PRE-PRINT
Smet, K. A. G. (2019). Tutorial: The LuxPy Python Toolbox for Lighting and Color Science. *LEUKOS*, 1–23.
doi.org/10.1080/15502724.2018.1518717

# Tutorial: The LuxPy Python toolbox for lighting and color science

Kevin A.G. SMET

ESAT/Light&Lighting Laboratory,
KU Leuven, Belgium

Kevin.Smet@kuleuven.be

**DOI:**
**10.1080/15502724.2018.1518717**

**ABSTRACT** LuxPy is a free and open source Python package that supports several common lighting, colorimetric, color appearance and other color science related calculations that should be useful to researchers and industry professionals. This paper describes the installation of the LuxPy toolbox, provides an overview of its basic design and functionality and gives several examples to demonstrate its basic utilities using a Jupyter notebook. LuxPy is available under a GPLv3 license at www.github.com/ksmet1977/luxpy/ or from the Python Package Index pypi.python.org/pypi/luxpy/.

**KEYWORDS** Python, toolbox, lighting, colorimetry, color appearance models, color science, cct, duv, tristimulus values, chromatic adaptation, color rendering, photobiological quantities, spectral optimization, hyper-spectral image simulation



Received 9 May 2018
Revised 28 August 2018
Accepted 29 August 2018

Address correspondence to

Kevin A.G. Smet, ESAT/Light&Lighting Laboratory, KU Leuven, Gebroeders Desmetstraat 1, 9000 Ghent, Belgium. E-mail: kevin.smet@kuleuven.be

Color versions of one or more of the figures in this article can be found online at http://www.tandfonline.com/ulks

## Learning Outcomes

It is the intent that, after studying this article and its associated Jupyter notebook, that the reader will:

1. be able to install the LuxPy package in a dedicated Python conda (virtual) environment;
2. be able to import the package in the Anaconda Spyder IDE or a Jupyter notebook;
3. have had an overview of the package's basic design and functionality;
4. have gained practical experience and knowledge on how to use the LuxPy package to perform a variety of basic, and some more advanced, calculations related to lighting and color science.

Note that this is a tutorial on the LuxPy package, not on Python itself. Good basic tutorials can be found online. For suggestions the reader is referred to the LuxPy github page.

## 1. Introduction

Researchers and industry professionals in lighting and color science related fields often need to make or share various kinds of common spectral or colorimetric calculations, such as interpolating spectral data following [CIE15-2004 2004] or determining its correlated color temperature (*CCT*), distance to the blackbody locus (*Duv*) and color rendering index (*CRI*).

Functions implementing such calculations are often implemented using Microsoft Excel (possibly in combination with Visual Basics for Applications, VBA) or Matlab [MathWorks Inc. 2018], each with their own advantages and disadvantages.

While Excel or some open-source variant (e.g. LibreCalc [www.libreoffice.org 2018]) are a great way to perform some simple calculations –with little or no learning curve, graphically represent the results and share them with colleagues or collaborators, Excel implementations can rapidly become very unwieldy and difficult to follow as the problem complexity and quantity of the data increases. In addition, spreadsheet type calculators lack flexibility when existing calculations need to be adjusted or need to work on different sized data sets. They also lack some basic functionality that often comes in handy when analyzing data or graphically representing results (for example, try making a 3-dimensional plot of some *L*a*b** data).

For more complex problems, code based calculations using a high-level – quite easy to learn– scripting language, such as Matlab or Python (www.python.org), are much more flexible and easier to follow and comprehend. In addition, Matlab and Python come with a large set of toolboxes that support a wide variety of functionality. However, that functionality and ease of use, comes at a cost, at least with Matlab, as it requires a quite expensive yearly renewable license. Python, on the other hand, is free and open source and has lots of toolboxes for analysis and numerical computing (e.g. Numpy), hence its growing popularity.

To facilitate lighting and color science related calculations and to foster scientific collaboration using a shared, free, easy to learn and powerful high-level scripting 'language', the LuxPy package was developed in Python 3. In addition, LuxPy could also be used in an academic or other teaching setting, to help students develop a basic understanding of and practical skills in lighting and color science, while simultaneously learning the basics of the Python programming language.

This paper describes the installation of the LuxPy toolbox, its basic design and functionality and provides several examples to demonstrate its usage. A Jupyter notebook (www.jupyter.org) – a document that contains live code, graphics and text and that can be run in a browser – has also been created to help the interested reader exploring the LuxPy toolbox in practice.

## 2. LuxPy installation

The easiest and recommended way of installing the LuxPy package is as follows.

- First, install *miniconda* for Python 3 by downloading the installer (from conda.io/miniconda.html) for your operating system. This installs the conda package manager and python. Make sure *conda.exe* can be found on the system path (if necessary, add it manually).
- Next, create a conda virtual environment for Python version 3.6 or higher with a full *Anaconda distribution* by opening up a (Anaconda) `command shell` and typing the following one line of commands after the $-prompt:

```
$ conda create --name py36
python=3.6 anaconda
```

  Note that Anaconda from Continuum Analytics is a free and open source distribution of Python for scientific computing that eases

package management and deployment (it contains a large set packages specifically aimed at scientific computation). *Enthought Canopy* is another option.

- Once installed, activate the virtual environment (this step must be done each time you want to use any of the packages installed in the environment) by typing:

```
$ activate py36
```

- Install *pip* – installer for packages listed on the Python Package Index (pypi.python.org) – to the *py36* conda virtual environment to ensure that any packages installed with pip will be installed to the *py36* environment and not globally:

```
(py36) $ conda install pip
```

- The LuxPy package can now be installed by typing:

```
(py36) $ pip install luxpy
```

The LuxPy packages has several dependencies (SciPy, NumPy, Pandas, Matplotlib, scikit-image and Setuptools). Should any errors show up, try and do a manual install of these dependencies, either using e.g. "*conda install scipy*" or "*pip install scipy*", and then try and reinstall luxpy using pip.

### 2.1 Use of LuxPy within the Spyder IDE

Spyder is an Interactive Development Environment (IDE) for scientific programming in the Python language. It supports IPython and popular Python libraries such as NumPy, SciPy, or Matplotlib. It should come installed with the Anaconda distribution in the conda virtual environment. If not, it can be installed by typing:

```
$ conda install -n py36 spyder
```

To start using LuxPy, open up the Spyder IDE in the py36 virtual environment:

```
       $ activate py36
(py36) $ spyder
```

The LuxPy package can now be imported into a module (.py script file) or the global namespace in the Spyder IDE (command line) by typing:

```
import luxpy as lx
```

### 2.2 Use of LuxPy within a Jupyter notebook

A Jupyter notebook (www.jupyter.org) is a document that can contain live code, graphics and text and that can be run in a browser. It is therefore an easy way of sharing calculations and their results, while giving the user access to edit the code and re-run the calculations, perhaps with different input or parameters. The Jupyter notebook should come installed with the Anaconda distribution in the *py36* virtual environment (if not, install it using "*pip install jupyter*") and can be run as follows:

```
(py36) $ jupyter notebook
```

To start using LuxPy, open up a new or existing notebook (e.g. the "*luxpy_basic_usage.ipynb*" notebook that can be downloaded from www.github.com/ksmet1977/luxpy) and as before, import the package by typing:

```
import luxpy as lx
```

## 3. LuxPy design and functionality

The LuxPy package functionality is distributed among several sub-packages and modules. Some sub-packages/modules load within the LuxPy namespace, others, dedicated to a specific functionality load within the sub-package/module namespace itself and should be accessed through that

name. A full description of all functions and their input arguments in the current version of LuxPy is beyond the scope of this paper, which only addresses the basic usage of the package. A detailed description of each function's input and output can be obtained in the usual way, by typing:

```
?lx.spd_to_xyz
```

to get help on for example the *spd_to_xyz()* function.

An overview with a short description of all functions in the LuxPy package can be found at: ksmet1977.github.io/luxpy/ or in the ***LuxPy_Documentation.pdf***.

## 3.1 Package structure and functionality

The LuxPy toolbox contains a variety **sub-packages** (**folder/subfolder/**) and *modules* (*.py*) providing support for a number of basic and more advanced calculation related to lighting and color science. They are listed in Table 1, on www.github.com/ksmet1977/luxpy and are briefly reviewed in the following subsections.

### 3.1.1 utils/helpers/

- *helpers*.py: Module with helper functions

### 3.1.2 utils/math/

- *math*.py: Module with math functions
- *optimizers*.py: Module for bounded optimization: *minimizebnd()*.

### 3.1.3 spectrum/spectral/

- *cmf*.py: Module with Color Matching Functions and related.
- *spectral*.py: Module with spectral data related functions: e.g. *spd_to_xyz()*, *spd_to_power()*, *cie_interp()*, *spd_normalize()*, *blackbody()*, *daylightphase()*, *cri_ref()*, …
- *spectral_databases*.py: Module with spectral data: e.g. CIE Reference Illuminants, CIE TCS reflectance functions, IESTM30 spectral functions, …

### 3.1.4 color/ctf/

- *colortransforms*.py: Module with color transformation functions: e.g. *xyz_to_Yxy()*, *Yxy_to_xyz()*, *xyz_to_lab()*, *lab_to_xyz()*,…
- *colortf*.py: Module enabling easy, short-hand, color transformation: e.g. *lx.colortf(spd, tf = 'spd>Yxy')* and *lx.xyz_to_Yxy(lx.spd_to_xyz(spd))* are equivalent.

### 3.1.5 color/cct/

- *cct*.py: Module with xyz tristimulus value conversions to *CCT* and *Duv*.

### 3.1.6 color/cat/

- *chromaticadaptation*.py: Module supporting chromatic adaptation transforms. (von Kries and RLAB).

### 3.1.7 color/cam/

- *colorappearancemodels*.py: Module with color appearance models.
  i. *cam_02_X.py:* Module with CIECAM02 based models. [Li 2017; Luo 2006; Moroney 2002] e.g. ciecam02, cam16, cam02ucs, cam16ucs/lcd/scd.
  ii. *cam15u*.py: Module with CAM for unrelated self-luminous stimuli. [Withouck 2015]
  iii. *sww2016*.py: Module with CAM based on mapping of Munsell color system. [Smet 2016]

### 3.1.8 color/deltaE/

- *colordifferences.py*: Module with color difference formulas: e.g. *DE_camucs()*, *DE2000()*, *DE_cspace()*

### 3.1.9 color/cri/

- *colorrendition*.py: Module related to color rendition calculations: e.g. *spd_to_cri()*, *spd_to_ciera()*, *spd_to_cierf()*, *spd_to_iesrf()*, …

**Table 1. Overview of structure of the subpackages / modules in the LuxPy package.**

| # | Sub package(s) | | Module(.py) | Namespace |
|---|---|---|---|---|
| **1** | **.utils** | **.helpers** | | **luxpy** |
| | | | *helpers.py* | **.** |
| | | **.math** | | **luxpy.math** |
| | | | *math.py* | **.** |
| | | | *optimizers.py* | **.** |
| **2** | **.spectrum** | **.spectral** | | **luxpy** |
| | | | *cmf.py* | **.** |
| | | | *spectral.py* | **.** |
| | | | *spectral_databases.py* | **.** |
| **3** | **.color** | **.ctf** | | **luxpy** |
| | | | *colortransformations.py* | **.** |
| | | | *colortf.py* | **.** |
| | | **.cct** | | **luxpy** |
| | | | *cct.py* | **.** |
| | | **.cat** | | **luxpy.cat** |
| | | | *chromaticadaptation.py* | **.** |
| | | **.cam** | | **luxpy.cam** |
| | | | *colorappearancemodels.py* | **.** |
| | | | *cam_02_X.py* | **.** |
| | | | *cam15u.py* | **.** |
| | | | *sww2016.py* | **.** |
| | | **.deltaE** | | **luxpy.deltaE** |
| | | | *colordifferences.py* | **.** |
| | | **.cri** | | **luxpy.cri** |
| | | | *colorrendition.py* | **.** |
| | | **.cri.utils** | *DE_scalers.py* | **.** |
| | | | *helpers.py* | **.** |
| | | | *init_cri_defaults_database.py* | **.** |
| | | | *graphics.py* | **.** |
| | | **.cri.indices** | *indices.py* | **.** |
| | | | *cie_wrappers.py* | **.** |
| | | | *ies_wrappers.py* | **.** |
| | | | *cri2012.py* | **.** |
| | | | *mcri.py* | **.** |
| | | | *cqs.py* | **.** |
| | | **.cri.iestm30** | *ies_tm30_metrics.py* | **.** |
| | | | *ies_tm30_graphics.py* | **.** |
| | | **.cri.VFPX** | *VF_PX_models.py* | **luxpy.cri.VFPX** |
| | | | *vectorshiftmodel.py* | **.** |
| | | | *pixelshiftmodel.py* | **.** |
| | | **.color.utils** | | **luxpy** |
| | | | *plotters.py* | **.** |
| **4** | **.classes** | | | **luxpy** |
| | | | *SPD.py (class SPD)* | **.** |
| | | | *CDATA.py (classes CDATA, XYZ, LAB)* | **.** |
| **5** | **.toolboxes** | **.photbiochem** | | **luxpy.photbiochem** |
| | | | *cie_tn003_2015.py* | **.** |
| | | | *ASNZS_1680_2_5_1997_COI.py* | **.** |
| | | | *circadian_CS_CLa_lrc.py* | **.** |
| | | **.indvcmf** | | **luxpy.indvcmf** |
| | | | *individual_observer_cmf_model.py* | **.** |
| | | **.spdbuild** | | **luxpy.spdbuild** |
| | | | *spd_builder.py* | **.** |
| | | **.hypspcim** | | **.luxpy.hypspcim** |
| | | | *hyperspectral_img_simulator.py* | |

i. *DE_scalers*.py: scaling functions for color difference *DE* to *CRI* conversion.

ii. *helpers*.py: basic functions and data for CRI calculations.

iii. *indices.py*: Module with wrapper functions for various cri measures.

   a. *cie_wrappers.py:* Module with CIE cri metrics *($R_a$, $R_f$)*. [CIE13.3-1995 1995; CIE224:2017 2017]

   b. *ies_wrappers.py:* Module with IES cri metrics *($R_f$, $R_g$)*. [David 2015; IES 2015]

   c. *cri2012*.py: Module with CRI2012 metrics.[Smet 2013]

   d. *mcri*.py: Module with memory color rendition index, $R_m$.[Smet 2012]

   e. *cqs*.py: Module supporting Color Quality Scale (v7.5 & v9.0).[Davis 2010]

iv. *graphics*.py: basic cri plotting functions.

v. *ies_tm30_metrics*.py: specific functionality for IES TM30 metrics.

vi. *ies_tm30_graphics*.py: specific graphic support for IES TM30.

vii. *VF_PX_models*.py

   a. *vectorshiftmodel*.py: support for vector field color shift and metameric uncertainty indices.

   b. *pixelshiftmodel*.py: supports color space pixel based color shift calculations.

#### 3.1.10 color/utils/

- *plotters*.py: Module with basic plotting functions: e.g. for spectrum, daylight and blackbody loci, …

#### 3.1.11 classes/

- *SPD*.py: Class wrapper (SPD) around basic spectral functionality.
- *CDATA*.py: Class wrappers around basic tristimulus value (class *XYZ*) and color space and color appearance model transformations (class *LAB*).

#### 3.1.12 toolboxes/ciephotbio/

- *cie_tn003_2015*.py: Module for CIE photobiological quantities (α-opic irradiance and equivalent illuminance)[CIE-TN003:2015 2015].
- *ASNZS_1680_2_5_1997_COI*.py: Module for the calculation of the Cyanosis Observation Index. [AS/NZS1680.2.5 1997]
- *circadian_CS_CLa_lrc.py*: Module implementing the Circadian Stimulus (CS) and Circadian Light (CLa) indices developed by the RPI Lighting Research Center. [Rea 2012].

#### 3.1.13 toolboxes/indvcmf/

- *individual_observer_cmf_model.py*: Module supporting Monte-Carlo generation of individual observer cone fundamentals (lms-CMFs)[Asano 2016].

#### 3.1.14 toolboxes/spdbuild/

- *spd_builder*.py: Module for building (simulating) and optimizing multi-component spectra.

#### 3.1.15 toolboxes/hypspcim/

- *hyperspectral_img_simulator*.py: Module for simulating hyper spectral images.

### 3.2 Data structure

LuxPy global constants (which should not be changed, so use `copy()` or `deepcopy()` when necessary!) can be recognized throughout in that they start with '_' and are in an all `_CAPITALS` format. These constants contain various parameters, settings or dictionaries with useful data, e.g.

*lx._CIE_ILLUMINANTS* is a Python dictionary (dict) containing spectral data of various CIE illuminants [CIE15-2004 2004].

Many LuxPy functions expect to receive their input data in specific formats. Two major data types can be distinguished. Spectral data and colorimetric data.

### 3.2.1 Spectral data using NumPy.ndarrays

Spectral data should be inputted to functions in a row-based format as NumPy.ndarrays, with the first row containing the wavelengths and subsequent rows the values of one or more spectra. When using the function *helpers.getdata* to read spectral data stored in .csv or other data file, the data in the file itself should be in a column-format.

A NumPy.ndarray variable '*spds*' containing *2* spectra and with wavelengths ranging from *380* nm to *780* nm and with a *5* nm wavelength spacing, thus should have a shape of (*3,81*).

As an example, see the following code which creates two spectra, a blackbody radiator and a CIE daylight phase, both with a CCT of *4500* K and with the wavelength range specified above, and queries its shape (whereby the code is split over multiple lines using the '\' line separator character):

```
import luxpy as lx
spds = lx.cri_ref([4500,4500],\
                  ref_type = ['BB','DL'],\
                   wl3 = [380,780,5])
print(spds.shape)
(3, 81)
```

### 3.2.2 Colorimetric data using NumPy.ndarrays

Colorimetric data is represented using 2 or 3 dimensional NumPy.ndarrays. The colorimetric dimension, e.g. X, Y, Z tristimulus values, are always on the last (highest) array axis.

For example, the tristimulus values of an equi-energy white point should have a shape (*1,3*) and can be created as follows:

```
import numpy as np # import NumPy package
xyzE = np.array([[100., 100., 100.]])
```

An array with tristimulus values for *N* stimuli should thus have a shape (*N,3*).

However, 3-dimensional arrays are also valid. In this case, the first axis is assumed to refer to different reflectance/transmittance samples, the second axis to the light sources illuminating those samples and the last axis, again to the colorimetric dimension.

The following code generates for example the CIE 1931 2° *X, Y, Z* tristimulus values for the *8* Munsell Test Colour Samples (*TCS*) used in the CIE 13.3-1995 color rendering index $R_a$ [CIE13.3-1995 1995], when illuminated by the 2 light source spectra generated earlier, i.e. the *4500* K blackbody and daylight phase spectra.

```
# gets the 8 CIE TCS from a database:
TCS8 = lx._CRI_RFL['cie-13.3-1995']['8']

# calculate xyz:
xyz8 = lx.spd_to_xyz(spds,\
                     cieobs = '1931_2',\
                     relative = True,\
                     rfl = TCS8)
print(xyz8.shape)
(8, 2, 3)
```

As can be seen, the shape of the xyz8 variable is (*8,2,3*), i.e. *(sample total, light source total, 3)*.

Many other LuxPy functions, e.g. *xyz_to_Yuv()*, take as input the either 2d or 3d Numpy.ndarray with the *X, Y, Z* tristimulus data to perform further calculations. These functions always assume that the array axes are specified as above! The advantage is that it is easy to do calculations on a

whole block of data at once, without the user having to specifically write loops to calculate the requested transformation for each sample and/or light source. For example:

```
# Calculate Y, CIE1976 u'v' chromaticity
# coordinates:
Yuv8 = lx.xyz_to_Yuv(xyz8)

# Convert back to tristimulus values:
xyz8_2 = lx.Yuv_to_xyz(Yuv8)

print(Yuv8.shape)
(8, 2, 3)
```

As a final example, consider the calculation of the *CCT* and *Duv* of a bunch of (in this case, the two specified above) light source spectra:

```
xyzw = lx.spd_to_xyz(spds,\
                     cieobs = '1931_2',\
                     relative = True,\
                     rfl = None)


cct, duv = lx.xyz_to_cct(xyzw,\
                    cieobs = '1931_2',\
                    mode = 'ohno',\
                    out = 2)
print(cct.shape)
(2, 1)
```

### 3.2.3 Spectral and colorimetric data using LuxPy classes

Spectral and colorimetric data also have their own dedicated classes which have basic functionality by providing *class method* wrappers to some of the LuxPy functions.

An instance of the class SPD can be created by initiating it directly with a NumPy.ndarray or by providing the path to a datafile as input. Using the 'spds' data array specified above (containing blackbody radiator and daylight phase spectra of 4500 K):

```
spds_inst = lx.SPD(spds)
```

The advantage of the creating such an SPD instance is that it allows for very readable code, when performing several operations in sequence:

```
xyz_inst = lx.SPD(spds).cie_interp(wl_new
= [400, 700, 5]).to_xyz()
```

The code above produced an instance of class XYZ, *xyz_inst*, which itself has several methods for easy use. It is also easy to store some additional specific info related to the data:

```
print("*Spectral data type in .dtype field:
{}".format(spds_inst.dtype))

print("*Wavelengths in .wl field:
{}".format(spds_inst.wl.shape))

print("*Values in .value field:
{}".format(spds_inst.value.shape))

print("*Number of spds in .N field:
{}".format(spds_inst.N))

print("*Shape of .value in .shape field:
{}".format(spds_inst.shape))
```

**Output:**
```
*Spectral data type in .dtype field: S
*Wavelengths in .wl field: (81,)
*Values in .value field: (2, 81)
*Number of spds in .N field: 2
*Shape of .value in .shape field: (2, 81)
```

The further conversion to CIELUV *L*u*v** coordinates, with assumed white point the equi-energy white, can for example be done as follows:

```
luv_inst = xyz_inst.ctf(dtype = 'luv',\
       xyzw = np.array([[100, 100, 100]]))
```

In each instance, the field '*value*' contains the actual data. For the SPD class, this is the spectral data stripped of its wavelengths (contained in a '*wl*' field), for the XYZ and *LAB* classes, this is an array with the tristimulus values and specified colorimetric data, respectively. For the latter a field called '*dtype*' keeps track of what type of data (e.g. '*xyz*', '*Yuv*', '*luv*', ...) the instance contains. In addition to a field '*cieobs*', which specifies the CMF set used, instances of the class *LAB* also contain a field '*cspace_par*' that contains a Python dictionary with a set of parameters (e.g. '*xyzw*') relevant to the calculation of the colorimetric values.

```
# Print data in parameter dictionary:
print(luv_inst.cspace_par)
```

```
Output:
{'cieobs': '1931_2',
'xyzw': array([[100, 100, 100]]),
'M': None,
'scaling': None,
'Lw': None,
'Yw': None,
'Yb': None,
'conditions': None,
'yellowbluepurplecorrect': None,
'mcat': None,
'ucstype': None,
'fov': None,
'parameters': None}
```

Each class also contains a basic plotting method for a quick look at the data contained in the instance:

```
spds_inst.plot() #class SPD

xyz_inst.plot() #class XYZ

xyz_inst.ctf(dtype = 'Yuv').plot(plt_type
= '2d', title = "CIE u'v'") #class LAB

lx.plotSL() # plots spectrum locus
```

The output of the first and last two commands is illustrated in respectively the top and bottom graphs of Fig. 1.

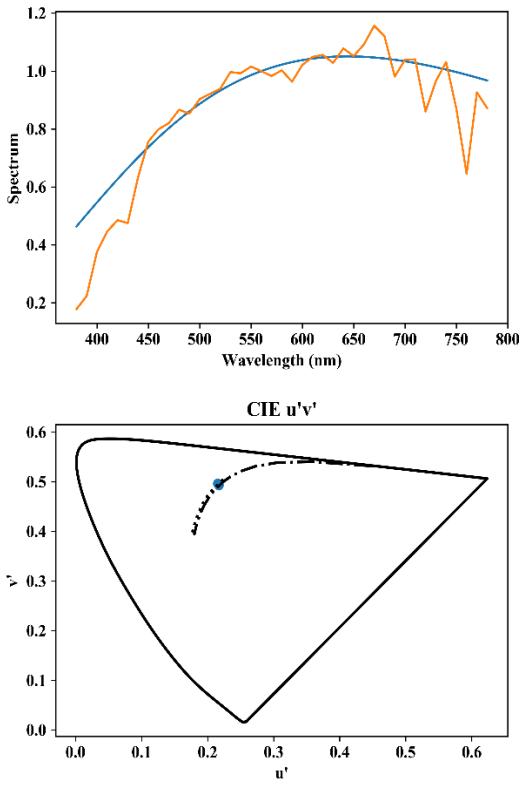


**Fig. 1. Output as generated by the plot() methods of classes SPD (top) and LAB (bottom).**

## 3.3 Versioning system

Each release of LuxPy is tagged with a *MAJOR.MINOR.PATCH* version number following the principles of semantic versioning whereby the *MAJOR* number reflects changes in the LuxPy package that break compatibility with previous versions, the *MINOR* number reflects changes that add functionality or that are backward compatible and with *PATCH* number reflecting bug fixes or minor additions.
The current version associated with this paper is *LuxPy_v1.3.8*.

## 4. Example usage

In this section, the basic use of the LuxPy package will be demonstrated with the help of a *Jupyter notebook*, "*luxpy_basic_usage.ipynb*", which can be downloaded from: www.github.com/ksmet1977/luxpy.

Alternatively, a live cloud based version can be accessed at Microsoft's Azure notebooks (best accessed using Internet Explorer): https://notebooks.azure.com/ksmet1977/libraries/luxpyhowtouse.

### 4.1 Importing LuxPy and other Python packages

Any session would start with importing the necessary packages.

```
# Package for color science calculations:
import luxpy as lx

# Package for scientific computing:
import numpy as np

# Package for plotting:
import matplotlib.pyplot as plt

# Package for timing functions:
import timeit

# Ensure inline plotting of figures:
%matplotlib inline
```

Note that getting help on a function or module is as easy as typing:

```
# get help on function arguments:
?lx.spd_to_xyz

# get help on cri sub-package:
?lx.cri
```

## 4.2 Working with spectral data

### *4.2.1 cri_ref()*

As a first example, let's get ($M = 4$) spectra normalized at wavelength ('lambda') 560 nm with respective CCTs of 3000 K, 4000 K, 4500 K and 6000 K and respective types: *'BB'* (blackbody radiator), *'DL'* (daylight phase), *'cierf'* (mixed reference illuminant as defined in [CIE224:2017 2017]) and *'BB'*:

```
# Set CIE observer, i.e. cmf set:
cieobs = '1964_10'

# Define M = 4 CCTs:
ccts = [3000,4000,4500, 6000]

# Define reference illuminant types
ref_types = ['BB','DL','cierf','DL']

# Calculate reference illuminants:
REF = lx.cri_ref(ccts, \
                 ref_type = ref_types,\
                 norm_type = 'lambda',\
                 norm_f = 600)

print('* REF.shape --> (M + 1 x number of
wavelengths): {}'.format(REF.shape))
```

```
Output:
* REF.shape --> (M + 1 x number of
wavelengths): (5, 471)
```

### *4.2.2 cie_interp()*

Secondly, let's interpolate the *REF* spectral data to a 400 nm – 700 nm range with a 5 nm spacing, and let the *cie_interp()* function determine the right kind of interpolation by defining the spectral data type as *kind = 'S'*:

```
REFi = lx.cie_interp(REF, \
         wl_new = np.arange(400,700+1,5),\
         kind = 'S')

print('* REFi.shape --> (M + 1 x number of
wavelengths): {}'.format(REFi.shape))
```

```
Output:
* REFi.shape --> (M + 1 x number of
wavelengths): (5, 61)
```

### *4.2.3 spd_to_xyz()*

Next, the tristimulus values (using the CIE 1964 10° CMFs) of those light source spectra can be obtained as follows:

```
# Get illuminant xyz:
xyz_REF = lx.spd_to_xyz(REF, \
                        cieobs = cieobs)
```

The tristimulus values of set of $N$ reflective samples illuminated by those $M$ sources can be obtained in the following way:

```
# Get 8 TCS from CIE 13.3-1995:
TCS8 = lx._CRI_RFL['cie-13.3-1995']['8']

# Get TCS xyz:
xyz_TCS8_REF = lx.spd_to_xyz(REF, \
                         cieobs = cieobs,\
                         rfl = TCS8, \
                         relative = True)
```

whereby *relative = True* (default) specifies that the output tristimulus values are normalized such that a *perfect white diffuse* sample has $Y = 100$.

The tristimulus values of the $N$ samples illuminated by the $M$ light sources and of the M light sources themselves can be obtained using:

```
xyz_TCS8_REF_2, xyz_REF_2 =
    lx.spd_to_xyz(REF,\
                  cieobs = cieobs,\
                  rfl = TCS8,\
                  relative = True,\
                  out = 2)

print('* xyz_TCS8_REF_2.shape -->
(NxMx3): {}'.format(xyz_TCS8_REF_2.shape))

print('* xyz_REF_2.shape -->
(Mx3):}'.format(xyz_REF_2.shape))
```

```
Output:
* xyz_TCS8_REF_2.shape --> (NxMx3):
(8, 4, 3)
* xyz_REF_2.shape --> (Mx3): (4, 3)
```

### *4.2.4 Plotting*

A simple plot (see Fig. 2) of the spectral data in REF can be made by the following lines of code:

```
plt.plot(REF[0],REF[1:].T)
plt.xlabel('wl (nm)')
plt.ylabel('spectral intensity (a.u)')
lgnd_str = ['{} K'.format(x) for x in ccts]
plt.legend(lgnd_str)
```

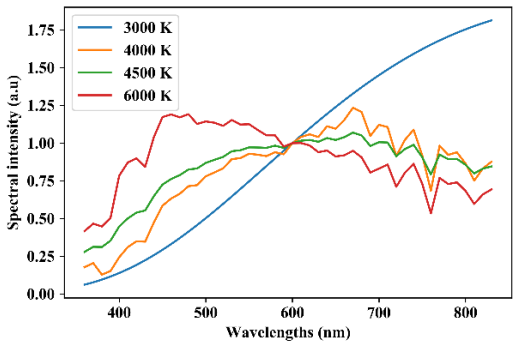


**Fig. 2. Plot of spectral data in REF.**

Figures can be easily exported for use in presentations and publications using the matplotlib function *figsave()*:

```
# Save figure in TIFF format at 600 dpi:
plt.figsave('Fig1_REF_spectra.tif',\
            format = 'tiff',\
            dpi = 600)
```

### *4.2.5 Individual Observer Color Matching Functions*

Color matching functions are stored in a dictionary that can be accessed as follows:

```
# Get CIE 1931 2° CMFs:
cmf_1931 = lx._CMF['1931_2']['bar']
# List all available CMF sets:
print(lx._CMF['types'])
```

```
Output:
['1931 2','1964 10','2006 2','2006 10',
'1931_2_judd1951',
'1931_2_juddvos1978','1951_20_scotopic'
]
```

There is also a module, called *'indvcmf'*, that supports Monte-Carlo generation of individual observer cone fundamentals and XYZ color matching functions [Asano 2016]. For example, a set of cone fundamentals (lms) for a stimulus with field size of 6° for 20 observers aged 32 can be generated as follows:

```
# Get 20 individual observer lms-CMFs:
lmsb =
  lx.indvcmf.genMonteCarloObs(n_obs = 20,\
                              list_Age = [32],\
                              fieldsize = 6)
# Or, using US 2010 population census to
generate Age Distribution,
# and output as XYZ CMF:
xyzb_us =
  lx.indvcmf.genMonteCarloObs(n_obs = 20,\
                    list_Age = 'us_census',\
                    fieldsize = 6,\
                    out ='XYZ')
```

The graph in Fig. 3 was generated by:

```
# Plot CMFs:
plt.figure()
plt.plot(xyzb_us[0],xyzb_us[1],\
         color ='r', linestyle ='-')
plt.plot(xyzb_us[0],xyzb_us[2],\
         color ='g', linestyle ='-')
plt.plot(xyzb_us[0],xyzb_us[3],\
         color ='b', linestyle ='-')
plt.xlabel('Wavelength (nm)')
plt.ylabel('XYZbar (area normalized)')
plt.title('Individual XYZ CMF (US census)')
```

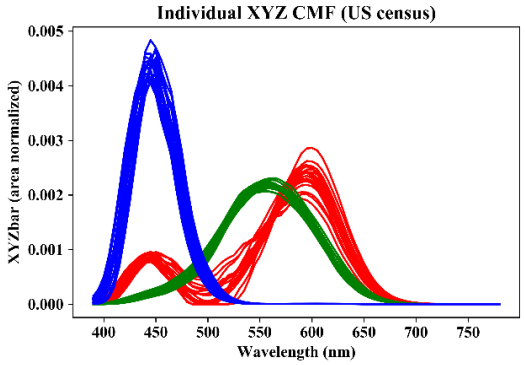


**Fig. 3. Monte-Carlo generated individual XYZ color matching functions.**

The XYZ values, using the Monte-Carlo generated CMFs, for spectral data in REF can be calculated as follows:

```
# Calculate XYZ values for REF:
xyz_ind = np.empty((REF.shape[0] - 1,\
                    xyzb_us.shape[-1],3))

for i in range(xyzb_us.shape[-1]):
    xyz_ind[:,i,:] = lx.spd_to_xyz(REF,\
                cieobs = xyzb_us[...,i],\
                relative = True)

print(xyz_ind.shape)
```

```
Output:
(4, 20, 3)
```

Categorical observer CMFs can be generated in a similar manner using the *indvcmf.genCatObs* function.

## 4.3 Working with colorimetric data

The tristimulus values XYZ obtained earlier can be further transformed to a range of color coordinates.

### *4.3.1 xyz_to_Yuv()*

As an example, the transformation to the CIE 1976 *u'v'* chromaticity coordinates is outlined below:

```
Yuv_REF_2 = lx.xyz_to_Yuv(xyz_REF_2)

Yuv_TCS8_REF_2 =
            lx.xyz_to_Yuv(xyz_TCS8_REF_2)

print("*   Yuv_REF_2.shape   -->   (Mx3):
{}".format(Yuv_REF_2.shape))
print("* Yuv_TCS8_REF_2.shape --> (NxMx3):
{}".format(Yuv_TCS8_REF_2.shape))
```

```
Output:
* Yuv_REF_2.shape --> (Mx3): (4, 3)
* Yuv_TCS8_REF_2.shape --> (NxMx3):
(8, 4, 3)
```

### *4.3.2 asplit(), plotSL() and plot_color_data()*

The colorimetric data can also be easily plotted. First the spectrum locus is drawn using *plotSL()*. This function has several input options (*cspace* and *cieobs* respectively determine the color space and CMF set to be used, *show = False* waits with showing the output graph until everything has been drawn in the *axh* axes, while *BBL = True* and *DL = True* regulate the plotting of a blackbody and daylight locus).

```
axh = lx.plotSL(cspace = 'Yuv',\
                cieobs = cieobs,\
                show = False,\
                BBL = True,\
                DL = True)
```

In a next step, the data is prepared for easy plotting by splitting the data array along the colorimetric dimension (last axis) and squeezing out any axes with a 1-dimensional size:

```
# Split array along last axis:
Y, u, v = np.squeeze(lx.asplit(Yuv_REF_2))
```

Finally, to plot the *u',v'* chromaticity coordinates (see Fig. 4):

```
lx.plot_color_data(u,v, formatstr = 'go',\
                   label = 'Yuv_REF_2')
```

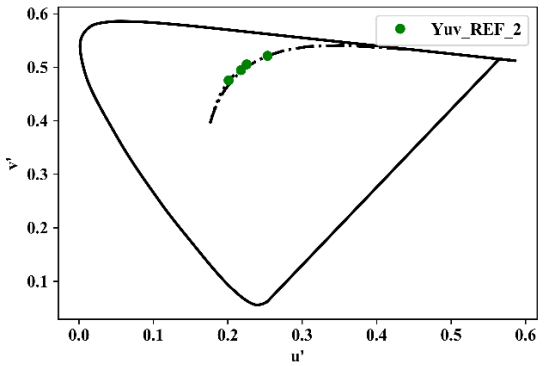


**Fig. 4 Output of `plot_color_data()`.**

### *4.3.3 colortf() and _CIE_ILLUMINANTS*

A short-hand color transformation function also exists. This function makes it easy to combine several transformations in one function call.

In the following example, tristimulus values XYZ are transformed to CIE 1976 *u'v'* chromaticity coordinates (*tf = 'Yuv'* defines the type of transform):

```
Yuv_REF = lx.colortf(xyz_REF_2, \
                     tf = 'Yuv')
```

Note that the input does not have to be an array with tristimulus values. One can also perform transformation between for example an array with spectral data and CIE 1976 *u'v'* chromaticity coordinates. Any combination of transformations for which the necessary forward (*xyz_to_...*) and backward (*..._to_xyz*) functions are defined can be made. For example, the spectral data in *REF* can be readily converted to colorimetric data by specifying the correct transform type (*tf = 'spd>Yuv'*):

```
Yuv_REF = lx.colortf(REF,\
                     tf = 'spd>Yuv')
```

Some color transformations require additional input parameters. This can be supplied as Python dictionary with the parameters names as keys. The forward and backward transform parameters should be supplied to the *colortf* function using keyword arguments, *fwtf* and *bwtf*, respectively.

For example, let's transform the *Yuv_REF* to CIE 1976 CIELAB *L*a*b** coordinates, with as white point tristimulus values those of CIE illuminant A. First, the dictionary with input parameters for the forward (i.e. *xyz_to_lab*) transform is defined:

```
xyzwA =
   lx.spd_to_xyz(lx._CIE_ILLUMINANTS['A'])
fwtf_userdefined = {'xyzw' : xyzwA}
```

whereby the spectrum of the CIE illuminant A is obtained from a LuxPy database with CIE illuminants, i.e. *_CIE_ILLUMINANTS*.

Next, the transform can be made as follows:

```
lab_REF2 = lx.colortf(Yuv_REF_2,\
                  tf = 'lab',\
                  fwtf = fwtf_userdefined)
```

Input parameters for the forward (only) transform can also be directly entered as keyword arguments to the *colortf* function:

```
lab_REF2 = lx.colortf(Yuv_REF_2, \
                      tf = 'lab',\
                      xyzw = xyzwA)
```

### 4.3.4 xyz_to_cct() and cct_to_xyz()

The correlated color temperature (CCT) and distance to the blackbody locus (Duv) can be obtained, using the Ohno's method [Ohno 2014] or using a brute-force search method in the CIE 1960 UCS color space, as follows:

```
# Ohno's approach with a Look-Up-Table:
cct_ohno, duv_ohno =
       lx.xyz_to_cct(xyz_REF_2,\
                     cieobs = cieobs,\
                     out = 'cct,duv',\
                     mode = 'lut')

# Brute-force search approach:
cct_search, duv_search =
       lx.xyz_to_cct(xyz_REF_2,\
                     cieobs = cieobs,\
                     out = 'cct,duv',\
                     mode = 'search')
```

whereby *cieobs* is a specifier for the type of CMFs that were used to calculate the tristimulus input values, *out* specifies the format of the output and *mode* determined the method used to obtain the *CCT* and *Duv*.

To perform the inverse calculation, i.e. going from *CCT* and *Duv* to tristimulus values:

```
# Ohno's approach with a Look-Up-Table:
xyz_REF_2_ohno =
               lx.cct_to_xyz(cct_ohno,\
                         duv = duv_ohno,\
                         cieobs = cieobs,\
                         mode = 'lut')

# Brute-force search approach:
xyz_REF_2_search =
               lx.cct_to_xyz(cct_search,\
                         duv = duv_search,\
                         cieobs = cieobs,\
                         mode = 'search')
```

Note that the brute-force method takes considerably longer than Ohno's method:

```
print('\nRun-time: cct_to_xyz_ohno():')
%timeit xyz_REF_2_ohno =
lx.cct_to_xyz(cct_ohno, duv = duv_ohno,\
            cieobs = cieobs, mode = 'lut')

print('\nRun-time: cct_to_xyz_search():')
%timeit xyz_REF_2_search =
          lx.cct_to_xyz(cct_search,
                         duv = duv_search,\
                         cieobs = cieobs,\
                         mode = 'search')
```

**Output:**

```
Run-time: cct to xyz ohno():
84.8 ms ± 1.14 ms per loop (mean ± std.
dev. of 7 runs, 10 loops each)

Run-time: cct_to_xyz_search():
10.1 s ± 167 ms per loop (mean ± std.
dev. of 7 runs, 1 loop each)
```

Please be aware that running the above code block could take a while, as the *timeit* macro needs to execute the *cct_to_xyz()* function several times!

## 4.4 Chromatic adaptation and corresponding colors (.cat)

LuxPy supports basic *von Kries* type chromatic adaptation transforms, with a wide choice of sensor spaces (e.g. CAT02, CMC2000, etc.; for an overview see *lx.cat._MCATS*) and several degree of adaptation formulas.
As an example, let's calculate the corresponding colors under CIE illuminant D65 (*xyzw2*) of the 8 CIE TCS illuminated by the spectra in *REF* (*xyzw1*) using a two-step (*catmode = '1>0>2'*) chromatic adaptation transform [Smet 2017] with CAT02 sensors and with an assumed adapting field luminance, *La* of resp. 100 cd/m² and 500 cd/m²:

```
# Original illumination condition:
xyzw1 = xyz_REF

# New illumination condition:
D65 = lx._CIE_ILLUMINANTS['D65']
xyzw2 = lx.spd_to_xyz(D65,\
                      cieobs = cieobs,\
                      relative = True)
# Apply von Kries CAT:
xyz_TCS8_REFc= lx.cat.apply(xyz_TCS8_REF,\
               xyzw1 = xyzw1,\
               xyzw2 = xyzw2,\
               catmode = '1>0>2',\
               cattype='vonkries',\
              D = None,\
               La = np.array([[100,500]]))
```

The original and corresponding colors can be plotted (see Fig. 5) as follows:

```
# convert to CIE 1976 u'v':
Yuv_TCS8_REF =
 lx.xyz_to_Yuv(xyz_TCS8_REF)
Yuv_TCS8_REFc =
 lx.xyz_to_Yuv(xyz_TCS8_REFc)

# create figure with 1 axes:
ax = plt.figure().add_subplot(111)
ax.plot(Yuv_TCS8_REF[:,0,1],\
        Yuv_TCS8_REF[:,0,2],\
        color ='r',\
        marker = 'o',\
        linestyle = 'none')
ax.plot(Yuv_TCS8_REFc[:,0,1],\
        Yuv_TCS8_REFc[:,0,2],\
        color ='b',\
        marker ='d',\
        linestyle = 'none')
ax.set_xlabel("u'") # set x-axis label
ax.set_ylabel("v'") # set y-axis label

# Plot spectrum, BB and DL loci:
lx.plotSL(cieobs = cieobs,\
          cspace = 'Yuv', \
          DL = True, \
          BBL = True)
```

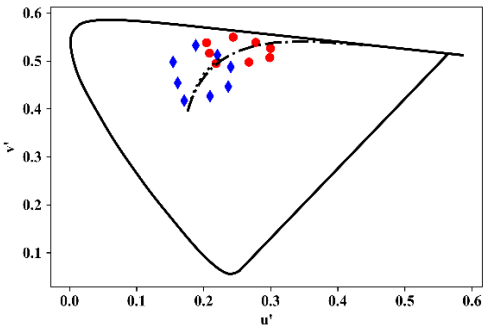


**Fig. 5. Plot u'v' coordinates of CIE TCS8 bef ore (red circles) and after (blue diamonds) a von Kries CAT from a 3000 K blackbody rad iator to CIE illuminant D65.**

## 4.5 Color appearance models (.cam)

Several color appearance models (CAM), such as CIECAM02 [Moroney 2002], CAM02-UCS/LCD/SCD [Luo 2006], CAM16 & CAM16-UCS/LCD/SCD[Li 2017], CAM15u [Withouck 2015], CAMSWW2016[Smet 2016], have been implemented. In this section, the use of the *xyz_to_jab…* type wrapper functions will be demonstrated.

For example, let's look at the conversion from XYZ tristimulus value to CIECAM02 $J,a_M,b_M$ coordinates and plot the results in Fig. 6. (Similar functionality is supported for the other CAMs):

```
# Use default conditions
# (see lx.cam._CAM_DEFAULT_CONDITIONS):
JabM_TCS8_REF =
     lx.xyz_to_jabM_ciecam02(xyz_TCS8_REF)

# Create a dictionary with user defined
viewing conditions:
usr_cnds = {'D': None,\
            'Dtype': None,\
            'La': 500.0,\
            'Yb': 40.0,\
            'surround': 'dim'}
```

```
JabM_TCS8_REF_user =
 lx.xyz_to_jabM_ciecam02(xyz_TCS8_REF,\
                     xyzw = xyz_REF,\
                     conditions = usr_cnds)

# Plot color coordinates in a 3d graph:
ax = plt.figure()
ax.add_subplot(111,projection = '3d')
ax.plot(JabM_TCS8_REF[:,0,1],\
        JabM_TCS8_REF[:,0,2],\
        JabM_TCS8_REF[:,0,0],\
         color = 'r', marker = 'o')
ax.plot(JabM_TCS8_REF_user[:,0,1], \
        JabM_TCS8_REF_user[:,0,2],\
        JabM_TCS8_REF_user[:,0,0], \
        color ='b', marker ='d')
ax.set_zlabel("J'")
ax.set_xlabel("a'")
ax.set_ylabel("b'")
```

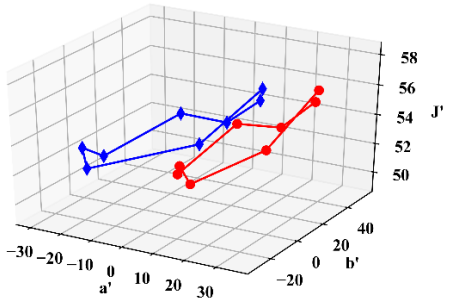


**Fig. 6. Plot of CIE TCS8 samples in CIECAM02 J,$a_M$,$b_M$. Red circles: default viewing conditions, blue diamonds: user defined conditions.**

## 4.6 Color difference calculation (.deltaE)

The *deltaE* sub-package supports calculation of color differences in various color spaces or color appearance spaces.

The widely used *DE2000* color difference formula [Sharma 2005] has been implemented and supports direct input of tristimulus values or CIE 1976 *L*a*b** color coordinates. For example, let's calculate the color difference between the color samples illuminated by the first and last spectrum in *REF*.

```
# Calculate DE2000:
DE00 =
   lx.deltaE.DE2000(xyz_TCS8_REF[:,0,:], \
                    xyz_TCS8_REF[:,-1,:],\
                    xyzwt = xyz_REF[:1,:],\
                    xyzwr = xyz_REF[-1:,:])

# Calculate DE2000,
# and also output rms average (DE00a),
# use default KLCH scaling factors:
DE00i, DE00a =
    lx.deltaE.DE2000(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     avg = lx.math.rms, \
                     out = 'DEi,DEa', \
                     KLCH = [1,1,1])

# Calculate DE2000,
# and also output rms average (DE00a),
# but ignore L*,
# and double the kC scaling factor:
DE00i_ab, DE00a_ab =
    lx.deltaE.DE2000(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     avg = lx.math.rms,\
                     out = 'DEi,DEa',\
                     KLCH = [1,2,1],\
                     DEtype = 'ab')
```

The function *DE_camucs* supports calculation of color differences in color appearance spaces (CIECAM02, CAM16, CAM02ucs, CAM16lcd, ...):

```
# Calculate DE in CIECAM02
# using default CAM viewing conditions:
DEi_ciecam02 =
 lx.deltaE.DE_camucs(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out ='DEi', \
                     camtype = 'ciecam02',\
                     ucstype = 'none')

# DE in CAM02-ucs (def. CAM viewing conds.)
DEi_cam02ucs =
 lx.deltaE.DE_camucs(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:], \
                     xyzwr = xyz_REF[-1:,:],\
                     out ='DEi', \
                     camtype = 'ciecam02', \
                     ucstype = 'ucs')

# DE in CAM16-lcd (def. CAM viewing conds.)
DEi_cam16lcd =
 lx.deltaE.DE_camucs(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out ='DEi',\
                     camtype = 'cam16',\
                     ucstype = 'lcd')
```

A handy function, *DE_cspace()*, exists that can calculate DE2000, CAMUCS-type color differences, but also color differences in any color space or chromaticity diagram for which the function *xyz_to_...* is defined. The input arguments are as before, with the additional argument '*tf*' specifying the color difference type (i.e. *'DE2000'* or *'camucs'*) or color space (e.g. *'Yuv'* or *'Lab'*). When '*tf*' is different from *'DE2000'* or *'camucs'*, there is the possibility of using the *KLCH* argument to specify different weights for the lightness, chroma and hue dimensions.

```
# DE2000:
DEi_2000 =
 lx.deltaE.DE_cspace(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out = 'DEi',\
                     tf = 'DE2000')

# DE CAM02-ucs:
DEi_cam02ucs =
 lx.deltaE.DE_cspace(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out = 'DEi',\
                     camtype = 'ciecam02',\
                     ucstype = 'ucs',\
                     tf = 'camucs')

# DE CIE 1976 L*a*b*:
DEi_lab =
 lx.deltaE.DE_cspace(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out = 'DEi',\
                     tf = 'lab')

# DE CIE 1976 L*a*b*
# with kL, kC and kH weights different from
# unity (KLCH = [2,3,4]):
DEi_lab_KLCH =
 lx.deltaE.DE_cspace(xyz_TCS8_REF[:,0,:],\
                     xyz_TCS8_REF[:,-1,:],\
                     xyzwt = xyz_REF[:1,:],\
                     xyzwr = xyz_REF[-1:,:],\
                     out = 'DEi',\
                     tf = 'lab',\
                     KLCH = [2,3,4])
```

## 4.7 Color rendition measures (.cri)

The cri sub-package supports several color rendition metrics, such as CIE13.3-1995 Ra [CIE13.3-1995 1995], CIE224-2017 Rf [CIE224:2017 2017], IESTM30 Rf and Rg [David 2015; IES 2015], CRI2012 [Smet 2013], MCRI [Smet 2012] and CQS [Davis 2010]. It also contains handy functions to construct 'new' color fidelity type metrics very easily by changing for example the sample set, averaging method, color space, CIE observer, ...

Finally, a sub-package *cri.VFPX* also supports the calculation of vector field based and pixel based color shifts, of a metameric uncertainty index, ...

Wrapper functions have been created specifically dedicated to the standard definition of the published color rendition measures:

```
# Get some spectra from the IESTM30 light
# source database:
N = 4
spds = lx._IESTM3018['S']['data'][:N+1]

# CIE13.3-1995 Ra:
ciera = lx.cri.spd_to_ciera(spds)

# CIE224-2017 Rf:
cierf = lx.cri.spd_to_cierf(spds)

# IES-TM30-15 Rf:
iesrf_15 = lx.cri.spd_to_iesrf(spds,\
               cri_type = 'iesrf-tm30-15')

# IES-TM30-15 Rg:
iesrg_15 = lx.cri.spd_to_iesrg(spds,\
               cri_type = 'iesrf-tm30-15')

# IES-TM30-18 Rf:
iesrf_18 = lx.cri.spd_to_iesrf(spds)

# IES-TM30-18 Rg:
iesrg_18 = lx.cri.spd_to_iesrg(spds)
```

The use of the more versatile *spd_to_cri* function allows the calculation of a color fidelity value using all the same standard parameters as those of the metric

defined in *cri_type*, but with for example a change of the sample set from the IES99 to the IES4880, using a blackbody radiator as reference illuminant and using the *rms* instead of the *mean* to average the color differences before calculating the general color fidelity index. Requested output can be specified by *out*:

```
# Get IES 4880 reference set:
R = lx._CRI_RFL['ies-tm30']['4880']['5nm']

# Calculate custom Rf & Rg
# and cct & duv:
Rf_custom, Rg_custom, cct, duv =
           lx.cri.spd_to_cri(spds,\
                    cri_type = 'ies-tm30',\
                    out = 'Rf,Rg,cct,duv',\
                    avg = lx.math.rms,\
                    sampleset = R,\
                    ref_type = 'BB')
```

The LuxPy.cri sub-package also supports a function that calculates several color rendition index values and provides TM30-like graphical output. For example, the code below generates the output in Fig. 7.

```
# Calculate and plot IES TM30
# color rendition measures:
SPD = lx._CIE_ILLUMINANTS['F4']
data,_,_ = lx.cri.plot_cri_graphics(SPD)
```

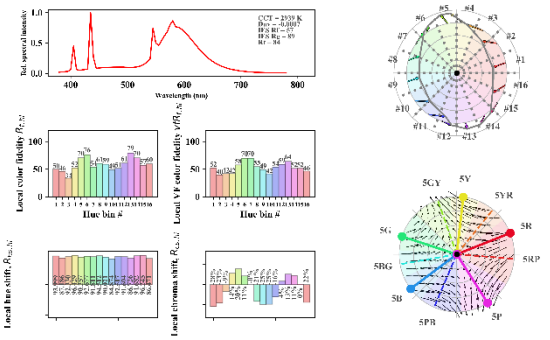

**Fig. 7. TM30 like graphic output of the color rendition properties of CIE illuminant F4. Top left: SPD with inline text of CCT, Duv, Rf, Rg and Rt (metameric uncertainty index). Mid row, left: Rfhi index values for hue bins 1-16. Mid row, right: same but base color shifts predicted by a vector field model. Bottom row left and right: local hue hue and chroma shifts. Top right: color graphic icon. Bottom right: base color shifts predicted by a vector field model together with the Munsell-5-Hue lines.**

The output format of and displayed information in some of the right hand graphs can be easily customized. For example, the graph in Fig. 8 was generated by the following code, which turns off the coloring of the right-hand side color icons, plots the vector field base color shifts in grey and plots the distortion of the color gamut using circle fields (the color of the distorted circles provides information on the size of the hue shift, red being larger):

```
data = lx.cri.plot_cri_graphics(SPD,\
                plot_bin_colors = False,\
                vf_plot_bin_colors = False,\
                vf_color = 'grey',\
                plot_CF = True)
```

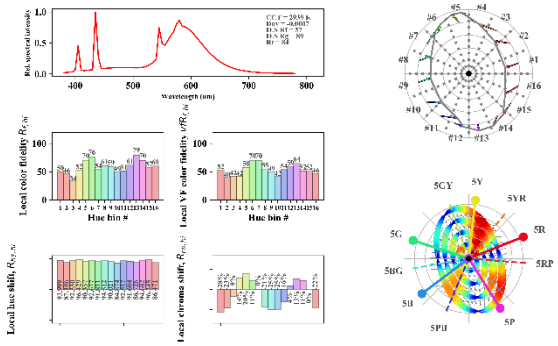

**Fig. 8. TM30 like graphic output of the color rendition properties of CIE illuminant F4. Similar as Fig. 7, but with hue coloring turned off in the right graphs, and with base color shift vectors overlayed with lines representing the distortion of concentric chroma circles due to the specific color rendition properties of the test light source. Hue shift information is represented by the width and coloring of the lines: wider, more reddish lines represent larger hue shifts.**

## 4.8 LuxPy classes

The basic use of the *SPD*, *XYZ* and *LAB* classes has already been described in section *3.2.3 Spectral and colorimetric data using LuxPy classes*. For further functionality the reader is referred to the help in the *__doc__* strings.

### 4.9 Calculation of photobiological quantities

Sub-package *.photbiochem* supports the calculation of various photobiological or related quantities, such as the Cyanosis Observation Index (COI)[AS/NZS1680.2.5 1997], the CIE photobiological quantities (α-opic irradiance and equivalent illuminance) defined in TN003:2015 [CIE-TN003:2015 2015; Lucas 2014] or the LRC's Circadian Stimulus (*CS*) and Circadian Lighting (*CLa*) [Rea 2012] from spectral data.

Let's start with the TN003:2015 α-actinic quantities, for which a single dedicated function has been implemented: *spd_to_aopicE()*. The input spectral irradiance can be in $W/m^2{\cdot}nm$ (default) or in $\mu W/cm^2{\cdot}nm$. The output is in SI units.

```
# Define a 4200 K blackbody radiator with
# a wavelength range between 378 nm and
# 782 nm, and a wavelength spacing of 1 nm:
BB = lx.blackbody(4200,\
                  lx.getwlr([378,782,1]))

# Calculate the a-opic irradiance Eea
# and equivalent illuminance Ee
# for the blackbody radiator spectrum
# normalized to 100 lx:
Ees,Es = lx.photbiochem.spd_to_aopicE(BB,\
                                      E = 100)
# Print results:
print("Photoreceptor cells:")
print(lx.photbiochem._PHOTORECEPTORS)
print("\na-opic irradiance symbols:")
print(lx.photbiochem._Ee_SYMBOLS)
print("a-opic irradiance values:")
print(Ees)
print("\na-opic illuminance symbols:")
print(lx.photbiochem._E_SYMBOLS)
print("a-opic illuminance values:")
print(Es)
```

```
Output:
Photoreceptor cells:
['l-cone', 'm-cone', 's-cone','rod',
'iprgc']

a-opic irradiance symbols:
['Ee,lc','Ee,mc','Ee,sc','Ee,r','Ee,z']
a-opic irradiance values:
[[0.17140925 0.15161959 0.04969815 0.12
215436 0.09792088]]

a-opic illuminance symbols:
['E,lc', 'E,mc', 'E,sc', 'E,r', 'E,z']
a-opic illuminance values:
[[99.22008705 94.01060178 65.86204629 8
6.16355622 81.51105241]]
```

The following code snippet illustrates the use of the *spd_to_COI_ASNZS1680* function that calculates the COI and can also output the CCT:

```
# Get a light source spectrum:
F4 = lx._CIE_ILLUMINANTS['F4']

# Calculate COI and CCT:
coi, cct =
   lx.photbiochem.spd_to_COI_ASNZS1680(F4)
```

The Circadian Stimulus (*CS*) and Circadian Light (*CLa*) quantities of a (set of) light source(s) can be calculated as follows:

```
# Import spd_to_CS_CLa_lrc for brevity:
from luxpy.toolboxes.photbiochem import
                          spd_to_CS_CLa_lrc

# Get a set of 3 light source spectra:
S = lx._IESTM3018['S']['data'][:4]

# Calculate CS and CLa:
CS1, CLa1 = spd_to_CS_CLa_lrc(El = S)

# The same, but this time with all
# 3 normalized to 100 lx:
CS2, CLa2 = spd_to_CS_CLa_lrc(El = S,\
                              E = 100)

# Or to different illuminance values:
CS3, CLa3 = spd_to_CS_CLa_lrc(El = S,\
                      E = [100, 200, 50])

# The input spectra can also be summed to
generate a composite spectrum for which the
CS and CLa will be calculated:
CS4, CLa4 = spd_to_CS_CLa_lrc(El = S,\
                      E = [100, 200, 50],\
                      sum_sources = True)
```

The output generated using *print* is given below:

```
Output:

CS1, CLa1:
[6.9688e-01 6.9545e-01 6.9309e-01]
[4.7973e+04 3.4060e+04 2.3242e+04]

CS2, CLa2:
[1.7518e-01 9.3232e-02 4.7315e-02]
[1.3149e+02 6.5061e+01 3.2918e+01]


CS3, CLa3:
```

```
[1.7518e-01 1.7478e-01 2.2740e-02]
[1.3149e+02 1.3113e+02 1.6379e+01]

CS4, CLa4:
[3.0622e-01] [2.8316e+02]
```

## 4.10 Spectrum creation and optimization (.spdbuild)

The *.spdbuild* sub_package supports creating Gaussian, monochromatic [Ohno 2005] and phosphor LED-type [Smet 2011] spectra, and combining them into multi-component spectra. It also supports automatic flux calculations to obtain a desired spectrum chromaticity. Additional objective functions can also be used to optimize the component fluxes to these additional constraints. This section will only briefly introduce some of its features. For more info the reader is referred to the *__doc__* strings of the module functions.

As a first example, let's start with the creation of 4 monochromatic LEDs and import the module into the global namespace for ease of use. The output is given in Fig. 9.

```
# Import module luxpy.spdbuild:
import luxpy.toolboxes.spdbuild as spb

# Set peak wavelengths:
peakwl = [450,530,560,610]

# Set Full-Width-Half-Maxima:
fwhm = [30,35,30,15]
S = spb.spd_builder(peakwl = peakwl,\
                    fwhm = fwhm)

# Plot component spds:
lx.SPD(S).plot()
```

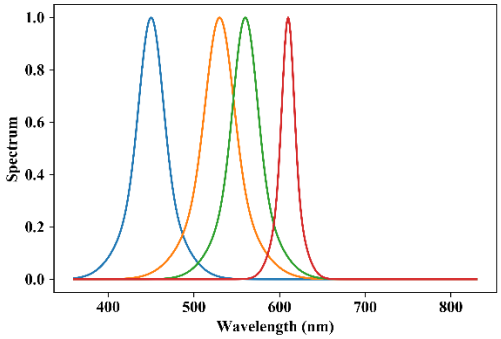


**Fig. 9. Four monochromatic leds generated with `spd_builder()`.**

Next, let's create 3 additional spectra: 2 phosphor type LED spectra (one with 1 phosphor and one with 2 phosphors) and one monochromatic LED-type spectrum (see Fig. 10 for output):

```
# peak wavelengths of monochromatic leds:
peakwl = [440,460,610]

# Full-Width-Half-Maximas:
fwhm = [30,35,15]

# Set phosphor strengths (i.e. alpha factor
# in Smet et al. 2011):
strength_ph = [1.5, 0.4, 0]

# Set phosphor 1 parameters:
strength_ph1 = [1, 1, 0] # beta factor
peakwl_ph1 = [530, 510, 510]
fwhm_ph1 = [60, 60, 60]

# Set phosphor 2 parameters:
strength_ph2 = [2, 1, 0] # beta factor
peakwl_ph2 = [600, 590, 590]
fwhm_ph2 = [70, 70, 70]

S = spb.spd_builder(peakwl = peakwl,\
                    fwhm = fwhm,\
                    strength_ph = strength_ph, \
                    strength_ph1 = strength_ph1,\
                    peakwl_ph1 = peakwl_ph1,\
                    fwhm_ph1 = fwhm_ph1,\
                    strength_ph2 = strength_ph2,\
                    peakwl_ph2 = peakwl_ph2,\
                    fwhm_ph2 = fwhm_ph2)

# Plot component spds:
lx.SPD(S).plot()
```

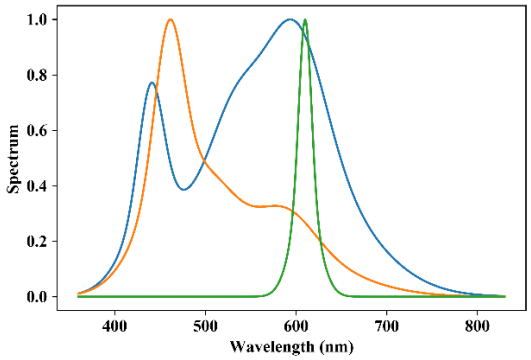


**Fig. 10. Two phosphor type and one monochromatic LED spectrum generated with `spd_builder()`.**

One can also set them a specific target chromaticity, specified by a *'target'* input argument (*'tar_type'* specifies the type of target, e.g. 'cct', 'Yxy', ... and *'cspace_bwtf'* is dict specifying the

parameters need for the backward transformation called in *colortf()* inside *spd_builder()*). The code below calculates the fluxes required for a *CCT* = 3500 K (*Duv* = 0). There are only two spectra plotted in Fig. 11 as any set of 3 spectral components, i.e. a monochromatic led and 2 phosphors, for which the target is out of gamut results in an array of *NaN*'s.

```
# CIE standard 2° observer:
cieobs = '1931_2'
cspace_bwtf = {'cieobs' : cieobs,\
               'mode' : 'search'}

S = spb.spd_builder(peakwl = peakwl,\
              fwhm = fwhm,\
              strength_ph = strength_ph,\
              strength_ph1 = strength_ph1,\
              peakwl_ph1 = peakwl_ph1,\
              fwhm_ph1 = fwhm_ph1,\
              strength_ph2 = strength_ph2,\
              peakwl_ph2 = peakwl_ph2,\
              fwhm_ph2 = fwhm_ph2,\
              target = 3500,\
              tar_type = 'cct',\
              cspace_bwtf = cspace_bwtf)

# Plot component spds:
lx.SPD(S).plot()
```

The package also supports spectra optimization beyond the target chromaticity when there are more than 3 spectral components provided.

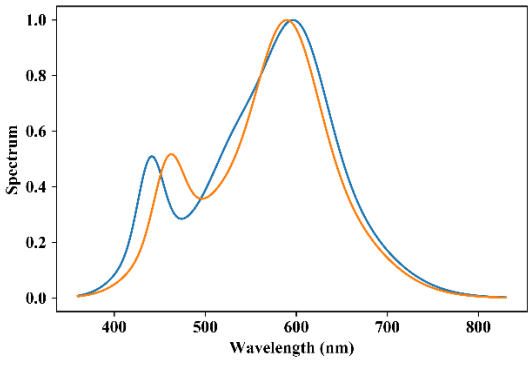


**Fig. 11. Two phosphor type LED spectra generated with spd_builder() for which the target chromaticity (CCT = 3500 K, Duv = 0) was inside the gamut spanned by its components.**

The following code block provides a practical example of flux optimization with fixed component spectra using objective functions for a target *CCT* of 4000 K (*Duv* = 0). The objective functions and their target values are specified as two lists. For this example the IES TM30 Rf and Rg color rendition measures [IES 2015] were chosen with target values of resp. 90 and 110. The output is plotted in Fig. 12. The optimization is done using the *'2mixer'* optimizer (another option is *'3mixer'*). For more info on the optimization algorithms the reader is referred to the *__doc__* string of the *lx.spdbuild.spd_optimizer* function.

```
# Set target chromaticity:
target = 3500

# Set objective functions and targets:
obj_fcn = [lx.cri.spd_to_iesrf,\
           lx.cri.spd_to_iesrg]
obj_tar_vals = [90,110]

# Set component model parameters:
peakwl = [450,530,560,610]
fwhm = [30,35,30,15]

# Perform optimization:
S,_ = spb.spd_optimizer(target = target, \
              tar_type = 'cct',\
              cspace_bwtf = cspace_bwtf,\
              optimizer_type = '2mixer',\
              peakwl = peakwl,\
              fwhm = fwhm,\
              obj_fcn = obj_fcn,\
              obj_tar_vals = obj_tar_vals)

# Check output agrees with target:
xyz = lx.spd_to_xyz(S, relative = False,\
                    cieobs = cieobs)
cct = lx.xyz_to_cct(xyz, cieobs = cieobs,\
                    mode = 'search')[0,0]
Rf, Rg = [fcn(S)[0,0] for fcn in obj_fcn]

print('\nResults (optim,target):')
print("cct(K):
({:1.1f},{:1.1f})".format(cct, target))

print("Rf:   ({:1.2f},{:1.2f})".format(Rf,
obj_tar_vals[0]))

print("Rg:  ({:1.2f},  {:1.2f})".format(Rg,
obj_tar_vals[1]))

#plot spd:
plt.figure()
lx.SPD(S).plot()
```

```
Output:
Optimization terminated successfully.
    Current function value: 0.077599
    Iterations: 23
    Function evaluations: 48

Results (optim,target):
cct(K): (3500.0,3500.0)
Rf: (85.98,90.00)
Rg: (103.02, 110.00)
```

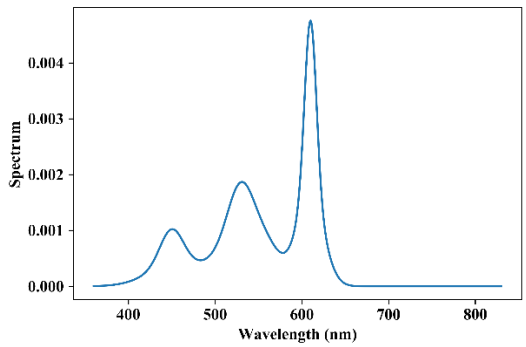


**Fig. 12. Spectrum optimized from predefined, fixed component spectra for a CCT = 3500 K and IES TM30 Rf and Rg objective functions.**

In addition to lux-only optimization, the peak wavelengths and full-width-half-maxima of the components can also be optimized. The example below calculates a spectrum with a *CCT* of 3500 K (*Duv* = 0) and optimized for IES TM30 Rf and Rg target values of resp. 90 and 110. The spectrum is built from 6 components with free peak wavelength, *FWHM* and fluxes. The *verbosity* argument toggles intermediate output of objective functions values on and off. The optimized spectrum is shown in Fig.13.

```
# Perform optimization:
S,_ = spb.spd_optimizer(target = target, \
            tar_type = 'cct',\
            cspace_bwtf = cspace_bwtf,\
            optimizer_type = '2mixer',\
            N_components = 6,\
            obj_fcn = obj_fcn,\
            obj_tar_vals = obj_tar_vals,\
            verbosity = 0)

# Check output agrees with target:
xyz = lx.spd_to_xyz(S, relative = False,\
                    cieobs = cieobs)
cct = lx.xyz_to_cct(xyz, cieobs = cieobs,\
                    mode = 'search')[0,0]
Rf, Rg = [fcn(S)[0,0] for fcn in obj_fcn]
```

```
print('\nResults (optim,target):')

print("cct(K):
({:1.1f},{:1.1f})".format(cct, target))

print("Rf:   ({:1.2f},{:1.2f})".format(Rf,
obj_tar_vals[0]))

print("Rg: ({:1.2f}, {:1.2f})".format(Rg,
obj_tar_vals[1]))

#plot spd:
plt.figure()
lx.SPD(S).plot()
```

```
Output:
Optimization terminated successfully.
    Current function value: 0.002494
    Iterations: 1356
    Function evaluations: 2037

Results (optim,target):
cct(K): (3500.0,3500.0)
Rf: (89.85,90.00)
Rg: (109.79, 110.00)
```

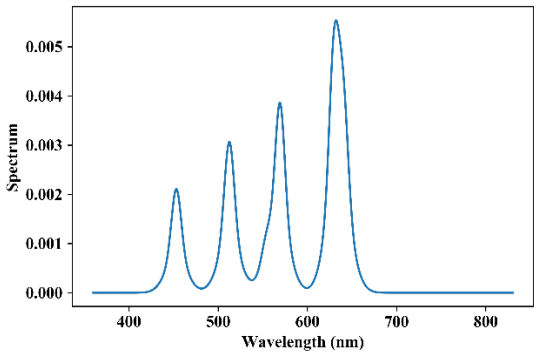


**Fig. 13. Spectrum optimized from predefined, fixed component spectra for a CCT = 3500 K and IES TM30 Rf and Rg objective functions.**

## 4.11 Simulation and rendering of hyper-spectral images (.hypspcim)

Luxpy contains a module that can simulate a hyper-spectral image of any regular sRGB image by mapping the color coordinates of each pixel in the image to a sample from a spectral reflectance set that is near-metameric under CIE D65 or some other user specified reference spectrum. The algorithm uses SciPy's cKDTree to find the closest neighbours to each pixel's color coordinates and then calculates a weighted mean of the nearest spectral reflectance functions in a set. The

*render_image* function can then simulate the effect of a light source on the color appearance of the imaged scene to explore the source's color rendition properties. Note though that no account is taken (yet) for out of gamut values (i.e. no extrapolation has been implemented and out of gamut color coordinates are simply clipped). The user can provide a set of reflectance functions, or else the IESTM30 4880 Reference set [IES 2015] is used as default. There is also the option to apply a von Kries chromatic adaptation to the reference spectrum chromaticity, whereby the user can specify the degree of adaptation by input argument *D*. The following illustrates the use of the *render_image* function:

```
from luxpy import hypspcim

# Generate 4000 K RGB LED spectrum:
S=spb.spd_builder(peakwl = [450,530,610],\
                  fwhm = [10,10,10],\
                  target = 4000,\
                  tar_type = 'cct')


# Specify filename (+path) to some image or
# read in ndarray with image data.
# In this case the filename of the default
# image was used for simplicity:
img = hypspcim._HYPSPCIM_DEFAULT_IMAGE



# Calculate hyper-spectral version
# of the img image:
img_hyp = hypspcim.render_image(img = img,\
                      cspace = 'ipt',\
                      spd = S, \
                      D = 1, \
                      show = True,\
                      stack_test_ref = 21)

print(img_hyp.shape)
```

```
Output:
(681, 1024, 401)
```

In the code above the *stack_test_ref* argument specifies how the rendered images under the test and reference light sources should be stacked. The output of the rendering is shown in Fig. 14.


**Fig. 14. Rendering results of simulated hyper-spectral images. Top: using the 4000 K RGB LED test light source spectrum. Bottom: using the reference (D65) spectrum. A von Kries chromatic adaptation with degree of adaptation set to *D* = 1 has been applied.**

## 5. Summary

This paper described the installation of the LuxPy toolbox and provided an overview of its basic design and functionality. It also demonstrated its basic usage by providing several code examples using a Jupyter notebook. It is hoped that the LuxPy package – written in an easy to learn open-source programming language – would be useful to many lighting and color researchers and professionals, and would stimulate collaborative research and ease sharing calculations and analyses.

## ORCID

Kevin A.G. Smet http://orcid.org/0000-0003-3825-6274

## Funding

The author reports no funding for the preparation of this manuscript.

## Disclosure Statement

The authors report no declarations of interest.

## REFERENCES

AS/NZS1680.2.5 (1997). INTERIOR LIGHTING PART 2.5: HOSPITAL AND MEDICAL TASKS.

Asano Y, Fairchild MD, and Blondé L (2016). Individual Colorimetric Observer Model. PLoS One *11*, 1–19.

CIE-TN003:2015 (2015). Report on the first international workshop on circadian and neurophysiological photometry, 2013 (Vienna, Austria).

CIE13.3-1995 (1995). Method of Measuring and Specifying Colour Rendering Properties of Light Sources (Vienna, Austria: CIE).

CIE15-2004 (2004). Colorimetry (Vienna, Austria: CIE).

CIE224:2017 (2017). CIE 2017 Colour Fidelity Index for accurate scientific use (Vienna, Austria).

David A, Fini PT, Houser KW, Ohno Y, Royer MP, Smet KAG, Wei M, and Whitehead L (2015). Development of the IES method for evaluating the color rendition of light sources. Opt. Express *23*, 15888–15906.

Davis W, and Ohno Y (2010). Color quality scale. Opt. Eng. *49*, 33602–33616.

IES (2015). IES-TM-30-15: Method for Evaluating Light Source Color Rendition (New York, NY: The Illuminating Engineering Society of North America).

Li C, Li Z, Wang Z, Xu Y, Luo MR, Cui G, Melgosa M, Brill MH, and Pointer M (2017). Comprehensive color solutions: CAM16, CAT16, and CAM16-UCS. Color Res. Appl. 1–16.

Lucas RJ, Peirson SN, Berson DM, Brown TM, Cooper HM, Czeisler CA, Figueiro MG, Gamlin PD, Lockley SW, O'Hagan JB, et al. (2014). Measuring and using light in the melanopsin age. Trends Neurosci. *37*, 1–9.

Luo MR, Cui G, and Li C (2006). Uniform colour spaces based on CIECAM02 colour appearance model. Color Res. Appl. *31*, 320–330.

MathWorks Inc. (2018). Matlab.

Moroney N, Fairchild MD, Hunt RWG, Li C, Luo MR, and Newman T (2002). The CIECAM02 color appearance model. IS&T/SID Tenth Color Imaging Conf. 23.

Ohno Y (2005). Spectral design considerations for white LED color rendering. Opt. Eng. *44*, 111302.

Ohno Y (2014). Practical Use and Calculation of CCT and Duv. LEUKOS *10*, 47–55.

Rea MS, Figueiro MG, Bierman A, and Hamner R (2012). Modelling the spectral sensitivity of the human circadian system. Light. Res. Technol. *44*, 386–396.

Sharma G, Wu W, and Dalal EN (2005). The CIEDE2000 color-difference formula: Implementation notes, supplementary test data, and mathematical observations. Color Res. Appl. *30*, 21–30.

Smet K, Ryckaert WR, Pointer MR, Deconinck G, and Hanselaer P (2011). Optimal colour quality of LED clusters based on memory colours. Opt. Express *19*, 6903–6912.

Smet K, Schanda J, Whitehead L, and Luo R (2013). CRI2012: A proposal for updating the CIE colour rendering index. Light. Res. Technol. *45*, 689–709.

Smet KAG, Ryckaert WR, Pointer MR, Deconinck G, and Hanselaer P (2012). A memory colour quality metric for white light sources. Energy Build. *49*, 216–225.

Smet KAG, Webster MA, and Whitehead LA (2016). A simple principled approach for modeling and understanding uniform color metrics. J. Opt. Soc. Am. A *33*, A319–A331.

Smet KAG, Zhai Q, Luo MR, and Hanselaer P (2017). Study of chromatic adaptation using memory color matches, Part I: neutral illuminants. Opt. Express *25*, 7732–7748.

Withouck M, Smet KAG, Ryckaert WR, and Hanselaer P (2015). Experimental driven modelling of the color appearance of unrelated self-luminous stimuli: CAM15u. Opt. Express *23*, 12045–12064.

www.libreoffice.org (2018). LibreCalc.