

CODA19 data - First 24h
Imputed dataset

2021-01-23 21:19:56

Data report overview

The dataset examined has the following dimensions:

Feature	Result
Number of observations	358
Number of variables	135

Checks performed

The following variable checks were performed, depending on the data type of each variable:

	character	factor	labelled	haven labelled	numeric	integer	logical	Date
Identify miscoded missing values	×	×	×	×	×	×		×
Identify prefixed and suffixed whitespace	×	×	×	×				
Identify levels with < 6 obs.	×	×	×	×				
Identify case issues	×	×	×	×				
Identify misclassified numeric or integer variables	×	×	×	×				
Identify outliers					×	×		×

Please note that all numerical values in the following have been rounded to 2 decimals.

Summary table

	Variable class	# unique values	Missing observations	Any problems?
patient_site_uid	numeric	351	0.00 %	
female	integer	2	0.00 %	
male	integer	2	0.00 %	
patient_age	numeric	75	0.00 %	
death	numeric	2	0.00 %	
x5_alpha_reductase_inhibitors	numeric	2	0.00 %	
acetaminophene	numeric	2	0.00 %	
adjuvants_anesthesia	numeric	2	0.00 %	×
adrenergic_alpha_1_receptor_antagonists	numeric	2	0.00 %	
adrenergic_beta_3_receptor_agonists	numeric	2	0.00 %	×
adrenergic_beta_antagonists	numeric	2	0.00 %	
adrenergic_uptake_inhibitors	numeric	2	0.00 %	×
analgesics	numeric	2	0.00 %	
analgesics_opioid	numeric	2	0.00 %	
androgens	numeric	2	0.00 %	×
anesthetics_local	numeric	2	0.00 %	×
anti_anxiety_agents	numeric	2	0.00 %	×
anti_arrhythmia_agents	numeric	2	0.00 %	
anti_asthmatic_agents	numeric	2	0.00 %	×
anti_bacterial_agents	numeric	2	0.00 %	
anti_infective_agents_local	numeric	2	0.00 %	
anti_inflammatory_agents	numeric	2	0.00 %	
anti_inflammatory_agents_non_steroidal	numeric	2	0.00 %	×
anti_ulcer_agents	numeric	2	0.00 %	
anticholesteremic_agents	numeric	2	0.00 %	
anticoagulants	numeric	2	0.00 %	
anticonvulsants	numeric	2	0.00 %	
antidepressive_agents	numeric	2	0.00 %	
antidepressive_agents_tricyclic	numeric	2	0.00 %	×
antidiarrheals	numeric	2	0.00 %	
antiemetics	numeric	2	0.00 %	
antifibrinolytic_agents	numeric	2	0.00 %	×
antifungal_agents	numeric	2	0.00 %	×
antihypertensive_agents	numeric	2	0.00 %	
antimalarials	numeric	2	0.00 %	×
antimetabolites	numeric	2	0.00 %	
antineoplastic_agents_hormonal	numeric	2	0.00 %	×
antiparkinson_agents	numeric	2	0.00 %	
antipruritics	numeric	2	0.00 %	×
antipsychotic_agents	numeric	2	0.00 %	
antithyroid_agents	numeric	2	0.00 %	×
antitubercular_agents	numeric	2	0.00 %	×

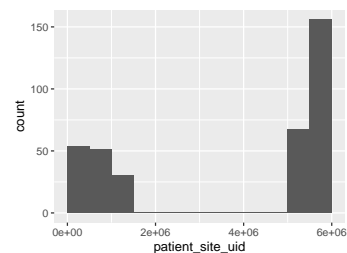
	Variable class	# unique values	Missing observations	Any problems?
antitussive_agents	numeric	2	0.00 %	×
antiviral_agents	numeric	2	0.00 %	×
benzodiazepines	numeric	2	0.00 %	
bicarbonate	numeric	2	0.00 %	
bone_density_conservation_agents	numeric	2	0.00 %	
bronchodilator_agents	numeric	2	0.00 %	
calcium_regulating_hormones_and_agents	numeric	2	0.00 %	
carbonic_anhydrase_inhibitors	numeric	2	0.00 %	
chelating_agents	numeric	2	0.00 %	
cholagogues_and_choleretics	numeric	2	0.00 %	×
cholinesterase_inhibitors	numeric	2	0.00 %	
contraceptive_agents_hormonal	numeric	2	0.00 %	×
diuretics	numeric	2	0.00 %	
factor_xa_inhibitors	numeric	2	0.00 %	
gastrointestinal_agents	numeric	2	0.00 %	×
glucocorticoids	numeric	2	0.00 %	
gout_suppressants	numeric	2	0.00 %	×
hematologic_agents	numeric	2	0.00 %	×
hemostatics	numeric	2	0.00 %	×
hiv_medication	numeric	2	0.00 %	×
hypoglycemic_agents	numeric	2	0.00 %	
immunosuppressive_agents	numeric	2	0.00 %	
laxatives	numeric	2	0.00 %	
levothyroxine	numeric	2	0.00 %	
miotics	numeric	1	0.00 %	×
muscarinic_antagonists	numeric	2	0.00 %	×
muscle_relaxants_central	numeric	2	0.00 %	×
narcotic_antagonists	numeric	2	0.00 %	×
neuromuscular_blocking_agents	numeric	2	0.00 %	
ophthalmic_solutions	numeric	2	0.00 %	
parasympatholytics	numeric	2	0.00 %	×
platelet_aggregation_inhibitors	numeric	2	0.00 %	
progestins	numeric	2	0.00 %	×
reverse_transcriptase_inhibitors	numeric	2	0.00 %	×
sedation	numeric	2	0.00 %	
serotonin_5_ht1_receptor_agonists	numeric	2	0.00 %	×
serotonin_uptake_inhibitors	numeric	2	0.00 %	
sleep_aids_pharmaceutical	numeric	2	0.00 %	×
smoking_cessation_agents	numeric	2	0.00 %	
vasodilator_agents	numeric	2	0.00 %	
vasopressors	numeric	2	0.00 %	
vitamin_b_complex	numeric	2	0.00 %	
vitamins	numeric	2	0.00 %	
hemoglobin_min	numeric	88	0.00 %	×
hemoglobin_max	numeric	86	0.00 %	×
hemoglobin_mean	numeric	142	0.00 %	×
plt_min	numeric	207	0.00 %	×
plt_max	numeric	207	0.00 %	×
plt_mean	numeric	242	0.00 %	×
wbc_min	numeric	115	0.00 %	×
wbc_max	numeric	124	0.00 %	×
wbc_mean	numeric	169	0.00 %	×
sodium_min	numeric	37	0.00 %	×
sodium_max	numeric	30	0.00 %	×

	Variable class	# unique values	Missing observations	Any problems?
sodium_mean	numeric	78	0.00 %	×
potassium_min	numeric	27	0.00 %	×
potassium_max	numeric	45	0.00 %	×
potassium_mean	numeric	79	0.00 %	×
creatinine_min	numeric	152	0.00 %	×
creatinine_max	numeric	144	0.00 %	×
creatinine_mean	numeric	187	0.00 %	×
eos_min	numeric	31	0.00 %	×
eos_max	numeric	35	0.00 %	×
eos_mean	numeric	33	0.00 %	×
lymph_min	numeric	153	0.00 %	×
lymph_max	numeric	165	0.00 %	×
lymph_mean	numeric	164	0.00 %	×
neutrophil_min	numeric	257	0.00 %	×
neutrophil_max	numeric	265	0.00 %	×
neutrophil_mean	numeric	258	0.00 %	×
mono_min	numeric	112	0.00 %	×
mono_max	numeric	117	0.00 %	×
mono_mean	numeric	114	0.00 %	×
baso_min	numeric	10	0.00 %	×
baso_max	numeric	14	0.00 %	×
baso_mean	numeric	12	0.00 %	×
sbp_min	numeric	79	0.00 %	×
sbp_max	numeric	85	0.00 %	×
sbp_mean	numeric	227	0.00 %	×
dbp_min	numeric	53	0.00 %	×
dbp_max	numeric	61	0.00 %	×
dbp_mean	numeric	203	0.00 %	×
temp_min	numeric	35	0.00 %	×
temp_max	numeric	46	0.00 %	×
temp_mean	numeric	139	0.00 %	×
so2_min	numeric	32	0.00 %	×
so2_max	numeric	11	0.00 %	×
so2_mean	numeric	157	0.00 %	×
rr_min	numeric	15	0.00 %	×
rr_max	numeric	25	0.00 %	×
rr_mean	numeric	104	0.00 %	×
mv	numeric	2	0.00 %	
icu	numeric	2	0.00 %	

Variable list

patient_site_uid

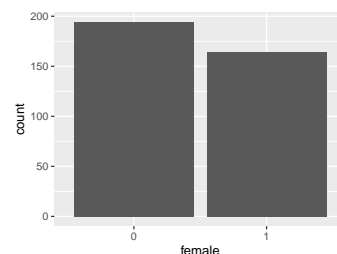
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	351
Median	5318271.5
1st and 3rd quartiles	847135.5; 5635039.75
Min. and max.	720; 5655546



female

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

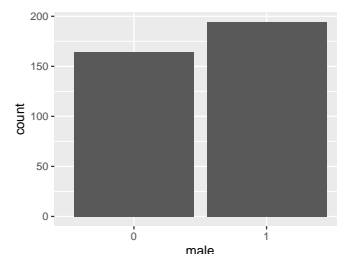
Feature	Result
Variable type	integer
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“0”
Reference category	0



male

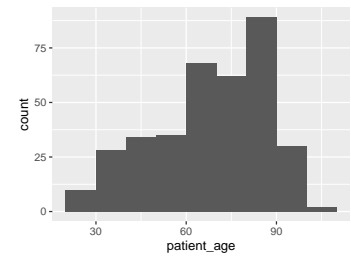
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	integer
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“1”
Reference category	0



patient_age

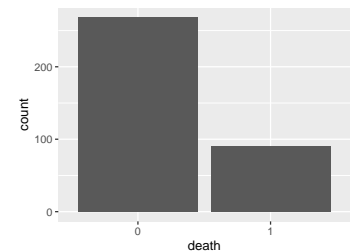
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	75
Median	71
1st and 3rd quartiles	55; 85
Min. and max.	24; 103



death

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

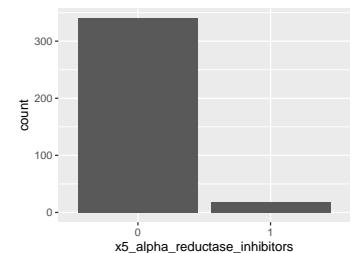
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“0”
Reference category	0



x5_alpha_reductase_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

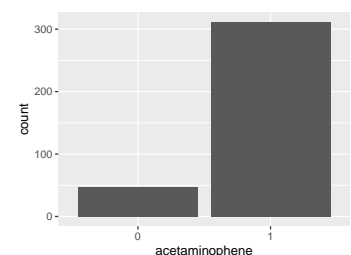
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“0”
Reference category	0



acetaminophene

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

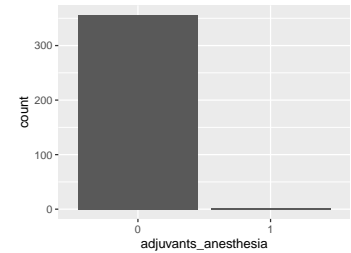
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“1”
Reference category	0



adjuvants_anesthesia

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

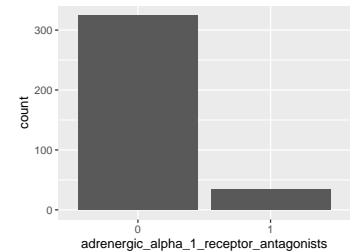


- Note that the following levels have at most five observations: "1".

adrenergic_alpha_1_receptor_antagonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

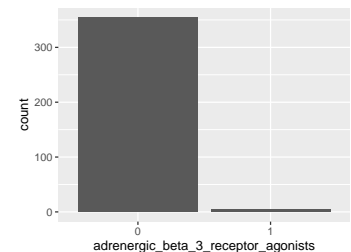
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic_beta_3_receptor_agonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

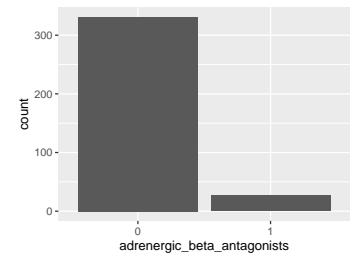


- Note that the following levels have at most five observations: "1".

adrenergic__beta__antagonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

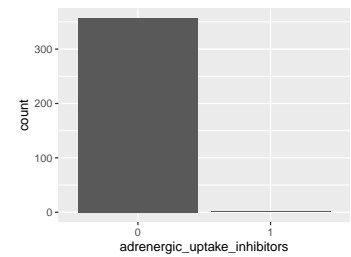
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic__uptake__inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

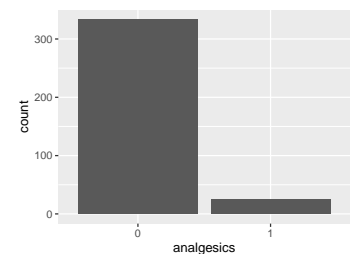


- Note that the following levels have at most five observations: "1".

analgesics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

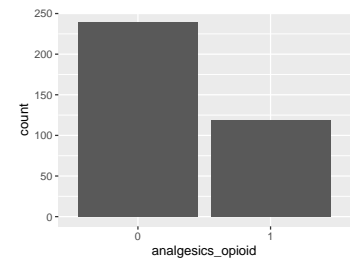
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



analgesics__opioid

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

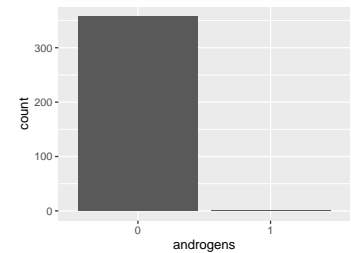
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



androgens

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

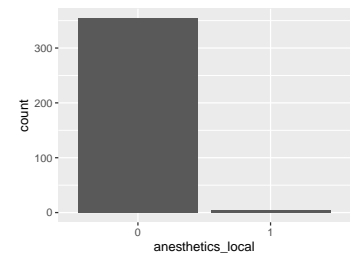


- Note that the following levels have at most five observations: "1".

anesthetics_local

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

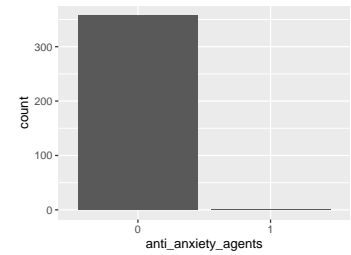


- Note that the following levels have at most five observations: "1".

anti_anxiety_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

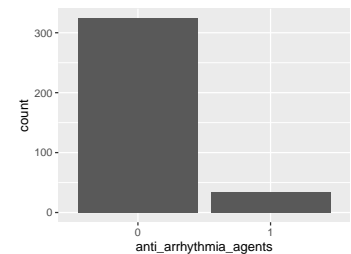


- Note that the following levels have at most five observations: "1".

anti_arrhythmia_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

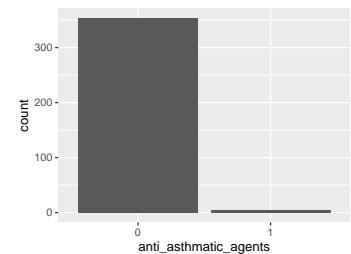
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_asthmatic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

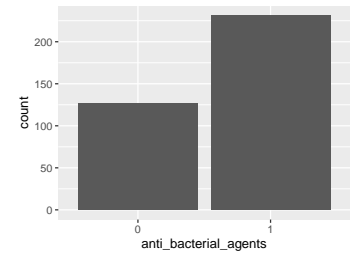


- Note that the following levels have at most five observations: "1".

anti_bacterial_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

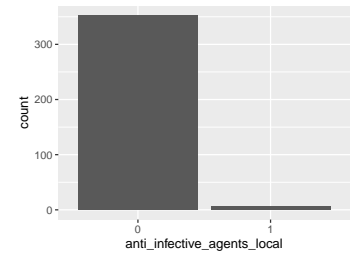
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"1"
Reference category	0



anti_infective_agents_local

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

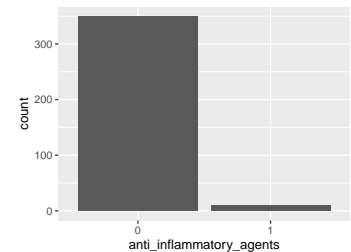
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

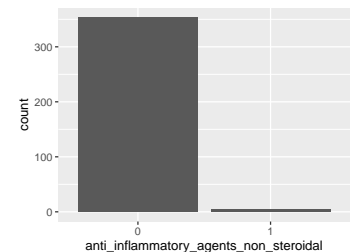
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents_non_steroidal

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

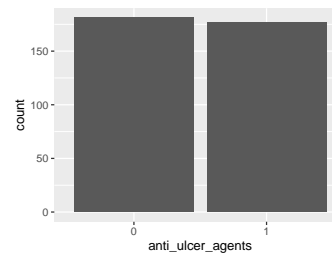


- Note that the following levels have at most five observations: "1".

anti_ulcer_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

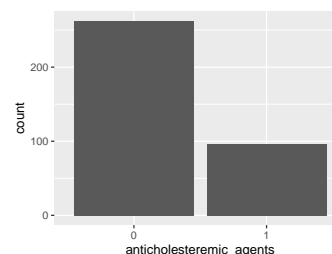
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



anticholesteremic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

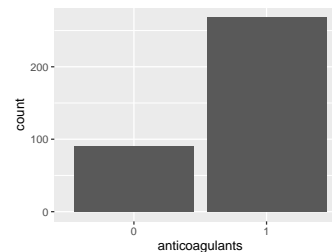
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



anticoagulants

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

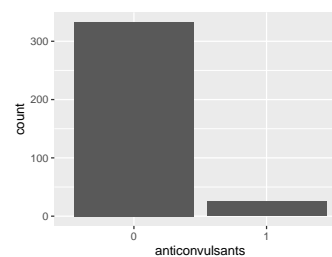
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"1"
Reference category	0



anticonvulsants

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

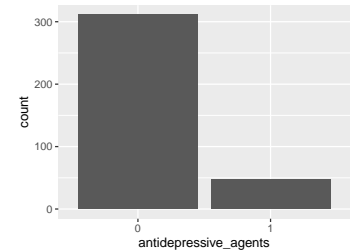
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antidepressive_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

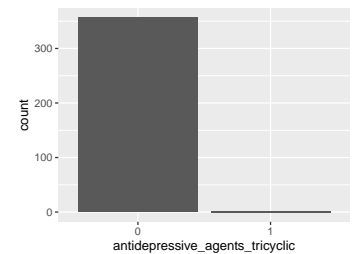
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antidepressive_agents_tricyclic

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

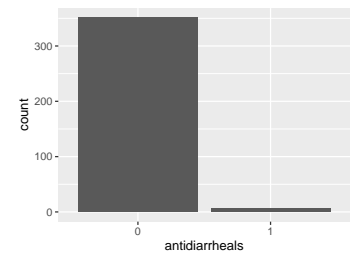


- Note that the following levels have at most five observations: "1".

antidiarrheals

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

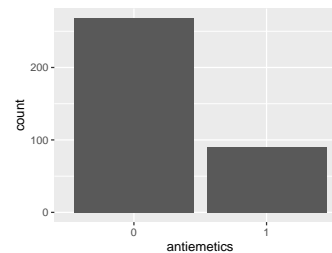
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antiemetics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

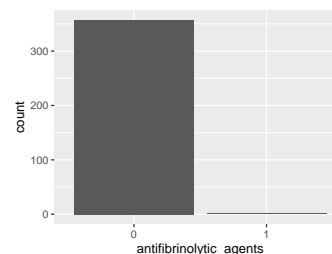
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antifibrinolytic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

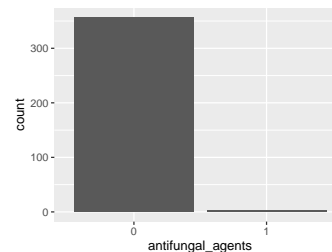


- Note that the following levels have at most five observations: "1".

antifungal_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

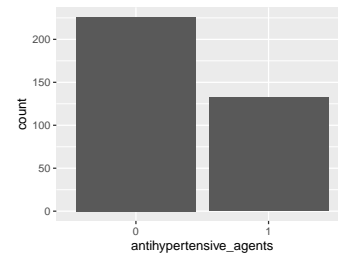


- Note that the following levels have at most five observations: "1".

antihypertensive__agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

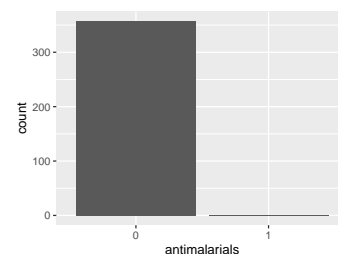
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antimalarials

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

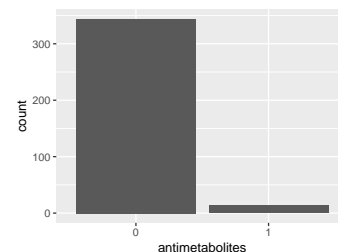


- Note that the following levels have at most five observations: "1".

antimetabolites

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

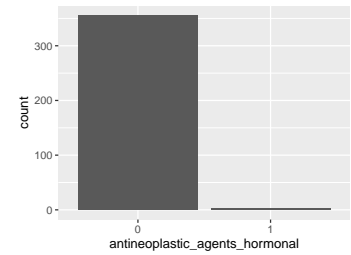
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antineoplastic__agents__hormonal

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

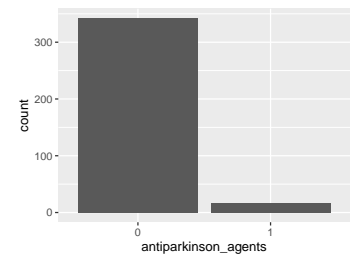


- Note that the following levels have at most five observations: "1".

antiparkinson_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

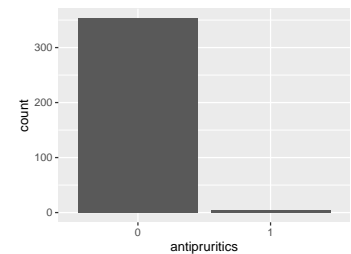
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antipruritics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

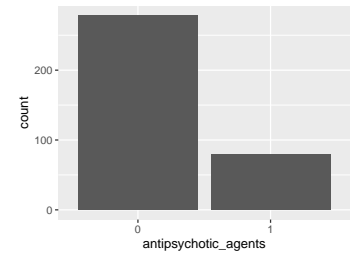


- Note that the following levels have at most five observations: "1".

antipsychotic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

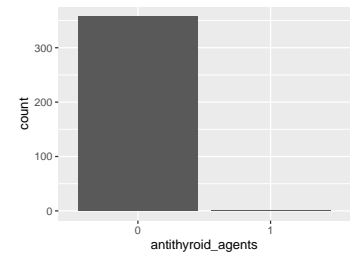
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



antithyroid_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

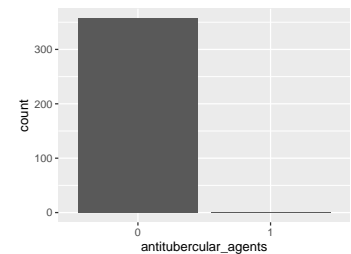


- Note that the following levels have at most five observations: "1".

antitubercular_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

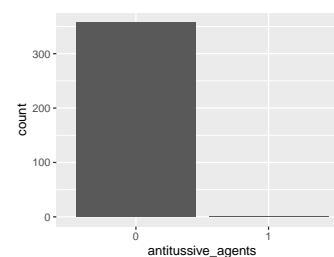


- Note that the following levels have at most five observations: "1".

antitussive_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

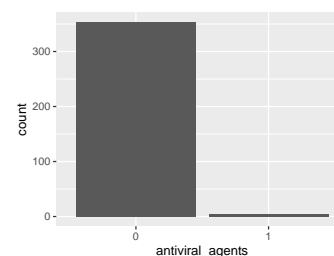


- Note that the following levels have at most five observations: "1".

antiviral_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

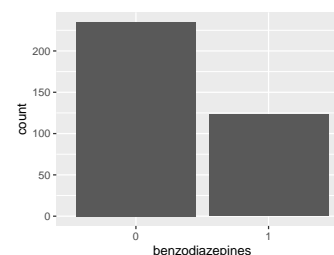


- Note that the following levels have at most five observations: "1".

benzodiazepines

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

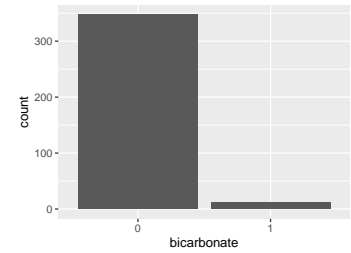
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



bicarbonate

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

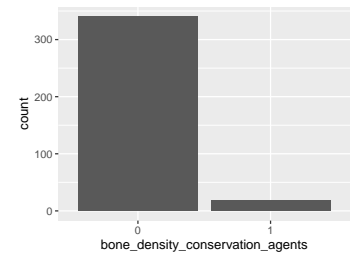
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



bone_density_conservation_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

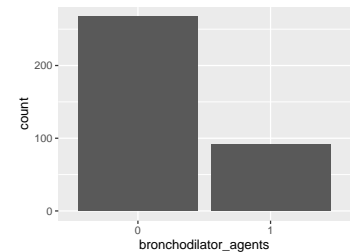
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



bronchodilator_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

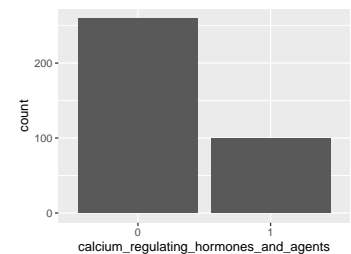
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



calcium_regulating_hormones_and_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

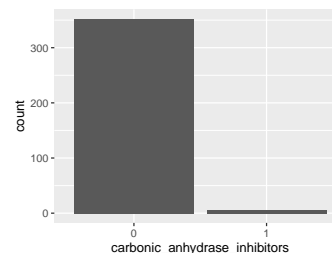
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



carbonic_anhydrase_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

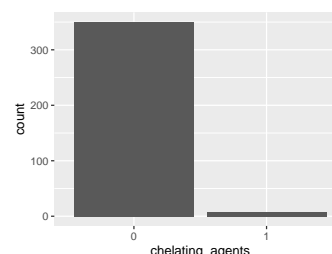
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



chelating_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

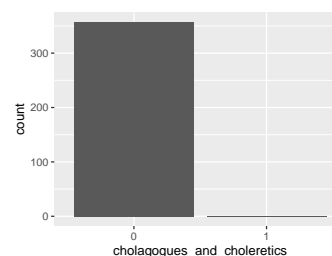
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



cholagogues_and_choleretics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

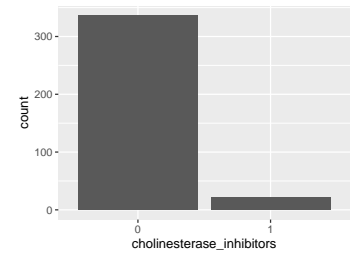


- Note that the following levels have at most five observations: "1".

cholinesterase_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

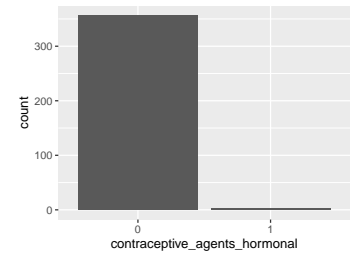
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



contraceptive_agents_hormonal

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

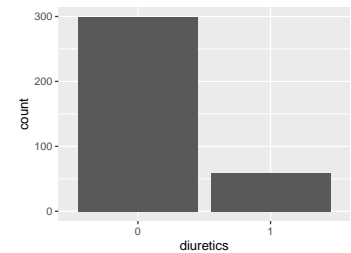


- Note that the following levels have at most five observations: "1".

diuretics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

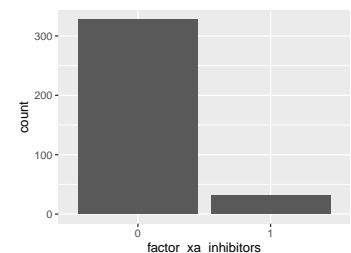
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



factor_xa_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

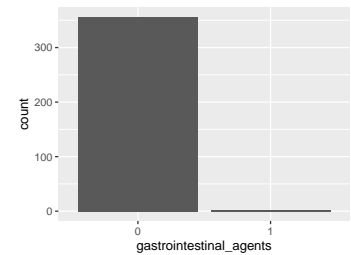
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



gastrointestinal_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

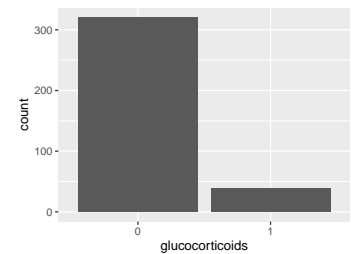


- Note that the following levels have at most five observations: "1".
-

glucocorticoids

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

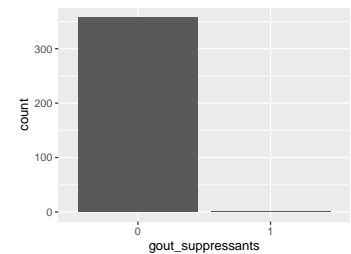
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



gout_suppressants

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

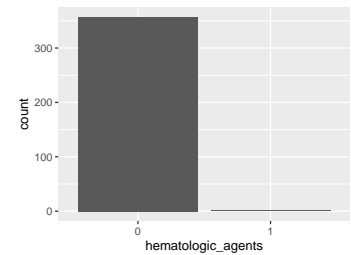


- Note that the following levels have at most five observations: "1".
-

hematologic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

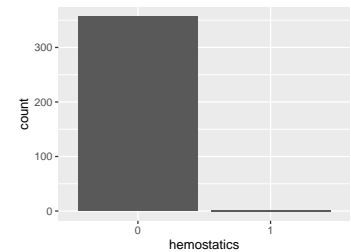


- Note that the following levels have at most five observations: "1".

hemostatics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

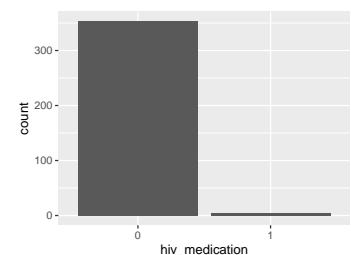


- Note that the following levels have at most five observations: "1".

hiv_medication

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

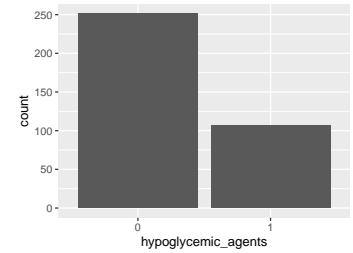


- Note that the following levels have at most five observations: "1".

hypoglycemic_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

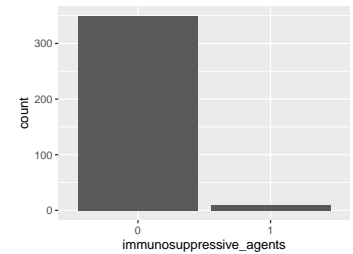
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



immunosuppressive_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

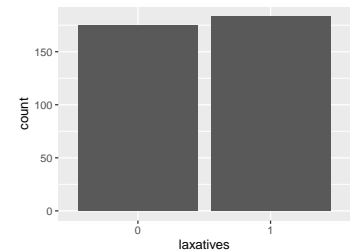
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



laxatives

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

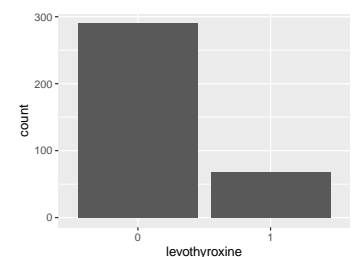
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"1"
Reference category	0



levothyroxine

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



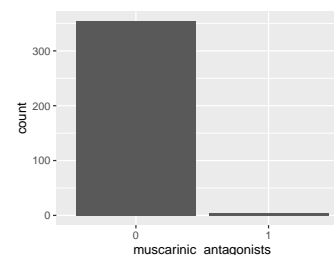
miotics

- The variable only takes one (non-missing) value: "0". The variable contains 0 % missing observations.
-

muscarinic_antagonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

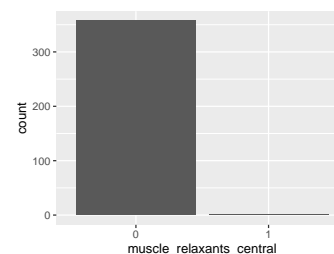


- Note that the following levels have at most five observations: "1".
-

muscle_relaxants_central

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

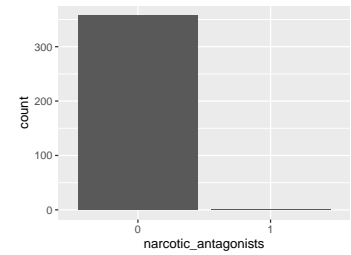


- Note that the following levels have at most five observations: "1".
-

narcotic_antagonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

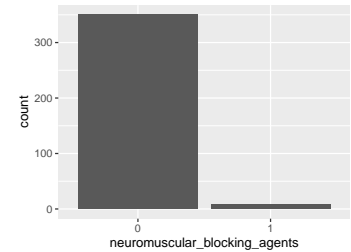


- Note that the following levels have at most five observations: "1".

neuromuscular_blocking_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

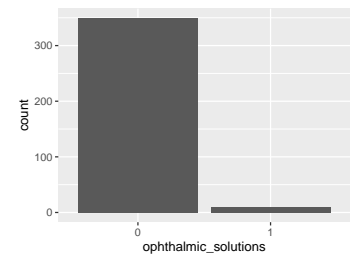
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



ophthalmic_solutions

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

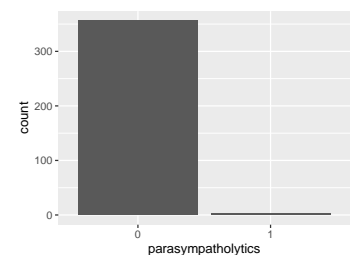
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



parasympatholytics

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

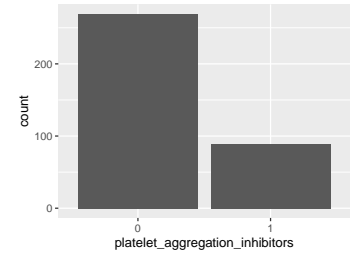


- Note that the following levels have at most five observations: "1".

platelet_aggregation_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

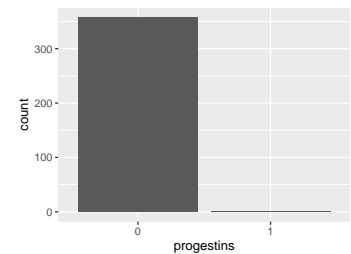
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



progestins

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

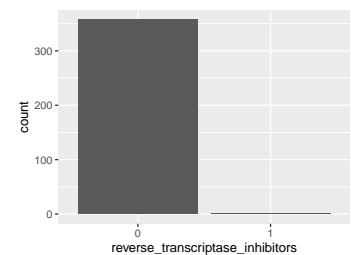


- Note that the following levels have at most five observations: "1".

reverse_transcriptase_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

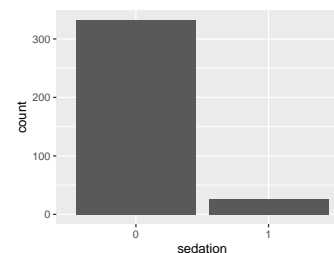


- Note that the following levels have at most five observations: "1".

sedation

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

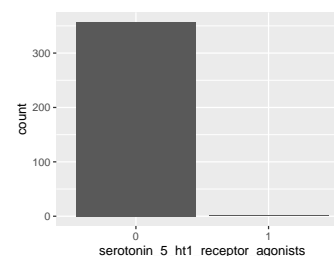
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



serotonin_5_ht1_receptor_agonists

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

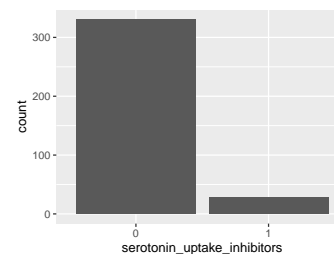


- Note that the following levels have at most five observations: "1".

serotonin_uptake_inhibitors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

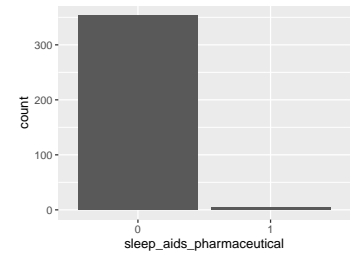
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



sleep_aids_pharmaceutical

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0

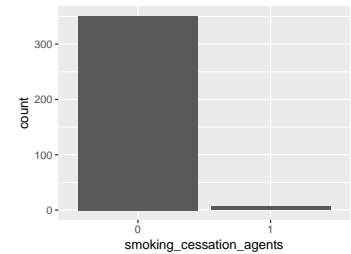


- Note that the following levels have at most five observations: "1".

smoking_cessation_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

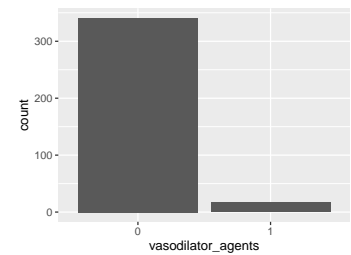
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasodilator_agents

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

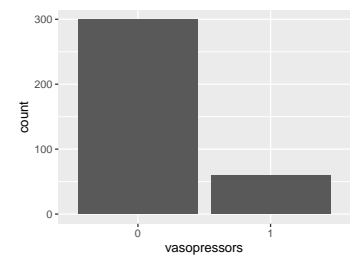
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasopressors

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

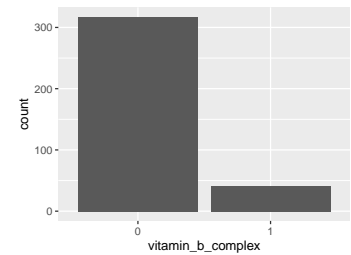
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamin_b_complex

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

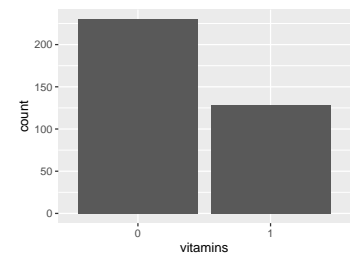
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamins

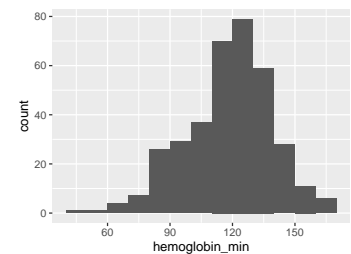
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



hemoglobin_min

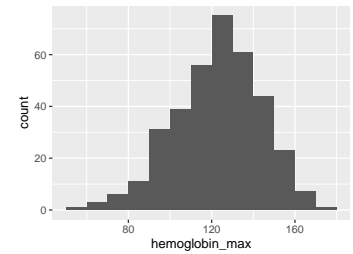
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	88
Median	121
1st and 3rd quartiles	108; 133
Min. and max.	41; 166



- Note that the following possible outlier values were detected: "41", "53", "158", "160", "161", "163", "165", "166".

hemoglobin__max

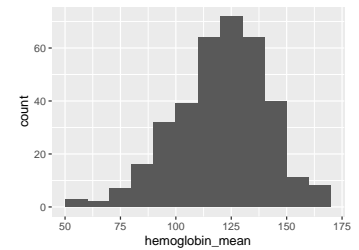
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	86
Median	125
1st and 3rd quartiles	110; 138.75
Min. and max.	59; 172



- Note that the following possible outlier values were detected: "172".

hemoglobin__mean

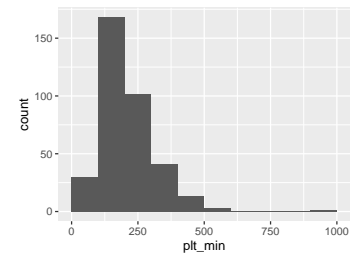
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	142
Median	123
1st and 3rd quartiles	109; 135.75
Min. and max.	56.4; 166.5



- Note that the following possible outlier values were detected: "56.4", "56.67".

plt__min

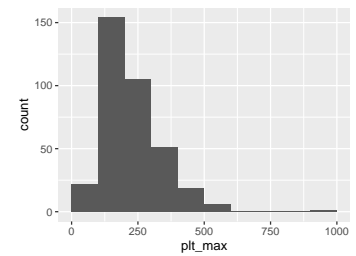
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	207
Median	187
1st and 3rd quartiles	141; 263.75
Min. and max.	21; 941



- Note that the following possible outlier values were detected: "21", "26", "37", "43", "47", "55", "56", "58", "59", "64" (2 additional values omitted).

plt_max

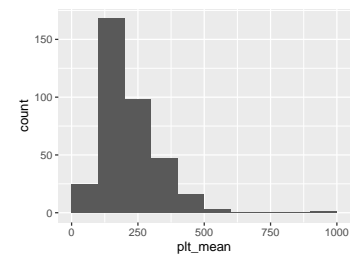
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	207
Median	202
1st and 3rd quartiles	152; 288
Min. and max.	26; 941



- Note that the following possible outlier values were detected: "26", "36", "43", "44", "55", "56", "58", "59", "61", "66" (1 additional values omitted).

plt_mean

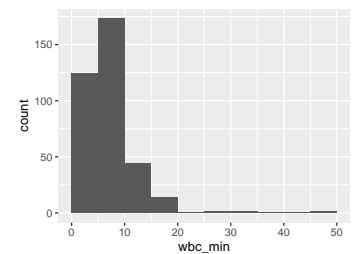
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	242
Median	194
1st and 3rd quartiles	146.25; 276.5
Min. and max.	23.6; 941



- Note that the following possible outlier values were detected: "23.6", "31", "40", "43.5", "54", "55", "56", "58", "59", "65" (2 additional values omitted).

wbc_min

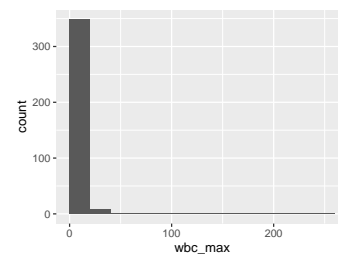
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	115
Median	6.2
1st and 3rd quartiles	4.5; 8.7
Min. and max.	1; 46.1



- Note that the following possible outlier values were detected: "1", "1.2", "1.3", "1.6", "26.9", "32", "46.1".

wbc__max

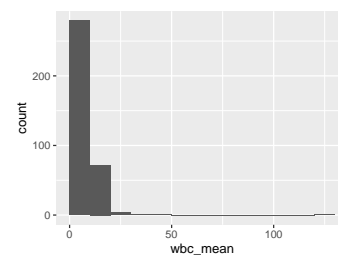
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	124
Median	6.85
1st and 3rd quartiles	4.9; 10.35
Min. and max.	1.3; 250



- Note that the following possible outlier values were detected: "1.3", "1.6", "1.7", "1.8", "2.1", "2.2", "32", "51.2", "250".

wbc__mean

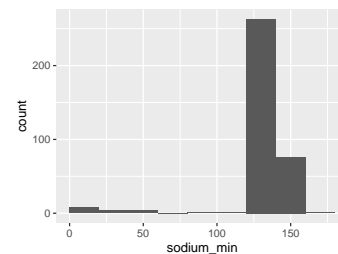
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	169
Median	6.6
1st and 3rd quartiles	4.75; 9.57
Min. and max.	1.3; 127



- Note that the following possible outlier values were detected: "1.3", "1.46", "1.6", "1.7", "2.05", "2.1", "26.9", "32", "48.65", "127".

sodium__min

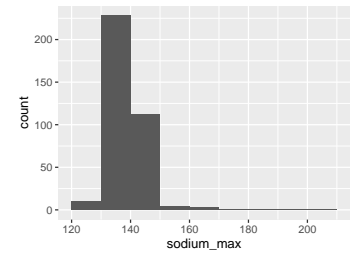
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	37
Median	138
1st and 3rd quartiles	135; 140
Min. and max.	10; 162



- Note that the following possible outlier values were detected: "10", "12", "13", "14", "22", "31", "39", "48", "53", "55" (9 additional values omitted).

sodium_max

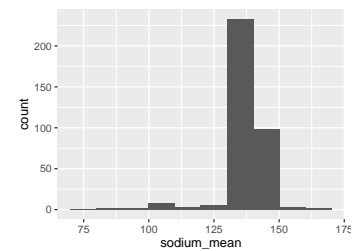
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	30
Median	139
1st and 3rd quartiles	137; 141
Min. and max.	126; 204



- Note that the following possible outlier values were detected: "126", "127", "129", "130", "148", "149", "156", "158", "159", "161" (3 additional values omitted).

sodium_mean

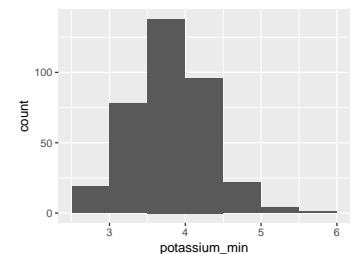
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	78
Median	138
1st and 3rd quartiles	136; 141
Min. and max.	72; 162



- Note that the following possible outlier values were detected: "72", "89.67", "90", "91.67", "92", "100.5", "103.67", "105.8", "108.5", "109.25" (10 additional values omitted).

potassium_min

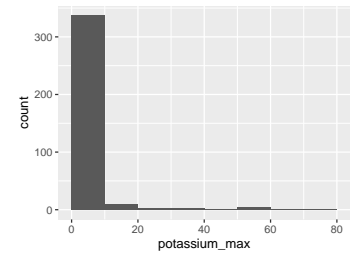
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	27
Median	3.9
1st and 3rd quartiles	3.5; 4.2
Min. and max.	2.6; 5.6



- Note that the following possible outlier values were detected: "4.9", "5", "5.1", "5.2", "5.6".

potassium__max

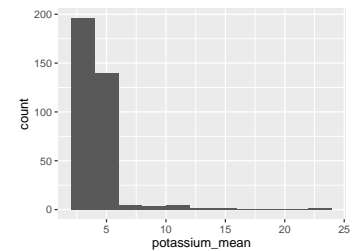
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	45
Median	4.1
1st and 3rd quartiles	3.8; 4.4
Min. and max.	2.9; 80



- Note that the following possible outlier values were detected: "2.9", "5.4", "5.5", "5.6", "5.7", "6.9", "14", "15", "16", "18" (11 additional values omitted).

potassium__mean

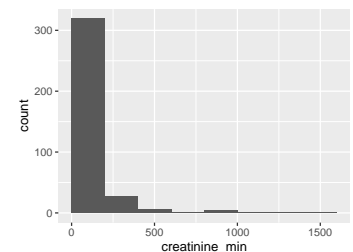
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	79
Median	4
1st and 3rd quartiles	3.65; 4.3
Min. and max.	2.9; 23.58



- Note that the following possible outlier values were detected: "2.9", "5.6", "6.02", "6.3", "7.33", "7.4", "8.57", "9.23", "9.45", "10.1" (10 additional values omitted).

creatinine__min

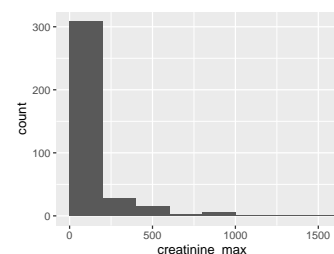
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	152
Median	73
1st and 3rd quartiles	55; 102.75
Min. and max.	1.3; 1455



- Note that the following possible outlier values were detected: "1.3", "1.9", "2", "2.4", "2.8", "3.4", "5", "5.4", "5.6", "5.8" (37 additional values omitted).

creatinine__max

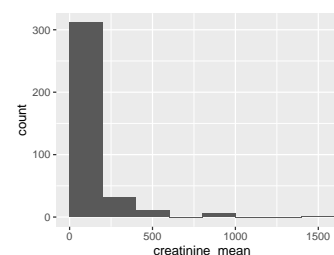
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	144
Median	82
1st and 3rd quartiles	63.25; 120.5
Min. and max.	27; 1457



- Note that the following possible outlier values were detected: "27", "29", "34", "36", "38", "39", "40", "41", "42", "43" (18 additional values omitted).

creatinine__mean

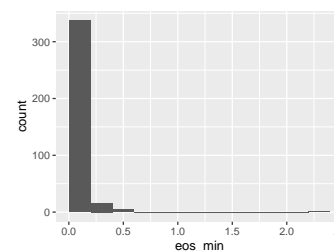
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	187
Median	76
1st and 3rd quartiles	60; 109.43
Min. and max.	21.45; 1456



- Note that the following possible outlier values were detected: "21.45", "25.77", "26.5", "27", "28.17", "29", "33", "36", "38.45", "38.5" (29 additional values omitted).

eos__min

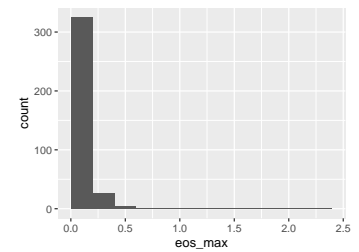
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	31
Median	0.01
1st and 3rd quartiles	0; 0.06
Min. and max.	0; 2.22



- Note that the following possible outlier values were detected: "2.22".

eos__max

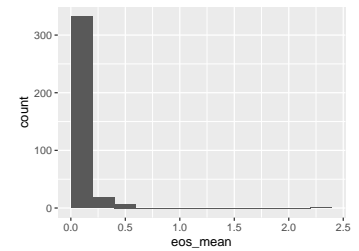
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	35
Median	0.01
1st and 3rd quartiles	0; 0.07
Min. and max.	0; 2.22



- Note that the following possible outlier values were detected: "2.22".

eos__mean

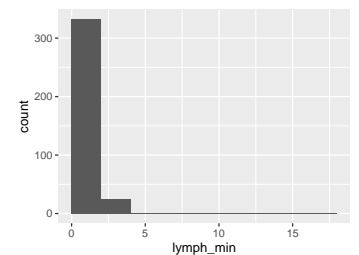
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	33
Median	0.01
1st and 3rd quartiles	0; 0.07
Min. and max.	0; 2.22



- Note that the following possible outlier values were detected: "2.22".

lymph__min

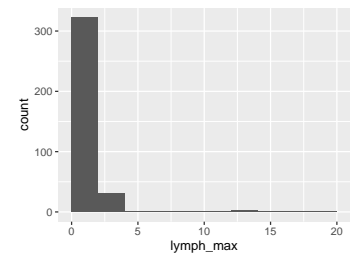
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	153
Median	0.92
1st and 3rd quartiles	0.57; 1.31
Min. and max.	0.1; 16.9



- Note that the following possible outlier values were detected: "3.07", "3.11", "3.14", "3.25", "3.81", "5.85", "16.9".

lymph_max

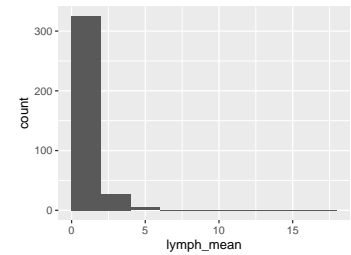
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	165
Median	1.07
1st and 3rd quartiles	0.72; 1.53
Min. and max.	0.1; 18.6



- Note that the following possible outlier values were detected: "0.1", "3.81", "6", "6.89", "14", "18.6".

lymph_mean

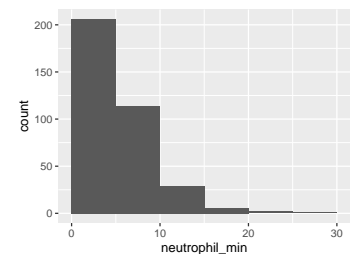
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	164
Median	0.97
1st and 3rd quartiles	0.69; 1.42
Min. and max.	0.1; 17.75



- Note that the following possible outlier values were detected: "0.1", "0.15", "0.18", "0.24", "0.25", "3.81", "4.85", "5.5", "5.93", "17.75".

neutrophil_min

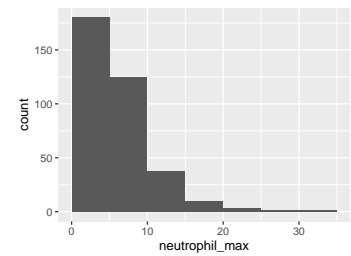
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	257
Median	4.47
1st and 3rd quartiles	2.98; 6.99
Min. and max.	0.09; 29.76



- Note that the following possible outlier values were detected: "0.09", "0.37", "0.7", "0.94", "1.07", "1.13", "1.16", "24.69", "24.83", "29.76".

neutrophil_max

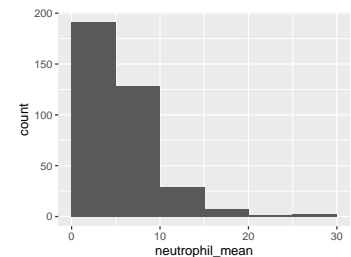
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	265
Median	4.98
1st and 3rd quartiles	3.34; 7.6
Min. and max.	0.18; 32.26



- Note that the following possible outlier values were detected: "0.18", "0.48", "0.7", "0.94", "1.07", "1.15", "1.16", "1.29", "1.36", "24.83" (2 additional values omitted).

neutrophil_mean

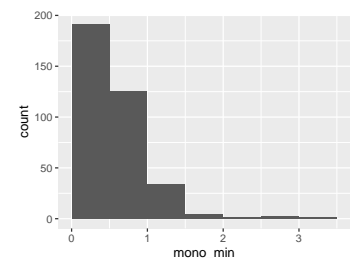
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	258
Median	4.72
1st and 3rd quartiles	3.18; 7.4
Min. and max.	0.14; 29.76



- Note that the following possible outlier values were detected: "0.14", "0.41", "0.7", "0.94", "1.07", "1.11", "1.16", "1.29", "1.36", "24.83" (2 additional values omitted).

mono_min

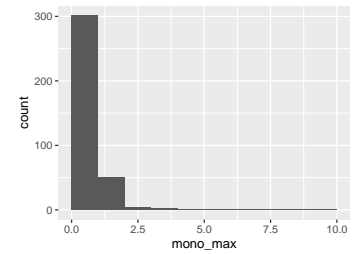
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	112
Median	0.48
1st and 3rd quartiles	0.34; 0.73
Min. and max.	0; 3.39



- Note that the following possible outlier values were detected: "0", "0.03", "0.06", "0.08", "0.1", "0.13", "0.14", "2.18", "2.51", "2.85" (1 additional values omitted).

mono__max

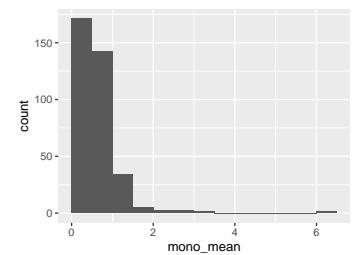
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	117
Median	0.55
1st and 3rd quartiles	0.4; 0.8
Min. and max.	0.06; 9.5



- Note that the following possible outlier values were detected: "0.06", "0.08", "0.09", "0.1", "0.13", "0.14", "0.15", "0.16", "0.18", "0.19" (7 additional values omitted).

mono__mean

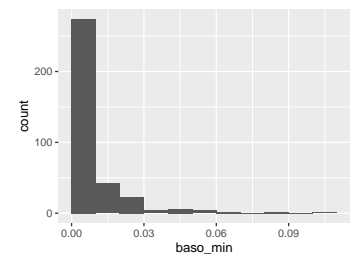
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	114
Median	0.53
1st and 3rd quartiles	0.38; 0.78
Min. and max.	0.05; 6.18



- Note that the following possible outlier values were detected: "0.05", "0.06", "0.08", "0.12", "0.13", "0.14", "2.08", "2.34", "2.51", "3.39" (1 additional values omitted).

baso__min

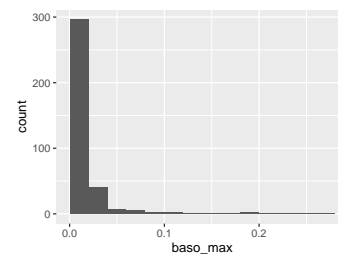
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	10
Median	0.01
1st and 3rd quartiles	0; 0.01
Min. and max.	0; 0.11



- Note that the following possible outlier values were detected: "0.03", "0.04", "0.05", "0.06", "0.07", "0.09", "0.11".

baso__max

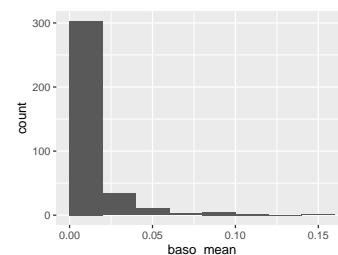
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	14
Median	0.01
1st and 3rd quartiles	0.01; 0.02
Min. and max.	0; 0.27



- Note that the following possible outlier values were detected: "0", "0.12", "0.19", "0.27".

baso__mean

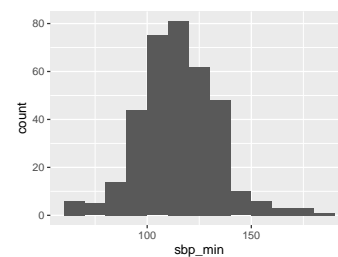
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	12
Median	0.01
1st and 3rd quartiles	0.01; 0.02
Min. and max.	0; 0.16



- Note that the following possible outlier values were detected: "0", "0.09", "0.1", "0.11", "0.16".

sbp__min

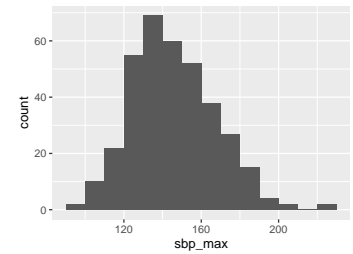
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	79
Median	115
1st and 3rd quartiles	102; 125
Min. and max.	60; 185



- Note that the following possible outlier values were detected: "60", "62", "70", "164", "171", "173", "179", "185".

sbp_max

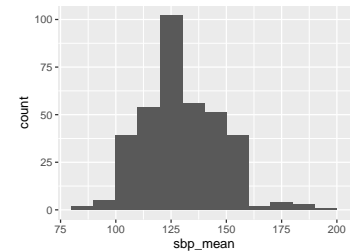
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	85
Median	144
1st and 3rd quartiles	131; 160
Min. and max.	97; 223



- Note that the following possible outlier values were detected: "97", "98", "221", "223".

sbp_mean

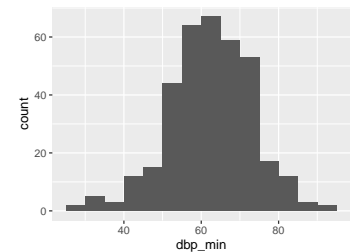
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	227
Median	127.88
1st and 3rd quartiles	118.09; 141
Min. and max.	87.57; 193.4



- Note that the following possible outlier values were detected: "87.57", "90.67", "90.71", "92.4", "94", "96.2", "193.4".

dbp_min

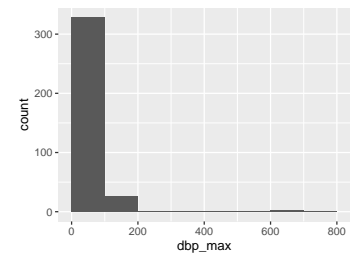
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	53
Median	63
1st and 3rd quartiles	56; 70
Min. and max.	29; 93



- Note that the following possible outlier values were detected: "29", "30", "93".

dbp__max

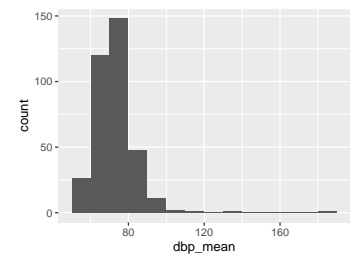
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	61
Median	80
1st and 3rd quartiles	75; 88
Min. and max.	57; 787



- Note that the following possible outlier values were detected: "57", "59", "60", "61", "62", "63", "64", "65", "66", "125" (5 additional values omitted).

dbp__mean

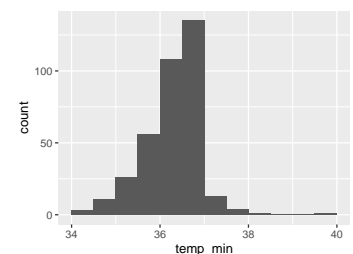
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	203
Median	72.15
1st and 3rd quartiles	66; 77
Min. and max.	50.4; 184.6



- Note that the following possible outlier values were detected: "91.27", "92", "92.4", "92.94", "93", "93.87", "99.2", "101.75", "104.6", "113.31" (2 additional values omitted).

temp__min

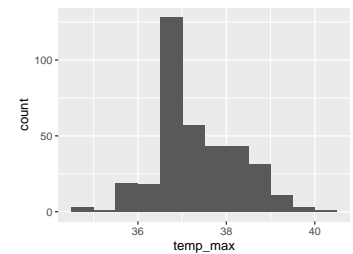
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	35
Median	36.5
1st and 3rd quartiles	36; 37
Min. and max.	34.2; 39.6



- Note that the following possible outlier values were detected: "38.5", "39.6".

temp_max

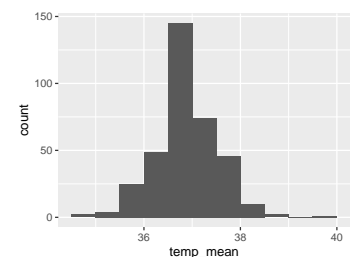
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	46
Median	37.1
1st and 3rd quartiles	37; 38
Min. and max.	34.5; 40.2



- Note that the following possible outlier values were detected: "34.5", "35", "35.2", "35.6", "35.7", "35.8", "35.9", "36", "36.1", "36.2" (6 additional values omitted).

temp_mean

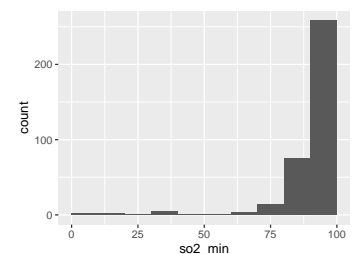
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	139
Median	36.9
1st and 3rd quartiles	36.56; 37.3
Min. and max.	34.5; 39.6



- Note that the following possible outlier values were detected: "34.5", "35", "35.1", "35.25", "38.5", "38.75", "39.6".

so2_min

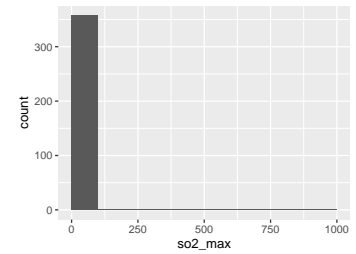
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	32
Median	93
1st and 3rd quartiles	90; 95
Min. and max.	0; 100



- Note that the following possible outlier values were detected: "0", "2", "18", "20", "32", "36", "63", "65", "69", "72" (4 additional values omitted).

so2__max

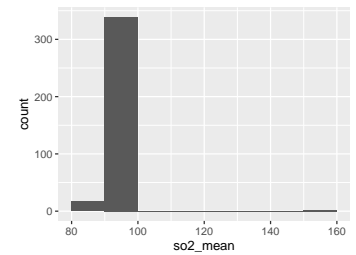
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	11
Median	97
1st and 3rd quartiles	96; 99
Min. and max.	91; 969



- Note that the following possible outlier values were detected: "91", "969".

so2__mean

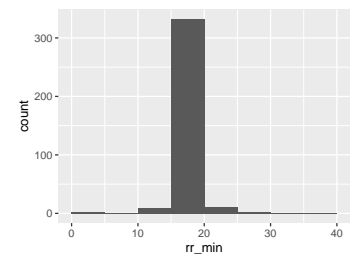
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	157
Median	95
1st and 3rd quartiles	93.67; 96.75
Min. and max.	80.71; 152.87



- Note that the following possible outlier values were detected: "80.71", "81.07", "85.38", "85.91", "86.78", "86.89", "88", "88.33", "88.64", "152.87".

rr__min

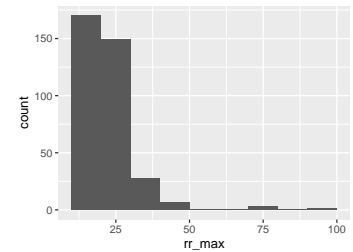
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	15
Median	18
1st and 3rd quartiles	18; 20
Min. and max.	0; 40



- Note that the following possible outlier values were detected: "0", "2", "12", "14", "24", "30", "32", "40".

rr_max

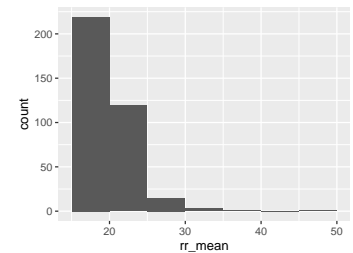
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	25
Median	22
1st and 3rd quartiles	20; 24
Min. and max.	16; 98



- Note that the following possible outlier values were detected: "16", "18", "42", "44", "48", "50", "80", "98".

rr_mean

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	104
Median	20
1st and 3rd quartiles	19; 21
Min. and max.	16; 46

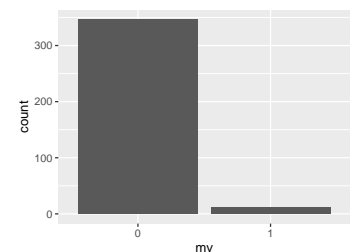


- Note that the following possible outlier values were detected: "16", "16.67", "16.86", "16.89", "17", "17.14", "17.5", "17.6", "17.78", "27.69" (8 additional values omitted).

mv

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

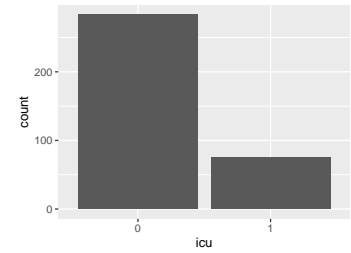
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



icu

- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



Report generation information:

- Created by: Eric Yamga (username: `eyamga`).
- Report creation time: Sat Jan 23 2021 21:19:59
- Report was run from directory: `/Users/eyamga/Documents/Médecine/Recherche/CODA19/code/r_eyamga`
- dataMaid v1.4.0 [Pkg: 2019-12-10 from CRAN (R 4.0.2)]
- R version 4.0.3 (2020-10-10).
- Platform: x86_64-apple-darwin17.0 (64-bit)(macOS Catalina 10.15.7).
- Function call: `dataMaid::makeDataReport(data = covid24h_imputed, render = FALSE, file = "coda19CHUM_imputed.rmd", replace = TRUE)`