

covid24h_notimputed

Autogenerated data summary from dataMaid

2021-02-01 18:39:24

Data report overview

The dataset examined has the following dimensions:

Feature	Result
Number of observations	1045
Number of variables	291

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	74.83 %	
chf	numeric	3	74.83 %	
pvd	numeric	3	74.83 %	×
cevd	numeric	3	74.83 %	
dementia	numeric	3	74.83 %	
copd	numeric	3	74.83 %	
rheumd	numeric	3	74.83 %	×
pud	numeric	3	74.83 %	×
mld	numeric	3	74.83 %	×
diab	numeric	3	74.83 %	
diabwc	numeric	3	74.83 %	
hp	numeric	3	74.83 %	×
rend	numeric	3	74.83 %	
canc	numeric	3	74.83 %	
msld	numeric	3	74.83 %	×
metacanc	numeric	3	74.83 %	
aids	numeric	2	74.83 %	×
score	numeric	5	74.83 %	
neuromuscular_blocking_agents	numeric	3	8.61 %	×
x5_alpha_reductase_inhibitors	numeric	3	8.61 %	
acetaminophene	numeric	3	8.61 %	
adjuvants_anesthesia	numeric	3	8.61 %	×
adrenergic_alpha_1_receptor_antagonists	numeric	3	8.61 %	
adrenergic_beta_3_receptor_agonists	numeric	3	8.61 %	
adrenergic_beta_antagonists	numeric	3	8.61 %	
adrenergic_uptake_inhibitors	numeric	3	8.61 %	×
alcohol_deterrents	numeric	3	8.61 %	×
analgesics	numeric	3	8.61 %	
analgesics_opioid	numeric	3	8.61 %	
androgens	numeric	3	8.61 %	×
anesthetics_local	numeric	3	8.61 %	
anti_anxiety_agents	numeric	3	8.61 %	×
anti_arrhythmia_agents	numeric	3	8.61 %	
anti_asthmatic_agents	numeric	3	8.61 %	
anti_bacterial_agents	numeric	3	8.61 %	
anti_infective_agents_local	numeric	3	8.61 %	
anti_inflammatory_agents	numeric	3	8.61 %	

	Variable class	# unique values	Missing observations	Any problems?
anti_inflammatory_agents_non_steroidal	numeric	3	8.61 %	
anti_ulcer_agents	numeric	3	8.61 %	
anticholesteremic_agents	numeric	3	8.61 %	
anticoagulants	numeric	3	8.61 %	
anticonvulsants	numeric	3	8.61 %	
antidepressive_agents	numeric	3	8.61 %	
antidepressive_agents_tricyclic	numeric	3	8.61 %	×
antidiarrheals	numeric	3	8.61 %	
antiemetics	numeric	3	8.61 %	
antifibrinolytic_agents	numeric	3	8.61 %	×
antifungal_agents	numeric	3	8.61 %	×
antihypertensive_agents	numeric	3	8.61 %	
antimalarials	numeric	3	8.61 %	×
antimetabolites	numeric	3	8.61 %	
antineoplastic_agents_hormonal	numeric	3	8.61 %	×
antiparkinson_agents	numeric	3	8.61 %	
antipruritics	numeric	3	8.61 %	
antipsychotic_agents	numeric	3	8.61 %	
antithyroid_agents	numeric	3	8.61 %	×
antitubercular_agents	numeric	3	8.61 %	×
antitussive_agents	numeric	3	8.61 %	
antiviral_agents	numeric	3	8.61 %	
benzodiazepines	numeric	3	8.61 %	
bicarbonate	numeric	3	8.61 %	
bone_density_conservation_agents	numeric	3	8.61 %	
bronchodilator_agents	numeric	3	8.61 %	
calcium_regulating_hormones_and_agents	numeric	3	8.61 %	
carbonic_anhydrase_inhibitors	numeric	3	8.61 %	
chelating_agents	numeric	3	8.61 %	
cholagogues_and_choleretics	numeric	3	8.61 %	
cholinesterase_inhibitors	numeric	3	8.61 %	
contraceptive_agents_hormonal	numeric	3	8.61 %	×
diuretics	numeric	3	8.61 %	
factor_xa_inhibitors	numeric	3	8.61 %	
fibrinolytic_agents	numeric	3	8.61 %	×
gastrointestinal_agents	numeric	3	8.61 %	
glucocorticoids	numeric	3	8.61 %	
gout_suppressants	numeric	3	8.61 %	
hematologic_agents	numeric	3	8.61 %	×
hemostatics	numeric	3	8.61 %	×
hiv_medication	numeric	3	8.61 %	
hypoglycemic_agents	numeric	3	8.61 %	
immunosuppressive_agents	numeric	3	8.61 %	
laxatives	numeric	3	8.61 %	
levothyroxine	numeric	3	8.61 %	
miotics	numeric	3	8.61 %	×
muscarinic_antagonists	numeric	3	8.61 %	×
muscle_relaxants_central	numeric	3	8.61 %	×
narcotic_antagonists	numeric	3	8.61 %	×
neuromuscular_blocking_agents_2	numeric	3	8.61 %	
ophthalmic_solutions	numeric	3	8.61 %	
parasympatholytics	numeric	3	8.61 %	
platelet_aggregation_inhibitors	numeric	3	8.61 %	
progestins	numeric	3	8.61 %	×

	Variable class	# unique values	Missing observations	Any problems?
reverse_transcriptase_inhibitors	numeric	3	8.61 %	×
sedation	numeric	3	8.61 %	
serotonin_5_ht1_receptor_agonists	numeric	3	8.61 %	×
serotonin_uptake_inhibitors	numeric	3	8.61 %	
sleep_aids_pharmaceutical	numeric	3	8.61 %	
smoking_cessation_agents	numeric	3	8.61 %	
vasodilator_agents	numeric	3	8.61 %	
vasopressors	numeric	3	8.61 %	
vitamin_b_complex	numeric	3	8.61 %	
vitamins	numeric	3	8.61 %	
hemoglobin_min	numeric	103	23.44 %	×
hemoglobin_max	numeric	102	23.44 %	×
hemoglobin_mean	numeric	212	23.44 %	×
plt_min	numeric	304	23.44 %	×
plt_max	numeric	320	23.44 %	×
plt_mean	numeric	428	23.44 %	×
wbc_min	numeric	153	23.44 %	×
wbc_max	numeric	175	23.44 %	×
wbc_mean	numeric	275	23.44 %	×
albumin_min	numeric	35	57.51 %	×
albumin_max	numeric	31	57.51 %	×
albumin_mean	numeric	75	57.51 %	×
globulin_min	numeric	4	99.71 %	×
globulin_max	numeric	4	99.71 %	×
globulin_mean	numeric	4	99.71 %	×
protein_min	numeric	33	92.73 %	×
protein_max	numeric	31	92.73 %	×
protein_mean	numeric	33	92.73 %	×
sodium_min	numeric	41	23.73 %	×
sodium_max	numeric	38	23.73 %	×
sodium_mean	numeric	107	23.73 %	×
chloride_min	numeric	36	36.94 %	×
chloride_max	numeric	36	36.94 %	×
chloride_mean	numeric	102	36.94 %	×
potassium_min	numeric	32	24.21 %	×
potassium_max	numeric	37	24.21 %	×
potassium_mean	numeric	105	24.21 %	×
bicarbonate_min	numeric	144	44.78 %	×
bicarbonate_max	numeric	135	44.78 %	×
bicarbonate_mean	numeric	260	44.78 %	×
bun_min	numeric	145	73.59 %	×
bun_max	numeric	150	73.59 %	×
bun_mean	numeric	178	73.59 %	×
calcium_min	logical	1	100.00 %	×
calcium_max	logical	1	100.00 %	×
calcium_mean	logical	1	100.00 %	×
magnesium_min	numeric	68	57.99 %	×
magnesium_max	numeric	74	57.99 %	×
magnesium_mean	numeric	66	57.99 %	×
phosphate_min	numeric	110	68.04 %	×
phosphate_max	numeric	122	68.04 %	×
phosphate_mean	numeric	115	68.04 %	×
creatinine_min	numeric	200	23.83 %	×
creatinine_max	numeric	200	23.83 %	×

	Variable class	# unique values	Missing observations	Any problems?
creatinine_mean	numeric	304	23.83 %	×
gfr_min	logical	1	100.00 %	×
gfr_max	logical	1	100.00 %	×
gfr_mean	logical	1	100.00 %	×
glucose_min	numeric	104	32.44 %	×
glucose_max	numeric	166	32.44 %	×
glucose_max_1	numeric	334	32.44 %	×
anion_gap_min	numeric	18	66.60 %	×
anion_gap_min_1	numeric	21	66.60 %	×
anion_gap_mean	numeric	31	66.60 %	×
eos_min	numeric	37	29.28 %	×
eos_max	numeric	44	29.28 %	×
eos_mean	numeric	41	29.28 %	×
lymph_min	numeric	216	24.11 %	×
lymph_max	numeric	225	24.11 %	×
lymph_mean	numeric	224	24.11 %	×
neutrophil_min	numeric	461	24.11 %	×
neutrophil_max	numeric	470	24.11 %	×
neutrophil_mean	numeric	473	24.11 %	×
mono_min	numeric	140	24.11 %	×
mono_max	numeric	152	24.11 %	×
mono_mean	numeric	146	24.11 %	×
baso_min	numeric	15	24.11 %	×
baso_max	numeric	18	24.11 %	×
baso_mean	numeric	16	24.11 %	×
stab_min	numeric	12	98.56 %	×
stab_max	numeric	12	98.56 %	×
stab_mean	numeric	12	98.56 %	×
pt_min	numeric	17	89.95 %	×
pt_max	numeric	17	89.95 %	×
pt_mean	numeric	26	89.95 %	×
ptt_min	numeric	30	65.55 %	×
ptt_max	numeric	39	65.55 %	×
ptt_mean	numeric	67	65.55 %	×
fibrinogen_min	numeric	130	83.25 %	×
fibrinogen_max	numeric	134	83.25 %	×
fibrinogen_mean	numeric	134	83.25 %	×
d_dimer_min	numeric	169	82.11 %	×
d_dimer_max	numeric	170	82.11 %	×
d_dimer_mean	numeric	170	82.11 %	×
alt_min	numeric	105	43.92 %	×
alt_max	numeric	108	43.92 %	×
alt_mean	numeric	157	43.92 %	×
ast_min	numeric	88	76.56 %	×
ast_max	numeric	91	76.56 %	×
ast_mean	numeric	114	76.56 %	×
palc_min	numeric	136	49.57 %	×
palc_max	numeric	140	49.57 %	×
palc_mean	numeric	169	49.57 %	×
ggt_min	numeric	59	91.29 %	×
ggt_max	numeric	59	91.29 %	×
ggt_mean	numeric	60	91.29 %	×
amylase_min	logical	1	100.00 %	×
amylase_max	logical	1	100.00 %	×

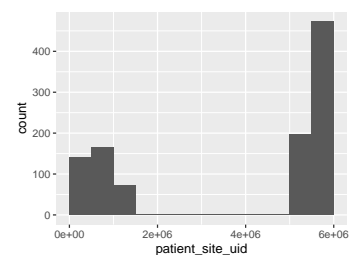
	Variable class	# unique values	Missing observations	Any problems?
amylase_mean	logical	1	100.00 %	×
lipase_min	numeric	96	68.42 %	×
lipase_max	numeric	96	68.42 %	×
lipase_mean	numeric	111	68.42 %	×
bili_tot_min	numeric	41	46.12 %	×
bili_tot_max	numeric	44	46.12 %	×
bili_tot_mean	numeric	72	46.12 %	×
bili_direct_min	numeric	36	96.17 %	×
bili_direct_max	numeric	37	96.17 %	×
bili_direct_mean	numeric	37	96.17 %	×
bili_indirect_min	numeric	35	96.17 %	×
bili_indirect_max	numeric	37	96.17 %	×
bili_indirect_mean	numeric	39	96.17 %	×
lipase_min_1	numeric	96	68.42 %	×
lipase_max_1	numeric	96	68.42 %	×
lipase_mean_1	numeric	111	68.42 %	×
ck_min	numeric	128	84.59 %	×
ck_max	numeric	128	84.59 %	×
ck_mean	numeric	134	84.59 %	×
ckmb_min	numeric	43	91.48 %	×
ckmb_max	numeric	49	91.48 %	×
ckmb_mean	numeric	57	91.48 %	×
ldh_min	numeric	182	75.41 %	×
ldh_max	numeric	187	75.41 %	×
ldh_mean	numeric	186	75.41 %	×
tropot_min	numeric	80	78.37 %	×
tropot_max	numeric	87	78.37 %	×
tropot_mean	numeric	109	78.37 %	×
lactate_min	numeric	30	75.12 %	×
lactate_max	numeric	40	75.12 %	×
lactate_mean	numeric	84	75.12 %	×
svo2sat_min	numeric	90	58.85 %	
svo2sat_max	numeric	86	58.85 %	
svo2sat_max_1	numeric	133	58.85 %	
pao2_min	numeric	68	92.34 %	×
pao2_max	numeric	62	92.34 %	×
pao2_mean	numeric	70	92.34 %	×
pvo2_min	numeric	273	58.76 %	×
pvo2_max	numeric	294	58.76 %	×
pvo2_mean	numeric	305	58.76 %	×
paco2_min	numeric	212	58.76 %	×
paco2_max	numeric	222	58.76 %	×
paco2_mean	numeric	252	58.76 %	×
pvco2_min	numeric	212	58.76 %	×
pvco2_max	numeric	222	58.76 %	×
pvco2_mean	numeric	252	58.76 %	×
tsh_min	numeric	87	90.43 %	×
tsh_max	numeric	87	90.43 %	×
tsh_mean	numeric	86	90.43 %	×
vitd_min	numeric	7	99.43 %	
vitd_max	numeric	7	99.43 %	
vitd_mean	numeric	7	99.43 %	
crp_min	numeric	405	49.19 %	
crp_max	numeric	399	49.19 %	

	Variable class	# unique values	Missing observations	Any problems?
crp_mean	numeric	407	49.19 %	
ferritin_min	numeric	76	92.34 %	×
ferritin_max	numeric	76	92.34 %	×
ferritin_mean	numeric	76	92.34 %	×
bnp_min	numeric	109	88.33 %	×
bnp_max	numeric	109	88.33 %	
bnp_mean	numeric	109	88.33 %	×
weight_min	numeric	215	72.73 %	×
weight_max	numeric	212	72.73 %	×
weight_mean	numeric	222	72.73 %	×
sbp_min	numeric	107	17.32 %	×
sbp_max	numeric	113	17.32 %	×
sbp_mean	numeric	518	17.32 %	×
dbp_min	numeric	70	17.32 %	×
dbp_max	numeric	79	17.32 %	×
dbp_mean	numeric	420	17.32 %	×
temp_min	numeric	42	20.67 %	×
temp_max	numeric	52	20.67 %	×
temp_mean	numeric	210	20.67 %	×
so2_min	numeric	42	12.15 %	×
so2_max	numeric	17	12.15 %	×
so2_mean	numeric	298	12.15 %	×
rr_min	numeric	21	17.70 %	×
rr_max	numeric	34	17.70 %	×
rr_mean	numeric	195	17.70 %	×
flow_min	numeric	15	72.06 %	×
flow_max	numeric	16	72.06 %	×
flow_mean	numeric	121	72.06 %	×
fio2_min	numeric	39	46.99 %	
fio2_max	numeric	39	46.99 %	
fio2_mean	numeric	195	46.99 %	
mv	numeric	2	0.00 %	
icu	numeric	2	0.00 %	

Variable list

patient_site_uid

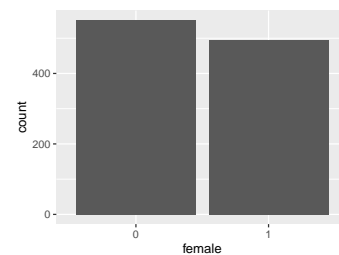
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	1009
Median	5351949
1st and 3rd quartiles	847996; 5637308
Min. and max.	720; 5683923



female

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

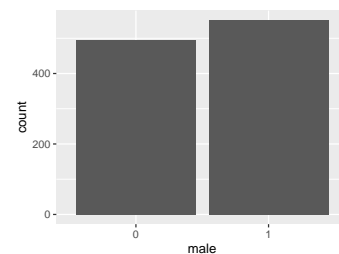
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	“0”
Reference category	0



male

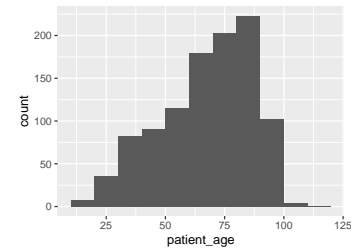
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

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	54; 84
Min. and max.	12; 120

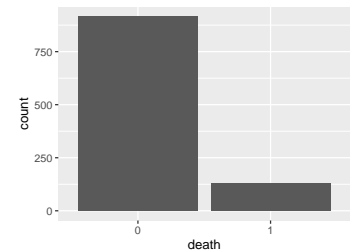


- Note that the following possible outlier values were detected: "120".

death

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

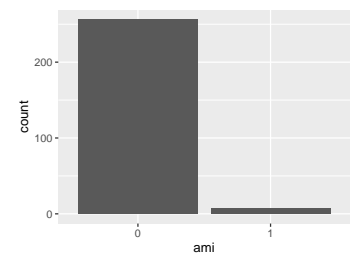
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



ami

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

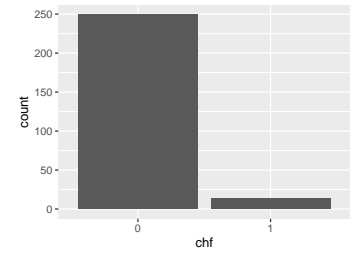
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



chf

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

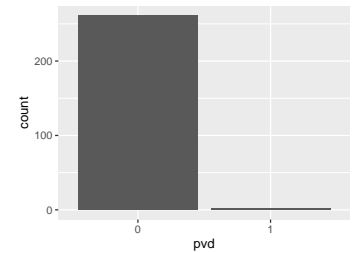
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

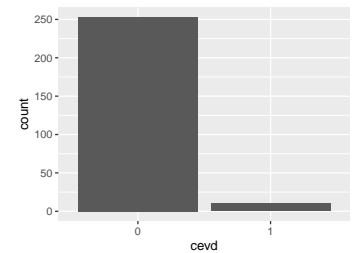


- Note that the following levels have at most five observations: "1".

cevd

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

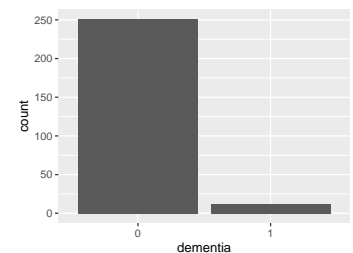
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



dementia

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

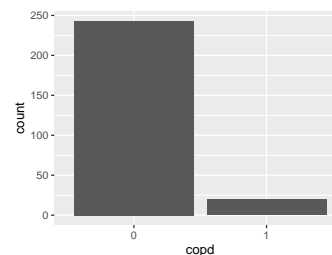
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



copd

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

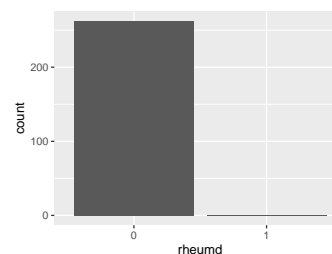
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

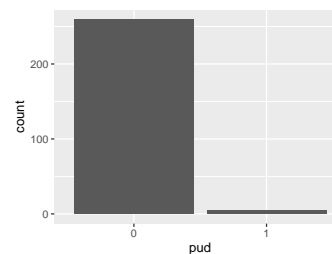


- Note that the following levels have at most five observations: "1".

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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

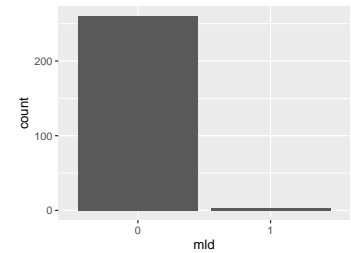


- Note that the following levels have at most five observations: "1".

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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

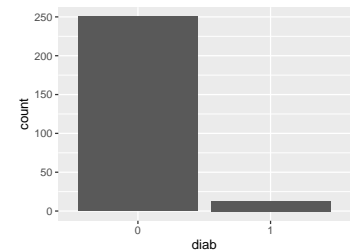


- Note that the following levels have at most five observations: "1".

diab

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

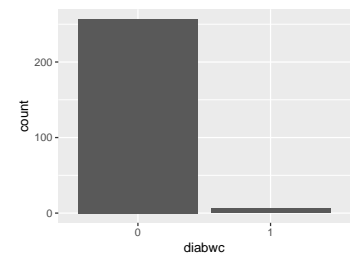
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



diabwc

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

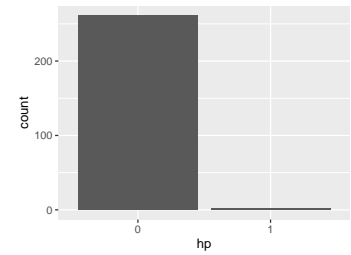
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

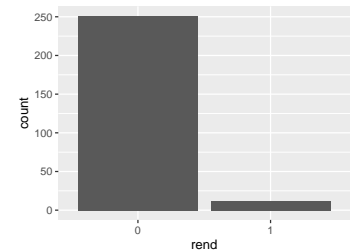


- Note that the following levels have at most five observations: "1".

rend

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

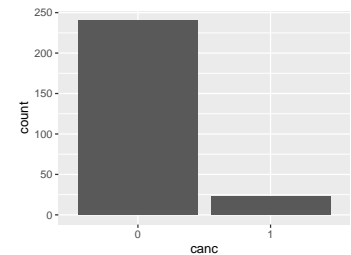
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



canc

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

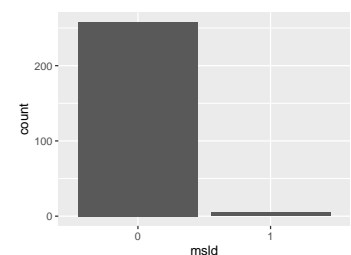
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0

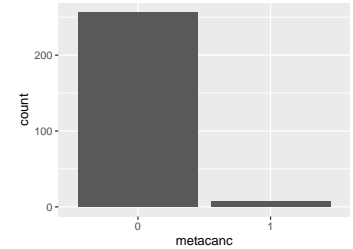


- Note that the following levels have at most five observations: "1".

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.	782 (74.83 %)
Number of unique values	2
Mode	"0"
Reference category	0



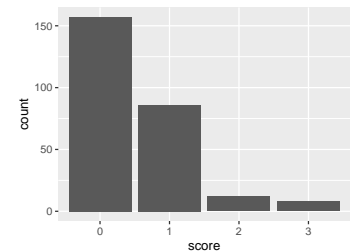
aids

- The variable only takes one (non-missing) value: "0". The variable contains 74.83 % missing observations.

score

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

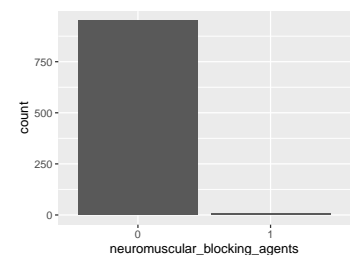
Feature	Result
Variable type	numeric
Number of missing obs.	782 (74.83 %)
Number of unique values	4
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

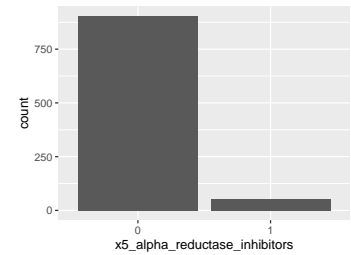


- Note that the following levels have at most five observations: "1".

x5_alpha_reductase_inhibitors

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

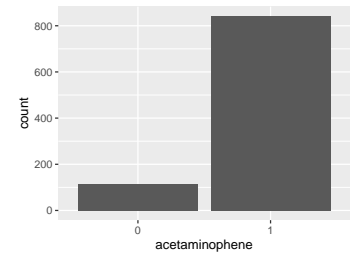
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



acetaminophene

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

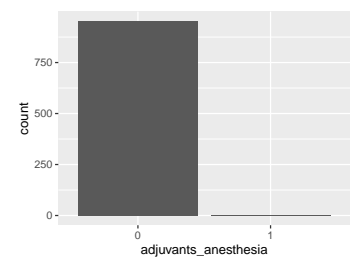
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"1"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

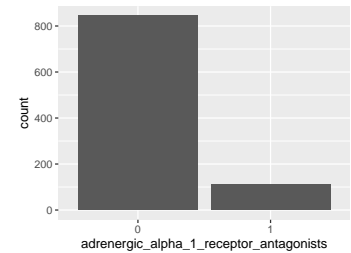


- Note that the following levels have at most five observations: "1".

adrenergic_alpha_1_receptor_antagonists

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

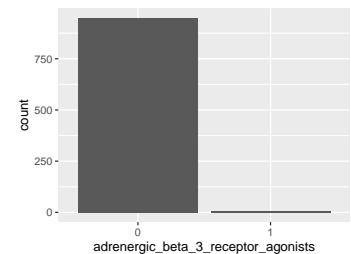
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic_beta_3_receptor_agonists

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

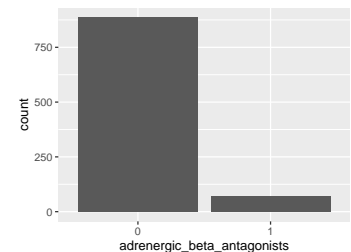
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



adrenergic_beta_antagonists

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

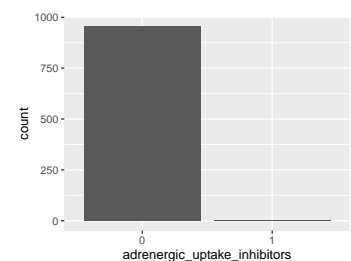
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

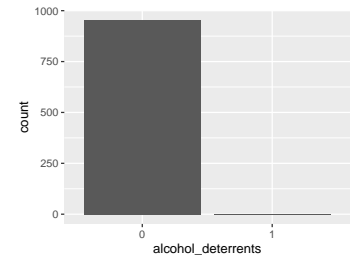


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

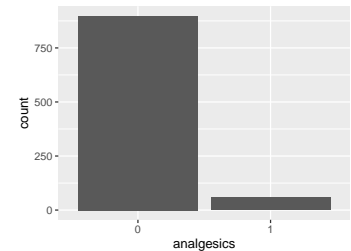


- Note that the following levels have at most five observations: "1".

analgesics

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

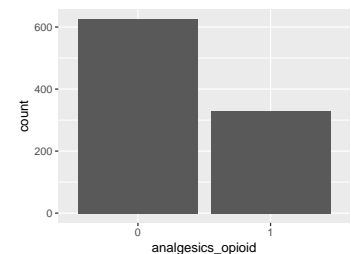
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



analgesics_opioid

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

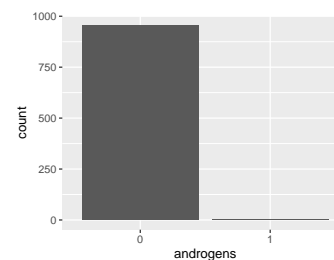
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

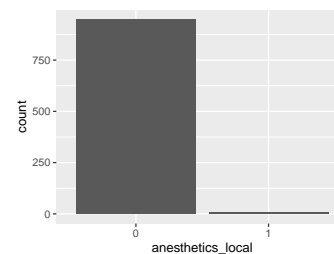


- Note that the following levels have at most five observations: "1".

anesthetics_local

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

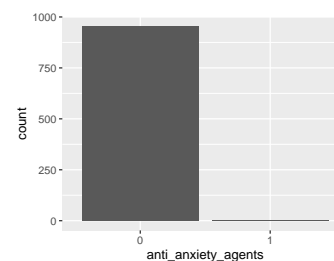
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

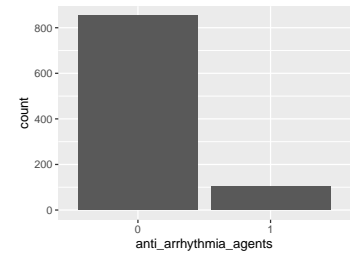


- Note that the following levels have at most five observations: "1".

anti_arrhythmia_agents

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

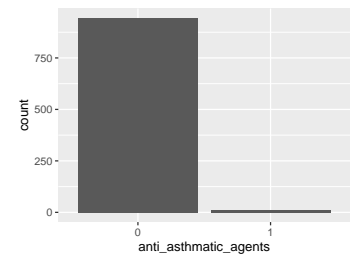
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_asthmatic_agents

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

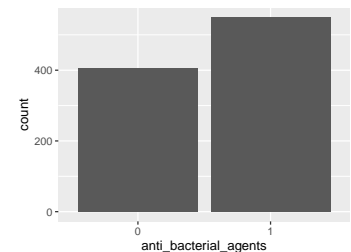
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_bacterial_agents

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

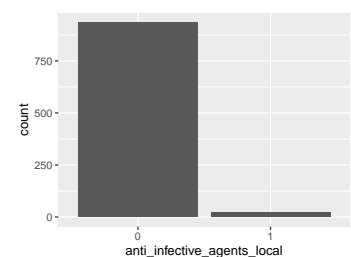
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"1"
Reference category	0



anti_infective_agents_local

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

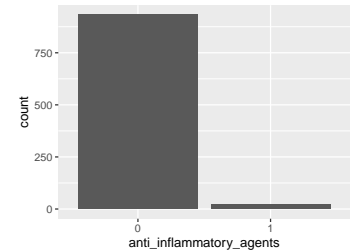
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents

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

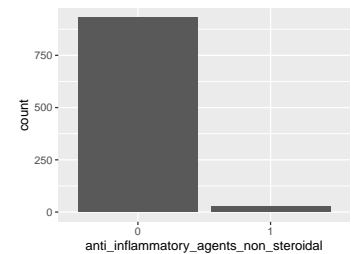
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_inflammatory_agents_non_steroidal

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

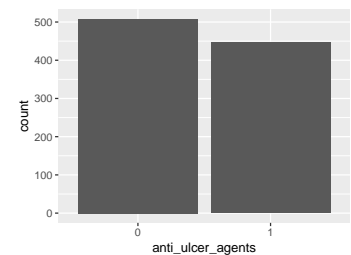
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anti_ulcer_agents

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

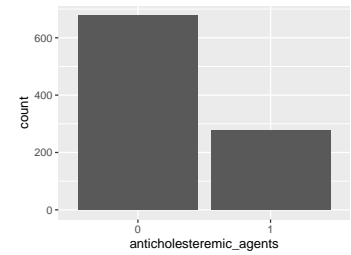
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anticholesteremic_agents

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

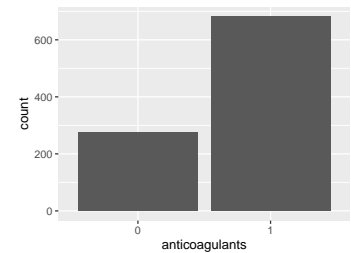
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



anticoagulants

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

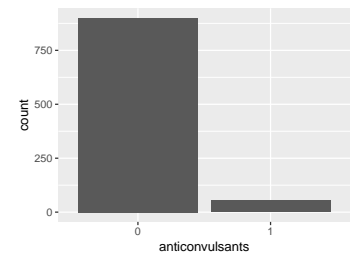
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"1"
Reference category	0



anticonvulsants

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

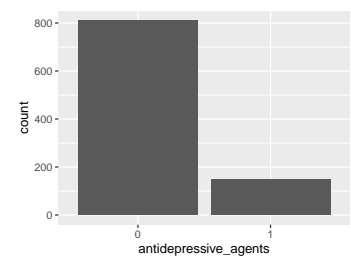
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



antidepressive_agents

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

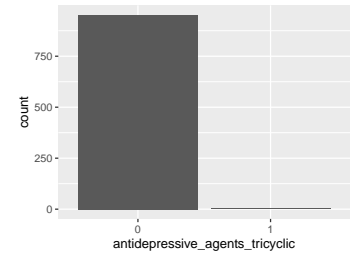
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

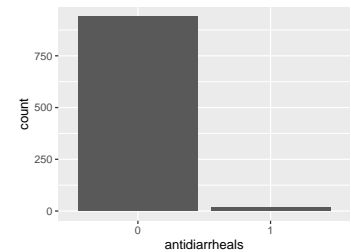


- Note that the following levels have at most five observations: "1".

antidiarrheals

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

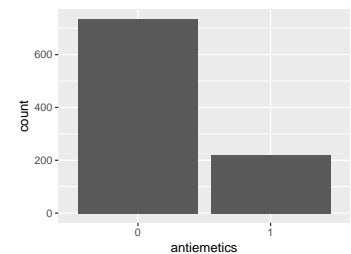
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



antiemetics

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

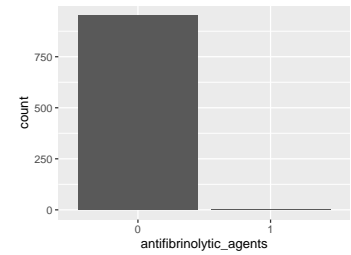
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

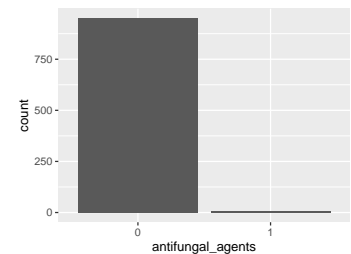


- Note that the following levels have at most five observations: "1".

antifungal_agents

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

Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

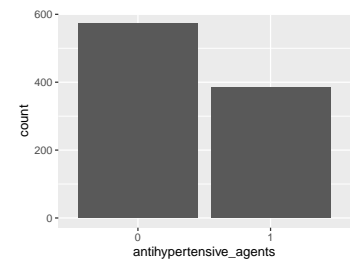


- Note that the following levels have at most five observations: "1".

antihypertensive_agents

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

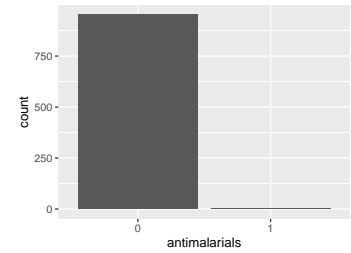
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

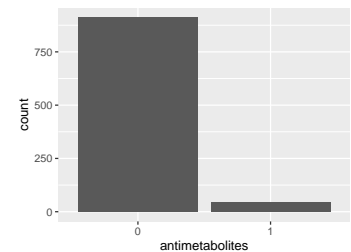


- Note that the following levels have at most five observations: "1".

antimetabolites

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

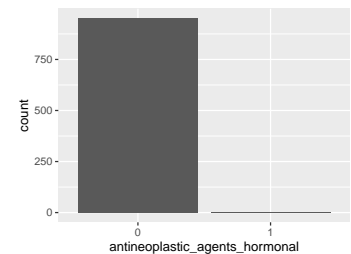
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

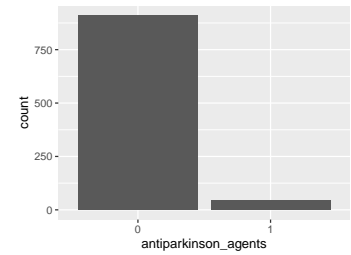


- Note that the following levels have at most five observations: "1".

antiparkinson_agents

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

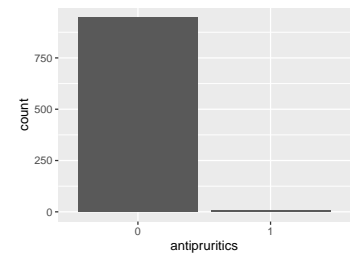
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



antipruritics

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

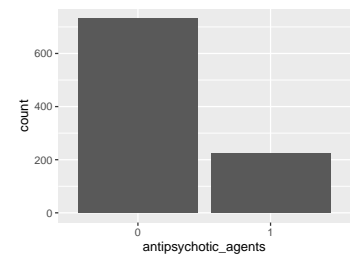
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



antipsychotic_agents

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

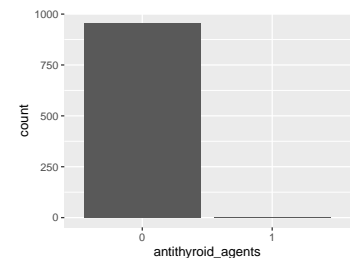
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

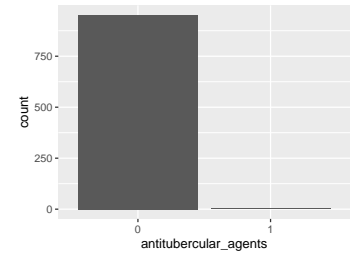


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

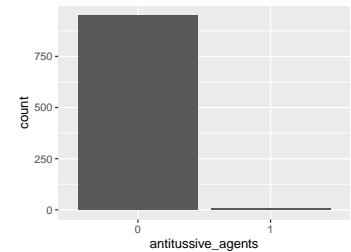


- Note that the following levels have at most five observations: "1".

antitussive_agents

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

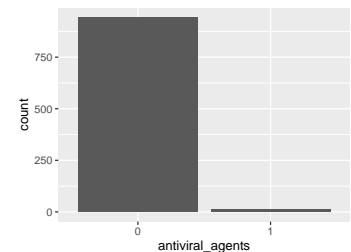
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
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.

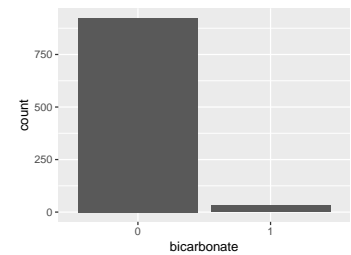
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



bicarbonate

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

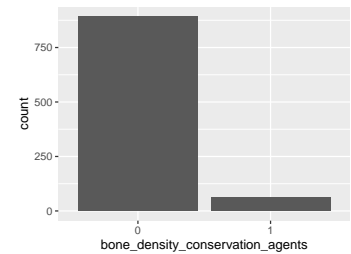
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



bone_density_conservation_agents

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

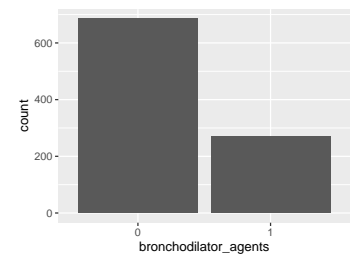
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



bronchodilator_agents

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

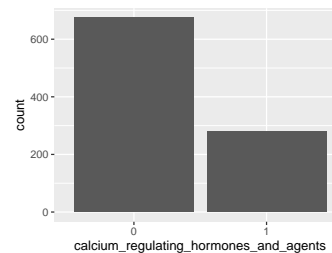
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



calcium_regulating_hormones_and_agents

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

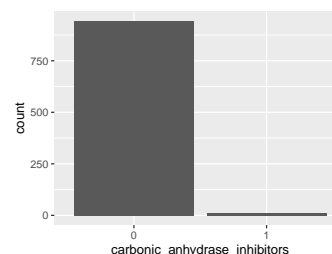
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



carbonic_anhydrase_inhibitors

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

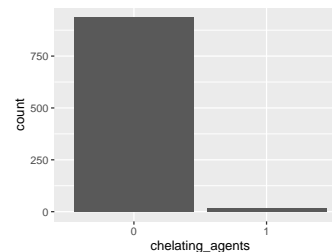
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



chelating_agents

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

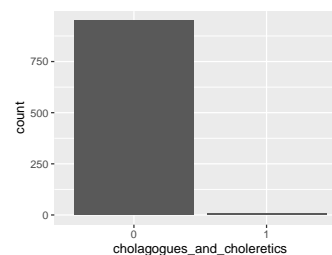
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



cholagogues_and_choleretics

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

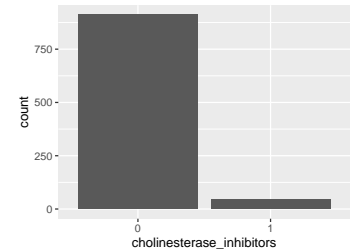
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



cholinesterase_inhibitors

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

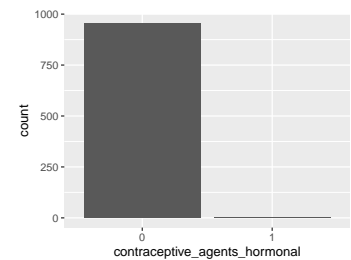
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

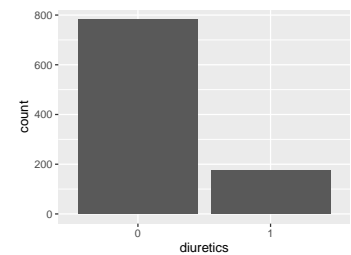


- Note that the following levels have at most five observations: "1".

diuretics

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

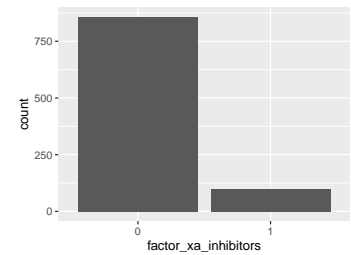
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



factor_xa_inhibitors

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

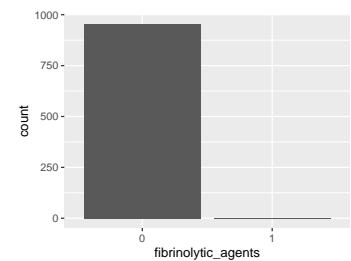
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

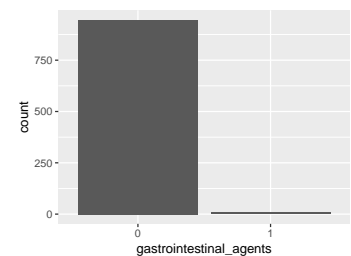


- Note that the following levels have at most five observations: "1".

gastrointestinal_agents

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

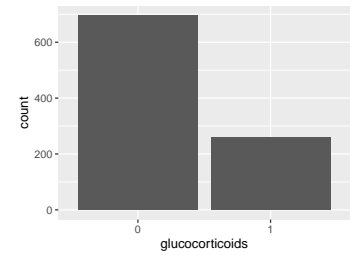
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



glucocorticoids

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

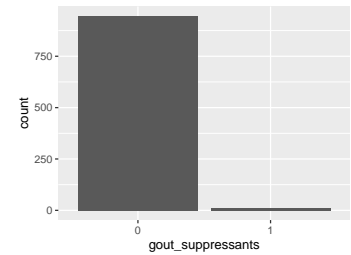
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



gout_suppressants

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

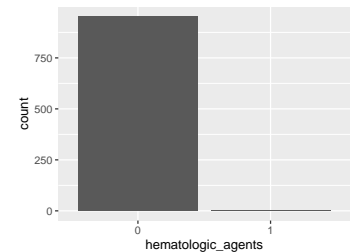
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

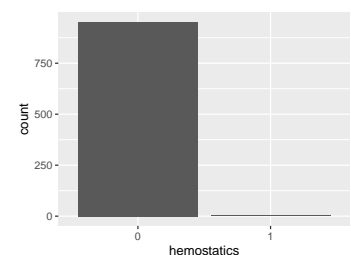


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

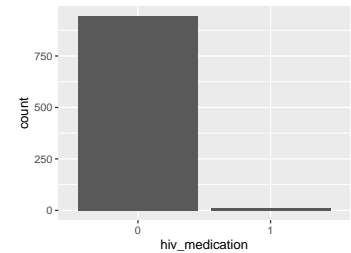


- Note that the following levels have at most five observations: "1".

hiv_medication

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

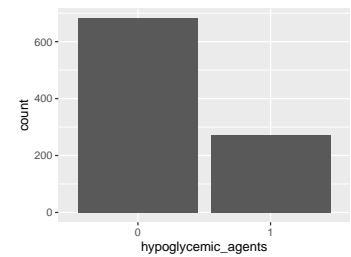
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



hypoglycemic_agents

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

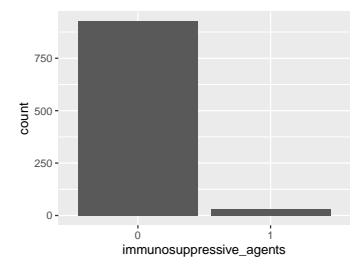
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



immunosuppressive_agents

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

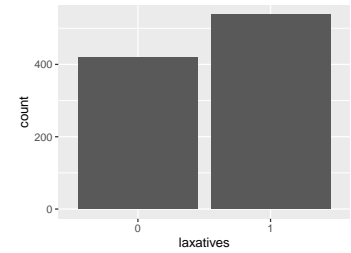
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



laxatives

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

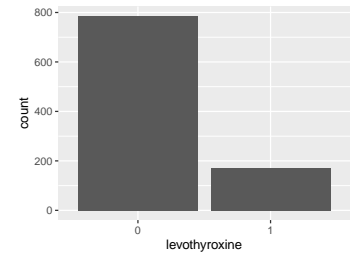
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"1"
Reference category	0



levothyroxine

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

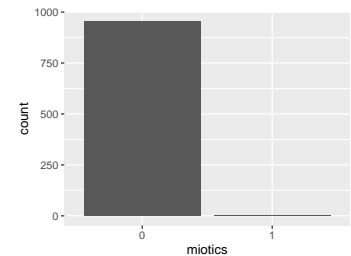
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

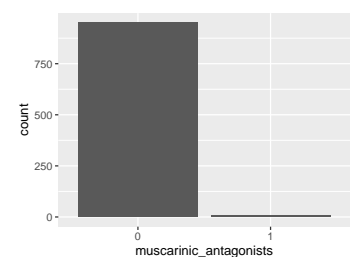


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

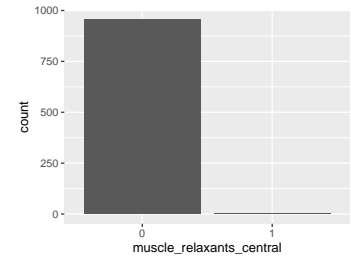


- Note that the following levels have at most five observations: "1".

muscle_relaxants_central

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

Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

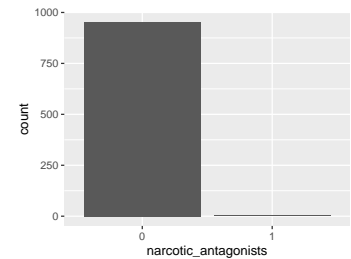


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

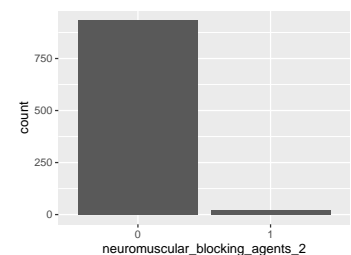


- Note that the following levels have at most five observations: "1".

neuromuscular_blocking_agents_2

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

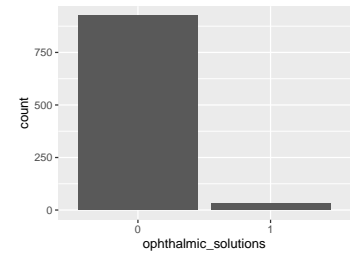
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



ophthalmic_solutions

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

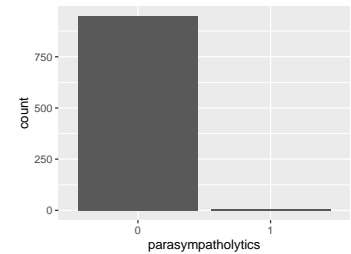
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



parasympatholytics

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

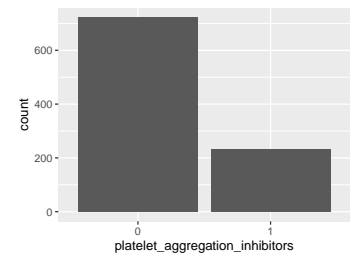
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



platelet_aggregation_inhibitors

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

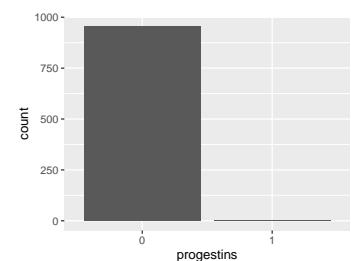
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

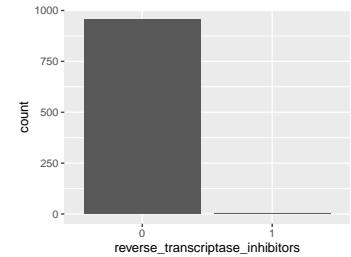


- Note that the following levels have at most five observations: "1".

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.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0

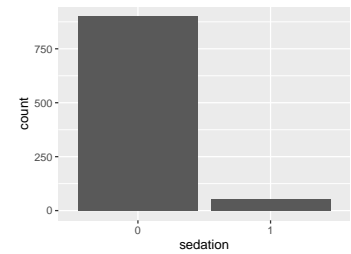


- Note that the following levels have at most five observations: "1".

sedation

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

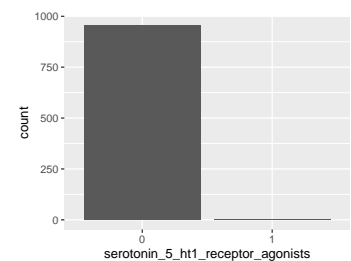
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



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.	90 (8.61 %)
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.

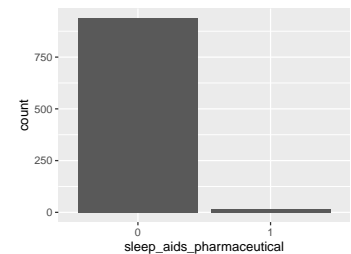
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



sleep_aids_pharmaceutical

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

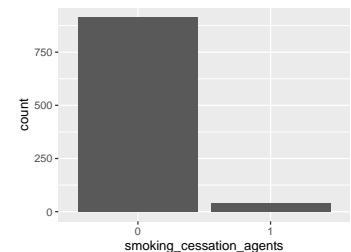
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



smoking_cessation_agents

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

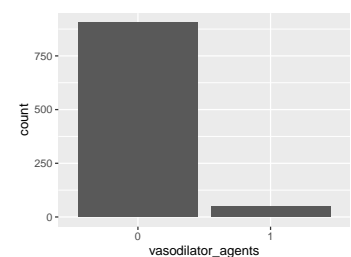
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasodilator_agents

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

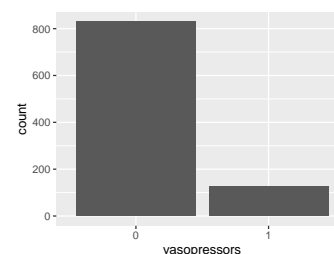
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



vasopressors

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

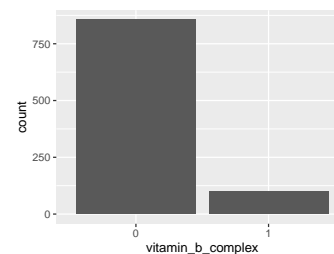
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamin_b_complex

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

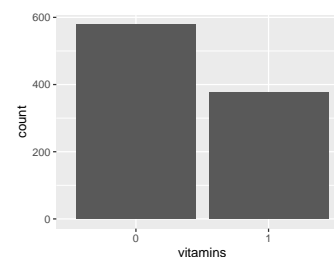
Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



vitamins

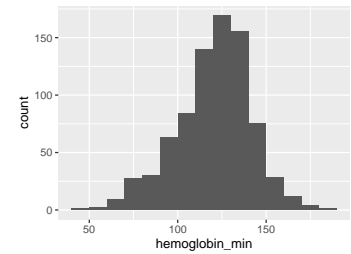
- Note that this variable is treated as a factor variable below, as it only takes a few unique values.

Feature	Result
Variable type	numeric
Number of missing obs.	90 (8.61 %)
Number of unique values	2
Mode	"0"
Reference category	0



hemoglobin_min

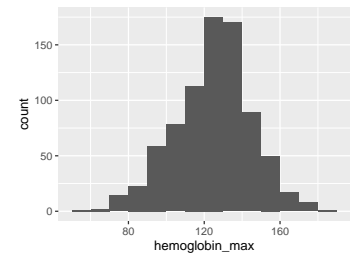
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	102
Median	123
1st and 3rd quartiles	109; 135
Min. and max.	41; 185



- Note that the following possible outlier values were detected: "41", "53", "161", "162", "163", "165", "166", "170", "172", "173" (3 additional values omitted).

hemoglobin_max

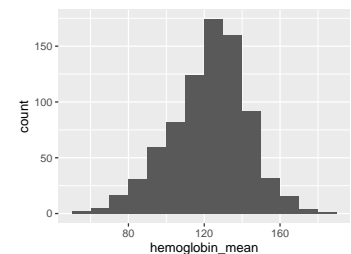
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	101
Median	127
1st and 3rd quartiles	112; 138
Min. and max.	59; 185



- Note that the following possible outlier values were detected: "162", "163", "165", "167", "169", "170", "171", "172", "173", "176" (2 additional values omitted).

hemoglobin_mean

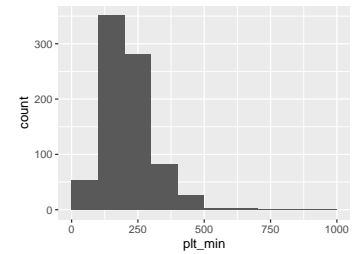
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	211
Median	125.75
1st and 3rd quartiles	111; 137
Min. and max.	56.4; 185



- Note that the following possible outlier values were detected: "157.5", "158", "160", "161", "161.5", "162", "162.5", "163", "164.33", "166" (8 additional values omitted).

plt_min

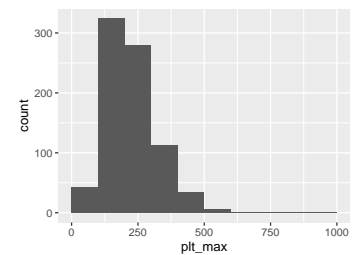
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	303
Median	199.5
1st and 3rd quartiles	148; 263
Min. and max.	21; 941



- Note that the following possible outlier values were detected: "21", "24", "26", "32", "34", "36", "37", "512", "519", "526" (3 additional values omitted).

plt_max

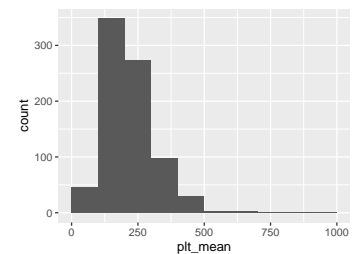
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	319
Median	207
1st and 3rd quartiles	157; 279
Min. and max.	24; 941



- Note that the following possible outlier values were detected: "24", "26", "36", "43", "44", "50", "55", "56", "58", "59" (8 additional values omitted).

plt_mean

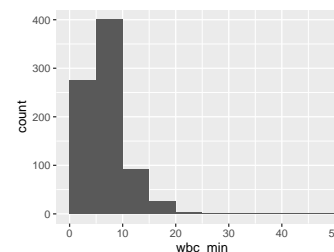
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	427
Median	203
1st and 3rd quartiles	152; 271.38
Min. and max.	23.6; 941



- Note that the following possible outlier values were detected: "23.6", "24", "31", "37.5", "40", "43.5", "46.5", "50", "50.63", "54" (8 additional values omitted).

wbc_min

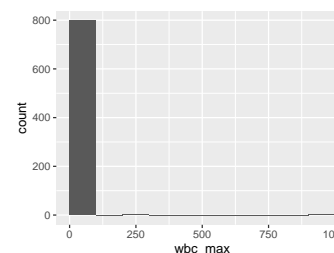
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	152
Median	6.2
1st and 3rd quartiles	4.5; 8.3
Min. and max.	0.7; 46.1



- Note that the following possible outlier values were detected: "0.7", "1", "1.2", "1.3", "1.5", "1.6", "1.9", "18.7", "18.8", "19.8" (7 additional values omitted).

wbc_max

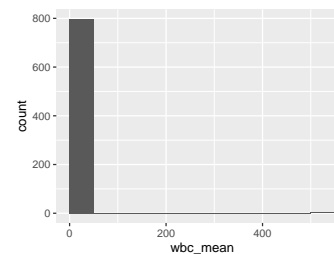
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	174
Median	6.7
1st and 3rd quartiles	5.07; 9.8
Min. and max.	0.8; 1000



- Note that the following possible outlier values were detected: "0.8", "1.3", "1.6", "1.7", "1.8", "2.1", "2.2", "2.4", "2.5", "2.6" (10 additional values omitted).

wbc_mean

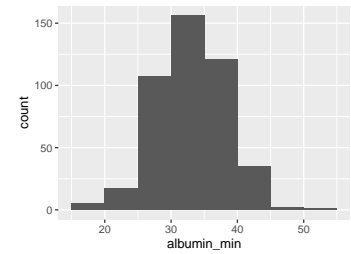
Feature	Result
Variable type	numeric
Number of missing obs.	245 (23.44 %)
Number of unique values	274
Median	6.5
1st and 3rd quartiles	4.75; 9.1
Min. and max.	0.75; 502.5



- Note that the following possible outlier values were detected: "0.75", "1.3", "1.46", "1.6", "1.7", "1.95", "2.05", "2.1", "2.2", "2.3" (7 additional values omitted).

albumin_min

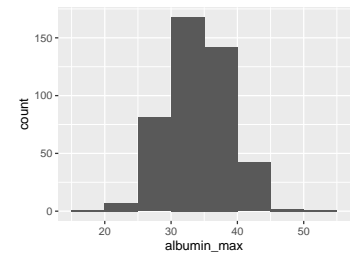
Feature	Result
Variable type	numeric
Number of missing obs.	601 (57.51 %)
Number of unique values	34
Median	34
1st and 3rd quartiles	30; 37
Min. and max.	18; 54



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

albumin_max

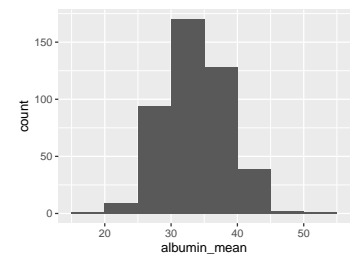
Feature	Result
Variable type	numeric
Number of missing obs.	601 (57.51 %)
Number of unique values	30
Median	34
1st and 3rd quartiles	31; 38
Min. and max.	18; 54



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

albumin_mean

Feature	Result
Variable type	numeric
Number of missing obs.	601 (57.51 %)
Number of unique values	74
Median	34
1st and 3rd quartiles	30.69; 37
Min. and max.	18; 54

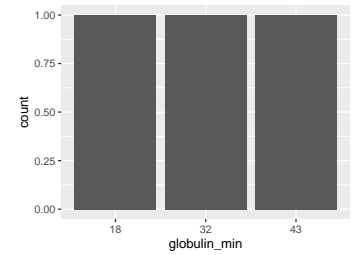


- Note that the following possible outlier values were detected: "18", "20.2", "48", "54".

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.	1042 (99.71 %)
Number of unique values	3
Mode	"18"
Reference category	18

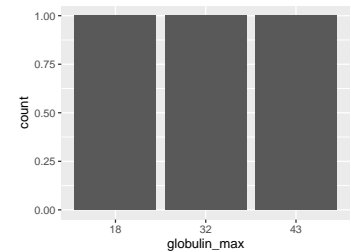


- Note that the following levels have at most five observations: "18", "32", "43".

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.	1042 (99.71 %)
Number of unique values	3
Mode	"18"
Reference category	18

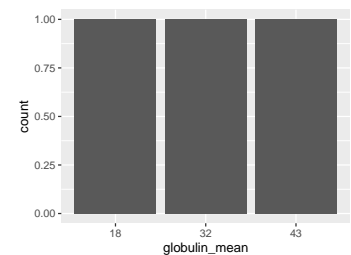


- Note that the following levels have at most five observations: "18", "32", "43".

globulin__mean

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

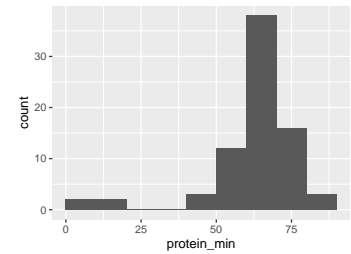
Feature	Result
Variable type	numeric
Number of missing obs.	1042 (99.71 %)
Number of unique values	3
Mode	"18"
Reference category	18



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

protein__min

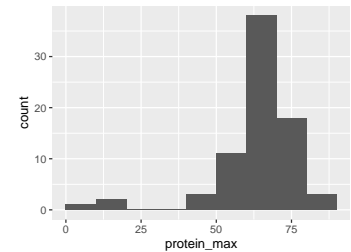
Feature	Result
Variable type	numeric
Number of missing obs.	969 (92.73 %)
Number of unique values	32
Median	66
1st and 3rd quartiles	60.75; 70.25
Min. and max.	0.47; 86



- Note that the following possible outlier values were detected: "0.47", "0.52", "20", "82", "84", "86".

protein__max

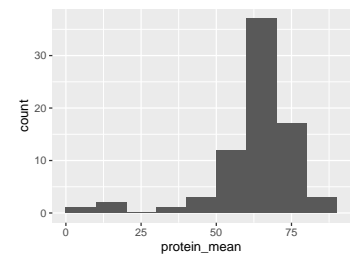
Feature	Result
Variable type	numeric
Number of missing obs.	969 (92.73 %)
Number of unique values	30
Median	66
1st and 3rd quartiles	61; 71
Min. and max.	0.47; 86



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

protein__mean

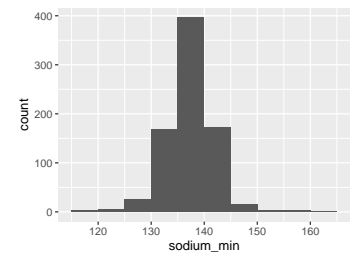
Feature	Result
Variable type	numeric
Number of missing obs.	969 (92.73 %)
Number of unique values	32
Median	66
1st and 3rd quartiles	60.75; 71
Min. and max.	0.47; 86



- Note that the following possible outlier values were detected: "0.47", "20", "39.26", "84", "86".

sodium__min

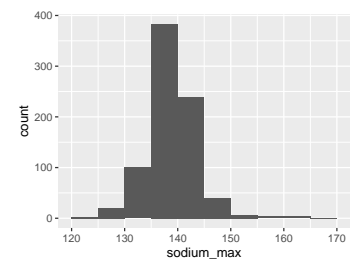
Feature	Result
Variable type	numeric
Number of missing obs.	248 (23.73 %)
Number of unique values	40
Median	138
1st and 3rd quartiles	135; 140
Min. and max.	115; 163



- Note that the following possible outlier values were detected: "115", "117", "120", "121", "122", "123", "125", "126", "127", "148" (10 additional values omitted).

sodium__max

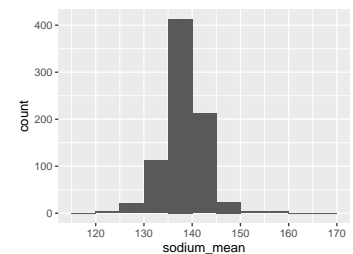
Feature	Result
Variable type	numeric
Number of missing obs.	248 (23.73 %)
Number of unique values	37
Median	139
1st and 3rd quartiles	137; 142
Min. and max.	122; 168



- Note that the following possible outlier values were detected: "122", "125", "126", "127", "128", "129", "150", "151", "152", "153" (7 additional values omitted).

sodium__mean

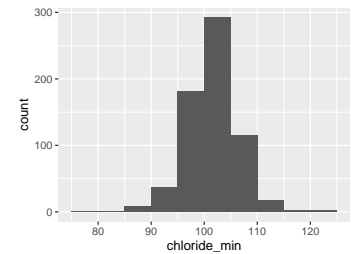
Feature	Result
Variable type	numeric
Number of missing obs.	248 (23.73 %)
Number of unique values	106
Median	138.5
1st and 3rd quartiles	136.33; 141
Min. and max.	119.67; 165.33



- Note that the following possible outlier values were detected: "119.67", "123.5", "123.67", "124", "125", "126", "126.38", "126.67", "127", "127.33" (15 additional values omitted).

chloride_min

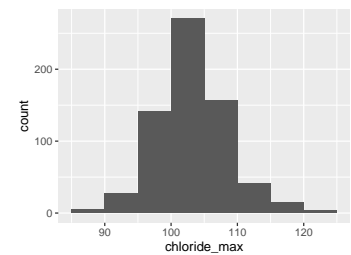
Feature	Result
Variable type	numeric
Number of missing obs.	386 (36.94 %)
Number of unique values	35
Median	102
1st and 3rd quartiles	99; 105
Min. and max.	76; 124



- Note that the following possible outlier values were detected: "76", "84", "87", "88", "89", "115", "116", "117", "123", "124".

chloride_max

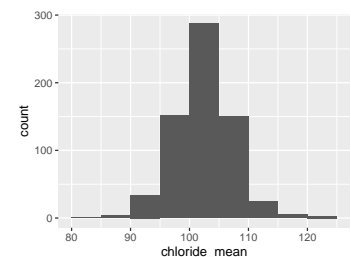
Feature	Result
Variable type	numeric
Number of missing obs.	386 (36.94 %)
Number of unique values	35
Median	103
1st and 3rd quartiles	100; 106.5
Min. and max.	88; 124



- Note that the following possible outlier values were detected: "88", "89", "90", "91", "119", "120", "122", "123", "124".

chloride_mean

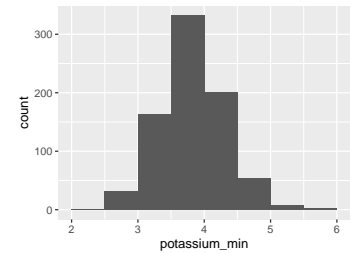
Feature	Result
Variable type	numeric
Number of missing obs.	386 (36.94 %)
Number of unique values	101
Median	103
1st and 3rd quartiles	100; 106
Min. and max.	84.17; 124



- Note that the following possible outlier values were detected: "84.17", "88", "89", "90", "90.67", "116", "116.25", "116.5", "117", "117.13" (3 additional values omitted).

potassium_min

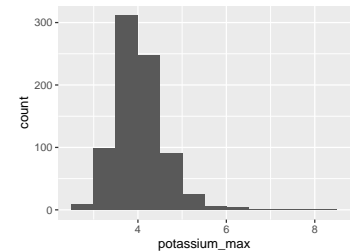
Feature	Result
Variable type	numeric
Number of missing obs.	253 (24.21 %)
Number of unique values	31
Median	3.9
1st and 3rd quartiles	3.6; 4.2
Min. and max.	2.4; 5.6



- Note that the following possible outlier values were detected: "2.4", "2.6", "5.2", "5.3", "5.4", "5.6".

potassium_max

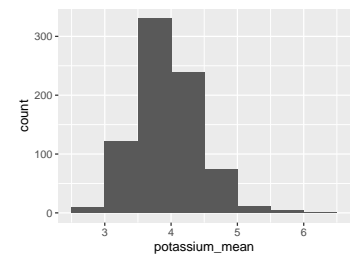
Feature	Result
Variable type	numeric
Number of missing obs.	253 (24.21 %)
Number of unique values	36
Median	4
1st and 3rd quartiles	3.8; 4.4
Min. and max.	2.7; 8.2



- Note that the following possible outlier values were detected: "2.7", "2.9", "3", "3.1", "3.2", "3.3", "6", "6.2", "6.3", "6.4" (2 additional values omitted).

potassium_mean

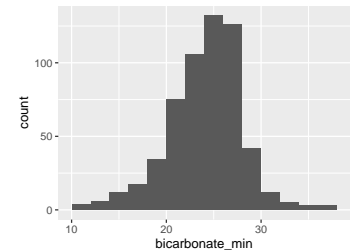
Feature	Result
Variable type	numeric
Number of missing obs.	253 (24.21 %)
Number of unique values	104
Median	3.93
1st and 3rd quartiles	3.67; 4.25
Min. and max.	2.7; 6.02



- Note that the following possible outlier values were detected: "2.7", "2.9", "2.95", "3", "3.05", "3.07", "3.1", "3.15", "5.55", "5.6" (4 additional values omitted).

bicarbonate_min

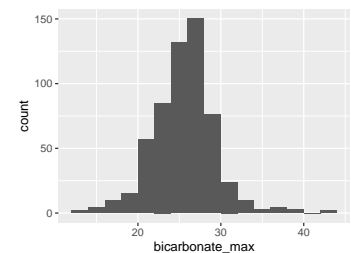
Feature	Result
Variable type	numeric
Number of missing obs.	468 (44.78 %)
Number of unique values	143
Median	24.9
1st and 3rd quartiles	22; 27
Min. and max.	10.4; 37.7



- Note that the following possible outlier values were detected: "30.7", "31", "31.9", "32.1", "32.4", "32.5", "32.9", "33.4", "34.3", "34.4" (3 additional values omitted).

bicarbonate_max

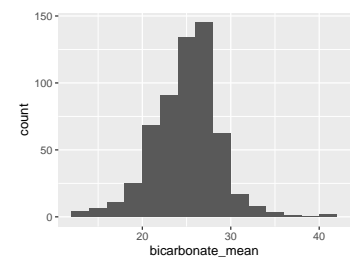
Feature	Result
Variable type	numeric
Number of missing obs.	468 (44.78 %)
Number of unique values	134
Median	26
1st and 3rd quartiles	23.5; 28
Min. and max.	13; 42.8



- Note that the following possible outlier values were detected: "13", "14", "32.9", "33.4", "34.3", "34.4", "35.3", "36.2", "37.2", "38" (3 additional values omitted).

bicarbonate_mean

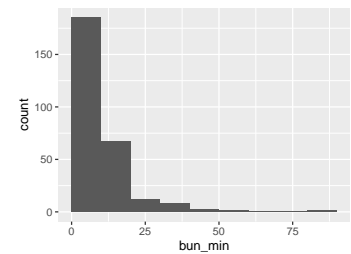
Feature	Result
Variable type	numeric
Number of missing obs.	468 (44.78 %)
Number of unique values	259
Median	25.47
1st and 3rd quartiles	23; 27.3
Min. and max.	12; 40.21



- Note that the following possible outlier values were detected: "12", "30.85", "30.97", "31", "31.07", "31.25", "31.6", "31.9", "32.4", "32.5" (10 additional values omitted).

bun__min

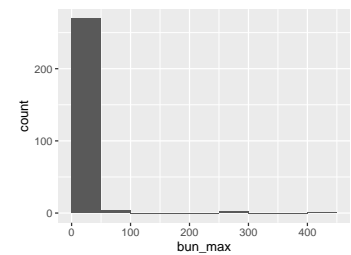
Feature	Result
Variable type	numeric
Number of missing obs.	769 (73.59 %)
Number of unique values	144
Median	7.3
1st and 3rd quartiles	4.88; 12.72
Min. and max.	0.8; 90



- Note that the following possible outlier values were detected: "0.8", "1.2", "1.3", "1.6", "1.7", "2", "2.4", "2.5", "90".

bun__max

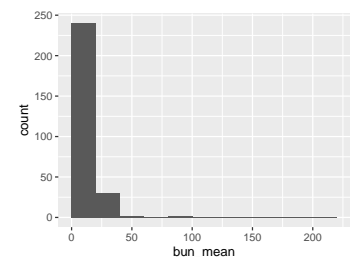
Feature	Result
Variable type	numeric
Number of missing obs.	769 (73.59 %)
Number of unique values	149
Median	7.8
1st and 3rd quartiles	5.2; 14.22
Min. and max.	0.8; 417



- Note that the following possible outlier values were detected: "0.8", "1.2", "1.3", "2", "2.4", "2.5", "2.6", "2.8", "69", "90" (3 additional values omitted).

bun__mean

Feature	Result
Variable type	numeric
Number of missing obs.	769 (73.59 %)
Number of unique values	177
Median	7.5
1st and 3rd quartiles	5.1; 13.76
Min. and max.	0.8; 211.8



- Note that the following possible outlier values were detected: "0.8", "1.2", "1.3", "2", "2.03", "2.25", "2.3", "2.4", "2.5", "2.57" (7 additional values omitted).

calcium_min

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

calcium_max

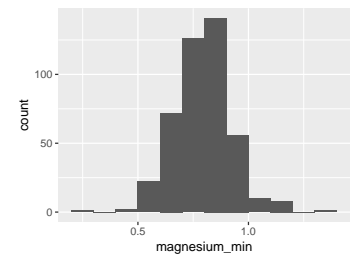
- The variable only takes one value: "NA".

calcium_mean

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

magnesium_min

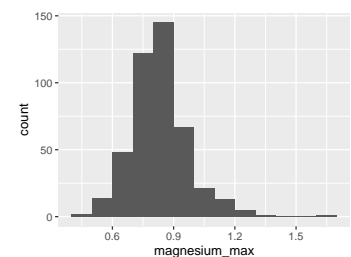
Feature	Result
Variable type	numeric
Number of missing obs.	606 (57.99 %)
Number of unique values	67
Median	0.8
1st and 3rd quartiles	0.71; 0.88
Min. and max.	0.23; 1.38



- Note that the following possible outlier values were detected: "0.23", "1.08", "1.09", "1.1", "1.11", "1.12", "1.16", "1.18", "1.19", "1.2" (1 additional values omitted).

magnesium_max

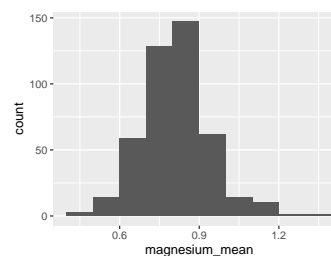
Feature	Result
Variable type	numeric
Number of missing obs.	606 (57.99 %)
Number of unique values	73
Median	0.83
1st and 3rd quartiles	0.74; 0.9
Min. and max.	0.42; 1.67



- Note that the following possible outlier values were detected: "0.42", "0.5", "1.15", "1.16", "1.17", "1.18", "1.19", "1.2", "1.21", "1.27" (3 additional values omitted).

magnesium_mean

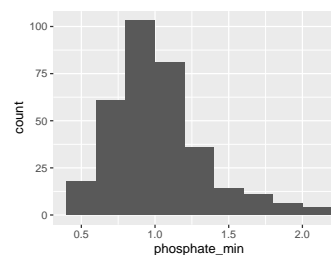
Feature	Result
Variable type	numeric
Number of missing obs.	606 (57.99 %)
Number of unique values	65
Median	0.81
1st and 3rd quartiles	0.73; 0.89
Min. and max.	0.42; 1.38



- Note that the following possible outlier values were detected: "0.42", "0.44", "1.14", "1.16", "1.18", "1.19", "1.2", "1.24", "1.38".

phosphate_min

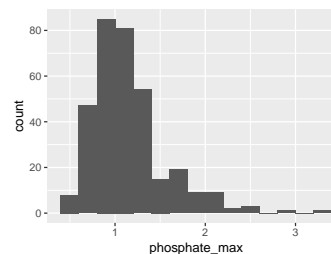
Feature	Result
Variable type	numeric
Number of missing obs.	711 (68.04 %)
Number of unique values	109
Median	0.97
1st and 3rd quartiles	0.81; 1.17
Min. and max.	0.47; 2.13



- Note that the following possible outlier values were detected: "0.47", "0.49", "0.51", "0.52", "0.53", "2.13".

phosphate_max

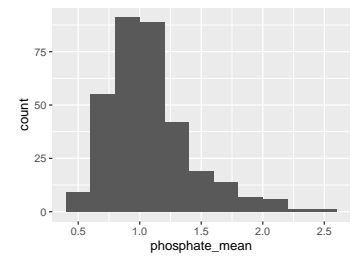
Feature	Result
Variable type	numeric
Number of missing obs.	711 (68.04 %)
Number of unique values	121
Median	1.06
1st and 3rd quartiles	0.89; 1.29
Min. and max.	0.47; 3.27



- Note that the following possible outlier values were detected: "0.47", "0.51", "0.52", "0.53", "0.56", "0.58", "0.63", "0.64", "0.65", "2.82" (1 additional values omitted).

phosphate__mean

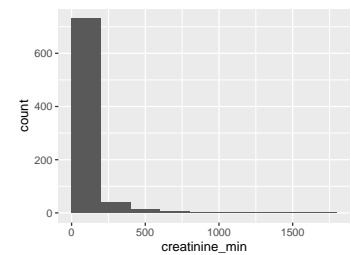
Feature	Result
Variable type	numeric
Number of missing obs.	711 (68.04 %)
Number of unique values	114
Median	1.02
1st and 3rd quartiles	0.85; 1.23
Min. and max.	0.47; 2.59



- Note that the following possible outlier values were detected: "0.47", "0.51", "0.52", "0.53", "2.2", "2.22", "2.59".

creatinine__min

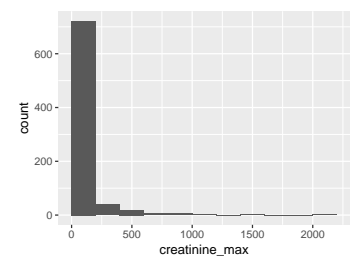
Feature	Result
Variable type	numeric
Number of missing obs.	249 (23.83 %)
Number of unique values	199
Median	74
1st and 3rd quartiles	58; 102
Min. and max.	20; 1762



- Note that the following possible outlier values were detected: "20", "23", "24", "26", "27", "29", "30", "32", "33", "34" (45 additional values omitted).

creatinine__max

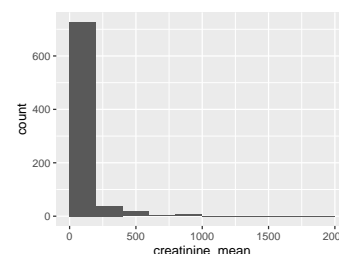
Feature	Result
Variable type	numeric
Number of missing obs.	249 (23.83 %)
Number of unique values	199
Median	79
1st and 3rd quartiles	60.75; 109
Min. and max.	20; 2094



- Note that the following possible outlier values were detected: "20", "27", "29", "32", "34", "35", "36", "37", "38", "39" (43 additional values omitted).

creatinine__mean

Feature	Result
Variable type	numeric
Number of missing obs.	249 (23.83 %)
Number of unique values	303
Median	76
1st and 3rd quartiles	59.5; 103.62
Min. and max.	20; 1952.25



- Note that the following possible outlier values were detected: "20", "26.5", "27", "29", "31", "32", "34.33", "34.5", "35", "36" (55 additional values omitted).

gfr__min

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

gfr__max

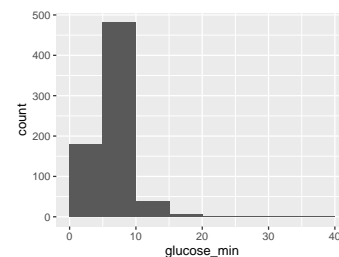
- The variable only takes one value: "NA".

gfr__mean

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

glucose__min

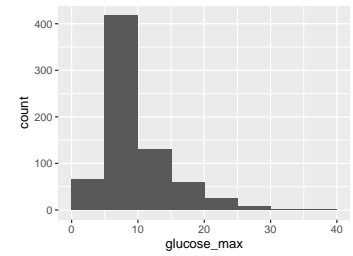
Feature	Result
Variable type	numeric
Number of missing obs.	339 (32.44 %)
Number of unique values	103
Median	5.8
1st and 3rd quartiles	5; 6.9
Min. and max.	1.9; 35.8



- Note that the following possible outlier values were detected: "1.9", "2.2", "2.4", "2.5", "2.7", "2.8", "2.9", "3", "3.1", "3.2" (19 additional values omitted).

glucose__max

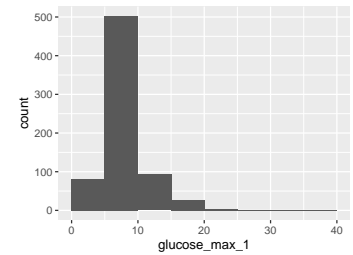
Feature	Result
Variable type	numeric
Number of missing obs.	339 (32.44 %)
Number of unique values	165
Median	7.8
1st and 3rd quartiles	5.9; 11.4
Min. and max.	3.7; 35.8



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

glucose__max__1

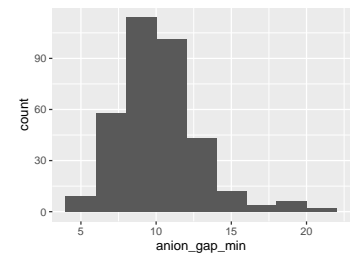
Feature	Result
Variable type	numeric
Number of missing obs.	339 (32.44 %)
Number of unique values	333
Median	6.75
1st and 3rd quartiles	5.6; 8.8
Min. and max.	3.7; 35.8



- Note that the following possible outlier values were detected: "3.7", "3.9", "4", "4.1", "4.2", "4.25", "24.9", "35.8".

anion__gap__min

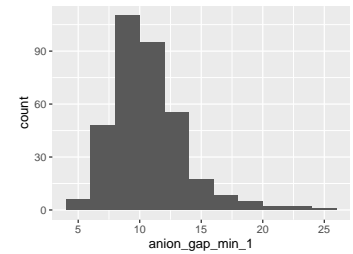
Feature	Result
Variable type	numeric
Number of missing obs.	696 (66.6 %)
Number of unique values	17
Median	10
1st and 3rd quartiles	9; 12
Min. and max.	4; 22



- Note that the following possible outlier values were detected: "4", "5", "6", "22".

anion_gap_min_1

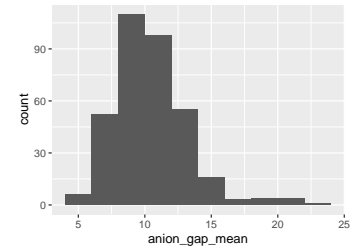
Feature	Result
Variable type	numeric
Number of missing obs.	696 (66.6 %)
Number of unique values	20
Median	11
1st and 3rd quartiles	9; 13
Min. and max.	5; 25



- Note that the following possible outlier values were detected: "20", "21", "22", "23", "25".

anion_gap_mean

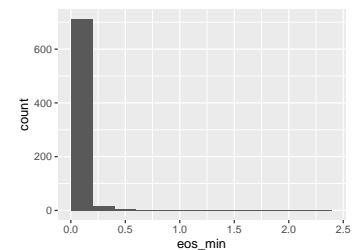
Feature	Result
Variable type	numeric
Number of missing obs.	696 (66.6 %)
Number of unique values	30
Median	11
1st and 3rd quartiles	9; 12
Min. and max.	4.5; 22.5



- Note that the following possible outlier values were detected: "18", "19", "20", "20.5", "21.33", "21.5", "22", "22.5".

eos_min

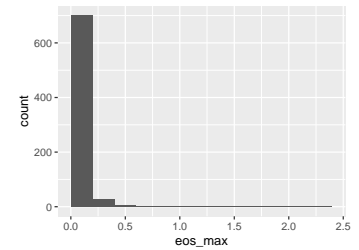
Feature	Result
Variable type	numeric
Number of missing obs.	306 (29.28 %)
Number of unique values	36
Median	0
1st and 3rd quartiles	0; 0.04
Min. and max.	0; 2.22



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

eos__max

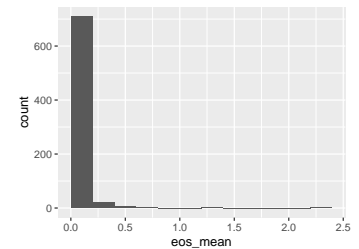
Feature	Result
Variable type	numeric
Number of missing obs.	306 (29.28 %)
Number of unique values	43
Median	0
1st and 3rd quartiles	0; 0.06
Min. and max.	0; 2.22



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

eos__mean

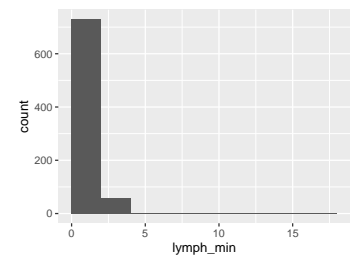
Feature	Result
Variable type	numeric
Number of missing obs.	306 (29.28 %)
Number of unique values	40
Median	0
1st and 3rd quartiles	0; 0.05
Min. and max.	0; 2.22



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

lymph__min

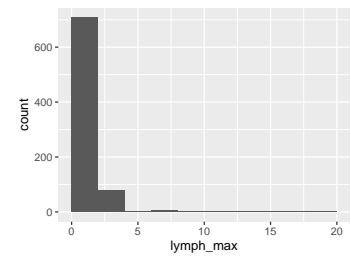
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	215
Median	0.9
1st and 3rd quartiles	0.6; 1.3
Min. and max.	0; 16.9



- Note that the following possible outlier values were detected: "0", "0.05", "0.08", "0.1", "3.19", "3.2", "3.25", "3.3", "3.5", "3.66" (6 additional values omitted).

lymph_max

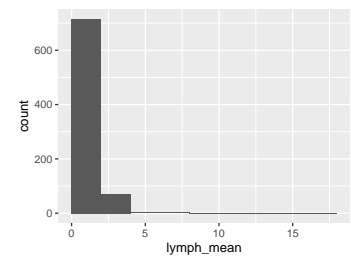
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	224
Median	1.04
1st and 3rd quartiles	0.71; 1.5
Min. and max.	0.1; 18.6



- Note that the following possible outlier values were detected: "0.1", "0.16", "0.18", "0.2", "3.81", "6", "6.25", "6.4", "6.89", "7.3" (2 additional values omitted).

lymph_mean

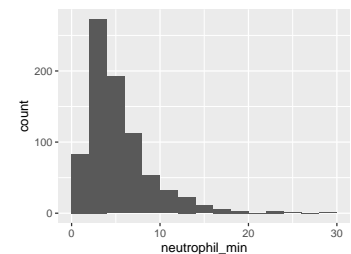
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	223
Median	0.96
1st and 3rd quartiles	0.68; 1.4
Min. and max.	0.1; 17.75



- Note that the following possible outlier values were detected: "0.1", "0.15", "0.16", "0.18", "0.2", "0.24", "0.25", "3.66", "3.81", "4.61" (6 additional values omitted).

neutrophil_min

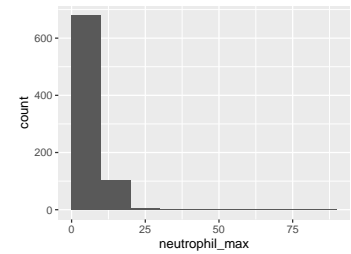
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	460
Median	4.29
1st and 3rd quartiles	2.93; 6.8
Min. and max.	0.09; 29.76



- Note that the following possible outlier values were detected: "0.09", "0.17", "0.37", "0.6", "0.64", "0.66", "0.7", "0.78", "0.8", "0.9" (19 additional values omitted).

neutrophil_max

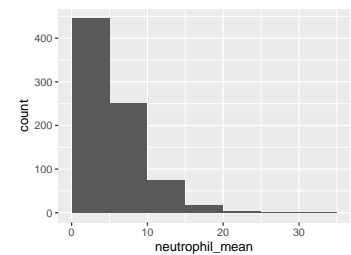
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	469
Median	4.83
1st and 3rd quartiles	3.38; 7.46
Min. and max.	0.18; 83



- Note that the following possible outlier values were detected: "0.18", "0.28", "0.48", "0.64", "0.7", "0.78", "0.8", "0.94", "0.95", "1" (20 additional values omitted).

neutrophil_mean

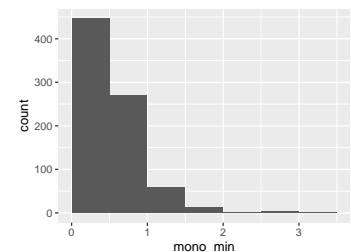
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	472
Median	4.56
1st and 3rd quartiles	3.16; 7.25
Min. and max.	0.14; 33.59



- Note that the following possible outlier values were detected: "0.14", "0.23", "0.41", "0.64", "0.7", "0.78", "0.8", "0.93", "0.94", "1" (19 additional values omitted).

mono_min

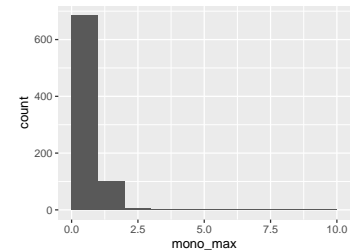
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	139
Median	0.49
1st and 3rd quartiles	0.31; 0.7
Min. and max.	0; 3.39



- Note that the following possible outlier values were detected: "1.59", "1.64", "1.71", "1.77", "1.8", "1.88", "1.98", "2.18", "2.51", "2.8" (2 additional values omitted).

mono__max

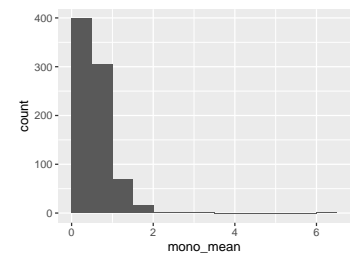
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	151
Median	0.55
1st and 3rd quartiles	0.39; 0.8
Min. and max.	0; 9.5



- Note that the following possible outlier values were detected: "0", "0.01", "0.04", "0.06", "0.07", "0.08", "0.09", "0.1", "0.11", "0.12" (10 additional values omitted).

mono__mean

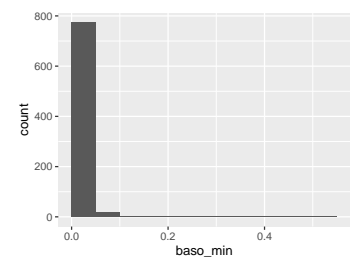
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	145
Median	0.5
1st and 3rd quartiles	0.36; 0.75
Min. and max.	0; 6.18



- Note that the following possible outlier values were detected: "0", "0.01", "0.04", "0.05", "0.06", "0.07", "0.08", "0.1", "0.11", "0.12" (8 additional values omitted).

baso__min

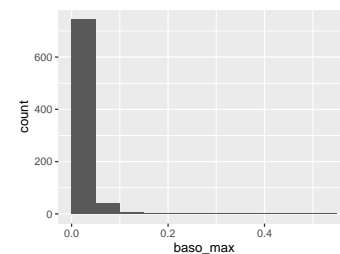
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	14
Median	0
1st and 3rd quartiles	0; 0.01
Min. and max.	0; 0.52



- Note that the following possible outlier values were detected: "0.4", "0.52".

baso_max

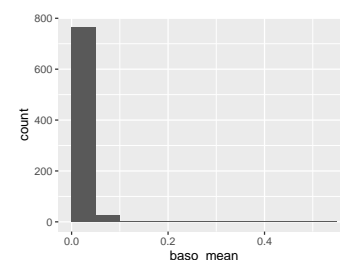
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	17
Median	0.01
1st and 3rd quartiles	0; 0.02
Min. and max.	0; 0.52



- Note that the following possible outlier values were detected: "0.06", "0.07", "0.08", "0.09", "0.1", "0.11", "0.12", "0.19", "0.27", "0.5" (1 additional values omitted).

baso_mean

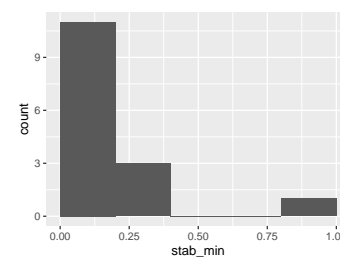
Feature	Result
Variable type	numeric
Number of missing obs.	252 (24.11 %)
Number of unique values	15
Median	0.01
1st and 3rd quartiles	0; 0.01
Min. and max.	0; 0.52



- Note that the following possible outlier values were detected: "0.03", "0.04", "0.05", "0.06", "0.07", "0.08", "0.09", "0.1", "0.11", "0.16" (2 additional values omitted).

stab_min

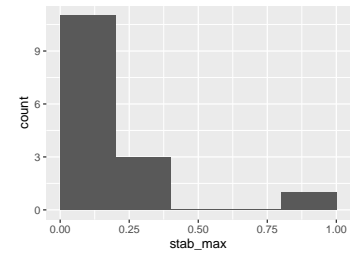
Feature	Result
Variable type	numeric
Number of missing obs.	1030 (98.56 %)
Number of unique values	11
Median	0.17
1st and 3rd quartiles	0.01; 0.22
Min. and max.	0; 0.85



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

stab_max

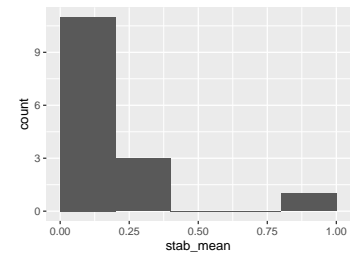
Feature	Result
Variable type	numeric
Number of missing obs.	1030 (98.56 %)
Number of unique values	11
Median	0.17
1st and 3rd quartiles	0.01; 0.22
Min. and max.	0; 0.85



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

stab_mean

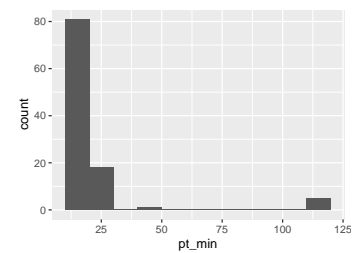
Feature	Result
Variable type	numeric
Number of missing obs.	1030 (98.56 %)
Number of unique values	11
Median	0.17
1st and 3rd quartiles	0.01; 0.22
Min. and max.	0; 0.85



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

pt_min

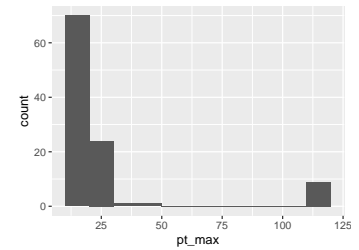
Feature	Result
Variable type	numeric
Number of missing obs.	940 (89.95 %)
Number of unique values	16
Median	18
1st and 3rd quartiles	17; 20
Min. and max.	13; 120



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

pt__max

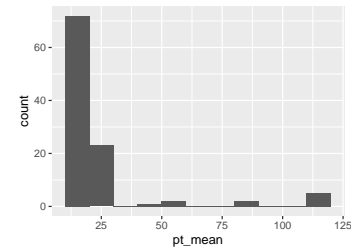
Feature	Result
Variable type	numeric
Number of missing obs.	940 (89.95 %)
Number of unique values	16
Median	19
1st and 3rd quartiles	17; 21
Min. and max.	14; 120



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

pt__mean

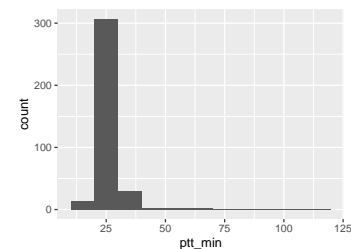
Feature	Result
Variable type	numeric
Number of missing obs.	940 (89.95 %)
Number of unique values	25
Median	19
1st and 3rd quartiles	17; 21
Min. and max.	14; 120



- Note that the following possible outlier values were detected: "45", "51.33", "88.33", "120".

ptt__min

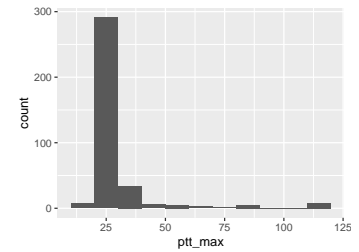
Feature	Result
Variable type	numeric
Number of missing obs.	685 (65.55 %)
Number of unique values	29
Median	25.5
1st and 3rd quartiles	23; 28
Min. and max.	19; 111



- Note that the following possible outlier values were detected: "36", "37", "38", "40", "41", "43", "50", "55", "56", "69" (2 additional values omitted).

ptt_max

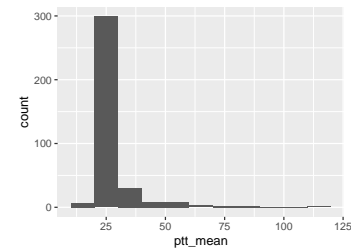
Feature	Result
Variable type	numeric
Number of missing obs.	685 (65.55 %)
Number of unique values	38
Median	26
1st and 3rd quartiles	24; 29
Min. and max.	19; 120



- Note that the following possible outlier values were detected: "19", "20", "43", "46", "47", "49", "51", "55", "56", "58" (8 additional values omitted).

ptt_mean

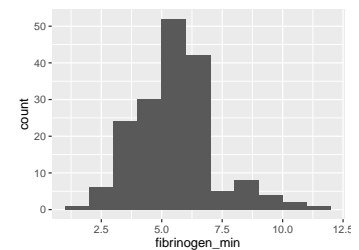
Feature	Result
Variable type	numeric
Number of missing obs.	685 (65.55 %)
Number of unique values	66
Median	26
1st and 3rd quartiles	23.5; 29
Min. and max.	19; 111



- Note that the following possible outlier values were detected: "19", "43", "44", "46", "48", "48.5", "49.33", "54", "55", "56" (10 additional values omitted).

fibrinogen_min

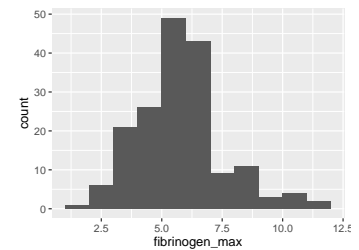
Feature	Result
Variable type	numeric
Number of missing obs.	870 (83.25 %)
Number of unique values	129
Median	5.58
1st and 3rd quartiles	4.42; 6.22
Min. and max.	1.17; 11.78



- Note that the following possible outlier values were detected: "8.25", "8.28", "8.31", "8.51", "8.76", "8.97", "9.13", "9.26", "9.42", "9.62" (2 additional values omitted).

fibrinogen_max

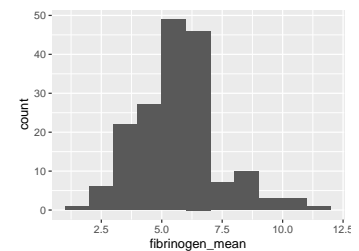
Feature	Result
Variable type	numeric
Number of missing obs.	870 (83.25 %)
Number of unique values	133
Median	5.66
1st and 3rd quartiles	4.52; 6.41
Min. and max.	1.17; 11.78



- Note that the following possible outlier values were detected: "1.17", "9.13", "9.26", "9.62", "10.57", "10.65", "10.7", "11.69", "11.78".

fibrinogen_mean

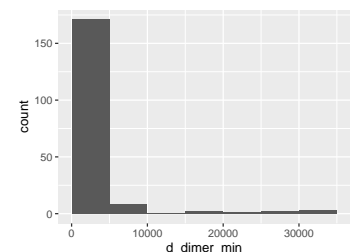
Feature	Result
Variable type	numeric
Number of missing obs.	870 (83.25 %)
Number of unique values	133
Median	5.6
1st and 3rd quartiles	4.47; 6.32
Min. and max.	1.17; 11.78



- Note that the following possible outlier values were detected: "1.17", "8.6", "8.76", "8.97", "9.13", "9.26", "9.62", "10.56", "10.65", "11.78".

d_dimer_min

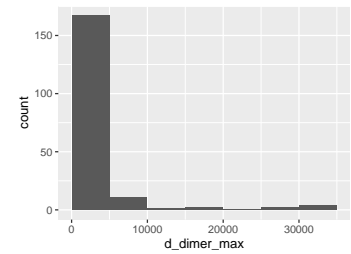
Feature	Result
Variable type	numeric
Number of missing obs.	858 (82.11 %)
Number of unique values	168
Median	1035
1st and 3rd quartiles	510; 1985.5
Min. and max.	170; 34255



- Note that the following possible outlier values were detected: "18871", "19574", "24707", "27322", "28525", "31118", "34255".

d_dimer_max

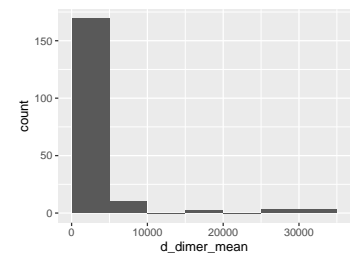
Feature	Result
Variable type	numeric
Number of missing obs.	858 (82.11 %)
Number of unique values	169
Median	1051
1st and 3rd quartiles	515; 1985.5
Min. and max.	170; 34255



- Note that the following possible outlier values were detected: "14725", "18871", "19574", "27322", "28525", "31118", "31651", "34255".

d_dimer_mean

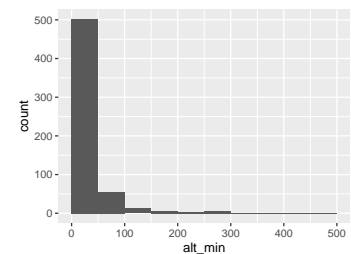
Feature	Result
Variable type	numeric
Number of missing obs.	858 (82.11 %)
Number of unique values	169
Median	1037
1st and 3rd quartiles	515; 1985.5
Min. and max.	170; 34255



- Note that the following possible outlier values were detected: "18871", "19574", "27322", "28179", "28525", "31118", "34255".

alt_min

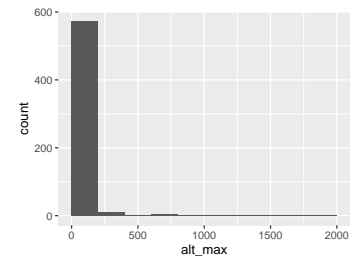
Feature	Result
Variable type	numeric
Number of missing obs.	459 (43.92 %)
Number of unique values	104
Median	22.5
1st and 3rd quartiles	15; 35
Min. and max.	5; 476



- Note that the following possible outlier values were detected: "5", "6", "7", "127", "140", "144", "175", "179", "183", "186" (11 additional values omitted).

alt_max

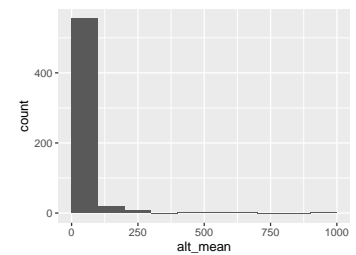
Feature	Result
Variable type	numeric
Number of missing obs.	459 (43.92 %)
Number of unique values	107
Median	23
1st and 3rd quartiles	15; 40
Min. and max.	5; 1861



- Note that the following possible outlier values were detected: "5", "6", "7", "175", "183", "186", "198", "200", "216", "230" (10 additional values omitted).

alt_mean

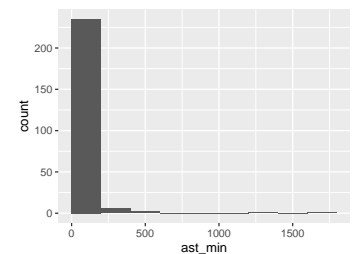
Feature	Result
Variable type	numeric
Number of missing obs.	459 (43.92 %)
Number of unique values	156
Median	23
1st and 3rd quartiles	15; 37.38
Min. and max.	5; 915.67



- Note that the following possible outlier values were detected: "5", "6", "6.5", "7", "140", "151", "175", "183", "186", "194" (12 additional values omitted).

ast_min

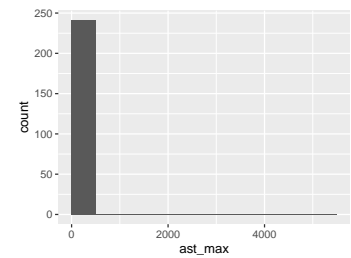
Feature	Result
Variable type	numeric
Number of missing obs.	800 (76.56 %)
Number of unique values	87
Median	34
1st and 3rd quartiles	23; 54
Min. and max.	9; 1735



- Note that the following possible outlier values were detected: "9", "10", "12", "211", "232", "234", "272", "280", "488", "547" (2 additional values omitted).

ast__max

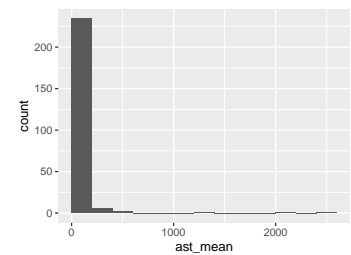
Feature	Result
Variable type	numeric
Number of missing obs.	800 (76.56 %)
Number of unique values	90
Median	36
1st and 3rd quartiles	24; 63
Min. and max.	10; 5486



- Note that the following possible outlier values were detected: "10", "12", "13", "14", "357", "488", "547", "1327", "2381", "5486".

ast__mean

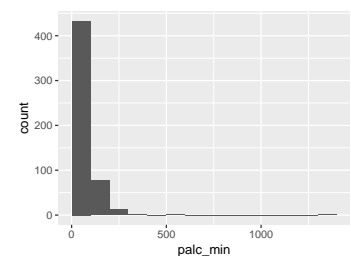
Feature	Result
Variable type	numeric
Number of missing obs.	800 (76.56 %)
Number of unique values	113
Median	34.5
1st and 3rd quartiles	23; 58.75
Min. and max.	10; 2493.33



- Note that the following possible outlier values were detected: "10", "11.5", "12", "14", "272", "280", "304.67", "488", "547", "1327" (2 additional values omitted).

palc__min

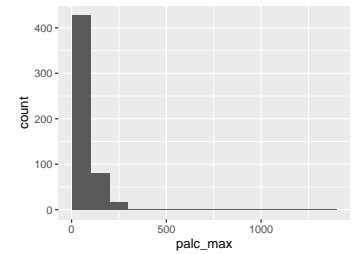
Feature	Result
Variable type	numeric
Number of missing obs.	518 (49.57 %)
Number of unique values	135
Median	63
1st and 3rd quartiles	51; 85
Min. and max.	20; 1358



- Note that the following possible outlier values were detected: "20", "22", "24", "26", "28", "29", "30", "31", "33", "34" (12 additional values omitted).

palc_max

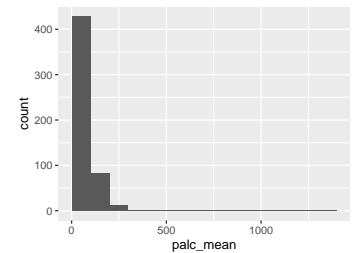
Feature	Result
Variable type	numeric
Number of missing obs.	518 (49.57 %)
Number of unique values	139
Median	64
1st and 3rd quartiles	52; 85
Min. and max.	20; 1358



- Note that the following possible outlier values were detected: "20", "24", "26", "28", "29", "30", "31", "32", "33", "34" (14 additional values omitted).

palc_mean

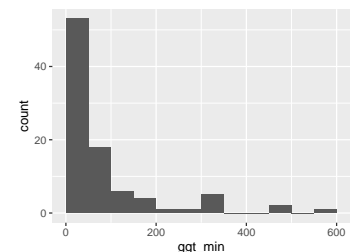
Feature	Result
Variable type	numeric
Number of missing obs.	518 (49.57 %)
Number of unique values	168
Median	63
1st and 3rd quartiles	52; 85
Min. and max.	20; 1358



- Note that the following possible outlier values were detected: "20", "24", "26", "26.75", "28", "29", "30", "31", "33", "34" (14 additional values omitted).

ggt_min

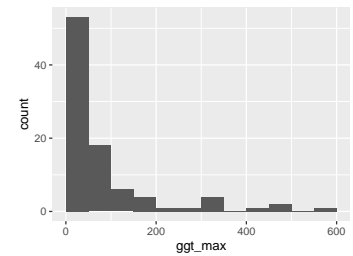
Feature	Result
Variable type	numeric
Number of missing obs.	954 (91.29 %)
Number of unique values	58
Median	41
1st and 3rd quartiles	23; 79.5
Min. and max.	8; 562



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

ggt_max

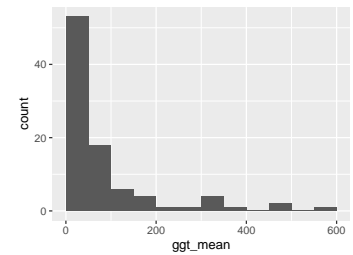
Feature	Result
Variable type	numeric
Number of missing obs.	954 (91.29 %)
Number of unique values	58
Median	41
1st and 3rd quartiles	23; 79.5
Min. and max.	8; 562



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

ggt_mean

Feature	Result
Variable type	numeric
Number of missing obs.	954 (91.29 %)
Number of unique values	59
Median	41
1st and 3rd quartiles	23; 79.5
Min. and max.	8; 562



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

amylase_min

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

amylase_max

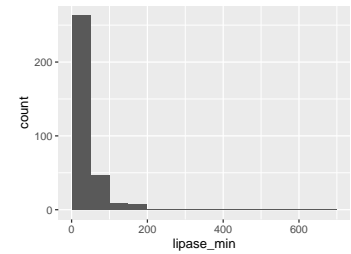
- The variable only takes one value: "NA".

amylase_mean

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

lipase__min

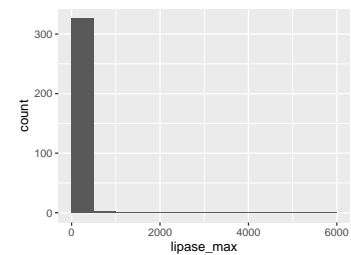
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	95
Median	25.5
1st and 3rd quartiles	16; 44.75
Min. and max.	5; 652



- Note that the following possible outlier values were detected: "5", "178", "182", "185", "194", "444", "466", "548", "652".

lipase__max

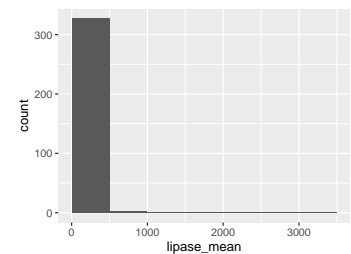
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	95
Median	26
1st and 3rd quartiles	16; 45.75
Min. and max.	5; 5709



- Note that the following possible outlier values were detected: "5", "6", "201", "274", "444", "522", "685", "736", "5709".

lipase__mean

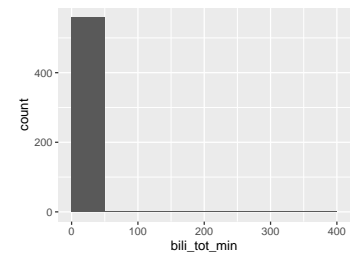
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	110
Median	26
1st and 3rd quartiles	16; 45
Min. and max.	5; 3180.5



- Note that the following possible outlier values were detected: "5", "6", "234", "352", "444", "575.5", "642", "3180.5".

bili_tot_min

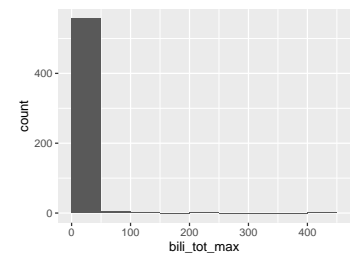
Feature	Result
Variable type	numeric
Number of missing obs.	482 (46.12 %)
Number of unique values	40
Median	8
1st and 3rd quartiles	6; 12
Min. and max.	3; 378



- Note that the following possible outlier values were detected: "3", "37", "40", "44", "48", "51", "70", "142", "226", "378".

bili_tot_max

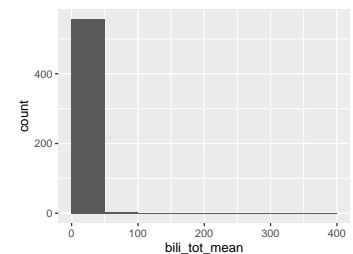
Feature	Result
Variable type	numeric
Number of missing obs.	482 (46.12 %)
Number of unique values	43
Median	9
1st and 3rd quartiles	7; 13
Min. and max.	3; 420



- Note that the following possible outlier values were detected: "3", "4", "38", "39", "40", "42", "45", "48", "57", "68" (4 additional values omitted).

bili_tot_mean

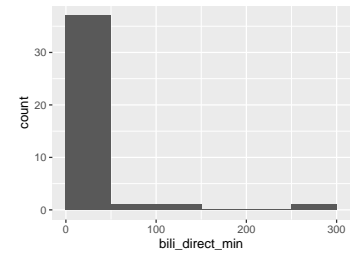
Feature	Result
Variable type	numeric
Number of missing obs.	482 (46.12 %)
Number of unique values	71
Median	8.5
1st and 3rd quartiles	6.5; 12
Min. and max.	3; 399



- Note that the following possible outlier values were detected: "3", "3.5", "3.75", "4", "4.5", "37", "40", "46.5", "48", "52.33" (5 additional values omitted).

bili_direct_min

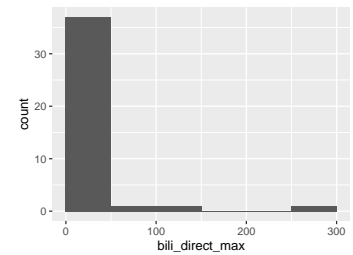
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	35
Median	12.4
1st and 3rd quartiles	7.15; 17.48
Min. and max.	4.5; 251.7



- Note that the following possible outlier values were detected: "91.8", "140.3", "251.7".

bili_direct_max

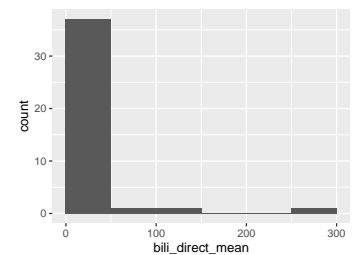
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	36
Median	13.25
1st and 3rd quartiles	7.97; 20.4
Min. and max.	4.5; 288.6



- Note that the following possible outlier values were detected: "91.8", "140.3", "288.6".

bili_direct_mean

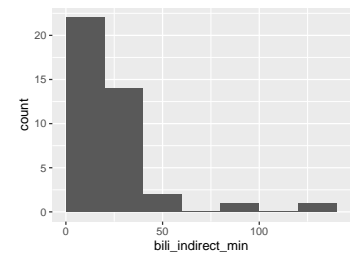
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	36
Median	13.25
1st and 3rd quartiles	7.22; 18.92
Min. and max.	4.5; 270.15



- Note that the following possible outlier values were detected: "91.8", "140.3", "270.15".

bili_indirect_min

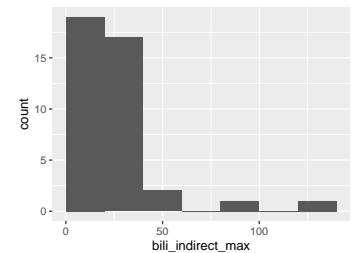
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	34
Median	19.4
1st and 3rd quartiles	14.8; 22.45
Min. and max.	7.5; 132



- Note that the following possible outlier values were detected: "31.4", "50.1", "50.6", "85.8", "132".

bili_indirect_max

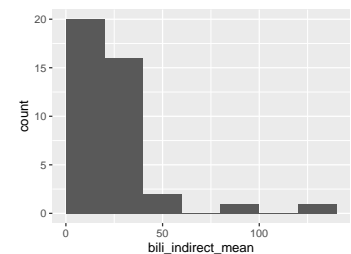
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	36
Median	20.35
1st and 3rd quartiles	16.8; 28.15
Min. and max.	7.5; 134.5



- Note that the following possible outlier values were detected: "7.5", "7.6", "9.5", "10.3", "11.5", "85.8", "134.5".

bili_indirect_mean

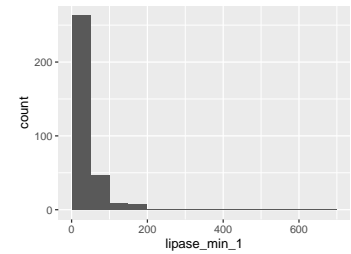
Feature	Result
Variable type	numeric
Number of missing obs.	1005 (96.17 %)
Number of unique values	38
Median	20.1
1st and 3rd quartiles	14.8; 23.62
Min. and max.	7.5; 133.25



- Note that the following possible outlier values were detected: "50.1", "50.6", "85.8", "133.25".

lipase__min__1

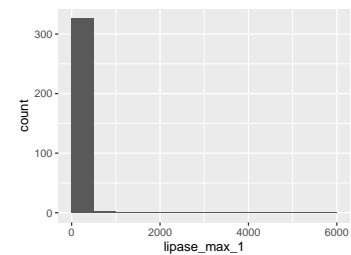
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	95
Median	25.5
1st and 3rd quartiles	16; 44.75
Min. and max.	5; 652



- Note that the following possible outlier values were detected: "5", "178", "182", "185", "194", "444", "466", "548", "652".

lipase__max__1

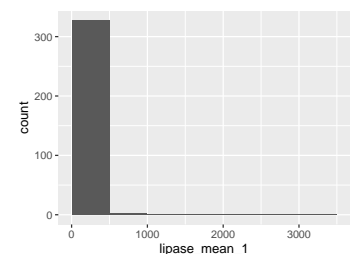
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	95
Median	26
1st and 3rd quartiles	16; 45.75
Min. and max.	5; 5709



- Note that the following possible outlier values were detected: "5", "6", "201", "274", "444", "522", "685", "736", "5709".

lipase__mean__1

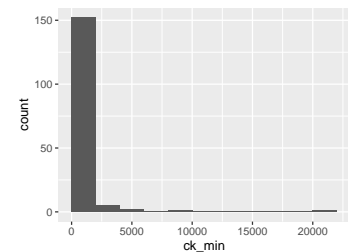
Feature	Result
Variable type	numeric
Number of missing obs.	715 (68.42 %)
Number of unique values	110
Median	26
1st and 3rd quartiles	16; 45
Min. and max.	5; 3180.5



- Note that the following possible outlier values were detected: "5", "6", "234", "352", "444", "575.5", "642", "3180.5".

ck_min

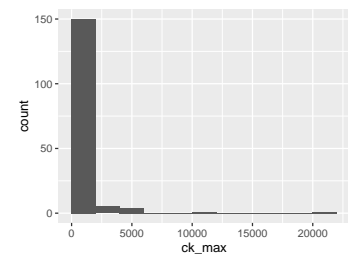
Feature	Result
Variable type	numeric
Number of missing obs.	884 (84.59 %)
Number of unique values	127
Median	156
1st and 3rd quartiles	73; 486
Min. and max.	16; 21926



- Note that the following possible outlier values were detected: "16", "4806", "8961", "21926".

ck_max

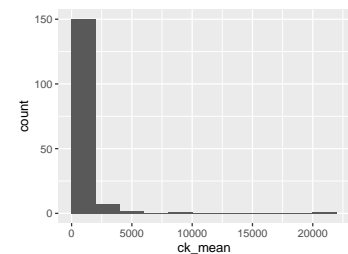
Feature	Result
Variable type	numeric
Number of missing obs.	884 (84.59 %)
Number of unique values	127
Median	193
1st and 3rd quartiles	93; 514
Min. and max.	16; 21926



- Note that the following possible outlier values were detected: "16", "25", "29", "30", "31", "5029", "5990", "10132", "21926".

ck_mean

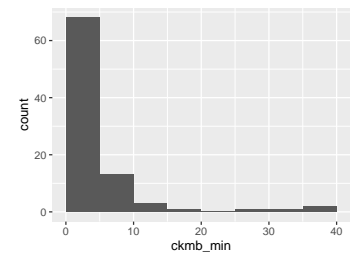
Feature	Result
Variable type	numeric
Number of missing obs.	884 (84.59 %)
Number of unique values	133
Median	173
1st and 3rd quartiles	90.25; 491.5
Min. and max.	16; 21926



- Note that the following possible outlier values were detected: "16", "25", "30", "31", "32", "33", "35", "35.67", "37", "5398" (2 additional values omitted).

ckmb_min

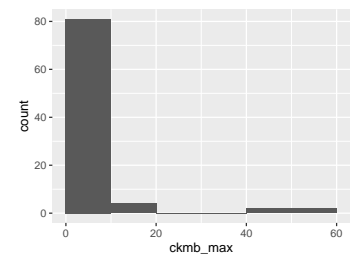
Feature	Result
Variable type	numeric
Number of missing obs.	956 (91.48 %)
Number of unique values	42
Median	2.2
1st and 3rd quartiles	1; 3.6
Min. and max.	0.4; 39.6



- Note that the following possible outlier values were detected: "28.2", "31.3", "39.6".

ckmb_max

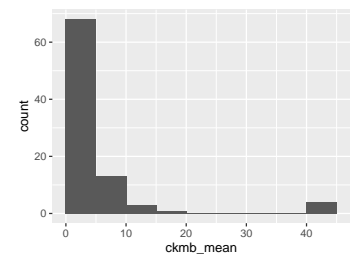
Feature	Result
Variable type	numeric
Number of missing obs.	956 (91.48 %)
Number of unique values	48
Median	2.6
1st and 3rd quartiles	1.2; 5.3
Min. and max.	0.4; 57.9



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

ckmb_mean

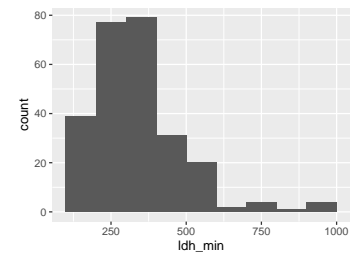
Feature	Result
Variable type	numeric
Number of missing obs.	956 (91.48 %)
Number of unique values	56
Median	2.33
1st and 3rd quartiles	1.15; 4.9
Min. and max.	0.4; 44.65



- Note that the following possible outlier values were detected: "42.4", "44.6", "44.65".

ldh_min

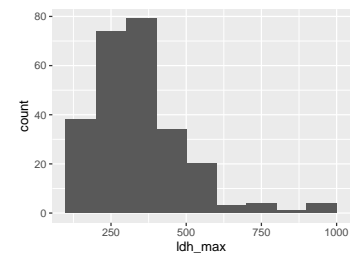
Feature	Result
Variable type	numeric
Number of missing obs.	788 (75.41 %)
Number of unique values	181
Median	318
1st and 3rd quartiles	238; 395
Min. and max.	107; 965



- Note that the following possible outlier values were detected: "690", "707", "709", "713", "812", "922", "926", "941", "965".

ldh_max

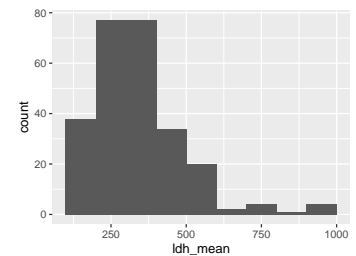
Feature	Result
Variable type	numeric
Number of missing obs.	788 (75.41 %)
Number of unique values	186
Median	321
1st and 3rd quartiles	240; 403
Min. and max.	107; 966



- Note that the following possible outlier values were detected: "707", "709", "713", "812", "922", "926", "965", "966".

ldh_mean

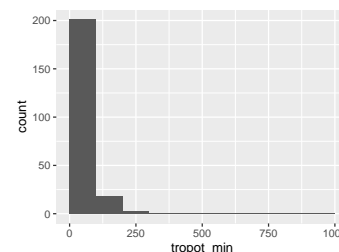
Feature	Result
Variable type	numeric
Number of missing obs.	788 (75.41 %)
Number of unique values	185
Median	318
1st and 3rd quartiles	240; 401
Min. and max.	107; 965



- Note that the following possible outlier values were detected: "709", "713", "812", "922", "926", "953.5", "965".

tropot__min

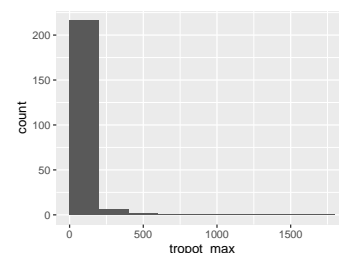
Feature	Result
Variable type	numeric
Number of missing obs.	819 (78.37 %)
Number of unique values	79
Median	22
1st and 3rd quartiles	11; 47.75
Min. and max.	10; 917



- Note that the following possible outlier values were detected: "342", "420", "561", "917".

tropot__max

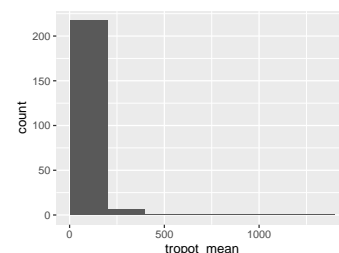
Feature	Result
Variable type	numeric
Number of missing obs.	819 (78.37 %)
Number of unique values	86
Median	24.5
1st and 3rd quartiles	12; 53
Min. and max.	10; 1617



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

tropot__mean

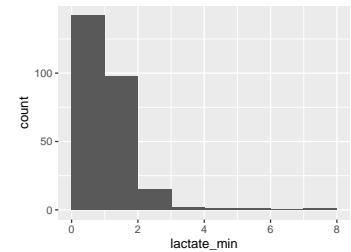
Feature	Result
Variable type	numeric
Number of missing obs.	819 (78.37 %)
Number of unique values	108
Median	22.75
1st and 3rd quartiles	11.5; 50.62
Min. and max.	10; 1274.5



- Note that the following possible outlier values were detected: "430", "706", "1274.5".

lactate__min

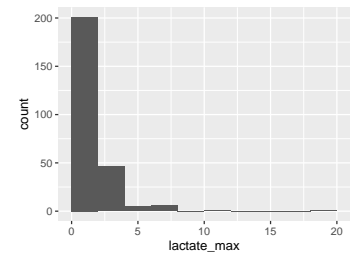
Feature	Result
Variable type	numeric
Number of missing obs.	785 (75.12 %)
Number of unique values	29
Median	1
1st and 3rd quartiles	0.8; 1.4
Min. and max.	0.2; 7.6



- Note that the following possible outlier values were detected: "0.2", "0.3", "0.4", "4.2", "5.6", "7.6".

lactate__max

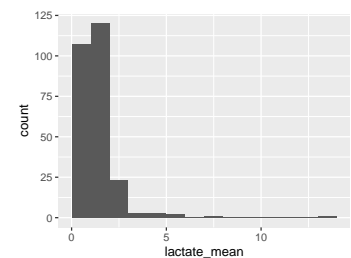
Feature	Result
Variable type	numeric
Number of missing obs.	785 (75.12 %)
Number of unique values	39
Median	1.3
1st and 3rd quartiles	0.98; 2
Min. and max.	0.5; 18.7



- Note that the following possible outlier values were detected: "0.5", "6.3", "7.5", "7.6", "11.4", "18.7".

lactate__mean

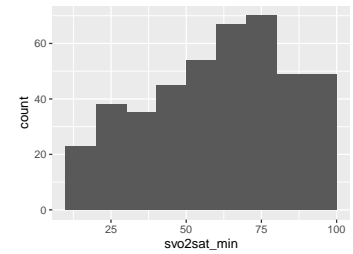
Feature	Result
Variable type	numeric
Number of missing obs.	785 (75.12 %)
Number of unique values	83
Median	1.19
1st and 3rd quartiles	0.9; 1.64
Min. and max.	0.35; 13.11



- Note that the following possible outlier values were detected: "0.35", "4.13", "4.15", "4.46", "5.24", "5.45", "7.6", "13.11".

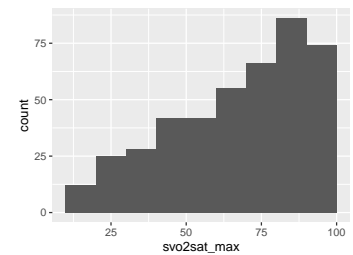
svo2sat__min

Feature	Result
Variable type	numeric
Number of missing obs.	615 (58.85 %)
Number of unique values	89
Median	65
1st and 3rd quartiles	43; 79
Min. and max.	12; 100



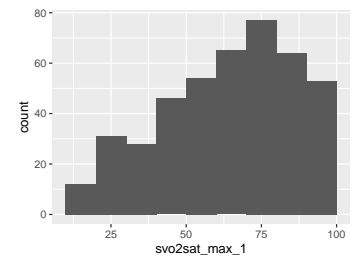
svo2sat__max

Feature	Result
Variable type	numeric
Number of missing obs.	615 (58.85 %)
Number of unique values	85
Median	73
1st and 3rd quartiles	51; 87
Min. and max.	12; 100



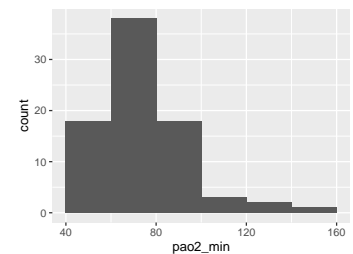
svo2sat__max_1

Feature	Result
Variable type	numeric
Number of missing obs.	615 (58.85 %)
Number of unique values	132
Median	68
1st and 3rd quartiles	48; 81
Min. and max.	12; 100



pao2__min

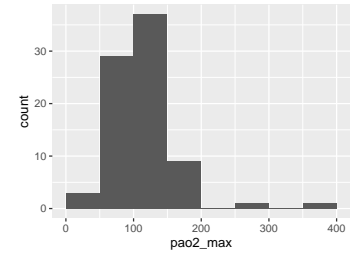
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	67
Median	70.6
1st and 3rd quartiles	60.93; 82.4
Min. and max.	43.7; 159



- Note that the following possible outlier values were detected: "159".

pao2__max

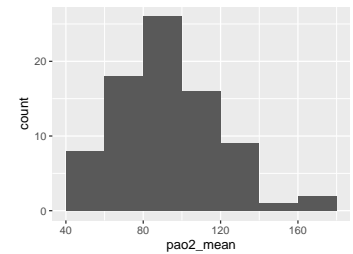
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	61
Median	118
1st and 3rd quartiles	85.83; 142.25
Min. and max.	43.7; 368



- Note that the following possible outlier values were detected: "187", "291", "368".

pao2__mean

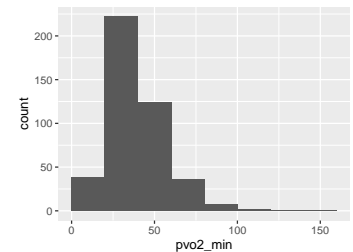
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	69
Median	88.77
1st and 3rd quartiles	73.4; 110.32
Min. and max.	43.7; 179.83



- Note that the following possible outlier values were detected: "43.7".

pvo2__min

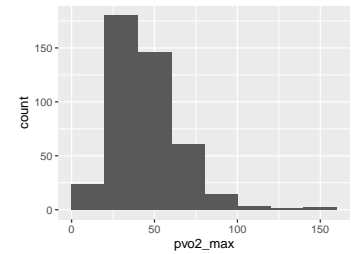
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	272
Median	36.4
1st and 3rd quartiles	26.7; 46.05
Min. and max.	12.2; 158



- Note that the following possible outlier values were detected: "84.3", "85.6", "95.3", "99.6", "108", "109", "158".

pvo2__max

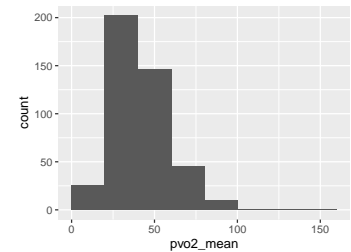
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	293
Median	40.9
1st and 3rd quartiles	30.6; 56.3
Min. and max.	13.7; 158



- Note that the following possible outlier values were detected: "149", "158".

pvo2__mean

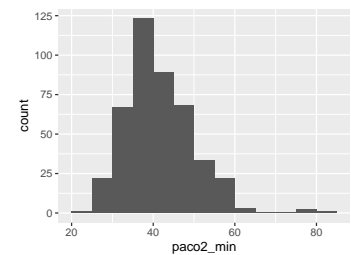
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	304
Median	39
1st and 3rd quartiles	29.33; 50.7
Min. and max.	13.7; 158



- Note that the following possible outlier values were detected: "99.6", "108", "129", "158".

paco2__min

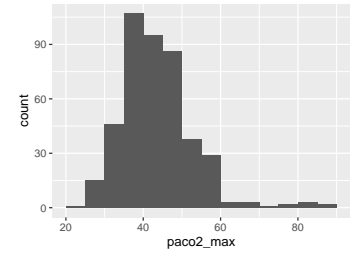
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	211
Median	40.2
1st and 3rd quartiles	35.8; 46.75
Min. and max.	24.9; 82



- Note that the following possible outlier values were detected: "24.9", "25.7", "25.8", "26", "26.3", "26.8", "27.6", "78.1", "82".

paco2__max

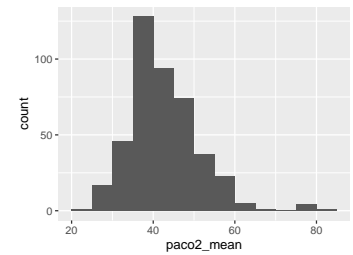
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	221
Median	42.2
1st and 3rd quartiles	37.25; 48.3
Min. and max.	24.9; 87



- Note that the following possible outlier values were detected: "24.9", "25.8", "26.3", "26.8", "27.6", "78.1", "82", "84.9", "87".

paco2__mean

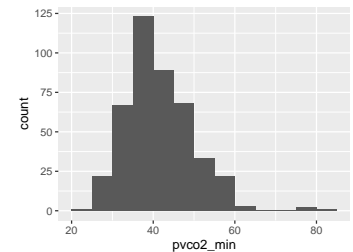
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	251
Median	41.3
1st and 3rd quartiles	36.75; 47.5
Min. and max.	24.9; 82



- Note that the following possible outlier values were detected: "24.9", "25.8", "26.3", "26.8", "27.6", "75.8", "78.1", "82".

pvco2__min

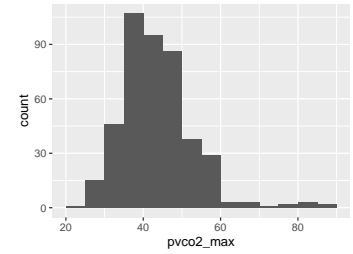
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	211
Median	40.2
1st and 3rd quartiles	35.8; 46.75
Min. and max.	24.9; 82



- Note that the following possible outlier values were detected: "24.9", "25.7", "25.8", "26", "26.3", "26.8", "27.6", "78.1", "82".

pvco2__max

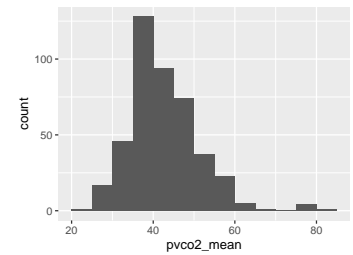
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	221
Median	42.2
1st and 3rd quartiles	37.25; 48.3
Min. and max.	24.9; 87



- Note that the following possible outlier values were detected: "24.9", "25.8", "26.3", "26.8", "27.6", "78.1", "82", "84.9", "87".

pvco2__mean

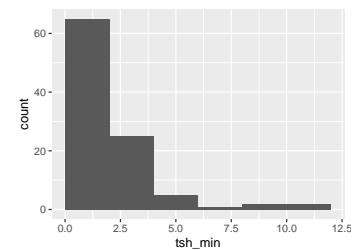
Feature	Result
Variable type	numeric
Number of missing obs.	614 (58.76 %)
Number of unique values	251
Median	41.3
1st and 3rd quartiles	36.75; 47.5
Min. and max.	24.9; 82



- Note that the following possible outlier values were detected: "24.9", "25.8", "26.3", "26.8", "27.6", "75.8", "78.1", "82".

tsh__min

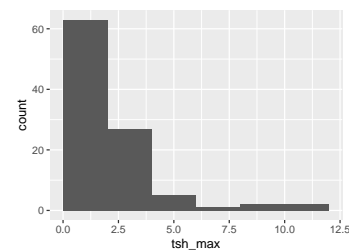
Feature	Result
Variable type	numeric
Number of missing obs.	945 (90.43 %)
Number of unique values	86
Median	1.33
1st and 3rd quartiles	0.86; 2.53
Min. and max.	0.1; 11.78



- Note that the following possible outlier values were detected: "0.1", "0.18", "0.25", "0.28", "0.3", "0.34", "0.35", "0.39", "0.4", "0.41".

tsh__max

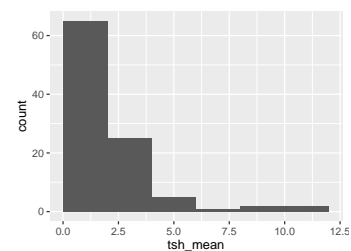
Feature	Result
Variable type	numeric
Number of missing obs.	945 (90.43 %)
Number of unique values	86
Median	1.33
1st and 3rd quartiles	0.86; 2.53
Min. and max.	0.1; 11.78



- Note that the following possible outlier values were detected: "0.1", "0.18", "0.25", "0.28", "0.3", "0.34", "0.35", "0.39", "0.4", "0.41".

tsh__mean

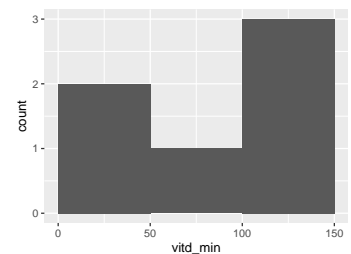
Feature	Result
Variable type	numeric
Number of missing obs.	945 (90.43 %)
Number of unique values	85
Median	1.33
1st and 3rd quartiles	0.86; 2.53
Min. and max.	0.1; 11.78



- Note that the following possible outlier values were detected: "0.1", "0.18", "0.25", "0.28", "0.3", "0.34", "0.35", "0.39", "0.4", "0.41".

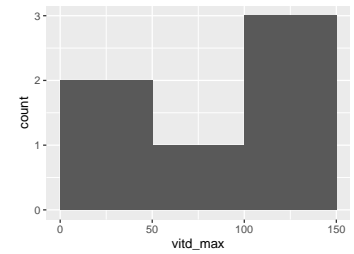
vitd__min

Feature	Result
Variable type	numeric
Number of missing obs.	1039 (99.43 %)
Number of unique values	6
Median	80
1st and 3rd quartiles	47.25; 110.5
Min. and max.	6; 123



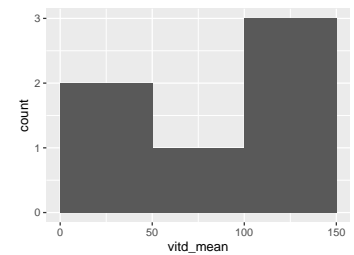
vitd_max

Feature	Result
Variable type	numeric
Number of missing obs.	1039 (99.43 %)
Number of unique values	6
Median	80
1st and 3rd quartiles	47.25; 110.5
Min. and max.	6; 123



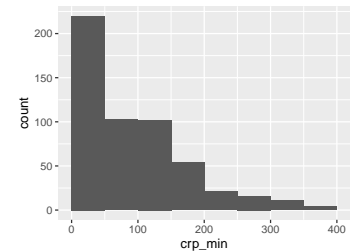
vitd_mean

Feature	Result
Variable type	numeric
Number of missing obs.	1039 (99.43 %)
Number of unique values	6
Median	80
1st and 3rd quartiles	47.25; 110.5
Min. and max.	6; 123



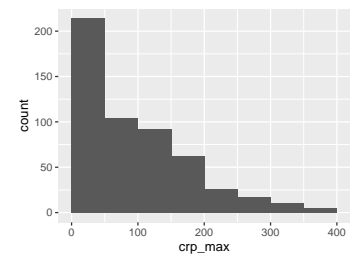
crp_min

Feature	Result
Variable type	numeric
Number of missing obs.	514 (49.19 %)
Number of unique values	404
Median	70.7
1st and 3rd quartiles	21.6; 134.75
Min. and max.	5; 384.6



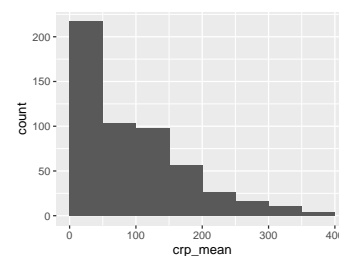
crp_max

Feature	Result
Variable type	numeric
Number of missing obs.	514 (49.19 %)
Number of unique values	398
Median	74.8
1st and 3rd quartiles	22.65; 141.1
Min. and max.	5; 384.6



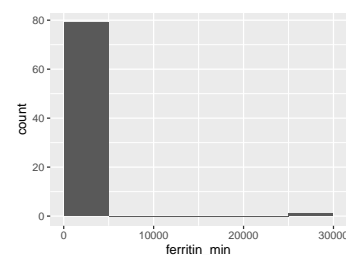
crp_mean

Feature	Result
Variable type	numeric
Number of missing obs.	514 (49.19 %)
Number of unique values	406
Median	73.9
1st and 3rd quartiles	22.65; 136.7
Min. and max.	5; 384.6



ferritin_min

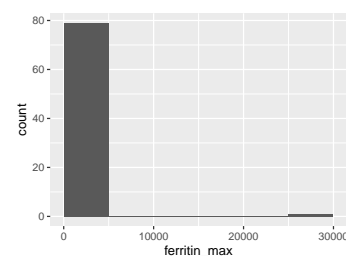
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	75
Median	318.5
1st and 3rd quartiles	183; 734.25
Min. and max.	10; 28696



- Note that the following possible outlier values were detected: "10", "24", "26", "32", "51", "28696".

ferritin_max

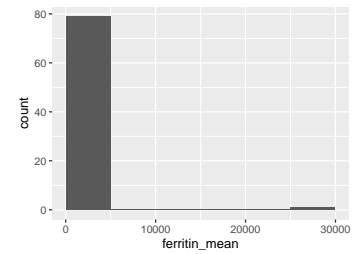
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	75
Median	318.5
1st and 3rd quartiles	183; 734.25
Min. and max.	10; 28696



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

ferritin_mean

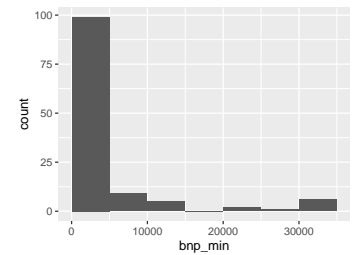
Feature	Result
Variable type	numeric
Number of missing obs.	965 (92.34 %)
Number of unique values	75
Median	318.5
1st and 3rd quartiles	183; 734.25
Min. and max.	10; 28696



- Note that the following possible outlier values were detected: "10", "24", "26.5", "32", "28696".

bnp_min

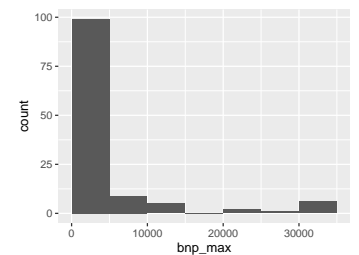
Feature	Result
Variable type	numeric
Number of missing obs.	923 (88.33 %)
Number of unique values	108
Median	1030
1st and 3rd quartiles	202.5; 3645.5
Min. and max.	6; 35000



- Note that the following possible outlier values were detected: "35000".

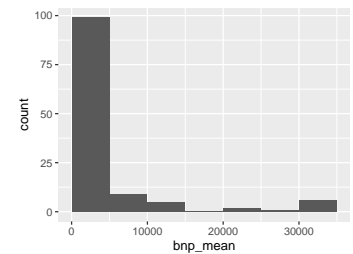
bnp_max

Feature	Result
Variable type	numeric
Number of missing obs.	923 (88.33 %)
Number of unique values	108
Median	1082.5
1st and 3rd quartiles	207; 3929.25
Min. and max.	6; 35000



bnp__mean

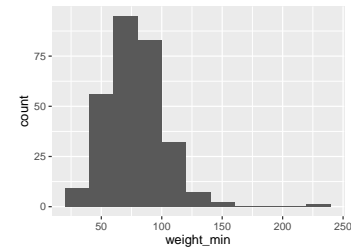
Feature	Result
Variable type	numeric
Number of missing obs.	923 (88.33 %)
Number of unique values	108
Median	1082.5
1st and 3rd quartiles	205.88; 3795
Min. and max.	6; 35000



- Note that the following possible outlier values were detected: "35000".

weight__min

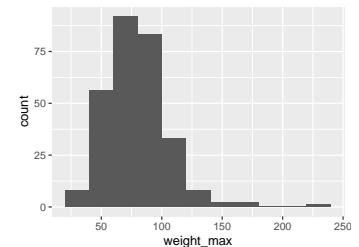
Feature	Result
Variable type	numeric
Number of missing obs.	760 (72.73 %)
Number of unique values	214
Median	77.4
1st and 3rd quartiles	62; 89.6
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "128", "128.9", "129.9", "135.7", "147.3", "150.4", "236".

weight__max

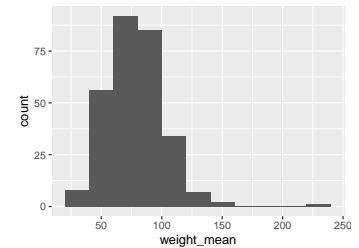
Feature	Result
Variable type	numeric
Number of missing obs.	760 (72.73 %)
Number of unique values	211
Median	78
1st and 3rd quartiles	62.4; 90
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "128", "128.9", "130.4", "135.7", "147.3", "150.4", "173", "236".

weight_mean

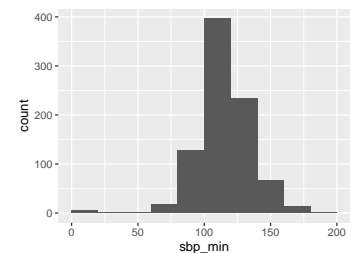
Feature	Result
Variable type	numeric
Number of missing obs.	760 (72.73 %)
Number of unique values	221
Median	78
1st and 3rd quartiles	62.2; 90
Min. and max.	30.4; 236



- Note that the following possible outlier values were detected: "128", "128.9", "130.15", "135.7", "147.3", "150.4", "236".

sbp_min

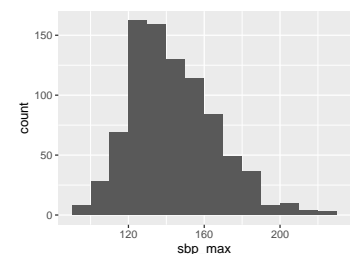
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	106
Median	115
1st and 3rd quartiles	104; 127
Min. and max.	11; 185



- Note that the following possible outlier values were detected: "11", "12", "16", "19", "60", "62", "66", "69", "70", "72" (10 additional values omitted).

sbp_max

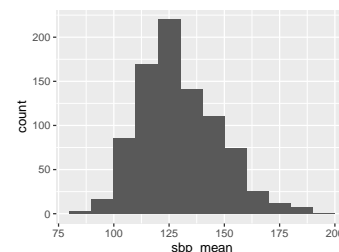
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	112
Median	141
1st and 3rd quartiles	128; 158
Min. and max.	95; 224



- Note that the following possible outlier values were detected: "95", "96", "97", "98", "100", "101", "102", "103".

sbp_mean

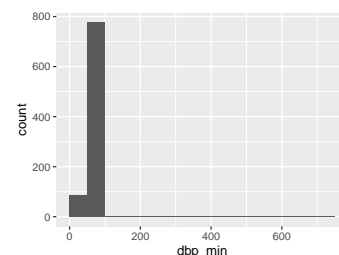
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	517
Median	126.9
1st and 3rd quartiles	117.4; 141
Min. and max.	83.33; 193.4



- Note that the following possible outlier values were detected: "83.33", "86.93", "87.57", "90.71", "92.4", "94", "95.63", "96", "96.2", "96.8" (7 additional values omitted).

dbp_min

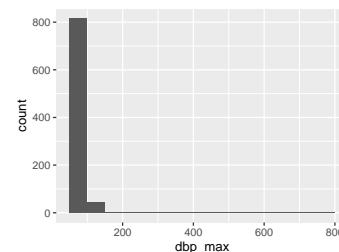
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	69
Median	64
1st and 3rd quartiles	56; 71
Min. and max.	6; 719



- Note that the following possible outlier values were detected: "6", "28", "29", "30", "96", "98", "106", "108", "109", "719".

dbp_max

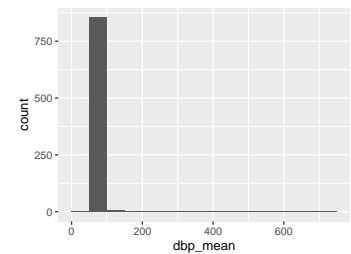
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	78
Median	81
1st and 3rd quartiles	74; 88
Min. and max.	52; 787



- Note that the following possible outlier values were detected: "52", "55", "56", "113", "114", "115", "116", "117", "118", "119" (13 additional values omitted).

dbp__mean

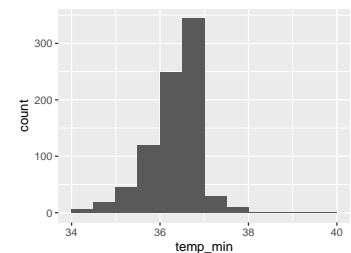
Feature	Result
Variable type	numeric
Number of missing obs.	181 (17.32 %)
Number of unique values	419
Median	72
1st and 3rd quartiles	66; 78.67
Min. and max.	49.5; 719



- Note that the following possible outlier values were detected: "49.5", "101.75", "104.6", "106", "108", "109", "119.36", "135.15", "184.6", "719".

temp__min

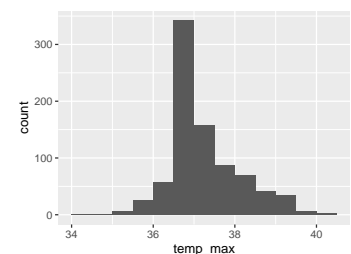
Feature	Result
Variable type	numeric
Number of missing obs.	216 (20.67 %)
Number of unique values	41
Median	36.5
1st and 3rd quartiles	36.1; 37
Min. and max.	34; 39.6



- Note that the following possible outlier values were detected: "34", "34.2", "34.4", "34.5", "34.6", "34.7", "38.5", "38.9", "39.3", "39.6".

temp__max

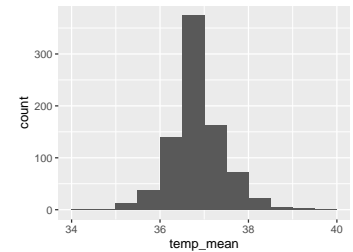
Feature	Result
Variable type	numeric
Number of missing obs.	216 (20.67 %)
Number of unique values	51
Median	37
1st and 3rd quartiles	37; 37.7
Min. and max.	34; 40.2



- Note that the following possible outlier values were detected: "34", "35.2", "35.3", "35.5", "35.6", "35.7", "35.8", "35.9", "36", "36.1" (8 additional values omitted).

temp__mean

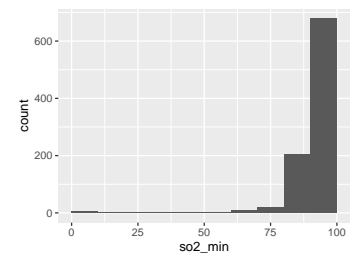
Feature	Result
Variable type	numeric
Number of missing obs.	216 (20.67 %)
Number of unique values	209
Median	36.9
1st and 3rd quartiles	36.55; 37.17
Min. and max.	34; 39.6



- Note that the following possible outlier values were detected: "34", "35.03", "35.1", "35.15", "35.2", "35.25", "37.78", "37.8", "37.83", "37.85" (34 additional values omitted).

so2__min

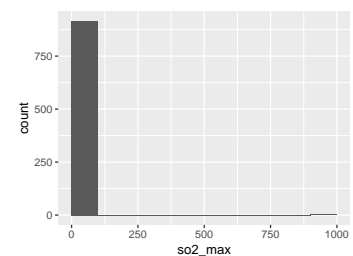
Feature	Result
Variable type	numeric
Number of missing obs.	127 (12.15 %)
Number of unique values	41
Median	93
1st and 3rd quartiles	90; 96
Min. and max.	0; 100



- Note that the following possible outlier values were detected: "0", "1", "2", "9", "18", "20", "25", "32", "63", "64" (8 additional values omitted).

so2__max

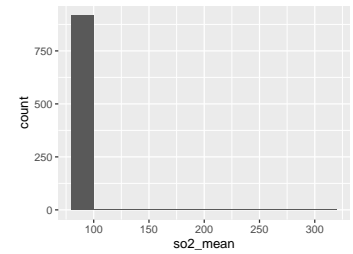
Feature	Result
Variable type	numeric
Number of missing obs.	127 (12.15 %)
Number of unique values	16
Median	98
1st and 3rd quartiles	96; 99
Min. and max.	84; 969



- Note that the following possible outlier values were detected: "84", "85", "966", "969".

so2_mean

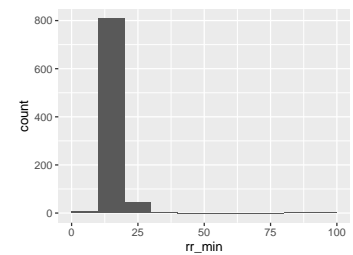
Feature	Result
Variable type	numeric
Number of missing obs.	127 (12.15 %)
Number of unique values	297
Median	95.32
1st and 3rd quartiles	93.67; 97
Min. and max.	80.57; 311.5



- Note that the following possible outlier values were detected: "80.57", "80.71", "81.07", "81.14", "82", "83.17", "84", "84.38", "84.88", "85.38" (9 additional values omitted).

rr_min

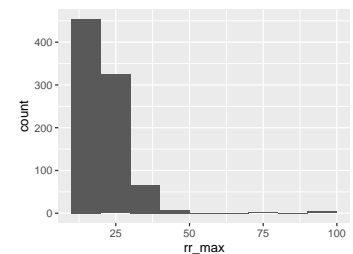
Feature	Result
Variable type	numeric
Number of missing obs.	185 (17.7 %)
Number of unique values	20
Median	18
1st and 3rd quartiles	18; 20
Min. and max.	0; 97



- Note that the following possible outlier values were detected: "0", "2", "8", "10", "12", "14", "15", "16", "17", "85" (1 additional values omitted).

rr_max

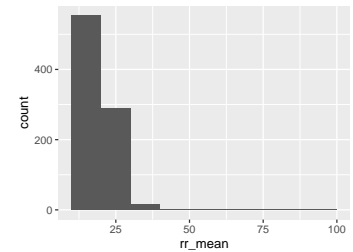
Feature	Result
Variable type	numeric
Number of missing obs.	185 (17.7 %)
Number of unique values	33
Median	20
1st and 3rd quartiles	20; 24
Min. and max.	10; 98



- Note that the following possible outlier values were detected: "10", "14", "16", "18", "19".

rr__mean

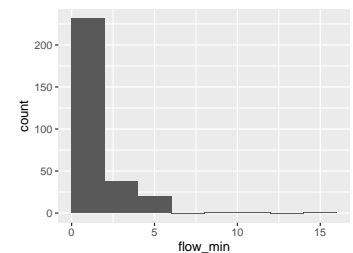
Feature	Result
Variable type	numeric
Number of missing obs.	185 (17.7 %)
Number of unique values	194
Median	20
1st and 3rd quartiles	19; 21
Min. and max.	10; 97



- Note that the following possible outlier values were detected: "10", "14", "15.67", "16", "16.67", "16.7", "16.86", "16.89", "17", "17.14" (31 additional values omitted).

flow__min

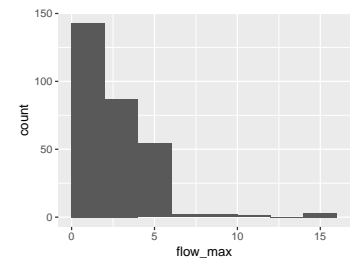
Feature	Result
Variable type	numeric
Number of missing obs.	753 (72.06 %)
Number of unique values	14
Median	1
1st and 3rd quartiles	1; 2
Min. and max.	0.5; 15



- Note that the following possible outlier values were detected: "0.5".

flow__max

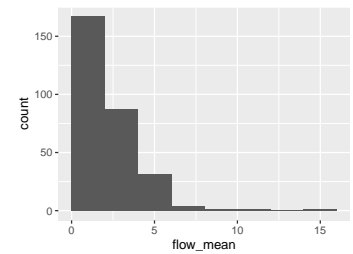
Feature	Result
Variable type	numeric
Number of missing obs.	753 (72.06 %)
Number of unique values	15
Median	2.5
1st and 3rd quartiles	1.5; 4
Min. and max.	0.5; 15



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

flow__mean

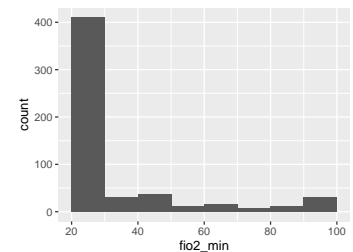
Feature	Result
Variable type	numeric
Number of missing obs.	753 (72.06 %)
Number of unique values	120
Median	1.94
1st and 3rd quartiles	1.08; 3
Min. and max.	0.5; 15



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

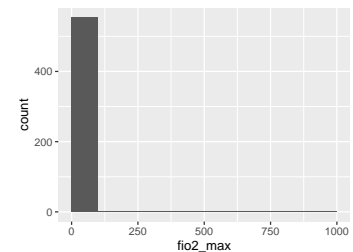
fio2__min

Feature	Result
Variable type	numeric
Number of missing obs.	491 (46.99 %)
Number of unique values	38
Median	21
1st and 3rd quartiles	21; 32
Min. and max.	21; 100



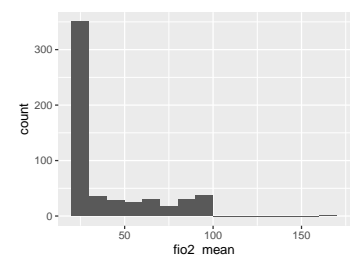
fio2__max

Feature	Result
Variable type	numeric
Number of missing obs.	491 (46.99 %)
Number of unique values	38
Median	24
1st and 3rd quartiles	21; 85
Min. and max.	21; 954



fio2__mean

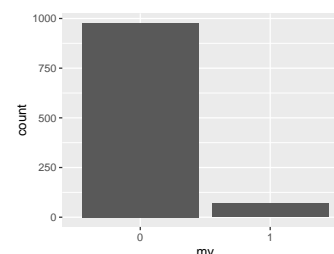
Feature	Result
Variable type	numeric
Number of missing obs.	491 (46.99 %)
Number of unique values	194
Median	22.9
1st and 3rd quartiles	21; 51.26
Min. and max.	21; 166.73



mv

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

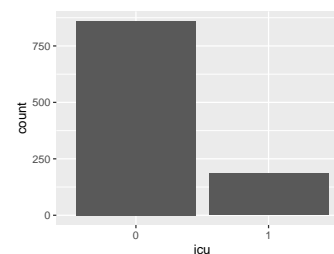
Feature	Result
Variable type	numeric
Number of missing obs.	0 (0 %)
Number of unique values	2
Mode	"0"
Reference category	0



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 18:39:30
- 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 = covid24h_notimputed, render = FALSE, replace = TRUE)`