

1 Dynamic changes in innate immune and T cell function and composition at the nasal
2 mucosa across the human lifespan

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24 **Abstract**

25 The very young and very old are at increased risk of serious infections, including
26 pneumonia. This may relate to changes in the immune system as young children have
27 limited immunological memory, while immunosenescence, inflammaging and a decreased
28 pool of naïve immune cells are described with advanced age. How the immune system
29 changes with age at mucosal surfaces, from where infections frequently develop, is not
30 very clear as access to human tissue samples is limited. Therefore, we aimed to assess
31 the composition and activation state of the immune system at the human mucosa. Here,
32 we profiled nasal immune cells from 207 individuals between 1 to 80 years old using flow
33 cytometry. Neutrophil and monocyte functionality were measured using whole blood
34 assays. Levels of thirty nasal cytokines were measured from nasal lining fluid.
35 Nasopharyngeal colonization by *Streptococcus pneumoniae* was assessed using
36 classical microbiology and associated with immune responses. We found that young
37 children have a striking paucity of granulocytes at the nasal mucosa compared to adults.
38 In addition, T cell numbers at the nasal mucosa decreased progressively with age and
39 were almost absent in older adults. While nasopharyngeal colonization by *Streptococcus*
40 *pneumoniae* was associated with elevated levels of inflammation it had a limited effect on
41 nasal immune composition, including levels of monocytes and neutrophils. These results
42 show that the immune system at the nasal mucosal surface changes drastically with age
43 and provides explanations for the increased susceptibility to infections in young and old
44 age.

45 **Significance statement**

46 How the immune system changes with age is an intensive area of research, but has been
47 primarily studied in blood. However, blood poorly reflects the immune system at the
48 mucosa, from where infections develop. This manuscript provides a first characterization
49 of how the composition and function of the immune system in the upper respiratory tract
50 changes with age, providing explanations for increased susceptibility to infection in the
51 very young and old. Furthermore, by linking mucosal and systemic measurements with
52 pneumococcal colonization, we observed that reduced monocyte and neutrophil
53 responses associate with the increased burden of pneumococcal colonization in children.
54 This study highlights the need to study the immune system also at other mucosal sites in
55 the context of aging.

56 **Abbreviations:**

57 Spn; *Streptococcus pneumoniae*

58 MPO; Myeloperoxidase

59 LPS; Lipopolysaccharide

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62 **Introduction**

63 Pneumonia is the most common infectious cause of death in children under 5 worldwide
64 (1). Individuals with advanced age are also at progressively increasing risk of acquiring
65 pneumonia (2, 3). Over the next two decades, the incidence of community-acquired
66 pneumonia is expected to double in the United States as the population ages (4). How
67 alterations in the mucosal immune system with age predispose to infections in the very
68 young and very old remains unclear as access to samples is limited. While many studies
69 have investigated how the immune system changes with age, most of these have been
70 conducted in blood, which poorly reflects the immune system at mucosal surfaces (5, 6).
71 Nonetheless, studies from blood and secondary lymphoid tissues have revealed
72 alterations in cell numbers in blood of young children and reduced immunological memory
73 in children, while increased inflammation, immunosenescence and reduced naïve
74 memory pools have been described with advanced age (5, 7-10). Interestingly, mouse
75 models have suggested that the mucosal immune system might age more rapidly than
76 the systemic compartment (11) .

77 *Streptococcus pneumoniae* (Spn) is the most common bacterial cause of pneumonia, but
78 usually colonizes the nasopharynx in absence of disease, with colonization frequency
79 decreasing with age (12, 13). Thus, a seeming paradox exists in individuals with advanced
80 age who are at increased risk of pneumococcal disease, but are rarely colonized. On the
81 other hand, children are frequently colonized and infected and are thus the main reservoir
82 for pneumococcal transmission. Mathematical modelling has suggested that the gradual
83 development of adaptive immunity leads to reduced colonization rates with advanced age
84 (14). Mouse models have suggested that Spn colonization is controlled by Th17 cells at

85 the nasal mucosa (15, 16). Using an experimental human pneumococcal challenge
86 model, we recently demonstrated that Spn colonization control in healthy young adults
87 was associated with responses of nasal-resident granulocytes and monocyte recruitment
88 (17, 18).

89 Here, we aimed to investigate whether immune cell composition at the nasal mucosa are
90 altered in young children and older adults compared to young adults, which could underlie
91 there differential susceptibility to respiratory tract infections. Therefore, we collected
92 minimally-invasive nasal microbiopsies from 207 individuals between 1 to 80 years old
93 and immunophenotyped immune cells using flow cytometry (19). We also investigated
94 circulating innate immune cell function and phenotype and correlated these with mucosal
95 findings. Finally, we investigated the effect of Spn colonization on nasal inflammation by
96 collecting nasal lining fluid and its effect on immune cell composition and activation (20).

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106 **Results**

107 **Composition and activation of nasal immune cells changes with age**

108 Here, we phenotyped nasal cells collected using minimally-invasive nasal microbiopsies
109 (Figure 1). Individuals were grouped as children (1-5 years old, n=43), young adults (18-
110 49 years old, n=121) or older adults (50-80 years old, n=43, Table 1). Samples from
111 children were obtained during planned procedures under general anaesthesia such as
112 dental extractions, while adults were awake for sample collection.

113 Nasal cell populations exhibited a significant shift with age (Figure 1A). The capacity to
114 maximally discriminate groups occurred at an age cut-off between young and older adults
115 at 50 years (see Figure S1 in the SI Appendix). Levels of granulocytes were 6.7x and 7.4x
116 lower in children than in young adults and older adults, respectively (Figure 1B).
117 Conversely, nasal T cell levels were 16.1x and 8.1x lower in older adults than in children
118 or young adults, respectively (Figure 1C). As a proportion of nasal immune cells,
119 neutrophils increased, and T cells decreased with age (Figure 1D, E). Monocytes were
120 rare at the nasal mucosa and did not change in frequency with age (Figure 1F). Among T
121 cells, CD8⁺ T cells were the most abundant subset in all age groups (Figure 1G). CD4⁻
122 CD8⁻ T cell numbers and CD8⁺ T cell numbers were 4.8x and 2.0x increased in children
123 compared to young adults, respectively (see Figure S2 in the SI Appendix). CD4⁺ T cell
124 numbers were similar between children and young adults and CD4⁺ T cells thus increased
125 in frequency among T cells in adults (Figure 1G and see Figure S2 in the SI Appendix).
126 In children, granulocytes were fewer, but exhibited higher expression of CD66b, a marker
127 of degranulation, than in adults (Figure 1H) (21). T cell activation status was also affected

128 by age, with young adults showing increased levels of human leukocyte antigen – DR
129 isotype (HLA-DR)⁺ T cells compared to children and older adults (Figure 1I).

130 **Circulating monocytes from young children have reduced CCL2 production upon**
131 ***Streptococcus pneumoniae* stimulation**

132 We then investigated how innate immune function was affected by age. As the numbers
133 of nasal cells collected using currettes precluded the conduction of functional assays,
134 blood was used. To measure monocyte function, we stimulated whole blood from 43
135 individuals for 4 hours with heat-killed Spn and assessed production of C-C motif
136 chemokine ligand 2 (CCL2), interleukin-10 (IL-10), IL-6 and tumor necrosis factor alpha
137 (TNF) (see Figure S3 in the SI Appendix). Monocytes from children displayed impaired
138 production of CCL2 upon stimulation compared to adults, while production of IL-6 and
139 TNF was similar (Figure 2A). Little or no of the anti-inflammatory cytokine IL-10 was
140 induced, demonstrating a pro-inflammatory response of blood monocytes upon Spn
141 stimulation.

142 **Blood neutrophils change functionally and phenotypically with age**

143 Neutrophil phagocytic and oxidative capacities were measured using a whole blood
144 reporter bead assay (see Figure S4 in the SI Appendix) (22). In parallel, neutrophils were
145 immunophenotyped using a panel of ten maturation and activation markers (see Figure
146 S4 in the SI Appendix). Neutrophils of older adults displayed increased uptake and
147 oxidation, while no significant differences were present between neutrophils of children
148 and young adults (Figure 2B). The oxidative capacity of neutrophils positively correlated
149 with expression of the activation and maturation markers CD10, CD11b, CD11c and

150 CD66b (Figure 2C). In addition, neutrophil surface levels of CD10, CD11b and CD33, but
151 not CD11c and CD66b, were significantly increased in older adults compared to children
152 (Figure 2D). Blood neutrophils in young adults had increased expression of CD62L
153 compared to children and older adults and increased expression of CD15 and CD16
154 compared to children and older adults, respectively. To compare blood and mucosal
155 neutrophils, paired neutrophils were phenotyped for 13 individuals (Figure 2E). Nasal
156 neutrophils had increased surface expression of CD10, CD11b, CD11c, CD54 and
157 CD66b, while CD62L and CD16 expression were lost. Neutrophils at the nasal mucosa
158 thus exhibit an activated phenotype as previously described for neutrophils in
159 bronchoalveolar fluid (23). Of all markers, only CD10 ($r=0.64$) and CD33 ($r=0.92$)
160 positively correlated between the two compartments, indicating that blood neutrophil
161 phenotype does not accurately reflect mucosal neutrophil phenotype on an individual level
162 (Figure 2F).

163 ***Streptococcus pneumoniae* colonization causes inflammation in children**

164 Finally, we investigated how colonization with Spn affects nasal immune populations and
165 responses by measuring levels of thirty cytokines and chemokines in nasal lining fluid
166 collected from adults ($n=37$, none colonized by Spn) or children ($n=49$, 22 of whom
167 colonized by Spn). Cytokine levels were similar between adults and children not colonized
168 with Spn, but different in children colonized with Spn (Figure 3A). Levels of granulocyte-
169 colony stimulating factor (G-CSF), IL-15, IL-6, CCL3, basic fibroblast growth factor (FGF-
170 basic), IL-17A, IL-10, CCL5, TNF and IL-5 were significantly elevated in colonized children
171 compared to adults (Figure 3B). Among these, CCL3 and IL-6 showed a positive
172 association with pneumococcal load (Figure 3C). G-CSF was the only protein also

173 increased in non-colonized children compared to adults. Interleukin-1 receptor antagonist
174 (IL-1RA) and IL-13 were decreased in both colonized and non-colonized children
175 compared to adults, while IL-7 was lower in only non-colonized children compared to
176 adults. There were no significant differences between colonized and non-colonized
177 children, but these groups showed a relatively large inter-individual variation compared to
178 adults. This is in accordance with previous findings that the gut microbiome shows greater
179 variation in young children than in adults (24). Subsequently, we assessed neutrophil
180 degranulation by measuring nasal levels of myeloperoxidase (MPO), which were
181 increased in both non-colonized (1.5x, $p = 0.02$) and colonized (4.6x, $p = 0.0005$) children
182 compared to adults (Figure 3D) (25). However, levels of MPO were not significantly
183 affected by Spn colonization status or load in children (Figure 3D, E). Nonetheless, the
184 increased levels of MPO in children compared to adults confirms the increased expression
185 of CD66b on nasal granulocytes in children. Indeed, MPO levels correlated on an
186 individual level with granulocyte activation but not granulocyte numbers (Figure 3F, G).

187 ***Streptococcus pneumoniae* colonization is not associated with altered nasal** 188 **immune cells in children**

189 Despite the increased cytokine production during Spn colonization, no clear differences
190 in immune cell levels were apparent between colonized and non-colonized children
191 (Figure 4A-C). Monocytes, which are recruited to the nose of adults experimentally
192 colonized with pneumococcus, were not affected by Spn colonization (Figure 4D) (17).
193 While Spn colonization was not associated with neutrophil activation levels, T cell
194 activation was increased in Spn-colonized children compared to non-colonized children
195 (29.3% versus 18.1% of HLA-DR⁺ T cells, Figure 4E, F).

196 **Discussion**

197 Here, we investigated how the composition of immune cells at the nasal mucosa is altered
198 with age, with only young adults having an immune profile with an abundance of both
199 granulocytes and T cells. Granulocytes were depleted in children, while there was a
200 paucity of T cells in older adults.

201 As neutrophils have a critical concentration threshold for effective bacterial killing, it is
202 possible the reduced number of nasal granulocytes is associated with the increased
203 susceptibility of children to respiratory tract colonization and infections (26). The reduced
204 expression of the adhesion molecules CD62L, CD11b and CD15, which are important for
205 extravasation and trafficking to tissues (27), on blood neutrophils from children could
206 explain their limited numbers at the nasal mucosa. Nasal IL-8 levels, which is important
207 for neutrophil migration, were not different between adults and children. We did not,
208 however, investigate expression of chemokine receptors as CXCR1 and 2, the receptors
209 for IL-8, on circulating neutrophils and it cannot be excluded that differences in surface
210 expression of these markers exist between children and adults (28).

211 Despite having reduced numbers of nasal granulocytes, the granulocytes in children were
212 activated to a higher degree than in young adults, as shown by increased levels of CD66b
213 expression and increased levels of MPO. Possible explanations for this include a different
214 effect of migration into tissues, which can activate neutrophils *per se*, and altered
215 activation by microbiota in children compared to adults. Indeed, surface marker
216 expression between blood and nasal neutrophils correlated poorly on an individual level,
217 with the exception of CD10 and CD33. This highlights that the study of immune cells in
218 the circulation potentially does not reflect the same cell type at mucosal sites.

219 Circulating neutrophil responses using the bead reporter assay were increased in older
220 adults compared to younger adults, in agreement with previous reports on neutrophils
221 from elderly individuals (29). Our observations also corroborate findings from mouse
222 models where aged neutrophils show an increased phagocytic capacity compared to non-
223 aged neutrophils, which was associated with elevated expression of CD11b (27). In
224 contrast, previous studies have shown that the antibody-mediated opsonophagocytic
225 capacity of neutrophils is reduced in the elderly with reduced observed responses to
226 *Staphylococcus aureus*, but not *Escherichia coli* (30, 31).

227 We observed that T cell levels were reduced in older adults, which was already apparent
228 around the age of 50, before the increased susceptibility to infections becomes apparent.
229 As tissue-resident memory T cells are crucial for protection against infections at mucosal
230 surfaces, this lack of mucosal T cells could provide an explanation for the increased
231 susceptibility to respiratory infections in the elderly (32). Although vaccine efficacy drops
232 with advanced age (33), the development of vaccines that increase tissue-resident
233 memory T cells might thus be particularly beneficial for the elderly. In that light it would be
234 interesting to further characterize these T cells at the mucosa to investigate which specific
235 cells are lost, although this would require access to large tissue samples such as biopsies.
236 Spn colonization led to increased nasal inflammation in children, in contrast to what we
237 previously observed in experimentally colonized adults (17). This could explain the
238 increased transmission potential of children, as inflammation was shown to augment
239 transmission in murine models (34). Although the molecular mechanisms that associate
240 with increased inflammation upon colonization in children remain unclear, it is likely not
241 uniquely due to higher pneumococcal density in children as many cytokines only poorly

242 correlated with Spn density. It is not impossible that the concurrent presence of other
243 bacterial and viral factor was associated inflammation but also predisposing to Spn
244 colonization.

245 Spn colonization had limited effect on the nasal immune cell composition or activation
246 status in children, with the exception of HLA-DR expression on T cells. This corroborates
247 the observed increase in nasal levels of prototypic T cell cytokines as IL-17A, IL-5, IL-10
248 and CCL-5 in colonized children. Levels of MPO and CD66b expression on neutrophils
249 were not affected by Spn colonization in children. This is in contrast with young adults
250 experimentally inoculated by Spn, who show an increase in MPO levels following
251 colonization (17). Possible explanations for this discrepancy are the reduced total number
252 of granulocytes in children, an increased baseline neutrophil activation in non-colonized
253 children compared to adults, or presence of low numbers of Spn that are not detected by
254 classical microbiology but could still activate neutrophils.

255 In addition, Spn colonization in children was not associated with increased levels of
256 monocytes, which are important for Spn clearance (17, 35). Indeed, nasal CCL2 levels,
257 which mediates monocyte recruitment (17, 35), were not increased in colonized children.
258 Interestingly, blood monocytes from children showed an impaired production of CCL2
259 upon Spn stimulation. This was specific as production of TNF and IL-6 was not affected.
260 Previously it was shown that infant mice also have a limited monocyte recruitment
261 following Spn colonization, leading to an inability to clear Spn colonization, although this
262 was associated with microbiota-driven increased baseline CCL2 levels (36).

263 In contrast to our findings, in one previous study in which nasal aspirates were collected
264 in children with acute otitis media, upper respiratory infections or without infection,

265 recruitment of neutrophils to the nasopharynx correlated with Spn density (37). However,
266 the previous study used qPCR for the genes *CD16*, *CD18* and *CD62L* on nasal aspirate
267 pellets to quantify neutrophils, which might not as accurately measure neutrophil counts
268 as flow cytometry, since qPCR reflects both the cellular composition and gene expression
269 levels of individual cells. Moreover, the previous study found increased neutrophil count
270 especially during otitis media, which also associated with increased Spn density, while we
271 studied immune composition in the absence of infection.

272 One previously postulated explanation for the increased susceptibility to Spn colonization
273 in children is an increased Treg/Th17 ratio in children compared to adults (38, 39).
274 However, IL-17A levels were elevated in nasal fluid of colonized children (Figure 3B). In
275 addition, levels of nasal Tregs were not affected by colonization state and were lower in
276 children than in young adults (Figure 4G).

277 A limitation of this cross-sectional observational study was that we focused on three
278 groups: young children, young adults and older adults. Consequently, we have no older
279 children from 6-17 years and we also have few adults between 30-50 years. This makes
280 it hard to detect at which ages nasal immune profiles start shifting.

281 In conclusion, we observed severe and dynamic alterations in mucosal immunity with age,
282 highlighting the need for measuring mucosal responses in target populations when
283 investigating host-pathogen interactions and vaccine-induced immunity, especially in the
284 young and elderly.

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287 **Methods**

288 **Study design**

289 We recruited individuals between 1–80 years of age in a series of studies
290 (ISRCTN85509051, ISRCTN10948363, ISRCTN16993271, ISRCTN68323432 and
291 ISRCTN76456378). Some of the subjects in this manuscript (the young adults cohort)
292 were originally described previously (40). For a subset of the young adults, cytokines and
293 nasal cell data have been used in another manuscript deposited on a pre-print server (18),
294 although the results do not overlap with those reported here. All adults were healthy and
295 inclusion criteria common to all adult studies were: capacity to give informed consent,
296 aged >18 years and speak fluent English. Children awaiting a procedure requiring general
297 anaesthesia: dental extraction (44%), MRI (36%), orthopaedic surgery (14%) and plastic
298 surgery (6%), were recruited. Inclusion criteria were: aged 1-5 years, capacity of a parent
299 of the participant to give informed consent and speak fluent English. Exclusion criteria and
300 sample collection details for adults and children are provided in an online data
301 supplement.

302 **Ethics statement**

303 All adult volunteers and a parent of children involved in the study gave written informed
304 consent and research was conducted in compliance with all relevant ethical regulations.
305 Ethical approval was obtained from the East Liverpool NHS Research Ethics Committee,
306 reference numbers: 17/NW/0663, 16/NW/0031, 17/NW/0029, 15/NW/0931 and
307 14/NW/1460.

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309 **Flow cytometry analysis**

310 All flow cytometry samples were acquired on a LSRII flow cytometer (BD) and analysed
311 using Flowjo X (Treestar). Compensation matrices were set using compensation beads
312 (BD Biosciences) and ArC™ Amine Reactive Compensation beads (Thermofisher) and
313 manually inspected for representative samples. All antibodies were titrated and
314 fluorescence minus one controls were used to verify specificity of signal. Additional detail
315 on immunophenotyping of nasal cells, neutrophil phenotyping and monocyte and
316 neutrophil functional assays is provided in an online data supplement.

317 **Luminex analysis of nasal lining fluid**

318 Cytokines were eluted from stored Nasosorption™ filters using 100µL of Luminex assay
319 buffer (ThermoFisher) by centrifugation, then the eluate was cleared by further
320 centrifugation at 16,000 x G, as described previously (17, 19). Concentrations of 30
321 cytokines were measured using the 30-plex magnetic human Luminex cytokine kit (all
322 using lot ID 1805187A, ThermoFisher). Samples were measured on a LX200 (Luminex)
323 and analysed with xPonent3.1 software (Luminex) following manufacturer's instructions.
324 Samples were analysed in duplicates and analytes with a CV > 50% were excluded.

325 **Myeloperoxidase (MPO) ELISA of nasal lining fluid**

326 Levels of myeloperoxidase were determined using the Human Myeloperoxidase DuoSet
327 ELISA Kit (R&D Systems) as per manufacturer's instructions. Plates were read on a
328 FLUOstar® Omega machine (BMG Labtech) and data was analysed with Mars data
329 analysis software version 3.1 following manufacturer's instructions. Samples were
330 analysed in duplicates and samples with a CV > 20% were excluded.

331 **Statistical analysis**

332 Statistical analyses were performed using R software (version 3.5.1). Two-tailed statistical
333 tests were used throughout the study. Mann-Whitney tests were used to compare groups
334 and multiple correction testing (Benjamin-Hochberg) was applied for Luminex analysis.
335 Correlations were assessed using Pearson's correlation test using either raw or log-
336 transformed values. Differences were considered significant at $p < 0.05$.

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459 **Tables**

460 **Table 1. Cohort characteristics.**

	Children	Young adults	Older adults
Nasal phenotyping sample size	43	121	43
Age in median years (range)	3 (1-5)	20 (18-49)	63 (50-80)
Number of females (%)	18 (41.2%)	64 (52.9%)	26 (60.5)
Number of Spn colonized (%)	19 (45.2%*)	9 (7.4%)	2 (4.7%)

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462 *Streptococcus pneumoniae (Spn) colonization was assessed by classical microbiology.

463 For one child no nasopharyngeal swab was collected and colonization was not assessed.

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474 **Figure legends**

475 **Figure 1. Nasal immune cell populations change drastically with age.** A) Multi-
476 dimensional scaling plot based on Euclidian distance, considering immune composition
477 (granulocytes, monocytes and CD4⁺ T, CD8⁺ T and double-negative (DN) T cells as
478 percentage of immune cells) and activation (HLA-DR⁺ and CD66b^{Hi} as percentage of T
479 cells and neutrophils, respectively). Individual children (red circles, n=42), young adults
480 (green triangles, n=86) and older adults (blue squares, n=36) are shown along with 50%
481 confidence intervals. R and p values represent analysis of similarity results and stress
482 depicts Kruskal stress. Violin plots with boxplots showing levels of B) granulocytes and C)
483 T cells normalized to epithelial cells for children (n=43), young adults (n=118) and older
484 adults (n=46). ****p=1.4 x 10⁻⁶ and p=7.2 x 10⁻⁵ by Mann-Whitney test comparing
485 granulocytes in children with young adults and older adults, respectively. ***p=0.0006 by
486 Mann-Whitney test comparing T cell levels between children and young adults. ****p=4.5
487 x 10⁻¹⁴ and p=2.4 x 10⁻¹¹ comparing T cell levels in older adults with those in children and
488 with young adults, respectively. Scatterplots showing percentage of D) granulocytes and
489 E) T cells among nasal immune cells. Individual subjects are depicted, a grey vertical
490 dashed line shows the cut-off between young and older adults at age 50 and a locally
491 estimated scatterplot smoothing (loess) curve with 95% confidence interval is plotted. F)
492 Violin plots with boxplots showing levels of monocytes normalized to epithelial cells. G)
493 Pie charts showing the mean levels of T cell subsets (CD4⁺, CD8⁺, double-negative (DN)
494 and double-positive (DP) per age group (children, n=43; young adults, n=109; older
495 adults, n=42). H) Violin plots with boxplots showing levels of CD66b^{Hi} granulocytes for
496 children (n=43), young adults (n=121) and older adults (n=41). ****p=2.3 x 10⁻⁶ and

497 *** $p=0.0004$ by Mann-Whitney test comparing children with young adults and older adults,
498 respectively. I) Violin plots with boxplots showing levels of HLA-DR⁺ T cells for children
499 (n=43), young adults (n=109) and older adults (n=38). ** $p=0.001$ and *** $p=0.0002$
500 comparing young adults with children and older adults by Mann-Whitney test, respectively.

501 **Figure 2. Changes in innate immune function with age.** A) Scatterplots showing
502 percentage of monocytes that are producing CCL2, IL-10, IL-6 or TNF following a two-
503 hour stimulation with heat-killed *Streptococcus pneumoniae*. Individual children (red
504 circles, n=23), young adults (green triangles, n=13) and older adults (blue squares, n=7)
505 are depicted. Black line and shaded area depict linear regression fit and 95% confidence
506 intervals. R and p represent rho and p-value from Pearson's correlation test, respectively.
507 B) Neutrophil oxidation capacity was assessed using a phagocytic bead assay at 4
508 timepoints, including a positive control where LPS was added for 45 minutes. Median and
509 interquartile range are shown for children (red circles, n=11), young adults (green
510 triangles, n=6) and older adults (blue squares, n=5). * $p<0.05$, ** $p<0.01$, *** $p<0.001$ by
511 Mann-Whitney test comparing two groups. C) Correlation matrix for oxidative capacity and
512 log-transformed blood neutrophil surface marker expression using Pearson correlation
513 test (n=21). Circle colour and size represent rho and absolute rho value, respectively.
514 * $p<0.05$, ** $p<0.01$ by Pearson correlation test D) Mean fluorescent intensity (MFI) of
515 surface markers on blood neutrophils per age group. Median and interquartile range are
516 shown for children (red circles, n=13), young adults (green triangles, n=8) and older adults
517 (blue squares, n=5). * $p<0.05$, *** $p<0.001$ by Mann-Whitney test. E) Comparison of marker
518 expression on neutrophils from blood (red) and nose (blue) for paired volunteers (n=11
519 children and 2 young adults). F) Correlation matrix for log-transformed marker expression

520 on blood and nasal neutrophils using Pearson correlation test (n=13). Circle colour and
521 size represent rho and absolute rho value, respectively. *p=0.018, ****p=1.1x10⁻⁵ by
522 Pearson correlation test.

523 **Figure 3. Nasal cytokine responses to pneumococcal colonization.** A) Multi-
524 dimensional scaling plot based on Euclidian distance, considering log-transformed
525 concentrations of 30 nasal cytokines. Individual Spn not-colonized children (Spn⁻ purple
526 circles, n=17), Spn colonized children (Spn⁺ red triangles, n=14) and young adults (green
527 squares, n=26) are shown along 50% confidence intervals. R and p values represent
528 analysis of similarity results and stress depicts Kruskal stress. B) Median and interquartile
529 range of the concentrations for each of the 30 cytokines are shown for not-colonized
530 children (purple, n=27), colonized children (red, n=22) and young adults (n=37). *p<0.05,
531 **p<0.01, ***p<0.001, ****p<0.0001 by Mann-Whitney test compared to young adults,
532 followed by Benjamini-Hochberg correction for multiple testing, with colour indicating
533 significantly altered group. C) Volcano plot showing Pearson correlation results between
534 log-transformed pneumococcal load for colonized children with log-transformed cytokine
535 concentration, not corrected for multiple testing. Significantly correlating cytokines are
536 depicted in red. D) Violin plots with boxplots showing concentration of myeloperoxidase
537 (MPO) for not-colonized children (purple, n=27), colonized children (red, n=22) and young
538 adults (green, n=36). *p=0.022, ***p=0.0005 by Mann-Whitney test. Scatterplots show
539 correlation between log-transformed concentration of nasal MPO with E) log-transformed
540 pneumococcal load, F) percentage of CD66b^{Hi} granulocytes and g) log-transformed levels
541 of nasal granulocytes (normalized to epithelial cells). Individuals are shown, and line and

542 shaded area represent linear regression and 95% confidence interval, respectively. R
543 (Pearson rho) and p (p-value) are shown for each correlation.

544 **Figure 4. Nasal immune cell populations are not affected by Spn colonization in**
545 **children.** A) Multi-dimensional scaling plot based on Euclidian distance, considering
546 immune composition (granulocytes, monocytes and CD4⁺ T, CD8⁺ T and double-negative
547 (DN) T cells as percentage of immune cells) and activation (HLA-DR⁺ and CD66b^{Hi} as
548 percentage of T cells and neutrophils, respectively). Individual non-colonized children
549 (Spn⁻ purple circles, n=23) and colonized children (Spn⁺, red triangles, n=19) are shown
550 along with 50% confidence intervals. R represents the analysis of similarity results and
551 stress depicts Kruskal stress. Violin plots with boxplots showing numbers of B)
552 granulocytes, C) T cells or D) monocytes normalized to epithelial cells for non-colonized
553 children (Spn⁻, purple, n=23) and colonized children (Spn⁺, red, n=19). Violin plots with
554 boxplots showing levels of E) CD66b^{Hi} granulocytes or F) HLA-DR⁺ T cells. **p=0.005 by
555 Mann-Whitney test. G) Violin plots with boxplots showing numbers of regulatory T cells
556 (Tregs) for non-colonized children (Spn⁻, purple, n=23), colonized children (Spn⁺, red,
557 n=19) and non-colonized young adults (green, n=11). *p=0.036, **p=0.0098 comparing
558 young adults with colonized and non-colonized children by Mann-Whitney test,
559 respectively.

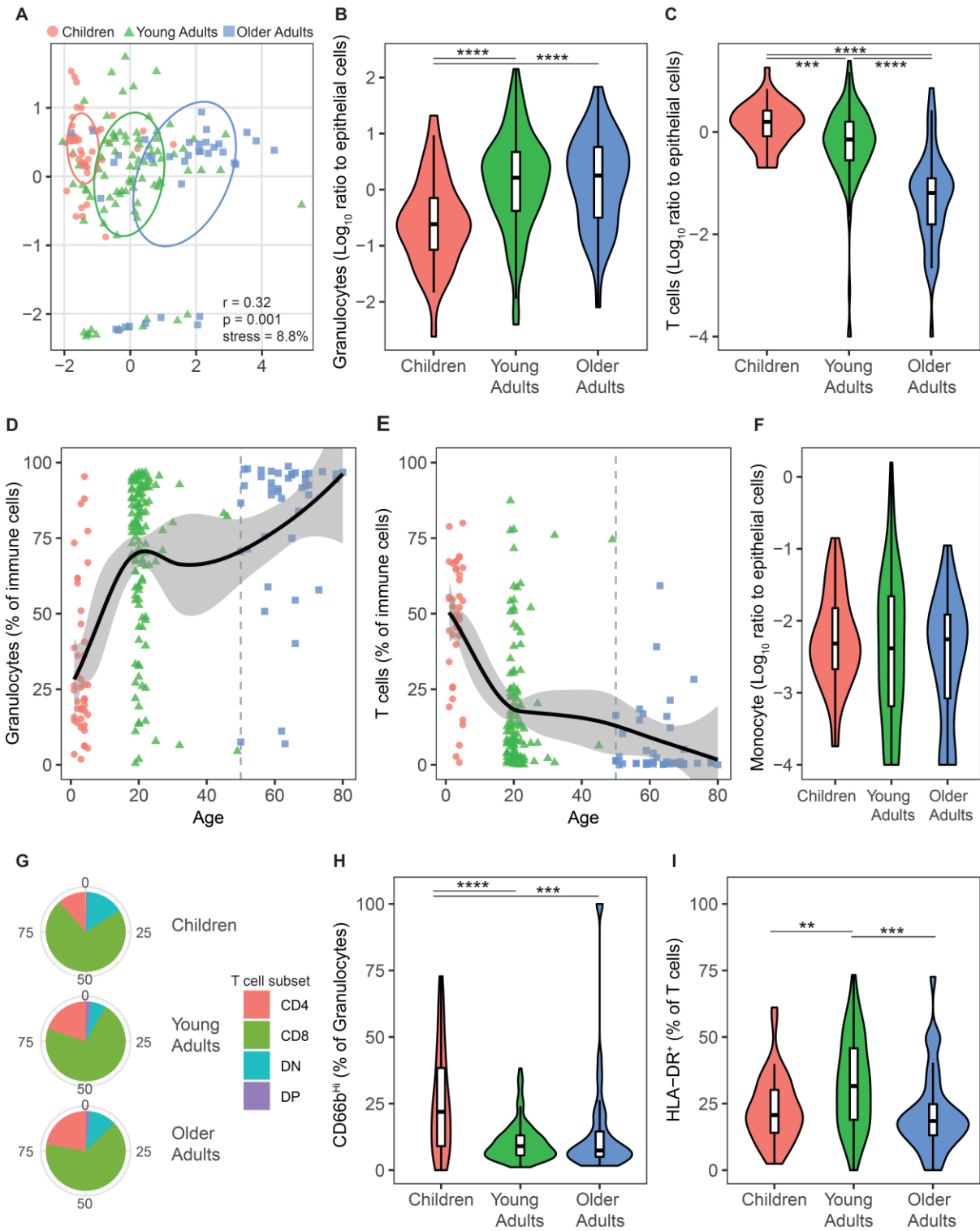
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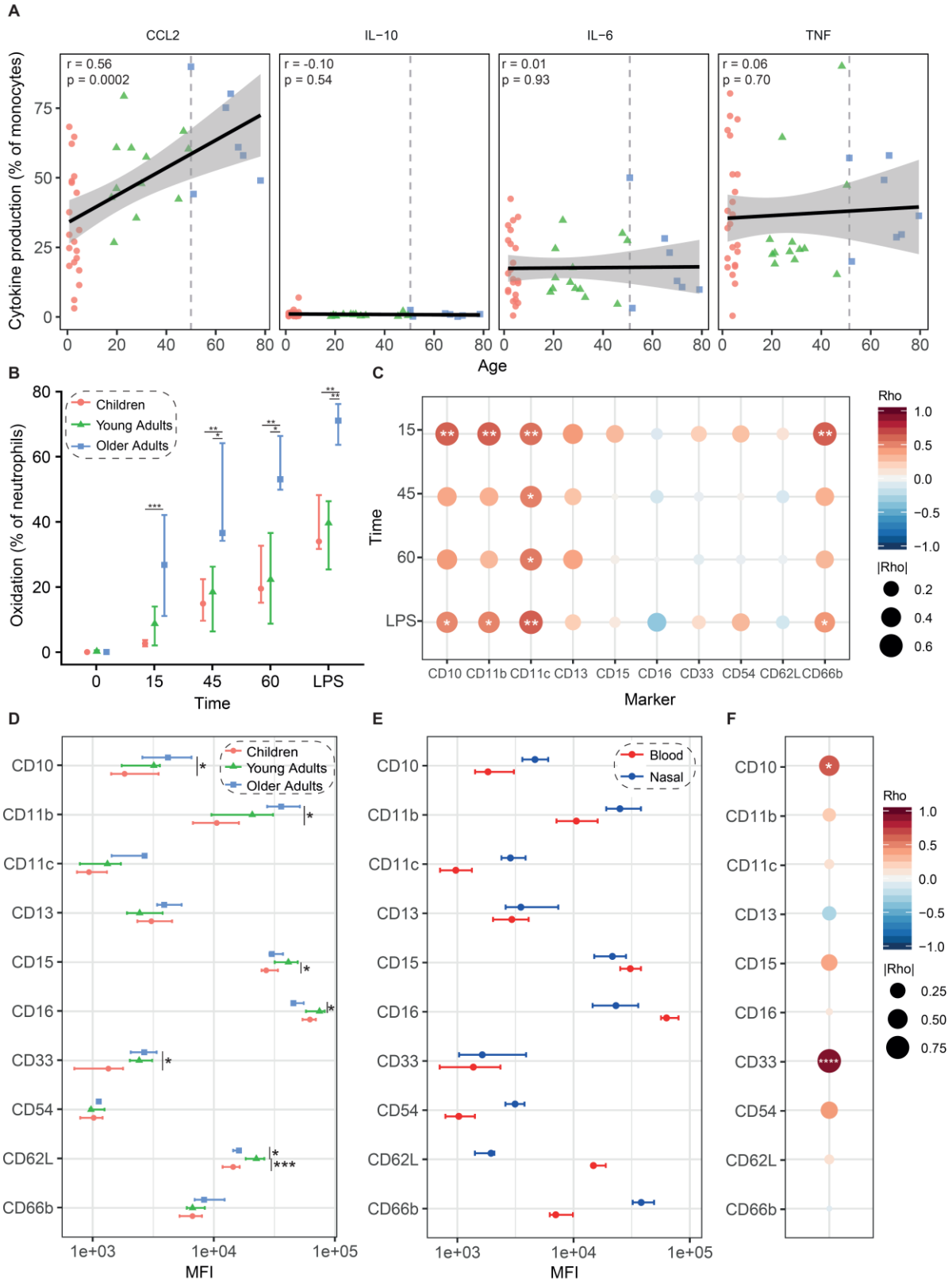
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564 **Figures**



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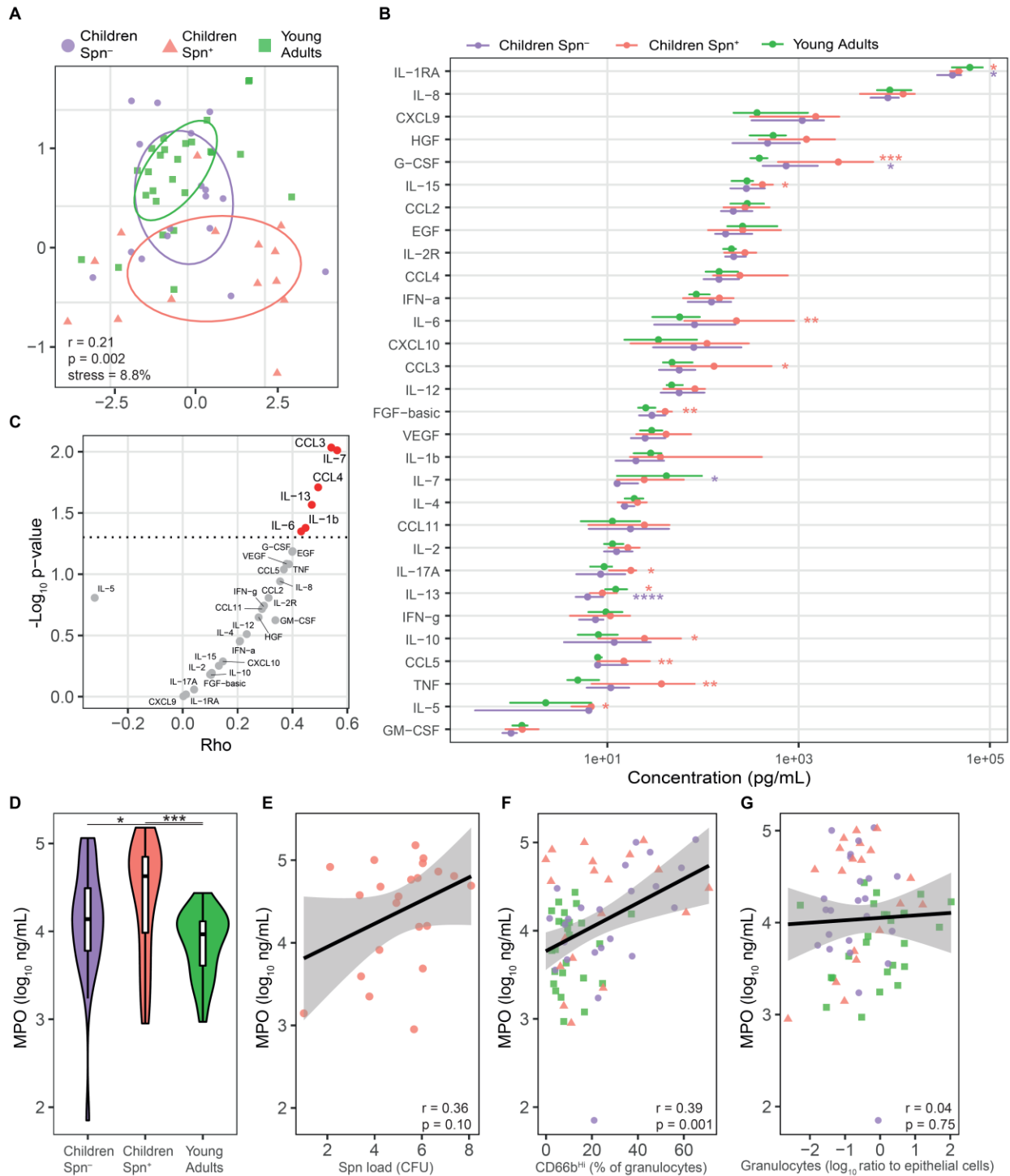
566 **Figure 1.**



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568 Figure 2.

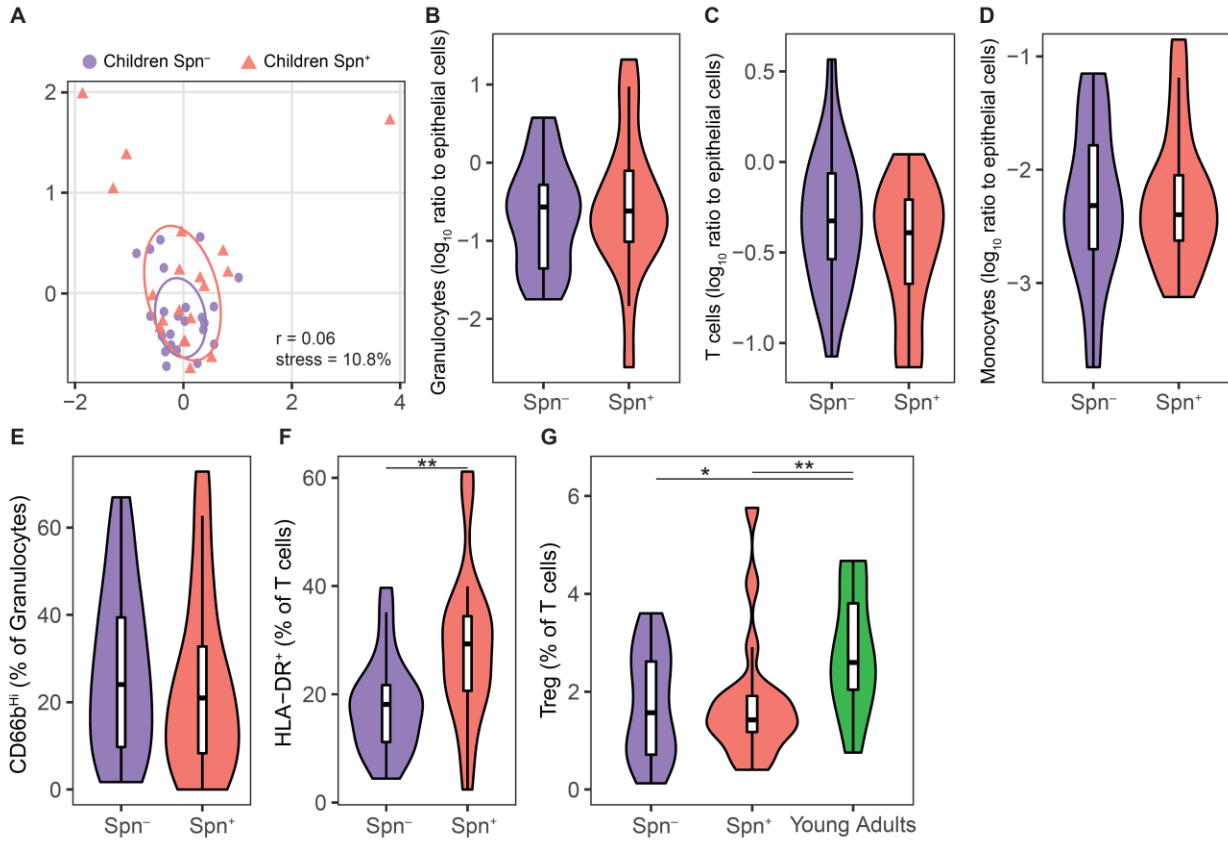
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571 Figure 3.

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