

# Exploration of FELLOW trait dataset

The objective of this document is to:

- visually explore the trait database
- understand trait coverage / data gaps
- check for possible inconsistencies

## Description of the species list

We compiled the species lists from 31 datasets. After cleaning and harmonization, there were 2102 unique taxa.

FAMILY	GENUS	SPECIES	SUBSPECIES	VARIETY
15	261	1694	121	11

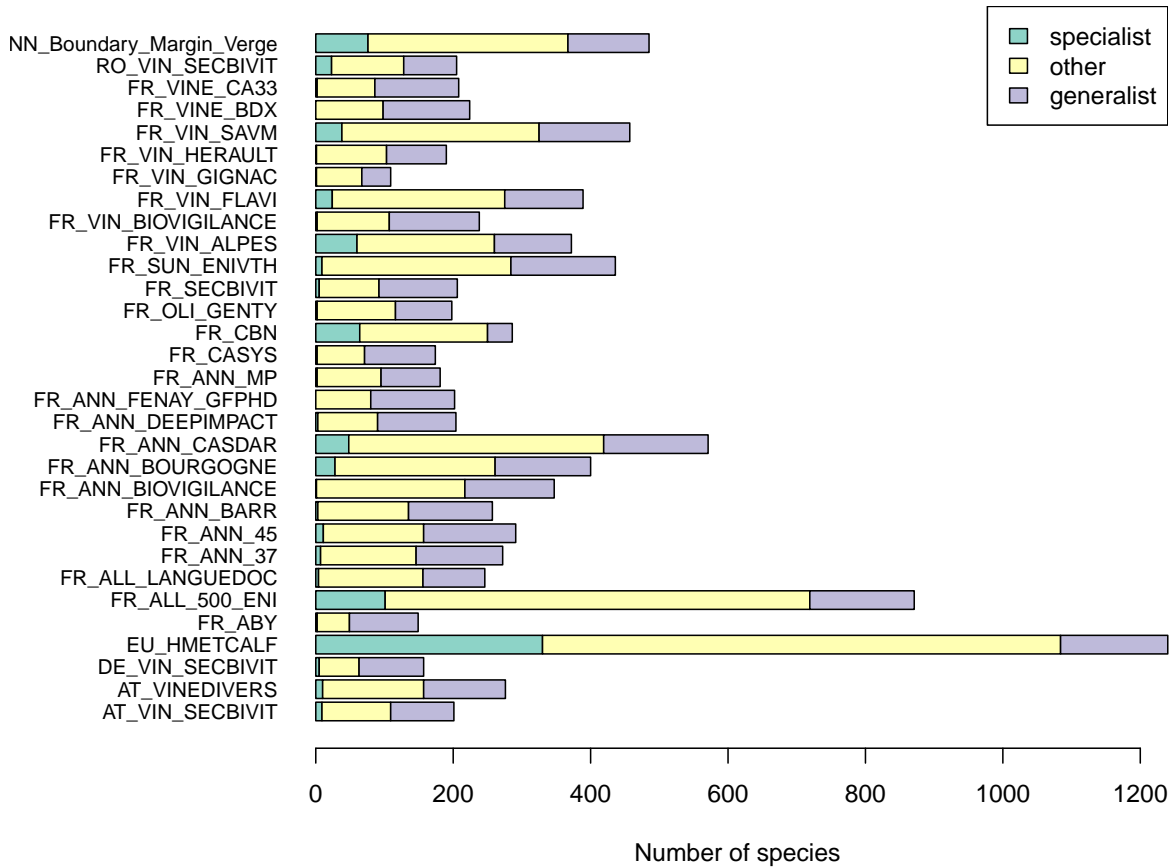
Let's define:

*specialist*: a taxa that occurred only in a single database

*generalist*: a taxa that is listed in 50% of the databases (16 out of 31)

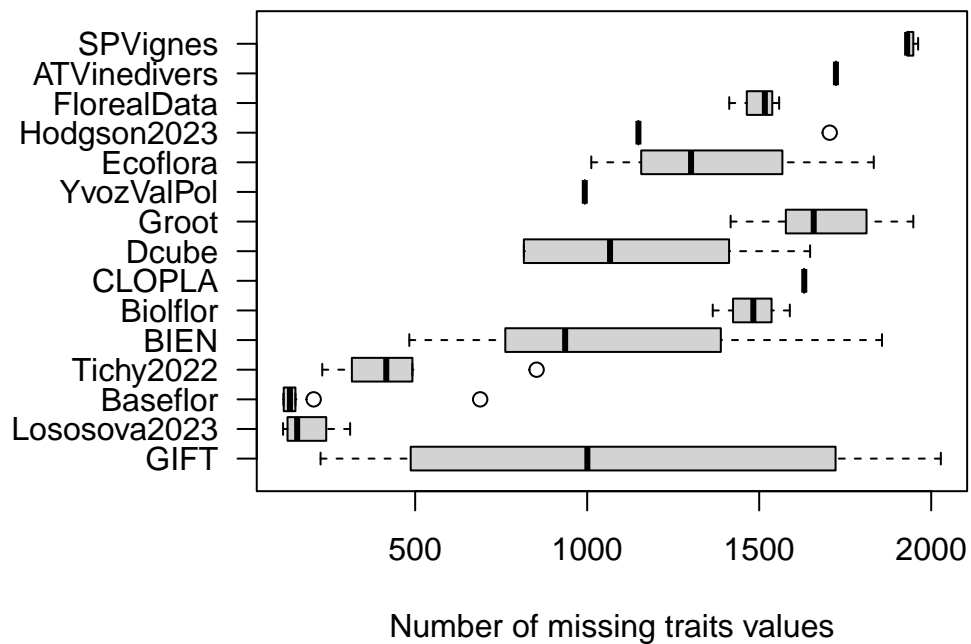
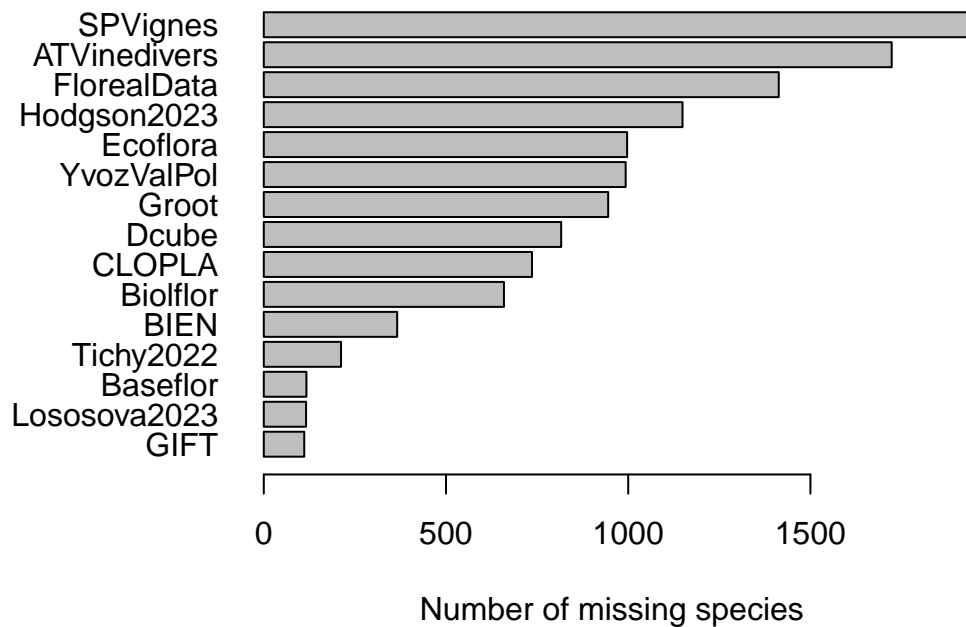
sp\_class

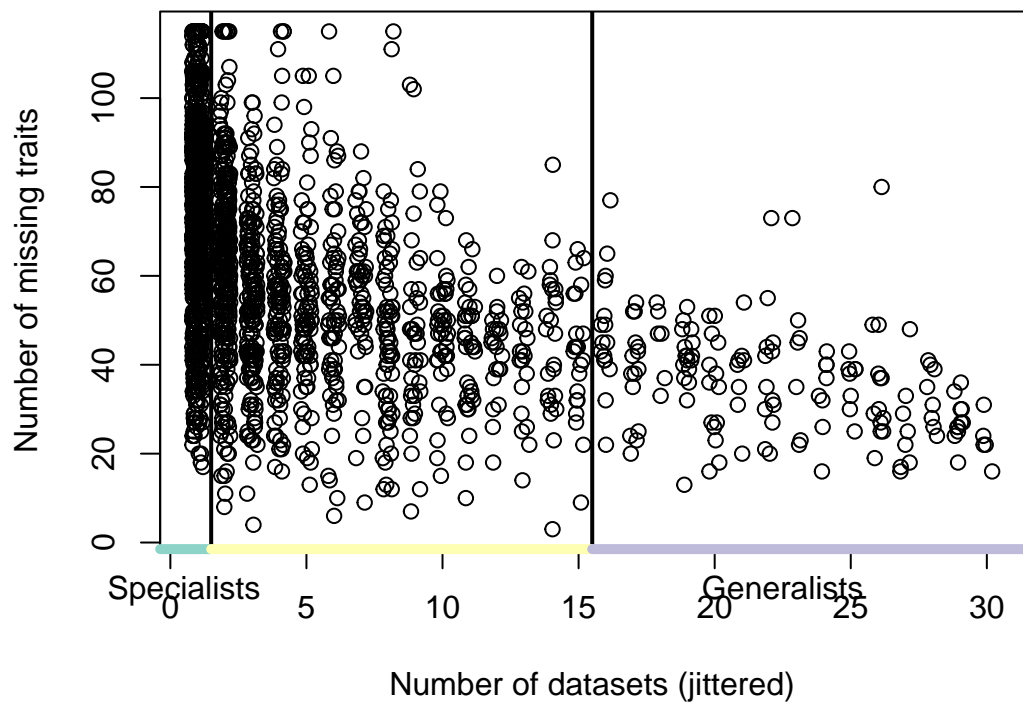
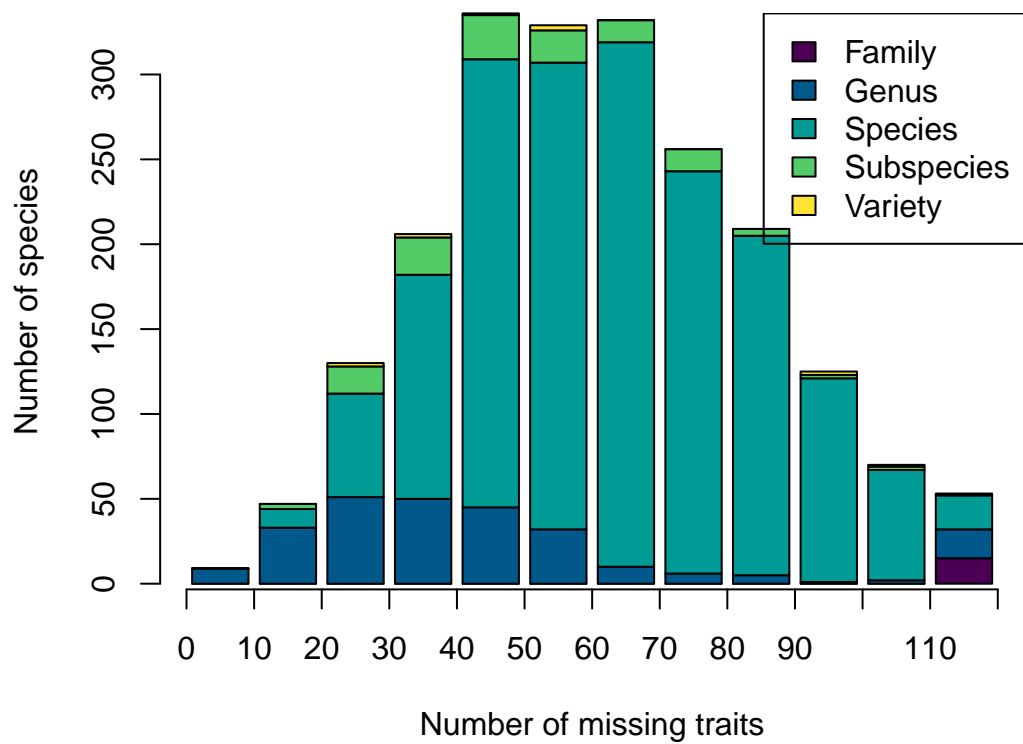
specialist	other	generalist
873	1072	157



## Description of trait databases

So far, we compiled 115 traits for 2102 taxa gathered from 15 trait databases. But there are many missing values.





Taxa with no or limited trait information (N=41).

[1] "Abies"	"Acacia"
[3] "Agrimonia agrimonoides"	"Agropyron"
[5] "Amaranthaceae"	"Apiaceae"
[7] "Asparagaceae"	"Aster"
[9] "Boraginaceae"	"Brassicaceae"
[11] "Bryum dichotomum"	"Caryophyllaceae"
[13] "Chaenomeles x superba"	"Chrysanthemum"
[15] "Circaea"	"Cochlearia"
[17] "Cosmos"	"Crambe abyssinica"
[19] "Cupressus"	"Dysphania aristata"
[21] "Geraniaceae"	"Imbriobryum subapiculatum"
[23] "Lamiaceae"	"Lavandula"
[25] "Leontodon autumnale"	"Liliaceae"
[27] "Matthiola"	"Miliun"
[29] "Moehringia"	"Orchis"
[31] "Paronychia"	"Piptatherum"
[33] "Poaceae"	"Primulaceae"
[35] "Pulmonaria"	"Rhizogemma staphylina"
[37] "Riccia sorocarpa"	"Riccia warnstorffii"
[39] "Roemeria hispida"	"Rosaceae"
[41] "Rubiaceae"	

Open question:

**How to deal with families taxa?**

**How to deal with missing trait values?** Trait imputation, discarding taxa, ...

## Summary of trait completeness

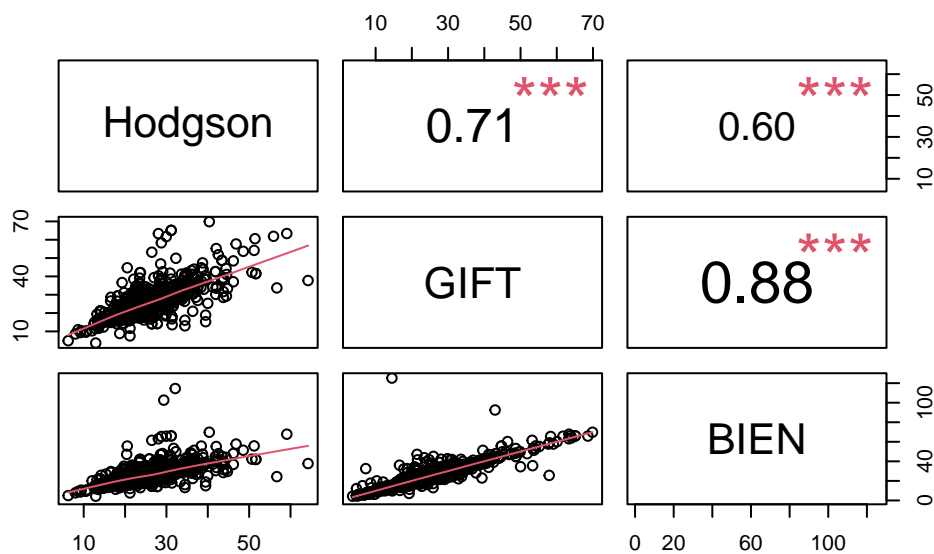
Trait	N database	N taxa	Completeness (%)
Growth.form	5	2047	97
Dispersal.mode	4	2034	97
Plant.height	8	1990	95
Chorology	1	1985	94
Habitat	1	1984	94
Sexuality	1	1975	94
Pollination	1	1974	94
Fruit.type	2	1973	94
Flower.color	5	1961	93
Dispersal.distance	1	1960	93
Inflorescence	2	1954	93
Lifecycle	2	1907	91
Ellenberg.Salinity	1	1872	89
Seed.mass	5	1869	89
Flowering	3	1805	86
Ellenberg.Light	1	1786	85
Photosynthetic.pathway	1	1761	84
Ellenberg.Reaction	1	1756	84
Ellenberg.Moisture	1	1616	77
Ellenberg.Nutrients	1	1610	77
SLA	3	1507	72
Pollination.syndrome	3	1469	70
Flower.UV.reflectance	3	1315	63
Diaspore.exposure	1	1286	61
Diaspore.type	1	1286	61
Leaf.area	2	1257	60
Ellenberg.Temperature	1	1249	59
Floral.symmetry	1	1109	53
Flower.class	1	1109	53
Flower.type	1	1109	53
Nectar.quantity	1	1109	53
Pollen.quantity	1	1109	53
PV.Bees	1	1109	53
PV.Bumblebees	1	1109	53
PV.butterflies	1	1109	53
PV.Hoverflies	1	1109	53
Leaf.dry.mass.content	1	1106	53
Anemochory	1	1099	52
Leaf.dry.mass	1	1055	50

Trait	N database	N taxa	Completeness (%)
Epizoochory	1	972	46
Canopy.diameter	1	953	45
Canopy.height	1	953	45
Diaspore.mass	1	895	43
Leaf.width	1	786	37
Lifeform	2	741	35
Root.mycorrhizal.colonization	1	685	33
Root.mass.fraction	1	632	30
Seed.length	1	619	29
Grassland.specialization	1	587	28
Specific.root.length	1	549	26
Flower.length	2	533	25
Root.lateral.spread	1	500	24
Root.diameter	1	496	24
Diaspore.height	1	484	23
lateral.spread	1	473	23
offspring	1	471	22
offspring.wsmall	1	471	22
clonal.index	1	470	22
Root.tissue.density	1	469	22
Hydrochory	1	454	22
Strategy	2	427	20
Root.N.concentration	1	418	20
Vegetative.propagation	1	397	19
Root.depth	2	384	18
Plant.lifespan	1	379	18
Leaf.nitrogen.content	1	372	18
Root.C.concentration	1	325	15
Leaf.length	1	306	15
Root.dry.matter	1	304	14
Root.C.N.ratio	1	276	13
Leaf.carbon.to.nitrogen.content	1	245	12
Germination	2	172	8
Root.length.density	1	154	7
Flower.width	2	148	7
Fruit.color	1	74	4

## Comparison

### SLA

There are three sources of information for Specific leaf area (SLA) : Hodgson et al. 2023 (in mm<sup>2</sup>/mg), GIFT (in cm<sup>2</sup>/g) and BIEN (in m<sup>2</sup>/kg = mm<sup>2</sup>/mg).



Values are highly correlated, so we could imagine filling the missing values (using preferred data sources or averaging them).

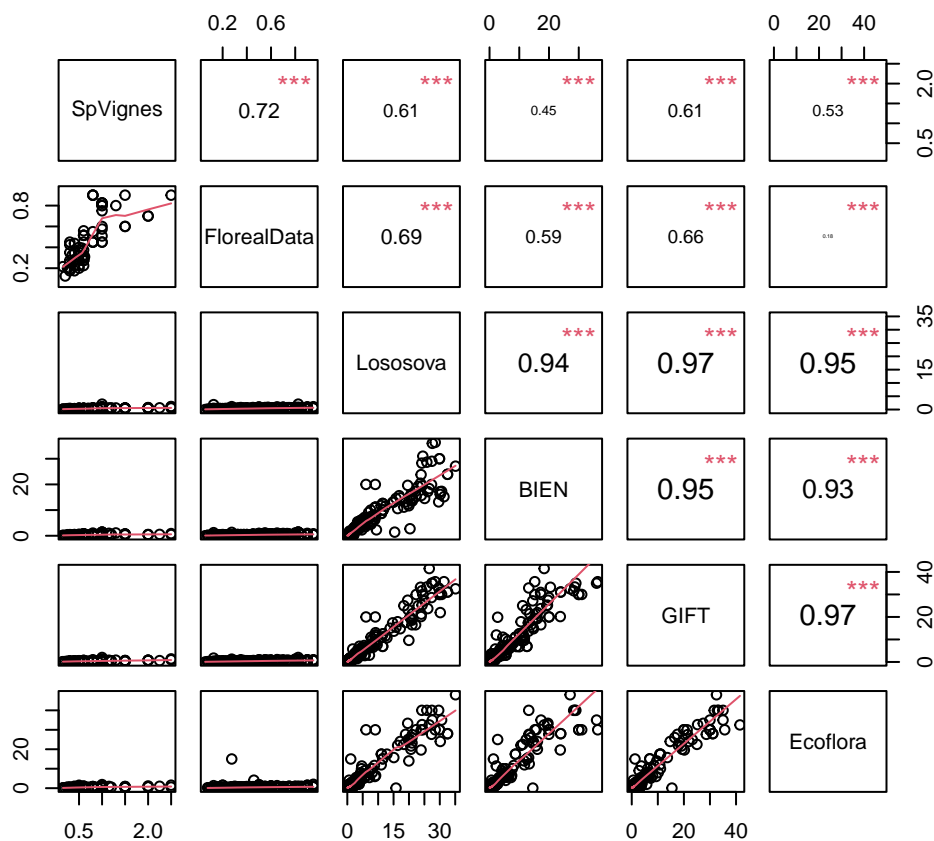
#### Number of NAs:

Hodgson	GIFT	BIEN	filled
1149	883	876	595



## Plant height

There are six sources of information for plant height.



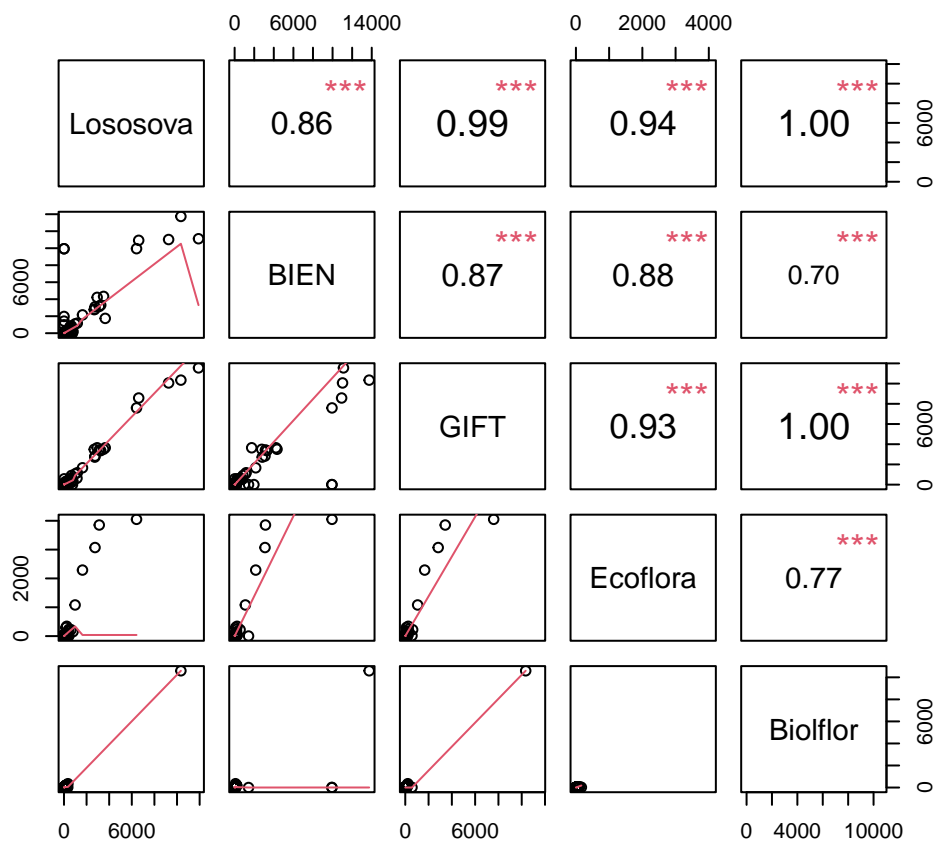
SPVignes and FlorealData are limited to small plants (<1m) (no trees) *but FlorealData>100cm must be clean.*

## Number of NAs:

SpVignes	FlorealData	Lososova	BIEN	GIFT	Ecoflora
1962	1641	172	744	601	1012
filled					
133					

## Seed mass

There are five sources of information for seed mass



## Number of NAs:

Lososova	BIEN	GIFT	Ecoflora	Biolflor	filled
311	741	373	1302	1589	233

## Flower colour

There are five sources of information for flower colour, but it must be cleaned

Baseflor	BIEN	GIFT FlorealData	YvozValPol
205	1825	1802	1518
993			
baseflor			
Blanc	Blanc, jaune	Blanc, jaune, bleu	Blanc, jaune, rose
362	33	3	4
Blanc, rose	Blanc, vert, rose	Bleu	Bleu, blanc
54	1	152	11
Bleu, blanc, rose	Bleu, jaune	Bleu, jaune, rose	Bleu, rose
11	5	2	10
Jaune	Jaune, rose	Jaune, vert	Marron
487	3	1	28
Noir	Rose	Vert	Vert, bleu
2	313	263	14
Vert, jaune, rose	Vert, rose		
1	36		