

covid72h_notimputed

Autogenerated data summary from dataMaid

2021-02-01 20:23:44

Data report overview

The dataset examined has the following dimensions:

Feature	Result
Number of observations	1061
Number of variables	293

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	1009	0.00 %	
female	numeric	2	0.00 %	
male	numeric	2	0.00 %	
patient_age	numeric	86	0.00 %	×
death	numeric	2	0.00 %	
ami	numeric	3	75.21 %	
chf	numeric	3	75.21 %	
pvd	numeric	3	75.21 %	×
cevd	numeric	3	75.21 %	
dementia	numeric	3	75.21 %	
copd	numeric	3	75.21 %	
rheumd	numeric	3	75.21 %	×
pud	numeric	3	75.21 %	×
mld	numeric	3	75.21 %	×
diab	numeric	3	75.21 %	
diabwc	numeric	3	75.21 %	
hp	numeric	3	75.21 %	×
rend	numeric	3	75.21 %	
canc	numeric	3	75.21 %	
msld	numeric	3	75.21 %	×
metacanc	numeric	3	75.21 %	
aids	numeric	2	75.21 %	×
score	numeric	5	75.21 %	
neuromuscular_blocking_agents	numeric	3	7.92 %	
x5_alpha_reductase_inhibitors	numeric	3	7.92 %	
acetaminophene	numeric	3	7.92 %	
adjuvants_anesthesia	numeric	3	7.92 %	×
adrenergic_alpha_1_receptor_antagonists	numeric	3	7.92 %	
adrenergic_beta_3_receptor_agonists	numeric	3	7.92 %	
adrenergic_beta_antagonists	numeric	3	7.92 %	
adrenergic_uptake_inhibitors	numeric	3	7.92 %	×
alcohol_deterrents	numeric	3	7.92 %	×
analgesics	numeric	3	7.92 %	
analgesics_opioid	numeric	3	7.92 %	
androgens	numeric	3	7.92 %	×
anesthetics_local	numeric	3	7.92 %	
anti_anxiety_agents	numeric	3	7.92 %	×
anti_arrhythmia_agents	numeric	3	7.92 %	
anti_asthmatic_agents	numeric	3	7.92 %	
anti_bacterial_agents	numeric	3	7.92 %	
anti_infective_agents_local	numeric	3	7.92 %	
anti_inflammatory_agents	numeric	3	7.92 %	

	Variable class	# unique values	Missing observations	Any problems?
anti_inflammatory_agents_non_steroidal	numeric	3	7.92 %	
anti_ulcer_agents	numeric	3	7.92 %	
anticholesteremic_agents	numeric	3	7.92 %	
anticoagulants	numeric	3	7.92 %	
anticonvulsants	numeric	3	7.92 %	
antidepressive_agents	numeric	3	7.92 %	
antidepressive_agents_tricyclic	numeric	3	7.92 %	×
antidiarrheals	numeric	3	7.92 %	
antiemetics	numeric	3	7.92 %	
antifibrinolytic_agents	numeric	3	7.92 %	
antifungal_agents	numeric	3	7.92 %	×
antihypertensive_agents	numeric	3	7.92 %	
antimalarials	numeric	3	7.92 %	×
antimetabolites	numeric	3	7.92 %	
antineoplastic_agents_hormonal	numeric	3	7.92 %	×
antiparkinson_agents	numeric	3	7.92 %	
antipruritics	numeric	3	7.92 %	
antipsychotic_agents	numeric	3	7.92 %	
antithyroid_agents	numeric	3	7.92 %	×
antitubercular_agents	numeric	3	7.92 %	×
antitussive_agents	numeric	3	7.92 %	
antiviral_agents	numeric	3	7.92 %	
benzodiazepines	numeric	3	7.92 %	
bicarbonate	numeric	3	7.92 %	
bone_density_conservation_agents	numeric	3	7.92 %	
bronchodilator_agents	numeric	3	7.92 %	
calcium_regulating_hormones_and_agents	numeric	3	7.92 %	
carbonic_anhydrase_inhibitors	numeric	3	7.92 %	
chelating_agents	numeric	3	7.92 %	
cholagogues_and_choleretics	numeric	3	7.92 %	
cholinesterase_inhibitors	numeric	3	7.92 %	
contraceptive_agents_hormonal	numeric	3	7.92 %	×
diuretics	numeric	3	7.92 %	
diuretics_osmotic	numeric	3	7.92 %	×
factor_xa_inhibitors	numeric	3	7.92 %	
fibrinolytic_agents	numeric	3	7.92 %	×
gastrointestinal_agents	numeric	3	7.92 %	
glucocorticoids	numeric	3	7.92 %	
gout_suppressants	numeric	3	7.92 %	
hematologic_agents	numeric	3	7.92 %	×
hemostatics	numeric	3	7.92 %	×
hiv_medication	numeric	3	7.92 %	
hypoglycemic_agents	numeric	3	7.92 %	
immunologic_factors	numeric	3	7.92 %	×
immunosuppressive_agents	numeric	3	7.92 %	
laxatives	numeric	3	7.92 %	
levothyroxine	numeric	3	7.92 %	
miotics	numeric	3	7.92 %	×
muscarinic_antagonists	numeric	3	7.92 %	
muscle_relaxants_central	numeric	3	7.92 %	×
narcotic_antagonists	numeric	3	7.92 %	×
neuromuscular_blocking_agents_2	numeric	3	7.92 %	
ophthalmic_solutions	numeric	3	7.92 %	
parasympatholytics	numeric	3	7.92 %	

	Variable class	# unique values	Missing observations	Any problems?
platelet_aggregation_inhibitors	numeric	3	7.92 %	
progestins	numeric	3	7.92 %	×
reverse_transcriptase_inhibitors	numeric	3	7.92 %	×
sedation	numeric	3	7.92 %	
serotonin_5_ht1_receptor_agonists	numeric	3	7.92 %	×
serotonin_uptake_inhibitors	numeric	3	7.92 %	
sleep_aids_pharmaceutical	numeric	3	7.92 %	
smoking_cessation_agents	numeric	3	7.92 %	
vasodilator_agents	numeric	3	7.92 %	
vasopressors	numeric	3	7.92 %	
vitamin_b_complex	numeric	3	7.92 %	
vitamins	numeric	3	7.92 %	
hemoglobin_min	numeric	107	17.34 %	×
hemoglobin_max	numeric	99	17.34 %	×
hemoglobin_mean	numeric	361	17.34 %	×
plt_min	numeric	316	17.34 %	×
plt_max	numeric	342	17.34 %	×
plt_mean	numeric	599	17.34 %	×
wbc_min	numeric	133	17.34 %	×
wbc_max	numeric	187	17.34 %	×
wbc_mean	numeric	402	17.34 %	×
albumin_min	numeric	37	41.66 %	×
albumin_max	numeric	32	41.66 %	×
albumin_mean	numeric	135	41.66 %	×
globulin_min	numeric	5	99.62 %	×
globulin_max	numeric	5	99.62 %	×
globulin_mean	numeric	5	99.62 %	×
protein_min	numeric	36	89.16 %	×
protein_max	numeric	34	89.16 %	×
protein_mean	numeric	44	89.16 %	×
sodium_min	numeric	39	18.00 %	×
sodium_max	numeric	37	18.00 %	×
sodium_mean	numeric	180	18.00 %	×
chloride_min	numeric	36	25.73 %	×
chloride_max	numeric	42	25.73 %	×
chloride_mean	numeric	180	25.73 %	×
potassium_min	numeric	33	18.19 %	×
potassium_max	numeric	35	18.19 %	×
potassium_mean	numeric	136	18.19 %	×
bicarbonate_min	numeric	154	39.21 %	×
bicarbonate_max	numeric	137	39.21 %	×
bicarbonate_mean	numeric	333	39.21 %	×
bun_min	numeric	162	52.87 %	×
bun_max	numeric	187	52.87 %	×
bun_mean	numeric	273	52.87 %	×
calcium_min	logical	1	100.00 %	×
calcium_max	logical	1	100.00 %	×
calcium_mean	logical	1	100.00 %	×
magnesium_min	numeric	67	42.04 %	×
magnesium_max	numeric	76	42.04 %	×
magnesium_mean	numeric	65	42.04 %	×
phosphate_min	numeric	113	50.61 %	×
phosphate_max	numeric	139	50.61 %	×
phosphate_mean	numeric	116	50.61 %	×

	Variable class	# unique values	Missing observations	Any problems?
creatinine_min	numeric	180	18.10 %	×
creatinine_max	numeric	206	18.10 %	×
creatinine_mean	numeric	443	18.10 %	×
gfr_min	logical	1	100.00 %	×
gfr_max	logical	1	100.00 %	×
gfr_mean	logical	1	100.00 %	×
glucose_min	numeric	99	27.80 %	×
glucose_max	numeric	178	27.80 %	×
glucose_max_1	numeric	390	27.80 %	×
anion_gap_min	numeric	19	55.42 %	×
anion_gap_min_1	numeric	23	55.42 %	×
anion_gap_mean	numeric	57	55.42 %	×
eos_min	numeric	43	21.11 %	×
eos_max	numeric	53	21.11 %	×
eos_mean	numeric	46	21.11 %	×
lymph_min	numeric	217	17.34 %	×
lymph_max	numeric	238	17.34 %	×
lymph_mean	numeric	230	17.34 %	×
neutrophil_min	numeric	441	17.34 %	×
neutrophil_max	numeric	506	17.34 %	×
neutrophil_mean	numeric	528	17.34 %	×
mono_min	numeric	128	17.34 %	×
mono_max	numeric	160	17.34 %	×
mono_mean	numeric	142	17.34 %	×
baso_min	numeric	15	17.34 %	
baso_max	numeric	26	17.34 %	×
baso_mean	numeric	20	17.34 %	×
stab_min	numeric	17	97.74 %	×
stab_max	numeric	18	97.74 %	×
stab_mean	numeric	17	97.74 %	×
pt_min	numeric	18	85.96 %	×
pt_max	numeric	26	85.96 %	×
pt_mean	numeric	43	85.96 %	×
ptt_min	numeric	36	59.10 %	×
ptt_max	numeric	51	59.10 %	×
ptt_mean	numeric	106	59.10 %	×
fibrinogen_min	numeric	155	78.04 %	×
fibrinogen_max	numeric	162	78.04 %	×
fibrinogen_mean	numeric	164	78.04 %	×
d_dimer_min	numeric	222	76.06 %	×
d_dimer_max	numeric	228	76.06 %	×
d_dimer_mean	numeric	229	76.06 %	×
alt_min	numeric	104	35.34 %	×
alt_max	numeric	123	35.34 %	×
alt_mean	numeric	229	35.34 %	×
ast_min	numeric	99	64.66 %	×
ast_max	numeric	107	64.66 %	×
ast_mean	numeric	169	64.66 %	×
palc_min	numeric	141	40.72 %	×
palc_max	numeric	151	40.72 %	×
palc_mean	numeric	236	40.72 %	×
ggt_min	numeric	105	80.40 %	×
ggt_max	numeric	106	80.40 %	×
ggt_mean	numeric	118	80.40 %	×

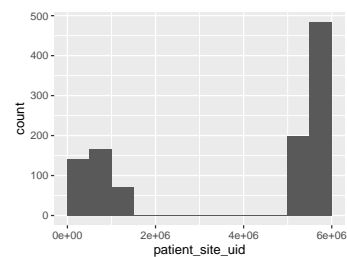
	Variable class	# unique values	Missing observations	Any problems?
amylase_min	logical	1	100.00 %	×
amylase_max	logical	1	100.00 %	×
amylase_mean	logical	1	100.00 %	×
lipase_min	numeric	99	63.24 %	×
lipase_max	numeric	114	63.24 %	×
lipase_mean	numeric	146	63.24 %	×
bili_tot_min	numeric	40	38.08 %	×
bili_tot_max	numeric	48	38.08 %	×
bili_tot_mean	numeric	121	38.08 %	×
bili_direct_min	numeric	42	95.19 %	×
bili_direct_max	numeric	44	95.19 %	×
bili_direct_mean	numeric	46	95.19 %	×
bili_indirect_min	numeric	47	95.19 %	×
bili_indirect_max	numeric	48	95.19 %	×
bili_indirect_mean	numeric	50	95.19 %	×
lipase_min_1	numeric	99	63.24 %	×
lipase_max_1	numeric	114	63.24 %	×
lipase_mean_1	numeric	146	63.24 %	×
ck_min	numeric	146	78.79 %	×
ck_max	numeric	157	78.79 %	×
ck_mean	numeric	169	78.79 %	×
ckmb_min	numeric	47	88.31 %	×
ckmb_max	numeric	57	88.31 %	×
ckmb_mean	numeric	72	88.31 %	×
ldh_min	numeric	219	67.30 %	×
ldh_max	numeric	226	67.30 %	×
ldh_mean	numeric	242	67.30 %	×
tropot_min	numeric	76	75.68 %	×
tropot_max	numeric	89	75.68 %	×
tropot_mean	numeric	130	75.68 %	×
lactate_min	numeric	29	70.03 %	×
lactate_max	numeric	45	70.03 %	×
lactate_mean	numeric	104	70.03 %	×
svo2sat_min	numeric	89	54.38 %	
svo2sat_max	numeric	84	54.38 %	
svo2sat_max_1	numeric	169	54.38 %	×
pao2_min	numeric	86	88.22 %	×
pao2_max	numeric	76	88.22 %	×
pao2_mean	numeric	96	88.22 %	×
pvo2_min	numeric	287	54.19 %	×
pvo2_max	numeric	322	54.19 %	×
pvo2_mean	numeric	347	54.19 %	×
paco2_min	numeric	221	54.19 %	×
paco2_max	numeric	237	54.19 %	×
paco2_mean	numeric	295	54.19 %	×
pvco2_min	numeric	221	54.19 %	×
pvco2_max	numeric	237	54.19 %	×
pvco2_mean	numeric	295	54.19 %	×
tsh_min	numeric	142	83.60 %	×
tsh_max	numeric	141	83.60 %	×
tsh_mean	numeric	142	83.60 %	×
vitd_min	numeric	17	98.49 %	×
vitd_max	numeric	17	98.49 %	×
vitd_mean	numeric	17	98.49 %	×

	Variable class	# unique values	Missing observations	Any problems?
crp_min	numeric	459	37.51 %	
crp_max	numeric	486	37.51 %	
crp_mean	numeric	511	37.51 %	
ferritin_min	numeric	120	87.37 %	×
ferritin_max	numeric	117	87.37 %	×
ferritin_mean	numeric	121	87.37 %	×
bnp_min	numeric	140	84.73 %	
bnp_max	numeric	142	84.73 %	
bnp_mean	numeric	142	84.73 %	
weight_min	numeric	301	59.47 %	×
weight_max	numeric	291	59.47 %	×
weight_mean	numeric	317	59.47 %	×
sbp_min	numeric	97	15.08 %	×
sbp_max	numeric	120	15.08 %	×
sbp_mean	numeric	649	15.08 %	×
dbp_min	numeric	67	15.08 %	×
dbp_max	numeric	78	15.08 %	×
dbp_mean	numeric	568	15.08 %	×
temp_min	numeric	37	15.36 %	×
temp_max	numeric	52	15.36 %	×
temp_mean	numeric	226	15.36 %	×
so2_min	numeric	53	9.05 %	×
so2_max	numeric	15	9.05 %	×
so2_mean	numeric	455	9.05 %	×
rr_min	numeric	21	15.08 %	×
rr_max	numeric	36	15.08 %	×
rr_mean	numeric	307	15.08 %	×
flow_min	numeric	14	66.73 %	
flow_max	numeric	17	66.73 %	×
flow_mean	numeric	163	66.73 %	×
fio2_min	numeric	40	43.36 %	
fio2_max	numeric	38	43.36 %	
fio2_mean	numeric	259	43.36 %	
mv	numeric	2	0.00 %	
icu	numeric	2	0.00 %	

Variable list

patient_site_uid

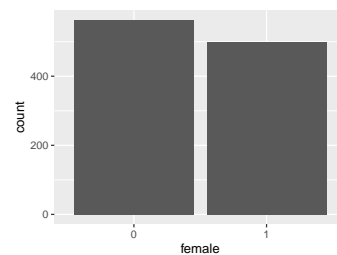
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	1009
Median	5361258
1st and 3rd quartiles	847996; 5637720
Min. and max.	720; 5683923



female

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

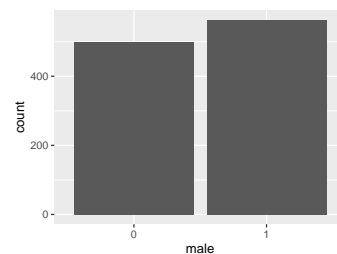
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“0”
Reference category	0



male

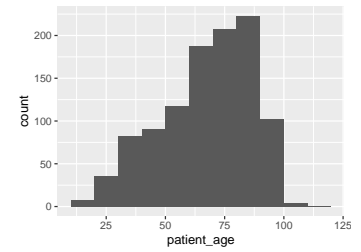
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

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



patient_age

Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	86
Median	71
1st and 3rd quartiles	55; 83
Min. and max.	12; 120

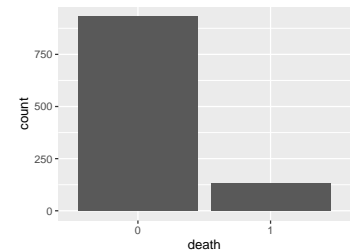


- Note that the following possible outlier values were detected: "120".

death

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

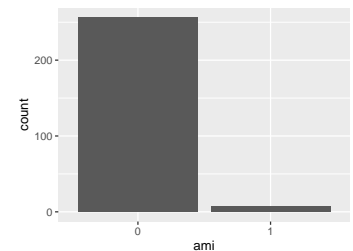
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



ami

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

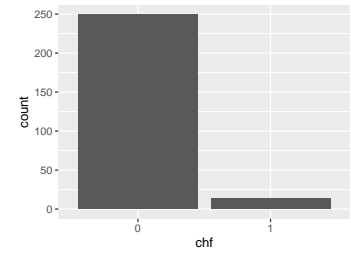
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



chf

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

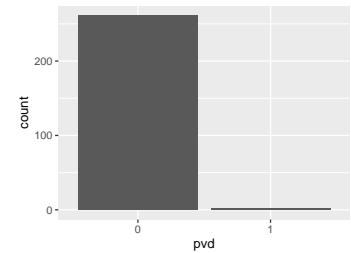
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



pvd

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

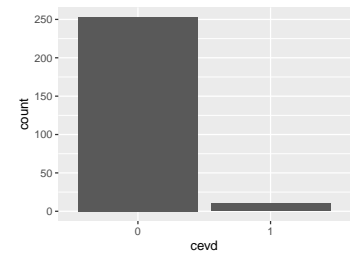


- Note that the following levels have at most five observations: "1".

cevd

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

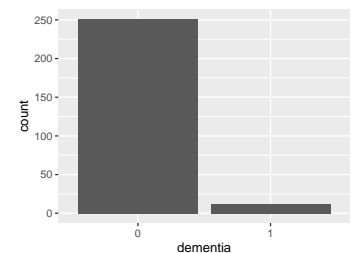
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



dementia

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

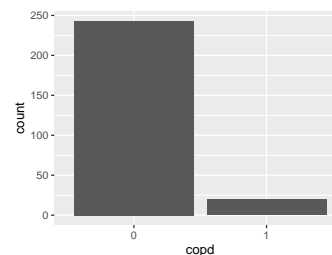
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



copd

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

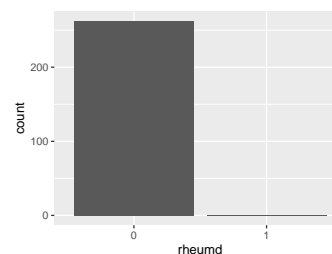
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



rheumd

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

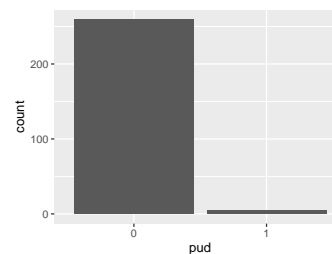


- Note that the following levels have at most five observations: "1".

pud

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

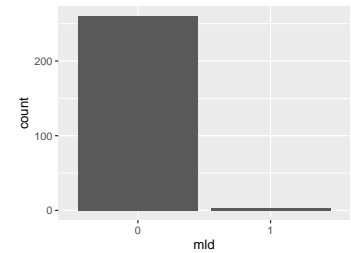


- Note that the following levels have at most five observations: "1".

mld

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

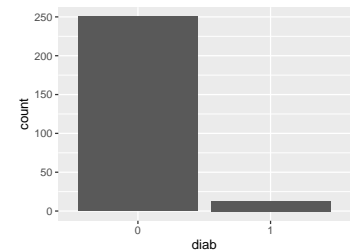


- Note that the following levels have at most five observations: "1".

diab

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

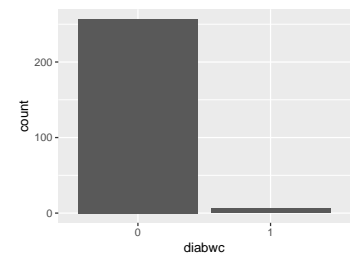
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



diabwc

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

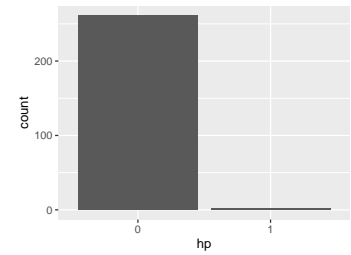
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



hp

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

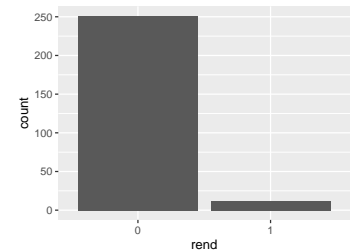


- Note that the following levels have at most five observations: "1".

rend

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

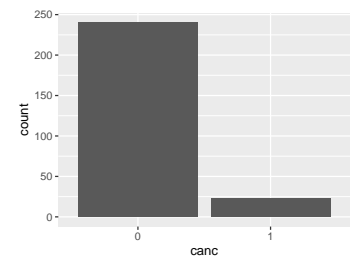
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



canc

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

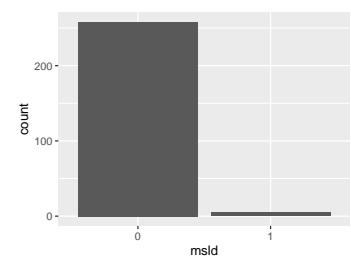
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



msld

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0

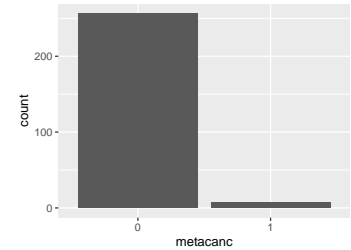


- Note that the following levels have at most five observations: "1".

metacanc

- 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.	798 (75.21 %)
Number of unique values	2
Mode	"0"
Reference category	0



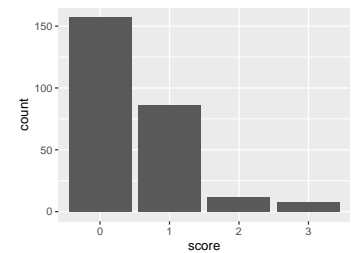
aids

- The variable only takes one (non-missing) value: "0". The variable contains 75.21 % missing observations.

score

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

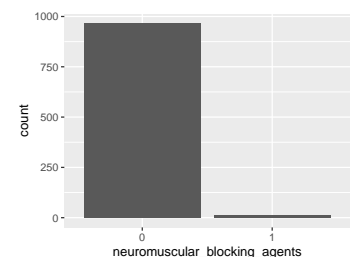
Feature	Result
Variable type	numeric
Number of missing obs.	798 (75.21 %)
Number of unique values	4
Mode	"0"
Reference category	0



neuromuscular_blocking_agents

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

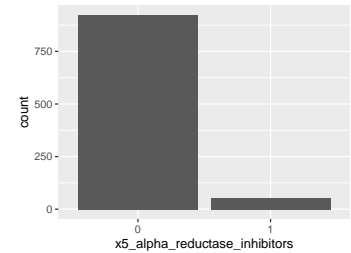
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



x5_alpha_reductase_inhibitors

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

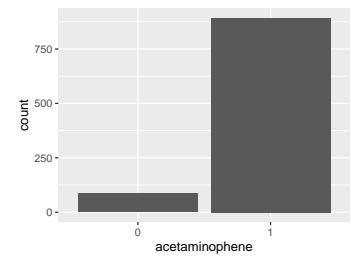
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



acetaminophene

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

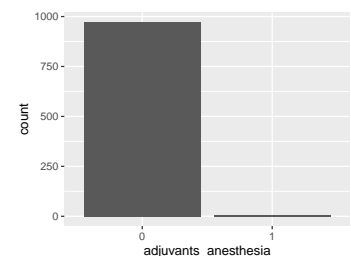
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"1"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

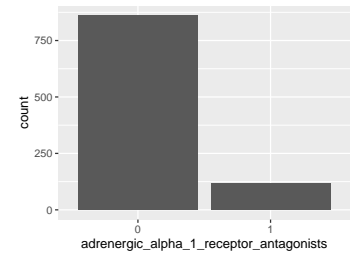


- Note that the following levels have at most five observations: "1".

adrenergic_alpha_1_receptor_antagonists

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

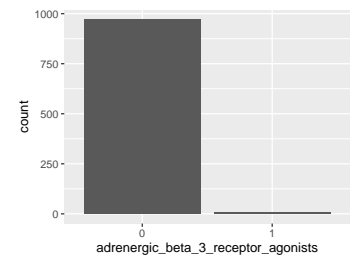
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic_beta_3_receptor_agonists

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

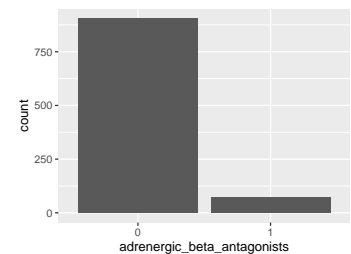
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic_beta_antagonists

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

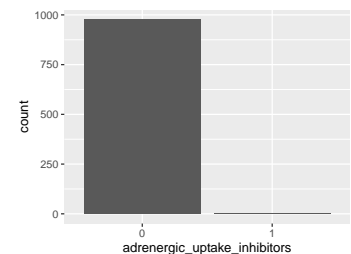
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

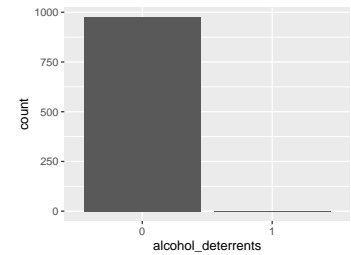


- Note that the following levels have at most five observations: "1".

alcohol_deterrents

- 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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

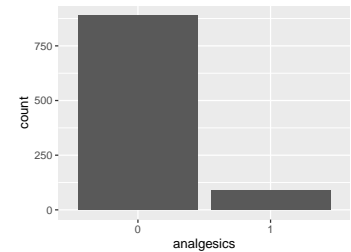


- Note that the following levels have at most five observations: "1".

analgesics

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

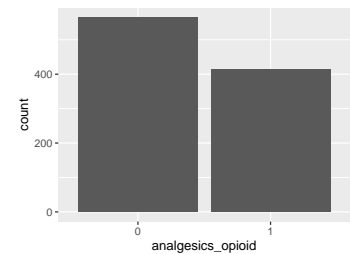
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



analgesics__opioid

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

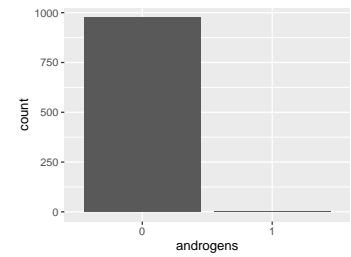
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

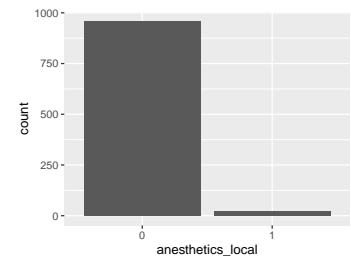


- Note that the following levels have at most five observations: "1".

anesthetics_local

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

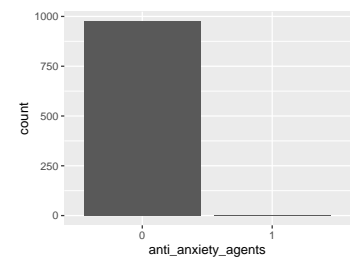
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

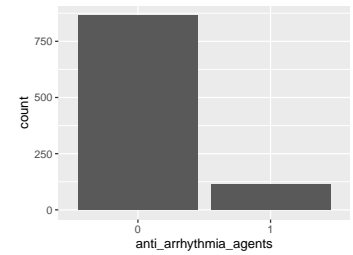


- Note that the following levels have at most five observations: "1".

anti_arrhythmia_agents

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

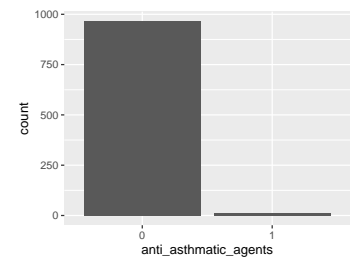
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti__asthmatic__agents

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

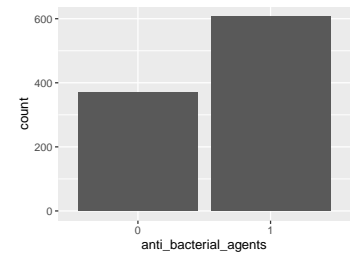
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti__bacterial__agents

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

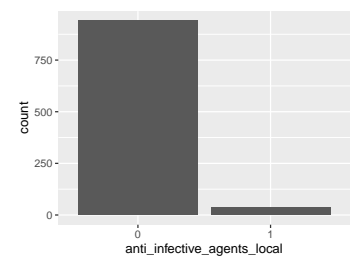
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"1"
Reference category	0



anti__infective__agents__local

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

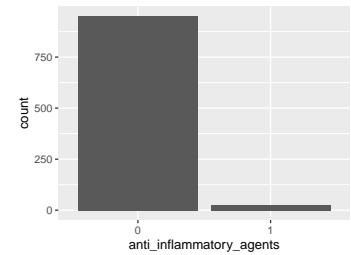
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents

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

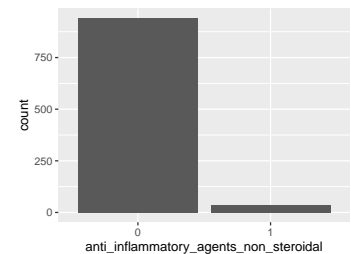
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents_non_steroidal

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

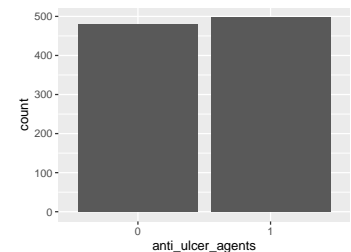
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_ulcer_agents

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

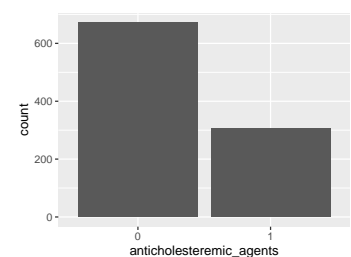
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"1"
Reference category	0



anticholesteremic_agents

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

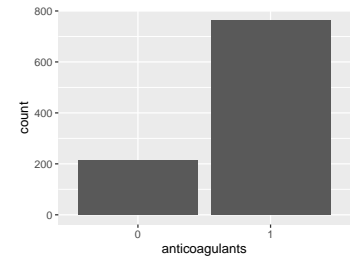
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



anticoagulants

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

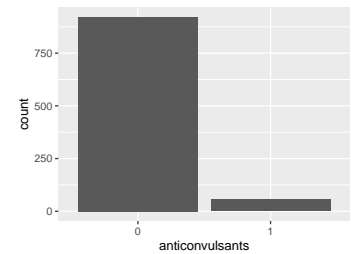
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"1"
Reference category	0



anticonvulsants

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

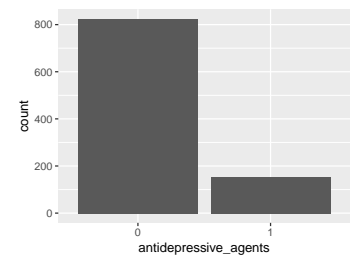
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antidepressive_agents

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

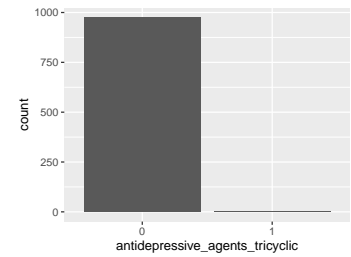
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

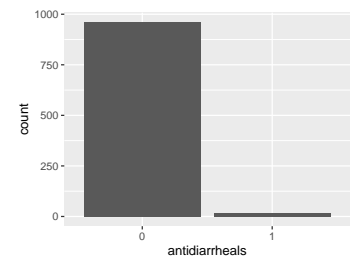


- Note that the following levels have at most five observations: "1".

antidiarrheals

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

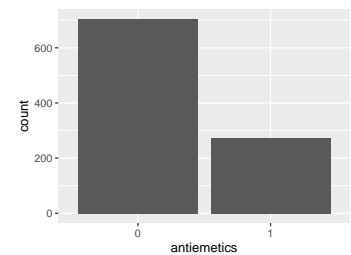
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antiemetics

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

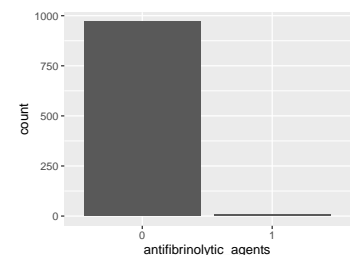
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antifibrinolytic_agents

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

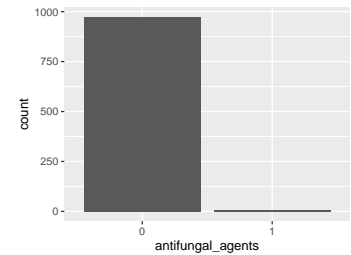
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

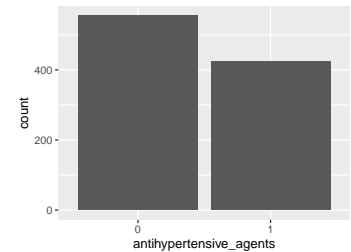


- Note that the following levels have at most five observations: "1".

antihypertensive_agents

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

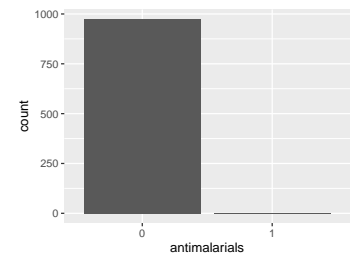
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

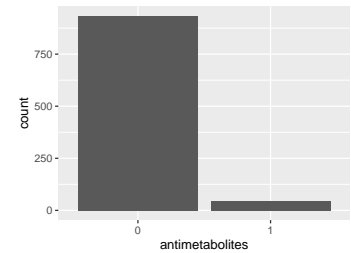


- Note that the following levels have at most five observations: "1".

antimetabolites

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

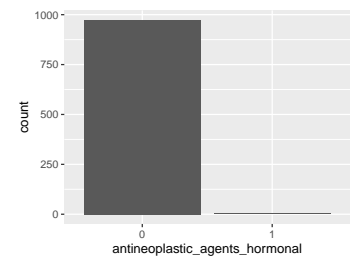
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

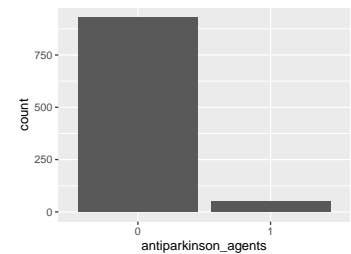


- Note that the following levels have at most five observations: "1".

antiparkinson_agents

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

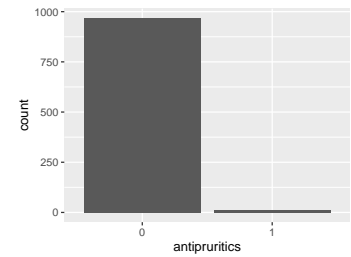
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antipruritics

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

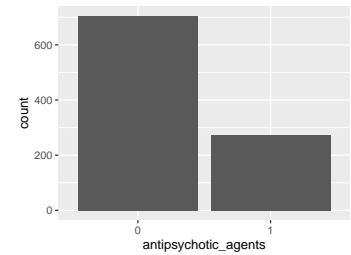
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antipsychotic_agents

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

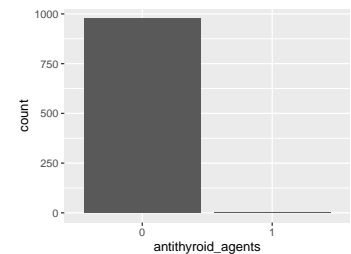
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

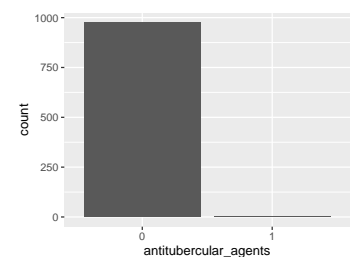


- Note that the following levels have at most five observations: "1".

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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

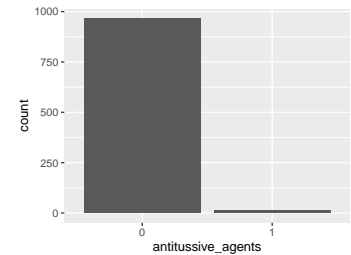


- Note that the following levels have at most five observations: "1".

antitussive__agents

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

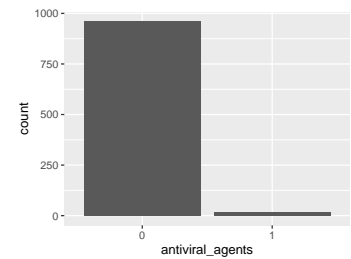
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



antiviral__agents

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

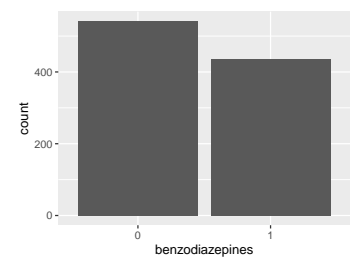
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



benzodiazepines

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

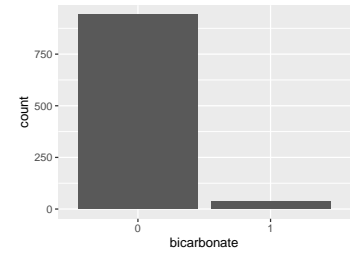
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



bicarbonate

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

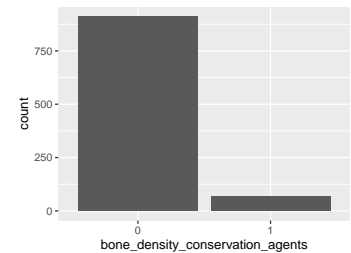
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



bone_density_conservation_agents

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

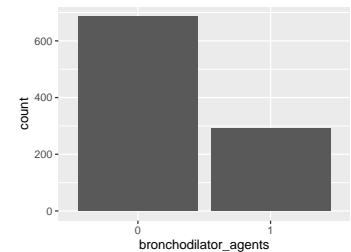
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



bronchodilator_agents

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

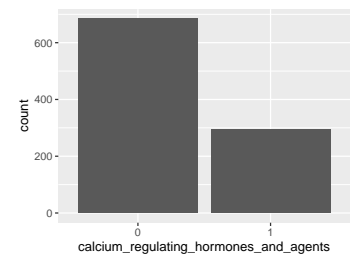
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



calcium_regulating_hormones_and_agents

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

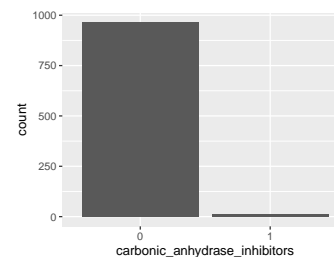
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



carbonic_anhydrase_inhibitors

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

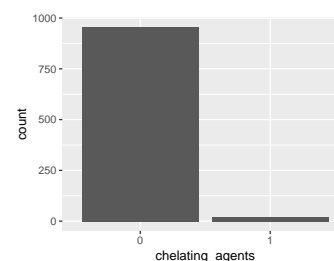
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



chelating_agents

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

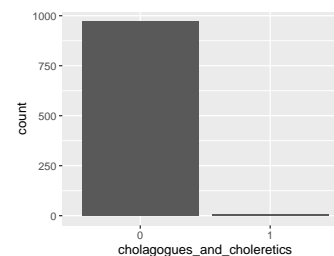
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



cholagogues_and_choleretics

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

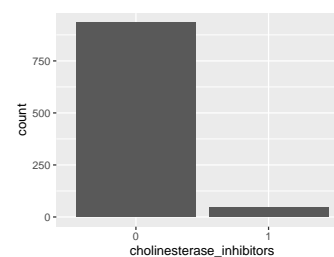
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



cholinesterase_inhibitors

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

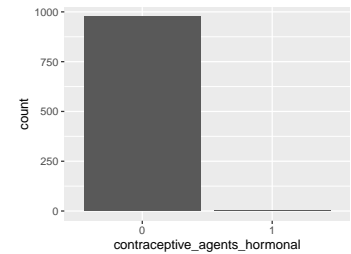
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

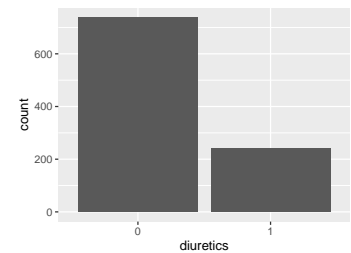


- Note that the following levels have at most five observations: "1".
-

diuretics

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

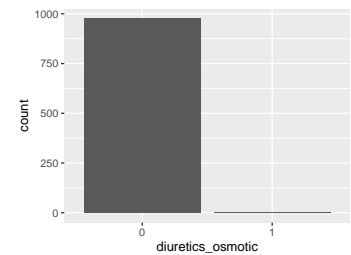
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



diuretics_osmotic

- 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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

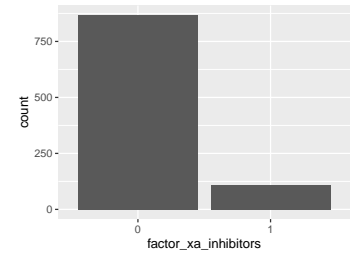


- Note that the following levels have at most five observations: "1".
-

factor__xa__inhibitors

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

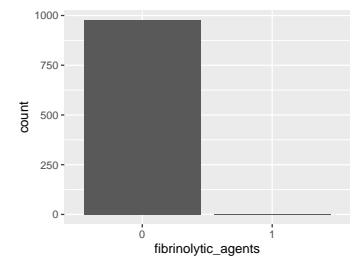
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



fibrinolytic__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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

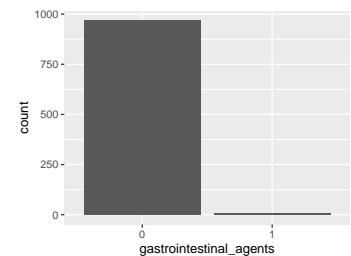


- Note that the following levels have at most five observations: "1".

gastrointestinal__agents

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

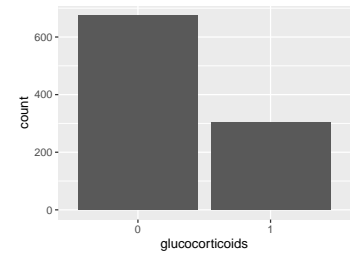
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



glucocorticoids

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

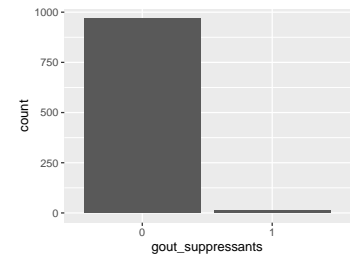
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



gout_suppressants

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

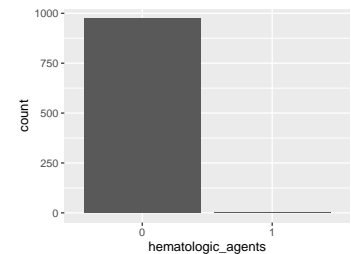
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

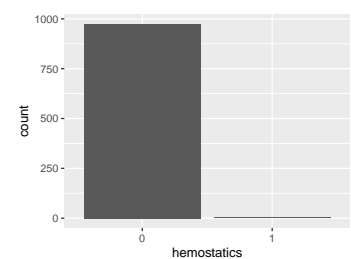


- Note that the following levels have at most five observations: "1".

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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

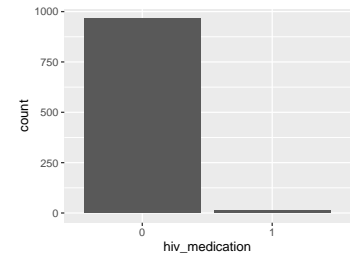


- Note that the following levels have at most five observations: "1".

hiv_medication

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

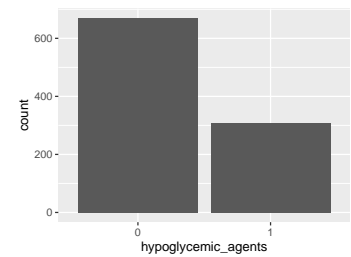
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



hypoglycemic_agents

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

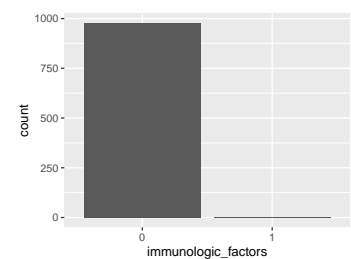
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



immunologic_factors

- 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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

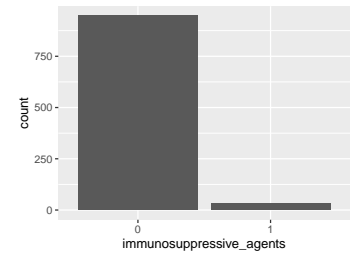


- Note that the following levels have at most five observations: "1".

immunosuppressive_agents

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

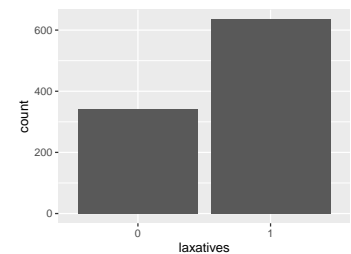
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



laxatives

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

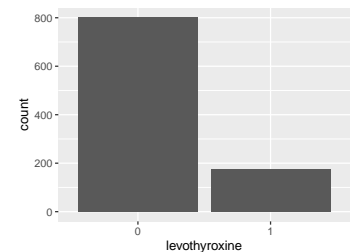
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"1"
Reference category	0



levothyroxine

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

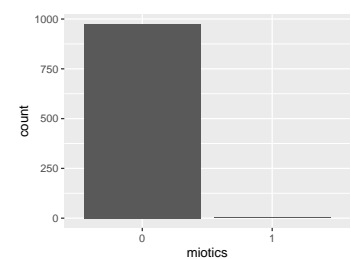
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



miotics

- 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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

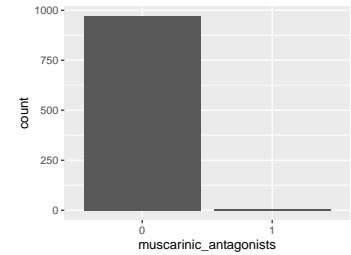


- Note that the following levels have at most five observations: "1".

muscarinic_antagonists

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

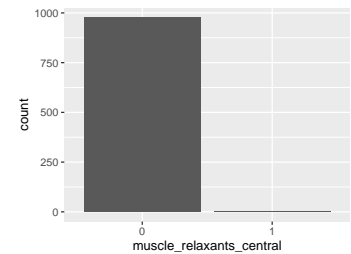
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

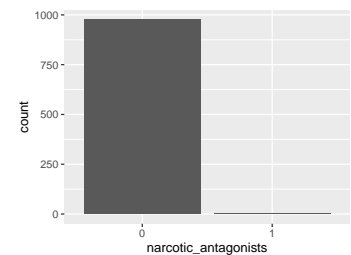


- Note that the following levels have at most five observations: "1".

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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

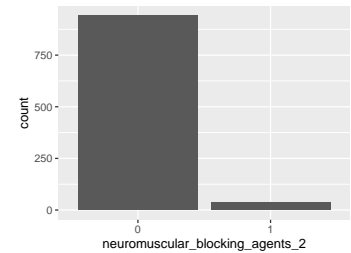


- Note that the following levels have at most five observations: "1".

neuromuscular_blocking_agents_2

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

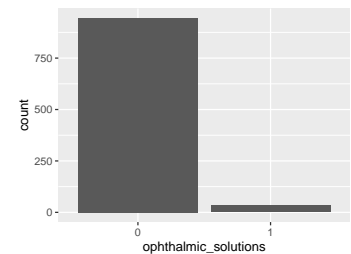
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



ophthalmic_solutions

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

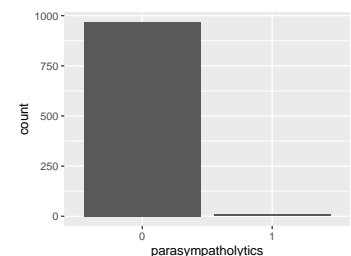
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



parasympatholytics

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

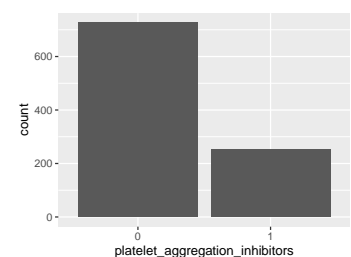
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



platelet_aggregation_inhibitors

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

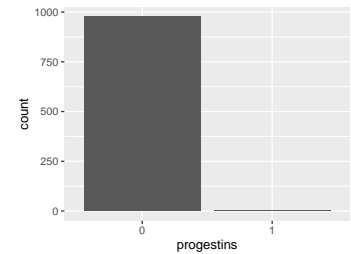
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

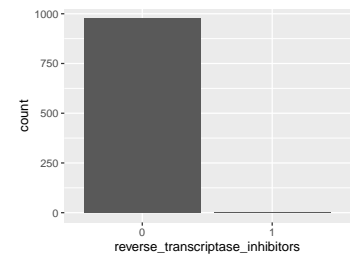


- Note that the following levels have at most five observations: "1".
-

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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

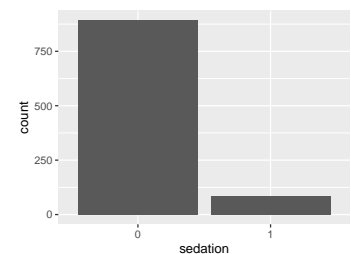


- Note that the following levels have at most five observations: "1".
-

sedation

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

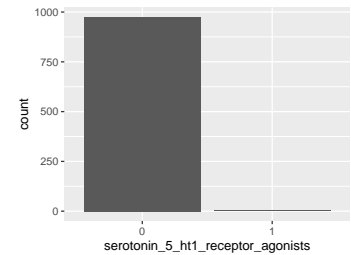
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0

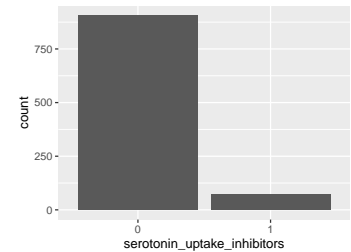


- Note that the following levels have at most five observations: "1".

serotonin_uptake_inhibitors

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

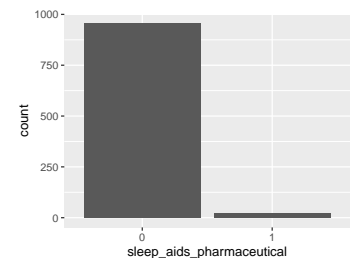
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



sleep_aids_pharmaceutical

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

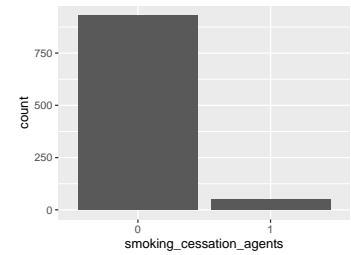
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



smoking_cessation_agents

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

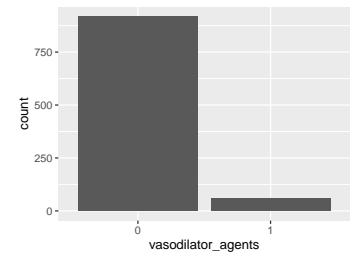
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasodilator__agents

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

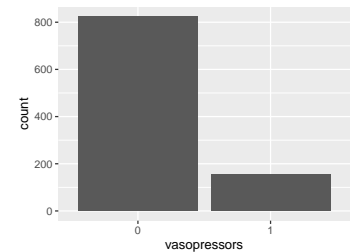
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasopressors

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

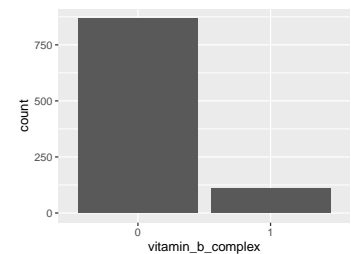
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamin_b_complex

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

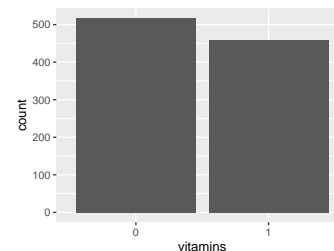
Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamins

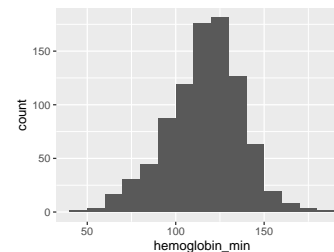
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	84 (7.92 %)
Number of unique values	2
Mode	"0"
Reference category	0



hemoglobin_min

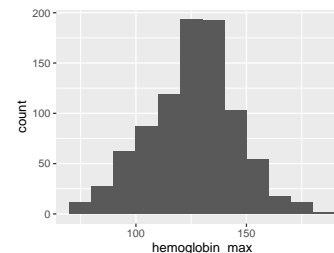
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	106
Median	119
1st and 3rd quartiles	104; 131
Min. and max.	41; 185



- Note that the following possible outlier values were detected: "41", "157", "158", "159", "161", "162", "163", "165", "170", "172" (3 additional values omitted).

hemoglobin_max

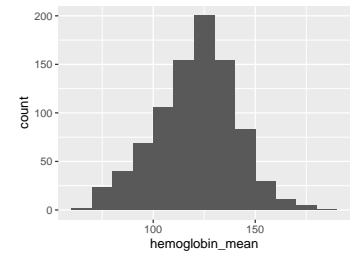
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	98
Median	128
1st and 3rd quartiles	113; 139
Min. and max.	71; 185



- Note that the following possible outlier values were detected: "165", "167", "169", "170", "171", "172", "173", "176", "178", "185".

hemoglobin_mean

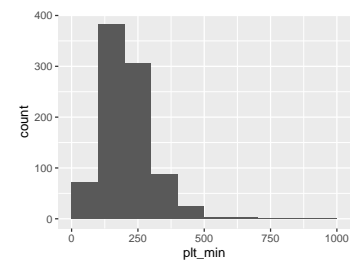
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	360
Median	122.67
1st and 3rd quartiles	108.75; 134
Min. and max.	61.71; 185



- Note that the following possible outlier values were detected: "161", "161.33", "162", "162.5", "162.75", "163", "168.33", "170.33", "171.5", "172" (3 additional values omitted).

plt_min

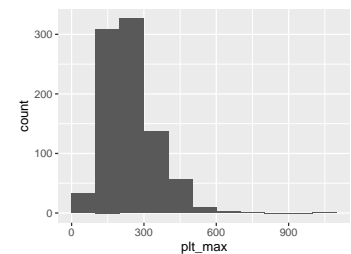
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	315
Median	196
1st and 3rd quartiles	144; 260
Min. and max.	21; 941



- Note that the following possible outlier values were detected: "21", "24", "26", "27", "519", "526", "609", "665", "941".

plt_max

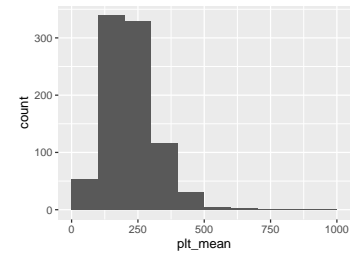
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	341
Median	228
1st and 3rd quartiles	168; 297
Min. and max.	24; 1052



- Note that the following possible outlier values were detected: "24", "26", "37", "44", "602", "608", "618", "713", "1052".

plt_mean

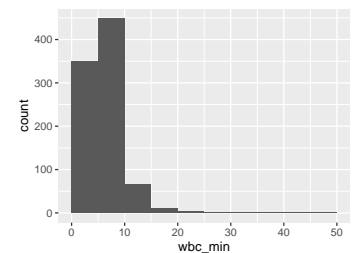
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	598
Median	209.67
1st and 3rd quartiles	158; 279
Min. and max.	23.86; 989.75



- Note that the following possible outlier values were detected: "23.86", "24", "33.75", "40.25", "43.5", "44.6", "47.33", "47.75", "48.4", "50" (6 additional values omitted).

wbc_min

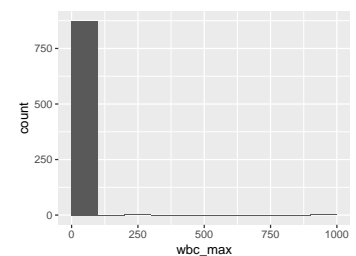
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	132
Median	5.6
1st and 3rd quartiles	4.1; 7.8
Min. and max.	0; 46.1



- Note that the following possible outlier values were detected: "0", "0.7", "1", "1.2", "1.3", "1.5", "1.6", "18.8", "20.1", "21.2" (3 additional values omitted).

wbc_max

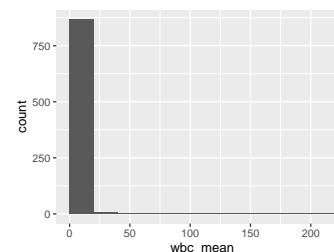
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	186
Median	7.5
1st and 3rd quartiles	5.4; 10.6
Min. and max.	1.3; 1000



- Note that the following possible outlier values were detected: "1.3", "1.6", "1.7", "1.8", "2.2", "2.4", "2.5", "2.6", "28.8", "29.7" (7 additional values omitted).

wbc__mean

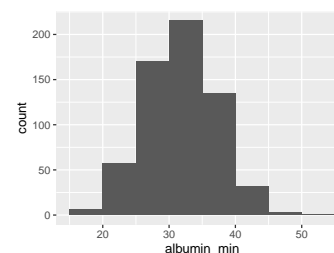
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	401
Median	6.63
1st and 3rd quartiles	4.9; 9.07
Min. and max.	1.3; 207.14



- Note that the following possible outlier values were detected: "1.3", "1.43", "1.6", "1.7", "1.97", "2.1", "2.2", "22.2", "22.55", "23.3" (6 additional values omitted).

albumin__min

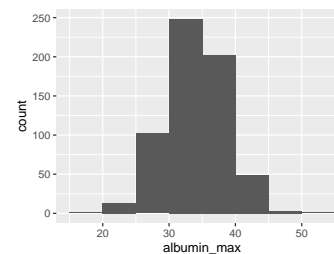
Feature	Result
Variable type	numeric
Number of missing obs.	442 (41.66 %)
Number of unique values	36
Median	33
1st and 3rd quartiles	28; 36
Min. and max.	17; 54



- Note that the following possible outlier values were detected: "44", "45", "45.5", "46.4", "48", "54".

albumin__max

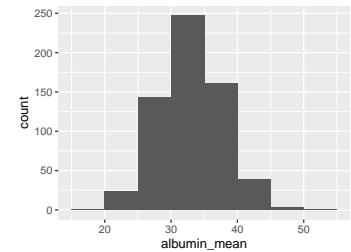
Feature	Result
Variable type	numeric
Number of missing obs.	442 (41.66 %)
Number of unique values	31
Median	35
1st and 3rd quartiles	32; 38
Min. and max.	18; 54



- Note that the following possible outlier values were detected: "18", "21", "22", "48", "54".

albumin__mean

Feature	Result
Variable type	numeric
Number of missing obs.	442 (41.66 %)
Number of unique values	134
Median	33.5
1st and 3rd quartiles	30; 36.67
Min. and max.	18; 54

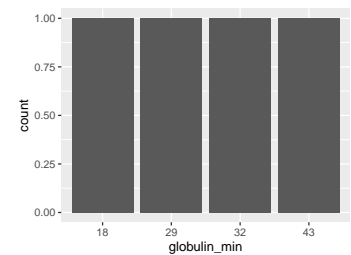


- Note that the following possible outlier values were detected: "18", "46.4", "48", "54".

globulin__min

- 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.	1057 (99.62 %)
Number of unique values	4
Mode	"18"
Reference category	18

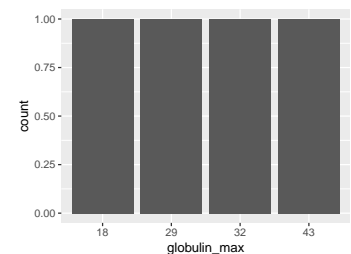


- Note that the following levels have at most five observations: "18", "29", "32", "43".

globulin__max

- 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.	1057 (99.62 %)
Number of unique values	4
Mode	"18"
Reference category	18

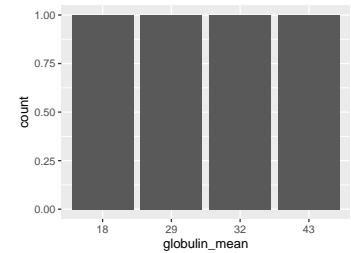


- Note that the following levels have at most five observations: "18", "29", "32", "43".

globulin__mean

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

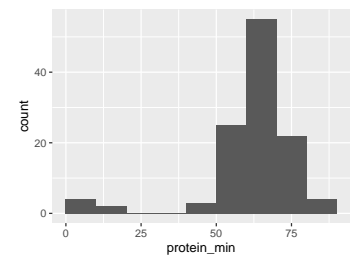
Feature	Result
Variable type	numeric
Number of missing obs.	1057 (99.62 %)
Number of unique values	4
Mode	"18"
Reference category	18



- Note that the following levels have at most five observations: "18", "29", "32", "43".

protein__min

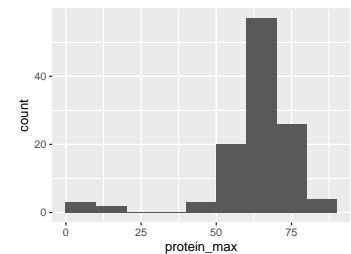
Feature	Result
Variable type	numeric
Number of missing obs.	946 (89.16 %)
Number of unique values	35
Median	65
1st and 3rd quartiles	59; 70
Min. and max.	0.47; 86



- Note that the following possible outlier values were detected: "0.47", "0.52", "0.62", "20", "84", "86".

protein__max

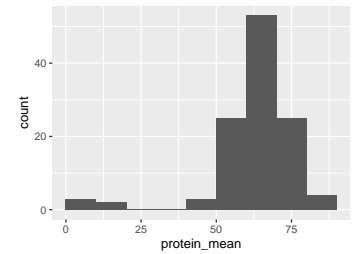
Feature	Result
Variable type	numeric
Number of missing obs.	946 (89.16 %)
Number of unique values	33
Median	65
1st and 3rd quartiles	61; 71
Min. and max.	0.47; 86



- Note that the following possible outlier values were detected: "0.47", "0.62", "20".

protein__mean

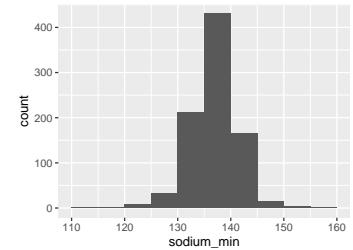
Feature	Result
Variable type	numeric
Number of missing obs.	946 (89.16 %)
Number of unique values	43
Median	65
1st and 3rd quartiles	60; 70.34
Min. and max.	0.47; 86



- Note that the following possible outlier values were detected: "0.47", "0.62", "20", "86".

sodium__min

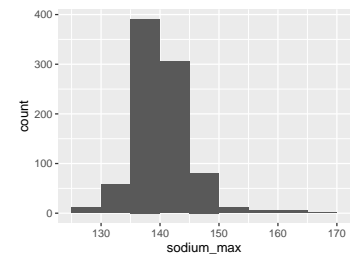
Feature	Result
Variable type	numeric
Number of missing obs.	191 (18 %)
Number of unique values	38
Median	138
1st and 3rd quartiles	135; 140
Min. and max.	114; 157



- Note that the following possible outlier values were detected: "114", "117", "120", "121", "122", "123", "124", "145", "146", "147" (8 additional values omitted).

sodium__max

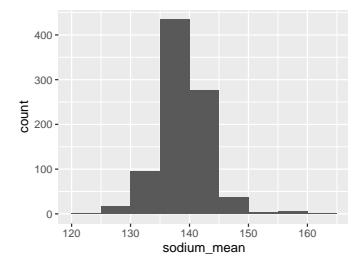
Feature	Result
Variable type	numeric
Number of missing obs.	191 (18 %)
Number of unique values	36
Median	140
1st and 3rd quartiles	138; 143
Min. and max.	127; 168



- Note that the following possible outlier values were detected: "127", "128", "129", "130", "131", "132", "133", "134", "158", "159" (4 additional values omitted).

sodium__mean

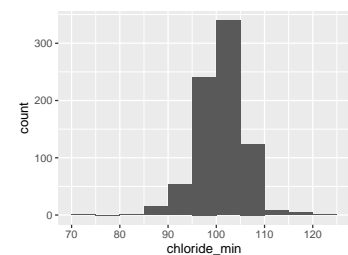
Feature	Result
Variable type	numeric
Number of missing obs.	191 (18 %)
Number of unique values	179
Median	139
1st and 3rd quartiles	136.8; 141.5
Min. and max.	120.13; 160.86



- Note that the following possible outlier values were detected: "120.13", "124.75", "125.67", "125.75", "125.86", "126.25", "126.5", "127", "127.56", "128" (12 additional values omitted).

chloride__min

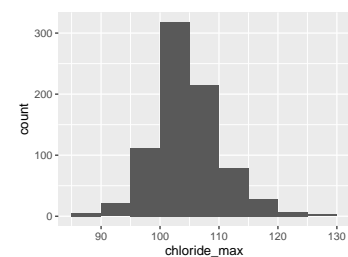
Feature	Result
Variable type	numeric
Number of missing obs.	273 (25.73 %)
Number of unique values	35
Median	102
1st and 3rd quartiles	99; 104
Min. and max.	74; 121



- Note that the following possible outlier values were detected: "74", "84", "86", "87", "88", "89", "90", "91", "112", "113" (5 additional values omitted).

chloride__max

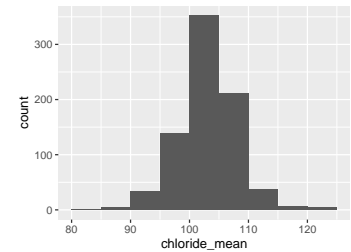
Feature	Result
Variable type	numeric
Number of missing obs.	273 (25.73 %)
Number of unique values	41
Median	104
1st and 3rd quartiles	102; 108
Min. and max.	86; 127



- Note that the following possible outlier values were detected: "86", "88", "89", "90", "91", "92", "93", "94", "95", "96" (6 additional values omitted).

chloride__mean

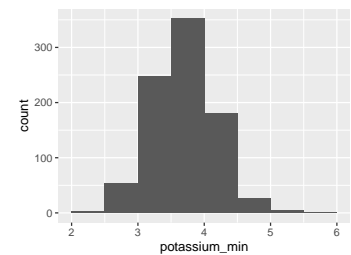
Feature	Result
Variable type	numeric
Number of missing obs.	273 (25.73 %)
Number of unique values	179
Median	103.33
1st and 3rd quartiles	100.8; 106
Min. and max.	82.75; 122.8



- Note that the following possible outlier values were detected: "82.75", "86", "88", "89.5", "89.67", "90", "90.67", "91", "91.5", "92" (16 additional values omitted).

potassium__min

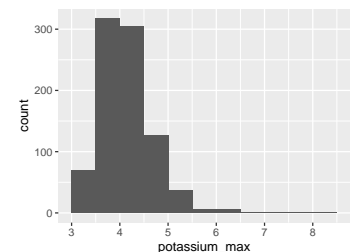
Feature	Result
Variable type	numeric
Number of missing obs.	193 (18.19 %)
Number of unique values	32
Median	3.7
1st and 3rd quartiles	3.4; 4
Min. and max.	2.4; 5.6



- Note that the following possible outlier values were detected: "2.4", "4.9", "5", "5.1", "5.2", "5.3", "5.4", "5.6".

potassium__max

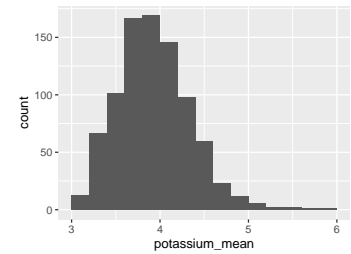
Feature	Result
Variable type	numeric
Number of missing obs.	193 (18.19 %)
Number of unique values	34
Median	4.1
1st and 3rd quartiles	3.8; 4.5
Min. and max.	3; 8.2



- Note that the following possible outlier values were detected: "3", "3.2", "6.2", "6.3", "6.4", "6.9", "8.2".

potassium__mean

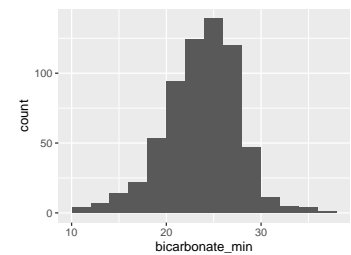
Feature	Result
Variable type	numeric
Number of missing obs.	193 (18.19 %)
Number of unique values	135
Median	3.93
1st and 3rd quartiles	3.68; 4.2
Min. and max.	3; 5.81



- Note that the following possible outlier values were detected: "3", "3.05", "5.17", "5.19", "5.2", "5.24", "5.35", "5.47", "5.6", "5.7" (1 additional values omitted).

bicarbonate__min

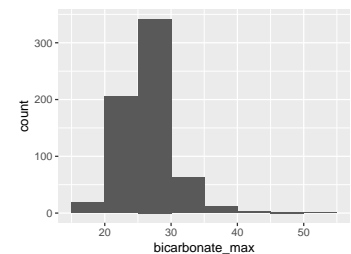
Feature	Result
Variable type	numeric
Number of missing obs.	416 (39.21 %)
Number of unique values	153
Median	24.1
1st and 3rd quartiles	21.5; 26.6
Min. and max.	10.4; 37.2



- Note that the following possible outlier values were detected: "10.4", "10.8", "11.6", "12.4", "12.5", "12.8", "13.1", "34", "34.3", "34.4" (2 additional values omitted).

bicarbonate__max

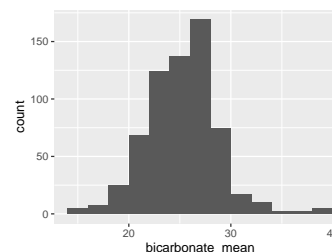
Feature	Result
Variable type	numeric
Number of missing obs.	416 (39.21 %)
Number of unique values	136
Median	26.6
1st and 3rd quartiles	24.1; 28.5
Min. and max.	15; 50.9



- Note that the following possible outlier values were detected: "15", "16", "33.8", "34", "34.3", "34.4", "35.7", "36", "36.1", "37.2" (9 additional values omitted).

bicarbonate__mean

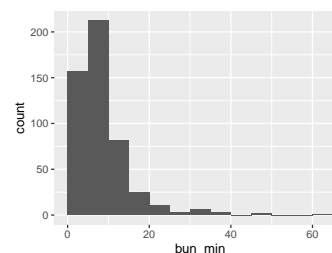
Feature	Result
Variable type	numeric
Number of missing obs.	416 (39.21 %)
Number of unique values	332
Median	25.61
1st and 3rd quartiles	23.1; 27.47
Min. and max.	14; 39.55



- Note that the following possible outlier values were detected: "30.97", "31", "31.3", "31.9", "32.1", "32.4", "32.7", "32.9", "33", "33.03" (10 additional values omitted).

bun__min

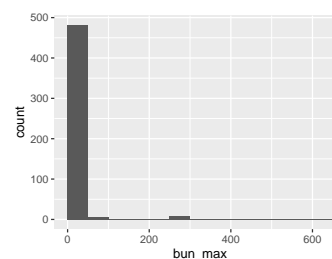
Feature	Result
Variable type	numeric
Number of missing obs.	561 (52.87 %)
Number of unique values	161
Median	6.6
1st and 3rd quartiles	4.5; 10.2
Min. and max.	0.7; 63.5



- Note that the following possible outlier values were detected: "0.7", "0.8", "1.1", "1.2", "1.4", "1.6", "1.7", "1.8", "1.9", "2" (6 additional values omitted).

bun__max

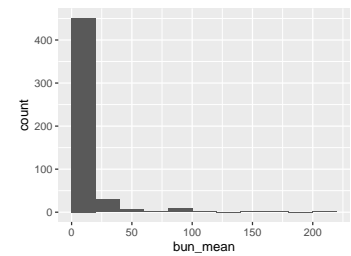
Feature	Result
Variable type	numeric
Number of missing obs.	561 (52.87 %)
Number of unique values	186
Median	7.95
1st and 3rd quartiles	5.38; 13.3
Min. and max.	0.7; 619



- Note that the following possible outlier values were detected: "0.7", "0.8", "1.2", "1.8", "1.9", "2", "2.2", "2.4", "2.5", "2.6" (15 additional values omitted).

bun__mean

Feature	Result
Variable type	numeric
Number of missing obs.	561 (52.87 %)
Number of unique values	272
Median	7.33
1st and 3rd quartiles	4.92; 11.62
Min. and max.	0.7; 211.8



- Note that the following possible outlier values were detected: "0.7", "0.8", "1.2", "1.6", "1.77", "1.89", "1.9", "2", "2.2", "2.25" (23 additional values omitted).

calcium__min

- The variable only takes one value: "NA".

calcium__max

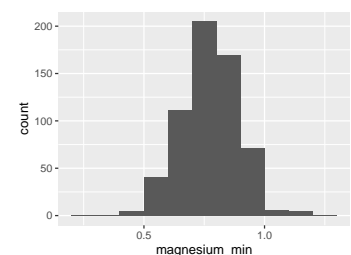
- The variable only takes one value: "NA".

calcium__mean

- The variable only takes one value: "NA".

magnesium__min

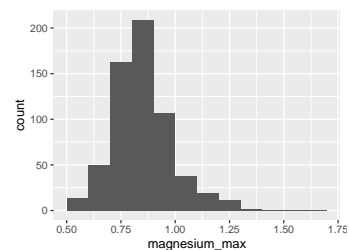
Feature	Result
Variable type	numeric
Number of missing obs.	446 (42.04 %)
Number of unique values	66
Median	0.78
1st and 3rd quartiles	0.7; 0.85
Min. and max.	0.23; 1.21



- Note that the following possible outlier values were detected: "0.23", "0.42", "0.43", "0.46", "0.47", "1.08", "1.12", "1.13", "1.16", "1.19" (1 additional values omitted).

magnesium_max

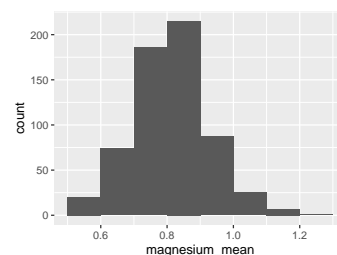
Feature	Result
Variable type	numeric
Number of missing obs.	446 (42.04 %)
Number of unique values	75
Median	0.84
1st and 3rd quartiles	0.76; 0.92
Min. and max.	0.52; 1.67



- Note that the following possible outlier values were detected: "0.52", "0.55", "0.56", "0.57", "1.27", "1.28", "1.29", "1.3", "1.38", "1.4" (1 additional values omitted).

magnesium_mean

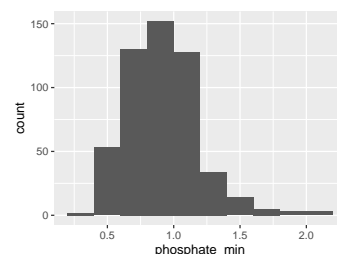
Feature	Result
Variable type	numeric
Number of missing obs.	446 (42.04 %)
Number of unique values	64
Median	0.81
1st and 3rd quartiles	0.74; 0.88
Min. and max.	0.51; 1.3



- Note that the following possible outlier values were detected: "0.51", "0.52", "1.1", "1.12", "1.13", "1.16", "1.17", "1.18", "1.19", "1.3".

phosphate_min

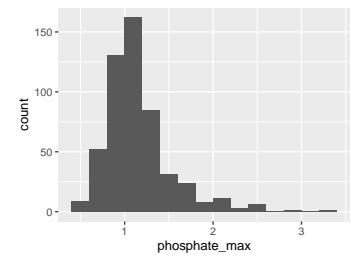
Feature	Result
Variable type	numeric
Number of missing obs.	537 (50.61 %)
Number of unique values	112
Median	0.91
1st and 3rd quartiles	0.74; 1.07
Min. and max.	0.32; 2.06



- Note that the following possible outlier values were detected: "1.52", "1.56", "1.67", "1.7", "1.73", "1.78", "1.82", "1.83", "1.88", "2.01" (1 additional values omitted).

phosphate_max

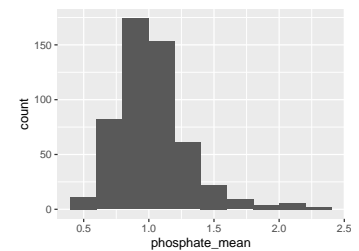
Feature	Result
Variable type	numeric
Number of missing obs.	537 (50.61 %)
Number of unique values	138
Median	1.08
1st and 3rd quartiles	0.92; 1.3
Min. and max.	0.44; 3.27



- Note that the following possible outlier values were detected: "0.44", "0.47", "0.51", "0.52", "0.56", "0.57", "0.58", "0.59", "0.61", "0.63" (9 additional values omitted).

phosphate_mean

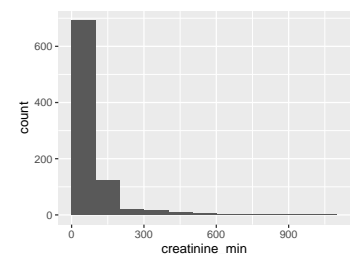
Feature	Result
Variable type	numeric
Number of missing obs.	537 (50.61 %)
Number of unique values	115
Median	1
1st and 3rd quartiles	0.84; 1.16
Min. and max.	0.44; 2.33



- Note that the following possible outlier values were detected: "0.44", "1.74", "1.78", "1.8", "1.82", "1.83", "1.88", "1.95", "2.01", "2.06" (4 additional values omitted).

creatinine_min

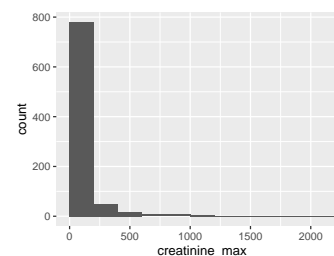
Feature	Result
Variable type	numeric
Number of missing obs.	192 (18.1 %)
Number of unique values	179
Median	69
1st and 3rd quartiles	55; 94
Min. and max.	20; 1049



- Note that the following possible outlier values were detected: "20", "23", "24", "25", "26", "27", "29", "31", "32", "33" (45 additional values omitted).

creatinine__max

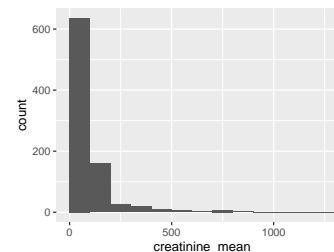
Feature	Result
Variable type	numeric
Number of missing obs.	192 (18.1 %)
Number of unique values	205
Median	81
1st and 3rd quartiles	63; 113
Min. and max.	25; 2094



- Note that the following possible outlier values were detected: "25", "27", "29", "31", "32", "34", "35", "37", "38", "39" (47 additional values omitted).

creatinine__mean

Feature	Result
Variable type	numeric
Number of missing obs.	192 (18.1 %)
Number of unique values	442
Median	74.25
1st and 3rd quartiles	58.5; 103.33
Min. and max.	22.5; 1209.7



- Note that the following possible outlier values were detected: "22.5", "26.33", "28.75", "29", "29.5", "31", "31.75", "35", "35.67", "36" (61 additional values omitted).

gfr__min

- The variable only takes one value: "NA".

gfr__max

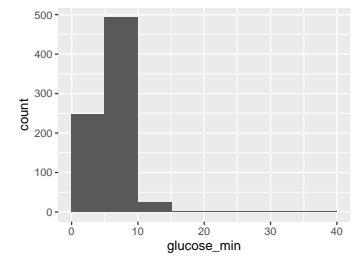
- The variable only takes one value: "NA".

gfr__mean

- The variable only takes one value: "NA".

glucose__min

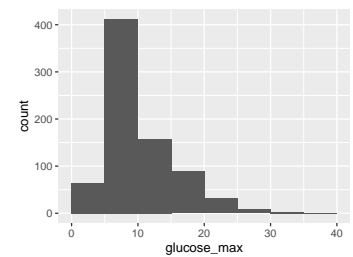
Feature	Result
Variable type	numeric
Number of missing obs.	295 (27.8 %)
Number of unique values	98
Median	5.5
1st and 3rd quartiles	4.8; 6.6
Min. and max.	1.2; 35.8



- Note that the following possible outlier values were detected: "1.2", "1.9", "2.1", "2.2", "2.3", "2.4", "2.5", "2.7", "2.8", "2.9" (20 additional values omitted).

glucose__max

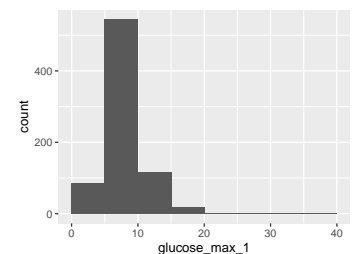
Feature	Result
Variable type	numeric
Number of missing obs.	295 (27.8 %)
Number of unique values	177
Median	8.4
1st and 3rd quartiles	6.1; 12.5
Min. and max.	3.7; 35.8



- Note that the following possible outlier values were detected: "3.7", "3.9".

glucose__max__1

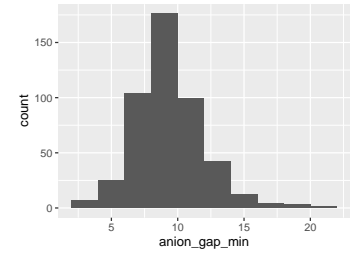
Feature	Result
Variable type	numeric
Number of missing obs.	295 (27.8 %)
Number of unique values	389
Median	6.85
1st and 3rd quartiles	5.63; 8.94
Min. and max.	3.7; 35.8



- Note that the following possible outlier values were detected: "3.7", "3.9", "3.95", "4", "4.1", "4.2", "35.8".

anion_gap_min

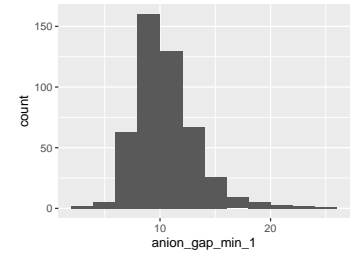
Feature	Result
Variable type	numeric
Number of missing obs.	588 (55.42 %)
Number of unique values	18
Median	10
1st and 3rd quartiles	8; 11
Min. and max.	2; 22



- Note that the following possible outlier values were detected: "2", "16", "18", "19", "20", "22".

anion_gap_min_1

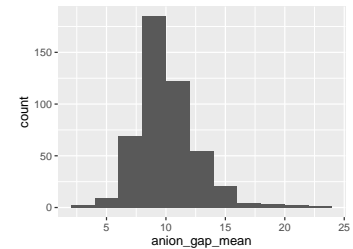
Feature	Result
Variable type	numeric
Number of missing obs.	588 (55.42 %)
Number of unique values	22
Median	11
1st and 3rd quartiles	9; 12
Min. and max.	2; 25



- Note that the following possible outlier values were detected: "2", "4", "17", "18", "19", "20", "21", "22", "23", "25".

anion_gap_mean

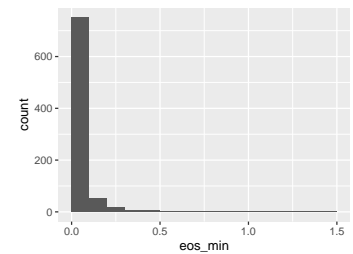
Feature	Result
Variable type	numeric
Number of missing obs.	588 (55.42 %)
Number of unique values	56
Median	10
1st and 3rd quartiles	9; 12
Min. and max.	2; 22.5



- Note that the following possible outlier values were detected: "2", "4", "5", "5.2", "5.67", "6", "6.4", "21.4", "21.5", "22.5".

eos__min

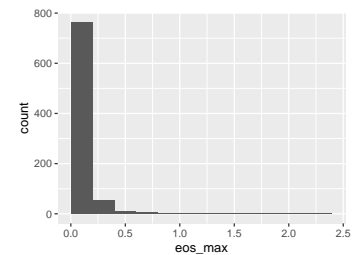
Feature	Result
Variable type	numeric
Number of missing obs.	224 (21.11 %)
Number of unique values	42
Median	0
1st and 3rd quartiles	0; 0.03
Min. and max.	0; 1.5



- Note that the following possible outlier values were detected: "1.5".

eos__max

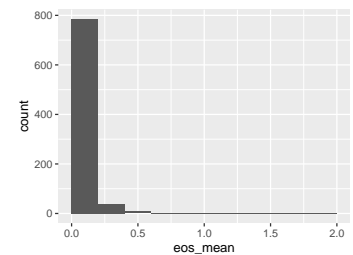
Feature	Result
Variable type	numeric
Number of missing obs.	224 (21.11 %)
Number of unique values	52
Median	0.01
1st and 3rd quartiles	0; 0.1
Min. and max.	0; 2.22



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

eos__mean

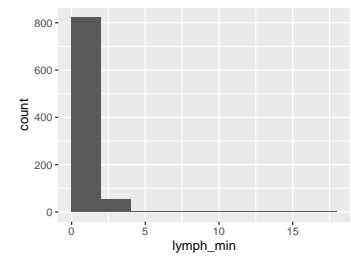
Feature	Result
Variable type	numeric
Number of missing obs.	224 (21.11 %)
Number of unique values	45
Median	0.01
1st and 3rd quartiles	0; 0.06
Min. and max.	0; 1.85



- Note that the following possible outlier values were detected: "0.97", "1.85".

lymph_min

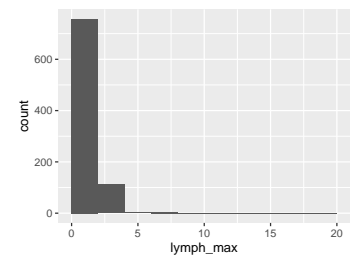
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	216
Median	0.8
1st and 3rd quartiles	0.51; 1.22
Min. and max.	0; 16.9



- Note that the following possible outlier values were detected: "0", "0.08", "3.66", "3.69", "5.85", "6", "7.3", "12.8", "16.9".

lymph_max

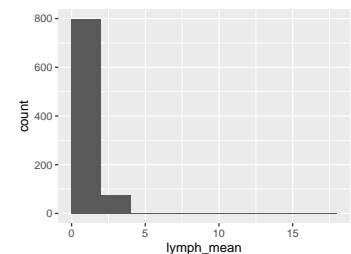
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	237
Median	1.15
1st and 3rd quartiles	0.84; 1.65
Min. and max.	0.16; 18.6



- Note that the following possible outlier values were detected: "0.16", "0.18", "0.24", "0.25", "0.28", "0.29", "0.3", "0.33", "0.34", "0.35" (9 additional values omitted).

lymph_mean

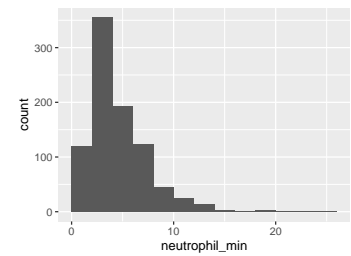
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	229
Median	0.97
1st and 3rd quartiles	0.7; 1.4
Min. and max.	0.11; 17.75



- Note that the following possible outlier values were detected: "0.11", "0.15", "0.16", "0.18", "0.19", "0.24", "0.25", "0.27", "0.28", "0.29" (11 additional values omitted).

neutrophil_min

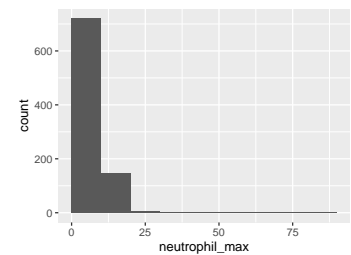
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	440
Median	3.9
1st and 3rd quartiles	2.6; 5.92
Min. and max.	0.09; 24.69



- Note that the following possible outlier values were detected: "0.09", "0.17", "0.35", "0.37", "0.39", "0.5", "0.59", "0.6", "0.66", "0.67" (5 additional values omitted).

neutrophil_max

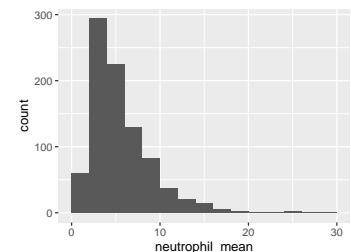
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	505
Median	5.4
1st and 3rd quartiles	3.6; 8.51
Min. and max.	0.64; 83



- Note that the following possible outlier values were detected: "0.64", "0.68", "0.8", "0.94", "0.95", "0.96", "1", "1.08", "1.1", "1.15" (10 additional values omitted).

neutrophil_mean

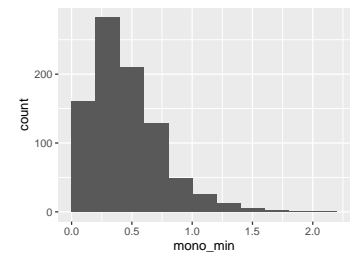
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	527
Median	4.72
1st and 3rd quartiles	3.2; 7.06
Min. and max.	0.47; 28.48



- Note that the following possible outlier values were detected: "0.47", "0.52", "0.61", "0.62", "0.8", "0.94", "0.96", "0.98", "1", "1.08" (5 additional values omitted).

mono__min

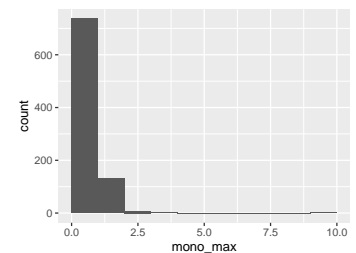
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	127
Median	0.4
1st and 3rd quartiles	0.28; 0.61
Min. and max.	0; 2.02



- Note that the following possible outlier values were detected: "0", "0.03", "0.04", "0.05", "0.06", "0.07", "0.08", "1.71", "1.72", "2.02".

mono__max

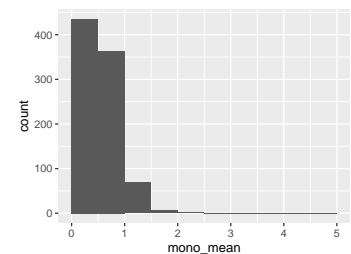
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	159
Median	0.6
1st and 3rd quartiles	0.4; 0.88
Min. and max.	0; 9.5



- Note that the following possible outlier values were detected: "0", "0.06", "0.08", "2.29", "2.49", "2.51", "2.61", "2.8", "3.39", "9.5".

mono__mean

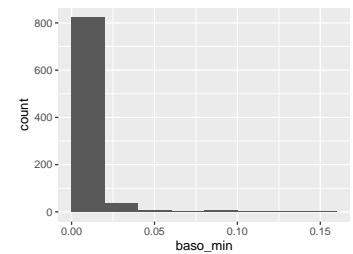
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	141
Median	0.51
1st and 3rd quartiles	0.38; 0.72
Min. and max.	0; 4.69



- Note that the following possible outlier values were detected: "0", "0.06", "0.08", "0.09", "0.1", "0.11", "0.12", "0.13", "0.14", "0.15" (8 additional values omitted).

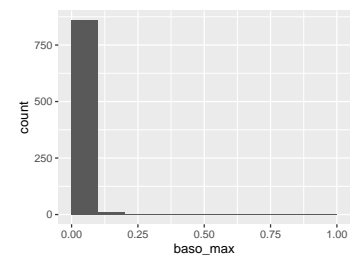
baso_min

Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	14
Median	0
1st and 3rd quartiles	0; 0.01
Min. and max.	0; 0.16



baso_max

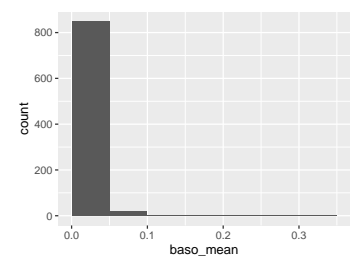
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	25
Median	0.01
1st and 3rd quartiles	0; 0.02
Min. and max.	0; 1



- Note that the following possible outlier values were detected: "0.11", "0.12", "0.14", "0.15", "0.16", "0.19", "0.23", "0.25", "0.27", "0.37" (4 additional values omitted).

baso_mean

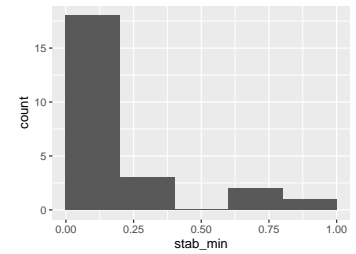
Feature	Result
Variable type	numeric
Number of missing obs.	184 (17.34 %)
Number of unique values	19
Median	0.01
1st and 3rd quartiles	0; 0.02
Min. and max.	0; 0.33



- Note that the following possible outlier values were detected: "0.06", "0.07", "0.08", "0.09", "0.1", "0.11", "0.13", "0.16", "0.21", "0.25" (3 additional values omitted).

stab_min

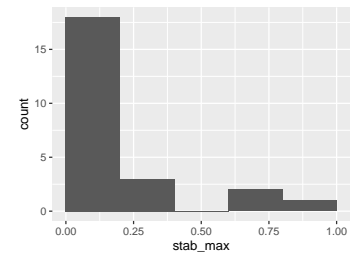
Feature	Result
Variable type	numeric
Number of missing obs.	1037 (97.74 %)
Number of unique values	16
Median	0.12
1st and 3rd quartiles	0.03; 0.2
Min. and max.	0; 0.85



- Note that the following possible outlier values were detected: "0.71", "0.73", "0.85".

stab_max

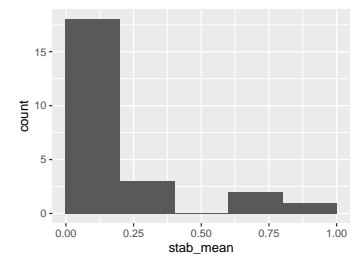
Feature	Result
Variable type	numeric
Number of missing obs.	1037 (97.74 %)
Number of unique values	17
Median	0.12
1st and 3rd quartiles	0.03; 0.21
Min. and max.	0; 0.85



- Note that the following possible outlier values were detected: "0.71", "0.73", "0.85".

stab_mean

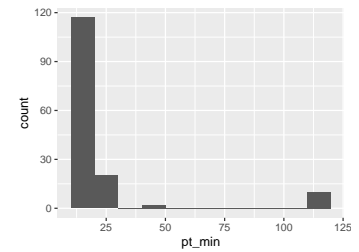
Feature	Result
Variable type	numeric
Number of missing obs.	1037 (97.74 %)
Number of unique values	16
Median	0.12
1st and 3rd quartiles	0.03; 0.2
Min. and max.	0; 0.85



- Note that the following possible outlier values were detected: "0.71", "0.73", "0.85".

pt__min

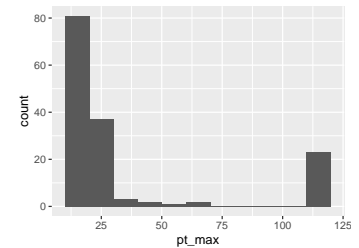
Feature	Result
Variable type	numeric
Number of missing obs.	912 (85.96 %)
Number of unique values	17
Median	18
1st and 3rd quartiles	17; 20
Min. and max.	13; 120



- Note that the following possible outlier values were detected: "13", "14", "15", "42", "45", "120".

pt__max

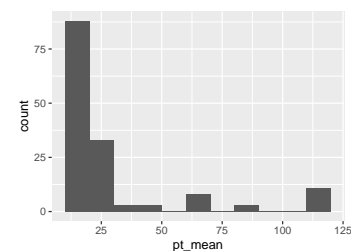
Feature	Result
Variable type	numeric
Number of missing obs.	912 (85.96 %)
Number of unique values	25
Median	20
1st and 3rd quartiles	18; 24
Min. and max.	14; 120



- Note that the following possible outlier values were detected: "14", "15", "16", "17", "118", "120".

pt__mean

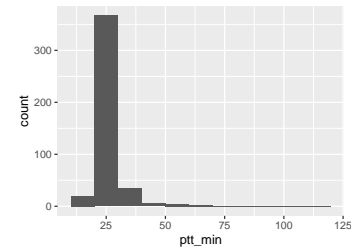
Feature	Result
Variable type	numeric
Number of missing obs.	912 (85.96 %)
Number of unique values	42
Median	19.67
1st and 3rd quartiles	18; 23
Min. and max.	14; 120



- Note that the following possible outlier values were detected: "14", "15", "16", "16.5", "62", "64.2", "67.5", "70", "82.25", "88.67" (2 additional values omitted).

ptt_min

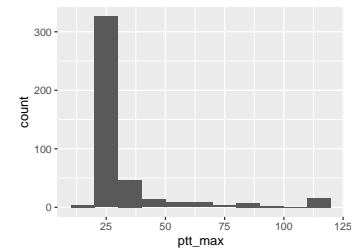
Feature	Result
Variable type	numeric
Number of missing obs.	627 (59.1 %)
Number of unique values	35
Median	25.5
1st and 3rd quartiles	23; 28
Min. and max.	15; 111



- Note that the following possible outlier values were detected: "15", "36", "37", "38", "39", "40", "41", "42", "43", "48" (8 additional values omitted).

ptt_max

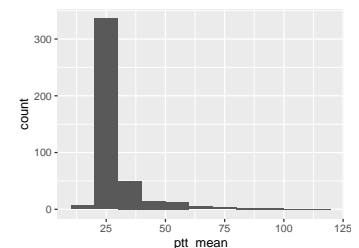
Feature	Result
Variable type	numeric
Number of missing obs.	627 (59.1 %)
Number of unique values	50
Median	26
1st and 3rd quartiles	24; 30
Min. and max.	19; 120



- Note that the following possible outlier values were detected: "19", "20", "21", "58", "62", "63", "64", "66", "67", "68" (9 additional values omitted).

ptt_mean

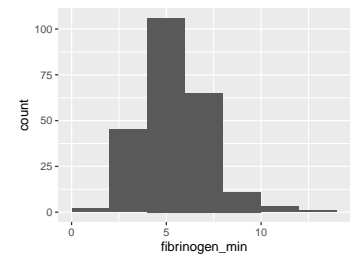
Feature	Result
Variable type	numeric
Number of missing obs.	627 (59.1 %)
Number of unique values	105
Median	26
1st and 3rd quartiles	24; 29.46
Min. and max.	19; 111



- Note that the following possible outlier values were detected: "19", "19.5", "19.75", "20", "20.25", "20.33", "20.5", "20.67", "21", "21.5" (21 additional values omitted).

fibrinogen_min

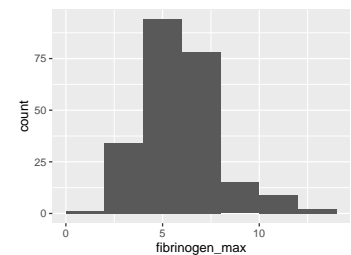
Feature	Result
Variable type	numeric
Number of missing obs.	828 (78.04 %)
Number of unique values	154
Median	5.6
1st and 3rd quartiles	4.34; 6.15
Min. and max.	1.17; 13.87



- Note that the following possible outlier values were detected: "6.97", "6.98", "7.03", "7.07", "7.08", "7.4", "7.57", "7.8", "8.25", "8.51" (10 additional values omitted).

fibrinogen_max

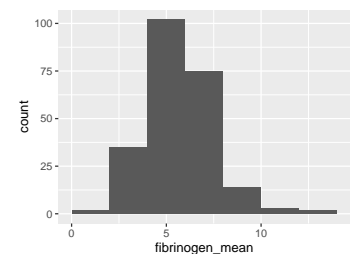
Feature	Result
Variable type	numeric
Number of missing obs.	828 (78.04 %)
Number of unique values	161
Median	5.88
1st and 3rd quartiles	4.54; 6.49
Min. and max.	1.17; 13.87



- Note that the following possible outlier values were detected: "8.15", "8.25", "8.28", "8.37", "8.51", "8.53", "8.76", "8.91", "8.97", "9.13" (9 additional values omitted).

fibrinogen_mean

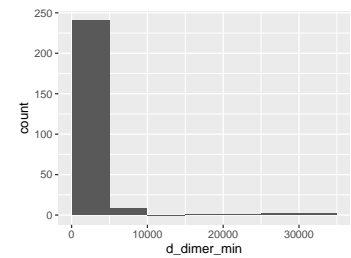
Feature	Result
Variable type	numeric
Number of missing obs.	828 (78.04 %)
Number of unique values	163
Median	5.75
1st and 3rd quartiles	4.52; 6.38
Min. and max.	1.17; 13.87



- Note that the following possible outlier values were detected: "7.48", "7.63", "7.68", "7.77", "7.83", "8.23", "8.25", "8.36", "8.51", "8.76" (9 additional values omitted).

d_dimer_min

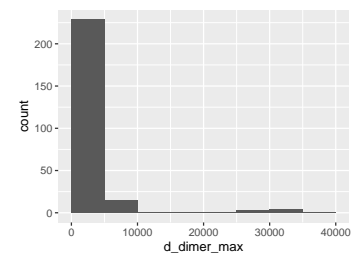
Feature	Result
Variable type	numeric
Number of missing obs.	807 (76.06 %)
Number of unique values	221
Median	951.5
1st and 3rd quartiles	513.5; 1725
Min. and max.	169; 34255



- Note that the following possible outlier values were detected: "7998", "9488", "9920", "18871", "24707", "27322", "28525", "31118", "34255".

d_dimer_max

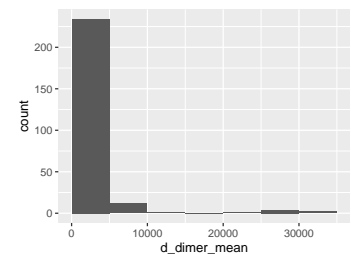
Feature	Result
Variable type	numeric
Number of missing obs.	807 (76.06 %)
Number of unique values	227
Median	1046.5
1st and 3rd quartiles	547; 2068.25
Min. and max.	169; 36480



- Note that the following possible outlier values were detected: "169", "170", "14725", "19574", "27322", "28525", "29297", "31118", "31651", "34255" (1 additional values omitted).

d_dimer_mean

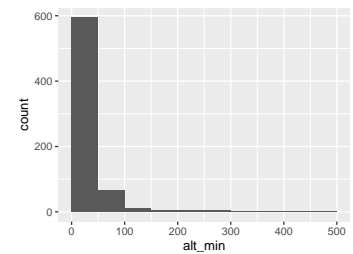
Feature	Result
Variable type	numeric
Number of missing obs.	807 (76.06 %)
Number of unique values	228
Median	1016.66
1st and 3rd quartiles	531.75; 1964.12
Min. and max.	169; 34255



- Note that the following possible outlier values were detected: "9444.5", "9488", "9920", "11307.33", "22990", "27322", "28179", "28525", "29868.67", "31118" (1 additional values omitted).

alt_min

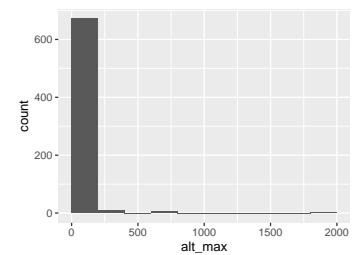
Feature	Result
Variable type	numeric
Number of missing obs.	375 (35.34 %)
Number of unique values	103
Median	21.5
1st and 3rd quartiles	14; 33
Min. and max.	5; 467



- Note that the following possible outlier values were detected: "5", "6", "7", "121", "126", "127", "133", "144", "174", "175" (11 additional values omitted).

alt_max

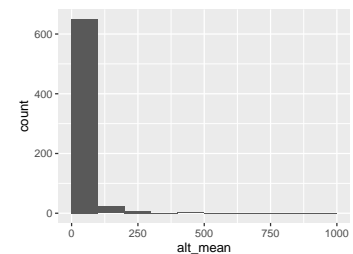
Feature	Result
Variable type	numeric
Number of missing obs.	375 (35.34 %)
Number of unique values	122
Median	25
1st and 3rd quartiles	15; 44
Min. and max.	5; 1861



- Note that the following possible outlier values were detected: "5", "183", "187", "198", "200", "223", "230", "245", "253", "258" (8 additional values omitted).

alt_mean

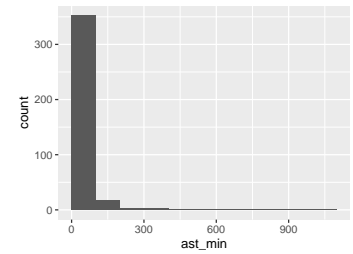
Feature	Result
Variable type	numeric
Number of missing obs.	375 (35.34 %)
Number of unique values	228
Median	23
1st and 3rd quartiles	15; 39.5
Min. and max.	5; 915.67



- Note that the following possible outlier values were detected: "5", "5.5", "5.67", "6", "6.5", "6.67", "7", "7.33", "179", "191.67" (12 additional values omitted).

ast__min

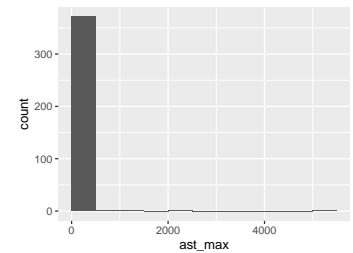
Feature	Result
Variable type	numeric
Number of missing obs.	686 (64.66 %)
Number of unique values	98
Median	30
1st and 3rd quartiles	22; 49
Min. and max.	3; 1096



- Note that the following possible outlier values were detected: "3", "7", "9", "10", "11", "12", "13", "14", "15", "16" (5 additional values omitted).

ast__max

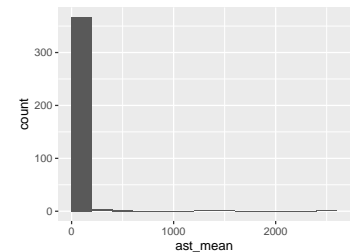
Feature	Result
Variable type	numeric
Number of missing obs.	686 (64.66 %)
Number of unique values	106
Median	37
1st and 3rd quartiles	25; 66
Min. and max.	3; 5486



- Note that the following possible outlier values were detected: "3", "7", "9", "12", "14", "15", "16", "357", "488", "547" (3 additional values omitted).

ast__mean

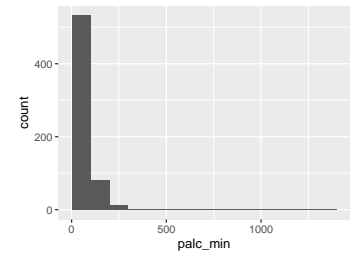
Feature	Result
Variable type	numeric
Number of missing obs.	686 (64.66 %)
Number of unique values	168
Median	33
1st and 3rd quartiles	24; 56.5
Min. and max.	3; 2493.33



- Note that the following possible outlier values were detected: "3", "7", "9", "12", "12.75", "14", "14.8", "15", "15.5", "16" (7 additional values omitted).

palc_min

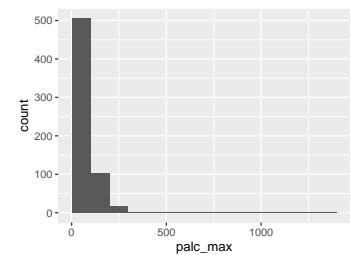
Feature	Result
Variable type	numeric
Number of missing obs.	432 (40.72 %)
Number of unique values	140
Median	61
1st and 3rd quartiles	49; 82
Min. and max.	18; 1363



- Note that the following possible outlier values were detected: "18", "20", "21", "22", "24", "26", "28", "29", "30", "31" (16 additional values omitted).

palc_max

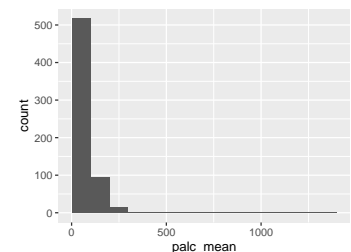
Feature	Result
Variable type	numeric
Number of missing obs.	432 (40.72 %)
Number of unique values	150
Median	65
1st and 3rd quartiles	53; 89
Min. and max.	20; 1363



- Note that the following possible outlier values were detected: "20", "22", "24", "26", "28", "29", "30", "31", "32", "33" (17 additional values omitted).

palc_mean

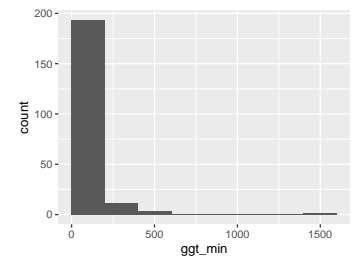
Feature	Result
Variable type	numeric
Number of missing obs.	432 (40.72 %)
Number of unique values	235
Median	63
1st and 3rd quartiles	51; 85.67
Min. and max.	19; 1363



- Note that the following possible outlier values were detected: "19", "21", "22.5", "24", "26", "27.6", "28", "29", "30", "31" (20 additional values omitted).

ggt_min

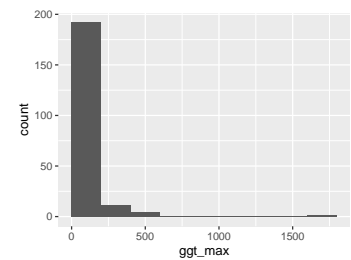
Feature	Result
Variable type	numeric
Number of missing obs.	853 (80.4 %)
Number of unique values	104
Median	44
1st and 3rd quartiles	24.75; 83.5
Min. and max.	5; 1508



- Note that the following possible outlier values were detected: "5", "8", "9", "464", "562", "1508".

ggt_max

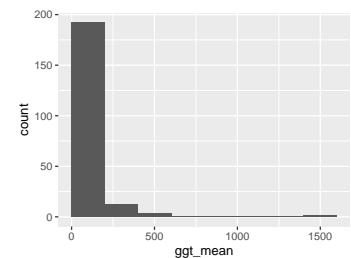
Feature	Result
Variable type	numeric
Number of missing obs.	853 (80.4 %)
Number of unique values	105
Median	44.5
1st and 3rd quartiles	27.75; 85.25
Min. and max.	5; 1658



- Note that the following possible outlier values were detected: "5", "8", "9", "11", "12", "13", "14", "479", "562", "1658".

ggt_mean

Feature	Result
Variable type	numeric
Number of missing obs.	853 (80.4 %)
Number of unique values	117
Median	44
1st and 3rd quartiles	25.88; 83.75
Min. and max.	5; 1583



- Note that the following possible outlier values were detected: "5", "8", "9", "11", "12", "471.5", "562", "1583".

amylase__min

- The variable only takes one value: "NA".

amylase__max

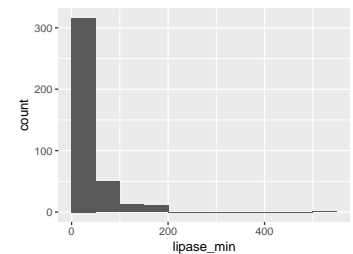
- The variable only takes one value: "NA".

amylase__mean

- The variable only takes one value: "NA".

lipase__min

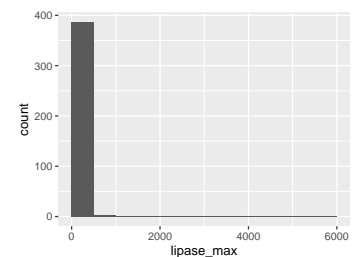
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	98
Median	26
1st and 3rd quartiles	16; 44
Min. and max.	5; 548



- Note that the following possible outlier values were detected: "5", "166", "174", "178", "181", "182", "185", "194", "548".

lipase__max

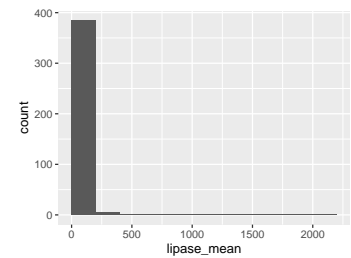
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	113
Median	29
1st and 3rd quartiles	17.25; 51
Min. and max.	5; 5709



- Note that the following possible outlier values were detected: "5", "6", "7", "229", "237", "253", "274", "444", "522", "685" (2 additional values omitted).

lipase__mean

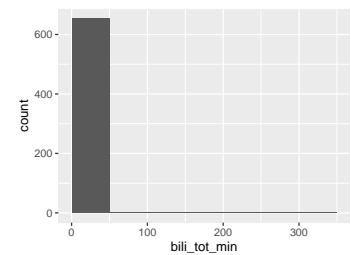
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	145
Median	27
1st and 3rd quartiles	17; 47.75
Min. and max.	5; 2178.33



- Note that the following possible outlier values were detected: "5", "6", "7", "7.5", "8", "8.33", "232.33", "234", "352", "354.4" (2 additional values omitted).

bili__tot__min

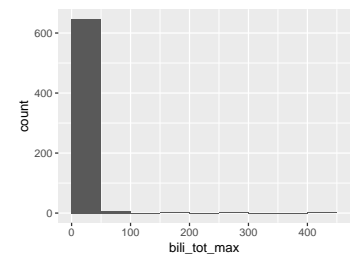
Feature	Result
Variable type	numeric
Number of missing obs.	404 (38.08 %)
Number of unique values	39
Median	8
1st and 3rd quartiles	6; 11
Min. and max.	2; 311



- Note that the following possible outlier values were detected: "2", "3", "4", "32", "36", "40", "44", "48", "50", "77" (3 additional values omitted).

bili__tot__max

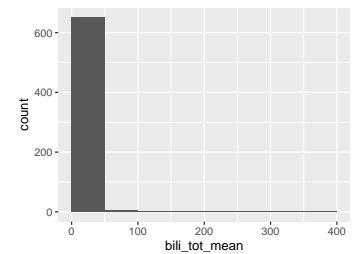
Feature	Result
Variable type	numeric
Number of missing obs.	404 (38.08 %)
Number of unique values	47
Median	9
1st and 3rd quartiles	7; 13
Min. and max.	3; 420



- Note that the following possible outlier values were detected: "3", "4", "38", "40", "42", "45", "46", "48", "51", "57" (7 additional values omitted).

bili_tot_mean

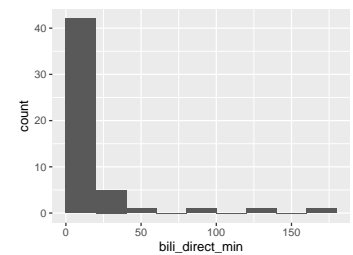
Feature	Result
Variable type	numeric
Number of missing obs.	404 (38.08 %)
Number of unique values	120
Median	8.5
1st and 3rd quartiles	6.5; 12
Min. and max.	3; 369.67



- Note that the following possible outlier values were detected: "3", "3.33", "3.5", "3.67", "4", "4.33", "4.5", "37", "37.67", "40" (9 additional values omitted).

bili_direct_min

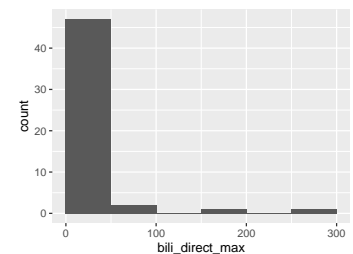
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	41
Median	12.4
1st and 3rd quartiles	7.3; 16.5
Min. and max.	3.4; 169.3



- Note that the following possible outlier values were detected: "57.4", "91.8", "136.3", "169.3".

bili_direct_max

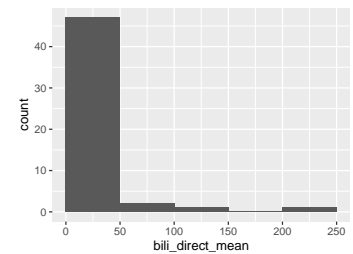
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	43
Median	13.8
1st and 3rd quartiles	7.75; 21.35
Min. and max.	4.5; 288.6



- Note that the following possible outlier values were detected: "98.9", "152.6", "288.6".

bili_direct_mean

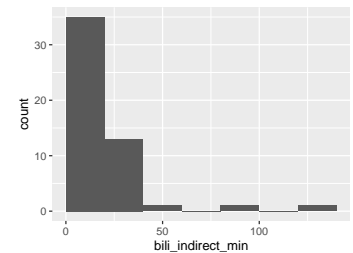
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	45
Median	13.1
1st and 3rd quartiles	7.3; 19.1
Min. and max.	4.5; 236.53



- Note that the following possible outlier values were detected: "95.35", "143.07", "236.53".

bili_indirect_min

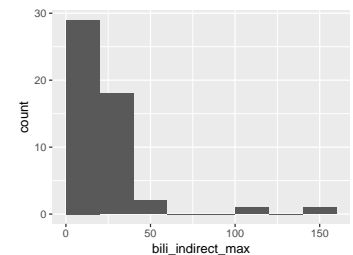
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	46
Median	17.4
1st and 3rd quartiles	13.3; 20.85
Min. and max.	5.8; 132



- Note that the following possible outlier values were detected: "31.4", "50.1", "85.8", "132".

bili_indirect_max

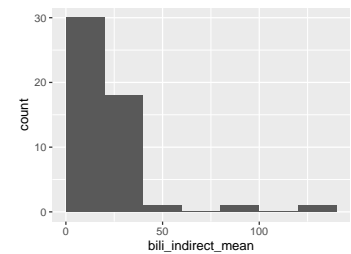
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	47
Median	19.3
1st and 3rd quartiles	15.05; 26.5
Min. and max.	7; 141.9



- Note that the following possible outlier values were detected: "7", "7.4", "7.5", "7.6", "104.3", "141.9".

bili_indirect_mean

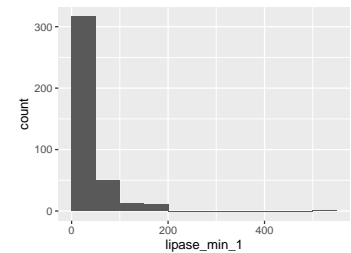
Feature	Result
Variable type	numeric
Number of missing obs.	1010 (95.19 %)
Number of unique values	49
Median	18.53
1st and 3rd quartiles	14.3; 22.87
Min. and max.	6.4; 136.13



- Note that the following possible outlier values were detected: "39.2", "51.1", "93.13", "136.13".

lipase_min_1

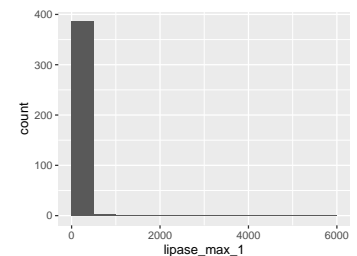
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	98
Median	26
1st and 3rd quartiles	16; 44
Min. and max.	5; 548



- Note that the following possible outlier values were detected: "5", "166", "174", "178", "181", "182", "185", "194", "548".

lipase_max_1

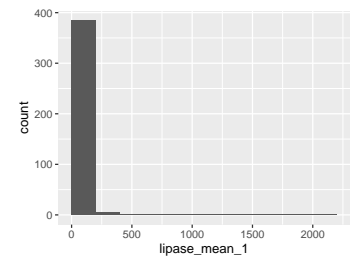
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	113
Median	29
1st and 3rd quartiles	17.25; 51
Min. and max.	5; 5709



- Note that the following possible outlier values were detected: "5", "6", "7", "229", "237", "253", "274", "444", "522", "685" (2 additional values omitted).

lipase__mean__1

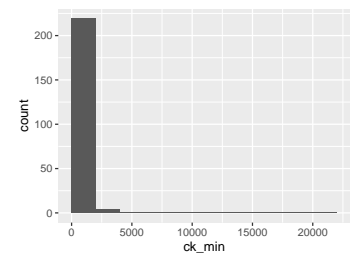
Feature	Result
Variable type	numeric
Number of missing obs.	671 (63.24 %)
Number of unique values	145
Median	27
1st and 3rd quartiles	17; 47.75
Min. and max.	5; 2178.33



- Note that the following possible outlier values were detected: "5", "6", "7", "7.5", "8", "8.33", "232.33", "234", "352", "354.4" (2 additional values omitted).

ck__min

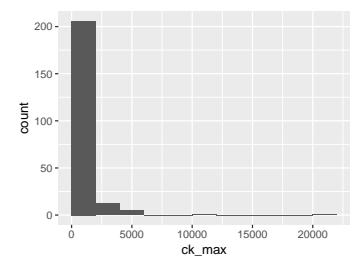
Feature	Result
Variable type	numeric
Number of missing obs.	836 (78.79 %)
Number of unique values	145
Median	130
1st and 3rd quartiles	70; 322
Min. and max.	13; 21926



- Note that the following possible outlier values were detected: "13", "14", "16", "18", "21", "22", "25", "29", "30", "32" (8 additional values omitted).

ck__max

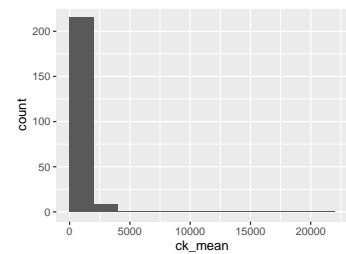
Feature	Result
Variable type	numeric
Number of missing obs.	836 (78.79 %)
Number of unique values	156
Median	232
1st and 3rd quartiles	97; 665
Min. and max.	14; 21926



- Note that the following possible outlier values were detected: "5990", "10132", "21926".

ck__mean

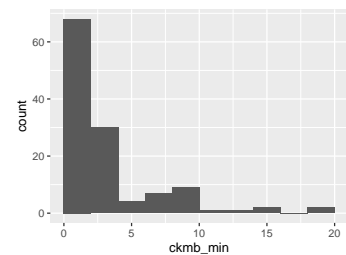
Feature	Result
Variable type	numeric
Number of missing obs.	836 (78.79 %)
Number of unique values	168
Median	197
1st and 3rd quartiles	88.2; 514
Min. and max.	14; 21926



- Note that the following possible outlier values were detected: "3723.33", "7708.5", "21926".

ckmb__min

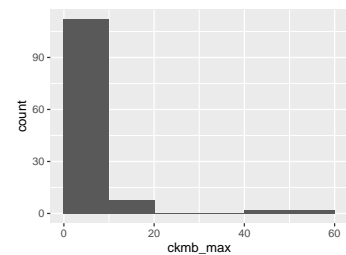
Feature	Result
Variable type	numeric
Number of missing obs.	937 (88.31 %)
Number of unique values	46
Median	1.9
1st and 3rd quartiles	1.08; 3.7
Min. and max.	0.4; 19.9



- Note that the following possible outlier values were detected: "19.9".

ckmb__max

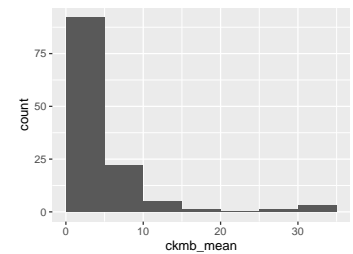
Feature	Result
Variable type	numeric
Number of missing obs.	937 (88.31 %)
Number of unique values	56
Median	2.8
1st and 3rd quartiles	1.5; 6.45
Min. and max.	0.4; 57.9



- Note that the following possible outlier values were detected: "0.4", "49.7", "56.6", "57.9".

ckmb__mean

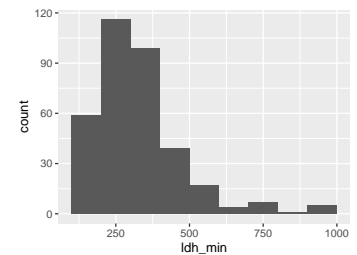
Feature	Result
Variable type	numeric
Number of missing obs.	937 (88.31 %)
Number of unique values	71
Median	2.47
1st and 3rd quartiles	1.37; 5.2
Min. and max.	0.4; 34.27



- Note that the following possible outlier values were detected: "0.4", "0.45", "29.87", "33.65", "34.27".

ldh__min

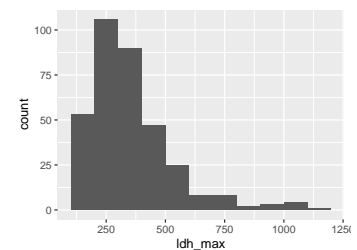
Feature	Result
Variable type	numeric
Number of missing obs.	714 (67.3 %)
Number of unique values	218
Median	299
1st and 3rd quartiles	220.5; 376
Min. and max.	107; 965



- Note that the following possible outlier values were detected: "707", "709", "713", "737", "745", "812", "922", "926", "941", "958" (1 additional values omitted).

ldh__max

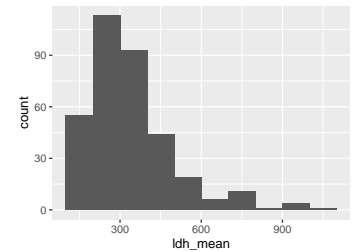
Feature	Result
Variable type	numeric
Number of missing obs.	714 (67.3 %)
Number of unique values	225
Median	317
1st and 3rd quartiles	233.5; 413
Min. and max.	107; 1121



- Note that the following possible outlier values were detected: "898", "922", "926", "965", "1009", "1093", "1121".

ldh_mean

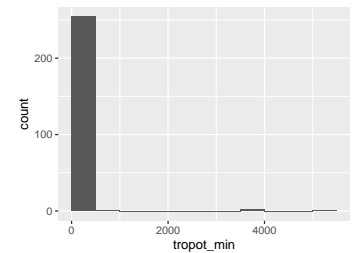
Feature	Result
Variable type	numeric
Number of missing obs.	714 (67.3 %)
Number of unique values	241
Median	305
1st and 3rd quartiles	228; 398.5
Min. and max.	107; 1039.5



- Note that the following possible outlier values were detected: "812", "922", "926", "965", "1000", "1039.5".

tropot_min

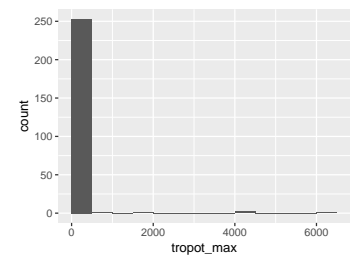
Feature	Result
Variable type	numeric
Number of missing obs.	803 (75.68 %)
Number of unique values	75
Median	21
1st and 3rd quartiles	11; 45
Min. and max.	10; 5389



- Note that the following possible outlier values were detected: "315", "342", "917", "3905", "5389".

tropot_max

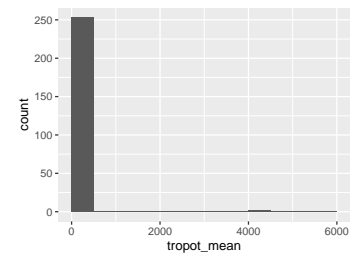
Feature	Result
Variable type	numeric
Number of missing obs.	803 (75.68 %)
Number of unique values	88
Median	26
1st and 3rd quartiles	13; 56
Min. and max.	10; 6098



- Note that the following possible outlier values were detected: "363", "440", "471", "851", "1617", "4340", "6098".

tropot__mean

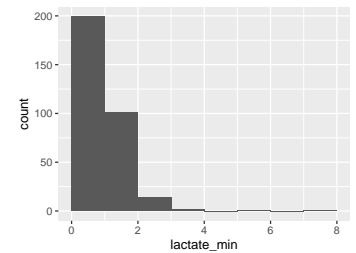
Feature	Result
Variable type	numeric
Number of missing obs.	803 (75.68 %)
Number of unique values	129
Median	23.75
1st and 3rd quartiles	12.23; 50.62
Min. and max.	10; 5743.5



- Note that the following possible outlier values were detected: "351.33", "542.5", "1214", "4056.33", "5743.5".

lactate__min

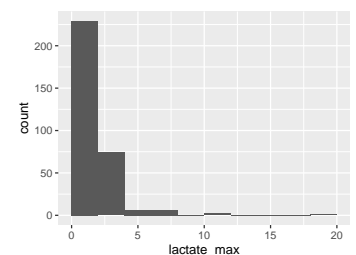
Feature	Result
Variable type	numeric
Number of missing obs.	743 (70.03 %)
Number of unique values	28
Median	0.9
1st and 3rd quartiles	0.7; 1.2
Min. and max.	0.2; 7.6



- Note that the following possible outlier values were detected: "0.2", "0.3", "0.4", "3.1", "3.4", "5.6", "7.6".

lactate__max

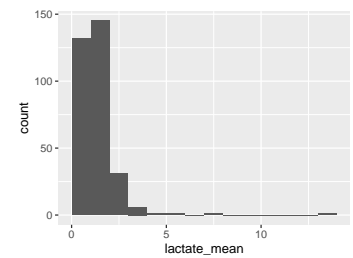
Feature	Result
Variable type	numeric
Number of missing obs.	743 (70.03 %)
Number of unique values	44
Median	1.4
1st and 3rd quartiles	1; 2.18
Min. and max.	0.5; 18.7



- Note that the following possible outlier values were detected: "0.5", "7.5", "7.6", "11.4", "18.7".

lactate__mean

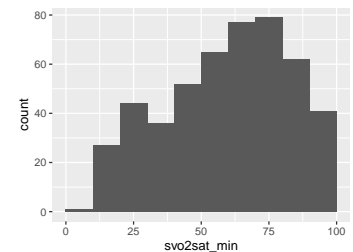
Feature	Result
Variable type	numeric
Number of missing obs.	743 (70.03 %)
Number of unique values	103
Median	1.17
1st and 3rd quartiles	0.9; 1.6
Min. and max.	0.41; 13.11



- Note that the following possible outlier values were detected: "0.41", "0.5", "4.07", "5.24", "7.6", "13.11".

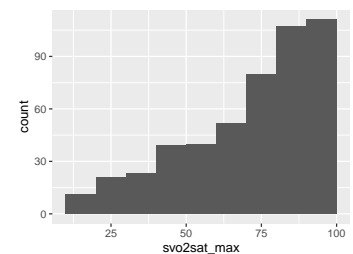
svo2sat__min

Feature	Result
Variable type	numeric
Number of missing obs.	577 (54.38 %)
Number of unique values	88
Median	63
1st and 3rd quartiles	43; 78.25
Min. and max.	9; 100



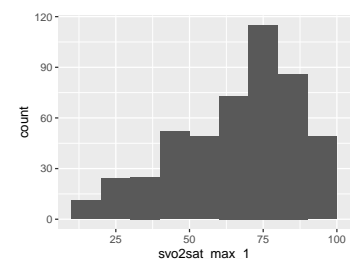
svo2sat__max

Feature	Result
Variable type	numeric
Number of missing obs.	577 (54.38 %)
Number of unique values	83
Median	78
1st and 3rd quartiles	57; 90
Min. and max.	12; 100



svo2sat__max_1

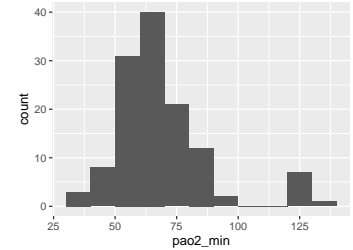
Feature	Result
Variable type	numeric
Number of missing obs.	577 (54.38 %)
Number of unique values	168
Median	71.5
1st and 3rd quartiles	53; 81
Min. and max.	12; 100



- Note that the following possible outlier values were detected: "94", "94.33", "95", "96", "98", "100".

pao2_min

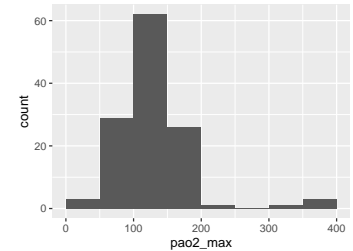
Feature	Result
Variable type	numeric
Number of missing obs.	936 (88.22 %)
Number of unique values	85
Median	63.8
1st and 3rd quartiles	57.5; 73.4
Min. and max.	33.4; 137



- Note that the following possible outlier values were detected: "33.4", "34.5", "40", "43.6", "43.7", "47.1", "47.2", "122", "123", "127" (1 additional values omitted).

pao2_max

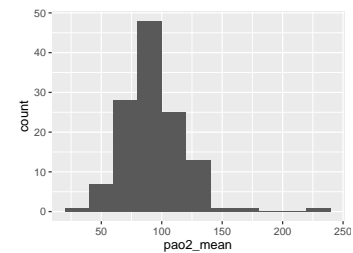
Feature	Result
Variable type	numeric
Number of missing obs.	936 (88.22 %)
Number of unique values	75
Median	133
1st and 3rd quartiles	99; 149
Min. and max.	33.4; 388



- Note that the following possible outlier values were detected: "194", "208", "336", "371", "388".

pao2_mean

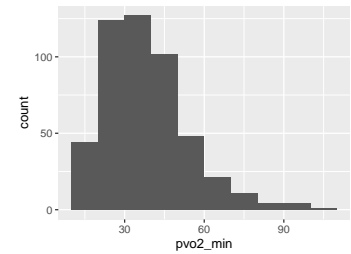
Feature	Result
Variable type	numeric
Number of missing obs.	936 (88.22 %)
Number of unique values	95
Median	92.7
1st and 3rd quartiles	77.85; 108.64
Min. and max.	33.4; 223.54



- Note that the following possible outlier values were detected: "170.97", "223.54".

pvo2__min

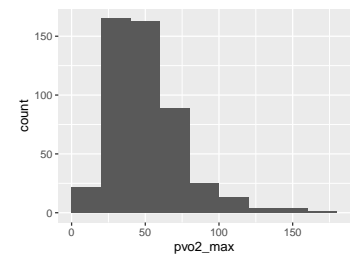
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	286
Median	36.2
1st and 3rd quartiles	26.52; 45.58
Min. and max.	12.1; 108



- Note that the following possible outlier values were detected: "77.6", "81.3", "83.6", "84.2", "84.3", "95.3", "99.3", "99.6", "100", "108".

pvo2__max

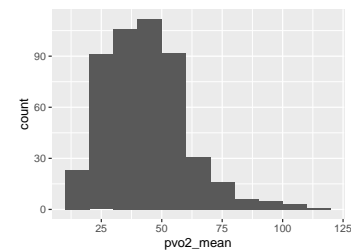
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	321
Median	45.85
1st and 3rd quartiles	32.3; 61.8
Min. and max.	14.6; 169



- Note that the following possible outlier values were detected: "137", "138", "149", "151", "158", "160", "169".

pvo2__mean

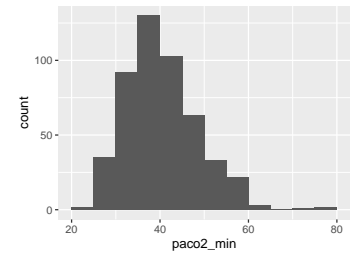
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	346
Median	42.05
1st and 3rd quartiles	30.68; 52.76
Min. and max.	14.6; 111.25



- Note that the following possible outlier values were detected: "94.92", "95.3", "99.3", "99.6", "100", "105.45", "107.15", "108", "111.25".

paco2__min

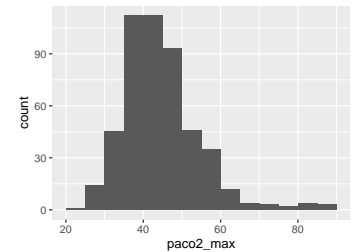
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	220
Median	39.4
1st and 3rd quartiles	34.7; 45.25
Min. and max.	23.7; 78.1



- Note that the following possible outlier values were detected: "23.7", "24.9", "74.2", "78.1".

paco2__max

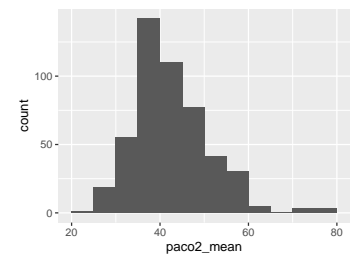
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	236
Median	42.7
1st and 3rd quartiles	37.9; 49.27
Min. and max.	23.7; 88.5



- Note that the following possible outlier values were detected: "23.7", "25.8", "26.3", "28.2", "28.6", "28.7", "28.9", "29", "29.3", "29.5" (8 additional values omitted).

paco2__mean

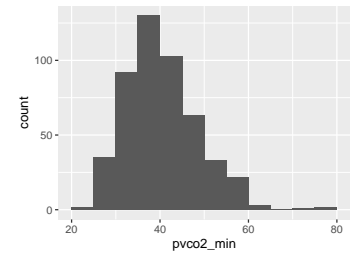
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	294
Median	41.2
1st and 3rd quartiles	36.76; 47.58
Min. and max.	23.7; 78.1



- Note that the following possible outlier values were detected: "23.7", "25.8", "26.3", "76.83", "78.1".

pvco2__min

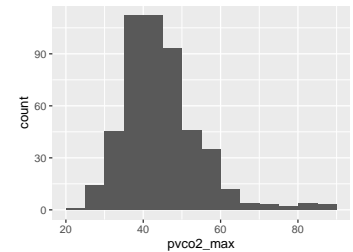
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	220
Median	39.4
1st and 3rd quartiles	34.7; 45.25
Min. and max.	23.7; 78.1



- Note that the following possible outlier values were detected: "23.7", "24.9", "74.2", "78.1".

pvco2__max

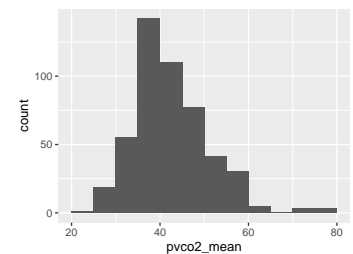
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	236
Median	42.7
1st and 3rd quartiles	37.9; 49.27
Min. and max.	23.7; 88.5



- Note that the following possible outlier values were detected: "23.7", "25.8", "26.3", "28.2", "28.6", "28.7", "28.9", "29", "29.3", "29.5" (8 additional values omitted).

pvco2__mean

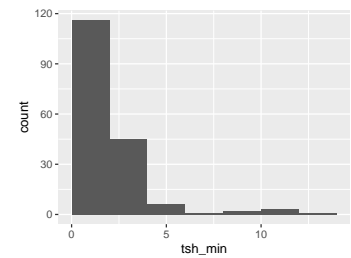
Feature	Result
Variable type	numeric
Number of missing obs.	575 (54.19 %)
Number of unique values	294
Median	41.2
1st and 3rd quartiles	36.76; 47.58
Min. and max.	23.7; 78.1



- Note that the following possible outlier values were detected: "23.7", "25.8", "26.3", "76.83", "78.1".

tsh__min

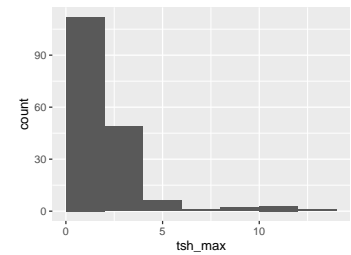
Feature	Result
Variable type	numeric
Number of missing obs.	887 (83.6 %)
Number of unique values	141
Median	1.29
1st and 3rd quartiles	0.76; 2.48
Min. and max.	0.05; 13.64



- Note that the following possible outlier values were detected: "0.05", "0.1", "0.14", "10.66", "11.78", "13.64".

tsh__max

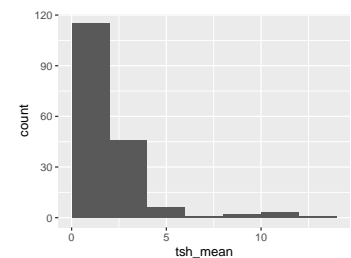
Feature	Result
Variable type	numeric
Number of missing obs.	887 (83.6 %)
Number of unique values	140
Median	1.33
1st and 3rd quartiles	0.76; 2.58
Min. and max.	0.05; 13.64



- Note that the following possible outlier values were detected: "0.05", "0.1", "0.14", "11.78", "13.64".

tsh__mean

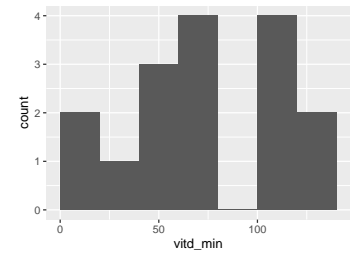
Feature	Result
Variable type	numeric
Number of missing obs.	887 (83.6 %)
Number of unique values	141
Median	1.31
1st and 3rd quartiles	0.76; 2.48
Min. and max.	0.05; 13.64



- Note that the following possible outlier values were detected: "0.05", "0.1", "0.14", "0.17", "10.66", "11.78", "13.64".

vitd_min

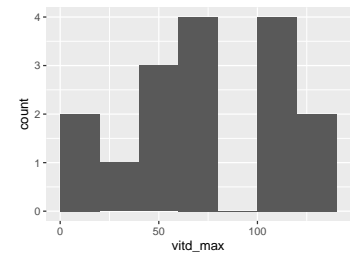
Feature	Result
Variable type	numeric
Number of missing obs.	1045 (98.49 %)
Number of unique values	16
Median	72.5
1st and 3rd quartiles	51.5; 105.5
Min. and max.	6; 130



- Note that the following possible outlier values were detected: "6".

vitd_max

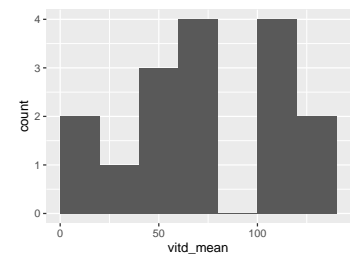
Feature	Result
Variable type	numeric
Number of missing obs.	1045 (98.49 %)
Number of unique values	16
Median	72.5
1st and 3rd quartiles	51.5; 105.5
Min. and max.	6; 130



- Note that the following possible outlier values were detected: "6".

vitd_mean

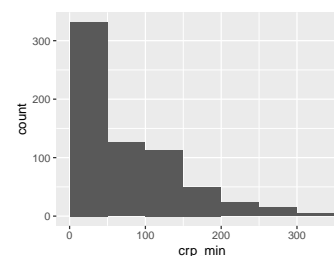
Feature	Result
Variable type	numeric
Number of missing obs.	1045 (98.49 %)
Number of unique values	16
Median	72.5
1st and 3rd quartiles	51.5; 105.5
Min. and max.	6; 130



- Note that the following possible outlier values were detected: "6".

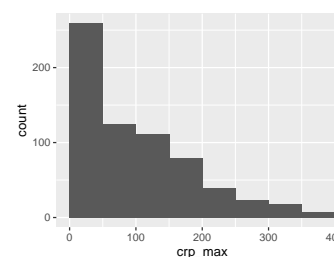
crp_min

Feature	Result
Variable type	numeric
Number of missing obs.	398 (37.51 %)
Number of unique values	458
Median	49.5
1st and 3rd quartiles	19.35; 115.95
Min. and max.	5; 344.5



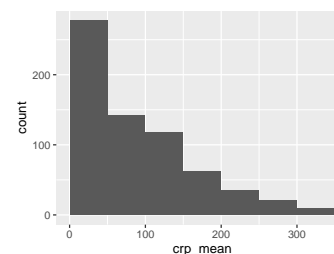
crp_max

Feature	Result
Variable type	numeric
Number of missing obs.	398 (37.51 %)
Number of unique values	485
Median	78.2
1st and 3rd quartiles	25.45; 150.15
Min. and max.	5; 384.6



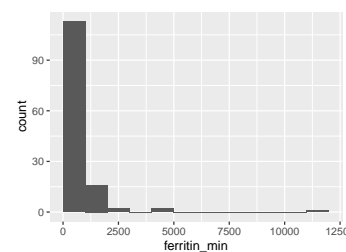
crp_mean

Feature	Result
Variable type	numeric
Number of missing obs.	398 (37.51 %)
Number of unique values	510
Median	65.5
1st and 3rd quartiles	24.7; 130.01
Min. and max.	5; 344.5



ferritin_min

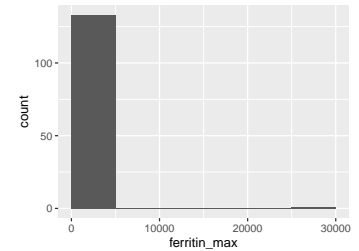
Feature	Result
Variable type	numeric
Number of missing obs.	927 (87.37 %)
Number of unique values	119
Median	301.5
1st and 3rd quartiles	160.25; 722.5
Min. and max.	10; 11540



- Note that the following possible outlier values were detected: "10", "20", "22", "26", "32", "51", "52", "4867", "11540".

ferritin_max

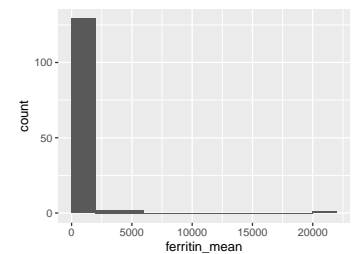
Feature	Result
Variable type	numeric
Number of missing obs.	927 (87.37 %)
Number of unique values	116
Median	320.5
1st and 3rd quartiles	165; 727.75
Min. and max.	10; 28696



- Note that the following possible outlier values were detected: "10", "20", "24", "27", "32", "4867", "28696".
-

ferritin_mean

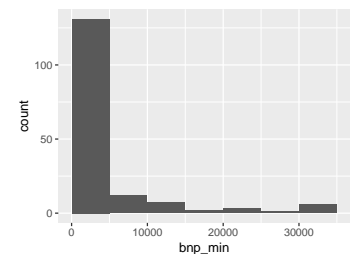
Feature	Result
Variable type	numeric
Number of missing obs.	927 (87.37 %)
Number of unique values	120
Median	315
1st and 3rd quartiles	162.5; 722.5
Min. and max.	10; 20118



- Note that the following possible outlier values were detected: "10", "20", "23", "26.5", "32", "4867", "20118".
-

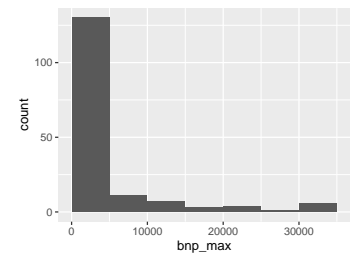
bnp_min

Feature	Result
Variable type	numeric
Number of missing obs.	899 (84.73 %)
Number of unique values	139
Median	835
1st and 3rd quartiles	198; 3576.5
Min. and max.	5; 35000



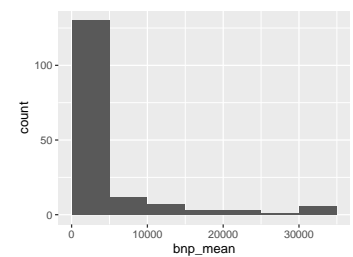
bnp__max

Feature	Result
Variable type	numeric
Number of missing obs.	899 (84.73 %)
Number of unique values	141
Median	894.5
1st and 3rd quartiles	204; 4068.5
Min. and max.	5; 35000



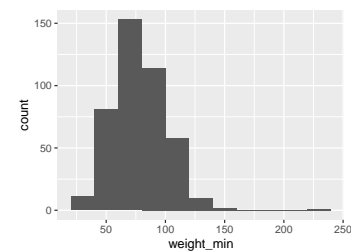
bnp__mean

Feature	Result
Variable type	numeric
Number of missing obs.	899 (84.73 %)
Number of unique values	141
Median	894.5
1st and 3rd quartiles	204; 3845
Min. and max.	5; 35000



weight__min

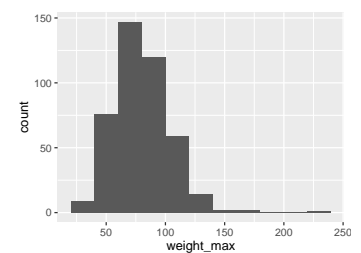
Feature	Result
Variable type	numeric
Number of missing obs.	631 (59.47 %)
Number of unique values	300
Median	77.5
1st and 3rd quartiles	62.52; 91.7
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "135.7", "147.3", "150.4", "236".

weight__max

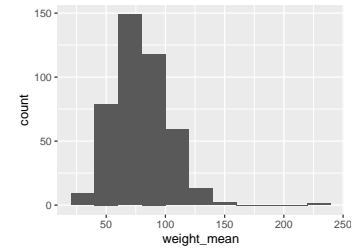
Feature	Result
Variable type	numeric
Number of missing obs.	631 (59.47 %)
Number of unique values	290
Median	78.95
1st and 3rd quartiles	64; 93.2
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "135.7", "147.3", "150.4", "173", "236".

weight_mean

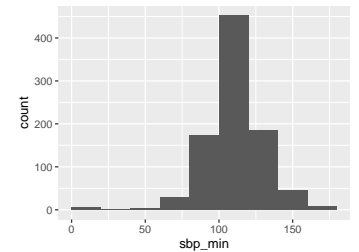
Feature	Result
Variable type	numeric
Number of missing obs.	631 (59.47 %)
Number of unique values	316
Median	78
1st and 3rd quartiles	62.96; 93.15
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "147.3", "150.4", "236".

sbp_min

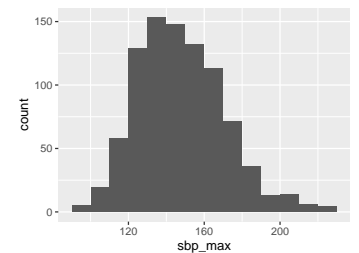
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	96
Median	111
1st and 3rd quartiles	101; 121
Min. and max.	11; 177



- Note that the following possible outlier values were detected: "11", "12", "16", "19", "54", "60", "62", "65", "66", "69" (13 additional values omitted).

sbp_max

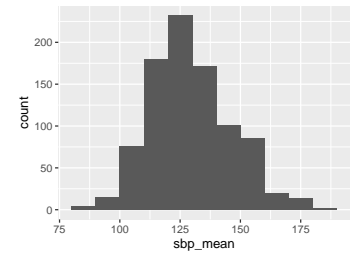
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	119
Median	147
1st and 3rd quartiles	132; 164
Min. and max.	95; 224



- Note that the following possible outlier values were detected: "221", "223", "224".

sbp_mean

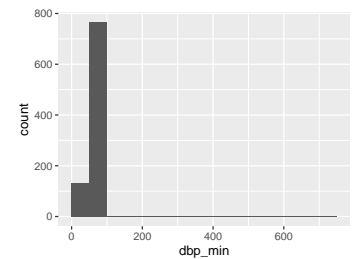
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	648
Median	127.22
1st and 3rd quartiles	117.78; 139.81
Min. and max.	86.83; 188.27



- Note that the following possible outlier values were detected: "86.83", "87.88", "89", "89.13", "92.4", "94", "95.17", "96", "96.2", "96.63" (9 additional values omitted).

dbp_min

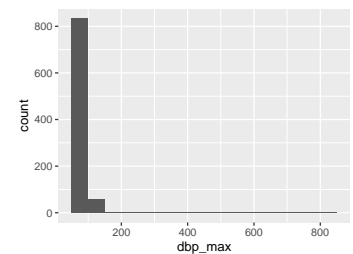
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	66
Median	61
1st and 3rd quartiles	55; 68
Min. and max.	6; 719



- Note that the following possible outlier values were detected: "6", "24", "28", "30", "34", "35", "89", "91", "92", "93" (4 additional values omitted).

dbp_max

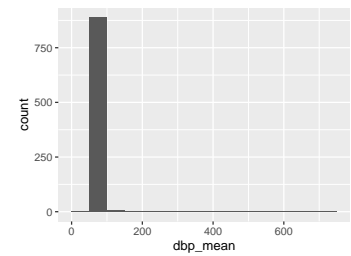
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	77
Median	83
1st and 3rd quartiles	77; 90
Min. and max.	55; 811



- Note that the following possible outlier values were detected: "55", "56", "59", "60", "61", "62", "63", "116", "117", "118" (16 additional values omitted).

dbp__mean

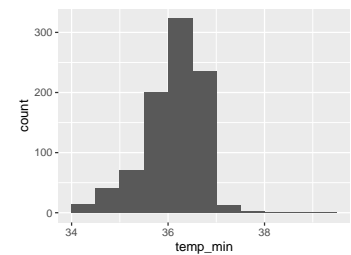
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	567
Median	72
1st and 3rd quartiles	66.4; 78.11
Min. and max.	48.19; 719



- Note that the following possible outlier values were detected: "48.19", "49.27", "50.4", "98", "99.1", "99.2", "101.27", "101.75", "103.67", "103.88" (6 additional values omitted).

temp__min

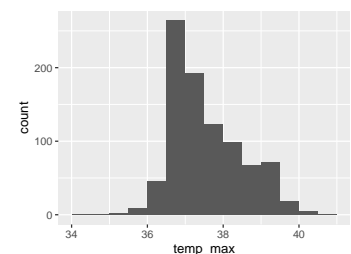
Feature	Result
Variable type	numeric
Number of missing obs.	163 (15.36 %)
Number of unique values	36
Median	36.3
1st and 3rd quartiles	35.9; 36.6
Min. and max.	34; 39.3



- Note that the following possible outlier values were detected: "34", "37.2", "37.3", "37.4", "37.5", "37.6", "39.3".

temp__max

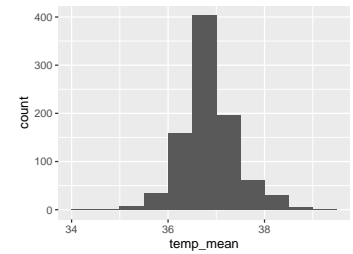
Feature	Result
Variable type	numeric
Number of missing obs.	163 (15.36 %)
Number of unique values	51
Median	37.3
1st and 3rd quartiles	37; 38.2
Min. and max.	34; 40.7



- Note that the following possible outlier values were detected: "34", "35.3", "35.4", "35.6", "35.8", "35.9", "36", "36.1", "36.2", "36.3" (3 additional values omitted).

temp_mean

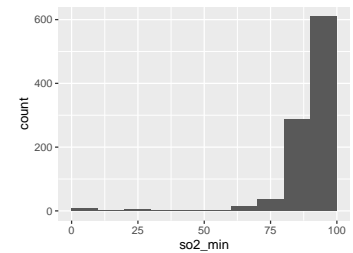
Feature	Result
Variable type	numeric
Number of missing obs.	163 (15.36 %)
Number of unique values	225
Median	36.86
1st and 3rd quartiles	36.54; 37.15
Min. and max.	34; 39.3



- Note that the following possible outlier values were detected: "34", "35.25", "35.33", "35.38", "35.4", "35.43", "35.45", "35.5", "35.53", "38.01" (28 additional values omitted).

so2_min

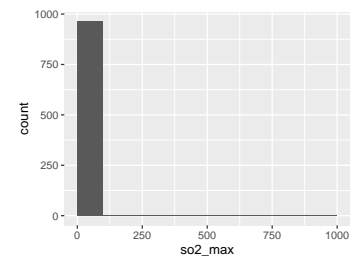
Feature	Result
Variable type	numeric
Number of missing obs.	96 (9.05 %)
Number of unique values	52
Median	92
1st and 3rd quartiles	88; 95
Min. and max.	0; 100



- Note that the following possible outlier values were detected: "0", "1", "2", "5", "8", "9", "18", "20", "23", "24" (13 additional values omitted).

so2_max

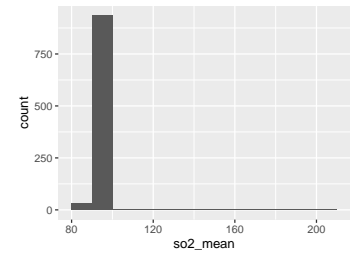
Feature	Result
Variable type	numeric
Number of missing obs.	96 (9.05 %)
Number of unique values	14
Median	98
1st and 3rd quartiles	97; 99
Min. and max.	85; 969



- Note that the following possible outlier values were detected: "85", "88", "90", "92", "93", "966", "969".

so2__mean

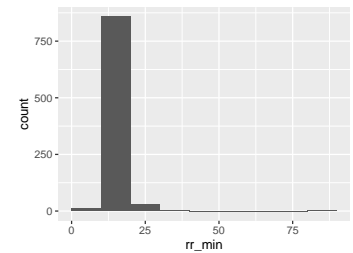
Feature	Result
Variable type	numeric
Number of missing obs.	96 (9.05 %)
Number of unique values	454
Median	95.17
1st and 3rd quartiles	93.59; 96.88
Min. and max.	81.07; 201.5



- Note that the following possible outlier values were detected: "81.07", "81.33", "82", "83.17", "84.29", "84.46", "85.57", "86.06", "86.89", "88" (3 additional values omitted).

rr__min

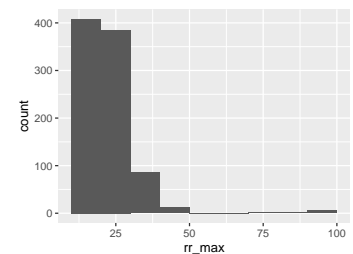
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	20
Median	18
1st and 3rd quartiles	17; 20
Min. and max.	0; 85



- Note that the following possible outlier values were detected: "0", "2", "8", "9", "10", "12", "26", "30", "32", "85".

rr__max

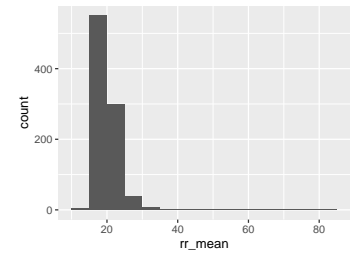
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	35
Median	22
1st and 3rd quartiles	20; 24
Min. and max.	10; 98



- Note that the following possible outlier values were detected: "10", "14", "16", "18", "42", "44", "45", "48", "50", "80" (5 additional values omitted).

rr_mean

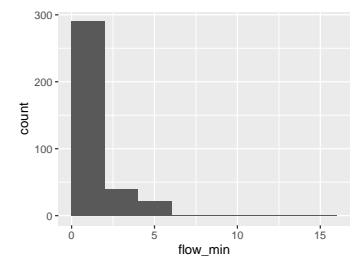
Feature	Result
Variable type	numeric
Number of missing obs.	160 (15.08 %)
Number of unique values	306
Median	19.9
1st and 3rd quartiles	19; 21
Min. and max.	10; 85



- Note that the following possible outlier values were detected: "10", "14", "14.57", "15.67", "15.8", "16", "16.67", "16.73", "16.75", "16.8" (47 additional values omitted).

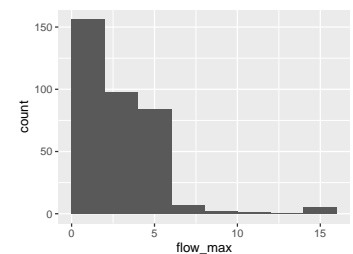
flow_min

Feature	Result
Variable type	numeric
Number of missing obs.	708 (66.73 %)
Number of unique values	13
Median	1
1st and 3rd quartiles	0.5; 2
Min. and max.	0.5; 15



flow_max

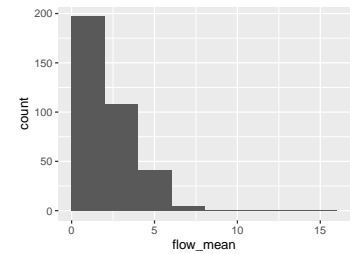
Feature	Result
Variable type	numeric
Number of missing obs.	708 (66.73 %)
Number of unique values	16
Median	3
1st and 3rd quartiles	2; 5
Min. and max.	0.5; 15



- Note that the following possible outlier values were detected: "10", "12", "15".

flow__mean

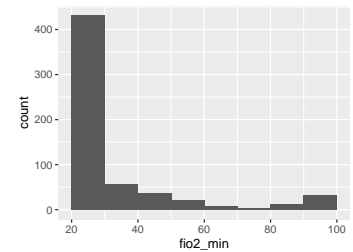
Feature	Result
Variable type	numeric
Number of missing obs.	708 (66.73 %)
Number of unique values	162
Median	1.9
1st and 3rd quartiles	1.11; 3
Min. and max.	0.5; 15



- Note that the following possible outlier values were detected: "10", "15".

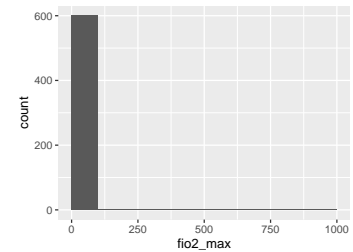
fio2__min

Feature	Result
Variable type	numeric
Number of missing obs.	460 (43.36 %)
Number of unique values	39
Median	21
1st and 3rd quartiles	21; 35
Min. and max.	21; 100



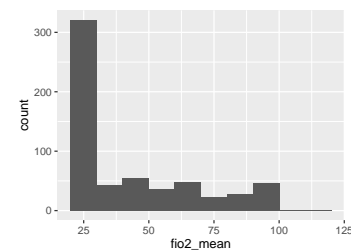
fio2__max

Feature	Result
Variable type	numeric
Number of missing obs.	460 (43.36 %)
Number of unique values	37
Median	32
1st and 3rd quartiles	21; 97
Min. and max.	21; 954



fio2__mean

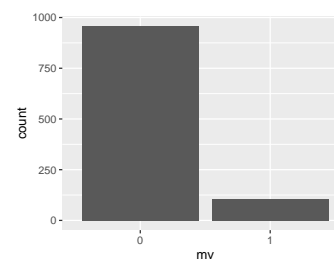
Feature	Result
Variable type	numeric
Number of missing obs.	460 (43.36 %)
Number of unique values	258
Median	27.67
1st and 3rd quartiles	21; 59.93
Min. and max.	21; 114.45



mv

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

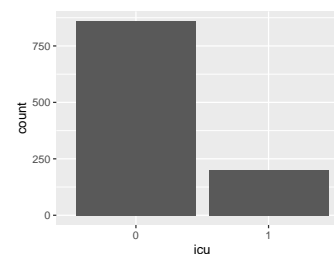
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



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: Mon Feb 01 2021 20:23:48
- Report was run from directory: /Users/eyamga/Documents/Médecine/Recherche/CODA19/git/CODA19-Phenotyper/r_ey
- 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 = covid72h_notimputed, render = FALSE, replace = TRUE)`