

1 **Expansion of different subpopulations of CD26^{-low} T cells in allergic**
2 **and non-allergic asthmatics.**

3 **Authors**

4 Juan José Nieto-Fontarigo¹, Francisco Javier Salgado*¹, María Esther San-José², María Jesús
5 Cruz³, Luis Valdés⁴, Amparo Pérez-Díaz⁵, Pilar Arias¹, Montserrat Nogueira¹, Francisco Javier
6 González-Barcala⁶.

7 **Affiliations:**

8 ¹Department of Biochemistry and Molecular Biology, Faculty of Biology-Biological Research
9 Centre (CIBUS), Universidade de Santiago de Compostela, Santiago de Compostela, Spain.

10 ²Clinical Analysis Service, USC University Hospital Complex (CHUS), Santiago de
11 Compostela, Spain.

12 ³Department of Respiratory Medicine-Hospital Vall d'Hebron, Universitat Autònoma de
13 Barcelona, Barcelona, Spain. Spanish Biomedical Research Networking Centre-CIBERES.

14 ⁴Department of Medicine-University of Santiago de Compostela; Department of Respiratory
15 Medicine-University Hospital of Santiago de Compostela; Health Research Institute of Santiago
16 de Compostela (IDIS).

17 ⁵Drug Screening Platform/Biofarma Research Group, Molecular Medicine and Chronic
18 Diseases Research Center, University of Santiago de Compostela, Santiago de Compostela,
19 Spain.

20 ⁶Department of Medicine-University of Santiago de Compostela, Department of Respiratory
21 Medicine-University Hospital of Santiago de Compostela, Spanish Biomedical Research
22 Networking Centre-CIBERES; Health Research Institute of Santiago de Compostela (IDIS).

23 **Corresponding author:**

24 Francisco Javier Salgado Castro. Biology of Lymphocyte Group (BioLympho). Department of
25 Biochemistry and Molecular Biology, Faculty of Biology-Biological Research Centre (CIBUS),

- 26 Universidade de Santiago de Compostela, Campus Vida, C/ Lope Gómez de marzoa s/n,
27 Santiago de Compostela, Spain. E-mail: Phone: +34 881816941.
- 28 **Supplementary Tables: 4**
- 29 **Supplementary Figures: 6**

Supplementary Table S1. Correlation of CD26 and CD126 expression with the % of different leukocyte subsets.

	Correlation between the % of cells and the % of CD26+ cells in each subpopulation		Correlation between the % of cells and the % of CD126+ cells in each subpopulation	
	r_s	P	r_s	P
Eosinophils	-.181	.003	.205	< .001
Monocytes	-.206	< .001	-.085	.168
Neutrophils	.041	.500	.035	.564
Lymphocytes	-.103	.094	-.057	.354
CD4⁻ cells	-.312	< .001	-.353	< .001
CD4⁺ cells	.367	< .001	.211	< .001
Teff cells	.469	< .001	.247	< .001
Treg cells	.023	.702	-.070	.254
Tlow cells	-.342	< .001	-.360	< .001

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Supplementary Table S2. Severity and control degree of the study patients.

	IMA	MSA		CA	UA
N	102	90		158	34
Age (mean (range))	44 (20-72)	41 (18-68)		45 (18-72)	45 (20-68)
Sex (M/F)	33/69	35/55		59/99	9/25
FEV1 (%)	105.0 (95.0-113.3)***	86.5 (67.0-100.0)		99.0 (87.0-111.0)*	82.0 (60.7-100.0)
FEV1/FVC (%)	77.5 (72.5-81.2)***	70.3 (59.5-78.5)		76.4 (69.0-80.5)*	70.0 (56.4-76.8)
Neutrophils (10³ cells/μL)	3.60 (3.09-4.23)	3.62 (2.88-4.40)		3.58 (3.07-4.23)	4.17 (2.78-5.00)
Lymphocytes (10³ cells/μL)	1.97 (1.65-2.25)	1.97 (1.64-2.54)		1.95 (1.60-2.26)	2.02 (1.86-2.74)
Monocytes (10³ cells/μL)	0.39 (0.30-0.48)	0.39 (0.32-0.49)		0.38 (0.30-0.49)	0.41 (0.32-0.48)
Eosinophils (10³ cells/μL)	0.26 (0.17-0.39)*	0.33 (0.21-0.48)		0.27 (0.18-0.40)*	0.38 (0.21-0.63)
Basophils (10³ cells/μL)	0.04 (0.02-0.05)	0.04 (0.03-0.05)		0.04 (0.02-0.05)	0.04 (0.03-0.06)
IgE (IU/mL)	45.5 (15.5-205.0)	84.0 (28.2-209.5)		79.0 (20.5-200.0)	62.0 (24.7-274.0)
<p>CA, controlled asthmatics; IMA, intermittent-mild asthmatics; MSA, moderate-severe asthmatics; UA, uncontrolled asthmatics. Data are presented as median value (IQR1-3), unless otherwise expressed. Mann-Whitney U test was used to compare IMA vs. MSA and CA vs. UA. * $P < 0.05$; *** $P < 0.001$</p>					

Supplementary Table S3. Average expression of CD26 and CD126 molecules in different leukocyte subsets and the half-life of these cells in peripheral circulation.

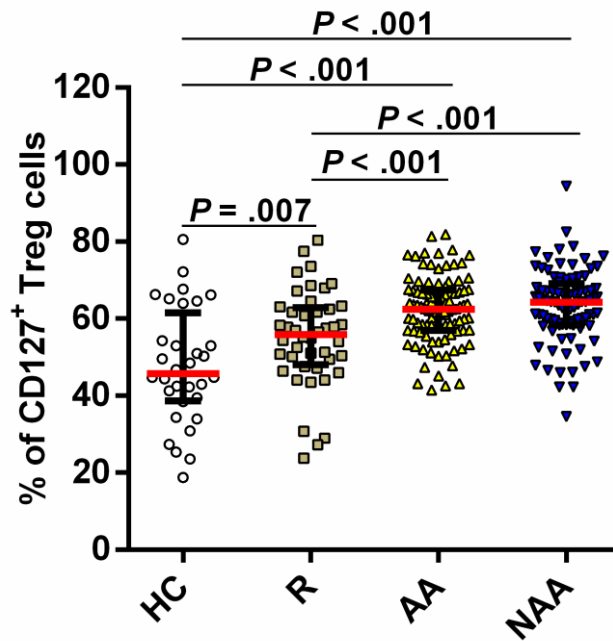
	Eosinophils	Monocytes	Neutrophils	Lymphocytes
Absolute count (x10³ cells/mL)	0.04-0.4	0.2-0.8	2.5-7.5	1.5-3.5
% of CD126⁺ cells*	63.4 (44.5-77.5)	78.5 (52.2-91.0)	99.5 (98.8-99.8)	55.3 (48.6-61.3)
% of CD26⁺ cells*	2.1 (1.4-3.2)	7.4 (3.5-17.6)	11.9 (5.0-27.7)	59.2 (54.4-66.4)
Half-life in peripheral blood[#]	8-18 h	22-24 h	6 h - 5 days	days-years

*Median value (IQR1-3).

[#]Data come from Pillay J, et al. Blood. 2010;116:625-7; Park YM, et al. Allergy Asthma Immunol Res. 2010;2:87-101; Patel AA, et al. J Exp Med. 2017;214:1913-1923; Robertson JM, et al. Immunol Rev. 2006;211:49-57.

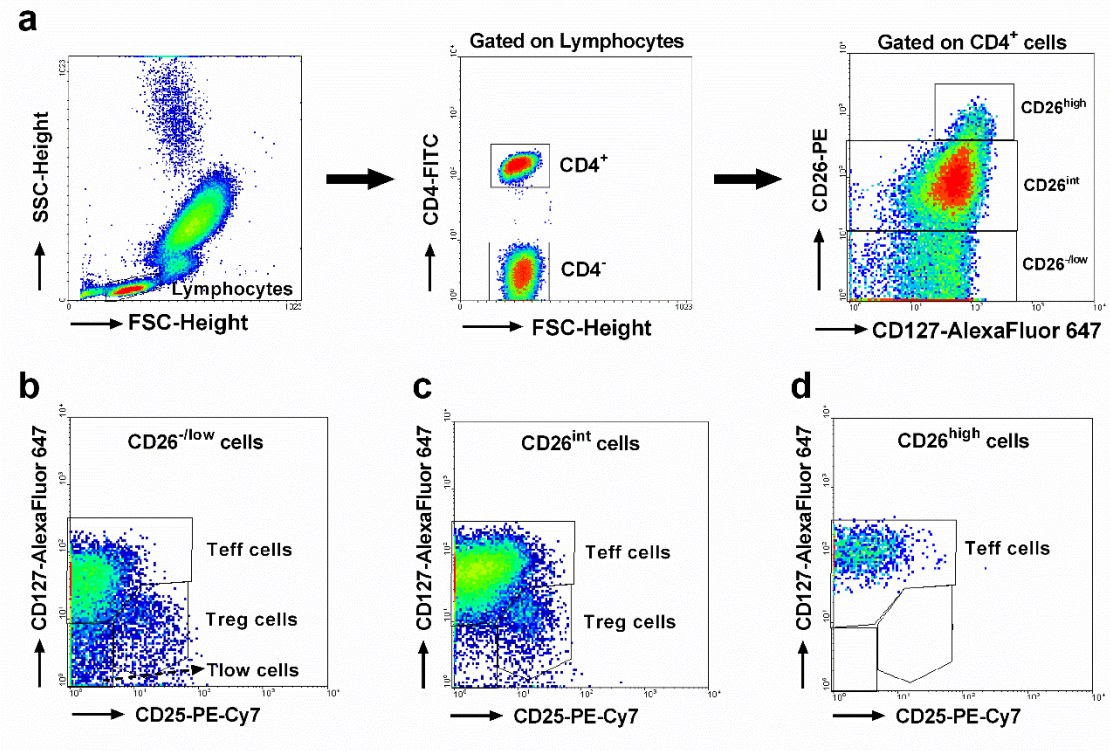
Supplementary Table S4. List of monoclonal antibodies used in flow cytometry assays.

Target	Clone	Fluorophore	Provider	Cat. no
CD3	UCHT1	FITC	BD Pharmigen™	555332
CD4	RPA-T4	FITC	BD Pharmigen™	555346
CD25	M-A251	PE-Cy7	BD Pharmigen™	5135905
CD127	HIL-7R-M21	AlexaFluor-647	BD Pharmigen™	558598
CD3	UCHT1	PerCP-Cy5.5	BD Pharmigen™	560835
CD8	RPA-T8	PerCP-Cy5.5	BD Pharmigen™	560662
CD19	HIB19	PerCP-Cy5.5	BD Pharmigen™	561295
CD27	M-T271	PerCP-Cy5.5	BD Pharmigen™	560612
CD28	CD28.2	PerCP-Cy5.5	BD Pharmigen™	560685
CD45RA	HI100	PerCP-Cy5.5	BD Pharmigen™	563429
CD56	B159	PerCP-Cy5.5	BD Pharmigen™	560842
CD197 (CCR7)	150503	PerCP-Cy5.5	BD Pharmigen™	561144
TCR-γ/δ	B1	FITC	Biolegend	331207
CD26	TP1/19	PE	Immunostep	PE26 100T
CD126 (IL-6Rα)	REA291	PE	MACS Miltenyi Biotec	130-104-101



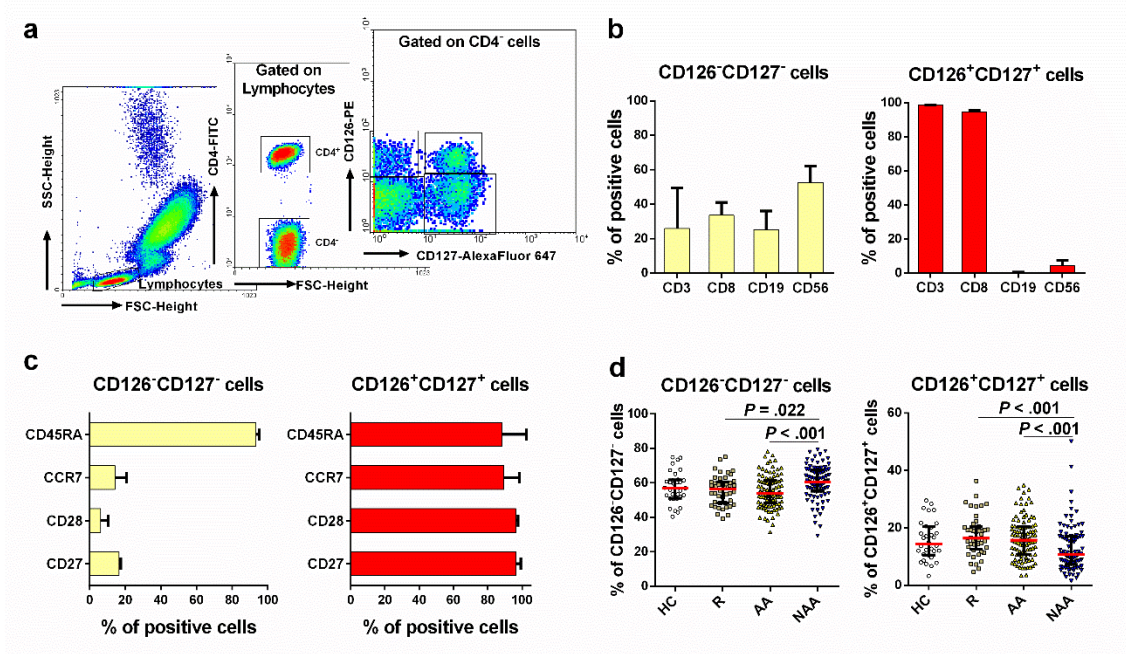
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36 **Supplementary Figure S1. Rhinitic and asthmatic patients show augmented levels of**
 37 **CD127⁺ regulatory T cells.** Lymphocytes from healthy controls (HC; N = 32) and patients with
 38 rhinitis (R; N = 44), allergic asthma (AA; N = 100), and non-allergic asthma (NAA; N = 92)
 39 were gated on FSC/SSC plots and further subdivided into CD4⁻ and CD4⁺ cells. Then,
 40 regulatory T cells (Treg) were identified amongst CD4⁺ T lymphocytes according to CD25 and
 41 CD127 markers. Afterwards, the percentage of CD127⁺ cells within the Treg compartment was
 42 measured. Statistically significant difference between the different groups of donors is indicated
 43 ($P < .05$).



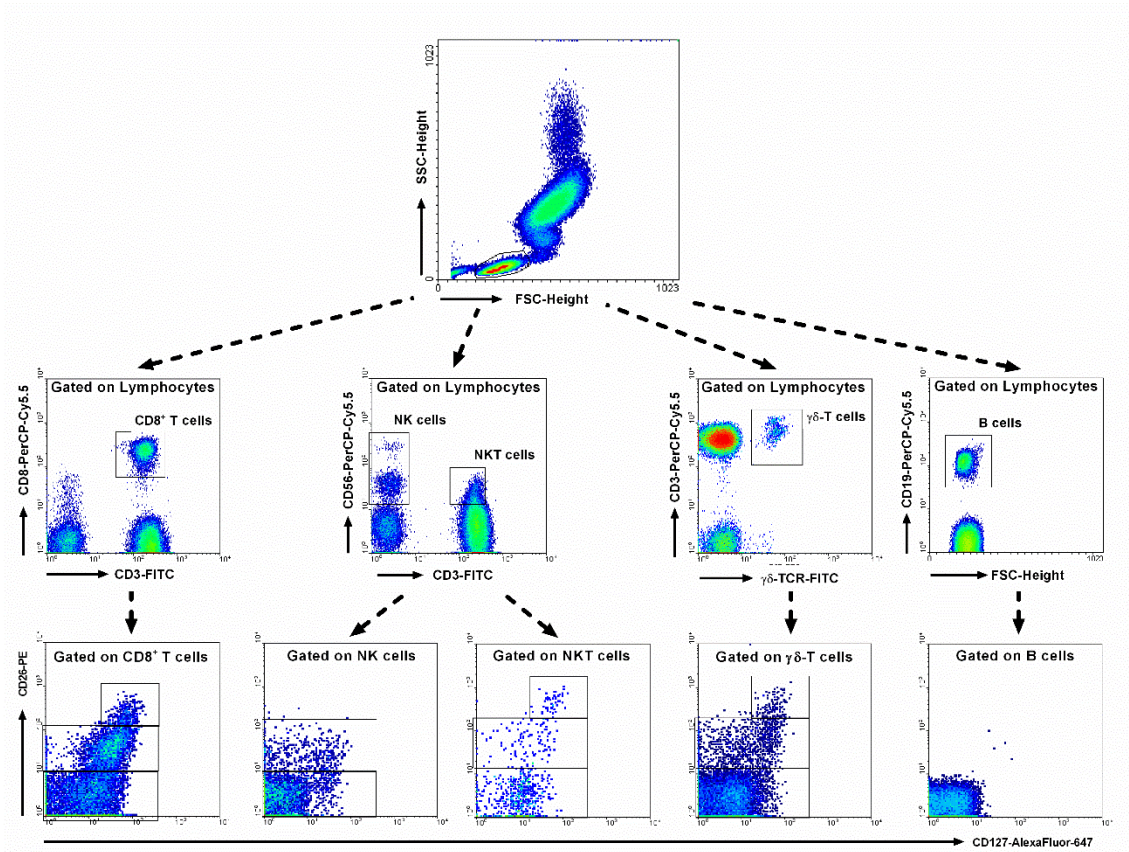
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45 **Supplementary Figure S2. Treg cells are distributed within CD26^{low} and CD26^{int} subsets**
 46 **of CD4⁺ T cells.** a) Lymphocytes from healthy controls (HC) and patients with rhinitis (R),
 47 allergic asthma (AA), and non-allergic asthma (NAA) were gated on FSC/SSC plots and further
 48 subdivided into CD4⁻ and CD4⁺ cells. Then, 3 subsets of CD4⁺ lymphocytes were identified
 49 based on CD26 and CD127 markers: CD26^{low}, CD26^{int} and CD26^{high}. As figures **b-d** point out,
 50 Treg cells (CD25^{high}CD127^{low}) are present in CD26^{low} (**b**) and CD26^{int} (**c**) subsets of CD4⁺ T
 51 lymphocytes, but not amongst the CD26^{high} compartment of T helper cells (**d**).



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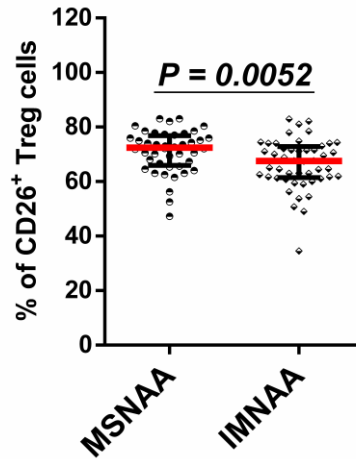
53 **Supplementary Figure S3. CD4⁺ lymphocytes with a CD126⁻CD127⁻ phenotype are**
 54 **increased in non-allergic asthmatics (NAA) compared to allergic asthmatics (AA).** a) CD4⁺
 55 lymphocytes were identified through the gating strategy depicted (left and middle dot-plots),
 56 and expression of CD127 and CD126 was used to identify four subsets (right dot plot). b)
 57 Lymphocyte subset composition of CD4⁺CD126⁻CD127⁻ and CD4⁺CD126⁺CD127⁺ cells based
 58 on CD3 (T), CD8 (Tc), CD19 (B), and CD56 (NK, NKT) antigens. c) Phenotype of CD4⁺
 59 CD126⁻CD127⁻ and CD4⁺CD126⁺CD127⁺ cells based on CD45RA, CCR7, CD127, CD28, and
 60 CD27 markers. We evidenced that the former CD26^{int}CD127⁺ and CD26^{high}CD127⁺ subsets
 61 were combined in one single CD4⁺ population (CD126⁺CD127⁺ cells), with a homogeneous
 62 naïve-memory CD3⁺ and CD8⁺ phenotype. In contrast, the CD126⁻CD127⁻ subset comprised
 63 T_{EMRA}-like cells (CD45RA^{high}CCR7^{low}CD27^{low}CD28^{low}), but a less-defined CD4⁺ lymphoid
 64 origin (B, NK, NKT, or $\gamma\delta$ -T cells). In b) and c), data were obtained from 3 representative
 65 donors. d) Percentage (median \pm IQR) of CD126⁻CD127⁻ and CD126⁺CD127⁺ cells in CD4⁺
 66 lymphocytes from healthy controls (HC; N = 32) and patients with rhinitis (R; N = 44), allergic
 67 asthma (AA; N = 100), and non-allergic asthma (NAA; N = 92). Like CD4⁺CD26⁺CD127⁺
 68 lymphocytes, CD4⁺CD126⁺CD127⁺ cells were reduced in NAA compared to AA, which was
 69 associated to the expansion of the double negative (CD126⁻CD127⁻) compartment in NAA.
 70 Only results from those subpopulations with significant differences between donor groups were
 71 shown. Statistically significant differences between groups of donors were indicated (Kruskal-
 72 Wallis test: $P < .05$).



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74 **Supplementary Figure S4. Gating strategy for CD8⁺ Tc, NK, NKT, $\gamma\delta$ -T, and B**
 75 **lymphocytes.** After gating, expression of CD127 and CD26 was used to identify different
 76 subsets (lower row of dot-plots). CD3⁺ lymphocytes (Tc, NKT, $\gamma\delta$ -T) were characterized by the
 77 presence of three subsets (CD26^{-/low}, CD26^{int}, and CD26^{high}), while CD3⁻ lymphocytes displayed
 78 a double (CD26^{-/low}, CD26^{int}; NK cells) or a single (CD26⁻; B cells) phenotype.

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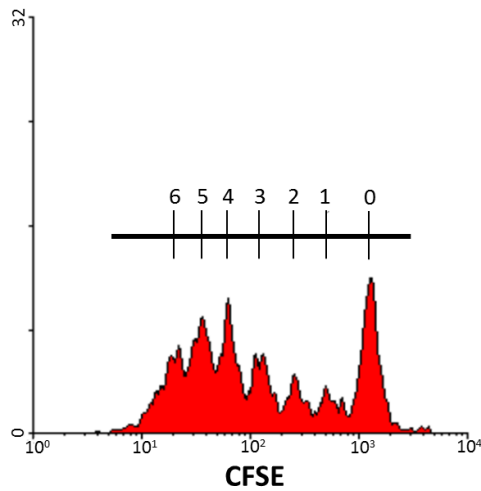


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81 **Supplementary Figure S5. Moderate-severe non-allergic asthmatics display higher**
82 **percentage of CD26⁺ regulatory T cells than intermittent-mild non-allergic asthmatics.**

83 Lymphocytes from either moderate-severe non-allergic (MSNAA; N = 90) or intermittent-mild
84 non-allergic (IMNAA; N = 102) asthmatics were gated on FSC/SSC plots and further
85 subdivided into CD4⁻ and CD4⁺ cells. Then, regulatory T cells (Treg) were identified amongst
86 CD4⁺ T lymphocytes according to CD25 and CD127 markers. Afterwards, the percentage of
87 CD26⁺ cells within the Treg compartment was measured. Statistically significant difference
88 between the different groups of donors is indicated (*P*-value).

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91 **Supplementary Figure S6. Analysis of the number of cell divisions by flow cytometry.**
 92 PBMCs were stained with 5 μ M CFSE and cultured at 0.25×10^6 cells/mL in complete medium
 93 (RPMI 1640, 10% FBS, 100 μ g/mL streptomycin, 100 U/mL penicillin) supplemented with 1
 94 μ g/ml PHA as a T-cell specific polyclonal stimulus. Upon 6 days of *in vitro* culture, CFSE
 95 fluorescence decay was measured by flow cytometry and allowed the calculation of the number
 96 of cell divisions. A representative histogram plot is shown, where numbers 1-6 represent the
 97 corresponding round of division, and 0 those lymphocytes that remained at a resting state.
 98 Responder frequency (Rf) was calculated as the percentage of cells that divided at least once
 99 (i.e., percentage of cells that divide from 1 to 6 times).

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