

detect_wells() — One-Page Spec

Purpose: Identify and label canonical wells (stable signal geometries) in per-primitive residual traces for NGF Stage-11 (Warp → Detect → Denoise).

API

Signature

detect_wells(traces: Dict[str, np.ndarray], *, jitter: int = 3, z_abs: float = 2.5, overlap_delta: float = 0.8) -> List[WellDescriptor]

Inputs

- **traces**: dict[primitive_id → 1D float array] residual energy signals (length T).
- **jitter**: number of stochastic resamples for stability checking (small input perturbations).
- **z_abs**: absolute z-score threshold to consider a peak significant.
- **overlap_delta**: minimum z-gap to suppress overlapping weaker peaks (lateral inhibition).

Outputs

- List of **WellDescriptor**: {primitive_id, well_type, center_t, width, z, stability, features, notes}.

Preconditioning

- 1) **Normalize** per channel (z-normalize or robust MAD).
- 2) **Smooth/bandlimit** (e.g., 5–11 sample moving average). Preserve both raw and smoothed versions.
- 3) **Spancomplement** to remove common-mode energy across channels if available.

Prototype kernels (matched filtering)

Use a compact family $K_\theta(t)$:

- **Unimodal** (half-sine / Gaussian) — axis pull.
- **Skewed unimodal** — quadrant well (off-center rise/decay).
- **Mexican hat** (two-lobe) — ring/oscillation well.
- **Broad hill** — shallow origin.
- **Narrow spike (edge-biased)** — deep edge.

For each channel, sweep θ (width, skew) and take the best correlation score and peak time.

Null calibration → z-scores

For each channel, build a **null** distribution via time permutations / phase shifts of the trace. Convert the best matched-filter score into a **z-score** relative to the null. Keep the tuple (best_kernel, z, t_peak).

Feature vector ϕ (measured at winning peak)

- **Height** (z), **width** (FWHM), **skewness**, **lobe ratio** (two-lobe vs one),
- **location** (early/mid/late window), **area** under bump, **pre/post slope symmetry**,
- **harmonic leakage** (energy outside passband). Concatenate into $\phi \in \mathbb{R}^m$.

Labeling rule (tiny decision tree)

If $z < z_{\text{abs}} \rightarrow$ **reject** (no well). Else:

- **Axis■Pull**: unimodal, low skew, mid width.
- **Quadrant**: unimodal, high skew, off■center peak.
- **Ring**: two lobes with central dip; balanced lobe ratio.
- **Shallow■Origin**: broad width, low height.
- **Deep■Edge**: very narrow, peak near boundary.

Lateral inhibition & stability

Overlap suppression: if two peaks (same channel or neighbors) overlap in time/shape and differ by $\geq \text{overlap_delta}$ in z , keep the stronger, suppress the weaker.

Jitter test: rerun detection on *jitter* perturbed inputs; compute stability = fraction of runs with same label $\pm \Delta t$. Require stability ≥ 0.67 – 0.8 .

Acceptance criteria

Criterion	Default
Significance (z)	$z \geq 2.5$ (tune 2.3–3.0 from validation)
Exclusivity (Δz to next overlap)	≥ 0.8
Stability (jitter)	≥ 0.67 (2/3) — prefer ≥ 0.8
False discovery rate on synthetic nulls	$\leq 1\%$

WellDescriptor schema

```
WellDescriptor = {  
  primitive_id: str,  
  well_type: Literal['Axis-Pull', 'Quadrant', 'Ring', 'Shallow-Origin', 'Deep-Edge'],  
  center_t: int, width: int, z: float, stability: float,  
  features: Dict[str, float],  
  notes: Optional[str]  
}
```