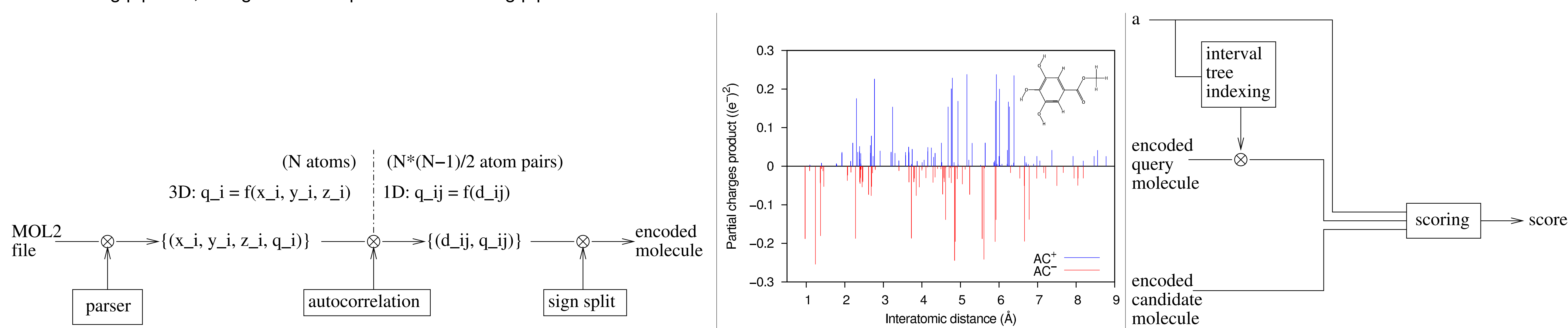


## METHOD

Overview of ACPC's encoding pipeline, along with its output and the scoring pipeline<sup>[1,4]</sup>. ACPC is written in OCaml<sup>[7]</sup>.

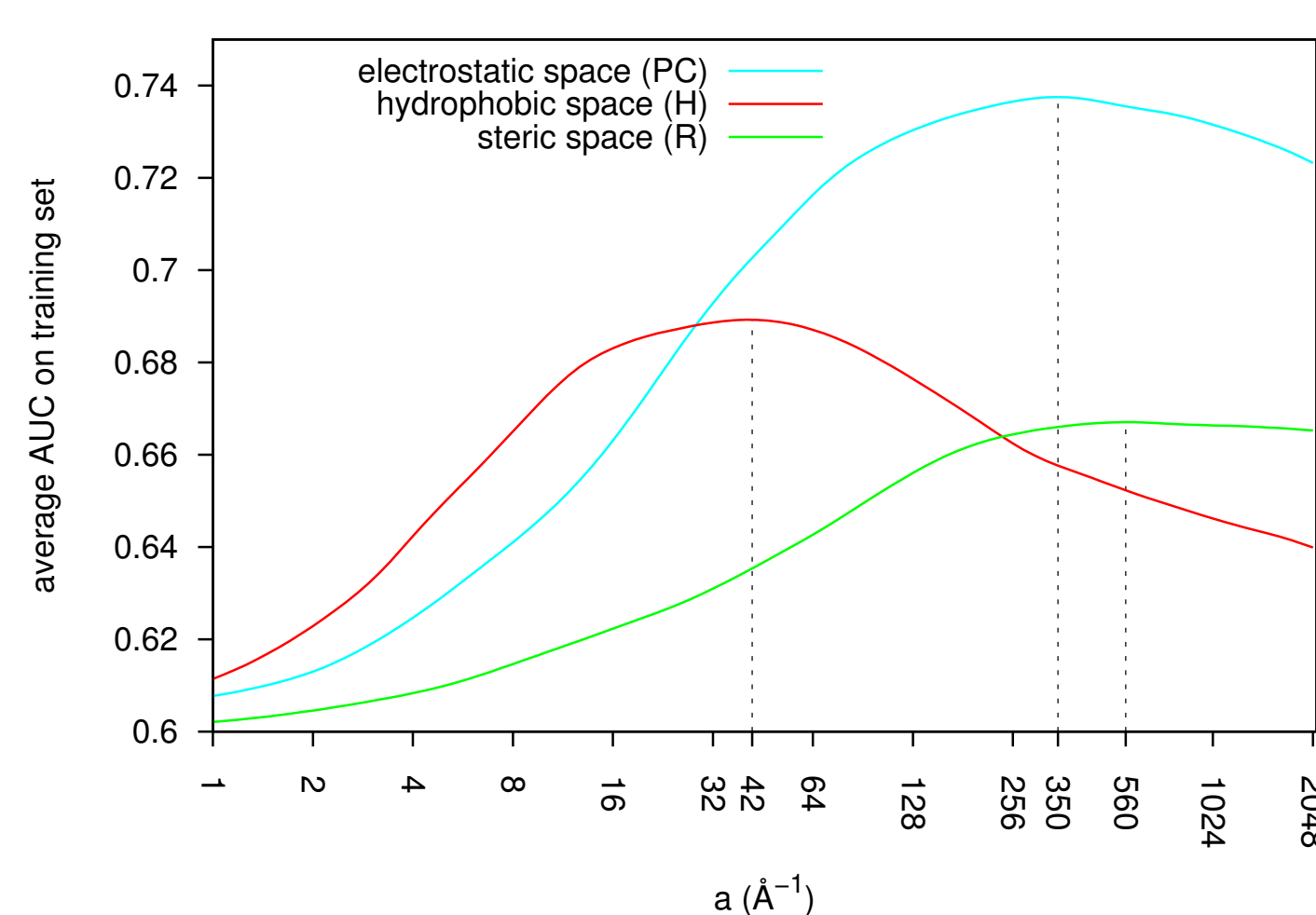


**Figure 1** : Left: encoding of a molecule. Middle: positive (in blue) and negative (in red) values of the autocorrelation of partial charges for the ligand molecule shown in 2D. Right: scoring of two encoded molecules (one query molecule and a candidate molecule from a database).  $\frac{1}{a}$  is the half-base length of a triangular kernel function used to transform the set of Kronecker deltas into a continuous function.

## TRAINING AND TEST DATASETS

DUDE<sup>[5]</sup> contains 102 targets, 25660 ligands and 1060858 decoys. Preparation protocol: i) Open Babel<sup>[6]</sup> 2.3.9 unique InChI filtering ii) OMEGA<sup>[3]</sup> 2.4.6 lowest energy conformer iii) partial charges assigned with MMFF94x of MOE 2013.08 iv) molecules renamed to follow the scheme [active]InChIKey[TheInChIKey]. Training set; "DUDE-training" = 51 targets chosen randomly and only the 1<sup>st</sup> ligand of each target used as a query. Test set; "DUDE-test" = all 51 remaining targets with all their ligands.

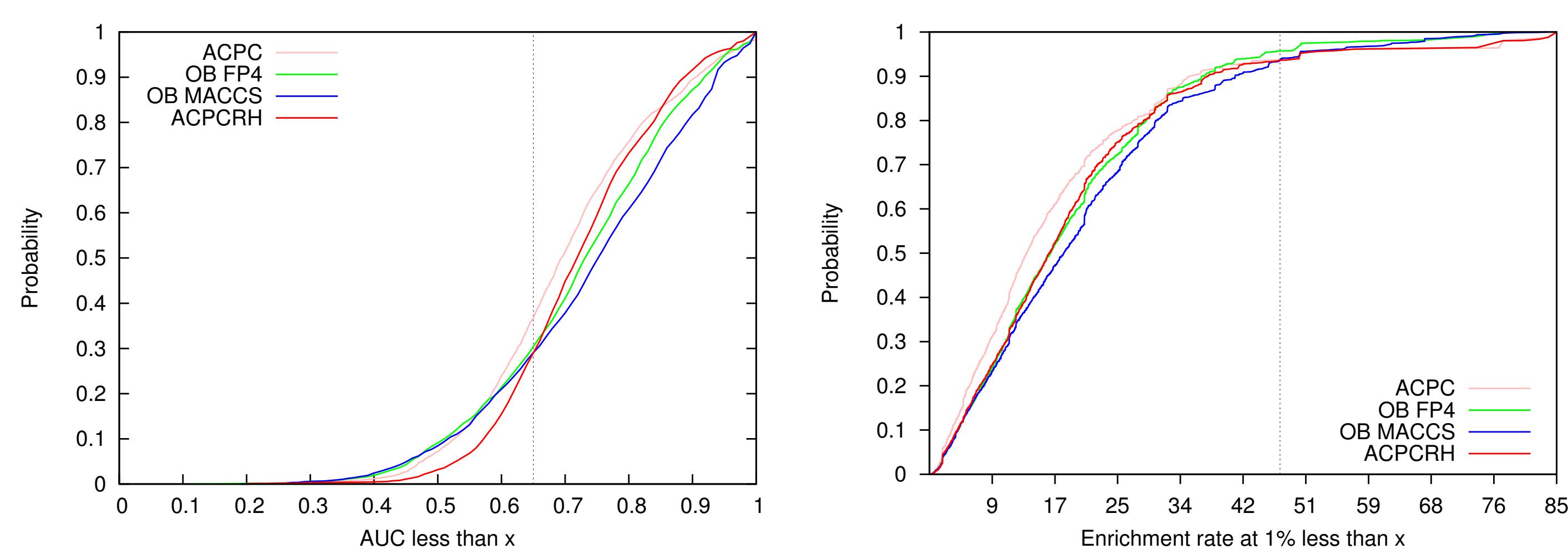
## PARAMETER SETUP



**Figure 2** : Using the training set, an optimal value for the  $a$  parameter was found for each feature space (dotted lines).

On the training set, scores from each feature space were also centered then scaled. Then, a combination of weights was found to improve the average AUC and ER at 1% reached by ACPC. The weights are 0.8 for the electrostatic space, 0.2 for the steric space and 0.5 for the hydrophobic space.

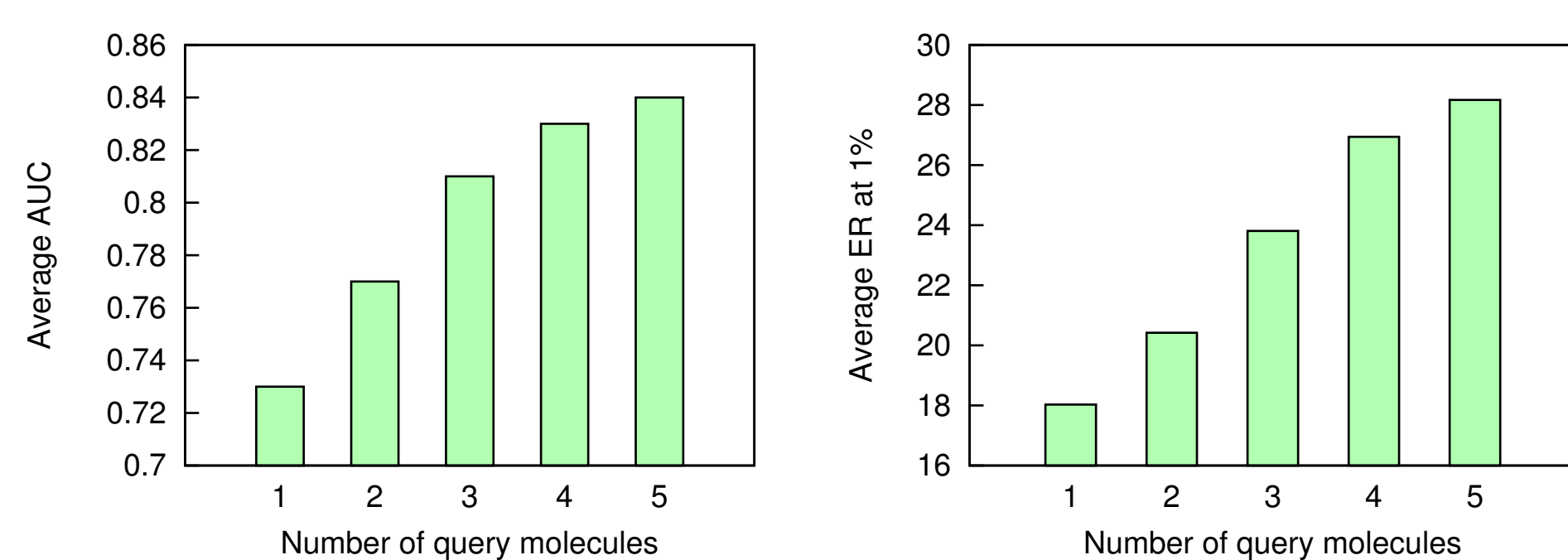
## VIRTUAL SCREENING PERFORMANCE



**Figure 3** : Left: AUCs' cumulative distribution functions for ACPC in the electrostatic feature space, Open Babel's (OB) FP4 and MACCS implementations and ACPCRH using three feature spaces (ACPCRH) on DUDE-test. Right: enrichment rates at 1% in the same experiment.

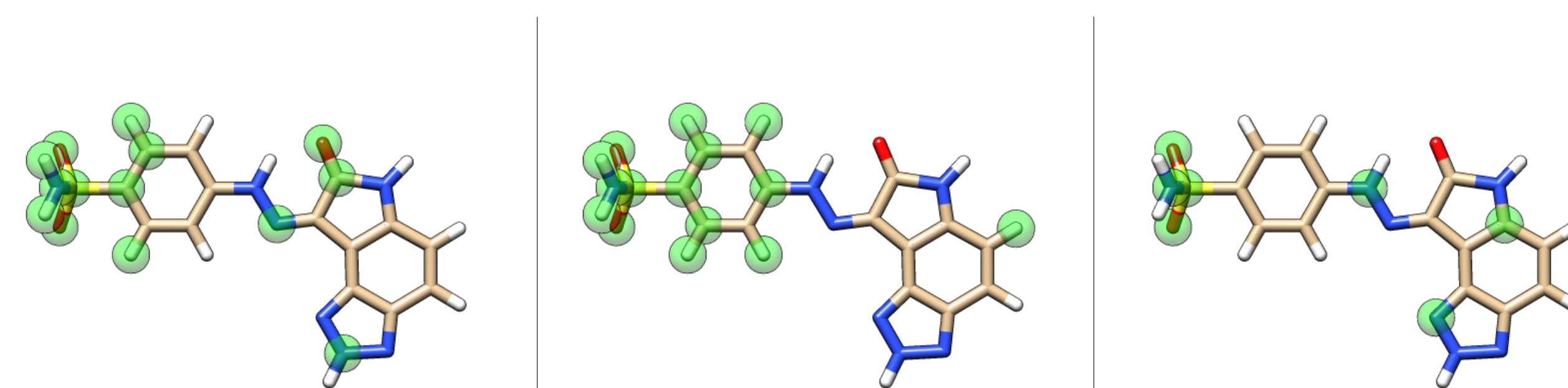
## CONSENSUS QUERIES

Five query molecules per target were chosen randomly on DUDE-training. Then, the min rank was used to rank-order molecules when using up to five query molecules for consensus in the electrostatic feature space.



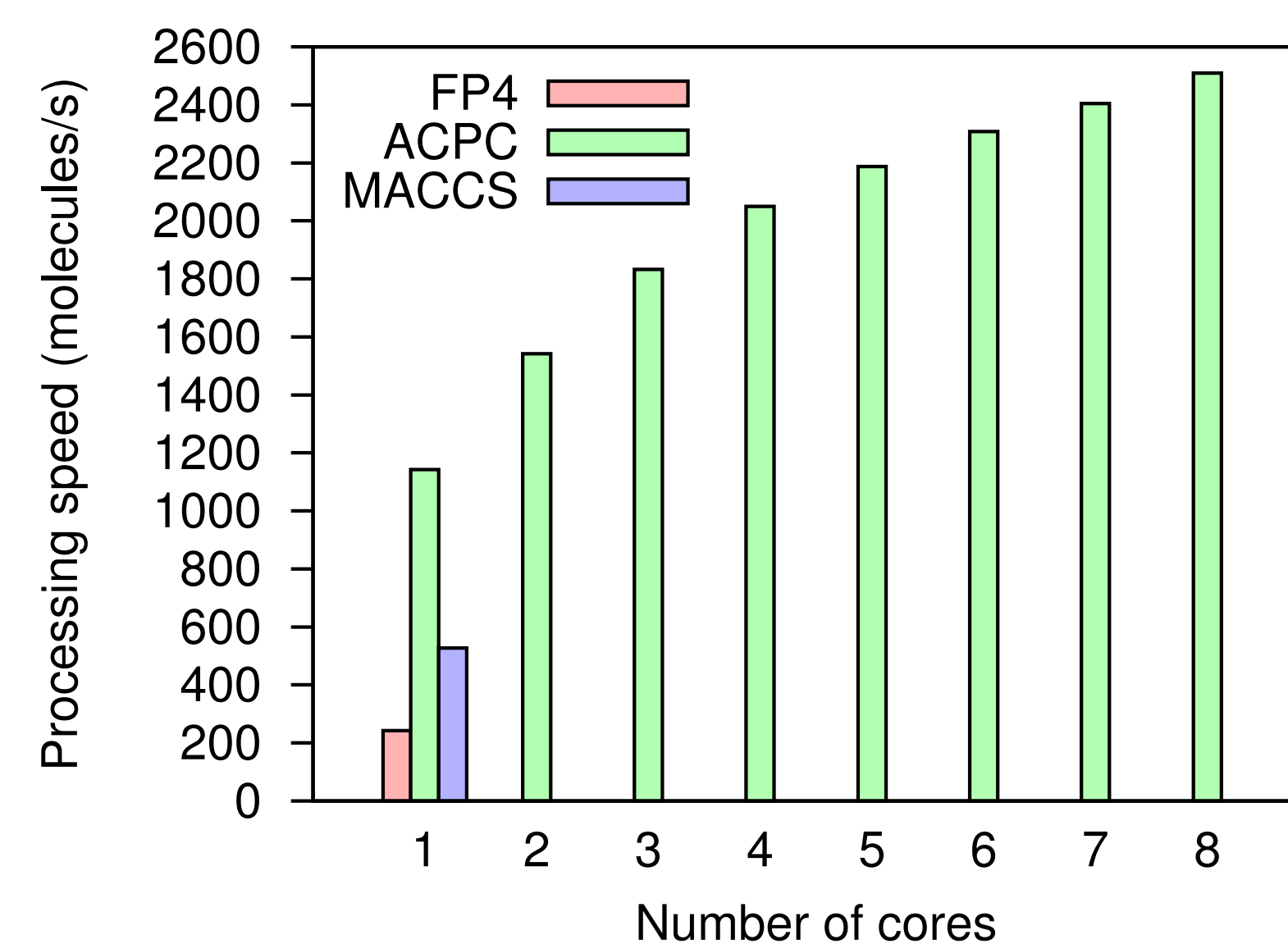
**Figure 4** : Left: average AUC as a function of the number of query molecules used during consensus. Right: average enrichment rate at 1% in the same experiment.

## QUERY MOLECULE ANNOTATION



**Figure 5** : In any feature space, the query molecule can be annotated based on atomic contribution to AUC (Left: electrostatic space, Middle: steric space, Right: hydrophobic space). Here, the 1<sup>st</sup> ligand of the cdk2 target has green balls highlighting atoms which if removed would decrease the AUC reached by the query molecule in the given feature space.

## SPEED



**Figure 6** : Speed comparison between ACPC in the electrostatic feature space (Parmap<sup>[2]</sup> is used for parallelization) and Open Babel 2.3.9's MACCS and FP4 implementations. For each method, runtimes were averaged over three runs after a warmup run. The target was hivpr from DUDE (26450 ligands and decoys).

## INSTALL ACPC

- 1 - Install and configure OPAM (cf. <http://opam.ocaml.org/>)
- 2 - Type: 'opam install acpc.1.2'

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