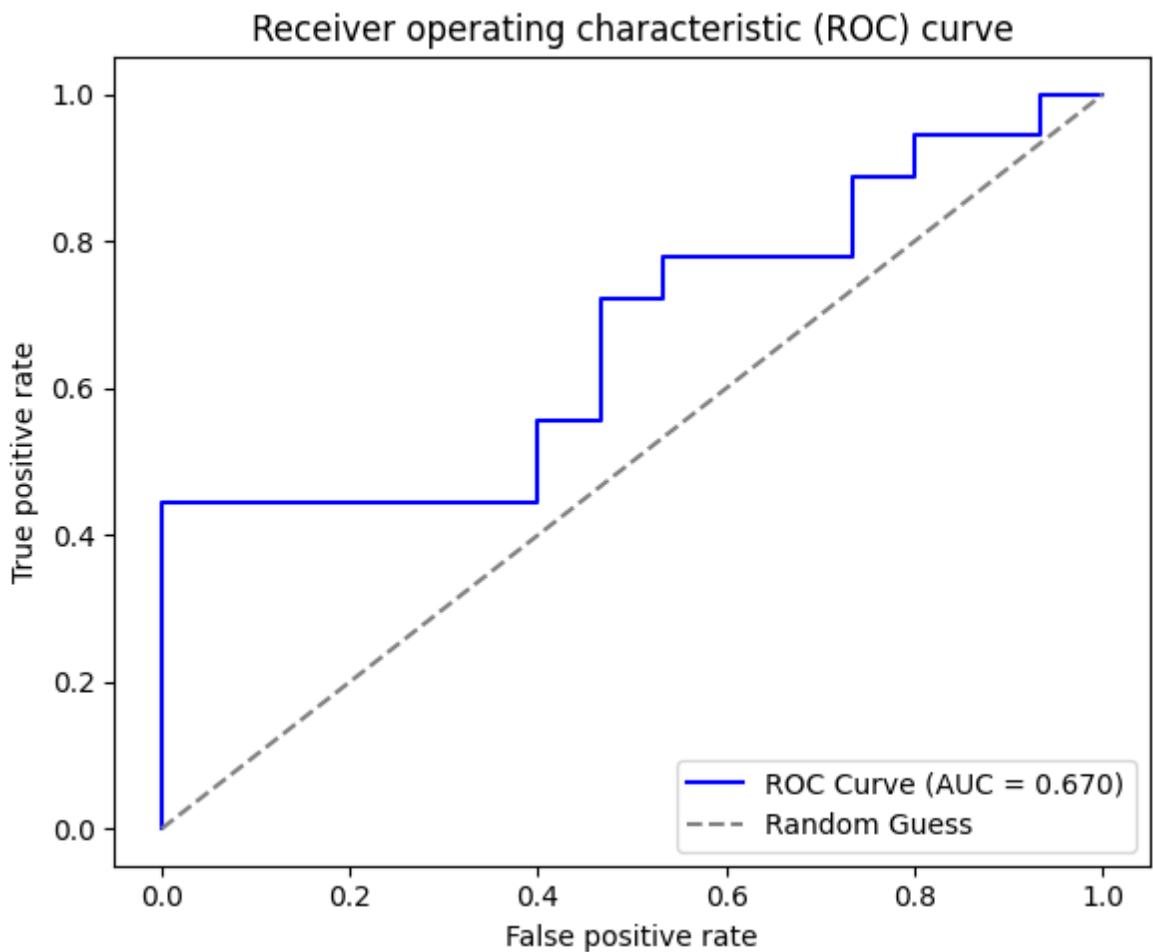
Accuracy: 0.5758
Precision: 0.6250
Recall: 0.5556
F1 Score: 0.5882
ROC AUC: 0.6704

Classification report:

	precision	recall	f1-score	support
0	0.529	0.600	0.562	15
1	0.625	0.556	0.588	18
accuracy			0.576	33
macro avg	0.577	0.578	0.575	33
weighted avg	0.582	0.576	0.577	33

Confusion matrix





```
In [68]: results = []

for n in range(5, 120, 1): # test top 5-119 features
    top_feats = feat_imp["feature"].head(n).tolist()

    model = xgb.XGBClassifier(
        n_estimators=400,
        learning_rate=0.05,
        max_depth=6,
        subsample=0.8,
        colsample_bytree=0.8,
        random_state=42,
        n_jobs=-1,
        tree_method="hist",
        eval_metric="logloss",
    )
    model.fit(X_train[top_feats], y_train)

    y_pred = model.predict(X_test[top_feats])
    y_prob = model.predict_proba(X_test[top_feats])[:, 1]

    acc = accuracy_score(y_test, y_pred)
    f1 = f1_score(y_test, y_pred)
    auc = roc_auc_score(y_test, y_prob)

    results.append((n, acc, f1, auc))

# Summary
res_df = pd.DataFrame(results, columns=["Top_N", "Accuracy", "F1", "ROC_A"])
```

```

# Find best by F1 score
best = res_df.iloc[res_df["F1"].idxmax()]
best_n = int(best.Top_N)

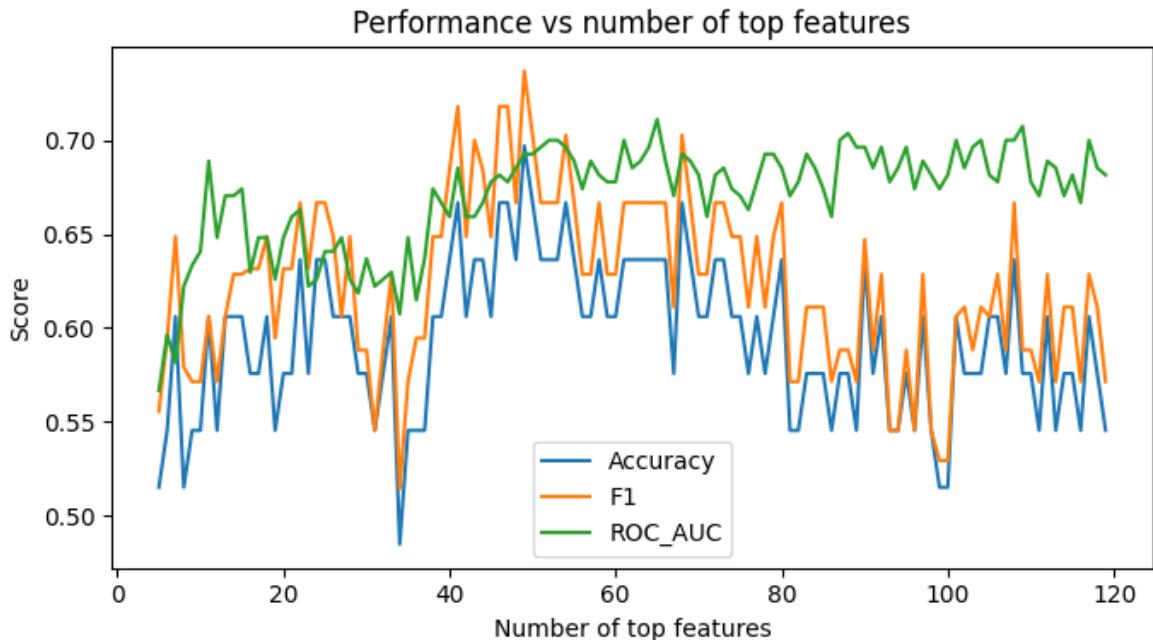
# Save the best top features
best_feats = feat_imp.head(best_n)
best_feats.to_csv(TOP_FEATURES_CSV, index=False)
print(f"Saved top {best_n} features -> {TOP_FEATURES_CSV}")

# All in one
plt.figure(figsize=(7,4))
plt.plot(res_df["Top_N"], res_df["Accuracy"], label="Accuracy")
plt.plot(res_df["Top_N"], res_df["F1"], label="F1")
plt.plot(res_df["Top_N"], res_df["ROC_AUC"], label="ROC_AUC")
plt.xlabel("Number of top features")
plt.ylabel("Score")
plt.title("Performance vs number of top features")
plt.legend()
plt.tight_layout()
plt.show()

print("\nBest number of features:", best_n)
print(best)

```

Saved top 49 features -> top_features_best.csv



Best number of features: 49
 Top_N 49.000000
 Accuracy 0.696970
 F1 0.736842
 ROC_AUC 0.692593
 Name: 44, dtype: float64

In [69]: # Best pick
 top_feats = feat_imp["feature"].head(best_n).tolist()
 final_model = xgb.XGBClassifier(
 n_estimators=400,
 learning_rate=0.05,
 max_depth=6,
 subsample=0.8,
 colsample_bytree=0.8,
 random_state=42,

```
n_jobs=-1,  
tree_method="hist",  
eval_metric="logloss",  
)  
final_model.fit(X_train[top_feats], y_train)  
  
# Predictions for ROC  
y_prob_best = final_model.predict_proba(X_test[top_feats])[:, 1]  
fpr, tpr, thresholds = roc_curve(y_test, y_prob_best)  
auc_best = roc_auc_score(y_test, y_prob_best)  
  
# Plot ROC  
plt.figure(figsize=(6,5))  
plt.plot(fpr, tpr, color="blue", label=f"Best Model (Top {best_n} Feats)  
plt.plot([0,1],[0,1],"--",color="gray",label="Random Guess")  
plt.xlabel("False positive Rate")  
plt.ylabel("True positive Rate")  
plt.title("ROC curve for best feature count")  
plt.legend(loc="lower right")  
plt.tight_layout()  
plt.show()
```

