

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 2109 unique taxa.

FAMILY	GENUS	SPECIES	SUBSPECIES	VARIETY
15	261	1701	122	10

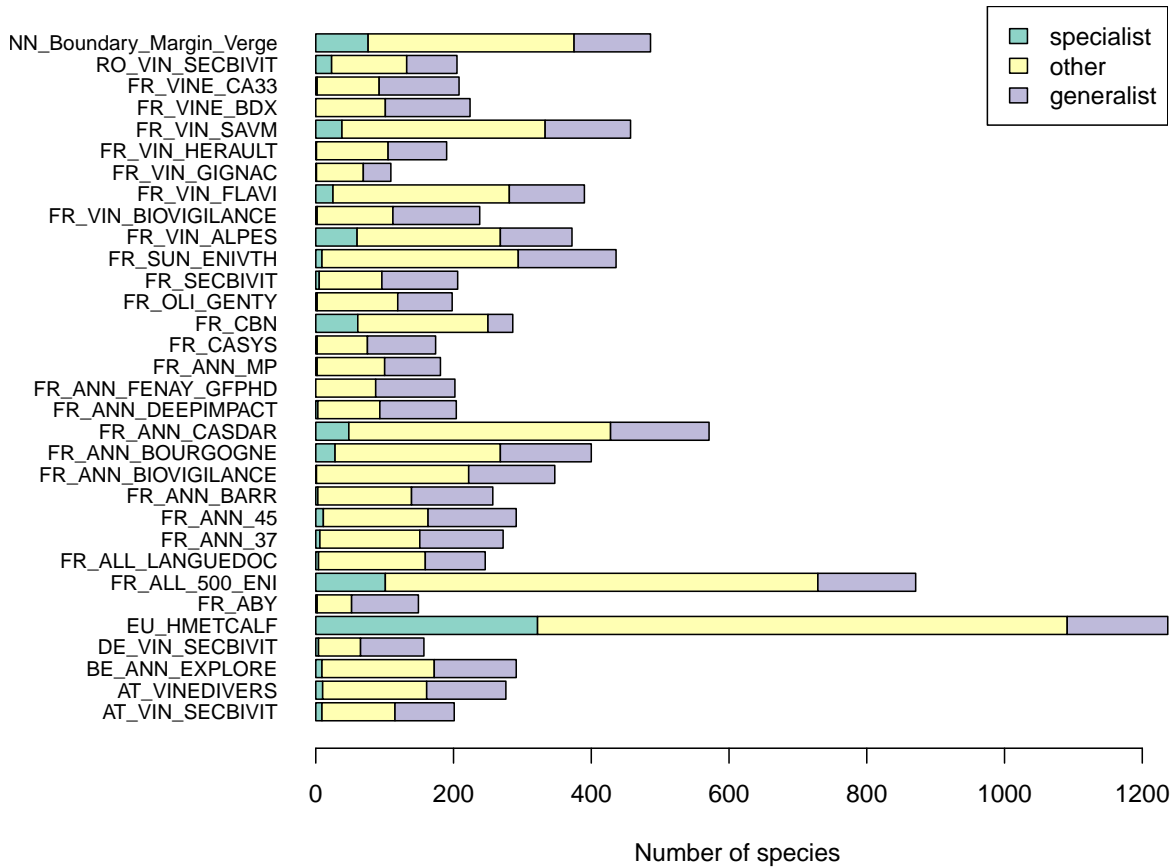
Let's define:

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)

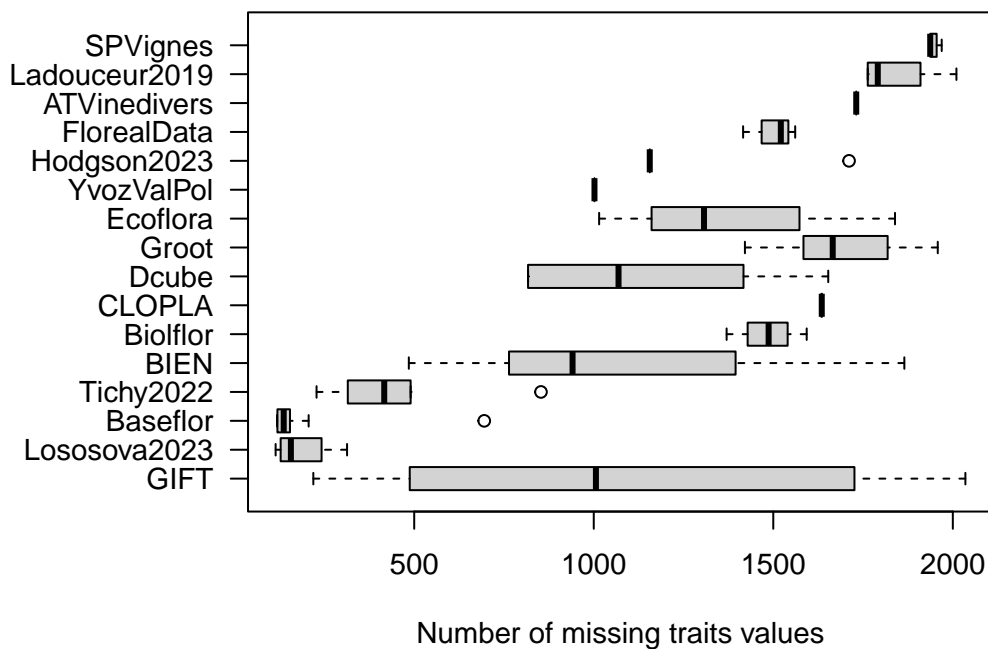
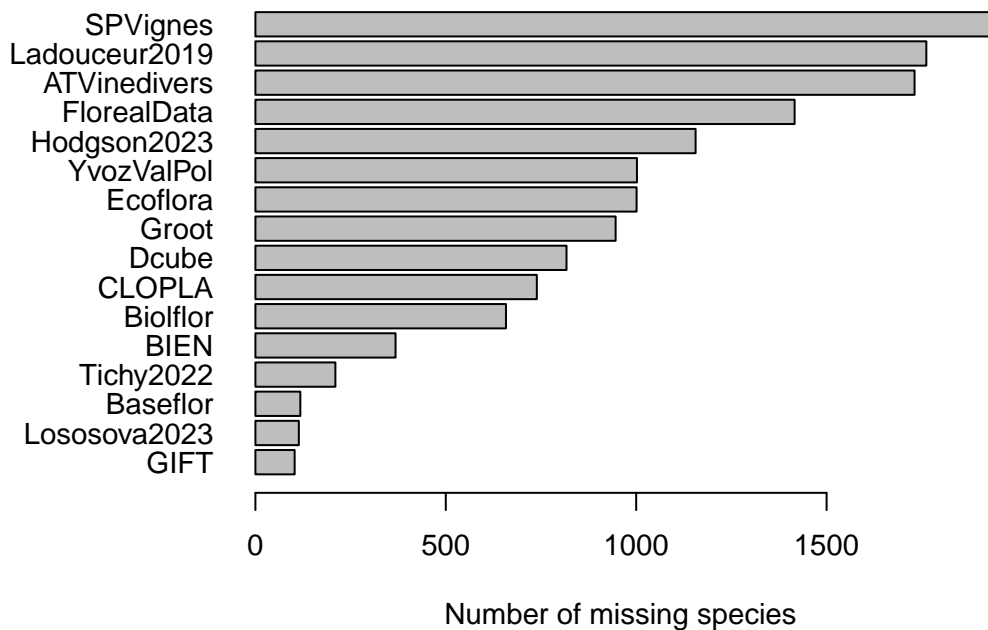
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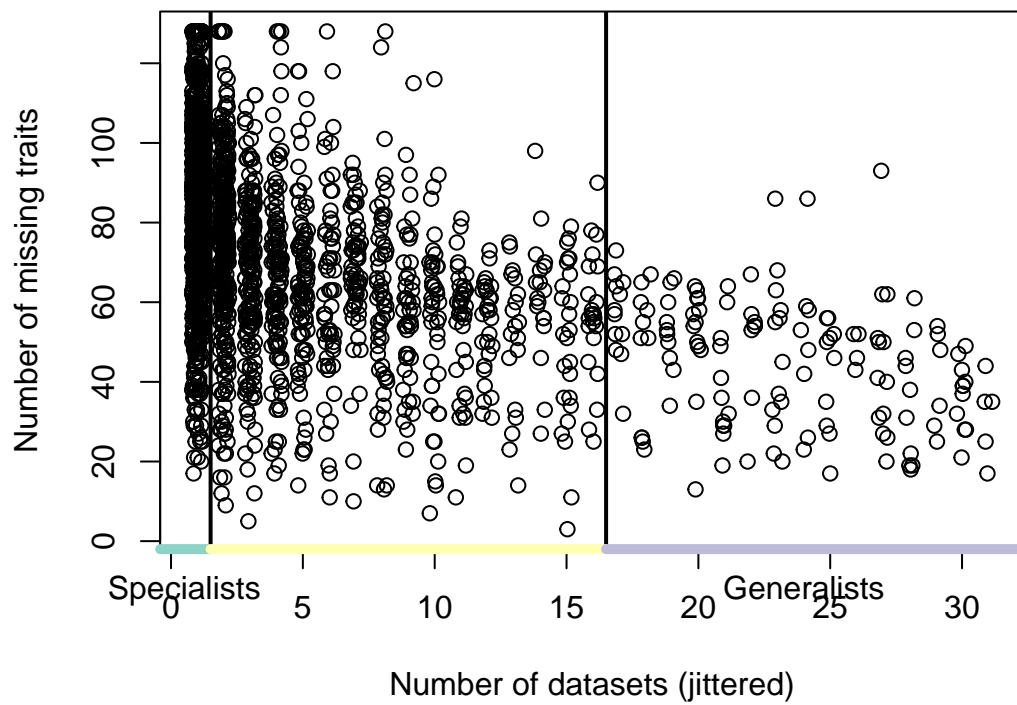
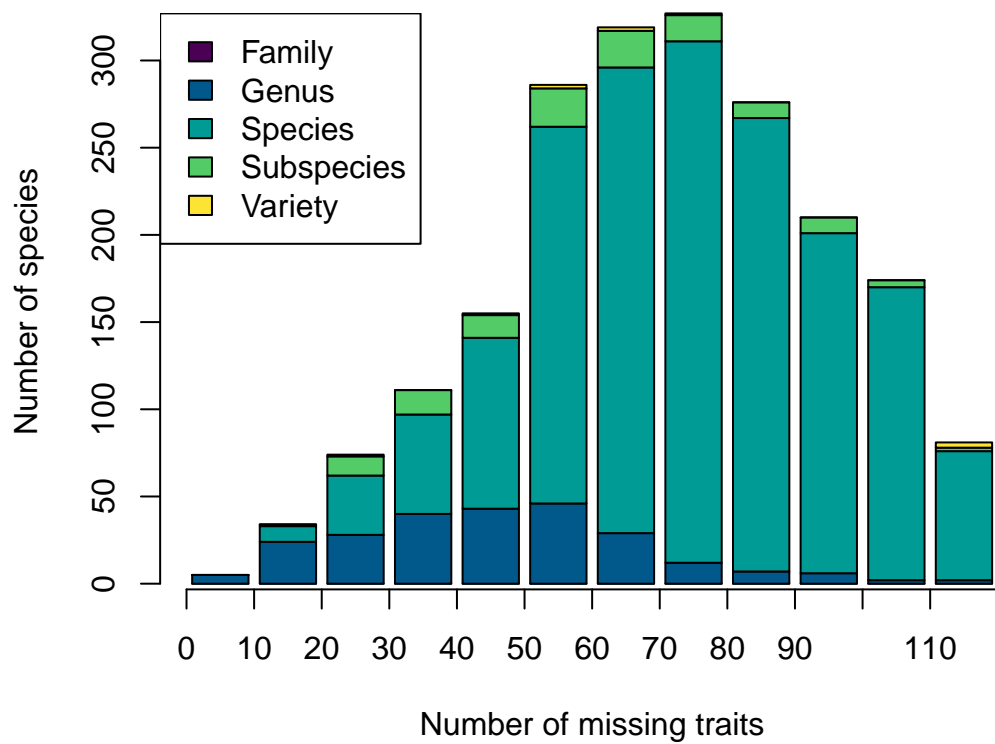
specialist	other	generalist
870	1092	147



Description of trait databases

So far, we compiled 128 traits for 2109 taxa gathered from 16 trait databases. But there are many missing values.





Taxa with no or limited trait information (N=38).

[1] "Abies"	"Acacia"
[3] "Agropyron"	"Amaranthaceae"
[5] "Apiaceae"	"Asparagaceae"
[7] "Aster"	"Boraginaceae"
[9] "Brassicaceae"	"Bryum dichotomum"
[11] "Caryophyllaceae"	"Chaenomeles x superba"
[13] "Chrysanthemum"	"Circaea"
[15] "Cochlearia"	"Cosmos"
[17] "Cupressus"	"Ephippiger perforatus"
[19] "Geraniaceae"	"Imbriobryum subapiculatum"
[21] "Lamiaceae"	"Lavandula"
[23] "Liliaceae"	"Matthiola"
[25] "Miliun"	"Moehringia"
[27] "Orchis"	"Paronychia"
[29] "Piptatherum"	"Poaceae"
[31] "Primulaceae"	"Pulmonaria"
[33] "Rhizogemma staphylina"	"Riccia sorocarpa"
[35] "Riccia warnstorffii"	"Roemeria hispida"
[37] "Rosaceae"	"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	2058	98
Dispersal.mode	4	2043	97
Plant.height	9	2000	95
Chorology	1	1991	94
Habitat	1	1990	94
Fruit.type	2	1982	94
Sexuality	1	1982	94
Pollination	1	1981	94
Flower.color	5	1968	93
Dispersal.distance	1	1967	93
Inflorescence	2	1959	93
Lifecycle	2	1918	91
Ellenberg.Salinity	1	1881	89
Seed.mass	6	1875	89
Flowering	2	1804	86
Ellenberg.Light	1	1794	85
Photosynthetic.pathway	1	1776	84
Ellenberg.Reaction	1	1763	84
Ellenberg.Moisture	1	1622	77
Ellenberg.Nutrients	1	1619	77
SLA	4	1514	72
Pollination.syndrome	3	1473	70
Flower.UV.reflectance	3	1316	62
Diaspore.exposure	1	1292	61
Diaspore.type	1	1292	61
Leaf.area	2	1259	60
Ellenberg.Temperature	1	1256	60
Leaf.dry.mass.content	1	1109	53
Floral.symmetry	1	1107	52
Flower.class	1	1107	52
Flower.type	1	1107	52
Nectar.quantity	1	1107	52
Pollen.quantity	1	1107	52
PV.Bees	1	1107	52
PV.Bumblebees	1	1107	52
PV.butterflies	1	1107	52
PV.Hoverflies	1	1107	52
Anemochory	1	1103	52
Leaf.dry.mass	1	1057	50

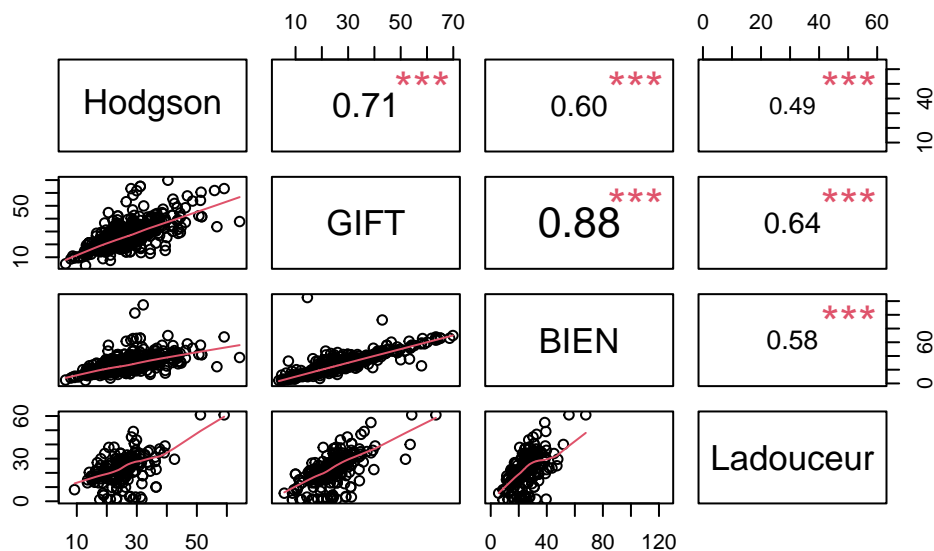
Trait	N database	N taxa	Completeness (%)
Epizoochory	1	977	46
Canopy.diameter	1	953	45
Canopy.height	1	953	45
Diaspore.mass	1	898	43
Leaf.width	1	785	37
Lifeform	2	745	35
Root.mycorrhizal.colonization	1	688	33
Root.mass.fraction	1	633	30
Seed.length	1	622	29
Grassland.specialization	1	590	28
Specific.root.length	1	548	26
Flower.length	2	534	25
Root.lateral.spread	1	502	24
Root.diameter	1	495	23
Diaspore.height	1	486	23
Clonal.index	2	483	23
lateral.spread	1	476	23
offspring	1	474	22
offspring.wsmall	1	474	22
Root.tissue.density	1	469	22
Flowering.onset	2	464	22
Hydrochory	1	456	22
Root.N.concentration	1	418	20
Vegetative.propagation	1	398	19
Root.depth	2	385	18
Plant.lifespan	1	378	18
Strategy.Grime	1	378	18
Leaf.nitrogen.content	1	371	18
Flowering.duration	1	346	16
Seeding.onset	1	346	16
Seeding.season	1	346	16
Root.C.concentration	1	325	15
Clonal.presence	1	318	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	244	12
Seed.dormancy	1	199	9
Clonal.spread	1	192	9
Germination.range	1	172	8

Trait	N database	N taxa	Completeness (%)
Germination.start	1	172	8
Strategy	1	154	7
Root.length.density	1	151	7
Flower.width	2	148	7
Radial.growth	1	99	5
Fruit.color	1	74	4

Comparison

SLA

There are four sources of information for Specific leaf area (SLA) : Hodgson et al. 2023 (in mm^2/mg), GIFT (in cm^2/g), BIEN (in $\text{m}^2/\text{kg} = \text{mm}^2/\text{mg}$), and Ladouceur et al. 2023 (in cm^2/g).



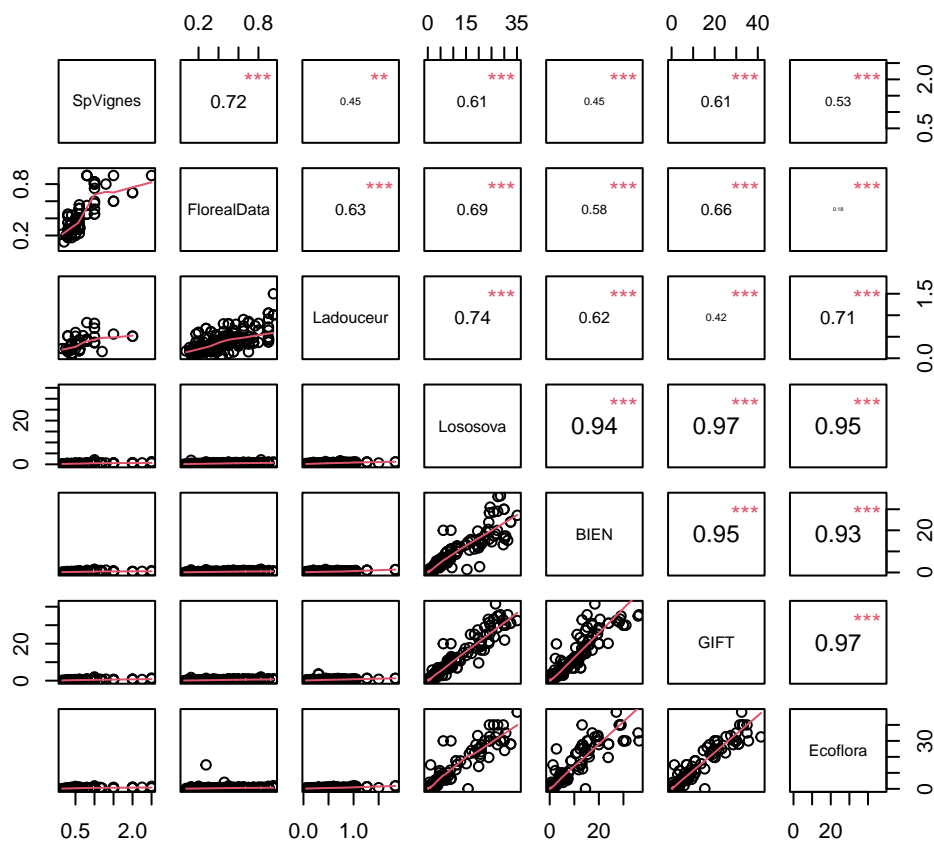
Values are highly correlated, so we could imagine filling the missing values (using preferred data sources or averaging them).

Number of NAs:

Hodgson	GIFT	BIEN	Ladouceur	filled
1156	888	880	1816	595

Plant height

There are seven sources of information for plant height.



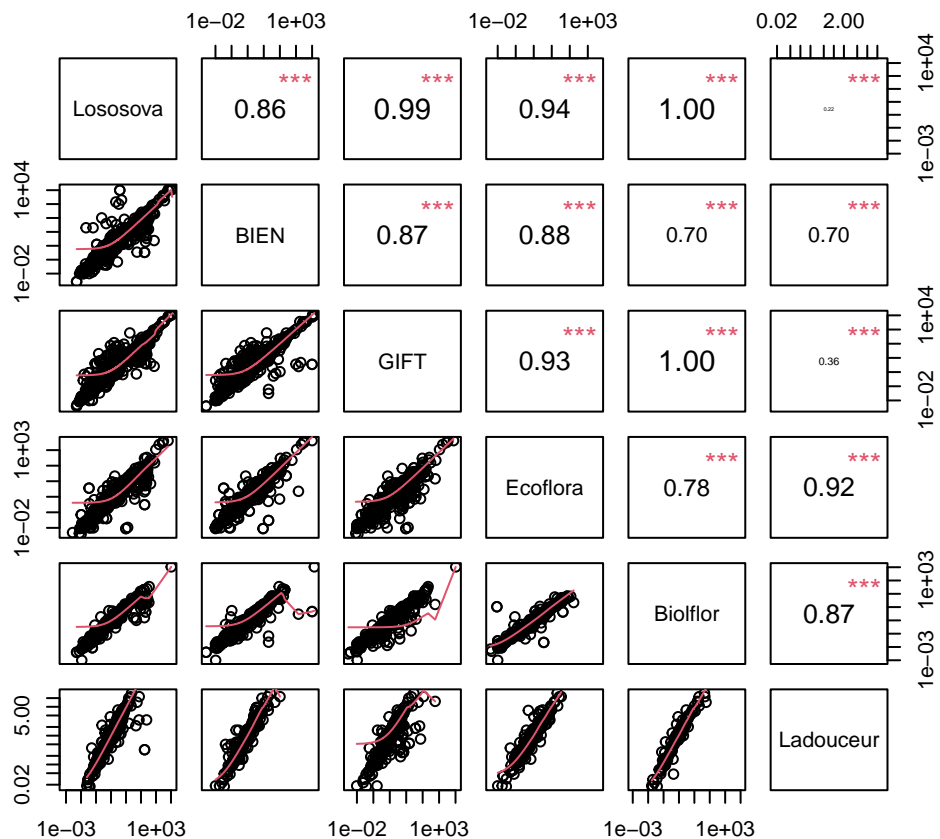
SPVignes, FlorealData, and Ladouceur_2019 are limited to small plants (<1m) (no trees) *but FlorealData>100cm must be clean.*

Number of NAs:

SpVignes	FlorealData	Ladouceur	Lososova	BIEN	GIFT
1969	1644	1779	171	745	600
Ecoflora	filled				
1015	131				

Seed mass

There are six sources of information for seed mass



Number of NAs:

Lososova	BIEN	GIFT	Ecoflora	Biolflor	Ladouceur	filled
313	744	375	1307	1593	1779	234

Flower colour

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

Baseflor	BIEN	GIFT FlorealData	YvozValPol
206	1832	1810	1523
1002			
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
488	3	1	28
Noir	Rose	Vert	Vert, bleu
2	314	264	14
Vert, jaune, rose	Vert, rose		
1	36		