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1 *For submission to Molecular Psychiatry*

2 **Neurogenesis is disrupted in human hippocampal progenitor cells upon exposure**  
3 **to serum samples from hospitalized COVID-19 patients**  
4 **with neurological symptoms**

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32 **Abstract**

33 Coronavirus disease 2019 (COVID-19), represents an enormous new threat to our healthcare  
34 system and particularly to the health of older adults. Although the respiratory symptoms of  
35 COVID-19 are well recognized, the neurological manifestations, and their underlying cellular  
36 and molecular mechanisms, have not been extensively studied yet. Our study is the first one to  
37 test the direct effect of serum from hospitalised COVID-19 patients on human hippocampal  
38 neurogenesis using a unique *in vitro* experimental assay with human hippocampal progenitor  
39 cells (HPC0A07/03C). We identify the different molecular pathways activated by serum from  
40 COVID-19 patients with and without neurological symptoms (i.e., delirium), and their effects  
41 on neuronal proliferation, neurogenesis, and apoptosis. We collected serum sample twice, at  
42 time of hospital admission and approximately 5 days after hospitalization. We found that  
43 treatment with serum samples from COVID-19 patients with delirium (n=18) decreased cell  
44 proliferation and neurogenesis, and increases apoptosis, when compared with serum samples  
45 of sex- and age-matched COVID-19 patients without delirium (n=18). This effect was due to  
46 a higher concentration of interleukin 6 (IL6) in serum samples of patients with delirium  
47 (mean±SD: 229.9±79.1pg/ml, vs. 32.5±9.5pg/ml in patients without delirium). Indeed,  
48 treatment of cells with an antibody against IL6 prevented the decreased cell proliferation and  
49 neurogenesis and the increased apoptosis. Moreover, increased concentration of IL6 in serum  
50 samples from delirium patients stimulated the hippocampal cells to produce IL12 and IL13,  
51 and treatment with an antibody against IL12 or IL13 also prevented the decreased cell  
52 proliferation and neurogenesis, and the increased apoptosis. Interestingly, treatment with the  
53 compounds commonly administered to acute COVID-19 patients (the Janus kinase inhibitors,  
54 baricitinib, ruxolitinib and tofacitinib) were able to restore normal cell viability, proliferation  
55 and neurogenesis by targeting the effects of IL12 and IL13. Overall, our results show that serum  
56 from COVID-19 patients with delirium can negatively affect hippocampal-dependent

57 neurogenic processes, and that this effect is mediated by IL6-induced production of the  
58 downstream inflammatory cytokines IL12 and IL13, which are ultimately responsible for the  
59 detrimental cellular outcomes.

60 **Keywords:** delirium; COVID-19; hippocampal neurogenesis; apoptosis; interleukin 6 (IL6);  
61 IL12; IL13

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90 **INTRODUCTION**

91 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), the novel coronavirus that  
92 causes coronavirus disease 2019 (COVID-19), represents an enormous new threat to our  
93 healthcare system and particularly to the health of older adults<sup>1, 2</sup>. Although the respiratory  
94 symptoms of COVID-19 are well recognized<sup>3</sup>, the neurological manifestations and their  
95 underlying mechanisms have not been extensively studied yet. So far, evidence has shown that  
96 COVID-19 patients can develop several neurological symptoms during the acute disease,  
97 including headaches, dizziness, loss of smell, cognitive problems, and in some cases even  
98 delirium symptoms<sup>4, 5</sup>. In older people, delirium, with the associated severe cognitive  
99 disturbances, including confusion and altered level of attention and consciousness, is associated  
100 with adverse outcomes, including prolonged hospitalization and death, even in the absence of  
101 COVID-19 infection<sup>6</sup>.

102

103 So far, studies indicate that 20–30% of COVID-19 patients will present with or develop  
104 delirium or mental status changes during the course of the disease, with rates of 60–70% in  
105 cases of severe illness, at all ages<sup>7-9</sup>. While admission in intensive therapy units (ITU) is itself  
106 associated with delirium symptoms<sup>10</sup>, a sub-group of COVID-19 patients start to show these  
107 symptoms early on before hospitalisation<sup>7, 8</sup>, suggesting that the infection itself, rather than the  
108 hospital or ITU admission, plays a role in the development and/or exacerbation of an already  
109 compromised neurological scenario, especially in older patients. However, the exact  
110 mechanisms underlying the association between COVID-19 and delirium (or other  
111 neurological symptoms) are still not well understood.

112

113

114 It is likely that a major contributing mechanism to the development of neurological symptoms  
115 in COVID-19 patients is a hyper-activated immune system. Indeed, the virus infection is now  
116 well-known for its ability to induce an over-reacting immune response (defined as “cytokine  
117 storm” syndrome) characterized by the production of multiple inflammatory cytokines and  
118 chemokines measurable in the periphery, including interleukin 2 (IL2), IL6, IL8, IL12, IL13,  
119 interferon-gamma (IFN- $\gamma$ ) and tumour necrosis factor-alpha (TNF- $\alpha$ )<sup>11-13</sup>. Once produced,  
120 these peripheral inflammatory cytokines can then penetrate the blood brain barrier (BBB)<sup>14-16</sup> and  
121 directly affect brain mechanisms and induce neurological symptoms affecting cognition, memory,  
122 alertness and emotional state, and leading to delirium<sup>17</sup>. Among these brain mechanisms,  
123 hippocampal neurogenesis is regarded as one of the most important cellular processes involved in  
124 the regulation of cognition<sup>18-20</sup>. However, which of the cytokines that are elevated by COVID-  
125 infection are also directly relevant to the development of neurological symptoms is currently  
126 unknown.

127

128 Using an *in vitro* model of human hippocampal progenitor cells, we have already demonstrated  
129 that treatment with high concentrations of IL6, similar to those found in peripheral blood of  
130 COVID-19 patients (~50-500pg/ml)<sup>12, 21</sup>, can reduce neurogenesis and increase apoptosis<sup>22, 23</sup>.  
131 Other proinflammatory cytokines, like IL1 $\beta$  and interferon-alpha (IFN- $\alpha$ ), have similar effects  
132 in this cellular model<sup>24-27</sup>. Moreover, we have shown that treatment of the same cellular model  
133 with serum samples from patients receiving IFN- $\alpha$ -treatment for Hepatitis C also increases  
134 apoptosis and reduces neurogenesis<sup>28</sup>, and that such changes are predictive of the development  
135 of IFN- $\alpha$ -induced depression. Of note, similarly to COVID-19 patients, patients receiving IFN-  
136  $\alpha$  treatment also can present with cognitive impairments, inattention, memory loss and  
137 confusion<sup>29</sup>. This therefore suggests that reduction of hippocampal neurogenesis and/or  
138 increased apoptosis is one of the mechanisms underpinning the inflammation-induced

139 neurological symptoms in COVID-19 infected patients, and that using serum samples from  
140 affected patients in this cellular model might be a mechanistically valid way to understand  
141 these effects.

142

143 Here we use this unique, well-established<sup>22-25, 30-36</sup>, *in vitro* model of human hippocampal  
144 progenitor cells to assess the direct effect of treatment with serum samples from COVID-19  
145 patients on hippocampal neurogenesis, and to investigate the molecular pathways activated by  
146 serum from patients with or without delirium. In particular, we tested the hypothesis that  
147 treatment of human hippocampal progenitor cells with serum from delirium patients would  
148 decrease cell proliferation and neuronal differentiation, and would increase apoptosis, when  
149 compared with serum from patients without delirium. We also identified the candidate  
150 inflammatory mechanisms that are involved in these effects of serum from patients.

151

## 152 **METHODS AND MATERIALS**

153 **Patients' cohort:** The study comprises serum samples collected from a total of 36 patients who  
154 were admitted to Guy's and St Thomas' NHS Foundation Trust during the first wave of the  
155 COVID-19 pandemic in UK (March-June 2020). Samples used in this study were collected at  
156 time of hospital admission (*Time point 1*) and during admission (~5 days after hospitalization;  
157 *Time point 2*). All patients (with and without clinical diagnosis of delirium) started to present  
158 COVID-19 symptoms ~5 days before hospitalisation. N=18 patients presenting with delirium  
159 symptoms at the time of hospital admission were age- and sex-matched with N=18 patients  
160 who did not present any symptoms of delirium, both at time of hospitalization and across  
161 disease progression (see Table 1). None of these patients have been admitted to ITU prior or at  
162 either time of samples collection. Delirium symptoms were documented in clinical notes by  
163 the treating physician around the time of blood collection (during admission, Time point 1; and

164 ~5 days (range: 2-7 days) after hospitalization, Time point 2). In both groups, the majority of  
165 patients presented with two or more medical comorbidities, with hypertension and type 2  
166 diabetes being the most frequently observed; there were no differences in respiratory symptoms  
167 severity, and survival rate, between the two groups (see Table 1). Levels of C reactive protein  
168 (CRP) in serum also did not significantly differ between the two groups, at both time points  
169 (group effect:  $F=1.3$ ,  $p=0.2$ ; time effect:  $F=0.4$ ,  $p=0.8$ , see Table 1). The study was approved  
170 by the King's College Hospital Research Ethics Committee (Ref: 20/SC/0310).

171

172 **Cell culture:** Multipotent human hippocampal progenitor cell line HPC0A07/03C (provided by  
173 ReNeuron, Surrey, UK) was used<sup>22-25, 30-34, 36</sup>. This model has been previously validated using  
174 a hippocampal newborn neuron specific marker, Prospero homeobox protein 1 (Prox1)<sup>31</sup>. Cells  
175 were left to proliferate in Dulbecco's Modified Eagle Medium: Nutrient Mixture F-12  
176 (DMEM/F-12) media to which we added the growth factors epidermal growth factor (EGF),  
177 basic fibroblast growth factor (bFGF) and 4-hydroxytamoxifen (4-OHT). Differentiation was  
178 initiated by removal of the growth factors and 4-OHT. Detailed information on this cell line  
179 can be found in our previous publication<sup>31</sup>.

180

181 **In vitro treatment with serum samples from COVID-19 patients with and without**  
182 **delirium:** Cells were plated on 96 well plates (Nunclon) at a density of  $1.2 \times 10^4$  cells per  
183 well. Following our established model<sup>28</sup>, after 1 day of proliferation, cells were treated with  
184 media containing 1% serum (from each COVID-19 patient, delirium and non-delirium) and  
185 0.5mg/ml penicillin for 2 days. At this stage cells were either fixed with 4% PFA for 20 min  
186 at room temperature and immunostained, or left to differentiate for additional 4 days, again  
187 in presence of 1% serum (from the same COVID-19 patient previously used in the  
188 proliferation phase) and 0.5mg/ml penicillin, and then again fixed and immunostained. In



189 addition, at the end of 1 day of serum treatment during proliferation, and of 1 day of serum  
190 treatment during differentiation, cell supernatant was collected for subsequent cytokines  
191 measurement (Figure 1a, b, c and d). For the experiment with the antibodies against  
192 cytokines, anti-IL6 antibody (0.1ug/ml) was added during both proliferation and  
193 differentiation, anti-IL12 antibody (0.3ug/ml) during proliferation, and anti-IL13 antibody  
194 (0.1ug/ml) during differentiation (Figure 1a-e), based on the results of supernatant cytokines  
195 measurement. Finally, for the experiments with the Janus kinase (JAK) inhibitors, baricitinib,  
196 ruxolitinib or tofacitinib (all 1nM) were added during proliferation or differentiation together  
197 with recombinant IL12 or IL13 at concentrations previously found in supernatant of cells  
198 exposed to serum from patients with delirium (IL12: 20pg/ml, IL13: 25pg/ml) or without  
199 delirium (IL12: 3pg/ml, IL13: 4pg/ml) (Figure 1f, g, Supplementary Figure 5a-e).

200

201 **Immunocytochemistry and quantification of immunofluorescence:** During the proliferation  
202 stage, fixed cells were stained for markers of stemness (sex determining region Y-box 2 (Sox2),  
203 Alexa 488 donkey anti-rabbit; 1:1000, Invitrogen; Nestin, Alexa 555 donkey anti-mouse, 1:1000,  
204 Invitrogen), proliferation (Ki67, Alexa 488 donkey anti-rabbit; 1:1000, Invitrogen) and apoptosis  
205 (CC3, Alexa 555 donkey anti-mouse, 1:1000, Invitrogen). During the differentiation stage, cells  
206 were stained for markers of neuroblasts and mature neurons (respectively, doublecortin (DCX),  
207 Alexa 555 donkey anti-rabbit; 1:1000, Invitrogen; microtubule-associated protein 2 (Map2),  
208 Alexa 488 donkey anti-mouse, 1:1000, Invitrogen), astrocytes (S100 calcium-binding protein  $\beta$   
209 (S100 $\beta$ ), Alexa 488 donkey anti-rabbit; 1:1000, Invitrogen), and again apoptosis (CC3; Alexa  
210 555 donkey anti-rabbit, 1:1000, Invitrogen). All cells were labelled using DAPI dye, as in  
211 previous publications<sup>22, 25, 28, 37-39</sup>. The percentage of Sox2, Nestin, Ki67, DCX, Map2, S100 $\beta$   
212 and CC3 positive cells over total DAPI positive cells was counted using an insight automated  
213 imaging platform (CellInsight) (Supplementary Figure 1a-h for representative images). At least

214 six independent experiments were conducted on independent biological cultures, and each  
215 sample was tested in quadruplicate.

216

217 **Multiplex cytokine measurement in cell supernatant and serum samples:** Cell supernatants  
218 and serum samples from COVID-19 patients with and without delirium (collected both at *Time*  
219 *point 1* and 2) were used for cytokines measurement (IL1 $\beta$ , IL2, IL4, IL6, IL8, IL10, IL12,  
220 IL13, TNF- $\alpha$ , IFN- $\gamma$ ), using the Human ProInflammatory Multiplex Very-Sensitive Kit from  
221 Meso Scale Discovery (MSD) (Gaithersburg, MD) and the SECTOR Imager MSD device,  
222 according to the manufacturers' instructions. Detailed information on the cytokines analyses  
223 procedure can be found in our previous publication<sup>22, 23, 40</sup>. At least six independent experiments  
224 were conducted on independent biological cultures, and each sample was tested in duplicates.

225

226 **Statistical Analysis:** Statistical analyses were performed with IBM SPSS statistical software  
227 version 25, StataCorp STATA version 16 and GraphPad Prism version 8 and consisted of two-  
228 way mixed analysis of variance (ANOVA), Chi-square  $\chi^2$  test, Mann-Whitney U-test, followed  
229 by Bonferroni's post hoc analyses where appropriate. Variance was tested using the Shapiro-  
230 Wilk test. Data are presented as mean $\pm$ SD, and p-values $\leq$ 0.05 were considered significant.

231

## 232 **RESULTS**

233 **Treatment with serum samples from COVID-19 patients with delirium decreased cell**  
234 **proliferation, neurogenesis, and increased apoptosis, when compared with serum from**  
235 **non-delirium patients.**

236 As in our established protocol<sup>28</sup>, we exposed cells to 2 days treatment during proliferation,  
237 with 1% serum sample from each COVID-19 patient with or without delirium, collected both  
238 at time of hospital admission (*Time point 1*) and during admission (*Time point 2*). We

239 measured markers of cell proliferation (Ki67), apoptosis (CC3) and stemness (Sox2, Nestin)  
240 (Figure 1a). Interestingly, we found that treatment of cells with serum samples from delirium  
241 patients decreased cell proliferation (group effect:  $p < 0.0001$ ; time effect:  $p = 0.6$ , Figure 2a,  
242 Supplementary Figure 1i) and increased apoptosis (group effect:  $p < 0.0001$ ; time effect:  $p = 0.5$ ,  
243 Figure 2b, Supplementary Figure 1j), with no differences between cell treated with serum  
244 samples collected at *Time point 1* or *Time point 2*.

245

246 We did not observe any difference in the marker of stemness (Sox2 or Nestin), when  
247 comparing the two groups (delirium vs non-delirium) and the two time points (Supplementary  
248 Figure 2a, b).

249

250 In the differentiation experiments, after the 2 days of treatment during proliferation, we left  
251 cells to differentiate for additional 4 days, again in presence of 1% serum. We measured  
252 markers of neurogenesis (DCX, Map2), astrogliogenesis (S100 $\beta$ ) and apoptosis (CC3)  
253 (Figure 1b). We found that treatment of cells with serum samples from delirium patients  
254 decreased neurogenesis (DCX: group effect:  $p < 0.0001$ ; time effect:  $p = 0.9$ , Figure 2c,  
255 Supplementary Figure 1k; Map2: group effect:  $p < 0.0001$ ; time effect:  $p = 0.9$ , Figure 2d,  
256 Supplementary Figure 1l) and increased apoptosis (group effect:  $p < 0.0001$ ; time effect:  $p = 0.9$ ,  
257 Figure 2e, Supplementary Figure 1m), again with no differences between the two time points.

258

259 Also, we did not observe any differences in the marker of astrogliogenesis (S100 $\beta$ ), when  
260 comparing the two groups (delirium vs non-delirium) and the two time points (Supplementary  
261 Figure 2c).

262

263 **Higher concentration of IL6 in serum samples from COVID-19 patients with delirium**  
264 **induces cells to produce IL12 and IL13, respectively during the proliferation and**  
265 **differentiation stage.**

266 In order to investigate the molecular mechanisms through which cell treatment with serum  
267 samples from delirium patients detrimentally affected cell proliferation, neurogenesis and  
268 apoptosis, we first measured levels of candidate cytokines (IL1 $\beta$ , IL2, IL4, IL6, IL8, IL10,  
269 IL12, IL13, TNF- $\alpha$ , IFN- $\gamma$ ) known to be modulated by the SARS-CoV-2 virus<sup>11-13</sup>, in serum  
270 samples of both patients with delirium and without delirium.

271

272 Interestingly, we found a significantly higher concentration of IL6 in serum samples of  
273 patients with delirium (229.9 $\pm$ 79.1pg/ml, Figure 3a-b), when compared with serum samples  
274 from patients without delirium (32.5 $\pm$ 9.5pg/ml, Figure 3a-b), again with no differences  
275 between serum samples collected at *Time point 1* or *Time point 2* (group effect: p<0.0001;  
276 time effect: p=0.4, Figure 3b). None of the other cytokines were differentially expressed  
277 between the two groups or the two time points (Supplementary Figure 3a-i).

278

279 We then measured the same panel of cytokines in supernatant of cells exposed to treatment  
280 with serum samples from patients with or without delirium (collected both at *Time point 1*  
281 and 2). In particular, cell supernatant was collected after 1 day of serum treatment during  
282 proliferation, and 1 day of serum treatment during differentiation (Figure 1a and b).

283

284 Results showed significantly higher levels of IL12 during the proliferation, and IL13 during  
285 the differentiation, in the supernatant of cells treated with serum from delirium patients vs  
286 non-delirium patients (IL12: 20.6 $\pm$ 2.7pg/ml vs 2.8 $\pm$ 2pg/ml, Figure 3c-d; IL13: 24.6 $\pm$ 3pg/ml  
287 vs 3.9 $\pm$ 3.1pg/ml, Figure 3e-f), with no differences between supernatant of cells treated with

288 serum samples collected at *Time point 1* or *Time point 2* (IL12: group effect:  $p < 0.0001$ ; time  
289 effect:  $p = 0.8$ , Figure 3d; IL13: group effect:  $p < 0.0001$ ; time effect:  $p = 0.9$ , Figure 3f).

290

291 Of note, concentrations of the same cytokines (IL12 and IL13) in the original serum samples  
292 were much lower, and did not differ between delirium and non-delirium patients (IL12:  
293  $2.1 \pm 1 \text{ pg/ml}$  vs  $2.05 \pm 1 \text{ pg/ml}$ , Supplementary Figure 3f; IL13:  $2.7 \pm 0.7 \text{ pg/ml}$  vs  $2.8 \pm 1 \text{ pg/ml}$ ,  
294 Supplementary Figure 3g), indicating that these differences in the supernatant levels are due  
295 to new production of these cytokines by the cells.

296

297 Finally, none of the other cytokines were differentially expressed in the supernatant of cells  
298 treated with serum samples from delirium or non-delirium patients, or from *Time point 1* or  
299 *Time point 2* (Supplementary Figure 3j-zz).

300

301 **During proliferation, treatment with an antibody against IL6 prevents the detrimental**  
302 **effect of serum from COVID-19 patients with delirium on both cell proliferation and**  
303 **apoptosis, and decreases IL12 production.**

304 In order to confirm that the detrimental effect of treatment with serum from delirium patients  
305 on cell proliferation and apoptosis was due to the higher levels of IL6 in the serum samples  
306 of the same patients, we exposed cells to treatment with serum from COVID-19 patients  
307 together with an antibody against IL6 (Figure 1c).

308

309 Interestingly, treatment with IL6 antibody prevented the decrease in proliferation (Ki67) and  
310 increase in apoptosis (CC3) caused by treatment with serum from delirium patients, when  
311 compared with treatment with serum from non-delirium patients (*Time point 1*, delirium vs  
312 delirium + IL6 antibody; for Ki67, 40.2% vs 61.3%,  $p < 0.0001$ ; for CC3, 10.2% vs. 3.4%,

313  $p < 0.0001$ ), with no differences between cells treated with serum samples collected at *Time*  
314 *point 2* (Figure 4a-d).

315

316 Moreover, treatment of cells with serum samples from delirium patients and IL6 antibody  
317 prevented the production of IL12 in cell supernatant (*Time point 1*, delirium vs delirium + IL6  
318 antibody; IL12:  $20.6 \pm 2.7$  pg/ml vs  $4.1 \pm 2.1$  pg/ml,  $p < 0.0001$ ), again with no differences  
319 between *Time point 1* and *Time point 2* (Figure 1c, Figure 4e, f).

320

321 This finding seems to suggest that, during proliferation, the detrimental effect on cell  
322 proliferation and apoptosis caused by IL6, which is produced in higher concentrations in  
323 serum from delirium patients, may be mediated by the production of IL12 in the brain  
324 environment. Indeed, treatment of cells with an antibody against IL12 prevented the decrease  
325 in cell proliferation (Ki67) and increase in apoptosis (CC3) caused by treatment with serum  
326 from delirium patients (*Time point 1*, delirium vs delirium + IL12 antibody; for Ki67, 40.2%  
327 vs 62.2%,  $p < 0.0001$ ; for CC3, 10.2% vs 3.2%,  $p < 0.0001$ ). Again, there were no differences  
328 between cells treated with serum samples collected at *Time point 1* or *Time point 2* (Figure  
329 1c, Figure 4g-j).

330

331 **During differentiation, treatment with an antibody against IL6 prevents the detrimental**  
332 **effect of serum from COVID-19 patients with delirium on both neurogenesis and**  
333 **apoptosis, and decreases IL13 production.**

334 In order to investigate whether the detrimental effect of treatment with serum from delirium  
335 patients on neurogenesis and apoptosis during *differentiation* was also mediated by a higher  
336 concentration of IL6 in the serum samples of the same patients, we exposed cells to treatment

337 with serum and an antibody against IL6, first during proliferation, then during differentiation,  
338 and finally during both proliferation and differentiation (Figure 1d, e).

339

340 Exposing cells to IL6 antibody during the proliferation stage did not prevent the effect of  
341 serum on the aforementioned cellular outcomes (Supplementary Figure 4a-f). However,  
342 treatment of cells with IL6 antibody during the differentiation stage prevented the reduction  
343 in neurogenesis (DCX, Map2) and increase in apoptosis (CC3) previously observed upon  
344 treatment with serum from delirium patients, when compared with serum from non-delirium  
345 patients (*Time point 1*, delirium vs delirium + IL6 antibody; for DCX, 19.1% vs 34.8%,  
346  $p<0.0001$ ; for Map2, 25.1% vs 40.7%,  $p<0.0001$ ; for CC3, 17.3% vs 3.4%,  $p<0.0001$ ). Again,  
347 there were no differences between cells *Time point 1* and *Time point 2* (Figure 5a-f). Results  
348 were confirmed when exposing cells to IL6 antibody during both the proliferation and  
349 differentiation stage (Supplementary Figure 4g-l).

350

351 Moreover, similar to the effects of IL6 antibody during the proliferation phase on the  
352 production of IL12, treatment of cells with serum samples from delirium patients and IL6  
353 antibody during the differentiation stage prevented the production of IL13 in cell supernatant,  
354 (*Time point 1*, delirium vs delirium + IL6 antibody; IL13:  $24.6\pm 3\text{pg/ml}$  vs  $5.1\pm 2.7\text{pg/ml}$ ,  
355  $p<0.0001$ ), again with no differences between *Time point 1* and *Time point 2* (Figure 1e,  
356 Figure 5g, h).

357

358 In this case, finding seems to suggest that, during differentiation, the detrimental effect on  
359 neurogenesis and apoptosis caused by the high IL6 concentrations in serum from delirium  
360 patients is mediated by production of IL13. Indeed, treatment of cells with an antibody against  
361 IL13 prevented the decrease in neurogenesis (DCX, Map2) and increase in apoptosis (CC3)

362 caused by treatment with serum from delirium patients (*Time point 1*, delirium vs delirium +  
363 IL13 antibody; for DCX, 19.1% vs 33.8%,  $p < 0.0001$ ; for Map2, 25.1% vs 41.4%,  $p < 0.0001$ ;  
364 for CC3, 17.3% vs 3.8%,  $p < 0.0001$ ). Again, there were no differences between *Time point 1*  
365 and *Time point 2* (Figure 1c, Figure 5i-n).

366

367 **Treatment with recombinant IL12 or IL13, at concentrations previously found in**  
368 **supernatant of cells exposed to serum from patients with delirium, decreases cell**  
369 **proliferation, neurogenesis and increase apoptosis, whereas co-treatment with the JAK**  
370 **inhibitors, baricitinib, ruxolitinib and tofacitinib, prevents these detrimental effects.**

371 Having identified the downstream production of IL12 and IL13 in supernatant of cells  
372 exposed to serum from patients with delirium as the mechanism responsible for the reduced  
373 cell proliferation and neurogenesis and increased apoptosis, we decided to test whether 1)  
374 these same effects could be replicated by treating cells directly with IL12 and IL13 at the  
375 same concentrations measured in the supernatant following treatment with serum; and 2) if  
376 treatment with effective and commonly administered therapeutic compounds for patients with  
377 acute COVID-19, the JAK inhibitors baricitinib, ruxolitinib and tofacitinib (all 1nM), could  
378 prevent these detrimental effects by IL12 and IL13.

379

380 For this purpose, we treated cells directly with recombinant IL12 and IL13, used at  
381 concentrations previously found in supernatant of cells exposed to serum from patients with  
382 delirium (IL12 delirium: 20pg/ml and IL13 delirium: 25pg/ml) or without delirium (IL12  
383 without delirium: 3pg/ml, IL13 without delirium: 4pg/ml, Figure 1f, g), with or without the  
384 JAK inhibitors.

385



386 As hypothesised, during proliferation, treatment with IL12 (20pg/ml) was able to cause a  
387 decrease in cell proliferation (Ki67) and increase in apoptosis (CC3), when compared with  
388 IL12 (3pg/ml) ( for Ki67, 42.8% vs 71.6%,  $p<0.0001$ ; for CC3, 11.5% vs 5.6%,  $p<0.0001$ ,  
389 Supplementary Figure 5a, b). This is similar to what previously shown upon exposure of cells  
390 to serum from patients. Interestingly, co-treatment with baricitinib, ruxolitinib or tofacitinib  
391 were able to prevent these detrimental effects (IL12 20pg/ml vs IL12 3pg/ml +  
392 vehicle/baricitinab/ruxolitinab/tofacitinab; for Ki67, 42.8% vs 68.4%,  $p<0.001$ , vs 69%,  
393  $p<0.001$ , vs 71.4%,  $p<0.001$ ; for CC3, 11.5% vs 5.7%,  $p<0.0001$ , vs 5%,  $p<0.0001$ , vs 5.4%,  
394  $p<0.0001$ , Supplementary Figure 5a, b).

395

396 Similarly, during differentiation, treatment with IL13 (25pg/ml) was able to cause a decrease  
397 in neurogenesis (DCX and Map2) and increase in apoptosis (CC3), when compared with IL13  
398 (4pg/ml) ( for DCX, 20.8% vs 31%,  $p<0.0001$ ; for Map2, 27.2% vs 41%,  $p<0.0001$ ; for CC3,  
399 15.8% vs 7.6%,  $p<0.0001$ , Supplementary Figure 5c-e). This is, again, similar to what  
400 previously shown upon exposure of cells to serum from patients. Again, co-treatment with  
401 baricitinib, ruxolitinib or tofacitinib were able to prevent these detrimental effects (IL13  
402 25pg/ml vs IL13 4pg/ml + vehicle/baricitinab/ruxolitinab/tofacitinab; for DCX, 20.8% vs  
403 30.4%,  $p<0.001$ , vs 29%,  $p<0.001$ , vs 30%,  $p<0.001$ ; for Map2, 27.2% vs 39.1%,  $p<0.001$ ,  
404 vs 39.7%,  $p<0.001$ , vs 39.7%,  $p<0.001$ ; for CC3, 15.8% vs 7.4%,  $p<0.0001$ , vs 8.3%,  
405  $p<0.0001$ , vs 8.1%,  $p<0.0001$ , Supplementary Figure 5c-e).

406

## 407 **DISCUSSION**

408 To our knowledge, this is the first study to identify the high IL6 levels in the serum of COVID-  
409 19 hospitalised patients with neurological symptoms (delirium) as the molecular mechanism  
410 leading to disruption in neurogenic processes, measured *in vitro* using a unique human model

411 of hippocampal precursors. Interestingly, IL-6-induced increased production by these  
412 hippocampal cells of IL12 (during proliferation) and IL13 (during differentiation) is the  
413 molecular mechanisms downstream to IL6, suggesting that brain production of cytokines in  
414 response to peripheral inflammation is important in neurological symptoms induced by  
415 COVID-19 infection. Indeed, treatment with an antibody against IL12 or IL13 prevented the  
416 effect of serum from delirium patients on cell proliferation, cell differentiation and apoptosis.  
417 Of note, treatment with the therapeutic compounds commonly administered to acute COVID-  
418 19 patients (the JAK inhibitors baricitinib, ruxolitinib and tofacitinib) were able to restore  
419 normal cell viability, proliferation and neurogenesis by targeting the effects of IL12 and IL13.  
420 The novelty of our study resides in the use of a well-established but, at the same time, unique  
421 model of human hippocampal neurogenesis<sup>25, 30-34</sup>, and a validated “blood-brain-axis”  
422 experimental assay<sup>28</sup> using serum samples from COVID-19 patients.

423

424 While there is evidence showing that patients with COVID-19 present with neurological  
425 symptoms both during and after the disease (the latter phenomenon is the so-called “long  
426 COVID”)<sup>7, 8, 41</sup>, the brain mechanisms through which this occur have not been identified yet.  
427 So far there is some indication that the SARS-CoV-2 can affect transcriptionally brain cells  
428 from the parenchymal and choroid plexus, and is able to reduce hippocampal neurogenesis in  
429 animal and human models of COVID-19<sup>42-44</sup>. Similarly, there is evidence for an association  
430 between neuroinflammation and neurological symptoms during the acute and mild phase of  
431 COVID-19<sup>45-47</sup>. Our study identifies IL6 and IL6-induced IL12 and IL13 as the putative  
432 mechanisms mediating these effects. Specifically, we found a 6-fold increase in the  
433 concentrations of IL6 in serum samples of patients with delirium ( $229.9 \pm 105.2$  pg/ml), when  
434 compared with patients without delirium ( $32.5 \pm 9.5$  pg/ml). In line with our findings, higher  
435 levels of IL6 have been reported also by other studies in the periphery and in the cerebrospinal

436 fluid (CSF) of COVID-19 patients<sup>12, 13, 21, 48, 49</sup>, as well as in children with acute encephalitis-  
437 like syndrome during infection with coronavirus-OC43<sup>50</sup>. Overall, these findings identify IL6  
438 an important therapeutic target not only for COVID-19 symptoms in general, but also for those  
439 neurological manifestations which are commonly observed as a consequence of the disease.  
440 Interestingly, this result is consistent with our recent study showing that exposing the same  
441 cells to concentrations of IL6 ranging from 50 to 5000pg/ml dramatically decrease  
442 neurogenesis and increase apoptosis<sup>22</sup>. Likewise, previous findings generated in other  
443 laboratories also demonstrated that exposure to concentrations of IL6 similar to those detected  
444 in the present study is associated with a significant reduction in neurogenesis and increase in  
445 apoptosis, both *in vitro* and *in vivo*<sup>51-53</sup>.

446

447 Of note, treatment with serum from delirium patients induced our cells to produce two well-  
448 known inflammatory cytokines, IL12 and IL13, in the supernatant; these cytokines were in  
449 very low quantities in serum, indicating a direct production by cells. The causal role of IL6  
450 and IL6-induced IL12 and IL13 in mediating the effect of serum from delirium patients on  
451 the neurogenic outcomes is confirmed by the use of an antibody against these cytokines in  
452 our experiments. Interestingly, IL12 was produced only during the proliferation stage,  
453 whereas IL13 only during the differentiation stage, but always as a consequence of IL6  
454 stimulation. Interestingly, while both IL12 and IL13 are expressed in the brain<sup>54, 55</sup>, IL12 is  
455 primarily known to regulate cell proliferation<sup>54</sup>, whereas IL13 is mainly involved in  
456 differentiation-related processes<sup>55</sup>. Indeed, IL12 reduces cell proliferation and migration in  
457 several neuronal cultures via activation of distinct pro-inflammatory signalling pathways,  
458 including janus-activated kinase and signal transducers and activators of transcription (JAK-  
459 STAT) pathway<sup>56, 57</sup>, which we have also previously identified to be involved in the regulation  
460 of neuronal apoptosis upon exposure of hippocampal cell to IFN- $\alpha$  treatment<sup>24</sup>. Accordingly,

461 previous evidence has reported an increase in the production of IL12 and IL13 in the CSF of  
462 COVID-19 patients with neurological manifestations<sup>58, 59</sup>. Similarly, in mice, studies have  
463 shown that increased cerebral expression of IL12 is able to induce neurological symptoms,  
464 and that these symptoms are worsen by an infectious disease<sup>54</sup>. Likewise, in humans, IL13  
465 expression has shown to be associated with the development of neurological disorders, and  
466 with the experience of more severe neuropsychiatric symptoms. In particular, individuals with  
467 high IL13 expression are more likely to develop neurological diseases, including  
468 Parkinson's<sup>60</sup>, whereas in those individuals who already suffer from a major depressive  
469 disorder, higher IL13 levels are associated with more severe depression and a higher number  
470 of suicide attempts<sup>61</sup>.

471

472 Our study identifies multiple pharmacological approaches that could be beneficial for  
473 treatment of neurological symptoms in the. context of COVID-19 infection, and thus  
474 potentially in other conditions associated with high levels of IL6. First, IL6 itself can be  
475 considered a valuable therapeutic target in treating COVID-19 patients. Indeed, treatment  
476 with IL6 receptor antagonists, such as tocilizumab or sarilumab, have so far shown a relatively  
477 good level of efficacy in reducing COVID-19 mortality rates, although these compounds have  
478 a high molecular weight, making them unable to act centrally<sup>62</sup>. Second, there are several  
479 IL12 inhibitors available, such as ustekinumab and briakinumab, which have already been  
480 tested for their efficacy and tolerability in clinical context<sup>63</sup>. However, these compounds are  
481 not exclusively targeting IL12, but rather the p40 subunit of both IL12 and IL23, and have a  
482 relatively high molecular weight, making them less likely to exert their properties centrally<sup>64</sup>.  
483 The situation is similar for IL-13: although a selective inhibitor, lebrikizumab, has already  
484 been developed and tested in a phase 2b trial, this compound also has a very high molecular  
485 weight, making it unsuitable for treatment of brain inflammatory neuropathologies<sup>65</sup>. Finally,

486 we have shown that the JAK inhibitors, baricitinib, ruxolitinib and tofacitinib, prevent the  
487 reduction in cell proliferation and neurogenesis, and the increase in apoptosis, caused by  
488 treatment with recombinant IL12 or IL13. Recently randomized, placebo-controlled trials of  
489 these have shown these compounds to be among the most effective treatment interventions  
490 for acute COVID-19 patients, especially for those with a high level of inflammation<sup>66-68</sup>.  
491 Moreover, these JAK inhibitors have low molecular weight (< 400Da) making them suitable  
492 to act centrally, and potentially on mechanisms linked to the neuronal production of IL12 and  
493 IL13 that we describe. All together, we propose that baseline stratification of patients based  
494 on their level of inflammatory biomarkers, including IL6, IL12 and IL13, or the presence of  
495 neurological symptoms, may represent the best therapeutic approach for those individuals  
496 experiencing severe COVID-19 infection.

497

498 We acknowledge several limitations associated with our study, including the use of an *in vitro*  
499 system of immortalized cell line. However, while theoretically this system may differ from  
500 the scenario of an adult *in vivo* environment and the adult neurogenic niche, in particular  
501 because of the absence of microglia cells and the fetal status of the immortalized cells, we  
502 have extensive confirmation from previous studies that our results are relevant to the human  
503 and animal brains. Indeed, over the years we have been able to replicate all our results  
504 obtained in this *in vitro* model in both the animal brain and the human blood, including  
505 changes in neurogenesis by cortisol, antidepressants, IL1 $\beta$ , IFN- $\alpha$ , and serum samples from  
506 depressed patients, and changes in stress-, antidepressants- and inflammation-regulated  
507 mRNA gene expression in both the hippocampal mRNA of animal models of depression and  
508 the whole blood mRNA of depressed patients<sup>22, 25, 28, 30-33, 40, 69, 70</sup>. Nevertheless, in future  
509 studies, we aim to expand this cellular model using both neurons and microglia co-cultures,  
510 in order to create a more comprehensive and realistic experimental scenario. Also, our study

511 does not investigate the direct effects of COVID-19 on these cells. However, considering that  
512 there is evidence showing that the virus has difficulty crossing the BBB in humans, and that  
513 its central effect is instead mediated by the activation of peripheral mechanisms<sup>71-73</sup>, we  
514 believe that our experimental approach, exposing brain cells to serum from COVID-19  
515 infected patients rather than to the virus itself, provides mechanistic understanding on how  
516 COVID-19 indirectly regulates brain function. Finally, the number of samples used in this  
517 study, although matched for age and sex, was also relatively small, 18 patients per group.  
518 Therefore, the results presented here should be replicated in a bigger cohort of COVID-19  
519 patients with and without delirium, including patients with long COVID, who are also known  
520 to experience persistent neurological manifestations<sup>41</sup>. However, while the sample size is  
521 small, this is the first study demonstrating for the first time that it is the elevated peripheral  
522 concentration of IL-6 (rather than of other pro-inflammatory cytokines) that clinically affects  
523 cognition and mechanistically affects neurogenesis in the same COVID-19 patients with  
524 delirium. In addition, the identification of the IL12 and IL13 as downstream mediators is a  
525 completely novel mechanism for the IL-6-induced reduction in neurogenesis, and the novel  
526 experiments using JAK inhibitors performed in the present study offer additional mechanistic  
527 insights into these pathways.

528

529 In summary, our *in vitro* study demonstrates that that serum from COVID-19 patients with  
530 delirium can detrimentally affect human hippocampal neurogenesis, and that this effect is  
531 mediated by IL6-induced production of the downstream inflammatory cytokines IL12 and  
532 IL13, which are ultimately responsible for the negative cellular outcomes. Understanding the  
533 mechanisms through which neurogenesis is altered in COVID-19 patients with neurological  
534 symptoms will ultimately contribute to the development of novel interventional and

535 preventative strategies for this particular sub-group of patients, and potentially for patients with  
536 cognitive and neurological symptoms in the context of inflammation.

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592

593 **Author Contributions**

594 AB, BM, JE and CMP designed research; AB performed research and analysed data; AB, BM,  
595 JE, GM, DS, ACV, GN, ST and CMP wrote the paper.

596

597 **Statement of Interest**

598 Dr Alessandra Borsini, Professor Carmine M. Pariante and Professor Sandrine Thuret have  
599 received research funding from Johnson & Johnson for research on depression and  
600 inflammation which included cellular work (2012-2018), but this work is unrelated to that  
601 funding; moreover, less than 10% of Professor Pariante’s support in the last 10 years derives  
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608 Dohme Corp. The work presented in this paper is unrelated to these awards. Dr Blair Merrick,  
609 Dr Jonathan Edgeworth, Dr Deepak Srivastava, Dr Anthony C. Vernon and Dr Gaia Nebbia  
610 have no conflicts of interest to declare.

611

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911 **Figures Caption**

912 **Table 1. Characteristics of COVID-19 patients with and without delirium.** Demographics of the  
913 COVID-19 patients' cohort. Legend: \* Mann-Whitney U-test; \*\* Chi squared test; # Severity Score:  
914 0=Asymptomatic or no requirement for supplemental oxygen, 1= Requirement for supplemental  
915 oxygen (fraction of inspired oxygen (FiO<sub>2</sub>) < 0.4) for at least 12 h, 2= Requirement for supplemental  
916 oxygen (FiO<sub>2</sub> ≥ 0.4) for at least 12 h, 3= Requirement for supplemental oxygen (FiO<sub>2</sub> > 0.6) for at least  
917 12 h, and not a candidate for escalation above level one (ward-based) care, 4= Requirement for  
918 supplemental oxygen (FiO<sub>2</sub> > 0.8) and peripheral oxygen saturations <90% (with no history of type 2  
919 respiratory failure (T2RF)) or <85% (with known T2RF) for at least 12 h, 5= Requirement for ECMO;  
920 ## Comorbidities: hypertension, type 2 diabetes mellitus, (paroxysmal) atrial fibrillation aortic stenosis,  
921 benign prostatic hypertrophy/ hyperplasia, benign paroxysmal positional vertigo, cancer, congestive  
922 cardiac failure, chronic kidney disease, chronic obstructive pulmonary disease, gastroesophageal reflux  
923 disease, ischaemic heart disease, monoclonal gammopathy of undetermined significance, osteoarthritis,  
924 osteoporosis, peripheral vascular disease, rheumatoid arthritis, transient ischaemic attack, venous  
925 thromboembolism.  
926

927 **Figure 1. Timeline of *in vitro* experiments with the same serum samples.** (a-e) Cells were treated  
928 with media containing 1% serum (from each COVID-19 patient, delirium and non-delirium) with or  
929 without IL6 Antibody (A) (0.1ug/ml) or IL12A (0.3ug/ml) for additional 2 days. At this stage cells were  
930 either fixed or left to differentiate for additional 4 days, again in presence of 1% serum (from the same  
931 COVID-19 patient previously used in the proliferation phase), with or without IL6A or IL13A  
932 (0.1ug/ml). At the end of day 2 of proliferation and day 4 of differentiation, cell supernatant was  
933 collected for subsequent cytokines measurement. Legend: Sox2, sex determining region Y-box 2; DCX,  
934 doublecortin; Map2, microtubule-associated protein 2; S100β, S100 calcium-binding protein β; CC3,  
935 caspase 3; interleukin, IL.  
936

937 **Figure 2. Cell treatment with serum samples from COVID-19 patients with delirium decreased  
938 cell proliferation, neurogenesis, and increased apoptosis, when compared with serum from non-  
939 delirium patients.** (a-b) Cell treatment with serum samples from delirium patients decreased  
940 proliferation (KI67+cells) and increased apoptosis (CC3+ cells), when compared with serum from non-  
941 delirium patients. There were no differences across Time point 1 and 2. (c-e) Treatment with serum  
942 samples from delirium patients decreased neurogenesis (DCX+ and Map2+cells) and increased  
943 apoptosis (CC3+ cells), when compared with serum from non-delirium patients. Again, there were no  
944 differences across Time point 1 and 2. Two-way mixed ANOVA was performed.  
945

946 **Figure 3. Higher concentration of IL6 in serum samples from COVID-19 patients with delirium  
947 induced cells to produce IL12 and IL13, respectively during the proliferation and differentiation  
948 stage.** (a-b) Serum samples from delirium patients showed increased levels of IL6, when compared  
949 with non-delirium patients, both at Time point 1 and 2. (c-d) During proliferation, exposure of cells  
950 with serum samples from delirium patients (collected both at Time point 1 and 2) induced the production  
951 of IL12 in the supernatant, when compared with treatment with serum samples from non-delirium

952 patients. (e-f) During differentiation, exposure of cells with serum samples from delirium patients  
953 (collected both at Time point 1 and 2) induced the production of IL13 in the supernatant, when  
954 compared with treatment with serum samples from non-delirium patients. Mann-Whitney U-test and  
955 two-way mixed ANOVA were performed. Data are shown as mean; \*\*\*\*p<0.0001 comparisons as  
956 indicated.

957  
958 **Figure 4. During proliferation, treatment with an antibody against IL6 prevented the detrimental**  
959 **effect of serum from COVID-19 patients with delirium on both cell proliferation and apoptosis**  
960 **and decreased IL12 production. (a-d)** Cell treatment with IL6A (0.1ug/ml) prevented the decrease in  
961 proliferation (Ki67+cells) and increase in apoptosis (CC3+cells) previously observed upon treatment  
962 with serum samples from patients with delirium. This is confirmed across treatment with serum samples  
963 collected at Time point 1 and 2. (e, f) Treatment with IL6A also prevented production of IL12 in cell  
964 supernatant. (g-j) Cell treatment with IL12A (0.3ug/ml) prevented the decrease in proliferation  
965 (Ki67+cells) and increase in apoptosis (CC3+cells) previously observed upon treatment with serum  
966 samples from patients with delirium. Again, this is confirmed across treatment with serum samples  
967 collected at Time point 1 and 2. Two-way mixed ANOVA with Bonferroni's post hoc test was  
968 performed. Data are shown as mean; \*\*\*\*p<0.0001 comparisons as indicated.

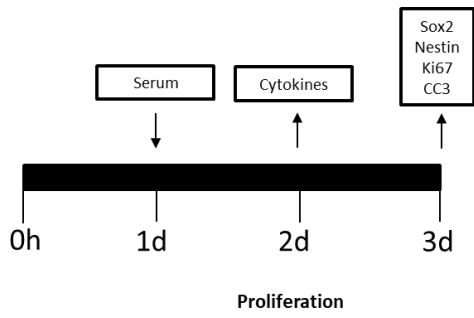
969  
970 **Figure 5. During differentiation, treatment with an antibody against IL6 prevented the**  
971 **detrimental effect of serum from COVID-19 patients with delirium on both neurogenesis and**  
972 **apoptosis and decreased IL13 production. (a-f)** Cell treatment with IL6A (0.1ug/ml) prevented the  
973 decrease in neurogenesis (DCX+ and Map2+cells) and increase in apoptosis (CC3+cells) previously  
974 observed upon treatment with serum samples from patients with delirium. This is confirmed across  
975 treatment with serum samples collected at Time point 1 and 2. (g, h) Treatment with IL6A also  
976 prevented production of IL13 in cell supernatant. (i-n) Cell treatment with IL13A (0.1ug/ml) prevented  
977 the decrease in neurogenesis (DCX+ and Map2+cells) and increase in apoptosis (CC3+cells) previously  
978 observed upon treatment with serum samples from patients with delirium. Again, this is confirmed  
979 across treatment with serum samples collected at Time point 1 and 2. Two-way mixed ANOVA with  
980 Bonferroni's post hoc test was performed. Data are shown as mean; \*\*\*\*p<0.0001 comparisons as  
981 indicated.

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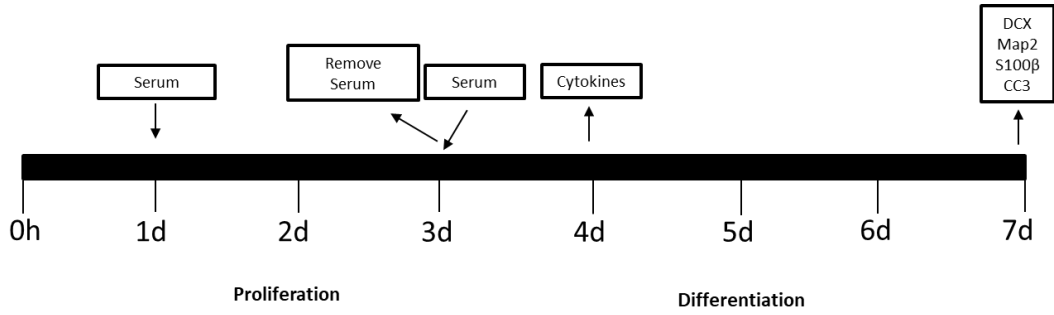


	<b>Delirium</b>	<b>Non Delirium</b>	
	<i>(n = 18)</i>	<i>(n = 18)</i>	<b><i>p value</i></b>
<b>Age (years)</b>			
Mean±SD	76.2±14.7	75.4±14.2	<i>p=0.9*</i>
<b>Gender</b>			
Male	12 (66.7%)	11 (61.1%)	<i>p=0.7**</i>
<b>Severity Score<sup>#</sup></b>			
Mean±SD	1.2±1.2	1.3±1.2	<i>p=0.9*</i>
<b>Survival</b>			
Yes	13 (72.2%)	16 (88.9%)	<i>p=0.2**</i>
<b>Two or more comorbidities<sup>##</sup></b>			
Baseline	15 (83.3%)	14 (77.8%)	<i>p=0.3**</i>
<b>Hypertension</b>			
Baseline	13 (72.2%)	12 (66.7%)	<i>p=0.7**</i>
<b>Type 2 Diabetes</b>			
Baseline	5 (27.8%)	8 (44.4%)	<i>p=0.3**</i>
<b>CRP Time point 1</b>			
Mean±SD	72±53.9 mg/L	115.5±94.4 mg/L	<i>p=0.2*</i>
<b>CRP Time point 2</b>			
Mean±SD	83.7±88.4 mg/L	97.5±102.7 mg/L	<i>p=0.9*</i>

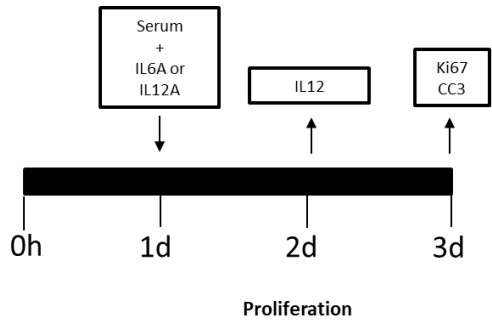
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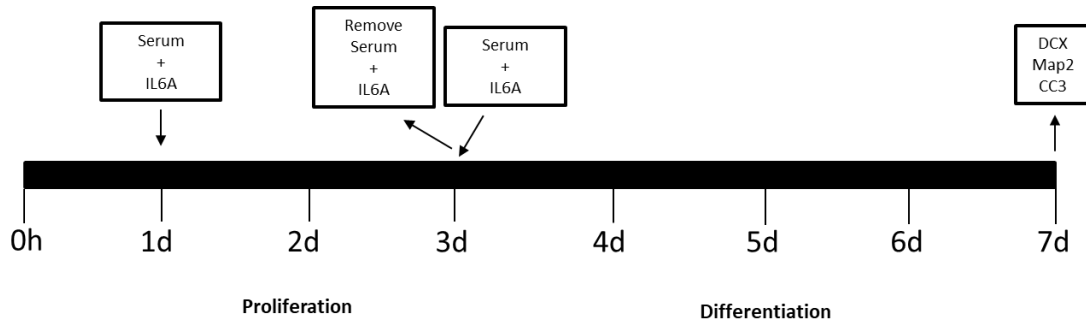
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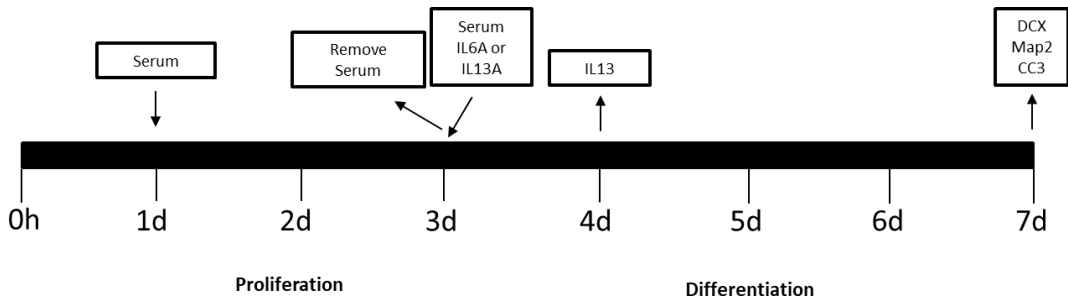
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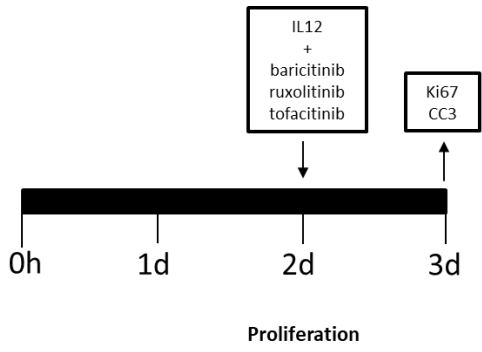
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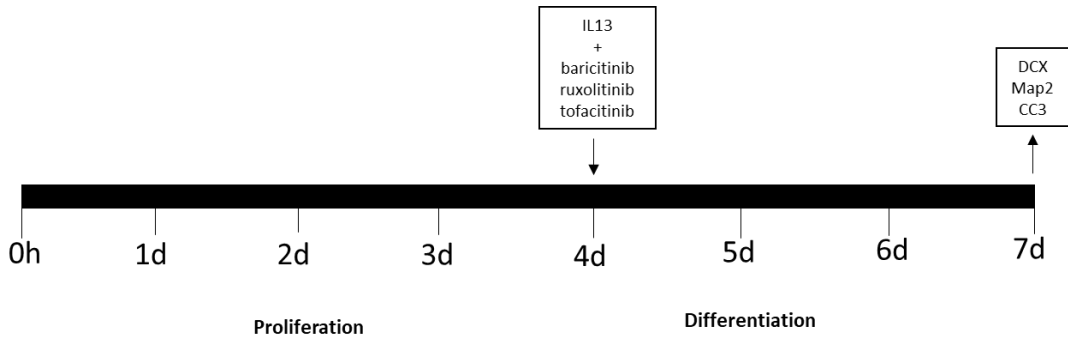
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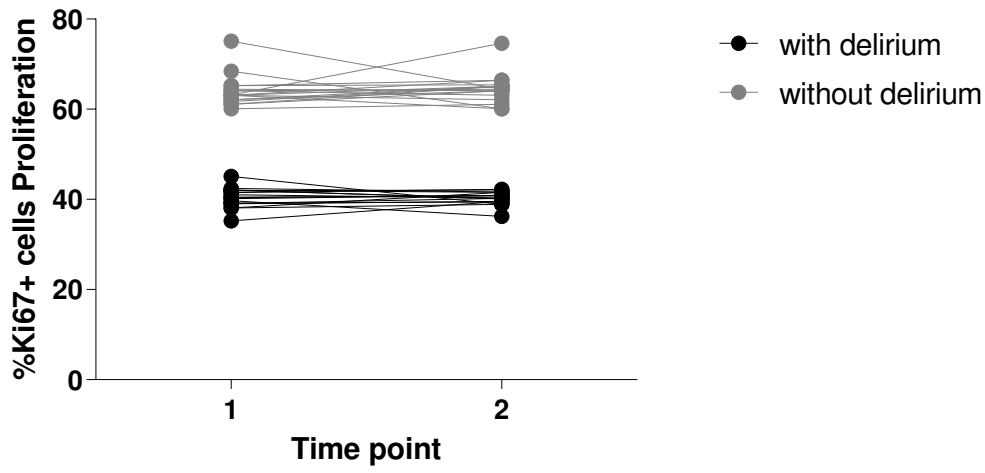
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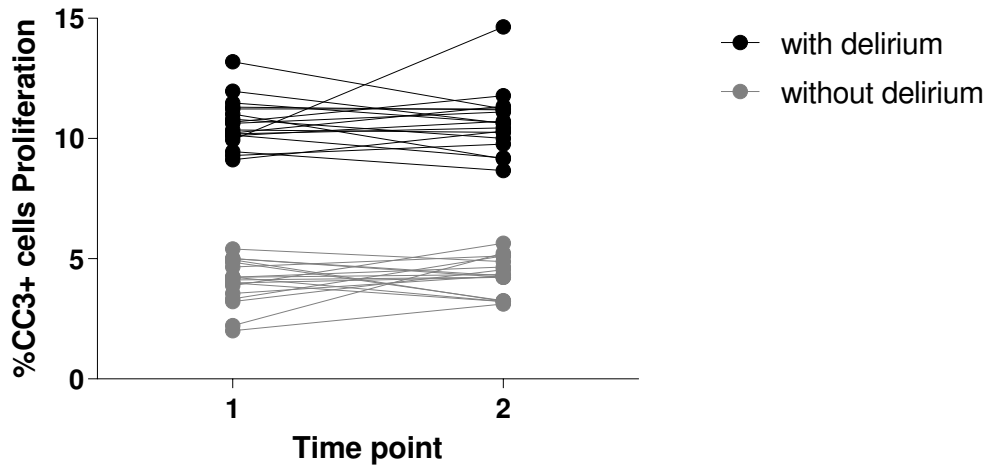
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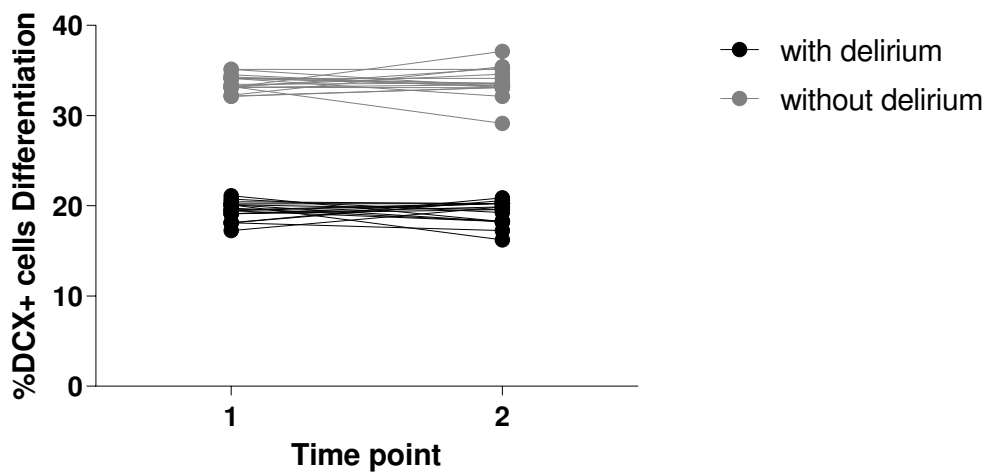
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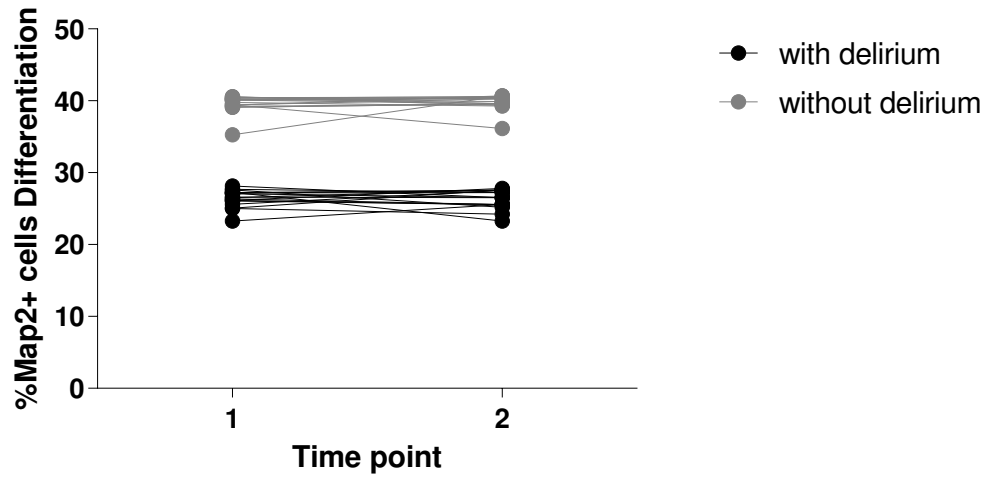
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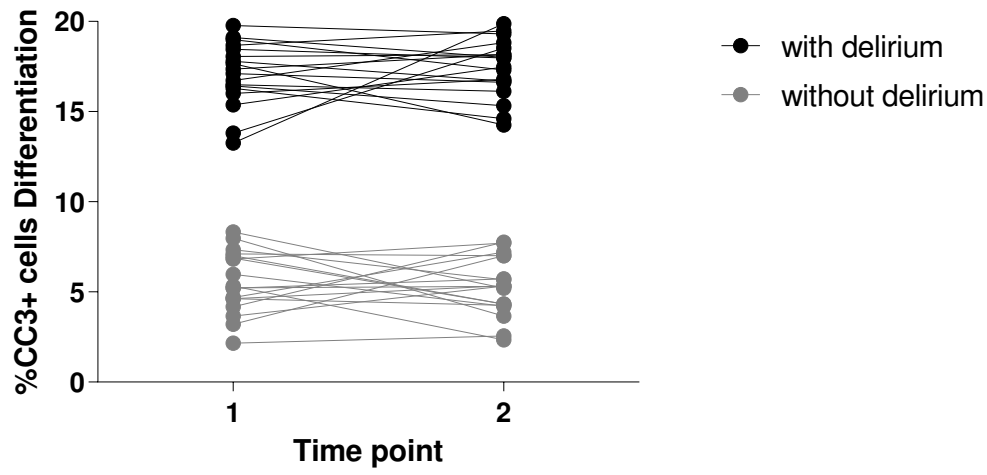
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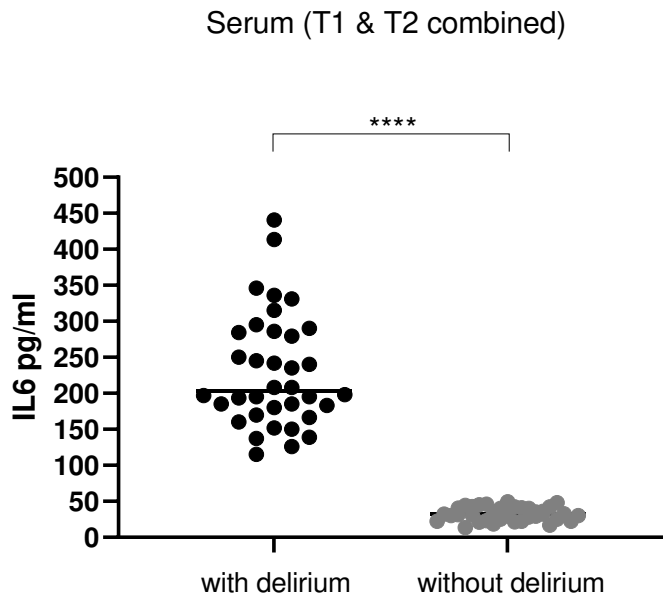
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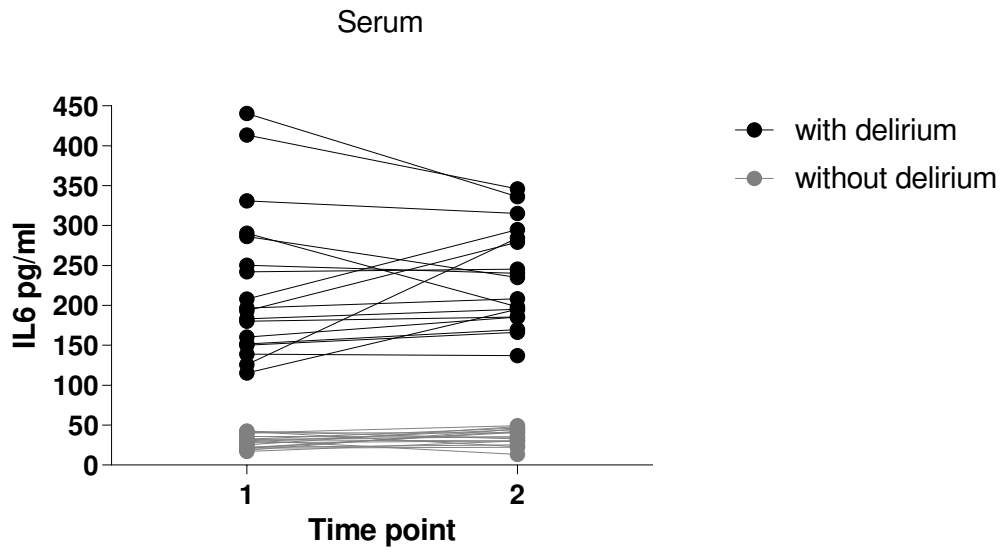
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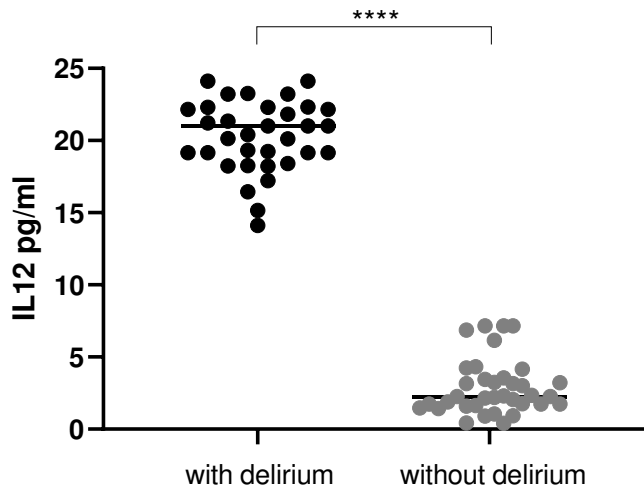


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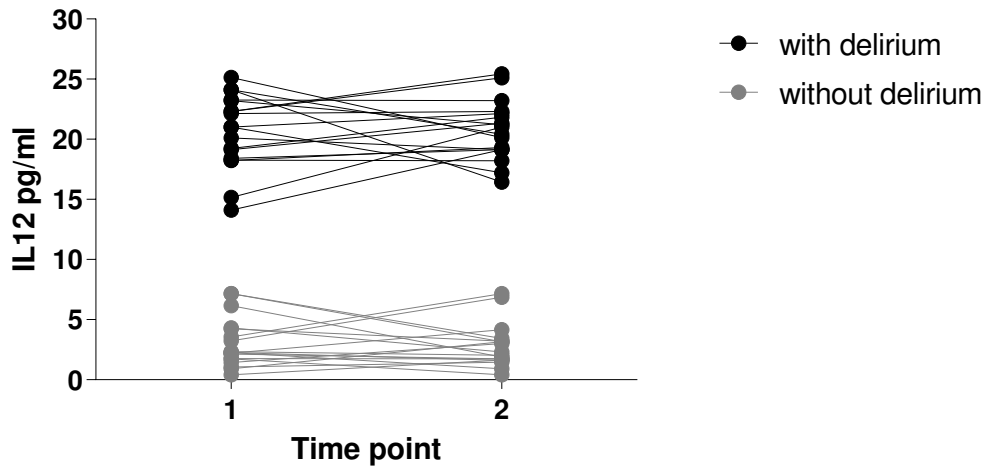
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Supernatant - Proliferation (T1 & T2 combined)



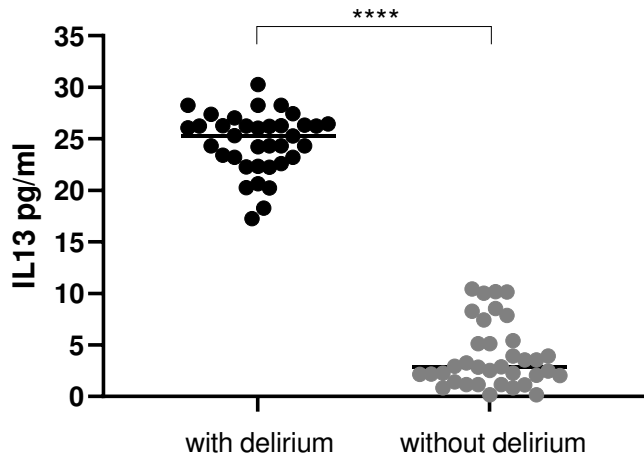
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Supernatant - Proliferation



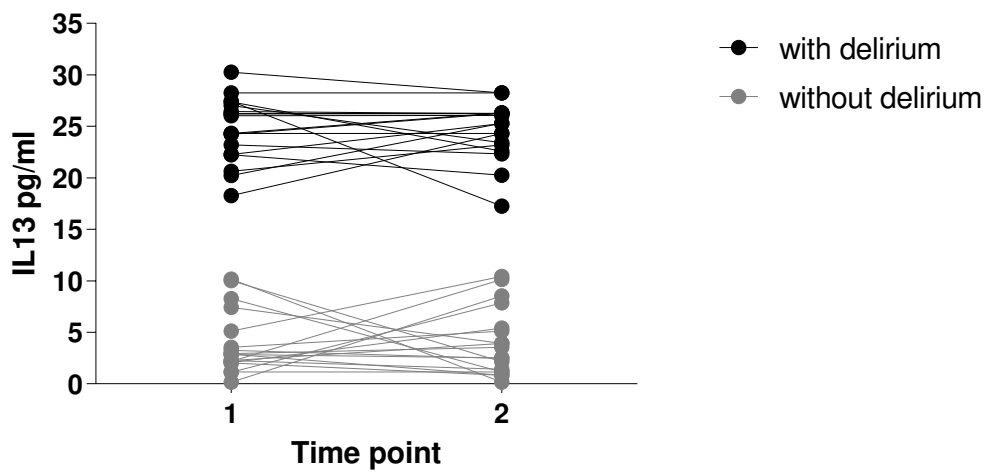
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Supernatant - Differentiation (T1 & T2 combined)



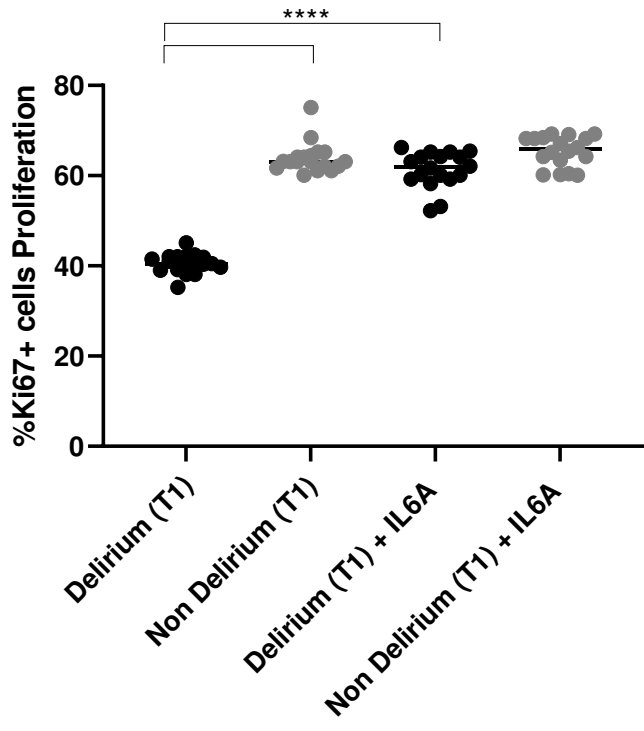
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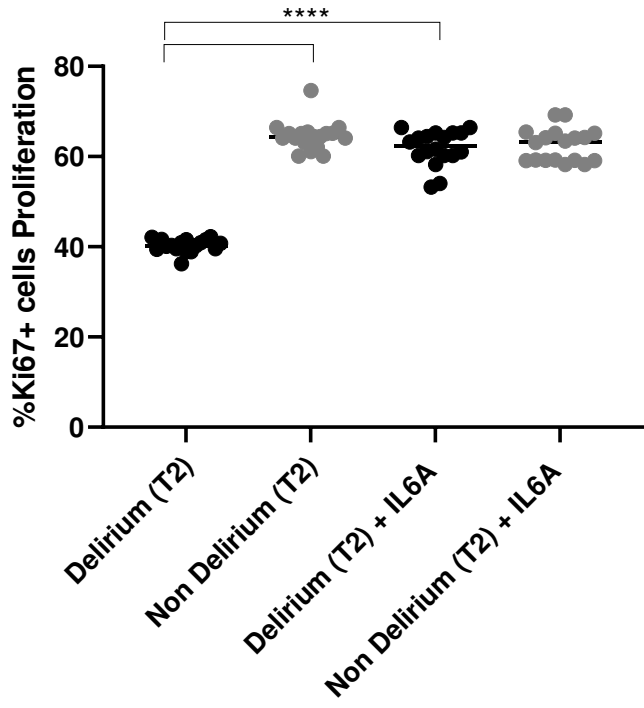




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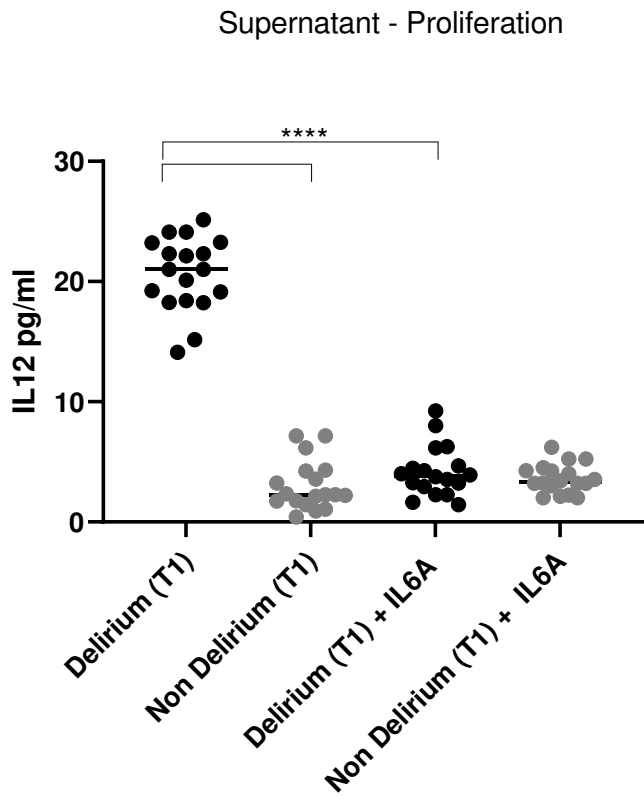


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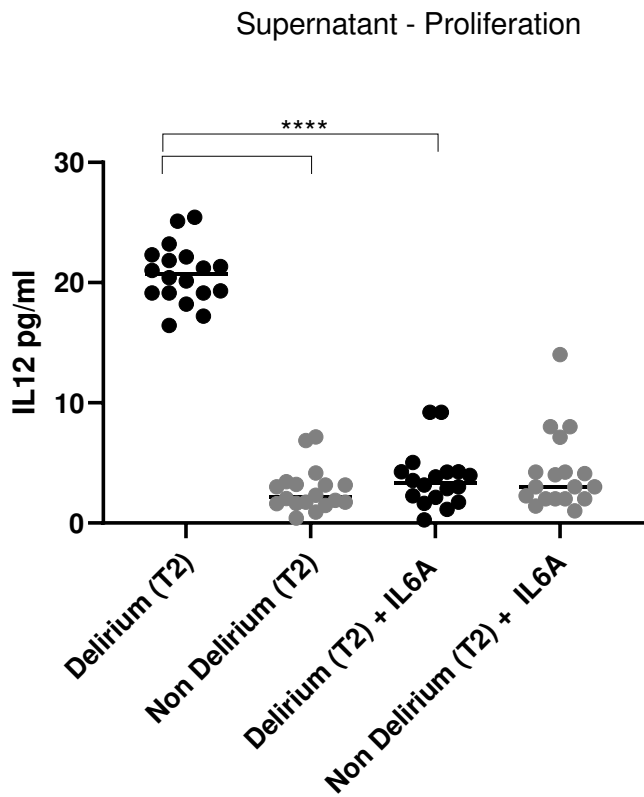




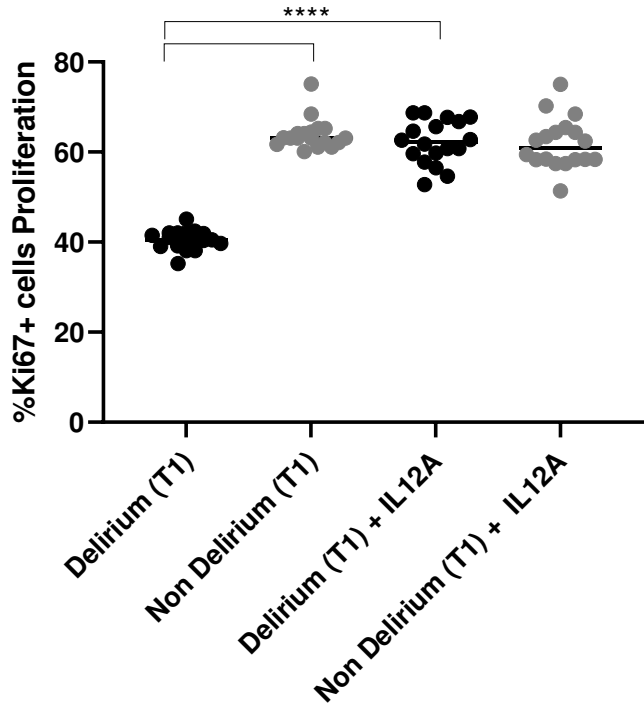
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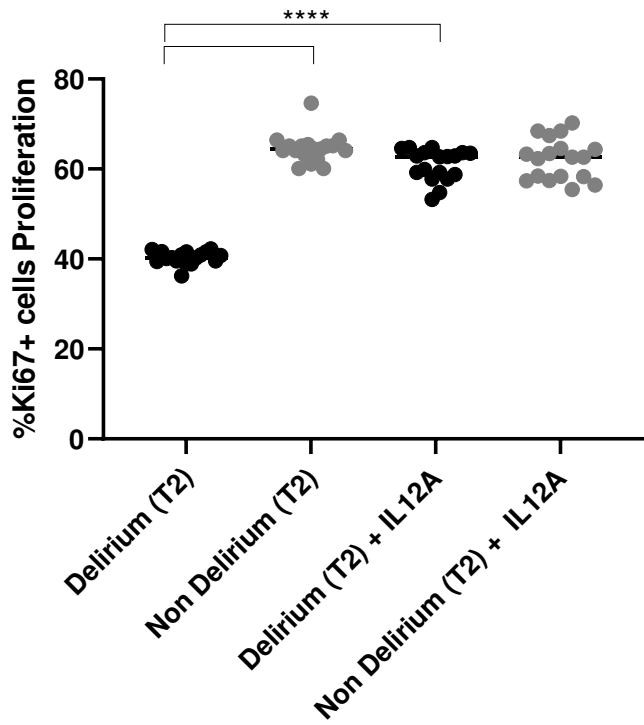
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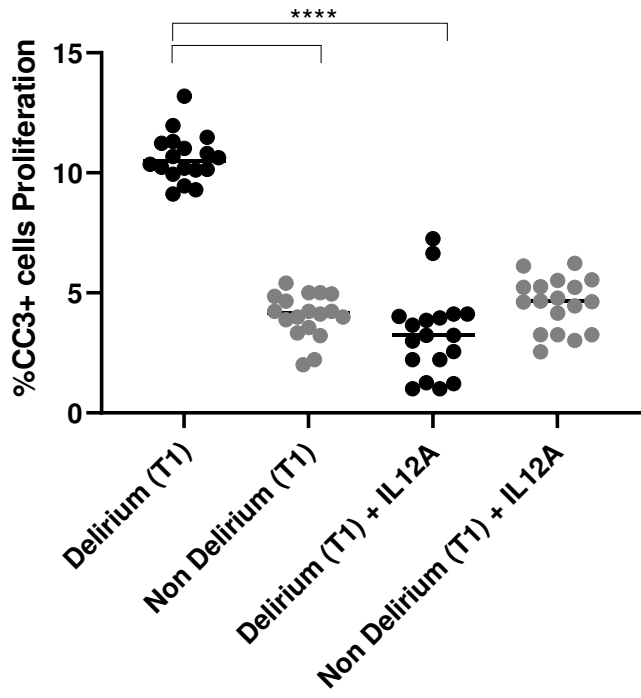
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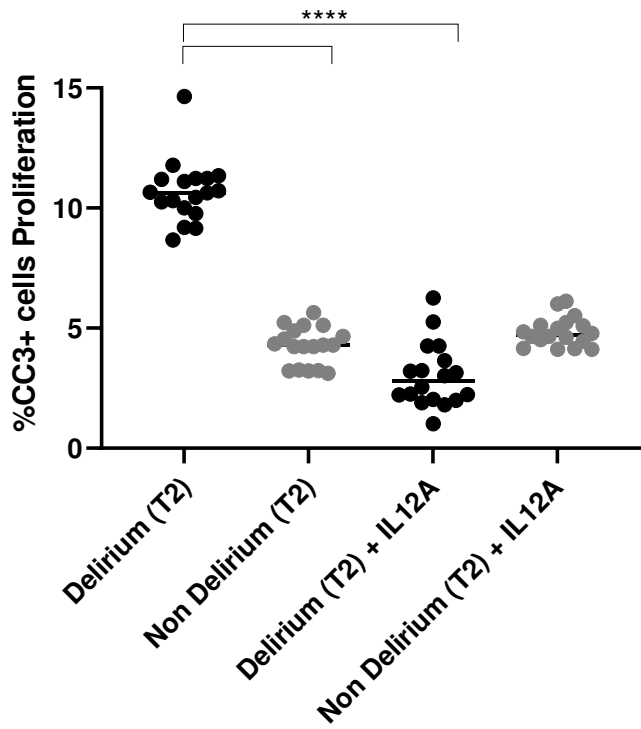
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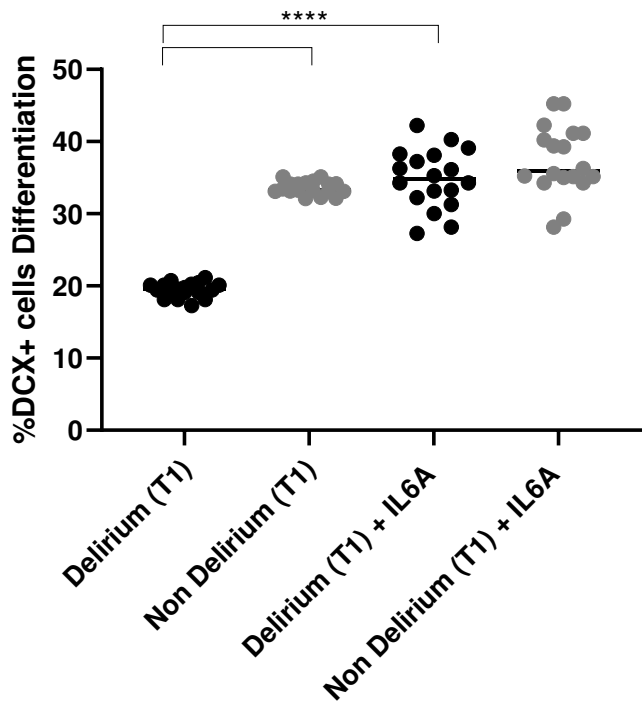
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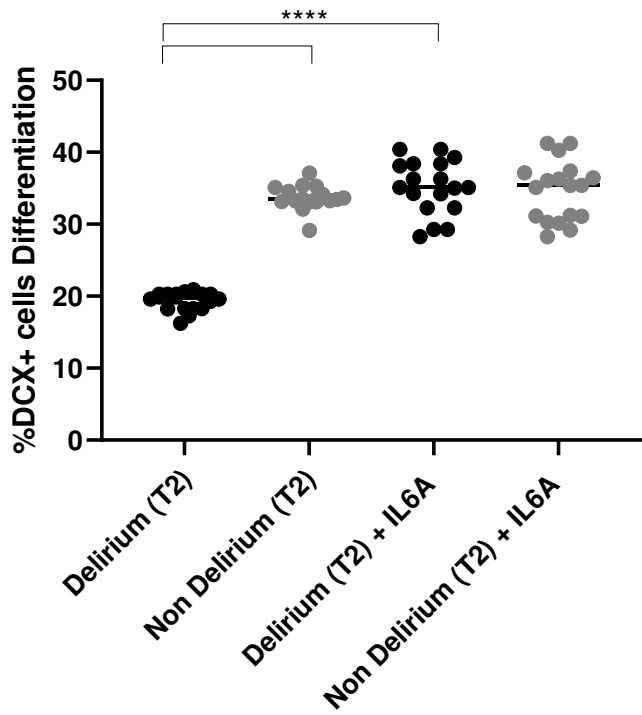
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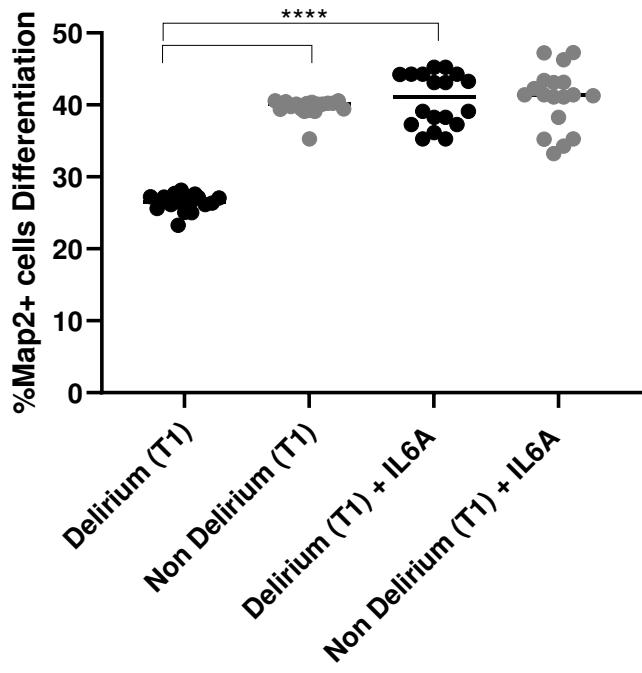
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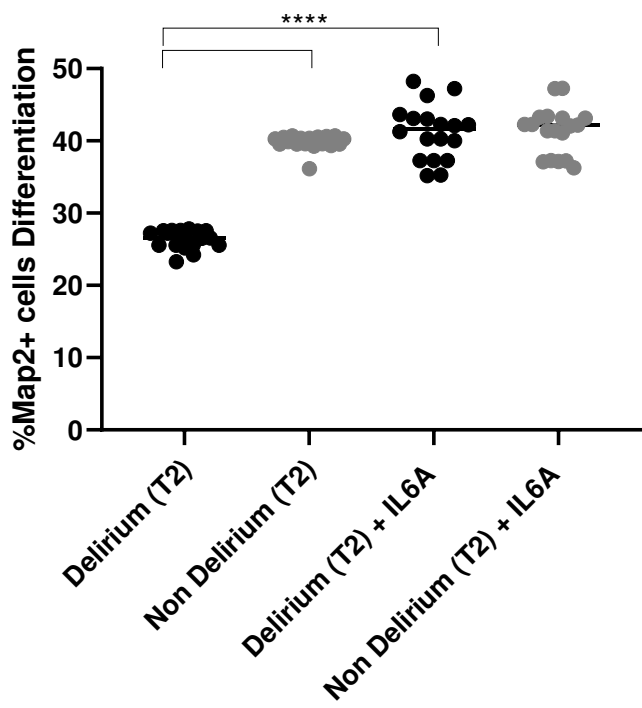
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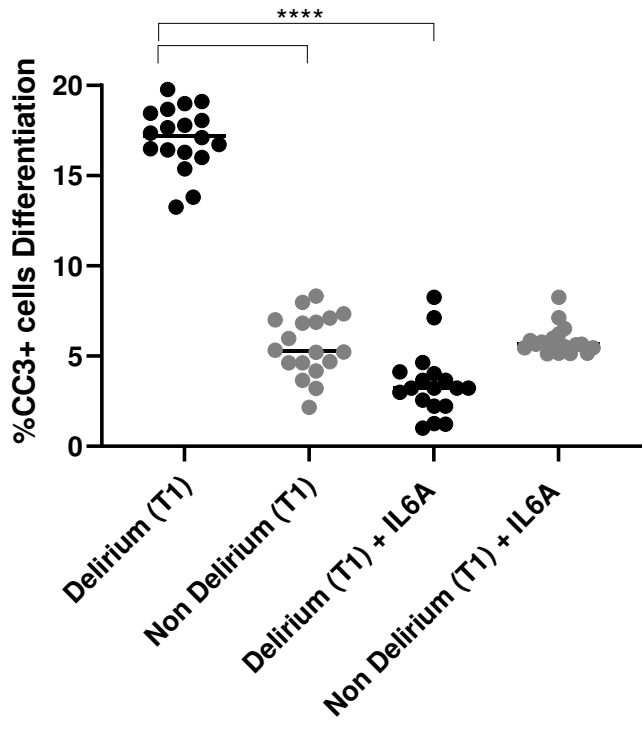
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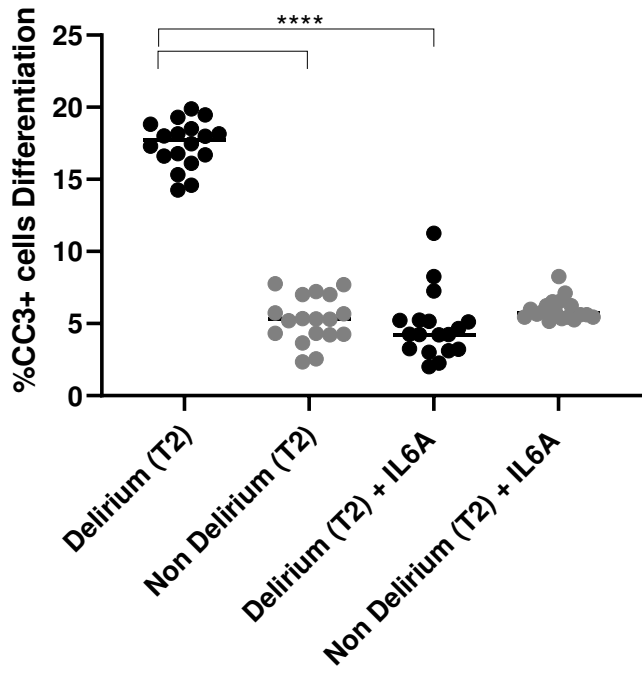
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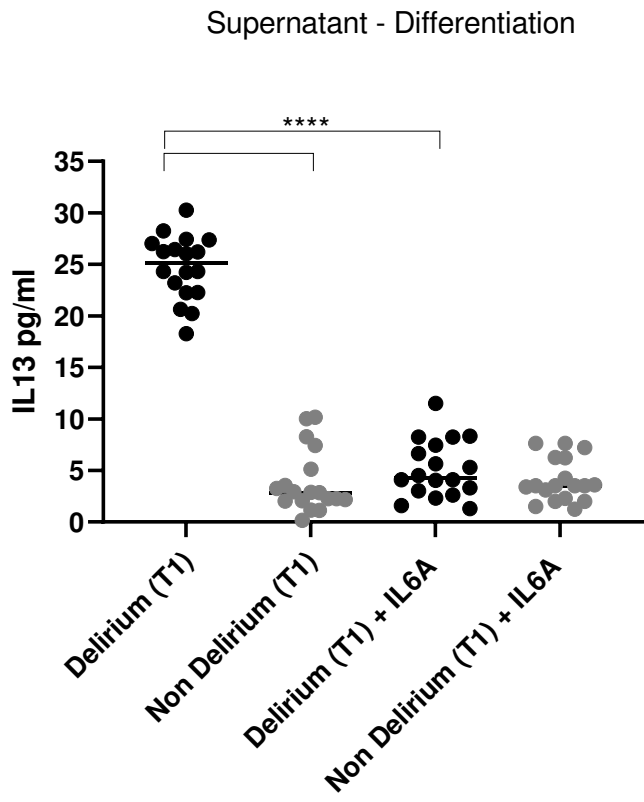


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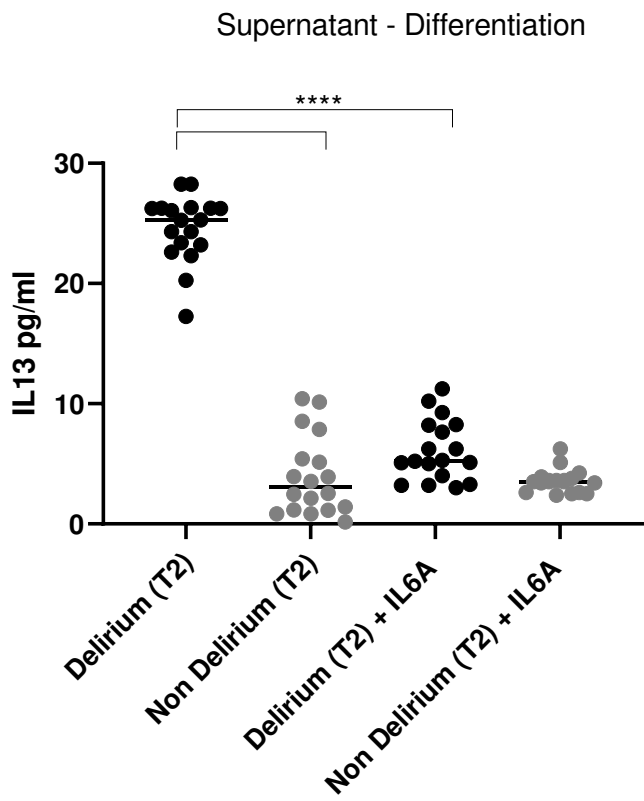




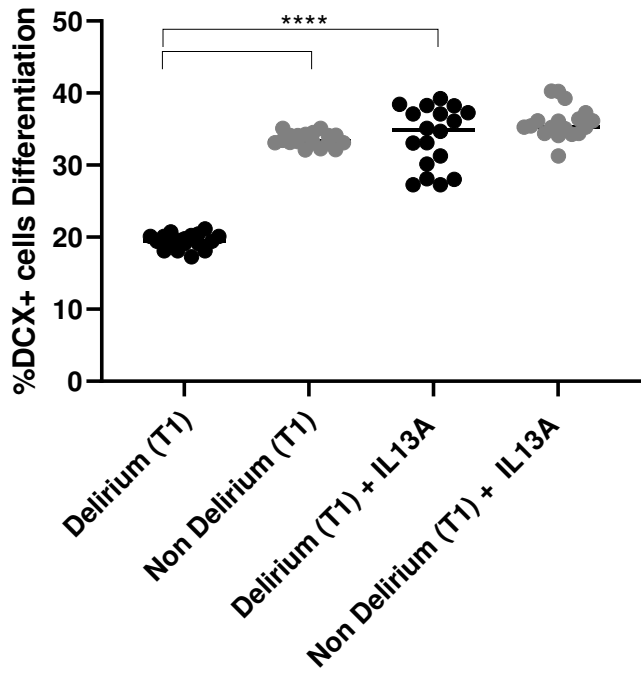
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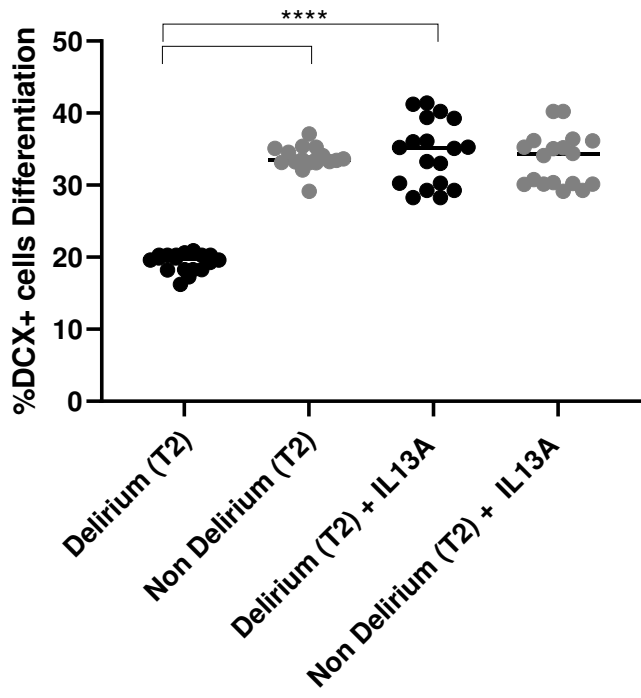
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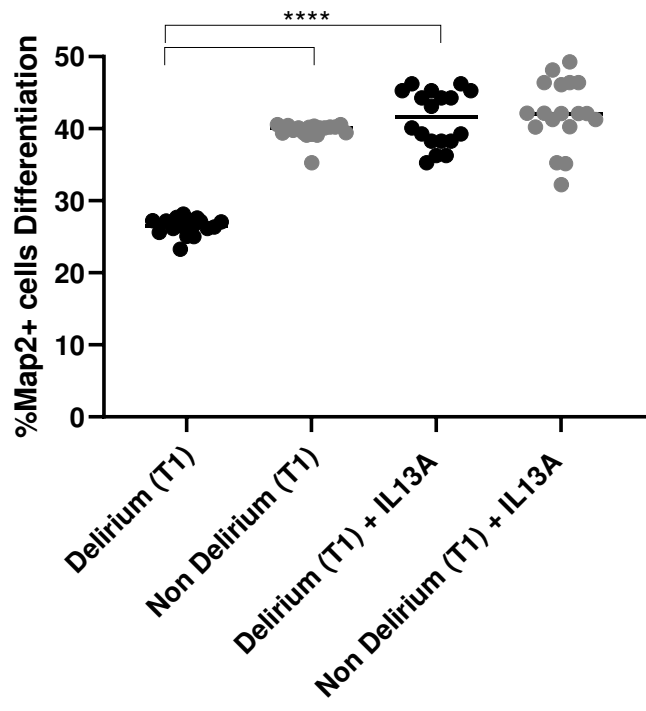
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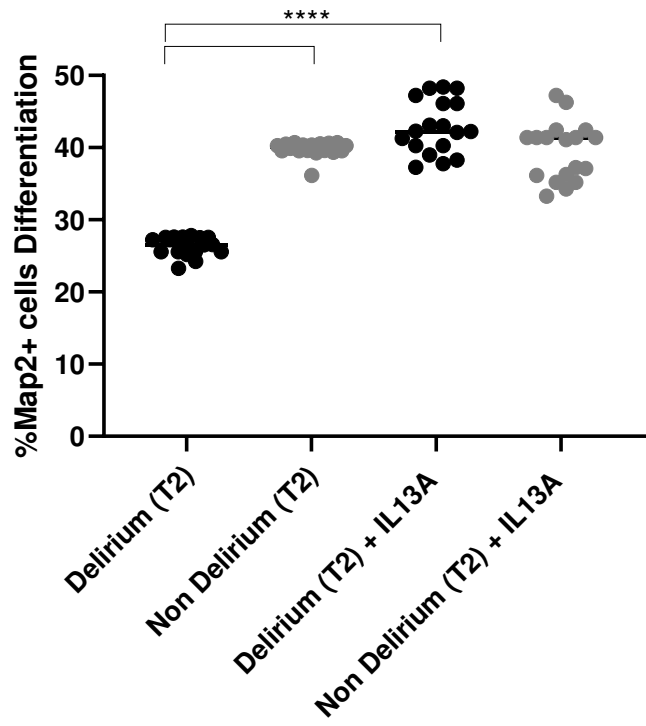
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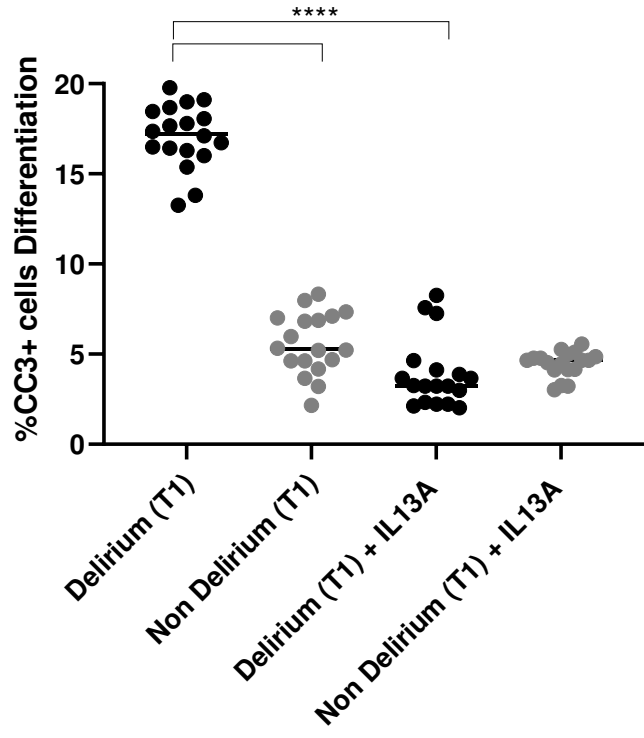
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