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CubeFit — LOSVD-convolved stellar population fitting for IFU cubes

CubeFit builds a **hypercube of convolved template spectra** for every (`spaxel S`, component `C`, population `P`) and solves for a global, non-negative mixture that best explains the observed IFU data cube. It is designed for large datasets ($10^3\text{--}10^5$ spaxels, $10^2\text{--}10^3$ populations), is robust to restarts, and uses HDF5 end-to-end with streaming, tile-aligned I/O.

- **Hypercube builder:** FFT-based LOSVD convolution on the template grid, followed by flux-conserving rebin to the observed grid.
- **Normalization:** choose “model” (LOSVD amplitude) or “data” (per-spaxel observed flux with LOSVD-proportional splits). Convert between modes later without a rebuild.
- **Line emphasis:** optional λ -weights so absorption features drive the fit.
- **Solver:** multiprocess Kaczmarz with diagonal preconditioning, trust region, backtracking, global step caps, near-zero column freezing, optional de-correlation nudge, and a tile-local **weighted NNLS polish**.
- **Tracking:** streaming fit metrics and optional snapshots to a SWMR-friendly sidecar.
- **Utilities:** chunk-aware normalization conversion; global column energy; reconstruction and plotting helpers that never touch the giant `/HyperCube/models` unless needed.

HDF5 layout (what lives where)

Core inputs:

- `/Templates` — (`P`, `T`) template spectra on the template grid (time domain).
- `/TemPix` — (`T`, `)` template grid in $\log\lambda$ (natural log).
- `/ObsPix` — (`L`, `)` observed wavelength grid ($\log\lambda$).

- $/R_T - (T, L)$ (or (L, T)) **flux-conserving** linear rebin operator mapping the template grid to the observed grid.
- $/LOSVD - (S, V, C)$ LOSVD histograms (per spaxel and component).
- $/VelPix - (V,)$ velocity grid in km/s (for LOSVD).

Optional inputs:

- $/Mask - (L,)$ boolean wavelength mask; used consistently by builder and solver.
- $/HyperCube/data_flux - (S,)$ masked mean data flux per spaxel (required for $\text{norm} = \text{"data"}$).

Built artifacts:

- $/HyperCube/models - (S, C, P, L)$ float32 convolved+rebinned spectra, chunked for streaming and resumable via a `_done` bitmap.
- $/HyperCube/col_energy - (C, P)$ float64 global column energy $E[c, p] = \sum_{\lambda} s_{\lambda} A^2$, for step-size control.
- $/HyperCube/norm/losvd_amp - (S, C)$ LOSVD amplitude (sum or trapz).
- $/HyperCube/norm/losvd_amp_sum - (S,)$ per-spaxel sum of amplitudes.
- $/HyperCube/lambda_weights - (L,)$ optional λ -weights in $[\text{floor}, 1]$ (generated by a median-DoG heuristic).

Fit outputs:

- $/X_global - (C*P,)$ solution vector, row-major (C, P) flattened.
-

Manager: creating and populating the HDF5 backbone

`hdf5_manager.py` provides:

- **Safe file open** with retries, optional SWMR, and lock-handling (`open_h5`).
- **Dataset creation** for the core arrays and dimensions (`H5Manager + H5Dims`).
- **Population** from NumPy arrays (`populate_from_arrays`), including grid checks.
- **Flux-conserving rebin** operator construction (`ensure_rebin_and_resample`).

Typical setup:

```
from CubeFit.hdf5_manager import H5Manager, H5Dims

mgr = H5Manager("galaxy.h5", tem_pix=tem_loglam, obs_pix=obs_loglam)
mgr.init_base(H5Dims(nSpat=S, nLSpec=L, nTSpec=T, nVel=V, nComp=C, nPop=P))
mgr.populate_from_arrays(losvd=H_SVC, datacube=Y_SL, templates=T_PT)
mgr.ensure_rebin_and_resample() # builds /R_T (and validates shapes)
```

Notes:

- $/R_T$ is built by exact bin overlap in log- λ , preserving flux.
 - $/Mask$ (if present) is used later by both builder and solver.
 - Manager helpers centralize chunk-cache setup so downstream is fast.
-

Hypercube builder

`hypercube_builder.py` constructs `/HyperCube/models` in streaming, tile-aligned passes:

1. **Kernel from LOSVD:** each (s, c) LOSVD row \square **unit-area** kernel on the template grid (linear interpolation to integer pixel shifts). A separate scalar **amplitude** is computed per (s, c) (either sum or trapezoidal integral).
2. **Convolution (per p):** FFT-multiply template by kernel and crop the linear convolution back to the template length.
3. **Rebin to observed grid:** multiply by $/R_T$ to get (P, L) for each (s, c) .
4. **Normalization:**
 - `norm="model"`: multiply each $(s, c, p, :)$ by its LOSVD amplitude.
 - `norm="data"`: compute per-spaxel mean observed flux (from `/HyperCube/data_flux`), then split across components in proportion to their LOSVD amplitudes, preserving the (c) ratios determined by LOSVD at that spaxel.
5. **Global column energy:** simultaneously accumulate $E[c, p] = \sum_s \sum_p$ in `mask` A^2 into `/HyperCube/col_energy` for solver preconditioning.

Resumability & chunking: the builder creates `/HyperCube/models` with chunks (`S_chunk`, `C_chunk`, `P_chunk`, `L`) and maintains a `_done` bitmap over the (`S_chunk`, `C_chunk`, `P_chunk`) tile grid; this lets you resume safely or inspect progress mid-build. Metadata is flushed regularly.

Normalization conversion (post-hoc)

Flip between `norm="data"` and `norm="model"` after the build:

```
from CubeFit.hypercube_builder import convert_hypercube_norm
convert_hypercube_norm("galaxy.h5", to_mode="model", recompute_energy=True)
```

This reads `/HyperCube/norm/losvd_amp_sum` (and `/HyperCube/data_flux` when needed), computes a **per-spaxel scalar** $F[s]$, and scales `/HyperCube/models[s, :, :, :] *= F[s]` in **S-tiles** (chunk-aligned). It updates `/HyperCube/col_energy` (optional) and flips the `/HyperCube` attribute `norm.mode`. It also respects `/Mask` when deriving or validating the spaxel flux vector.

This is a safe, **no-FFT** operation. Use it to avoid rebuilding the hypercube when only a normalization flip is required.

λ -feature emphasis (line weights)

Absorption features can be down-weighted in plain least squares. To emphasize them, the solver supports **λ -weights**:

- Build weights once via a **median spectrum** and a two-scale smoothing **difference-of-Gaussians** (implemented with boxcars), then map to $[min_w, 1]$. Store at `/HyperCube/lambda_weights`.
- During solving, multiply both **residuals** and **design matrix columns** by \sqrt{w} — equivalent to solving in a diagonal metric where line pixels matter more.

Helpers in `cube_utils.py`:

```
from CubeFit.cube_utils import ensure_lambda_weights, read_lambda_weights

w = ensure_lambda_weights("galaxy.h5") # writes /HyperCube/lambda_weights
w2 = read_lambda_weights("galaxy.h5") # reads, with floor & mask handling
```

At solver startup you'll see a banner printing min/max/mean of the λ -weights used (after masking).

Solver: multiprocess batched Kaczmarz (+ weighted NNLS polish)

Entry point:

```
from CubeFit.pipeline_runner import PipelineRunner

runner = PipelineRunner("galaxy.h5")
x_global, stats = runner.solve_all_mp_batched(
    epochs=E,
    pixels_per_aperture=...,                      # row batching (if used)
    lr=...,                                         # base learning-rate
    project_nonneg=True,                           # x >= 0
    processes=..., blas_threads=...,                # MP and BLAS knobs
    orbit_weights=...,                            # optional (C,) ratio guidance
    # ...other optional arguments...
)
```

High-level flow (`pipeline_runner.py` → `kaczmarz_solver_cchunk_mp.py`):

1. **Tile scheduling:** sort S-tiles by data norm (coarse “hardest first” ordering).
2. **Workers:** each process takes a contiguous **c-band** for the active S-tile.
3. **Inside a worker (per band):**
 - **Sanitize** non-finite values in A and R to zeros.
 - **Apply λ -weights** by computing gradients and denominators on $\|w\| \cdot A$ and $\|w\| \cdot R$ (but updates and residuals are returned in the unweighted space).
 - **Diagonal preconditioning:** per-population step $dx_p \leftarrow g_p / \|A_w\|^2$, blended with global column energy $E[c, p]$ for stability.
 - **Near-zero column freeze** (relative & absolute thresholds).
 - **Trust region** on ΔR in weighted space ($\|R_w\| \leq \epsilon \|R\|$) + **backtracking** for monotone weighted RMSE drop.
 - **Optional de-correlation nudge** to avoid chasing flat null-space directions across P.
 - **Candidate set for tile-local weighted NNLS polish** (top-K by score).
4. **Parent process:**
 - Aggregates ΔR over bands and runs **tile-level backtracking**.
 - Applies a **global step cap** using $\|dx\|^2 \cdot E[c, p]$ vs. $\|Y\|$.
 - Optionally runs **NNLS polish** on the proposed (c,p) set using a λ -subsample (rows) and accepts only if it reduces **weighted RMSE** enough.
 - Commits the single global α for the tile: update R and x, enforce $x \geq 0$.
 - Optional **ratio penalty** step nudges the component totals toward **orbit_weights** (if provided).
 - Streams metrics to **FitTracker** and snapshots (optional).

The result is written to `/X_global` (flattened ($C \times P$)), along with run stats.

Reconstruction & plots (without storing a full ModelCube)

You can reconstruct $\hat{Y} = A \cdot x$ for diagnostics without materializing a full /ModelCube:

- **Reconstruction utilities** read A in the same chunk order and contract against x in λ -bands. They write a small target array you name (e.g., `/ReconTile`) or return the result.
- **Parallel spectral plots** pull just the selected rows from `/DataCube` and the reconstructed array and produce side-by-side data/model spectral plots for worst/best spaxels.

Both are tile-aligned and cache-aware; they never read /HyperCube/models beyond the needed slices.

Fit tracking & snapshots

`fit_tracker.py` provides a **non-blocking** sidecar writer process:

- Destination: a sidecar H5 next to the main file (`*.fit.<pid>.<ts>.h5`).
- Bounded `mp.Queue` so the solver never blocks on I/O.
- SWMR enabled for dashboards.
- Streams **RMSE history**, **EWMA**, **progress**, and optional **x snapshots** (best/last/history).

Use `tracker=FitTracker(...)` in the solver; the runner wires this up for you.

Utilities

- **Normalization conversion:** `convert_hypertable_norm(h5, to_mode="model"|"data", recompute_energy=True)` — chunk-aligned, mask-aware.
 - **Global column energy:** created during build; can be recomputed later if you changed normalization.
 - **Done-bitmap control:** functions to inspect and invalidate `_done` to resume or redo specific tiles.
 - **Recompression:** rewrite /HyperCube/models with different compression/filter options (tile-wise copy).
-

Environment variables (knobs)

These can be exported to tune performance, stability, or behavior. Defaults shown are what the code uses if you don't set them.

HDF5 & dataset caching

Variable	Default	Purpose
<code>HDF5_USE_FILE_LOCKING</code>	<code>FALSE</code>	Avoid HDF5 file locking issues on shared filesystems.
<code>CUBEFIT_RDCC_NSLOTS</code>	<code>400003</code>	File-level raw chunk cache: number of slots.
<code>CUBEFIT_RDCC_NBYTES</code>	<code>4294967296</code> (4 GiB)	File-level raw chunk cache size in bytes.
<code>CUBEFIT_RDCC_W0</code>	<code>0.9</code>	File-level raw chunk cache preemption policy.
<code>CUBEFIT_RDCC_SLOTS</code>	<code>1000003</code>	Per-dataset cache slots used by some readers.
<code>CUBEFIT_RDCC_BYTES</code>	<code>268435456</code> (256 MiB)	Per-dataset cache size used by some readers.

Multiprocessing & BLAS/OpenMP

Variable	Default	Purpose
<code>CUBEFIT_MP_CTX</code>	<code>forkserver</code> (fallback <code>spawn</code>)	MP start method for the solver pool.
<code>OMP_NUM_THREADS</code>	set by code	BLAS/OpenMP threads per worker (also OPENBLAS_NUM_THREADS)
<code>KMP_INIT_AT_FORK</code>	<code>FALSE</code>	Avoid OpenMP deadlocks with <code>fork</code> .

Variable	Default	Purpose
SLURM_CPUS_PER_TASK	(if present)	Used by reconstruction helpers to set BLAS threads.
PYTHONUNBUFFERED	1 (recommended)	Unbuffered stdout for real-time logs on HPC.

Solver: worker-level stability & steps (in `kaczmarz_solver_cchunk_mp.py`)

Variable	Default	Purpose
CUBEFIT_BT_STEPS	3	Backtracking steps per band (worker) for monotone weighted RMSE.
CUBEFIT_BT_FACTOR	0.5	Multiplicative α shrink in backtracking.
CUBEFIT_TRUST_TAU	0.7	Trust-region cap: ' τ '.
CUBEFIT_EPS	1e-12	Numerical floor in denominators and divisions.
CUBEFIT_ZERO_COL_REL	1e-12	Relative threshold (vs median energy) to freeze near-zero columns.
CUBEFIT_ZERO_COL_ABS	1e-24	Absolute threshold to freeze columns.
CUBEFIT_DEBUG_SAFE	0	Print sanitization/freeze stats per tile if 1.
CUBEFIT_NNLS_PROP_PER_BAND	6	How many (c,p) candidates to propose for NNLS polish per band.

Solver: aggregation & NNLS polish (in `kaczmarz_solver_cchunk_mp.py`)

Variable	Default	Purpose
CUBEFIT_TILE_BT_STEPS	6	Backtracking steps for the aggregated ΔR across bands.
CUBEFIT_TILE_BT_FACTOR	0.5	Aggregated α shrink factor.
CUBEFIT_GLOBAL_TAU	0.5	Global step cap using $\ \Delta R \ _2^2 \cdot E[c, p]$ vs ' τ '.
CUBEFIT_GLOBAL_ENERGY_BLEND	1e-3	Blend local $\ A_w \ _2^2$ with global column energy $E[c, p]$.
CUBEFIT_NNLS_ENABLE	1	Enable tile-local weighted NNLS polish.
CUBEFIT_NNLS_EVERY	1	Run NNLS polish every N tiles.
CUBEFIT_NNLS_MAX_COLS	128	Max columns K in the NNLS system.
CUBEFIT_NNLS_MAX_BYTES	1000000000	Approx memory cap for NNLS system (rows×8 per column).
CUBEFIT_NNLS_SUB_L	0	λ -subsample size for NNLS (0 → use all).
CUBEFIT_NNLS_SOLVER	nnls	One of <code>nnls</code> , <code>lsq</code> , <code>mu</code> , <code>fista</code> (FISTA recommended for NNLS).
CUBEFIT_NNLS_MAX_ITER	50	Iteration cap for iterative NNLS (<code>mu</code> , <code>fista</code> , <code>lsq</code> 's TRF).
CUBEFIT_NNLS_MIN_IMPROVE	0.999	Required weighted-RMSE ratio to accept polish (< this accept).

Solver: λ -weights (in `kaczmarz_solver_cchunk_mp.py` / `cube_utils.py`)

Variable	Default	Purpose
CUBEFIT_LAMBDA_WEIGHTS_ENABLE	1	Enable λ -weights for solver (0 → disable).
CUBEFIT_LAMBDA_WEIGHTS_DSET	/HyperCube/lambda_weights	Dataset to read.
CUBEFIT_LAMBDA_MIN_W	1e-6	Floor on λ -weights; protects vs zero.
CUBEFIT_LAMBDA_WEIGHTS_AUTO	1	Auto-generate weights if missing.

Fit tracking (in `fit_tracker.py`)

Variable	Default	Purpose
FITTRACKER_START	spawn (or best available)	Start method for the tracker process (spawn).

Variable	Default	Purpose
CUBEFIT_TRACKER_QSIZE	8192	Max queue size; drops messages when full if
CUBEFIT_TRACKER_FLUSH_EVERY	128	Flush to disk after this many messages.
CUBEFIT_TRACKER_FLUSH_SEC	5.0	Or flush if this many seconds have passed.
CUBEFIT_RMSE_STRIDE	16	Only enqueue every Nth batch RMSE sample

Typical end-to-end workflow

```

# 1) Build HDF5 backbone & rebin operator
mgr = H5Manager("galaxy.h5", tem_pix=tem_loglam, obs_pix=obs_loglam)
mgr.init_base(H5Dims(nSpat=S, nLSpec=L, nTSpec=T, nVel=V, nComp=C, nPop=P))
mgr.populate_from_arrays(losvd=H_SVC, datacube=Y_SL, templates=T_PT)
mgr.ensure_rebin_and_resample()

# 2) Build hypercube
from CubeFit.hypercube_builder import build_hypercube
build_hypercube("galaxy.h5", norm_mode="data") # or "model"

# (Optional) Flip normalization later without rebuild
from CubeFit.hypercube_builder import convert_hypercube_norm
convert_hypercube_norm("galaxy.h5", to_mode="model", recompute_energy=True)

# 3) Create λ-weights
from CubeFit.cube_utils import ensure_lambda_weights
ensure_lambda_weights("galaxy.h5")

# 4) Solve
from CubeFit.pipeline_runner import PipelineRunner
runner = PipelineRunner("galaxy.h5")
x, stats = runner.solve_all_mp_batched(
    epochs=6, lr=0.6, project_nonneg=True, processes=8, blas_threads=2,
    orbit_weights=orbit_w # optional (C, )
)

```

Notes on normalization & physics

- `norm="model"` keeps each (s, c) model's scale proportional to the LOSVD amplitude — closest to “physical mass/flux” carried by LOSVD at that spaxel.
- `norm="data"` ties the per-spaxel model sum to the observed mean flux (masked), splitting across components in proportion to LOSVD amplitudes. This is convenient for direct data-scale comparisons but can blur absolute differences across datasets. You can switch modes post-hoc with `convert_hypercube_norm`.

The solver respects /Mask and λ -weights uniformly, so **absorption features** can dominate the fit when λ -weights are present (verify the printed min/max/mean at startup to ensure they are not all ones).

Troubleshooting & sanity checks

- **Hypercube completeness:** use the `_done` bitmap helpers to inspect or reset tiles if needed.
 - **Normalization flips:** use `convert_hypercube_norm` (streams in S-tiles and handles /Mask); do **not** read the entire dataset into RAM.
 - **Line weights:** ensure `/HyperCube/lambda_weights` exists and spans a range (e.g., min ≈ 1 , max ≈ 1); the solver prints min/max/mean.
 - **Population mixture looks uniform across components:**
 - Confirm λ -weights are applied (banner shows they're not all 1).
 - Increase epochs modestly; keep `GLOBAL_ENERGY_BLEND` small but nonzero.
 - Enable NNLS polish with a moderate `CUBEFIT_NNLS_MAX_COLS`, a λ -subsample (`CUBEFIT_NNLS_SUB_L`), and `CUBEFIT_NNLS_SOLVER=fista` for sharper per-tile updates.
-

Where to look in the code

- **HDF5 backbone:** `hdf5_manager.py` (safe open, grids, rebin operator).
 - **Builder:** `hypercube_builder.py` (FFT conv, normalization, col_energy, chunking + resume).
 - **Solver:** `kaczmarz_solver_cchunk_mp.py` (MP loop, λ -weights, trust/backtracking, NNLS polish).
 - **Runner:** `pipeline_runner.py` (orchestration, tracking, `/X_global`).
 - **Tracking:** `fit_tracker.py` (sidecar writer, RMSE history, progress, snapshots).
 - **Recon/plots:** utilities in `kz_fitSpec.py` (tile-aligned recon) and safe plotting helpers.
-

Building the docs

Place this file at `docs/CubeFit.md` and run:

```
make          # builds docs/CubeFit.html and docs/CubeFit.pdf via pandoc  
mkdocs serve  # live preview with Material theme (served from docs/)
```

Pandoc targets (from your Makefile):

- `docs/CubeFit.html` via:
`pandoc -s -f gfm -t html5 docs/CubeFit.md -o docs/CubeFit.html --metadata title="CubeFit" --toc`
- `docs/CubeFit.pdf` via XeLaTeX:
`pandoc -s -f gfm docs/CubeFit.md -o docs/CubeFit.pdf --pdf-engine=xelatex -V geometry:margin=1in -V mainfont="Latin Modern Roman" -V monofont="Latin Modern Mono" --toc`

MkDocs (from your `mkdocs.yml`):

```
site_name: CubeFit  
theme:  
  name: material  
nav:  
  - Home: index.md  
  - CubeFit: CubeFit.md  
docs_dir: docs
```