

Slide 9: Fialuridine (FIAU)

Initial Task: DILI

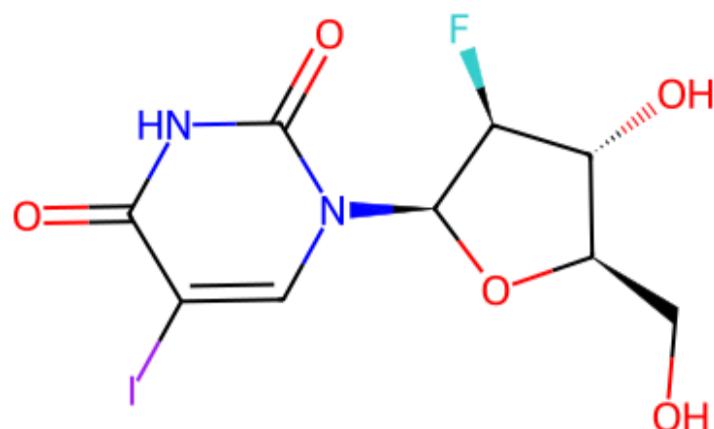
SMILES: C1=C(C(=O)NC(=O)N1[C@H]2[C@H](C[C@H]([C@@H](CO)O2)O)F)I

Fialuridine (FIAU) - Optimization Results

Optimization Path

 Overall Comparison - Initial → Final

Initial Molecule `C1=C(C(=O)NC(=O)N1[C@H]2[C@H]
([C@@H])([C@@H](CO)O2)O)F`

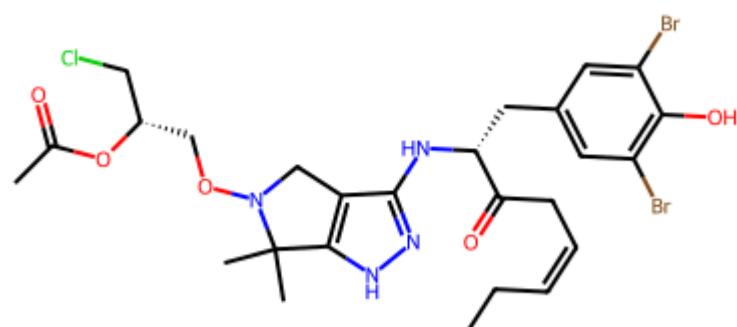


QED (Drug-likeness): 0.5621 **Number of Blocks:** 1

► Show ADMET Scores

Task	Score
AMES	0.114162
BBBP	1.000000
CYP3A4	0.000000
DILI	0.145744
HIA	1.000000
PGP	0.940474

Final Optimized `CC/C=C\CC(=O)[C@@H]
(Cc1cc(Br)c(O)c(Br)c1)Nc1n[nH]c2c1CN(OC[C@H](CCl)OC(C)=O)C2(C)C`



QED (Drug-likeness): 0.1398 (-0.4222) ✗ **Number of Blocks:** 3 (+2) ↑
Total Block Changes: 10

► Show ADMET Scores

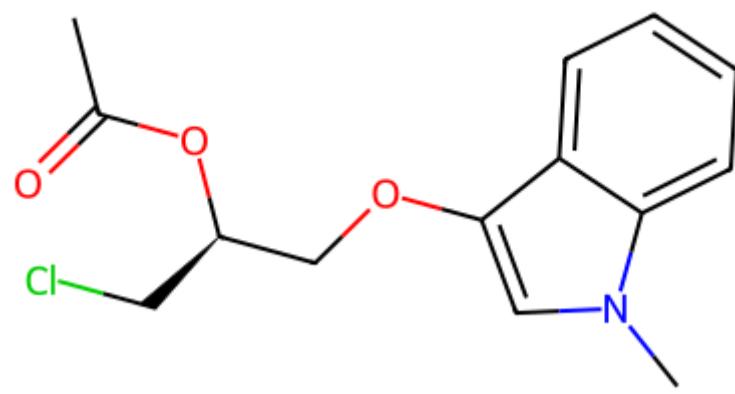
Task	Score	Change	Rel. Improvement	% Change
AMES ✗	0.266793	+0.152631	-1.3370	+133.70%
BBBP ✗	1.000000	+0.000000	+0.0000	+0.00%
CYP3A4 ✗	0.000067	+0.000067	-5806.1212	+580612.12%
DILI ✗	0.259323	+0.113579	-0.7793	+77.93%
HIA ✗	0.999987	-0.000013	-0.0000	-0.00%
PGP ✓	0.866124	-0.074351	+0.0791	-7.91%

Optimization Steps:

DETAILS PLACEHOLDER2

After (Step 1)

CC(=O)O[C@H](CCl)C0c1cn(C)c2ccccc12



QED: 0.6249 (+0.0629)

Number of Blocks: 2 (+1) ↑

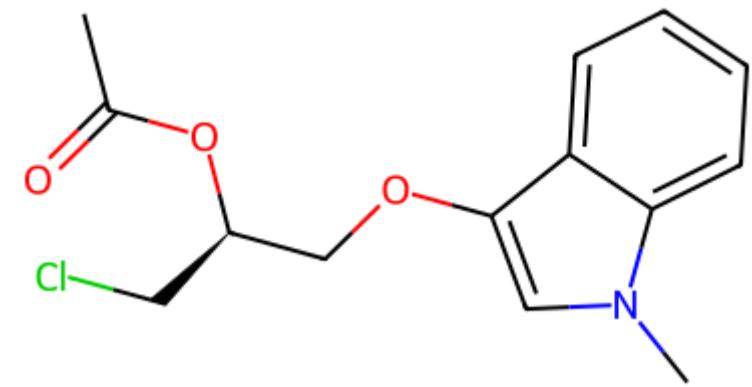
Block Changes: 3 (+2, -1)

DILI Score: 0.145744 → 0.641619 (+0.495875)

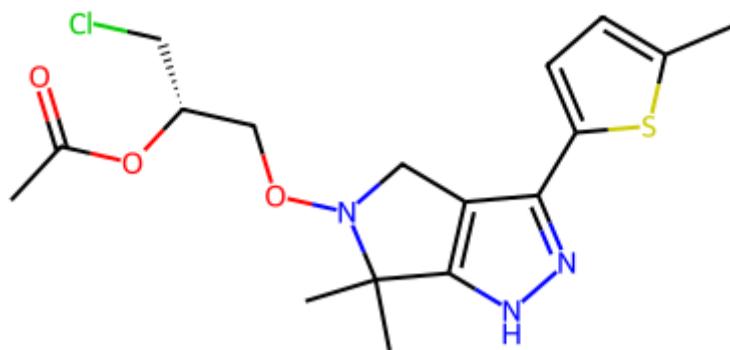
DETAILS *PLACEHOLDER3*

► Step 2: CYP3A4 (-0.1500 ↓)

****Before (Step 1)**** `CC(=O)O[C@H](CCl)COc1cn(C)c2ccccc12`



****After (Step 2)**** `CC(=O)O[C@H](CCl)CON1Cc2c(-c3ccc(C)s3)n[nH]c2C1(C)C`



****QED:**** 0.6249 ****Number of Blocks:**** 2

► All ADMET Scores

Task	Score	Direction
AMES	0.376149	↓ lower
BBBP	0.999958	↑ higher
CYP3A4	0.159087	↓ lower
DILI	0.641619	↓ lower
HIA	0.976337	↑ higher
PGP	0.594567	↓ lower

QED: 0.6094 (-0.0155) X

Number of Blocks: 2 (+0) ➡

Block Changes: 4 (+2, -2)

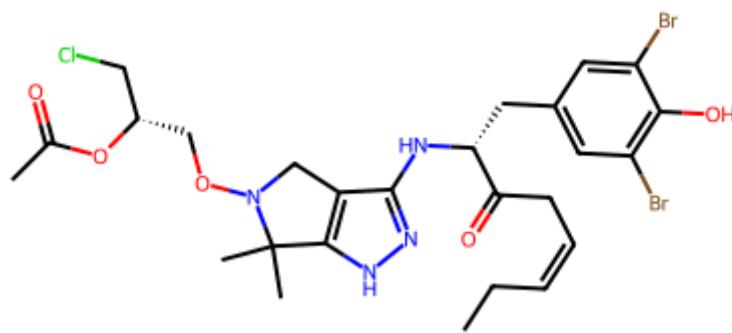
CYP3A4 Score: 0.159087 → 0.009106 (-0.149981)

DETAILSPLACEHOLDER5

DETAILSPLACEHOLDER6

After (Step 3)

CC/C=C\CC(=O)[C@@H](CCl)C(O)C(Br)C1Nc2n[nH]c2C1CN(OC[C@@H](CCl)OC(C)=O)C2(C)C



QED: 0.1398 (-0.4696) X

Number of Blocks: 3 (+1) ↑

Block Changes: 3 (+2, -1)

DILI Score: 0.576163 → 0.259323 (-0.316840)

DETAILS PLACEHOLDER7

Step Details

Step 1: DILI ⚠

Original	New	Change
0.145744	0.641619	+0.495875 ↓

CC(=O)O[C@H](CCl)COC1CN(C)C2CCCCC12

Step 2: CYP3A4 ✓

Original	New	Change
0.159087	0.009106	-0.149981 ↓

CC(=O)O[C@H](CCl)CON1Cc2c(-c3ccc(C)s3)n[nH]c2C1(C)C

Step 3: DILI ✓

Original	New	Change
0.576163	0.259323	-0.316840 ↓

CC/C=C\CC(=O)[C@@H](CC1CC(Br)C(O)C(Br)C1)Nc1n[nH]c2c1CN(OC[C@@H](CCl)OC(C)=O)C2(C)C

ADMET Comparison

Task	Direction	Initial	Final	Change	Rel. Improvement	% Change	Status
AMES	↓ lower	0.114162	0.266793	+0.152631	-1.3370	+133.70%	X Declined
BBBP	↑ higher	1.000000	1.000000	+0.000000	+0.0000	+0.00%	→ Unchanged
CYP3A4	↓ lower	0.000000	0.000067	+0.000067	-5806.1212	+580612.12%	X Declined
DILI	↓ lower	0.145744	0.259323	+0.113579	-0.7793	+77.93%	X Declined
HIA	↑ higher	1.000000	0.999987	-0.000013	-0.0000	-0.00%	X Declined
PGP	↓ lower	0.940474	0.866124	-0.074351	+0.0791	-7.91%	✓ Improved

Improved: 1/6 (16.7%) | Molecules: 271 | Paths: 19870

🔍 Safety Threshold Analysis

Status: 5/6 meet thresholds

⚠ Below threshold: 1

Task	Score	Threshold	Gap
PGP	0.8661	↓ 0.3	0.5661

✓ Passing: 5

Task	Score	Threshold
AMES	0.2668	↓ 0.3
BBBP	1.0000	↑ 0.5
CYP3A4	0.0001	↓ 0.55
DILI	0.2593	↓ 0.4
HIA	1.0000	↑ 0.2