







Data Frame Summary

csv_input

Dimensions: 2265 x 10

Duplicates: 0

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing																																																
1	X [integer]	Mean (sd) : 1132 (653.994) min < med < max: 0 < 1132 < 2264 IQR (CV) : 1132 (0.578)	2265 distinct values (Integer sequence)		2265 (100.0%)	0 (0.0%)																																																
2	case_id [character]	1. 0067a0e0-63d8-11e8-bcf1-0 2. 00dc49de-1fba-11e9-b7f8-0 3. 01072f05-118a-11e9-afb9-0 4. 010df72d-63d9-11e8-bcf1-0 5. 01b2691b-63d8-11e8-bcf1-0 6. 0232701d-6d00-440c-af6c-5 7. 024cad7d-63d9-11e8-bcf1-0 8. 02b84fb5-1fba-11e9-b7f8-0 9. 02f4db2f-63d8-11e8-bcf1-0 10. 032d27ba-118a-11e9-afb9-0 [2255 others]	<table><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>1</td><td>(</td><td>0.0%</td><td>)</td></tr><tr><td>2255</td><td>(</td><td>99.6%</td><td>)</td></tr></table>	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%)	1	(0.0%)	2255	(99.6%)		2265 (100.0%)	0 (0.0%)
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2255	(99.6%)																																																			
3	case_submitter_id [character]	1. pooled sample 2. QC1 3. QC2 4. QC3 5. QC4 6. QC5 7. QC6 8. 1 9. 10 10. 11 [2157 others]	<table><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>3</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>2</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>2</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>2</td><td>(</td><td>0.1%</td><td>)</td></tr><tr><td>2238</td><td>(</td><td>98.8%</td><td>)</td></tr></table>	3	(0.1%)	3	(0.1%)	3	(0.1%)	3	(0.1%)	3	(0.1%)	3	(0.1%)	3	(0.1%)	2	(0.1%)	2	(0.1%)	2	(0.1%)	2238	(98.8%)		2265 (100.0%)	0 (0.0%)				
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4	project_submitter_id [character]	1. CPTAC3-Discovery 2. CPTAC-TCGA 3. CPTAC-2 4. Proteogenomic Analysis of 5. Integrated Proteogenomic 6. Academia Sinica LUAD-100 7. Human Early-Onset Gastric 8. Oral Squamous Cell Carcin 9. CPTAC2 Retrospective 10. Georgetown Lung Cancer Pr [2 others]	<table><tr><td>884</td><td>(</td><td>39.0%</td><td>)</td></tr><tr><td>371</td><td>(</td><td>16.4%</td><td>)</td></tr><tr><td>350</td><td>(</td><td>15.5%</td><td>)</td></tr><tr><td>207</td><td>(</td><td>9.1%</td><td>)</td></tr><tr><td>171</td><td>(</td><td>7.5%</td><td>)</td></tr><tr><td>110</td><td>(</td><td>4.9%</td><td>)</td></tr><tr><td>80</td><td>(</td><td>3.5%</td><td>)</td></tr><tr><td>39</td><td>(</td><td>1.7%</td><td>)</td></tr><tr><td>32</td><td>(</td><td>1.4%</td><td>)</td></tr><tr><td>11</td><td>(</td><td>0.5%</td><td>)</td></tr><tr><td>10</td><td>(</td><td>0.4%</td><td>)</td></tr></table>	884	(39.0%)	371	(16.4%)	350	(15.5%)	207	(9.1%)	171	(7.5%)	110	(4.9%)	80	(3.5%)	39	(1.7%)	32	(1.4%)	11	(0.5%)	10	(0.4%)		2265 (100.0%)	0 (0.0%)				
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5	tissue_source_site_code [character]	1. SKHS-UM 2. 2 3. CRE17 4. SCHC 5. 1 6. NCI-Ua 7. STJOE 8. Cureline-35 9. CRE16 10. TJU [22 others]	<table><tr><td>62</td><td>(</td><td>19.2%</td><td>)</td></tr><tr><td>34</td><td>(</td><td>10.5%</td><td>)</td></tr><tr><td>33</td><td>(</td><td>10.2%</td><td>)</td></tr><tr><td>30</td><td>(</td><td>9.3%</td><td>)</td></tr><tr><td>21</td><td>(</td><td>6.5%</td><td>)</td></tr><tr><td>17</td><td>(</td><td>5.3%</td><td>)</td></tr><tr><td>17</td><td>(</td><td>5.3%</td><td>)</td></tr><tr><td>16</td><td>(</td><td>5.0%</td><td>)</td></tr><tr><td>10</td><td>(</td><td>3.1%</td><td>)</td></tr><tr><td>8</td><td>(</td><td>2.5%</td><td>)</td></tr><tr><td>75</td><td>(</td><td>23.2%</td><td>)</td></tr></table>	62	(19.2%)	34	(10.5%)	33	(10.2%)	30	(9.3%)	21	(6.5%)	17	(5.3%)	17	(5.3%)	16	(5.0%)	10	(3.1%)	8	(2.5%)	75	(23.2%)		323 (14.3%)	1942 (85.7%)				
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75	(23.2%)																																																			
6	days_to_lost_to_followup [numeric]	1 distinct value	<table><tr><td>0</td><td>:</td><td>947</td><td>(</td><td>100.0%</td><td>)</td></tr></table>	0	:	947	(100.0%)		947 (41.8%)	1318 (58.2%)																																										
0	:	947	(100.0%)																																																	

No	Variable	Stats / Values	Freqs (% of Valid)	Graph	Valid	Missing
7	disease_type [character]	1. Ovarian Serous Cystadenoc 2. Breast Invasive Carcinoma 3. Lung Adenocarcinoma 4. Pediatric/AYA Brain Tumor 5. Hepatocellular Carcinoma 6. Colon Adenocarcinoma 7. Other 8. Pancreatic Ductal Adenoca 9. Lung Squamous Cell Carcin 10. Clear Cell Renal Cell Car [8 others]	<div>283 (12.5%)</div> <div>233 (10.3%)</div> <div>216 (9.5%)</div> <div>199 (8.8%)</div> <div>170 (7.5%)</div> <div>164 (7.2%)</div> <div>161 (7.1%)</div> <div>140 (6.2%)</div> <div>118 (5.2%)</div> <div>116 (5.1%)</div> <div>465 (20.5%)</div>		2265 (100.0%)	0 (0.0%)
8	index_date [character]	1. Diagnosis	<div>768 (100.0%)</div>		768 (33.9%)	1497 (66.1%)
9	lost_to_followup [character]	1. N/A 2. No 3. Yes	<div>1 (0.2%)</div> <div>395 (83.7%)</div> <div>76 (16.1%)</div>		472 (20.8%)	1793 (79.2%)
10	primary_site [character]	1. Brain 2. Ovary 3. Breast 4. Bronchus and lung 5. Liver 6. Colon 7. Head and Neck 8. Pancreas 9. Lung 10. Kidney [3 others]	<div>309 (14.6%)</div> <div>283 (13.3%)</div> <div>233 (11.0%)</div> <div>222 (10.5%)</div> <div>170 (8.0%)</div> <div>164 (7.7%)</div> <div>148 (7.0%)</div> <div>140 (6.6%)</div> <div>120 (5.7%)</div> <div>119 (5.6%)</div> <div>214 (10.1%)</div>		2122 (93.7%)	143 (6.3%)