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A gradient-orthogonal Q-polynomial representation of axially symmetric optical surfaces has been used for several years by designers. It improves upon the standard polynomial form by being simpler to interpret, using fewer terms to adequately define a surface, and sometimes even offering quicker convergence in design optimization. Q-polynomials were first introduced with the publication of:


The use of Q-polynomials was extended, by the original author, to address freeform shapes through the following articles:


The implementation of this package follows the description in “Fitting freeform shapes with orthogonal bases”.
The Q-spectrum offers a natural way to quantify and investigate surface structure on parts with circular apertures. For example, spoke structure on the surface appears in specific columns of the spectrum whereas raster patterns appear as diagonal bands with a slope of one half. Rotationally symmetric structure, such as rings etc., are all contained in the $m=0$ column. The spectrum is computed with an FFT-like step that suffers similarly from aliasing unless a sufficiently large range of frequencies is computed.
1.1 Generating maps

Interesting features in the spectrum can be extracted to create topography layers that quantify and highlight flaws in the surface. The same process can be used to monitor residuals in the data after subtracting fitted polynomial terms. In the example below the radial-azimuthal spectrum shows quite distinct columns at particular azimuthal orders that are even more evident by summing the squares of the coefficients to produce a spectral line plot. The last image shows the contribution of just a sample of the spectral lines in the original data map. The article “Fitting freeform shapes with orthogonal bases” provides more details on the types of spectral analysis that can be performed and the characteristic flaws associated with polishing techniques that employ either raster or rotational modes.
The above images were generated from data supplied by Mahr GmbH.
To generate a Q-freeform spectrum from a data map, pass the coordinate axis x and y and 2-D array of data with shape (x.size, y.size) as arguments to the method qspec(). The azimuthal and radial spectrum limits are set by m_max and n_max respectively.

```python
>>> import skqfit.qspectre as qf
>>> ...
>>> qspec = qf.qspec(x, y, zmap, m_max=500, n_max=500)
```

To observe the contribution of spectrum components it is necessary to use the sine and cosine elements in the forward processing step. After modifying the spectrum, an inverse process creates a data map along with optional x and y derivatives.

```python
>>> qs = qf.QSpectrum()
>>> qs.data_map(x, y, zmap)
>>> a_nm, b_nm = qs.q_fit(mmax, nmax)
>>> ...
>>> nmap, dfdx, dfdy = qs.build_map(x, y, a_nm, b_nm, inc_deriv=True)
```

A 1D trace across the fitted data can also be generated by passing a set of xv, yv arrays that represent the (x, y) coordinates.

```python
>>> zv, dfdx, dfdy = qs.build_profile(xv, yv, a_nm, b_nm, inc_deriv=True)
```
CHAPTER 3

Code structure

The package only contains two modules. The Jacobi module provides support for normalised Jacobi polynomials that extend the range of usable parameters before an overflow condition is encountered. Even though the Q-fitting algorithm is procedural the implementation is via a Q-spectrum class as it allowed the spectrum to be iterated with different parameters without having to reload the data. The algorithm implementation refers to the relevant section of the reference documents.

3.1 The skqfit.asmjacp Module

References:


class skqfit.asmjacp.AsymJacobiP (nmax)

    Bases: object

    Generate asymmetric Jacobi like polynomials needed for the freeform fit as defined in the reference document [A.1]

    The Jacobi P polynomials can generate large values that lead to a double overflow for large m and n values. It is tested to (1500, 1500) for (m,n). Exceeding these values may lead to overflow events.

    The scipi.special.jacobi function is has a 1.5:1 performance advantage over the current implementation but it doesn’t support the normalization and can lead to an overflow condition for n or m over 500.

    build_recursion (m)

        Build the recursion coefficients and saves them as a sequence of tuples. These are the coefficients to build the m type polynomials up to order n.

    jmat_u_x (jmat, uv, xv)

        Builds the asymmetric Jacobi P polynomial as defined in [2] A.1 for all of the x values with the scaling factor of u**m which extends the range of usable (m,n) values before an overflow condition occurs.

    jmat_x (jmat, xv)

        Builds the asymmetric Jacobi P polynomial as defined in [2] A.1 for all a vector of x values.

    jvec_u_x (jvec, u, x)

        Builds the asymmetric Jacobi P polynomial for a single x as defined in [2] A.1 for all of the x values with the scaling factor of u**m which extends the range of usable (m,n) values.

    jvec_x (jvec, x)

        Builds the sequence of Jacobi polynomials for the value x based on [A.2-5]
3.2 The skqfit.qspectre Module

Reference documents


class skqfit.qspectre.QSpectrum(m_max=None, n_max=None)

Bases: object

Performs precomputation if Q spectrum limits are passed, otherwise it is delayed until the data is loaded. The class supports processing a data map or a pointer to a sag function that can be used for analytic testing.

Parameters: m_max, n_max: int

The azimuthal and radial spectrum order. Setting values above 1500 may lead to overflow events.

bfs_param()

Returns the fitted radius, curvature and centre.

Returns: radius, curvature, centre: float, float, (x,y) tuple

build_map(x, y, a_nm, b_nm, curv=None, radius=None, centre=None, extend=1.0, interpolated=True, inc_deriv=False)

Creates a 2D topography map and optional x and y derivate maps using the x and y axis vectors and the Q-freeform parameters.

Parameters:

x, y: array X, and Y axis values for the map to be created. The arrays must be sorted to increasing order.

a_nm, b_nm: 2D array The cosine and sine terms for the Q freeform polynomial

curv: float Nominal curvature for the part. If None uses the estimated value from the previous fit.

radius: float Defines the circular domain from the centre. If None uses the estimated value from the previous fit.

centre: (cx, cy) The centre of the part in axis coordinates. If None uses the estimated value from previous fit.

extend: float Generate a map over extend * radius from the centre

interpolated: boolean If True uses a high resolution regular polar grid to build the underlying data and a spline interpolation to extract the (x, y) grid, otherwise it evaluates each (x, y) point exactly. The non-interpolated solution is significantly slower and only practical for smaller array sizes.

inc_deriv: boolean Return the X and Y derivatives as additional maps

Returns:

zmap: 2-D array Data map with shape (x.size, y.size)

xder: 2-D array X derivative map with shape (x.size, y.size) if inc_deriv is True, else None

yder: 2-D array Y derivative map with shape (x.size, y.size) if inc_deriv is True, else None

build_profile(xv, yv, a_nm, b_nm, curv=None, radius=None, centre=None, extend=1.0, incderiv=False)

Returns the nominal sag and optional x and y derivatives along a 1D trajectory of (x, y) coordinates.

Parameters:
x, y: arrays  Arrays of values representing the (x, y) coordinates.

a_nm, b_nm: 2D array  The cosine and sine terms for the Q freeform polynominal

curv: float  Nominal curvature for the part. If None uses the estimated value from the previous fit.

radius: float  Defines the circular domain from the centre. If None uses the estimated value from the previous fit.

centre: (cx, cy)  The centre of the part in axis coordinates. If None uses the estimated value from previous fit.

extend: float  Generate a map over extend * radius from the centre

inc_deriv: boolean  Return the X and Y derivatives as additional maps

Returns:

zval: array  Sag values for the (x, y) sequence

xder: array  X derivative map for the (x, y) sequence if inc_deriv is True, else None

yder: array  Y derivative map for the (x, y) sequence if inc_deriv is True, else None

build_q_spectrum(m_max=None, n_max=None)  Fits the departure from a best fit sphere to the Q-polynominals as defined in [1](1.1) and returns the root sum square of the azimuthal terms.

Parameters:

m_max, n_max: int  The azimuthal and radial spectrum order. If None it uses the previous values and if not defined it matches the values to the pixel resolution.

Returns:

2D array  The (m,n) matrix representation of the spectrum (sqrt(cos^2 + sin^2)) terms

data_map(x, y, zmap, centre=None, radius=None, shrink_pixels=7, bfs_curv=None)  Creates the spline interpolator for the map, determines the best fit sphere and minimum valid radius.

Parameters:

x, y: arrays  The arrays are the X and Y axis values for the data map. The arrays must be sorted to increasing order.

zmap: array_like  2-D array of data with shape (x.size,y.size).

centre: (cx, cy)  The centre of the part in axis coordinates. If None the centre is estimated by a centre of mass calculation

radius: float  Defines the circular domain from the centre. If None it determines the maximum radius from the centre that contains no invalids (NAN).

shrink_pixels: int  The estimated radius is reduced by 7 pixels to avoid edge effects with the spline interpolation. Ignored if the radius is specified.

q_fit(m_max=None, n_max=None)  Fits the departure from a best fit sphere to the Q-freeform polynominals as defined in [1](1.1) and returns the individual sine and cosine terms.

Parameters:

m_max, n_max: int  The azimuthal and radial spectrum order. If None it uses the previous values and if not defined it matches the values to the pixel resolution. The maximum resolution supported is (1500, 1500)

Returns:
a_nm, b_nm: 2D array  The (n,m) matrix representation of the cosine and sine terms

set_sag_fn (sag_fn, radius, bfs_curv=None)
  A vectorized polar sag function that takes rho and theta as arguments.

This function is used to pass an analytic sag function to test the performance of the algorithm to a higher
precision but can also be used to define the input map and bypass data_map().

skqfit.qspectre.qspec (x, y, zmap, m_max=None, n_max=None, centre=None, radius=None,
  shrink_pixels=7)
  A wrapper function that creates the Q-spectrum object, loads and performs the fit of the data to the Q polynomials.

Parameters:

  x, y: array_like  The interpolator uses grid points defined by the coordinate arrays x, y. The arrays must
    be sorted to increasing order.

  zmap: array_like  2-D array of data with shape (x.size,y.size)

  m_max, n_max: int  The azimuthal and radial spectrum order. If None, it uses the previous values and if
    not defined it matches the values to the pixel resolution.

  centre: (cx, cy)  The centre of the part in axis coordinates. If None the centre is estimated by a centre of
    mass calculation

  radius: float  Defines the circular domain from the centre. If None it determines the maximum radius
    from the centre that contains no invalids (NAN).

  shrink_pixels: int  The estimated radius is reduced by 7 pixels to avoid edge effects with the spline inter-
    polation. Ignored if the radius is specified.

Returns:

  2D array  The (m,n) matrix representation of the spectrum (sqrt(cos^2 + sin^2)) terms
4.1 Quick Installation

If you have pip installed, you should be able to install the latest stable release of scikit-qfit by running the following:

```bash
pip install scikit-qfit
```

4.2 Latest Software

The latest software can be downloaded from GitHub

4.3 Installation Dependencies

scikit-qfit requires that the following software packages to be installed:

- Python 2.7.6 or later.
- NumPy 1.8.2 or later.
- SciPy 0.13.3 or later.
5.1 Modules

5.1.1 skqfit package

Submodules

skqfit.asmjacp module

References:

class skqfit.asmjacp.AsymJacobiP (nmax)

Bases: object

Generate asymmetric Jacobi like polynomials needed for the freeform fit as defined in the reference document [A.1]

The Jacobi P polynomials can generate large values that lead to a double overflow for large m and n values. It is tested to (1500, 1500) for (m,n). Exceeding these values may lead to overflow events.

The scipy.special.jacobi function is has a 1.5:1 performance advantage over the current implementation but it doesn’t support the normalization and can lead to an overflow condition for n or m over 500.

build_recursion (m)

Build the recursion coefficients and saves them as a sequence of tuples. These are the coefficients to build the m type polynomials up to order n.

jmat_u_x (jmat, uv, xv)

Builds the asymmetric Jacobi P polynomial as defined in [2] A.1 for all of the x values with the scaling factor of u**m which extends the range of usable (m,n) values before an overflow condition occurs.

jmat_x (jmat, xv)

Builds the asymmetric Jacobi P polynomial as defined in [2] A.1 for all a vector of x values.

jvec_u_x (jvec, u, x)

Builds the asymmetric Jacobi P polynomial for a single x as defined in [2] A.1 for all of the x values with the scaling factor of u**m which extends the range of usable (m,n) values.

jvec_x (jvec, x)

Builds the sequence of jacobi polynomials for the value x based on [A.2-5]
skfit.qspectre module

Reference documents


class skfit.qspectre.QSpectrum(m_max=None, n_max=None)
    Bases: object

    Performs precomputation if Q spectrum limits are passed, otherwise it is delayed until the data is loaded. The class supports processing a data map or a pointer to a sag function that can be used for analytic testing.

    Parameters: m_max, n_max: int
        The azimuthal and radial spectrum order. Setting values above 1500 may lead to overflow events.

    bfs_param()
        Returns the fitted radius, curvature and centre.

    Returns: radius, curvature, centre: float, float, (x,y) tuple

build_map(x, y, a_nm, b_nm, curv=None, radius=None, centre=None, extend=1.0, interpolated=True, inc_deriv=False)

    Creates a 2D topography map and optional x and y derivate maps using the x and y axis vectors and the Q-freeform parameters.

    Parameters:

        x, y: array  X, and Y axis values for the map to be created. The arrays must be sorted to increasing order.

        a_nm, b_nm: 2D array  The cosine and sine terms for the Q freeform polynominal

        curv: float  Nominal curvature for the part. If None uses the estimated value from the previous fit.

        radius: float  Defines the circular domain from the centre. If None uses the estimated value from the previous fit.

        centre: (cx, cy)  The centre of the part in axis coordinates. If None uses the estimated value from previous fit.

        extend: float  Generate a map over extend * radius from the centre

        interpolated: boolean  If True uses a high resolution regular polar grid to build the underlying data and a spline interpolation to extract the (x, y) grid, otherwise it evaluates each (x, y) point exactly.

        The non-interpolated solution is significanatly slower and only practical for smaller array sizes.

        inc_deriv: boolean  Return the X and Y derivatives as additional maps

    Returns:

        zmap: 2-D array  Data map with shape (x.size, y.size)

        xder: 2-D array  X derivative map with shape (x.size, y.size) if inc_deriv is True, else None

        yder: 2-D array  Y derivative map with shape (x.size, y.size) if inc_deriv is True, else None

build_profile(xv, yv, a_nm, b_nm, curv=None, radius=None, centre=None, extend=1.0, inc_deriv=False)

    Returns the nominal sag and optional x and y derivatives along a 1D trajectory of (x, y) coordinates.

    Parameters:
Data Map

**x, y:** arrays
Arrays of values representing the (x, y) coordinates.

**a_nm, b_nm:** 2D array
The cosine and sine terms for the Q freeform polynominal

**curv:** float
Nominal curvature for the part. If None uses the estimated value from the previous fit.

**radius:** float
Defines the circular domain from the centre. If None uses the estimated value from the previous fit.

**centre:** (cx, cy)
The centre of the part in axis coordinates. If None uses the estimated value from previous fit.

**extend:** float
Generate a map over extend * radius from the centre

**inc_deriv:** boolean
Return the X and Y derivatives as additional maps

**Returns:**

**zval:** array
Sag values for the (x, y) sequence

**xder:** array
X derivative map for the (x, y) sequence if inc deriv is True, else None

**yder:** array
Y derivative map for the (x, y) sequence if inc deriv is True, else None

**build_q_spectrum** *(m_max=None, n_max=None)*

Fits the departure from a best fit sphere to the Q-polynominals as defined in \[1\](1.1) and returns the root sum square of the azimuthal terms.

**Parameters:**

**m_max, n_max:** int
The azimuthal and radial spectrum order. If None it uses the previous values and if not defined it matches the values to the pixel resolution.

**Returns:**

**2D array**
The (m,n) matrix representation of the spectrum (sqrt(cos^2 + sin^2)) terms

**data_map** *(x, y, zmap, centre=None, radius=None, shrink_pixels=7, bfs_curv=None)*

Creates the spline interpolator for the map, determines the best fit sphere and minimum valid radius.

**Parameters:**

**x, y:** arrays
The arrays are the X and Y axis values for the data map. The arrays must be sorted to increasing order.

**zmap:** array_like
2-D array of data with shape (x.size, y.size).

**centre:** (cx, cy)
The centre of the part in axis coordinates. If None the centre is estimated by a centre of mass calculation.

**radius:** float
Defines the circular domain from the centre. If None it determines the maximum radius from the centre that contains no invalids (NAN).

**shrink_pixels:** int
The estimated radius is reduced by 7 pixels to avoid edge effects with the spline interpolation. Ignored if the radius is specified.

**q_fit** *(m_max=None, n_max=None)*

Fits the departure from a best fit sphere to the Q-freeform polynomials as defined in \[1\](1.1) and returns the individual sine and cosine terms.

**Parameters:**

**m_max, n_max:** int
The azimuthal and radial spectrum order. If None it uses the previous values and if not defined it matches the values to the pixel resolution. The maximum resolution supported is (1500, 1500)

**Returns:**
**a_nm, b_nm: 2D array** The (n,m) matrix representation of the cosine and sine terms

**set_sag_fn (sag_fn, radius, bfs_curv=None)**
A vectorized polar sag function that takes rho and theta as arguments.

This function is used to pass an analytic sag function to test the performance of the algorithm to a higher precision but can also be used to define the input map and bypass data_map().

**skqfit.qspectre.qspec (x, y, zmap, m_max=None, n_max=None, centre=None, radius=None, shrink_pixels=7)**
A wrapper function that creates the Q-spectrum object, loads and performs the fit of the data to the Q polynomials.

**Parameters:**

**x, y: array_like** The interpolator uses grid points defined by the coordinate arrays x, y. The arrays must be sorted to increasing order.

**zmap: array_like** 2-D array of data with shape (x.size,y.size)

**m_max, n_max: int** The azimuthal and radial spectrum order. If None, it uses the previous values and if not defined it matches the values to the pixel resolution.

**centre: (cx, cy)** The centre of the part in axis coordinates. If None the centre is estimated by a centre of mass calculation

**radius: float** Defines the circular domain from the centre. If None it determines the maximum radius from the centre that contains no invalids (NAN).

**shrink_pixels: int** The estimated radius is reduced by 7 pixels to avoid edge effects with the spline interpolation. Ignored if the radius is specified.

**Returns:**

**2D array** The (m,n) matrix representation of the spectrum (sqrt(cos^2 + sin^2)) terms

**Module contents**

**5.2 License**

The MIT License (MIT)

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5.3 Developers

- npdata <npdata@bigpond.com>

5.4 Changelog

5.4.1 Version 0.1.4

- Adds spectrum to topography functionality
- Updated documentation and minor fixes

5.4.2 Version 0.1.3

- Fixed error when data has invalids

5.4.3 Version 0.1.2

- Fixed error in interpolator call
- Added documentation to readthedocs

5.4.4 Version 0.1.1

- Corrections and addition to PyPi documentation
- Added example file

5.4.5 Version 0.1

- Initial public pre-alpha release
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