The Boots Company synthesis of ibuprofen (1960's)

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 \begin{array}{lll} & \text{CC}(C) \, \text{Cc1ccccc1.CC}(=0) \, \text{DC}(C) = 0 > \text{[A1]} \, \text{(C1)} \, \text{(C1)} > \text{CC}(=0) \, \text{c1ccc} \, \text{(CC}(C) \, \text{C)} \, \text{cc1} \\ & \text{CC}(=0) \, \text{c1ccc} \, \text{(CC}(C) \, \text{C)} \, \text{cc1.} \, \text{[Na+].CC} \, \text{[0-].CCOC}(=0) \, \text{CC1} > \text{CCOC}(=0) \, \text{C10C1} \, \text{(C)} \, \text{c2ccc} \, \text{(CC}(C) \, \text{C)} \, \text{cc2} \\ & \text{CCOC}(=0) \, \text{C10C1} \, \text{(C)} \, \text{c2ccc} \, \text{(CC}(C) \, \text{C)} \, \text{cc2.} \, \text{[DH3+]} > \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=0)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=0)} \, \text{cc1.} \, \text{NO} > \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} > \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C\#N)} \, \text{cc1.} \, \text{\{2\}0} > \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C(=0))} \, \text{oc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{C=NO)} \, \text{cc1} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{(C(C)} \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{(C(C)} \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{CC)} \, \text{Cc1ccc} \, \text{(C(C)} \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{(C(C)} \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{CC}(C) \, \text{Cc1ccc} \, \text{CC}(C) \, \text{Cc1ccc} \, \text{CC)} \\ & \text{CC}(C) \, \text{Cc1ccc} \, \text{CC}(C) \,
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```
#use triple quotes (""") to define a multiline string
reactions_smiles = """CC(C)Cc1ccccc1.CC(=0)OC(C)=0>A1(C1)C1>CC(=0)c1ccc(CC(C)C)cc1
CC(=0)c1ccc(CC(C)C)cc1.[Na+].CC[0-].CCOC(=0)CC1>>CCOC(=0)C10C1(C)c2ccc(CC(C)C)cc2
CCOC(=0)C10C1(C)c2ccc(CC(C)C)cc2.[OH3+]>>CC(C)Cc1ccc(C(C)C=0)cc1
CC(C)Cc1ccc(C(C)C=0)cc1.NO>>CC(C)Cc1ccc(C(C)C=N0)cc1
CC(C)Cc1ccc(C(C)C=N0)cc1>>CC(C)Cc1ccc(C(C)C#N)cc1
CC(C)Cc1ccc(C(C)C#N)cc1.{2}O>>CC(C)Cc1ccc(C(C)C=0)0)cc1"""
atom_economy(reactions_smiles)
```

REACTANTS SMILES: CC(C)Cc1ccccc1 Molecular Formula: C10H14 Molecular Weight: 134.22 g/mol Coefficient: 1.0 SMILES: CC(=0)OC(C)=0

Molecular Formula: C4H6O3
Molecular Weight: 102.09 g/mol

Coefficient: 1.0

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SMILES: [Na+] Molecular Formula: Na+

Molecular Weight: 22.99 g/mol

Coefficient: 1.0

SMILES: CC[0-]
Molecular Formula: C2H50-

Molecular Weight: 45.06 g/mol

Coefficient: 1.0

SMILES: CCOC(=0)CC1
Molecular Formula: C4H7ClO2
Molecular Weight: 122.55 g/mol

Coefficient: 1.0

SMILES: [OH3+]
Molecular Formula: H3O+

Molecular Weight: 19.02 g/mol

Coefficient: 1.0

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SMILES: NO Molecular Formula: H3NO

Molecular Weight: 33.03 g/mol

Coefficient: 1.0

SMILES: 0
Molecular Formula: H20

Molecular Weight: 18.02 g/mol

Coefficient: 2.0

PRODUCTS

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SMILES: CC(C)Cc1ccc(C(C)C(=0)0)cc1

Molecular Formula: C13H1802 Molecular Weight: 206.28 g/mol

Coefficient: 1.0

Atom Economy: 40.1%

Calculate atom economy for individual synthesis steps

```
from rxnSMILES4AtomEco import get_atom_economy
# value = get_atom_economy(reactions_smiles)
# print(value)
# Split reactions_smiles into lines
lines = reactions_smiles.splitlines()
# Create a dictionary to store the results
results_individual = {}
# Loop through each line and call get_atom_economy for each one
for i, line in enumerate(lines):
    # Create Step 1, Step 2, ..., Step n
    step_key = f"Step {i+1}"
    # Call the function and store the result
    results_individual[step_key] = get_atom_economy(line)
# Now, 'results' contains the results for each line, \
# e.g., results["Step 1"], results["Step 2"], etc.
# Example: Print all results
for step, result_individual in results_individual.items():
    print(f"{step}: {result_individual:.1f}%")
```

```
Step 1: 74.6%
Step 2: 71.5%
Step 3: 67.6%
Step 4: 91.9%
Step 5: 91.2%
Step 6: 92.4%
```

Calculate atom economy for cumulative synthesis steps

```
# Split reactions_smiles into lines
lines = reactions_smiles.splitlines()

# Create a dictionary to store the results
```

```
results_cumulative = {}
# Loop through each incremental combination of lines and call get_atom_economy
# Start from 1 to n
for i in range(1, len(lines) + 1):
    # Get the first i lines
    combined_reactions = "\n".join(lines[:i])
    # Call get_atom_economy with the combined lines up to the current step
    step_key = f"Step_{i}"
   # Store the result
   results_cumulative[step_key] = get_atom_economy(combined_reactions)
    # Example: Print the current step's result
    print(f"{step_key}: {results_cumulative[step_key]:.1f}%")
Step_1: 74.6%
Step_2: 61.5%
Step_3: 42.7%
Step_4: 42.9%
Step_5: 39.1%
Step_6: 40.1%
import matplotlib.pyplot as plt
# Extract the steps (x-axis) and Atom Economy values (y-axis) \
# for both individual and cumulative
# Step names: ['step_1', 'step_2', ...]
steps = list(results_individual.keys())
# Atom Economy values for individual reactions
individual_values = list(results_individual.values())
# Atom Economy values for cumulative reactions
cumulative_values = list(results_cumulative.values())
# Calculate the Byproduct(s) as the difference between 100% \
# and the desired product
byproduct_values = [100 - value for value in individual_values]
# Create a single plot
```

```
# Single axis for both plots
fig, ax = plt.subplots(figsize=(10, 7))
plt.rcParams.update({'font.size': 16})
# Plot the stem plot for cumulative Atom Economy values in black
ax.stem(steps, cumulative_values, linefmt='k:', markerfmt='k>:', basefmt=" ", \
        label="Cumulative Atom Economy").markerline.set_markersize(10)
# Bar plot for individual Atom Economy values
# Green bars
bars = ax.bar(steps, individual_values, color='#009E73', \
              label="Desired product", alpha=0.8)
# Orange stacked bars
ax.bar(steps, byproduct_values, bottom=individual_values, color='#E69F00', \
       label="Byproduct(s)", alpha=0.8)
# Add the text (value of desired product) in white in the middle of each bar
for i, bar in enumerate(bars):
   height = bar.get_height()
    ax.text(bar.get_x() + bar.get_width() / 2, height / 2, f'{height:.1f}%',
             ha='center', va='center', color='white', fontweight='bold')
# Add labels and titles
ax.set_title('Boots Atom Economy: Individual- and Cumulative Steps\n\n\n')
ax.set_xlabel('\nSteps')
ax.set_ylabel('Atom Economy (%)')
# Set Y-axis from 0 to 100%
ax.set_ylim(0, 100)
# if used, uncomment for better readability \
#ax.tick_params(axis='x', rotation=45) # Rotate x-axis labels
# Add the legend
ax.legend(loc='upper center', bbox_to_anchor=(0.5, 1.2), ncol=3,reverse=True)
# Improve layout to avoid label overlap
plt.tight_layout()
# Show the plot
plt.show()
# Save the plot
# To save the plot, if used, comment line above and uncomment lines below
```

```
#save_path = '/tmp/Boots_histogram.png' # Writable on many systems
#plt.draw() # Force render
#print(f"Saving to: {save_path}")
#plt.savefig(save_path, dpi=300, bbox_inches='tight')
#plt.close()
#print("Save completed (check directory).")
```

Boots Atom Economy: Individual- and Cumulative Steps

