

Data Frame Summary

csv_input


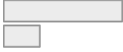


Dimensions: 246719 x 27

Duplicates: 0

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing
1	project_id [factor]	1. BEATAML1.0-COHORT 2. BEATAML1.0-CRENOLANIB 3. CGCI-BLGSP 4. CGCI-HTMCP-CC 5. CMI-ASC 6. CMI-MBC 7. CMI-MPC 8. CPTAC-2 9. CPTAC-3 10. CTSP-DLBCL1 [58 others]	1481 (0.6%) 119 (0.0%) 470 (0.2%) 861 (0.3%) 94 (0.0%) 645 (0.3%) 60 (0.0%) 1024 (0.4%) 4077 (1.7%) 175 (0.1%) 237713 (96.3%)		246719 (100.0%)	0 (0.0%)
2	case_id [factor]	1. 00016c8f-a0be-4319-9c42-4 2. 00021f80-e72d-4c18-9e36-7 3. 00026454-8225-4395-b536-4 4. 0002f311-f263-4816-95f5-f 5. 00039740-17c1-4430-a112-5 6. 00041061-3fe2-417a-a333-2 7. 00048fa6-4318-42ef-9709-7 8. 0004bacb-64f9-4823-a962-6 9. 0004d251-3f70-4395-b175-c 10. 000520f2-c081-4757-8f5c-0 [84559 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 7 (0.0%) 1 (0.0%) 246703 (100.0%)		246719 (100.0%)	0 (0.0%)
3	case_submitter_id [factor]	1. 01BR001 2. 01BR008 3. 01BR009 4. 01BR010 5. 01BR015 6. 01BR017 7. 01BR018 8. 01BR020 9. 01BR023 10. 01BR025 [84481 others]	3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 3 (0.0%) 246689 (100.0%)		246719 (100.0%)	0 (0.0%)
4	sample_id [factor]	1. 000002fc-53a0-420e-b2aa-a 2. 00020a53-a03e-4806-8cf6-a 3. 000225dc-7d01-504a-967f-f 4. 0002a8e8-b820-47d7-a45f-3 5. 0003bde8-b7af-46d7-908f-2 6. 0003bfc0-c5aa-4087-8ebe-2 7. 0004971d-f77b-4116-a49a-e 8. 00069694-c259-4b10-8353-5 9. 0006a6a4-bdec-4942-960c-c 10. 00072b50-23e0-4437-89cc-3 [107613 others]	4 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 2 (0.0%) 1 (0.0%) 1 (0.0%) 246705 (100.0%)		246719 (100.0%)	0 (0.0%)

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing
5	sample_submitter_id [factor]	1. 0065bfd9-e1b9-43a8-9379-9 2. 012baf0d-2c53-4409-91ec-2 3. 012ff650-0db2-41a2-a309-3 4. 01bf1906-9bdc-41c3-89a0-b 5. 01d1ffca-4235-4dfe-bf62-5 6. 01de44a7-81fa-4e25-aba5-9 7. 01f203fd-e664-4d78-bea0-7 8. 02345909-96ed-4f40-a623-7 9. 02ac4d5b-a5e7-4e2c-8ede-f 10. 02f6afea-1e5b-4151-bafd-c [107566 others]	2 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 2 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 246707 (100.0%)		246719 (100.0%)	0 (0.0%)
6	portion_id [factor]	1. 00012d1d-f6ca-4133-9a82-b 2. 0001ab72-8fa0-4b52-9202-5 3. 00026551-91bc-49da-abb1-1 4. 0002b969-6506-46fb-9b6a-e 5. 0002c07e-726f-4b58-9d55-1 6. 0002f5a0-c16b-5810-b0d7-1 7. 00034baa-35a9-5d56-89e1-7 8. 0003c41c-27c6-53cb-bac0-d 9. 0004689e-43d0-4a55-95d3-4 10. 000490e7-cf60-40af-b463-2 [118404 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 246709 (100.0%)		246719 (100.0%)	0 (0.0%)
7	portion_submitter_id [factor]	1. BLGSP-71-06-00001-01A-11 2. BLGSP-71-06-00001-99A-01 3. BLGSP-71-06-00002-01C-01 4. BLGSP-71-06-00002-01D-01 5. BLGSP-71-06-00002-99A-01 6. BLGSP-71-06-00004-01A-11 7. BLGSP-71-06-00004-99A-01 8. BLGSP-71-06-00007-01A-11 9. BLGSP-71-06-00007-01B-01 10. BLGSP-71-06-00007-99A-01 [25451 others]	3 (0.0%) 1 (0.0%) 3 (0.0%) 1 (0.0%) 1 (0.0%) 3 (0.0%) 1 (0.0%) 3 (0.0%) 1 (0.0%) 1 (0.0%) 153247 (100.0%)		153265 (62.1%)	93454 (37.9%)
8	analyte_id [factor]	1. 000109ad-4b03-5ddb-a308-e 2. 00011725-8ad4-509d-8fe7-9 3. 00014e23-c05c-4a2c-9c19-f 4. 0001c4de-9a36-44b8-9db9-f 5. 0001c7b1-381d-4143-92b9-f 6. 00025eb5-d192-4e33-8809-8 7. 000286ef-5d06-42f7-9abd-4 8. 0002e4c6-6d59-59d7-af2b-4 9. 00041799-fd3d-41b5-bd5f-7 10. 00041c1d-47fa-421e-8dde-6 [157500 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 4 (0.0%) 1 (0.0%) 3 (0.0%) 3 (0.0%) 1 (0.0%) 4 (0.0%) 1 (0.0%) 246699 (100.0%)		246719 (100.0%)	0 (0.0%)
9	analyte_submitter_id [factor]	1. ANALYTE1 2. ANALYTE10 3. ANALYTE100 4. ANALYTE101 5. ANALYTE102 6. ANALYTE103 7. ANALYTE104 8. ANALYTE105 9. ANALYTE106 10. ANALYTE107 [64919 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 153629 (100.0%)		153639 (62.3%)	93080 (37.7%)

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing
10	aliquot_id [factor]	1. 00003a00-d06e-4142-b106-8 2. 00012aa3-295d-4e23-ab5b-1 3. 00013730-00f6-445a-b95a-c 4. 00014f01-24a5-4226-9ed6-4 5. 000161c2-e342-4140-bb44-f 6. 0001891b-675c-4fde-911b-e 7. 00025585-b50a-421e-a96c-5 8. 000297b0-854d-492c-8ebf-c 9. 0002a1e0-73ea-42dc-9eb0-8 10. 00035d8f-9716-4a4a-948e-6 [246210 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 246709 (100.0%)		246719 (100.0%)	0 (0.0%)
11	aliquot_submitter_id [factor]	1. 0065bfd9-e1b9-43a8-9379-9 2. 0065bfd9-e1b9-43a8-9379-9 3. 012baf0d-2c53-4409-91ec-2 4. 012ff650-0db2-41a2-a309-3 5. 01bf1906-9bdc-41c3-89a0-b 6. 01d1ffca-4235-4dfe-bf62-5 7. 01d1ffca-4235-4dfe-bf62-5 8. 01de44a7-81fa-4e25-aba5-9 9. 01f203fd-e664-4d78-bea0-7 10. 02345909-96ed-4f40-a623-7 [246163 others]	1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 1 (0.0%) 246709 (100.0%)		246719 (100.0%)	0 (0.0%)
12	aliquot_quantity [numeric]	Mean (sd) : 23.273 (110.469) min < med < max: 0 < 2.4 < 900 IQR (CV) : 2 (4.747)	207 distinct values		7911 (3.2%)	238808 (96.8%)
13	aliquot_volume [numeric]	Mean (sd) : 25.293 (31.273) min < med < max: 0 < 13 < 400 IQR (CV) : 33.3 (1.236)	400 distinct values		9735 (3.9%)	236984 (96.1%)
14	amount [integer]	1 distinct value	65 : 48 (100.0%)		48 (0.0%)	246671 (100.0%)
15	analyte_type [factor]	1. cfDNA 2. DNA 3. FFPE DNA 4. FFPE RNA 5. Nuclei RNA 6. RNA	1 (0.0%) 24534 (83.4%) 352 (1.2%) 276 (0.9%) 18 (0.1%) 4249 (14.4%)		29430 (11.9%)	217289 (88.1%)
16	analyte_type_id [factor]	1. D 2. E 3. R 4. S	1184 (41.2%) 38 (1.3%) 1617 (56.2%) 38 (1.3%)		2877 (1.2%)	243842 (98.8%)
17	concentration [numeric]	Mean (sd) : 0.227 (0.913) min < med < max: 0 < 0.15 < 62 IQR (CV) : 0.03 (4.021)	630 distinct values		149530 (60.6%)	97189 (39.4%)
18	no_matched_normal_low_pass_wgs [logical]	All NA's			0 (0.0%)	246719 (100.0%)
19	no_matched_normal_targeted_sequencing [factor]	1. True	623 (100.0%)		623 (0.3%)	246096 (99.7%)
20	no_matched_normal_wgs [logical]	All NA's			0 (0.0%)	246719 (100.0%)

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing
21	no_matched_normal_wxs [factor]	1. False 2. True	1256 (51.0%) 1207 (49.0%)		2463 (1.0%)	244256 (99.0%)
22	selected_normal_low_pass_wgs [logical]	All NA's			0 (0.0%)	246719 (100.0%)
23	selected_normal_targeted_sequencing [logical]	All NA's			0 (0.0%)	246719 (100.0%)
24	selected_normal_wgs [logical]	All NA's			0 (0.0%)	246719 (100.0%)
25	selected_normal_wxs [factor]	1. False 2. True	1137 (76.8%) 344 (23.2%)		1481 (0.6%)	245238 (99.4%)
26	source_center [factor]	1. 22 2. 23 3. None	68289 (41.3%) 83891 (50.7%) 13193 (8.0%)		165373 (67.0%)	81346 (33.0%)
27	state [factor]	1. released	246719 (100.0%)		246719 (100.0%)	0 (0.0%)

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