# Data collection and preprocessing 

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## Playing with multi-dimensional data

- Part 1: some real-life datasets, surveys and queries.
- Part 2: visualising and dimensionality reduction, PCA, kPCA



## Part 1: surveys, databases, queries \& thereabouts

General problem: we have big heaps of data produced by surveys/experiments and need to make sense of them.

## Example from astro: spectra, fluxes, colours.



Spectrum: blueprint of an object (more or less). Magnitudes: what we get most of the time.


From big data to science: discover, classify, characterise. NB: light-curve data (left) don't always have the same number of points!

Various magnitude systems for different uses ${ }^{1}$. Each magnitude has a central wavelength and a width.


Bessell. MS. 2005
Amu. Rev. Astron Astrophys. 43: 293-336
${ }^{1}$ If you're really, really curious: Bessel, M. S. 2005, ARA\&A, 43, 293


From big data to science: finding rare objects/events. (these ones are very rare)

## OK, but where do we begin???

Different experiments//surveys gather different kinds of info. We "just" need to grab it...


## Telescope//experiment (pipelines) $\mapsto$ data, various formats (database) $\mapsto$ catalog tables

| \|ID ra dec class | subClass |
| :--- | :--- | :--- |
| 1237678661968265435 | 16.878845 |
| 1237678623308578947 | 17.145415 |
| 1237678623308644624 | 17.274179 |
| 12376786277177328 | 17.297309 |
| 1237678661968396491 | 1.230792 |
| 1237678661968331270 | 17.112495 |
| 1237678661968265425 | 16.932224 |
| 1237678622771642595 | 16.988994 |
| 12376786196826378 | 16.918275 |
| 1237678622771642499 | 1.963018 |
| 1237678622771707950 | 17.151402 |
| 1237669702124241089 | 15.152699 |
| 1237669702124109952 | 14.874624 |
| 1237669702124175820 | 15.049748 |

$z \quad$ zerr
5.0594924
5.2240461
5.1563299
4.7099285
4.9492185
4.895233
4.9781829
4.8418012
5.009444
4.7222274
4.8186359
7.2441582
7.3149651
7.194844


## Queries

Sometimes you can do a bulk download of a catalog table, sometimes it's unfeasible or unnecessary.
SQL: Structured Query Language. Basic syntax:
SELECT \{fields\} FROM \{table\} WHERE \{conditions\}

```
SELECT TOP 100
    objID, ra ,dec
FROM
    PhotoPrimary
WHERE
    ra > 185 and ra < 185.1
    AND dec > 15 and dec < 15.1
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    ra > 185 and ra < 185.1
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## Slightly more complicated:

```
    SELECT D.coadd_object_id, W.cntr, D.alphawin_j2000 as
desra, D.deltawin_j2000 as desdec, D.mag_auto_i,
W.w1mpro, W.w2mpro
FROM des_dr1.main AS D
JOIN des_dr1.des_allwise AS W on
W.coadd_object_id=D.coadd_object_id
WHERE ( D.galactic_b<-20.0 AND D.mag_auto_i>8.0 AND
D.deltawin_j2000>-60.0 AND D.deltawin_j2000<-55.0 )
Q: how many differences can you spot with the simplest query?
```


## Many examples here:

http://skyserver.sdss.org/dr8/en/help/docs/realquery.asp
Quote of the day:
"Most of the Al you may need is an SQL SELECT followed by an ORDER BY clause"

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## Exercise

To familiarise with it a bit: Let's have a look at the SDSS

- Have a look at the Schema Browser for the PhotoObj and SpecPhoto tables.
- Query coordinates (ra, dec) and PSF magnitudes in $u-, g-$, $r-, i-, z$-bands, plus spectroscopic redshift, for ten thousand object with CLASS==' QSO', ten thousand with CLASS==' GALAXY', ten thousand with CLASS==' STAR'. You can use the web query page here . ${ }^{2}$
- Q: how well can you fit the redshift using only the magnitudes above? How well can you fit the class, given only the magnitudes?
- Repeat but also using magnitudes w1mpro and w2mpro from AllWISE.

Various examples of SDSS queries here

[^0]Can't we do it in python?

- For access to SQL servers, you can use sqlite (ask Carl!).
- For astronomical surveys, you can use astroqueries. Some examples given in ExampleQueries.txt , courtesy of Zoe Ansari and Sofie H. Bruun (DARK-NBI).


## Part 2: handling

OK, I have my table: now what?

- First things first: look at it!

Do the entries make sense? Are there any missing entries? Are some lines redundant?

- Second: plot familiar (and unfamiliar) stuff.




## Easy things first: plot feature vs feature:



Python tips and tricks: you should do it yourselves, but someone has already done it for you...

1. Pair plots (with seaborn)
https://seaborn.pydata.org/generated/seaborn.pairplot.html
import seaborn as sns; sns.set(style="ticks", color_codes=True) iris = sns.load_dataset("iris")
g = sns.pairplot(iris, hue="species", palette="husl")
2. Corner plots (with corner)
https://corner.readthedocs.io/en/latest/pages/quickstart.html
import corner
fig = corner.corner(samples, labels=["\$m\$", "\$b\$", "\$\ln<br>,f\$"])
fig.show()

[^1]

But how do I decide which features are important? Should I plot all of them?!
What if l'm dealing with collections of pictures instead of tables with some columns?

Common issue, 1: the dataset may be easier to crunch in a different
coordinate system.
Common issue, 2: are there any combinations of features that
maximize information?

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Sometimes you don't need hundreds of features:


This is actually done with something more advanced (Kingma \& Welling 2014), but still...

## Linear: Principal Component Analysys (PCA)

The maths: we want to transform our feature vectors $\left\{\mathbf{x}_{i} \in \mathbb{R}^{p}\right\}_{i=1, \ldots, N}$ into others $\left\{\mathbf{f}_{i} \in \mathbb{R}^{p}\right\}_{i=1, \ldots, N}$ that are uncorrelated. How to? Find eigenvectors of the covariance matrix:

$$
\begin{equation*}
C_{k, l}=\frac{1}{N} \sum_{i=1}^{N} x_{i, k} x_{i, l} \tag{1}
\end{equation*}
$$

$$
\begin{equation*}
\mathbf{C} \mathbf{v}_{k}=\lambda_{k} \mathbf{v}_{k} \tag{2}
\end{equation*}
$$

The eigenvectors are the principal components.
Fraction of explained variance:

$$
\begin{equation*}
\operatorname{var}_{(r)}:=\frac{\sum_{k=1}^{r} \lambda_{k}}{\sum_{k=1}^{p} \lambda_{k}} \tag{3}
\end{equation*}
$$

NB do you need to standardize your dataset?

## Example on (simple stuff) images: ${ }^{3}$



[^2]
## Example (from scikit-learn):4

```
>>> import numpy as np
>>> from sklearn.decomposition import PCA
>> X = np.array([[-1, -1], [-2, -1], [-3, -2], [1, 1], [2, 1], [3, 2]])
>> pca = PCA(n_components=2)
>>> pca.fit(X)
PCA(copy=True, iterated_power='auto', n_components=2, random_state=None,
    svd_solver='auto', tol=0.0, whiten=False)
>>> print(pca.explained_variance_ratio_)
[0.9924... 0.0075...]
>>> print(pca.singular_values_)
[6.30061... 0.54980...]
```


## Methods

| fit (X[, y]) | Fit the model with X. |
| :--- | :--- |
| $\mathbf{f i t \_ t r a n s f o r m ~ ( X [ , ~ y ] ) ~}$ | Fit the model with X and apply the dimensionality reduction on X. |
| get_covariance () | Compute data covariance with the generative model. |
| get_params ([deep]) | Get parameters for this estimator. |
| get_precision () | Compute data precision matrix with the generative model. |
| inverse_transform (X) | Transform data back to its original space. |
| score (X[, y]) | Return the average log-likelihood of all samples. |
| score_samples (X) | Return the log-likelihood of each sample. |
| set_params (**params) | Set the parameters of this estimator. |
| transform (X) | Apply dimensionality reduction to X. |

## Q: Run a PCA on the quark data table, see where the ' 1 ' and ' 0 ' subsamples lie. <br> 

## Bonus track: kPCA

How it works: ${ }^{5}$


Projection by KPCA


1st principal component in space induced by $\phi$


Original spate after inver inversent transform

${ }^{5}$ You can find code for this example on the scikit-learn website.

How the 'kernel trick' works: map feature space $\Phi: \mathbb{R}^{p} \mapsto \mathcal{H}$ to very-high-dimensional space with its own scalar product $k\left(\mathbf{x}_{i}, \mathbf{x}_{j}\right)=\left\langle\Phi\left(\mathbf{x}_{i}\right), \Phi\left(\mathbf{x}_{j}\right)\right\rangle$. Diagonalize a *big* matrix

$$
\begin{array}{r}
K_{i, j}=(1 / N) k\left(\mathbf{x}_{i}, \mathbf{x}_{j}\right) \\
K \mathbf{a}=\lambda \mathbf{a} \tag{5}
\end{array}
$$

Then the components of a given feature vector $\Phi(\mathbf{f})$ in this space, relative to $r$-th component, are

$$
\begin{equation*}
t_{r}=\left\langle\mathbf{a}_{r}, \Phi(\mathbf{f})\right\rangle=\sum_{i=1}^{N} a_{r, i} k\left(\mathbf{x}_{i}, \mathbf{f}\right) \tag{6}
\end{equation*}
$$

Theorem: everything exists if $k(\bullet, \bullet)$ is semi-positive definite. Q: Run a (k)PCA on the b-quark data table, try to separate the jets. Q: Run a (k)PCA on the SDSS data table, try to separate the classes.

## Summary

So to sum it up:
© data are ugly.
(3) know where your data come from!
© inspect your data tables, plot stuff.
( One method does not necessarily fit every purpose.
(0) there is already technology to parse tables, if needed (SQL and thereabouts).
(C) datasets can be very-high-dimensional

Linear: PCA; non-linear: kPCA (and tSNE, and UMAP...)

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[^0]:    ${ }^{2}$ To query and save heavier stuff, have a loot at CasJobs!

[^1]:    >> $g$ = sns.pairplot(iris, hue="species", palette="hus(")

[^2]:    ${ }^{3}$ That's from an old paper of mine, you don't really need to know-about it.

