

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 32 datasets. After cleaning and harmonization, there were 2113 unique taxa.

FAMILY	GENUS	SPECIES	SUBSPECIES	VARIETY
15	261	1705	121	11

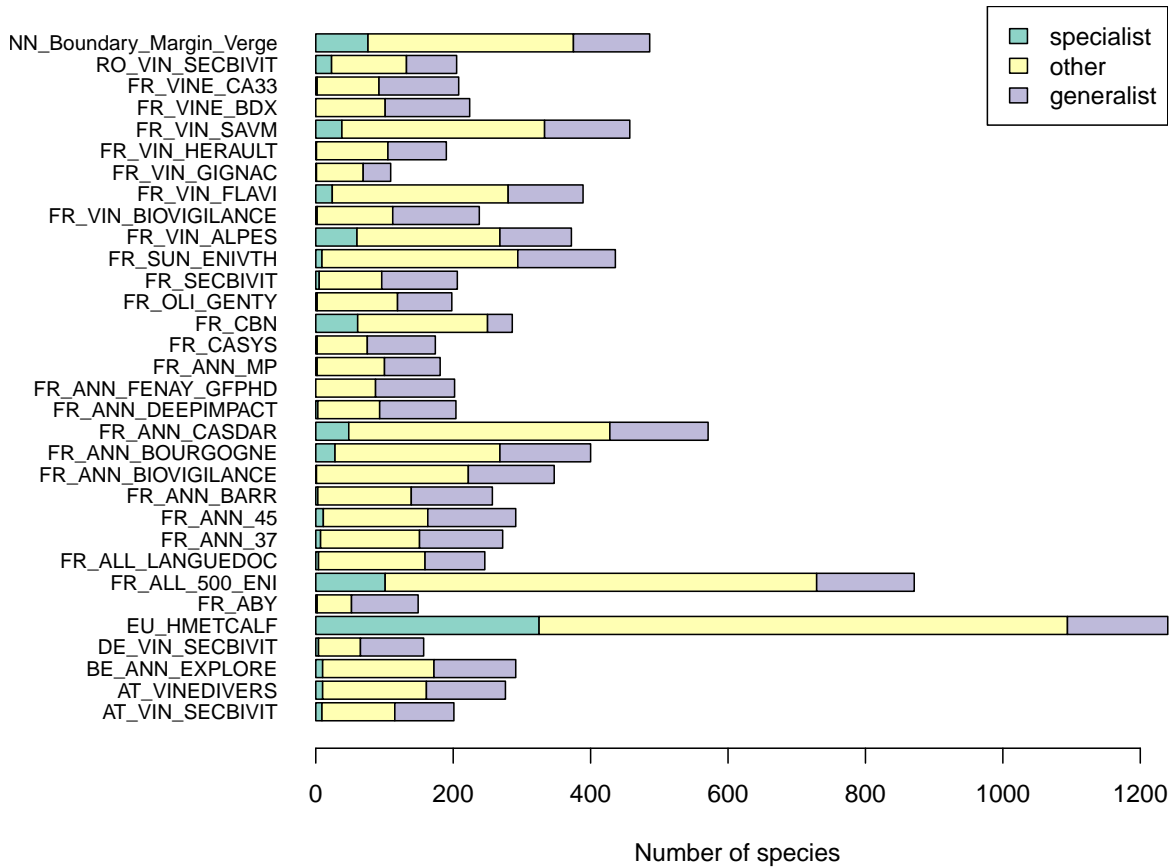
Let's define:

specialist: a taxa that occurred only in a single database

generalist: a taxa that is listed in 50% of the databases (17 out of 32)

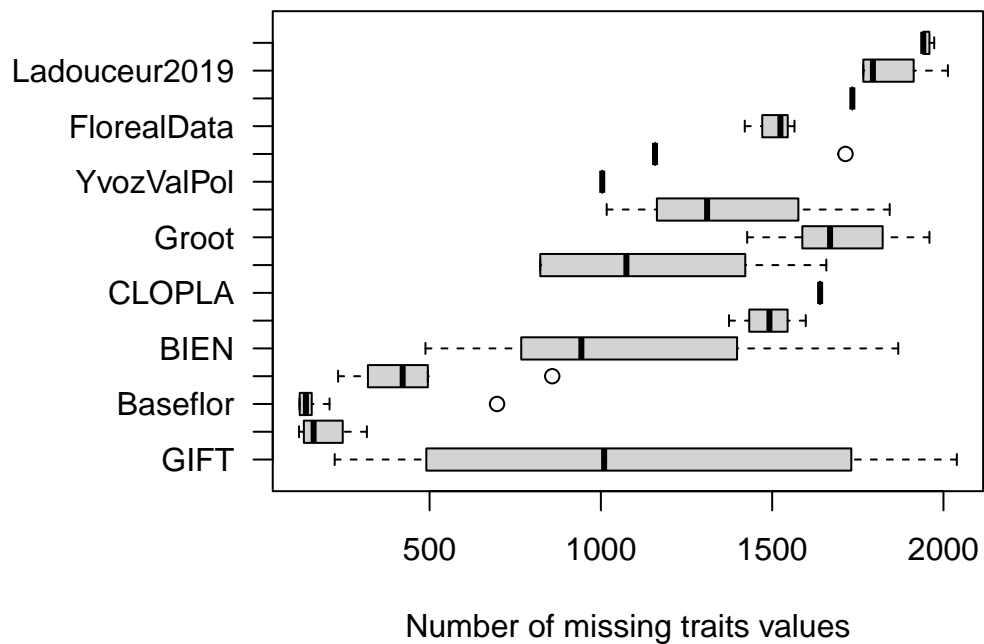
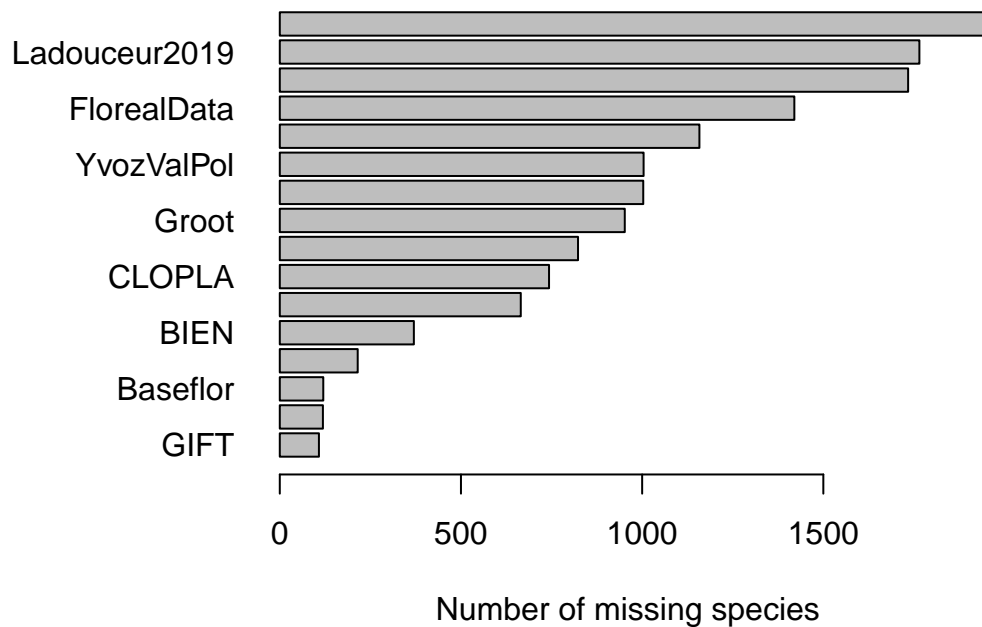
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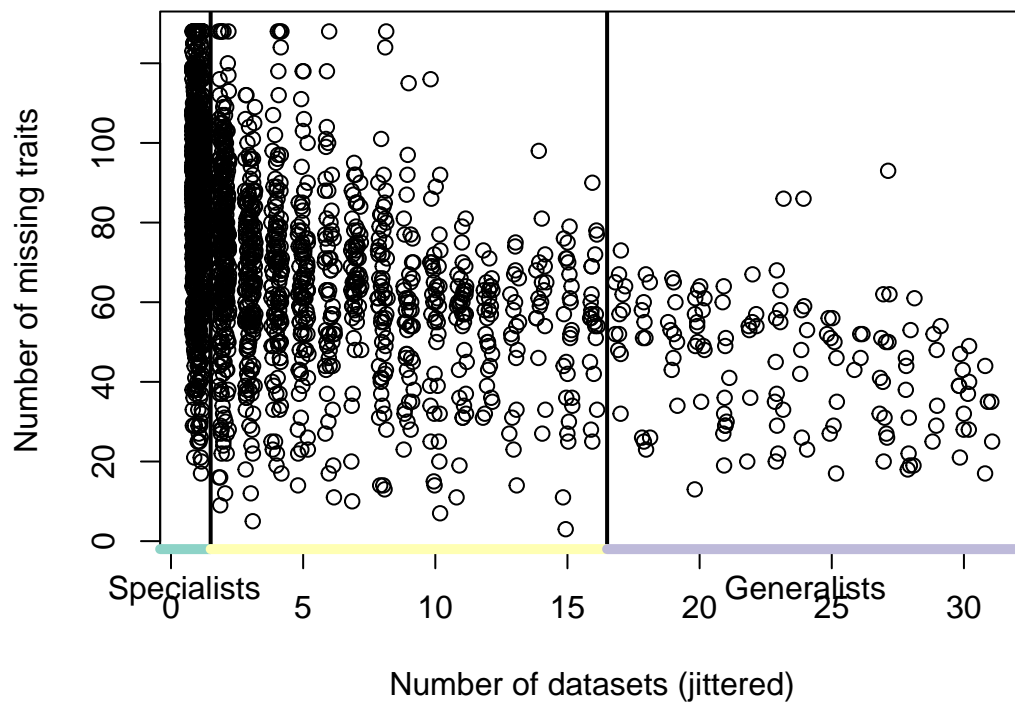
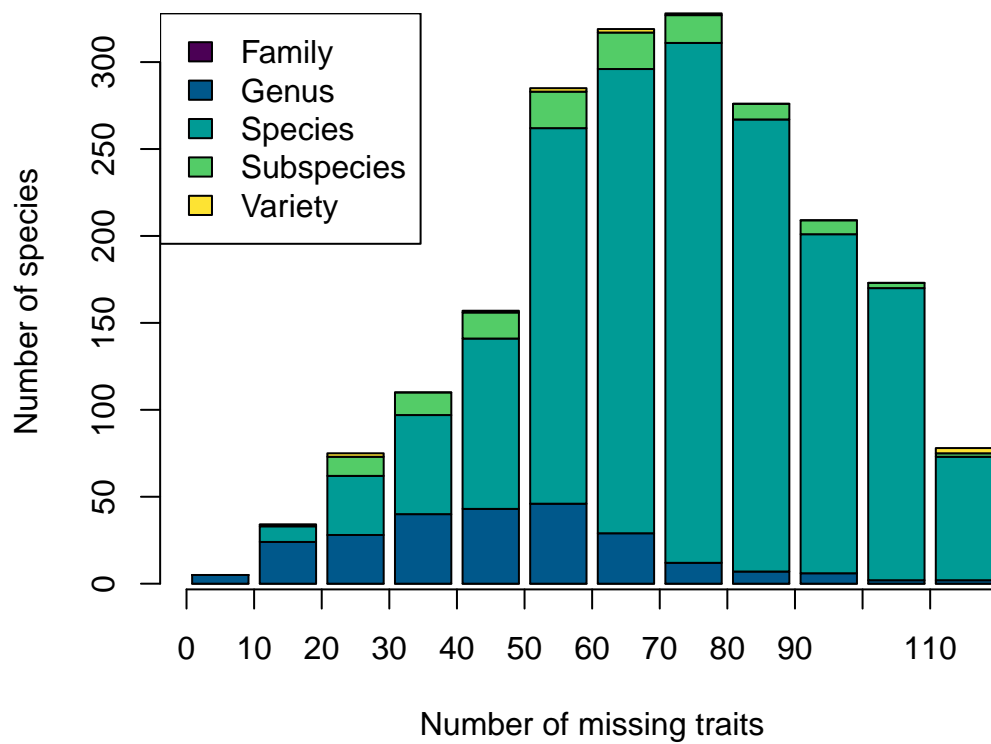
specialist	other	generalist
874	1092	147



Description of trait databases

So far, we compiled 128 traits for 2113 taxa gathered from 16 trait databases. But there are many missing values.





Taxa with no or limited trait information (N=43).

[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] "Ehippiger perforatus"	"Geraniaceae"
[23] "Imbriobryum subapiculatum"	"Lamiaceae"
[25] "Lavandula"	"Leontodon autumnale"
[27] "Liliaceae"	"Matthiola"
[29] "Miliun"	"Moehringia"
[31] "Orchis"	"Paronychia"
[33] "Piptatherum"	"Poaceae"
[35] "Primulaceae"	"Pulmonaria"
[37] "Rhizogemma staphylina"	"Riccia sorocarpa"
[39] "Riccia warnstorffii"	"Roemeria hispida"
[41] "Rosaceae"	"Rubiaceae"
[43] "Vicia nigra"	

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	2056	97
Dispersal.mode	4	2042	97
Plant.height	9	1998	95
Chorology	1	1993	94
Habitat	1	1992	94
Sexuality	1	1983	94
Pollination	1	1982	94
Fruit.type	2	1981	94
Flower.color	5	1969	93
Dispersal.distance	1	1966	93
Inflorescence	2	1961	93
Lifecycle	2	1918	91
Ellenberg.Salinity	1	1880	89
Seed.mass	6	1875	89
Flowering	2	1805	85
Ellenberg.Light	1	1793	85
Photosynthetic.pathway	1	1776	84
Ellenberg.Reaction	1	1762	83
Ellenberg.Moisture	1	1621	77
Ellenberg.Nutrients	1	1618	77
SLA	4	1516	72
Pollination.syndrome	3	1474	70
Flower.UV.reflectance	3	1317	62
Diaspore.exposure	1	1290	61
Diaspore.type	1	1290	61
Leaf.area	2	1261	60
Ellenberg.Temperature	1	1255	59
Leaf.dry.mass.content	1	1110	53
Floral.symmetry	1	1109	52
Flower.class	1	1109	52
Flower.type	1	1109	52
Nectar.quantity	1	1109	52
Pollen.quantity	1	1109	52
PV.Bees	1	1109	52
PV.Bumblebees	1	1109	52
PV.butterflies	1	1109	52
PV.Hoverflies	1	1109	52
Anemochory	1	1101	52
Leaf.dry.mass	1	1059	50

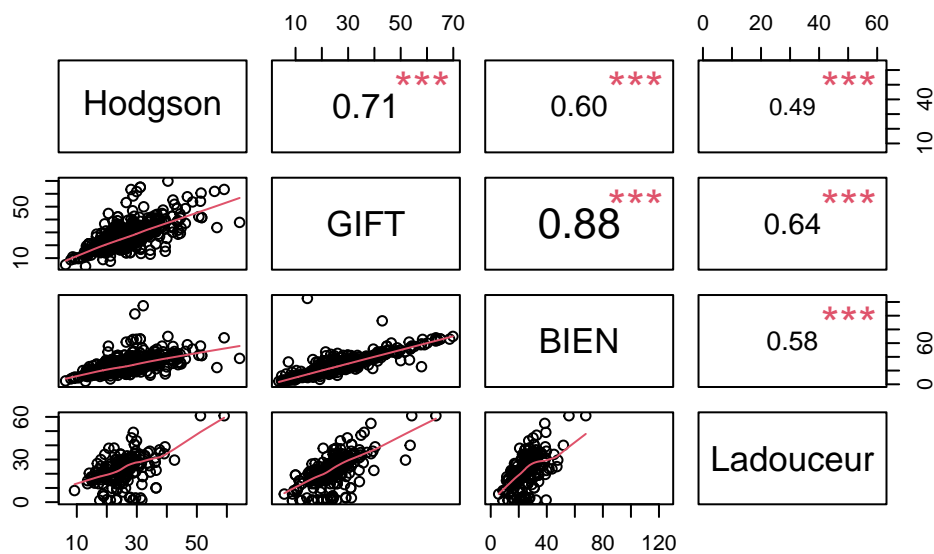
Trait	N database	N taxa	Completeness (%)
Epizoochory	1	976	46
Canopy.diameter	1	955	45
Canopy.height	1	955	45
Diaspore.mass	1	899	43
Leaf.width	1	787	37
Lifeform	2	745	35
Root.mycorrhizal.colonization	1	686	32
Root.mass.fraction	1	634	30
Seed.length	1	621	29
Grassland.specialization	1	591	28
Specific.root.length	1	549	26
Flower.length	2	535	25
Root.lateral.spread	1	501	24
Root.diameter	1	496	23
Diaspore.height	1	484	23
Clonal.index	2	482	23
lateral.spread	1	475	22
offspring	1	473	22
offspring.wsmall	1	473	22
Root.tissue.density	1	470	22
Flowering.onset	2	465	22
Hydrochory	1	455	22
Root.N.concentration	1	418	20
Vegetative.propagation	1	399	19
Root.depth	2	385	18
Plant.lifespan	1	379	18
Strategy.Grime	1	379	18
Leaf.nitrogen.content	1	372	18
Flowering.duration	1	347	16
Seeding.onset	1	347	16
Seeding.season	1	347	16
Root.C.concentration	1	325	15
Clonal.presence	1	319	15
Leaf.length	1	310	15
Root.dry.matter	1	305	14
Clonal.cyclicity	1	303	14
Root.C.N.ratio	1	276	13
Leaf.carbon.to.nitrogen.content	1	245	12
Seed.dormancy	1	200	9
Clonal.spread	1	192	9
Germination.range	1	172	8

Trait	N database	N taxa	Completeness (%)
Germination.start	1	172	8
Root.length.density	1	154	7
Strategy	1	154	7
Flower.width	2	148	7
Radial.growth	1	100	5
Fruit.color	1	74	4

Comparison

SLA

There are three sources of information for Specific leaf area (SLA) : Hodgson et al. 2023 (in mm²/mg), GIFT (in cm²/g) and BIEN (in m²/kg = mm²/mg).



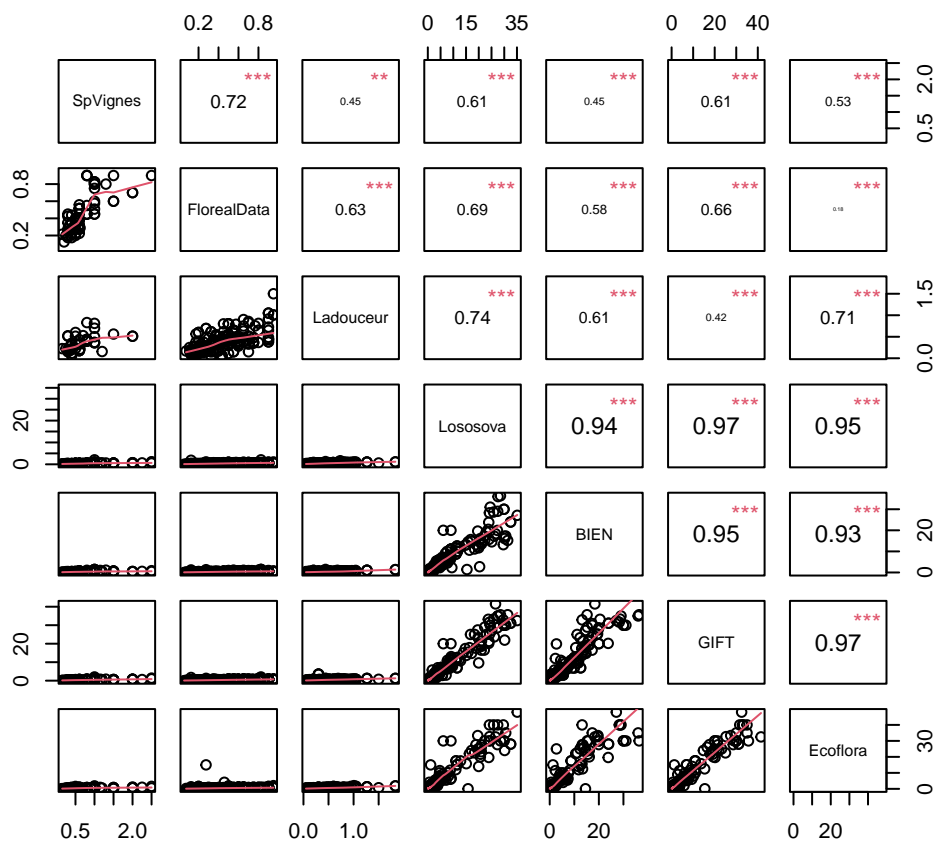
Values are highly correlated, so we could imagine filling the missing values (using preferred data sources or averaging them).

Number of NAs:

Hodgson	GIFT	BIEN	Ladouceur	filled
1158	890	882	1819	597

Plant height

There are six sources of information for plant height.



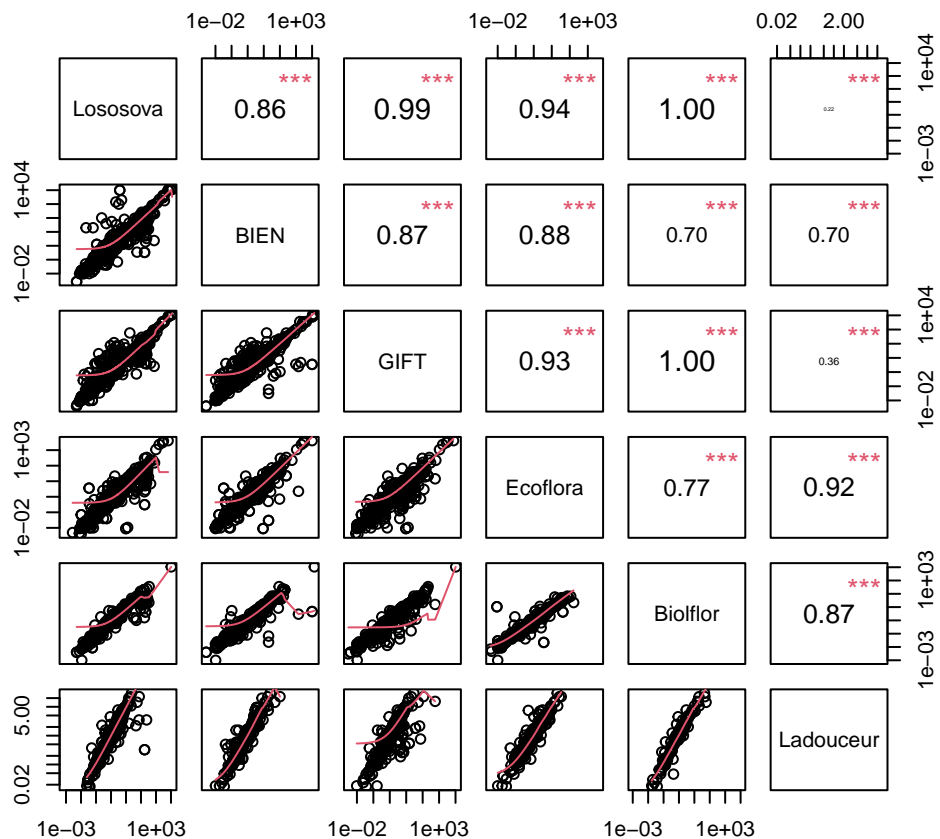
SPVignes, FlorealData, and Ladouceur_2019 are limited to small plants (<1m) (no trees) *but FlorealData>100cm must be clean.*

Number of NAs:

SpVignes	FlorealData	Ladouceur	Lososova	BIEN	GIFT
1973	1648	1782	176	749	604
Ecoflora	filled				
1017	136				

Seed mass

There are five sources of information for seed mass



Number of NAs:

Lososova	BIEN	GIFT	Ecoflora	Biolflor	Ladouceur	filled
317	745	377	1310	1598	1782	238

Flower colour

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

Baseflor	BIEN	GIFT FlorealData	YvozValPol
208	1836	1814	1526
			1004
baseflor			
Blanc	Blanc, jaune	Blanc, jaune, bleu	Blanc, jaune, rose
364	33	3	4
Blanc, rose	Blanc, vert, rose	Bleu	Bleu, blanc
54	1	154	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	315	265	14
Vert, jaune, rose	Vert, rose		
1	36		