

Introduction

Stylus is a Cadence tool designed to help designer define a proper CAD tool flow from behavioral to GDSII

The **flowtool** is the core function for

- Automatically generating a set of scripts, with PLACE_HOLDERS, for designers to configure with design and foundry references
- Executing the tool flow, with many knobs for controlling which elements are run, and for inspecting and debugging the sequence of Cadence tools that are run

Cadence tools include:

- Genus: Behavioral synthesis tool
- Innovus: Place&Route tool
- Tempus: Static timing analysis tool
- Voltus: Full-chip electromigration, IR drop and power analysis tool
- Quantus: Parasitic extraction tool
- Conformal: Formal verification via equivalence checking tool

FlowTool Definitions

Flows are defined with

- **Flow steps (*flow_step*):** Associate a set of (tcl) commands to a label

Flow steps are created with *create_flow_step*

- **Flow objects:** Flow objects specify a set of actions for a CAD tool to execute
Typically, one of the Cadence tools are referred with '-tool genus'
Non-Cadence tools are allowed using the '-tool_options' switch

- **Flow scheduling:** Adds additional flow actions to the current flow

The keyword *schedule_flow* is used inside a *create_flow_step* definition

Stylus supports generating a generic flow environment with *write_flow_template*

write_flow_template -list gives a list of templates that can be generated for configuring new flows for a design and/or foundry

The command *write_flow_template* MUST be run within a Cadence UI

It generates YAML files, e.g., *flow.yaml* and TCL files, e.g., *design_config.tcl*

FlowTool Components

You MUST run the command within Cadence *genus*

```
@genus:root: 1> write_flow_template -tools "genus" -describe stylus
Features for flow templates matching 'stylus'
```

```
Template stylus
```

```
-----
Provider      : tool
Max version   : 1
Description   : Standard flow for block implementation defined using YAML and TCL
This is the default template
```

Feature	Description	Default	Valid
--- the following features are mutually exclusive (dft_style group)			
dft_compressor	Add flow support for scan chains with compression insertion		0 1 {}
dft_simple	Add flow support for scan chain insertion		0 1 {}

dynamic_view	single dynamic analysis_view to activate		
ff_setup	Enable reading design config from legacy FF flow setup.tcl file		0 1 {}
flow_express	Enable express synthesis and implementation flow		0 1 {}
hold_views	list of hold analysis_views to activate		
leakage_view	single leakage analysis_view to activate		
report_clp	Add CLP dofile generation and checks to the flow		0 1 {}
--- the following features are mutually exclusive (report_style group)			
report_defer	Defer report generation		0 1 {}
report_inline	Run report generation as part of parent flow versus schedule_flow		0 1 {}
report_none	Disable report generation		0 1 {}

report_lec	Add LEC dofile generation and checks to the flow		0 1 {}
setup_views	list of setup analysis_views to activate		
--- the following features are mutually exclusive (synth_style group)			
synth_hybrid	Physically aware synthesis flow with logical final optimization		0 1 {}
synth_ispatial	Physically aware synthesis flow with ispatial final optimization		0 1 {}
synth_physical	Full physically aware synthesis flow		0 1 {}
synth_spatial	Physically aware synthesis flow with spatial final optimization		0 1 {}

use_common_db	Enable using common DB format for synth and implementation flows		0 1 {}

YAML

YAML is a standard string processing language for describing data serialization, i.e., a series of data processing tasks, using high-level, user-friendly constructs

It describes the flow process in an **outline** style format

- Comments are indicated with '#'
- Key-value pairs are created using ':'
- Collections of actions are created using '-'

Note that YAML uses spaces to create dependencies among statements, like python

Stylus creates flow control extensions in YAML using:

- **CMD_<NAME>**: Automatically encapsulates content in a flow_step with NAME
CMD_write_def: write_def out.def
- **FILE_<NAME>**: Automatically encapsulates file contents in a flow_step via NAME
FILE_floorplan: floorplan.tcl
- **SCHEDULE**: Method to schedule new flows in YAML, similar to schedule_flow
SCHEDULE: -flow report_prechts

YAML

Portion of *flow.yaml* in scripts directory generated by *write_flow_template*

```
flows:
#-----
# synthesis
#-----
  synthesis:
    args: -tool genus -owner cadence -skip_metric -tool_options -disable_user_startup
    features:
    steps:
      - syn_generic:
        args: -owner cadence
        features:
        steps:
          - block_start:
          - init_elaborate:
          - init_design:
            args: -owner cadence
            features:
            steps:
              - read_mmmc:
              - read_physical:
              - read_hdl:
              - read_power_intent:
              - run_init_design:
              - read_def:
                enabled: "synth_spatial || synth_ispatial || synth_physical || synth_hybrid"
          - init_genus:
          - set_dont_use:
          - init_dft:
          - commit_power_intent:
          - create_cost_group:
          - run_syn_generic:
          - block_finish:
          - SCHEDULE:
            args: -flow report_synth -include_in_metrics
            enabled: "!report_none && !report_inline && !report_defer"
      - syn_map:
        ...
```

YAML

Note that **syn_generic** and **syn_map** are *genus* commands

The items '-' are collections of actions to execute for these *genus* commands

The scripts file, *setup.yaml*, *flow_config.tcl* and *design_config.tcl* contain most of the **PLACEHOLDER** keywords that need to be filled in for a new design/foundry

In *setup.yaml*, the following are needed:

- **library_sets**: List of *xxx.lib* files
- **opconds**: Process, temperature, voltage files
- **timing_conditions**: *library_set* objects to associate to a *timing_condition*
- **rc_corners**: *qrc_tech* files for RC corners
- **delay_corners**: *early* and *late* RC corner and *timing_condition* names
- **constraint_modes**: Constraint files to associate with a *constraint_mode*
- **analysis_views**: name of *constraint_mode* and *delay_corner* + others for view
- **lef_files/oa_ref_libs/oa_search_libs**: Files for *read_physical* command
- **init_power_nets/init_ground_nets** + others: Net names to assign
- **design_name/design_process_node/design_flow_effort**, etc: Design attributes
- **add_fillers_cells/add_tieoffs_cells**, etc: Names of std cells

FlowTool Commands

Validate the *setup.yaml* by running flowtool, but only for *init_design*

flowtool -predict summary -flow init_design

flowtool takes many arguments:

- **flowtool -run_tag synthesis_only -to syn_opt.block_finish**
- **flowtool -from prects.block_start -to prects.block_finish -predict verbose**

From within the Cadence UIs, you can run flows:

- **run_flow -step init_design**

To run the entire flow, just use:

- **flowtool**

Viewing the results:

reports directory: Run your browser and point it to the *qor.html*

reports directory: For *genus*, text reports in *syn_generic*, *syn_map* and *syn_opt*

json: Use *write_metric -format json -file run1.json*

logs directory: Use *vim* to view, look for 'errors' and 'warnings'