

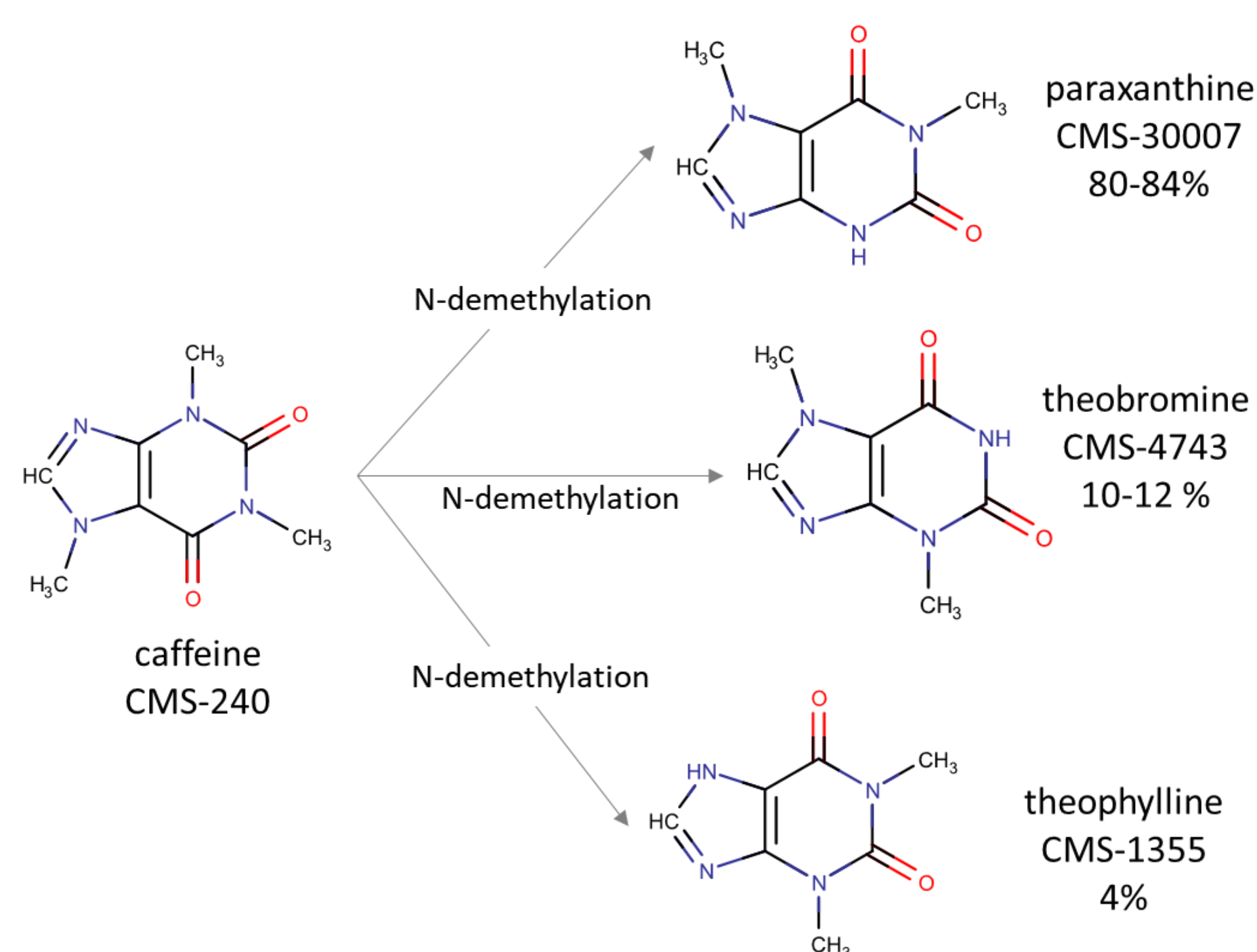
MOTIVATION

- ❖ Read-across approaches for parents and metabolites are considered from a mixture perspective.
- ❖ Compilation of evidence to support read-across includes Absorption, Distribution, Metabolism, and Excretion (ADME) profiles and biological assays, in addition to the conventional structure and molecular/physicochemical properties.
- ❖ Biological activities relevant to the toxicity of interest are captured by either assay activities or in silico methods such as QSAR or ML-hybrid structure rules.
- ❖ Various metabolites with estimated abundance are combined to give a quantitative read-across outcome with NOAEL bounds and uncertainties.



METABOLISM

Partial Metabolic Pathway of Caffeine: Human Oral



CONCLUSION

- ❖ Systemic/target organ NOAEL bounds were estimated to be 13-47 mg/kg-bw/day (at 95% confidence) for caffeine by read-across from three metabolites treated as a mixture.
- ❖ Analogue selection used structure, molecular properties, ADME properties, ER/AR assays, and ML-hybrid rules to quantify similarity to target.
- ❖ ML-hybrid rules allow for estimation of assay-based similarities when assay data for structures of interested is limited or unavailable. [Yang et al. submitted to Chem. Res. Toxicol. 2023]



ANALOG IDENTIFICATION, CHARACTERIZATION & ANALOG QUALITY (AQ)

$$AQ = \sqrt{\prod_{i=1}^N (SimilarityMeasure)_i}$$

ToxPrints, MACCS Keys, RDKit Molecular fingerprints (Tanimoto coefficients)

Rotational bonds, H-ACC, H-DON, Lipinski rule violations, MW, Complexity, TPSA, logP, Asphericity, Eccentricity.

Caco-2 Papp (cm/h), eFlux ratio, CLint_hepatic, Fraction bound, BBB, Oral intestinal abs %, Skin perm (logKp)

AR, AR BLA, AR LUC, ER, ERa, ERb, ERaERa, ERa BLA, ERb BLA, ERa LUC, ERa TRANS, Ahr, PR, PPAR

ML-Hybrid Rule-based Similarity for ER/AR activities

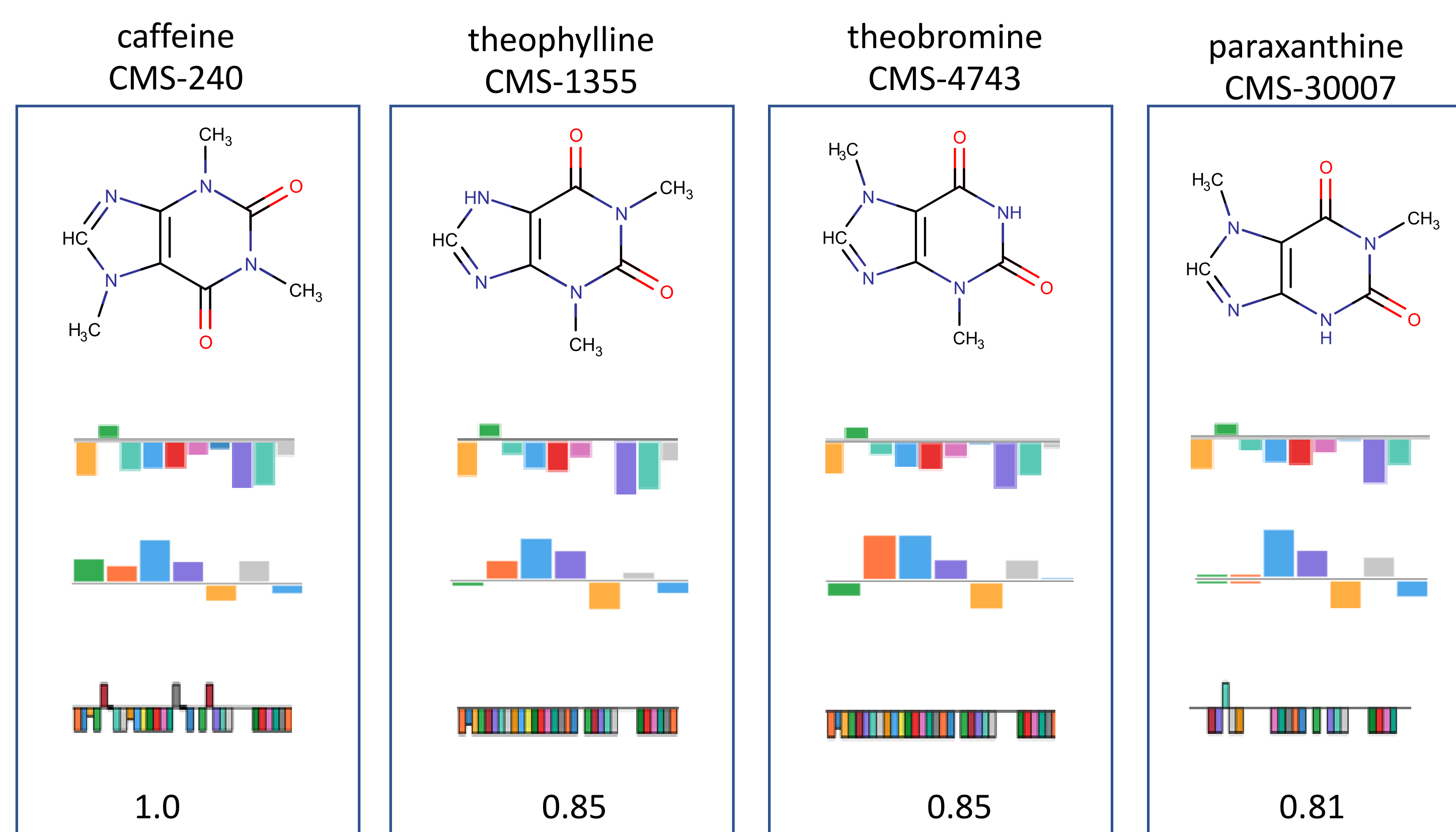
Structures (ToxPrints)

Molecular Properties

ADME Properties

ER/AR Activity

Rule Similarity



MACHINE LEARNING (ML) – HYBRID RULES

Benefits of ML-Hybrid Rules

- ❖ Rules can be easily developed for a particular set of inventories or datasets with specific chemical classes
- ❖ ML-Hybrid rules can be developed for both positive and negative outcomes against the endpoints.

ER/AR Rules based on ToxCast/Tox21 Assays

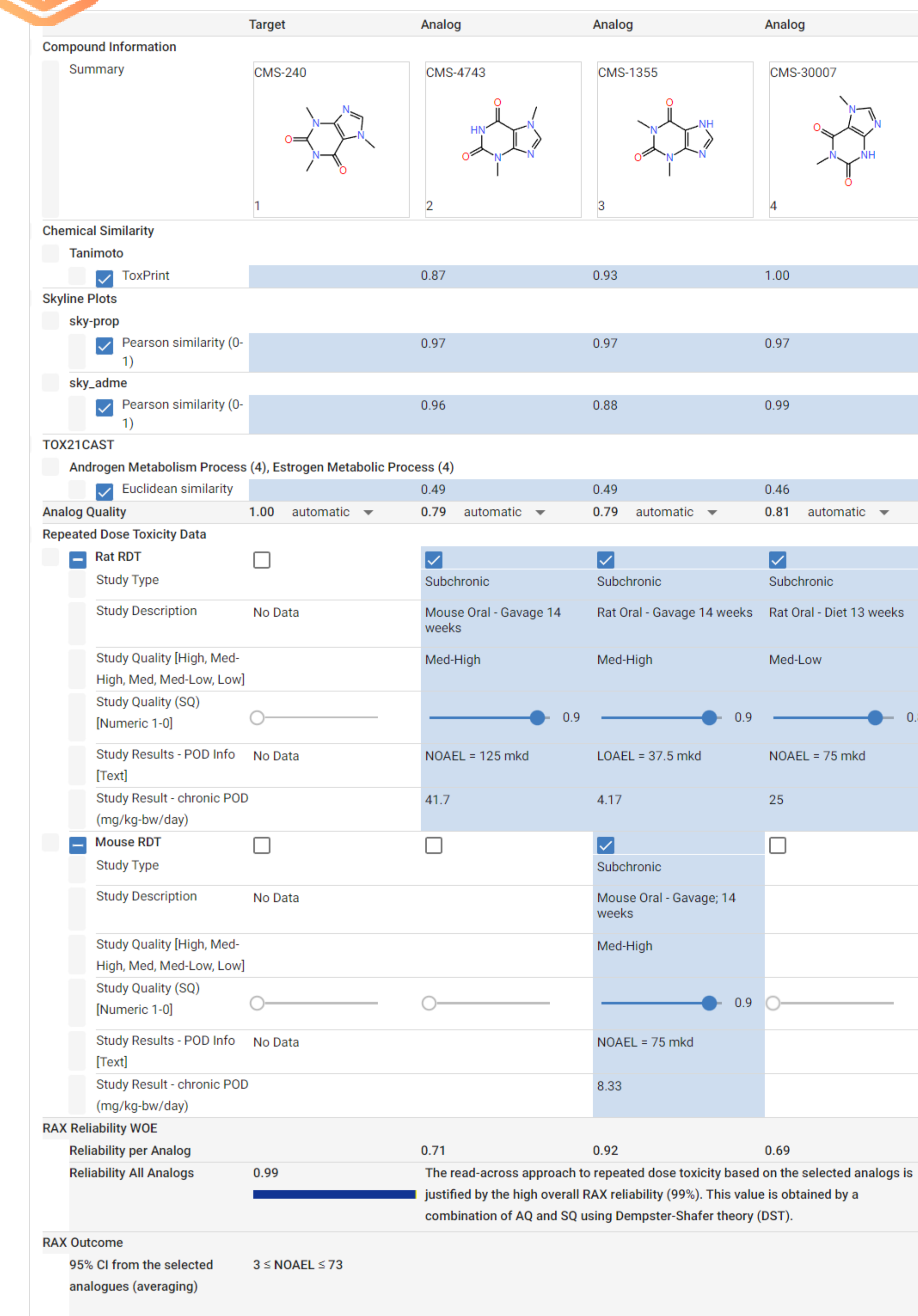
- ❖ In vitro dataset contains 8593 compounds with more than 1400 assays
- ❖ Grouped by 24 modes of action and 43 mechanistic targets
- ❖ AR and ER assays were aggregated according to the MOA groups [Yang et al. Comp. Toxicol. 2023]
- ❖ In this study, 103 structures contain nucleobase, 23 structures are related to caffeine (purinedione class)
- ❖ Other 20 ML-Hybrid rules were also developed for ER/AR activities. [Yang et al. Chem. Res. Toxicol. 2023]

Rule-based Similarity

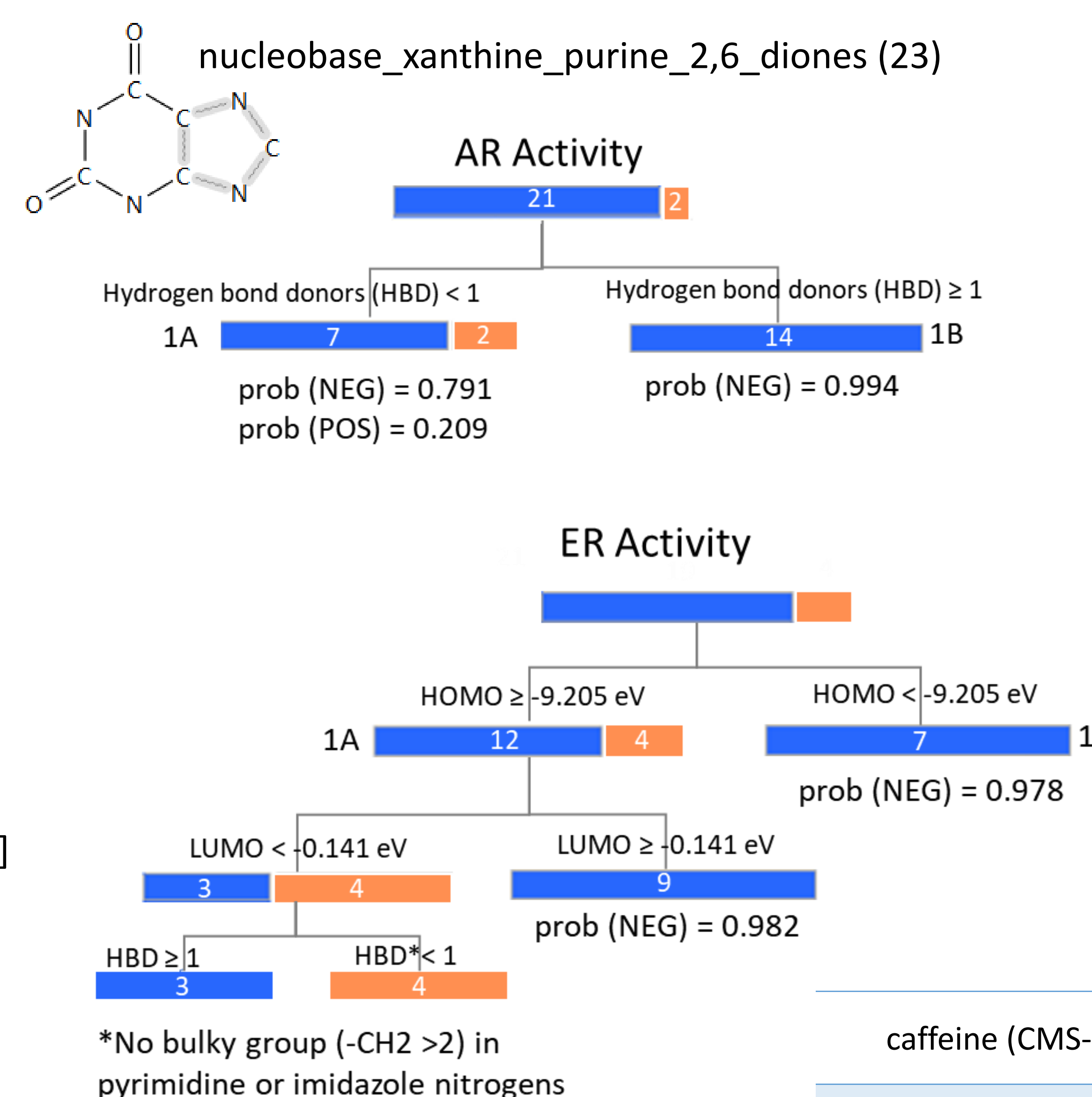
- ❖ ML-hybrid rules were also matched with structures. Each structure was represented by a vector of Bayes Factors, from which pairwise Euclidean similarities were calculated to be used in read-across. [Yang et al. Chem. Res. Toxicol. 2023]



ASSESSMENT SUMMARY REPORT



Recursive Partitioning Tree



*No bulky group (-CH₂ >2) in pyrimidine or imidazole nitrogens
HOMO: Highest Occupied Molecular Orbital Energy (eV)
LUMO: Lowest Occupied Molecular Orbital Energy (eV)

	Target	Relative amount	chronic NOAEL (mg/kg-bw/day)
caffeine (CMS-240)	Target		
theophylline (CMS-1355)	Analog-1	4%	4.17, 8.33
theobromine (CMS-4743)	Analog-2	10-12%	41.7
paraxanthine (CMS-30007)	Analog-3	80-84%	25

Weighted 95% CI NOAEL:
13.3 – 47.4 mg/kg-bw/day
3 – 73 mg/kg-bw/day (without weighting by abundance)